

Fragment Ion View

Project name: 20200330COV19
Sample name: 20200416_COV19-double_1
Created date: 2020-04-18 15:22:56.0
Search name:
Activation type: CID

Peptide sequence: R.EGVFVS(203.079373)NGTHWF.V
Scan number: 52184
Calculated M+H: 1582.7173
Measured M+H: 1582.723
Charge state: 2
File name: 20200416_COV19-double_1
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

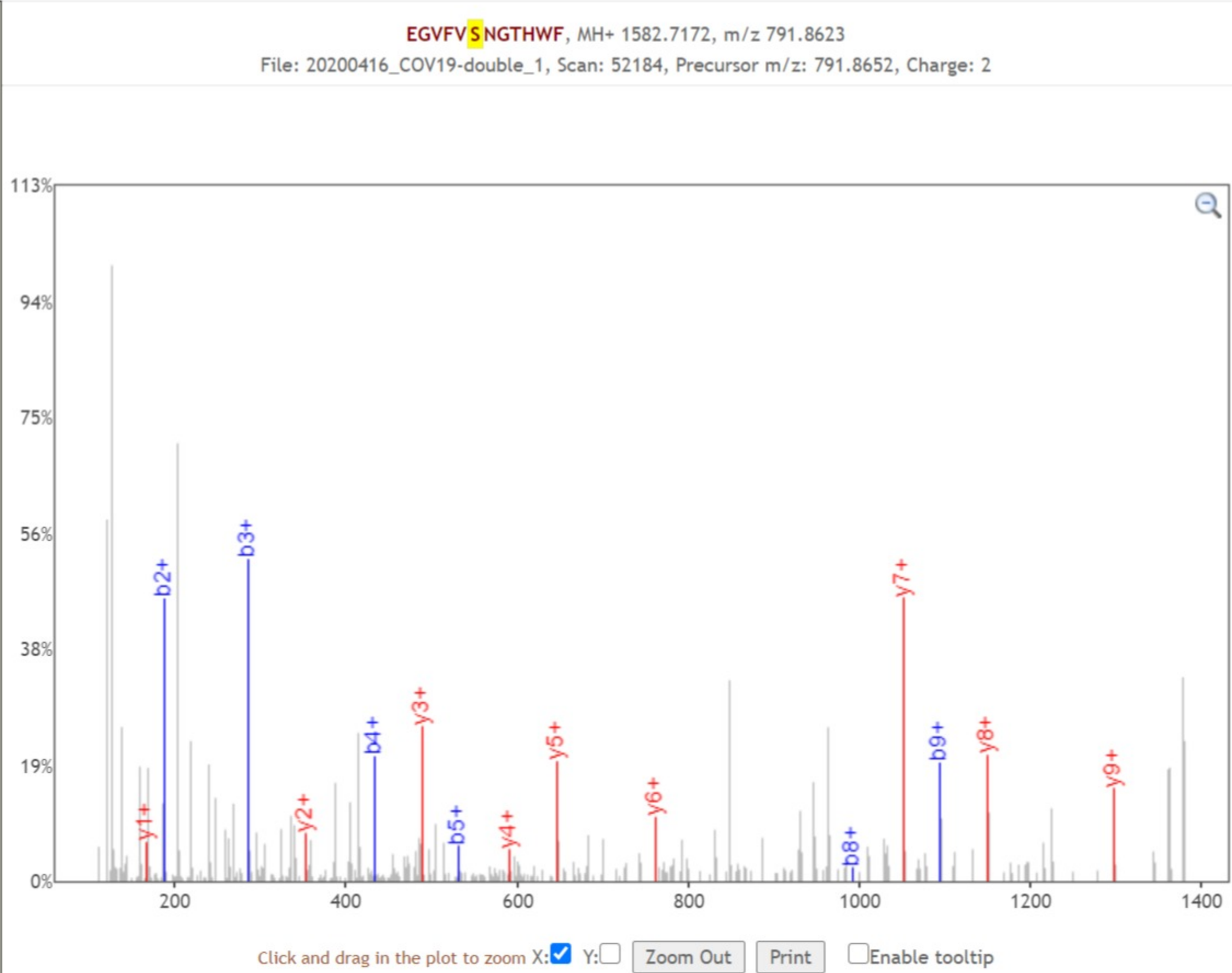
Mass Type:
☒ Mono ☐ Avg
Mass Tol:
☐ Dalton ☒ PPM

Update

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

Peak Labels:
☒ Ion ☐ m/z
☐ None

Width:
Height:



b+	#	Seq	#	y+
130.0499	1	E	12	
187.0713	2	G	11	1453.6747
286.1397	3	V	10	1396.6532
433.2082	4	F	9	1297.5848
532.2766	5	V	8	1150.5164
822.3880	6	S	7	1051.4480
936.4309	7	N	6	761.3365
993.4524	8	G	5	647.2936
1094.5000	9	T	4	590.2722
1231.5590	10	H	3	489.2245
1417.6383	11	W	2	352.1656
	12	F	1	166.0863

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