Supplementary material for

DispHScan: a multi-sequence web tool for predicting protein disorder as a function of pH

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Figure S1. Median DispH scores obtained for the human proteome (UP000005640_9606) analyzed with DispHScan at pH 7 for (A) transitioning (0.014) and non-transitioning proteins (0.240) and (B) multi-transitioning (-0.034) and single-transitioning proteins (0.024).