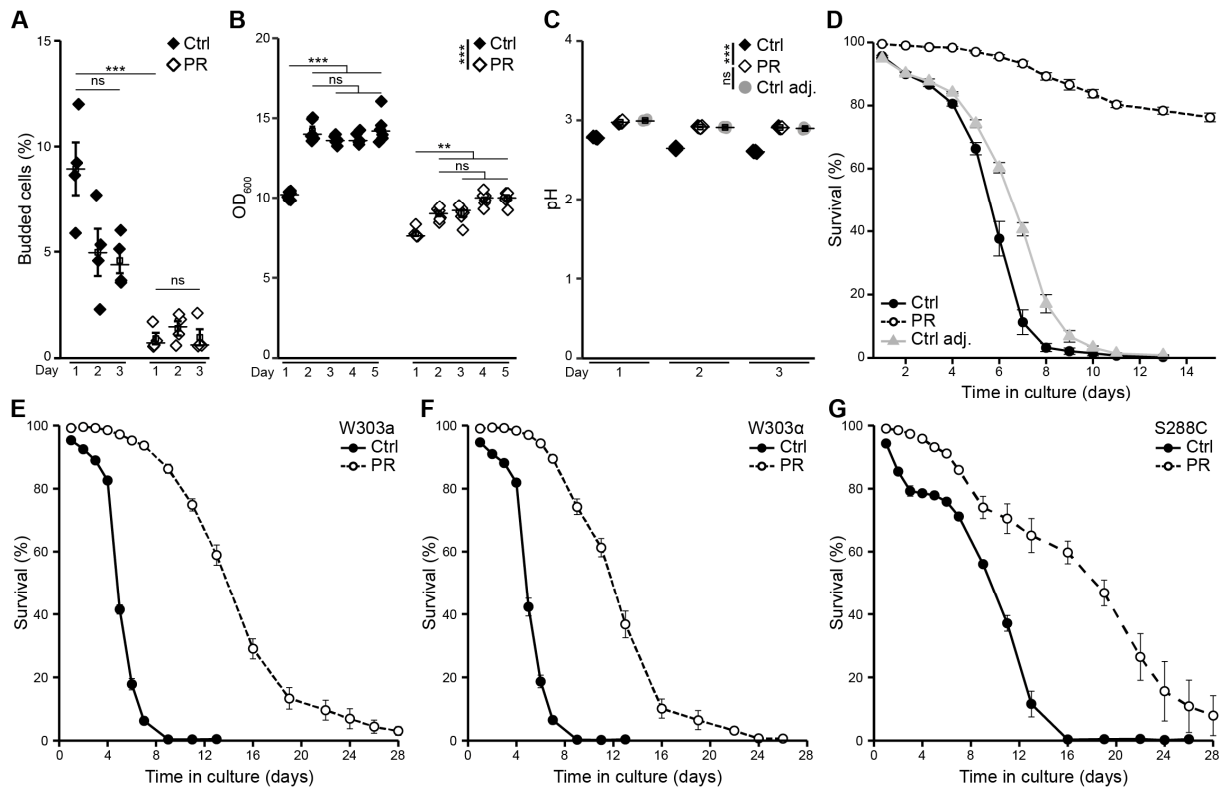
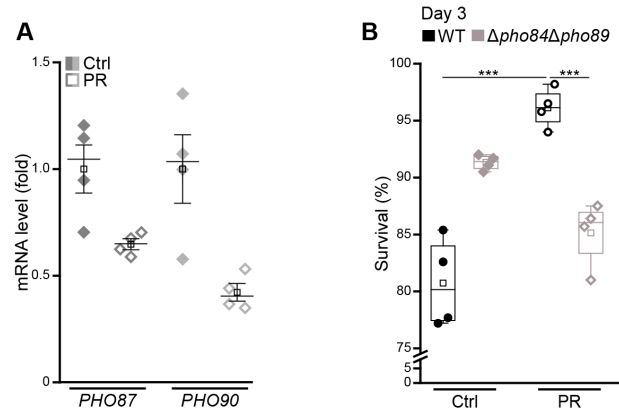


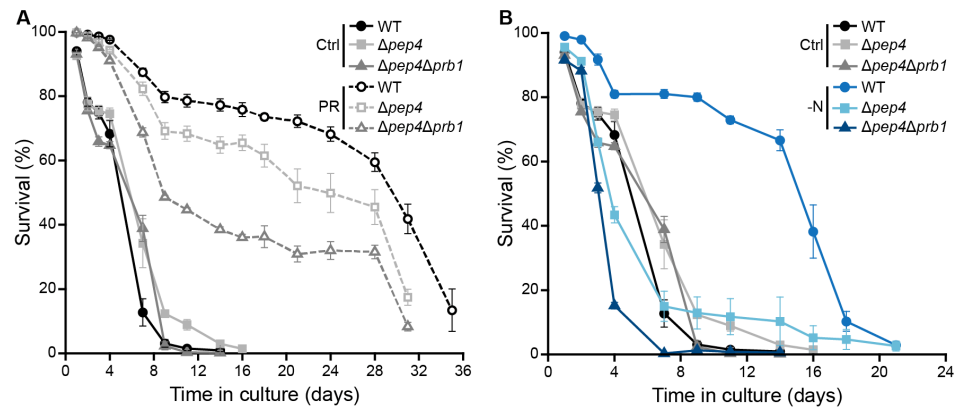
# Supplemental Material



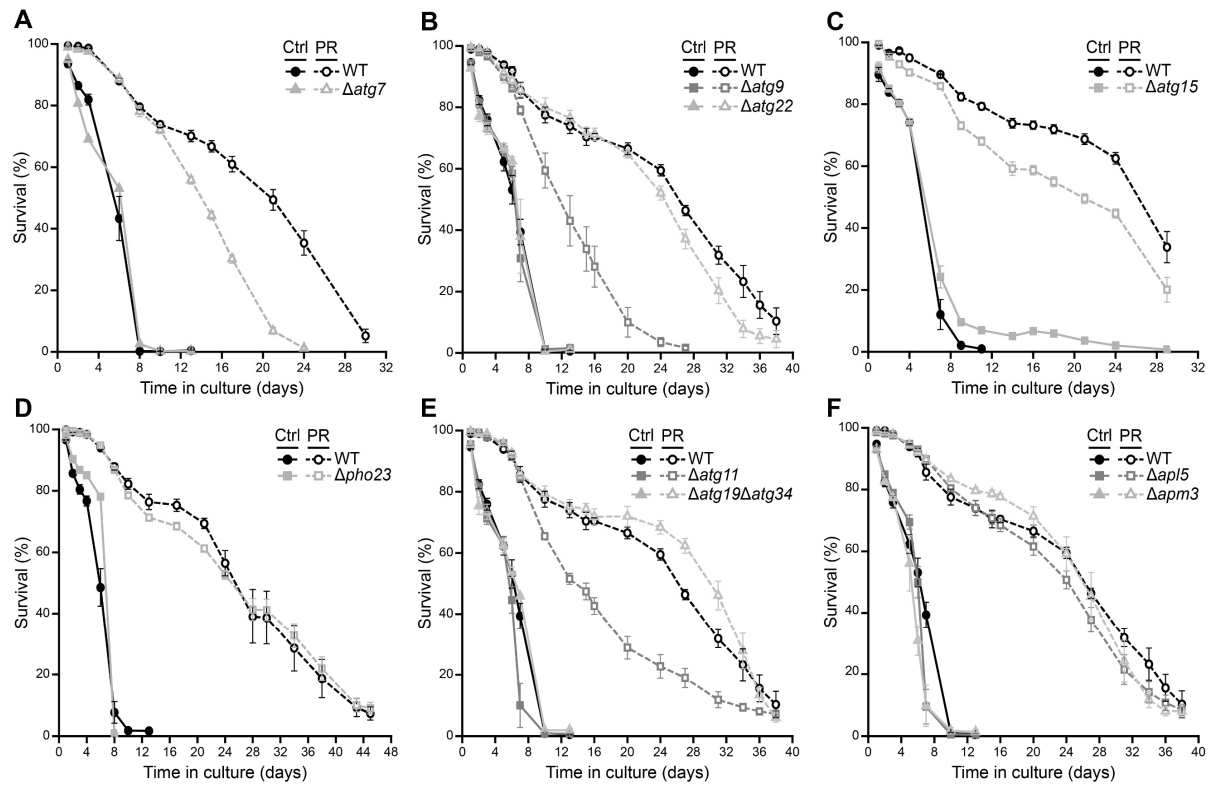
**Figure S1.** Phosphate exhaustion results in efficient entry into quiescence and longevity in different genetic backgrounds. (A) Quantification of budded WT cells cultured on standard medium (Ctrl) and phosphate-restricted medium (PR) for indicated days. Cells were manually counted, and percentage of budded cells is shown;  $n = 4$ , with at least 100 cells per experiment and condition. (B) Optical density (OD<sub>600</sub>) of WT cells cultured in Ctrl and PR media, determined at indicated time points during aging;  $n = 6$ . (C) Determination of culture media pH after WT cells have been cultured for indicated days on standard medium (Ctrl), phosphate-restricted medium (PR) and standard medium with pH adjustment to the media pH of PR cultures (Ctrl adj.). (D) Survival during chronological aging of WT cells cultured as described in (C). Survival was determined by flow cytometric analysis of PI staining;  $n = 5$ . (E-G) Survival of the yeast wild type strains W303a (E), W303α (F), and S288c (G) during chronological aging on Ctrl and PR medium. Survival was determined by flow cytometric analysis of PI staining at indicated days during aging;  $n=3-6$ . For statistical analysis, two-way ANOVA (A, B, C) was used. \*\* $p < 0.01$  and \*\*\* $p < 0.001$ ; ns not significant.



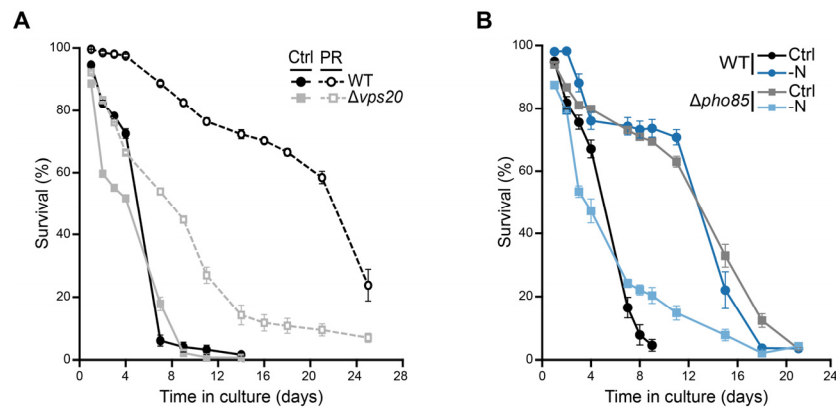
**Figure S2.** Gradual phosphate exhaustion leads to transcriptional downregulation of low-affinity phosphate transporters. **(A)** Quantification of *PHO87* and *PHO90* mRNA levels via RT-qPCR in WT cells grown for 24 h on standard (Ctrl) or phosphate restriction (PR) medium. Comparative CT method ( $\Delta\Delta$ CT method) was used to calculate the relative gene expression using *UBC6* as house-keeping gene;  $n = 4$ . **(B)** Survival of WT cells and cells lacking both high-affinity phosphate transporters ( $\Delta pho84\Delta pho89$ ) on day 3 of chronological aging on Ctrl or PR medium, corresponding to the survival curve shown in Fig. 1L;  $n = 4$ . For statistical analysis, one-way ANOVA followed by Tukey post hoc for multiple comparisons **(B)** was used. \*\*\* $p < 0.001$ .



**Figure S3.** Simultaneous lack of the vacuolar proteases Pep4 and Prb1 impairs phosphate restriction-induced longevity. Survival during chronological aging, determined via flow cytometric quantification of PI staining, of WT,  $\Delta pep4$  and  $\Delta pep4\Delta prb1$  cells grown under different nutrient conditions: phosphate restriction (PR), nitrogen starvation (-N) and standard conditions (Ctrl). For clarity, survival curves are shown in two separate graphs for phosphate restriction **(A)** and nitrogen starvation **(B)**;  $n \geq 4$ .



**Figure S4.** The core machinery of autophagy is required for full lifespan extension via phosphate restriction. Survival during chronological aging, determined via flow cytometric quantification of PI staining, of WT,  $\Delta atg7$  (A),  $\Delta atg9$ ,  $\Delta atg22$  (B),  $\Delta atg15$  (C),  $\Delta pho23$  (D),  $\Delta atg11$ ,  $\Delta atg19\Delta atg34$  (E), and  $\Delta apl5$  and  $\Delta apm3$  (F) cells grown on standard (Ctrl) or phosphate restriction (PR) medium;  $n \geq 4$ .



**Figure S5.** The MVB pathway and Pho85 are critical for survival upon nutrient limitation. Survival during chronological aging, determined via flow cytometric quantification of PI staining, of WT and  $\Delta vps20$  cells grown on standard (Ctrl) or phosphate-restricted (PR) medium;  $n \geq 4$  (A), as well as of WT and  $\Delta pho85$  cells grown on Ctrl or -N media;  $n = 4$  (B).

**Table S1.** Yeast strains used in this study.

Strain	Genotype	Source
BY4742	MAT $\alpha$ , <i>his3</i> 1, <i>leu2</i> 0, <i>lys2</i> 0, <i>ura3</i> 0	Euroscarf
W303a	MATa, <i>ade2-1</i> , <i>ura3-1</i> , <i>his3-11</i> , <i>trp1-1</i> , <i>leu2-3</i> , <i>leu2-112</i> , <i>can1-100</i>	
W303 $\alpha$	MAT $\alpha$ , <i>ade2-1</i> , <i>ura3-1</i> , <i>his3-11</i> , <i>trp1-1</i> , <i>leu2-3</i> , <i>leu2-112</i> , <i>can1-100</i>	
S288C	MAT $\alpha$ , <i>SUC2</i> , <i>gal2</i> , <i>mal2</i> , <i>mel</i> , <i>flo1</i> , <i>flo8-1</i> , <i>hap1</i> , <i>ho</i> , <i>bio1</i> , <i>bio6</i>	
BY4742 Vph1 <sup>mCherry</sup>	BY4742 <i>VPH1</i> -mCherry-natNT2	This study
<i>pho84</i>	BY4742 <i>pho84</i> ::hphNT1	This study
<i>pho89</i>	BY4742 <i>pho89</i> ::hphNT1	This study
<i>pho84 pho89</i>	BY4742 <i>pho84</i> ::hphNT1, <i>pho89</i> ::natNT2	This study
<i>pho87</i>	BY4742 <i>pho87</i> ::hphNT1	Euroscarf
<i>pho90</i>	BY4742 <i>pho90</i> ::hphNT1	Euroscarf
<i>vtc1</i>	BY4742 <i>vtc1</i> ::hphNT1	This study
<i>vtc2</i>	BY4742 <i>vtc2</i> ::hphNT1	This study
<i>vtc3</i>	BY4742 <i>vtc3</i> ::hphNT1	This study
<i>vtc4</i>	BY4742 <i>vtc4</i> ::hphNT1	This study
<i>vtc5</i>	BY4742 <i>vtc5</i> ::natNT2	This study
<i>vtc4 vtc5</i>	BY4742 <i>vtc4</i> ::hphNT1, <i>vtc5</i> ::natNT2	This study
BY4742 <sup>GFP</sup> Atg8	BY4742 pATG8 ::natNT2-pATG8-EGFP	This study
BY4742 <sup>GFP</sup> Atg8 Vph1 <sup>Redstar</sup>	BY4742 pATG8 ::natNT2-pATG8-EGFP, <i>VPH1</i> -Red- star-kanMX	This study
<i>pep4</i>	BY4742 <i>pep4</i> ::hphNT1	This study
<i>pep4 prb1</i>	BY4742 <i>pep4</i> ::hphNT1, <i>prb1</i> ::kanMX	This study
<i>atg1</i>	BY4742 <i>atg1</i> ::hphNT1	This study
<i>atg4</i>	BY4742 <i>atg4</i> ::hphNT1	This study
<i>atg7</i>	BY4742 <i>atg7</i> ::hphNT1	This study
<i>atg8</i>	BY4742 <i>atg8</i> ::hphNT1	This study
<i>atg9</i>	BY4742 <i>atg9</i> ::hphNT1	This study
<i>atg11</i>	BY4742 <i>atg11</i> ::hphNT1	This study
<i>atg15</i>	BY4742 <i>atg15</i> ::hphNT1	This study
<i>atg22</i>	BY4742 <i>atg22</i> ::hphNT1	This study
<i>atg19 atg34</i>	BY4742 <i>atg19</i> ::hphNT1, <i>atg34</i> ::kanMX	This study
<i>atg39</i>	BY4742 <i>atg39</i> ::hphNT1	This study
<i>apl5</i>	BY4742 <i>apl5</i> ::hphNT1	This study
<i>apm3</i>	BY4742 <i>apm3</i> ::hphNT1	This study
<i>pho23</i>	BY4742 <i>pho23</i> ::hphNT1	This study
BY4742 Sna3 <sup>GFP</sup>	BY4742 <i>SNA3</i> -EGFP-kanMX	This study
BY4742 Sna3 <sup>GFP</sup> Vph1 <sup>mCherry</sup>	BY4742 <i>SNA3</i> -EGFP-KanMX, <i>VPH1</i> -mCherry-natNT2	This study
<i>vps24</i>	BY4742 <i>vps24</i> ::hphNT1	This study
<i>vps24 Sna3<sup>GFP</sup></i>	BY4742 <i>vps24</i> ::hphNT1, <i>SNA3</i> -yeGFP-kanMX	This study
<i>vps24 Sna3<sup>GFP</sup></i> Vph1 <sup>mCherry</sup>	BY4742 <i>vps24</i> ::hphNT1, <i>SNA3</i> -yeGFP-kanMX, <i>VPH1</i> - mCherry-natNT2	This study
<i>vps4</i>	BY4742 <i>vps4</i> ::hphNT1	This study
<i>vps20</i>	BY4742 <i>vps20</i> ::hphNT1	This study
<i>atg1 vps24</i>	BY4742 <i>atg1</i> ::kanMX, <i>vps24</i> ::kanMX	Ralf Braun
<i>mon1</i>	BY4742 <i>mon1</i> ::natNT2	This study
<i>ypt7</i>	BY4742 <i>ypt7</i> ::hphNT1	This study
<i>rim15</i>	BY4742 <i>rim15</i> ::hphNT1	This study

<i>pho80</i>	BY4742 <i>pho80</i> ::hphNT1	This study
<i>pho81</i>	BY4742 <i>pho81</i> ::hphNT1	This study
<i>pho85</i>	BY4742 <i>pho85</i> ::hphNT1	This study
BY4742 <sup>GFP</sup> Atg8	BY4742 pATG8 ::natNT2-pATG8-EGFP,	This study
<i>pho85</i>	<i>pho85</i> ::hphNT1	
BY4742 <sup>GFP</sup> Atg8	BY4742 pATG8 ::natNT2-pATG8-EGFP,	This study
<i>pho85</i> Vph1 <sup>Redstar</sup>	<i>pho85</i> ::hphNT1, <i>VPH1</i> -Redstar-kanMX	

**Table S2.** Oligonucleotides used for gene deletion, chromosomal tagging and RT-qPCR.

Modification	Oligonucleotides	PCR template
Deletion and tagging of genes		
C-terminal tagging of <i>VPH1</i>	5'-GAAGTACTTAAATGTTTCGCTTTTTTTTAAAGTCCTCAAAAT TTAATCGATGAATTCGAGCTCG-3' 5'-GACATGGAAGTCGCTGTTGCTAGTGCAAGCTCTTCCGCTTC AAGCCGTACGCTGCAGGTCGAC-3'	pYM42-mCherry-natNT2 [1] pYM38 [2]
Control PCR <i>VPH1</i> tagging	5'-CGAGCTCGAATTCATCGAT-3' 5'-GTATTCGAGGCCAATACTTG-3'	
Deletion of <i>PHO84</i>	5'-CGTAGATCACCAGGGCACACAACAAACAAAACCTCCACGAA TACAATCCAA ATG CGTACGCTGCAGGTCGAC-3' 5'-GTATTATTTGTTCTAGTTTACAAGTTTTAGTGCATCTTTG AG- GCTTTTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>PHO84</i> deletion	5'-CGTTCCTCTCACTGCCGCAC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>PHO89</i>	5'-AATACTGGAGCGTATCTAATCGAATCAATATAAACAAAGATT AAGCAAAA ATG CGTACGCTGCAGGTCGAC-3' 5'-TACCTAATGTTGTAACAACTG TATTCATAACCGAAACACATT ATGATTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2] pFA6a-natNT2 [2]
Control PCR <i>PHO89</i> deletion	5'-GAATATACCGTACAAAGGAG-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>VTC1</i>	5'-CGATTTACTACATTATCGAATACGATTAACACTACGCC AGAT- TTCCACAATATGCGTACGCTGCAGGTCGAC-3' 5'-GAAAATACAGTTTGTGCGTAACCCACGCTTACGATATT GGAATTACAATTTCAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>VTC1</i> deletion	5'-GGACCAGTTTATAAGAAGGT-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>VTC2</i>	5'-GTAGAAAGAACGACTACACCTCAACATAACGACACTT TTTT- GACATG CGTACGCTGCAGGTCGAC-3' 5'-GCTACACATTACAAACATAAAAAACACATGGTCTCAGTAGA TA- GAGTACATATTCTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>VTC2</i> deletion	5'-CTGGAAGGTATTTTCATCAG-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>VTC3</i>	5'-GGCTATTAGAGCGAACAGCAGAATTTGTCCTTGGTTTTCA GAG- TTTGAAA ATGCGTACGCTGCAGGTCGAC-3' 5'-GATAGTAACTGGTACTTGTGTAATATATGTGTATATAAAAA AT- ATACATGTTCTTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>VTC3</i> deletion	5'-GGATGTTAATGAGTAACTTG-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>VTC4</i>	5'-GCTAACAATCAAATCGGCCAATAAAAGAGCATAACAAGGC AG- GAACAGCTATGCGTACGCTGCAGGTCGAC-3'	pFA6a-hphNT1 [2]

Control PCR <i>VTC4</i> deletion	5'-CTAATATGATTATTACTTAATTATACAGTAAAAAAACA CGCTGTGTATTCAATCGATGAATTCGAGCTCG-3' 5'-TGTAAGCCAGTACCACAAG-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>VTC5</i>	5'-TATCTCCATCTCAGCCCTAGGTAAGGATAGATACGAAGGG TTGTACAATCATGCGTACGCTGCAGGTCGAC-3' 5'-CATATTTTATATAGACTGTACACATGAAAACTTTTATC GTCCAAAATAATTAATCGATGAATTCGAGCTCG-3'	pFA6a-natNT2 [2]
Control PCR <i>VTC5</i> deletion	5'-CATTACACCCGTACGTAATC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>ATG8</i>	5'-CTAATAATTGTAAAGTTGAGAAAATCATAATAAAATAATTACTA GAGACATGCGTACGCTGCAGGTCGAC-3' 5'-CTATAATTTTCGATTTTAGATGTTAACGCTTCATTTCTTTT ATA- TAAAAGACTAATCGATGAATTCGAGCTCG-3' 5'-CTAATAATTGTAAAGTTGAGAAAATCATAATAAAATAATTAC TAGAGACATGCGTACGCTGCAGGTCGAC-3'	pFA6a-hphNT1 [2]
N-terminal tagging of <i>ATG8</i>	5'-GACTCCGCCTTCCTTTTTTCAAATGGATATTCAGACTTAA ATGTAGACTTCATCGATGAATTCTCTGTGCG-3'	pYM-pATG8 [3]
Control PCR <i>ATG8</i> deletion	5'-CTAACTGTCTCCACCGATAATG-3' 5'-GTCGACCTGCAGCGTACG-3'	
Control PCR <i>ATG8</i> tagging	5'-GAACAATAGATGGCTAATGAGTCC-3' 5'-GGTTATGGTGTCAATGTTTTGC-3'	
Deletion of <i>PEP4</i>	5'-GTGACCTAGT ATTTAATCCA AATAAAATTC AAACAAAAAC CAAACTAAC ATG CGTACGCTGCAGGTCGAC-3' 5'-CTAGATGGCAGAAAAGGATAGGGCGGAGAAGTAAGAAAAGT TTAGCTCA ATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>PEP4</i> deletion	5'-CATAATCCTGCTTGATGTGG-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>PRB1</i>	5'-ATCGCCAATAAAAAAACAACTAAACCTAATTCTAACAA GCAAAGCAGCTGAAGCTTCGTACGC-3' 5'-CGCCTAAGGAAAGAAAAAGAAAAAAAGCAGCTGAA ATTTTCTAAATGAAGAAGCATAGGCCACTAGTGGATCTG-3'	pUG6 [4]
Control PCR <i>PRB1</i> tagging	5'-GAGATGGACGAGAAACAGGG-3' 5'-GCGTACGAAGCTTCAGCTG-3'	
Deletion of <i>ATG1</i>	5'-ACCCCATATTTTCAAATCTCTTTTACAACACCAGACGAGA AATTAAGAAAATGCGTACGCTGCAGGTCGAC-3' 5'-ATATAGCAGGTCATTTGTACTTAATAAGAAAACCATATTATG CATCACTTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>ATG1</i> deletion	5'-GTAATGTAAGGAAAACCCAC-3' 5'-GTCGACCTGCAGCGTACG-3' '	
Deletion of <i>ATG4</i>	5'-GTTAGTAGATGAAGAATGGACGACTTCTTATCACGTATAGG AG- TGATATACATGCGTACGCTGCAGGTCGAC-3' 5'-GAATATATTAAAAACAAGTATATATGCTTATGAACTAGTGAA TTCCTTACACTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>ATG4</i> deletion	5'-CGCCTACATCATCAAGATCAC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>ATG7</i>	5'-ATAACTAAAGTTCATTATATTTCAACAAATATAAGATAAT CAA- GAATAAAATGCGTACGCTGCAGGTCGAC-3' 5'-CGGAAAGTGGCACCACAATATGTACCAATGCTATTATATGC AAAATATTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]

Control PCR <i>ATG7</i> deletion	5'-TGGAACGTGCAGAAAAGGAC -3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>ATG9</i>	5'-GACATTTAAGAACAGCCTGAAATATCAAAATCACGGAATTAT TAGGTTATGCGTACGCTGCAGGTCGAC-3' 5'-CAGTTATATATATAGTTATATTGGATGATGTACACGACAC AG- TCTGCCTTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>ATG9</i> deletion	5'-CGAGCTCGAATTCATCGAT-3' 5'-TGCAAAGATGGCTCATATTG-3'	
Deletion of <i>ATG11</i>	5'-GTGTACTGTTGTTGTTTCGGAAAGTACTTCTTTATTTTCT TTTATA- CATCATGCGTACGCTGCAGGTCGAC-3' 5'-GATACATAATTAATAATCTTGTCAATTTGTGACAAACGTTTAGCA CTGTTCAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>ATG11</i> deletion	5'-GCTAGCATTTCCTATATATCC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>ATG15</i>	5'-AACTGATCTAGGCATTACAATTAAGGAAACAAGGGAAAT ATTCTATTGAATGCGTACGCTGCAGGTCGAC-3' 5'-CGCATAGGCCCTAAAACAACACTAGGGTCATAATAGATGT ATGGGTCTTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>ATG15</i> deletion	5'-TATAATAAGCATAACATCGG-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>ATG22</i>	5'-GCCCTTATAAGCAATAGTTTGCTCATAACATATTCTCTA CATT- AGATATG CGTACGCTGCAGGTCGAC-3' 5'-CCTTATATATATACGTAAGCAAAGTTTTATGTAACAAA AAAAAAAAAAGAATAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>ATG22</i> deletion	5'-GTTAATCATCCTCCAGCAAAGC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>ATG19</i>	5'-GGTAATTGCGGCGGCACCTTGCTTCAGTAACGCCCAA AGGAGAG- TTCTGGTAAATGCGTACGCTGCAGGTCGAC-3' 5'-GTGTATGTGAAAAGGTACTCATTGCTGTATAAAAA TAGAGTTT- GACCTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>ATG19</i> deletion	5'-CGCGTTCCATTTGAGAAAGC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>ATG34</i>	5'-GTTAAATAAGTACTATAGCCAAAGAACTGGAAGAATA TAAAAAAGCATGCATAGGCCACTAGTGGATCTG-3' 5'-GAAACTAGTTCCTATAGGTTGAGTGTCTATCAAA AATTTACGGA- GACGCGCAGCTGAAGCTTCGTACGC-3'	pUG6 [4]
Control PCR <i>ATG34</i> deletion	5'-GGTCGTATCCTTTTCGACAGTG-3' 5'-GCGTACGAAGCTTCAGCTG-3'	
Deletion of <i>ATG39</i>	5'-GCAGTGACGATAATAGAGACTAGTAAAACAGTCGAGTTGT CGGACCTAAAATGCGTACGCTGCAGGTCGAC-3' 5'-CGTTTTTTTTTCTTTTGTTAATTTTCATTCTTCATGCTGGGTT TTGGATGATCTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>ATG39</i> deletion	5'-GCTGCATATTTGCTTTTCGCCG-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>APL5</i>	5'-CTGAAGTGGG AAAGGCAGTTGCCACAACCAGAAGAGCATA ACATATTGTCATGCGTACGCTGCAGGTCGAC-3' 5'-TACGTACGTAATCGCTGTCATTGAGAATTCACGACTTCG ATTTTTTTATCGTTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>APL5</i> deletion	5'-CGAAGCTTTTCATCCTAACGAAC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>APM3</i>	5'-GACTTGGGCA ACAACAGAGGCTGTAAACCTTACCAACCCA AC- CAAAATAG ATGCGTACGCTGCAGGTCGAC-3'	pFA6a-hphNT1 [2]

Control PCR <i>APM3</i> deletion	5'-TCATTATATTCTATTTAGTTTCGCATGGAATTTCAAGTA CACATA-TAACTAATCGATGAATTCGAGCTCG-3' 5'-CAGTAATTGCATTACAAACCTGTG-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>PHO23</i>	5'-CTTGCAGTATCACAAACAGCAACCACGTTCATAGTCTGAAATACGTACAAATGCGTACGCTGCAGGTCGAC-3' 5'-GTATTTAGTAGATATTTTCCTTAAGGCGATATTCGCCCCGAG AGC-TATTTCAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>PHO23</i> deletion	5'-ATCAGGCGCCTTAGACCGCT-3' 5'-GTCGACCTGCAGCGTACG-3'	
C-terminal tagging of <i>SNA3</i>	5'-CGATGTGCCCTTGATGGACAACAAACAACAGCTCTCTTCCGGCCGTACTCGTACGCTGCAGGTCGAC-3' 5'-CTCTATGTGTTATTTAGAACTACGGTGATAAAGAGCTCGTTCCGATCACTAATCGATGAATTCGAGCTCG-3'	pYM27 [2]
Control PCR <i>SNA3</i> tagging	5'-GTCGACCTGCAGCGTACG-3' 5'-GCACGTGATG AAGGCTCGTT-3'	
Deletion of <i>VPS24</i>	5'-ACCTTTAGTAGTTTGGGGGGCAGTTTTCTGGGCAATACAAAGTTTA CTTTTGATGCGTACGCTGCAGGTCGAC-3' 5'-CATTATTTATTCACTTATTTATTTATTTTCTTTGTACAGTCACAGTAACACTCTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>VPS24</i> deletion	5'-CGCAACTCCCACTCCCAATA-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>VPS4</i>	5'-ATGGAAGACAAAAATAAAGCAGCATAGAGTGCCTATAGTAGATGGG GTACAAATGCGTACGCTGCAGGTCGAC-3' 5'-TTTTTTATTTTCATGTACACAAGAAATCTACATTAGCACGTTAATCAATTGACTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>VPS4</i> deletion	5'-CAGTCGCGCCAACGACCAGT-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>VPS20</i>	5'-ATAGTACAGACTGCTGAATTA ACTCCACTT GGTGCTTTTG TATA-TATCGA ATG CGTACGCTGCAGGTCGAC-3' 5'-CTATTTACATTCCCTTTATTTTTTAATTTTGAAGCTACATACAGACATTCA ATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>VPS20</i> deletion	5'-GAGGGGTTCTCTAACGCTGT-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>MON1</i>	5'-GAATAAAAAAGTCTATCAAAGTACACAAACGTAGAATCAGTACATCGGAACTATGCGTACGCTGCAGGTCGAC-3' 5'-GATATATTTACAAATTAAGGAAAAAATAATAAAATAACCTCCCTGTACAAAGTTAATCGATGAATTCGAGCTCG-3'	pFA6a-natNT2 [2]
Control PCR <i>MON1</i> deletion	5'-CGAAAAGCAGTTCAAGCTCAC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>YPT7</i>	5'-CCACTTCTTATCCATATAGA AACCCCTTCTGTATCAATCAAATTAAGTGATGCGTACGCTGCAGGTCGAC-3' 5'-AAGGATACGCTATAAAGGATTACATAATAGAAGATACAATAAGTAGTACAGCTCAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>YPT7</i> deletion	5'-TATATACGGGCAGTCCGTA-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>RIM15</i>	5'-TTTCTCTTGCCTCATTGATAGAATAGATAAGCCCAGTAGAGGAAGACAGATGCGTACGCTGCAGGTCGAC-3' 5'-TTTCAAGCAAAGTTTTTATTCAGTTATTTTTTTTAATTATCTTTATCTTAAAATTTATCAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]



Control PCR <i>RIM15</i> deletion	5'-CTAATATATAGTGAATGTAGGGAGC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>PHO80</i>	5'-CAAATTTAAGTCTGCAAGCTATCATAAGACGAGGATATCC TTT- GGAGACTCATAGAAATCATGCGTACGCTGCAGGTCGAC-3' 5'-TGGAAAGTTTTAATTTTGTCTCAATCATGATTGCTTTCA TAA- TACCCACGAAAAATCATTATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>PHO80</i> deletion	5'-TTAACGAATTATGGTATTCT-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>PHO81</i>	5'-CTTTTTGAGAGAATAACCCCTTTGGAGGCAACATAGATAGA TAAACGTGCAATGCGTACGCTGCAGGTCGAC-3' 5'-AAAAGTTTTTCCTAAATAATGTATAAGATTTCAAA AC- TACATATTACAGAACTTTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>PHO81</i> deletion	5'-CTAATTGCATCAGCCTATCTTTC-3' 5'-GTCGACCTGCAGCGTACG-3'	
Deletion of <i>PHO85</i>	5'-CAGATAAGGGATATATAGCGCGGCAAAC TTAGAG- CAATACCAATGCGTACGCTGCAGGTCGAC-3' 5'-ATTATTATCATTATATATACATGGCTACGGTTTTTCGCT GAC- GGGCTGCGTTAATCGATGAATTCGAGCTCG-3'	pFA6a-hphNT1 [2]
Control PCR <i>PHO85</i> deletion	5'-GTCATTGTTGAGAGAAAGGAG-3' 5'-GTCGACCTGCAGCGTACG-3'	
<b>RT-qPCR</b>		
<i>PHO84</i>	5'-AGCTTGGGGTCAAATCTCCG-3' 5'-TCAGCACCAGAGTTTGCGTA-3'	
<i>PHO89</i>	5'-CCAGTGCTTTGATTGGCTCG-3' 5'-GCGCCAATTGTACCACCAAC-3'	
<i>PHO87</i>	5'-ATGGGGTTGGCTTCATCTGG-3' 5'-AGCGCCTACAGTTAGCCATC-3'	
<i>PHO90</i>	5'-GTGGGTACTTTTCGTCTCGCA-3' 5'-TAGGAGCAGCTTTGGGGTTG-3'	
<i>UBC6</i>	5'-GGACGTTTCAAGCCCAACAC-3' 5'-TGAGACAGACCAGCCAGGAT-3'	

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