

miRSM

dumbbell microbial MetaCyto miRspongeR rnmamod FlowSOM HierDpart **SPONGE** FRK MQmetrics epidecodeR plotGrouper entropar\$COUTer UCSCXenaShinyScITD grapesAgri1 MethReg iglu TestGardener BNPmixspatialTIME digitalDLSorteR cageminer GeneAccord IRISFGM LinkHD vmeasur crctStepdown promotionImpact FuncNN APCtools Bion dasper ggquickeda FuncNN ShellChron insane
ENMTools MetaIntegrator APCI GenomicSuperSignature metaviz

dataquieR epigraHMM SEIRfansy UniprotR
metaviz

MSPRT
maseqSampleSize ggpubr
factorMerger ShellChron missingHE MSPRT SiteAdapt factorMerger dfoliatR viewpoly convergEU ParBayesianOptimization tipsae orthogene miceRanger PhosR coda4microbiome methylclock TADCompare rrtable ODER dPCP greatR shinyTempSignal PUMP InTAD fastaicvCovEst ptairMS ClustEff processR gfoRmula CeTF clustEff bayMDS popstudy diversityForest MERO CovidMutations vsd bdl SHAPforxgboost iCellR rPACI ADAMgui iPath BinMat ELMER VALERIE TNBC.CMS conformalInference.fd RVA volcano3DhdpGLM^{TargetDecoy} **TCGAbiolinksGUI** riskCommunicator AnnoProbe sigminer glmSparseNet pathwayTMB **PDATK** reconstructKM survtype GDCRNATools survminer MetabolicSurv CondiS LongCART ivygapSE concurve SurvMetrics tinyarray GFDsurv omicsViewer PMAPscore survival666 survivalAnalysis RTCGA.mutations RTCGA RTCGA.miRNASeq RTCGA.rnaseq RTCGA.CNV RTCGA.mRNA

RTCGA.PANCAN12 RTCGA.RPPA