Correlation (pearson's r)

Variance explained

Covariates correlation

Row labels (factors): 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

Covariate row labels: 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

Variance explained column headers: crisprcas9, drugresponse, metabolomics, methylation, proteomics, transcriptomics

Bottom axis labels: 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