README

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| a_semi_model_final.py
| a_train_final.py
| a_train_final_sim.py
                                                 # model running script
| a_Transformer.py
| a_utilize_semi.py
                                                 # Sigma generating script
| generate_Sigma.py
                                                 # SimDat generating script
| gen_Data_SIM.R
| get_Folds.R
| pack_prepare.R
                                                 # package preparation for R
scripts
| ReadMe_Simulation.txt
| run_example.sh
                                                 # Example for the whole
pipeline
                                                 # Principle of simulation data
| Simdat_generation.pdf
generation
  simulation_data_functions_v4.R
⊢.idea
|—data
      embedding_selected.csv
                                                 # embedding_selected for
simulation
      embedding_selected.npy
      Norms.csv
      Sigma.csv
                                                 # sigma generated for SimDat
genration
      Sigma.npy
                                                 # example for embeddings and
|—example
input data
      embedding_example.csv
      Input_example.csv
      input_example_sim.csv
⊢Results
|-Results_example
|—Simulation
```

```
∟SimDat
                                            # Simdat generated by me
    \vdashSimDat.1
                                            # Simdat.1 is generated using
                                            # All of SimDat.1
    | | SimDat.1.csv
    | | SimDat.1.Rds
       | SimDat.1_labeled.csv
                                            # labeled part of SimDat.1
      | SimDat.1_label_patient_info.csv
      | SimDat.1_unlabeled.csv
                                            # unlabeled part of SimDat.1
       | SimDat.1_unlabeled.csv6116_ALL.pkl
    | | SimDat.1_unlabeled.csv9061_ALL.pkl
       | test_patients.csv
                                           # test patients' numbers
                                        # train patients' numbers
      | train_patients_450.csv
       | unlabeled_patients.csv
                                          # unlabeled patients' numbers
    | <del>|</del>test
              test_data.csv
      ∟train
                                            # Simdat.2 is generated using
              train_data.csv
    └─SimDat.2
```

Input Data & Data Generation

Input Structure

- ID: patient_num
- Y: the label 0 or 1
- T: integers that flags the dates
- other columns for the covariant(columns length is flexible)

```
"","ID","Y","T","S.1","S.2","S.3","S.4","S.5","S.6","S.7","S.8","S.9","S.10"
"100",6,0,0,0,0,0,0,0,0,0,0,0,0,0
"101",6,0,1,0,0,0,0,0,0,0,0,0,0
"102",6,0,2,0,0,0,0,0,0,0,0,0,0,0
"104",6,0,3,0,0,0,0,0,0,0,0,0,0
"105",6,0,5,0,0,0,0,0,0,0,0,0,0
"106",6,0,6,0,0,0,0,0,0,0,0,0,0
"107",6,0,7,0,0,0,0,0,0,0,0,0,0
"108",6,0,8,0,0,0,0,0,0,0,0,0,0
"109",6,1,9,13,5,6,7,7,10,11,18,4,4
"110",6,1,10,8,4,3,4,7,12,14,11,2,4
"111",6,1,11,3,0,4,2,3,6,6,3,4,2
"112",6,1,12,0,3,2,0,3,2,10,7,2,4
"113",6,1,13,2,1,0,1,1,1,2,2,0,0
"114",6,1,14,1,1,2,0,0,4,2,0,1
"115",6,1,15,4,1,2,4,3,5,4,8,4,2
"116",6,1,16,3,0,4,1,2,9,9,9,4,1
"117",6,1,17,7,0,5,2,1,2,4,7,1,0
"118",6,1,18,0,0,0,0,0,0,0,0,0,0,0
```

Data Generation

you need to download R and run pack_prepare.R before generating simulation data.

See run_example.py for an example of the whole pipeline.

Principle of simulation data generation.

See detailed description at /Simdat_generation.pdf

Pipeline of data generation

1. First generate Sigma for multivariant norm distribution.

```
run genrate_Sigma.py
```

this python script **generate simulation embeddings** by cutting example_embedding.csv. **Then generate Sigma matrix** using dot production of different concepts' embeddings.

- selected embeddings saved at /data/embeddings_selected.csv
- Sigma saved at `/data/Sigma.csv

The **embedding dimension** and **covariant column number** is both set to 10 initially, change the d or 1 in the script if needed. Remember to **adjust the column names** at line.35 if changing the column numbers

```
$\(\s.1, \s.2, \s.3, \s.4, \s.5, \s.6, \s.7, \s.8, \s.9, \s.10\)

$\(\s.2, \s.3, \s.4, \s.5, \s.6, \s.7, \s.8, \s.9, \s.10\)

$\(\s.2, \s.3, \s.4, \s.5, \s.6, \s.7, \s.8, \s.9, \s.10\)

$\(\s.2, \s.3, \s.4, \s.6, \s.7, \s.8, \s.9, \s.10\)

$\(\s.2, \s.3, \s.4, \s.6, \s.7, \s.8, \s.9, \s.10\)

$\(\s.2, \s.2, \s.3, \s.4, \s.6, \s.7, \s.6, \s.7, \s.7, \s.8, \s.9, \s.10\)

$\(\s.2, \s.2, \s.3, \s.4, \s.6, \s.7, \s.6, \s.7, \s.7, \s.6, \s.7, \s.
```

Example of selected embeddings

Example of generated Sigma

2. Generate simulation data

run Rscript gen_Data_SIM.R 2000 500 100 200 at terminal

```
Rscript gen_Data_SIM.R {1} {2} {3} {4}
# ARGS: 1.total number 2.labeled number 3.train(labeled) num 4. test(labeled)
num
#Rscript gen_Data_SIM.R 2000 500 100 200
```

Generated data is saved at /Simulation/SimDat

- SimDat.1 & SimDat.2 are generated using different Hazard Models.
- Remember to adjust the line.213-line.222 of simulation_data_functions_v4.R if changing the column numbers

Run Model

run a_train_final_sim.py

Look into Simulation_runmodel_descript.txt for description of different ARGS.

Result Structure

- Results are saved in /Results/, you can see example results at /Results_Example/
- Document Tree of Results directory:

```
C:.
Incident_epoch29_818__results_RETTAIN.csv
                                             # important
Incident_epoch30_818__results_RETTAIN.csv
                                              # important
Prevalence__818__results_RETTAIN.csv
                                              # important
results_RETTAIN.csv1_Attenation_value_patient_visit_code_prediction_label_we
ight_test.pkl
results_RETTAIN.csv1__embedding_patient_ori_codeRw_codeVistRw_hiddenFCN_labe
results_RETTAIN.csv1__embedding_patient_ori_hiddenFCN_label_train.pkl
results_RETTAIN.csv_incident_evaluation.txt # important
| results_RETTAIN.csv_prevalence_evaluation.txt # important
results_RETTAIN.csv_code_weights.csv
└results_RETTAIN.csv_model
   | keras_metadata.pb
   | saved_model.pb
   -assets
   ∟variables
           variables.data-00000-of-00001
           variables.index
```

- Incident_epoch{}_results_RETTAIN.csv: incident phenotyping(prediction on every time spot) results
- Prevalence_results_RETTAIN.csv: Prevalence(EVER/NEVER) binary phenotyping results
- .txt: the calculated AUC, ACC, F1, Speci etc.