

# NIST Tandem Mass Spectral Library

## 2023 Release

**51,501** Compounds, **60%** More than 2020  
**400 K** Precursor Ions – **2.4 M** Spectra

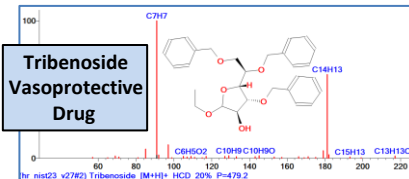
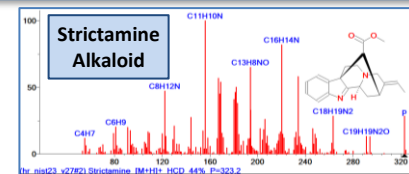
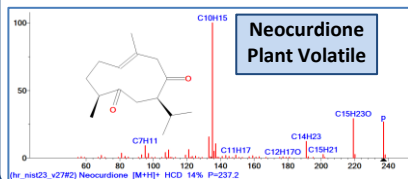
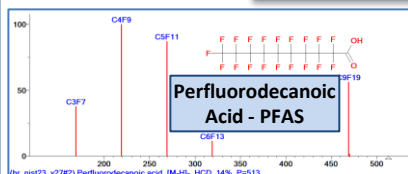
### Fragmentation Methods

49,590 HRAM (High Res Accurate Mass) Compounds  
50,071 QTOF, HCD, IT-HRAM, QqQ Compounds  
49,561 Ion Trap Compounds (Low Res., up to MS<sup>4</sup>)  
561 APCI HRAM Compounds

### Precursor Ion Types

44,191 Protonated  
19,620 Deprotonated  
14,318 Water/Ammonia Loss  
44,547 Other In-Source Generated

## ALL COMPOUNDS SELECTED FOR RELEVANCE MEASURED AT NIST, THOROUGHLY EVALUATED

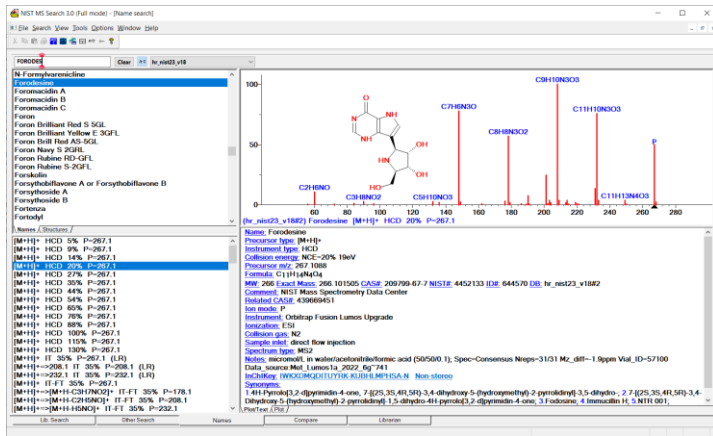


Citation	New	Total
Wikipedia	1618	6424
EPA Tox	3181	8146
Food DB	602	4491
EU Contaminants	6553	15818
Protein Data Bank	1246	4945
Human Metabolite DB	2071	9686
PFAS	90	116

### All Spectra Shown for Each Compound

1. Compound Name  
Many Synonyms

2. All Related Spectra  
All Energies  
Fragmentation Types  
In-Source Ions, ...



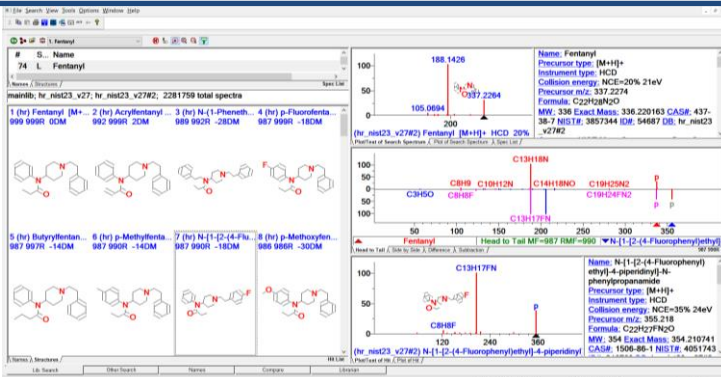
Full Spectrum  
Selected from List  
at Lower Left

All Spectrum Text  
Information

Hybrid Search  
Recommended for All  
Tandem MS Searches

Find Related  
Compounds Even if  
Query is not in Library

Differences in masses  
are 'Modifications'



Query Spectrum

Shifted Library Peaks  
(gray->red)  
Contain Modification

Library Spectrum

# NIST TANDEM LIBRARY SOFTWARE

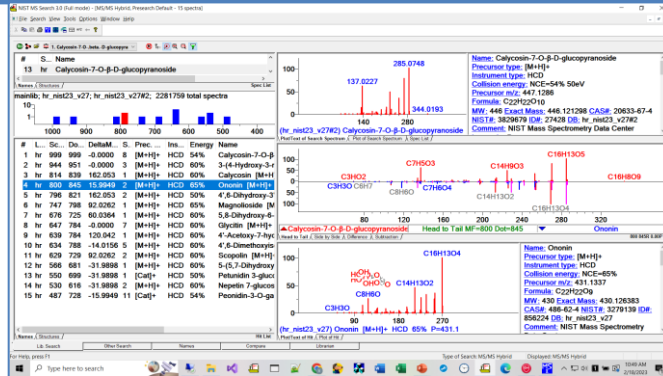
## 2023 Tandem Release

### TANDEM (MS/MS) ANALYSIS TOOLS

#### NISTMS

ALL SPECTRA FOR EACH COMPOUND IN ONE LIST

MULTIPLE ENERGIES ION MODES FRAGMENTATION TYPE IN SOURCE IONS



#### FILTER IDS BY SPECTRUM TYPE

M/SMS Hit List Filter

Enable Filtering (Tandem Only)

Instrument Type:  Ion Trap,  Ion Trap + FT,  Other

Allow Filtering:  C/H,  C/H + Na, K, Cl, HCO,  HCO, H<sub>2</sub>O,  All other (dimers/losses),  Exclude isotopic precursors

Preferred HCE/Voltage: 20

Polarity: Any

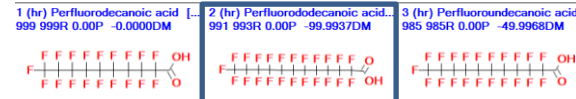
Buttons: OK, Cancel, Help

#### Hybrid Search

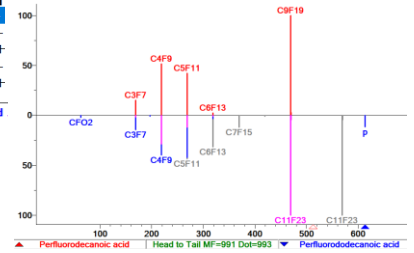
-FINDS COMPOUNDS NOT IN LIBRARY AND CONFIRMS IDS  
-USES PEAKS AND LOSSES  
-SHOWS SHIFTED PEAKS

RECOMMENDED FOR ROUTINE USE

#	L.	S...	D...	DeltaMa...	DBs	Prec...	Inst...	En...	Name	
1	hr	935	999	-0.0000	20	EGM	[M-H] <sup>-</sup>	HCD	20%	Perfluoroheptanoic acid [M-H] <sup>-</sup>
2	hr	891	976	-49.9968	27	WC...	[M-H] <sup>-</sup>	HCD	20%	Perfluoroheptanoic acid [M-H] <sup>-</sup>
3	hr	866	938	-99.9936	24	WC...	[M-H] <sup>-</sup>	HCD	14%	Perfluorononanoic acid [M-H] <sup>-</sup>
4	hr	511	796	-249.984	18	CE...	[M-H] <sup>-</sup>	HCD	27%	Perfluorododecanoic acid [M-H] <sup>-</sup>
5	hr	466	714	-149.990	22	WC...	[M-H] <sup>-</sup>	HCD	9%	Perfluorodecanoic acid [M-H] <sup>-</sup>
6	hr	460	828	-199.987	18	CE...	[M-H] <sup>-</sup>	HCD	44%	Perfluoroundecanoic acid [M-H] <sup>-</sup>



PFAS ILLUSTRATION: APPLICABLE TO MANY CLASSES



#### MS Interpreter

DIRECT ACCESS FROM NISTMS

CONNECT PEAKS TO STRUCTURE

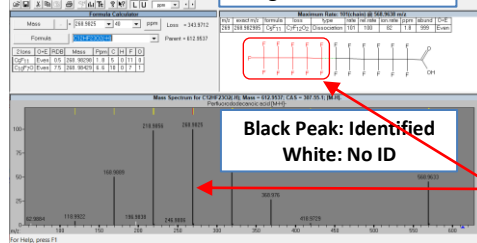
#### Chemical Formula Calculator

#### Isotope Calculator

Any resolution +/- Charge

Multiple Display Options

#### Fragmentation Details



Formula Calculator

Mass: 1.031634, mDa: 500

Formula: C-2 H-3 N-1 O-1 S-1

1 Ion	O+E	RDB	Mass	mDa	C	H	N	O	S
H <sub>2</sub> NO-1	Odd	0	1.03109	-0.5	0	3	1	1	0

Find Formula from Delta Mass

Select Peak Show Structure Fragment

#### NIST Website Chemdata.nist.gov

FREELY AVAILABLE LIBRARIES AND SOFTWARE

#### Downloadable Libraries

Annotated Recurrent Unidentified Spectra  
Urine, Plasma/Serum (ARUS)  
Tryptic Peptides (Human, ...)  
Oligosaccharide Libraries (Milk)  
Glycopeptides (mAb, Glycan Distributions)  
Acyl Carnitines

#### Software

NISTMS (2019)  
MS Piano (Peptide a\Annotation)  
MS Pepsearch – General Search Utility for NIST Libraries  
MS Interpreter  
Lib2NIST – Library Conversion



<http://chemdata.nist.gov>