

Fragment Ion View

Project name: 20200330COV19
Sample name: 20200416_COV19-double_2
Created date: 2020-04-18 15:22:56.0
Search name:
Activation type: CID

Peptide sequence: R.SSVLHSTQDLFLPFFSNVT(203.079373)WF.H
Scan number: 56810
Calculated M+H: 2675.3027
Measured M+H: 2675.3083
Charge state: 2
File name: 20200416_COV19-double_2
search type: light

Ions:
a ☐1+ ☐2+ ☐3+
b ☒1+ ☐2+ ☐3+
c ☐1+ ☐2+ ☐3+
x ☐1+ ☐2+ ☐3+
y ☒1+ ☐2+ ☐3+
z ☐1+ ☐2+ ☐3+
[\[Deselect All\]](#)

Neutral Loss:
☐ H₂O (o)
☐ NH₃ (*)
☐ H₃PO₄ (^)
☐ Precursor (#)
☐ Precursor with reduced charges

Mass Type:
☒ Mono ☐ Avg
Mass Tol:
☐ Dalton ☒ PPM

Update

Peak Assignment:
☒ Most Intense
☐ Nearest Match
☐ Peak Detect
☐ Precursor Removal

Peak Labels:
☒ Ion ☐ m/z
☐ None

Width: 750

Height: 450

SSVLHSTQDLFLPFFSNVTWF, MH+ 2675.3028, m/z 1338.1550
File: 20200416_COV19-double_2, Scan: 56810, Precursor m/z: 1338.1578, Charge: 2

Click and drag in the plot to zoom X: ☒ Y: ☐

Zoom Out

Print

☐ Enable tooltip

b+	#	Seq	#	y+
88.0393	1	S	21	
175.0713	2	S	20	2588.2708
274.1397	3	V	19	2501.2387
387.2238	4	L	18	2402.1703
524.2827	5	H	17	2289.0863
611.3148	6	S	16	2152.0274
712.3624	7	T	15	2064.9953
840.4210	8	Q	14	1963.9476
955.4480	9	D	13	1835.8891
1068.5320	10	L	12	1720.8621
1215.6004	11	F	11	1607.7781
1328.6845	12	L	10	1460.7096
1425.7373	13	P	9	1347.6256
1572.8057	14	F	8	1250.5728
1719.8741	15	F	7	1103.5044
1806.9061	16	S	6	956.4360
1920.9490	17	N	5	869.4040
2020.0175	18	V	4	755.3610
2324.1445	19	T	3	656.2926
2510.2238	20	W	2	352.1656
	21	F	1	166.0863

[\[Click\]](#) to move table
Static Modifications:
C: 57.02146
Variable Modifications:
T: 203.079373