Supplementary Table 5  Prediction for the first group of 60 “ambiguous” annotated proteins.

In this table, “Annotation” denotes the experimentally observed subcellular localizations in Yeast Gtp Fusion Localization Database [17]. Predictions in different color regions correspond to different types: blue (correct), orange (partial correct), red (mismatch) and white (unknown).

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<th>SGD cellular component</th>
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