

Online Supporting Information S2a. The degenerate testing dataset used for a head-to-head comparison between **MultiLoc2**, an updated version of **MultiLoc** (Hoglund et al., Bioinformatics 2006, 22, 1158-1165), and **iLoc-Hum** of this paper. The dataset contains 270 single-location proteins classified into the following nine subcellular location sites: (1) cytoplasmic, (2) endoplasmic reticulum, (3) extracellular, (4) Golgi apparatus, (5) lysosomal, (6) mitochondrion, (7) nuclear, (8) peroxisomal, and (9) plasma membrane, with each containing 30 protein sequences that were randomly picked from Swiss-Prot database (Release 2011_06 of 31-May-11). To avoid bias, none of the proteins included here occurs in the training dataset of **MultiLoc2**, nor in the training dataset of **iLoc-Hum**. See the text of the paper for further explanation.

(1) 30 cytoplasmic proteins

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>060259

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>060759

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>075190

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>P14136

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>P35573

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>P53370

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>P78362

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>O43567

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>P07237

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>O75900

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>O55232

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>O77588

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>094491

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>P17598

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