

Resource Summary Report

Generated by [Kravitz](#) on Sep 10, 2026

UniSave

RRID:SCR_004946

Type: Tool

Proper Citation

UniSave (RRID:SCR_004946)

Resource Information

URL: <http://www.ebi.ac.uk/uniprot/unisave/?help=0&session=>

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Description: The UniProtKB Sequence/Annotation Version Archive (UniSave) is a repository of UniProtKB/Swiss-Prot and UniProtKB/TrEMBL entry versions. Entries can be retrieved by entering a primary accession number or an entry name and pressing the Go! button. The result of the query is a list of entry versions with the UniProtKB database name, entry status, primary accession number, entry name, entry version, sequence version, release number and the release date, ordered by the release date, the latest version first. The entry version status can be ""incorporated"", ""active"", ""changed"", ""replaced"" or ""deleted"". An incorporated entry version is the first entry version added into UniProtKB, an active entry version is part of the latest public release, a changed entry version has been superseded by a newer entry version, a replaced entry has become secondary to another entry, and a deleted entry has been removed from the UniProtKB without becoming secondary to any other entry. For replaced entry versions, the status ""Replaced"" can be clicked to return all entries, which have the given entry as a secondary entry. If a date is provided as part of the query then only the version of the entry that was current at that date is displayed. Entries can be viewed by clicking ""View"" in the query results table. The "" Earlier"" and ""Later >>"" links can be used to access the earlier and later entry versions. The ""Back to List"" link returns the user to the query results table. Selecting ""UniProtKB"" or ""Fasta"" and pressing ""Save"" downloads the entry in flat file or fasta format. Comparison between entry versions is straightforward: selecting two entries and clicking the ""Compare Selected"" button will show the differences between the two entries. Whenever comparisons are made a Smith-Waterman sequence alignment is computed using SSEARCH, and displayed at the bottom of the entry. The actual alignment is displayed only when the sequences are not identical.

Abbreviations: UniSave

Synonyms: UniProtKB Sequence/Annotation Version Archive

Resource Type: data or information resource, database

Defining Citation: [PMID:16551660](#)

Keywords: gold standard

Resource Name: UniSave

Resource ID: SCR_004946

Alternate IDs: nlx_91568, r3d100011244

Alternate URLs: <https://doi.org/10.17616/R3MP6J>

Record Creation Time: 20220129T080227+0000

Record Last Update: 20260905T133127+0000

Ratings and Alerts

No rating or validation information has been found for UniSave.

No alerts have been found for UniSave.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 2 mentions in open access literature.

Listed below are recent publications. The full list is available at [Kravitz](#) .

Birman D, et al. (2023) Pinpoint: trajectory planning for multi-probe electrophysiology and injections in an interactive web-based 3D environment. bioRxiv : the preprint server for biology.

Gómez-Gálvez Y, et al. (2020) Quantitative proteomic profiling of the rat substantia nigra places glial fibrillary acidic protein at the hub of proteins dysregulated during aging: Implications for idiopathic Parkinson's disease. Journal of neuroscience research, 98(7), 1417.