1. ProcessDataTPM: Formats input TPM data.
2. 5FoldCrossVal\_ReducedFeaturesTPM: Predicts population label on basis of TPM
3. 5FoldCrossVal\_ReducedFeaturesNewlabel: Predicts Sequencing Center on basis of TPM
4. FinalSubmissionDecisionTreeMultilabel: Predicts both on Sequencing Center and population.
5. DecisionTreeEqClass: Predicts population label on basis of Equivalence Classes
6. ProcessEqClass: Formats input Eq Class data.
7. predict\_experimentLbl\_2: Final submission file
8. FinalPickle: pickle file submitted