

1. Install calculix, gfortran (for UMAT) and python 3 with numpy and scipy module;
2. Generate microstructure using matlab
 - Open generate_synthetic_microstructures.m in the folder “Microstructure_Generation_Code”
 - Input specimen size, fiber radius, fiber volume fraction in generate_synthetic_microstructures.m ;
 - Run generate_synthetic_microstructures.m and it will automatically generate the folder “fibercentroid_rxfydomainz”
 - ❖ x,y,z respectively stand for the value of fiber radius, volume fraction and domain size; The output information like fiber centroid coordinates, amount of fibers are stored in fibercentroid_info.dat and fibercoord.dat within the folder;
3. Generate inp. file for calculix solver
 - Copy the folder “fibercentroid_rxfydomainz” to the folder “abaquscae_pythonscripting”;
 - Run casefiles_edit.py and generate composites_cae_forrun.py
 - Open abaqus cae module:
 - 1) For creating geometry, copy the scripts in composites_cae_forrun.py from the first line to the line “del modelObject.sketches['__profile__]” to Kernel Command Line Interface in abaqus cae module and run.
 - 2) Generate the mesh.
 - 3) For creating the sets of fiber centroid nodes, copy the scripts from the line “allNodes = mdb.models[model].parts[part].nodes” to the last line in composites_cae_forrun.py;
 - 4) Create the element sets for matrix, fiber; Create node sets for boundaries;
 - 5) The inp. File format should be modified for calculix usage; The format can be referenced based on the existing inp. file;

3. Run the reference simulation using calculix

- Create the folder `jobname_results` (“jobname” is the name of the case) and subdirectory “`jobname_results/ref`” and “`jobname_results/opt`”.
 - ❖ “`jobname_results/ref`” will store the measurement data, which in this case is generated by numerical simulation.
 - “`jobname_results/opt`” stores the information for optimization;
- Open `fem2Dmechload_expmod.py` and input the parameters;
 - Input `jobname`, `solver_name`, `solver_directory1`, `parent_dir`.
 - ❖ The whole solver directory for executing calculix program is “`solver_directory1 + solver_name + solver_directory2`”. `solver_directory1` is the home directory of calculix solver. `solver_directory2` is fixed. Since calculix solver cannot deal with multiple UMAT tasks simultaneously, different copies of calculix solver (contained in the folder named “`solver_name`”) are made in the folder `solver_directory1`.
 - ❖ `parent_dir` is the home directory where reference simulation and optimization locate;
 - Input `measind1`, `measind2` (measurement number from `measind1` to `measind2`), `errorbnd` (measurement error bound), `Em_bar`, `Em_inter`, `alpha`, `num`(reference material parameters) and `fiber_centroid_number` (number of fiber centroid , extracted from number of nodes in the nodes set `fibercentroid` in `inp.` file)
- Copy the folder “source” (containing `umat` file), `compile.sh` (employed for compiling UMAT for calculix), `initial.sh` (for submitting job to the cluster), `fibercoord.dat` and `inp.` file in the folder “`fibercentroid_rxvfydomainz`” to the home directory;
- Run `fem2Dmechload_expmod.py`
 - ❖ Reference reaction force files (`reactionforce_ex_my.dat`), fiber centroid displacement files (`fiber_centroid_disp_ex_my.dat`), `error_reactionforce_ex_my.dat`, `error_ux_ex_my.dat`, `error_uy_ex_my.dat` stand for exact value of error added to the reference reaction force, `ux` and `uy`;
 - ❖ `x` stands for the error bound (`x=1` for 1% errorbound), `y` stands for sequence number of measurement data (`y=30` for measurement no. 37);

4. Execute optimization using scipy;

- Make multiple copies of calculix solver with the name “Calculix_*X*” in the directory solver_directory1, where *X* values from 1 to the total number of copies;
- Open initializationsetting_rferr.py and input parameters:
 - job_name, solver_directory1, parent_dir, errorbnd, measind1, measind2, fiber_centroid_number;
 - level (total levels of stratified sampling for initialization, One set of optimization is executed using multiple initialization, which is set using stratified sampling);
 - num_cpu (total number of cpus employed for the same set of optimizations with multiple initializations);
- Run initializationsetting_rferr.py and create all the files for optimization settings;
 - The optimization setting files, including the bash files (initial.sh) for submitting the jobs in the cluster are stored under the directory ex_cpuy, where *x* stands for error bound, *y* stands for sequence number of cpu;
- Run submit.sh to submit the jobs, the results will be stored in the directory “jobname_results/opt”;
 - The results file name format is objfunc_exEijaknuml.dat, where *x* stands for error bound, *j*, *k*, *l* stand for number of stratified sampling level;
- Running errordetect.py will detect whether there is error information (stored in the file ex_cpuy.e* under the directory ex_cpuy) occurring during the optimization;
- Run plot_optdata.m will plot the final results;

- Note:
 - collectoptinfo.sh can copy all the cluster job information files `ex_cpuy.e*` and `ex_cpuy.o*` to one place;
 - jobdeletion.sh can delete all of the cluster jobs;
 - Running errordetect.py will show the optimization which exceeds the maximum iteration step. Those optimizations need to be run again. Running initializationsetting_supplementary.py will create the initialization for those optimizations.
 - initializationsetting.py and optmechload_expmod.py are for the optimizations in which the reaction force has no measurement error;
 - The final optimization results are stored in the directory “optresults_sample`x`_microns`y`_vfz_`xxx`”; The numerical reference measurement data are stored in microns`y`vfz_results, where `x` stands for total amount of samples, `y` stands for specimen size, `z` stands for fiber volume fraction;
 - initializationsetting*.py edit the information of line 107 to 120 in optmechload_expmod*.py, line 15 in compile.sh , line 56 in source/umat.f90 and initial.sh. Please be careful about the corresponding modifications of line number if there is any modifications in those files