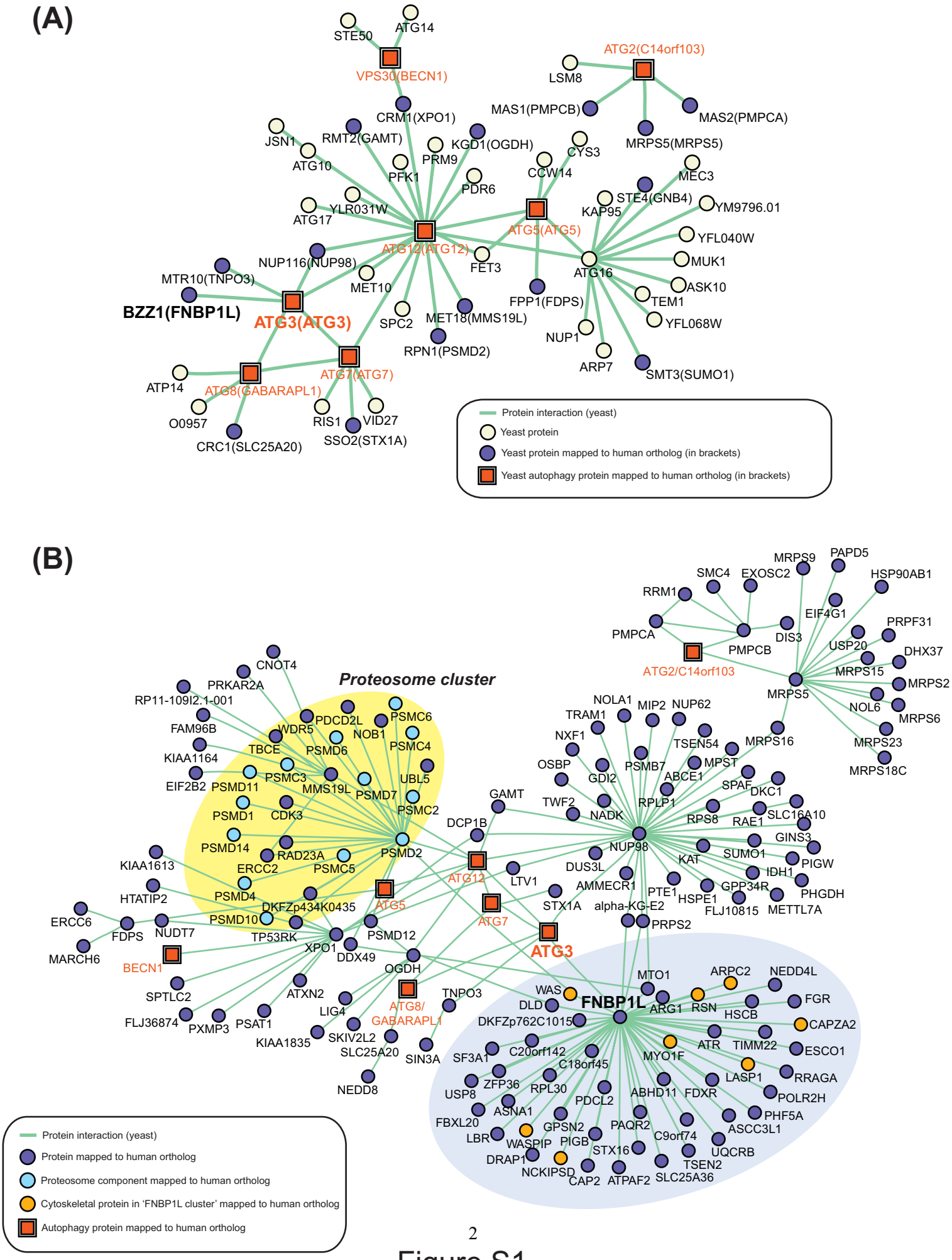
(A)
STE50
ATG14
VPS30(BECN1)
CRM1(XPO1)
ATG2(C14orf103)
LSM8
MAS1(PMPCB)
MAS2(PMPCA)
MRPS5(MRPS5)
JSN1
RMT2(GAMT)
KGD1(OGDH)
CYS3
ATG10
PRM9
PFK1
CCW14
MEC3
PDR6
STE4(GNB4)
YM9796.01
ATG17
YLR031W
KAP95
ATG5(ATG5)
YFL040W
ATG12(ATG12)
FET3
MUK1
NUP116(NUP98)
ATG16
MTR10(TNPO3)
MET10
ASK10
FPP1(FDPS)
TEM1
BZZ1(FNBP1L)
ATG3(ATG3)
SPC2
MET18(MMS19L)
YFL068W
NUP1
RPN1(PSMD2)
ARP7
ATG7(ATG7)
SMT3(SUMO1)
ATP14
ATG8(GABARAPL1)
O0957
RIS1
VID27
SSO2(STX1A)
CRC1(SLC25A20)
Protein interaction (yeast)
Yeast protein
Yeast protein mapped to human ortholog (in brackets)
Yeast autophagy protein mapped to human ortholog (in brackets)
(B)
MRPS9
PAPD5
SMC4
EXOSC2
HSP90AB1
RRM1
EIF4G1
PRPF31
PMPCA
DIS3
PMPCB
USP20
DHX37
ATG2/C14orf103
MRPS15
MRPS2
MRPS5
NOL6
MRPS6
MRPS23
MRPS18C
CNOT4
Proteosome cluster
PRKAR2A
RP11-109I2.1-001
PDCD2L
PSMC6
FAM96B
WDR5
NOB1
PSMC4
TBCE
PSMD6
KIAA1164
PSMC3
UBL5
EIF2B2
PSMD11
MMS19L
PSMD7
PSMD1
CDK3
PSMC2
PSMD14
RAD23A
PSMD2
KIAA1613
ERCC2
PSMC5
HTATIP2
PSMD4
DKFZp434K0435
ERCC6
PSMD10
FDPS
NUDT7
TP53RK
PSMD12
MARCH6
XPO1
DDX49
BECN1
OGDH
SPTLC2
ATXN2
TNPO3
FLJ36874
PXMP3
PSAT1
LIG4
KIAA1835
SKIV2L2
SLC25A20
SIN3A
NEDD8
ATG8/
GABARAPL1
ATG12
ATG5
ATG7
ATG3
NOLA1
MIP2
NUP62
TRAM1
NXF1
TSEN54
MRPS16
OSBP
PSMB7
MPST
GDI2
ABCE1
SPAF
DKC1
GAMT
TWF2
RPLP1
NADK
RPS8
RAE1
SLC16A10
DCP1B
GINS3
NUP98
SUMO1
PIGW
DUS3L
KAT
IDH1
LTV1
AMMECR1
PTE1
GPP34R
PHGDH
STX1A
alpha-KG-E2
HSPE1
FLJ10815
METTL7A
PRPS2
FNBP1L
MTO1
ARPC2
NEDD4L
WAS
DLD
ARG1
RSN
FGR
HSCB
DKFZp762C1015
CAPZA2
ATR
TIMM22
ESCO1
MYO1F
SF3A1
C20orf142
C18orf45
LASP1
RRAGA
ZFP36
USP8
RPL30
ABHD11
FDXR
POLR2H
PDCL2
ASNA1
PHF5A
FBXL20
PAQR2
GPSN2
ASCC3L1
LBR
WASPIP
PIGB
C9orf74
STX16
UQCRB
DRAP1
TSEN2
NCKIPSD
CAP2
ATPAF2
SLC25A36
Protein interaction (yeast)
Protein mapped to human ortholog
Proteosome component mapped to human ortholog
Cytoskeletal protein in 'FNBP1L cluster' mapped to human ortholog
Autophagy protein mapped to human ortholog

Figure S1