IM Calculation Refactor

Please read the readme @ https://github.com/ucgmsim/IM_calculation/blob/master/README.md for instructions on how to run the code.

DONE

- IM calculations have been separated from the "old post-processing" repository, extracting the relevant functions and classes.
- IM values validated on Hypocentre and Kupe against "old post-processing" on the same data.
- Two types of workflows: text based (most likely observations) and binary based (simulations once the binary workflow is in place).
 - If binary workflow does not happen soon, the text based option will be used in both cases.
- Outputs match the formats requested in File Formats Used On GM and should therefore be usable on the upcoming Non-ergodic codes.
- Tested on very simple multi-process on Kupe with good speed-up using 40 and 80 cores. For the sample, 2228 stations were used

Machine	Cores	Time
Hypocentre	1	132m
Hypocentre	8	8.7m
Kupe	40	27m
Kupe	80	

OUTPUT STRUCTURE

With command: python calculate_ims.py ../BB.bin b -o /home/yzh231/ -i Albury_666_999 -r Albury -t s -v 18p3 -n 112A -m PGV pSA -p $0.02\ 0.03$ -e -c geom -np 2

- input file path: ../BB.bin
- b: input file type is binary
- -o: output result csvs location is /home/yzh231, default is /home/\$user
- -i: unique identifier/runname of the simrun and output folder name are Albury_666_999, default is 'all_station_ims'. This attribute will be stored in the meta data file.
- -r: rupture name is Albury, default is unknown. This attribute will be stored in the meta data file.
- -t: type of simrun is simulated, default is unknown. This attribute will be stored in the meta data file.
- -v: version of simrun is v18p3, default is XXpY. This attribute will be stored in the meta data file.
- -n: station names used to perform im claculation are 112A, default is all the stations in the binary file
- -m: measures used to perform im calculation are PGV and pSA, default is all the measures
- -p: period of pSA used to perform im calculation are **0.02 0.03**, default is Karim's 15 periods
- -e: In addition to the period specified by -p option, use extended 100 period of pSA, default not using
- -c: component of waveform acceleration used to perform im calculation is geom, default is '090, 000, ver'
- -np: number of processors used to perform im calculation is 2, default is 2

The result is outputted to the following location, where:

- 'Albury_666_999' is the folder that contains all outputs. The folder name 'Albury_666_999' is made of the string specified by the '-i' argument. Default is 'all_station_ims' if not specified.
- 'Albury_666_999.csv' is the summary csv file that contains all stations' im calculations. The summary file name is made of the string specified by the '-i' argument.
- 'Albury_666_999_imcalc.info' is the meta data file. The meta data file name is made of the string specified by the '-i' argument.
- 'station' is the folder that contains all individual station's im_calculations. The folder name is defaulted and cannnot be specified by the user.
- '112A_geom.csv' is the individual csv file that contains geom component im calculation for station 112A. Each name of the individual station csv file name is made of station_name + component

```
yzh231@hypocentre ~ % tree Albury_666_999
Albury_666_999
— Albury_666_999.csv
— Albury_666_999_imcalc.info
— stations
— 112A_geom.csv
```

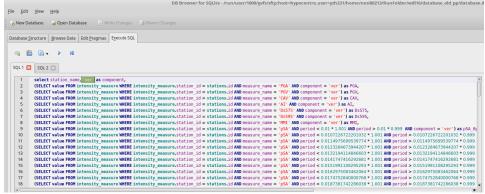
TEST FOR CALCUALTE_IMS.PY

All the steps below are to be carried out in hypocentre

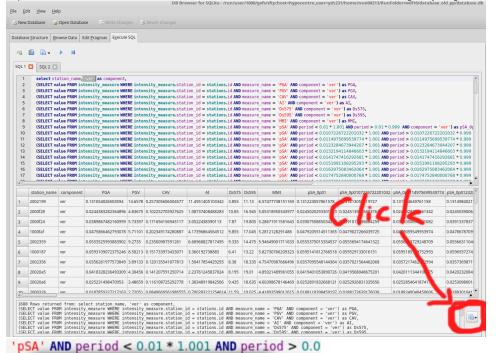
1.Generate summary benchmark:

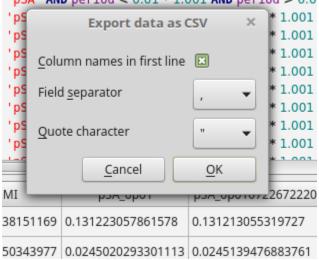
The following steps should only be performed once for each selected binary file

- Select a source binary file: /nesi/transit/nesi00213/RunFolder/daniel.lagrava/Kelly_VMSI_Kelly-h0p4_EMODv3p0p4_180531/BB/Cant1D_v2-midQ_leer_hfnp2mm+_rvf0p8_sd50_k0p045/Kelly_HYP01-03_S1244/Acc/BB_with_siteamp.bin
- 2. Identify corresponding databse for the selected source binary file: /home/nesi00213/RunFolder/wdl16/database_old_pp/database.db
- 3. Find the script to extract benchmark im value files from the database in step 2: /nesi/projects/nesi00213/dev/impp_datasets/extract_ims.sql
- 4. Create a folder to store benchmark files. eg benchmark_im_sims
- 5. Execute extract_ims.sql in database.db 4 times with specified components. eg: 'ver'



6. Export results to benchmark_im_sims/benchmark_im_sim_ver.csv. Clik OK and don't change anything when 'Export data as csv' window prompts





- 7. Repeat step 4 and 5 with different components: '090', '000', 'geom'
- 8. Now you have 4 summary benchmark files benchmark_im_sim_090/000/ver/geom.csv

2.Generate test input files

1. Follow the instruction in Binary Workflow FAQ, we can generate single waveform files. These waveforms are intended for the testing of ascii functionality of calculate_ims.py. Open a python cell

```
from qcore.timeseries import BBSeis
bb = BBSeis('/nesi/transit/nesi00213/RunFolder/daniel.lagrava/Kelly_VMSI_Kelly-h0p4_EMODv3p0p4_180531/BB
/Cant1D_v2-midQ_leer_hfnp2mm+_rvf0p8_sd50_k0p045/Kelly_HYP01-03_S1244/Acc/BB_with_siteamp.bin')
bb.all2txt(self, prefix='/home/$user/benchmark_im_sim_waveforms/', f='acc'):
```

Now we have all the waveforms. 000281†.ver 0002820.000 0002015.000 0002307.090 0002540.ver 0002bce.090 10002b7.ver 1000871.000 0002015.090 0002307.ver 0002541.000 0002820.090 0002bce.ver 10002ca.000 1000871.090 0002541.090 0002541.ver 0002bcf.000 0002bcf.090 0002015.ver 0002308.000 0002820.ver 10002ca.090 1000871.ver 0002821.000 10002ca.ver 0002016.000 0002308.090 1000882.000 00025e0.000 0002821.090 0002bcf.ver 10002cc.000 0002016.090 0002308.ver 1000882.090 0002016.ver 0002309.000 00025e0.090 0002821.ver 0002bd0.000 10002cc.090 1000882.ver 0002017.000 0002309.090 00025e0.ver 0002822.000 0002bd0.090 10002cc.ver 10008cc.000 0002017.090 0002bd0.ver 0002bd1.000 10002d9.000 10002d9.090 10008cc.090 0002309.ver 00025e1.000 0002822.090 0002017.ver 000230a.000 00025e1.090 0002822.ver 10008cc.ver 0002018.000 000230a.090 00025e1.ver 0002823.000 10002d9.ver 10008cf.000 0002bd1.090 0002018.090 000230a.ver 00025e2.000 0002823.090 0002bd1.ver 10002e3.000 10008cf.090 0002c82.000 0002c82.090 0002018.ver 00023ab.000 00025e2.090 0002823.ver 10002e3.090 10008cf.ver 00025e2.ver 0002824.000 10008d3.000 00023ab.090 10002e3.ver 00020c1.000 00020c1.090 00023ab.ver 00025e3.000 0002824.090 0002c82.ver 10002f2.000 10008d3.090 00020c1.ver 00023ac.000 00025e3.090 0002824.ver 0002c83.000 10002f2.090 10008d3.ver 00020c2.000 00020c2.090 10002f2.ver 10002ff.000 00023ac.090 00025e3.ver 0002825.000 0002c83.090 10008de.000 0002825.090 10008de.090 00025e4.000 0002c83.ver 00023ac.ver 00020c2.ver 00023ad.000 00025e4.090 0002825.ver 0002c84.000 10002ff.090 10008de.ver 00020c3.000 00023ad.090 00025e4.ver 0002826.000 0002c84.090 10002ff.ver 10008e4.000 0002826.090 0002826.ver 0002827.000 00020c3.090 00023ad.ver 00023ae.000 00025e5.000 00025e5.090 0002c84.ver 0002c85.000 1000302.000 1000302.090 10008e4.090 10008e4.ver 00020c3.ver 00020c4.000 00025e5.ver 10008e8.000 00023ae.090 0002c85.090 1000302.ver 00020c4.090 00023ae.ver 00025e6.000 0002827.090 0002c85.ver 100030e.000 10008e8.090 00020c4.ver 0002827.ver 10008e8.ver 00023af.000 00025e6.090 0002c86.000 100030e.090

3. Create Test Folder

- 1. Create The test folder structure follows Testing Standards for ucgmsim Git repositories
- 2. Select 10 stations you want to test and cp corresponding waveforms files to the singel_files directory as below
- 3. Copy the source binary file 'BB_with_siteamp.bin' to the input folder
- 4. Run 'write_benchmark_csv(sample_bench_path)' function inside test_calculate_ims.py to generate 'new_im_sim_benchmark.csv', where 'sample_bench_path' is the folder we created in 1.4 Generate summary_benchmark: benchmark_im_sims. This function should only be run once for each binary file.

NOW you have all the input files ready

```
test
       - README
       - sample1
            input
                 BB_with_siteamp.bin
                new_im_sim_benchmark.csv
              — single_files
                    - 00020d3.000
                    - 00020d3.090
                    - 00020d3.ver
                    - 2002199.000
                    2002199.090
                    2002199.ver
                    - CASH.000
                    CASH.090
                    - CASH.ver
                    - CFW.000
                    - CFW.090

    CFW.ver

                    - DLX.000
                    - DLX.090
                    - DLX.ver
                    - EWZ.000
                    - EWZ.090
                    - EWZ.ver
                    - GRY.000
                    - GRY.090
                    - GRY.ver
                     LSRC.000
                    - LSRC.090

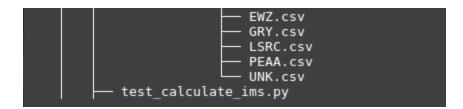
    LSRC.ver

                    - PEAA.000
                    - PEAA.090
                    - PEAA.ver
                    - UNK.000
                    - UNK.090
                 UNK.ver
            output
                 ascii darfield im sim
                   ascii_darfield_im_sim.csv
                    - ascii darfield im sim.info
                    - stations
                        - 00020d3.csv
                         2002199.csv
                         CASH.csv
                         CFW.csv
                        - DLX.csv
                        - EWZ.csv
                        - GRY.csv
                         LSRC.csv
                         PEAA.csv
                      — UNK.csv
               binary_darfield_im_sim
                   binary_darfield_im_sim.csv

    binary darfield im sim.info

    stations

                         - 00020d3.csv
                         2002199.csv
                         CASH.csv
                         CFW.csv
                         DLX.csv
```



4. Run Pytest

Make sure you are currently under the test_calculate_ims folder, run:

```
$ pytest -v -s test_calculate_ims.py
```

CHECKPOINTING & SPLITTING A BIG SLURM

Responsible scripts

- 1. slurn header template: https://github.com/ucgmsim/slurm_gm_workflow/blob/master/templates/slurm_header.cfg
- im_calc_slurm template: https://github.com/ucgmsim/slurm_gm_workflow/blob/master/templates/im_calc_sl.template
- 3. submit_hf.py that generates the slurm files: https://github.com/ucgmsim/slurm_gm_workflow/blob/master/scripts/submit_hf.py
- checkpointing functions: https://github.com/ucgmsim/slurm_gm_workflow/blob/master/scripts/checkpoint.py

Checkpointing

Checkpointing is needed for IM_calculation due to large job size and limited running time on Kupe. Therefore, we implemented checkpointing to track the current progress of an im_calculation job, and carry on from where the job was interrupted by slurm.

Note, the checkpointing code relies on the input/output directory structure specified in the im_calc_al.template in the checkpoint branch. Failure to match the dir structure will result in runtime error. A guick fix would be modifying the template to suit your own dir structure.

Example:

(1) Simulation

Input/output structure defined in im_calc_al.template

```
echo __calculating simulations__
{% for sim dir, sim_name, fault_name in sim dirs %}
    time python2 $IMPATH/calculate_ims.py {{sim_dir}}/Acc/BB.bin b -o {{sim_dir}}/../../IM_calc/ -np 40 -i {{sim_name}} -r {{fault_name}} -t s -s
{% endfor %}
```

Actual input data structure:

```
melody.zhu@kupe01:/nesi/nobackup/nesi00213/RunFolder/Cybershake/v18p6_batched/v18p6_1k_under2p0G_ab/Runs> ls
Aratiatia ArielBank ArielEast Astrolabe01 Astrolabe02 Astrolabe03 Astrolabe05 Astrolabe07 Awakeri Barefell Bidwill Billys BlueLk BlueMtn BooBooEAST
```

The input binary file is under:

/nesi/nobackup/nesi00213/RunFolder/Cybershake/v18p6_batched/v18p6_lk_under2p0G_ab/Runs/BlueMtn/BB/Cant1D_v3-midQ_oneRay_hfnp2mm+_rvf0p8_sd50_k0p045/BlueMtn_HYP28-31_S1514/Acc/BB.bin

The output IM_calc folder is under:

```
melody, zhugkupe01:/nesi/nobackup/nesi0e213/RunFolder/Cybershake/v18p6 batche/v18p6 1k under2p06 ab/Runs> 1s. /Bluelk/IM calc
Bluelk HYP01-29 51244 Bluelk HYP03-29 51244 Bluelk HYP06-29 51294 Bluelk HYP09-29 51324 Bluelk HYP03-29 51344 Bluelk HYP15-29 51384 Bluelk HYP19-29 51424 Bluelk HYP22-29 51454 Bluelk HYP23-29 51464 Bluelk HYP23-29 Bluelk HYP23-29 S1464 Bluelk HYP23-29 S1464 Bluelk HYP23-29 Bl
```

(2) Observed

Input/output structure defined in im_calc_al.template

Actual input data structure:

melody. Zhu@kupe01:-> ls test obs/JMKaltExample/ 2012p713691 2012p764736 2012p604509 2013p046977 2013p368016 2013p653666 2013p7086602 2013p817946 2013p868761 2014p237547 2014p933966 2014p965622 2016p119534 2016p158394 2016p355041 2122842 3631755

The output IM_calc folder is under:

```
melody.zhu@kupe01:~> ls test_obs/IMCalcExample/IM_calc/
2012p713691 2012p764736 2012p801609 2013p049577 2013p368016 2013p653606 2013p708602 2013p817946 2013p868761
```

Splitting a big slurm

Splitting a big slurm script into several smaller slurms is needed due to the maximum number of lines allowed in a slurm script on Kupe.

Inside **submit_imcalc.py** The **-ml** argument specifies the maximum number of lines of python call to calculate_ims.py/caculate_rrups.py. **Header** and **foot er** like '#SBATCH --time=15:30:00', 'date' etc are **NOT** included.

Say if the max number of lines allowed in a slurm script is 1000, and your (header + footer) is 30 lines, then the number \mathbf{n} that you pass to -mI should be 0 < \mathbf{n} <=967. eq. -mI 967.

Example:

We have 250 simulation dirs to run, by specifying -ml 100 (100 python calls to calculate_ims.py per slurm script), we expect 3 sim slurm scripts to be outputted.(1-100, 100-200, 200-250)

We have 3 observed dirs to run, by specifying -ml 100 (100 python calls to calculate_ims.py per slurm script), we expect 1 sim slurm scripts to be outputted.

We have 61 rrup files to run, by specifying -ml 100 (100 python calls to calcualte_rrups.py per slurm script), we expect 1 sim slurm scripts to be outputted.

Command to run checkpointing and splitting:

```
python submit_imcalc.py -obs ~/test_obs/IMCalcExample/ -sim runs/Runs -srf /nesi/nobackup/nesi00213/RunFolder /Cybershake/v18p6_batched/v18p6_exclude_1k_batch_6/Data/Sources -l1 /scale_akl_nobackup/filesets/transit /nesi00213/StationInfo/non_uniform_whole_nz_with_real_stations-hh400_v18p6.ll -o ~/rrup_out -ml 1000 -e -s -i OtaraWest02_HYP01-21_S1244 Pahiatua_HYP01-26_S1244 -t 24:00:00
```

Output:

```
-rw-r--r-- 1 melody.zhu melody.zhu
                                    48890 Jul
                                                1 03:47 sim im calc 0.sl
-rw-r--r-- 1 melody.zhu melody.zhu
                                    48674 Jul
                                                1 03:47 sim im calc 100.sl
-rw-r--r-- 1 melody.zhu melody.zhu
                                    31922
                                           Jul
                                                1 03:47 sim im calc 200.sl
                                     1329
           1 melody.zhu melody.zhu
                                           Jul
                                                1
                                                  03:47 obs im calc 0.sl
-rw-r--r-- 1 melody.zhu melody.zhu
                                     23246
                                           Jul
                                                  03:47
                                                        rrup im calc 0.sl
```

To submit the slurm script:

```
$cp test.sl /nesi/nobackup/nesi00213/tmp/auto_preproc
$sbatch test.sl
```

The reason that we have to run 'test.sl' under '/nesi/nobackup/nesi00213/tmp/auto_preproc' is otherwise slurm cannot find machine.env specified by the test.sl script:

source machine_env.sh

TODO

- Creation of semi-automatic slurm generation that will have all the calls to produce the results as needed.
- Progress printing statements
 Rrup calculation on a smaller station list currently when generating the slurm script it does the full grid even for stations outside the domain

Notes

• Extensive re-writing of code needs to have smaller deliverables in the future, as this simplifies the integration.