

Supplemental Figure 2. Sequence logos for the expanded curated proteomes of stroma (86), thylakoid membrane (95 proteins), inner envelope membrane (43 proteins) and lumen (53 proteins). Arrows indicated the predicted cleavage site (between the -1 and +1 position). Accession numbers for each subproteome are given in the Supplemental Table 1. These sequence logos are graphic representation of sequence conservation and amino acid frequency (Schneider, 1990).

