SEM-DSM hybrid method User Manual

Wenbo Wu^{1,2}, Sidao Ni¹, Zhongwen Zhan³, Shengji Wei⁴
¹State Key Laboratory of Geodesy and Earth's Dynamics, Institute of Geodesy and Geophysics, Chinese Academy of Sciences. Email: shaguan@gmail.com

² Department of Geosciences, Princeton University.

³Seismological Laboratory, California Institute of Technology.

⁴Earth Observatory of Singapore, Nanyang Technological University.

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Introduction

Despite computational capability of High Performance Computing (HPC) was being dramatically enhanced, simulating high frequency (e.g. 1Hz or higher) seismic wave propagation at the global scale is still prohibitive. Thus, we develop this software package SEM-DSM-hybrid to efficiently compute teleseismic synthetics with a local 3D structure limited in the source-side, mid-way or receiver-side. Currently, only the source-side SEM-DSM coupling is implemented and carefully benchmarked in this package. A receiver-side SEM-DSM coupling has been implemented by Monteiller et al. [2012]. In the future, we may add the receiver-side and mid-way coupling in this package.

1.1 General background

In this hybrid method, we combine the advantages of SPECFEM3D_Cartesian and DSM to efficiently compute teleseismic synthetics. As a 1D synthetic simulation tool, DSM has the advantage of high accuracy and cheap computational cost [Geller and Ohminato, 1994, Geller and Takeuchi, 1995, Takeuchi et al., 1996, Kawai et al., 2006]; As a 3D wave propagation simulation tool, SEM can handle full 3D velocity structures, topography, fluid/solid interfaces, and complex geometries accurately [Komatitsch and Tromp, 1999, 2002]. If you want to know more about SPECFEM3D and DSM, please read the relevant literature.

Our source-side SEM-DSM coupling is composed of three steps, SEM simulation, DSM computation and the final step of coupling (Fig. 1.1). More details about the coupling theory and more benchmark examples are described in Wu et al. [2018]. Our hybrid method is useful for a series of problems in seismology, such as imaging 3D structures of a subducting slab or a mid-ocean ridge and studying source parameters with 3D source-side complexities using teleseismic waveforms.

1.2 Citation

If you use this DSM-SEM hybrid method, please cite Wu et al. [2018].

We have used this method to investigate some earthquakes and you may be also interested in the following articles: Wang et al. [2017, 2018]

1.3 Support

This work was supported by funding from National Basic Research Program of China (973 Program) through grant 2014CB845901. Shengji Wei is supported by the internal grant (M4430255) of Earth Observatory of Singapore.

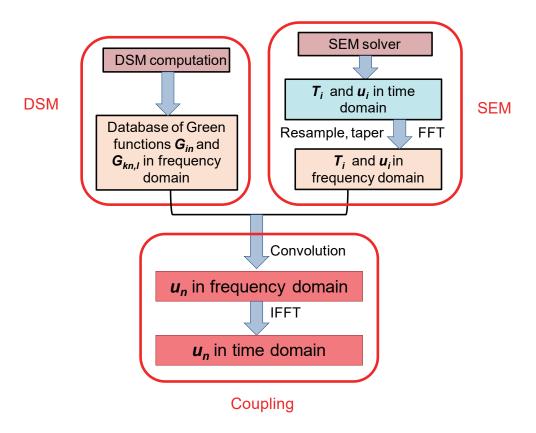


Figure 1.1: Work flow of SEM-DSM coupling at source-side.

Getting Started

To download the SEM_DSM_hybrid software package, type this:

```
git clone --recursive --branch devel https://github.com/wenbowu-geo/SEM-DSM-coupling
```

Before compiling the packages, you may need to load all the needed modules (e.g. openmpi, intel et al.). We have successfully compiled this software package on Tiger supercomputer in Princeton University, Fram cluster at SeismoLab of Caltech and TianHe-2 supercomputer in China.

2.1 Compile SPECFEM3D_Cartesian packages

The SPECFEM3D_Cartesian manual provides detailed information of compiling the codes. We modified some source codes, but most tricks introduced there should also work here. First, you need to go to the SEM root directory SEM_DSM_hybrid/specfem3d and run the configure shell script to produce the file Makefile:

```
>cd SEM_DSM_hybrid/specfem3d/
>./configure FC=ifort CC=icc MPIFC=mpif90 --with-mpi
```

You can replace the Intel compilers above (ifort and icc) with other compilers if you want to; for instance use GNU compilers FC=gfortran and FC=gcc instead.

Finally, type make all to compile the codes

```
>make all
```

To be consistent with DSM, we use cubed-sphere coordinates in the internal mesher. We provide tools to convert geographical topography, as a function of latitude and longitude, to that in cubed-sphere coordinates (ξ,η) . The first tool xcubedsphere_topo help you convert topography for interfaces. The topography is tapered near the boundaries of SEM box and some extremely high or low topography/bathymetry could be truncated if necessary. To compile it, type

```
>cd SEM_DSM_hybrid/specfem3d/src/cubedsphere_topo/
>make clean
>make
```

Another tool xcubedsphere_topo_FindOcean does a similar job, but it does not taper and/or truncate the topography. The tool is needed when incorporate ocean. To compile this tool, type

```
>cd SEM_DSM_hybrid/specfem3d/src/cubedsphere_topo_FindOcean/
>make clean
>make
```

2.2 Compile DSM codes

To compile, type

```
>cd SEM_DSM_hybrid/DSM/src/
>make
```

Again, you may need to replace the Intel compiler ifort with gfortran in the Makefile.

2.3 Compile coupling codes

We provide two different interpolation schemes to accomplish the final step of coupling (see 5). To compile the codes of linear interpolation scheme, type

```
>cd SEM_DSM_hybrid/coupling/Linear_piecewise_interpolation/
>make
```

To compile the codes of Lagrange interpolation scheme, type

```
>cd SEM_DSM_hybrid/coupling/Lagrange_interpolation/
>make
```

SEPCFEM3D

The same as the original SEPCFEM3D-Cartesian, running SEM in our coupling mode is composed of three steps: meshing, generating databases and running solver. The usage of SEPCFEM3D-Cartesian in method is almost the same as the original code, so we recommend users to read the SPECFEM3D-Cartesian Manual before running this package. However, some changes are made to implement the coupling and the following subsections list these changes.

The first step in running a spectral-element simulation consists of constructing a high-quality mesh for the region under consideration. We provide two possibilities to do so: (1) relying on the external, hexahedral mesher CUBIT, or (2) using the provided, internal mesher xmeshfem3D. In the following, we explain these two approaches.

3.1 Meshing with xmeshfem3D in a cubed-sphere coordinate system

Meshing is the first step of running SPECFEM3D-Cartesian. The original package provides two ways to do meshing: (1) A professional meshing tool, such as Cubit, Gmesh or other meshing tools, or (2) internal meshing package xmeshfem3D. The current version of our SEM-DSM coupling only allows users to use the internal meshing xmeshfem3D. In order to couple SEM with DSM, we made the below changes to the original SPECFEM3D-Cartesian.

First, we use cubed-sphere coordinates here, instead of UTM coordinates in the original code. Using the original xmeshfem3D, the meshed model must be a regular box, whose edges are aligned with the X, Y and Z axises. In our SEM-DSM coupling method, we modify the subroutines to create a small chunk in a cubed-sphere coordinate system, instead of a X-Y-Z box. By doing that, the curvature of interfaces within the Earth are reserved (see Figure ??). Another advantage of cubed-sphere coordinates is making elements at the coupling boundary have regular shape, that results in a small number of depths of GLL points and therefore significantly simplifies DSM computation.

Secondly, ocean water, if existing, is incorporated via xmeshfem3D. Investigating effects of ocean water on teleseismic waves is one of the major motivations for developing this code, so we write a subroutine named

```
redefine ocean land.f90
```

to add ocean water. Running xmeshfem3D reads the file real_bathymetry_topography, which specifies the real topography/bathymetry as a function of cubed-sphere coordinates (ξ, η) , and then define elements as either a solid media type or a fluid media type.

The two main parameter files in the meshing step are Par_file and Mesh_Par_file. In the file Par_file, all the previously used parameters are retained and we add the below new parameters:

CUBED_SPHERE_PROJECTION You must set this flag to .true. when the SEM-DSM coupling mode is enabled.

COUPLING_TYPE 1 for source-side coupling, 2 for station-side coupling, 3 for mid-way coupling and 4 for no coupling. The current version of our code only allows source-side coupling mode or no coupling mode, so do not set it to 2 and 3.

Most parameters needed by xmeshfem3D are specified in the file Mesh_Par_file. The functions of most parameters in this file does not change in our coupling method, except the below ones:

LATITUDE_MIN is not functional. Instead, the dimension of block is defined in another file SEMtoTele_Par_file.

LATITUDE_MAX is not functional.

LONGITUDE MIN is not functional.

LONGITUDE_MAX is not functional.

DEPTH_BLOCK_KM has the same meaning, but its unit is km.

UTM PROJECTION ZONE is not functional.

SUPPRESS_UTM_PROJECTION is not functional.

- **INTERFACES_FILE** This file contains the information of all interfaces. Its format is not changed, but the topography of each individual interface is given in the cubed-sphere coordinates. We provide two executable files xcubedsphere_topo and xcubedsphere_topo_FindOcean to convert topography (lat,lon) into topography (ξ,η) .
- **CUBED_SPHERE_PROJECTION** is a new parameter in our SEM-DSM code. You must set this flag to .true. when the SEM-DSM coupling mode is enabled.
- NDOUBLINGS The number of horizontal doubling layers. In our SEM-DSM coupling, it must be set either 1 or 2 if USE_REGULAR_MESH is set to .true.. In the original code, it can be larger than 2. Multiple mesh doublings can be chosen, for which each an NZ_DOUBLING_** entry must be given. By default, we only provide two possible entries in the Mesh_Par_file. For higher numbers of doubling layers, additional entries must be added.
- NZ_DOUBLING_1 The position of the first doubling layer (only interpreted if USE_REGULAR_MESH is set to .true.).
- NZ_DOUBLING_2 The position of the second doubling layer (only interpreted if USE_REGULAR_MESH is set to .true. and if NDOUBLINGS is set to 2). Doubling layers must be at least 2 layers apart. The layer count starts from the bottom layer. More entries must be listed by the user if NDOUBLINGS is larger than 2.
- **THICKNESS_OF_X_PML** The thickness of PML layer at max_{ξ} and min_{ξ} . Different from the original code, the unit in the SEM-DSM mode is degree (1 degree \cong 111.2 km at the free surface of the Earth). Typically, a 0.1 degrees (roughly 11 km) thick PML layer works well for seismic waves with a period range of 0.5s-100s and 0.2 degrees is sufficiently high. However, you must set it thicker than the size of at least one element in the ξ direction. Other wise, it PML layer does work well. If irregular meshes are used and NDOUBLINGS=1, the final PML thickness must be $2\times$ a multiple of finer_element_size on the ξ direction, so the value of this parameter must be greater than $2\times M\times$ finer_element_size and smaller than $(2\times M+1)\times$ finer_element_size. If NDOUBLINGS=2, $2\times M\times$ finer_element_size is replaced by $4\times M\times$ finest_element_size.
- **THICKNESS_OF_Y_PML** The thickness of PML layer at $max_{-}\eta$ and $max_{-}\eta$, in unit of degree. More details see the above parameter <code>THICKNESS_OF_X_PML</code>.
- **THICKNESS_OF_Z_PML** The thickness of PML layer at bottom, in unit of meter. Usually, a value of 10000m-20000m is proper. The same as THICKNESS_OF_X_PML and THICKNESS_OF_Y_PML, you must guarantee the PML layer to work on at least one element size. If irregular meshes are used, the transition from fine to coarse grids usually happens well above the bottom boundary of SEM box, so just set this parameter to a thickness of a few elements at the most bottom.
- NREGIONS Do not specify any fluid block here. The ocean water is defined by the file

real_bathymetry_topography

In addition to the file Mesh_Par_file and Par_file, another file SEMtoTele_Par_file is new and only works for SEM-DSM coupling. It includes the following parameters:

ANGULAR_WIDTH_XI_IN_DEGREES The dimension of block on the ξ direction, in unit of degree. If

```
GAMMA_ROTATION_AZIMUTH=0.0,
```

positive ξ at the central position of block points to the East or longitude direction.

ANGULAR_WIDTH_ETA_IN_DEGREES The dimension of block on the η direction, in unit of degree. If

```
GAMMA_ROTATION_AZIMUTH=0.0,
```

positive η at the central position of block points to the North direction.

- **CENTER_LATITUDE_IN_DEGREES** Latitude of the central position of the block. Usually, it is close to the latitude of the earthquake hypocenter.
- **CENTER_LONGITUDE_IN_DEGREES** Longitude of the central position of the block. Usually, it is close to the longitude of the earthquake hypocenter.
- **GAMMA_ROTATION_AZIMUTH** How much the block is counterclockwise rotated, in unit of degree. Sometimes, rotating the block would save computational resources or simplify the model setup.
- **nx_TopographyTaper** The number of elements used to make the topography tapering in the ξ direction. A number between 4-20 is usually proper.
- **ny_TopographyTaper** The number of elements used to make the topography tapering in the η direction. A number between 4-20 is usually proper
- nx_NoTopography The number of elements on left side max_ξ and right side min_ξ of the block, where topography is tapered to zero. In the current version, topography is not allowed in the PML absorbing layer, if PML layer used, and on the coupling boundary, nx_NoTopography must be sufficiently large to ensure that. A number between 4-20 is usually proper, but it depends on the parameters THICKNESS_OF_X_PML, ix_LowBound and ix_HighBound.
- ny_NoTopography The number of elements on left side $max_{-}\eta$ and right side $min_{-}\eta$ of the block, where topography is tapered to zero. In the current version, topography is not allowed in the PML absorbing layer, if PML layer used, and on the coupling boundary, ny_NoTopography must be sufficiently large to ensure that. A number between 4-20 is usually proper, but it depends on the parameters THICKNESS_OF_X_PML, iy_LowBound and iy_HighBound.
- **ix_LowBound** The left position of coupling boundary, in unit of number of elements. If PML layer used, this parameter be sufficiently large to ensure the coupling boundary not within the PML layer. If regular meshes are used, the number must be a multiple of 2 for NDOUBLINGS=1 and a multiple of 4 for NDOUBLINGS=2.
- **ix_HighBound** The right position of coupling boundary, in unit of number of elements. If PML layer used, this parameter be sufficiently small to ensure the coupling boundary not within the PML layer. If regular meshes are used, the number must be a multiple of 2 for NDOUBLINGS=1 and a multiple of 4 for NDOUBLINGS=2.
- **iy_LowBound** The upper position of coupling boundary, in unit of number of elements. If PML layer used, this parameter be sufficiently large to ensure the coupling boundary not within the PML layer. If regular meshes are used, the number must be a multiple of 2 for NDOUBLINGS=1 and a multiple of 4 for NDOUBLINGS=2.
- **iy_HighBound** The lower position of coupling boundary, in unit of number of elements. If PML layer used, this parameter be sufficiently small to ensure the coupling boundary not within the PML layer. If regular meshes are used, the number must be a multiple of 2 for NDOUBLINGS=1 and a multiple of 4 for NDOUBLINGS=2.
- **ir_BoundTop** The top of coupling boundary, in unit of number of elements. For source-side SEM-DSM coupling, this number must be equal to NZ, the total number of elements on the vertical direction.

- **ir_BoundBot** The bottom position of coupling boundary, in unit of number of elements. a number between 3-5 is usually good, but you must ensure the coupling boundary is not with PML layer.
- **LOW_RESOLUTION** Choose low or high spatial resolution coupling mode. If .True., only the central GLL points of coupling faces are used. If .True., the displacement and strain on all the coupling GLL points are saved to get more accurate results, but it usually needs a large size of disk storage and slows down the SEM computation.
- **NSTEP_BETWEEN_OUTPUTBOUND** How many time steps we save the displacement and strain of coupling GLL points. Note that 'time step' here is after decimating.
- **DECIMATE_COUPLING** Resampling rate of the saved displacement and strain. Resampling the displacement and strain helps us save disk storage, but resampling them down could bring in large errors. Thus, you should avoid too much information missed in resampling, e.g. ensure at least eight points per source duration. However, when you use Dirac delta source time function or the specified source duration is smaller than the minimum period resolved, resampling rate must be 1. Otherwise, DECIMATE_COUPLING>1 could cause large errors.
- **NPOINTS_PER_PACK** The number of GLL points in one output package. In order to avoid large size files, we divide the coupling GLL points into a number of groups and the output results of each group are saved as one package. A number between 200-1000 is usually proper.

```
Some output files after running xmeshfem3D are:
```

```
OUTPUT_FILES/DATABASES_MPI/proc??????_Database
OUTPUT_FILES/DATABASES_MPI/proc??????_TeleEle_rBot.info
OUTPUT_FILES/DATABASES_MPI/proc??????_TeleEle_rTop.info
OUTPUT_FILES/DATABASES_MPI/proc??????_TeleEle_xHigh.info
OUTPUT_FILES/DATABASES_MPI/proc??????_TeleEle_xLow.info
OUTPUT_FILES/DATABASES_MPI/proc??????_TeleEle_yHigh.info
OUTPUT_FILES/DATABASES_MPI/proc??????_TeleEle_yLow.info
```

Among above files, the first one is from the original code and other files contain the information of coupling elements.

3.2 Generating databases

This step of running xgenerate_databases is almost the same as the original package, except the process of assigning seismic structure. In the current version of our code, a 1D velocity model provided in the file dsm_model_input is read and assigned to each GLL points. In the next step of running DSM, you need to use this file as the input of DSM computation. For more complex velocity structure, you need to modify the code, basically the subroutine model_tomography in the file model_tomography.f90.

After running xgenerate_databases, the depths of all the coupling GLL points are collected and saved in the files

```
{\tt OUTPUT\_FILES/DATABASES\_MPI/depth\_table\_acoustic\_innerbound} \\ {\tt and} \\
```

OUTPUT_FILES/DATABASES_MPI/depth_table_elastic_innerbound. The depths of in the former file (elastic coupling GLL points) will be used in the DSM computation. The depths in the later file might be useful in mid-way coupling, but not in the source-side coupling.

The parameters depth_id, media type, normal vector, Jacobian and Cartesian coordinates associated each coupling GLL point are saved in the file

```
OUTPUT_FILES/DATABASES_MPIiproc????/disp_pack??????
```

These parameters are used in the final step of coupling.

3.3 Running the solver

This step is the same as the original package. The manual of SPECFEM3D_Cartesian explains the input parameters in the file DATA/Par_file in detail, but you should be careful to set up NSTEP. A complete 3D SEM wavefield is critical for accurate source-side coupling [see Wu et al., 2018], so you need to use a sufficiently large NSTEP to ensure that all waves are well absorbed. Once the solver starts, the displacement and strain on the coupling GLL points are continuously saved at every NSTEP_BETWEEN_OUTPUTBOUND time steps. The displacement and strain are saved in the files

```
OUTPUT_FILES/DATABASES_MPIiproc????/disp_pack?????
and
OUTPUT_FILES/DATABASES_MPIiproc????/traction_pack?????
```

DSM computation

After running SEM, the next step is to compute the Green's functions using DSM.

4.1 Main parameter file

The main input parameters are specified a file (e.g. dsm_model_input). Most of these parameters are the same as the original DSM code, except the ones listed at the end of the file. For example, the file dsm_model_input used in the example 1D_IASP91 is as follow:

4096.0 512 #time series length, number of frequency							
1.1d-3 #imaginary omega for artificial attenuation							
60000 0 28000 #number of grids on the radius direction, minimum angular order,							
maximum angular order							
11 #number	of structure zone						
0.0 1217.1 13	.0885 0.0000 -8	8.8381 0.0000	#zone1 inner core: r_min, r_max, coef1_density, coef2_density, coef3_density, coef4_density				
11.2409 0.0000	-4.0969 0.0000		#coef1_Vpv, coef2_Vpv, coef3_Vpv, coef4_Vpv				
11.2409 0.0000	-4.0969 0.0000		#coef1_Vph, coef2_Vph, coef3_Vph, coef4_Vph				
3.56454 0.0000	-3.4524 0.0000		#coef1_Vsv, coef2_Vsv, coef3_Vsv, coef4_Vsv				
3.56454 0.0000	-3.4524 0.0000		#coef1_Vsh, coef2_Vsh, coef3_Vsh, coef4_Vsh				
1.0000 0.0000	0.0000 0.0000	84.6 1327.7	#coef1_eta, coef2_eta, coef3_eta, coef4_eta,				
			Qmu, Qkappa				
1217.1 3482.0	12.5815 -1.2638	-3.6426 -5.528	#zone2 outer core: r_min, r_max, coef1_density, coef2_density, coef3_density, coef4_density				
10.0390 3.7567	-13.6705 0.0000	1	#coef1_Vpv, coef2_Vpv, coef3_Vpv, coef4_Vpv				
10.0390 3.7567	-13.6705 0.0000	1	#coef1_Vph, coef2_Vph, coef3_Vph, coef4_Vph				
0.0000 0.0000	0.0000 0.0000		#coef1_Vsv, coef2_Vsv, coef3_Vsv, coef4_Vsv				
0.0000 0.0000	0.0000 0.0000		#coef1_Vsh, coef2_Vsh, coef3_Vsh, coef4_Vsh				
1.0000 0.0000	0.0000 0.0000	-1.0 57823.0	#coef1_eta, coef2_eta, coef3_eta, coef4_eta, Qmu, Qkappa				
•••							
0.01 0.00 90.00 2 #source depth, source latitude, source longitude, source type (1 for moment tensor and 2 for single force)*							
20 1.0 0.0	0.0 0.0 0.0		noment tensor (six components) or force (only				
the first three components f_r , f_θ and f_ϕ are functional)*							
depth_solid_list # output of SEM*							
dist_solid_list # output of SEM*							
	-						

```
depth_fluid_list # output of SEM*
dist_fluid_list # output of SEM*
1 #save velocity (1) or displacement seismograms (2)*
end
```

The parameters marked with "*" are new variables in our SEM-DSM method.

DSM code uses the cubic polynomials to describe the 1D model, the same as PREM. For example, the Vp of the inner core in the above IASP91 model is $V_p=11.2409+0.0*a-4.0969*a^2+0.0*a^3$ km/s, where $a=r/R_{Earth}=r/6371.0$.

For source-side coupling,

- 1. The source must be a single force (source type=1).
- 2. The unit of force is dyne. For example, the line "20 1.0 0.0 0.0 0.0 0.0 0.0 0.0" in the above file dsm_model_input represents a vertical single force $f_r = 1.0 * 10^{20} dyn = 1.0 * 10^{15} N$.
 - Note that, Green's function corresponds an unit force of $1 N = 10^5 dyn$. However, we usually use a proper number of nexp (i.e. 20) to keep high numerical precision. Otherwise, too small variables in the code would give rise large numerical errors. Consequently, the final hybrid synthetics needs to be corrected back (i.e. multiply them with 10^{-15}).
- 3. The files "depth_solid_list" and "depth_fluid_list" are automatically generated when running SEM. Just copy them to under the DSM work folder.
- 4. the files "dist_solid_list" and "dist_fluid_list" contains the distances, where we want to compute Green's functions. Each target depth has a evenly spaced table.
- 5. save Green's functions of velocity seismograms.

4.1.1 The parameters you may need to change for source-side coupling

- 1. time series length: We have tested many cases and found a time length of 4096s works well for the direct P, S saves and other body wave phases (e.g. PcP, ScP, PKP, ScS, SKKS etc). If you really want to change it, do not use a too small number (i.e. <1024 s), because the artificial and real attenuation needs sufficiently long time to attenuate the late arrival seismic waves. Otherwise, the wrap-around effects of these late waves would give rise to large errors. A longer time length allows attenuation to attenuate late waves better, but it increases the computation cost.
- 2. number of frequency: This parameter together with time series length determines the maximum frequency of DSM synthetics. For example, a combination of number of frequency=512 and time series length=4096.0 means that the maximum frequency of DSM synthetics is 0.125 Hz. Personally, I prefer to use a power of two (i.e. 256, 512, 1024, 2048, and so on) for this parameter.
- 3. 1D Earth model: If you want to use other 1D model, you need to convert it to DSM format.
- 4. f_r , f_θ and f_ϕ : If only vertical components of hybrid synthetics are of concern, you only need to run the case of $f_r = 1.0$, $f_\theta = 0.0$ and $f_\phi = 0.0$. Similarly, radial and transpose components correspond to $f_\theta = 1.0$ and $f_\phi = 1.0$ respectively. If all the three components are of concern, you have to run each case individually and save their Green's functions in separate folders (refer to the xxxxx example).
- 5. dist_solid_list: refer to subsection 4.3

4.2 The depth tables depth_solid_list and depth_fluid_list

These two depth tables contain the depths of the coupling grids of SEM elements. They are automatically generated when you run SEM, so you can just simply copy them from the SEM output folders into DSM work folder. This depth table file is like:

```
2.000000E-02
                     35
                               #depth_shift_add_nodes, shift this small depth to add new radial nodes, number of
depths
   4.999999E-03 # depth_separation_tolerance.
   5 #number of GLL points on a coupling face of SEM element
   0.00000000000000E+000
                                11
                                          #depth0, izone_this_depth
   1.72673164646048
                         11
                                   #depth1, izone_this_depth
                                   #depth2, izone_this_depth
   5.000000000000000
                         11
                                   #depth3, izone this depth
   8.27326835353952
                         11
                                   #depth4, izone this depth
   10.00000000000000
                         11
   70.00000000000000
                        9
                                 #depth33, izone this depth
   73.2732683535395
                        9
                                 #depth34, izone_this_depth
                        9
   74.999999999991
                                 #depth35, izone_this_depth
```

For most cases, you can simply copy them from the SEM output and do not need to change the above parameters. The only two parameters you can change are depth_shift_add_nodes and depth_separation_tolerance. In order to accurately compute stress, we add one new radial nodes above and another one below the target depth, that give us more accurate strain on the radial direction. The small parameter depth_shift_add_nodes is the depth shift of the two new nodes relative to the target depth. In addition to that, we also need to add a new node for the target depth itself. However, if there is already an preexisting node sufficiently close to the node awaiting to be created, it is not necessary to add it. The criteria is depth_separation_tolerance, so depth_separation_tolerance must be larger than depth_shift_add_nodes. Again, the values of these two parameters in the above table have been confirmed to work well for many test cases, so you may just use them.

Similarly, the table depth_fluid_list corresponds to fluid media. Basically, this table is designed for mid-way coupling across core mantle boundary or inner core boundary. Thus, they are not used in source-side or receiver-side coupling. In the future, it could be used in a source-side or receiver-side coupling, in which ocean water is fully incorporated in both local and global simulation (e.g. hybrid method combing SPECFEM3D_GLOBE and SPECFEM3D_Cartesian).

4.3 The distance table dist solid list and dist fluid list

These two files contain evenly spaced distance tables, where Green's function are computed and stored. The file dist_solid_list is like:

```
100 #number of distances for depth0
25 #distance0
25.1 #distance1
25.2 #distanc2
...
34.8 #distacne99
34.9 #distance100
100 #number of distances for depth1
25 #distance0
```

25.1 #distance1
25.2 #distance2
...
34.8 #distacne99
34.9 #distance100
...
100 #number of distances for depth35
25 #distance0
25.1 #distance1
25.2 #distance2
...
34.8 #distacne99

34.9 #distance100

In this table, each depth has the same distance table. You may notice that the information in the above table is highly redundant and they can be easily simplified to just four parameters: number of depths, number of distances, minimum distance and distance space. However, we still use this redundant table, because a future version of SEM-DSM hybrid method may use unevenly spaced distance table.

Similarly, dist_fluid_list contains the distance table corresponding to fluid media (the Earth's outer core). For source-side coupling, the current version of code does not support coupling SEM elements in the ocean, so the table is not used.

4.3.1 How to generate a distance table?

The key to make the distance table is using a proper distance range and distance space. The distance range should be sufficiently large to covers all the distances used in the final step of coupling. For example, supposing that the SEM box has dimensions of $5^o \times 5^o$ and its horizontal center is 20^o away from the target teleseismic receiver, a distance table spanning a range of $16^o - 24^o$ must cover all the distances used in coupling. Of course, a very large distance range is always safe, but it takes more computational costs.

We use interpolation to obtain the Green's function at target distances, so a small distance space is important for accurate interpolation. However, a very small distance space may causes a large dimension of distance table and therefore significantly increases computational costs. Basically, a higher maximum frequency of concern requires a smaller distance space. More details about this issue are discussed in [Wu et al., 2018].

4.4 Output of DSM computation

If you successfully run DSM, you will see a file "Green_Par_forConvolution" and other subdirectories in the directory "OUTPUT_GREENS/". For source-side coupling, the relevant subdirectories are only "velo_solid/" and "sigma/", where Green's functions of velocity and stress in solid media are stored.

Running the Coupling

Now that you have successfully obtained the boundary information (displacement and traction) from SEM simulations and Green's function from DSM computation, you are ready to couple them and compute hybrid teleseismic synthetics. We provide two different interpolation schemes for the coupling. You can choose either linear interpolation or Lagrange interpolation scheme for the Green's functions. The Lagrange polynomial interpolation is computationally slower, but has higher accuracy than linear interpolation. These two schemes use the same input parameters, so you can keep the same input setups, run both of them and compare their difference. If significant differences between them are seen, you should be ware that large errors must be present in the results of linear interpolation and possibly also in Lagrange interpolation.

The following subsections explain the input files for a source-side coupling. Only the main parameter file DATA/Par_file needs to be manually edited. All other input files described in subsections 5.2 and 5.3 are automatically produced by SEM and DSM, you can simply copy them to the right directories.

5.1 Main parameter file

The main parameter file DATA/Par_file is as follow

```
npoints_taper_SEM = 100
vertical_component= .true.
radial_component= .false.
transverse_component= .false.
```

npoints_taper_SEM gives the width of the taper applied on the SEM displacement and traction, in an unit of deltat_SEM. Note that the deltat_SEM here is the time step after decimating (see 5.2), rather than the original time step in the SEM solver. Ideally, the displacement and traction of SEM coupling grids should be very close to zero at the very late time. However, real cases are not always ideal. For example, some SEM coupling grids might be close to the source and therefore have large static displacement; Absorbing boundaries do not absorb waves well, due to too short time length (see 3.3), too thin PML layers (see 3.1) or other reasons. In this situation, tapering the ends of traction and displacement are useful.

vertical_component Do coupling for vertical component if set it to .true.. If .true., make sure you have computed the Green's functions associated with vertical single force f_r .

radial_component Do coupling for radial component if set it to .true.. If .true., make sure you have computed the Green's functions associated with vertical single force f_{θ} and f_{ϕ} .

transverse_component Do coupling for transverse component if set it to .true.. If .true., make sure you have computed the Green's functions associated with vertical single force f_{θ} and f_{ϕ} .

If only vertical component are of concern (e.g. direct P-waves), you do not need to compute the Green's functions associated with f_{θ} and f_{ϕ} , because vertical component is independent on them [see the eq. 9 in Wu et al., 2018]. Similarly, the two horizontal components, radial and transverse, are independent on Green's functions associated with f_r . However, you have to compute Green's functions of both f_{θ} and f_{ϕ} , even if you are only interested in one of the two horizontal components.

5.2 input files from SEM

The input information from SEM simulation includes package_list, SEM_Par_Coupling and other variables (i.e. geometry information, media type, traction and displacement) on coupling grids. These files are automatically generated when you run SEM, so you can simply copy them from SEM outputs into the directory SEM_input/.

5.2.1 package_list

```
The structure of package_list is as follow
```

```
# ipackage_global, iproc_SEM, ipackage_this_proc, npoints_per_pack_SEM
         1
             300
2
         2
    0
             300
3
    0
         3
             300
    0
         4
             300
          27
                300
27
     0
28
     0
          28
               300
29
     1
          1
              300
30
     1
          2
              300
      3
           23
                 300
105
106
       3
           24
                 300
107
       3
           25
                 225
```

5.2.2 SEM_Par_Coupling

The file SEM_Par_Coupling includes the below parameters

npackage_SEM is the number of lines of the file SEM_input/package_list.

npoints_per_pack_SEM corresponds to NPOINTS_PER_PACK in the SEM input file SEMtoTele_Par_file
 (see section 3.1).

nstep_each_section_SEM is NSTEP_BETWEEN_OUTPUTBOUND in the **SEM** input file SEMtoTele_Par_file (see section 3.1).

total_nstep_SEM is the number of time steps of SEM displacement and traction. It equals nstep_SEM_original/DECIMATE_COUNTED_COUNTED_SEM_original is the NSTEP in the SEM main parameter file Par_file and DECIMATE_COUPLING can be found in SEMtoTele_Par_file (see section 3.1)

nsection_SEM is the number of sections in each package of SEM displacement and traction. It is equal to right-round(total_nstep_SEM/nstep_each_section_SEM), where 'right-round' means the nearest integer larger than total_nstep_SEM/nstep_each_section_SEM.

deltat_SEM equals delta_SEM_original × DECIMATE_COUPLING. delta_SEM_original is the DT in the SEM main parameter file Par_file and DECIMATE_COUPLING can be found in SEMtoTele_Par_file (see section 3.1).

5.2.3 Variables associated with each coupling grid

The geometry information (i.e normal vector, Jacobian, X-Y-Z coordinates), media type, traction and displacement on coupling grids from each CPU processor are stored in the corresponding directories SEM_input/iproc?????.

5.3 input files from DSM

Input information from DSM computation includes the Green's functions and parameter file Green_Par. They are stored in the directory DSM_input.

The structure of file DSM_input/Green_par is as below 512 #number of frequency

```
0.0 0.1 0 #min_dep_acoustic, ddep_acoustic, ndepth_acoustic
0.0 0.1 35 #min_dep_solid, ddep_solid, ndepth_solid
25.0 0.1 100 #min_distance, distance_space, number of distances
4096.0 #time length
1.1E-003 #artificial attenuation coefficient - imaginary_omega In the above parameters, min_dep_acoustic,
```

ddep acoustic, min dep solid and ddep solid are not functional.

The Green's function corresponding to f_r , f_θ , f_ϕ are stored in the directories DATA/zcomp, DATA/rcomp and DATA/tcomp respectively.

5.4 Output hybrid synthetics

When the coupling is completed, hybrid synthetics are stored in the directory seismograms. Note that

- (1) The seismograms are displacement and the unit is 10^{-15} meters (see subsection 4.1). Thus, you need to multiply them with 10^{-15} to get an unit of meter.
- (2) There might be large errors at very low frequencies (i.e. <0.01 Hz), due to errors from series of sources, including SEM computation, local wavefield approximation and tapering the SEM synthetics.
- (3) The maximum resolved frequency of hybrid synthetic seismograms is the lower one between that of SEM and DSM. Usually, applying a band-pass filter to synthetics is necessary when comparing synthetic seismograms to data.
- (4) If ocean is present and truncated at the boundaries of SEM box, you must be ware that only the direct P-waves are valid. Do not use the waves after the direct P-waves, unless you have carefully checked them. Because the ocean is truncated at the boundaries of SEM box, large errors might be present in the late waves (see the example in subsection 6.4).

Examples

For a source-side coupling, you need to run the three steps of SEM, DSM and coupling one by one, because the following step is dependent on the previous one. We provide four examples to show how to run a SEM-DSM source-side coupling. In each example, you can see an README file, which contains some useful information and descriptions of how to run that example. Among the three steps, creating a good SEM model needs user-specified input parameters, so we mostly focus on how to set up SEM model in the below subsections.

6.1 1D synthetic (vertical component) benchmark using the IASP91 model

In this example, we simply run a 1D model without any 3D heterogeneities. The 1D model is IASP91 velocity model and PREM attenuation model. In the SEM part, part of the parameters (see more detailed descriptions in subsection 3.1) are

Mesh_Par_file:

```
= 105.d0
DEPTH_BLOCK_KM
                                = 30
NEX_XI
                                = 30
NEX_ETA
                                = 0.24
THICKNESS_OF_X_PML
THICKNESS_OF_Y_PML
                                = 0.24
THICKNESS_OF_Z_PML
                                = 25000.0
NREGIONS
                                = 1
# define the different regions of the model as :
#NEX_XI_BEGIN #NEX_XI_END #NEX_ETA_BEGIN #NEX_ETA_END #NZ_BEGIN #NZ_END #material_id
  30 1 30 1 11 2
```

SEMtoTele_Par_file:

```
ANGULAR_WIDTH_XI_IN_DEGREES = 3.0d0
ANGULAR_WIDTH_ETA_IN_DEGREES = 3.0d0
...
nx_NoTopography=5
ny_NoTopography=5
ix_LowBound=4
ix_HighBound=26
iy_LowBound=4
iy_HighBound=26
```

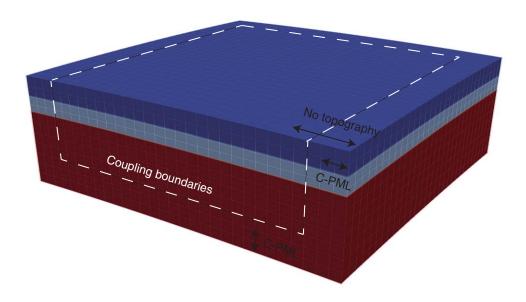


Figure 6.1: Screenshot of mesh of the 1D synthetic benchmark. The white dashed lines indicate the coupling interfaces, where seismic displacement and traction are stored and used in the final step of SEM-DSM coupling. The black arrow on the top interface shows the width of no topography region on the right boundary of SEM box, which equals five elements length. The other two arrows show the C-PML absorbing boundaries with a width of two elements on the right and bottom boundaries of the SEM box.

```
ir_BoundTop=11
ir_BoundBot=4
```

Based on the above parameters, the internal mesh tool xmeshfem3D produces meshes shown in Fig. 6.1. The dimensions of one element are $9.55~km~(105~km/11, {\rm vertical}) \times 0.1^o~(3.0^o/30, {\rm horizontal}~\xi) \times 0.1^o~(3.0^o/30, {\rm horizontal}~\eta)$. Thus, <code>THICKNESS_OF_X_PML=0.24</code> produces C-PML absorbing boundaries with the same width of two elements at the sides of $\xi = min_\xi$ and $\xi = min_\xi$. The widths of C-PML layers at the bottom, $\eta = min_\eta$ and $\eta = max~\eta$ are also two elements (Fig. 6.1).

Although there is no topography in this 1D SEM box, we still need to specify the parameters nx_NoTopography and ny_NoTopography and ny_NoTopography to 5. It means that the most outside five elements must be flat (Fig. 6.1). As we mentioned in subsection 3.1, the coupling interface must be inside the no topography region and outside the C-PML band (Fig. 6.1). If this condition is not satisfied, error report are raised.

Please read the three README files in the corresponding SEM, DSM and coupling directories, which help you how to run this example. Basically, you just follow the below steps to run the executable bash command files.

Note that only vertical components are valid, because we only compute the Green's functions of vertical single force to save computational cost.

6.2 1D synthetic (three components) benchmark using the IASP91 model

In the above example of 1D synthetic benchmark, we only compute the vertical components. In this case, the radial and transverse components are also computed. To accomplish that, you need to run DSM for all the three single forces f_r , f_θ and f_ϕ .

How to run this example?

(5) check the hybrid synthetics in the directory SEM_DSM_hybrid/coupling/example/iasp91_1D_benchmark_three_components/seismograms.

6.3 A simple sinusoidal shape topography model (no attenuation)

In this example, we duplicate the case shown in Fig.5 of Wu et al. [2018], where a sinusoidal shape topography is introduced in the SEM box. Because we use cubed-sphere coordinates in SEM simulation, topography as a function of latitude and longitude needs to be converted to that in cubed-sphere coordinates (ξ , η). The below line in run_SEM.sh does coordinate conversion and save the output results in DATA/meshfem3D_files/real_bathymetry_topography.

```
../../bin/xcubedsphere_topo_FindOcean ./DATA/meshfem3D_files/latlon_surf_topo.dat 320 320 -0.1 -0.1 0.01d0 0.01d0
```

The first parameter ./DATA/meshfem3D_files/topo_surf.dat is the file name of topography model in the longitude-latitude coordinates.

The second and third parameters (320 and 320) are the numbers of points along ξ and η respectively.

The fourth and fifth parameters (-0.1 and -0.1) are minimum ξ and η . Note that, these two parameters must be smaller than 0.0 to fully cover the simulated region.

The last two parameters (0.01d0 and 0.01d0) are spacing between points.

The SEM model of this example contains three interfaces (from bottom to top: Moho, upper crust and free surface discontinuities). We assume no topography on the upper crust and Moho interfaces and it is easy to produce these interfaces (see details in run_SEM.sh). The topography of free surface in the is formed by the bellow command.

```
../../bin/xcubedsphere_topo 3 ./DATA/meshfem3D_files/latlon_surf_topo.dat 0.0 -20000.0 30000.0
```

The first argument "3" indicates that we are working on the third interface listed in the file INTERFACES_FILE (see subsection cha:Running-the-Mesher-cubed-sphere). The other two interfaces are mid-crust and Moho discontinuity.

The second parameter ./DATA/meshfem3D_files/topo_surf.dat is the file name of topography model in the longitude-latitude coordinates.

The third parameter 0.0 is the basement value of the free surface topography. The topography on the boundaries of SEM box are tapered to the basement value (see more interpretations in the next example in subsection 6.4).

The last two parameters -20000.0 and 30000.0 are the minimum and maximum allowed topography. Any topography out of this range will be truncated. Here, the peak topography is 10000 meters, so max_allowed_topography=30000.0 does not truncate any topography.

Executing the above command produces the file DATA/meshfem3D_files/topo_surf.dat. You can see the above commands in the file run_SEM.sh in the corresponding SEM directory.

xmeshfem3D reads topo_surf.dat to create the free surface. Then the file real_bathymetry_topography is read to classy the SEM elements to either acoustic ocean or solid media. Any element with a central GLL point above the bathymetry given in real_bathymetry_topography is classified to acoustic media.

Note that the dimensions of region described in real_bathymetry_topography and all the interfaces listed in INTERFACES_FILE should be sufficiently large to cover the SEM region. Otherwise, the codes cannot properly build up the model you want. For example, the ranges of dimensions in the above command are -0.1 deg to 3.1 deg along ξ and η , which fully cover the SEM box (0 deg to 3.0 deg). There is no ocean in this example, so the geometries described in real_bathymetry_topography and topo_surf.dat are the same. In the example of next section 6.4, we will discuss how to use the real topography/bathymetry data to incorporate the ocean water.

How to run this example?

(5) check the hybrid synthetics in the directory SEM_DSM_hybrid/coupling/example/iasp91_sinusoidal_topography_no_attenuation/seismograms.

6.4 Simulating the P-waves from the 2016/10/27 Mw6.0 South America earthquake

This example duplicates the synthetics in Fig. 11b in Wu et al. [2018]. We use the 30 arc-second resolution to-pography/bathymetry data to incorporate the ocean and land topography. First, the topography/bathymetry data in geographical coordinates needs to be converted that in the cubed-sphere coordinates.

```
../../bin/xcubedsphere_topo_FindOcean ./DATA/meshfem3D_files/SA_real_topo.dat 540\ 540\ -0.2\ -0.2\ 0.01\ 0.01
```

The above command is executed when you run the script run_SEM. sh in the corresponding SEM directory. The converted topography/bathymetry is stored in real_bathymetry_topography (Fig. 6.2).

Then we need to prepare the topography files of the four interfaces (Moho, upper crust, ocean bottom and free surface). Comparing to the example of sinusoidal topography in subsection 6.3, another interface of ocean bottom is needed to incorporate the ocean. The ocean bottom interface is formed using the below command.

```
../../bin/xcubedsphere_topo 3 ./DATA/meshfem3D_files/SA_real_topo.dat -3878.0 -20000.0 -1500.0
```

We have explained what each parameter does in subsection 6.3. The minimum allowed bathymetry of -20000.0 meters is deeper than any ocean bottom in the world, that means no deep bathymetry is truncated. The output results are stored in the file topo_botoC.dat (Fig. 6.3a). Similarly, the topography of the free surface is formed by the below command and stored in topo_surf.dat (Fig. 6.3b).

```
../../bin/xcubedsphere_topo 4 ./DATA/meshfem3D_files/SA_real_topo.dat 0.0 0.0 30000.0
```

Based on these two interface, xmeshfem3d creates and meshes the top layer, where the ocean is incorporated (Fig. 6.4). This layer is meshed using three elements along vertical direction and the element size are proportional to the layer's thickness. Thus, using proper input parameters to make ocean bottom and free surface is important to generate high quality elements. For example, we use a maximum allowed bathymetry of -1500 m to produce the ocean bottom. This value allow us to keep most deep bathymetry features exactly honored (Fig. 6.4) and generate elements with good qualities. In some extreme cases (e.g. rapidly varying topography/bathymetry), properly truncating and/or smoothing the topography/bathymetry is necessary to fix highly skewed elements.

Running this example needs moderate computational costs (e.g. 10-20 hours using hundreds of CPU processors),

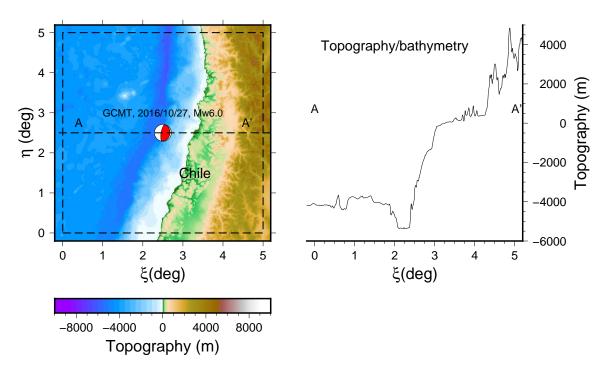


Figure 6.2: Topography/bathymetry in the cubed-sphere coordinates (ξ , η). The left figure shows the topography/bathymetry described in real_bathymetry_topography. The beach ball represents the GCMT solution [Global Centroid Moment Tensor, Ekström et al., 2012] of this earthquake. The dashed line box shows the dimension of the simulated SEM box (5 deg \times 5 deg). The topography/bathymetry on the profile A-A' is plotted in the right figure. Note the different scales on x- and y-axis (1 degree is equivalent to \sim 111.19 km).

that cannot be accomplished by a few CPUs. If you successfully run this example, the synthetic seismograms are stored in the directory SEM_DSM_hybrid/coupling/example/SouthAmerica_Earthquake_161027/seismograms. We note that only the direct P-waves in the first ~60 seconds (maybe a little longer, e.g. 120 seconds) has been checked. Because the ocean is truncated at the boundaries of SEM box, large errors might be present in the very late waves [see Wu et al., 2018, , especially Fig. S9].

How to run this example?

- (1) Read section 2 and compile the software packages.
- (2) run SEM
- >cd SEM_DSM_hybrid/specfem3d/EXAMPLES_COUPLING/SouthAmerica_Earthquake_161027
- >./run_SEM.sh
- (3) run DSM
- >cd SEM_DSM_hybrid/DSM/example/SouthAmerica_Earthquake_161027
- >./run_DSM.sh
- (4) run the coupling step
- >cd SEM_DSM_hybrid/coupling/example/SouthAmerica_Earthquake_161027
- >./run_coupling.sh
- (5) check the hybrid synthetics in the directory
- SEM_DSM_hybrid/coupling/example/SouthAmerica_Earthquake_161027/seismograms.

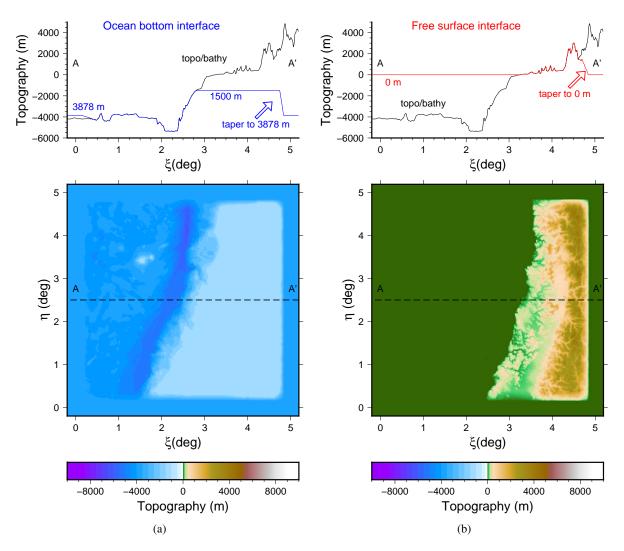


Figure 6.3: The topography of the free surface and ocean bottom interfaces in the SEM box. (a) The ocean bottom interface described in topo_botOC.dat. This interface extends to -1500 m below the land and tapered to 3480 m near the boundaries. The upper panel shows the bathymetry on the A-A' profile. (b) The free surface topography described in topo_surf.dat. The topography is fixed at 0 m in the ocean. The upper panel shows the topography on the A-A' profile.

6.5 Run your own example

In the above examples, we show how to run modes of 1D structure, 3D topography and 3D ocean. You can create your own example using even more complex model. If you want to model effects of topography/bathymetry and ocean, the 30 arc-second resolution global topography/bathymetry model can be freely downloaded from General Bathymetric Chart of the Oceans (GEBCO, https://www.gebco.net/); To incorporate 3D velocity model in SEM (e.g. sediment, slab or tomography 3D model), you may need to write some new routines or modify the subroutine model_tomography in the Fortran file SEM_DSM_hybrid/specfem3d/src/generate_databases/model_tomography.f90.

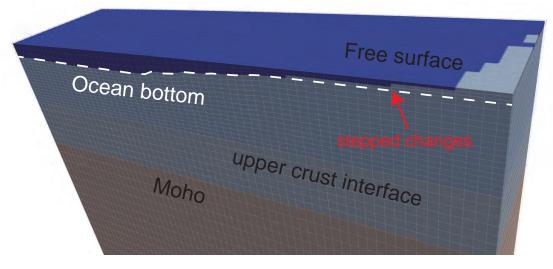


Figure 6.4: Part of meshes. The dark blue region represents the ocean. The white dashed line shows the ocean bottom interface. The deep bathymetry (i.e. <-1500 m) is exactly honoured and the shallower bathymetry is approximated by stepped grids.

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