0.0 0.2 0.0 0.0 0.6 0.2 0.0 0.0 0.0 0.5 0.0 0.0 0.1 0.2 0.0 0.1 0.1 0.1 0.0 0.0 0.2 0.0 0.0 0.3 0.0 0.1 0.0 0.0 0.0 0.1 0.0 0.4 0.0 0.1 0.0 0.0 0.0 0.0 0.0 0.3

crisprcas9 drugresponse netabolomics methylation proteomics

 $0.0010\,0.0040\,$ -44-08-21 (0.01 (0 0.000 67 68 69 71 

GeneExpressionCorrelation CopyNumberInstability ploids mutational burden growth size RensCorrelation VIM prot CDH1\_gexp VIM\_gexp D/F12 sanger Adherent MeanProteomics MeanDrugResponse

