CopyNumberAttenuation GeneExpressionCorrelation CopyNumberInstability ploidy mutational burden growth size RepsCorrelation CDH1 prot VIM prot CDH1 gexp VIM_gexp D/F12 sanger Adherent sanger_Semi-Adherent sanger_Suspension sanger_Unknown broad_Adherent

F3

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0.9 0.0 0.4 0.3 0.0 0.1 0.1 0.0 0.0 0.1 0.0 1.4 01 01 01 03 04 05 00 01 01 00 07 05 0.0 0.1 0.2 0.7 0.2 0.3 0.0 0.0 0.0 1.0 0.1 0.2 00 01 01 02 04 04 00 01 00 00 03 05 0.2 0.0 0.0 0.0 0.0 0.7 0.0 0.0 0.2 0.1 0.0 0.3 proteomics crisprcas9 ngresponse ranscriptomics

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1.2 0.0 0.0 0.0 0.6 0.1

0.1 0.1 0.7 0.9 0.1 0.2

0.1 0.2 0.2 0.4 0.4 0.6

1.1 0.1 0.0 0.3 0.1 0.2

0.1 0.0 0.0 0.0 0.2 1.5

0.0 0.0 1.3 0.2 0.2 0.1

0.2 0.1 0.2 0.6 0.2 0.4

0.1 0.2 0.3 0.2 0.3 0.6

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