

Haem cell lines	drespo	-	0.2	0.1	0.2	0.4	0.3
	methylation	-	-0.4	0.4	10.5	6.5	0.1
	proteomics	-	1.3	1.3	0.0	4.0	3.3
	proteomics	-	0.2	0.4	0.1	0.4	6.1
	transcriptomics	-	0.3	0.9	1.6	1.7	0.0
Other cell lines	drespo	-	2.5	4.3	0.1	0.4	0.4
	methylation	-	11.1	3.3	8.2	3.9	1.0
	proteomics	-	6.3	4.3	0.0	2.7	2.7
	proteomics	-	6.7	4.6	0.0	0.4	6.0
	transcriptomics	-	12.3	9.3	1.8	1.1	1.3
Corr_CopyNumberAttenuation		-	0.26	-0.09	-0.14	0.19	-0.02
Corr_GeneExpressionAttenuation		-	-0.24	0.30	-0.00	-0.04	0.01
Corr_EMT		-	-0.15	-0.59	-0.05	0.08	0.05
Corr_Proteasome		-	-0.17	-0.03	0.06	-0.02	-0.06
Corr_TranslationInitiation		-	-0.17	-0.15	0.15	-0.09	-0.03
Corr_CopyNumberInstability		-	0.17	-0.06	-0.05	0.14	-0.09
Corr_CDH1_prot		-	0.59	-0.22	0.23	-0.02	0.19
Corr_VIM_prot		-	-0.54	-0.58	0.04	0.05	0.01
Corr_CDH1_gexp		-	0.72	0.19	-0.04	0.14	-0.10
Corr_VIM_gexp		-	-0.57	-0.56	0.01	0.09	0.06
Corr_D/F12		-	-0.02	-0.12	0.00	0.02	0.08
Corr_R		-	0.02	0.12	-0.00	-0.02	-0.08
Corr_Adherent		-	0.12	-0.19	0.07	0.05	0.03
Corr_Semi-Adherent		-	-0.01	0.17	-0.05	0.00	0.02
Corr_Suspension		-	-0.07	0.28	-0.04	-0.11	-0.07
Corr_Unknown		-	-0.07	-0.12	-0.03	0.05	0.03
Corr_Haematopoietic and Lymphoid		-	0.01	-0.00	0.01	-0.01	-0.01
Corr_Lung		-	-0.04	0.24	-0.20	0.06	-0.11
Corr_ploidy		-	0.06	-0.08	0.04	0.01	-0.03
Corr_mutational_burden		-	0.12	0.15	0.19	-0.04	-0.07
Corr_growth		-	0.03	-0.17	0.23	-0.10	-0.01
Corr_size		-	-0.00	-0.35	0.06	0.15	-0.03
Corr_RepsCorrelation		-	0.13	0.14	-0.10	0.09	0.01
Corr_MeanProteomics		-	-0.03	-0.16	-0.04	-0.04	-0.04
Corr_MeanProteomicsBroad		-	0.02	0.03	-0.01	0.04	-0.10
Corr_MeanMethylation		-	-0.00	-0.01	0.41	0.07	0.03
Corr_MeanDrugResponse		-	0.13	-0.06	-0.02	0.05	-0.03
			F1	F2	F3	F4	F5