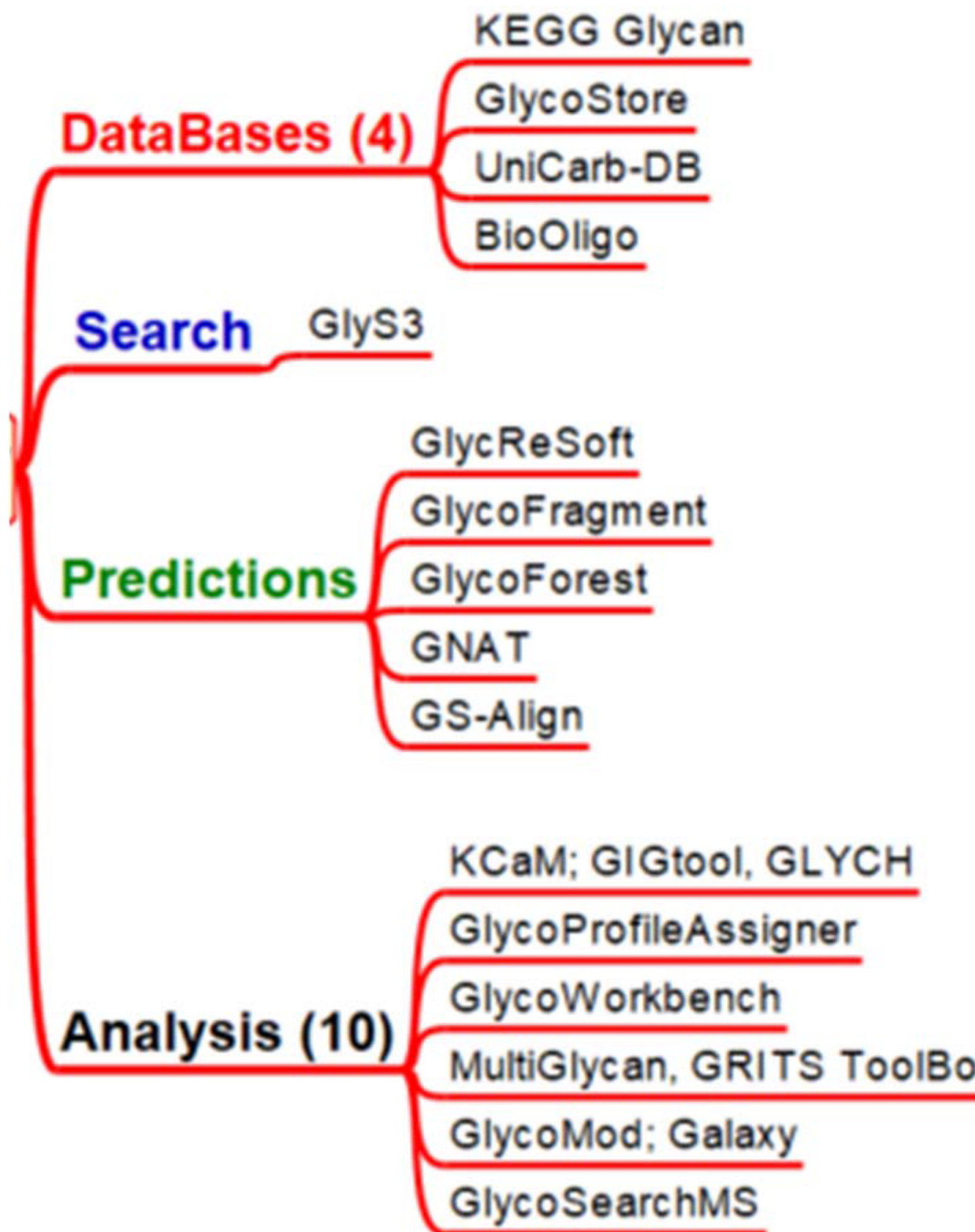






DATABASES

- **BioOligo** <http://glyco3d.cermav.cnrs.fr/search.php?type=bioligo> As part of the suite of interlinked Databases of 3D Structures of Glycan, BioOligo. contains representations, 3D structures and NMR spectra of most occurring glycans
- **GlycoStore** <http://www.glycostore.org> Provides a centralised resource that combines glycan structure information with chromatographic separation and electrophoretic data.
- **KEGG Glycan** <https://www.genome.jp/kegg/glycan/> The KEGG GLYCAN structure database is a collection of experimentally determined glycan structures. It contains all unique structures taken from CarbBank, structures entered from recent publications, and structures present in KEGG pathways
- **UniCarb-DB** <http://unicarb-db.expasy.org/> UniCarb-DB is a structural and mass spectrometric database used in glycomics. UniCarb-DB provides over 1000 LC-MS/MS spectra for N- and O-linked glycans released from glycoproteins that were manually annotated.

SEARCH

- **GlyS3** <https://glycoproteome.expasy.org/substructuresearch/> GlyS3 matches any substructure such as glycan determinants to a large collection of structures recorded in GlyConnect and SugarBindDB.

PREDICTIONS

- **GS Align** <http://www.glycanstructure.org/gsalign> A computational method for glycan structure alignment and similarity measurement. GS-align generates possible alignments between two glycan structures
- **GlycoFragment** <http://www.glycosciences.de/tools/GlycoFragments/manual.pdf> A web tool to support the interpretation of mass spectra of complex carbohydrates.
- **GlycoForest** <https://glycoforest.expasy.org/index.html> Glycoforest is a partial de-novo algorithm for sequencing glycan structures based on MS/MS spectra.
- **GNAT** <http://gnatmatlab.sourceforge.net/> Glycosylation Network Analysis Toolbox contains a variety of classes to describe glycans and glycosylation reaction networks. It also provides various methods to manipulate these classes
- **GlyReSoft** <https://github.com/GlycReSoft2> GlyReSoft is a modular software tool for assigning site specific glycosylation from bottom-up mass spectrometry data sets.

ANALYSIS

- **Galaxy** http://www.glycoanalysis.info/galaxy2/manual/User_Manual.pdf Provides a useful method for an analytical procedure for N-glycan structures. Galaxy is a 2D/3D mapping method developed for the structural determination of asparagine-linked oligosaccharides (N-glycans) in glycoproteins.
- **GIGTool** <https://www.genome.jp/tools/kcam/> GIG Tool is an application that extracts (i) precursor masses, (ii) oxonium ions and glycan fragments from tandem (liquid chromatography (LC)-MS/MS) mass spectra for glycan identification, and (iii) reporter ions from quaternary amine containing isobaric tag for glycan (QUANTITY) isobaric tags.
- **GLYCH** GLYCH (GLYcan CHaracterization) is package for glycan characterization using MS/MS.

- **GlycoMod** <https://web.expasy.org/glycomod/> GlycoMod is a tool that can predict the possible oligosaccharide structures that occur on proteins from their experimentally determined masses
- **GlycoProfileAssigner** <http://glycananalyzer.neb.com> Automated structural assignment with error estimation for glycan LC data
- **GlycoSearchMS** http://www.glycosciences.de/database/start.php?action=form_ms_search GlycoSearchMS takes a list of mass spectra peak values as input and searches for matches with the calculated fragments of SweetDB structures.
- **GlycoWorkbench** <https://glycoworkbench.software.informer.com/2.1/> A suite of software tools designed for the rapid drawing of glycan structures and for assisting the process of structure determination from mass spectrometry data.
- **GRITS ToolBox** www.grits-toolbox.org/ GRITS Toolbox combines the analytical power of glycan permethylation with glycopeptide analysis, preserving site-specific information for comprehensive glycoprotein analysis
- **KCaM** <https://www.genome.jp/tools/kcam/> KEGG Carbohydrate Matcher s a tool for the analysis of carbohydrate sugar chains, or glycans. It consists of a web-based graphical user interface that allows users to enter glycans easily with the mouse
- **MultiGlycan** <https://bio.tools/MultiGlycan> To help user to gather glycan profile information from LC-MS Spectra. It also reports quantity of specific glycan composition

Category

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