Classifier Package

Modules in the **biolockj.module.classifier** package categorize micbrobial samples into Operational Taxonomic Units (OTUs) either by reference or with clustering algorithms.

This package contains 2 sub-packages:

1. [[module.classifier.r16s]] contains modules designed to classify 16S data.
2. [[module.classifier.wgs]] contains modules designed to classify whole genome sequence data.

Modules in these sub-packages extend the ClassifierModuleImpl class.

ClassifierModuleImpl

Description: Abstract implementation of the ClassifierModule interface that the other classifier modules extend to inherit standard functionality.
Abstract modules cannot be included in the pipeline run order.