0.1 0.2 0.2 0.4 0.4 0.6

11 01 00 03 01 02

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

0.0 0.0 0.2 0.1 0.0 0.3

crisprcas9 ngresponse. methylation proteomics ranscriptomics

 $-0.00.1\ 0.0 - 0$ 

 $0.1 \cdot 0.0 \cdot 0.0$ 

growth size RepsCorrelation CDH1\_prot VIM prot D/F12 R sanger Adherent sanger Semi-Adherent sanger Suspension sänger Unknown broad Adherent broad Mixed broad Unknown MeanProteomics MeanDrugResponse

CopyNumberAttenuation

CopyNumberInstability

mutational burden

GeneExpressionCorrelation

F2

F3

F5

F6

E7

F8

FQ

F13

F14

F15

F16

F17

F18

F19

F21

F22

F23

F24

F25

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F49

ploidy

0.103 $-0.0 \\ \hline -0.0 \\ -0.0 \\ \hline -0.0 \\ -0.0 \\ \hline -0.0 \\ -0.0 \\ \hline -0.0 \\ -0.0 \\ \hline -0.0 \\ -0.0 \\ \hline -0.0 \\ -0.0 \\ \hline -0.0$ broad Suspension 0.1 0.3 -0.0 -0.0 -0.1 0.1 0.0 -0.0 0.1 -0.2 -0.1 0.1 -0.2 0.2 0.1 0.0 -0.1 0.1 0.0 -0.1 0.2 0.1 -0.1 0.0 0.0 0.1 -0.2 0.0 -0.1 0.1 0.0 -0.1 -0.2 0.0 -0.1 0.1 0.0 -0.1 -0.2 0.0 0.2 -0.1 0.1 -0.2 -0.1 0.1 -0.2 -0.0 -0.1 -0.1 0.2 -0.0 0.1 -0.1 0.0  $-0.0-0.0 \ 0.0 \$