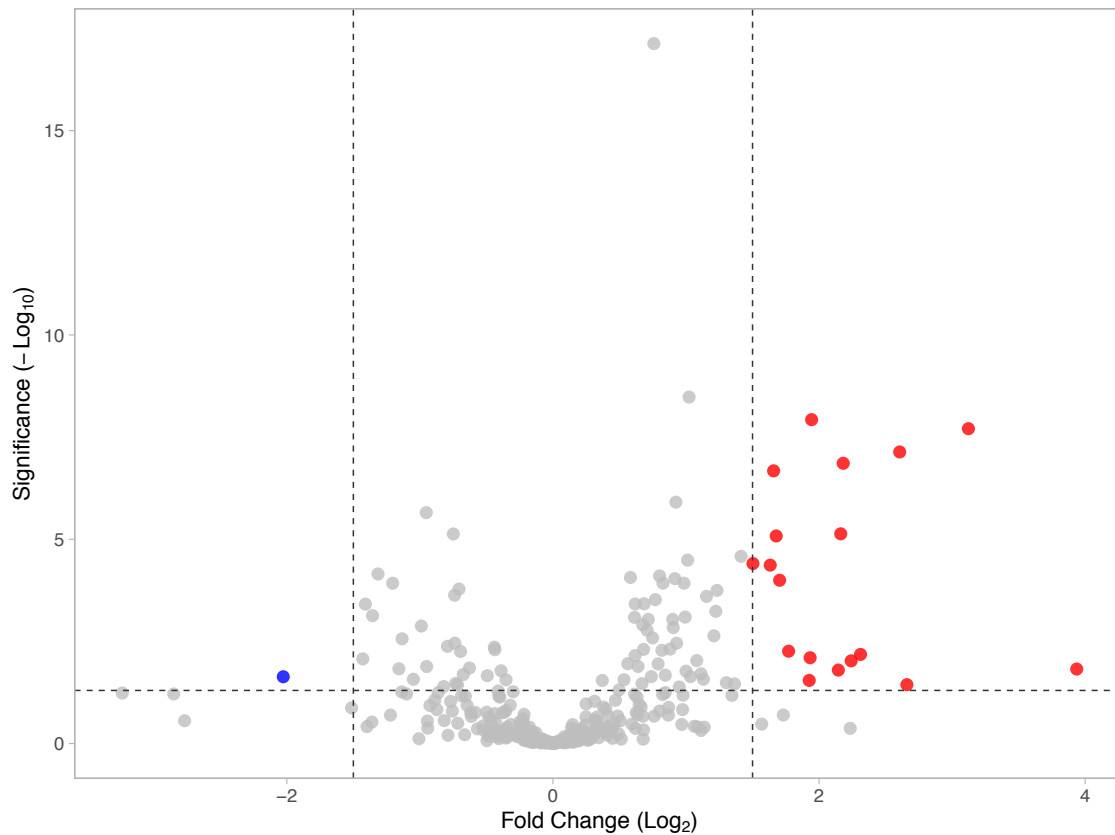


Figure S3

Distribution of 336 *GCN5*-regulated genes from Shivarathri et al. in the *tra1Q3* dataset

a



b

Gene ID	Log ₂ (Fold-Change) <i>gcn5</i> Δ/Δ	Log ₂ (Fold-Change) <i>tra1Q3</i>
CR_05270W_A	1.74	-2.03
C3_06660C	2.43	1.50
RBR1	1.73	1.63
HAK1	2.2	1.66
FTR1	3.01	1.68
C2_04910W_A	2.03	1.70
FET99	3.35	1.77
CFL5	2.51	1.93
CR_04990C_A	1.91	1.93
PGA13	3.88	1.94
RBT5	2.48	2.14
CFL2	3.35	2.16
CRP1	-2.64	2.18
ATX1	-3.64	2.24
C7_04070C_A	2.29	2.31
CFL4	6.99	2.61
CFL11	2.63	2.66
CR_05190W_A	1.97	3.12
C7_04080C_A	1.58	3.94