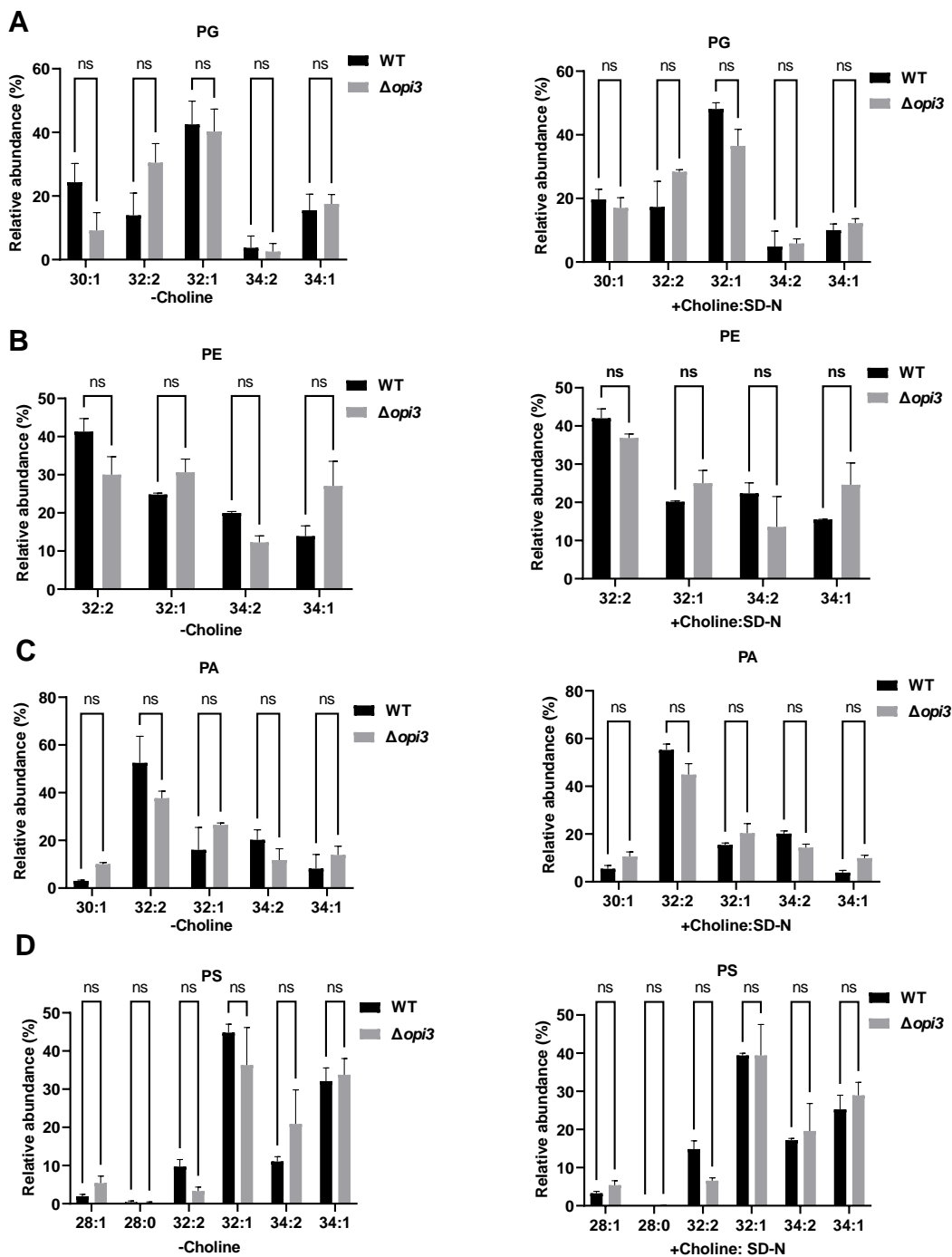
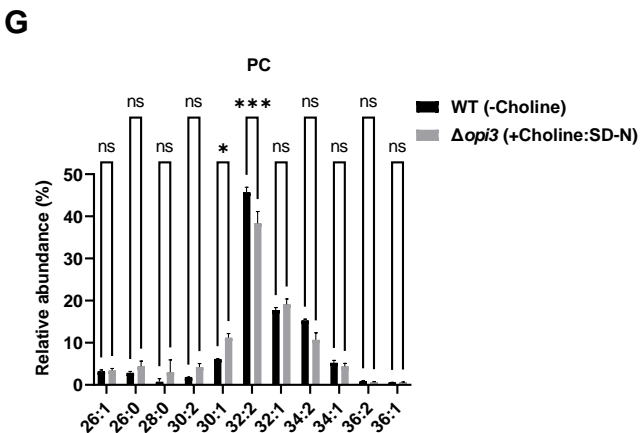
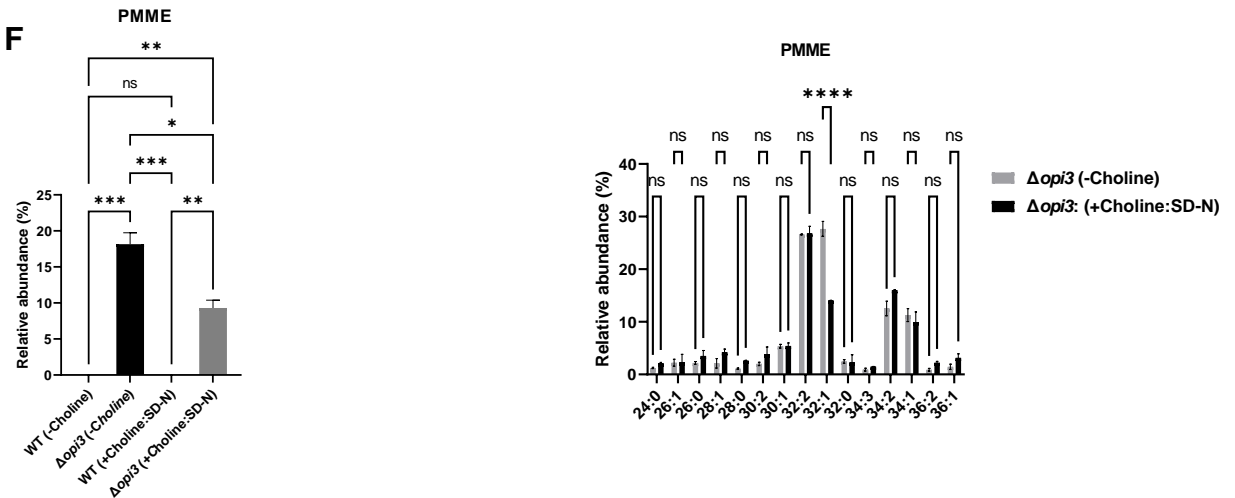
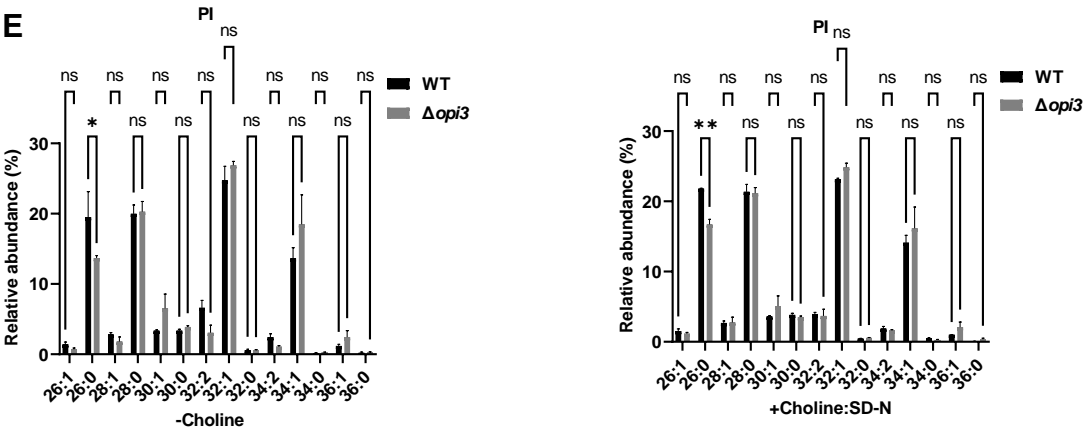


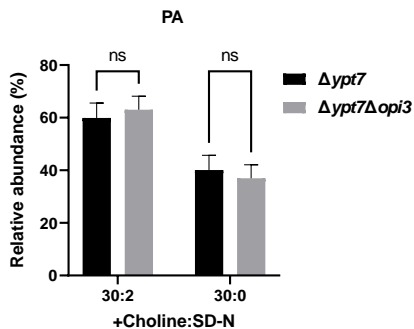
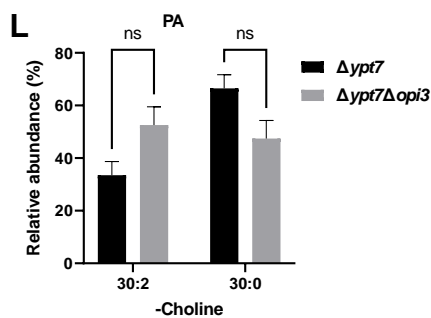
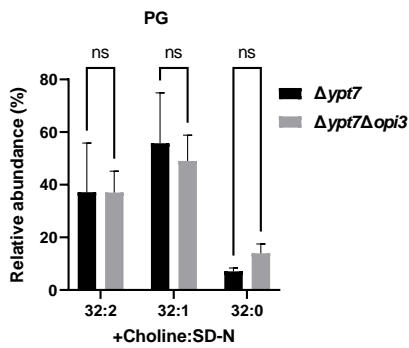
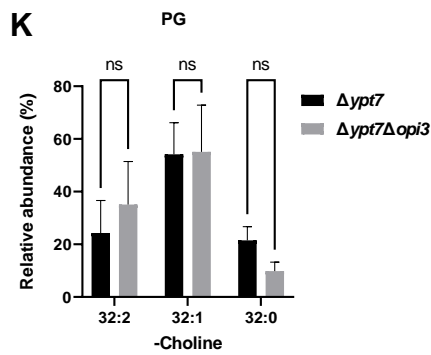
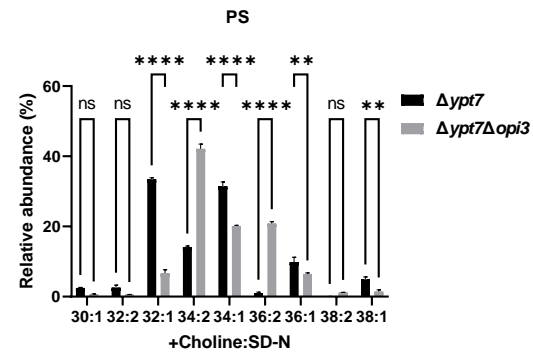
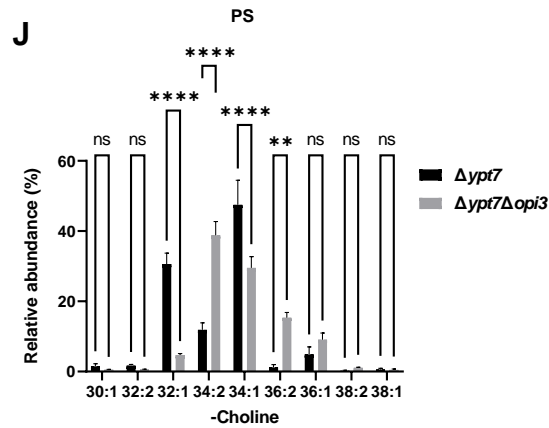
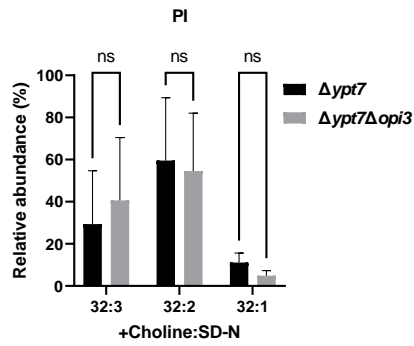
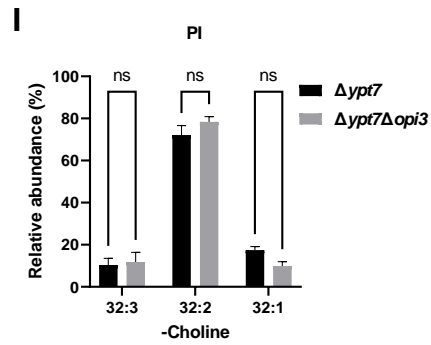
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H Yeast cells → Lysis → Pellet → IP GFP magnetic beads → Lipidomics of Atg8 membranes



Appendix figure S1: Addition of choline to $\Delta opi3$ cells specifically restores PC levels

A-G. Phospholipids species of WT and $\Delta opi3$ cells. Shown are acyl-chain length and composition of PG (A), PE (B), PA (C), PS (D), PI (E), PMME (F), and PC (G). Cells were grown in SD-URA, and starved for 4 h in SD-N, with choline (1 mM) supplementation excluded (-choline) or during starvation (+choline: SD-N) as indicated, harvested and analyzed by shotgun lipidomics. Statistical analysis was done by Anova multiple comparisons test- Sidak's (****, $p \leq 0.0001$, **, $p \leq 0.005$, *, $p \leq 0.001$, ns- not significant), error bars represent SEM (n=3, except for PE, PMME and WT with choline, n=2). For PMME (F- left panel)- Tucky's multiple comparison test.

H. Schematics of the procedure for lipidomic analysis of autophagic membranes. GFP-Atg8 expressing cells were lysed, Atg8-positive membranes were isolated from membrane fraction by GFP magnetic beads, and analyzed by shotgun lipidomics for lipid composition.

I-L: Phospholipids species of $\Delta ypt7$ and $\Delta ypt7 \Delta opi3$ cells. Shown are acyl chain length and composition of PI (I), PS (J), PG (K), PA (L). Cells expressing GFP-Atg8 were grown to log phase in SD-URA, and shifted to starvation medium. Cells were harvested after 3 h of starvation, lysed and membrane fraction was immunoprecipitated with GFP-magnetic beads. Atg8 enriched membranes were eluted and analyzed by shotgun lipidomics. Statistical analysis was done by Anova multiple comparisons test- Sidak's (****, $p \leq 0.0001$, **, $p \leq 0.005$, ns- not significant), error bars represent SEM of 3 independent experiments.