

## 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.EGVFVSN(2.988261)GTHWF.V

Scan number: 53387

Calculated M+H: 1382.6261

Measured M+H: 1382.631

Charge state: 2

File name: 20200416\_COV19-double\_1

search type: light

**Ions:**

a ☐ 1<sup>+</sup> ☐ 2<sup>+</sup> ☐ 3<sup>+</sup>

b ☒ 1<sup>+</sup> ☐ 2<sup>+</sup> ☐ 3<sup>+</sup>

c ☐ 1<sup>+</sup> ☐ 2<sup>+</sup> ☐ 3<sup>+</sup>

x ☐ 1<sup>+</sup> ☐ 2<sup>+</sup> ☐ 3<sup>+</sup>

y ☒ 1<sup>+</sup> ☐ 2<sup>+</sup> ☐ 3<sup>+</sup>

z ☐ 1<sup>+</sup> ☐ 2<sup>+</sup> ☐ 3<sup>+</sup>

[\[Deselect All\]](#)

**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

**Peak Assignment:**

☒ Most Intense

☐ Nearest Match

☐ Peak Detect

☐ Precursor Removal

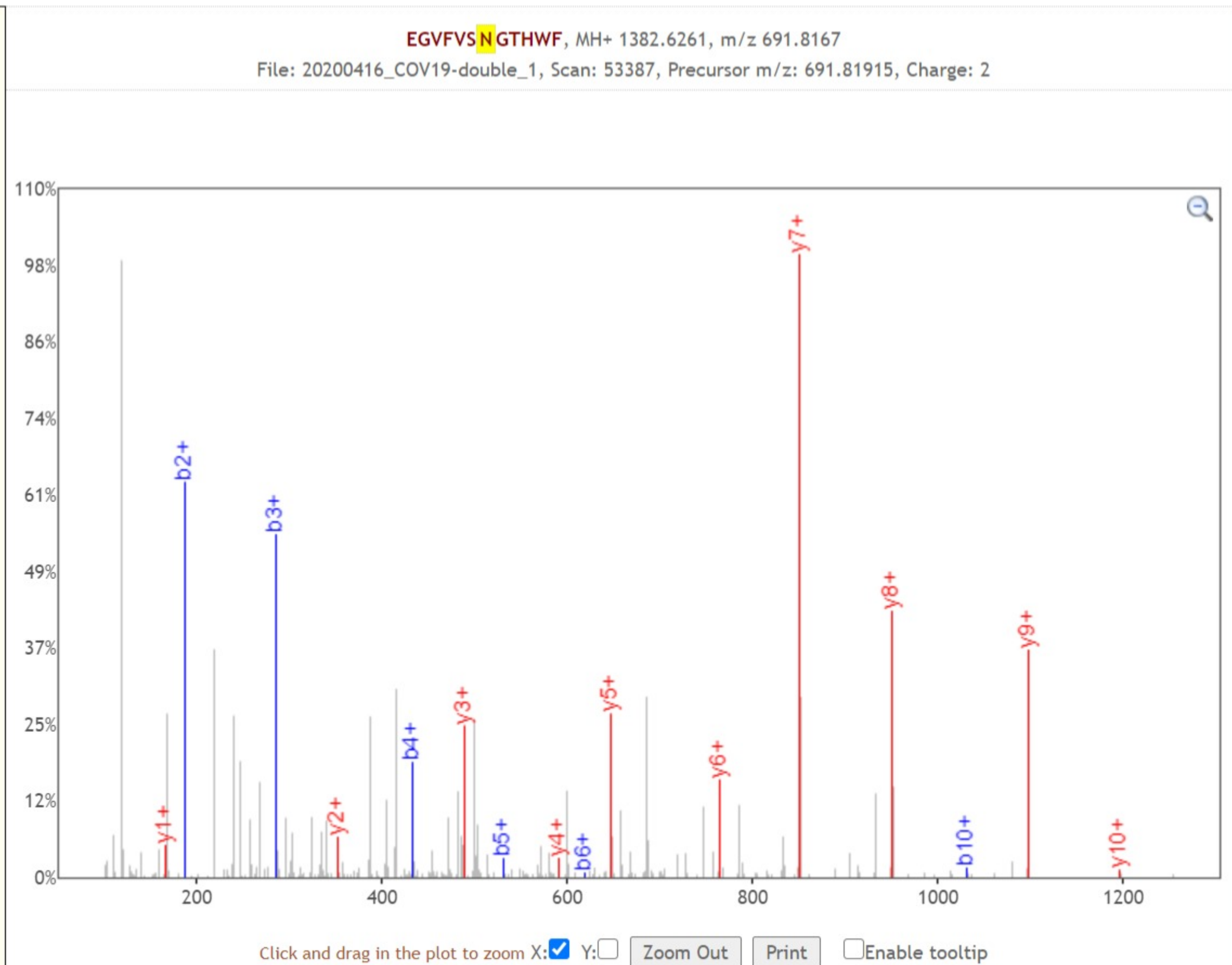
**Peak Labels:**

☒ Ion ☐ m/z

☐ None

Width:

Height:



b <sup>+</sup>	#	Seq	#	y <sup>+</sup>
130.0499	1	E	12	
<b>187.0713</b>	2	G	11	1253.5835
<b>286.1397</b>	3	V	10	<b>1196.5621</b>
<b>433.2082</b>	4	F	9	<b>1097.4937</b>
<b>532.2766</b>	5	V	8	<b>950.4253</b>
<b>619.3086</b>	6	S	7	<b>851.3568</b>
736.3398	7	N	6	<b>764.3248</b>
793.3613	8	G	5	<b>647.2936</b>
894.4089	9	T	4	<b>590.2722</b>
<b>1031.4678</b>	10	H	3	<b>489.2245</b>
1217.5472	11	W	2	<b>352.1656</b>
	12	F	1	<b>166.0863</b>

[Click] to move table

### Static Modifications:

C: 57.02146

**Variable Modifications:**

N: 2.988261