



SUPPLEMENTARY FIGURE S1

Supplementary Figure S1. Protein sequences were deduced from cDNA or EST sequences obtained from the Phytozome interface of the Joint Genome Institute (JGI) and from the National Center for Biotechnology Information (NCBI) for *Arabidopsis thaliana* (*Arabidopsis*), *Glycine max* (soybean), *Nicotiana tabacum* (tobacco), *Oryza sativa* (rice), *Physcomitrella patens* (*Physcomitrella*), *Populus trichocarpa* (poplar), *Saccharomyces cerevisiae* (yeast1), *Saccharomyces bayanus* (yeast2), *Solanum esculentum* (tomato), *Solanum tuberosum* (potato) and *Sorghum bicolor* (sorghum). The sources for the sequences are as follows: the JGI gene names are given for *Physcomitrella*, poplar, and sorghum sequences; the genome annotations are given for the *Arabidopsis* and rice sequences; the NCBI accession numbers are given for the tobacco, soybean, yeast, potatoPKCYT1 and tomato7 sequences; the TIGR Transcript Contig (TC) codes are given for the rest of the potato and tomato sequences. Regions of the deduced protein sequences that were judged to be incorrectly annotated or missing from the current gene models are highlighted in yellow.

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>tobacco1 Q40546

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>tobacco2 CK294819

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>tobacco3 CN747592 and DW001363

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>tobacco4 Q40545

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>tobacco5 Z29492

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>tobacco6 CK290844

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>tomato1 TC170021

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>tomato2 TC170352

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>tomato3 TC172280

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>tomato4 TC179493

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>tomato5 TC181248

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>tomato6 TC182827
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>tomato7 AW933570
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>tomato9 TC183199
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>yeast1 NP_009362
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>yeast2 AA032371
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