

Week 1

- Complete script for data parsing
- Complete script for data preprocessing
- Read scikit learn documentation on Support Vector Machines (1.4)
- Read scikit learn documentation on cross validation (3.1)
- Read scikit learn documentation on model evaluation (3.3)
- Work on script for k-fold cross-validation

Week 2

- Complete script for k-fold cross-validation
- Plot learning curve of predictor to diagnose bias and variance
- Brainstorm more features to include in predictor
- Use cross validation to optimize C and gamma parameters
- Use the optimum parameters to train the SVM
- Test the SVM on test set

Week 3

- Read the paper on using PSSM to predict protein secondary structure by Jones DT (1999)
- Use psi-blast to extract evolutionary information for my SVM
- Train my SVM based on multiple sequence alignment
- Use cross validation to optimize parameters for the improved SVM
- Use the optimum parameters to train the improved SVM
- Test the SVM on test set
- Prepare and present paper

Week 4

- Read scikit learn documentation on random forest and decision tree
- Create random forest and decision tree predictors
- Compare these predictors to my SVM predictor
- Test performance of my predictors on 50 new proteins
- Review my predictor based on current literature
- Write report for the project

Week 5

- Submit report