NIST MS Search with Glycopeptide Library

This free product is for searching and browsing NIST glycopeptide fragmentation library (MS/MS).
It includes:

o NIST MASS SPECTRAL SEARCH PROGRAM Version 2.0g for use with Microsoft(R) Windows(TM)

o The IgG Glycopeptide library (11/6/2015, HCD).

System requirements

This MS search program is a 32-bit Unicode application and thus requires
Windows NT 4.0, 2000, XP, Vista, or Windows 7.

MS Search is compatible with MGF, DTA, and PKL formats, all common to proteomics workflows.

Instructions:

1. To install, run the program "NIST_IgG_Glycopeptide.exe" and then follow instructions.
2. To run, execute "nistms.exe" for searching and browsing NIST glycopeptide fragmentation library .
3. To search glycopeptide spectra, go to "names" option at the bottom as follows: