

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: F.SALEPLVDLPIGINIT(203.079373).R
Scan number: 56315
Calculated M+H: 1868.0262
Measured M+H: 1868.0323
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

Peak Assignment:

- ☒ Most Intense
☐ Nearest Match
☐ Peak Detect
☐ Precursor Removal

Peak Labels:

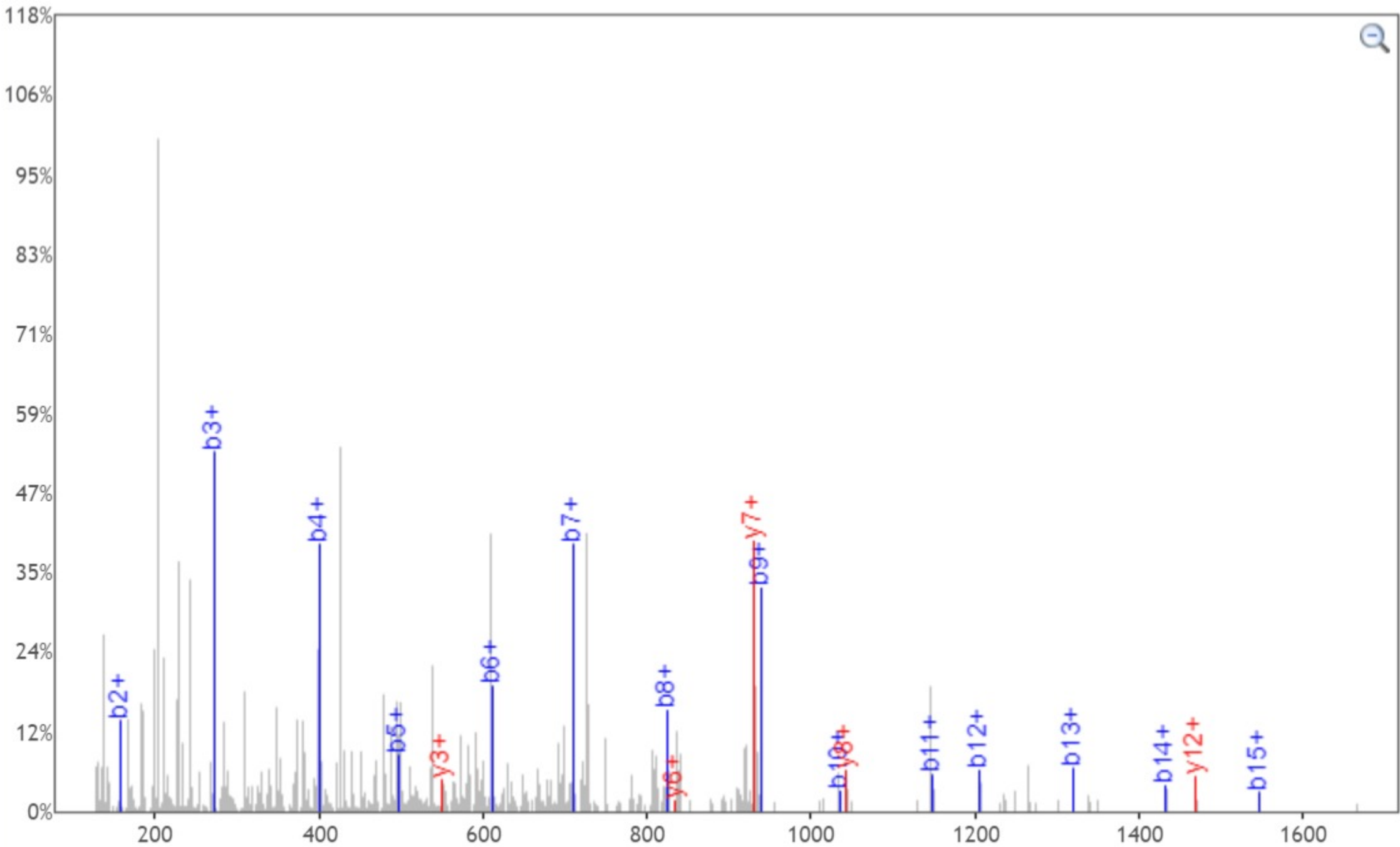
☒ Ion ☐ m/z
☐ None

Width:

Height:

SALEPLVDLPIGINIT, MH+ 1868.0263, m/z 934.5168

File: 20200416_COV19-double_2, Scan: 56315, Precursor m/z: 934.51984, Charge: 2



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

☐ Enable tooltip

b+	#	Seq	#	y+
88.0393	1	S	16	
159.0764	2	A	15	1780.9943
272.1605	3	L	14	1709.9571
401.2031	4	E	13	1596.8731
498.2558	5	P	12	1467.8305
611.3399	6	L	11	1370.7777
710.4083	7	V	10	1257.6937
825.4353	8	D	9	1158.6252
938.5193	9	L	8	1043.5983
1035.5721	10	P	7	930.5142
1148.6562	11	I	6	833.4615
1205.6776	12	G	5	720.3774
1318.7617	13	I	4	663.3559
1432.8046	14	N	3	550.2719
1545.8887	15	I	2	436.2290
	16	T	1	323.1449

[\[Click\]](#) to move table

Static Modifications:

C: 57.02146

Variable Modifications:

T: 203.079373