GNOme – Glycan Naming and Subsumption Ontology

~ A GlyGen Project ~

http://gnome.glyomics.org/

Wenjin Zhang and Nathan Edwards

Clinical and Translational Glycoscience Research Center, Georgetown

Introduction

GlyTouCan provides stable glycan accessions

...at *any* level of characterization

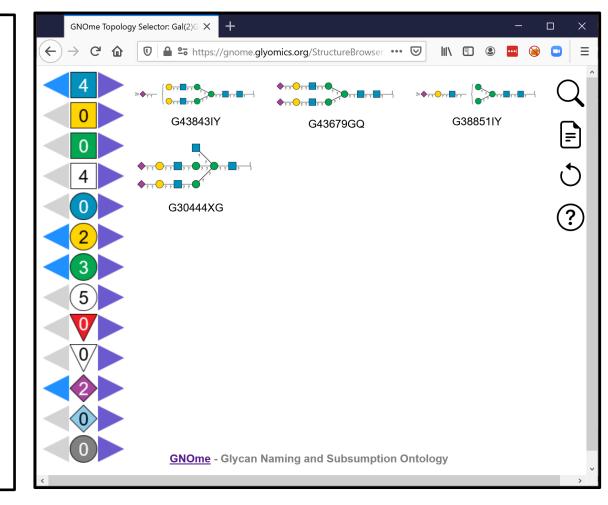
GNOme organizes the stable accessions:

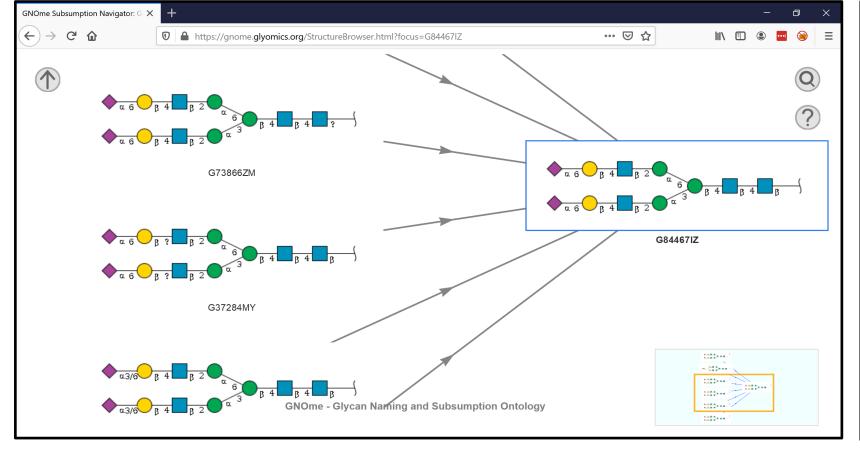
...for interactive browsing by subsumption

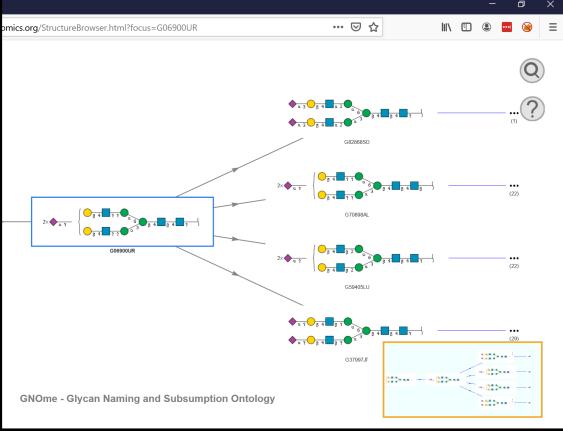
...for automated subsumption reasoning

...for text-based lookup

...with well-defined characterization levels Integrated with GlyGen as Related Glycans





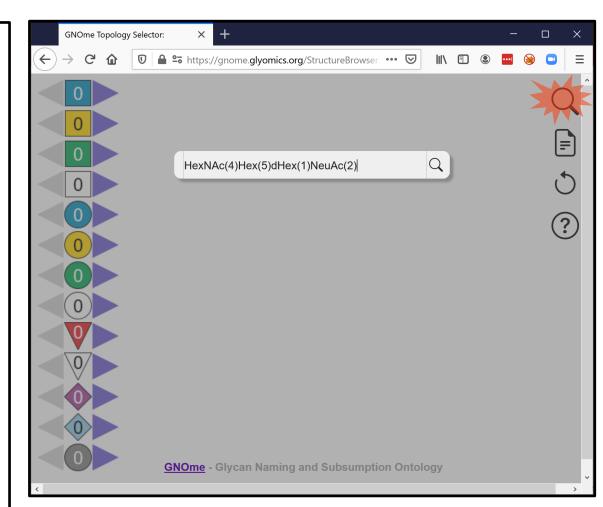


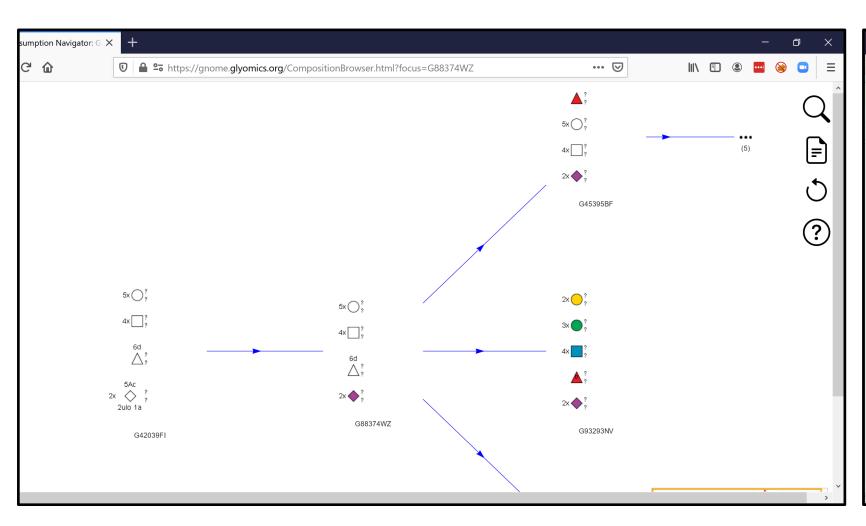
GNOme – Glycan Naming and Subsumption Ontology

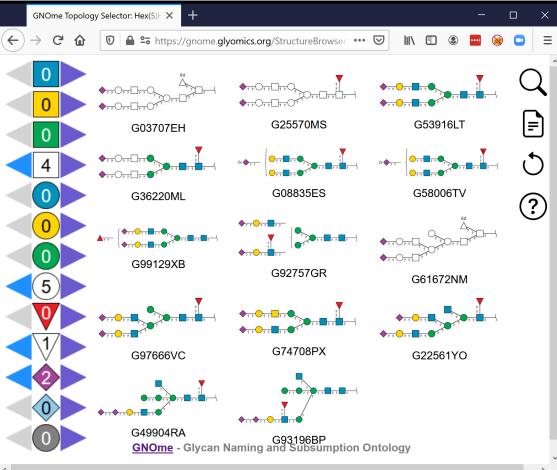
http://gnome.glyomics.org/

Glycan Lookup & Semantic Names

- Lookup glycans by accession or semantic names ("Find" (a))
 - Byonic format compositions:
 - HexNAc(4)Hex(5)dHex(1)NeuAc(2)
 - Short form: H5N3FS2
- The applet supports deep linking by GlyTouCan accession, semantic name, or monosaccharide composition
 - ?focus=G62109NW; ?HexNAc=5&Hex=4
- Explore monosaccharide compositions too







GNOme – Glycan Naming and Subsumption Ontology

http://gnome.glyomics.org/

On-Demand Alignment

- Align novel glycans using GlycoCT or WURCS sequence ("Align")
- Subsumption webservice computes alignment against GlyTouCan glycans
 - https://subsumption.glyomics.org/
 - Query directly, or run docker locally.
 - Interactive applet uses this webservice.
- Focus on known glycan, if registered
 - Otherwise, create node + related glycans
- Also used by Glycan Image Extractor
 - https://extractor.glyomics.org/

