

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.EGVFVSN(2.988261)GTHWF.V  
Scan number: 53355  
Calculated M+H: 1382.6261  
Measured M+H: 1382.6302  
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:  
☐ H<sub>2</sub>O (o)  
☐ NH<sub>3</sub> (\*)  
☐ H<sub>3</sub>PO<sub>4</sub> (^)  
☐ Precursor (#)  
☐ Precursor with reduced charges

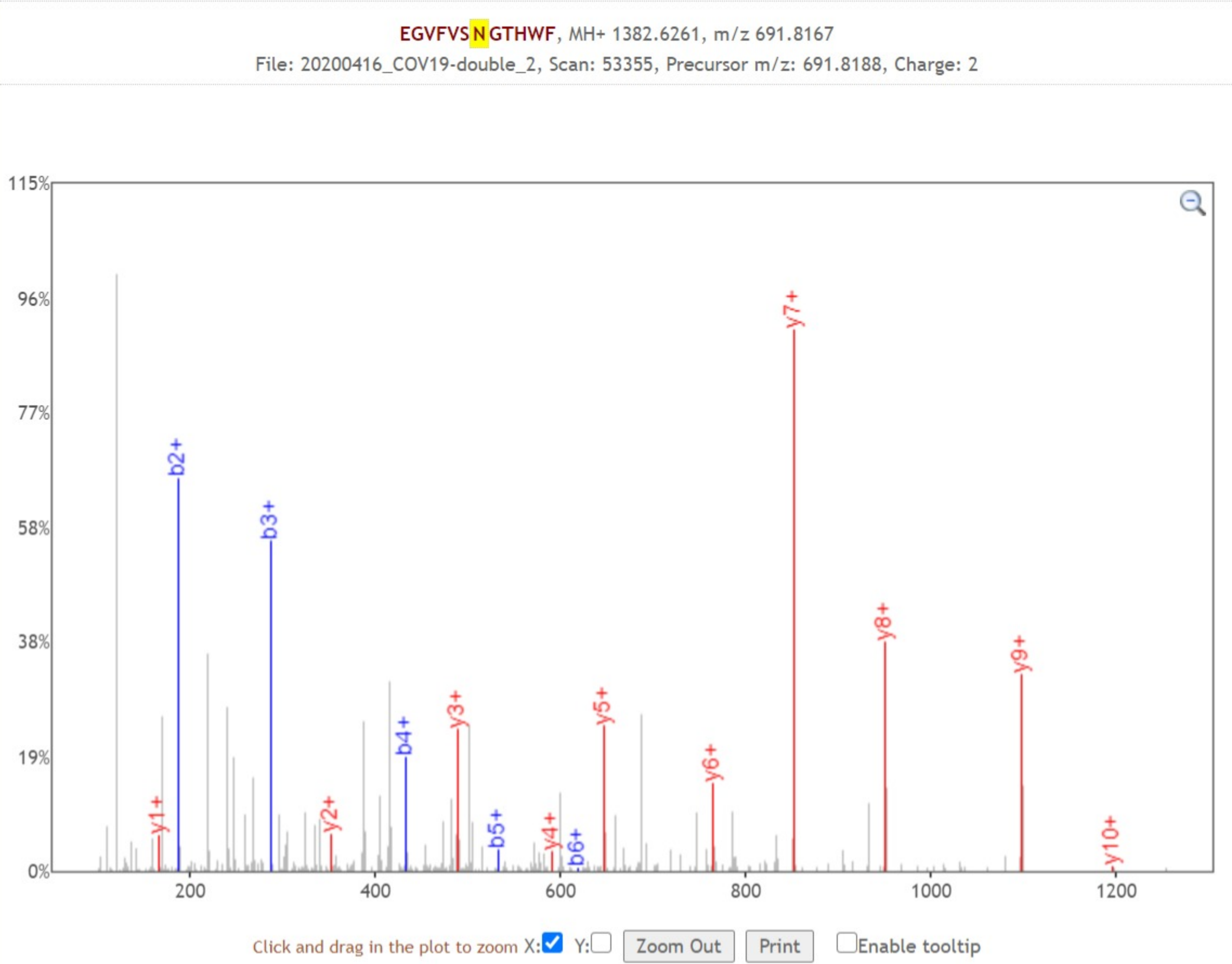
Mass Type:  
☒ Mono ☐ Avg  
Mass Tol:   
☐ Dalton ☒ PPM  

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	1253.5835
286.1397	3	V	10	1196.5621
433.2082	4	F	9	1097.4937
532.2766	5	V	8	950.4253
619.3086	6	S	7	851.3568
736.3398	7	N	6	764.3248
793.3613	8	G	5	647.2936
894.4089	9	T	4	590.2722
1031.4678	10	H	3	489.2245
1217.5472	11	W	2	352.1656
	12	F	1	166.0863

[\[Click\]](#) to move table  
Static Modifications:  
C: 57.02146  
Variable Modifications:  
N: 2.988261