

Before UID conversion (enter your data in the box below)



SUPPORTED FORMATS for unique identifiers (UID)

List: UIDs separated by spaces, tabs, ...

Alignment: fasta, msf, dustalw, nexus, phytip, mega & stockholm

Tree: newick, nexus

To see an example for the name template shown below, press

List or Alignment or Tree ?

CAB04658.1 NP_248318.1 AAB99427.1 AAC73542.1 BAA36038.1 ZWECAP P03033 133353 AAA85499.1

or upload it from file:

Browse...

Send results to e-mail:

Reset

Names/format designer (select name template, refine if needed, and Submit the query)



Select name template Complex protein name

Refine name template ☒

Template-building facility

#	Delimiter	Characteristic	Length			Location
			min	max	abbreviate	
01		Primary ID	9	9		↑ ↓ + -
02	space	Organism=>G[enus].species	6	6		↑ ↓ + -
03	[	Protein, product	17	17		↑ ↓ + -
04	]	CDS, locus tag				↑ ↓ + -

tabs ☐

Show * if characteristic is not available

Select database source GenBank

Submit

OUTPUT:

Alignment/tree options

☐ alignment format
☐ show sequence sizes

☐ tree format
☐ draw as cladogram

Interactive display options

☐ names only
☐ annotation table
☒ id-to-name log

substitute in names

with

After UID conversion (see results below and in separate window as Text or Excel formats)

• Info: Number of IDs: submitted = 9, converted = 9. Time total: 0.87 sec (0.10 per ID).

- Warning: the input was processed as an ID list. If it was not ID list, please modify your input and try again.
- Warning: some IDs were recognized as non-unique IDs (see ID-to-name log for details). These IDs can be linked to more than one entry that could result in misprocessing. For unambiguous ID processing, either switch to UIDs or search another DB.

3879212 C.eleg[C. elegans protei]R13H4.6
15669508 M.jann[hypothetical prot]MJ1318
1592066 M.jann[ATP-dependent pro]MJ_1417
1786643 E.coli[DNA-binding ATP-d]b0439
1651586 E.coli[UmuD protein.]*
2144964 E.coli[mucA protein]*
126224 {...} [LexA repressor]*
133353 E.phag[Repressor protein]*
763050 S.phag[repressor protein]*

Primary ID,
9 positions

Organism=>G[enus].species,
6 positions

Protein, product,
17 positions

CDS, locus tag