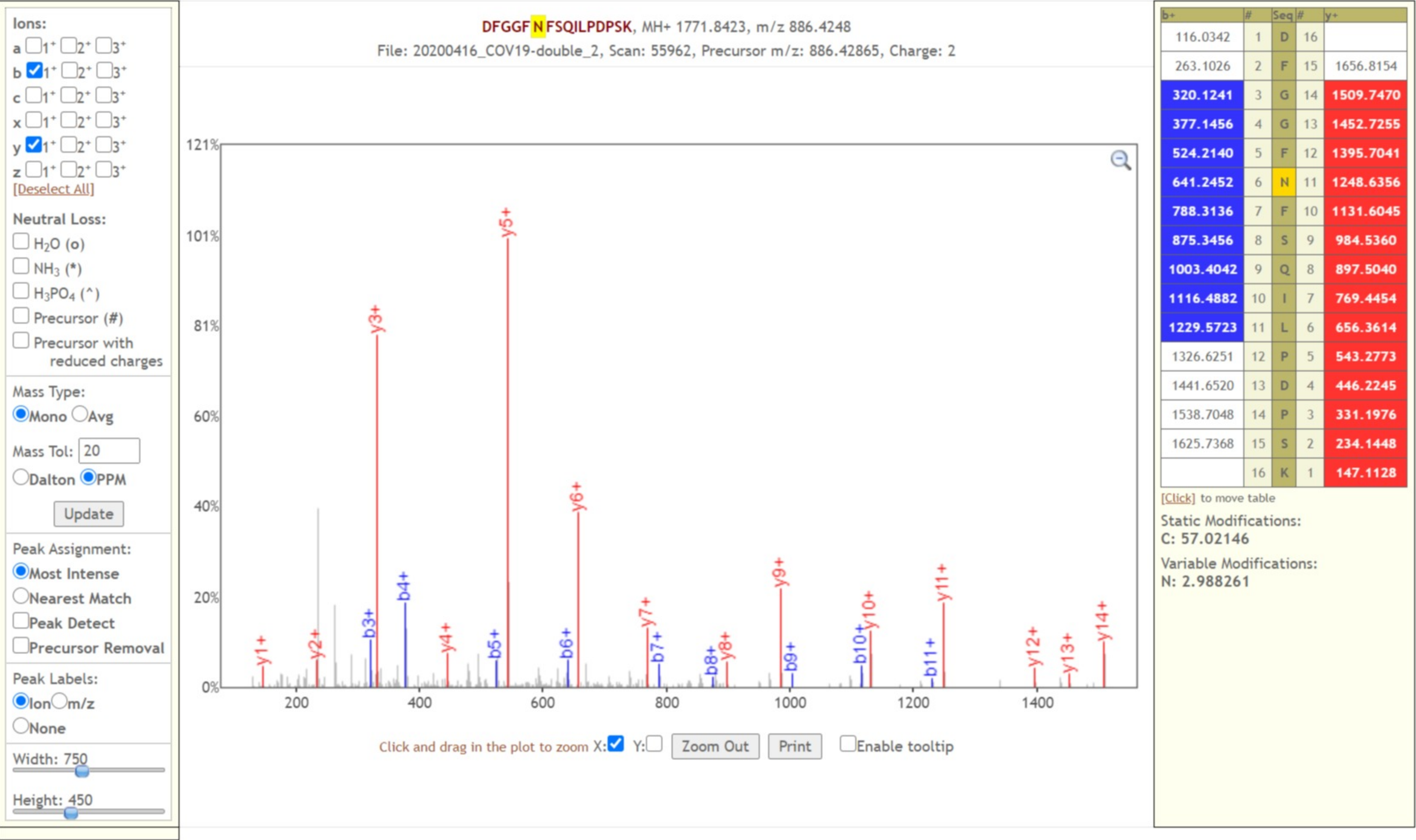


Fragment Ion View

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: K.DFGGFN(2.988261)FSQILPDPSK.P
Scan number: 55962
Calculated M+H: 1771.8424
Measured M+H: 1771.85
Charge state: 2
File name: 20200416_COV19-double_2
search type: light



z

☐

1⁺

☐

2⁺

☐

3⁺

[Deselect All]

Neutral Loss:

☐ H₂O (o)

☐ NH₃ (*)

☐ H₃PO₄ (^)

☐ Precursor (#)

☐ Precursor with reduced charges

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

DFGGF**N**FSQILPDPSK, MH+ 1771.8423, m/z 886.4248

File: 20200416_COV19-double_2, Scan: 55962, Precursor m/z: 886.42865, Charge: 2

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

Zoom Out

Print

☐ Enable tooltip

b+	#	Seq	#	y+
116.0342	1	D	16	
263.1026	2	F	15	1656.8154
320.1241	3	G	14	1509.7470
377.1456	4	G	13	1452.7255
524.2140	5	F	12	1395.7041
641.2452	6	N	11	1248.6356
788.3136	7	F	10	1131.6045
875.3456	8	S	9	984.5360
1003.4042	9	Q	8	897.5040
1116.4882	10	I	7	769.4454
1229.5723	11	L	6	656.3614
1326.6251	12	P	5	543.2773
1441.6520	13	D	4	446.2245
1538.7048	14	P	3	331.1976
1625.7368	15	S	2	234.1448
	16	K	1	147.1128

[Click] to move table

Static Modifications:
C: 57.02146

Variable Modifications:
N: 2.988261