

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.EGVFVS(203.079373)NGTHWF.V
Scan number: 52170
Calculated M+H: 1582.7173
Measured M+H: 1582.7219
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

☒☐☐

z

☐☐☐

[Deselect All]

Neutral Loss:

☐

 H₂O (o)

☐

 NH₃ (*)☐☐☐

Mass Type:

☒

 Mono

☐

 Avg

Mass Tol:

20

☐

 Dalton

☒

 PPM

Update

Peak Assignment:

☒

 Most Intense

☐

 Nearest Match

☐

 Peak Detect

☐

 Precursor Removal

Peak Labels:

☒

 Ion

☐

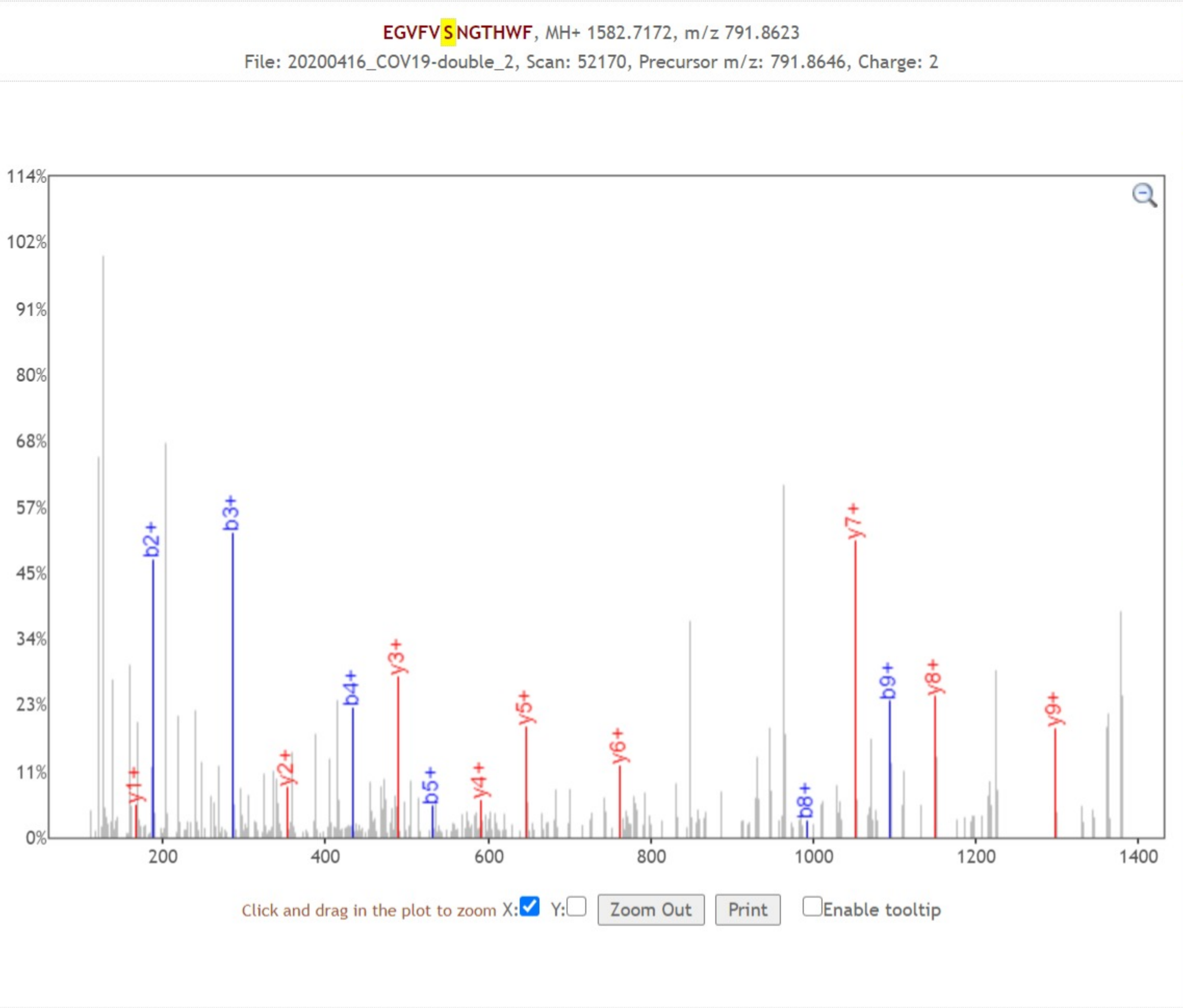
 m/z

☐

 None

Width: 750

Height: 450



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