Supporting online material for:

LocText: relation extraction of protein localizations to assist database curation

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1. Short description of Supporting Online Material

Some results not shown in main paper but supporting some described findings.

2. Material

(starts in next page; one Table/Figure per page)

Fig. S1. Parse Tree Features. *LocText* derives features from parsed syntax trees: dependencies between tokens (e.g. "to" prepositions "the trans-Golgi network"), noun phrases (e.g. the AP-1 μ adaptin subunit"), part of speech tags (e.g. "localized" is a verb, past participle), linear distance between tokens (e.g. "adaptin" and "Golgi" are 10-tokens away from each other), or dependency distances between tokens (e.g. "adapting" and "Golgi" are 4-dependencies away).

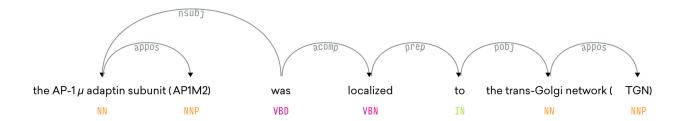


Table S1. Sample features after L1 regularization selection.

Feature name	Explanation	
Total count of entities	Float (scaled [0, 1]), count of entities (proteins + localizations + organisms) in sentence-based instance.	
Is Protein Marker	(Binary) test whether the marked protein text was equal to a list of manually defined protein markers: GFP, CYH2, ALG2, MSB2, KSS1, KRE11, SER2.	
PROTEIN localize	Binary, Linear Dependency (LD) feature. Test if the sentence-based instance contains two consecutive tokens (2-gram), where the first token is part of an protein name and the following has lemma "localize" (e.g. localized).	
SOURCE localization at	Binary, LD. Test if the first entity (source) in the sentence-based instance (either protein or location; likely in this case, a protein), is followed by the lemmas "localization" and "at".	
SOURCE proliferation TARGET	Binary, Parsing Dependency (PD). Test if the source entity is connected to the second entity (target) by the lemma "proliferation" in the dependency parse tree.	
in the TARGET	Binary, LD. Test if the lemmas "in" and "the" follow the target entity (likely a location).	
SOURCE VERB DET	Binary, LD. Test if the source entity is followed by two tokens, the first a VERB, the next a determiner (DET)	
VERB NOUN TARGET	Binary, PD. Test if, in the dependency parse tree, a VERB token connects a NOUN token that connects the target entity.	
NSUBJ DOBJ PREP	Binary, PD. Test if, in the dependency parse tree, three tokens are connected with the dependencies NSUBJ (nominal subject) to (direct object) to PREP (prepositional modifier).	

Table S2. Examples of novel GO annotations text-mined by *LocText*.

Protein	Localization	Text Source
P61586 human	GO:0005634 (nucleus)	Recent studies have revealed the localization of RhoA protein in the cell nucleus , in addition to its distribution in the cytosol and cell membrane. <i>PMID</i> 26622605, 2015
P09936 human	GO:0008021 (synaptic vesicle)	Both synaptic vesicle markers co-localized with the neuronal marker PGP 9.5 and exhibited granular accumulation patterns in the human and rat ENS. <i>PMID</i> 24025431, 2013
P56817 hu- man	GO:0005764 (lysosome)	Here, we report that lack of ubiquitination at Lys-501 (BACE1K501R) does not affect the rate of endocytosis but produces BACE1 stabilization and accumulation of BACE1 in early and late endosomes/ lysosomes as well as at the cell membrane. <i>PMID 23109336</i> , 2012
P01149 yeast	GO:0005773 (vacuole)	[] we provided evidence for the existence of an endocytic intermediate(s) from the yeast Saccharomyces cerevisiae that is responsible for the transport of the pheromone alpha-factor from the plasma membrane to the vacuole . PMID 8314797, 1993
P13134 yeast	GO:0005768 (endo- some)	However, Kex2 localization is not static, and its itinerary apparently involves transiting out of the late Golgi and cycling back from post-Golgi endosomal compartments during its lifetime. We tested whether the endocytic pathway could deliver small molecules to Kex2 from the extracellular medium. Here we report that intramolecularly quenched fluorogenic substrates taken up into intact yeast revealed fluorescence due to specific cleavage by Kex2 protease in endosomal compartments. <i>PMID 10393104, 1999</i>
Q9FE59 cress	GO:0009705 (plant-type vacuole membrane)	We demonstrate that this motif can reroute other proteins, such as INT4, SU-CROSE TRANSPORTER2 (SUC2) , or SWEET1, to the tonoplast and that the position of the motif relative to the transmembrane helix is critical. <i>PMID</i> 22253225, 2012
O82533 cress	GO:0005829 (cytosol)	Here, we report the identification of a second nuclear-encoded FtsZ-type protein from Arabidopsis that does not contain a chloroplast targeting sequence or other obvious sorting signals and is not imported into isolated chloroplasts, which strongly suggests that it is localized in the cytosol . <i>PMID 9836740</i> , <i>1998</i>

Fig. S2. List of all finally selected features. Python-readable list of descriptive feature names.

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    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<be ~~ protein>_[0]", # 196
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<induce>_[0]", # 197
    DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<e_2 ~~ and>_[0]", # 198"
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<and ~~ inhibit>_[0]", # 199
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<retention ~~ inhibit>_[0]", # 200
    "DependencyFeatureGenerator::23_PD_pos_N_gram_PD_3_<ADP ~~ ADP ~~ TARGET]>_[0]", # 201
    "DependencyFeatureGenerator::26 PD undirected edges N gram PD 3 <nsubj ~~ dobj ~~ compound> [0]",
# 202
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_2_<compound ~~ conj>_[0]", # 203
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_2_<dobj ~~ advcl>_[0]", # 204
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_3_<pobj ~~ prep ~~ dobj>_[0]", #
205
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<function ~~ TARGET]>_[0]", # 206
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<VERB ~~ ADJ ~~ TARGET]>_[0]", # 207
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_2_<nmod ~~ pobj>_[0]", # 208
    "DependencyFeatureGenerator::23_PD_pos_N_gram_PD_3_<ADP ~~ PROPN ~~ TARGET]>_[0]", # 209
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_3_<[SOURCE ~~ be ~~ require> [0]", # 210
    "DependencyFeatureGenerator::19 LD pos_N_gram_LD_3_<PUNCT ~~ ADV ~~ PUNCT>_[0]", # 211
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<function>_[0]", # 212
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<of ~~ e_1>_[0]", # 213
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_3_<- ~~ independent ~~ function>_[0]", # 214
    "DependencyFeatureGenerator::22 PD bow N gram PD 3 <[SOURCE ~~ function ~~ require> [0]", # 215
    "DependencyFeatureGenerator::22 PD bow N gram PD 2 <of ~~ function> [0]", # 216
    "DependencyFeatureGenerator::22 PD bow N gram PD 3 < [SOURCE ~~ of ~~ function> [0]", # 217
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_3_<pobj ~~ prep ~~ nsubjpass>_[0]",
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_2_<pobj ~~ nmod>_[0]", # 219
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_3_<prep ~~ pobj ~~ nmod>_[0]",
220
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<PUNCT ~~ ADP ~~ VERB>_[0]", # 221
    "DependencyFeatureGenerator::18 LD bow N gram LD 1 <signal> [0]", # 222
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<[SOURCE ~~ signal>_[0]", # 223
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_2_<amod ~~ nsubj>_[0]", # 224
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_1_<show>_[0]", # 225
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_2 <dobj ~~ ccomp> [0]", # 226
    "DependencyFeatureGenerator::18 LD bow N gram LD 1 <ubiquitination> [0]", # 227
    "DependencyFeatureGenerator::22 PD bow N gram PD 1 <ubiquitination> [0]", # 228
    "DependencyFeatureGenerator::23_PD_pos_N_gram_PD_3_<[SOURCE ~~ ADP ~~ TARGET]>_[0]", # 229
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<activation>_[0]", # 230
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<NOUN ~~ VERB ~~ PROPN>_[0]", # 231
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<NUM ~~ NOUN ~~ ADP>_[0]", # 232
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_1_<region>_[0]", # 233
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<ADJ ~~ ADP ~~ PROPN>_[0]", # 234
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_3_<nsubj ~~ advcl ~~
nsubjpass>_[0]", # 235
    "DependencyFeatureGenerator::18 LD bow N gram LD 3 <e 1 ~~ , ~~ the> [0]", # 236
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_2_<advcl ~~ prep>_[0]", # 237
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<[SOURCE ~~ from>_[0]", # 238
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<into ~~ TARGET]>_[0]", # 239
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<ADP ~~ ADJ ~~ TARGET]>_[0]", # 240
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_2_<ADV ~~ TARGET]>_[0]", # 241
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<this>_[0]", # 242
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<DET ~~ ADJ ~~ ADJ> [0]", # 243
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<ADP ~~ PROPN ~~ PUNCT>_[0]", # 244
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<compromise>_[0]", # 245
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_1_<fuse>_[0]", # 246
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_2_<acl ~~ acomp>_[0]", # 247
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<NOUN ~~ PUNCT ~~ PUNCT>_[0]", # 248
```

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"DependencyFeatureGenerator::18 LD bow N gram LD 1 <vesicle> [0]", # 249
    "DependencyFeatureGenerator::18 LD_bow_N_gram_LD_1_<dispensable>_[0]", # 250
    "DependencyFeatureGenerator::18 LD bow N gram LD 2 <be ~~ dispensable> [0]", # 251
   "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<dispensable ~~ for>_[0]", # 252
   "DependencyFeatureGenerator::18\_LD\_bow\_N\_gram\_LD\_2\_<e\_1 \sim\sim to>\_[0]", \# 254
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<location>_[0]", # 255
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<[SOURCE ~~ location>_[0]", # 256
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<[SOURCE ~~ location>_[0]", # 257
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<, ~~ to>_[0]", # 258
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<, ~~ while>_[0]", # 259
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_2_<appos ~~ compound>_[0]", # 260
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<PUNCT ~~ VERB ~~ TARGET]>_[0]", # 261
   "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<important>_[0]", # 262
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_1_<morphology>_[0]", # 263
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_3_<pobj ~~ amod ~~ npadvmod>_[0]",
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<defect>_[0]", # 265
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<cohesin>_[0]", # 266
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<assembly>_[0]", # 267
    DependencyFeatureGenerator::22 PD bow N gram PD 3 <[SOURCE ~~ determine ~~ localize> [0]", #
268
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<[SOURCE ~~ TARGET]>_[0]", # 269
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<significantly>_[0]", # 270
    "DependencyFeatureGenerator::18 LD bow N gram LD 2 <be ~~ significantly> [0]",
   "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<at ~~ have>_[0]", # 272
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_3_<[SOURCE ~~ at ~~ have>_[0]", # 273
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_3_<[SOURCE ~~ recruitment ~~ of>_[0]", # 274
    DependencyFeatureGenerator::26 PD undirected edges N gram PD 3 <prep ~~ nsubj ~~ prep> [0]", #
275
   "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_3_<[SOURCE ~~ domain ~~ of>_[0]", # 276
    "DependencyFeatureGenerator::22 PD bow N gram PD 3 < [SOURCE ~~ domain ~~ of> [0]", # 277
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_1_<reconfiguring>_[0]", # 278
   "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2 <division ~~ TARGET]>_[0]", # 279
   "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<abundance ~~ TARGET]>_[0]", # 280
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<eliminate>_[0]", # 281
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2_<from ~~ through>_[0]", # 282
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_3_<from ~~ through ~~ TARGET]>_[0]", # 283
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_1_<stress>_[0]", # 284
    DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<[SOURCE ~~ stress>_[0]", # 285"
    "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_1_<stress>_[0]", # 286
    "DependencyFeatureGenerator::26_PD_undirected_edges_N_gram_PD_3_<appos ~~ nsubj ~~ dobj>_[0]", #
287
   "DependencyFeatureGenerator::22 PD bow N gram PD 3 < localization ~~ disrupt ~~ by> [0]", # 288
    "DependencyFeatureGenerator::18 LD bow N gram LD 3 <e 1 ~~ to ~~ TARGET]> [0]", # 289
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<that ~~ u>_[0]", # 290
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_2_<require ~~ e_1>_[0]", # 291
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_2_<PRON ~~ NUM>_[0]", # 292
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_3_<show ~~ that ~~ u>_[0]", # 293
    "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_3_<first ~~ require ~~ e_1>_[0]", # 294
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<ADP ~~ PRON ~~ NUM>_[0]", # 295
    "DependencyFeatureGenerator::19_LD_pos_N_gram_LD_3_<NUM ~~ ADV ~~ VERB>_[0]", # 296
   "DependencyFeatureGenerator::22_PD_bow_N_gram_PD_2 <show ~~ require>_[0]", # 297
   "DependencyFeatureGenerator::18_LD_bow_N_gram_LD_3_<[SOURCE ~~ to ~~ mediate>_[0]", # 298
    "IsSpecificProteinType::40_is_marker_[0]", # 299
    "LocalizationRelationsRatios::50 corpus unnormalized total background loc rels ratios [0]", #
300
    "LocalizationRelationsRatios::58_SwissProt_normalized_exists_relation_[0]", # 301
```

Fig. S3. PR-curve analysis. *LocText* vs. *Baseline*, using manually-annotated entities. The maximum recall for both methods is 74%. The *Baseline* is shown as a single point (no decision value). A two-sample two-tailed t-test was performed to determine whether the methods' difference in F-Measure, $F(LocText)=79\%\pm3$ vs. $F(Baseline)=74\%\pm3$, was significant. The t-statistic was significant at the 99% confidence level, t(3998)=28.04, p=3.99e-165.

