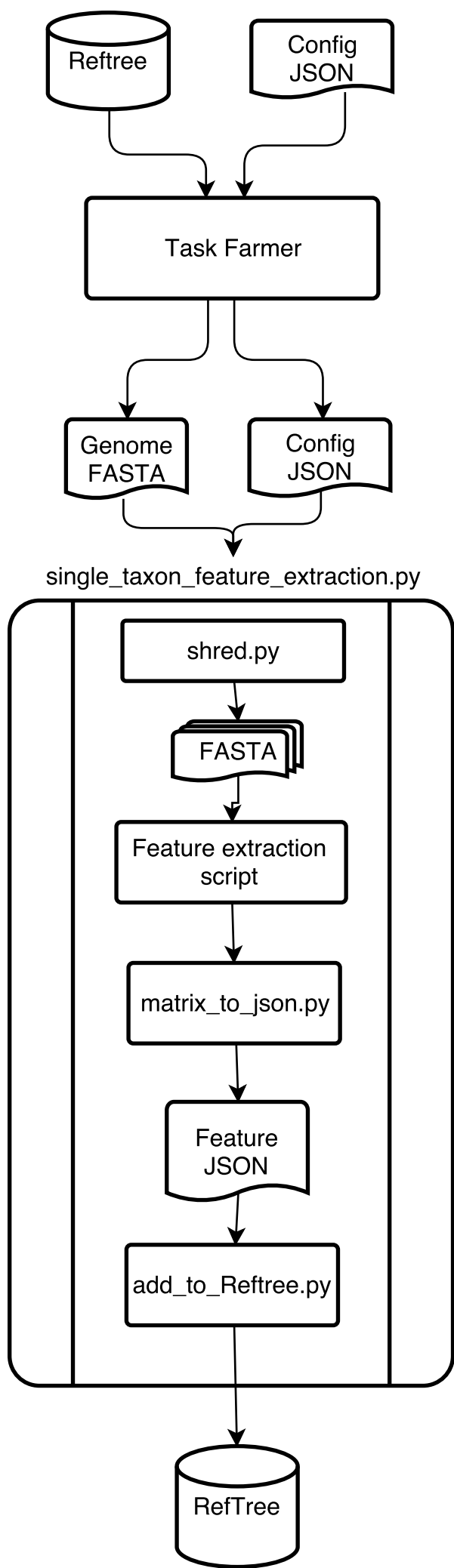
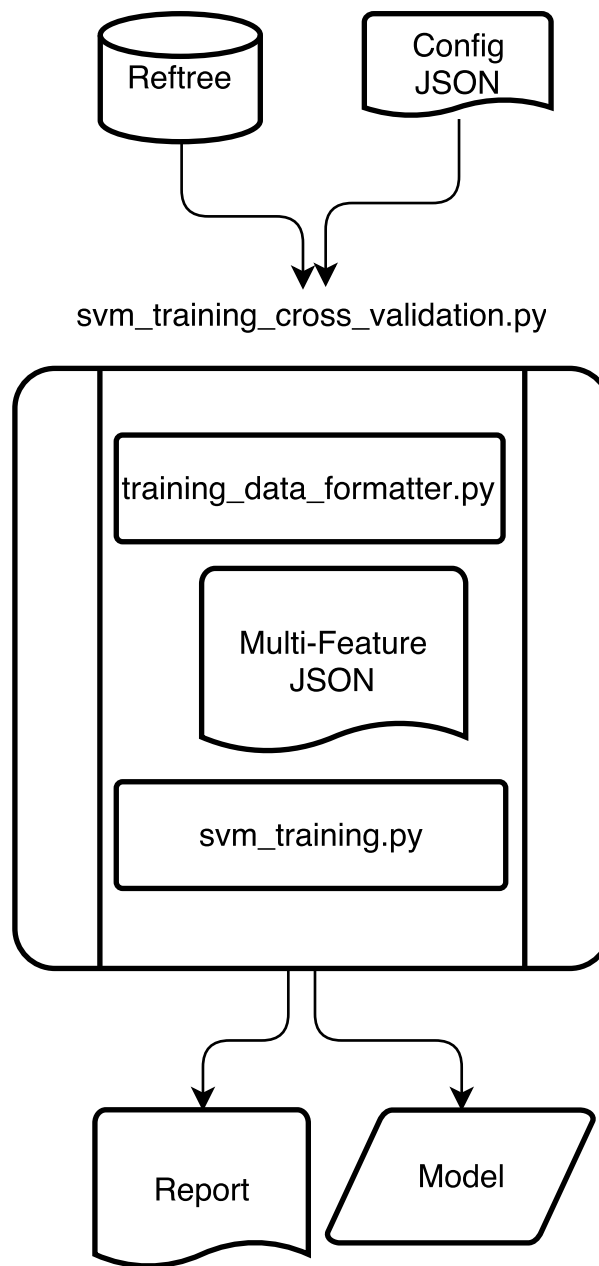


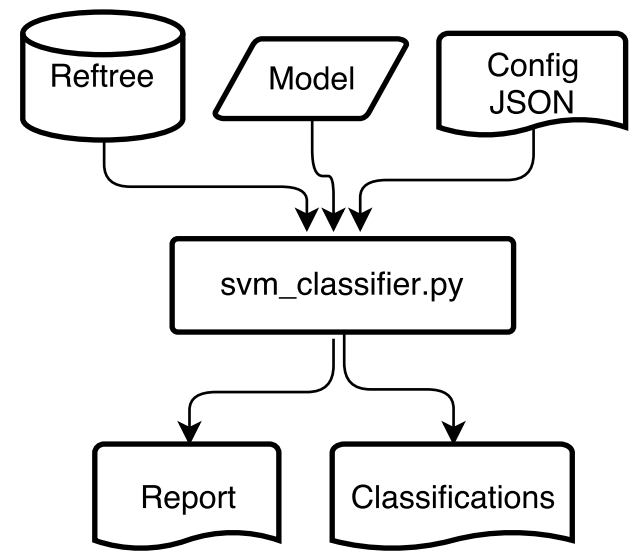
Getting initial data for training



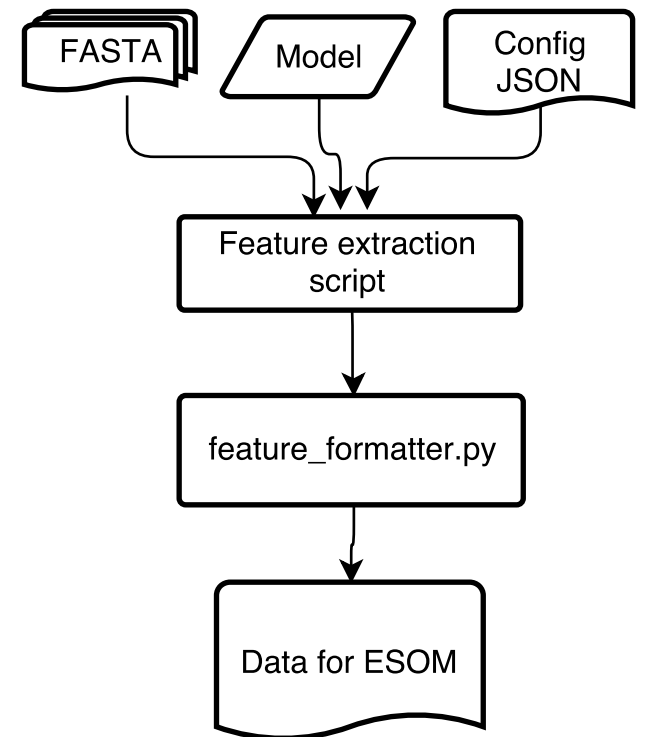
model training and cross-validation



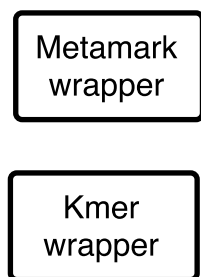
Sample classification



Metagenome binning



Feature Extraction Scripts



Utilities

