F49 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 broad Mixed broad Suspension broad Unknown MeanProteomics MeanMethylation MeanDrugResponse

F2

F3

F5

F6

E7 F8

FQ

E11

F13

F14

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-0.01 -0.0

0.1 + 0.0 + 0.0 + 1.0 + 0.0 + 1.0 + 0.0

0.0 0.0 0.2 0.1 0.0 0.3 proteomics ngresponse methylatior ranscriptomics

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

01 02 03 02 03 06

08 00 04 02 01 02

0.9 0.0 0.4 0.3 0.0 0.1

01 00 00 01 00 14

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

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