

Note2: how to install normalizer module-

<https://github.com/bio-ontology-research-group/el-embeddings/tree/37ea9ded9e1b0eef6ac38632863bc7e4ead29708> README.md

Step 5: Run **AddRelations.ipynb** to integrate *hasAnnotation* and *Dir* axioms to ontology.

Change the input and output files as-

df_train – the train g-d pair file

norm_ontology- normalized ontology

annot_file – the annotation file

output_finalized_ontology- finalized ontology integrating all data that will be passed to EIE model.

Step6: Run **Train_Test_Split.ipynb** to generate the 4 train-test sets

Change the input as:

pair file- the name of file that contains the g-d associations

n- number of instances each of positive and negative association you wish to see in test sets. If n=2046 then there will be 2046 positive and negative pairs in the test set I.e, 4092 total pairs.

Step 7: Run **EIE_Algorithm.ipynb** to generate the embeddings.

Change input as

FILE_NAME – name of finalized ontology(.owl) file.

Follow the instructions given in the notebook.

Step 8: Run **Rank_Distribution_Plot.ipynb** to calculate rank distribution of each model.

Change input, output as:

test_data- g-d test file

all_pairs- full g-d pair

lists- lists of all the embedding files. Note the token for each file ('EIE', 'TransE1', 'rdf2vec') should be same.

rel_file- name of relation embedding model. Required only for EIE model.

output_rank_distribution- the file where rank distribution will be saved.

Step9: Run **Evaluation.ipynb** to generate performance statistics of KGE models.

input_rank_file- rank distribution file from prev. step.

output_stats_file_name- output file where the statistics are saved.

Note: for other models like TransE, OWL2Vec etc.,- refer-

https://github.com/liseda-lab/KGE_Predictions_GD