54 notable communities found by walktrap clustering with Gcn5 Targets

Peptide:N-glycanase–Rad23 complex|1|2;CDC48–CDC23–UFD2 complex|1|2;Nucleotide excision repair factor 2 complex|1|2

Ribosome quality control complex|1|6;CDC48–NPL4–UFD1 AAA ATPase complex|1|5;CDC48–NPL4–VMS1 AAA ATPase complex|1|3;CDC48–CDC23–UFD2 complex|1|1

HMC complex|6|7;HAP1 transcriptional repressor complex, SSA1 variant|3|4;HAP1 transcriptional repressor complex, SSA2 variant|3|4

TOM40 mitochondrial outer membrane translocase holocomplex|6|7;TOM40 mitochondria outer membrane translocase core complex|1|5

INO80 chromatin remodeling complex|2|15;NuA4 histone acetyltransferase complex|1|15;Swr1 chromatin remodelling complex|1|14;NuA3 histone acetyltransferase complex|1| 5;Transcription factor TFIIF complex|1| 3;SWI/SNF chromatin remodelling complex|1| 1;General transcription factor complex

Box H/ACA ribonucleoprotein complex;Transcription factor 3 core complex|4|5

TREX complex|2|7

RSC complex variant RSC2|1|16;RSC complex variant RSC1|1|16

mRNA cleavage factor complex CF I|1|9

RAP1–GCR1–GCR2 transcription activation complex|1|3;Telosome|1|2;RAP1–GCR1 transcription activation complex|1|2

SAGA complex|1|20;SLIK (SAGA–like) complex|0| 0

Trehalose–6–phosphate synthase/phosphatase variant 2|1|3

Exomer complex|1|5

Spindle pole body central plaque complex|1|4;Myosin class I complex, MYO5 variant|1|2;Myosin class I complex, MYO3 variant|1|2;Myosin class V complex, MYO4 variant|1|1;Myosin class V complex, MYO2 variant|1|1

PMT1–PMT2 dolichyl–phosphate–mannose–protein mannosyltransferase complex|1|3;PMT1–PMT3 dolichyl–phosphate–mannose–protein

cAMP–dependent protein kinase complex variant 4|1|3;cAMP–dependent protein kinase complex variant 5|1|3;cAMP–dependent protein kinase complex variant 6|1|3;cAMP–dependent protein kinase complex variant 1|1|2;cAMP–dependent protein kinase

Myosin class II complex|1|3;Myosin class V complex, MYO4 variant|1|2;MIH complex|1|2;Myosin class V complex, MYO2 variant|1|1

COPI vesicle coat complex|4|7

PAN1 actin cytoskeleton–regulatory complex|1|3;SLAC complex|1|2

RENT complex|3|3

Luminal surveillance complex|1|3

Elongation Factor eEF1 complex, variant TEF4|3|3;Elongation Factor eEF1 complex, variant CAM1|3|3

SET3C histone deacetylase complex|1|7

...ate dehydrogenase complex|3|5;Glycine decarboxylase multienzyme complex|1| ...;Mitochondrial oxoglutarate dehydrogenase complex|1|4

...|4|U7|U5 snRNP complex|2|28;U4/U6 snRNP|2|17;U6 snRNP complex|2| 8;LSM1–7 PAT1 complex|2| 8;LSM2–8 complex|2| 7;LSM1–7 complex|2| 7;LSM2–7 complex|2| 6

ESCRT–III complex|1|4

1,3–beta–D–glucan synthase complex, FKS1–RHO1 variant|1|2;1,3–beta–D–glucan synthase complex, FKS2–RHO1 variant|1|2;1,3–beta–D–glucan synthase complex, FKS3–RHO1 variant|1|2

PAT1 complex|2|5

RNA polymerase I upstream activating factor complex|1|6;Nucleosome, variant HTA2–HTB2|1|4;Nucleosome, variant HTA2–HTB1|1|4;Nucleosome, variant HTA1–HTB1|1|4;Nucleosome, variant HTZ1–HTB1|1|4;Nucleo...

TIM2 mitochondrial inner membrane twin–pore carrier translocase complex|2|6;TIM9–TIM10–TIM12 mitochondrial intermembrane space protein transporter complex|2|1...

Small ribosomal subunit processome, variant 1|8|47;Small ribosomal subunit processome, variant 2|8|47;Small ribosomal subunit processome, variant 3|8|47;Box C/D snoRNP complex|3| 4;U4/U6.U5 tri–snRNP complex|1| 1;U4 snRNP|1| 1;U4/U6 snRNP|1...

GET4–GET5 transmembrane domain recognition complex|3|5

Nucleotide excision repair factor 4 complex|1|3 40S cytosolic small ribosomal subunit|8|9

SDS22–GLC7–YPI1 phosphatase complex|1|3;SDS22–GLC7 phosphatase complex|1|2

SCF–Ydr131c ubiquitin ligase complex|1|5;SCF–Cdc4 ubiquitin ligase complex|1|5;SCF–Grr1 ubiquitin ligase complex|1|5;SCF–Mdm30 ubiquitin ligase complex|1|5;SCF–Ufo1 ubiquitin ligase complex|1|5;SCF–Das1 ubiquitin ligase complex|1|5;SCF–Met30 ubiquitin ligase complex|1|5;SCF–Dia2 ubiquitin ligase complex|1|5;SCF–Ylr352w ubiquitin ligase complex

Prefoldin co–chaperone complex|4|6

TIM23 mitochondrial inner membrane pre–sequence translocase complex, motor variant|3|9;TIM23 mitochondrial inner membrane pre–sequence translocase complex, sort variant

...complex|1|2;...TRM112 methyltransferase ...;...TRM112 ...;tRNA (m2G10) methyltransferase complex|0|0

Mitochondrial respiratory chain complex III|2|10

Ribonucleoside–diphosphate reductase variant 2|2|4;Ribonucleoside–diphosphate reductase variant 1|2|3

Nuclear pore complex|3|32

Chaperonin–containing T–complex|4|8

Vacuolar proton translocating ATPase complex, vacuole variant|7|14;Vacuolar proton translocating ATPase complex, Golgi variant|7|14

COPII vesicle coat complex|3|4;SEC23–GST1 COPII cargo recruitment complex|1|2

Mitochondrial proton–transporting ATP synthase complex|2|17

26S Proteasome complex|13|33

Vesicular SNARE complex SSO2–SEC9–SNC2|1|3;Vesicular SNARE complex SSO2–SEC9–SNC1|1|3

DNA–directed RNA polymerase III complex|4|17;DNA–directed RNA Polymerase I complex|4|14;DNA–directed RNA polymerase II complex|2|10

Gin4 serine/threonine kinase complex|2|8;Septin complex|1|5