

Online Supporting Information A. This benchmark dataset \mathcal{S} for **Plant-mPloc** includes 1,055 plant protein sequences (978 different proteins), classified into 12 plant subcellular locations. Among the 978 different proteins, 904 belong to one subcellular location, 71 to two locations, and 3 to three locations. Both the accession numbers and sequences are given. None of the proteins has $\geq 25\%$ sequence identity to any other in the same subset (subcellular location). See the text of the paper for further explanation.

(1) \mathcal{S}_1 : 56 cell membrane proteins

>P25848

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>P42804

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>O20162

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>Q96291

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>P46269

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(11) S_{11} : 39 plastid proteins

>P28643

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>P28645

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>Q85UK6

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>P31562

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>Q2EEW0

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>Q2EEW1

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>Q2EEW2

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>Q2EEW4

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>Q2EEW7

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>P30060

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>Q2EEW8

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>Q9TJQ6

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>P30061

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>P30401

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>Q9TJQ7

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>Q9TIR8

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>Q9TJQ9

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>P82412

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>P30067

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>P30068

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>P25699

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>A7M988

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>O47032

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>Q2EEX0

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>Q2EEX2

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>Q2EEX3

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>Q2EEV8

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>P27068

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>Q2EEV9

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>P30070

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>Q2EEX9

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>P32034

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>P32035

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>A7M934

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(12) S_{12} : 52 vacuole proteins

>P08031

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>Q9LDC0

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>P15797

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>P07052

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>P04347

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>P07053

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>P23472

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>O22213

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>Q00326

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>P18417

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>P06673

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>P15478

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