1.3 0.8 2.1 0.1 1.4 7.4 4.5 10.0 0.8 1.0 3.6 -0.8 5.3 2.4 5.9 6.5 0.2 0.0 22 1 0.0 0.1 0.0 0.2 0.1 1.7 0.0 0.0 0.0 0.6 0.0 14.9 0.2 0.2 0.0 0.0 0.0 0.0 8.0 0.1 0.7 1.6 0.4 1.0 0.1 0.6 1.5 1.6 2.2 0.6 0.3 1.5 0.2 0.6 1.4 1.0 0.2 0.1 0.2 0.0 0.1 5.3 0.1 1.1 0.6 0.2 0.8 0.0 0.9 1.3 1.1 1.4 0.5 0.2 1.0 0.1 1.0 0.8 0.9 1.1 0.0 4.7 0.0 -0.0 0.0 0.0 0.2 0.1 0.1 0.2 0.6 0.0 0.9 1.3 0.5 1.3 0.1 0.0 0.0 -0.9 0.1 4.3 0.1 1.0 0.5 0.2 0.5 0.1 0.3 1.3 0.5 1.4 0.1 3.2 0.1 0.0 0.4 0.1 0.1 0.0 0.5 0.0 1.1 0.0 0.5 0.0 1.7 0.1 1.2 0.7 0.5 0.1 0.2 0.1 0.4 0.3 0.7 0.1 0.2 0.0 0.2 0.5 0.5 0.8 0.7 0.1 0.2 0.0 0.2 0.5 0.5 0.6 2.2 0.1 0.1 0.0 0.1 0.0 0.3 0.2 0.1 0.1 0.4 0.0 0.2 0.9 0.4 0.9 0.3 0.1 0.2 0.0 0.3 0.7 0.4 0.9 0.2 0.1 0.0 -0.1 0.1 0.0 0.0 2.5 0.1 2.5 0.0 0.0 0.0 0.1 0.0 0.0 0.2 0.1 0.3 0.0 0.2 0.7 0.3 1.0 0.2 0.1 0.6 0.1 0.1 0.6 0.3 0.7 1.9 0.0 0.1 0.0 0.1 0.1 0.3 0.2 0.5 0.1 0.2 0.0 0.3 0.4 0.4 0.6 1.4 0.0 0.1 0.0 0.1 0.1 0.3 0.3 0.1 0.0 0.0 0.0 0.2 0.0 1.9 0.1 0.2 0.0 0.0 0.0 0.1 0.3 0.4 0.0 1.5 0.0 0.0 0.0 0.1 0.1 0.2 0.2 1.6 0.0 0.0 0.0 0.1 1.2 0.1 0.1 0.0 0.0 0.1 0.2 0.1 1.2 0.0 0.1 0.0 0.1 0.0 0.2 0.1 0.3 0.0 0.0 0.0 0.2 0.6 0.1 0.5 0.9 0.0 0.1 0.0 0.3 0.1 0.2 0.1 0.1 0.1 0.2 0.0 0.4 0.2 0.2 0.5 0.0 0.1 0.1 0.0 0.1 0.7 0.3 0.4 0.0 1.2 0.0 0.0 0.2 0.1 0.2 0.2 0.2 0.0 0.1 0.0 0.0 0.4 0.3 0.4 0.0 0.2 0.0 0.0 0.0 0.1 0.1 0.8 0.0 0.7 0.0 0.0 0.3 0.0 0.0 0.0 0.5 0.0 0.1 0.0 0.3 0.0 0.1 0.1

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 methylation proteomics

crisprcas9 drugresponse netabolomics

 $\begin{smallmatrix} 0.01 & 0.01$ $\begin{smallmatrix} 0.5, 0.05, 0.02, 0.02, 0.03, 0.04, 0.03, 0.0$ $\frac{1}{100} + \frac{1}{100} + \frac{1}$ $0.641 \pm 0.01 \pm$ 67 68 69 71 $\frac{1}{11}\frac{1}$

CopyNumberInstability ploids mutational burden growth size CDH1 prot VIM prot CDH1_gexp VIM_gexp D/F12 sanger Adherent MeanProteomics MeanDrugResponse

 $-0.33 \pm 0.063 \pm 0.033 \pm 0.040 \pm 0.04$ Reps Correlation and an analysis and an analys $-0.12 \cdot 0.0 \cdot 0.$ $-0.130 \cdot 0.0 \cdot 0$ sanger Sem Adherent - 0.000 0. $-0.23 \cdot 0.000 \cdot 0.0023 \cdot 0.020 \cdot 0.0000 \cdot 0.000 \cdot 0.000 \cdot 0.002 \cdot 0.002 \cdot 0.0022 \cdot 0.0000 \cdot 0.0020 \cdot$ -thania-ado-thania-ado-thania-ado-thania-ado-thania-ado-thania-ado-thania-ado-thania-ado-thania-ado-thania-ado