SUPPLEMENTAL MATERIAL: SUPPLEMENTAL FIGURES AND LEGENDS OF SUPPLEMENTAL TABLES

Quantitative Proteomics Reveals that Plasma Membrane Microdomains from Poplar Cell Suspension Cultures are Enriched in Markers of Signal Transduction, Molecular Transport and Callose Biosynthesis

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Supplemental Figure 1:

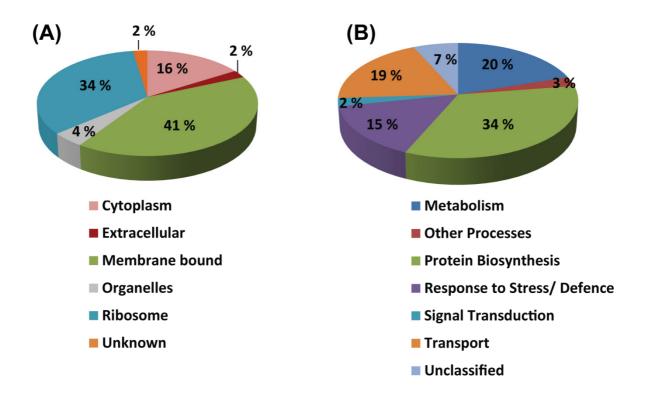


Fig. S1. Predicted cellular localization (A) and functional classification (B) of proteins identified after SDS-PAGE analysis of the TE fraction. Gene ontology information was retrieved from TAIR (www.arabidopsis.org).

Supplemental Figure 2:

Callose staining and measurement: Cells from suspension cultures were washed with 0.1 M K_2HPO_4 (pH 8.5) and stained with 0.01 % aniline blue for 12 min. Confocal imaging was performed using an inverted Zeiss LSM 780 microscope fitted with a 63 x 1.15 NA water immersion objective. A 405 diode laser (30 mW) was set at 5.5 % for excitation of aniline blue and emission was detected between 420 and 515 nm. The 405 laser was set to 0.6 % for bright field imaging. A maximum intensity projection was generated from 47 images acquired with an interval of 0.8 μ m (Fig. S2, G). The ZEN 2011 (Zeiss) software was used for image acquisition and processing.

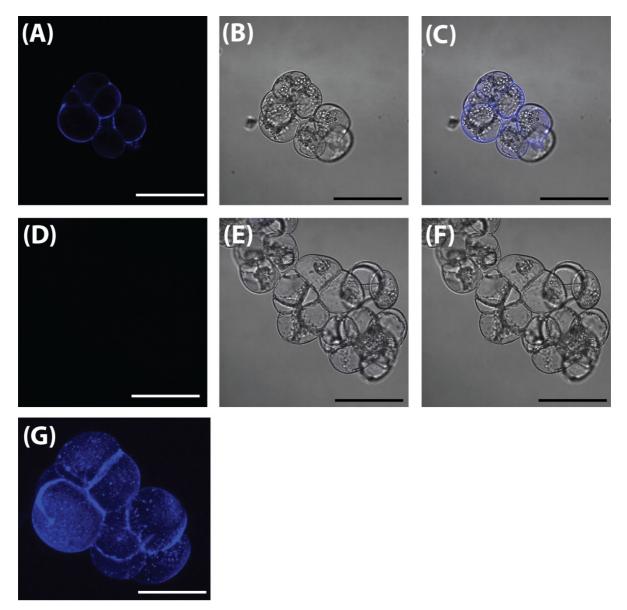


Fig. S2. Callose detection in poplar cells. Cells incubated in K₂HPO₄ buffer with (A-C and G) or without (D-F) aniline blue. A, D, and G: fluorescence image; B and E, corresponding transmission image of A and D, respectively; C, overlaid image of A and B; F, overlaid image of D and E; G, confocal maximum intensity projection image of poplar cells stained with aniline blue. Scale bars represent 20 μm.

Legends of supplemental Tables

Supplemental Table S1 - List of unique proteins and their corresponding peptides obtained from in-gel and solution (iTRAQ) proteomics analysis. Results are shown in the following sheets:

- **1. DRM-Proteins from SDS-PAGE** List of unique DRM proteins and their corresponding peptides identified from SDS-PAGE gels.
- **2. PMF- Proteins from SDS-PAGE** List of unique PMF proteins and their corresponding peptides identified from SDS-PAGE gels.
- **3. TE- Proteins from SDS-PAGE** List of unique TE proteins and their corresponding peptides identified from SDS-PAGE gels.
- **4.** Total-InGel+iTRAQidentification Combined list of unique proteins and their corresponding peptides obtained from in-gel and solution (iTRAQ) proteomics analysis.

Supplemental Table S2 - List of unique PMF and DRM proteins and their corresponding peptides quantified by iTRAQ. Results from each biological replicate are shown in the following sheets:

- **1. BR1-iTRAQquant -** List of unique PMF and DRM proteins quantified by iTRAQ (first biological replicate).
- **2. BR1-iTRAQquant+peptides** List of unique PMF and DRM proteins and their corresponding peptides quantified by iTRAQ (first biological replicate).
- **3. BR2-iTRAQquant** List of unique PMF and DRM proteins quantified by iTRAQ (second biological replicate).
- **4. BR2-iTRAQquant+peptides** List of unique PMF and DRM proteins and their corresponding peptides quantified by iTRAQ (second biological replicate).
- **5. BR3-iTRAQquant** List of unique PMF and DRM proteins quantified by iTRAQ (third biological replicate).
- **6. BR3-iTRAQquant+peptides -** List of unique PMF and DRM proteins and their corresponding peptides quantified by iTRAQ (third biological replicate).

Supplemental Table S3 - MS/MS spectra of peptides used for single-peptide based protein identification in DRM samples analyzed by SDS-PAGE.

Supplemental Table S4 - MS/MS spectra of peptides used for single-peptide based protein identification in PMF samples analyzed by SDS-PAGE.

Supplemental Table S5 - MS/MS spectrum of peptides used for single-peptide based protein identification in TE samples analyzed by SDS-PAGE.

Supplemental Table S6 - MS/MS spectra of peptides from the DRM and PMF samples used for single-peptide based protein identification in the 1st biological replicate (iTRAQ experiments).

Supplemental Table S7 - MS/MS spectra of peptides from the DRM and PMF samples used for single-peptide based protein identification in the 2^{nd} biological replicate (iTRAQ experiments).

Supplemental Table 8 - MS/MS spectra of peptides from the DRM and PMF samples used for single-peptide based protein identification in the 3rd biological replicate (iTRAQ experiments).