

Online Supporting Information A. This benchmark dataset \mathcal{S} includes 523 Gram-positive bacterial protein sequences (519 different proteins), classified into 4 subcellular locations. Among the 519 different proteins, 515 belong to one location; and 4 to two 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 : 174 cell membrane proteins

>P23453

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>P07373

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>P54104

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>P50002

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>O34853

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>P54146

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>Q02114

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>P38536

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>P09333

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>P54422

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