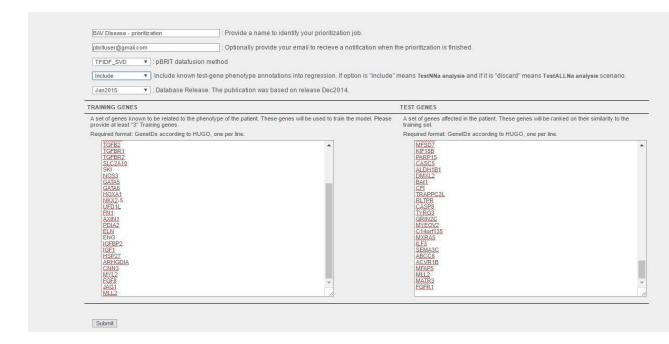# Running pBRIT:

1. **Prioritization Name**: Provide a name to identify your prioritization job. (optional)
2. **Email** : Optional
3. **Data Fusion method**: TF-IDF or TFIDF_SVD
4. **Phenotype Info**: Include or Discard
5. **Database Release**: Internal database of pBRIT (Currently January 2015 version)
6. **Training Genes**: Set of training genes (only HUGO Symbols)
7. **Test Genes**: Set of Candidate genes (only HUGO Symbols)
8. **Submit**: Click submit button for prioritization.

Waiting time for finishing the prioritization is average 90-120 seconds depending upon the queue in the CMG cluster network.