- 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\_rxvfydomainz"
    - \* 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\_rxvfydomainz" to the folder "abaquscae\_pythonscripting";
  - Run casefiles\_edit.py and generate composites\_cae\_forrun.py
  - Open abagus 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
  - > Select a home directory for optimization. Create the folder jobname\_results ("jobname" is the name of the case) and subdirectory "jobname\_results/ref" and "jobname\_results/opt" in the home directory.
    - "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:

- $\triangleright$  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 needs 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\_samplex\_micronsy\_vfz\_xxx"; The numerical reference measurement data are stored in micronsyvfz\_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