TargetExperiment

+ bedFile : GRanges
+ bamFile : BamFile
+ fastaFile : FaFile
+ featurePanel : GRanges
+ genePanel : GRanges
+ scanBamP : ScanBamParam
+ pileupP : PileupParam
+ attribute : character
+ feature : character

+ TargetExperiment()
+ pileupCounts()
+ buildFeaturePanel()
+ summarizePanel()
+ show()
+ print()
+ summary()
+ summaryFeatureLev()
+ summaryGeneLev()
+ summaryIntervals()
+ getLowCtsFeatures()
+ plot()
+ plotAttrExpl()
+ plotFeatPerform()
+ plotFeature()
+ plotGeneAttrPerFeat()
+ plotNtdPercentage()
+ getRegion()
+ plotRegion()
+ buildReport()

GRanges

FaFile

BamFile

ScanBamParam

PileupParam