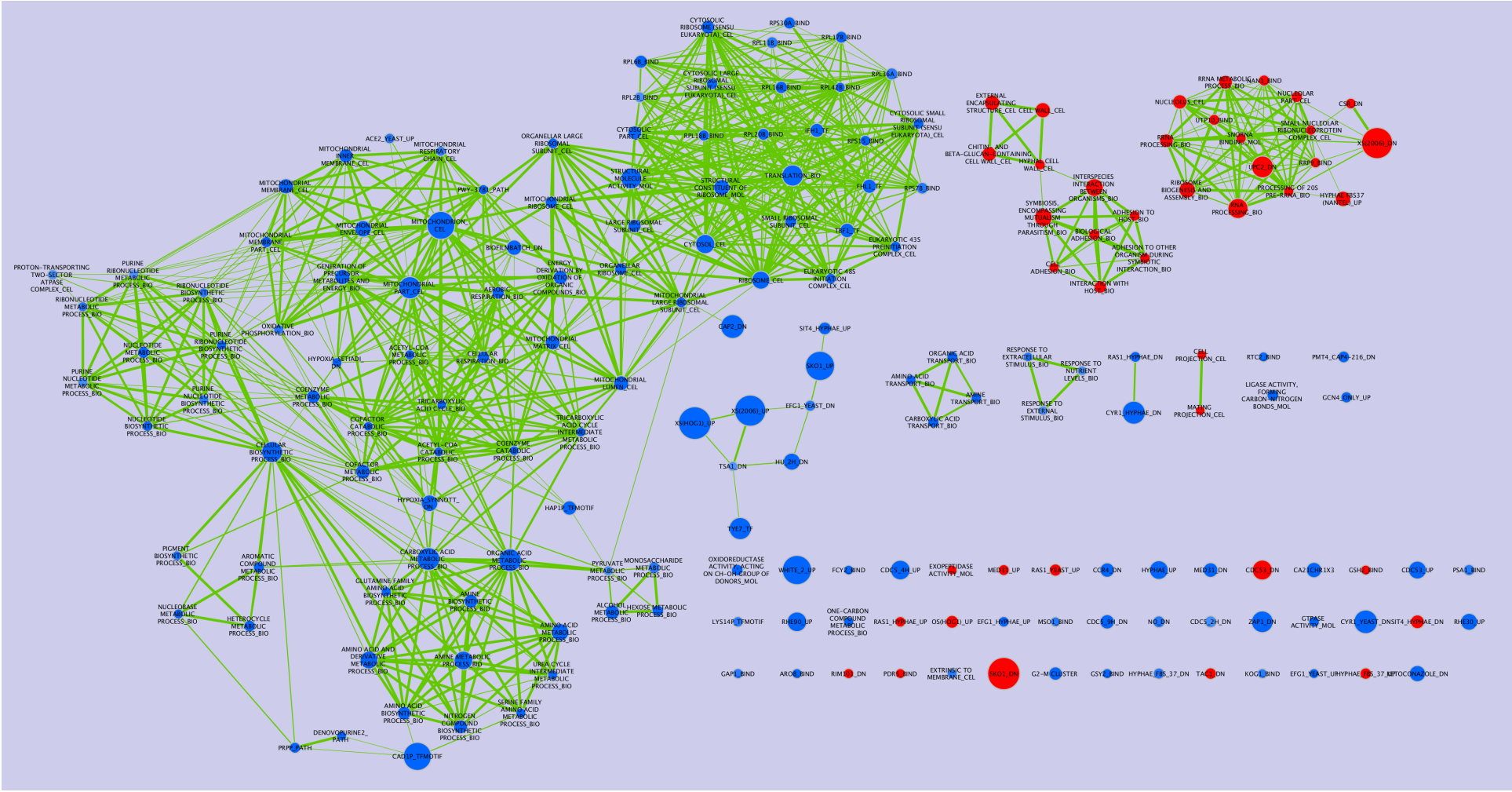
Figure S3


CYTOSOLIC RIBOSOMAL SUBUNIT (SENSU EUKARYOTA)_CEL
RPS3A_BIND
RPL11B_BIND
RPL17B_BIND
RPL10B_BIND
RPL2B_BIND
RPL13A_BIND
CYTOSOLIC LARGE RIBOSOMAL SUBUNIT (SENSU EUKARYOTA)_CEL
RPL18_BIND
RPL12B_BIND
CYTOSOLIC SMALL RIBOSOMAL SUBUNIT (SENSU EUKARYOTA)_CEL
CYTOSOLIC PART_CEL
RPL16B_BIND
RPL20B_BIND
IFH1_TF
RPS13_BIND
ORGANELLAR LARGE RIBOSOMAL SUBUNIT_CEL
ACE2_YEAST_UP
MITOCHONDRIAL INNER MEMBRANE_CEL
MITOCHONDRIAL RESPIRATORY CHAIN_CEL
STRUCTURAL MOLECULE ACTIVITY_MOL
STRUCTURAL CONSTITUENT OF RIBOSOME_MOL
TRANSLATION_BIO
FHL1_TF
RPS7B_BIND
MITOCHONDRIAL MEMBRANE_CEL
PWY-781_PATH
MITOCHONDRIAL RIBOSOME_CEL
MITOCHONDRION_CEL
LARGE RIBOSOMAL SUBUNIT_CEL
SMALL RIBOSOMAL SUBUNIT_CEL
IFH1_TF
MITOCHONDRIAL ENVELOPE_CEL
BIOGENESIS_DN
CYTOSOL_CEL
EUKARYOTIC 43S PREINITIATION COMPLEX_CEL
MITOCHONDRIAL MEMBRANE PART_CEL
ENERGY DERIVATION BY OXIDATION OF ORGANIC COMPOUNDS_BIO
ORGANELLAR RIBOSOME_CEL
RIBOSOME_CEL
EUKARYOTIC 48S INITIATION COMPLEX_CEL
PROTON-TRANSPORTING TWO-SECTOR ATPASE COMPLEX_CEL
PURINE RIBONUCLEOTIDE METABOLIC PROCESS_BIO
GENERATION OF PRECURSOR METABOLITES AND ENERGY_BIO
MITOCHONDRIAL PART_CEL
AEROBIC RESPIRATION_BIO
MITOCHONDRIAL LARGE RIBOSOMAL SUBUNIT_CEL
RIBONUCLEOTIDE BIOSYNTHETIC PROCESS_BIO
RIBONUCLEOTIDE METABOLIC PROCESS_BIO
OXIDATIVE PHOSPHORYLATION_BIO
CAP2_DN
SIT4_HYPHAE_UP
NUCLEOTIDE METABOLIC PROCESS_BIO
PURINE RIBONUCLEOTIDE BIOSYNTHETIC PROCESS_BIO
HYPOXIA_STEIDL_DN
ACETYL-COA METABOLIC PROCESS_BIO
CELLULAR RESPIRATION_BIO
MITOCHONDRIAL MATRIX_CEL
SKO1_UP
PURINE NUCLEOTIDE METABOLIC PROCESS_BIO
MITOCHONDRIAL LUMEN_CEL
COENZYME METABOLIC PROCESS_BIO
PURINE NUCLEOTIDE BIOSYNTHETIC PROCESS_BIO
TRICARBOXYLIC ACID CYCLE_BIO
COFACTOR CATABOLIC PROCESS_BIO
XS(2006)_UP
EFG1_YEAST_DN
XS(HOG1)_UP
NUCLEOTIDE BIOSYNTHETIC PROCESS_BIO
TRICARBOXYLIC ACID CYCLE INTERMEDIATE METABOLIC PROCESS_BIO
CELLULAR BIOSYNTHETIC PROCESS_BIO
ACETYL-COA CATABOLIC PROCESS_BIO
COENZYME CATABOLIC PROCESS_BIO
TSA1_DN
HU_2H_DN
COFACTOR METABOLIC PROCESS_BIO
HYPOXIA_SYNNOTT_DN
HAP1P_TFMOTIF
TYE7_TF
PIGMENT BIOSYNTHETIC PROCESS_BIO
AROMATIC COMPOUND METABOLIC PROCESS_BIO
CARBOXYLIC ACID METABOLIC PROCESS_BIO
ORGANIC ACID METABOLIC PROCESS_BIO
PYRUVATE METABOLIC PROCESS_BIO
MONOSACCHARIDE METABOLIC PROCESS_BIO
GLUTAMINE FAMILY AMINO ACID BIOSYNTHETIC PROCESS_BIO
AMINE BIOSYNTHETIC PROCESS_BIO
NUCLEOBASE METABOLIC PROCESS_BIO
HETEROCYCLE METABOLIC PROCESS_BIO
ALCOHOL METABOLIC PROCESS_BIO
HEXOSE METABOLIC PROCESS_BIO
AMINO ACID METABOLIC PROCESS_BIO
AMINO ACID AND DERIVATIVE METABOLIC PROCESS_BIO
AMINE METABOLIC PROCESS_BIO
UREA CYCLE INTERMEDIATE METABOLIC PROCESS_BIO
AMINO ACID BIOSYNTHETIC PROCESS_BIO
NITROGEN COMPOUND BIOSYNTHETIC PROCESS_BIO
SERINE FAMILY AMINO ACID METABOLIC PROCESS_BIO
DENOVOPURINE2_PATH
PRPP_PATH
CAD1P_TFMOTIF
EXTERNAL ENCAPSULATING STRUCTURE_CEL
CELL WALL_CEL
CHITIN- AND BETA-GLUCAN-CONTAINING CELL WALL_CEL
HYPHAE CELL WALL_CEL
INTERSPECIES INTERACTION BETWEEN ORGANISMS_BIO
SYMBIOSIS, ENCOMPASSING MUTUALISM THROUGH PARASITISM_BIO
ADHESION TO HOST_BIO
BIOLOGICAL ADHESION_BIO
ADHESION TO OTHER ORGANISM DURING SYMBIOTIC INTERACTION_BIO
CELL ADHESION_BIO
INTERACTION WITH HOST_BIO
RRNA METABOLIC PROCESS_BIO
NAN1_BIND
NUCLEOLAR PART_CEL
NUCLEOLUS_CEL
CSE_DN
UTP10_BIND
SMALL NUCLEOLAR RIBONUCLEOPROTEIN COMPLEX_CEL
RRN3 PROCESSING_BIO
SNR/NHA BINDING_MOL
XS(2000)_DN
UPF2_DN
KRR1_BIND
RIBOSOME BIOGENESIS AND ASSEMBLY_BIO
PROCESSING OF 20S PRE-RNA_BIO
HYPHAE_RBS37 (NANTEL)_UP
RNA PROCESSING_BIO
ORGANIC ACID TRANSPORT_BIO
RESPONSE TO EXTRACELLULAR STIMULUS_BIO
RESPONSE TO NUTRIENT LEVELS_BIO
RAS1_HYPHAE_DN
CEN PROJECTION_CEL
RTC2_BIND
PMT4_CAP6-216_DN
AMINO ACID TRANSPORT_BIO
AMINE TRANSPORT_BIO
LIGASE ACTIVITY, FORMING CARBON-NITROGEN BONDS_MOL
GCN4_ONLY_UP
RESPONSE TO EXTERNAL STIMULUS_BIO
CYR1_HYPHAE_DN
MATING PROJECTION_CEL
CARBOXYLIC ACID TRANSPORT_BIO
OXIDOREDUCTASE ACTIVITY, ACTING ON CH-OH GROUP OF DONORS_MOL
WHITE_2_UP
FCY2_BIND
CDC5_4H_UP
EXOPEPTIDASE ACTIVITY_MOL
MED31_UP
RAS1_YEAST_UP
CCN4_DN
HYPHAE_UP
MED31_DN
CDC53_DN
CA2(CHR1X3
GSH2_BIND
CDC53_UP
PSA1_BIND
LYS14P_TFMOTIF
RHE90_UP
ONE-CARBON COMPOUND METABOLIC PROCESS_BIO
RAS1_HYPHAE_UP
OS(HOG1)_UP
EFG1_HYPHAE_UP
MSO1_BIND
CDC5_9H_DN
NO_DN
CDC5_2H_DN
ZAP1_DN
GTPASE ACTIVITY_MOL
CYR1_YEAST_DN
SIT4_HYPHAE_DN
RHE30_UP
GAP1_BIND
AROC_BIND
RIM101_DN
PDR5_BIND
EXTRINSIC TO MEMBRANE_CEL
SKO1_DN
G2-M CLUSTER
GSY2_BIND
HYPHAE_FBS_37_DN
TAC1_DN
KOG1_BIND
EFG1_YEAST_UP
HYPHAE_FBS_37_UP
ITRACONAZOLE_DN