

Figure S1. Assessment of darkinduced senescence in Arabidopsis.
(A) Change of chlorophyll content (mg $\mathrm{chl} / \mathrm{mg} \mathrm{FW}$ ) during dark-induced senescence of intact plants.
(B) Exemplary tray of detached Arabidopsis plants after 48-h darkinduced senescence.


| B CFP-PTS1 ${ }^{\circ}$ | YFP-FBA1.2 (SFL>) | Merged |
| :---: | :---: | :---: |
| C Chlorophyll | FBA1.2-YFP | Merged |
| D COX4-CFP | FBA1.2-YFP |  |
| E CFP-PTS1 | FBA1.2-YFP $2^{2}$ | Merged $\begin{aligned} & 8, p+8 \\ & =x^{2} \end{aligned}$ |



Figure S3. Subcellular targeting analysis of ACO3.
Confocal images were taken in tobacco leaf epidermal/mesophyll cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1 (A, B), YFP fusion alone (C), or YFP fusion and the mitochondrial marker COX4-CFP (D). Scale bars $=10 \mu \mathrm{~m}$. Predicted PTS1 tripeptide is in parenthesis. ACO3 contains the C-terminal 390 aa of the 990-aa protein.


Figure S4. Subcellular targeting analysis of RPL19A and GSTU20.

Confocal images were taken in tobacco leaf epidermal cells
transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1 or YFP fusion and the mitochondrial marker COX4-CFP. Scale bars $=10 \mu \mathrm{~m}$. Predicted PTS1 peptides are in parentheses.
A

B


Figure S5. Subcellular targeting analysis of RH37.
(A) Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1 (upper), or YFP fusion and the mitochondrial marker COX4-CFP (lower). Scale bar $10 \mu \mathrm{~m}$.
(B) Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1 (upper), or YFP fusion and the mitochondrial marker COX4-CFP (lower). Scale bar, $10 \mu \mathrm{~m}$.


Figure $\mathbf{S 6}$.
Subcellular targeting
analysis of GRP2 and
GLX2-3.
Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1 or YFP fusion and the mitochondrial marker COX4-CFP. Scale bars, $10 \mu \mathrm{~m}$.


Figure S7. Subcellular targeting analysis of APX2.
(A) Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1 (upper), or YFP fusion and the mitochondrial marker COX4-CFP (lower). Scale bar, $10 \mu \mathrm{~m}$.
(B) Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1 (upper), or YFP fusion and the mitochondrial marker COX4-CFP (lower). Scale bar, $10 \mu \mathrm{~m}$.


Figure S8. Subcellular targeting analysis of ATL65.
(A) Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1. Scale bar, $10 \mu \mathrm{~m}$.
(B) Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1 (upper), YFP fusion and the mitochondrial marker COX4-CFP (middle), or YFP fusion alone (lower). Scale bars, $10 \mu \mathrm{~m}$.


Figure $\mathbf{S 9}$.
Subcellular targeting analysis of
AT3G47560 and PGL4.

Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1, or YFP fusion and the mitochondrial marker COX4-CFP. Scale bars, $10 \mu \mathrm{~m}$.


B


Figure S10. Subcellular targeting analysis of OARP1
(A) Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the peroxisome marker CFP-PTS1 (upper), or YFP fusion and the mitochondrial marker COX4-CFP (lower). Scale bars, 10 $\mu \mathrm{m}$.
(B) Confocal images were taken in tobacco leaf epidermal cells transiently expressing the YFP fusion protein and the mitochondrial marker COX4-CFP (upper), or in mesophyll cells expressing YFP fusion alone (lower). Scale bars, $10 \mu \mathrm{~m}$.


Figure S11. Gene expression data from Arabidopsis eFP browser (bar.utoronto.ca).
Data was normalized via the GeneChip Operating Software (GCOS). T, target intensity.
Bkg, background. Developmental stages from left to right: Rosette Leaf 2, Rosette Leaf 4, Rosette Leaf 6, Rosette Leaf 8, Rosette Leaf 10, Rosette Leaf 12, Senescing Leaf.

