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# **emg3d Documentation**

***Release 0.14.3***

**The emg3d Developers**

**20 November 2020**



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A multigrid solver for 3D electromagnetic diffusion with tri-axial electrical anisotropy. The matrix-free solver can be used as main solver or as preconditioner for one of the Krylov subspace methods implemented in *scipy.sparse.linalg*, and the governing equations are discretized on a staggered Yee grid. The code is written completely in Python using the NumPy/SciPy-stack, where the most time- and memory-consuming parts are sped up through jitted numba-functions.



# CHAPTER 1

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## More information

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For more information regarding installation, usage, contributing, roadmap, bug reports, and much more, see

- **Website:** <https://empymod.github.io>,
- **Documentation:** <https://emg3d.readthedocs.io>,
- **Source Code:** <https://github.com/empymod/emg3d>,
- **Examples:** <https://empymod.github.io/emg3d-gallery>.





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### Features

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- Multigrid solver for 3D electromagnetic (EM) diffusion with regular grids (where source and receiver can be electric or magnetic).
- Compute the 3D EM field in the complex frequency domain or in the real Laplace domain.
- Includes also routines to compute the 3D EM field in the time domain.
- Can be used together with the [SimPEG](#)-framework.
- Can be used as a standalone solver or as a pre-conditioner for various Krylov subspace methods implemented in SciPy, e.g., BiCGSTAB (*scipy.sparse.linalg.bicgstab*) or CGS (*scipy.sparse.linalg.cgs*).
- Tri-axial electrical anisotropy.
- Isotropic magnetic permeability.
- Semicoarsening and line relaxation.
- Grid-size can be anything.
- As a multigrid method it scales with the number of unknowns  $N$  and has therefore optimal complexity  $O(N)$ .



## CHAPTER 3

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### Installation

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You can install `emg3d` either via `conda` (preferred):

```
conda install -c conda-forge emg3d
```

or via `pip`:

```
pip install emg3d
```

Minimum requirements are Python version 3.7 or higher and the modules `scipy` and `numba`. Various other packages are recommended or required for some advanced functionalities (`xarray`, `discretize`, `matplotlib`, `h5py`, `empymod`, `scooby`). Consult the installation notes in the [manual](#) for more information regarding installation, requirements, and soft dependencies.



## CHAPTER 4

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### Citation

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If you publish results for which you used *emg3d*, please give credit by citing [Werthmüller et al. \(2019\)](#):

Werthmüller, D., W. A. Mulder, and E. C. Slob, 2019, emg3d: A multigrid solver for 3D electromagnetic diffusion: *Journal of Open Source Software*, 4(39), 1463; DOI: [10.21105/joss.01463](#).

All releases have a Zenodo-DOI, which can be found on [10.5281/zenodo.3229006](#).

See *CREDITS* for the history of the code.



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## License information

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## 5.1 Getting started

The code `emg3d` ([WeMS19]) is a three-dimensional modeller for electromagnetic (EM) diffusion as used, for instance, in controlled-source EM (CSEM) surveys frequently applied in the search for, amongst other, groundwater, hydrocarbons, and minerals.

The core of the code is primarily based on [Mul06], [Mul07], and [Mul08]. You can read more about the background of the code in the chapter *Credits*. An introduction to the underlying theory of multigrid methods is given in the chapter *Theory*, and further literature is provided in the *References*.

### 5.1.1 Installation

You can install `emg3d` either via `conda`:

```
conda install -c conda-forge emg3d
```

or via `pip`:

```
pip install emg3d
```

Minimum requirements are Python version 3.7 or higher and the modules `scipy` and `numba`. Various other packages are recommended or required for some advanced functionalities, namely:

- `xarray`: For the `Survey` class (many sources and receivers at once).
- `discretize`: For advanced meshing tools (fancy mesh-representations and plotting utilities).
- `matplotlib`: To use the plotting utilities within `discretize`.
- `h5py`: Save and load data in the HDF5 format.
- `empyod`: Time-domain modelling (`utils.Fourier`).
- `scooby`: For the version and system report (`emg3d.Report()`).

If you are new to Python we recommend using a Python distribution, which will ensure that all dependencies are met, specifically properly compiled versions of NumPy and SciPy; we recommend using [Anaconda](#). If you install Anaconda you can simply start the *Anaconda Navigator*, add the channel `conda-forge` and `emg3d` will appear in the package list and can be installed with a click.

Using NumPy and SciPy with the Intel Math Kernel Library (*mkl*) can significantly improve computation time. You can check if `mkl` is used via `conda list`: The entries for the BLAS and LAPACK libraries should contain something with `mkl`, not with `openblas`. To enforce it you might have to create a file `pinned`, containing the line `libblas[build=*mkl]` in the folder `path-to-your-conda-env/conda-meta/`.

## 5.1.2 Basic Example

Here we show a *very* basic example. To see some more realistic models have a look at the [gallery](#). This particular example is also there, with some further explanations and examples to show how to plot the model and the data; see «[Minimum working example](#)». It also contains an example without using `discretize`.

First, we load `emg3d` and `discretize` (to create a mesh), along with `numpy`:

```
>>> import emg3d
>>> import discretize
>>> import numpy as np
```

First, we define the mesh (see `discretize.TensorMesh` for more info). In reality, this task requires some careful considerations. E.g., to avoid edge effects, the mesh should be large enough in order for the fields to dissipate, yet fine enough around source and receiver to accurately model them. This grid is too small, but serves as a minimal example.

```
>>> grid = discretize.TensorMesh(
>>>     [[(25, 10, -1.04), (25, 28), (25, 10, 1.04)],
>>>      [(50, 8, -1.03), (50, 16), (50, 8, 1.03)],
>>>      [(30, 8, -1.05), (30, 16), (30, 8, 1.05)]],
>>>     x0='CCC')
>>> print(grid)
```

TensorMesh: 49,152 cells

dir	nC	MESH EXTENT		CELL WIDTH		FACTOR
		min	max	min	max	
x	48	-662.16	662.16	25.00	37.01	1.04
y	32	-857.96	857.96	50.00	63.34	1.03
z	32	-540.80	540.80	30.00	44.32	1.05

Next we define a very simple fullspace model with  $\rho_x = 1.5 \Omega \text{ m}$ ,  $\rho_y = 1.8 \Omega \text{ m}$ , and  $\rho_z = 3.3 \Omega \text{ m}$ . The source is an x-directed dipole at the origin, with a 10 Hz signal of 1 A.

```
>>> model = emg3d.models.Model(
>>>     grid, property_x=1.5, property_y=1.8, property_z=3.3)
>>> sfield = emg3d.fields.get_source_field(
>>>     grid, src=[0, 0, 0, 0, 0], freq=10.0)
```

Now we can compute the electric field with `emg3d`:

```
>>> efield = emg3d.solve(grid, model, sfield, verb=4)

:: emg3d START :: 15:24:40 :: v0.13.0

MG-cycle      : 'F'                sslsolver : False
semicoarsening : False [0]          tol        : 1e-06
linerelaxation : False [0]          maxit      : 50
nu_{i,1,c,2}  : 0, 2, 1, 2         verb       : 3
```

(continues on next page)



(continued from previous page)

```

Original grid : 48 x 32 x 32    => 49,152 cells
Coarsest grid : 3 x 2 x 2      => 12 cells
Coarsest level : 4 ; 4 ; 4

[hh:mm:ss]  rel. error          [abs. error, last/prev]  1 s

      h_
     2h_ \
    4h_  \  / \
   8h_   \ / \ / \
  16h_    \/  \/  \/

[11:18:17]  2.623e-02  after  1 F-cycles  [1.464e-06, 0.026]  0 0
[11:18:17]  2.253e-03  after  2 F-cycles  [1.258e-07, 0.086]  0 0
[11:18:17]  3.051e-04  after  3 F-cycles  [1.704e-08, 0.135]  0 0
[11:18:17]  5.500e-05  after  4 F-cycles  [3.071e-09, 0.180]  0 0
[11:18:18]  1.170e-05  after  5 F-cycles  [6.531e-10, 0.213]  0 0
[11:18:18]  2.745e-06  after  6 F-cycles  [1.532e-10, 0.235]  0 0
[11:18:18]  6.873e-07  after  7 F-cycles  [3.837e-11, 0.250]  0 0

> CONVERGED
> MG cycles      : 7
> Final rel. error : 6.873e-07

:: emg3d END      :: 15:24:42 :: runtime = 0:00:02

```

So the computation required seven multigrid F-cycles and took just a bit more than 2 seconds. It was able to coarsen in each dimension four times, where the input grid had 49,152 cells, and the coarsest grid had 12 cells.

### 5.1.3 Related ecosystem

To create advanced meshes it is recommended to use [discretize](#) from the SimPEG framework. It also comes with some neat plotting functionalities to plot model parameters and resulting fields. Furthermore, it can serve as a link to use [PyVista](#) to create nice 3D plots even within a notebook.

Projects which can be used to compare or validate the results are, e.g., [empymod](#) for layered models or [SimPEG](#) for 3D models. It is also possible to create a geological model with [GemPy](#) and, again via [discretize](#), move it to [emg3d](#) to compute CSEM responses for it.

Have a look at the [gallery](#) for many examples of how to use [emg3d](#) together with the mentioned projects and more!

### 5.1.4 Tips and Tricks

The function `emg3d.solver.solve()` is the main entry point, and it takes care whether multigrid is used as a solver or as a preconditioner (or not at all), while the actual multigrid solver is `emg3d.solver.multigrid()`. Most input parameters for `emg3d.solver.solve()` are sufficiently described in its docstring. Here a few additional information.

- You can input any three-dimensional tensor mesh into `emg3d`. However, the implemented multigrid technique works with the existing nodes, meaning there are no new nodes created as coarsening is done by combining adjacent cells. The more times the grid dimension can be divided by two the better it is suited for MG. Ideally, the number should be dividable by two a few times and the dimension of the coarsest grid should be a low prime number  $p$ , for which good sizes can then be computed with  $p2^n$ . Good grid sizes (in each direction) up to 1024 are
  - $22^{3,4,\dots,9}$ : 16, 32, 64, 128, 256, 512, 1024,
  - $32^{3,4,\dots,8}$ : 24, 48, 96, 192, 384, 768,
  - $52^{3,4,\dots,7}$ : 40, 80, 160, 320, 640,

–  $72^{3,4,\dots,7}$ : 56, 112, 224, 448, 896,

and preference decreases from top to bottom row (stick to the first two or three rows if possible). Good grid sizes in sequential order, excluding  $p=7$ : 16, 24, 32, 40, 48, 64, 80, 96, 128, 160, 192, 256, 320, 384, 512, 640, 768, 1024. You can get this list via `emg3d.meshes.good_mg_cell_nr()`.

- The multigrid method can be used as a solver or as a preconditioner, for instance for BiCGSTAB. Using multigrid as a preconditioner for BiCGSTAB together with semicoarsening and line relaxation is the most stable version, but expensive, and therefore only recommended on highly stretched grids. Which combination of solver is best (fastest) depends to a large extent on the grid stretching, but also on anisotropy and general model complexity. See «Parameter tests» in the gallery for an example how to run some tests on your particular problem.

### 5.1.5 Contributing and Roadmap

New contributions, bug reports, or any kind of feedback is always welcomed! Have a look at the [Roadmap-project](#) to get an idea of things that could be implemented. The GitHub [issues](#) and [PR's](#) are also a good starting point. The best way for interaction is at <https://github.com/empymod> or by joining the [Slack channel](#) «em-x-d» of SimPEG. If you prefer to get in touch outside of GitHub/Slack use the contact form on <https://werthmuller.org>.

To install emg3d from source, you can download the latest version from GitHub and install it in your python distribution via:

```
python setup.py install
```

Please make sure your code follows the pep8-guidelines by using, for instance, the python module `flake8`, and also that your code is covered with appropriate tests. Just get in touch if you have any doubts.

### 5.1.6 Tests and benchmarks

The modeller comes with a test suite using `pytest`. If you want to run the tests, just install `pytest` and run it within the `emg3d-top`-directory.

```
> pytest --cov=emg3d --flake8
```

It should run all tests successfully. Please let us know if not!

Note that installations of `em3gd` via `conda` or `pip` do not have the test-suite included. To run the test-suite you must download `emg3d` from GitHub.

There is also a benchmark suite using *airspeed velocity*, located in the [empymod/emg3d-asv](#)-repository. The results of my machine can be found in the [empymod/emg3d-bench](#), its rendered version at [empymod.github.io/emg3d-asv](#).

### 5.1.7 License

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<https://www.apache.org/licenses/LICENSE-2.0>

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## 5.2 Multi-what?

If you have never heard of the multigrid method before you might ask yourself “*multi-what?*” The following is an intent to describe the multigrid method without the maths; just some keywords and some figures. **It is a heavily simplified intro, using a 2D grid for simplicity.** Have a look at the *Theory*-section for more details. A good, four-page intro with some maths is given by [Mul11]. More in-depth information can be found, e.g., in [BrHM00], [Hack85], and [Wess91].

The multigrid method ([Fed064])

- is an iterative solver;
- scales almost linearly (CPU & RAM);
- can serve as a pre-conditioner or as a solver on its own.

The main driving motivation to use multigrid is the part about linear scaling.

### 5.2.1 Matrix-free solver

The implemented multigrid method is a matrix free solver, it never constructs the full matrix. This is how it achieves its relatively low memory consumption. To solve the system, it solves for all fields adjacent to one node, moves then to the next node, and so on until it reaches the last node, see Figure 5.1, where the red lines indicate the fields which are solved simultaneously per step (the fields on the boundaries are never computed, as they are assumed to be 0).

Figure 5.1:: The multigrid solver solves by default on a node-by-node basis.

Normally, you would have to do this over and over again to achieve a good approximate solution. multigrid typically does it only a few times per grid, typically 2 times (one forward, one backward). This is why it is called **smoother**, as it only smoothes the error, it does not solve it. The implemented method for this is the *Gauss-Seidel* method.

Iterative solver which work in this matrix-free manner are typically **very fast at solving for the local problem**, hence at reducing the **high frequency error**, but **very slow at solving the global problem**, hence at reducing the **low frequency error**. High and low frequency errors are meant relatively to cell-size here.

### 5.2.2 Moving between different grids

The main thinking behind multigrid is now that we move to coarser grids. This has two advantages:

- Fewer cells means faster computation and less memory.
- Coarser grid size transforms lower frequency error to higher frequency error, relatively to cell size, which means faster convergence.

The implemented multigrid method simply joins two adjacent cells to get from finer to coarser grids, see Figure 5.2 for an example coarsening starting with a 16 cells by 16 cells grid.

Figure 5.2:: Example of the implemented coarsening scheme.

There are different approaches how to cycle through different grid sizes, see Figures 5.7 to 5.9. The downsampling from a finer grid to a coarser grid is often termed **restriction**, whereas the interpolation from a coarser grid to a finer grid is termed **prolongation**.

### 5.2.3 Specialities

The convergence rate of the multigrid method suffers on severely stretched grids or by models with strong anisotropy. Two techniques are implemented, **semicoarsening** (Figure 5.3) and **line relaxation** (Figure 5.4). Both require more CPU and higher RAM per grid than the standard multigrid, but they can improve the convergence rate, which then in turn improves the overall CPU time.

Figure 5.3:: Example of semicoarsening: The cell size is kept constant in one direction. The direction can be alternated between iterations.

Figure 5.4:: Example of line relaxation: The system is solved for all fields adjacent to a whole line of nodes simultaneously in some direction. The direction can be alternated between iterations.

## 5.3 Theory

The following provides an introduction to the theoretical foundation of the solver *emg3d*. More specific theory is covered in the docstrings of many functions, have a look at the [Other functions](#)-section or follow the links to the corresponding functions here within the theory. If you just want to use the solver, but do not care much about the internal functionality, then the function `emg3d.solver.solve()` is the only function you will ever need. It is the main entry point, and it takes care whether multigrid is used as a solver or as a preconditioner (or not at all), while the actual multigrid solver is `emg3d.solver.multigrid()`.

---

**Note:** This section is not an independent piece of work. Most things are taken from one of the following sources:

- [\[Mul06\]](#), pages 634-639:
  - The *Maxwell's equations* and *Discretisation* sections correspond with some adjustments and additions to pages 634-636.
  - The start of *The Multigrid Method* corresponds roughly to page 637.
  - Pages 638 and 639 are in parts reproduced in the code-docstrings of the corresponding functions.
- [\[BrHM00\]](#): This book is an excellent introduction to multigrid methods. Particularly the *Iterative Solvers* section is taken to a big extent from the book.

**Please consult these original resources for more details, and refer to them for citation purposes and not to this manual.** More in-depth information can also be found in, e.g., [\[Hack85\]](#) and [\[Wess91\]](#).

---

### 5.3.1 Maxwell's equations

Maxwell's equations in the presence of a current source  $\mathbf{J}_s$  are

$$\begin{aligned}\partial_t \mathbf{B}(\mathbf{x}, t) + \nabla \times \mathbf{E}(\mathbf{x}, t) &= 0, \\ \nabla \times \mathbf{H}(\mathbf{x}, t) - \partial_t \mathbf{D}(\mathbf{x}, t) &= \mathbf{J}_c(\mathbf{x}, t) + \mathbf{J}_s(\mathbf{x}, t),\end{aligned}\tag{5.1}$$

where the conduction current  $\mathbf{J}_c$  obeys Ohm's law,

$$\mathbf{J}_c(\mathbf{x}, t) = \sigma(\mathbf{x})\mathbf{E}(\mathbf{x}, t).\tag{5.2}$$

Here,  $\sigma(\mathbf{x})$  is the conductivity.  $\mathbf{E}(\mathbf{x}, t)$  is the electric field and  $\mathbf{H}(\mathbf{x}, t)$  is the magnetic field. The electric displacement  $\mathbf{D}(\mathbf{x}, t) = \varepsilon(\mathbf{x})\mathbf{E}(\mathbf{x}, t)$  and the magnetic induction  $\mathbf{B}(\mathbf{x}, t) = \mu(\mathbf{x})\mathbf{H}(\mathbf{x}, t)$ . The dielectric constant or permittivity  $\varepsilon$  can be expressed as  $\varepsilon = \varepsilon_r \varepsilon_0$ , where  $\varepsilon_r$  is the relative permittivity and  $\varepsilon_0$  is the vacuum value. Similarly, the magnetic permeability  $\mu$  can be written as  $\mu = \mu_r \mu_0$ , where  $\mu_r$  is the relative permeability and  $\mu_0$  is the vacuum value.

The magnetic field can be eliminated from Equation (5.1), yielding the second-order parabolic system of equations,

$$\varepsilon \partial_{tt} \mathbf{E} + \sigma \partial_t \mathbf{E} + \nabla \times \mu^{-1} \nabla \times \mathbf{E} = -\partial_t \mathbf{J}_s. \quad (5.3)$$

To transform from the time domain to the frequency domain, we substitute

$$\mathbf{E}(\mathbf{x}, t) = \frac{1}{2\pi} \int_{-\infty}^{\infty} \hat{\mathbf{E}}(\mathbf{x}, \omega) e^{-i\omega t} d\omega, \quad (5.4)$$

and use a similar representation for  $\mathbf{H}(\mathbf{x}, t)$ . The resulting system of equations is

$$-s\mu_0(\sigma + s\varepsilon)\hat{\mathbf{E}} - \nabla \times \mu_r^{-1} \nabla \times \hat{\mathbf{E}} = s\mu_0 \hat{\mathbf{J}}_s, \quad (5.5)$$

where  $s = -i\omega$ . The multigrid method converges in the case of the diffusive approximation (with its smoothing and approximation properties), but not in the high-frequency range (at least not in the implemented form of the multigrid method in `emg3d`). The code `emg3d` assumes therefore the diffusive approximation, hence only low frequencies are considered that obey  $|\omega\varepsilon| \ll \sigma$ . In this case we can set  $\varepsilon = 0$ , and Equation (5.5) simplifies to

$$-s\mu_0\sigma\hat{\mathbf{E}} - \nabla \times \mu_r^{-1} \nabla \times \hat{\mathbf{E}} = s\mu_0 \hat{\mathbf{J}}_s, \quad (5.6)$$

From here on, the hats are omitted. We use the perfectly electrically conducting boundary

$$\mathbf{n} \times \mathbf{E} = 0 \quad \text{and} \quad \mathbf{n} \cdot \mathbf{H} = 0, \quad (5.7)$$

where  $\mathbf{n}$  is the outward normal on the boundary of the domain.

The Maxwell's equations and Ohm's law are solved in the **frequency domain**. The **time-domain** solution can be obtained by taking the inverse Fourier transform.

---

**Note:** [Mul06] uses the time convention  $e^{-i\omega t}$ , see Equation (5.4), with  $s = -i\omega$ . However, the code `emg3d` uses the convention  $e^{i\omega t}$ , hence  $s = i\omega$ . This is the same convention as used in `empymod`, and commonly in CSEM.

---

## Laplace domain

It is also possible to solve the problem in the **Laplace domain**, by using a real value for  $s$  in Equation (5.6), instead of the complex value  $-i\omega$ . This simplifies the problem from complex numbers to real numbers, which accelerates the computation. It also improves the convergence rate, as the solution is a smoother function. The solver `emg3d.solver.solve()` is agnostic to the data type of the provided source field, and can solve for real and complex problems, hence frequency and Laplace domain. See the documentation of the functions `emg3d.fields.get_source_field()` and `emg3d.models.Model()` to see how you can use `emg3d` for Laplace-domain computations.

## 5.3.2 Discretisation

Equation (5.6) can be discretised by the finite-integration technique ([Wei77], [CIWe01]). This scheme can be viewed as a finite-volume generalization of [Yee66]'s scheme for tensor-product Cartesian grids with variable grid spacings. An error analysis for the constant-coefficient case ([MoSu94]) showed that both the electric and magnetic field components have second-order accuracy.

Consider a tensor-product Cartesian grid with nodes at positions  $(x_k, y_l, z_m)$ , where  $k = 0, \dots, N_x, l = 0, \dots, N_y$  and  $m = 0, \dots, N_z$ . There are  $N_x \times N_y \times N_z$  cells having these nodes as vertices. The cell centres are located at

$$\begin{aligned} x_{k+1/2} &= \frac{1}{2} (x_k + x_{k+1}), \\ y_{l+1/2} &= \frac{1}{2} (y_l + y_{l+1}), \\ z_{m+1/2} &= \frac{1}{2} (z_m + z_{m+1}). \end{aligned} \quad (5.8)$$

The material properties,  $\sigma$  and  $\mu_r$ , are assumed to be given as cell-averaged values. The electric field components are positioned at the edges of the cells, as shown in Figure 5.5, in a manner similar to Yee's scheme. The first component of the electric field  $E_{1,k+1/2,l,m}$  should approximate the average of  $E_1(x, y_l, z_m)$  over the edge from  $x_k$  to  $x_{k+1}$  at given  $y_l$  and  $z_m$ . Here, the average is defined as the line integral divided by the length of the integration interval. The other components,  $E_{2,k,l+1/2,m}$  and  $E_{3,k,l,m+1/2}$ , are defined in a similar way. Note that these averages may also be interpreted as point values at the midpoint of edges:

$$\begin{aligned} E_{1,k+1/2,l,m} &\simeq E_1(x_{k+1/2}, y_l, z_m), \\ E_{2,k,l+1/2,m} &\simeq E_2(x_k, y_{l+1/2}, z_m), \\ E_{3,k,l,m+1/2} &\simeq E_3(x_k, y_l, z_{m+1/2}). \end{aligned} \quad (5.9)$$

The averages and point-values are the same within second-order accuracy.

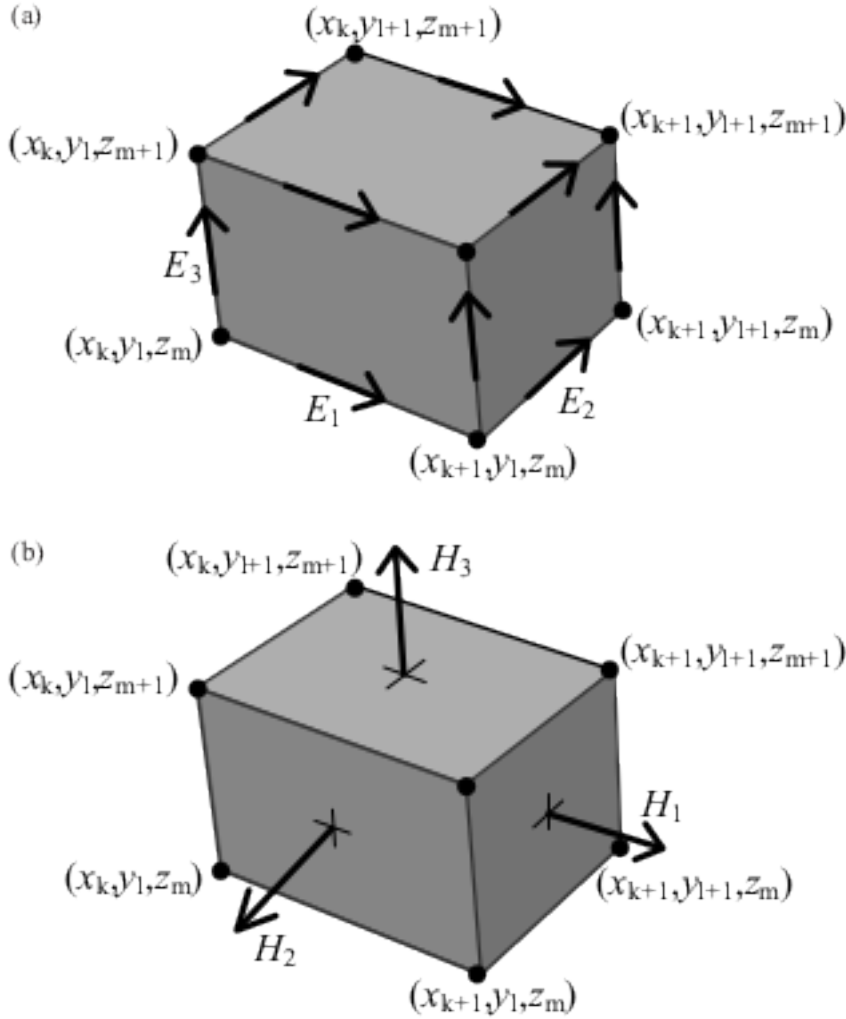


Figure 5.5:: (a) A grid cell with grid nodes and edge-averaged components of the electric field. (b) The face-averaged magnetic field components that are obtained by taking the curl of the electric field.

For the discretisation of the term  $-s\mu_0\sigma\mathbf{E}$  related to Ohm's law, dual volumes related to edges are introduced. For a given edge, the dual volume is a quarter of the total volume of the four adjacent cells. An example for  $E_1$  is shown in Figure 5.6(b). The vertices of the dual cell are located at the midpoints of the cell faces.

The volume of a normal cell is defined as

$$V_{k+1/2,l+1/2,m+1/2} = h_{k+1/2}^x h_{l+1/2}^y h_{m+1/2}^z, \quad (5.10)$$

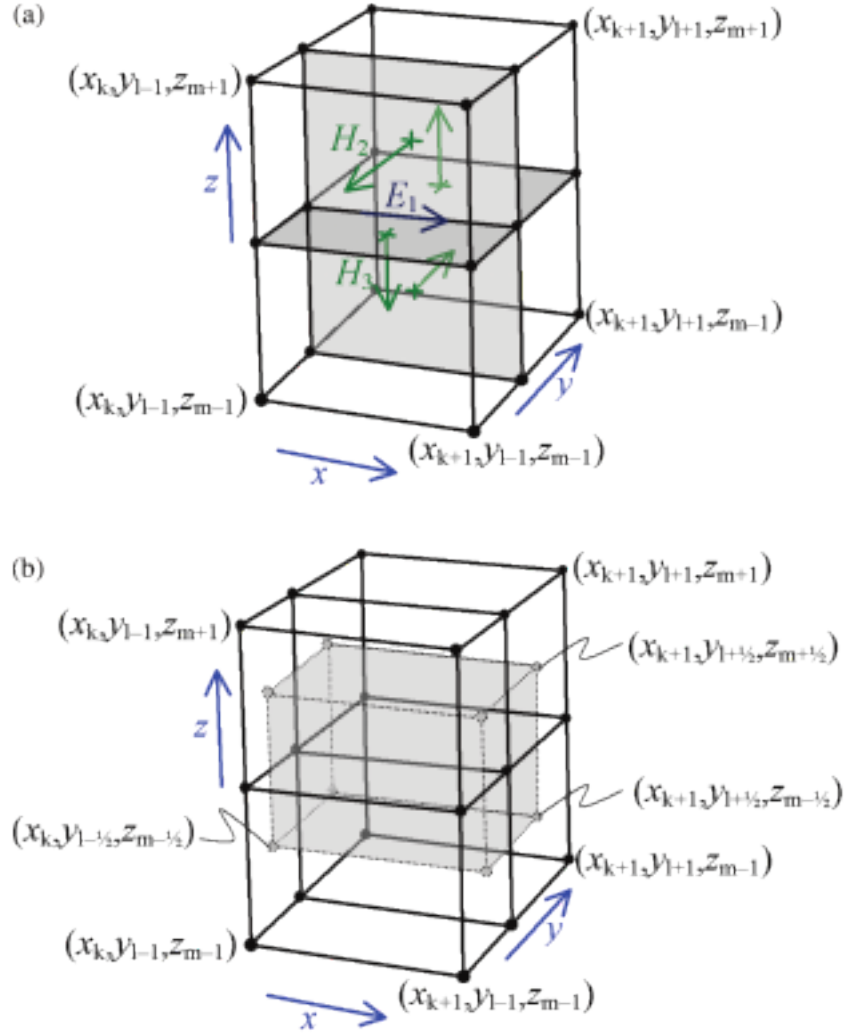


Figure 5.6:: The first electric field component  $E_{1,k,l,m}$  is located at the intersection of the four cells shown in (a). Four faces of its dual volume are sketched in (b). The first component of the curl of the magnetic field should coincide with the edge on which  $E_1$  is located. The four vectors that contribute to this curl are shown in (a). They are defined as normals to the four faces in (a). Before computing their curl, these vectors are interpreted as tangential components at the faces of the dual volume shown in (b). The curl is evaluated by taking the path integral over a rectangle of the dual volume that is obtained for constant  $x$  and by averaging over the interval  $[x_k, x_{k+1}]$ .

where

$$\begin{aligned} h_{k+1/2}^x &= x_{k+1} - x_k, \\ h_{l+1/2}^y &= y_{l+1} - y_l, \\ h_{m+1/2}^z &= z_{m+1} - z_m. \end{aligned} \quad (5.11)$$

For an edge parallel to the x-axis on which  $E_{1,k+1/2,l,m}$  is located, the dual volume is

$$V_{k+1/2,l,m} = \frac{1}{4} h_{k+1/2}^x \sum_{m_2=0}^1 \sum_{m_3=0}^1 h_{l-1/2+m_2}^y h_{m-1/2+m_3}^z. \quad (5.12)$$

With the definitions,

$$\begin{aligned} d_k^x &= x_{k+1/2} - x_{k-1/2}, \\ d_l^y &= y_{l+1/2} - y_{l-1/2}, \\ d_m^z &= z_{m+1/2} - z_{m-1/2}, \end{aligned} \quad (5.13)$$

we obtain

$$\begin{aligned} V_{k+1/2,l,m} &= h_{k+1/2}^x d_l^y d_m^z, \\ V_{k,l+1/2,m} &= d_k^x h_{l+1/2}^y d_m^z, \\ V_{k,l,m+1/2} &= d_k^x d_l^y h_{m+1/2}^z. \end{aligned} \quad (5.14)$$

Note that Equation (5.13) does not define  $d_k^x$ , etc., at the boundaries. We may simply take  $d_0^x = h_{1/2}^x$  at  $k = 0$ ,  $d_{N_x}^x = h_{N_x-1/2}^x$  at  $k = N_x$  and so on, or use half of these values as was done by [MoSu94].

The discrete form of the term  $-s\mu_0\sigma\mathbf{E}$  in Equation (5.6), with each component multiplied by the corresponding dual volume, becomes  $\mathcal{S}_{k+1/2,l,m} E_{1,k+1/2,l,m}$ ,  $\mathcal{S}_{k,l+1/2,m} E_{2,k,l+1/2,m}$  and  $\mathcal{S}_{k,l,m+1/2} E_{3,k,l,m+1/2}$  for the first, second and third components, respectively. Here  $\mathcal{S} = -s\mu_0\sigma V$  is defined in terms of cell-averages. At the edges parallel to the x-axis, an averaging procedure similar to (5.12) gives

$$\begin{aligned} \mathcal{S}_{k+1/2,l,m} &= \frac{1}{4} (\mathcal{S}_{k+1/2,l-1/2,m-1/2} + \mathcal{S}_{k+1/2,l+1/2,m-1/2} \\ &\quad + \mathcal{S}_{k+1/2,l-1/2,m+1/2} + \mathcal{S}_{k+1/2,l+1/2,m+1/2}). \end{aligned} \quad (5.15)$$

$\mathcal{S}_{k,l+1/2,m}$  and  $\mathcal{S}_{k,l,m+1/2}$  are defined in a similar way.

The curl of  $\mathbf{E}$  follows from path integrals around the edges that bound a face of a cell, drawn in Figure 5.5(a). After division by the area of the faces, the result is a face-averaged value that can be positioned at the centre of the face, as sketched in Figure 5.5(b). If this result is divided by  $i\omega\mu$ , the component of the magnetic field that is normal to the face is obtained. In order to find the curl of the magnetic field, the magnetic field components that are normal to faces are interpreted as tangential components at the faces of the dual volumes. For  $E_1$ , this is shown in Figure 5.6. For the first component of Equation (5.6) on the edge  $(k+1/2, l, m)$  connecting  $(x_k, y_l, z_m)$  and  $(x_{k+1}, y_l, z_m)$ , the corresponding dual volume comprises the set  $[x_k, x_{k+1}] \times [y_{l-1/2}, y_{l+1/2}] \times [z_{m-1/2}, z_{m+1/2}]$  having volume  $V_{k+1/2,l,m}$ .

The scaling by  $\mu_r^{-1}$  at the face requires another averaging step because the material properties are assumed to be given as cell-averaged values. We define  $\mathcal{M} = V\mu_r^{-1}$ , so

$$\mathcal{M}_{k+1/2,l+1/2,m+1/2} = h_{k+1/2}^x h_{l+1/2}^y h_{m+1/2}^z \mu_{r,k+1/2,l+1/2,m+1/2}^{-1} \quad (5.16)$$

for a given cell  $(k+1/2, l+1/2, m+1/2)$ . An averaging step in, for instance, the z-direction gives

$$\mathcal{M}_{k+1/2,l+1/2,m} = \frac{1}{2} (\mathcal{M}_{k+1/2,l+1/2,m-1/2} + \mathcal{M}_{k+1/2,l+1/2,m+1/2}) \quad (5.17)$$

at the face  $(k+1/2, l+1/2, m)$  between the cells  $(k+1/2, l+1/2, m-1/2)$  and  $(k+1/2, l+1/2, m+1/2)$ .

Starting with  $\mathbf{v} = \nabla \times \mathbf{E}$ , we have

$$\begin{aligned} v_{1,k,l+1/2,m+1/2} &= e_{l+1/2}^y (E_{3,k,l+1,m+1/2} - E_{3,k,l,m+1/2}) \\ &\quad - e_{m+1/2}^z (E_{2,k,l+1/2,m+1} - E_{2,k,l+1/2,m}), \\ v_{2,k+1/2,l,m+1/2} &= e_{m+1/2}^z (E_{1,k+1/2,l,m+1} - E_{1,k+1/2,l,m}) \\ &\quad - e_{k+1/2}^x (E_{3,k+1,l,m+1/2} - E_{3,k,l,m+1/2}), \\ v_{3,k+1/2,l+1/2,m} &= e_{k+1/2}^x (E_{2,k+1/2,l+1,m} - E_{1,k+1/2,l,m}) \\ &\quad - e_{l+1/2}^y (E_{1,k+1/2,l+1,m} - E_{1,k+1/2,l,m}). \end{aligned} \quad (5.18)$$



Here,

$$e_{k+1/2}^x = 1/h_{k+1/2}^x, \quad e_{l+1/2}^y = 1/h_{l+1/2}^y, \quad e_{m+1/2}^z = 1/h_{m+1/2}^z. \quad (5.19)$$

Next, we let

$$\begin{aligned} u_{1,k,l+1/2,m+1/2} &= \mathcal{M}_{k,l+1/2,m+1/2} v_{1,k,l+1/2,m+1/2}, \\ u_{2,k+1/2,l,m+1/2} &= \mathcal{M}_{k+1/2,l,m+1/2} v_{2,k+1/2,l+1/2,m}, \\ u_{3,k+1/2,l+1/2,m} &= \mathcal{M}_{k+1/2,l+1/2,m} v_{3,k+1/2,l+1/2,m}. \end{aligned} \quad (5.20)$$

Note that these components are related to the magnetic field components by

$$\begin{aligned} u_{1,k,l+1/2,m+1/2} &= i\omega\mu_0 V_{k,l+1/2,m+1/2} H_{1,k+1/2,l,m+1/2}, \\ u_{2,k+1/2,l,m+1/2} &= i\omega\mu_0 V_{k+1/2,l,m+1/2} H_{2,k+1/2,l,m+1/2}, \\ u_{3,k+1/2,l+1/2,m} &= i\omega\mu_0 V_{k+1/2,l+1/2,m} H_{3,k+1/2,l+1/2,m}, \end{aligned} \quad (5.21)$$

where

$$\begin{aligned} V_{k,l+1/2,m+1/2} &= d_k^x h_{l+1/2}^y h_{m+1/2}^z, \\ V_{k+1/2,l,m+1/2} &= h_{k+1/2}^x d_l^y h_{m+1/2}^z, \\ V_{k+1/2,l+1/2,m} &= h_{k+1/2}^x h_{l+1/2}^y d_m^z. \end{aligned} \quad (5.22)$$

The discrete representation of the source term  $i\omega\mu_0 \mathbf{J}_s$ , multiplied by the appropriate dual volume, is

$$\begin{aligned} s_{1,k+1/2,l,m} &= i\omega\mu_0 V_{k+1/2,l,m} J_{1,k+1/2,l,m}, \\ s_{2,k,l+1/2,m} &= i\omega\mu_0 V_{k,l+1/2,m} J_{2,k,l+1/2,m}, \\ s_{3,k,l,m+1/2} &= i\omega\mu_0 V_{k,l,m+1/2} J_{3,k,l,m+1/2}. \end{aligned} \quad (5.23)$$

Let the residual for an arbitrary electric field that is not necessarily a solution to the problem be defined as

$$\mathbf{r} = V \left( i\omega\mu_0 \mathbf{J}_s + -s\mu_0 \sigma \mathbf{E} - \nabla \times \mu_r^{-1} \nabla \times \mathbf{E} \right). \quad (5.24)$$

Its discretisation is

$$\begin{aligned} r_{1,k+1/2,l,m} &= s_{1,k+1/2,l,m} + \mathcal{S}_{k+1/2,l,m} E_{1,k+1/2,l,m} \\ &\quad - \left[ e_{l+1/2}^y u_{3,k+1/2,l+1/2,m} - e_{l-1/2}^y u_{3,k+1/2,l-1/2,m} \right] \\ &\quad + \left[ e_{m+1/2}^z u_{2,k+1/2,l,m+1/2} - e_{m-1/2}^z u_{2,k+1/2,l,m-1/2} \right], \\ r_{2,k,l+1/2,m} &= s_{2,k,l+1/2,m} + \mathcal{S}_{k,l+1/2,m} E_{2,k,l+1/2,m} \\ &\quad - \left[ e_{m+1/2}^z u_{1,k,l+1/2,m+1/2} - e_{m-1/2}^z u_{1,k,l+1/2,m-1/2} \right] \\ &\quad + \left[ e_{k+1/2}^x u_{3,k+1/2,l+1/2,m} - e_{k-1/2}^x u_{3,k-1/2,l+1/2,m} \right], \\ r_{3,k,l,m+1/2} &= s_{3,k,l,m+1/2} + \mathcal{S}_{k,l,m+1/2} E_{3,k,l,m+1/2} \\ &\quad - \left[ e_{k+1/2}^x u_{2,k+1/2,l,m+1/2} - e_{k-1/2}^x u_{2,k-1/2,m+1/2} \right] \\ &\quad + \left[ e_{l+1/2}^y u_{1,k,l+1/2,m+1/2} - e_{l-1/2}^y u_{1,k,l-1/2,m+1/2} \right]. \end{aligned} \quad (5.25)$$

The weighting of the differences in  $u_1$ , etc., may appear strange. The reason is that the differences have been multiplied by the local dual volume. As already mentioned, the dual volume for  $E_{1,k,l,m}$  is shown in [Figure 5.6\(b\)](#).

For further details of the discretisation see [\[Mul06\]](#) or [\[Yee66\]](#). The actual meshing is done using `discretize` (part of the `SimPEG`-framework). The coordinate system of `discretize` uses a coordinate system where positive  $z$  is upwards.

The method is implemented in a matrix-free manner: the large sparse linear matrix that describes the discretised problem is never explicitly formed, only its action is evaluated on the latest estimate of the solution, thereby reducing storage requirements.

### 5.3.3 Iterative Solvers

The multigrid method is an iterative (or relaxation) method and shares as such the underlying idea of iterative solvers. We want to solve the linear equation system

$$A\mathbf{x} = \mathbf{b}, \quad (5.26)$$

where  $A$  is the  $n \times n$  system matrix and  $x$  the unknown. If  $v$  is an approximation to  $x$ , then we can define two important measures. One is the error  $e$

$$\mathbf{e} = \mathbf{x} - \mathbf{v}, \quad (5.27)$$

which magnitude can be measured by any standard vector norm, for instance the maximum norm and the Euclidean or 2-norm defined respectively, by

$$\|\mathbf{e}\|_{\infty} = \max_{1 \leq j \leq n} |e_j| \quad \text{and} \quad \|\mathbf{e}\|_2 = \sqrt{\sum_{j=1}^n e_j^2}.$$

However, as the solution is not known the error cannot be computed either. The second important measure, however, is a computable measure, the residual  $r$  (computed in `emg3d.solver.residual()`)

$$\mathbf{r} = \mathbf{b} - A\mathbf{v}. \quad (5.28)$$

Using Equation (5.27) we can rewrite Equation (5.26) as

$$A\mathbf{e} = \mathbf{b} - A\mathbf{v},$$

from which we obtain with Equation (5.28) the *Residual Equation*

$$A\mathbf{e} = \mathbf{r}. \quad (5.29)$$

The *Residual Correction* is given by

$$\mathbf{x} = \mathbf{v} + \mathbf{e}. \quad (5.30)$$

### 5.3.4 The Multigrid Method

---

**Note:** If you have never heard of multigrid methods before you might want to read through the [Multi-what?](#)-section.

---

Multigrid is a numerical technique for solving large, often sparse, systems of equations, using several grids at the same time. An elementary introduction can be found in [BrHM00]. The motivation for this approach follows from the observation that it is fairly easy to determine the local, short-range behaviour of the solution, but more difficult to find its global, long-range components. The local behaviour is characterized by oscillatory or rough components of the solution. The slowly varying smooth components can be accurately represented on a coarser grid with fewer points. On coarser grids, some of the smooth components become oscillatory and again can be easily determined.

The following constituents are required to carry out multigrid. First, a sequence of grids is needed. If the finest grid on which the solution is to be found has a constant grid spacing  $h$ , then it is natural to define coarser grids with spacings of  $2h$ ,  $4h$ , etc. Let the problem on the finest grid be defined by  $A^h \mathbf{x}^h = \mathbf{b}^h$ . The residual is  $\mathbf{r}^h = \mathbf{b}^h - A^h \mathbf{x}^h$  (see the corresponding function `emg3d.solver.residual()`, and for more details also the function `emg3d.core.amat_x()`). To find the oscillatory components for this problem, a smoother or relaxation scheme is applied. Such a scheme is usually based on an approximation of  $A^h$  that is easy to invert. After one or more smoothing steps (see the corresponding function `emg3d.solver.smoothing()`), say  $\nu_1$  in total, convergence will slow down because it is generally difficult to find the smooth, long-range components of the solution. At this point, the problem is mapped to a coarser grid, using a restriction operator  $I_h^{2h}$  (see

the corresponding function `emg3d.solver.restriction()`, and for more details, the functions `emg3d.core.restrict_weights()` and `emg3d.core.restrict()`. On the coarse-grid,  $\mathbf{b}^{2h} = \tilde{I}_h^{2h} \mathbf{r}^h$ . The problem  $\mathbf{r}^{2h} = \mathbf{b}^{2h} - A^{2h} \mathbf{x}^{2h} = 0$  is now solved for  $\mathbf{x}^{2h}$ , either by a direct method if the number of points is sufficiently small or by recursively applying multigrid. The resulting approximate solution needs to be interpolated back to the fine grid and added to the solution. An interpolation operator  $I_{2h}^h$ , usually called prolongation in the context of multigrid, is used to update  $\mathbf{x}^h := \mathbf{x}^h + I_{2h}^h \mathbf{x}^{2h}$  (see the corresponding function `emg3d.solver.prolongation()`). Here  $I_{2h}^h \mathbf{x}^{2h}$  is called the coarse-grid correction. After prolongation,  $\nu_2$  additional smoothing steps can be applied. This constitutes one multigrid iteration.

So far, we have not specified the coarse-grid operator  $A^{2h}$ . It can be formed by using the same discretisation scheme as that applied on the fine grid. Another popular choice,  $A^{2h} = \tilde{I}_h^{2h} A^h I_{2h}^h$ , has not been considered here. Note that the tilde is used to distinguish restriction of the residual from operations on the solution, because these act on elements of different function spaces.

If multigrid is applied recursively, a strategy is required for moving through the various grids. The simplest approach is the V-cycle shown in Figure 5.7 for the case of four grids. Here, the same number of pre- and post-smoothing steps is used on each grid, except perhaps on the coarsest. In many cases, the V-cycle does not solve the coarse-grid equations sufficiently well. The W-cycle, shown in Figure 5.8, will perform better in that case. In a W-cycle, the number of coarse-grid corrections is doubled on subsequent coarser grids, starting with one coarse-grid correction on the finest grid. Because of its cost, it is often replaced by the F-cycle (Figure 5.9). In the F-cycle, the number of coarse-grid corrections increases by one on each subsequent coarser grid.

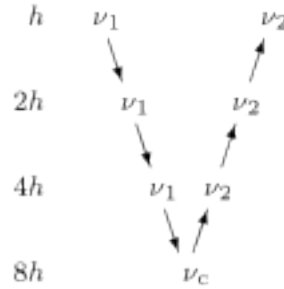


Figure 5.7:: V-cycle with  $\nu_1$  pre-smoothing steps and  $\nu_2$  post-smoothing steps. On the coarsest grid,  $\nu_c$  smoothing steps are applied or an exact solver is used. The finest grid has a grid spacing  $h$  and the coarsest  $8h$ . A single coarse-grid correction is computed for all grids but the coarsest.

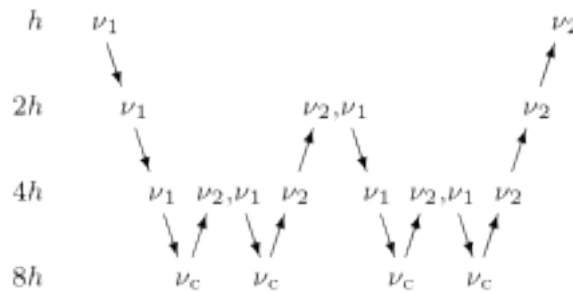


Figure 5.8:: W-cycle with  $\nu_1$  pre-smoothing steps and  $\nu_2$  post-smoothing steps. On each grid except the coarsest, the number of coarse-grid corrections is twice that of the underlying finer grid.

One reason why multigrid methods may fail to reach convergence is strong anisotropy in the coefficients of the governing partial differential equation or severely stretched grids (which has the same effect as anisotropy). In that case, more sophisticated smoothers or coarsening strategies may be required. Two strategies are currently implemented, *semicoarsening* and *line relaxation*, which can be used on their own or combined. Semicoarsening is when the grid is only coarsened in some directions. Line relaxation is when in some directions the whole gridlines of values are found simultaneously. If slow convergence is caused by just a few components of the solution, a Krylov subspace method can be used to remove them. In this way, multigrid is accelerated by a Krylov method. Alternatively, multigrid might be viewed as a preconditioner for a Krylov method.

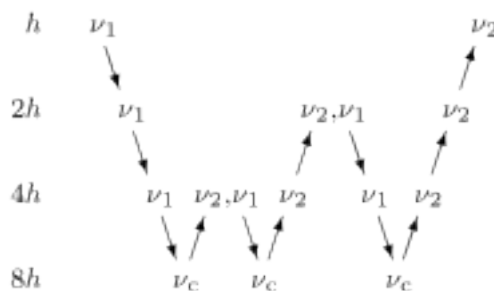


Figure 5.9:: F-cycle with  $\nu_1$  pre-smoothing steps and  $\nu_2$  post-smoothing steps. On each grid except the coarsest, the number of coarse-grid corrections increases by one compared to the underlying finer grid.

## Gauss-Seidel

The smoother implemented in `emg3d` is a Gauss-Seidel smoother. The Gauss-Seidel method solves the linear equation system  $A\mathbf{x} = \mathbf{b}$  iteratively using the following method:

$$\mathbf{x}^{(k+1)} = L_*^{-1} (\mathbf{b} - U\mathbf{x}^{(k)}) , \quad (5.31)$$

where  $L_*$  is the lower triangular component, and  $U$  the strictly upper triangular component,  $A = L_* + U$ . On the coarsest grid it acts as direct solver, whereas on the finer grid it acts as a smoother with only few iterations.

See the function `emg3d.solver.smoothing()`, and for more details, the functions `emg3d.core.gauss_seidel()`, `emg3d.core.gauss_seidel_x()`, `emg3d.core.gauss_seidel_y()`, `emg3d.core.gauss_seidel_z()`, and also `emg3d.core.blocks_to_amat()`.

## Choleski factorisation

The actual solver of the system  $A\mathbf{x} = \mathbf{b}$  is a non-standard Cholesky factorisation without pivoting for a symmetric, complex matrix  $A$  tailored to the problem of the multigrid solver, using only the main diagonal and five lower off-diagonals of the banded matrix  $A$ . The result is the same as simply using, e.g., `numpy.linalg.solve()`, but faster for the particular use-case of this code.

See `emg3d.core.solve()` for more details.

## 5.4 CPU & RAM

The multigrid method is attractive because it shows optimal scaling for both runtime and memory consumption. In the following are a few notes regarding memory and runtime requirements. It also contains information about what has been tried and what still could be tried in order to improve the current code.

### 5.4.1 Runtime

The [gallery](#) contains a script to do some testing with regards to runtime, see the [Tools Section](#). An example output of that script is shown in [Figure 5.10](#).

Figure 5.10:: Runtime as a function of cell size, which shows nicely the linear scaling of multigrid solvers (using a single thread).

The costliest functions (for big models) are:

- >90 %: `emg3d.solver.smoothing()` (`emg3d.core.gauss_seidel()`)
- <5 % each, in decreasing importance:

- `emg3d.solver.prolongation()` (`emg3d.solver.RegularGridProlongator`)
- `emg3d.solver.residual()` (`emg3d.core.amat_x()`)
- `emg3d.solver.restriction()`

Example with 262,144 / 2,097,152 cells (`nu_{i,1,c,2}=0,2,1,2`; `sslsolver=False`; `semicoarsening=True`; `linerelaxation=True`):

- 93.7 / 95.8 % smoothing
- 3.6 / 2.0 % prolongation
- 1.9 / 1.9 % residual
- 0.6 / 0.4 % restriction

The rest can be ignored. For small models, the percentage of smoothing goes down and of prolongation and restriction go up. But then the modeller is fast anyway.

`emg3d.core.gauss_seidel()` and `emg3d.core.amat_x()` are written in numba; jitting `emg3d.solver.RegularGridProlongator` turned out to not improve things, and many functions used in the restriction are jitted too. The costliest functions (RAM- and CPU-wise) are therefore already written in numba.

**Any serious attempt to improve the speed will have to tackle the smoothing itself.**

#### Things which could be tried

- Not much has been tested with the numba-options `parallel`; `prange`; and `nogil`.
- There might be an additional gain by making `emg3d.meshes.TensorMesh`, `emg3d.models.Model`, and `emg3d.fields.Field` instances jitted classes.

#### Things which have been tried

- One important aspect of the smoothing part is the memory layout. `emg3d.core.gauss_seidel()` and `emg3d.core.gauss_seidel_x()` are ideal for F-arrays (loop z-y-x, hence slowest to fastest axis). `emg3d.core.gauss_seidel_y()` and `emg3d.core.gauss_seidel_z()`, however, would be optimal for C-arrays. But copying the arrays to C-order and afterwards back is costlier in most cases for both CPU and RAM. The one possible and therefore implemented solution was to swap the loop-order in `emg3d.core.gauss_seidel_y()`.
- Restriction and prolongation information could be saved in a dictionary instead of recomputing it every time. Turns out to be not worth the trouble.
- Rewrite `emg3d.RegularGridInterpolator` as jitted function, but the iterator approach seems to be better for large grids.

## 5.4.2 Memory

Most of the memory requirement comes from storing the data itself, mainly the fields (source field, electric field, and residual field) and the model parameters (resistivity, eta, mu). For a big model, they sum up; e.g., almost 3 GB for an isotropic model with 256x256x256 cells.

The [gallery](#) contains a script to do some testing with regards to the RAM usage, see the [Tools Section](#). An example output of that script is shown in [Figure 5.11](#).

Figure 5.11:: RAM usage, showing the optimal behaviour of multigrid methods. “Data RAM” is the memory required by the fields (source field, electric field, residual field) and by the model parameters (resistivity; and eta, mu). “MG Base” is for solving one Gauss-Seidel iteration on the original grid. “MG full RAM” is for solving one multigrid F-Cycle.

The theory of multigrid says that in an ideal scenario, multigrid requires 8/7 (a bit over 1.14) the memory requirement of carrying out one Gauss-Seidel step on the finest grid. As can be seen in the figure, for models up to 2 million cells that holds pretty much, afterwards it becomes a bit worse.

However, for this estimation one has to run the model first. Another way to estimate the requirement is by starting from the RAM used to store the fields and parameters. As can be seen in the figure, for big models one is on the save side estimating the required RAM as 1.35 times the storage required for the fields and model parameters.

The figure also shows nicely the linear behaviour of multigrid; for twice the number of cells twice the memory is required (from a certain size onwards).

**Attempts at improving memory usage should focus on the difference between the red line (actual usage) and the dashed black line (1.14 x base usage).**

## 5.5 CLI interface

Command-line interface for certain specific tasks, such as forward modelling and gradient computation of the misfit function. The command is `emg3d`, consult the inbuilt help to get started:

```
emg3d --help
```

The CLI is driven by command-line parameters and a configuration file. The default configuration file is `emg3d.cfg`, but another name can be provided as first positional argument to `emg3d`. Note that arguments provided in the command line overwrite the settings in the configuration file.

### 5.5.1 Format of the config file

The shown values are the defaults. All values are commented out in this example; remove the comment signs to use them.

```
# Files
# -----
# If the files are provided without ending the suffix `.h5` will be appended.
# The log has the same name as `output`, but with the suffix `.log`.
[files]
# path = .                # Path (absolute or relative) to the data
# survey = survey.h5      # Also via `--survey`
# model = model.h5        # Also via `--model`
# output = emg3d_out.h5   # Also via `--output`
# store_simulation = False # Stores entire simulation in output if True

# Simulation parameters
# -----
# Input parameters for the `Simulation` class, except for `solver_opts`
# (defined in their own section), but including the parameter `min_offset`
# for `compute()`.
[simulation]
# max_workers = 4        # Also via `-n` or `--nproc`.
# gridding = single      # One grid for all sources and frequencies.
# min_offset = 0.0        # Only relevant if `observed=True` ( $r < r_{\min}$  set to NaN).

# Solver options
# -----
# Input parameters for the solver.
# See https://emg3d.readthedocs.io/en/stable/api/emg3d.solver.solve.html
# for a list of all parameters. The only parameters that cannot be provided
# here are grid, model, sfield, efield, and return_info.
#
# Note that currently sslsolver, semicoarsening, and linerelaxation only
# accept True/False through the CLI.
[solver_opts]
# sslsolver = True
# semicoarsening = True
```

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```

# linerelaxation = True
# verb = 0

# Gridding options
# -----
# Input parameters for the automatic gridding.
# See the description of `gridding_opts` and link therein in
# https://emg3d.readthedocs.io/en/stable/api/emg3d.simulations.Simulation.html
# for more details.
#
# List of lists: lists are comma-separated values, lists are separated by
# semi-colons.
#
# One of the limitation of the CLI is that `vector` has to be a string.
[gridding_opts]
# properties =          # list, e.g.: 0.3, 1, 1e5
# center =              # list, e.g.: 0, 0, 0
# cell_number =         # list, e.g.: 8, 16, 32, 64, 128
# min_width_pps =       # list, e.g.: 5, 3, 3
# expand =              # list, e.g.: 0.3, 1e8
# domain =              # list of lists, e.g.: -10000, 10000; None; None
# stretching =          # list of lists, e.g.: None; None; 1.05, 1.5
# min_width_limits =    # list of lists, e.g.: 10, 100; None; 50
# mapping =             # string, e.g.: Resistivity
# vector =              # string, e.g.: xy
# frequency =           # float, e.g.: 1.0
# searface =            # float, e.g.: 0.0
# max_buffer =          # float, e.g.: 100000.0
# lambda_factor =       # float, e.g.: 1.0
# verb =                # int, e.g.: 0
# lambda_from_center =  # bool, e.g.: False

# Data
# ----
# Select which sources, receivers, and frequencies of the survey are used. By
# default all data is used. These are comma-separated lists.
[data]
# sources = Tx02, Tx08, Tx14
# receivers = Rx01, Rx10
# frequencies = 0.5, 0.75

```

## 5.6 Gallery

The gallery with many examples can be found at [empymod.github.io/emg3d-gallery](https://empymod.github.io/emg3d-gallery).

## 5.7 References

## 5.8 Credits

This project was started by **Dieter Werthmüller**. Every contributor will be listed here and is considered to be part of «The emg3d Developers»:

- Dieter Werthmüller

Various bits got improved through discussions on Slack at **SWUNG** and at **SimPEG**, thanks to both communities. Special thanks to @jokva (general), @banesullivan (visualization), @joferkington (interpolation), and @jcapriot

(volume averaging).

### 5.8.1 Historical credits

The core of *emg3d* is a complete rewrite and redesign of the multigrid code by **Wim A. Mulder** ([Muld06], [Muld07], [Muld08], [MuWS08]), developed at Shell and at TU Delft. Various authors contributed to the original code, amongst others, **Tom Jönsthövel** ([JoOM06]; improved solver for strongly stretched grids), **Marwan Wirianto** ([WiMS10], [WiMS11]; computation of time-domain data), and **Evert C. Slob** ([SIHM10]; analytical solutions). The original code was written in Matlab, where the most time- and memory-consuming parts were sped up through mex-files (written in C). It included multigrid with or without BiCGSTAB, VTI resistivity, semi-coarsening, and line relaxation; the number of cells had to be powers of two, and coarsening was done only until the first dimension was at two cells. As such it corresponded roughly to *emg3d* v0.3.0.

See the *References* in the manual for the full citations and a more extensive list.

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**Note:** This software was initially (till 05/2021) developed at *Delft University of Technology* (<https://www.tudelft.nl>) within the **Gitaro.JIM** project funded through MarTERA as part of Horizon 2020, a funding scheme of the European Research Area (ERA-NET Cofund, <https://www.martera.eu>).

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## 5.9 Changelog

### 5.9.1 recent versions

#### v0.14.3 : Bug fix

2020-11-19

- Bug fix for `discretize`  $\geq 0.6.0$ .

#### v0.14.2 : Bug fix

2020-11-18

- Bug fix for Windows affecting `good_mg_cell_nr` (int32 issue).

#### v0.14.1 : Bug fix

2020-11-14

- Fix for `h5py`  $\geq 3.0$ .
- Improved docs re automatic gridding.

#### v0.14.0 : Automatic gridding

2020-11-07

The simulation class comes new with an automatic gridding functionality, which should make it much easier to compute CSEM data. With that the entire optimization routine was improved too. See the API docs for more info of the relevant implementation.

- `simulation`:
  - `Simulation`: New gridding options `'single'`, `'frequency'` `'source'`, and `'both'`; new default is `'single'`.



- `compute()` takes a new argument, `min_offset`. If `observed=True`, it will add Gaussian random noise according to the standard deviation of the data; it will set receivers responses below the minimum offset to NaN.
  - There is no longer a reference model.
  - `misfit` and `gradient` can now handle observations with NaN's.
- `survey`: A `Survey` has new attributes `standard_error`, `noise_floor`, and `relative_error`.
- `optimize`: Completely changed `misfit` and data-weighting to more sensible functions.
- `cli`:
  - As a consequence of the changes the `data_weight_opts` got removed.
  - New sections `[data]` to select the wanted data and `[gridding_opts]` for options of the automatic gridding.
  - Section `[simulation]` has a new parameter `min_offset` (for creating observed data).
  - Output has a new parameter `n_observations` if `misfit` or `gradient` were called, which is the number of observations that were used to compute the misfit.
- `meshes`:
  - New functions `construct_mesh`, `get_origin_widths`, `good_mg_cell_nr` and other, smaller helper routines.
  - Deprecated the old meshing routines `get_hx_h0`, `get_cell_numbers`, `get_stretched_h`, `get_domain`, `get_hx`; they will be removed in the future.
  - Default of `good_mg_cell_nr` changed, and the documentation (and verbosity) with regards to «good» number of cells was improved.
- Bug fixes:
  - `maps`: Fixed the mapping of the gradients (Conductivity is the only mapping that was not affected by this bug).
- Removed deprecated features:
  - `models.Model`: Removed parameters `res_{x;y;z}`.
  - `io.save`: Removed deprecated parameter `backend`.
  - `io.save`: Removed default, file extension has to be provided.

### v0.13.0 : CLI

2020-09-22

- New Module `cli` for command-line interaction:

The command-line interface can currently be used to forward model an entire `Simulation`, and also to compute the misfit of it with respect to some data and the gradient of the misfit function. See the section “CLI interface” in the documentation for more info.

*Note that, while `cli` (v0.13.0) and `optimize` (v0.12.0) are implemented, they are still in development and are likely going to change throughout the next two minor releases or so.*

- Other changes:
  - `solver`: Changes in verbosity for `emg3d.solve`:
    - \* New default verbosity is 1 (only warnings; before it was 2).
    - \* Verbosities `{-1;0;1}` remain unchanged.
    - \* Verbosities `{2;3;4}` => `{3;4;5}`.

- \* New verbosity 2: Only shows a one-liner at the end (plus warnings).
- survey and simulation: `to_file` and `from_file` have new a parameter name, to store and load with a particular name instead of the default `survey/simulation` (useful when storing, e.g., many surveys in one file).
- survey: stores new also the reference-data; different data (observed, reference) is contained in a data-dict when storing.
- simulation: takes new a `verb` parameter.
- optimize:
  - \* Gradient now possible for arbitrarily rotated sources and receivers.
  - \* Falls back to `synthetic` instead of `observed` now if reference not found.
- io: `np.bool_` are converted back to `bool` when loading.
- Re-arrange, improve, and update documentation.

## v0.12.0 : Survey & Simulation

2020-07-25

This is a big release with many new features, and unfortunately not completely backwards compatible. The main new features are the new **Survey** and **Simulation** classes, as well as some initial work for **optimization** (misfit, gradient). Also, a **Model** can now be a resistivity model, a conductivity model, or the logarithm (natural or base 10) thereof. Receivers can now be arbitrarily rotated, just as the sources. In addition to the existing **soft-dependencies** `empymod`, `discretize`, and `h5py` there are the new soft-dependencies `xarray` and `tqdm`; `discretize` is now much tighter integrated. For the new survey and simulation classes `xarray` is a required dependency. However, the only hard dependency remain `scipy` and `numba`, if you use `emg3d` purely as a solver. Data reading and writing has new a JSON-backend, in addition to the existing HDF5 and NumPy-backends.

In more detail:

- Modules:
  - surveys (**new**; requires `xarray`):
    - \* Class `surveys.Survey`, which combines sources, receivers, and data.
    - \* Class `surveys.Dipole`, which defines electric or magnetic point dipoles and finite length dipoles.
  - simulations (**new**; requires `xarray`; soft-dependency `tqdm`):
    - \* Class `simulations.Simulation`, which combines a survey with a model. A simulation computes the e-field (and h-field) asynchronously using `concurrent.futures`. This class will include automatic, source- and frequency-dependent gridding in the future. If `tqdm` is installed it displays a progress bar for the asynchronous computation. Note that the simulation class has still some limitations, consult the class documentation.
  - models:
    - \* Model instances take new the parameters `property_{x;y;z}` instead of `res_{x;y;z}`. The properties can be either resistivity, conductivity, or `log_{e;10}` thereof. What is actually provided has to be defined with the parameter mapping. By default, it remains resistivity, as it was until now. The keywords `res_{x;y;z}` are **deprecated**, but still accepted at the moment. The attributes `model.res_{x;y;z}` are still available too, but equally **deprecated**. However, it is **no longer possible to assign values to these attributes**, which is a **backwards incompatible** change.
    - \* A model knows now how to interpolate itself from its grid to another grid (`interpolate2grid`).
  - maps:

- \* **New** mappings for `models.Model` instances: The mappings take care of how to transform the investigation variable to conductivity and back, and how it affects its derivative.
- \* **New** interpolation routine `edges2cellaverages`.
- `fields`:
  - \* Function `get_receiver_response` (**new**), which returns the response for arbitrarily rotated receivers.
  - \* Improvements to `Field` and `SourceField`:
    - `_sval` and `_smu0` not stored any longer, derived from `_freq`.
    - `SourceField` is now using the `copy()` and `from_dict()` from its parents class `Field`.
- `io`:
  - \* File-format `json` (**new**), writes to a hierarchical, plain json file.
  - \* **Deprecated** the use of `backend`, it uses the file extension of `fname` instead.
  - \* This means `.npz` (instead of `numpy`), `.h5` (instead of `h5py`), and new `.json`.
  - \* New parameter `collect_classes`, which can be used to switch-on collection of the main classes in root-level dictionaries. By default, they are no longer collected (**changed**).
- `meshes`:
  - \* `meshes.TensorMesh` **new** inherits from `discretize` if installed.
  - \* Added `__eq__` to `models.TensorMesh` to compare meshes.
- `optimize` (**new**)
  - \* Functionalities related to inversion (data misfit, gradient, data weighting, and depth weighting). This module is in an early stage, and the API will likely change in the future. Current functions are `misfit`, `gradient` (using the adjoint-state method), and `data_weighting`. These functionalities are best accessed through the `Simulation` class.
- Dependencies:
  - `empymod` is now a soft dependency (no longer a hard dependency), only required for `utils.Fourier` (time-domain modelling).
  - Existing soft dependency `discretize` is now baked straight into `meshes`.
  - New soft dependency `xarray` for the `Survey` class (and therefore also for the `Simulation` class and the `optimize` module).
  - New soft dependency `tqdm` for nice progress bars in asynchronous computation.
- Deprecations and removals:
  - Removed deprecated functions `data_write` and `data_read`.
  - Removed all deprecated functions from `utils`.
- Miscellaneous:
  - Re-organise API-docs.
  - Much bookkeeping (improve error raising and checking; chaining errors, numpy types, etc).

## v0.11.0 : Refactor

2020-05-05

Grand refactor with new internal layout. Mainly splitting-up `utils` into smaller bits. Most functionalities (old names) are currently retained in `utils` and it should be mostly backwards compatible for now, but they are deprecated and will eventually be removed. Some previously deprecated functions were removed, however.

- Removed deprecated functions:
  - `emg3d.solver.solver` (use `emg3d.solver.solve` instead).
  - Aliases of `emg3d.io.data_write` and `emg3d.io.data_read` in `emg3d.utils`.
- Changes:
  - `SourceField` has now the same signature as `Field` (this might break your code if you called `SourceField` directly, with positional arguments, and not through `get_source_field`).
  - More functions and classes in the top namespace.
  - Replaced `core.l2norm` with `scipy.linalg.norm`, as SciPy 1.4 got the following PR: <https://github.com/scipy/scipy/pull/10397> (reason to raise minimum SciPy to 1.4).
  - Increased minimum required versions of dependencies to
    - \* `scipy>=1.4.0` (raised from 1.1, see note above)
    - \* `empymod>=2.0.0` (no min requirement before)
    - \* `numba>=0.45.0` (raised from 0.40)
- New layout
  - `njitted` -> `core`.
  - `utils` split in `fields`, `meshes`, `models`, `maps`, and `utils`.
- Bugfixes:
  - Fixed `to_dict`, `from_dict`, and `copy` for the `SourceField`.
  - Fixed `io` for `SourceField`, that was not implemented properly.

## 5.9.2 v0.8.0 - v0.10.x

### v0.10.1 : Zero Source

2020-04-29

- Bug fixes:
  - Checks now if provided source-field is zero, and exists gracefully if so, returning a zero electric field. Until now it failed with a division-by-zero error.
- Improvements:
  - Warnings: If `verb=1` it prints a warning in case it did not converge (it finished silently until now).
  - Improvements to docs (figures-scaling; intersphinx).
  - Adjust `Fields.pha` and `Fields.amp` in accordance with `empymod v2`: `.pha` and `.amp` are now methods; uses directly `empymod.utils.EMArray`.
  - Adjust tests for `empymod v2` (`Fields`, `Fourier`).

### v0.10.0 : Data persistence

2020-03-25

- New:

- New functions `emg3d.save` and `emg3d.load` to save and load all sort of `emg3d` instances. The currently implemented backends are `h5py` for `.h5`-files (default, but requires `h5py` to be installed) and `numpy` for `.npz`-files.
- Classes `emg3d.utils.Field`, `emg3d.utils.Model`, and `emg3d.utils.TensorMesh` have new methods `.copy()`, `.to_dict()`, and `.from_dict()`.
- `emg3d.utils.Model`: Possible to create new models by adding or subtracting existing models, and comparing two models (`+`, `-`, `==` and `!=`). New attributes `shape` and `size`.
- `emg3d.utils.Model` does not store the volume any longer (just `vnC`).
- Deprecations:
  - Deprecated `data_write` and `data_read`.
- Internal and bug fixes:
  - All I/O-related stuff moved to its own file `io.py`.
  - Change from `NUMBA_DISABLE_JIT` to use `py_func` for testing and coverage.
  - Bugfix: `emg3d.njitTED.restrict` did not store the `{x;y;z}`-field if `sc_dir` was `{4;5;6}`, respectively.

### v0.9.3 : Sphinx gallery

#### 2020-02-11

- Rename `solver.solver` to `solver.solve`; load `solve` also into the main namespace as `emg3d.solve`.
- Adjustment to termination criterion for *STAGNATION*: The current error is now compared to the last error of the same cycle type. Together with this the workaround for `sslsolver` when called with an initial `efield` introduced in v0.8.0 was removed.
- Adjustment to `utils.get_hx_h0` (this might change your boundaries): The computation domain is now computed so that the distance for the signal travelling from the source to the boundary and back to the most remote receiver is at least two wavelengths away. If this is within the provided domain, then now extra buffer is added around the domain. Additionally, the function has a new parameter `max_domain`, which is the maximum distance from the center to the boundary; defaults to 100 km.
- New parameter `log` for `utils.grid2grid`; if `True`, then the interpolation is carried out on a `log10`-scale.
- Change from the notebook-based `emg3d-examples-repo` to the sphinx-based `emg3d-gallery-repo`.

### v0.9.2 : Complex sources

#### 2019-12-26

- Strength input for `get_source_field` can now be complex; it also stores now the source location and its strength and moment.
- `get_receiver` can now take entire `Field` instances, and returns in that case `(fx, fy, fz)` at receiver locations.
- Krylov subspace solvers:
  - Solver now finishes in the middle of preconditioning cycles if tolerance is reached.
  - Solver now aborts if solution diverges or stagnates also for the SSL solvers; it fails and returns a zero field.
  - Removed `gmres` and `lgmres` from the supported SSL solvers; they do not work nice for this problem. Supported remain `bicgstab` (default), `cgs`, and `gcrotmk`.

- Various small things:
  - New attribute `Field.is_electric`, so the field knows if it is electric or magnetic.
  - New `verb`-possibility: `verb=-1` is a continuously updated one-liner, ideal to monitor large sets of computations or in inversions.
  - The returned `info` dictionary contains new keys:
    - \* `runtime_at_cycle`: accumulated total runtime at each cycle;
    - \* `error_at_cycle`: absolute error at each cycle.
  - Simple `__repr__` for `TensorMesh`, `Model`, `Fourier`, `Time`.
- Bugfixes:
  - Related to `get_hx_h0`, `data_write`, printing in `Fourier`.

### **v0.9.1 : VolumeModel**

**2019-11-13**

- New class `VolumeModel`; changes in `Model`:
  - `Model` now only contains resistivity, magnetic permeability, and electric permittivity.
  - `VolumeModel` contains the volume-averaged values `eta` and `zeta`; called from within `emg3d.solver.solver`.
  - Full wave equation is enabled again, via `epsilon_r`; by default it is set to `None`, hence diffusive approximation.
  - `Model` parameters are now internally stored as 1D arrays.
  - An {isotropic, VTI, HTI} initiated model can be changed by providing the missing resistivities.
- Bugfix: Up and till version 0.8.1 there was a bug. If resistivity was set with slices, e.g., `model.res[:, :, :5]=1e10`, it DID NOT update the corresponding `eta`. This bug was unintentionally fixed in 0.9.0, but only realised now.
- Various:
  - The log now lists the version of `emg3d`.
  - PEP8: internal imports now use absolute paths instead of relative ones.
  - Move from conda-channel `prisae` to `conda-forge`.
  - Automatic deploy for PyPi and `conda-forge`.

### **v0.9.0 : Fourier**

**2019-11-07**

- New routine:
  - `emg3d.utils.Fourier`, a class to handle Fourier-transform related stuff for time-domain modelling. See the example notebooks for its usage.
- Utilities:
  - Fields and returned receiver-arrays (`EMArray`) both have `amplitude (.amp)` and `phase (.pha)` attributes.
  - Fields have attributes containing frequency-information (`freq`, `smu0`).
  - New class `SourceField`; a subclass of `Field`, adding `vector` and `v{x, y, z}` attributes for the real valued source vectors.

- The `Model` is not frequency-dependent any longer and does NOT take a `freq`-parameter any more (currently it still takes it, but it is deprecated and will be removed in the future).
- `data_write` automatically removes `_vol` from `TensorMesh` instances and `_eta_{x,y,z}`, `_zeta` from `Model` instances. This makes the archives smaller, and they are not required, as they are simply reconstructed if needed.
- Internal changes:
  - The multigrid method, as implemented, only works for the diffusive approximation. Nevertheless, we always used `\sigma-i\omega\epsilon`, hence a complex number. This is now changed and `\epsilon` set to 0, leaving only `\sigma`.
  - Change time convention from `exp(-iwt)` to `exp(iwt)`, as used in `empymod` and commonly in CSEM. Removed the parameter `conjugate` from the solver, to simplify.
  - Change own private class variables from `__` to `_`.
  - `res` and `mu_r` are now checked to ensure they are  $>0$ ; `freq` is checked to ensure  $\neq 0$ .
- New dependencies and maintenance:
  - `empymod` is a new dependency.
  - Travis now checks all the url's in the documentation, so there should be no broken links down the road. (Check is allowed to fail, it is visual QC.)
- Bugfixes:
  - Fixes to the `setuptools_scm`-implementation (`MANIFEST.in`).

### v0.8.1 : setuptools\_scm

2019-10-22

- Implement `setuptools_scm` for versioning (adds git hashes for dev-versions).

### v0.8.0 : Laplace

2019-10-04

- Laplace-domain computation: By providing a negative `freq`-value to `utils.get_source_field` and `utils.Model`, the computation is carried out in the real Laplace domain  $s = \text{freq}$  instead of the complex frequency domain  $s = 2i\pi\text{freq}$ .
- New meshing helper routines (particularly useful for transient modelling where frequency-dependent/adaptive meshes are inevitable):
  - `utils.get_hx_h0` to get cell widths and origin for given parameters including a few fixed interfaces (center plus two, e.g. top anomaly, sea-floor, and sea-surface).
  - `utils.get_cell_numbers` to get good values of number of cells for given primes.
- Speed-up `njitted.volume_average` significantly thanks to @jcapriot.
- Bugfixes and other minor things:
  - Abort if l2-norm is NaN (only works for MG).
  - Workaround for the case where a `sslsolver` is used together with a provided initial `efield`.
  - Changed parameter `rho` to `res` for consistency reasons in `utils.get_domain`.
  - Changed parameter `h_min` to `min_width` for consistency reasons in `utils.get_stretched_h`.

### 5.9.3 v0.1.0 - v0.7.x

#### v0.7.1 : JOSS article

2019-07-17

- Version of the JOSS article, <https://doi.org/10.21105/joss.01463>.
- New function `utils.grid2grid` to move from one grid to another. Both functions (`utils.get_receiver` and `utils.grid2grid`) can be used for fields and model parameters (with or without extrapolation). They are very similar, the former taking coordinates (x, y, z) as new points, the latter one another `TensorMesh` instance.
- New jitted function `njitted.volume_average` for interpolation using the volume-average technique.
- New parameter `conjugate` in `solver.solver` to permit both Fourier transform conventions.
- Added `exit_status` and `exit_message` to `info_dict`.
- Add section `Related ecosystem` to documentation.

#### v0.7.0 : H-field

2019-07-05

- New routines:
  - `utils.get_h_field`: Small routine to compute the magnetic field from the electric field using Faraday's law.
  - `utils.get_receiver`: Small wrapper to interpolate a field at receiver positions. Added 3D spline interpolation; is the new default.
- Re-implemented the possibility to define isotropic magnetic permeabilities in `utils.Model`. Magnetic permeability is not tri-axially included in the solver currently; however, it would not be too difficult to include if there is a need.
- CPU-graph added on top of RAM-graph.
- Expand `utils.Field` to work with pickle/shelve.
- Jit `np.linalg.norm(njitted.l2norm)`.
- Use `scooby` (soft dependency) for versioning, rename `Version` to `Report` (backwards incompatible).
- Bug fixes:
  - Small bugfix introduced in ebd2c9d5: `sc_cycle` and `lr_cycle` was not updated any longer at the end of a cycle (only affected `sslsolver=True`).
  - Small bugfix in `utils.get_hx`.

#### v0.6.2 : CPU & RAM

2019-06-03

Further speed and memory improvements:

- Add *CPU & RAM*-page to documentation.
- Change loop-order from x-z-y to z-x-y in Gauss-Seidel smoothing with line relaxation in y-direction. Hence reversed lexicographical order. This results in a significant speed-up, as x is the fastest changing axis.
- Move total residual computation from `solver.residual` into `njitted.amat_x`.
- Simplifications in `utils`:
  - Simplify `utils.get_source_field`.



- Simplify `utils.Model`.
- Removed unused timing-stuff from early development.

### v0.6.1 : Memory

2019-05-28

Memory and speed improvements:

- Only compute residual and l2-norm when absolutely necessary.
- Inplace computations for `np.conjugate` in `solver.solver` and `np.subtract` in `solver.residual`.

### v0.6.0 : RegularGridInterpolator

2019-05-26

- Replace `scipy.interpolate.RegularGridInterpolator` with a custom tailored version of it (class:`emg3d.solver.RegularGridProlongator`); results in twice as fast prolongation.
- Simplify the fine-grid computation in prolongation without using `gridE*`; memory friendlier.
- Submission to JOSS.
- Add *Multi-what?*-page to documentation.
- Some major refactoring, particularly in `solver`.
- Removed `discretize` as hard dependency.
- Rename `rdir` and `ldir` (and related `p*dir`; `*cycle`) to the more descriptive `sc_dir` and `lr_dir`.

### v0.5.0 : Accept any grid size

2019-05-01

- First open-source version.
- Include RTD, Travis, Coveralls, Codacy, and Zenodo. No benchmarks yet.
- Accepts now *any* grid size (warns if a bad grid size for MG is provided).
- Coarsens now to the lowest level of each dimension, not only to the coarsest level of the smallest dimension.
- Combined `restrict_rx`, `restrict_ry`, and `restrict_rz` to `restrict`.
- Improve speed by passing pre-allocated arrays to jitted functions.
- Store `res_y`, `res_z` and corresponding `eta_y`, `eta_z` only if `res_y`, `res_z` were provided in initial call to `utils.model`.
- Change `zeta` to `v_mu_r`.
- Include rudimentary `TensorMesh`-class in `utils`; removes hard dependency on `discretize`.
- Bugfix: Take a provided `efield` into account; don't return if provided.

### v0.4.0 : Cholesky

2019-03-29

- Use `solve_chol` for everything, remove `solve_zlin`.
- Moved `mesh.py` and some functionalities from `solver.py` into `utils.py`.

- New mesh-tools. Should move to `discretize` eventually.
- Improved source generation tool. Might also move to `discretize`.
- `printversion` is now included in `utils`.
- Many bug fixes.
- Lots of improvements to tests.
- Lots of improvements to documentation. Amongst other, moved docs from `__init__.py` into the docs `rst`.

## v0.3.0 : Semicoarsening

2019-01-18

- Semicoarsening option.
- Number of cells must still be  $2^n$ , but  $n$  can be different in the x-, y-, and z-directions.
- Many other iterative solvers from `scipy.sparse.linalg` can be used. It seems to work fine with the following methods:
  - `scipy.sparse.linalg.bicgstab()`: BIConjugate Gradient STABilize;
  - `scipy.sparse.linalg.cgs()`: Conjugate Gradient Squared;
  - `scipy.sparse.linalg.gmres()`: Generalized Minimal RESidual;
  - `scipy.sparse.linalg.lgmres()`: Improvement of GMRES using alternating residual vectors;
  - `scipy.sparse.linalg.gcrotmk()`: GCROT: Generalized Conjugate Residual with inner Orthogonalization and Outer Truncation.
- The SciPy-solver or MG can be used all in combination or on its own, hence only MG, SciPy-solver with MG preconditioning, only SciPy-solver.

## v0.2.0 : Line relaxation

2019-01-14

- Line relaxation option.

## v0.1.0 : Initial

2018-12-28

- Standard multigrid with or without BiCGSTAB.
- Tri-axial anisotropy.
- Number of cells must be  $2^n$ , and  $n$  has to be the same in the x-, y-, and z-directions.

## 5.10 Maintainers Guide

### 5.10.1 Making a release

1. Update `CHANGELOG.rst`.
2. Push it to GitHub, create a release tagging it.

3. Tagging it on GitHub will automatically deploy it to PyPi, which in turn will create a PR for the conda-forge [feedstock](#). Merge that PR.
4. Check that:
  - [PyPi](#) deployed;
  - [conda-forge](#) deployed;
  - [Zenodo](#) minted a DOI;
  - [emg3d.rtfd.io](#) created a tagged version.

### 5.10.2 Useful things

- If there were changes to README, check it with:

```
python setup.py --long-description | rst2html.py --no-raw > index.html
```

- If unsure, test it first on testpypi (requires ~/.pypirc):

```
~/anaconda3/bin/twine upload dist/* -r testpypi
```

- If unsure, test the test-pypi for conda if the skeleton builds:

```
conda skeleton pypi --pypi-url https://test.pypi.io/pypi/ emg3d
```

- If it fails, you might have to install python3-setuptools:

```
sudo apt install python3-setuptools
```

### 5.10.3 CI

#### Automatic bits

- Testing on [Travis](#), includes:
  - Tests using `pytest`
  - Linting / code style with `pytest-flake8`
  - Ensure all http(s)-links work (`sphinx linkcheck`)
- Line-coverage with `pytest-cov` on [Coveralls](#)
- Code-quality on [Codacy](#)
- Manual on [ReadTheDocs](#)
- DOI minting on [Zenodo](#)

#### Manual things

- Benchmarks with [Airspeed Velocity](#) (`asv`)
- Gallery in [emg3d-gallery](#) (`sphinx-gallery`)

#### Automatically deploys if tagged

- [PyPi](#)
- `conda -c conda-forge`

## 5.11 Solver

Electromagnetic modeller in the diffusive limit (low frequencies) for 3D media with tri-axial electrical anisotropy. The matrix-free multigrid solver can be used as main solver or as preconditioner for one of the Krylov subspace methods implemented in `scipy.sparse.linalg`, and the governing equations are discretized on a staggered Yee grid. The code is written completely in Python using the `numpy/scipy`-stack, where the most time-consuming parts are sped-up through jitted `numba`-functions.

### 5.11.1 emg3d.solver Module

The actual multigrid solver routines. The most computationally intensive parts, however, are in the `emg3d.core` as `numba`-jitted functions.

#### Functions

<code>solve(grid, model, sfield[, efield, cycle, ...])</code>	Solver for 3D CSEM data with tri-axial electrical anisotropy.
<code>multigrid(grid, model, sfield, efield, var, ...)</code>	Multigrid solver for 3D controlled-source electromagnetic (CSEM) data.
<code>smoothing(grid, model, sfield, efield, nu, ...)</code>	Reducing high-frequency error by smoothing.
<code>restriction(grid, model, sfield, residual, ...)</code>	Downsampling of grid, model, and fields to a coarser grid.
<code>prolongation(grid, efield, cgrid, cefield, ...)</code>	Interpolating the electric field from coarse grid to fine grid.
<code>residual(grid, model, sfield, efield[, norm])</code>	Computing the residual.
<code>krylov(grid, model, sfield, efield, var)</code>	Krylov Subspace iterative solver for 3D CSEM data.

#### solve

`emg3d.solver.solve(grid, model, sfield, efield=None, cycle='F', sslsolver=False, semicoarsening=False, linerelaxation=False, verb=1, **kwargs)`  
Solver for 3D CSEM data with tri-axial electrical anisotropy.

The principal solver of `emg3d` is using the multigrid method as presented in [Mul06]. Multigrid can be used as a standalone solver, or as a preconditioner for an iterative solver from the `scipy.sparse.linalg`-library, e.g., `scipy.sparse.linalg.bicgstab()`. Alternatively, these Krylov subspace solvers can also be used without multigrid at all. See the `cycle` and `sslsolver` parameters.

Implemented are the *F*-, *V*-, and *W*-cycle schemes for multigrid (`cycle` parameter), and the amount of smoothing steps (initial smoothing, pre-smoothing, coarsest-grid smoothing, and post-smoothing) can be set individually (`nu_init`, `nu_pre`, `nu_coarse`, and `nu_post`, respectively). The maximum level of coarsening can be restricted with the `clevel` parameter.

Semicoarsening and line relaxation, as presented in [Mul07], are implemented, see the `semicoarsening` and `linerelaxation` parameters. Using the BiCGSTAB solver together with multigrid preconditioning with semicoarsening and line relaxation is slow but generally the most robust. Not using BiCGSTAB nor semicoarsening nor line relaxation is fast but may fail on stretched grids.

#### Parameters

- grid** [`emg3d.meshes.TensorMesh`] The grid. See `emg3d.meshes.TensorMesh`.
- model** [`emg3d.models.Model`] The model. See `emg3d.models.Model`.
- sfield** [`emg3d.fields.SourceField`] The source field. See `emg3d.fields.get_source_field()`.
- efield** [`emg3d.fields.Field`, optional] Initial electric field. It is initiated with zeroes

if not provided. A provided `efield` MUST have frequency information (initiated with `emg3d.fields.Field(..., freq)`).

If an initial `efield` is provided nothing is returned, but the final `efield` is directly put into the provided `efield`.

If an initial field is provided and a `sslsolver` is used, then it first carries out one multigrid cycle without semicoarsening nor line relaxation. The `sslsolver` is at times unstable with an initial guess, carrying out one MG cycle helps to stabilize it.

**cycle** [str; optional] Type of multigrid cycle. Default is 'F'.

- 'V': V-cycle, simplest version;
- 'W': W-cycle, most expensive version;
- 'F': F-cycle, sort of a compromise between 'V' and 'W';
- None: Does not use multigrid, only `sslsolver`.

If None, `sslsolver` must be provided, and the `sslsolver` will be used without multigrid pre-conditioning.

Comparison of V (left), F (middle), and W (right) cycles for the case of four grids (three relaxation and prolongation steps):



**sslsolver** [str, optional] A `scipy.sparse.linalg`-solver, to use with MG as pre-conditioner or on its own (if `cycle=None`). Default is False.

Current possibilities:

- True or 'bicgstab': BIConjugate Gradient STABILized `scipy.sparse.linalg.bicgstab()`;
- 'cgs': Conjugate Gradient Squared `scipy.sparse.linalg.cgs()`;
- 'gcrotmk': GCROT: Generalized Conjugate Residual with inner Orthogonalization and Outer Truncation `scipy.sparse.linalg.gcrotmk()`.

It does currently not work with 'cg', 'bicg', 'qmr', and 'minres' for various reasons (e.g., some require `rmatvec` in addition to `matvec`).

**semicoarsening** [int; optional] Semicoarsening. Default is False.

- True: Cycling over 1, 2, 3.
- 0 or False: No semicoarsening.
- 1: Semicoarsening in x direction.
- 2: Semicoarsening in y direction.
- 3: Semicoarsening in z direction.
- Multi-digit number containing digits from 0 to 3. Multigrid will cycle over these values, e.g., `semicoarsening=1213` will cycle over [1, 2, 1, 3].

**linerelaxation** [int; optional] Line relaxation. Default is False.

This parameter is not respected on the coarsest grid, except if it is set to 0. If it is bigger than zero line relaxation on the coarsest grid is carried out along all dimensions which have more than 2 cells.

- True: Cycling over [4, 5, 6].
- 0 or False: No line relaxation.

- 1: line relaxation in x direction.
- 2: line relaxation in y direction.
- 3: line relaxation in z direction.
- 4: line relaxation in y and z directions.
- 5: line relaxation in x and z directions.
- 6: line relaxation in x and y directions.
- 7: line relaxation in x, y, and z directions.
- Multi-digit number containing digits from 0 to 7. Multigrid will cycle over these values, e.g., `linerelaxation=1213` will cycle over [1, 2, 1, 3].

Note: Smoothing is generally done in lexicographical order, except for line relaxation in y direction; the reason is speed (memory access).

**verb** [int; optional] Level of verbosity (the higher the more verbose). Default is 1.

- 0: Nothing.
- 1: Warnings.
- 2: One-liner at the end.
- 3: Runtime and information about the method.
- 4: Additional information for each MG-cycle.
- 5: Everything (slower due to additional error computations).
- -1: One-liner (dynamically updated).

**\*\*kwargs** [Optional solver options:]

- *tol* : float

Convergence tolerance. Default is 1e-6.

Iterations stop as soon as the norm of the residual has decreased by this factor, relative to the residual norm obtained for a zero electric field.

- *maxit* : int

Maximum number of multigrid iterations. Default is 50.

If *sslsolver* is used, this applies to the *sslsolver*.

In the case that multigrid is used as a pre-conditioner for the *sslsolver*, the maximum iteration for multigrid is defined by the maximum length of the *linerelaxation* and *semicoarsening*-cycles.

- *nu\_init* : int

Number of initial smoothing steps, before MG cycle. Default is 0.

- *nu\_pre* : int

Number of pre-smoothing steps. Default is 2.

- *nu\_coarse* : int

Number of smoothing steps on coarsest grid. Default is 1.

- *nu\_post* : int

Number of post-smoothing steps. Default is 2.

- *clevel* : int

The maximum coarsening level can be different for each dimension and is, by default, automatically determined (*clevel*=-1). The parameter *clevel* can be used to restrict the maximum coarsening level in any direction by its value. Default is -1.

- *return\_info* : bool

If True, a dictionary is returned with runtime info (final norm and number of iterations of MG and the sslsolver).

## Returns

**efield** [*emg3d.fields.Field*] Resulting electric field. Is not returned but replaced in-place if an initial efield was provided.

**info\_dict** [dict] Dictionary with runtime info; only if *return\_info*=True.

Keys:

- *exit*: Exit status, 0=Success, 1=Failure;
- *exit\_message*: Exit message, check this if *exit*=1;
- *abs\_error*: Absolute error;
- *rel\_error*: Relative error;
- *ref\_error*: Reference error [*norm(sfield)*];
- *tol*: Tolerance (*abs\_error*<*ref\_error*\**tol*);
- *it\_mg*: Number of multigrid iterations;
- *it\_ssl*: Number of SSL iterations;
- *time*: Runtime (s).
- *runtime\_at\_cycle*: Runtime after each cycle (s).
- *error\_at\_cycle*: Absolute error after each cycle.

## Examples

```
>>> import emg3d
>>> import numpy as np
>>> # Create a simple grid, 8 cells of length 1 in each direction,
>>> # starting at the origin.
>>> grid = emg3d.TensorMesh(
>>>     [np.ones(8), np.ones(8), np.ones(8)],
>>>     x0=np.array([0, 0, 0]))
>>> # The model is a fullspace with tri-axial anisotropy.
>>> model = emg3d.Model(grid, property_x=1.5, property_y=1.8,
>>>     property_z=3.3, mapping='Resistivity')
>>> # The source is a x-directed, horizontal dipole at (4, 4, 4)
>>> # with a frequency of 10 Hz.
>>> sfield = emg3d.fields.get_source_field(
>>>     grid, src=[4, 4, 4, 0, 0], freq=10)
>>> # Compute the electric signal.
>>> efield = emg3d.solve(grid, model, sfield, verb=4)
>>> # Get the corresponding magnetic signal.
>>> hfield = emg3d.fields.get_h_field(grid, model, efield)
.
:: emg3d START :: 10:27:25 :: v0.9.1
.
MG-cycle      : 'F'                      sslsolver : False
semicoarsening : False [0]                tol       : 1e-06
```

(continues on next page)

(continued from previous page)

```

linerelaxation : False [0]          maxit      : 50
nu_{i,1,c,2}   : 0, 2, 1, 2        verb       : 4
Original grid   : 8 x 8 x 8          => 512 cells
Coarsest grid   : 2 x 2 x 2          => 8 cells
Coarsest level  : 2 ; 2 ; 2

.
[hh:mm:ss] rel. error                [abs. error, last/prev] 1 s
.
    h_
  2h_ \   /
    4h_ \/\/

.
[10:27:25] 2.284e-02 after 1 F-cycles [1.275e-06, 0.023] 0 0
[10:27:25] 1.565e-03 after 2 F-cycles [8.739e-08, 0.069] 0 0
[10:27:25] 1.295e-04 after 3 F-cycles [7.232e-09, 0.083] 0 0
[10:27:25] 1.197e-05 after 4 F-cycles [6.685e-10, 0.092] 0 0
[10:27:25] 1.233e-06 after 5 F-cycles [6.886e-11, 0.103] 0 0
[10:27:25] 1.415e-07 after 6 F-cycles [7.899e-12, 0.115] 0 0

.
> CONVERGED
> MG cycles      : 6
> Final rel. error : 1.415e-07

.
:: emg3d END      :: 10:27:25 :: runtime = 0:00:00

```

## multigrid

`emg3d.solver.multigrid(grid, model, sfield, efield, var, **kwargs)`

Multigrid solver for 3D controlled-source electromagnetic (CSEM) data.

Multigrid solver as presented in [Muld06], including semicoarsening and line relaxation as presented in and [Muld07].

- The electric field is stored in-place in *efield*.
- The number of multigrid cycles is stored in *var.it*.
- The current error (l2-norm) is stored in *var.l2*.
- The reference error (l2-norm of *sfield*) is stored in *var.l2\_refe*.

This function is called by `solve()`.

### Parameters

**grid** [`emg3d.meshes.TensorMesh`] The grid. See `emg3d.meshes.TensorMesh`.

**model** [`emg3d.models.VolumeModel`] The Model. See `emg3d.models.VolumeModel`.

**sfield** [`emg3d.fields.SourceField`] The source field. See `emg3d.fields.get_source_field()`.

**efield** [`emg3d.fields.Field`] The electric field. See `emg3d.fields.Field`.

**var** [`MGParameters` instance] As returned by `multigrid()`.

**\*\*kwargs** [Recursion parameters.] Do not use; only used internally by recursion; *level* (current coarsening level) and *new\_cycmax* (new maximum of MG cycles, takes care of V/W/F-cycling).



## smoothing

`emg3d.solver.smoothing(grid, model, sfield, efield, nu, lr_dir)`

Reducing high-frequency error by smoothing.

Solves the linear equation system  $Ax = b$  iteratively using the Gauss-Seidel method. This acts as smoother or, on the coarsest grid, as a direct solver.

This is a simple wrapper for the jitted computation in `emg3d.core.gauss_seidel()`, `emg3d.core.gauss_seidel_x()`, `emg3d.core.gauss_seidel_y()`, and `emg3d.core.gauss_seidel_z()` (@*njit* can not [yet] access class attributes). See these functions for more details and corresponding theory.

The electric fields are updated in-place.

This function is called by `multigrid()`.

### Parameters

- grid** [`emg3d.meshes.TensorMesh`] Input grid.
- model** [`emg3d.models.VolumeModel`] Input model.
- sfield** [`emg3d.fields.SourceField`] Input source field.
- efield** [`emg3d.fields.Field`] Input electric field.
- nu** [int] Number of Gauss-Seidel steps; odd numbers are forward, even numbers are reversed. E.g., `nu=2` is one symmetric Gauss-Seidel iteration, with a forward and a backward step.
- lr\_dir** [int] Direction of line relaxation {0, 1, 2, 3, 4, 5, 6, 7}.

## restriction

`emg3d.solver.restriction(grid, model, sfield, residual, sc_dir)`

Downsampling of grid, model, and fields to a coarser grid.

The restriction of the residual is used as source term for the coarse grid.

Corresponds to Equations 8 and 9 in [Mul06] and surrounding text. In the case of the restriction of the residual, this function is a wrapper for the jitted functions `emg3d.core.restrict_weights()` and `emg3d.core.restrict()` (@*njit* can not [yet] access class attributes). See these functions for more details and corresponding theory.

This function is called by `multigrid()`.

### Parameters

- grid** [`emg3d.meshes.TensorMesh`] Input grid.
- model** [`emg3d.models.VolumeModel`] Input model.
- sfield** [`emg3d.fields.SourceField`] Input source field.
- sc\_dir** [int] Direction of semicoarsening (0, 1, 2, or 3).

### Returns

- cgrid** [`emg3d.meshes.TensorMesh`] Coarse grid.
- cmodel** [`emg3d.models.VolumeModel`] Coarse model.
- csfield** [`emg3d.fields.SourceField`] Coarse source field. Corresponds to restriction of fine-grid residual.
- cefield** [`emg3d.fields.Field`] Coarse electric field, complex zeroes.

## prolongation

`emg3d.solver.prolongation(grid, efield, cgrid, cefield, sc_dir)`

Interpolating the electric field from coarse grid to fine grid.

The prolongation from a coarser to a finer grid is the inverse process of the restriction (`restriction()`) from a finer to a coarser grid. The interpolated values of the coarse grid electric field are added to the fine grid electric field, in-place. Piecewise constant interpolation is used in the direction of the field, and bilinear interpolation in the other two directions.

See Equation 10 in [Mul06] and surrounding text.

This function is called by `multigrid()`.

### Parameters

- grid, cgrid** [`emg3d.meshes.TensorMesh`] Fine and coarse grids.
- efield, cefield** [`emg3d.fields.Field`] Fine and coarse grid electric fields.
- sc\_dir** [int] Direction of semicoarsening (0, 1, 2, or 3).

## residual

`emg3d.solver.residual(grid, model, sfield, efield, norm=False)`

Computing the residual.

Returns the complete residual as given in [Mul06], page 636, middle of the right column:

$$\mathbf{r} = V \left( i\omega\mu_0\mathbf{J}_s + i\omega\mu_0\tilde{\sigma}\mathbf{E} - \nabla \times \mu_r^{-1} \nabla \times \mathbf{E} \right).$$

This is a simple wrapper for the jitted computation in `emg3d.core.amat_x()` (@*njit* can not [yet] access class attributes). See `emg3d.core.amat_x()` for more details and corresponding theory.

This function is called by `multigrid()`.

### Parameters

- grid** [`emg3d.meshes.TensorMesh`] Input grid.
- model** [`emg3d.models.VolumeModel`] Input model.
- sfield** [`emg3d.fields.SourceField`] Input source field.
- efield** [`emg3d.fields.Field`] Input electric field.
- norm** [bool] If True, the error (l2-norm) of the residual is returned, not the residual.

### Returns

- residual** [Field] Returned if `norm=False`. The residual field; `emg3d.fields.Field` instance.
- norm** [float] Returned if `norm=True`. The error (l2-norm) of the residual

## krylov

`emg3d.solver.krylov(grid, model, sfield, efield, var)`

Krylov Subspace iterative solver for 3D CSEM data.

Using a Krylov subspace iterative solver (defined in `var.ssolver`) implemented in SciPy with or without multigrid as a pre-conditioner ([Mul06]).

- The electric field is stored in-place in `efield`.
- The current error (l2-norm) is stored in `var.l2`.

- The reference error (l2-norm of sfield) is stored in `var:l2_refe`.

This function is called by `solve()`.

#### Parameters

**grid** [`emg3d.meshes.TensorMesh`] The grid. See `emg3d.meshes.TensorMesh`.

**model** [`emg3d.models.VolumeModel`] The Model. See `emg3d.models.VolumeModel`.

**sfield** [`emg3d.fields.SourceField`] The source field. See `emg3d.fields.get_source_field()`.

**efield** [`emg3d.fields.Field`] The electric field. See `emg3d.fields.Field`.

**var** [`MGParameters` instance] As returned by `multigrid()`.

## Classes

<code>MGParameters(verb, cycle, sslsolver, ...)</code>	Collect multigrid solver settings.
<code>RegularGridProlongator(x, y, cxy)</code>	Prolongate field from coarse to fine grid.

## MGParameters

**class** `emg3d.solver.MGParameters` (*verb: int, cycle: str, sslsolver: str, linerelaxation: int, semi-coarsening: int, vnC: tuple, tol: float = 1e-06, maxit: int = 50, nu\_init: int = 0, nu\_pre: int = 2, nu\_coarse: int = 1, nu\_post: int = 2, clevel: int = -1, return\_info: bool = False*)

Bases: `object`

Collect multigrid solver settings.

This dataclass is used by the main `solve()`-routine. See `solve()` for a description of the mandatory and optional input parameters and more information .

#### Returns

**var** [`class:MGParameters`] As required by `multigrid()`.

#### Attributes Summary

<code>clevel</code>	
<code>max_level</code>	Sets dimension-dependent level variable <code>clevel</code> .
<code>maxit</code>	
<code>nu_coarse</code>	
<code>nu_init</code>	
<code>nu_post</code>	
<code>nu_pre</code>	
<code>return_info</code>	
<code>tol</code>	

#### Methods Summary

<code>cprint(info, verbosity, **kwargs)</code>	Conditional printing.
<code>one_liner(l2_last[, last])</code>	Print continuously updated one-liner.

## Attributes Documentation

**clevel** = -1

**max\_level**

Sets dimension-dependent level variable *clevel*.

Requires at least two cells in each direction (for *nCx*, *nCy*, and *nCz*).

**maxit** = 50

**nu\_coarse** = 1

**nu\_init** = 0

**nu\_post** = 2

**nu\_pre** = 2

**return\_info** = False

**tol** = 1e-06

## Methods Documentation

**cprint** (*info*, *verbosity*, *\*\*kwargs*)

Conditional printing.

Prints *info* if *self.verb* > *verbosity*.

### Parameters

**info** [str] String to be printed.

**verbosity** [int] Verbosity of info.

**kwargs** [optional] Arguments passed to *print*.

**one\_liner** (*l2\_last*, *last=False*)

Print continuously updated one-liner.

### Parameters

**l2\_last** [float] Current error.

**last** [bool] If True, adds *exit\_message* and finishes line.

## RegularGridProlongator

**class** emg3d.solver.RegularGridProlongator (*x*, *y*, *cxy*)

Bases: object

Prolongate field from coarse to fine grid.

This is a heavily modified and adapted version of `scipy.interpolate.RegularGridInterpolator`.

The main difference (besides the pre-sets) is that this version allows to initiate an instance with the coarse and fine grids. This initialize will compute the required weights, and it has therefore only to be done once.

After this, interpolating values from the coarse to the fine grid can be carried out much faster.

Simplifications in comparison to `scipy.interpolate.RegularGridInterpolator`:

- No sanity checks what-so-ever.
- Only 2D data;
- `method='linear'`;

- `bounds_error=False`;
- `fill_value=None`.

It results in a speed-up factor of about 2, independent of grid size, for this particular case. The prolongation is the second-most expensive part of multigrid after the smoothing. Trying to improve this further might therefore be useful.

#### Parameters

**x, y** [ndarray] The x/y-coordinates defining the coarse grid.

**cxy** [ndarray of shape  $(\dots, 2)$ ] The  $([x], [y])$ .T-coordinates defining the fine grid.

### Methods Summary

<code>__call__</code> (values)	Return values of coarse grid on fine grid locations.
--------------------------------	------------------------------------------------------

### Methods Documentation

`__call__`(values)

Return values of coarse grid on fine grid locations.

#### Parameters

**values** [ndarray] Values corresponding to x/y-coordinates.

#### Returns

**result** [ndarray] Values corresponding to cxy-coordinates.

## 5.11.2 emg3d.core Module

The core functionalities, the most computationally demanding parts, of the `emg3d.solver` as just-in-time (jit) compiled functions using `numba`.

### Functions

<code>amat_x(rx, ry, rz, ex, ey, ez, eta_x, eta_y, ...)</code>	Residual without or with source term.
<code>blocks_to_amat(amat, bvec, middle, left, ...)</code>	Insert middle, left, and rhs into main arrays amat and bvec.
<code>gauss_seidel(ex, ey, ez, sx, sy, sz, eta_x, ...)</code>	Gauss-Seidel method.
<code>gauss_seidel_x(ex, ey, ez, sx, sy, sz, ...)</code>	Gauss-Seidel method with line relaxation in x-direction.
<code>gauss_seidel_y(ex, ey, ez, sx, sy, sz, ...)</code>	Gauss-Seidel method with line relaxation in y-direction.
<code>gauss_seidel_z(ex, ey, ez, sx, sy, sz, ...)</code>	Gauss-Seidel method with line relaxation in z-direction.
<code>restrict(crx, cry, crz, rx, ry, rz, wx, wy, ...)</code>	Restriction of residual from fine to coarse grid.
<code>restrict_weights(vectorN, vectorCC, h, ...)</code>	Restriction weights for the coarse-grid correction operator.
<code>solve(amat, bvec)</code>	Solve $Ax = b$ using a non-standard Cholesky factorisation.

#### amat\_x

`emg3d.core.amat_x(rx, ry, rz, ex, ey, ez, eta_x, eta_y, eta_z, zeta, hx, hy, hz)`  
Residual without or with source term.

Compute the residual as given in [Mul06] in middle of the right column on page 636, but without the source term:

$$\mathbf{r} = V \left( i\omega\mu_0\tilde{\sigma}\mathbf{E} - \nabla \times \mu_r^{-1} \nabla \times \mathbf{E} \right).$$

The computation is carried out in a matrix-free manner; on said page 636 (or in the *Theory* of the manual) are the various steps laid out to discretise the different parts, for instance involved curls. This can also be understood as the left-hand-side of  $Ax = b$ , as given in Equation 2 in [Mul06] (here without the cell volumes  $V$ ),

$$i\omega\mu_0\tilde{\sigma}\mathbf{E} - \nabla \times \zeta^{-1} \nabla \times \mathbf{E} = -i\omega\mu_0\mathbf{J}_s.$$

It can therefore be used as *matvec* to create a *LinearOperator*, which can be passed to a solver.

It is assumed that *ex*, *ey*, and *ez* have PEC boundaries; otherwise the output will not have PEC boundaries.

The residuals are subtracted in-place from *rx*, *ry*, and *rz*. That means that if *rx*, *ry*, and *rz* contain the source field, they will contain the total residual afterwards; if they are empty fields, they will contain the negative partial residual afterwards.

### Parameters

**rx, ry, rz** [ndarray] Source field or pre-allocated zero residual field in x-, y-, and z-directions.

**ex, ey, ez** [ndarray] Electric fields in x-, y-, and z-directions, as obtained from *emg3d.fields.Field*.

**eta\_x, eta\_y, eta\_z, zeta** [ndarray] VolumeModel parameters (multiplied by volumes) as obtained from *emg3d.models.VolumeModel()*.

**hx, hy, hz** [ndarray] Cell widths in x-, y-, and z-directions.

### blocks\_to\_amat

*emg3d.core.blocks\_to\_amat* (*amat*, *bvec*, *middle*, *left*, *rhs*, *im*, *nC*)

Insert *middle*, *left*, and *rhs* into main arrays *amat* and *bvec*.

The banded matrix *amat* contains the main diagonal and the first five lower off-diagonals. They are stored one column after the other, in a 6\*n ndarray.

The complete main matrix *amat* and the *middle* and *left* blocks are given by:

```
.-0
|X|\  0
0-.-0      left:  middle:  right:
|X|\      (not used)
0-.-0      0-      .-      0
|X|\      \      |X      |\
0-.-0
0  |X|
0-.
```

. 1\*1, - 4\*1, | 1\*4, X 4\*4, \ 4\*4 upper or lower

Both, *middle* and *left*, are 5x5 matrices. The corresponding right-hand-side *rhs* is filled into *bvec*. The matrices *left* and *middle* provided in a single call are horizontally aligned (not vertically). The sorting of *amat* (banded matrix) and *bvec* are given by:

amat (66,)	example: n = 11	bvec (11,)
-----		--
01	FIRST CALL	01
02 07	Only `middle` and `rhs`	02

(continues on next page)

(continued from previous page)

03 08 13		are used, not `left`.	03
04 09 14 19			04
05 10 15 20 25			05
-----			
0 11 16 21 26 31		SUBSEQUENT CALLS	06
12 17 22 27 32 37		(normal case)	07
18 23 28 33 38 43		Complete `left`,	08
24 29 34 39 44 49		`middle` and `rhs`	09
30 35 40 45 50 55		are used.	10
-----			
	0 41 46 51 56 61	LAST CALL	11
	0 0 0 0  0	Only top row of `left`	
	0 0 0  0	and the first elements	
	0 0  0	of `middle` and `rhs`	
	0  0	are used.	
-----			
	0		

Single zeros (0) show elements in amat which are 0, hence not used.  
Their location in amat can be deduced from their neighbours.

### Parameters

**amat** [ndarray] Main banded matrix (stored as array) of length 6\*n.

**bvec** [ndarray] Main right-hand-side of length n.

**middle** [ndarray] Middle block of size 5x5, as ndarray of length 25. Only the diagonal and the lower triangular part are used.

**left** [ndarray] Left block of size 5x5, as ndarray of length 25. Only the diagonal and the first row are used.

**rhs** [ndarray] Corresponding right-hand-side of length 5.

**im** [int] Current minus-index of direction of line relaxation, {ixm, iym, izm}.

**nC** [int] Total number of cells in direction of line relaxation, {nCx, nCy, nCz}.

### gauss\_seidel

emg3d.core.gauss\_seidel (ex, ey, ez, sx, sy, sz, eta\_x, eta\_y, eta\_z, zeta, hx, hy, hz, nu)  
Gauss-Seidel method.

Solves the linear equation system  $Ax = b$  iteratively using the following method:

$$\mathbf{x}^{(k+1)} = L_*^{-1} \left( \mathbf{b} - U\mathbf{x}^{(k)} \right),$$

where  $L_*$  is the lower triangular component, and  $U$  the strictly upper triangular component,  $A = L_* + U$ :

$$L_* = \begin{bmatrix} a_{11} & 0 & \cdots & 0 \\ a_{21} & a_{22} & \cdots & 0 \\ \vdots & \vdots & \ddots & \vdots \\ a_{n1} & a_{n2} & \cdots & a_{nn} \end{bmatrix}, \quad U = \begin{bmatrix} 0 & a_{12} & \cdots & a_{1n} \\ 0 & 0 & \cdots & a_{2n} \\ \vdots & \vdots & \ddots & \vdots \\ 0 & 0 & \cdots & 0 \end{bmatrix}.$$

On the coarsest grid it acts as direct solver, whereas on the fine grid it acts as a smoother with only few iterations, defined by  $\nu$  (nu). Odd numbers of nu use forward ordering, even numbers use backwards ordering. nu=2 is therefore one symmetric Gauss-Seidel iteration, one forward ordered iteration followed by one backward ordered iteration.

From [Muldo06]: The method proposed by [ArFW00] is chosen as a smoother. It selects one node of the grid and simultaneously solves for the six degrees of freedom on the six edges attached to the node. If

node  $(x_k, y_l, z_m)$  is selected, the six equations,  $r_{x;k\pm 1/2,l,m} = 0$ ,  $r_{y;k,l\pm 1/2,m} = 0$  and  $r_{z;k,l,m\pm 1/2} = 0$ , are solved for  $e_{x;k\pm 1/2,l,m}$ ,  $e_{y;k,l\pm 1/2,m}$  and  $e_{z;k,l,m\pm 1/2}$ . Here, this smoother is applied in a symmetric Gauss-Seidel fashion, following the lexicographical ordering of the nodes  $(x_k, y_l, z_m)$ , with fastest index  $k = 1, \dots, N_x - 1$ , intermediate index  $l = 1, \dots, N_y - 1$ , and slowest index  $m = 1, \dots, N_z - 1$ .

To actually solve the system of six equations a non-standard Cholesky factorisation is used, `solve()`.

Tangential components at the boundaries are assumed to be zero (PEC boundaries).

The result is stored in the provided electric fields `ex`, `ey`, and `ez`.

#### Parameters

**ex, ey, ez** [ndarray] Electric fields in x-, y-, and z-directions, as obtained from `emg3d.fields.Field`.

**sx, sy, sz** : Source fields in x-, y-, and z-directions, as obtained from `emg3d.fields.Field`.

**eta\_x, eta\_y, eta\_z, zeta** : VolumeModel parameters (multiplied by volumes) as obtained from `emg3d.models.VolumeModel()`.

**hx, hy, hz** [ndarray] Cell widths in x-, y-, and z-directions.

**nu** [int] Number of Gauss-Seidel iterations.

### gauss\_seidel\_x

`emg3d.core.gauss_seidel_x(ex, ey, ez, sx, sy, sz, eta_x, eta_y, eta_z, zeta, hx, hy, hz, nu)`

Gauss-Seidel method with line relaxation in x-direction.

This is the equivalent to `gauss_seidel()`, but with line relaxation in the x-direction. See `gauss_seidel()` for more details.

The resulting system  $Ax = b$  to solve consists of  $n$  unknowns (x-vector), and the corresponding matrix  $A$  is a banded matrix with the main diagonal and five upper and lower diagonals:

```
.-0
|X|\  0
0-.-0      left:  middle:  right:
\|X|\      (not used)
0-.-0      0-      .-      0
\|X|\      \|      |X      |\
0-.-0
0      \|X|
0-.
```

. 1\*1, - 4\*1, | 1\*4, X 4\*4, \ 4\*4 upper **or** lower

The matrix  $A$  is complex and symmetric ( $A = A^T$ ), and therefore only the main diagonal and the lower five off-diagonals are required.

- The right-hand-side  $b$  has length  $5*n_{Cx}-4$  ( $n_{Cx}$  even).
- The matrix  $A$  has length of  $b$  and  $1+2*5$  diagonals; we use for it an array of length  $6*\text{len}(b)$ .

The values are computed in rows of 5 lines, with the indicated middle and left matrices as indicated in the above scheme. These blocks are filled into the main matrix  $A$  and vector  $b$ , and subsequently solved with a non-standard Cholesky factorisation, `solve()`.

Tangential components at the boundaries are assumed to be 0 (PEC boundaries).

The result is stored in the provided electric fields `ex`, `ey`, and `ez`.

#### Parameters



**ex, ey, ez** [ndarray] Electric fields in x-, y-, and z-directions, as obtained from `emg3d.fields.Field`.

**sx, sy, sz** : Source fields in x-, y-, and z-directions, as obtained from `emg3d.fields.Field`.

**eta\_x, eta\_y, eta\_z, zeta** : VolumeModel parameters (multiplied by volumes) as obtained from `emg3d.models.VolumeModel()`.

**hx, hy, hz** [ndarray] Cell widths in x-, y-, and z-directions.

**nu** [int] Number of Gauss-Seidel iterations.

## gauss\_seidel\_y

`emg3d.core.gauss_seidel_y(ex, ey, ez, sx, sy, sz, eta_x, eta_y, eta_z, zeta, hx, hy, hz, nu)`

Gauss-Seidel method with line relaxation in y-direction.

This is the equivalent to `gauss_seidel()`, but with line relaxation in the y-direction. See `gauss_seidel()` for more details.

The resulting system  $Ax = b$  to solve consists of  $n$  unknowns ( $x$ -vector), and the corresponding matrix  $A$  is a banded matrix with the main diagonal and five upper and lower diagonals:

```
.-0
|X|\  0
0-.-0      left:  middle:  right:
\|X|\      (not used)
0-.-0      0-      .-      0
\|X|\      \|      |X      |\
0-.-0
0  \|X|
0-.
```

. 1\*1, - 4\*1, | 1\*4, X 4\*4, \ 4\*4 upper or lower

The matrix  $A$  is complex and symmetric ( $A = A^T$ ), and therefore only the main diagonal and the lower five off-diagonals are required.

- The right-hand-side  $b$  has length  $5*n_{Cy}-4$  ( $n_{Cy}$  even).
- The matrix  $A$  has length of  $b$  and  $1+2*5$  diagonals; we use for it an array of length  $6*\text{len}(b)$ .

The values are computed in rows of 5 lines, with the indicated middle and left matrices as indicated in the above scheme. These blocks are filled into the main matrix  $A$  and vector  $b$ , and subsequently solved with a non-standard Cholesky factorisation, `solve()`.

Note: The smoothing with linerelaxation in y-direction is carried out in reversed lexicographical order, in order to improve speed (memory access). All other smoothers (`gauss_seidel()`, `gauss_seidel_x()`, and `gauss_seidel_z()`) use lexicographical order.

Tangential components at the boundaries are assumed to be 0 (PEC boundaries).

The result is stored in the provided electric fields `ex`, `ey`, and `ez`.

### Parameters

**ex, ey, ez** [ndarray] Electric fields in x-, y-, and z-directions, as obtained from `emg3d.fields.Field`.

**sx, sy, sz** : Source fields in x-, y-, and z-directions, as obtained from `emg3d.fields.Field`.

**eta\_x, eta\_y, eta\_z, zeta** : VolumeModel parameters (multiplied by volumes) as obtained from `emg3d.models.VolumeModel()`.

**hx, hy, hz** [ndarray] Cell widths in x-, y-, and z-directions.

**nu** [int] Number of Gauss-Seidel iterations.

## gauss\_seidel\_z

`emg3d.core.gauss_seidel_z(ex, ey, ez, sx, sy, sz, eta_x, eta_y, eta_z, zeta, hx, hy, hz, nu)`

Gauss-Seidel method with line relaxation in z-direction.

This is the equivalent to `gauss_seidel()`, but with line relaxation in the z-direction. See `gauss_seidel()` for more details.

The resulting system  $Ax = b$  to solve consists of  $n$  unknowns ( $x$ -vector), and the corresponding matrix  $A$  is a banded matrix with the main diagonal and five upper and lower diagonals:

```
.-0
|X|\  0
0-.-0      left:  middle:  right:
\|X|\      (not used)
0-.-0      0-      .-      0
\|X|\      \|      |X      |\
0-.-0
0  \|X|
0-.
```

. 1\*1, - 4\*1, | 1\*4, X 4\*4, \| 4\*4 upper or lower

The matrix  $A$  is complex and symmetric ( $A = A^T$ ), and therefore only the main diagonal and the lower five off-diagonals are required.

- The right-hand-side  $b$  has length  $5*nCz-4$  ( $nCz$  even).
- The matrix  $A$  has length of  $b$  and  $1+2*5$  diagonals; we use for it an array of length  $6*\text{len}(b)$ .

The values are computed in rows of 5 lines, with the indicated middle and left matrices as indicated in the above scheme. These blocks are filled into the main matrix  $A$  and vector  $b$ , and subsequently solved with a non-standard Cholesky factorisation, `solve()`.

Tangential components at the boundaries are assumed to be 0 (PEC boundaries).

The result is stored in the provided electric fields  $ex$ ,  $ey$ , and  $ez$ .

### Parameters

**ex, ey, ez** [ndarray] Electric fields in x-, y-, and z-directions, as obtained from `emg3d.fields.Field`.

**sx, sy, sz** : Source fields in x-, y-, and z-directions, as obtained from `emg3d.fields.Field`.

**eta\_x, eta\_y, eta\_z, zeta** : VolumeModel parameters (multiplied by volumes) as obtained from `emg3d.models.VolumeModel()`.

**hx, hy, hz** [ndarray] Cell widths in x-, y-, and z-directions.

**nu** [int] Number of Gauss-Seidel iterations.

## restrict

`emg3d.core.restrict(crx, cry, crz, rx, ry, rz, wx, wy, wz, sc_dir)`

Restriction of residual from fine to coarse grid.

Corresponds to Equation 8 in [Mul06]. The equation for the x-direction, using the notation  $\{x, y, z\}$  instead of  $\{1, 2, 3\}$ , is given by

$$r_{x,K+1/2,L,M}^{2h} = \sum_{j_y=-1}^1 \sum_{j_z=-1}^1 w_{L,j_y}^y w_{M,j_z}^z \times \left( r_{x,k+1/2,l+j_y,m+j_z}^h + r_{x,k+3/2,l+j_y,m+j_z}^h \right).$$

The superscripts  $h, 2h$  indicate quantities defined on the coarse grid and on the fine grid, respectively. The indices  $\{K, L, M\}$  on the coarse grid correspond to  $\{k, l, m\} = 2\{K, L, M\}$  on the fine grid. The weights  $w$  are obtained from `restrict_weights()`.

The restrictions of  $rx$ ,  $ry$ , and  $rz$  are stored directly in `crx`, `cry`, and `crz`.

#### Parameters

- crx, cry, crz** [ndarray] Coarse grid  $\{x,y,z\}$ -directed residual (pre-allocated empty arrays).
- rx, ry, rz** [ndarray] Fine grid  $\{x,y,z\}$ -directed residual.
- wx, wy, wz: tuple** Tuples containing the weights ( $w_l, w_0, w_r$ ) as returned from `restrict_weights()` for the x-, y-, and z-directions.
- sc\_dir** [int] Direction of semicoarsening; 0 for no semicoarsening.

### restrict\_weights

`emg3d.core.restrict_weights(vectorN, vectorCC, h, cvectorN, cvectorCC, ch)`

Restriction weights for the coarse-grid correction operator.

Corresponds to Equation 9 in [Mul06]. A generalized version of that equation is given by

$$\begin{aligned} w_{Q,-1}^v &= \left( v_{q-1/2}^h - v_{Q-1/2}^{2h} \right) / d_{q-1}^v, \\ w_{Q,0}^v &= 1, \\ w_{Q,1}^v &= \left( v_{Q+1/2}^{2h} - v_{q+1/2}^h \right) / d_{q+1}^v, \end{aligned}$$

where  $d$  are the dual grid cell widths,  $v$  is one of  $\{x, y, z\}$ , and  $Q, q$  the corresponding entries of  $\{K, L, M\}, \{k, l, m\}$ . The superscripts  $h, 2h$  indicate quantities defined on the coarse grid and on the fine grid, respectively. The indices  $\{K, L, M\}$  on the coarse grid correspond to  $\{k, l, m\} = 2\{K, L, M\}$  on the fine grid.

For the dual volume cell widths at the boundaries the scheme of [MoSu94] is applied, where  $d_0^x = h_{1/2}^x/2$  at  $k = 0$ ,  $d_{N_x}^x = h_{N_x-1/2}^x$  at  $k = N_x$ , and so on.

The following parameters must all be in the same direction, hence, all must be either for the x, the y, or the z direction. The returned weights are for this direction.

#### Parameters

- vectorN, cvectorN** [ndarray] Cell edges of the fine (vectorN) and coarse (cvectorN) grids.
- vectorCC, cvectorCC** [ndarray] Cell centers of the fine (vectorCC) and coarse (cvectorCC) grids.
- h, ch** [ndarray] Cell widths of the fine (h) and coarse (ch) grids.

#### Returns

- wl, w0, wr** [ndarray] Left, central, and right weights in the direction provided in the input.

## solve

`emg3d.core.solve (amat, bvec)`

Solve  $Ax = b$  using a non-standard Cholesky factorisation.

Solve the system  $Ax = b$  using a non-standard Cholesky factorisation without pivoting for a symmetric, complex matrix  $A$  tailored to the problem of the multigrid solver. The matrix  $A$  (`amat`) is an array of length  $6*n$ , containing the main diagonal and the first five lower off-diagonals (ordered so that the first element of the main diagonal is followed by the first elements of the off diagonals, then the second elements and so on). The vector `bvec` has length  $b$ .

The solution is placed in `b` (`bvec`), and  $A$  (`amat`) is replaced by its decomposition.

### 1. Non-standard Cholesky factorisation.

From [Mul07]: We use a non-standard Cholesky factorisation. The standard factorisation factors a hermitian matrix  $A$  into  $LL^H$ , where  $L$  is a lower triangular matrix and  $L^H$  its complex conjugate transpose. In our case, the discretisation is based on the Finite Integration Technique ([Weil77]) and provides a matrix  $A$  that is complex-valued and symmetric:  $A = A^T$ , where the superscript  $T$  denotes the transpose. The line relaxation scheme takes a matrix  $B$  that is a subset of  $A$  along the line.  $B$  is a complex symmetric band matrix with eleven diagonals. The non-standard Cholesky factorisation factors the matrix  $B$  into  $LL^T$ . Because of the symmetry, only the main diagonal and five lower diagonal elements of  $B$  need to be computed. The Cholesky factorisation replaces this matrix by  $L$ , containing six diagonals, after which the line relaxation can be carried out by simple back-substitution.

$A = LDL^T$  factorisation without pivoting:

$$D(j) = A(j, j) - \sum_{k=1}^{j-1} L(j, k)^2 D(k), \quad j = 1, \dots, n;$$
$$L(i, j) = \frac{1}{D(j)} \left[ A(i, j) - \sum_{k=1}^{j-1} L(i, k) L(j, k) D(k) \right], \quad i = j + 1, \dots, n.$$

$A$  and  $L$  are in this case arrays, where  $A(i, j) \rightarrow A(i + 5j)$ .

### 2. Solve $Ax = b$ .

Solve  $Ax = b$ , given  $L$  which is the result from the factorisation in the first step (and stored in  $A$ ), hence, solve  $Lx = b$ , where  $x$  is stored in `b`:

$$b(j) = b(j) - \sum_{k=1}^{j-1} L(j, k)x(k), \quad j = 2, \dots, n.$$

The result is equivalent with simply using `numpy.linalg.solve()`, but faster for the particular use-case of this code.

Note that in this custom solver there is no pivoting, and the diagonals of the matrix cannot be zero.

### Parameters

**amat** [ndarray] Banded matrix  $A$  provided as a vector of length  $6*n$ , containing main diagonal plus first five lower diagonals.

**bvec** [ndarray] Right-hand-side vector  $b$  of length  $n$ .

## 5.12 Meshes, Models, and Fields

### 5.12.1 emg3d.meshes Module

Everything related to meshes appropriate for the multigrid solver.

## Functions

<code>construct_mesh(frequency, properties, center)</code>	Return a TensorMesh for given parameters.
<code>get_origin_widths(frequency, properties, center)</code>	Return origin and cell widths for given parameters.
<code>skin_depth(frequency, conductivity[, mu, ...])</code>	Return skin depth for provided frequency and conductivity.
<code>wavelength(skin_depth[, precision])</code>	Return the wavelength.
<code>good_mg_cell_nr([max_nr, max_prime, min_div])</code>	Returns 'good' cell numbers for the multigrid method.
<code>min_cell_width(skin_depth[, pps, limits, ...])</code>	Return the minimum cell width.
<code>get_hx_h0(freq, res, domain[, fixed, ...])</code>	Return cell widths and origin for given parameters.
<code>get_cell_numbers(max_nr[, max_prime, min_div])</code>	
<code>get_stretched_h(min_width, domain, nx[, x0, ...])</code>	Return cell widths for a stretched grid within the domain.
<code>get_domain([x0, freq, res, limits, ...])</code>	Get domain extent and minimum cell width as a function of skin depth.
<code>get_hx(alpha, domain, nx, x0[, resp_domain])</code>	Return cell widths for given input.

## construct\_mesh

`emg3d.meshes.construct_mesh(frequency, properties, center, domain=None, vector=None, sea-surface=None, **kwargs)`

Return a TensorMesh for given parameters.

The constructed mesh is frequency- and conductivity-dependent, where `properties` are turned into conductivities through the provided mapping, which is 'Resistivity' by default. Some details are explained in other functions:

- The minimum cell width  $\Delta_{\min}$  is a function of `frequency`, `properties[0]`, `min_width_pps`, and `min_width_limits`, see Equation (5.34).
- The skin depth  $\delta$  is a function of `frequency` and `properties`, see Equation (5.32).
- The wavelength  $\lambda$  is a function of `frequency` and `properties`, see Equation (5.33).

The relation of the survey domain, computational domain, and buffer zone is shown in Figure 5.12 for a x-z-section; the y-direction behaves the same as the x-direction (the figures are only visible in the web version on <https://emg3d.rtfd.io>).

By default, the buffer zone around the survey domain is one wavelength. This means that the signal has to travel two wavelengths to get from the end of the survey domain to the end of the computational domain and back. This approach is quite conservative and on the safe side. You can reduce the buffer thickness if you know what you are doing. There are three parameters which influence the thickness of the buffer for a given frequency: `properties`, which is used to calculate the skin depth and the wavelength, `lambda_factor` (default is 1) which sets how many times the wavelength is the thickness of the buffer (relative factor), and `max_buffer`, which is an absolute maximum for the buffer thickness. A graphical illustration is given in Figure 5.13.

### Parameters

**frequency** [float] Frequency (Hz) to calculate skin depth; both the minimum cell width and the extent of the buffer zone, and therefore of the computational domain, are a function of skin depth.

**properties** [float or list] Properties to calculate the skin depths. The properties can be either resistivities, conductivities, or the logarithm (natural or base 10) thereof. By default it assumes resistivities, but it can be changed with the parameter `mapping`.

Four formats are recognized:

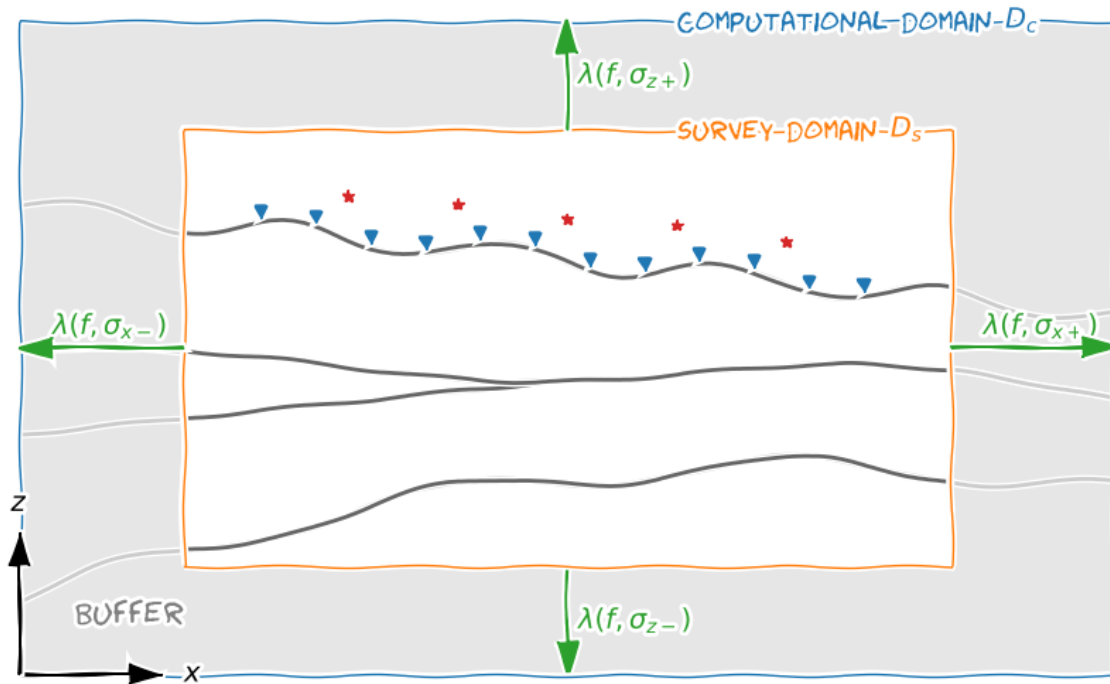


Figure 5.12:: Relation between survey domain ( $D_s$ , domain), computational domain ( $D_c$ ), and buffer zone. The survey domain should contain all sources and receivers as well as any important feature that should be represented in the data. The buffer zone is calculated as a function of wavelength with the provided property in the given direction.

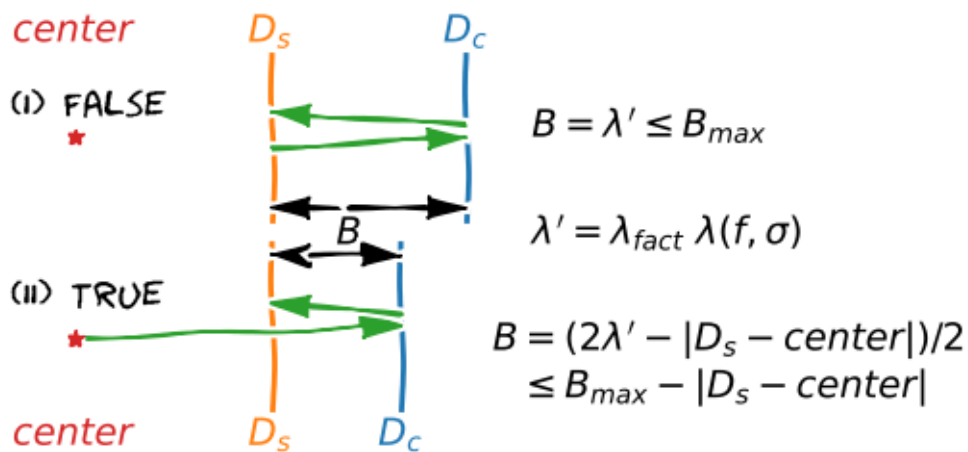


Figure 5.13:: The thickness of the buffer zone ( $B$ ) for (I) `lambda_from_center=False` (default) and for (II) `lambda_from_center=True`. The `lambda_factor` ( $\lambda_{fact}$ ) is a simple scaling factor for the wavelength  $\lambda$ . The `max_buffer` is an absolute limitation.

- 1: Same property for everything;
- 2: [min\_width, buffer (+/-)] for all directions;
- 4: [min\_width, xy-buffer (+/-), z-, z+];
- 7: [min\_width, x-, x+, y-, y+, z-, z+].

The property `min_width` is usually the property at the center, hence at the source location. The other properties are used to define the extent of the buffer zone around the survey domain in the respective directions.

**center** [tuple] Tuple (or list, ndarray) of three floats for (x, y, z). The mesh is centered around this point, which means that here is the smallest cell. Usually this is the source location.

**domain** [tuple of lists, list, or None, optional] Contains the survey-domain limits. This domain should include all source and receiver positions as well as any important feature of the model. Format: ([xmin, xmax], [ymin, ymax], [zmin, zmax]).

It can be None, or individual lists can be None (e.g., (None, None, [zmin, zmax])), in which case you have to provide a `vector`, which is then assumed to span exactly the domain. If only one list is provided it is applied to all dimensions.

**vector** [tuple of three ndarrays, ndarray, or None, optional] Contains vectors of mesh-edges that should be used. If provided, the vector MUST at least include all of the survey domain. If `domain` is not provided, it is defined as the minimum/maximum of the provided vector. Format: (xvector, yvector, zvector).

It can be None, or individual ndarrays can be None (e.g., (xvector, yvector, None)), in which case you have to provide a `domain`. If only one ndarray is provided it is applied to all dimensions.

**seasurface** [float, optional] Air-sea interface. This has only to be set in the marine case, when the mesh in z-direction is sought for (and the interface is not contained in `vector`). If set, it will ensure that at the sea surface is an actual boundary. It has to be bigger then the lower limit of the survey domain. Default is None.

**stretching** [list or tuple of lists, optional] Maximum stretching factors in the form of [max Ds, max Dc]: the first value is the maximum stretching for the survey domain (default is 1.0), the second value is the maximum stretching for the buffer zone (default is 1.5). If a list is provided the same is used for all three dimension. Alternatively a tuple of three lists can be provided, (x, y, z). Note that the first value has no influence on dimensions where a `vector` is provided.

**min\_width\_limits** [float, list or None, optional] Passed through `min_cell_width()` as `limits`. A tuple of three can be provided for direction dependent values. Note that this value has no influence on dimensions where a `vector` is provided.

Default is None.

**min\_width\_pps** [float or int, optional] Passed through `min_cell_width()` as `pps`. A tuple of three can be provided for direction dependent values. Note that this value has no influence on dimensions where a `vector` is provided.

Default is 3.

**lambda\_factor** [float, optional] The buffer is taken as one wavelength from the survey domain. This can be regarded as quite conservative (but safe). The parameter `lambda_factor` can be used to reduce (or increase) this factor. Default is 1.0.

**max\_buffer** [float, optional] Maximum thickness of the buffer zone around survey domain. If `lambda_from_center=True`, this is the maximum distance from the center to the end of the computational domain. Default is 100,000 (100 km).

**lambda\_from\_center** [bool, optional] Flag how to compute the extent of the computational mesh as a function of wavelength:

- False (default): The distance from the edge of the survey domain to the edge of the computational domain is one wavelength.
- True: The distance from the center to the edge of the computational domain and back to the end of the survey domain is two wavelengths.

**mapping** [str or map, optional] Defines what type the input `property_{x;y;z}`-values correspond to. By default, they represent resistivities (Ohm.m). The implemented mappings are:

- ‘Conductivity’;  $\sigma$  (S/m),
- ‘LgConductivity’;  $\log_{10}(\sigma)$ ,
- ‘LnConductivity’;  $\log_e(\sigma)$ ,
- ‘Resistivity’;  $\rho$  (Ohm.m); Default,
- ‘LgResistivity’;  $\log_{10}(\rho)$ ,
- ‘LnResistivity’;  $\log_e(\rho)$ .

**cell\_numbers** [list, optional] List of possible numbers of cells. See `good_mg_cell_nr()`. Default is `good_mg_cell_nr(1024, 5, 3)`, which corresponds to numbers 16, 24, 32, 40, 48, 64, 80, 96, 128, 160, 192, 256, 320, 384, 512, 640, 768, 1024.

**verb** [int, optional] Verbosity, -1 (error); 0 (warning), 1 (info), 2 (verbose). Default = 0 (Warnings only).

#### Returns

**origin** [float] Origin of the mesh.

**widths** [ndarray] Cell widths of mesh.

### get\_origin\_widths

`emg3d.meshes.get_origin_widths(frequency, properties, center, domain=None, vector=None, searface=None, **kwargs)`

Return origin and cell widths for given parameters.

This function works in one dimension only, and is used by `construct_mesh()` once in each direction. It is recommended to use directly function `construct_mesh()`, which returns a `TensorMesh`.

All the parameters are described in `construct_mesh()`. Described here are only the differences.

#### Parameters

**All** [description] All parameters are described in `construct_mesh()`. The only difference is that here only variables for one direction are accepted.

**raise\_error** [bool, optional] If True, an error is raised if no suitable grid is found. Otherwise it just prints a message and returns None's. Default is True.

#### Returns

**origin** [float] Origin of the mesh.

**widths** [ndarray] Cell widths of mesh.

### skin\_depth

`emg3d.meshes.skin_depth(frequency, conductivity, mu=1.25663706212e-06, precision=0)`

Return skin depth for provided frequency and conductivity.



The skin depth  $\delta$  (m) is given by

$$\delta = \sqrt{\frac{2}{\omega \sigma \mu}}, \quad (5.32)$$

where  $\omega = 2\pi f$  is angular frequency of frequency  $f$  (Hz),  $\sigma$  is conductivity (S/m), and  $\mu$  is magnetic permeability (H/m).

#### Parameters

**frequency** [float] Frequency (Hz).

**conductivity** [float] Conductivity (S/m).

**mu** [float, optional] Magnetic permeability (H/m); default is  $\mu_0$ .

**precision** [int, optional] Precision of the return skin depth. Default is 0, hence meters.

#### Returns

**skindepth** [float] Skin depth (m).

### wavelength

`emg3d.meshes.wavelength(skin_depth, precision=0)`

Return the wavelength.

The wavelength  $\lambda$  (m) is given by

$$\lambda = 2\pi\delta. \quad (5.33)$$

The skin depth  $\delta$  is a function of frequency and conductivity and is given by `skin_depth()`, Equation (5.32).

#### Parameters

**skin\_depth** [float or ndarray.] Skin depth (m).

**precision** [int, optional] Precision of the returned wave length. Default is 0, hence meters.

#### Returns

**wavelength** [float or ndarray] Wavelength (m).

### good\_mg\_cell\_nr

`emg3d.meshes.good_mg_cell_nr(max_nr=1024, max_prime=5, min_div=3)`

Returns ‘good’ cell numbers for the multigrid method.

‘Good’ cell numbers are numbers which can be divided by 2 as many times as possible. At the end there will be a low prime number.

The function adds all numbers  $p2^n \leq M$  for  $p = 2, 3, \dots, p_{\max}$  and  $n = n_{\min}, n_{\min} + 1, \dots, \infty$ ;  $M, p_{\max}, n_{\min}$  correspond to `max_nr`, `max_prime`, and `min_div`, respectively.

#### Parameters

**max\_nr** [int, optional] Maximum number of cells. Default is 1024.

**max\_prime** [int, optional] Highest permitted prime number  $p$  for  $p \cdot 2^n$ . {2, 3, 5, 7} are good upper limits in order to avoid too big lowest grids in the multigrid method. Default is 5.

**min\_div** [int, optional] Minimum times the number can be divided by two. Default is 3.

#### Returns

**numbers** [array] Array containing all possible cell numbers from lowest to highest.

## min\_cell\_width

`emg3d.meshes.min_cell_width(skin_depth, pps=3, limits=None, precision=0)`

Return the minimum cell width.

The minimum cell width is defined by the desired points per skin depth,

$$\Delta_{\min} = \text{limits}[0] \leq \frac{\delta}{\text{pps}} \leq \text{limits}[1] . \quad (5.34)$$

The skin depth  $\delta$  is a function of frequency and conductivity and is given by `skin_depth()`, Equation (5.32).

### Parameters

**skin\_depth** [float] Skin depth (m).

**pps** [int] Points per skin depth.

**limits** [None, float, or list of two floats] Limits on minimum width:

- None: No limits.
- float: Returns limits as dmin.
- [min, max]: dmin is limited to this range.

**precision** [int, optional] Precision of the cell width. Provided limits are not rounded. Default is 0, hence meters.

### Returns

**dmin** [float] Minimum cell width (m).

## get\_hx\_h0

`emg3d.meshes.get_hx_h0(freq, res, domain, fixed=0.0, possible_nx=None, min_width=None, pps=3, alpha=None, max_domain=100000.0, raise_error=True, verb=1, return_info=False)`

Return cell widths and origin for given parameters.

Returns cell widths for the provided frequency, resistivity, domain extent, and other parameters using a flexible amount of cells. See input parameters for more details. A maximum of three hard/fixed boundaries can be provided (one of which is the grid center).

The minimum cell width is computed through  $\delta/\text{pps}$ , where the skin depth is given by  $\delta = 503.3\sqrt{\rho/f}$ , and the parameter *pps* stands for ‘points-per-skindepth’. The minimum cell width can be restricted with the parameter *min\_width*.

The actual computation domain adds a buffer zone around the (survey) domain. The thickness of the buffer is six times the skin depth. The field is basically zero after two wavelengths. A wavelength is  $2\pi\delta$ , hence roughly 6 times the skin depth. Taking a factor 6 gives therefore almost two wavelengths, as the field travels to the boundary and back. The actual buffer thickness can be steered with the *res* parameter.

One has to take into account that the air is very resistive, which has to be considered not just in the vertical direction, but also in the horizontal directions, as the airwave will bounce back from the sides otherwise. In the marine case this issue reduces with increasing water depth.

### Parameters

**freq** [float] Frequency (Hz) to compute the skin depth. The skin depth is a concept defined in the frequency domain. If a negative frequency is provided, it is assumed that the computation is carried out in the Laplace domain. To compute the skin depth, the value of *freq* is then multiplied by  $-2\pi$ , to simulate the closest frequency-equivalent.

**res** [float or list] Resistivity (Ohm m) to compute the skin depth. The skin depth is used to compute the minimum cell width and the boundary thicknesses. Up to three resistivities can be provided:

- float: Same resistivity for everything;
- [min\_width, boundaries];
- [min\_width, left boundary, right boundary].

**domain** [list] Contains the survey-domain limits [min, max]. The actual computation domain consists of this domain plus a buffer zone around it, which depends on frequency and resistivity.

**fixed** [list, optional] Fixed boundaries, one, two, or maximum three values. The grid is centered around the first value. Hence it is the center location with the smallest cell. Two more fixed boundaries can be added, at most one on each side of the first one. Default is 0.

**possible\_nx** [list, optional] List of possible numbers of cells. See `good_mg_cell_nr()`. Default is `good_mg_cell_nr(1024, 5, 3)`, which corresponds to numbers 16, 24, 32, 40, 48, 64, 80, 96, 128, 160, 192, 256, 320, 384, 512, 640, 768, 1024.

**min\_width** [float, list or None, optional] Minimum cell width restriction:

- None : No restriction;
- float : Fixed to this value, ignoring skin depth and *pps*.
- list [min, max] : Lower and upper bounds.

Default is None.

**pps** [int, optional] Points per skindepth; minimum cell width is computed via  $dmin = \text{skindepth}/pps$ . Default = 3.

**alpha** [list, optional] Maximum alpha and step size to find a good alpha. The first value is the maximum alpha of the survey domain, the second value is the maximum alpha for the buffer zone, and the third value is the step size. Default = [1, 1.5, .01], hence no stretching within the survey domain and a maximum stretching of 1.5 in the buffer zone; step size is 0.01.

**max\_domain** [float, optional] Maximum computation domain from fixed[0] (usually source position). Default is 100,000.

**raise\_error** [bool, optional] If True, an error is raised if no suitable grid is found. Otherwise it just prints a message and returns None's. Default is True.

**verb** [int, optional] Verbosity, 0 or 1. Default = 1.

**return\_info** [bool] If True, a dictionary is returned with some grid info (min and max cell width and alpha).

## Returns

**hx** [ndarray] Cell widths of mesh.

**x0** [float] Origin of the mesh.

**info** [dict] Dictionary with mesh info; only if `return_info=True`.

Keys:

- *dmin*: Minimum cell width;
- *dmax*: Maximum cell width;
- *amin*: Minimum alpha;
- *amax*: Maximum alpha.

See also:

`get_stretched_h` Get  $h_x$  for a fixed number  $nx$  and within a fixed domain.

## get\_cell\_numbers

`emg3d.meshes.get_cell_numbers` (*max\_nr*, *max\_prime*=5, *min\_div*=3)

## get\_stretched\_h

`emg3d.meshes.get_stretched_h` (*min\_width*, *domain*, *nx*, *x0*=0, *x1*=None, *resp\_domain*=False)

Return cell widths for a stretched grid within the domain.

Returns  $nx$  cell widths within *domain*, where the minimum cell width is *min\_width*. The cells are not stretched within  $x_0$  and  $x_1$ , and outside uses a power-law stretching. The actual stretching factor and the number of cells left and right of  $x_0$  and  $x_1$  are found in a minimization process.

The domain is not completely respected. The starting point of the domain is, but the endpoint of the domain might slightly shift (this is more likely the case for small  $nx$ , for big  $nx$  the shift should be small). The new endpoint can be obtained with `domain[0]+np.sum(hx)`. If you want the domain to be respected absolutely, set `resp_domain=True`. However, be aware that this will introduce one stretch-factor which is different from the other stretch factors, to accommodate the restriction. This one-off factor is between the left- and right-side of  $x_0$ , or, if  $x_1$  is provided, just after  $x_1$ .

### Parameters

**min\_width** [float] Minimum cell width. If  $x_1$  is provided, the actual minimum cell width might be smaller than *min\_width*.

**domain** [list] [start, end] of model domain.

**nx** [int] Number of cells.

**x0** [float] Center of the grid.  $x_0$  is restricted to *domain*. Default is 0.

**x1** [float] If provided, then no stretching is applied between  $x_0$  and  $x_1$ . The non-stretched part starts at  $x_0$  and stops at the first possible location at or after  $x_1$ .  $x_1$  is restricted to *domain*. This will *min\_width* so that an integer number of cells fit within  $x_0$  and  $x_1$ .

**resp\_domain** [bool] If False (default), then the domain-end might shift slightly to assure that the same stretching factor is applied throughout. If set to True, however, the domain is respected absolutely. This will introduce one stretch-factor which is different from the other stretch factors, to accommodate the restriction. This one-off factor is between the left- and right-side of  $x_0$ , or, if  $x_1$  is provided, just after  $x_1$ .

### Returns

**hx** [ndarray] Cell widths of mesh.

See also:

`get_hx_x0` Get  $h_x$  and  $x_0$  for a flexible number of  $nx$  with given bounds.

## get\_domain

`emg3d.meshes.get_domain` (*x0*=0, *freq*=1, *res*=0.3, *limits*=None, *min\_width*=None, *fact\_min*=0.2, *fact\_neg*=5, *fact\_pos*=None)

Get domain extent and minimum cell width as a function of skin depth.

Returns the extent of the computation domain and the minimum cell width as a multiple of the skin depth, with possible user restrictions on minimum computation domain and range of possible minimum cell widths.

$$\delta = 503.3 \sqrt{\frac{\rho}{f}},$$

$$x_{\text{start}} = x_0 - k_{\text{neg}} \delta,$$

$$x_{\text{end}} = x_0 + k_{\text{pos}} \delta,$$

$$h_{\text{min}} = k_{\text{min}} \delta.$$

### Parameters

**x0** [float] Center of the computation domain. Normally the source location. Default is 0.

**freq** [float] Frequency (Hz) to compute the skin depth. The skin depth is a concept defined in the frequency domain. If a negative frequency is provided, it is assumed that the computation is carried out in the Laplace domain. To compute the skin depth, the value of *freq* is then multiplied by  $-2\pi$ , to simulate the closest frequency-equivalent.

Default is 1 Hz.

**res** [float, optional] Resistivity (Ohm m) to compute skin depth. Default is 0.3 Ohm m (sea water).

**limits** [None or list] [start, end] of model domain. This extent represents the minimum extent of the domain. The domain is therefore only adjusted if it has to reach outside of [start, end]. Default is None.

**min\_width** [None, float, or list of two floats] Minimum cell width is computed as a function of skin depth:  $\text{fact\_min} \times \text{sd}$ . If *min\_width* is a float, this is used. If a list of two values [min, max] are provided, they are used to restrain *min\_width*. Default is None.

**fact\_min, fact\_neg, fact\_pos** [floats] The skin depth is multiplied with these factors to estimate:

- Minimum cell width (*fact\_min*, default 0.2)
- Domain-start (*fact\_neg*, default 5), and
- Domain-end (*fact\_pos*, defaults to *fact\_neg*).

### Returns

**h\_min** [float] Minimum cell width.

**domain** [list] Start- and end-points of computation domain.

## get\_hx

`emg3d.meshes.get_hx(alpha, domain, nx, x0, resp_domain=True)`

Return cell widths for given input.

Find the number of cells left (*n<sub>l</sub>*) and right (*n<sub>r</sub>*) of the center *x0* for the provided alpha. For this, we solve

$$\frac{x_{\text{max}} - x_0}{x_0 - x_{\text{min}}} = \frac{a^{n_r} - 1}{a^{n_l} - 1}$$

where  $a = 1 + \alpha$ .

### Parameters

**alpha** [float] Stretching factor *a* is given by  $a=1+\text{alpha}$ .

**domain** [list] [*x\_min*, *x\_max*] of model domain.

**nx** [int] Number of cells.

**x0** [float] Center of the grid. *x0* is restricted to *domain*.

**resp\_domain** [bool] If False, then the domain-end might shift slightly to assure that the same stretching factor is applied throughout. If set to True (default), however, the domain is respected absolutely. This will introduce one stretch-factor which is different from the other stretch factors, to accommodate the restriction. This one-off factor is between the left- and right-side of  $x_0$ , or, if  $x_l$  is provided, just after  $x_l$ .

#### Returns

**hx** [ndarray] Cell widths of mesh. All points are given by `np.r_[xmin, xmin+np.cumsum(hx) ]`

## Classes

---

<code>TensorMesh([h, origin])</code>	A slightly modified <code>discretize.TensorMesh</code> .
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---

### TensorMesh

**class** `emg3d.meshes.TensorMesh` (*h=None, origin=None, \*\*kwargs*)

Bases: `discretize.tensor_mesh.TensorMesh`, `emg3d.meshes._TensorMesh`

A slightly modified `discretize.TensorMesh`.

Adds a few attributes (`__eq__`, `copy`, `to_dict`, and `from_dict`) to the `discretize.TensorMesh`.

Falls back to a minimal `TensorMesh` if `discretize` is not installed. Nothing fancy is possible with the minimal `TensorMesh`, particularly NO plotting nor nice repr-functions.

#### Parameters

**h** [list of three ndarrays]

Cell widths in [x, y, z] directions.

**x0** [tuple of length 3] Origin (x, y, z).

**\*\*Required Properties:\*\***

\* **axis\_u (Vector3):** Vector orientation of u-direction. For more details see the docs for the `rotation_m`

\* **axis\_v (Vector3):** Vector orientation of v-direction. For more details see the docs for the `rotation_m`

\* **axis\_w (Vector3):** Vector orientation of w-direction. For more details see the docs for the `rotation_m`

\* **h (a tuple of Array):** h is a list containing the cell widths of the tensor mesh in each dimension., a tuple

\* **origin (Array):** origin of the mesh (dim, ), a list or numpy array of <class 'float'>, <class 'int'> with sha

\* **reference\_system (String):** The type of coordinate reference frame. Can take on the values cartesian, c

#### Methods Summary

---

<code>copy()</code>	Return a copy of the <code>TensorMesh</code> .
<code>from_dict(inp)</code>	Convert dictionary into <code>TensorMesh</code> instance.

---

Continued on next page

Table 5.9 – continued from previous page

<code>to_dict([copy])</code>	Store the necessary information of the <code>TensorMesh</code> in a dict.
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## Methods Documentation

`copy()`

Return a copy of the `TensorMesh`.

`classmethod from_dict(inp)`

Convert dictionary into `TensorMesh` instance.

### Parameters

**inp** [dict] Dictionary as obtained from `TensorMesh.to_dict()`. The dictionary needs the keys `hx`, `hy`, `hz`, and `x0`.

### Returns

**obj** [`TensorMesh` instance]

`to_dict(copy=False)`

Store the necessary information of the `TensorMesh` in a dict.

## 5.12.2 emg3d.models Module

Everything to create model-properties for the multigrid solver.

## Classes

<code>Model(grid[, property_x, property_y, ...])</code>	Create a model instance.
<code>VolumeModel(grid, model, sfield)</code>	Return a volume-averaged version of provided model.

## Model

**class** `emg3d.models.Model` (*grid*, *property\_x=1.0*, *property\_y=None*, *property\_z=None*, *mu\_r=None*, *epsilon\_r=None*, *mapping='Resistivity'*, *\*\*kwargs*)

Bases: `object`

Create a model instance.

Class to provide model parameters (x-, y-, and z-directed properties [resistivity or conductivity; linear or on  $\log_{10}/\log_e$ -scale], electric permittivity and magnetic permeability) to the solver. Relative magnetic permeability  $\mu_r$  is by default set to one and electric permittivity  $\epsilon_r$  is by default set to zero, but they can also be provided (isotropically). Keep in mind that the multigrid method as implemented in *emg3d* only works for the diffusive approximation. As soon as the displacement-part in the Maxwell's equations becomes too dominant it will fail (high frequencies or very high electric permittivity).

### Parameters

**grid** [`TensorMesh`] Grid on which to apply model.

**property\_{x;y;z}** [float or ndarray; default to 1.] Material property in x-, y-, and z-directions. If ndarray, they must have the shape of `grid.vnC` (F-ordered) or `grid.nC`.

By default, property refers to electrical resistivity. However, this can be changed with an appropriate map. For more info, see the description of the parameter *mapping*. The internals of *emg3d* work, irrelevant of the map, with electrical conductivities.

Resistivities and conductivities have to be bigger than zero and smaller than infinity (if provided on a linear scale; not on logarithmic scales).

**mu\_r** [None, float, or ndarray] Relative magnetic permeability (isotropic). If ndarray it must have the shape of grid.vnC (F-ordered) or grid.nC. Default is None, which corresponds to 1., but avoids the computation of zeta. Magnetic permeability has to be bigger than zero and smaller than infinity.

**epsilon\_r** [None, float, or ndarray] Relative electric permittivity (isotropic). If ndarray it must have the shape of grid.vnC (F-ordered) or grid.nC. The displacement part is completely neglected (diffusive approximation) if set to None, which is the default. Electric permittivity has to be bigger than zero and smaller than infinity.

**mapping** [str] Defines what type the input *property\_{x;y;z}*-values correspond to. By default, they represent resistivities (Ohm.m). The implemented types are:

- ‘Conductivity’;  $\sigma$  (S/m),
- ‘LgConductivity’;  $\log_{10}(\sigma)$ ,
- ‘LnConductivity’;  $\log_e(\sigma)$ ,
- ‘Resistivity’;  $\rho$  (Ohm.m); Default,
- ‘LgResistivity’;  $\log_{10}(\rho)$ ,
- ‘LnResistivity’;  $\log_e(\rho)$ .

### Attributes Summary

<i>epsilon_r</i>	Electric permittivity.
<i>mu_r</i>	Magnetic permeability.
<i>property_x</i>	Property in x-direction.
<i>property_y</i>	Property in y-direction.
<i>property_z</i>	Property in z-direction.

### Methods Summary

<i>copy()</i>	Return a copy of the Model.
<i>from_dict</i> (inp)	Convert the dictionary into a Model instance.
<i>interpolate2grid</i> (grid, new_grid, ...)	Interpolate <i>Model</i> located on <i>grid</i> to <i>new_grid</i> .
<i>to_dict</i> ([copy])	Store the necessary information of the Model in a dict.

### Attributes Documentation

**epsilon\_r**  
Electric permittivity.

**mu\_r**  
Magnetic permeability.

**property\_x**  
Property in x-direction.

**property\_y**  
Property in y-direction.

**property\_z**  
Property in z-direction.



## Methods Documentation

**copy()**

Return a copy of the Model.

**classmethod from\_dict(inp)**

Convert the dictionary into a Model instance.

### Parameters

**inp** [dict] Dictionary as obtained from `Model.to_dict()`. The dictionary needs the keys `property_x`, `property_y`, `property_z`, `mu_r`, `epsilon_r`, `vnC`, and `mapping`.

### Returns

**obj** [`Model` instance]

**interpolate2grid(grid, new\_grid, \*\*grid2grid\_opts)**

Interpolate `Model` located on `grid` to `new_grid`.

### Parameters

**grid, new\_grid** [`TensorMesh`] Input and output model grids; `emg3d.meshes.TensorMesh` instances.

**grid2grid\_opts** [dict] Passed through to `maps.grid2grid()`. Defaults are `method='volume'`, `log=True`, and `extrapolate=True`.

### Returns

**NewModel** [`Model`] New `Model` instance on `new_grid`.

**to\_dict(copy=False)**

Store the necessary information of the Model in a dict.

## VolumeModel

**class** `emg3d.models.VolumeModel(grid, model, sfield)`

Bases: `object`

Return a volume-averaged version of provided model.

Takes a `Model` instance and returns the volume averaged values. This is used by the solver internally.

$$\eta_{\{x,y,z\}} = -V i \omega \mu_0 \left( \rho_{\{x,y,z\}}^{-1} + i \omega \varepsilon \right)$$

$$\zeta = V \mu_r^{-1}$$

### Parameters

**grid** [`TensorMesh`] Grid on which to apply model.

**model** [`Model`] Model to transform to volume-averaged values.

**sfield** [`SourceField`] A `VolumeModel` is frequency-dependent. The frequency-information is taken from the provided source field.

## Attributes Summary

<code>eta_x</code>	eta in x-direction.
<code>eta_y</code>	eta in y-direction.
<code>eta_z</code>	eta in z-direction.
<code>zeta</code>	zeta.

## Methods Summary

<code>calculate_eta(name, grid, model, field)</code>	eta: volume multiplied with conductivity.
<code>calculate_zeta(name, grid, model)</code>	zeta: volume divided by mu_r.

## Attributes Documentation

**eta\_x**  
eta in x-direction.

**eta\_y**  
eta in y-direction.

**eta\_z**  
eta in z-direction.

**zeta**  
zeta.

## Methods Documentation

**static** `calculate_eta(name, grid, model, field)`  
eta: volume multiplied with conductivity.

**static** `calculate_zeta(name, grid, model)`  
zeta: volume divided by mu\_r.

## 5.12.3 emg3d.maps Module

Interpolation routines mapping grids to grids, grids to fields, and fields to grids.

### Functions

<code>grid2grid(grid, values, new_grid[, method, ...])</code>	Interpolate <i>values</i> located on <i>grid</i> to <i>new_grid</i> .
<code>interp3d(points, values, new_points, method, ...)</code>	Interpolate values in 3D either linearly or with a cubic spline.
<code>edges2cellaverages(ex, ey, ez, vol, out_x, ...)</code>	Interpolate fields defined on edges to volume-averaged cell values.

### grid2grid

`emg3d.maps.grid2grid(grid, values, new_grid, method='linear', extrapolate=True, log=False)`  
Interpolate *values* located on *grid* to *new\_grid*.

**Note 1:** The default method is 'linear', because it works with fields and model parameters. However, recommended are 'volume' for model parameters and 'cubic' for fields.

**Note 2:** For model parameters with *method*='volume' the result is quite different if you provide resistivity, conductivity, or the logarithm of any of the two. The recommended way is to provide the logarithm of resistivity or conductivity, in which case the output of one is indeed the inverse of the output of the other.

#### Parameters

**grid, new\_grid** [TensorMesh] Input and output model grids; `TensorMesh` instances.

**values** [ndarray] Model parameters; `emg3d.fields.Field` instance, or a particular field (e.g. `field.fx`). For fields the method cannot be 'volume'.

**method** [{<'linear'>, 'volume', 'cubic'}, optional] The method of interpolation to perform. The volume averaging method ensures that the total sum of the property stays constant.

Volume averaging is only implemented for model parameters, not for fields. The method 'cubic' requires at least three points in any direction, otherwise it will fall back to 'linear'.

Default is 'linear', because it works with fields and model parameters. However, recommended are 'volume' for model parameters and 'cubic' for fields.

**extrapolate** [bool] If True, points on *new\_grid* which are outside of *grid* are filled by the nearest value (if *method*='cubic') or by extrapolation (if *method*='linear'). If False, points outside are set to zero.

For *method*='volume' it always uses the nearest value for points outside of *grid*.

Default is True.

**log** [bool] If True, the interpolation is carried out on a log10-scale; hence the same as `10**grid2grid(grid, np.log10(values), ...)`. Default is False.

#### Returns

**new\_values** [ndarray] Values corresponding to *new\_grid*.

See also:

**get\_receiver** Interpolation of model parameters or fields to (x, y, z).

## interp3d

`emg3d.maps.interp3d(points, values, new_points, method, fill_value, mode)`

Interpolate values in 3D either linearly or with a cubic spline.

Return *values* corresponding to a regular 3D grid defined by *points* on *new\_points*.

This is a modified version of `scipy.interpolate.interpn()`, using `scipy.interpolate.RegularGridInterpolator` if *method*='linear' and a custom-wrapped version of `scipy.ndimage.map_coordinates()` if *method*='cubic'. If speed is important then choose 'linear', as it can be significantly faster.

#### Parameters

**points** [tuple of ndarray of float, with shapes ((nx, ), (ny, ), (nz, ))] The points defining the regular grid in three dimensions.

**values** [array\_like, shape (nx, ny, nz)] The data on the regular grid in three dimensions.

**new\_points** [tuple (rec\_x, rec\_y, rec\_z)] Coordinates (x, y, z) of new points.

**method** [{<'cubic', 'linear'>}, optional] The method of interpolation to perform, 'linear' or 'cubic'. Default is 'cubic' (forced to 'linear' if there are less than 3 points in any direction).

**fill\_value** [float or None] Passed to `scipy.interpolate.RegularGridInterpolator` if *method*='linear': The value to use for points outside of the interpolation domain. If None, values outside the domain are extrapolated.

**mode** [{<'constant', 'nearest', 'mirror', 'reflect', 'wrap'>}] Passed to `scipy.ndimage.map_coordinates()` if *method*='cubic': Determines how the input array is extended beyond its boundaries.

#### Returns

**new\_values** [ndarray] Values corresponding to *new\_points*.

## edges2cellaverages

`emg3d.maps.edges2cellaverages` (*ex*, *ey*, *ez*, *vol*, *out\_x*, *out\_y*, *out\_z*)

Interpolate fields defined on edges to volume-averaged cell values.

### Parameters

**ex, ey, ez** [ndarray] Electric fields in x-, y-, and z-directions, as obtained from `emg3d.fields.Field`.

**vol** [ndarray] Volumes of the grid, as obtained from `emg3d.meshes.TensorMesh`.

**out\_x, out\_y, out\_z** [ndarray] Arrays where the results are placed (per direction).

## Classes

<code>MapConductivity()</code>	Maps $\sigma$ to computational variable $\sigma$ (conductivity).
<code>MapLgConductivity()</code>	Maps $\log_{10}(\sigma)$ to computational variable $\sigma$ (conductivity).
<code>MapLnConductivity()</code>	Maps $\log_e(\sigma)$ to computational variable $\sigma$ (conductivity).
<code>MapResistivity()</code>	Maps $\rho$ to computational variable $\sigma$ (conductivity).
<code>MapLgResistivity()</code>	Maps $\log_{10}(\rho)$ to computational variable $\sigma$ (conductivity).
<code>MapLnResistivity()</code>	Maps $\log_e(\rho)$ to computational variable $\sigma$ (conductivity).

## MapConductivity

**class** `emg3d.maps.MapConductivity`

Bases: `emg3d.maps._Map`

Maps  $\sigma$  to computational variable  $\sigma$  (conductivity).

- forward:  $x = \sigma$
- backward:  $\sigma = x$

### Methods Summary

<code>backward(mapped)</code>	Mapping to conductivity.
<code>derivative_chain</code> (gradient, mapped)	Chain rule to map gradient from conductivity to mapping space.
<code>forward</code> (conductivity)	Conductivity to mapping.

### Methods Documentation

**backward** (*mapped*)

Mapping to conductivity.

**derivative\_chain** (*gradient*, *mapped*)

Chain rule to map gradient from conductivity to mapping space.

**forward** (*conductivity*)

Conductivity to mapping.

## MapLgConductivity

**class** emg3d.maps.**MapLgConductivity**

Bases: emg3d.maps.\_Map

Maps  $\log_{10}(\sigma)$  to computational variable  $\sigma$  (conductivity).

- forward:  $x = \log_{10}(\sigma)$
- backward:  $\sigma = 10^x$

### Methods Summary

<i>backward</i> (mapped)	Mapping to conductivity.
<i>derivative_chain</i> (gradient, mapped)	Chain rule to map gradient from conductivity to mapping space.
<i>forward</i> (conductivity)	Conductivity to mapping.

### Methods Documentation

**backward** (*mapped*)

Mapping to conductivity.

**derivative\_chain** (*gradient*, *mapped*)

Chain rule to map gradient from conductivity to mapping space.

**forward** (*conductivity*)

Conductivity to mapping.

## MapLnConductivity

**class** emg3d.maps.**MapLnConductivity**

Bases: emg3d.maps.\_Map

Maps  $\log_e(\sigma)$  to computational variable  $\sigma$  (conductivity).

- forward:  $x = \log_e(\sigma)$
- backward:  $\sigma = \exp(x)$

### Methods Summary

<i>backward</i> (mapped)	Mapping to conductivity.
<i>derivative_chain</i> (gradient, mapped)	Chain rule to map gradient from conductivity to mapping space.
<i>forward</i> (conductivity)	Conductivity to mapping.

### Methods Documentation

**backward** (*mapped*)

Mapping to conductivity.

**derivative\_chain** (*gradient*, *mapped*)

Chain rule to map gradient from conductivity to mapping space.

**forward** (*conductivity*)

Conductivity to mapping.

## MapResistivity

**class** emg3d.maps.**MapResistivity**

Bases: emg3d.maps.\_Map

Maps  $\rho$  to computational variable  $\sigma$  (conductivity).

- forward:  $x = \rho = \sigma^{-1}$
- backward:  $\sigma = \rho^{-1} = x^{-1}$

### Methods Summary

<i>backward</i> (mapped)	Mapping to conductivity.
<i>derivative_chain</i> (gradient, mapped)	Chain rule to map gradient from conductivity to mapping space.
<i>forward</i> (conductivity)	Conductivity to mapping.

### Methods Documentation

**backward** (*mapped*)

Mapping to conductivity.

**derivative\_chain** (*gradient*, *mapped*)

Chain rule to map gradient from conductivity to mapping space.

**forward** (*conductivity*)

Conductivity to mapping.

## MapLgResistivity

**class** emg3d.maps.**MapLgResistivity**

Bases: emg3d.maps.\_Map

Maps  $\log_{10}(\rho)$  to computational variable  $\sigma$  (conductivity).

- forward:  $x = \log_{10}(\rho) = \log_{10}(\sigma^{-1})$
- backward:  $\sigma = \rho^{-1} = 10^{-x}$

### Methods Summary

<i>backward</i> (mapped)	Mapping to conductivity.
<i>derivative_chain</i> (gradient, mapped)	Chain rule to map gradient from conductivity to mapping space.
<i>forward</i> (conductivity)	Conductivity to mapping.

### Methods Documentation

**backward** (*mapped*)

Mapping to conductivity.

**derivative\_chain** (*gradient*, *mapped*)

Chain rule to map gradient from conductivity to mapping space.

**forward** (*conductivity*)

Conductivity to mapping.

## MapLnResistivity

**class** `emg3d.maps.MapLnResistivity`

Bases: `emg3d.maps._Map`

Maps  $\log_e(\rho)$  to computational variable  $\sigma$  (conductivity).

- forward:  $x = \log_e(\rho) = \log_e(\sigma^{-1})$
- backward:  $\sigma = \rho^{-1} = \exp(-x)$

### Methods Summary

<code>backward(mapped)</code>	Mapping to conductivity.
<code>derivative_chain(gradient, mapped)</code>	Chain rule to map gradient from conductivity to mapping space.
<code>forward(conductivity)</code>	Conductivity to mapping.

### Methods Documentation

**backward** (*mapped*)

Mapping to conductivity.

**derivative\_chain** (*gradient, mapped*)

Chain rule to map gradient from conductivity to mapping space.

**forward** (*conductivity*)

Conductivity to mapping.

## 5.12.4 emg3d.fields Module

Everything related to the multigrid solver that is a field: source field, electric and magnetic fields, and fields at receivers.

### Functions

<code>get_source_field(grid, src, freq[, strength])</code>	Return the source field.
<code>get_receiver(grid, values, coordinates[, ...])</code>	Return values corresponding to grid at coordinates.
<code>get_receiver_response(grid, field, rec)</code>	Return the field (response) at receiver coordinates.
<code>get_h_field(grid, model, field)</code>	Return magnetic field corresponding to provided electric field.

### get\_source\_field

`emg3d.fields.get_source_field(grid, src, freq, strength=0)`

Return the source field.

The source field is given in Equation 2 in [Mul06],

$$s\mu_0\mathbf{J}_s,$$

where  $s = i\omega$ . Either finite length dipoles or infinitesimal small point dipoles can be defined, whereas the return source field corresponds to a normalized (1 Am) source distributed within the cell(s) it resides (can be changed with the *strength*-parameter).

The adjoint of the trilinear interpolation is used to distribute the point(s) to the grid edges, which corresponds to the discretization of a Dirac ([PIDM07]).

### Parameters

**grid** [TensorMesh] Model grid; a `emg3d.meshes.TensorMesh` instance.

**src** [list of floats] Source coordinates (m). There are two formats:

- Finite length dipole: `[x0, x1, y0, y1, z0, z1]`.
- Point dipole: `[x, y, z, azimuth, dip]`.

**freq** [float] Source frequency (Hz), used to compute the Laplace parameter  $s$ . Either positive or negative:

- $freq > 0$ : Frequency domain, hence  $s = -i\omega = -2i\pi f$  (complex);
- $freq < 0$ : Laplace domain, hence  $s = f$  (real).

**strength** [float or complex, optional] Source strength (A):

- If 0, output is normalized to a source of 1 m length, and source strength of 1 A.
- If  $\neq 0$ , output is returned for given source length and strength.

Default is 0.

### Returns

**sfield** [`SourceField()` instance] Source field, normalized to 1 A m.

## get\_receiver

`emg3d.fields.get_receiver(grid, values, coordinates, method='cubic', extrapolate=False)`

Return values corresponding to grid at coordinates.

Works for electric fields as well as magnetic fields obtained with `get_h_field()`, and for model parameters.

### Parameters

**grid** [`emg3d.meshes.TensorMesh`] The model grid.

**values** [ndarray] Field instance, or a particular field (e.g. `field.fx`); Model parameters.

**coordinates** [tuple (x, y, z)] Coordinates (x, y, z) where to interpolate *values*; e.g. receiver locations.

**method** [str, optional] The method of interpolation to perform, 'linear' or 'cubic'. Default is 'cubic' (forced to 'linear' if there are less than 3 points in any direction).

**extrapolate** [bool] If True, points on *new\_grid* which are outside of *grid* are filled by the nearest value (if `method='cubic'`) or by extrapolation (if `method='linear'`). If False, points outside are set to zero.

Default is False.

### Returns

**new\_values** [ndarray or `utils.EMArray`] Values at *coordinates*.

If input was a field it returns an `EMArray`, which is a subclassed ndarray with `.pha` and `.amp` attributes.

If input was an entire Field instance, output is a tuple (fx, fy, fz).

See also:

**grid2grid** Interpolation of model parameters or fields to a new grid.

**get\_receiver\_response** Get response for arbitrarily rotated receivers.



## get\_receiver\_response

`emg3d.fields.get_receiver_response(grid, field, rec)`

Return the field (response) at receiver coordinates.

### Parameters

**grid** [`emg3d.meshes.TensorMesh`] The model grid.

**field** [`Field`] The electric or magnetic field.

**rec** [tuple (x, y, z, azimuth, dip)] Receiver coordinates and angles (m, °).

All values can either be a scalar or having the same length as number of receivers.

Angles:

- azimuth (°): horizontal deviation from x-axis, anti-clockwise.
- dip (°): vertical deviation from xy-plane up-wards.

### Returns

**responses** [`utils.EMArray`] Responses at receiver.

---

**Note:** Currently only implemented for point receivers, not for finite length dipoles.

---

See also:

`get_receiver` Get values at coordinates (fields and models).

## get\_h\_field

`emg3d.fields.get_h_field(grid, model, field)`

Return magnetic field corresponding to provided electric field.

Retrieve the magnetic field **H** from the electric field **E** using Farady's law, given by

$$\nabla \times \mathbf{E} = i\omega\mu\mathbf{H}.$$

Note that the magnetic field in x-direction is defined in the center of the face defined by the electric field in y- and z-directions, and similar for the other field directions. This means that the provided electric field and the returned magnetic field have different dimensions:

E-field:	x:	[ <code>grid.vectorCCx</code> , <code>grid.vectorNy</code> , <code>grid.vectorNz</code> ]
	y:	[ <code>grid.vectorNx</code> , <code>grid.vectorCCy</code> , <code>grid.vectorNz</code> ]
	z:	[ <code>grid.vectorNx</code> , <code>grid.vectorNy</code> , <code>grid.vectorCCz</code> ]
H-field:	x:	[ <code>grid.vectorNx</code> , <code>grid.vectorCCy</code> , <code>grid.vectorCCz</code> ]
	y:	[ <code>grid.vectorCCx</code> , <code>grid.vectorNy</code> , <code>grid.vectorCCz</code> ]
	z:	[ <code>grid.vectorCCx</code> , <code>grid.vectorCCy</code> , <code>grid.vectorNz</code> ]

### Parameters

**grid** [`TensorMesh`] Model grid; `TensorMesh` instance.

**model** [`Model`] Model; `Model` instance.

**field** [`Field`] Electric field; `Field` instance.

### Returns

**hfield** [`Field`] Magnetic field; `Field` instance.

## Classes

<i>Field</i>	Create a Field instance with x-, y-, and z-views of the field.
<i>SourceField</i>	Create a Source-Field instance with x-, y-, and z-views of the field.

## Field

**class** emg3d.fields.**Field**

Bases: `numpy.ndarray`

Create a Field instance with x-, y-, and z-views of the field.

A *Field* is an *ndarray* with additional views of the x-, y-, and z-directed fields as attributes, stored as *fx*, *fy*, and *fz*. The default array contains the whole field, which can be the electric field, the source field, or the residual field, in a 1D array. A *Field* instance has additionally the property *ensure\_pec* which, if called, ensures Perfect Electric Conductor (PEC) boundary condition. It also has the two attributes *amp* and *pha* for the amplitude and phase, as common in frequency-domain CSEM.

A *Field* can be initiated in three ways:

1. `Field(grid, dtype=np.complex128)`: Calling it with a *emg3d.meshes.TensorMesh* instance returns a *Field* instance of correct dimensions initiated with zeroes of data type *dtype*.
2. `Field(grid, field)`: Calling it with a *emg3d.meshes.TensorMesh* instance and an *ndarray* returns a *Field* instance of the provided *ndarray*, of same data type.
3. `Field(fx, fy, fz)`: Calling it with three *ndarray*'s which represent the field in x-, y-, and z-direction returns a *Field* instance with these views, of same data type.

Sort-order is 'F'.

### Parameters

**fx\_or\_grid** [*emg3d.meshes.TensorMesh* or *ndarray*] Either a *TensorMesh* instance or an *ndarray* of shape `grid.nEx` or `grid.vnEx`. See explanations above. Only mandatory parameter; if the only one provided, it will initiate a zero-field of *dtype*.

**fy\_or\_field** [*Field* or *ndarray*, optional] Either a *Field* instance or an *ndarray* of shape `grid.nEy` or `grid.vnEy`. See explanations above.

**fz** [*ndarray*, optional] An *ndarray* of shape `grid.nEz` or `grid.vnEz`. See explanations above.

**dtype** [*dtype*, optional] Only used if `fy_or_field=None` and `fz=None`; the initiated zero-field for the provided *TensorMesh* has data type *dtype*. Default: `complex`.

**freq** [*float*, optional] Source frequency (Hz), used to compute the Laplace parameter *s*. Either positive or negative:

- *freq* > 0: Frequency domain, hence  $s = -i\omega = -2i\pi f$  (complex);
- *freq* < 0: Laplace domain, hence  $s = f$  (real).

Just added as info if provided.

### Attributes Summary

<i>ensure_pec</i>	Set Perfect Electric Conductor (PEC) boundary condition.
<i>field</i>	Entire field, 1D [ <i>fx</i> , <i>fy</i> , <i>fz</i> ].
<i>freq</i>	Return frequency.

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Table 5.25 – continued from previous page

<i>fx</i>	View of the x-directed field in the x-direction (nC <sub>x</sub> , nN <sub>y</sub> , nN <sub>z</sub> ).
<i>fy</i>	View of the field in the y-direction (nN <sub>x</sub> , nC <sub>y</sub> , nN <sub>z</sub> ).
<i>fz</i>	View of the field in the z-direction (nN <sub>x</sub> , nN <sub>y</sub> , nC <sub>z</sub> ).
<i>is_electric</i>	Returns True if Field is electric, False if it is magnetic.
<i>smu0</i>	Return $s \cdot \mu_0$ ; $\mu_0 = \text{Magn.}$
<i>sval</i>	Return $s$ ; $s=iw$ in frequency domain; $s=\text{freq}$ in Laplace domain.

### Methods Summary

<i>amp()</i>	Amplitude of the electromagnetic field.
<i>copy()</i>	Return a copy of the Field.
<i>from_dict(inp)</i>	Convert dictionary into <i>Field</i> instance.
<i>pha([deg, unwrap, lag])</i>	Phase of the electromagnetic field.
<i>to_dict([copy])</i>	Store the necessary information of the Field in a dict.

### Attributes Documentation

#### **ensure\_pec**

Set Perfect Electric Conductor (PEC) boundary condition.

#### **field**

Entire field, 1D [fx, fy, fz].

#### **freq**

Return frequency.

#### **fx**

View of the x-directed field in the x-direction (nC<sub>x</sub>, nN<sub>y</sub>, nN<sub>z</sub>).

#### **fy**

View of the field in the y-direction (nN<sub>x</sub>, nC<sub>y</sub>, nN<sub>z</sub>).

#### **fz**

View of the field in the z-direction (nN<sub>x</sub>, nN<sub>y</sub>, nC<sub>z</sub>).

#### **is\_electric**

Returns True if Field is electric, False if it is magnetic.

#### **smu0**

Return  $s \cdot \mu_0$ ;  $\mu_0 = \text{Magn. permeability of free space [H/m]}$ .

#### **sval**

Return  $s$ ;  $s=iw$  in frequency domain;  $s=\text{freq}$  in Laplace domain.

### Methods Documentation

#### **amp()**

Amplitude of the electromagnetic field.

#### **copy()**

Return a copy of the Field.

#### **classmethod from\_dict(inp)**

Convert dictionary into *Field* instance.

**Parameters**

**inp** [dict] Dictionary as obtained from `Field.to_dict()`. The dictionary needs the keys `field`, `freq`, `vnEx`, `vnEy`, and `vnEz`.

**Returns**

**obj** [`Field` instance]

**pha** (`deg=False`, `unwrap=True`, `lag=True`)  
Phase of the electromagnetic field.

**Parameters**

**deg** [bool] If True the returned phase is in degrees, else in radians. Default is False (radians).

**unwrap** [bool] If True the returned phase is unwrapped. Default is True (unwrapped).

**lag** [bool] If True the returned phase is lag, else lead defined. Default is True (lag defined).

**to\_dict** (`copy=False`)  
Store the necessary information of the Field in a dict.

**SourceField**

**class** `emg3d.fields.SourceField`

Bases: `emg3d.fields.Field`

Create a Source-Field instance with x-, y-, and z-views of the field.

A subclass of `Field`. Additional properties are the real-valued source vector (`vector`, `vx`, `vy`, `vz`), which sum is always one. For a `SourceField` frequency is a mandatory parameter, unlike for a `Field` (recommended also for `Field` though),

**Parameters**

**fx\_or\_grid** [`emg3d.meshes.TensorMesh` or ndarray] Either a `TensorMesh` instance or an ndarray of shape `grid.nEx` or `grid.vnEx`. See explanations above. Only mandatory parameter; if the only one provided, it will initiate a zero-field of `dtype`.

**fy\_or\_field** [`Field` or ndarray, optional] Either a `Field` instance or an ndarray of shape `grid.nEy` or `grid.vnEy`. See explanations above.

**fz** [ndarray, optional] An ndarray of shape `grid.nEz` or `grid.vnEz`. See explanations above.

**dtype** [dtype, optional] Only used if `fy_or_field=None` and `fz=None`; the initiated zero-field for the provided `TensorMesh` has data type `dtype`. Default: `complex`.

**freq** [float] Source frequency (Hz), used to compute the Laplace parameter  $s$ . Either positive or negative:

- $freq > 0$ : Frequency domain, hence  $s = -i\omega = -2i\pi f$  (complex);
- $freq < 0$ : Laplace domain, hence  $s = f$  (real).

In difference to `Field`, the frequency has to be provided for a `SourceField`.

**Attributes Summary**

<code>vector</code>	Entire vector, 1D [ <code>vx</code> , <code>vy</code> , <code>vz</code> ].
<code>vx</code>	View of the x-directed vector in the x-direction ( <code>nCx</code> , <code>nNy</code> , <code>nNz</code> ).

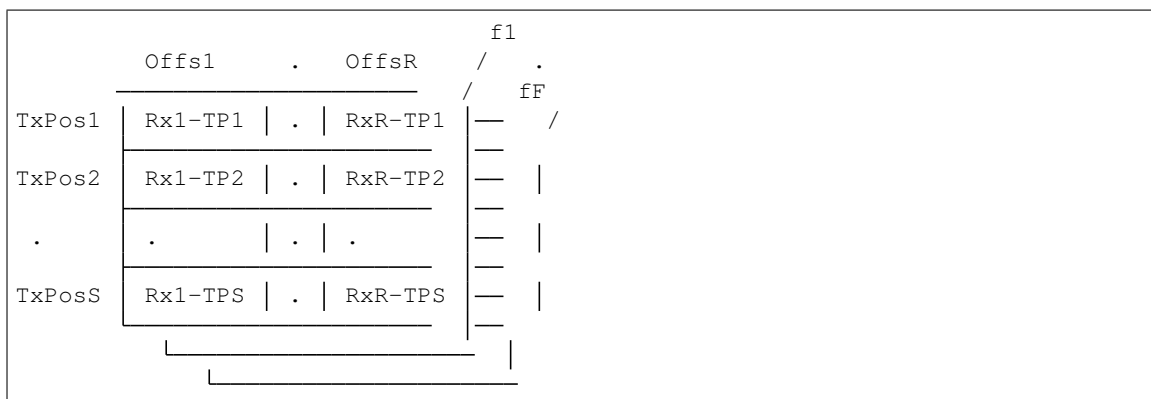
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However, the class can also be used for a CSEM streamer-style survey layout (by setting `fixed=True`), where there is a moving source with one or several receivers at a fixed offset. The layout of the data is then also (S, R, F), but here *S* is the number of locations of the only source, *R* is the number of receiver-offsets, and *F* is the number of frequencies:



This means that even though there is only one source, there are actually *S* source dipoles, as each position is treated as a different dipole. The number of receiver dipoles in this case is *SxR*. This setup can also be used for airborne EM.

### Parameters

**name** [str] Name of the survey

**sources, receivers** [tuple, list, or dict] Sources and receivers.

- Tuples: Coordinates in one of the two following formats:

- $(x, y, z, azimuth, dip)$  [m, m, m, °, °];
- $(x0, x1, y0, y1, z0, z1)$  [m, m, m, m, m, m].

Dimensions will be expanded (hence, if *n* dipoles, each parameter must have length 1 or *n*). These dipoles will be named sequential with *Tx###* and *Rx###*.

The tuple can additionally contain an additional element at the end (after *dip* or *z1*), *electric*, a boolean of length 1 or *n*, that indicates if the dipoles are electric or magnetic.

- List: A list of *Dipole*-instances. The names of all dipoles in the list must be unique.
- Dictionary: A dict of de-serialized *Dipole*-instances; mainly used for loading from file.

**frequencies** [ndarray] Source frequencies (Hz).

**data** [ndarray or None] The observed data (dtype=np.complex128); must have shape (nsr, nrec, nfreq) or, if `fixed=True`, (nsr, noff, nfreq). If None, it will be initiated with NaN's.

**fixed** [bool] Node-based CSEM survey (`fixed=False`; default) or streamer-type CSEM survey (`fixed=True`). In the streamer-type survey, the number of *receivers* supplied must be a multiple of the source positions. In this case, the receivers are grouped into offsets.

**noise\_floor, relative\_error** [float] Noise floor and relative error of the data. Default to None. See *Survey.standard\_deviation* for more info.

**std** [ndarray or None] Standard deviation of the data, same shape as data. Default to None.  
See [\*Survey.standard\\_deviation\*](#) for more info.

### Attributes Summary

<i>data</i>	Data, a <code>xarray.DataSet</code> instance.
<i>frequencies</i>	Frequency array.
<i>noise_floor</i>	Returns the noise floor of the data.
<i>observed</i>	Returns the observed data.
<i>rec_coords</i>	Return receiver coordinates.
<i>receivers</i>	Receiver dict containing all receiver dipoles.
<i>relative_error</i>	Returns the relative error of the data.
<i>shape</i>	Return <code>nsrc x nrec x nfreq</code> .
<i>size</i>	Return actual data size (does NOT equal <code>nsrc x nrec x nfreq</code> ).
<i>sources</i>	Source dict containing all source dipoles.
<i>src_coords</i>	Return source coordinates.
<i>standard_deviation</i>	Returns the standard deviation of the data.

### Methods Summary

<i>copy()</i>	Return a copy of the Survey.
<i>from_dict(inp)</i>	Convert dictionary into <a href="#"><i>Survey</i></a> instance.
<i>from_file(fname[, name])</i>	Load Survey from a file.
<i>to_dict([copy])</i>	Store the necessary information of the Survey in a dict.
<i>to_file(fname[, name])</i>	Store Survey to a file.

### Attributes Documentation

#### **data**

Data, a `xarray.DataSet` instance.

Contains the `xarray.DataArray` element *observed*, but other data can be added. E.g., [\*emg3d.simulations.Simulation\*](#) adds the *synthetic* array.

#### **frequencies**

Frequency array.

#### **noise\_floor**

Returns the noise floor of the data.

See [\*emg3d.surveys.Survey.standard\\_deviation\*](#) for more info.

#### **observed**

Returns the observed data.

#### **rec\_coords**

Return receiver coordinates.

The returned format is *(x, y, z, azm, dip)*, a tuple of 5 tuples. If *fixed=True* it returns a dict with the offsets as keys, and for each offset it returns the corresponding receiver coordinates as just outlined.

#### **receivers**

Receiver dict containing all receiver dipoles.

#### **relative\_error**

Returns the relative error of the data.

See [\*emg3d.surveys.Survey.standard\\_deviation\*](#) for more info.

**shape**

Return  $n_{\text{src}} \times n_{\text{rec}} \times n_{\text{freq}}$ .

Note that not all source-receiver-frequency pairs do actually have data. Check *size* to see how many data points there are.

**size**

Return actual data size (does NOT equal  $n_{\text{src}} \times n_{\text{rec}} \times n_{\text{freq}}$ ).

**sources**

Source dict containing all source dipoles.

**src\_coords**

Return source coordinates.

The returned format is  $(x, y, z, \text{azm}, \text{dip})$ , a tuple of 5 tuples.

**standard\_deviation**

Returns the standard deviation of the data.

The standard deviation can be set by providing an array of the same dimension as the data itself:

```
survey.standard_deviation = ndarray # (nsrc, nrec, nfreq)
```

Alternatively, one can set the *noise\_floor*  $\epsilon_{\text{nf}}$  and the *relative\_error*  $\epsilon_{\text{r}}$ :

```
survey.noise_floor = float or ndarray # (> 0 or None)
survey.relative_error = float or ndarray # (> 0 or None)
```

They must be either floats, or three-dimensional arrays of shape  $([n_{\text{src}} \text{ or } 1], [n_{\text{rec}} \text{ or } 1], [n_{\text{freq}} \text{ or } 1])$ ; dimensions of one will be broadcasted to the corresponding size. E.g., for a dataset of arbitrary amount of sources and receivers with three frequencies you can define a purely frequency-dependent relative error via `relative_error=np.array([err_f1, err_f2, err_f3])[None, None, :]`.

The standard deviation  $\varsigma_i$  of observation  $d_i$  is then given in terms of the noise floor  $\epsilon_{\text{nf};i}$  and the relative error  $\epsilon_{\text{re};i}$  by

$$\varsigma_i = \sqrt{\epsilon_{\text{nf};i}^2 + (\epsilon_{\text{re};i}|d_i|)^2}. \quad (5.35)$$

Note that a set standard deviation is prioritized over potentially also defined noise floor and relative error. To use the noise floor and the relative error after defining standard deviation directly you would have to reset it like

```
survey.standard_deviation = None
```

after which Equation (5.35) would be used again.

## Methods Documentation

**copy()**

Return a copy of the Survey.

**classmethod from\_dict(inp)**

Convert dictionary into *Survey* instance.

**Parameters**

**inp** [dict] Dictionary as obtained from *Survey.to\_dict()*. The dictionary needs the keys *name*, *sources*, *receivers*, *frequencies*, *data*, and *fixed*.

**Returns**

**obj** [*Survey* instance]



**classmethod** `from_file` (*fname*, *name*='survey', *\*\*kwargs*)

Load Survey from a file.

#### Parameters

**fname** [str] File name including extension. Used backend depends on the file extensions:

- '.npz': numpy-binary
- '.h5': h5py-binary (needs *h5py*)
- '.json': json

**name** [str] Name under which the survey is stored within the file.

**kwargs** [Keyword arguments, optional] Passed through to `io.load()`.

#### Returns

**survey** [*Survey*] The survey that was stored in the file.

**to\_dict** (*copy*=False)

Store the necessary information of the Survey in a dict.

**to\_file** (*fname*, *name*='survey', *\*\*kwargs*)

Store Survey to a file.

#### Parameters

**fname** [str] File name inclusive ending, which defines the used data format. Implemented are currently:

- *.h5* (default): Uses *h5py* to store inputs to a hierarchical, compressed binary hdf5 file. Recommended file format, but requires the module *h5py*. Default format if ending is not provided or not recognized.
- *.npz*: Uses *numpy* to store inputs to a flat, compressed binary file. Default format if *h5py* is not installed.
- *.json*: Uses *json* to store inputs to a hierarchical, plain text file.

**name** [str] Name under which the survey is stored within the file.

**kwargs** [Keyword arguments, optional] Passed through to `io.save()`.

## Dipole

**class** `emg3d.surveys.Dipole` (*name*, *coordinates*, *electric*=True, *\*\*kwargs*)

Bases: `emg3d.surveys.PointDipole`

Finite length dipole or point dipole.

Expansion of the basic `PointDipole` to allow for finite length dipoles, and to provide coordinate inputs in the form of (x, y, z, azimuth, dip) or (x0, x1, y0, y1, z0, z1).

Adds attributes *is\_finite*, *electrode1*, *electrode2*, *length*, and *coordinates* to the class.

For *point dipoles*, this gives it a length of unity (1 m), takes its coordinates as center, and computes the two electrode positions.

For *finite length dipoles* it sets the coordinates to its center and computes its length, azimuth, and dip.

Finite length dipoles and point dipoles have therefore the exactly same signature, and can only be distinguished by the attribute *is\_finite*.

#### Parameters

**name** [str] Dipole name.

**coordinates** [tuple of floats] Source coordinates, one of the following:

- (x0, x1, y0, y1, z0, z1): finite length dipole,
- (x, y, z, azimuth, dip): point dipole.

The coordinates x, y, and z are in meters (m), the azimuth and dip in degree (°).

Angles (coordinate system is right-handed with positive z up; East-North-Depth):

- azimuth (°): horizontal deviation from x-axis, anti-clockwise.
- +/-dip (°): vertical deviation from xy-plane down/up-wards.

**electric** [bool] Electric dipole if True, magnetic dipole otherwise. Default is True.

### Attributes Summary

<i>azm</i>
<i>dip</i>
<i>electric</i>
<i>name</i>
<i>xco</i>
<i>yco</i>
<i>zco</i>

### Attributes Documentation

**azm**

**dip**

**electric**

**name**

**xco**

**yco**

**zco**

### PointDipole

**class** emg3d.surveys.**PointDipole** (*name: str, xco: float, yco: float, zco: float, azm: float, dip: float, electric: bool*)

Bases: object

Infinitesimal small electric or magnetic point dipole.

Defined by its coordinates (xco, yco, zco), its azimuth (azm), its dip, and its type (electric).

Not meant to be used directly. Use *Dipole* instead.

#### Parameters

**name** [str] Dipole name.

**xco, yco, zco** [float] x-, y-, and z-coordinates (m).

**azm, dip** [float] Angles (in degrees °); coordinate system is right-handed with positive z up; East-North-Depth:

- azimuth (°): horizontal deviation from x-axis, anti-clockwise.
- +/-dip (°): vertical deviation from xy-plane down/up-wards.

**electric** [bool] Electric dipole if True, magnetic dipole otherwise. Default is True.

## Attributes Summary

<i>azm</i>
<i>dip</i>
<i>electric</i>
<i>name</i>
<i>xco</i>
<i>yco</i>
<i>zco</i>

## Attributes Documentation

**azm**

**dip**

**electric**

**name**

**xco**

**yco**

**zco**

## 5.13.2 emg3d.simulations Module

A simulation is the computation (modelling) of electromagnetic responses of a resistivity (conductivity) model for a given survey.

In its heart, *emg3d* is a multigrid solver for 3D electromagnetic diffusion with tri-axial electrical anisotropy. However, it contains most functionalities to also act as a modeller. The simulation module combines all these things by combining surveys with computational meshes and fields and providing high-level, specialised modelling routines.

## Functions

<i>expand_grid_model</i> (grid, model, expand, interface)	Expand model and grid according to provided parameters.
<i>estimate_gridding_opts</i> (gridding_opts, grid, ...)	Estimate parameters for automatic gridding.

## expand\_grid\_model

`emg3d.simulations.expand_grid_model (grid, model, expand, interface)`

Expand model and grid according to provided parameters.

Expand the grid and corresponding model in positive z-direction from the end of the grid to the interface with `expand[0]`, and a 100 m thick layer above the interface with `expand[1]`.

The provided properties are taken as isotropic; `mu_r` and `epsilon_r` are expanded with ones, if necessary.

The `interface` is usually the sea-surface, and `expand` is therefore `[property_sea, property_air]`.

### Parameters

**grid** [*emg3d.meshes.TensorMesh*] The grid.

**model** [*emg3d.models.Model*] The model.

**expand** [list] The two properties below and above the interface: [below\_interface, above\_interface].

**interface** [float] Interface between the two properties in expand.

#### Returns

**grid** [*emg3d.meshes.TensorMesh*] Expanded grid.

**model** [*emg3d.models.Model*] Expanded model.

### estimate\_gridding\_opts

`emg3d.simulations.estimate_gridding_opts` (*gridding\_opts*, *grid*, *model*, *survey*, *input\_nCz=None*)

Estimate parameters for automatic gridding.

Automatically determines the required gridding options from the provided grid, model, and survey, if they are not provided in *gridding\_opts*.

The dict *gridding\_opts* can contain any input parameter taken by *emg3d.meshes.construct\_mesh()*, see the corresponding documentation for more details with regards to the possibilities.

Different keys of *gridding\_opts* are treated differently:

- The following parameters are estimated from the *model* if not provided:
  - *properties*: volume averages (of x, y, and z-directed properties) on log10 scale of the outermost layer in a given direction.
  - *mapping*: taken from *model*.
- The following parameters are estimated from the *survey* if not provided:
  - *frequency*: average (on log10-scale) of all frequencies.
  - *center*: center of all sources.
  - *domain*: from *vector*, if provided, or
    - \* in x/y-directions: extent of sources and receivers plus 10% on each side, ensuring ratio of 3.
    - \* in z-direction: extent of sources and receivers, ensuring ratio of 2 to horizontal dimension; 1/10 tenth up, 9/10 down.

The ratio means that it is enforced that the survey dimension in x or y-direction is not smaller than a third of the survey dimension in the other direction. If not, the smaller dimension is expanded symmetrically. Similarly in the vertical direction, which must be at least half the dimension of the maximum horizontal dimension or 5 km, whatever is smaller. Otherwise it is expanded in a ratio of 9 parts downwards, one part upwards.

- The following parameter is taken from the *grid* if provided as a string:
  - *vector*: This is the only real “difference” to the inputs of *emg3d.meshes.construct\_mesh()*. The normal input is accepted, but it can also be a string contain any combination of ‘x’, ‘y’, and ‘z’. All directions contained in this string are then taken from the provided grid. E.g., if *gridding\_opts['vector']='xz'* it will take the x- and z-directed vectors from the grid.
- The following parameters are simply passed along if they are provided, nothing is done otherwise:
  - *vector*
  - *stretching*
  - *seasurface*
  - *cell\_numbers*

- `lambda_factor`
- `lambda_from_center`
- `max_buffer`
- `min_width_limits`
- `min_width_pps`
- `verb`

### Parameters

**gridding\_opts** [dict] Containing input parameters to provide to `emg3d.meshes.construct_mesh()`. See the corresponding documentation and the explanations above.

**grid** [`emg3d.meshes.TensorMesh`] The grid.

**model** [`emg3d.models.Model`] The model.

**survey** [`emg3d.surveys.Survey`] The survey.

**input\_nCz** [int, optional] If `expand_grid_model()` was used, `input_nCz` corresponds to the original `grid.nCz`.

### Returns

**gridding\_opts** [dict] Dict to provide to `emg3d.meshes.construct_mesh()`.

## Classes

---

<code>Simulation(name, survey, grid, model[, ...])</code>	Create a simulation for a given survey on a given model.
-----------------------------------------------------------	----------------------------------------------------------

---

## Simulation

**class** `emg3d.simulations.Simulation(name, survey, grid, model, max_workers=4, gridding='single', **kwargs)`

Bases: `object`

Create a simulation for a given survey on a given model.

A simulation can be used to compute responses for an entire survey, hence for an arbitrary amount of sources, receivers, and frequencies. The responses can be computed in parallel over sources and frequencies. It can also be used to compute the misfit with the data and to compute the gradient of the misfit function.

The computational grid(s) can either be provided, or automatic gridding can be used; see the description of the parameters `gridding` and `gridding_opts` for more details.

**Warning:** The automatic gridding does its best to generate meshes that are suitable for the provided model and survey. However, CSEM spans a wide range of acquisition layouts, and frequencies as well as conductivities or resistivities span many orders of magnitude. This makes it hard to have a function that fits all purposes. Check the meshes with your expert knowledge. Also, the automatic gridding is conservative in its estimate, in order to be on the safe side (correct results over speed). This means, however, that often smaller grids could be used by providing the appropriate options in `gridding_opts`.

---

**Note:** The Simulation-class has currently a few limitations:

- `survey.fixed`: must be *False*;

- sources and receivers must be electric;
- 

### Parameters

**survey** [*emg3d.surveys.Survey*] The survey layout, containing sources, receivers, frequencies, and optionally the measured data.

The survey-data will be modified in place. Provide `survey.copy()` if you want to avoid this.

**grid** [*emg3d.meshes.TensorMesh*] The grid. See *emg3d.meshes.TensorMesh*.

**model** [*emg3d.models.Model*] The model. See *emg3d.models.Model*.

**gridding** [str, optional] Method how the computational grids are computed. Default is 'single'. The different methods are:

- 'same': Same grid as for the input model.
- 'single': A single grid for all sources and frequencies.
- 'frequency': Frequency-dependent grids.
- 'source': Source-dependent grids.
- 'both': Frequency- and source-dependent grids.
- 'input': Same as 'single', but instead of automatically generate the mesh it has to be provided in `gridding_opts`.
- 'dict': Same as 'both', but instead of automatically generate the meshes they have to be provided as a dict [source] [frequency] in `gridding_opts`.

See the parameter `gridding_opts` for more details.

**gridding\_opts** [dict or *TensorMesh*, optional] Input format depends on `gridding`:

- 'same': Nothing, `gridding_opts` is not permitted.
- 'single', 'frequency', 'source', 'both': Described below.
- 'input': A *emg3d.meshes.TensorMesh*.
- 'dict': Dictionary of the format dict[source][frequency] containing a *emg3d.meshes.TensorMesh* for each source-frequency pair.

The dict in the case of 'single', 'frequency', 'source', 'both' is passed to *emg3d.meshes.construct\_mesh()*; consult the corresponding documentation for more information. Parameters that are not provided are estimated from the model, grid, and survey using *estimate\_gridding\_opts()*, which documentation contains more information too.

There are two notable differences to the parameters described in *emg3d.meshes.construct\_mesh()*:

- `vector`: besides the normal possibility it can also be a string containing one or several of 'x', 'y', and 'z'. In these cases the corresponding dimension of the input mesh is provided as vector. See *estimate\_gridding\_opts()*.
- `expand`: in the format of [property\_sea, property\_air]; if provided, the input model is expanded up to the searface with sea water, and an air layer is added. The actual height of the searface can be defined with the key `searface`. See *expand\_grid\_model()*.

**solver\_opts** [dict, optional] Passed through to *emg3d.solver.solve()*. The dict can contain any parameter that is accepted by the *emg3d.solver.solve()* except for `grid`, `model`, `sfield`, and `efield`. If not provided the following defaults are used:

- `sslsolver=True`;

- `semicoarsening=True`;
- `linerelaxation=True`;
- `verb=0` (yet warnings are capture and shown).

Note that these defaults are different from the defaults in `emg3d.solver.solve()`. The defaults chosen here will be slower in many cases, but they are the most robust combination at which you can throw most things.

**max\_workers** [int] The maximum number of processes that can be used to execute the given calls. Default is 4.

**verb** [int; optional] Level of verbosity. Default is 1.

- -1: Error.
- 0: Warning.
- 1: Info.
- 2: Debug.

### Attributes Summary

<code>data</code>	Shortcut to <code>survey.data</code> .
<code>gradient</code>	Return the gradient of the misfit function.
<code>misfit</code>	Return the misfit function.

### Methods Summary

<code>clean([what])</code>	Clean part of the data base.
<code>compute([observed])</code>	Compute efields asynchronously for all sources and frequencies.
<code>copy([what])</code>	Return a copy of the Simulation.
<code>from_dict(inp)</code>	Convert dictionary into <code>Simulation</code> instance.
<code>from_file(fname[, name])</code>	Load Simulation from a file.
<code>get_efield(source, frequency, **kwargs)</code>	Return electric field for given source and frequency.
<code>get_efield_info(source, frequency)</code>	Return the solver information of the corresponding computation.
<code>get_grid(source, frequency)</code>	Return computational grid of the given source and frequency.
<code>get_hfield(source, frequency, **kwargs)</code>	Return magnetic field for given source and frequency.
<code>get_model(source, frequency)</code>	Return model on the grid of the given source and frequency.
<code>get_sfield(source, frequency)</code>	Return source field for given source and frequency.
<code>to_dict([what, copy])</code>	Store the necessary information of the Simulation in a dict.
<code>to_file(fname[, what, name])</code>	Store Simulation to a file.

### Attributes Documentation

#### **data**

Shortcut to `survey.data`.

#### **gradient**

Return the gradient of the misfit function.

See `emg3d.optimize.gradient()`.

**misfit**

Return the misfit function.

See `emg3d.optimize.misfit()`.

## Methods Documentation

**clean** (*what*='computed')

Clean part of the data base.

**Parameters**

**what** [str] What to clean. Currently implemented:

- 'computed' (default): Removes all computed properties: electric and magnetic fields and responses at receiver locations.
- 'keepresults': Removes everything except for the responses at receiver locations.
- 'all': Removes everything (leaves it plain as initiated).

**compute** (*observed*=False, *\*\*kwargs*)

Compute efields asynchronously for all sources and frequencies.

**Parameters**

**observed** [bool] If True, it stores the current result also as observed model. This is usually done for pure forward modelling (not inversion). It will as such be stored within the survey. If the survey has either *relative\_error* or *noise\_floor*, random Gaussian noise with std will be added to the *data.observed* (not to *data.synthetic*). Also, data below the noise floor will be set to NaN.

**min\_offset** [float] Default is 0.0. Data in *data.observed* where the offset < min\_offset are set to NaN.

**copy** (*what*='computed')

Return a copy of the Simulation.

See *to\_file* for more information regarding *what*.

**classmethod from\_dict** (*inp*)

Convert dictionary into *Simulation* instance.

**Parameters**

**inp** [dict] Dictionary as obtained from *Simulation.to\_dict()*.

**Returns**

**obj** [*Simulation* instance]

**classmethod from\_file** (*fname*, *name*='simulation', *\*\*kwargs*)

Load Simulation from a file.

**Parameters**

**fname** [str] File name including extension. Used backend depends on the file extensions:

- '.npz': numpy-binary
- '.h5': h5py-binary (needs *h5py*)
- '.json': json

**name** [str] Name under which the simulation is stored within the file.

**kwargs** [Keyword arguments, optional] Passed through to `emg3d.io.load()`.



**Returns**

**simulation** [*Simulation*] The simulation that was stored in the file.

**get\_efield** (*source*, *frequency*, *\*\*kwargs*)

Return electric field for given source and frequency.

**get\_efield\_info** (*source*, *frequency*)

Return the solver information of the corresponding computation.

**get\_grid** (*source*, *frequency*)

Return computational grid of the given source and frequency.

**get\_hfield** (*source*, *frequency*, *\*\*kwargs*)

Return magnetic field for given source and frequency.

**get\_model** (*source*, *frequency*)

Return model on the grid of the given source and frequency.

**get\_sfield** (*source*, *frequency*)

Return source field for given source and frequency.

**to\_dict** (*what*='computed', *copy*=False)

Store the necessary information of the Simulation in a dict.

See *to\_file* for more information regarding *what*.

**to\_file** (*fname*, *what*='computed', *name*='simulation', *\*\*kwargs*)

Store Simulation to a file.

**Parameters**

**fname** [str] File name inclusive ending, which defines the used data format. Implemented are currently:

- *.h5* (default): Uses *h5py* to store inputs to a hierarchical, compressed binary hdf5 file. Recommended file format, but requires the module *h5py*. Default format if ending is not provided or not recognized.
- *.npz*: Uses *numpy* to store inputs to a flat, compressed binary file. Default format if *h5py* is not installed.
- *.json*: Uses *json* to store inputs to a hierarchical, plain text file.

**what** [str] What to store. Currently implemented:

- 'computed' (default): Stores all computed properties: electric fields and responses at receiver locations.
- 'results': Stores only the response at receiver locations.
- 'all': Stores everything.
- 'plain': Only stores the plain Simulation (as initiated).

**name** [str] Name under which the survey is stored within the file.

**kwargs** [Keyword arguments, optional] Passed through to *emg3d.io.save()*.

## 5.14 Optimize

### 5.14.1 emg3d.optimize Module

#### Inversion

Functionalities related to optimization (inversion), e.g., misfit function, gradient of the misfit function, or data- and depth-weighting.

Currently it follows the implementation of [PiMu08], using the adjoint-state technique for the gradient.

## Functions

<code>gradient(simulation)</code>	Compute the discrete gradient using the adjoint-state method.
<code>misfit(simulation)</code>	Return the misfit function.

## gradient

`emg3d.optimize.gradient(simulation)`

Compute the discrete gradient using the adjoint-state method.

The discrete gradient for a single source at a single frequency is given by Equation (10) in [PiMu08],

$$\nabla_p \phi(\mathbf{p}) = - \sum_{k,l,m} \bar{\lambda}_x^E \frac{\partial S}{\partial \mathbf{p}} \mathbf{E}_x - \sum_{k,l,m} \bar{\lambda}_y^E \frac{\partial S}{\partial \mathbf{p}} \mathbf{E}_y - \sum_{k,l,m} \bar{\lambda}_z^E \frac{\partial S}{\partial \mathbf{p}} \mathbf{E}_z ,$$

where the grid notation ( $\{k, l, m\}$  and its  $\{+1/2\}$  equivalents) have been omitted for brevity (except for the sum symbols).

---

**Note:** The gradient is currently implemented only for electric sources and receivers; only for isotropic models; and not for electric permittivity nor magnetic permeability.

---

### Parameters

**simulation** [`emg3d.simulations.Simulation`] The simulation.

### Returns

**grad** [ndarray] Adjoint-state gradient (same shape as `simulation.model`).

## misfit

`emg3d.optimize.misfit(simulation)`

Return the misfit function.

The data misfit or weighted least-squares functional using an  $l_2$  norm is given by

$$\phi = \frac{1}{2} \sum_f \sum_s \sum_r \left\{ \left\| W_{s,r,f} \left( \mathbf{d}_{s,r,f}^{\text{pred}} - \mathbf{d}_{s,r,f}^{\text{obs}} \right) \right\|^2 \right\} + R . \quad (5.36)$$

Here,  $f, s, r$  stand for frequency, source, and receiver, respectively;  $\mathbf{d}^{\text{obs}}$  are the observed electric and magnetic data, and  $\mathbf{d}^{\text{pred}}$  are the synthetic electric and magnetic data. Finally,  $R$  is a regularization term.

The data weight of observation  $d_i$  is given by  $W_i = \varsigma_i^{-1}$ , where  $\varsigma_i$  is the standard deviation of the observation (see `emg3d.surveys.Survey.standard_deviation`).

---

**Note:** This is an early implementation of the misfit function. Currently not yet implemented are:

- Magnetic data;
  - Regularization term.
- 

### Parameters

**simulation** [`emg3d.simulations.Simulation`] The simulation.

**Returns**

**misfit** [float] Value of the misfit function.

## 5.15 I/O and Utils

### 5.15.1 emg3d.io Module

Utility functions for writing and reading data.

**Functions**

<code>save(fname, **kwargs)</code>	Save surveys, meshes, models, fields, and more to disk.
<code>load(fname, **kwargs)</code>	Load meshes, models, fields, and other data from disk.

**save**

`emg3d.io.save(fname, **kwargs)`

Save surveys, meshes, models, fields, and more to disk.

Serialize and save data to disk in different formats (see parameter description of *fname* for the supported file formats). The main emg3d-classes (type `emg3d.io.KNOWN_CLASSES` to get a list) can be collected in corresponding root-folders by setting `collect_classes=True`.

Any other (non-emg3d) object can be added too, as long as it knows how to serialize itself.

The serialized instances will be de-serialized if loaded with `load()`.

**Parameters**

**fname** [str] File name inclusive ending, which defines the used data format. Implemented are currently:

- *.h5*: Uses *h5py* to store inputs to a hierarchical, compressed binary hdf5 file. Recommended file format, but requires the module *h5py*.
- *.npz*: Uses *numpy* to store inputs to a flat, compressed binary file.
- *.json*: Uses *json* to store inputs to a hierarchical, plain text file.

**compression** [int or str, optional] Passed through to *h5py*, default is 'gzip'.

**json\_indent** [int or None] Passed through to *json*, default is 2.

**collect\_classes** [bool] If True, input data is collected in folders for the principal emg3d-classes (type `emg3d.io.KNOWN_CLASSES` to get a list) and everything else collected in a *Data*-folder. Defaults to False.

**verb** [int] If 1 (default) verbose, if 0 silent.

**kwargs** [Keyword arguments, optional] Data to save using its key as name. The following instances will be properly serialized: `emg3d.meshes.TensorMesh`, `emg3d.fields.Field`, and `emg3d.models.Model` and serialized again if loaded with `load()`. These instances are collected in their own group if *h5py* is used.

Note that the provided data cannot contain the before described parameters as keys.

## load

`emg3d.io.load(fname, **kwargs)`

Load meshes, models, fields, and other data from disk.

Load and de-serialize `emg3d.meshes.TensorMesh`, `emg3d.fields.Field`, and `emg3d.models.Model` instances and add arbitrary other data that were saved with `save()`.

### Parameters

**fname** [str] File name including extension. Possibilities:

- `‘.npz’`: numpy-binary
- `‘.h5’`: h5py-binary (needs `h5py`)
- `‘.json’`: json

**verb** [int] If 1 (default) verbose, if 0 silent.

### Returns

**out** [dict] A dictionary containing the data stored in `fname`; `emg3d.meshes.TensorMesh`, `emg3d.fields.Field`, and `emg3d.models.Model` instances are de-serialized and returned as instances.

## 5.15.2 emg3d.utils Module

Utility functions for the multigrid solver.

### Classes

<code>Fourier(time, fmin, fmax[, signal, ft, ftarg])</code>	Time-domain CSEM computation.
<code>Time()</code>	Class for timing (now; runtime).
<code>Report([add_pkg, ncol, text_width, sort])</code>	Print date, time, and version information.
<code>EMArray</code>	Create an EM-ndarray: add <i>amplitude</i> <amp> and <i>phase</i> <pha> methods.

### Fourier

**class** `emg3d.utils.Fourier` (*time, fmin, fmax, signal=0, ft='dlf', ftarg=None, \*\*kwargs*)

Bases: `object`

Time-domain CSEM computation.

Class to carry out time-domain modelling with the frequency-domain code *emg3d*. Instances of the class take care of computing the required frequencies, the interpolation from coarse, limited-band frequencies to the required frequencies, and carrying out the actual transform.

Everything related to the Fourier transform is done by utilising the capabilities of the 1D modeller *empymod*. The input parameters *time*, *signal*, *ft*, and *ftarg* are passed to the function `empymod.utils.check_time()` to obtain the required frequencies. The actual transform is subsequently carried out by calling `empymod.model.tem()`. See these functions for more details about the exact implementations of the Fourier transforms and its parameters. Note that also the *verb*-argument follows the definition in *empymod*.

The mapping from computed frequencies to the frequencies required for the Fourier transform is done in three steps:

- Data for  $f > f_{\max}$  is set to  $0+0j$ .
- Data for  $f < f_{\min}$  is interpolated by adding an additional data point at a frequency of 1e-100 Hz.

The data for this point is `data.real[0]+0j`, hence the real part of the lowest computed frequency and zero imaginary part. Interpolation is carried out using PCHIP `scipy.interpolate.pchip_interpolate()`.

- Data for  $f_{\min} \leq f \leq f_{\max}$  is computed with cubic spline interpolation (on a log-scale) `scipy.interpolate.InterpolatedUnivariateSpline`.

Note that *fmin* and *fmax* should be chosen wide enough such that the mapping for  $f > f_{\max}$   $f < f_{\min}$  does not matter that much.

### Parameters

**time** [ndarray] Desired times (s).

**fmin, fmax** [float] Minimum and maximum frequencies (Hz) to compute:

- Data for  $\text{freq} > f_{\max}$  is set to  $0+0j$ .
- Data for  $\text{freq} < f_{\min}$  is interpolated, using an extra data-point at  $f = 1\text{e-}100$  Hz, with value `data.real[0]+0j`. (Hence zero imaginary part, and the lowest computed real value.)

**signal** [{0, 1, -1}, optional]

**Source signal, default is 0:**

- None: Frequency-domain response
- -1 : Switch-off time-domain response
- 0 : Impulse time-domain response
- +1 : Switch-on time-domain response

**ft** [{‘sin’, ‘cos’, ‘fftlog’}, optional] Flag to choose either the Digital Linear Filter method (Sine- or Cosine-Filter) or the FFTLog for the Fourier transform. Defaults to ‘sin’.

**ftarg** [dict, optional] Depends on the value for *ft*:

- If *ft*=‘dlf’:
  - *dlf*: string of filter name in `empymod.filters` or the filter method itself. (Default: `empymod.filters.key_201_CosSin_2012()`)
  - *pts\_per\_dec*: points per decade; (default: -1)
    - \* If 0: Standard DLF.
    - \* If < 0: Lagged Convolution DLF.
    - \* If > 0: Splined DLF
- If *ft*=‘fftlog’:
  - *pts\_per\_dec*: sampels per decade (default: 10)
  - *add\_dec*: additional decades [left, right] (default: [-2, 1])
  - *q*: exponent of power law bias (default: 0);  $-1 \leq q \leq 1$

**freq\_inp** [array] Frequencies to use for computation. Mutually exclusive with *every\_x\_freq*.

**every\_x\_freq** [int] Every *every\_x\_freq*-th frequency of the required frequency-range is used for computation. Mutually exclusive with *freq\_calc*.

### Attributes Summary

<i>every_x_freq</i>	If set, <i>freq_coarse</i> is <i>every_x_freq</i> -frequency of <i>freq_req</i> .
<i>fmax</i>	Maximum frequency (Hz) to compute.
<i>fmin</i>	Minimum frequency (Hz) to compute.
<i>freq_calc</i>	Frequencies at which the model has to be computed.
<i>freq_calc_i</i>	Indices of <i>freq_coarse</i> which have to be computed.
<i>freq_coarse</i>	Coarse frequency range, can be different from <i>freq_req</i> .
<i>freq_extrapolate</i>	These are the frequencies to extrapolate.
<i>freq_extrapolate_i</i>	Indices of the frequencies to extrapolate.
<i>freq_inp</i>	If set, <i>freq_coarse</i> is set to <i>freq_inp</i> .
<i>freq_interpolate</i>	These are the frequencies to interpolate.
<i>freq_interpolate_i</i>	Indices of the frequencies to interpolate.
<i>freq_req</i>	Frequencies required to carry out the Fourier transform.
<i>ft</i>	Type of Fourier transform.
<i>ftarg</i>	Fourier transform arguments.
<i>signal</i>	Signal in time domain {0, 1, -1}.
<i>time</i>	Desired times (s).

### Methods Summary

<i>fourier_arguments</i> (ft, ftarg)	Set Fourier type and its arguments.
<i>freq2time</i> (fdata, off)	Compute corresponding time-domain signal.
<i>interpolate</i> (fdata)	Interpolate from computed data to required data.

### Attributes Documentation

#### **every\_x\_freq**

If set, *freq\_coarse* is *every\_x\_freq*-frequency of *freq\_req*.

#### **fmax**

Maximum frequency (Hz) to compute.

#### **fmin**

Minimum frequency (Hz) to compute.

#### **freq\_calc**

Frequencies at which the model has to be computed.

#### **freq\_calc\_i**

Indices of *freq\_coarse* which have to be computed.

#### **freq\_coarse**

Coarse frequency range, can be different from *freq\_req*.

#### **freq\_extrapolate**

These are the frequencies to extrapolate.

In fact, it is done via interpolation, using an extra data-point at  $f = 1e-100$  Hz, with value `data.real[0]+0j`. (Hence zero imaginary part, and the lowest computed real value.)

#### **freq\_extrapolate\_i**

Indices of the frequencies to extrapolate.

#### **freq\_inp**

If set, *freq\_coarse* is set to *freq\_inp*.

**freq\_interpolate**

These are the frequencies to interpolate.

If `freq_req` is equal `freq_coarse`, then this is equal to `freq_calc`.

**freq\_interpolate\_i**

Indices of the frequencies to interpolate.

If `freq_req` is equal `freq_coarse`, then this is equal to `freq_calc_i`.

**freq\_req**

Frequencies required to carry out the Fourier transform.

**ft**

Type of Fourier transform. Set via `fourier_arguments(ft, ftarg)`.

**ftarg**

Fourier transform arguments. Set via `fourier_arguments(ft, ftarg)`.

**signal**

Signal in time domain {0, 1, -1}.

**time**

Desired times (s).

## Methods Documentation

**fourier\_arguments** (*ft, ftarg*)

Set Fourier type and its arguments.

**freq2time** (*fdata, off*)

Compute corresponding time-domain signal.

Carry out the actual Fourier transform.

**Parameters**

**fdata** [ndarray] Frequency-domain data corresponding to *freq\_calc*.

**off** [float] Corresponding offset (m).

**Returns**

**tdata** [ndarray] Time-domain data corresponding to `Fourier.time`.

**interpolate** (*fdata*)

Interpolate from computed data to required data.

**Parameters**

**fdata** [ndarray] Frequency-domain data corresponding to *freq\_calc*.

**Returns**

**full\_data** [ndarray] Frequency-domain data corresponding to *freq\_req*.

## Time

**class** `emg3d.utils.Time`

Bases: `object`

Class for timing (now; runtime).

## Attributes Summary

<i>elapsed</i>	Return runtime in seconds since time zero.
<i>now</i>	Return string of current time.
<i>runtime</i>	Return string of runtime since time zero.
<i>t0</i>	Return time zero of this class instance.

## Attributes Documentation

### **elapsed**

Return runtime in seconds since time zero.

### **now**

Return string of current time.

### **runtime**

Return string of runtime since time zero.

### **t0**

Return time zero of this class instance.

## Report

**class** emg3d.utils.**Report** (*add\_pkg=None, ncol=3, text\_width=80, sort=False*)

Bases: `scooby.report.Report`

Print date, time, and version information.

Use *scooby* to print date, time, and package version information in any environment (Jupyter notebook, IPython console, Python console, QT console), either as html-table (notebook) or as plain text (anywhere).

Always shown are the OS, number of CPU(s), *numpy*, *scipy*, *emg3d*, *numba*, *sys.version*, and time/date.

Additionally shown are, if they can be imported, *IPython* and *matplotlib*. It also shows MKL information, if available.

All modules provided in *add\_pkg* are also shown.

---

**Note:** The package *scooby* has to be installed in order to use *Report*: `pip install scooby`.

---

## Parameters

**add\_pkg** [packages, optional] Package or list of packages to add to output information (must be imported beforehand).

**ncol** [int, optional] Number of package-columns in html table (no effect in text-version); Defaults to 3.

**text\_width** [int, optional] The text width for non-HTML display modes

**sort** [bool, optional] Sort the packages when the report is shown

## Examples

```
>>> import pytest
>>> import dateutil
>>> from emg3d import Report
>>> Report()                                # Default values
>>> Report(pytest)                          # Provide additional package
>>> Report([pytest, dateutil], ncol=5)      # Set nr of columns
```



## EMArray

**class** emg3d.utils.EMArray

Bases: `numpy.ndarray`

Create an EM-ndarray: add *amplitude* <amp> and *phase* <pha> methods.

### Parameters

**data** [array] Data to which to add *.amp* and *.pha* attributes.

### Examples

```
>>> import numpy as np
>>> from empymod.utils import EMArray
>>> emvalues = EMArray(np.array([1+1j, 1-4j, -1+2j]))
>>> print(f"Amplitude      : {emvalues.amp()}")
Amplitude      : [1.41421356  4.12310563  2.23606798]
>>> print(f"Phase (rad)    : {emvalues.pha()}")
Phase (rad)     : [ 0.78539816 -1.32581766 -4.24874137]
>>> print(f"Phase (deg)    : {emvalues.pha(deg=True)}")
Phase (deg)     : [ 45.          -75.96375653 -243.43494882]
>>> print(f"Phase (deg; lag) : {emvalues.pha(deg=True, lag=False)}")
Phase (deg; lag) : [-45.          75.96375653  243.43494882]
```

### Methods Summary

<code>amp()</code>	Amplitude of the electromagnetic field.
<code>pha([deg, unwrap, lag])</code>	Phase of the electromagnetic field.

### Methods Documentation

**amp()**

Amplitude of the electromagnetic field.

**pha** (*deg=False, unwrap=True, lag=True*)

Phase of the electromagnetic field.

#### Parameters

**deg** [bool] If True the returned phase is in degrees, else in radians. Default is False (radians).

**unwrap** [bool] If True the returned phase is unwrapped. Default is True (unwrapped).

**lag** [bool] If True the returned phase is lag, else lead defined. Default is True (lag defined).

## 5.16 Command Line Interface

Functions related to the command-line interface (CLI) of *emg3d*.

Consult the *CLI interface* section in the documentation for more information.

### 5.16.1 emg3d.cli.main Module

Entry point for the command-line interface (CLI).

#### Functions

---

<code>main([args])</code>	Parsing command line inputs of CLI interface.
---------------------------	-----------------------------------------------

---

#### main

`emg3d.cli.main.main(args=None)`  
Parsing command line inputs of CLI interface.

### 5.16.2 emg3d.cli.parser Module

Parser for the configuration file of the command-line interface.

#### Functions

---

<code>parse_config_file(args_dict)</code>	Read and parse the configuration file and set defaults.
-------------------------------------------	---------------------------------------------------------

---

#### parse\_config\_file

`emg3d.cli.parser.parse_config_file(args_dict)`  
Read and parse the configuration file and set defaults.

##### Parameters

**args\_dict** [dict] Arguments from terminal, see `emg3d.cli.main()`.

##### Returns

**conf** [dict] Configuration-dict.

### 5.16.3 emg3d.cli.run Module

Functions that actually call emg3d within the CLI interface.

#### Functions

---

<code>initiate_logger(cfg, runtime, verb)</code>	Initiate logger for CLI of emg3d.
<code>simulation(args_dict)</code>	Run <i>emg3d</i> invoked by CLI.

---

#### initiate\_logger

`emg3d.cli.run.initiate_logger(cfg, runtime, verb)`  
Initiate logger for CLI of emg3d.

## simulation

`emg3d.cli.run.simulation(args_dict)`

Run *emg3d* invoked by CLI.

Run and log *emg3d* given the settings stored in the config file, overruled by settings passed in *args\_dict* (which correspond to command-line arguments).

Results are saved to files according to provided settings.

### Parameters

**args\_dict** [dict] Arguments from terminal, see `emg3d.cli.main()`. Parameters passed in *args\_dict* overrule parameters in the *config*.



---

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