Online Supporting Information A. This benchmark dataset includes 3,681 locative protein sequences (3,106 different proteins), classified into 14 human subcellular locations. Among the 3,106 different proteins, 2,580 belong to one location; 480 to two locations; 43 to three locations and 3 to 4 locations. Both the accession numbers and sequences are given. None of the proteins has ≥25% sequence identity to any other in the same subset (subcellular location). See the text of the paper for further explanation.

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(1) 77 centrosome proteins

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