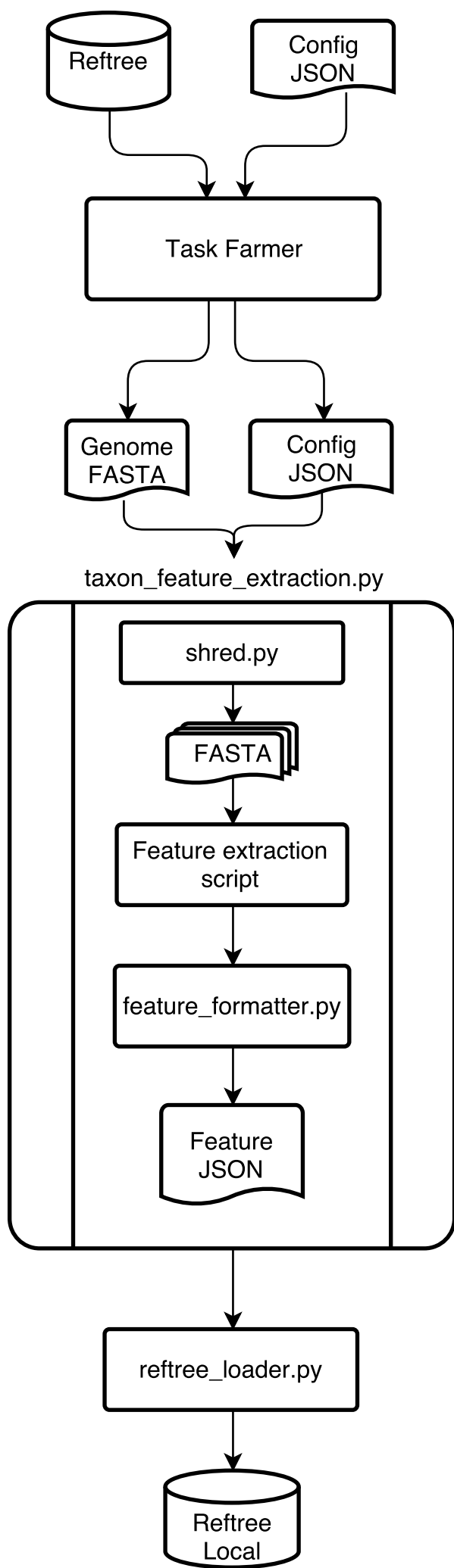
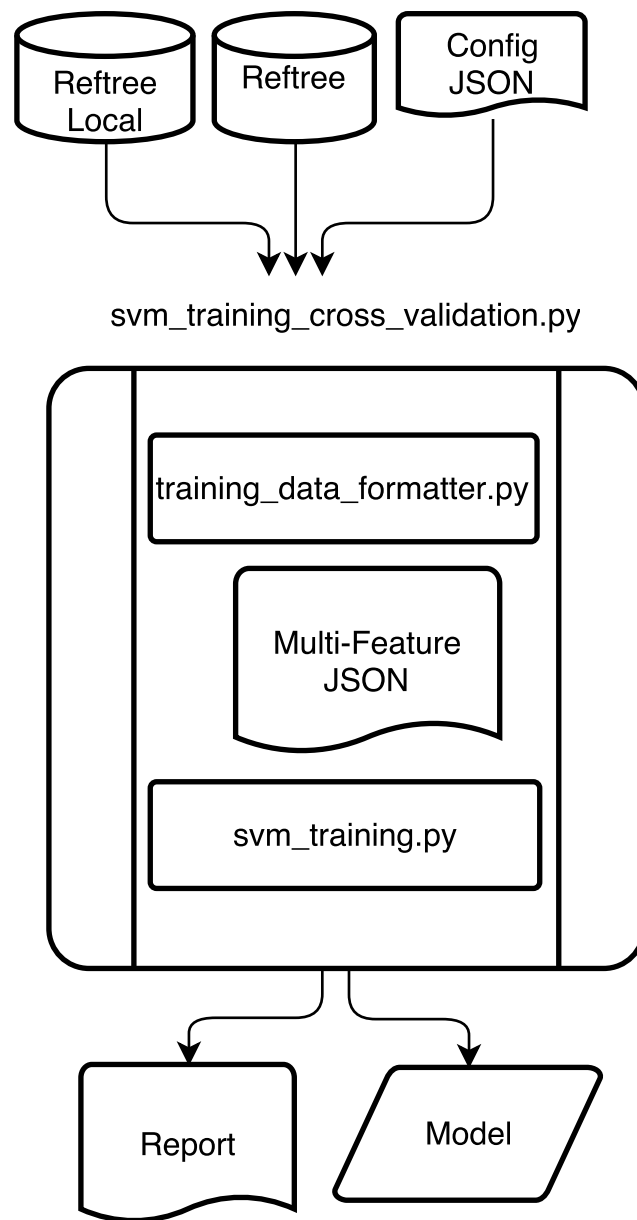


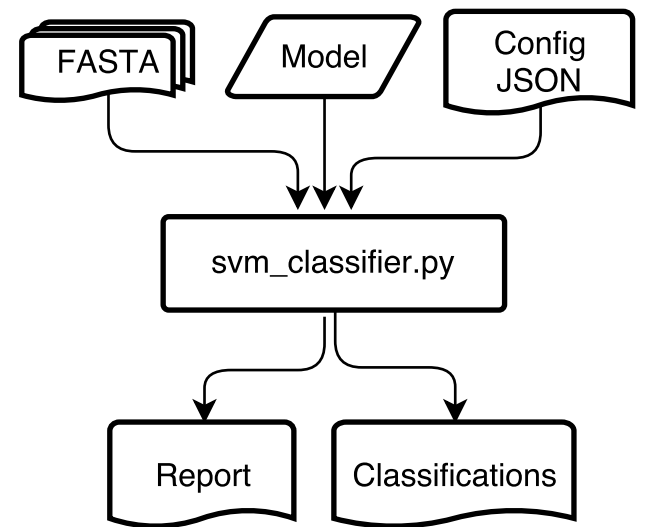
Getting initial data for training



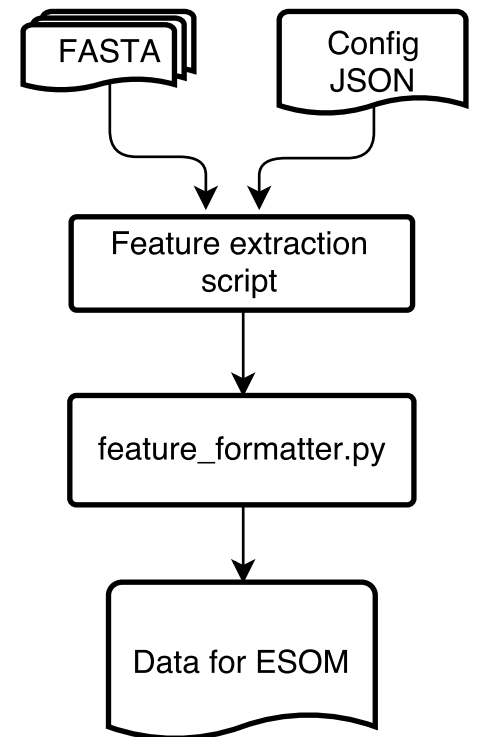
Model training and cross-validation



Sample classification



Metagenome binning



Feature Extraction Scripts



Utilities

