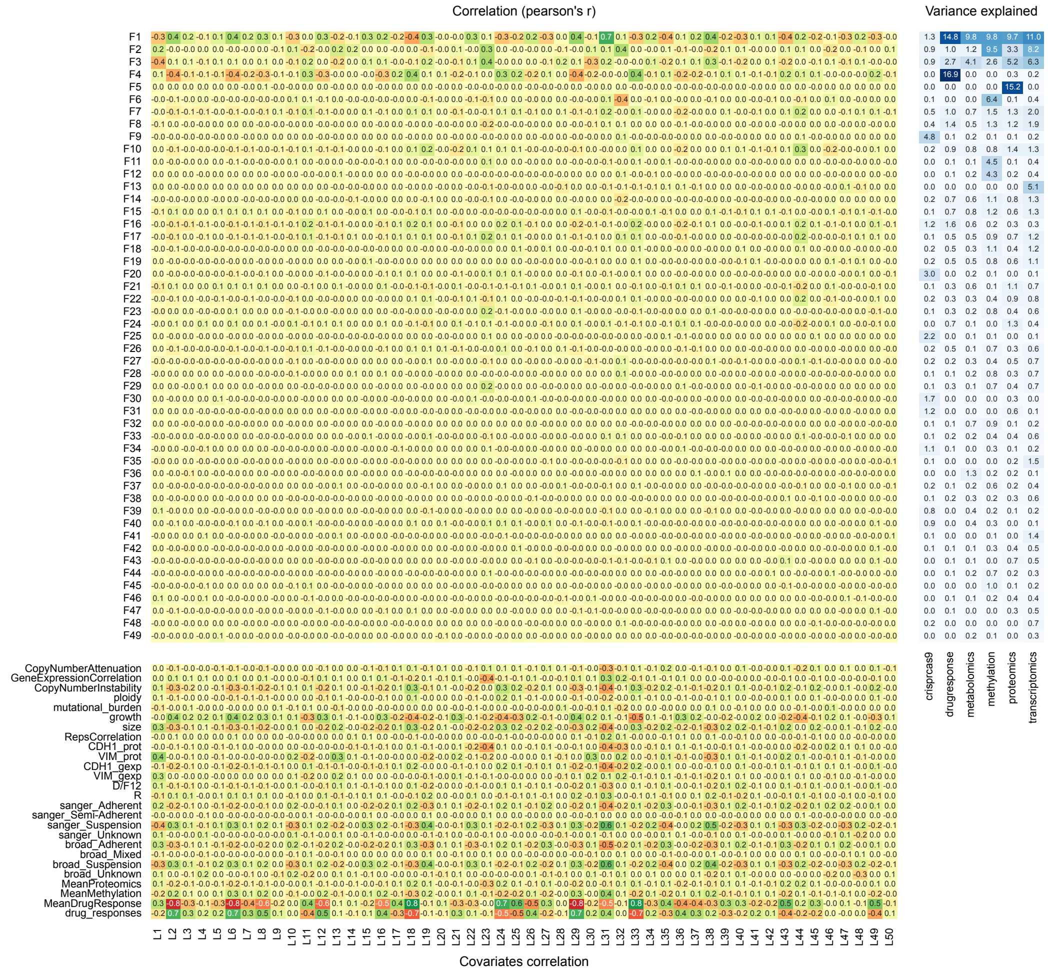
Correlation (pearson's r)
Variance explained
Covariates correlation
F1
F2
F3
F4
F5
F6
F7
F8
F9
F10
F11
F12
F13
F14
F15
F16
F17
F18
F19
F20
F21
F22
F23
F24
F25
F26
F27
F28
F29
F30
F31
F32
F33
F34
F35
F36
F37
F38
F39
F40
F41
F42
F43
F44
F45
F46
F47
F48
F49
CopyNumberAttenuation
GeneExpressionCorrelation
CopyNumberInstability
ploidy
mutational_burden
growth
size
RepsCorrelation
CDH1_prot
VIM_prot
CDH1_gexp
VIM_gexp
D/F12
R
sanger_Adherent
sanger_Semi-Adherent
sanger_Suspension
sanger_Unknown
broad_Adherent
broad_Mixed
broad_Suspension
broad_Unknown
MeanProteomics
MeanMethylation
MeanDrugResponse
drug_responses
L1 L2 L3 L4 L5 L6 L7 L8 L9 L10 L11 L12 L13 L14 L15 L16 L17 L18 L19 L20 L21 L22 L23 L24 L25 L26 L27 L28 L29 L30 L31 L32 L33 L34 L35 L36 L37 L38 L39 L40 L41 L42 L43 L44 L45 L46 L47 L48 L49 L50
crisprcas9
drugresponse
metabolomics
methylation
proteomics
transcriptomics
1.3 14.8 9.8 9.8 9.7 11.0
0.9 1.0 1.2 9.5 3.3 8.2
0.9 2.7 4.1 2.6 5.2 6.3
0.0 16.9 0.0 0.0 0.3 0.2
0.0 0.0 0.0 0.0 15.2 0.0
0.1 0.0 0.0 6.4 0.1 0.4
0.5 1.0 0.7 1.5 1.3 2.0
0.4 1.4 0.5 1.3 1.2 1.9
4.8 0.1 0.2 0.1 0.1 0.2
0.2 0.9 0.8 0.8 1.4 1.3
0.0 0.1 0.1 4.5 0.1 0.4
0.0 0.1 0.2 4.3 0.2 0.4
0.0 0.0 0.0 0.0 0.0 5.1
0.2 0.7 0.6 1.1 0.8 1.3
0.1 0.7 0.8 1.2 0.6 1.3
1.2 1.6 0.6 0.2 0.3 0.3
0.1 0.5 0.5 0.9 0.7 1.2
0.2 0.5 0.3 1.1 0.4 1.2
0.2 0.5 0.5 0.8 0.6 1.1
3.0 0.0 0.2 0.1 0.0 0.1
0.1 0.3 0.6 0.1 1.1 0.7
0.2 0.3 0.3 0.4 0.9 0.8
0.1 0.3 0.2 0.8 0.4 0.6
0.0 0.7 0.1 0.0 1.3 0.4
2.2 0.0 0.1 0.1 0.0 0.1
0.2 0.5 0.1 0.7 0.3 0.6
0.2 0.2 0.3 0.4 0.5 0.7
0.1 0.1 0.2 0.8 0.3 0.7
0.1 0.3 0.1 0.7 0.4 0.7
1.7 0.0 0.0 0.0 0.3 0.0
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
0.8 0.0 0.4 0.2 0.1 0.2
0.9 0.0 0.4 0.3 0.0 0.1
0.1 0.0 0.0 0.1 0.0 1.4
0.1 0.1 0.1 0.3 0.4 0.5
0.0 0.1 0.1 0.0 0.7 0.5
0.0 0.1 0.2 0.7 0.2 0.3
0.0 0.0 0.0 1.0 0.1 0.2
0.0 0.1 0.1 0.2 0.4 0.4
0.0 0.1 0.0 0.0 0.3 0.5
0.2 0.0 0.0 0.0 0.0 0.7
0.0 0.0 0.2 0.1 0.0 0.3