

**Search parameter**

Instrument type: MALDI-TOF-TOF

MS Tol.: 200 ppm

MSMS Tol.: 0.4 Da

Charge=1+

Mass value: Monoisotopic

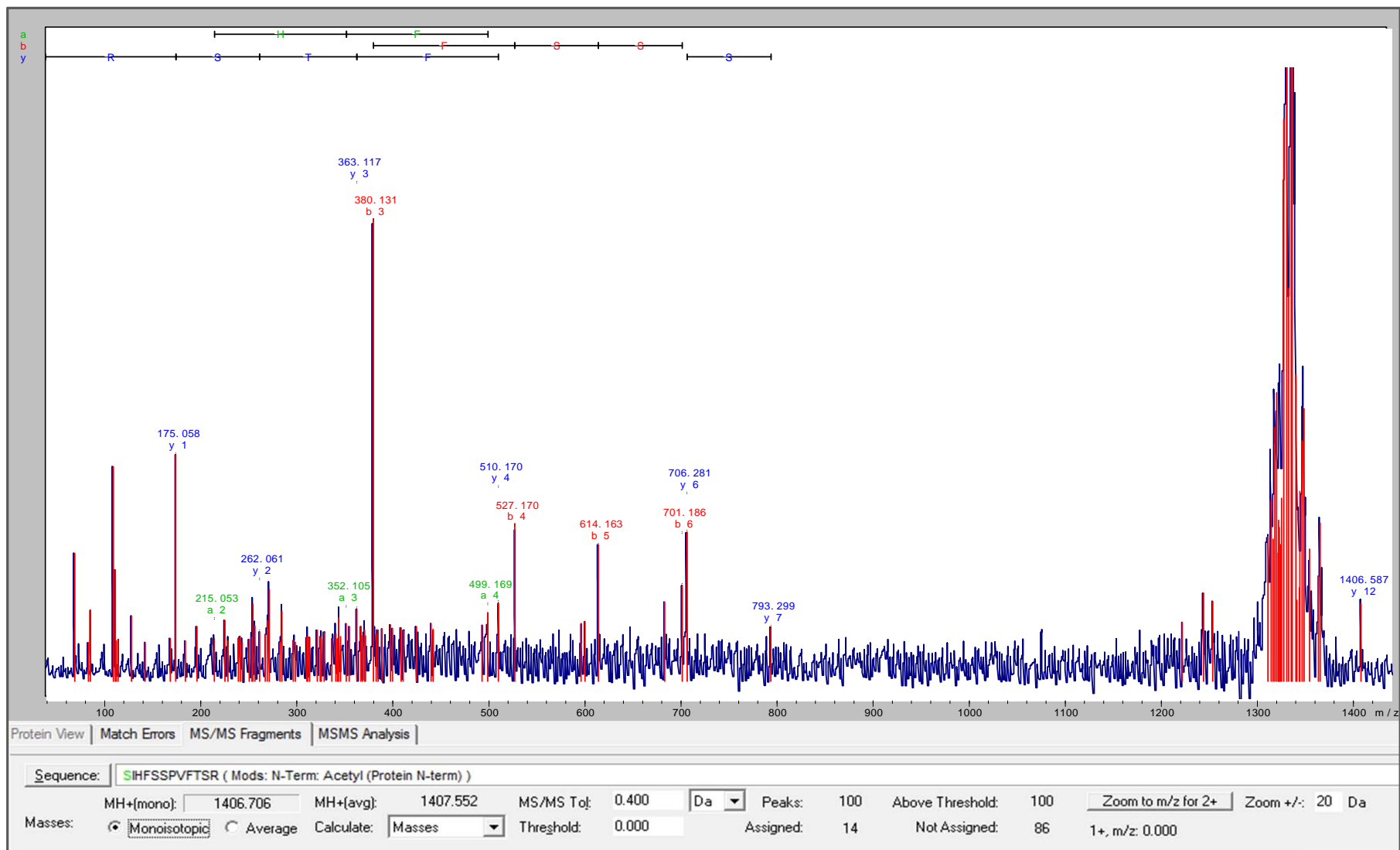
Mascot 2.4

SwissProt: SwissProt 2014\_10

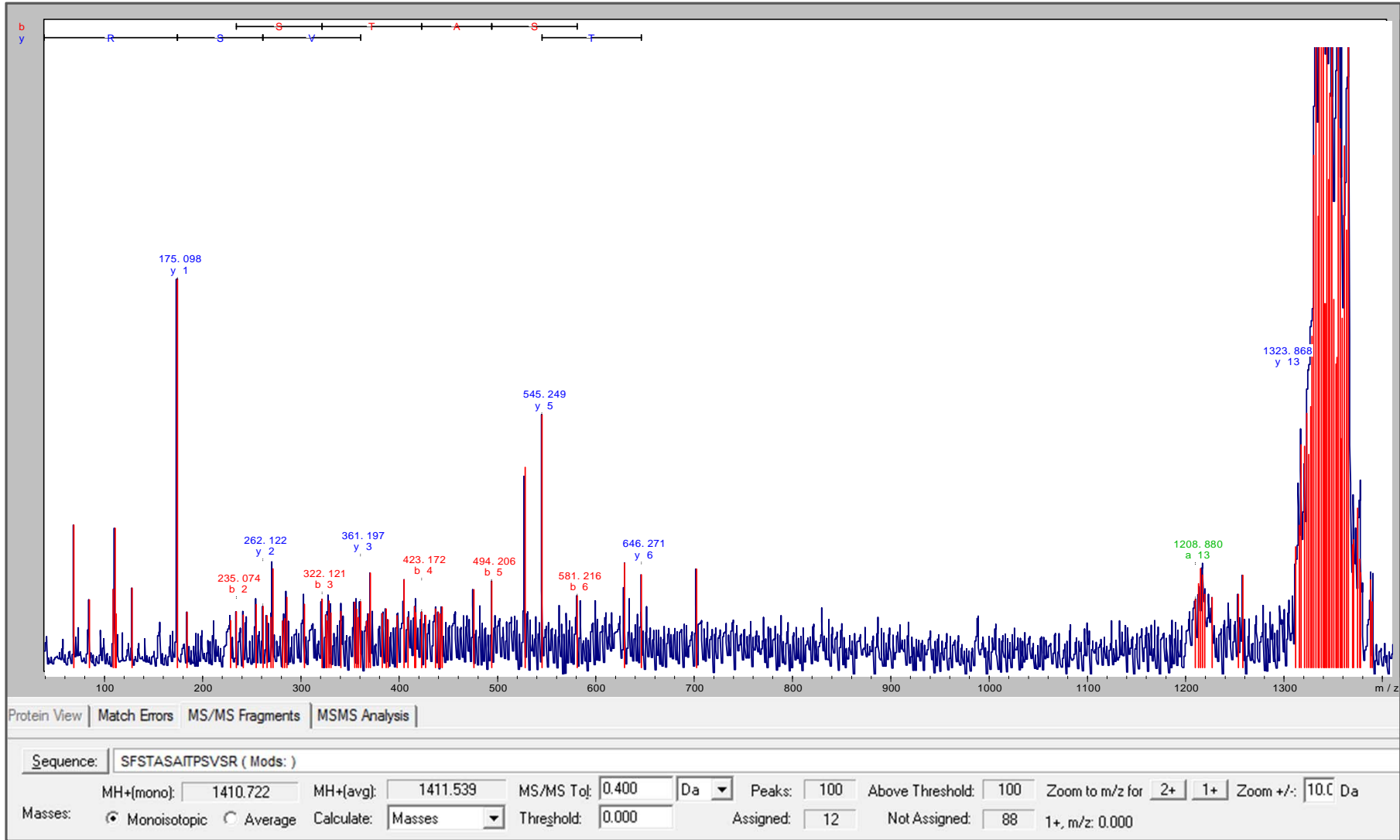
	SwissProt	Decoy	False discovery rate
Peptide matches above identity threshold	4	0	0.00%
Peptide matches above homology or identity threshold	5	0	0.00%

Protein	Calc. MH+	Meas. Mr	Calc. Mr	Dev.(Da)	Score	MascotScore	Sequence
Keratin type II cytoskeletal 7	1406.706	1405.593	1405.699	-0.106	40	32	SIHFSSPVFTSR N-Term: Acetyl (Protein N-term)
Keratin type II cytoskeletal 5	1410.722	1409.693	1409.715	-0.022	41	28	SFSTASAITPSVSR
Keratin type I cytoskeletal 15	1821.884	1820.833	1820.877	-0.044	308	128	GGSLLAGGGGFGGGSLSGGGSR
Keratin type I cytoskeletal 15	1877.888	1876.843	1876.88	-0.038	274	67	TTTFLQTSSTFGGGSTR N-Term: Acetyl (Protein N-term)
Heat shock protein beta-1	1905.992	1904.933	1904.984	-0.052	56	64	LATQSNITIPVTFSR

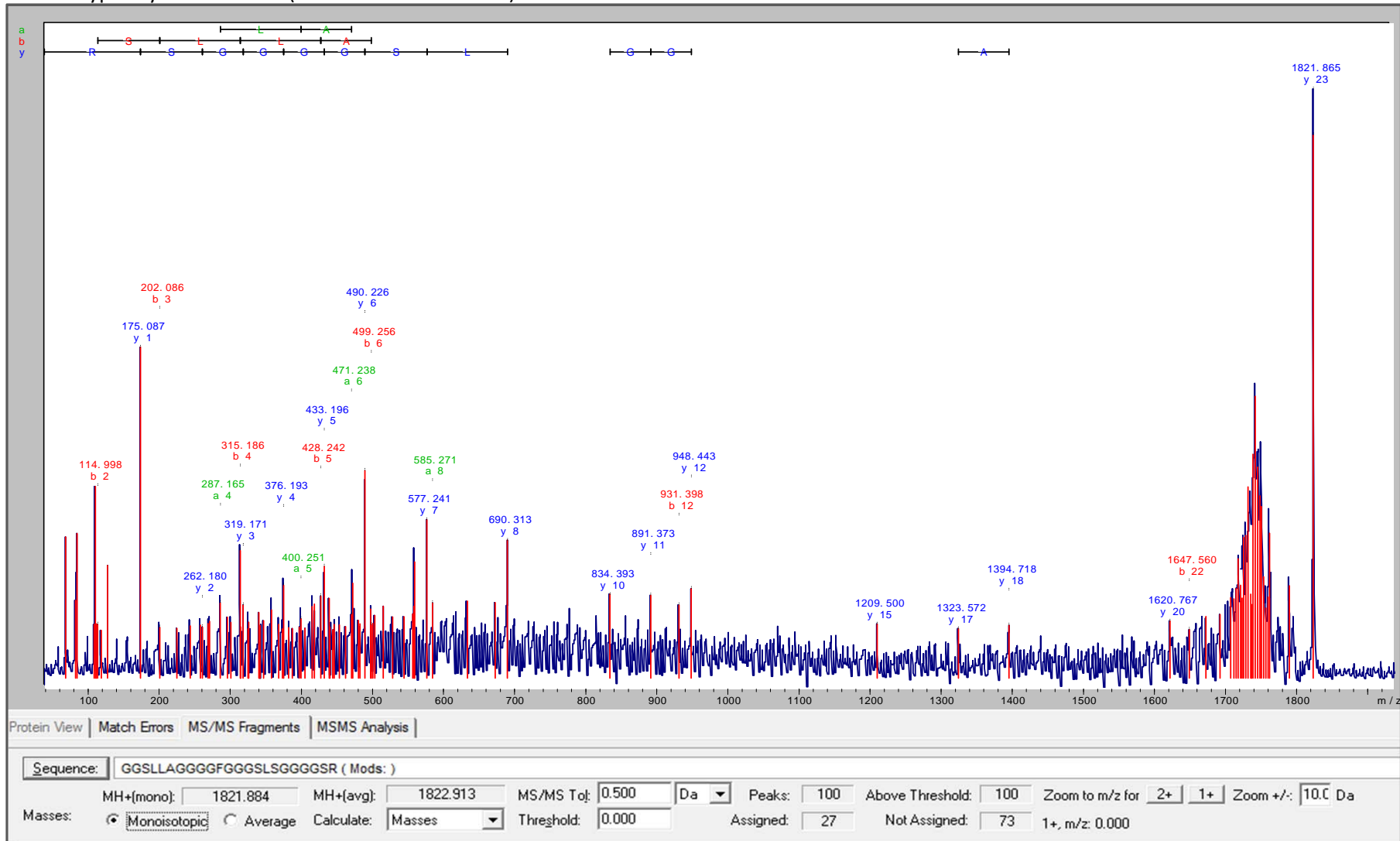
Keratin type II cytoskeletal 7 (observed mass: 1406.6)



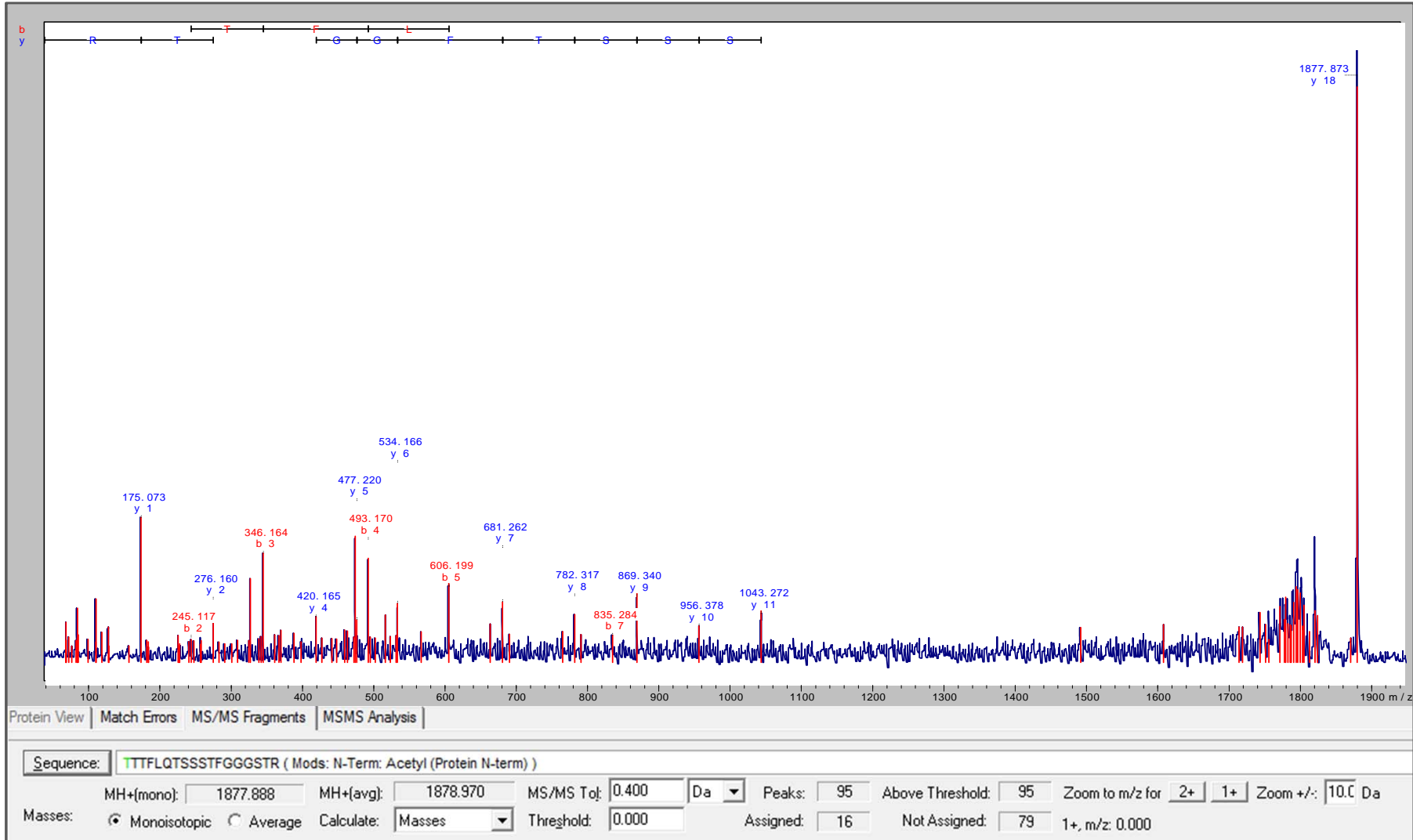
Keratin type II cytoskeletal 5 (observed mass: 1410.7)



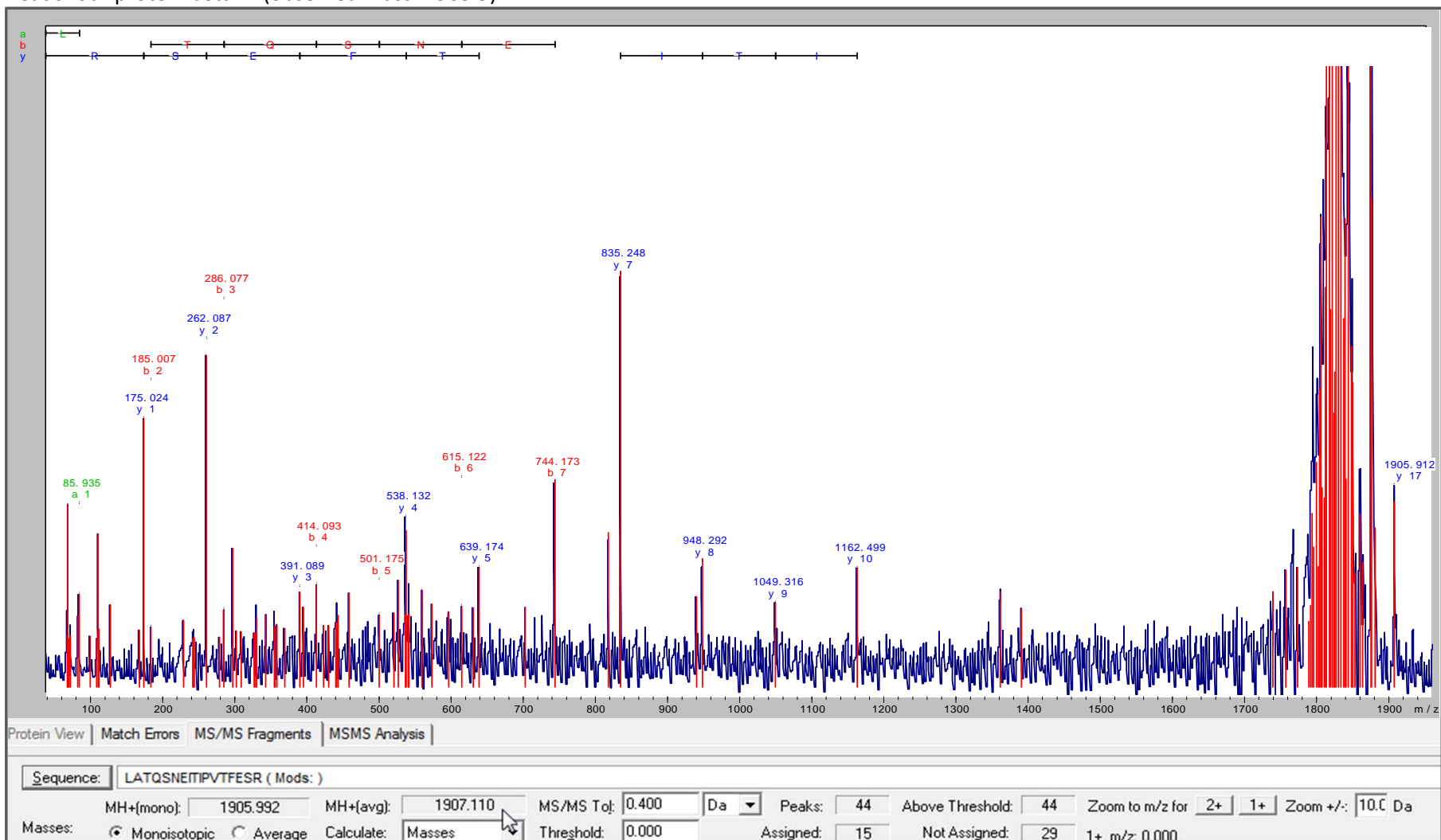
Keratin type I cytoskeletal 15 (observed mass: 1821.8)



Keratin type I cytoskeletal 15 (observed mass: 1877.8)



Heat shock protein beta-1 (observed mass: 1905.9)



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