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Worm Annotation in UniProtKB/Swiss-Prot

Duncan Legge and the UniProt Consortium.

Introduction to UniProtKB/Swiss-Prot

UniProt (Universal Protein Resource; http://www.uniprot.org) provides a central resource on protein sequences and functional annotation. The UniProt Consortium is comprised of the European Bioinformatics Institute (EBI), the Swiss Institute of Bioinformatics (SIB) and the Protein Information Resource (PIR). The UniProt Knowledgebase (UniProtKB) contains the manually annotated UniProtKB/Swiss-Prot section and the automatically annotated UniProtKB/TrEMBL section.

UniProtKB/TrEMBL
Computer-annotated protein sequences
Sequences with 100% identity from the same organism are merged

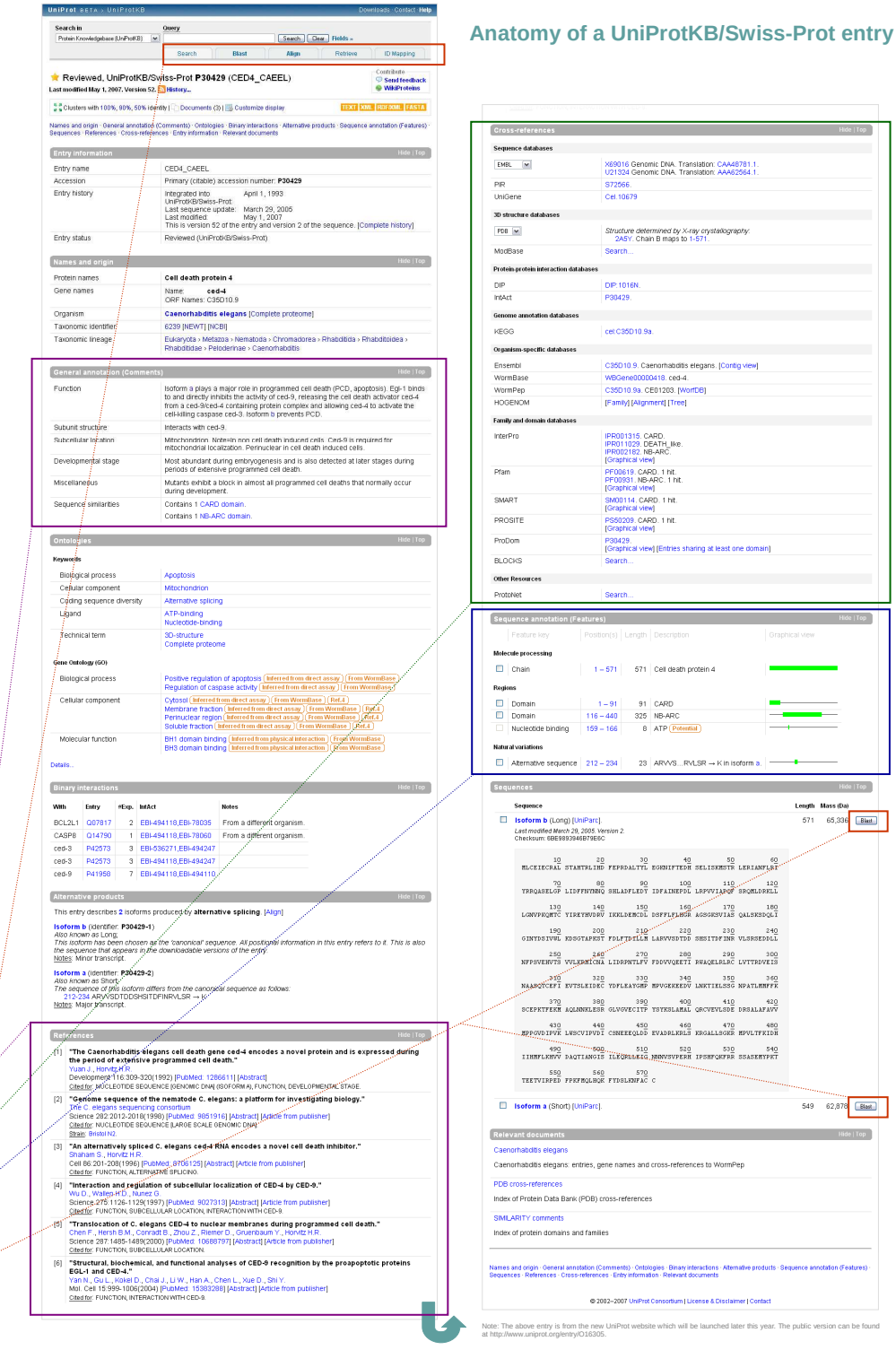
UniProtKB/Swiss-Prot
Manually annotated protein sequences
Non-redundant and high level of accuracy

The UniProt Consortium liaises with many of the research communities that concentrate on model organisms to ensure that we are supplying the best service.

Useful Features of UniProtKB/Swiss-Prot

- Manual referenced annotation
- Extensive cross-linking to other databases
- Feature table showing domains/sites of interest with illustrations
- Wide range of bioinformatics tools available for analysis

Anatomy of a UniProtKB/Swiss-Prot entry



Note: The above entry is from the new UniProt website which will be launched later this year. The public version can be found at http://www.uniprot.org/entry/O16305.

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