

Cellular Process (Assessed by Gene Ontology)	Genes Down-regulated 2 fold or more
Metabolic process	AAC3, ADH7, ANB1, ATG15, BIO2, BTT1, CPS1, CSN9, CUS1, DAK2, DAL80, ECM11, EDC1, FAL1, GPD2, HEM13, HIS1, HPR1, IRC4, IRC7, LYS9, MET14, MET16, MET17, MET2, MET32, MFT1, MIG2, MIG3, MKC7, NAT5, NDJ1, NRK1, PAU8, PDC5, PHO11, PHO5, PUF3, RBL2, RCK1, RPS22A, SAE2, SAP30, SMP1, SOR2, SPE3, SRN2, TAF3, THI4, THI80, UTR2, YAP6, YEH1, YGK3, YGR287C, YNL108C, ZAP1
transport	AAC3, APL6, BIK1, COG4, COG8, DAL5, DAN1, GIT1, HOT13, HPR1, HXT1, HXT11, HXT2, MFT1, OAC1, RIX7, RUD3, SLO1, RN2, SRP40, YMC2, ZRT1
transcription	DAL80, HPR1, MFT1, MIG2, MIG3, PAU8, SAP30, SMP1, TAF3, YAP6, ZAP1
cell cycle	BFA1, BIK1, BRN1, DAD2, ECM11, HUG1, MAM1, NDJ1, RCK1, SAE2, SCM3, SHE1, SPO22, TEM1, VIK1
cellular amino acid metabolic process	HIS1, LYS9, MET14, MET16, MET17, MET2, MET32, PDC5
signal transduction	BFA1
Not yet annotated	YDL038C
cannot be mapped to a GO slim term	PRM7, SPL2, TIR1, YCR102C