



PeakView软件手动拟合化学式

张海燕

zhy2009@ustc.edu.cn

中国科学技术大学生命科学实验中心

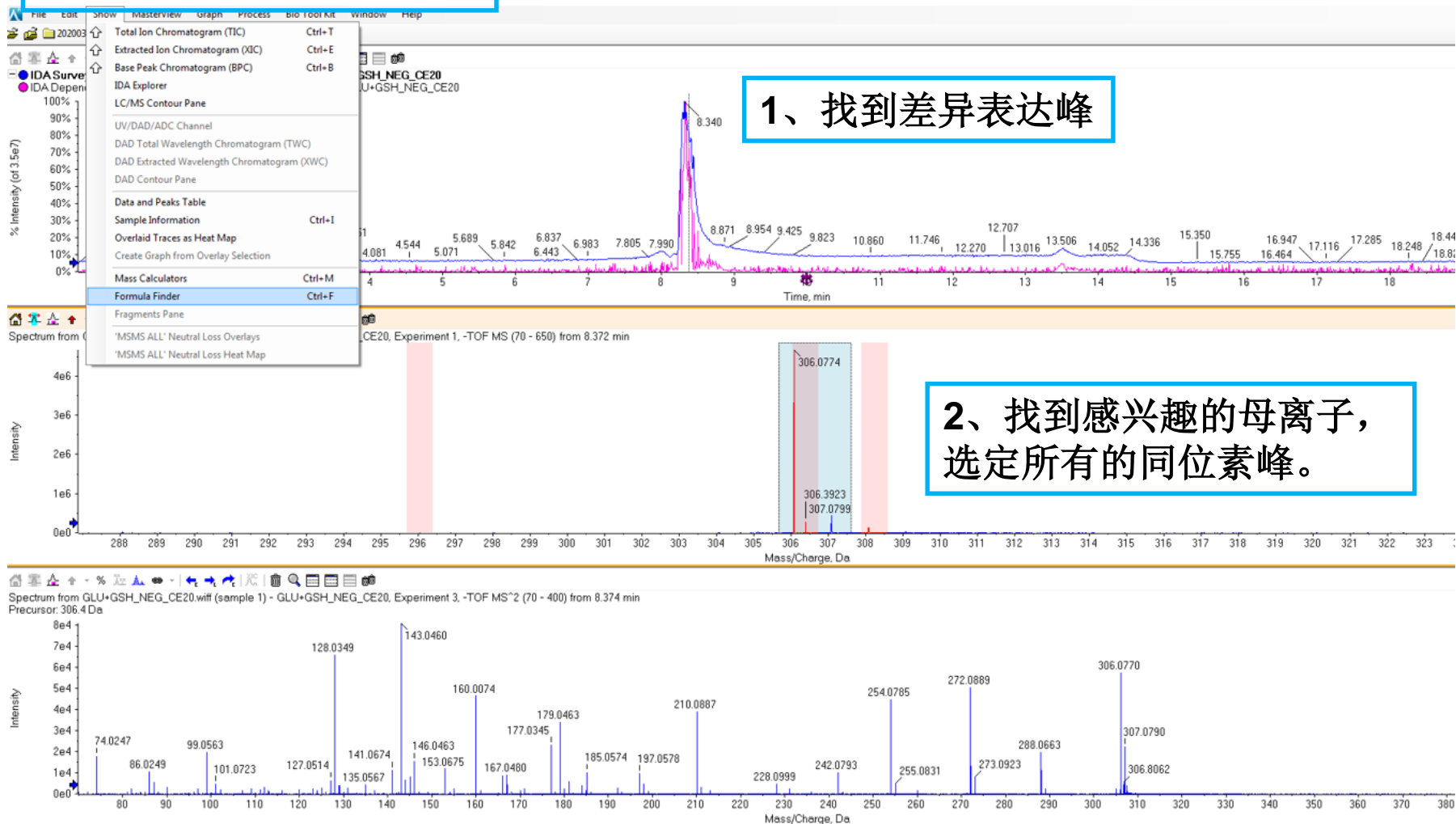
<http://biotech.ustc.edu.cn>

2020.04

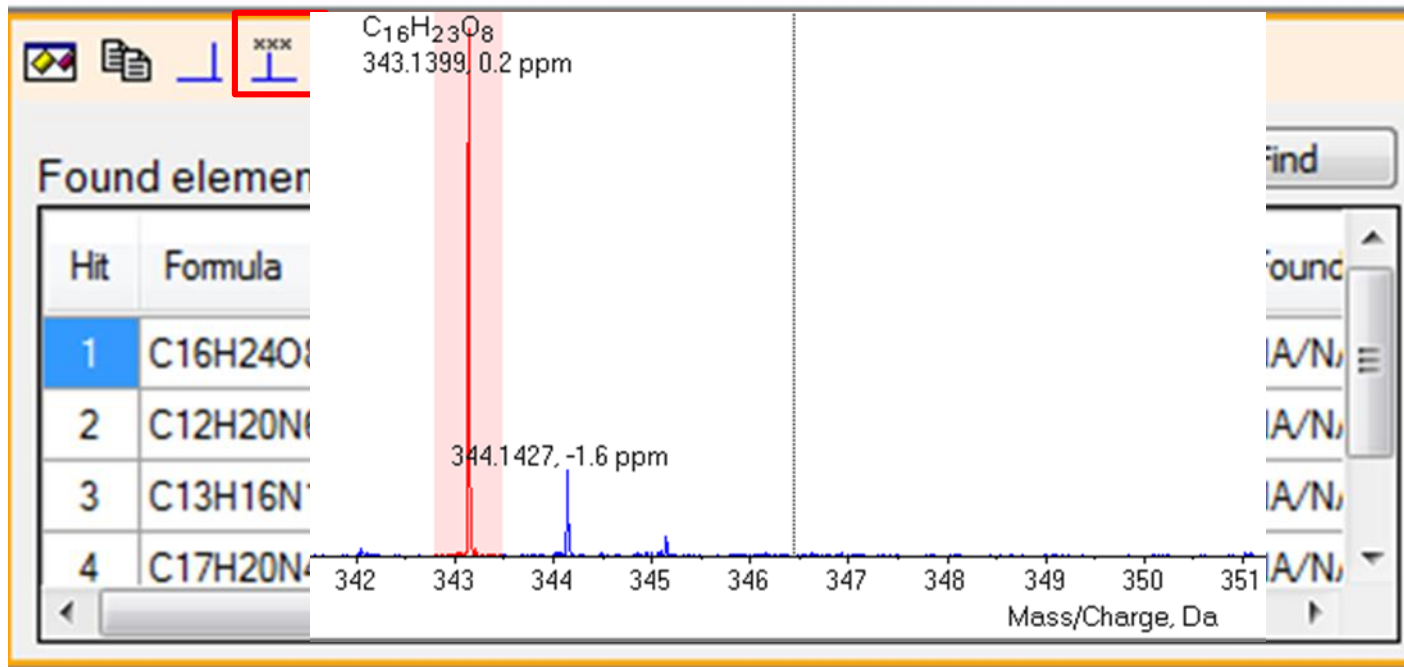


打开一个数据

3、Show-formula finder



拟合结果



点击红色图标，可在一级质谱上标记选中的formula及error

拟合无结果

Found elemental compositions

Hit	Formula	m/z	RDB	ppm	MS Rank	MSMS ppm	MSMS Rank	Found
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MS Details MSMS Details Compound Details

Isotope cluster details Charge -1

Peak	File	m/z	% Intensity	Width
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Elements from

Elements to C100H80N20O10

Mass tolerance (ppm) 10

Intensity tolerance (%) 30

#C/#heteroatoms greater than 0

PeakView

The requested action could not be completed. Make sure your data is complete and all fields contain appropriate values.

OK

100%
50%

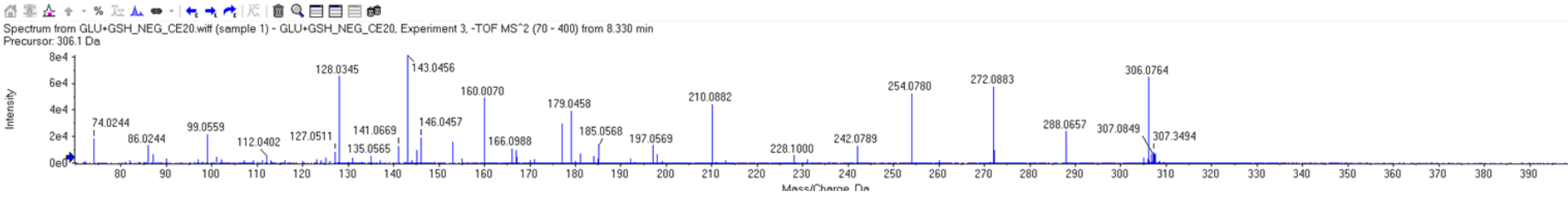
343.1399

Ion type: 4 additional ions...

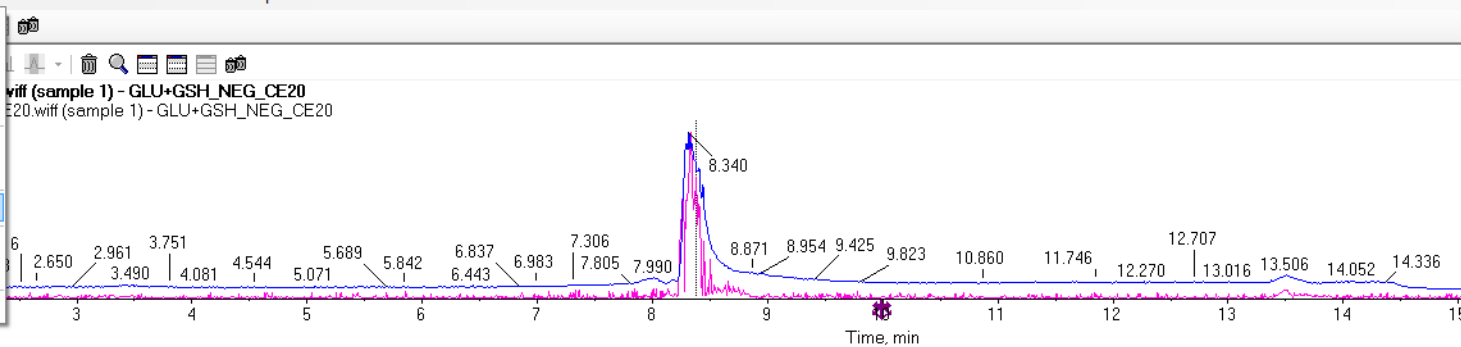
需要告诉离子加和形式



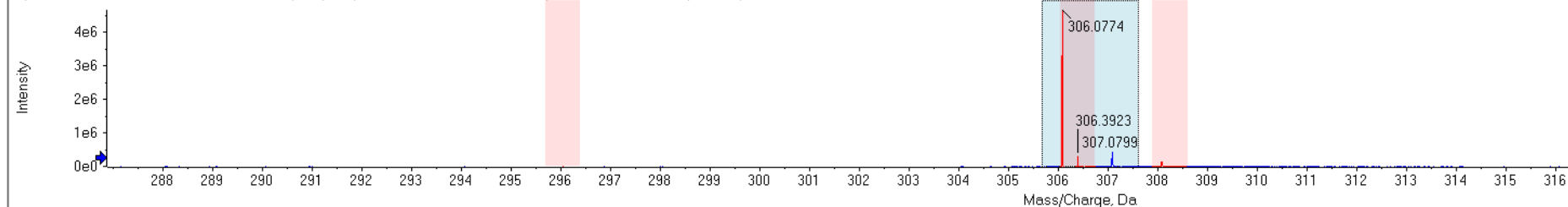
绿色代表吻合的碎片



- Open Sample... Ctrl+O
- Open Multiple Samples... Ctrl+Shift+O
- Open Heat Map TICs from Wiff...
- Find Wiff Samples Ctrl+L
- Open T2D Data
- Open Text Data
- Open Mol File...**
- Export
- Print
- Exit Alt+F4



Spectrum from GLU+GSH_NEG_CE20.wiff (sample 1) - GLU+GSH_NEG_CE20, Experiment 1, -TOF MS (70 - 650) from 8.372 min



Found elemental compositions

Hit	Formula	m/z	RDB	ppm	MS Rank	MSMS ppm	MSMS Rank	Found
1	C11H13N7O2S	306.0779	9.0	-1.5	1			NA/NA
2	C10H17N3O6S	306.0765	4.0	2.8	2			NA/NA
3	C18H13NO4	306.0772	13.0	0.7	3			NA/NA
4	C7H13N7O7	306.0804	5.0	-9.7	4 (2)			NA/NA
5	C14H9N7O2	306.0745	14.0	9.5	4 (2)			NA/NA
6	C19H9N5	306.0785	18.0	-3.7	6			NA/NA

MS Details MSMS Details Compound Details

Isotope cluster details Charge: -1

Peak Use m/z % Intensity Width

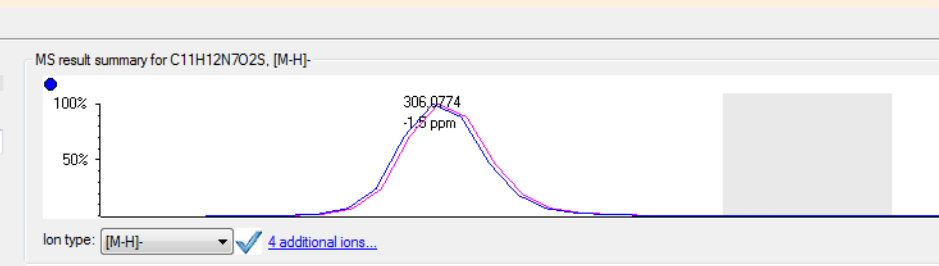
Elements from

Elements to C20H30N10O10S2

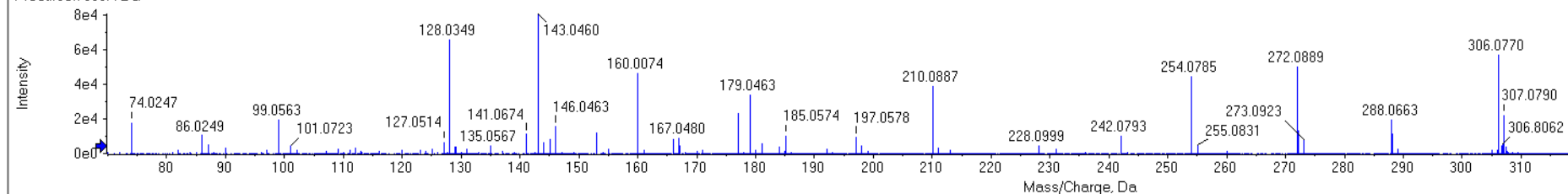
Mass tolerance (ppm) 10

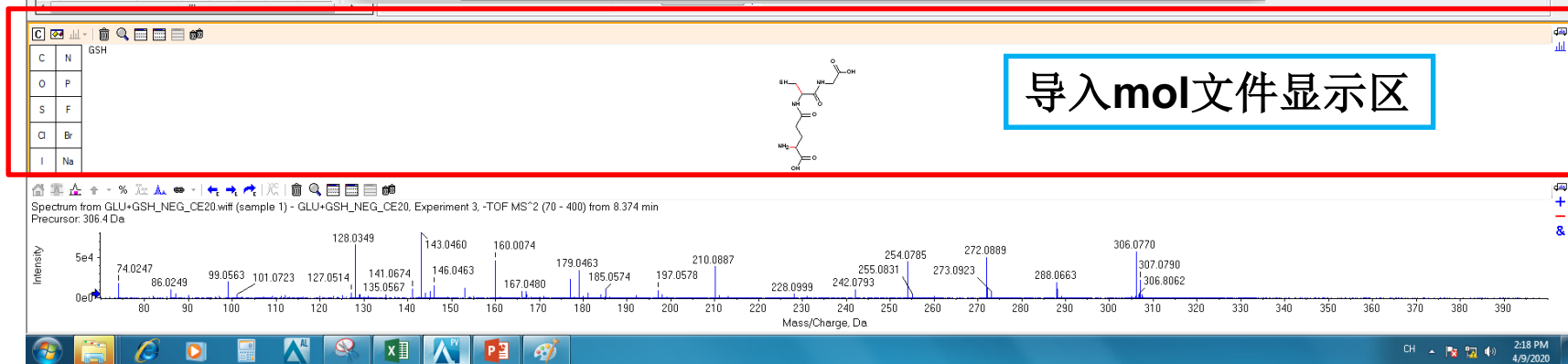
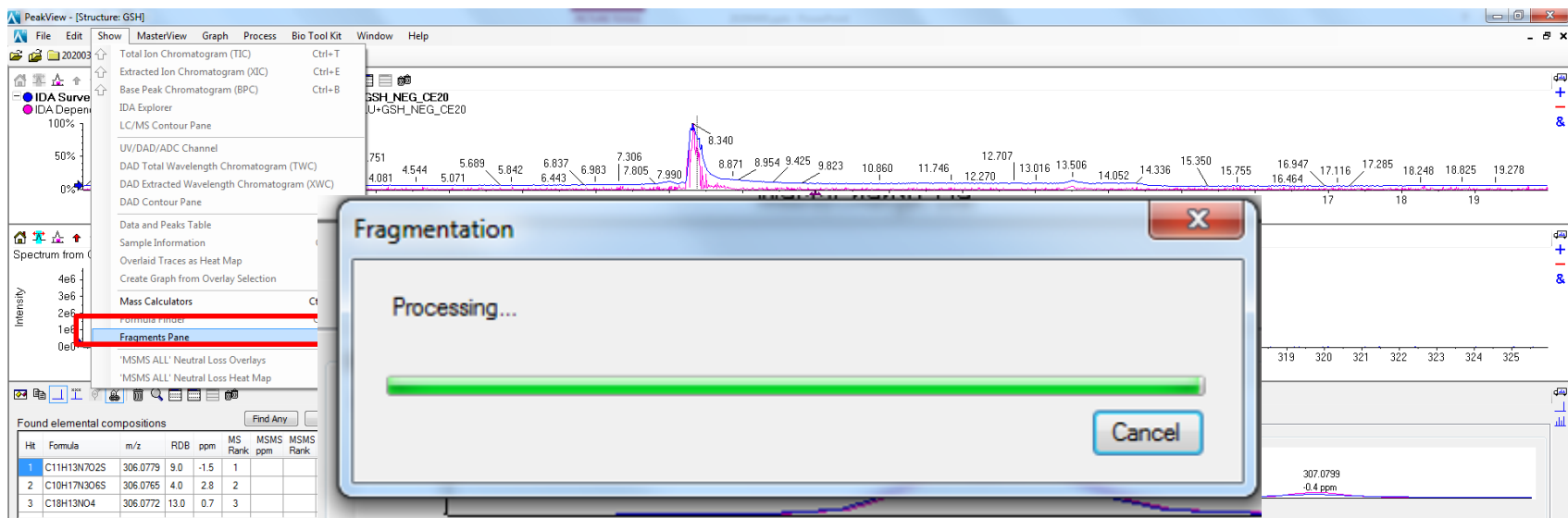
Intensity tolerance (%) 30

#C/#heteroatoms greater than 0



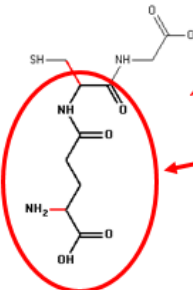
Spectrum from GLU+GSH_NEG_CE20.wiff (sample 1) - GLU+GSH_NEG_CE20, Experiment 3, -TOF MS^2 (70 - 400) from 8.374 min
Precursor: 306.4 Da





实测二级与导入化合物理论二级的匹配

GSH, selected composition: $C_3H_5N_2O_4^-$ (197.0568 Da)



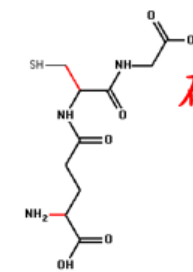
碎片加粗显示

碎片信息

m/z	Num H	Broken Bonds	Bond Closure	Rad	Rearr.	Error (ppm)	Composition
185.0568	1	3				0.0	C7H9N2O4-
185.0568	-1	3				0.0	C7H9N2O4-
197.0568	-2	2				0.8	C8H9N2O4-
197.0568	-2	2				0.8	C8H9N2O4-
198.0884	1	3				3.4	C8H12N3O3-
198.0884	1	3				3.4	C8H12N3O3-
198.0884	1	3				3.4	C8H12N3O3-
210.0884	-1	3				0.9	C9H12N3O3-
210.0884	-1	3				0.9	C9H12N3O3-
228.0990	0	2				4.3	C9H14N3O4-
228.0990	0	2				4.3	C9H14N3O4-
272.0660	-1	1				1.1	C10H14N3O5-

Num. fragments: 84

GSH, selected composition: $C_{10}H_{14}N_3O_6^-$ (272.0660 Da)



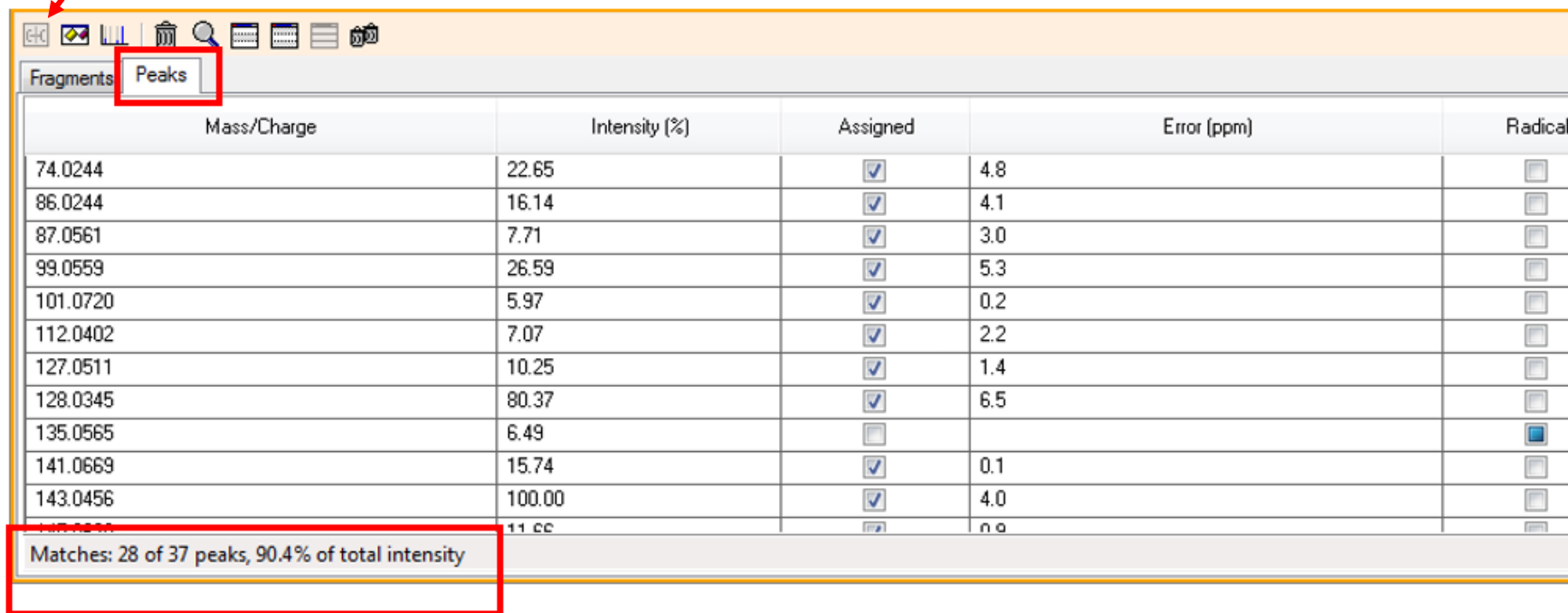
碎片加粗显示

碎片信息

m/z	Num H	Broken Bonds	Bond Closure	Rad	Rearr.	Error (ppm)	Composition
228.0990	0	2				4.3	C9H14N3O4-
228.0990	3	3				4.3	C9H14N3O4-
242.0782	0	2				2.9	C9H12N3O5-
242.0782	0	2				2.9	C9H12N3O5-
242.0782	-3	3				2.9	C9H12N3O5-
254.0782	-2	2				0.8	C10H12N3O5-
254.0782	-2	2				0.8	C10H12N3O5-
272.0660	-1	1				1.1	C10H14N3O5-
288.0660	-1	1				1.1	C10H14N3O5S-
288.0660	-1	1				1.1	C10H14N3O5S-
306.0755	0	0				0.4	C10H16N3O6S-

Num. fragments: 84

点击进行峰匹配



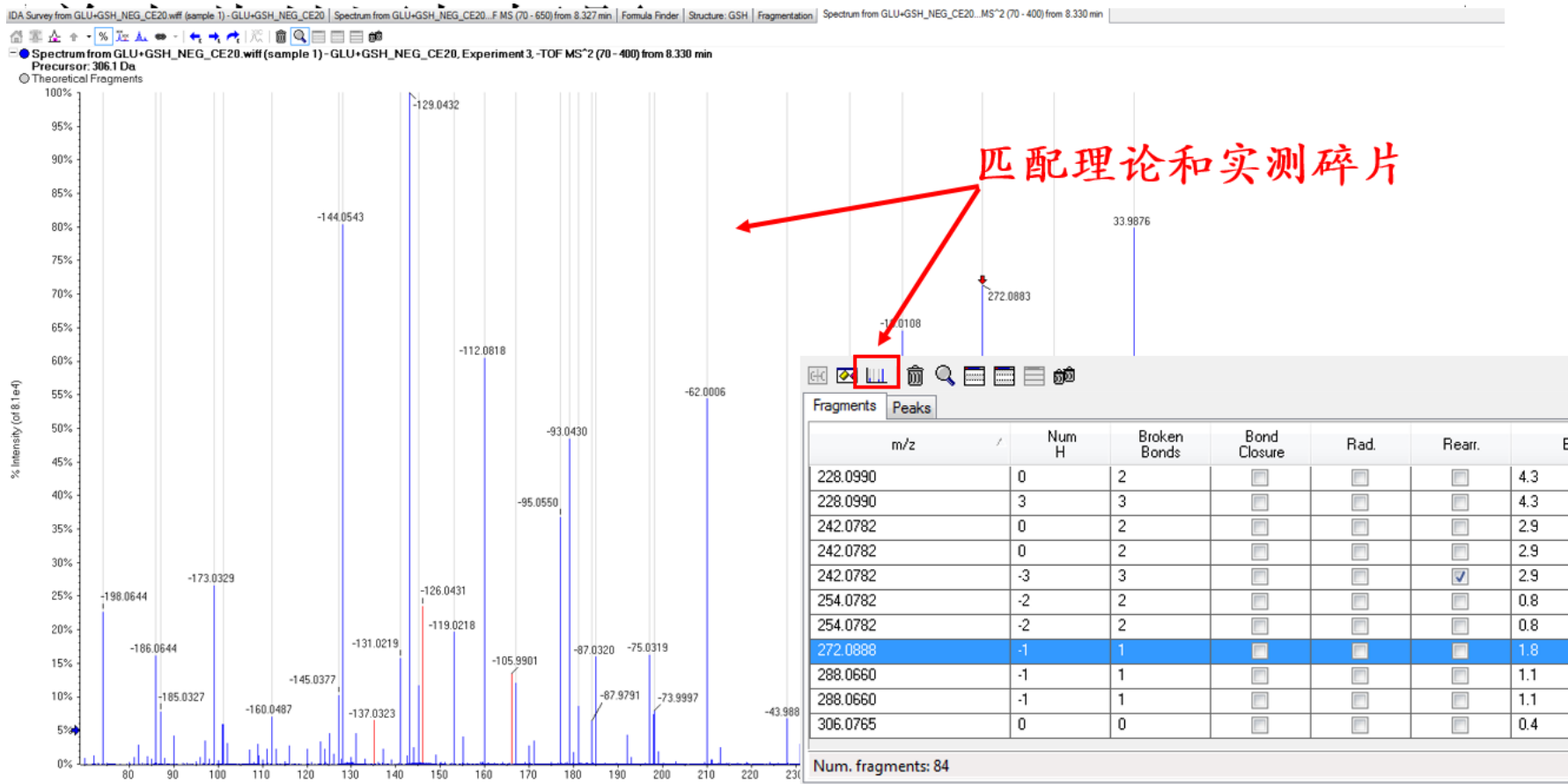
Mass/Charge	Intensity (%)	Assigned	Error (ppm)	Radical
74.0244	22.65	<input checked="" type="checkbox"/>	4.8	<input type="checkbox"/>
86.0244	16.14	<input checked="" type="checkbox"/>	4.1	<input type="checkbox"/>
87.0561	7.71	<input checked="" type="checkbox"/>	3.0	<input type="checkbox"/>
99.0559	26.59	<input checked="" type="checkbox"/>	5.3	<input type="checkbox"/>
101.0720	5.97	<input checked="" type="checkbox"/>	0.2	<input type="checkbox"/>
112.0402	7.07	<input checked="" type="checkbox"/>	2.2	<input type="checkbox"/>
127.0511	10.25	<input checked="" type="checkbox"/>	1.4	<input type="checkbox"/>
128.0345	80.37	<input checked="" type="checkbox"/>	6.5	<input type="checkbox"/>
135.0565	6.49	<input type="checkbox"/>		<input checked="" type="checkbox"/>
141.0669	15.74	<input checked="" type="checkbox"/>	0.1	<input type="checkbox"/>
143.0456	100.00	<input checked="" type="checkbox"/>	4.0	<input type="checkbox"/>
145.0236	11.66	<input checked="" type="checkbox"/>	0.0	<input type="checkbox"/>

Matches: 28 of 37 peaks, 90.4% of total intensity

匹配分值查看



实测二级与导入化合物理论二级的匹配



化合物断裂规律参数

m/z	Num H	Broken Bonds	Bond Closure
228.0990	0	2	<input type="checkbox"/>
228.0990	3	3	<input type="checkbox"/>
242.0782	0	2	<input type="checkbox"/>
242.0782	0	2	<input type="checkbox"/>
242.0782	-3	3	<input type="checkbox"/>
254.0782	-2	2	<input type="checkbox"/>
254.0782	-2	2	<input type="checkbox"/>
272.0888	-1	1	<input type="checkbox"/>
288.0660	-1	1	<input type="checkbox"/>
288.0660	-1	1	<input type="checkbox"/>
306.0765	0	0	<input type="checkbox"/>

Fragment Options

Fragmentation

- Only break single bonds
- Break ring bonds
- Maximum number of bonds to break: 4
- Maximum number of C-C bonds to break: 4
- Also break C-C bond if either carbon is bonded to a hetero atom
- Allow one bond closure (double bond formation)
- Include brute force rearrangements
- Allow radicals

Peak List

Mass tolerance: 20.0 ppm

- Constrain using peak list
- Require evidence for previous step when breaking bond

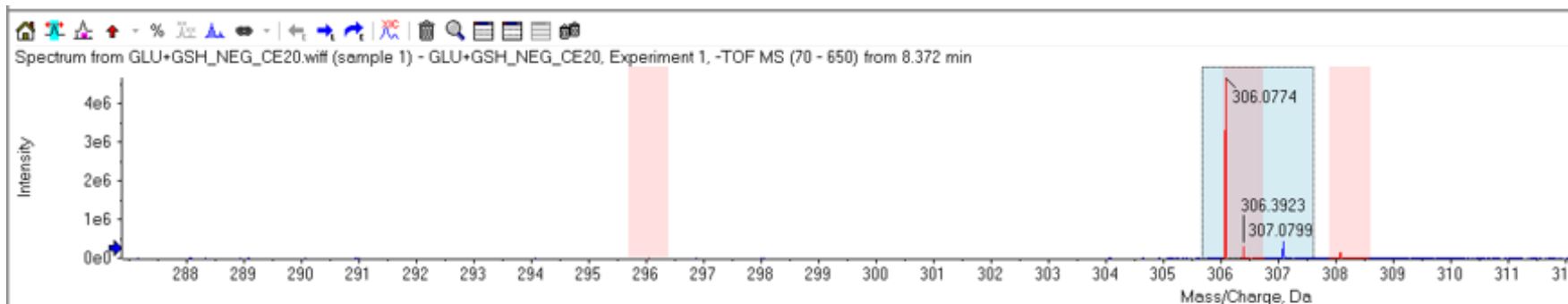
Display

Do not show fragments with m/z less than 40.0 Da

- Automatically recalculate on-the-fly

OK Cancel

注意事项



Found elemental compositions

Hit	Formula	m/z	RDB	ppm	MS Rank	MSMS ppm	MSMS Rank	Found
1	C11H13N7O2S	306.0779	9.0	-1.5	1			NA/NA
2	C10H17N3O6S	306.0765	4.0	2.8	2			NA/NA
3	C18H13NO4	306.0772	13.0	0.7	3			NA/NA
4	C7H13N7O7	306.0804	5.0	-9.7	4 (2)			NA/NA
5	C14H9N7O2	306.0745	14.0	9.5	4 (2)			NA/NA

MS Details

Isotope cluster details Charge: -1

Elements to: **C20H30N10O10S2**

Mass tolerance (ppm): 10

Intensity tolerance (%): 30

MS result summary for C11H13N7O2S, [M-H]⁻

Found elemental compositions

Hit	Formula	m/z	RDB	ppm	MS Rank	MSMS ppm	MSMS Rank	Found
1	C18H13NO4	306.0772	13.0	-0.3	1 (2)	15....	1	NA/NA
2	C14H9N7O2	306.0745	14.0	8.5	1 (2)	17....	2	NA/NA
3	C19H9N5	306.0785	18.0	-4.6	3	38....	3	NA/NA

MS Details

Isotope cluster details Charge: -1

Peak	Use	m/z	% Intensity	Width
0	<input checked="" type="checkbox"/>	306.0771	100.0	0.008
1	<input checked="" type="checkbox"/>	307.0796	17.5	0.011

Elements to: **C20H30N10O10**

Mass tolerance (ppm): 10

Intensity tolerance (%): 30

#C/#heteroatoms greater than: 0

MS result summary for C18H12NO4, [M-H]⁻

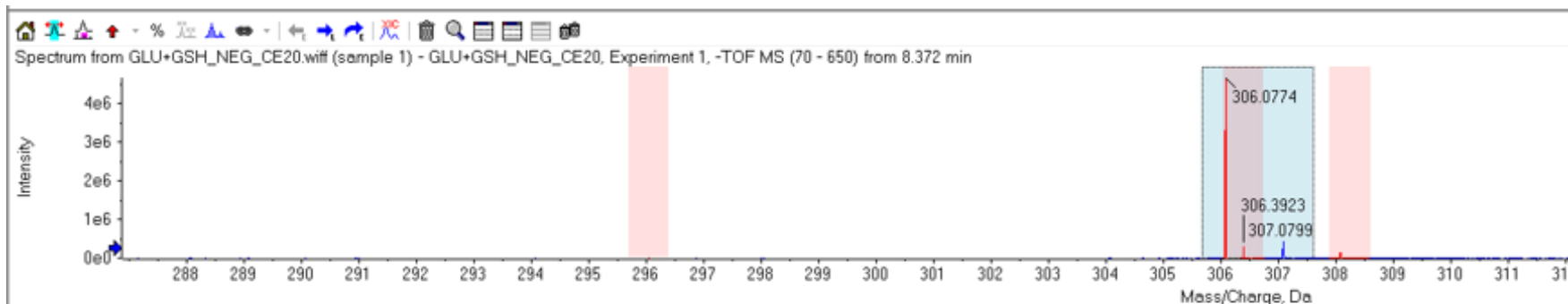
靶标

拟合formula不同

给定元素组成不同



注意事项



Found elemental compositions

Hit	Formula	m/z	RDB	ppm	MS Rank	MSMS ppm	MSMS Rank	Found
1	C11H13N7O2S	306.0779	9.0	-1.5	1			NA/NA
2	C10H17N3O6S	306.0765	4.0	2.8	2			NA/NA
3	C18H13NO4	306.0772	13.0	0.7	3			NA/NA
4	C7H13N7O7	306.0804	5.0	-9.7	4 (2)			NA/NA
5	C14H9N7O2	306.0745	14.0	9.5	4 (2)			NA/NA

MS Details

Isotope cluster details Charge: -1

Elements to: C20H30N10O10S2

Mass tolerance (ppm): 10

Intensity tolerance (%): 30

MS result summary for C11H13N7O2S, [M-H]⁻

Found elemental compositions

Hit	Formula	m/z	RDB	ppm	MS Rank	MSMS ppm	MSMS Rank	Found
1	C18H13NO4	306.0772	13.0	-0.3	1 (2)	15....	1	NA/NA
2	C14H9N7O2	306.0745	14.0	8.5	1 (2)	17....	2	NA/NA
3	C19H9N5	306.0785	18.0	-4.6	3	38....	3	NA/NA

MS Details

Isotope cluster details Charge: -1

Peak	Use	m/z	% Intensity	Width
0	<input checked="" type="checkbox"/>	306.0771	100.0	0.008
1	<input checked="" type="checkbox"/>	307.0796	17.5	0.011

Elements to: C20H30N10O10

Mass tolerance (ppm): 10

Intensity tolerance (%): 30

#C/#heteroatoms greater than: 0

MS result summary for C18H12NO4, [M-H]⁻

靶标

拟合formula不同

给定元素组成不同



