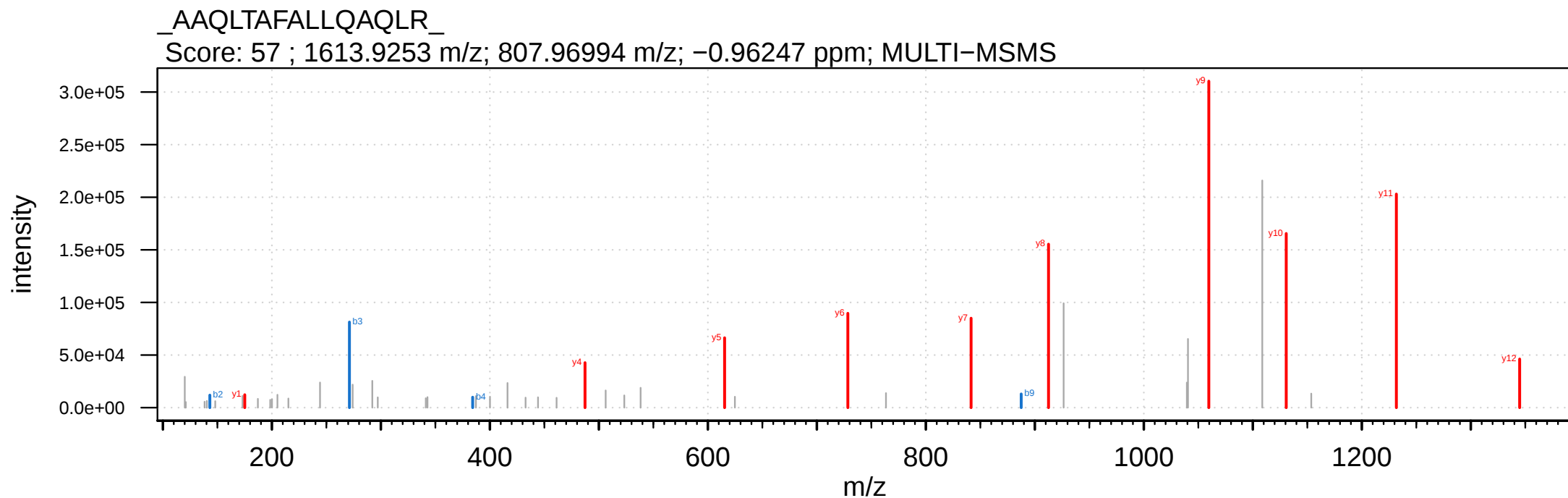
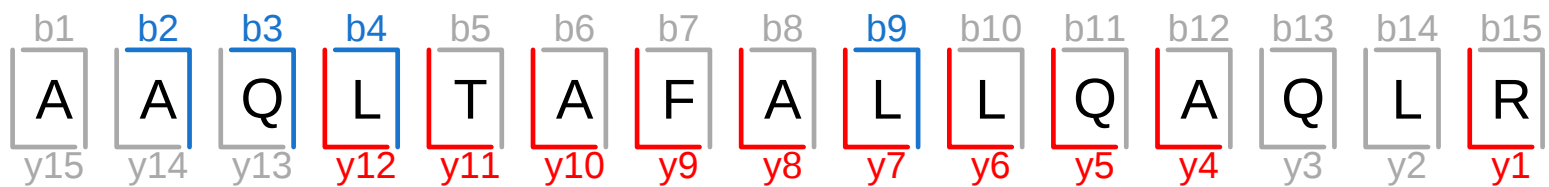
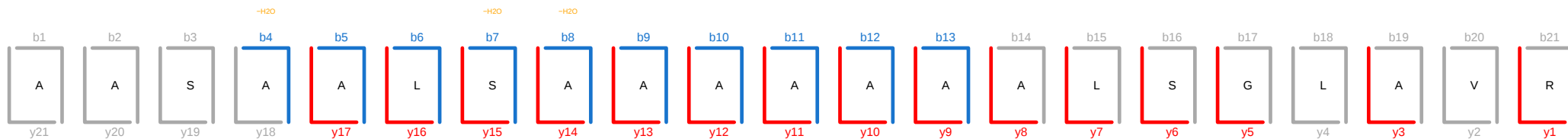


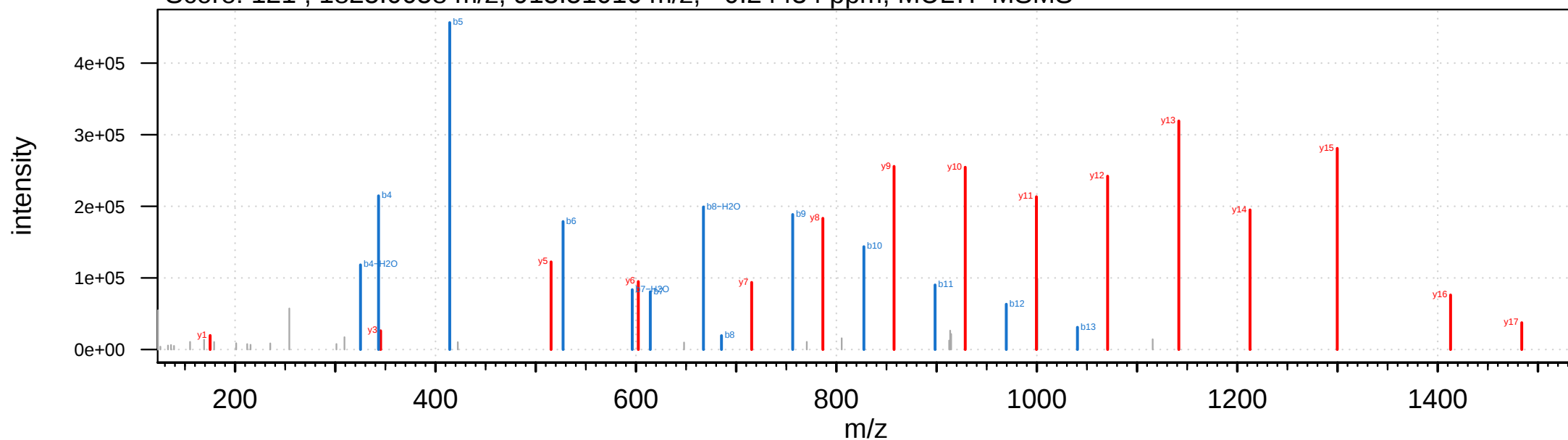


Raw File: Bibo_20130110_CHS_IEF100_20121129_3-10linear_Serva_7cm_250ug_06
 Scan Number: 45895
 Proteins:
 sp|Q69YL0|NCAS2_HUMAN
 ENST00000602845_chr3:196669588-196669887:+



(ac)AASAALSAAAAAALSGLAVR

Score: 121 ; 1825.0058 m/z; 913.51016 m/z; -0.24434 ppm; MULTI-MSMS



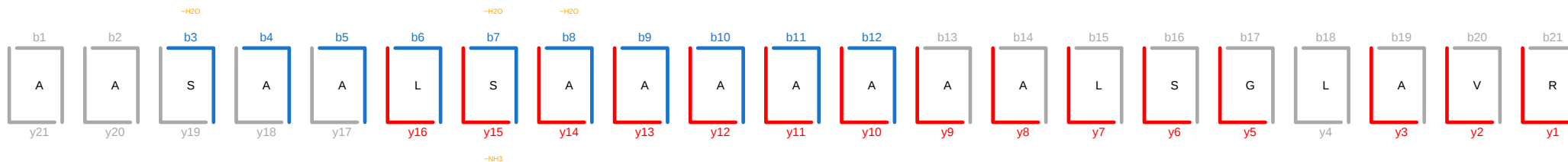
Raw File: Bibo_20130621_CHS_IEF_3-10linear_24slices_08

Scan Number: 16939

Proteins:

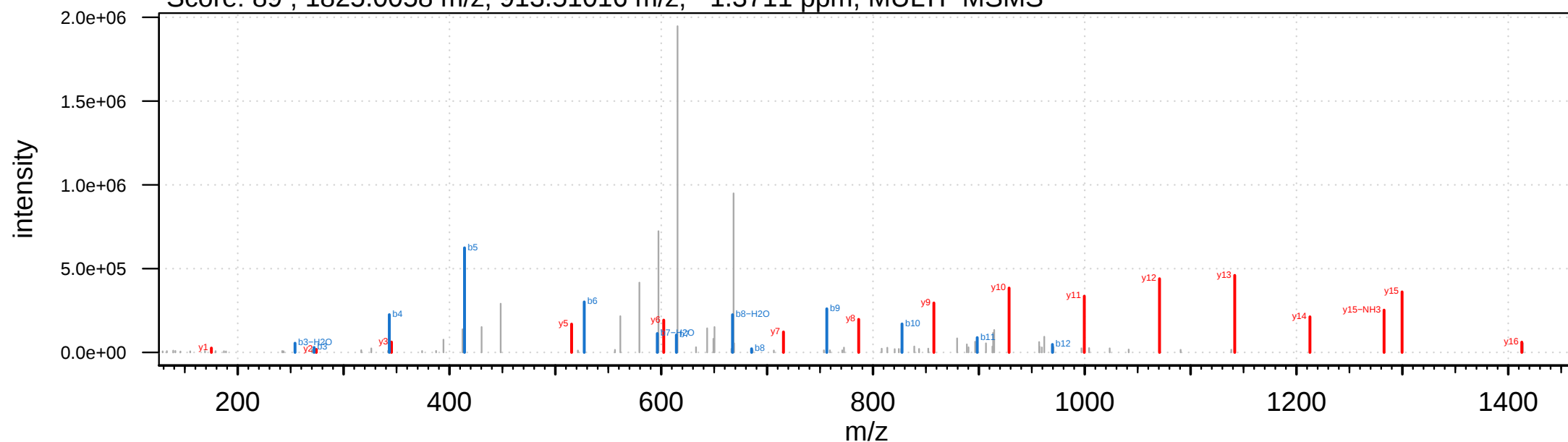
ENST00000433425_chrX:134232427-134232663:-

ENST00000417443_chrX:134556057-134556293:+



(ac)AASAALSAAAAAALSGLAVR

Score: 89 ; 1825.0058 m/z; 913.51016 m/z; -1.3711 ppm; MULTI-MSMS



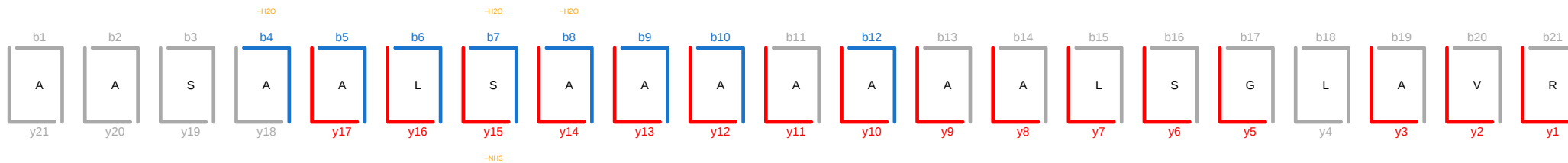
Raw File: Bibo_20130621_CHS_IEF_3-10linearpep_24slices_11

Scan Number: 17553

Proteins:

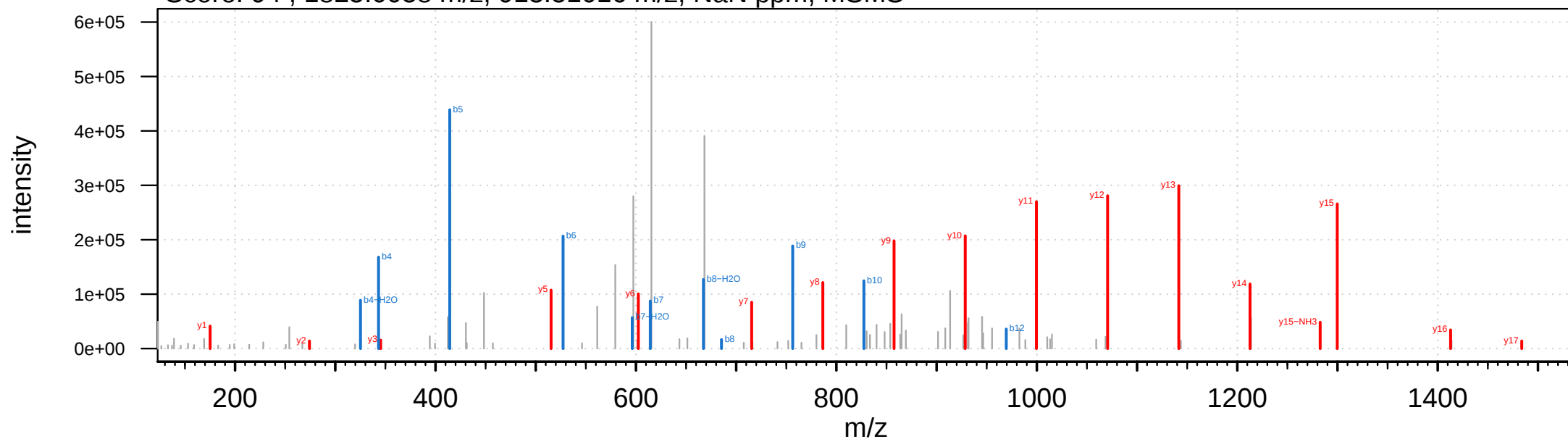
ENST00000433425_chrX:134232427-134232663:-

ENST00000417443_chrX:134556057-134556293:+



(ac)AASAALSAAAAAALSGLAVR

Score: 94 ; 1825.0058 m/z; 913.51016 m/z; NaN ppm; MSMS



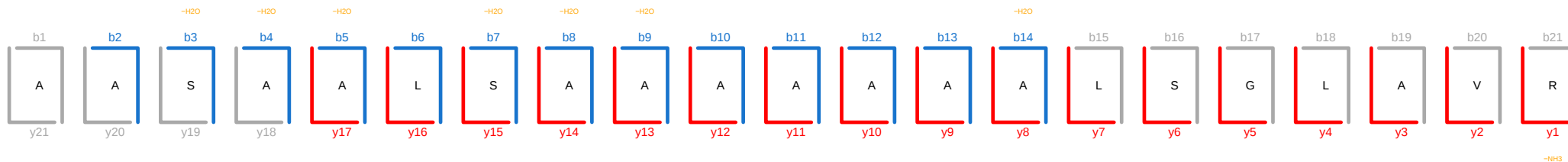
Raw File: Bibo_20130621_CHS_IEF_3-10linearpep_24slices_12

Scan Number: 18159

Proteins:

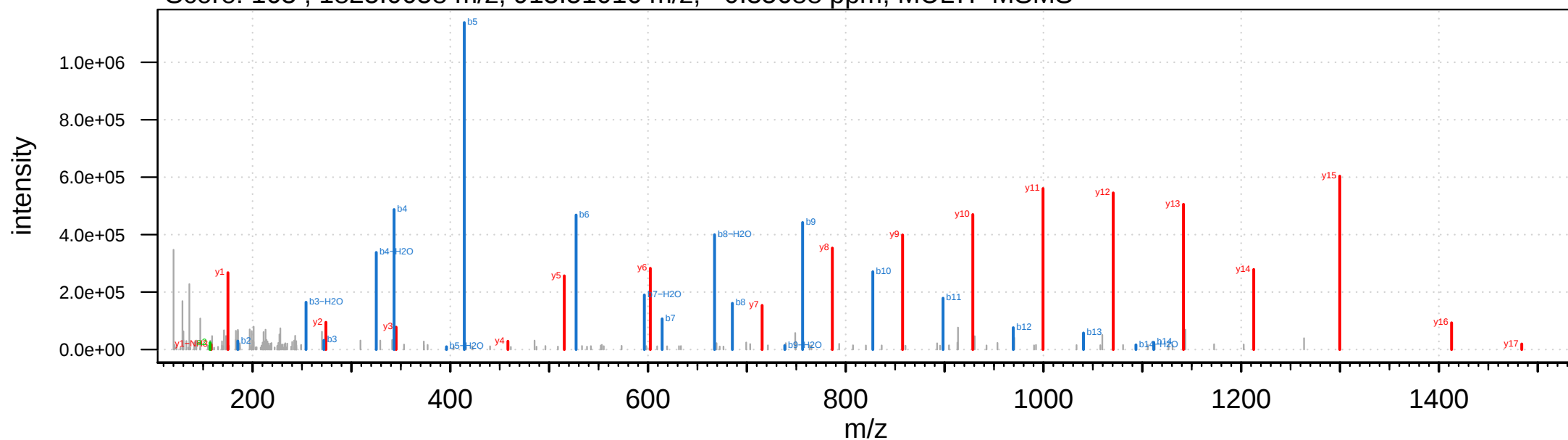
ENST00000433425_chrX:134232427-134232663:-

ENST00000417443_chrX:134556057-134556293:+



(ac)AASAALSAAAAAALSGLAVR

Score: 163 ; 1825.0058 m/z; 913.51016 m/z; -0.35688 ppm; MULTI-MSMS



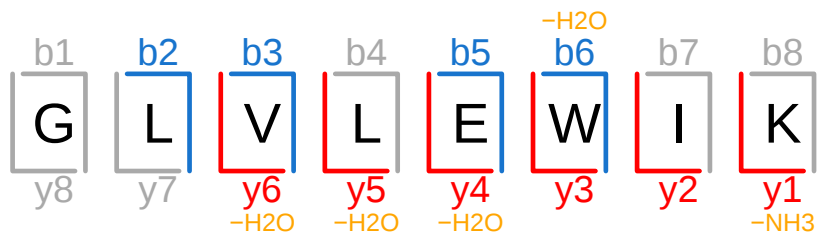
Raw File: Grobi_20121019_CHS_PIECEIEF_20121017_3-10_7cm_250ug_05

Scan Number: 102781

Proteins:

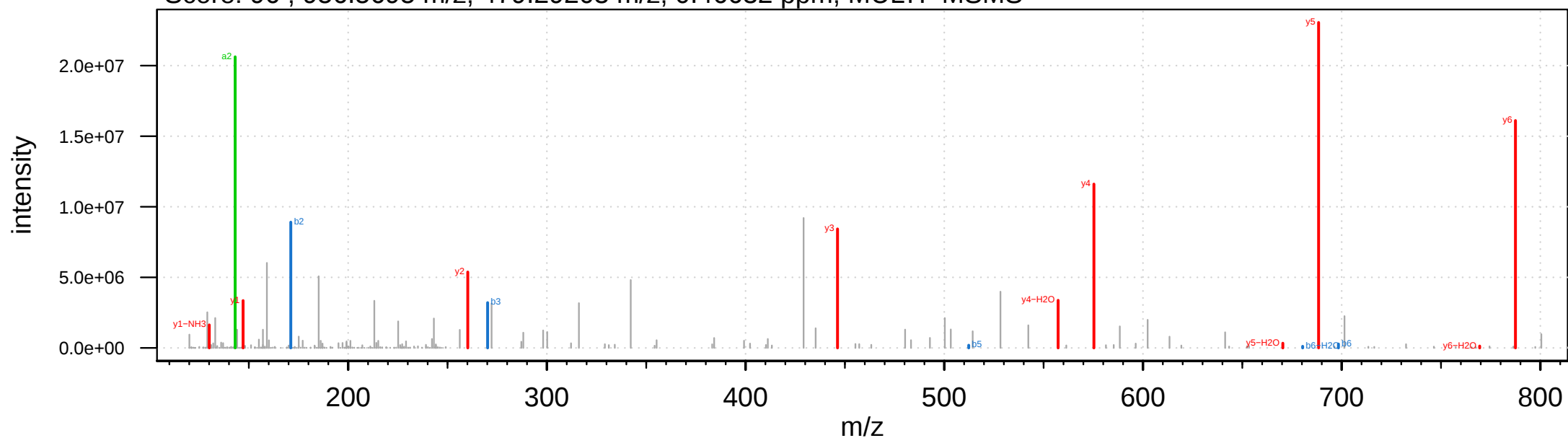
ENST00000433425_chrX:134232427-134232663:-

ENST00000417443_chrX:134556057-134556293:+



GLVLEWIK

Score: 96 ; 956.5695 m/z; 479.29203 m/z; 0.46632 ppm; MULTI-MSMS

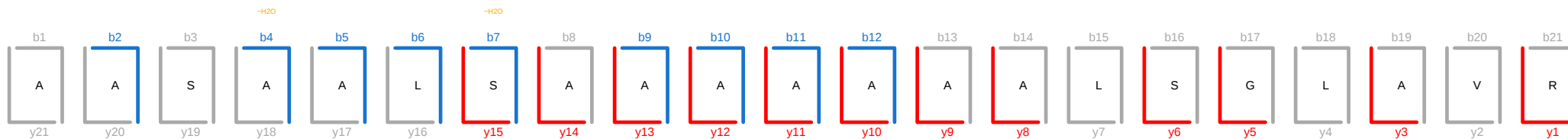


Raw File: Grobi_20121019_CHS_PIECEIEF_20121017_3-10_7cm_250ug_05

Scan Number: 63753

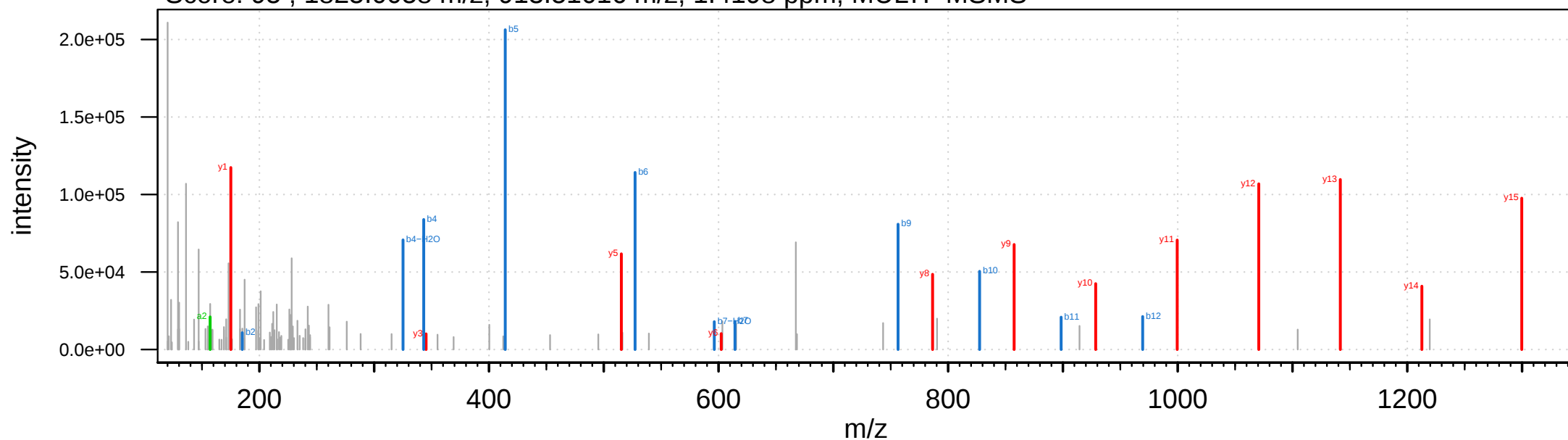
Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-



(ac)AASAALSAAAAAALSGLAVR

Score: 95 ; 1825.0058 m/z; 913.51016 m/z; 1.4198 ppm; MULTI-MSMS



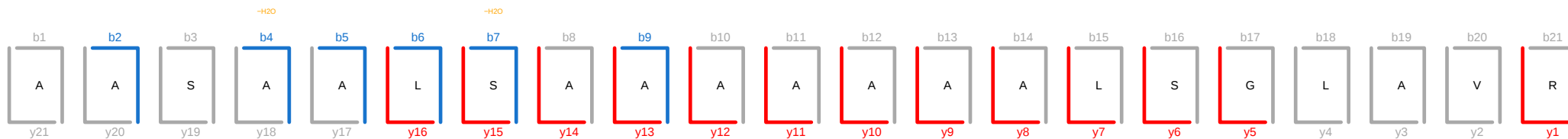
Raw File: Grobi_20130312_CHS_IEF100_20130309_3-10linear_Serva_7cm_1mg_1elution_02

Scan Number: 93471

Proteins:

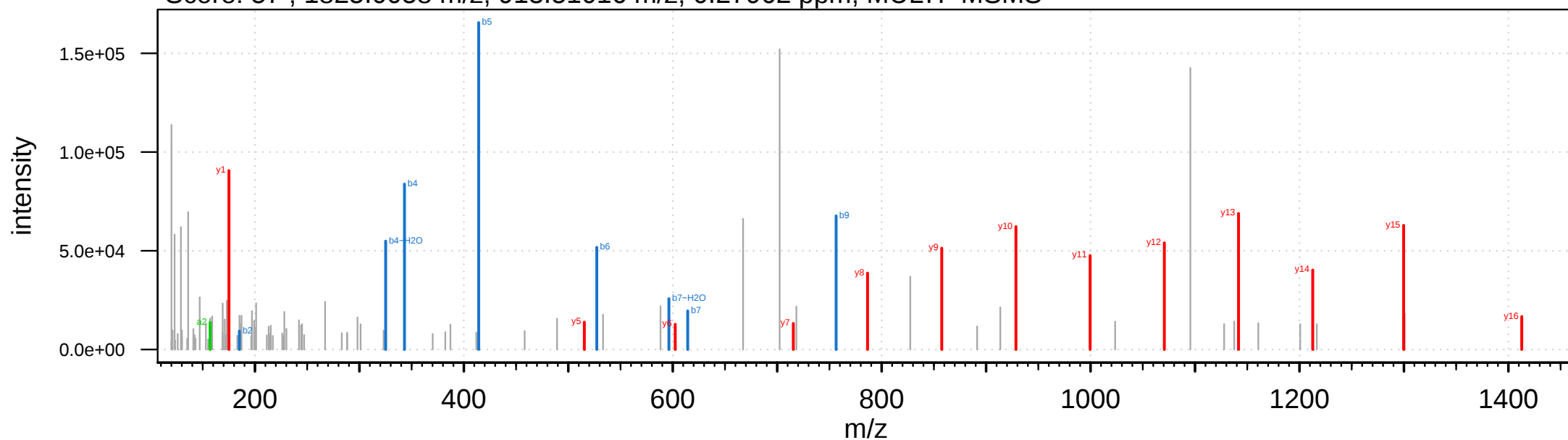
ENST00000433425_chrX:134232427-134232663:-

ENST00000417443_chrX:134556057-134556293:+



(ac)AASAALSAAAAAALSGLAVR

Score: 57 ; 1825.0058 m/z; 913.51016 m/z; 0.27962 ppm; MULTI-MSMS



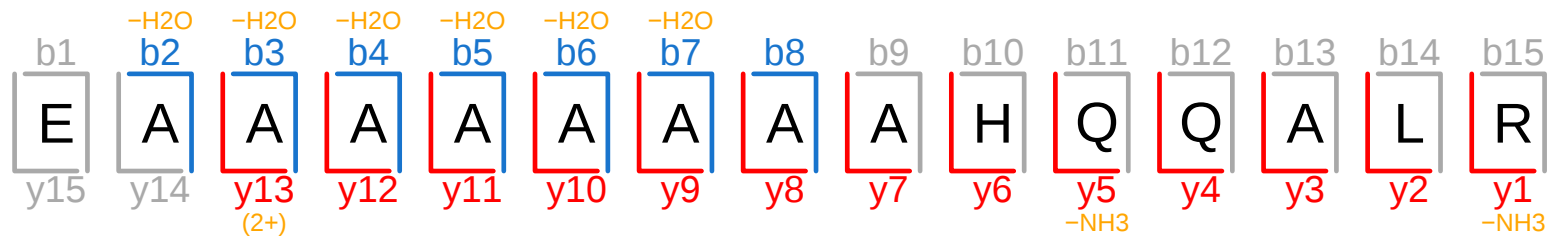
Raw File: Grobi_20130312_CHS_IEF100_20130309_3-10linear_Serva_7cm_1mg_1elution_04

Scan Number: 95039

Proteins:

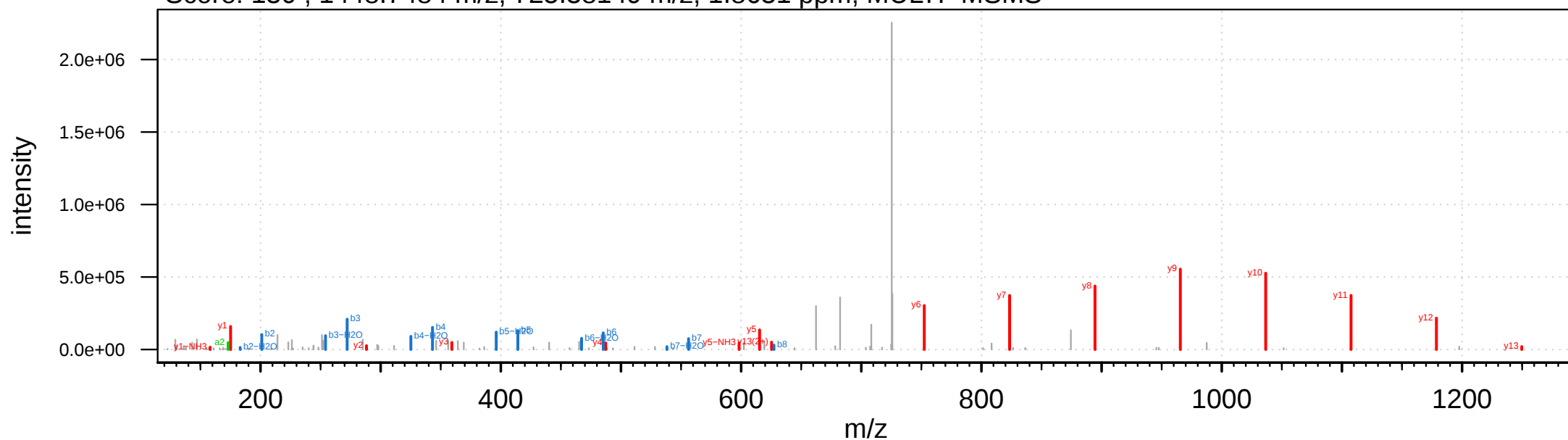
ENST00000433425_chrX:134232427-134232663:-

ENST00000417443_chrX:134556057-134556293:+



EAAAAAAAAAHQQALR

Score: 130 ; 1448.7484 m/z; 725.38149 m/z; 1.8631 ppm; MULTI-MSMS

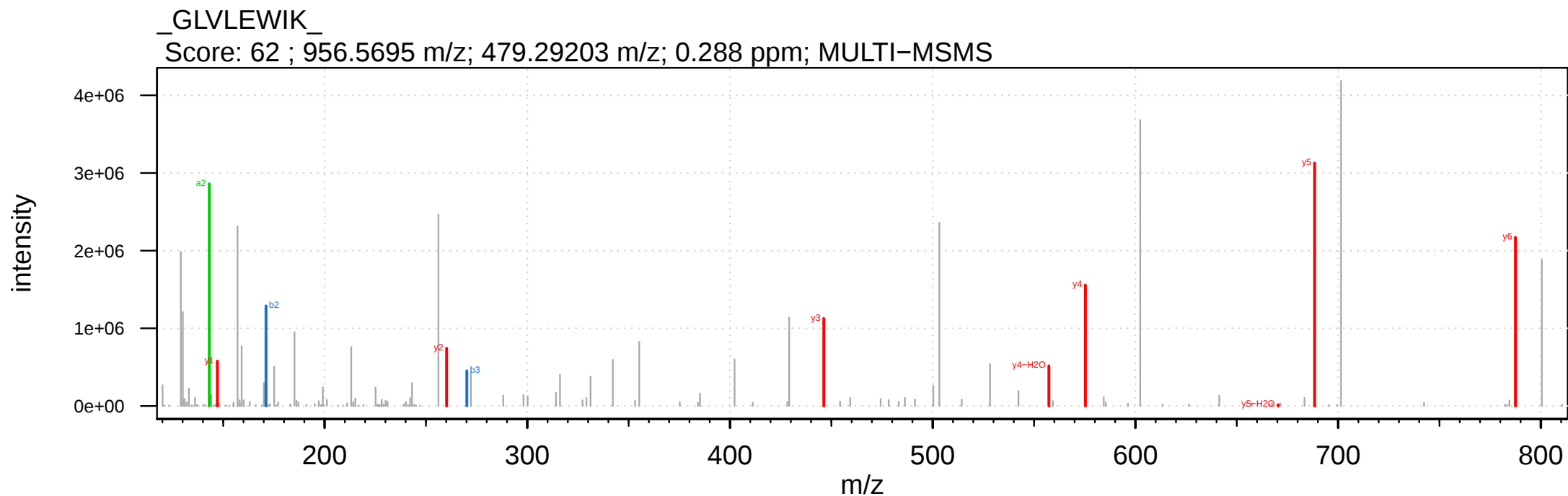
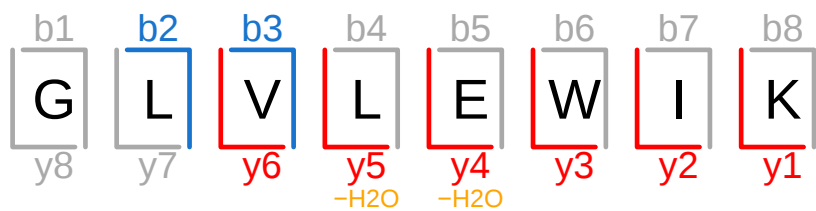


Raw File: Grobi_20130312_CHS_IEF100_20130309_3-10linear_Serva_7cm_1mg_1elution_04

Scan Number: 3952

Proteins:

ENST00000394071_chr14:74185546-74185677:-

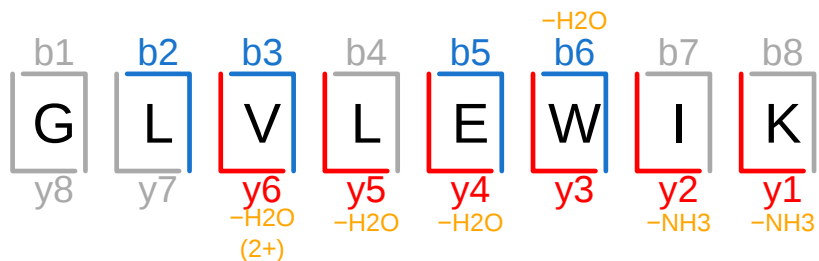


Raw File: Grobi_20130312_CHS_IEF100_20130309_3-10linear_Serva_7cm_1mg_1elution_04

Scan Number: 38409

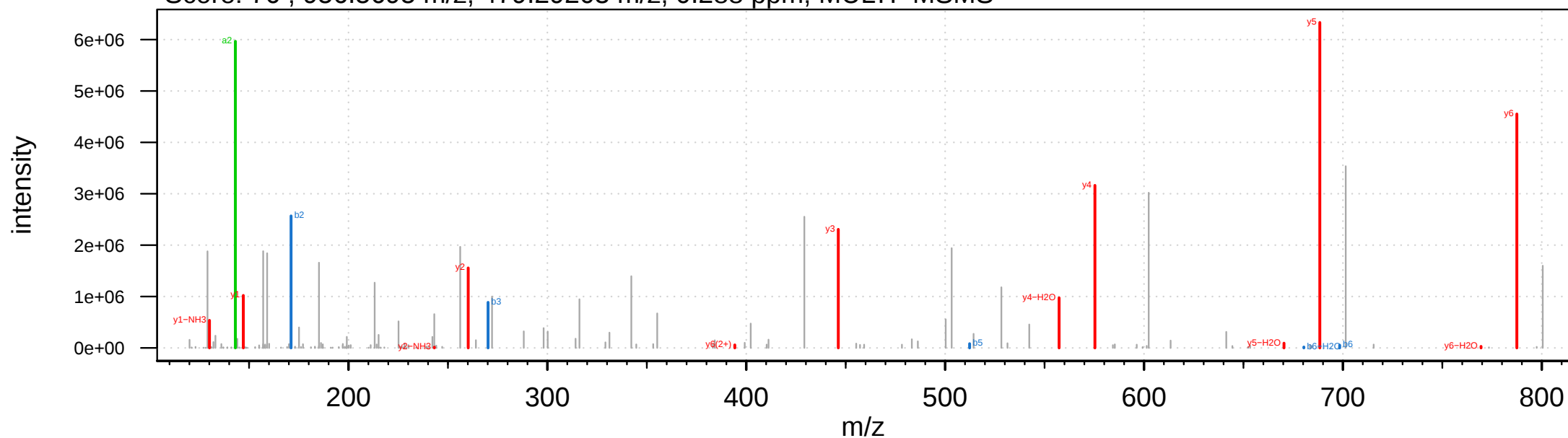
Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-



GLVLEWIK

Score: 76 ; 956.5695 m/z; 479.29203 m/z; 0.288 ppm; MULTI-MSMS

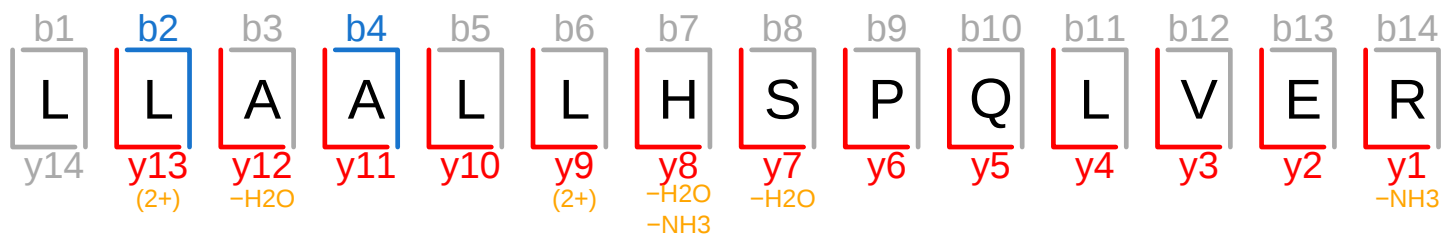


Raw File: Grobi_20130312_CHS_IEF100_20130309_3-10linear_Serva_7cm_1mg_1elution_04

Scan Number: 38654

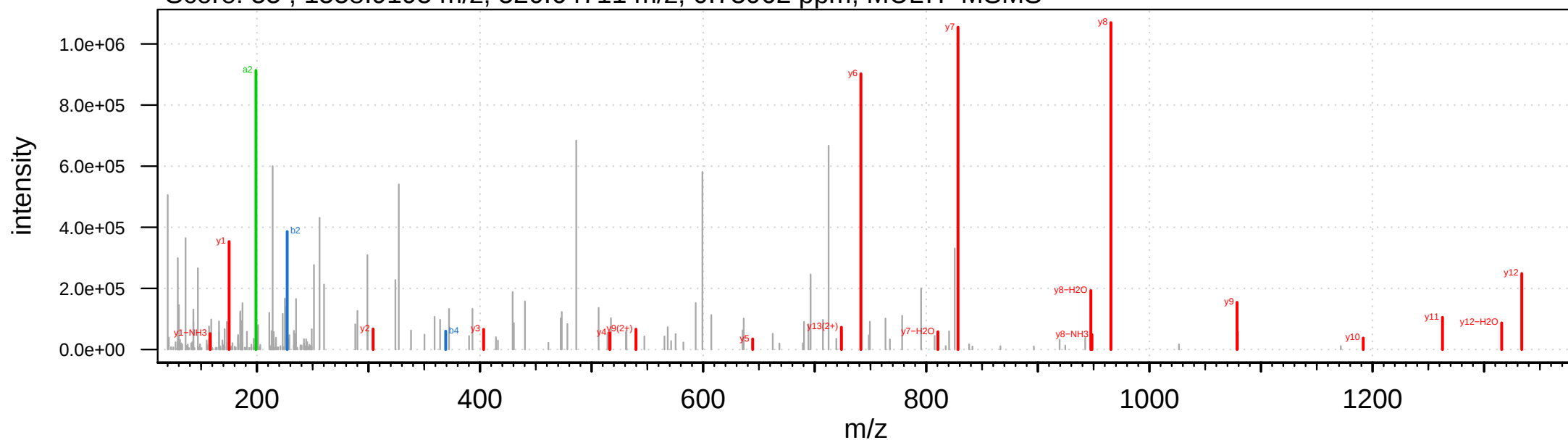
Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-



LLAALLHSPQLVER

Score: 55 ; 1558.9195 m/z; 520.64711 m/z; 0.73962 ppm; MULTI-MSMS



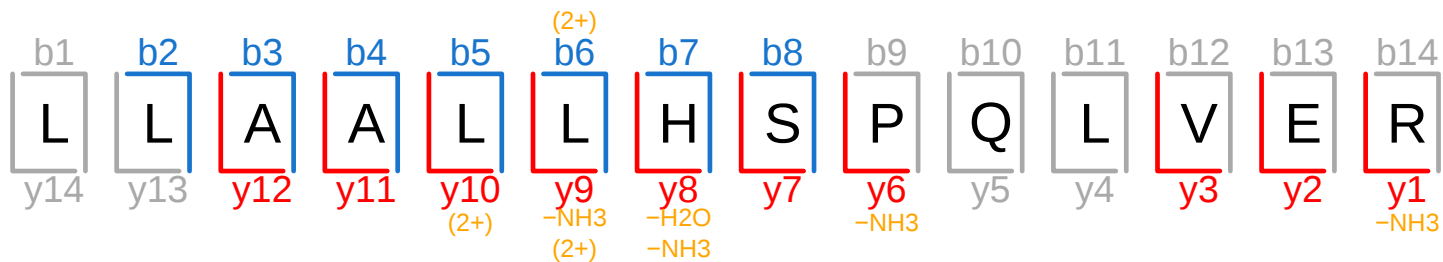
Raw File: Grobi_20130312_CHS_IEF100_20130309_3-10linear_Serva_7cm_1mg_1elution_04

Scan Number: 33266

Proteins:

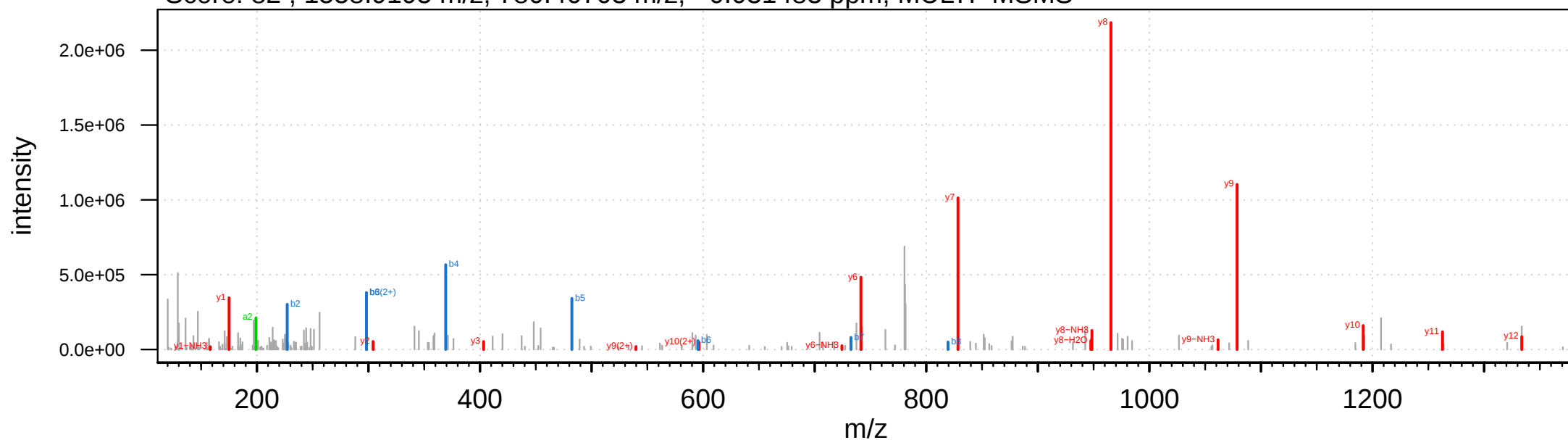
sp|Q69YL0|NCAS2_HUMAN

ENST00000602845_chr3:196669588-196669887:+



LLAALLHSPQLVER

Score: 82 ; 1558.9195 m/z; 780.46703 m/z; -0.031483 ppm; MULTI-MSMS



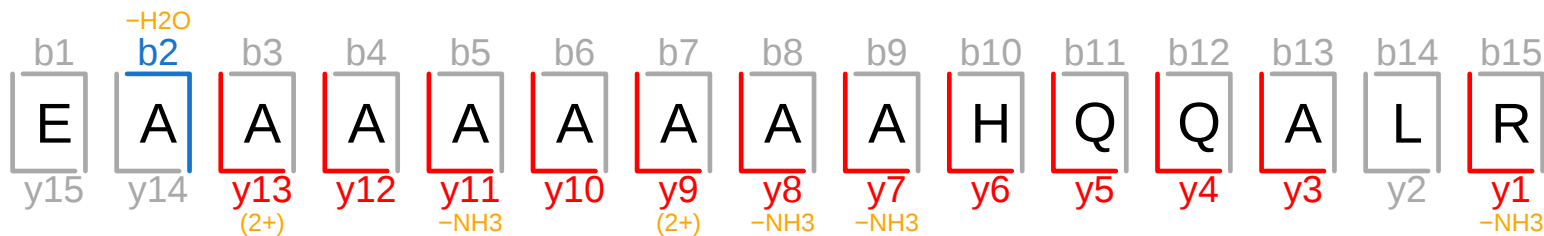
Raw File: Grobi_20130312_CHS_IEF100_20130309_3-10linear_Serva_7cm_1mg_1elution_04

Scan Number: 33305

Proteins:

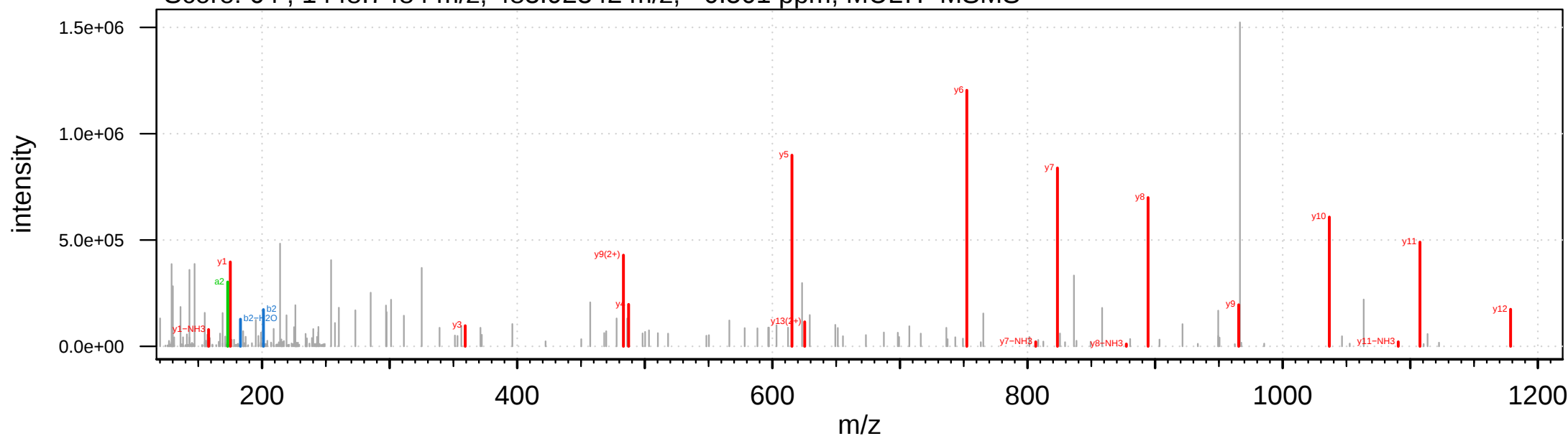
sp|Q69YL0|NCAS2_HUMAN

ENST00000602845_chr3:196669588-196669887:+



EAAAAAAAAAHQQALR

Score: 64 ; 1448.7484 m/z; 483.92342 m/z; -0.501 ppm; MULTI-MSMS

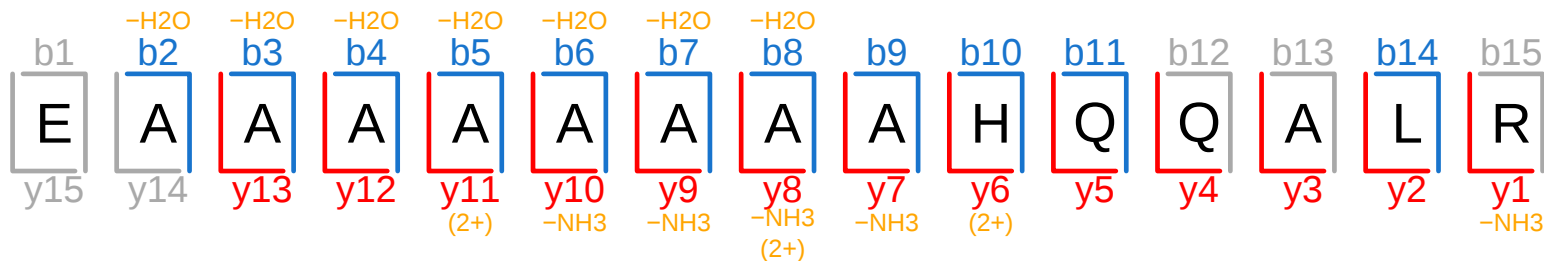


Raw File: Grobi_20121019_CHS_PIECEIEF_20121017_3-10_7cm_250ug_06

Scan Number: 17358

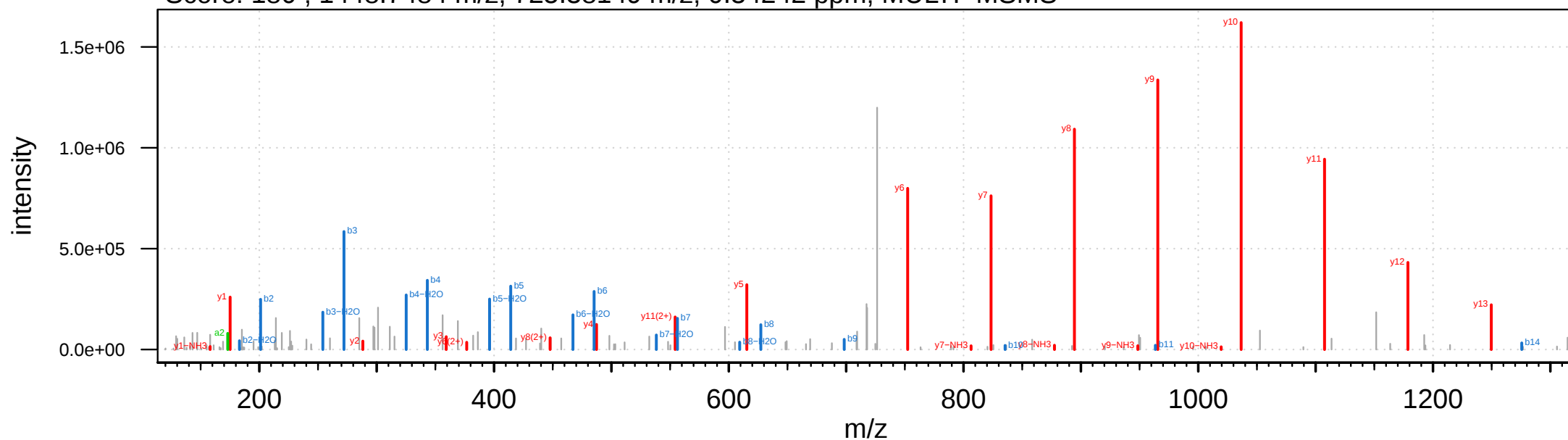
Proteins:

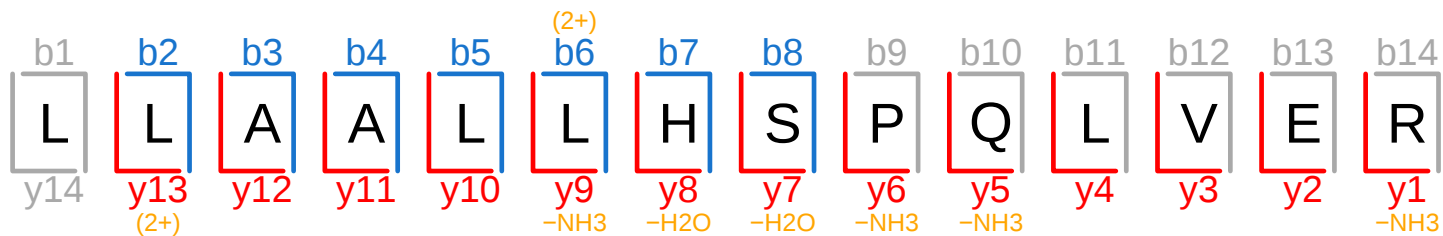
ENST00000394071_chr14:74185546-74185677:-



EAAAAAAAAAHQQALR

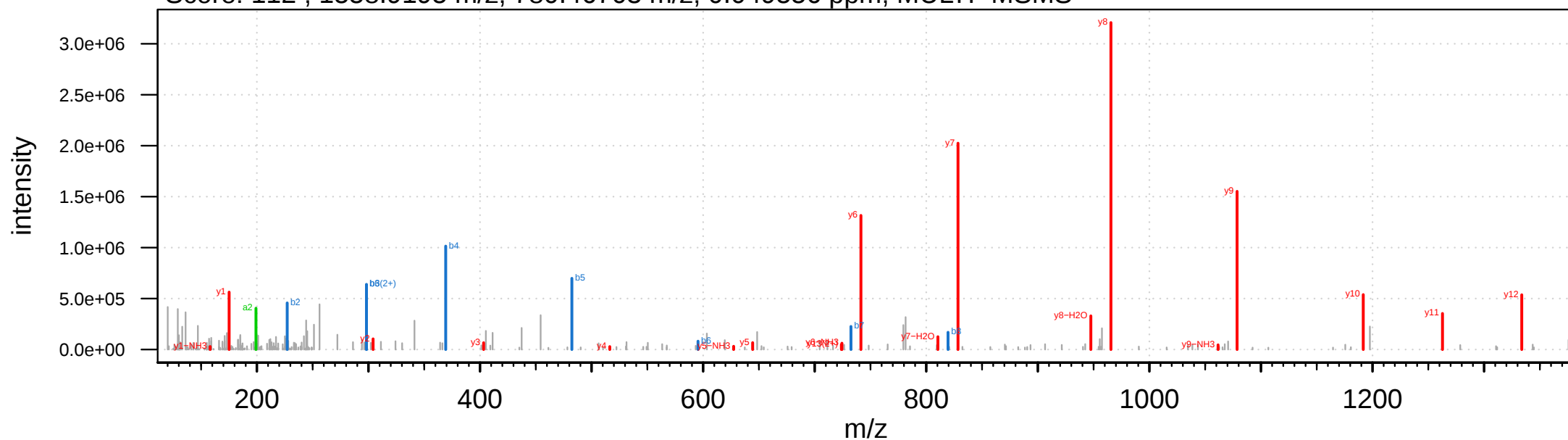
Score: 180 ; 1448.7484 m/z; 725.38149 m/z; 0.34242 ppm; MULTI-MSMS





LLAALLHSPQLVER

Score: 112 ; 1558.9195 m/z; 780.46703 m/z; 0.049556 ppm; MULTI-MSMS



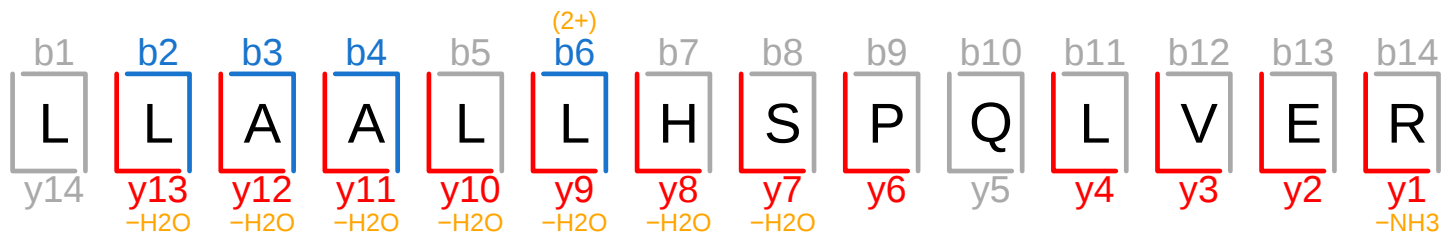
Raw File: Grobi_20121019_CHS_PIECEIEF_20121017_3-10_7cm_250ug_06

Scan Number: 57689

Proteins:

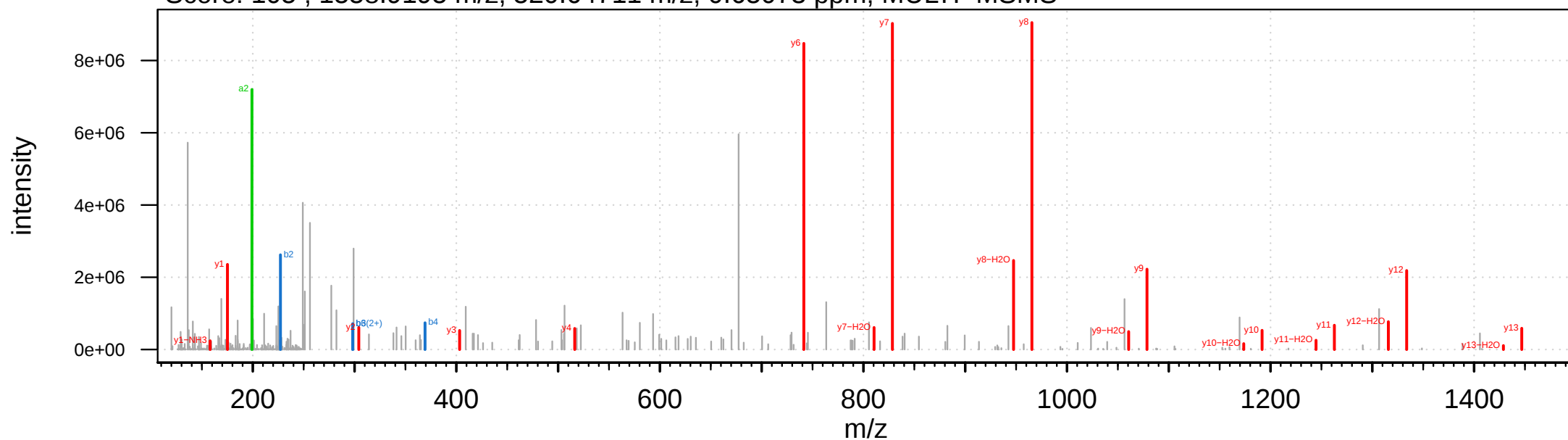
sp|Q69YL0|NCAS2_HUMAN

ENST00000602845_chr3:196669588-196669887:+



LLAALLHSPQLVER

Score: 105 ; 1558.9195 m/z; 520.64711 m/z; 0.65073 ppm; MULTI-MSMS



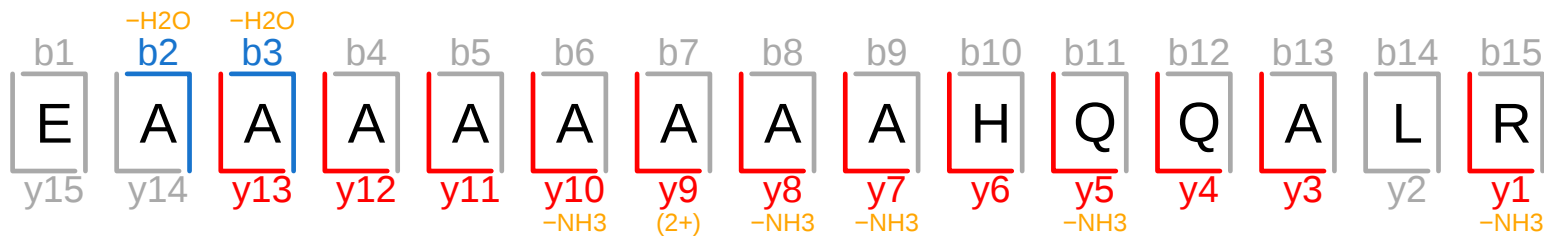
Raw File: Grobi_20121019_CHS_PIECEIEF_20121017_3-10_7cm_250ug_06

Scan Number: 57748

Proteins:

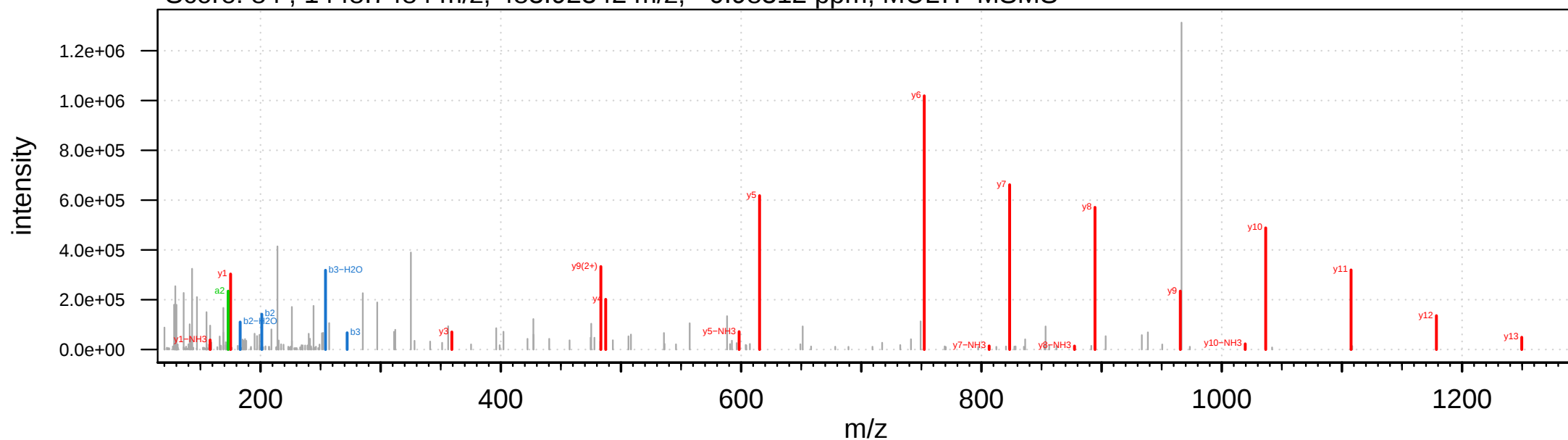
sp|Q69YL0|NCAS2_HUMAN

ENST00000602845_chr3:196669588-196669887:+



EAAAAAAAAAHQQALR

Score: 84 ; 1448.7484 m/z; 483.92342 m/z; -0.98312 ppm; MULTI-MSMS

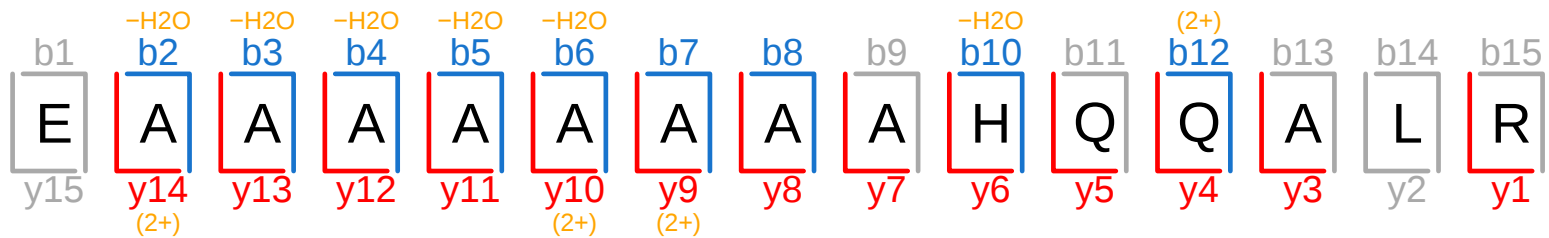


Raw File: Grobi_20121105_CHS_IEF_201210125_pep_gradient_7cm_250ug_06

Scan Number: 22066

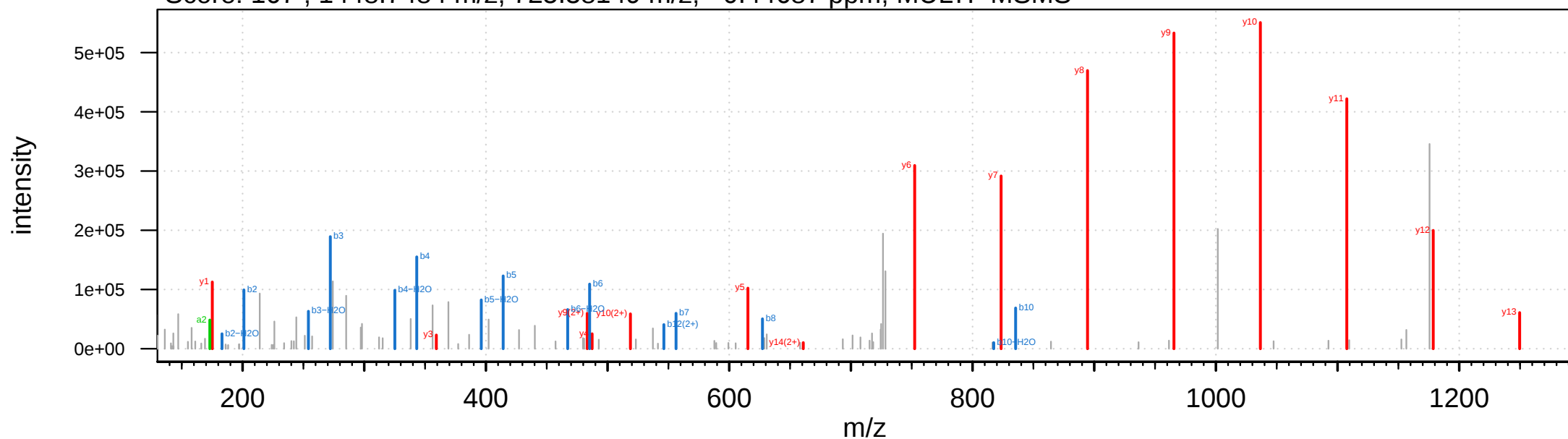
Proteins:

ENST00000394071_chr14:74185546-74185677:-



EAAAAAAAAAHQQALR

Score: 167 ; 1448.7484 m/z; 725.38149 m/z; -0.44687 ppm; MULTI-MSMS

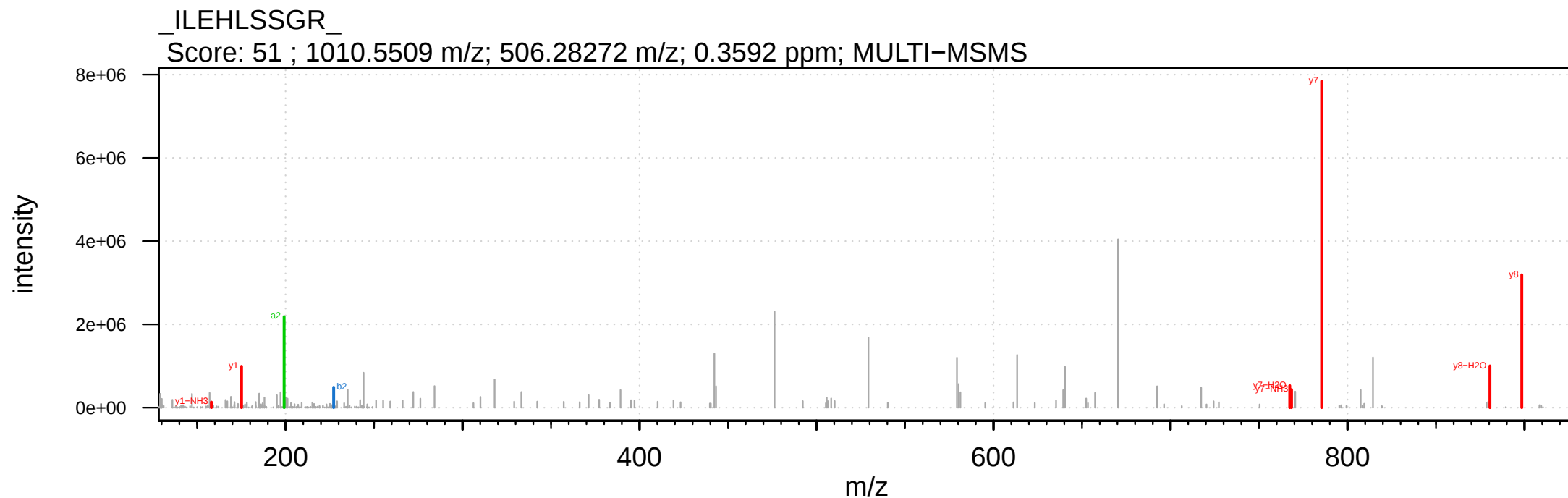
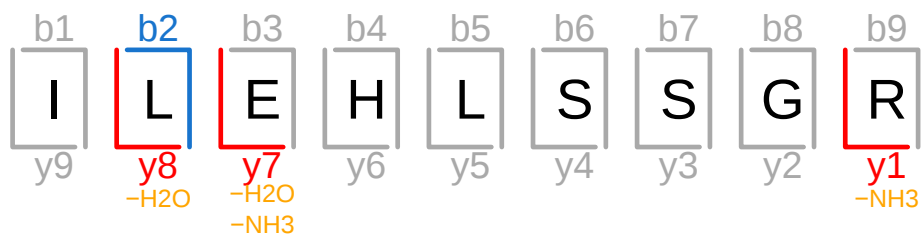


Raw File: Grobi_20121105_CHS_IEF_201210125_pep_gradient_7cm_250ug_06

Scan Number: 22080

Proteins:

ENST00000394071_chr14:74185546-74185677:-

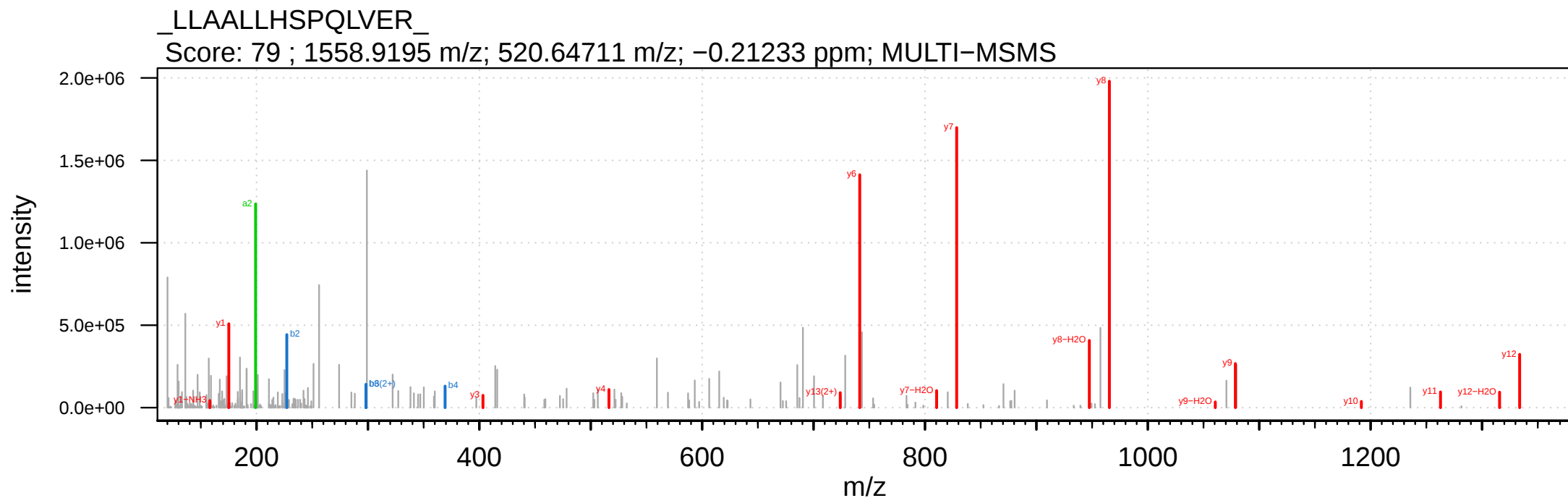
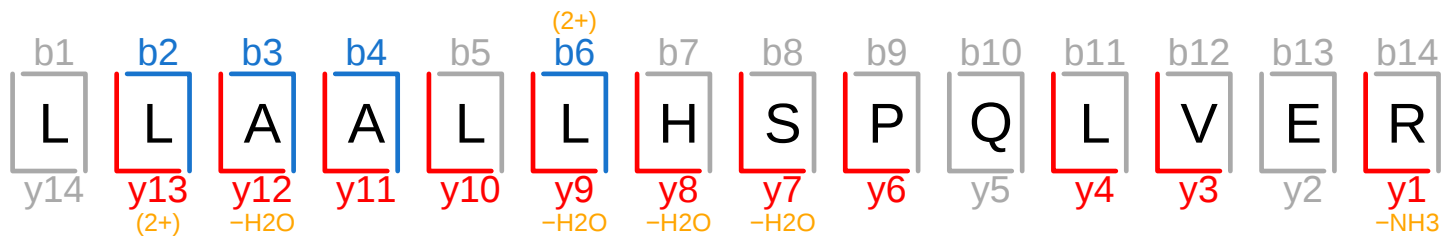


Raw File: Grobi_20121105_CHS_IEF_201210125_pep_gradient_7cm_250ug_06

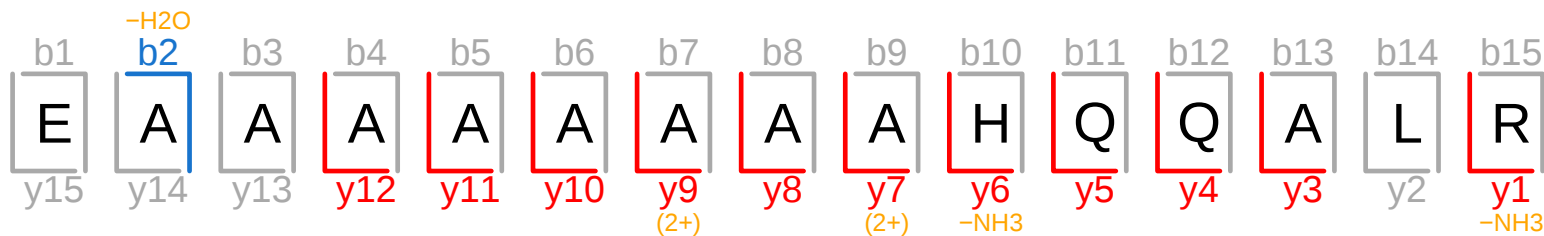
Scan Number: 26406

Proteins:

ENST00000452079_chr1:3663088-3663306:-

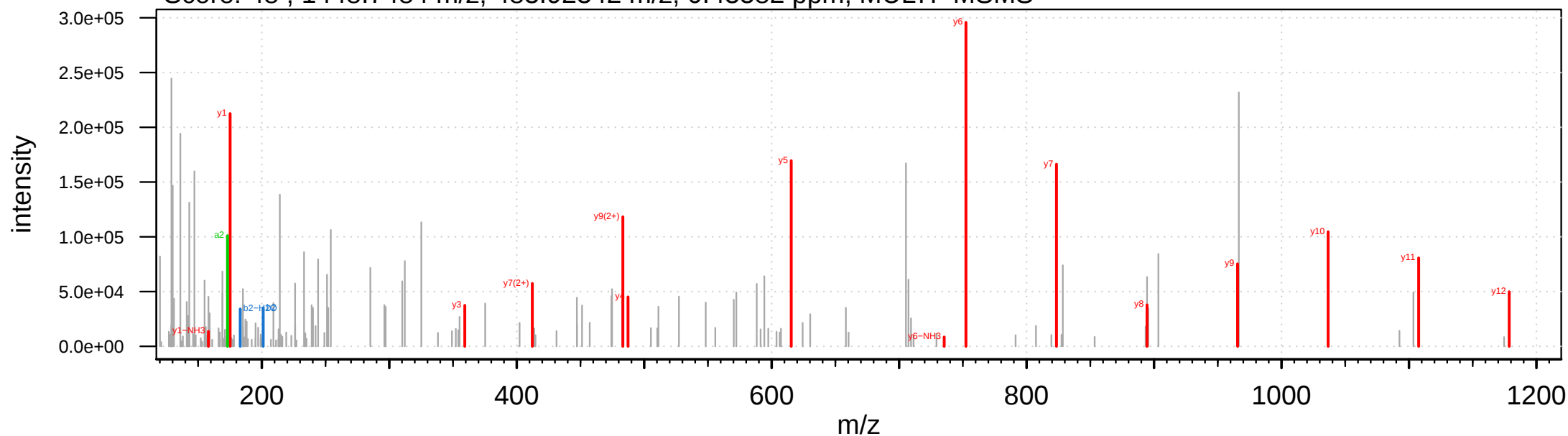


Raw File: Grobi_20121105_CHS_IEF_201210125_pep_gradient_7cm_250ug_06
Scan Number: 67878
Proteins:
sp|Q69YL0|NCAS2_HUMAN
ENST00000602845_chr3:196669588-196669887:+



EAAAAAAAAAHQQALR

Score: 48 ; 1448.7484 m/z; 483.92342 m/z; 0.43582 ppm; MULTI-MSMS

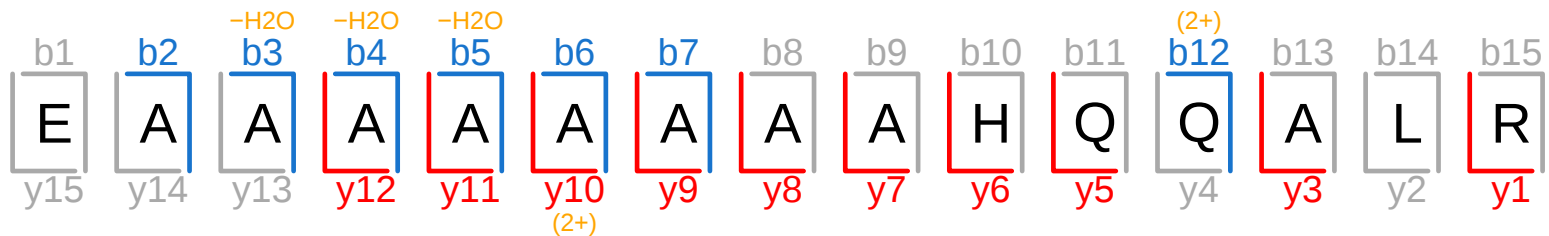


Raw File: Grobi_20121105_CHS_IEF100_20121025_pep_gradient_7cm_250ug_06

Scan Number: 21177

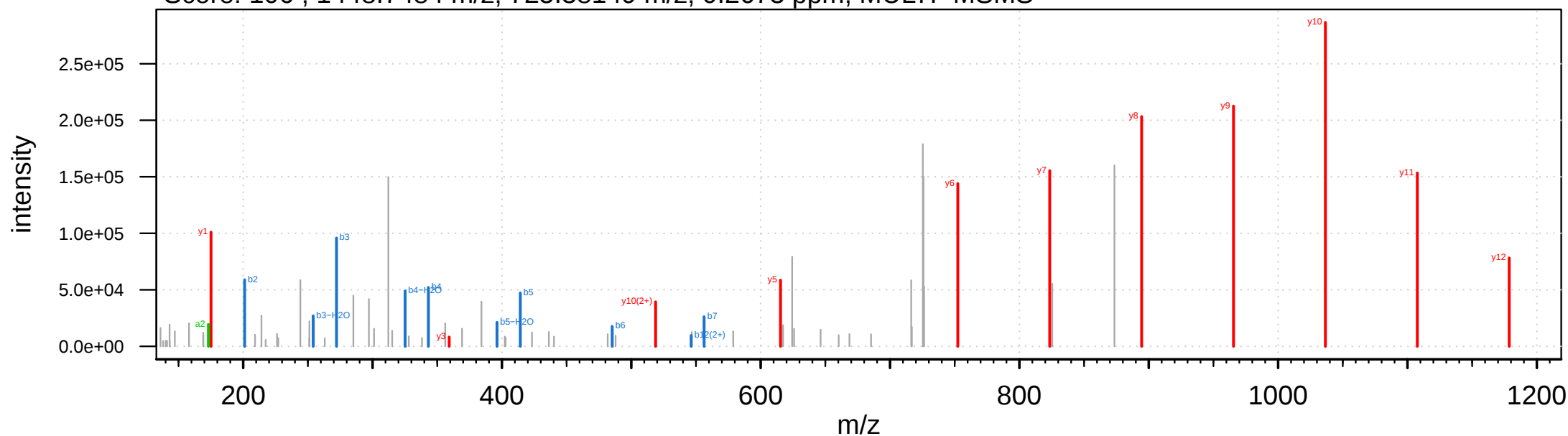
Proteins:

ENST00000394071_chr14:74185546-74185677:-



EAAAAAAAAAHQQALR

Score: 106 ; 1448.7484 m/z; 725.38149 m/z; 0.2673 ppm; MULTI-MSMS

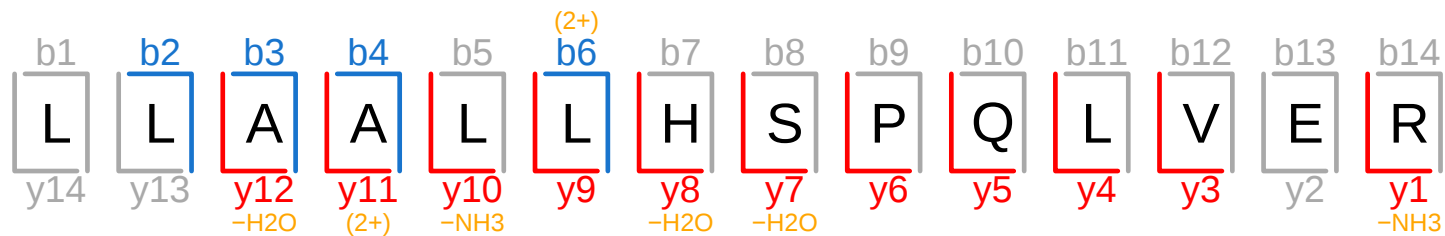


Raw File: Grobi_20121105_CHS_IEF100_20121025_pep_gradient_7cm_250ug_06

Scan Number: 21249

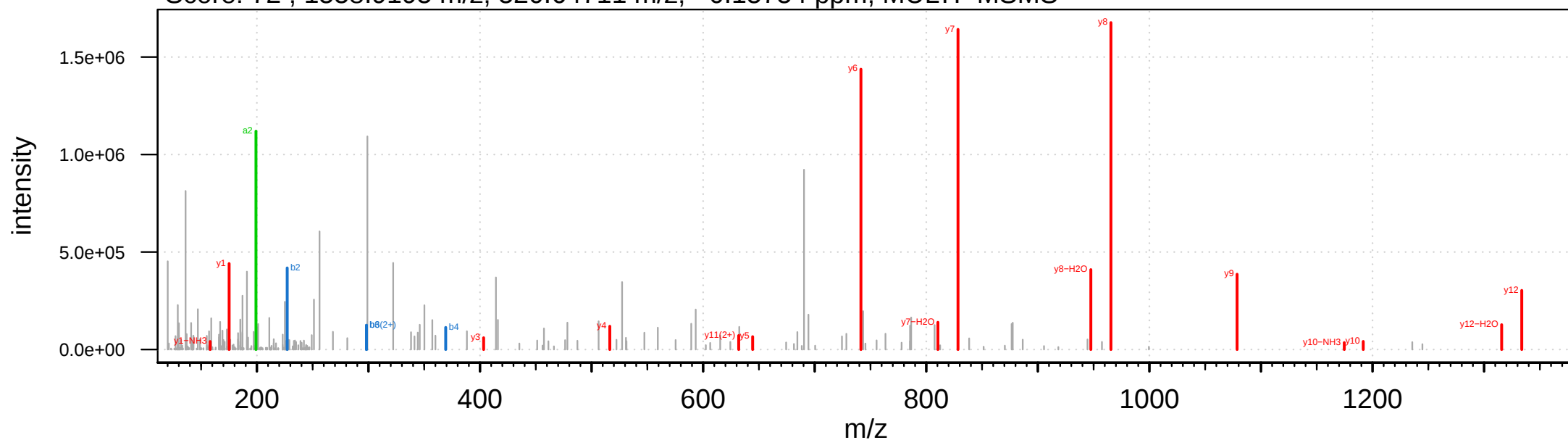
Proteins:

ENST00000394071_chr14:74185546-74185677:-



LLAALLHSPQLVER

Score: 72 ; 1558.9195 m/z; 520.64711 m/z; -0.15754 ppm; MULTI-MSMS



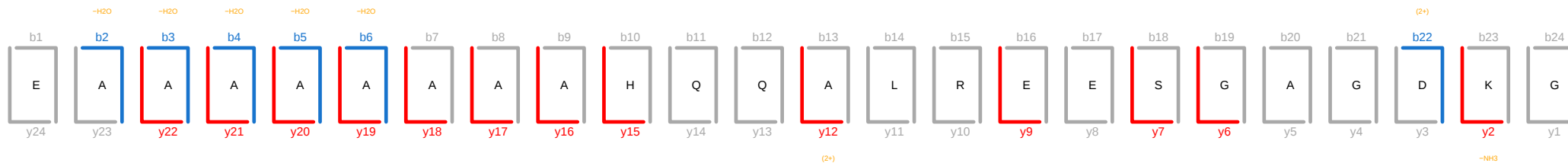
Raw File: Grobi_20121105_CHS_IEF100_20121025_pep_gradient_7cm_250ug_06

Scan Number: 67161

Proteins:

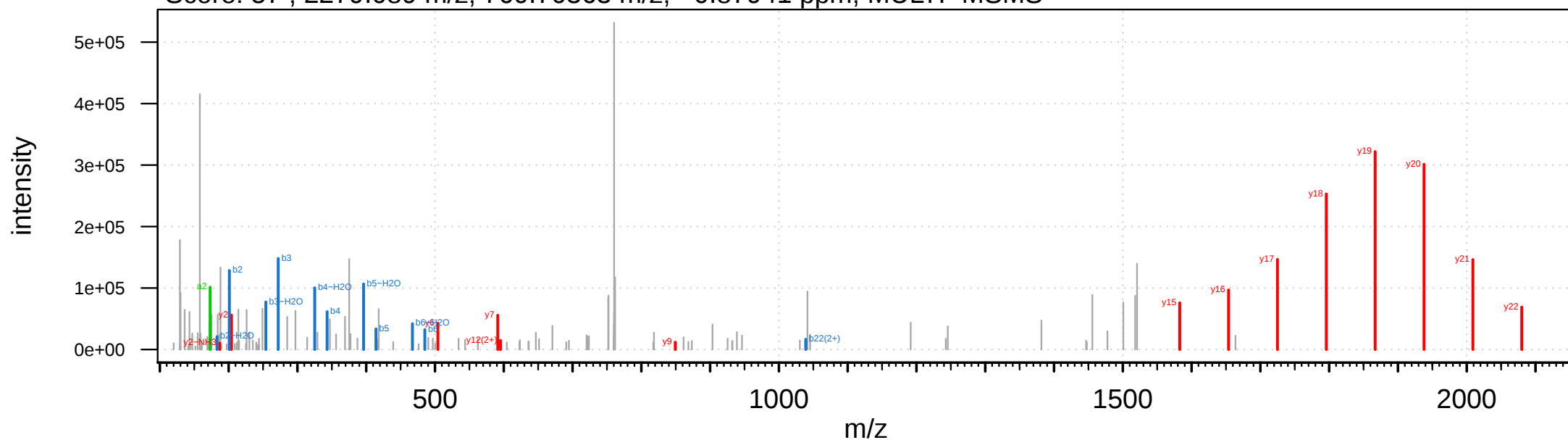
sp|Q69YL0|NCAS2_HUMAN

ENST00000602845_chr3:196669588-196669887:+



EAAAAAAAAAHQQALREESGAGDKG

Score: 57 ; 2279.089 m/z; 760.70363 m/z; -0.87941 ppm; MULTI-MSMS

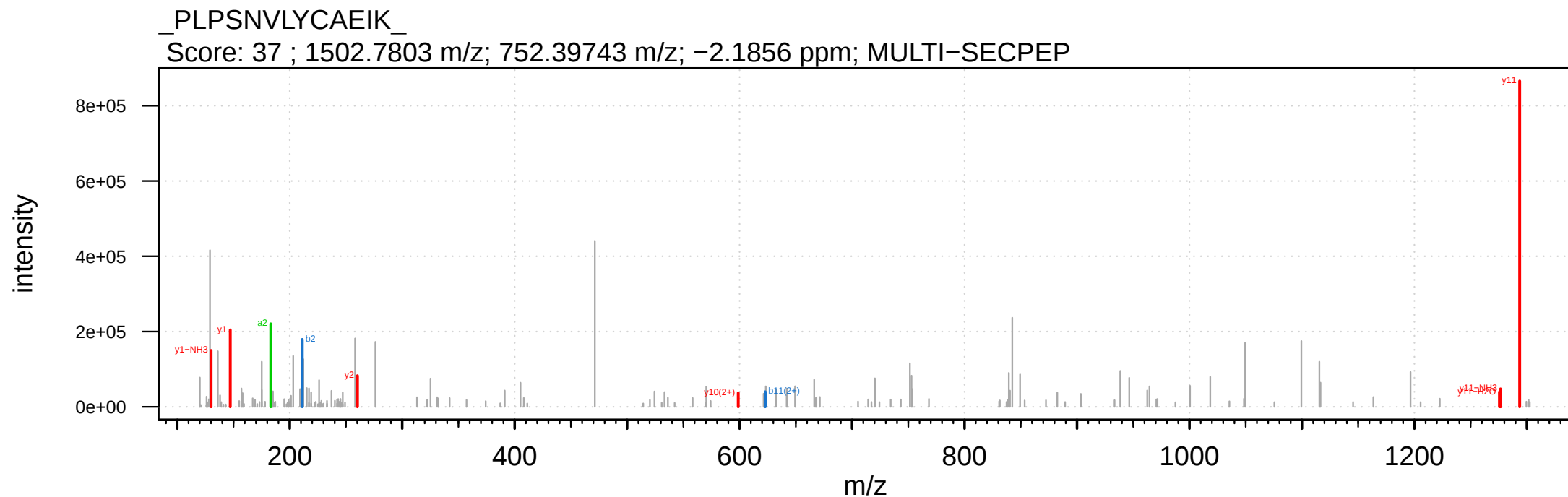
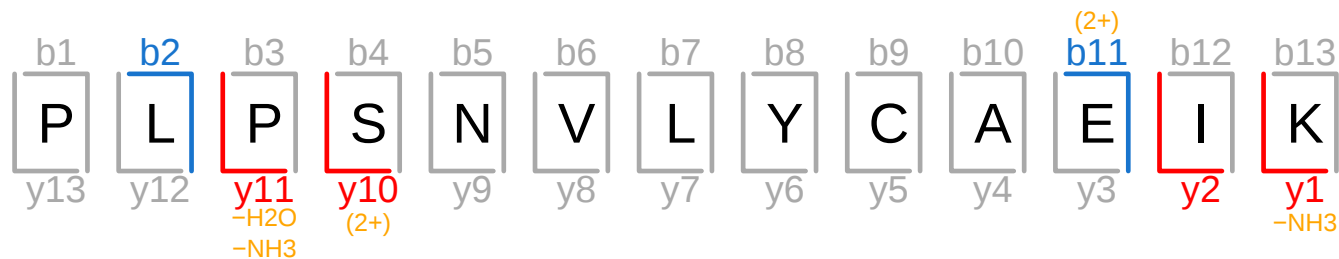


Raw File: Grobi_20121105_CHS_IEF100_20121025_pep_gradient_7cm_250ug_03

Scan Number: 25732

Proteins:

ENST00000394071_chr14:74185546-74185677:-



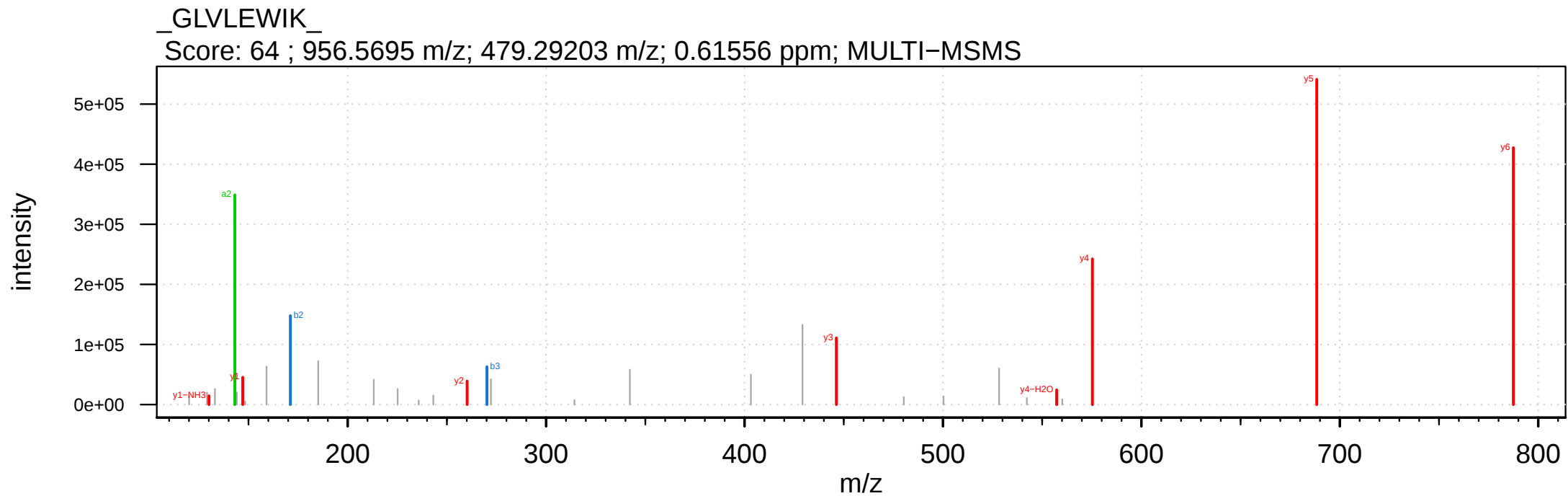
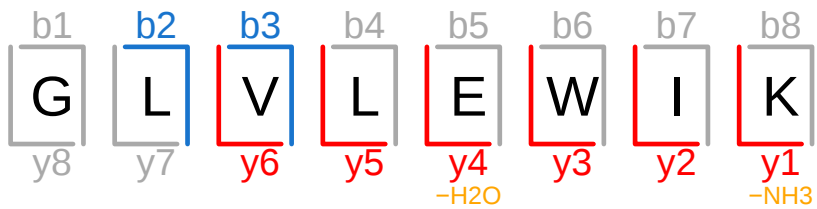
Raw File: Grobi_20121105_CHS_IEF100_20121025_pep_gradient_7cm_250ug_03

Scan Number: 38703

Proteins:

sp|A6NIN4|YQ014_HUMAN

ENST00000324348_chr17:7818840-7819234:-

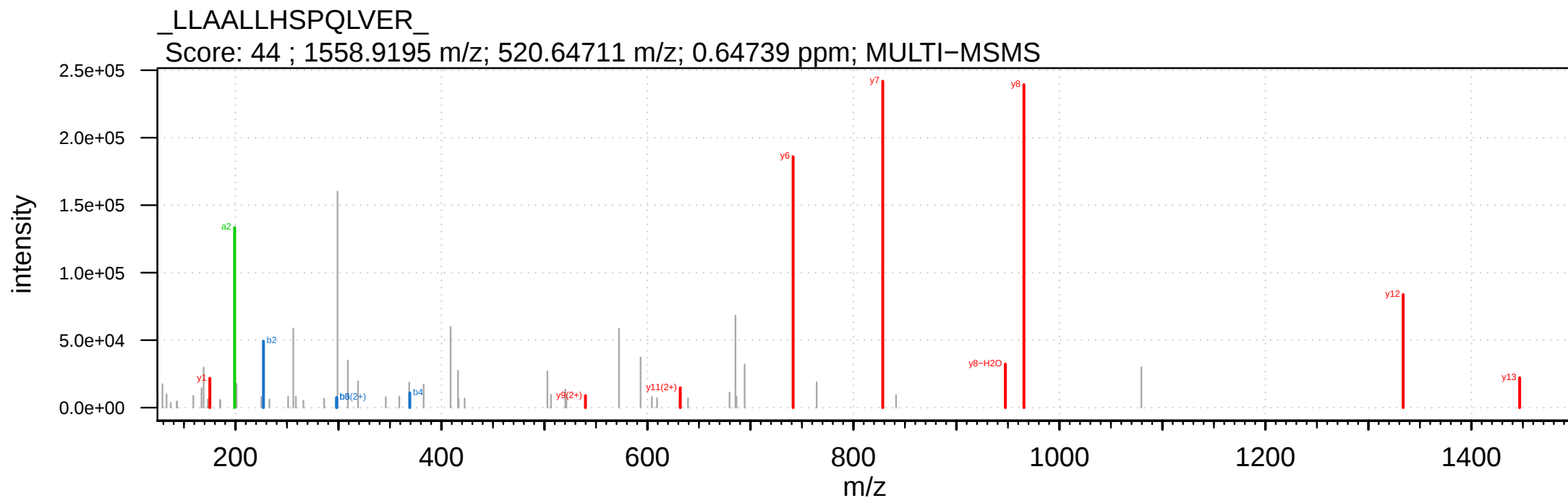
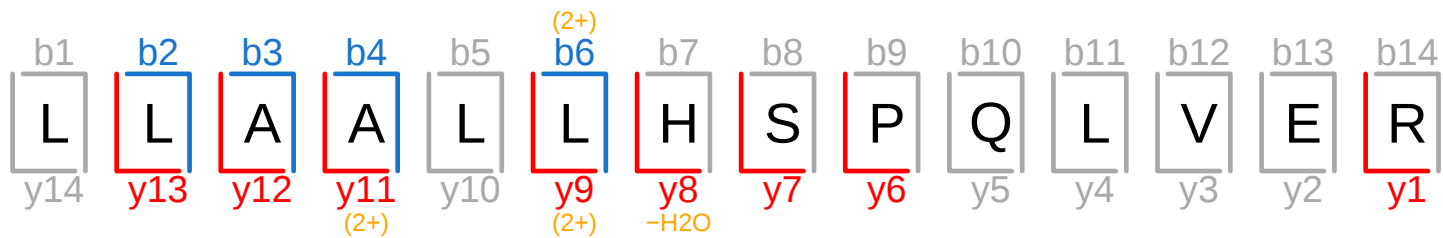


Raw File: Bibo_20121204_CHS_IEF100_20121129_3-10linear_Serva_7cm_250ug_03

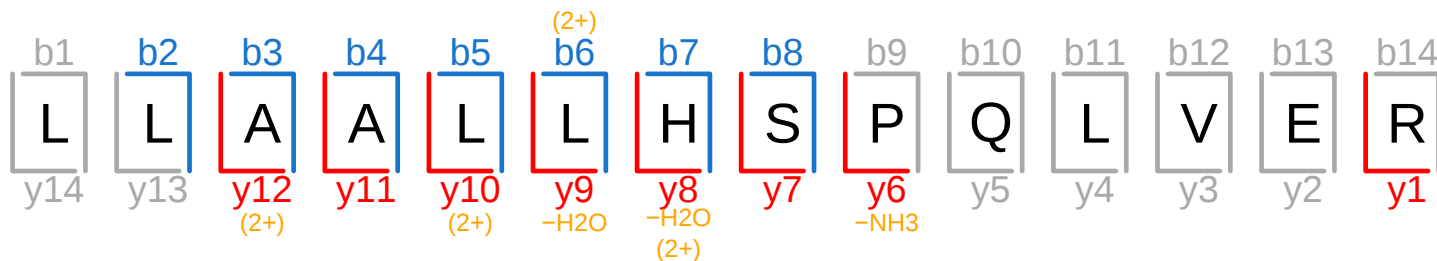
Scan Number: 42970

Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-

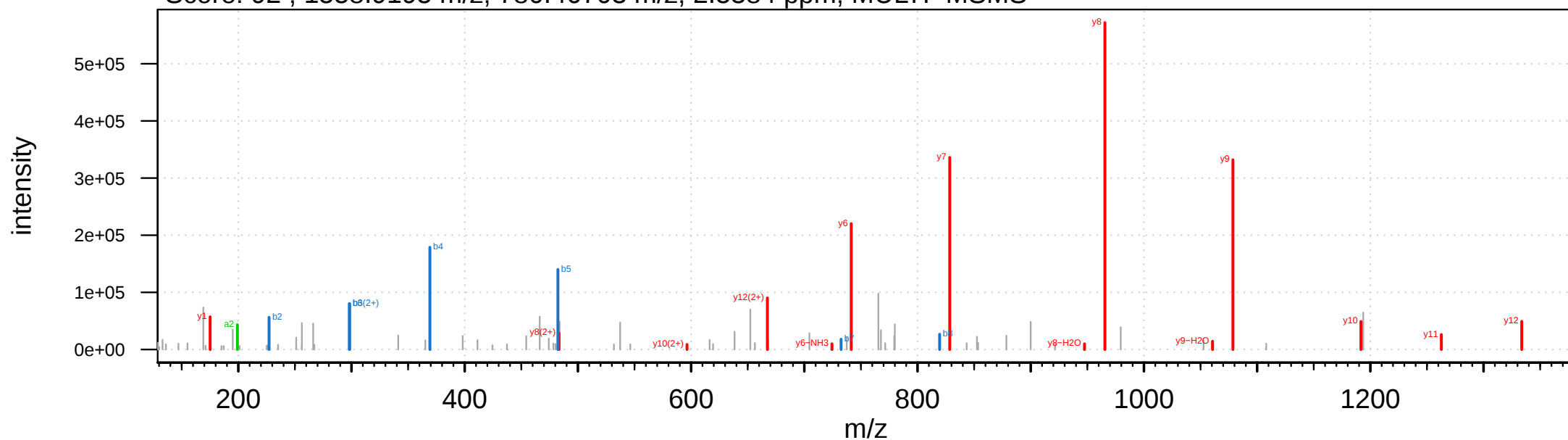


Raw File: Bibo_20130621_CHS_IEF_3-10linear_24slices_15
 Scan Number: 12012
 Proteins:
 sp|Q69YL0|NCAS2_HUMAN
 ENST00000602845_chr3:196669588-196669887:+



LLAALLHSPQLVER

Score: 92 ; 1558.9195 m/z; 780.46703 m/z; 2.5384 ppm; MULTI-MSMS



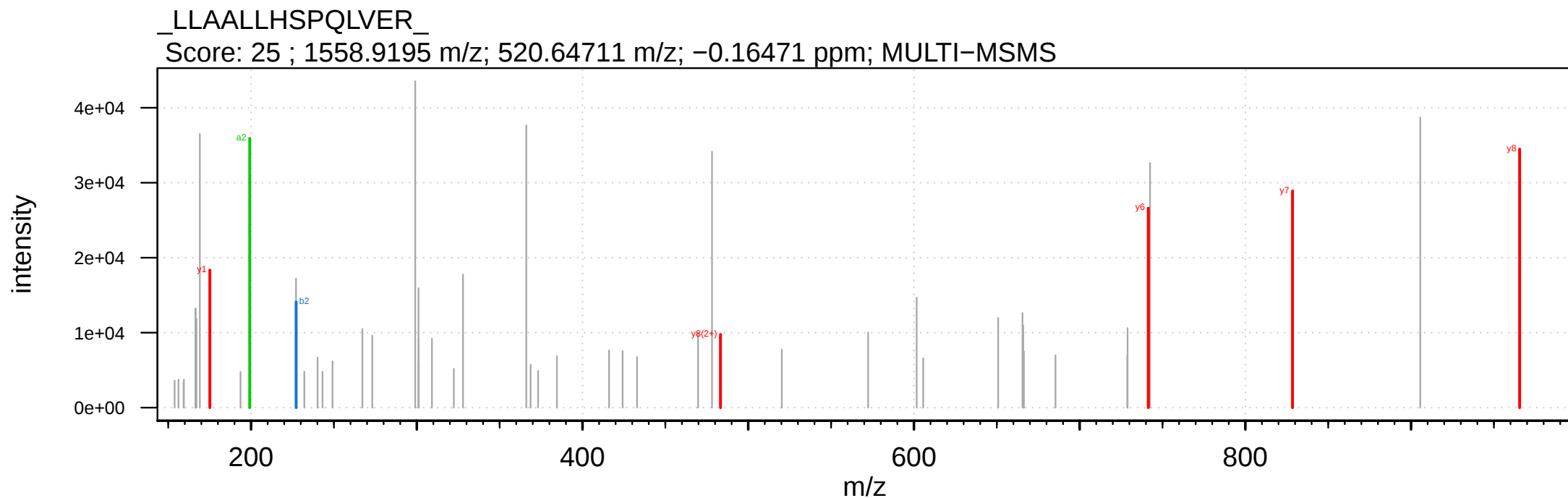
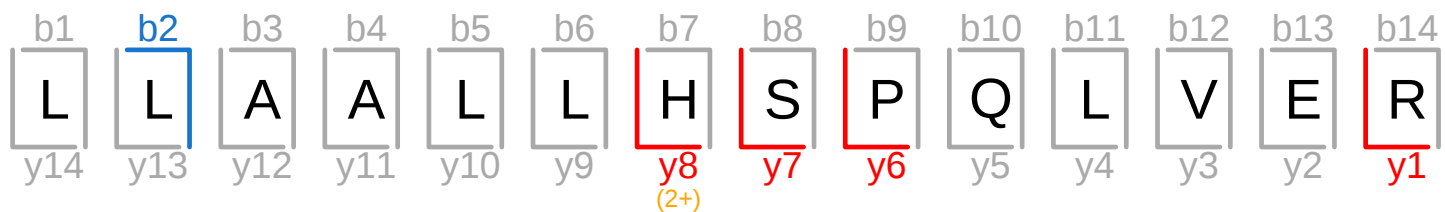
Raw File: Bibo_20130621_CHS_IEF_3-10linearpep_24slices_19

Scan Number: 11681

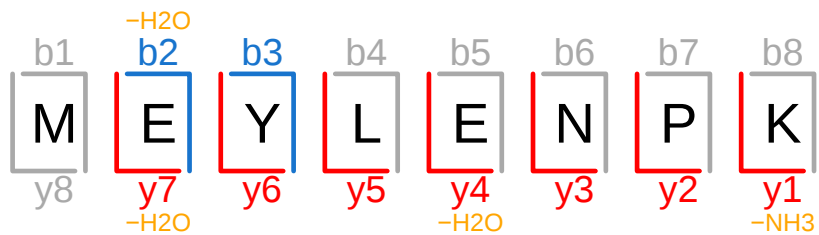
Proteins:

sp|Q69YL0|NCAS2_HUMAN

ENST00000602845_chr3:196669588-196669887:+

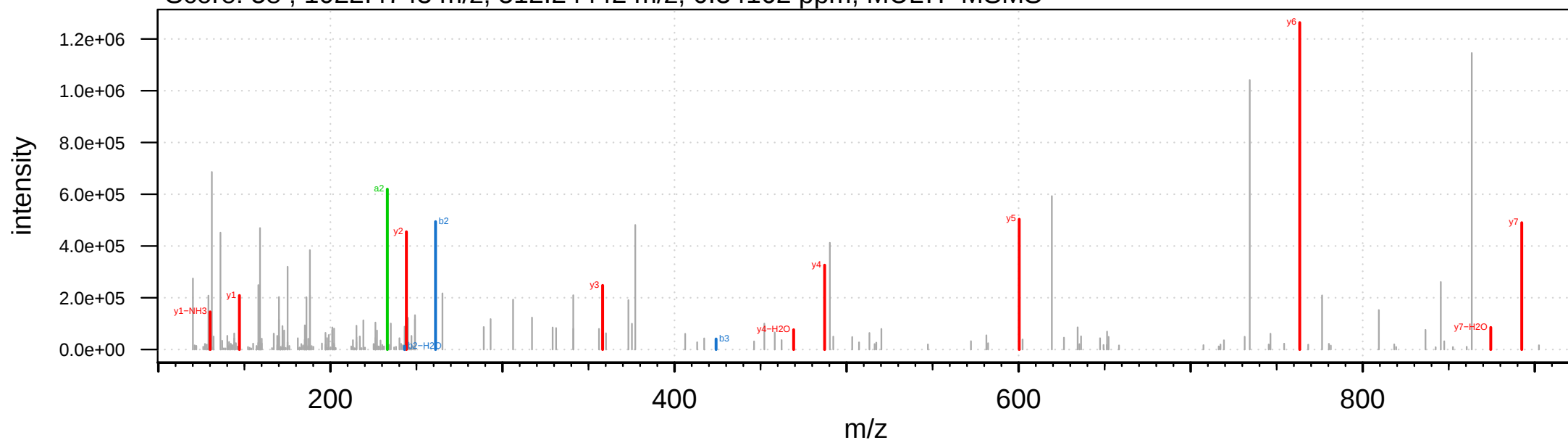


Raw File: Kermit_20130417_CHS_20130123_IEF100_201303xx_3-10linear_Serva_7cm_25ug_04
 Scan Number: 36545
 Proteins:
 sp|Q69YL0|NCAS2_HUMAN
 ENST00000602845_chr3:196669588-196669887:+



MEYLENPK

Score: 58 ; 1022.4743 m/z; 512.24442 m/z; 0.34162 ppm; MULTI-MSMS

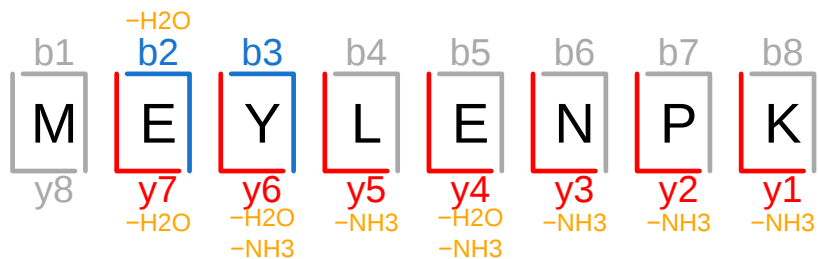


Raw File: Grobi_20121105_CHS_IEF_20121025_pep_gradient_7cm_250ug_01

Scan Number: 26967

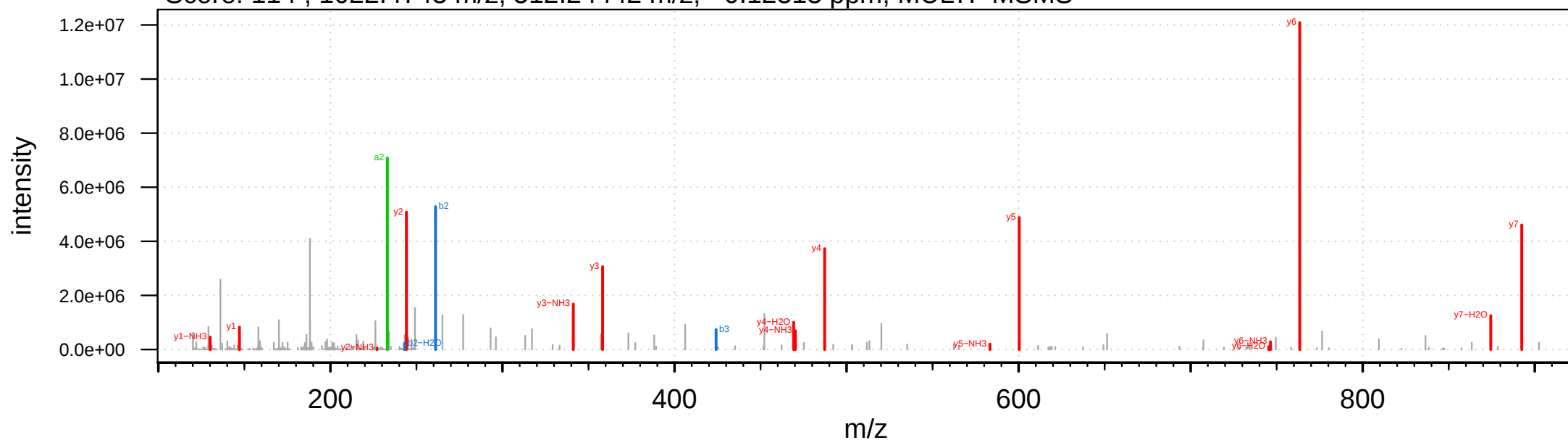
Proteins:

ENST00000518958_chr8:120630400-120630519:-



MEYLENPK

Score: 114 ; 1022.4743 m/z; 512.24442 m/z; -0.12313 ppm; MULTI-MSMS

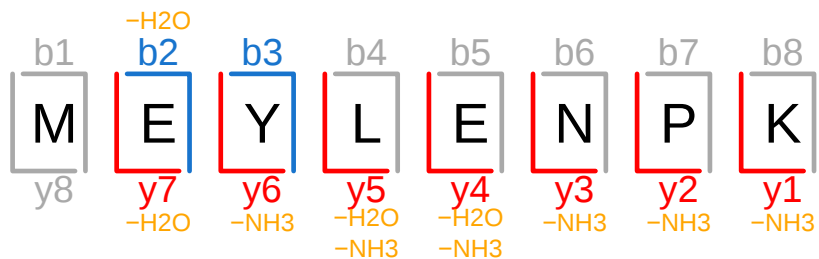


Raw File: Grobi_20121105_CHS_IEF_20121025_pep_gradient_7cm_250ug_02

Scan Number: 25840

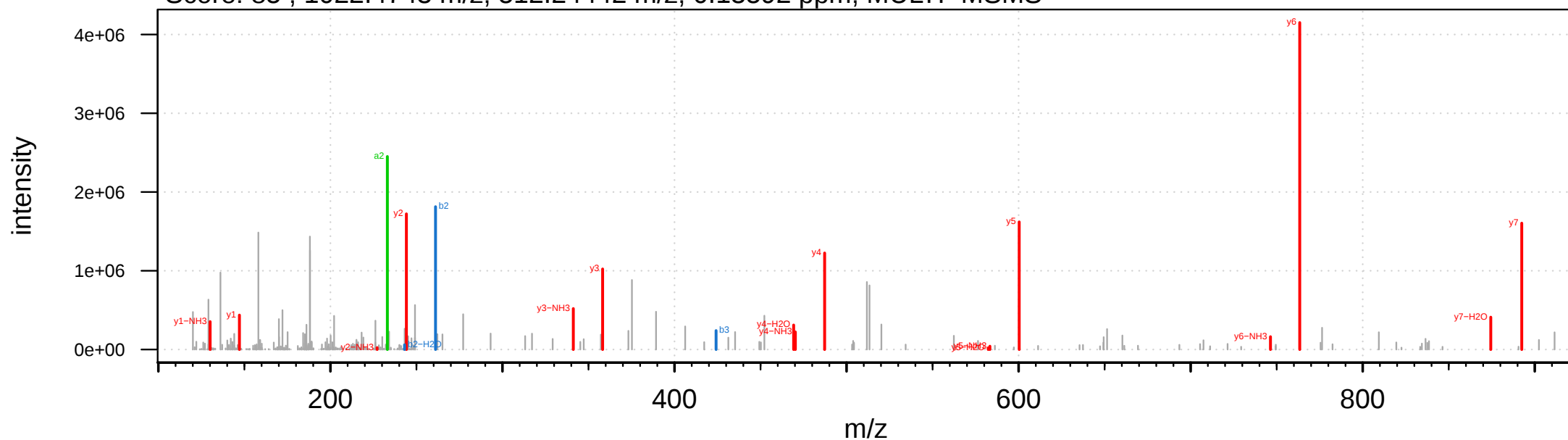
Proteins:

ENST00000518958_chr8:120630400-120630519:-



MEYLENPK

Score: 85 ; 1022.4743 m/z; 512.24442 m/z; 0.13592 ppm; MULTI-MSMS

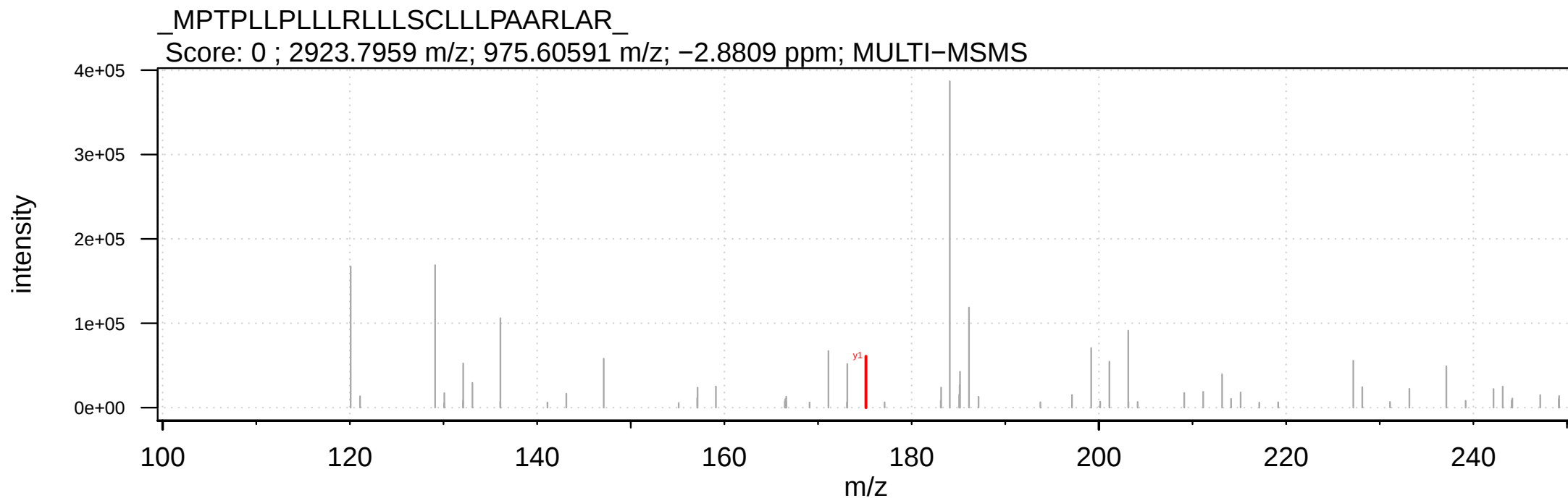
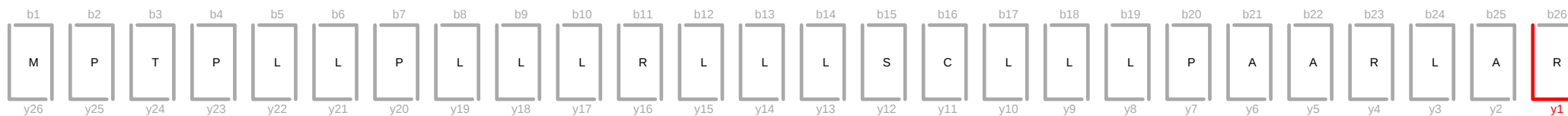


Raw File: Grobi_20121105_CHS_IEF_20121025_pep_gradient_7cm_250ug_03

Scan Number: 26964

Proteins:

ENST00000518958_chr8:120630400-120630519:-

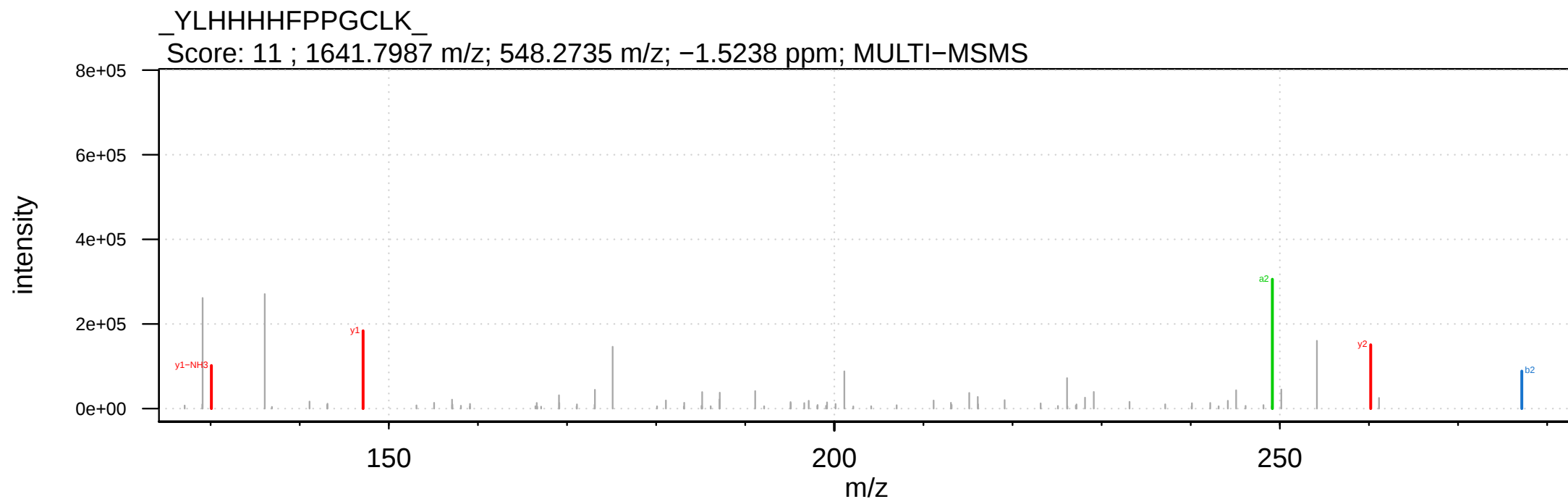
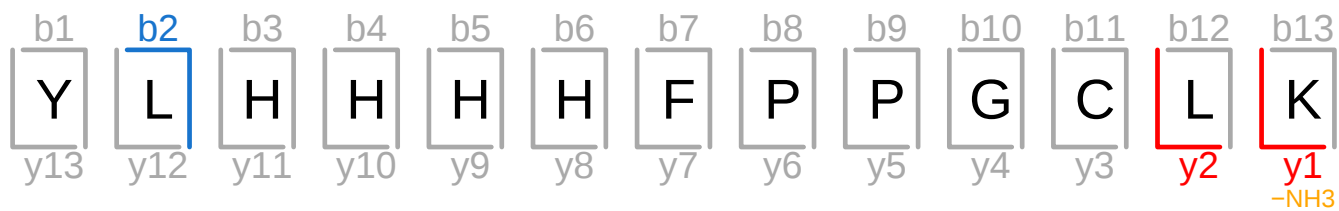


Raw File: Kermit_20130417_CHS_20130123_IEF100_201303xx_25ug

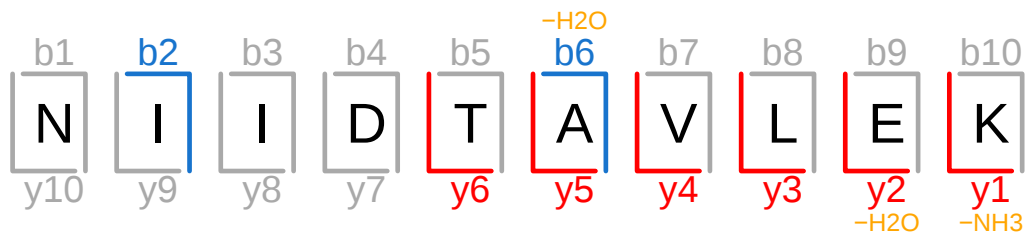
Scan Number: 103661

Proteins:

ENST00000573382_chr6:44185209-44185463:+

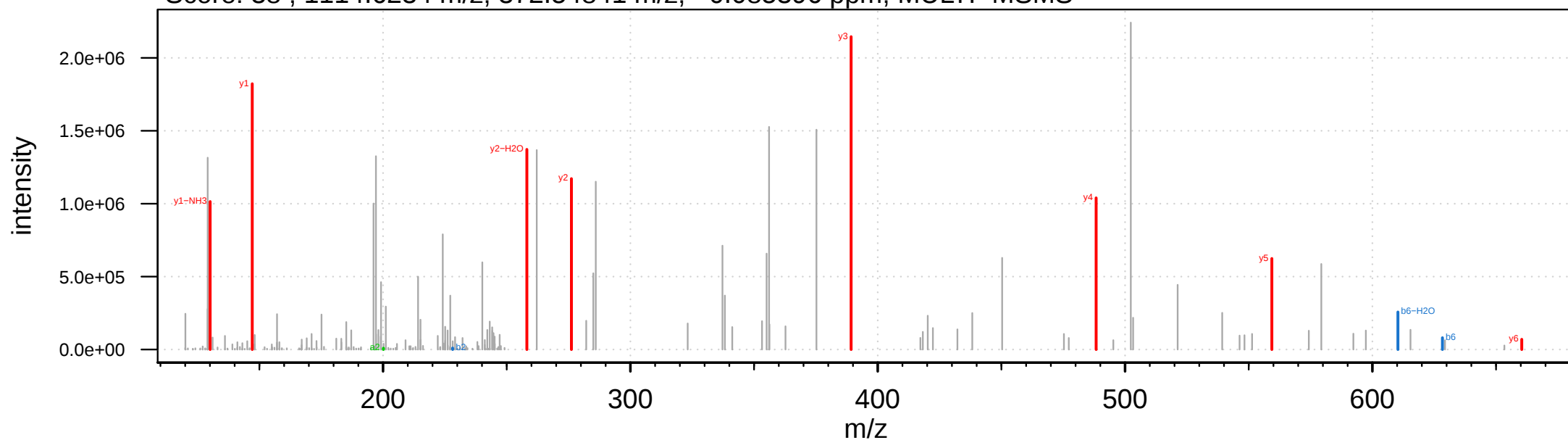


Raw File: Kermit_20130417_CHS_20130123_IEF100_201303xx_25ug
Scan Number: 13176
Proteins:
ENST00000525328_chr11:76423734-76423856:-



NIIDTAVLEK

Score: 38 ; 1114.6234 m/z; 372.54841 m/z; -0.085396 ppm; MULTI-MSMS

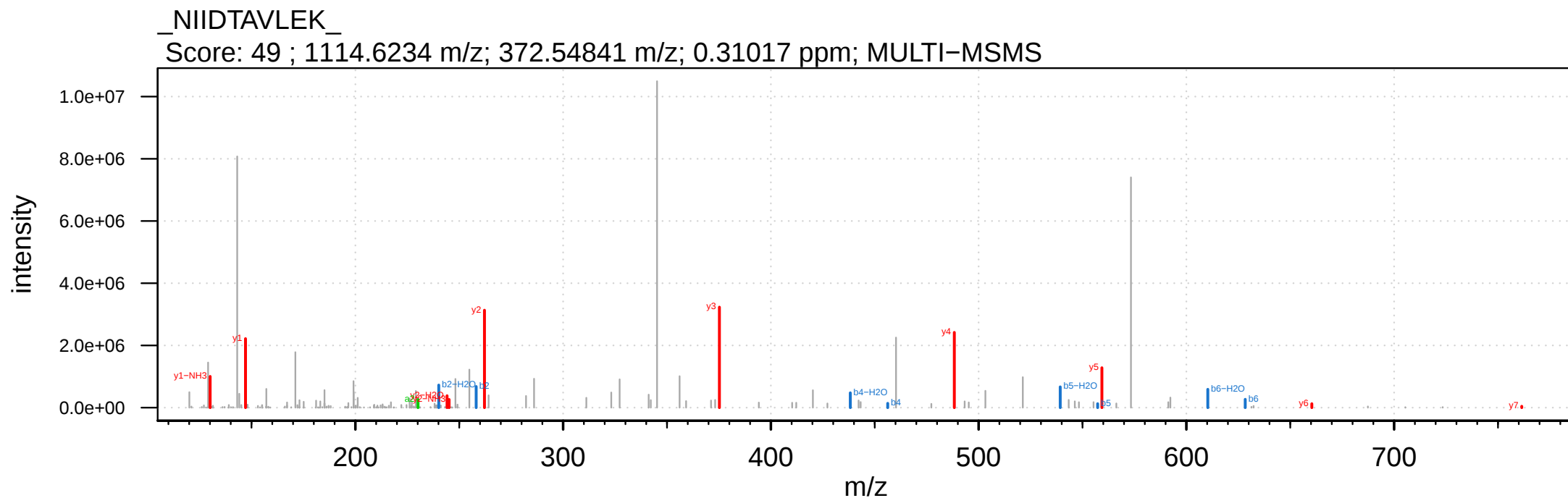
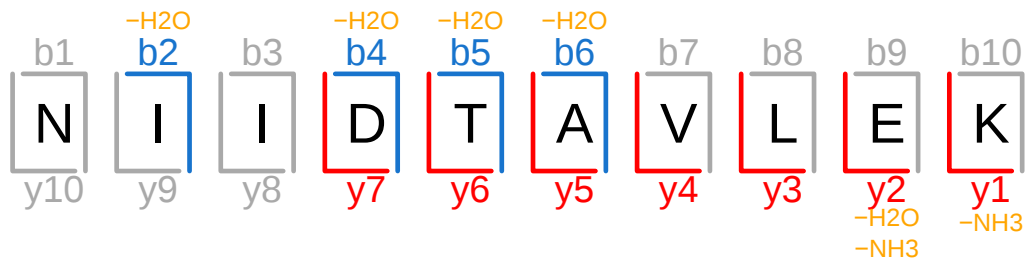


Raw File: Grobi_20121019_CHS_PIECEIEF_20121017_3-10_7cm_250ug_04

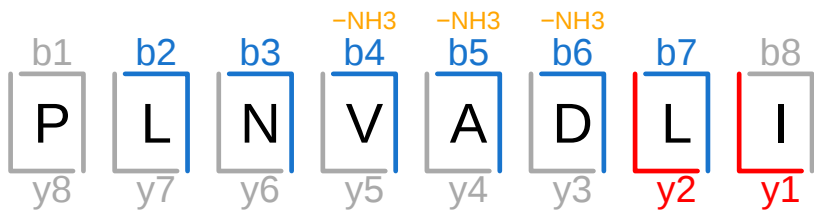
Scan Number: 19556

Proteins:

ENST00000497138_chr20:56806826-56807846:-

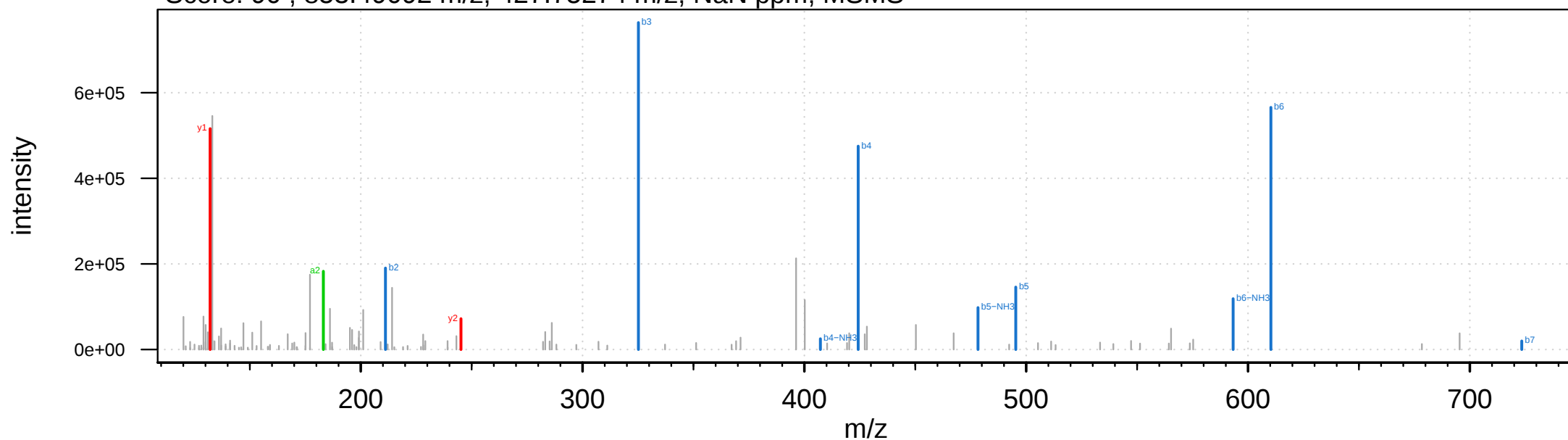


Raw File: Grobi_20130312_CHS_IEF100_20130309_3-10linear_Serva_7cm_1mg_1elution_03
Scan Number: 4098
Proteins:
ENST00000497138_chr20:56806826-56807846:-



PLNVADLI

Score: 90 ; 853.49092 m/z; 427.75274 m/z; NaN ppm; MSMS

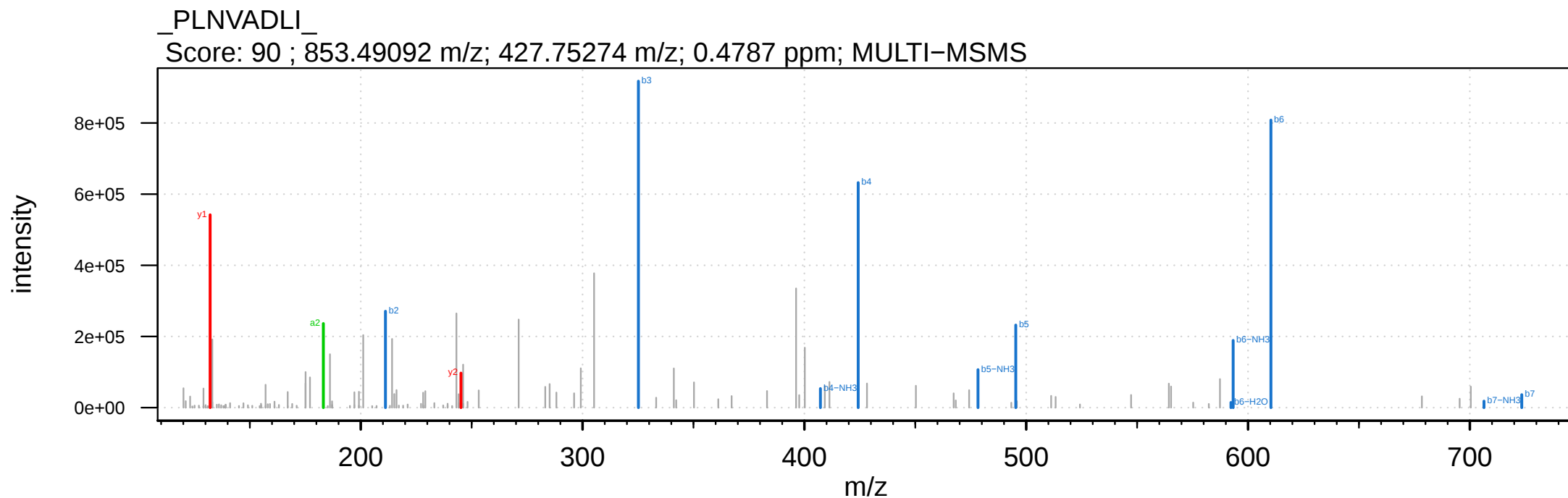
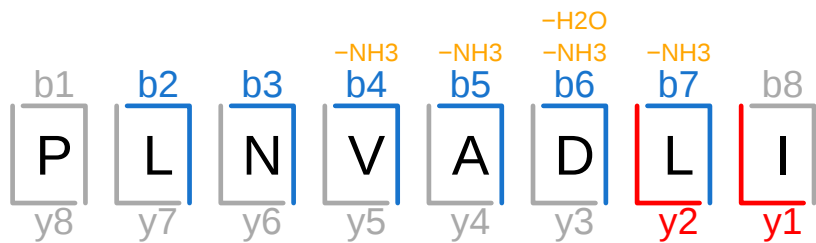


Raw File: Grobi_20121019_CHS_PIECEIEF_20121017_3-10_7cm_250ug_02

Scan Number: 75988

Proteins:

TCONS_I2_00005685_chr12:53548245-53548274:+

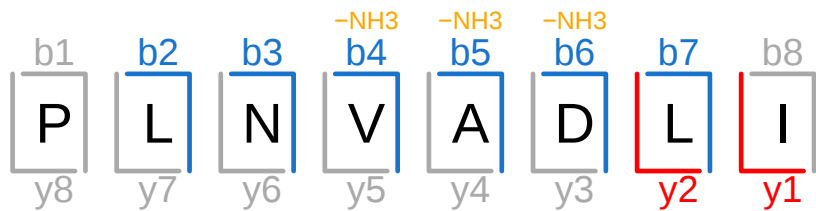


Raw File: Grobi_20121105_CHS_IEF100_20121025_pep_gradient_7cm_250ug_01

Scan Number: 83975

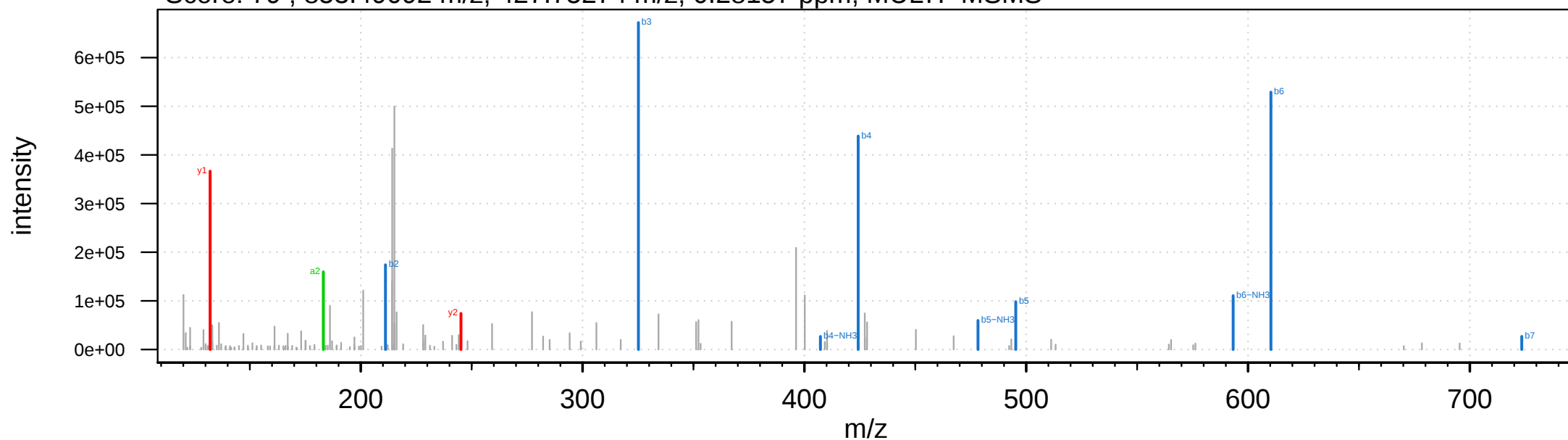
Proteins:

TCONS_I2_00005685_chr12:53548245-53548274:+



PLNVADLI

Score: 79 ; 853.49092 m/z; 427.75274 m/z; 0.28157 ppm; MULTI-MSMS

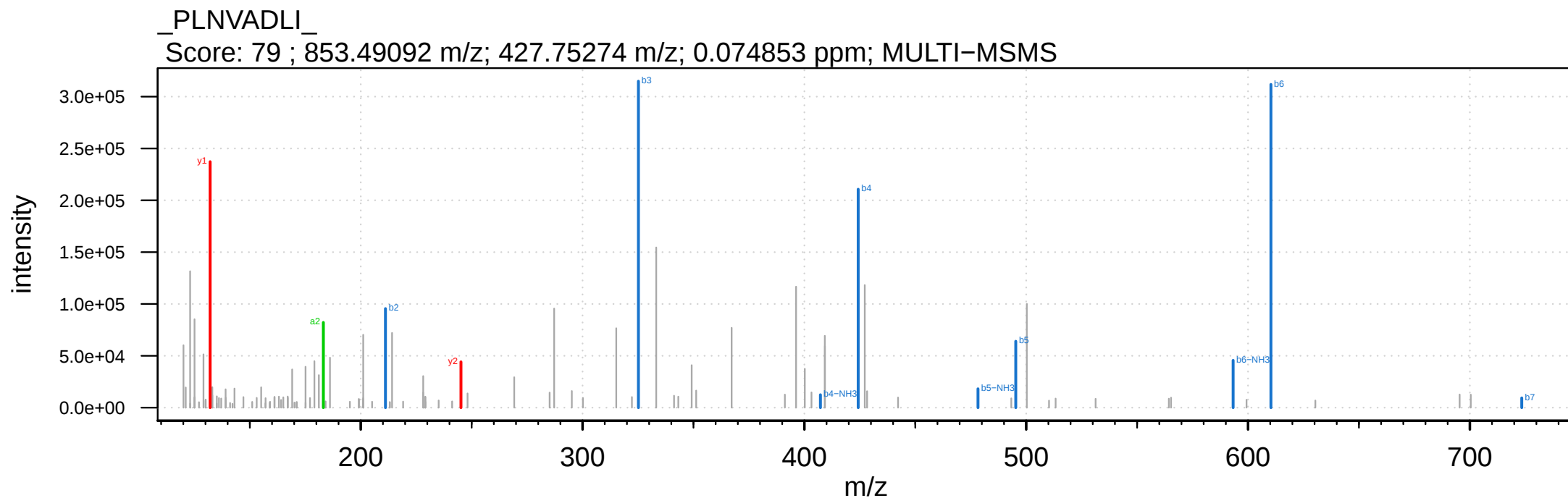
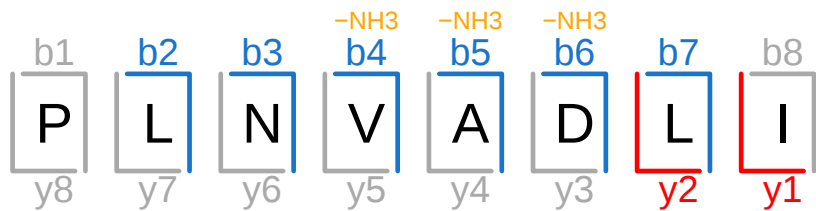


Raw File: Grobi_20121105_CHS_IEF100_20121025_pep_gradient_7cm_250ug_02

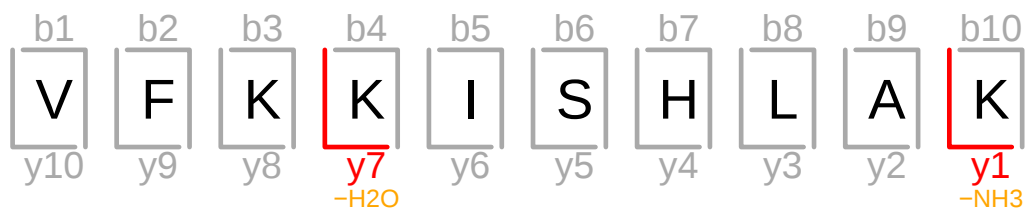
Scan Number: 84162

Proteins:

TCONS_I2_00005685_chr12:53548245-53548274:+

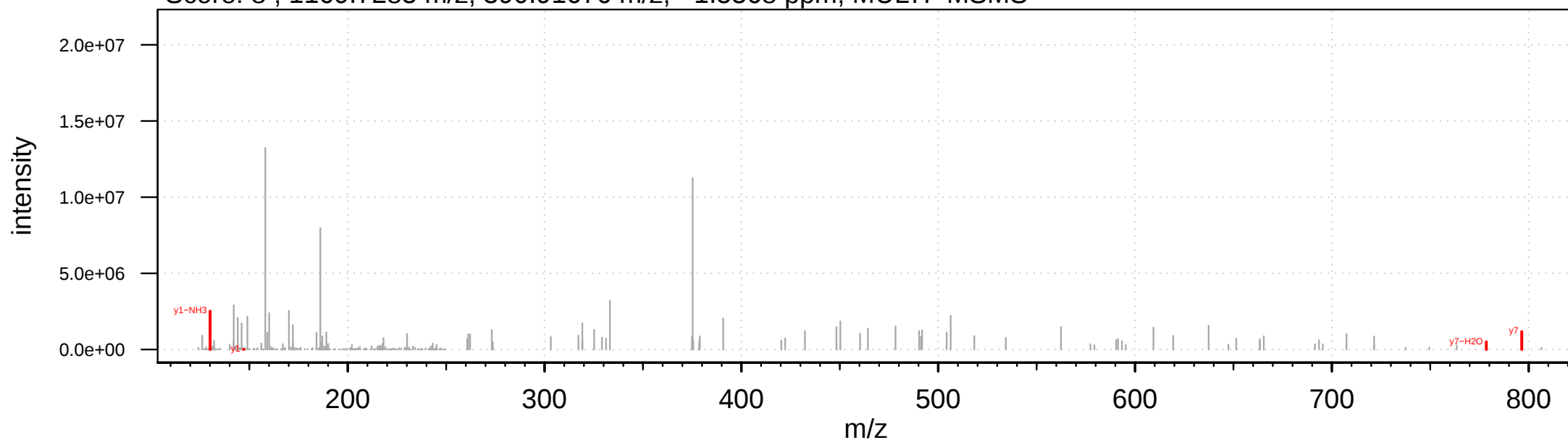


Raw File: Grobi_20130312_CHS_IEF100_20130309_3-10linear_Serva_7cm_1mg_1elution_01
 Scan Number: 52743
 Proteins:
 TCONS_I2_00005685_chr12:53548245-53548274:+



VFKKISHLAK

Score: 8 ; 1169.7285 m/z; 390.91676 m/z; -1.5368 ppm; MULTI-MSMS

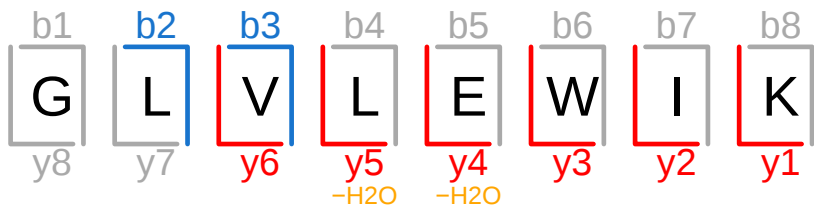


Raw File: Grobi_20121019_CHS_PIECEIEF_20121017_3-10_7cm_250ug_03

Scan Number: 8071

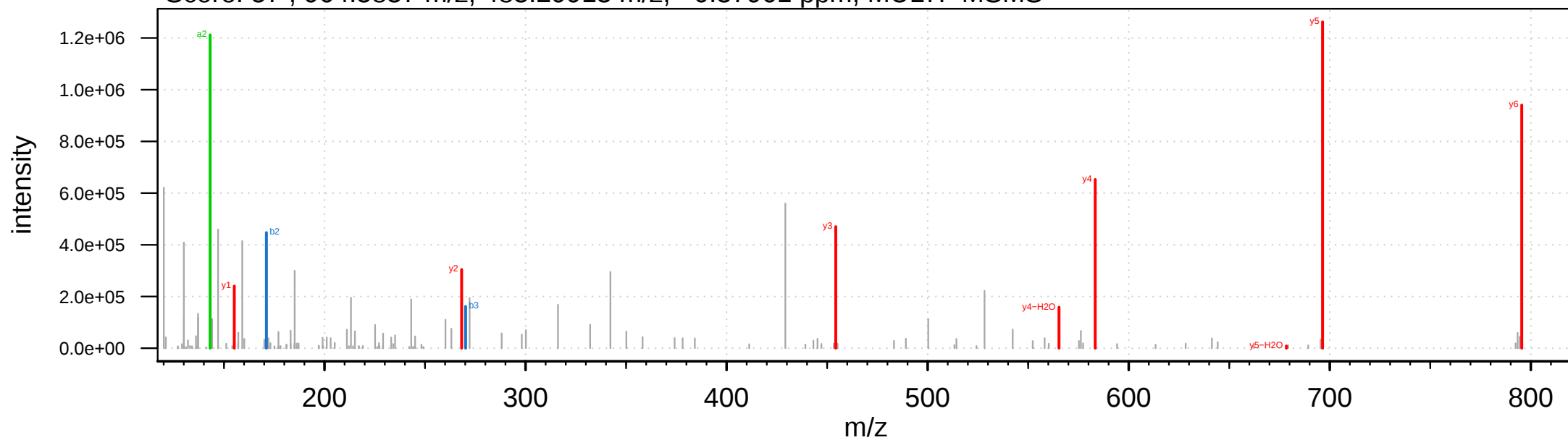
Proteins:

ENST00000490272_chr19:52826295-52826555:+



GLVLEWIK

Score: 57 ; 964.5837 m/z; 483.29913 m/z; -0.37961 ppm; MULTI-MSMS

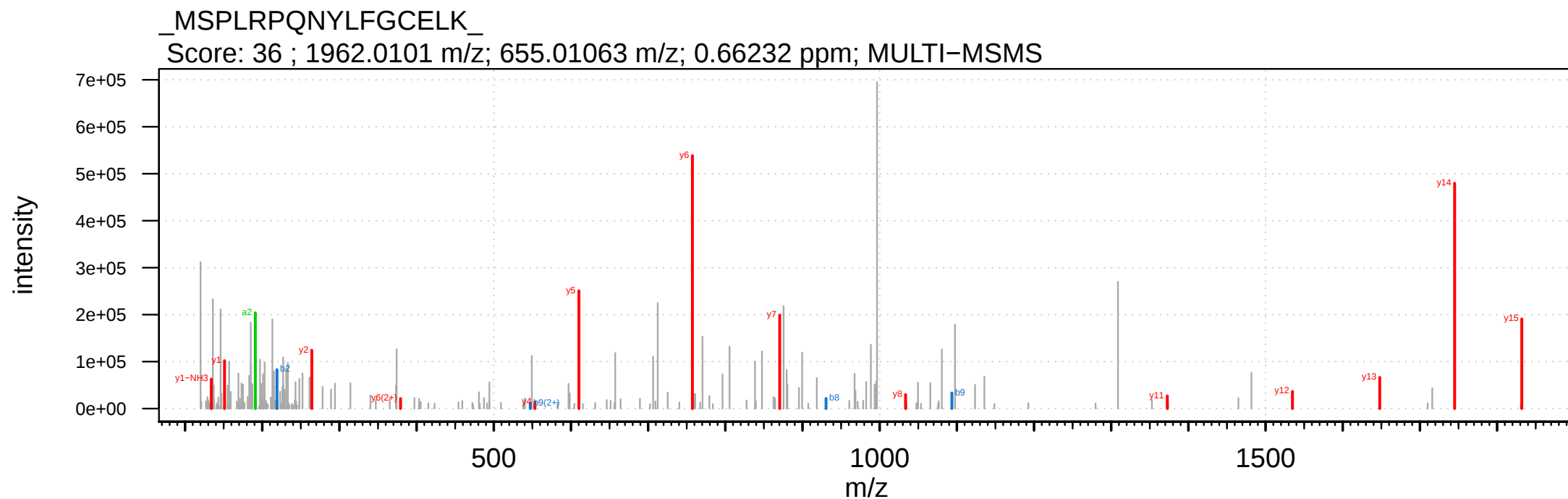
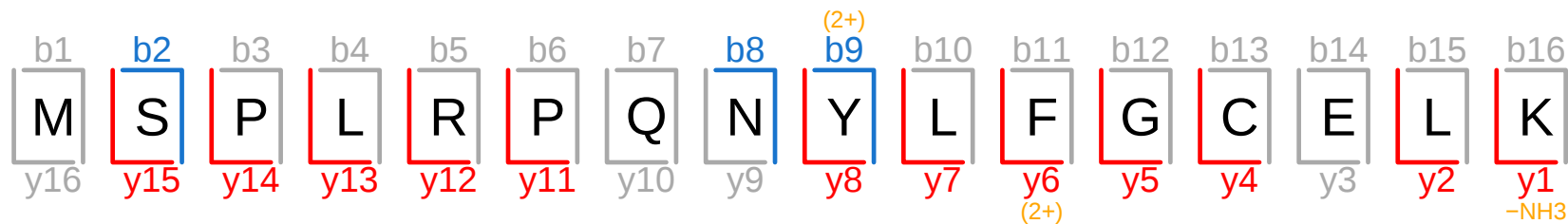


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

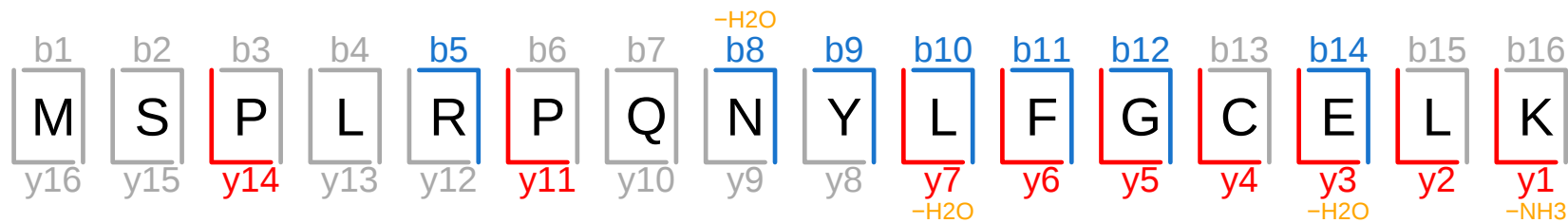
Scan Number: 73510

Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-

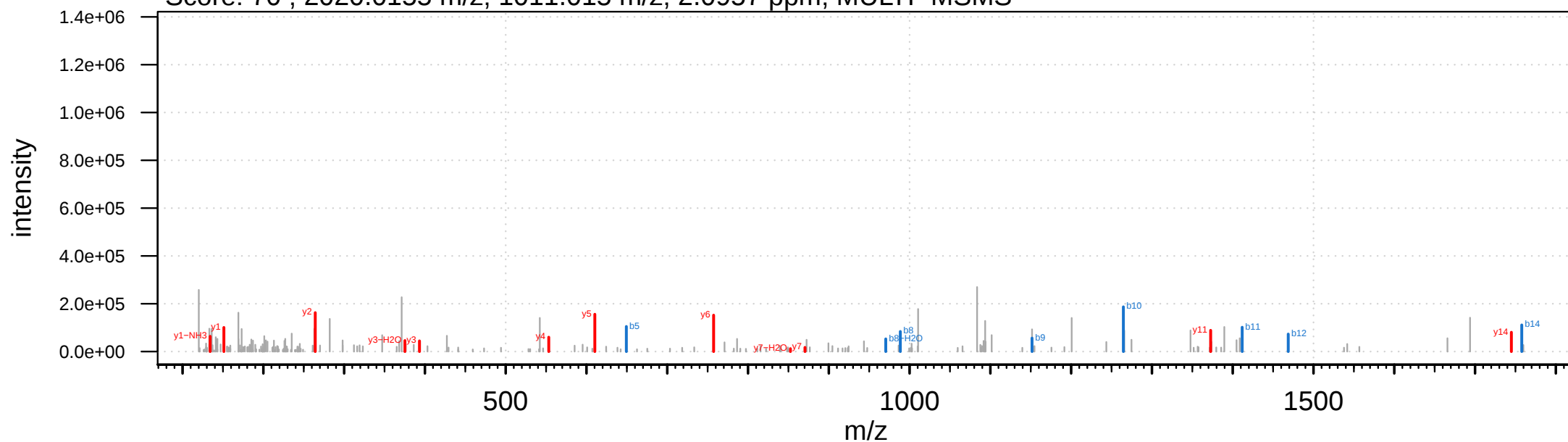


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac
Scan Number: 64401
Proteins:
TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)M(ox)SPLRPQNYLFGCELK

Score: 70 ; 2020.0155 m/z; 1011.015 m/z; 2.0957 ppm; MULTI-MSMS

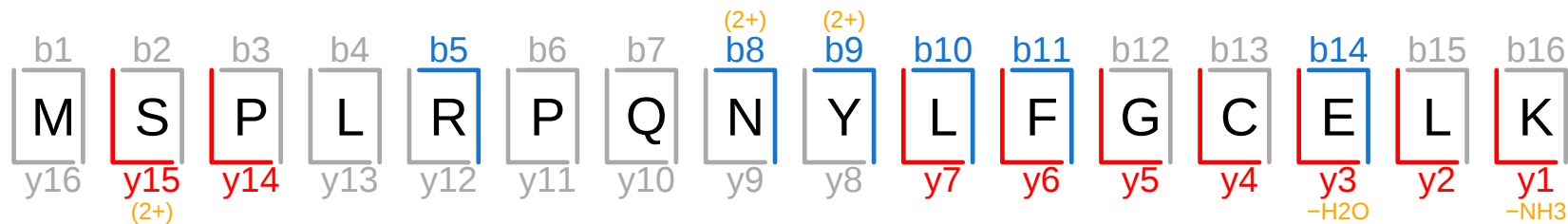


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

Scan Number: 72860

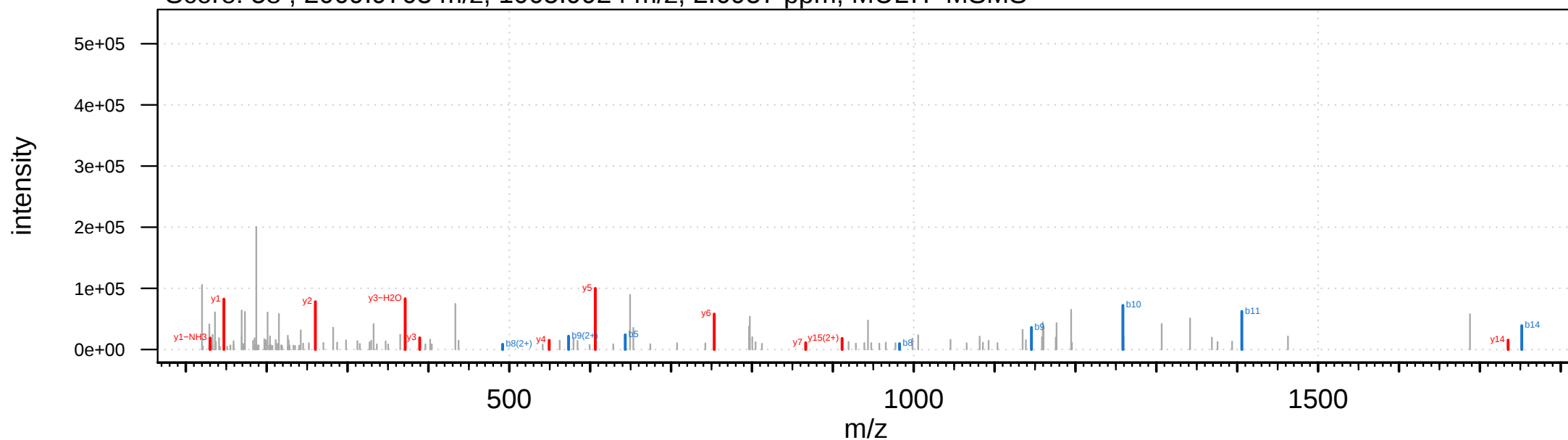
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)M(ox)SPLRPQNYLFGCELK

Score: 58 ; 2009.9703 m/z; 1005.9924 m/z; 2.0957 ppm; MULTI-MSMS

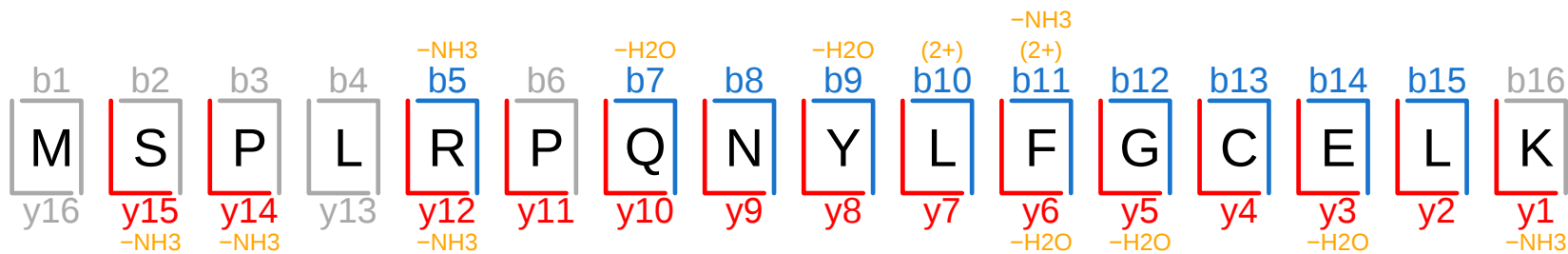


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

Scan Number: 72915

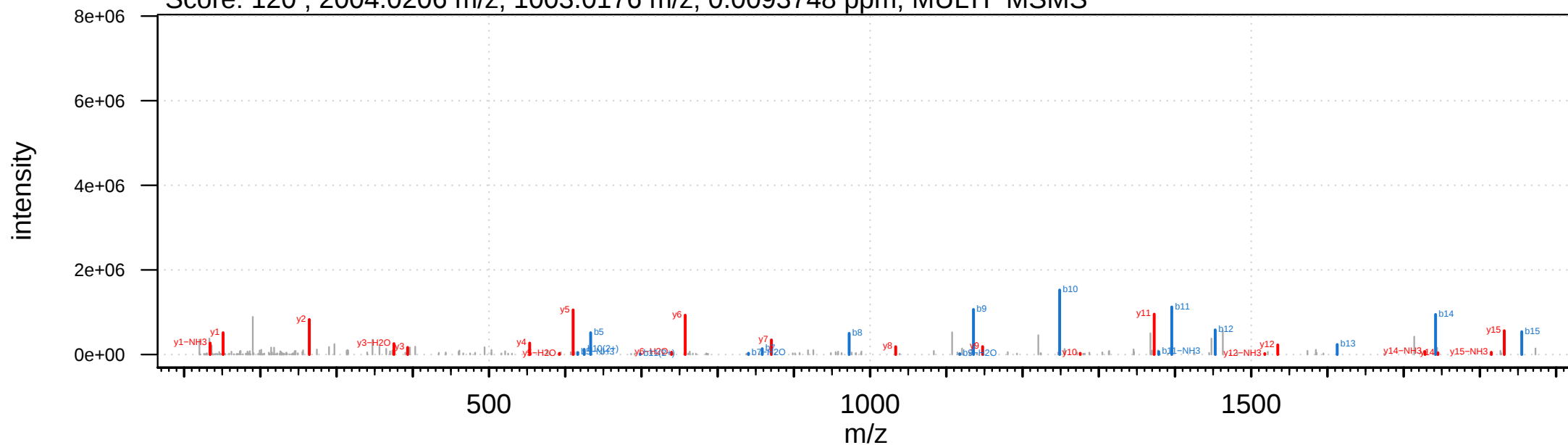
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 120 ; 2004.0206 m/z; 1003.0176 m/z; 0.0093748 ppm; MULTI-MSMS

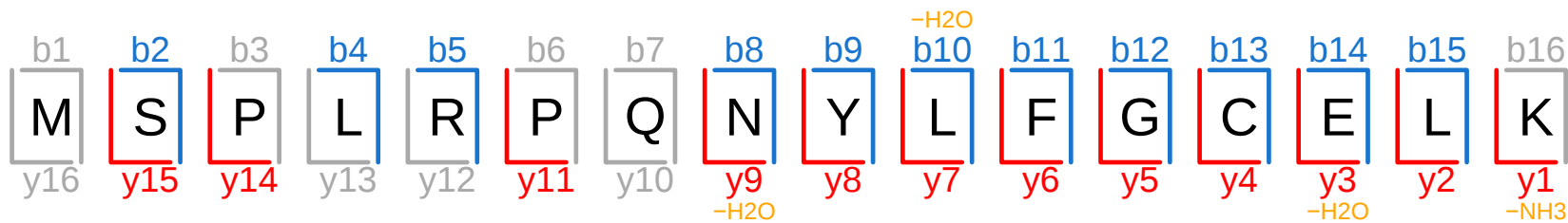


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

Scan Number: 79460

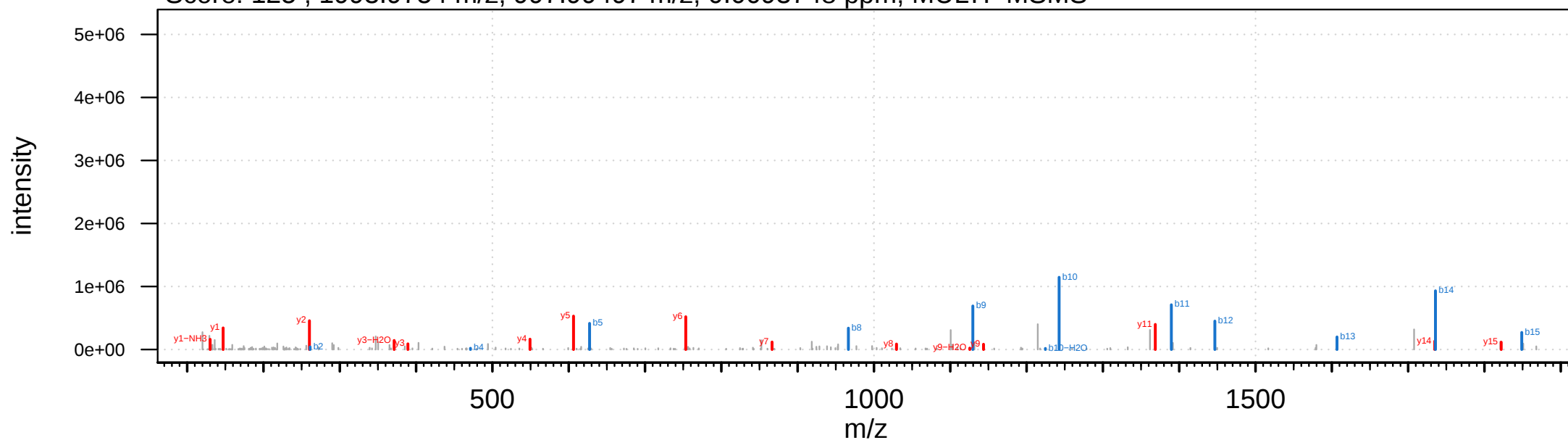
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 125 ; 1993.9754 m/z; 997.99497 m/z; 0.0093748 ppm; MULTI-MSMS

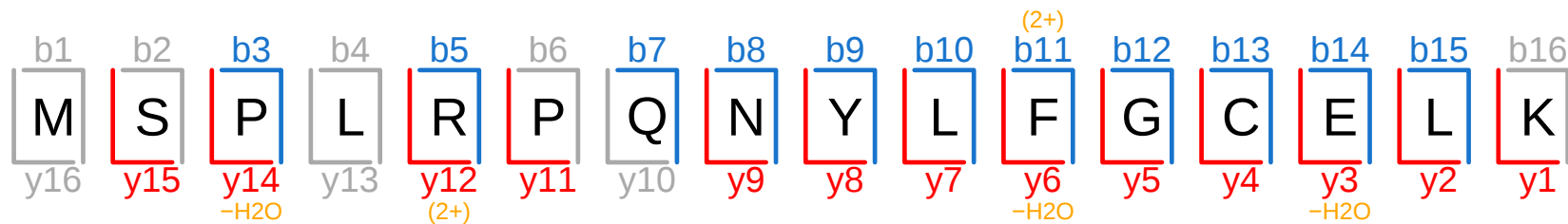


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

Scan Number: 79472

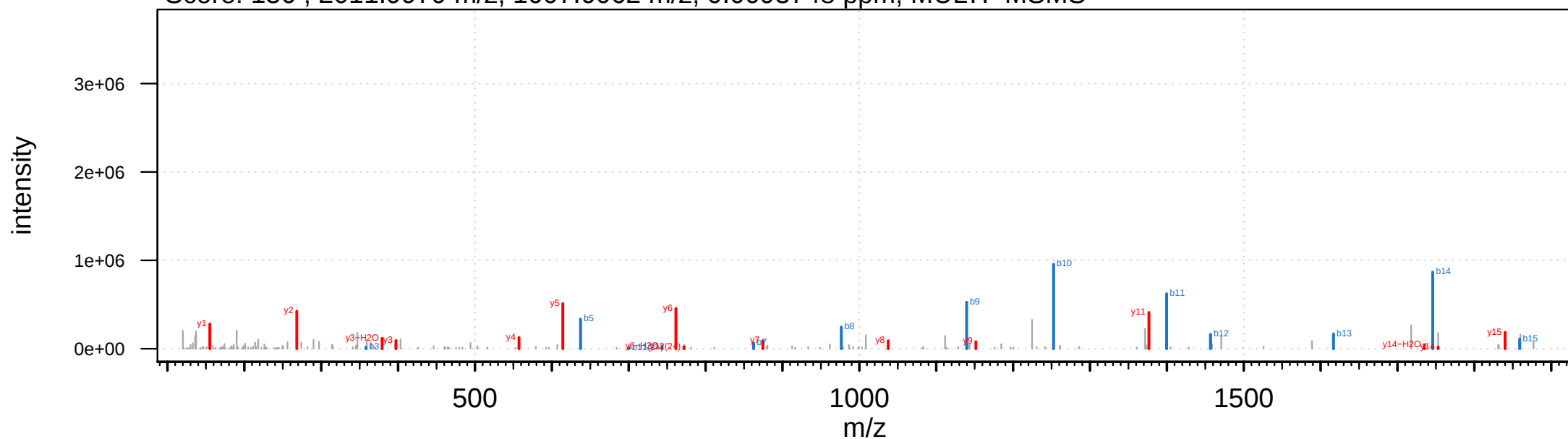
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 136 ; 2011.9979 m/z; 1007.0062 m/z; 0.0093748 ppm; MULTI-MSMS

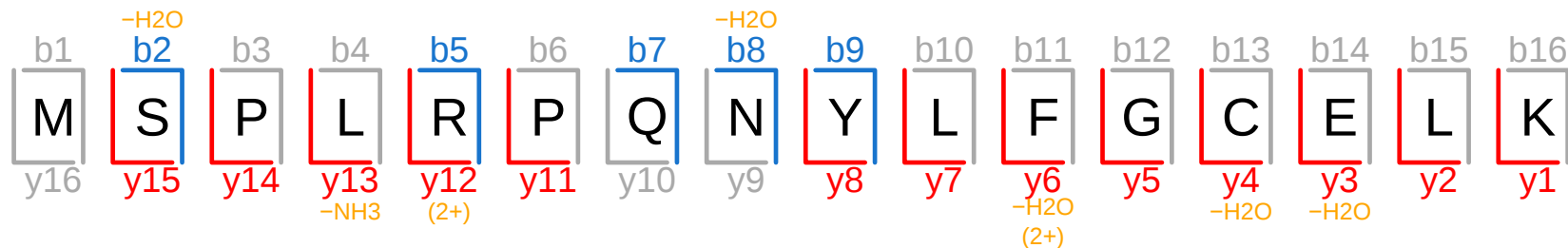


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

Scan Number: 79493

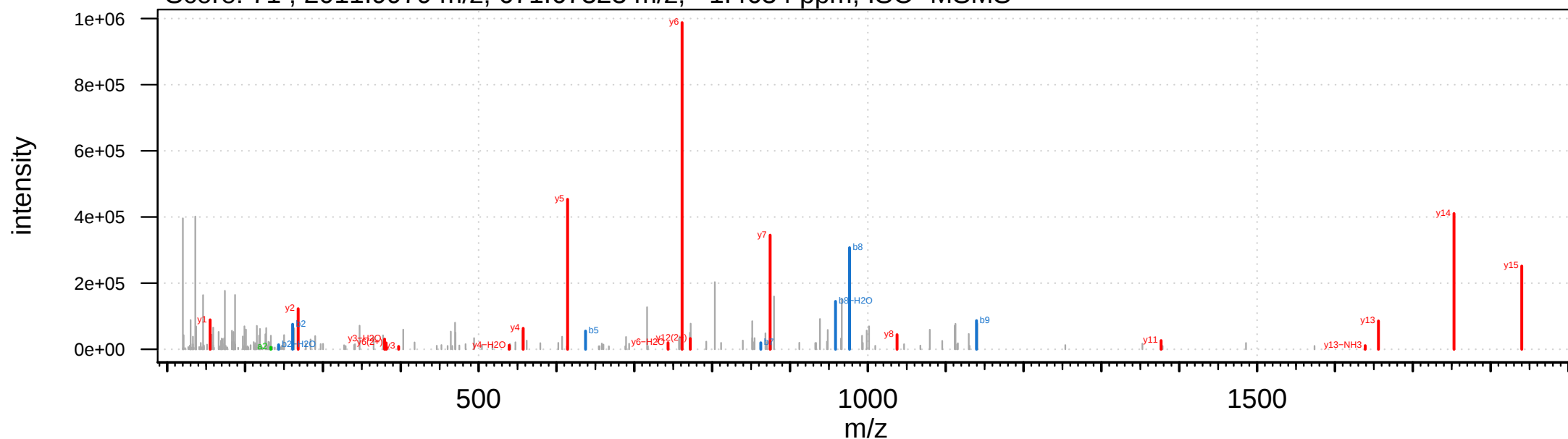
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 71 ; 2011.9979 m/z; 671.67323 m/z; -1.4634 ppm; ISO-MSMS

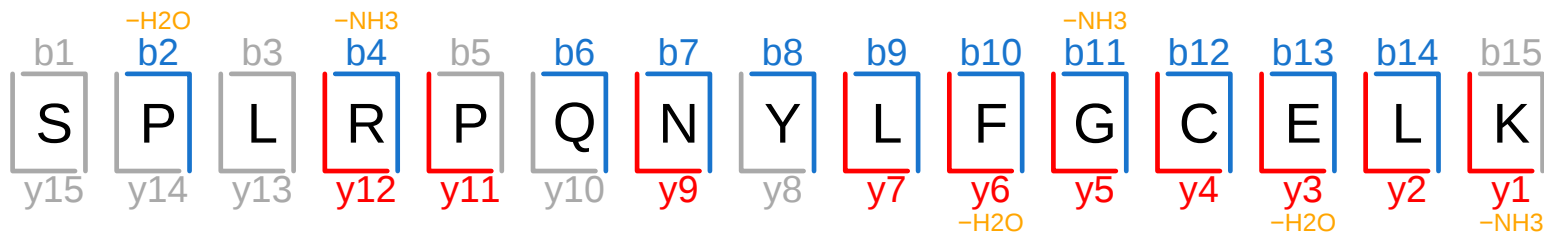


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

Scan Number: 79520

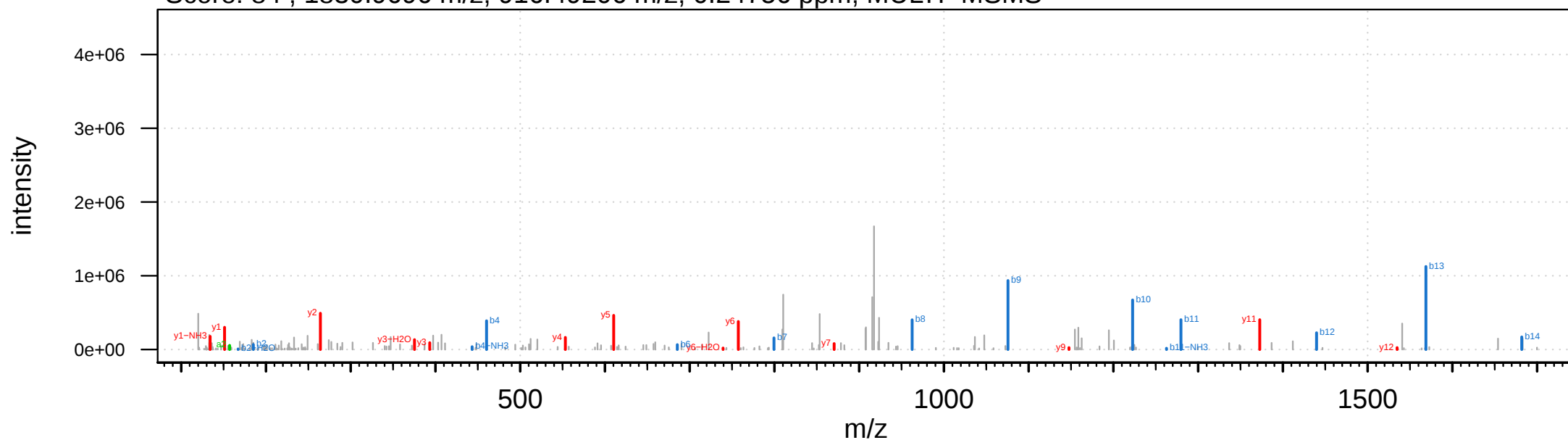
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 84 ; 1830.9696 m/z; 916.49206 m/z; 0.24756 ppm; MULTI-MSMS

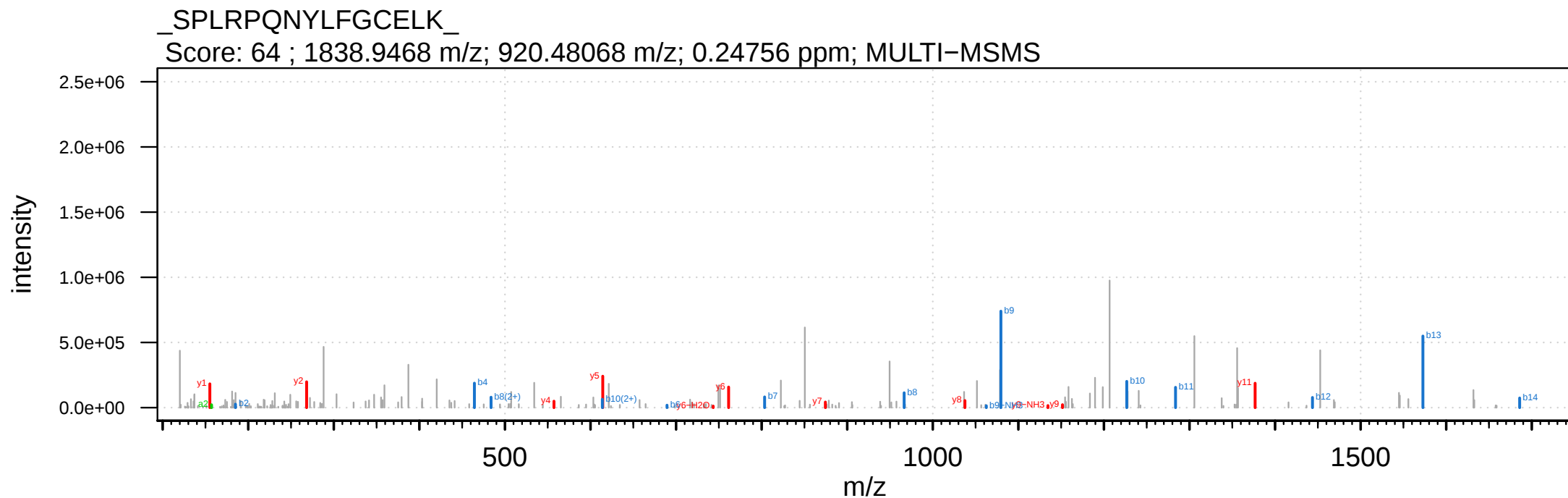
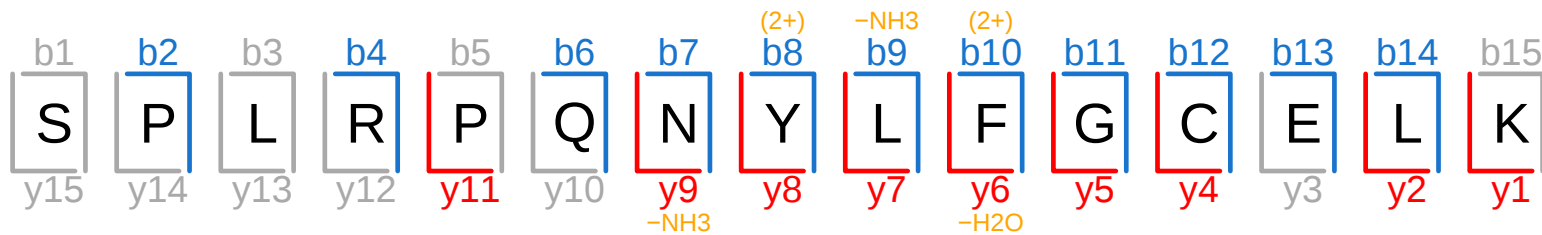


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

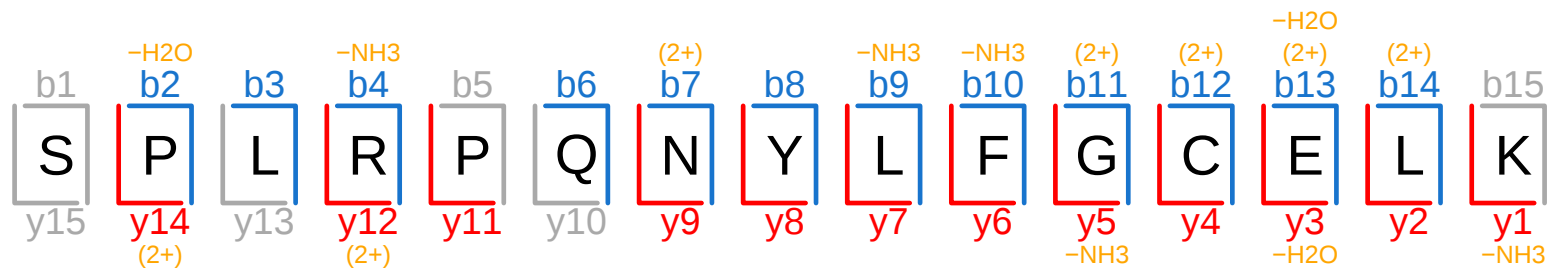
Scan Number: 60226

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

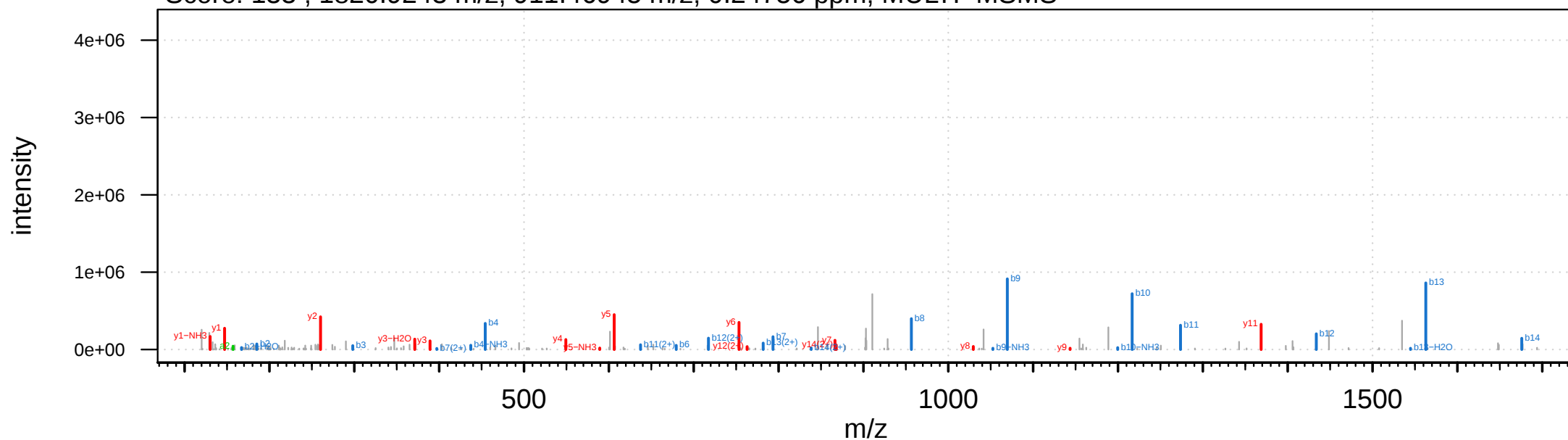


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac
 Scan Number: 60231
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 133 ; 1820.9243 m/z; 911.46945 m/z; 0.24756 ppm; MULTI-MSMS

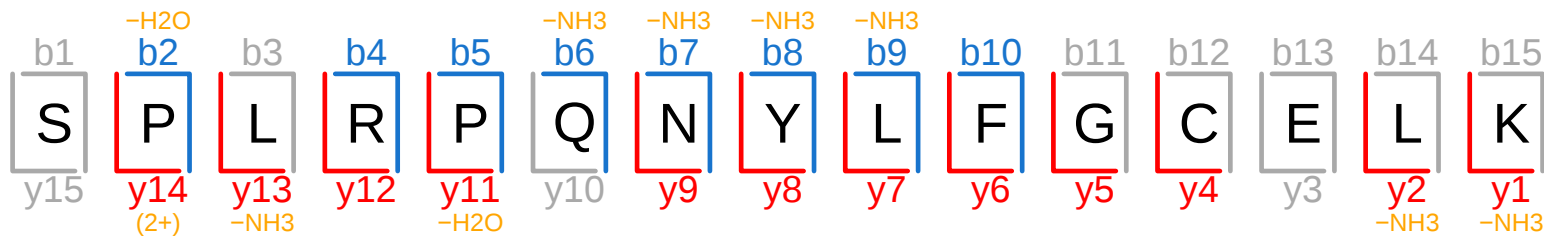


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

Scan Number: 60271

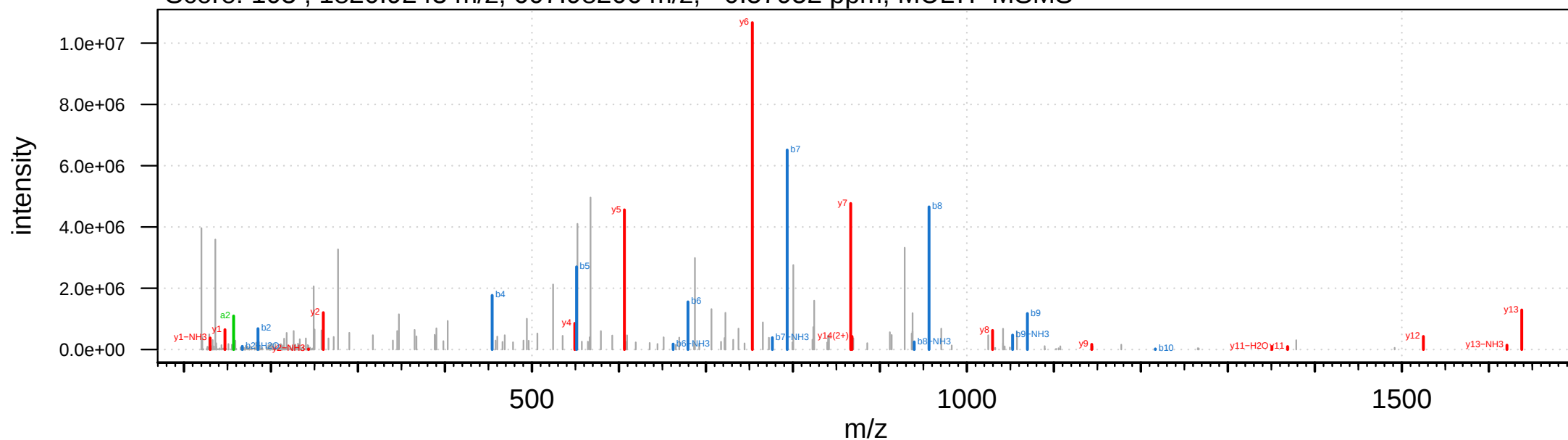
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 103 ; 1820.9243 m/z; 607.98206 m/z; -0.57952 ppm; MULTI-MSMS

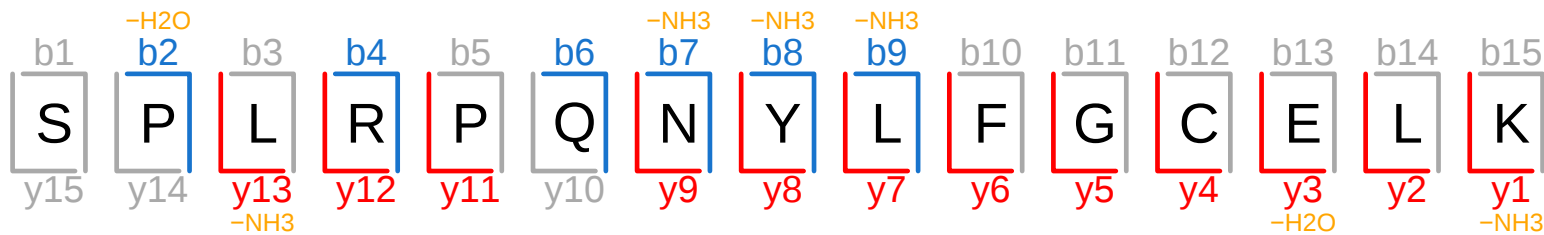


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

Scan Number: 60270

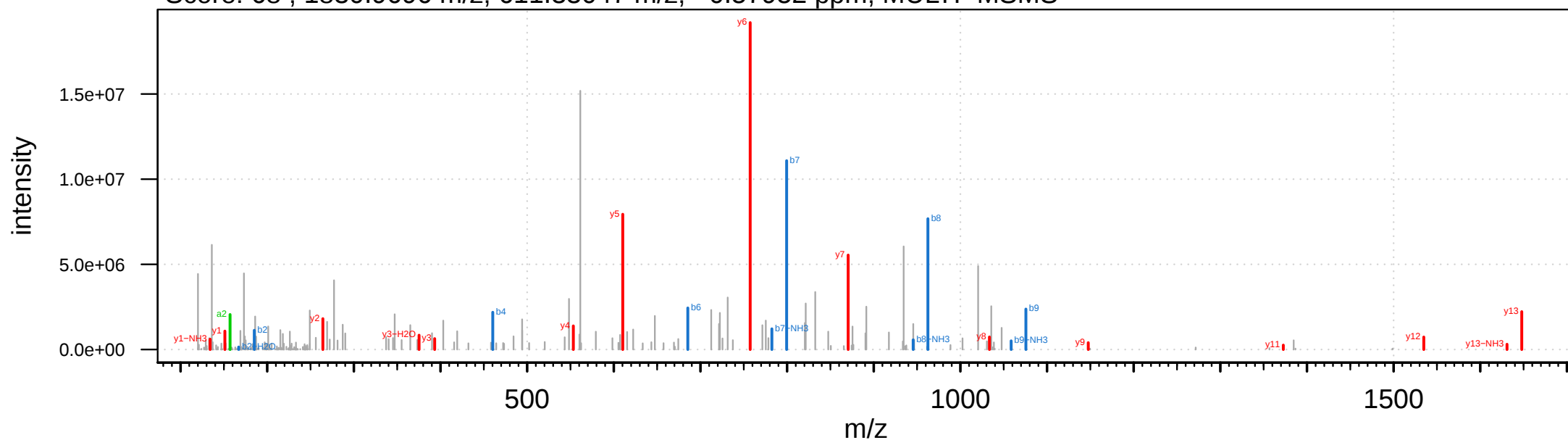
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 68 ; 1830.9696 m/z; 611.33047 m/z; -0.57952 ppm; MULTI-MSMS

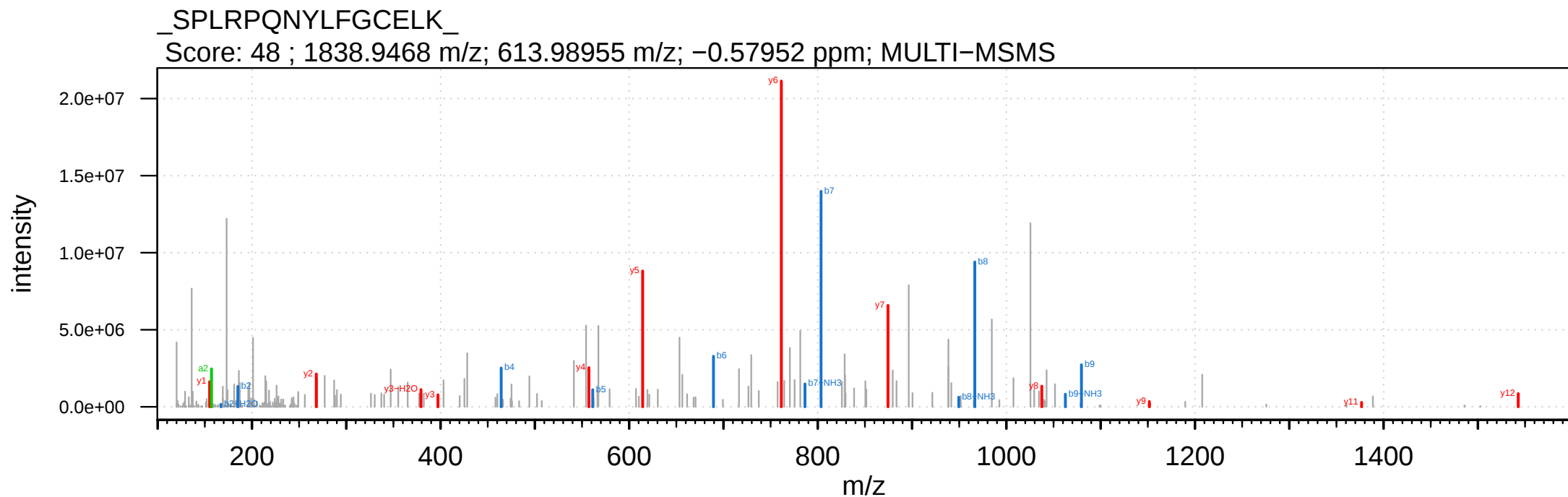
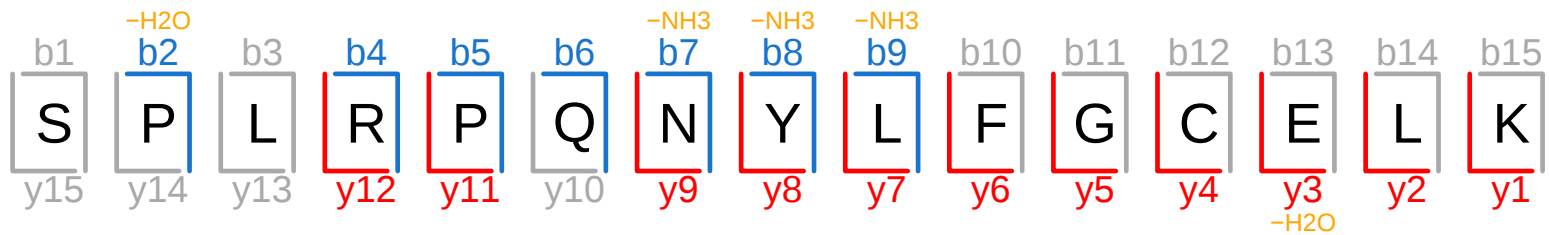


Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac

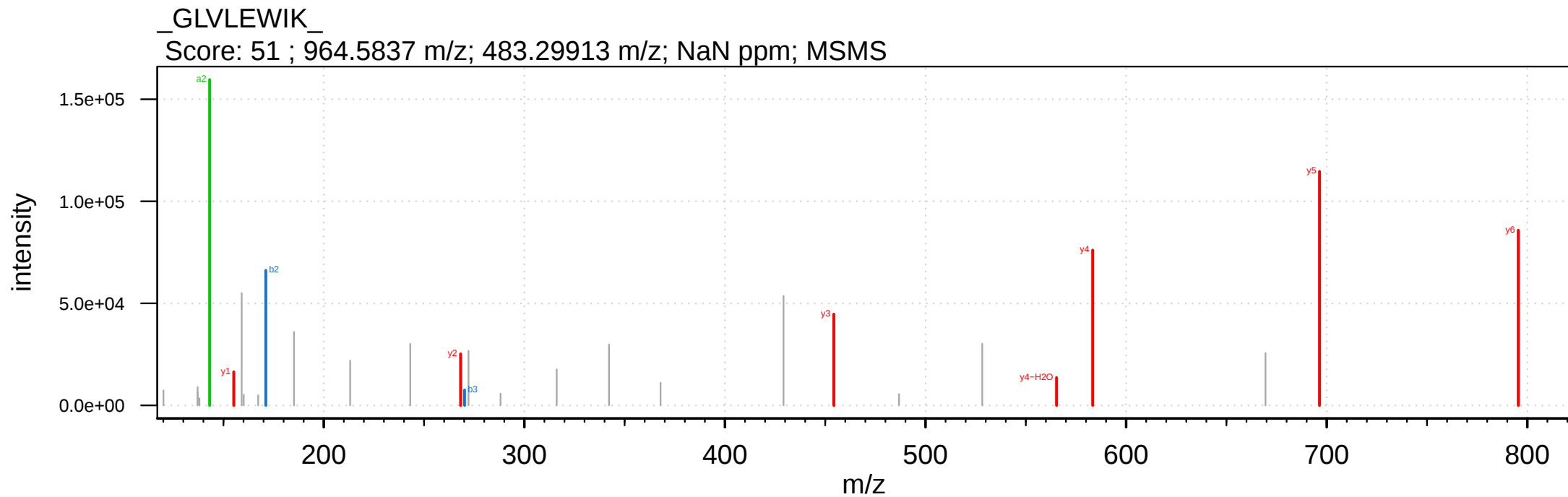
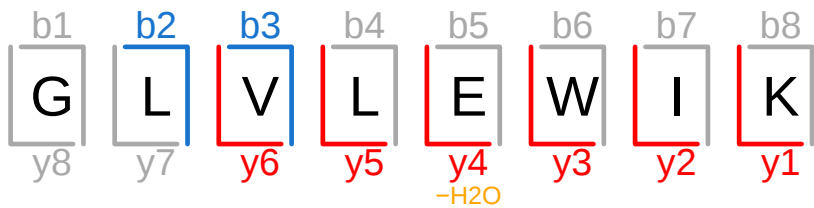
Scan Number: 60303

Proteins:

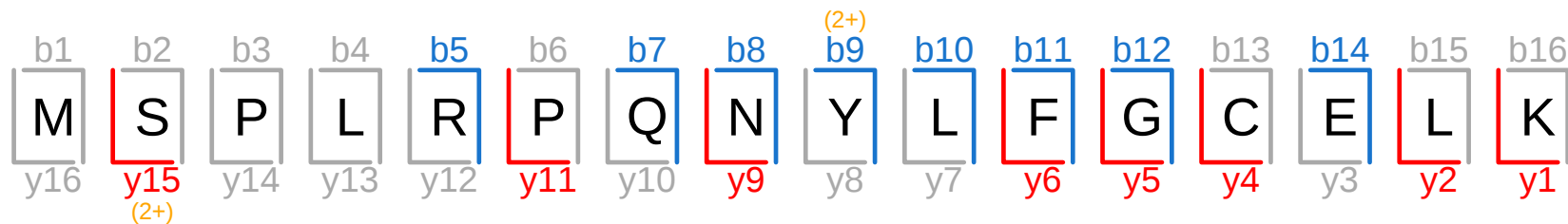
TCONS_I2_00008829_chr15:92829088-92829258:+



Raw File: Grobi_20140814_HZ_HS_A1L_MicroPeps_TriSilac
 Scan Number: 60304
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

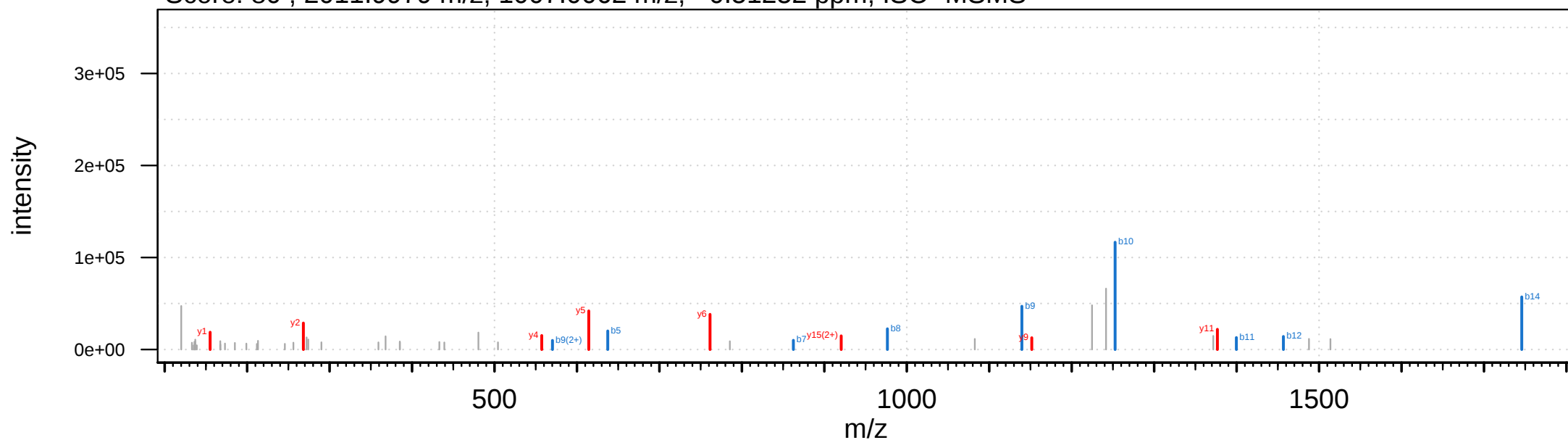


Raw File: Kermit_20140724_KK_HS_B1large
Scan Number: 39837
Proteins:
TCONS_I2_00001296_chr1:79520703-79520992:-



(ac)MSPLRPQNYLFGCELK

Score: 80 ; 2011.9979 m/z; 1007.0062 m/z; -0.51252 ppm; ISO-MSMS

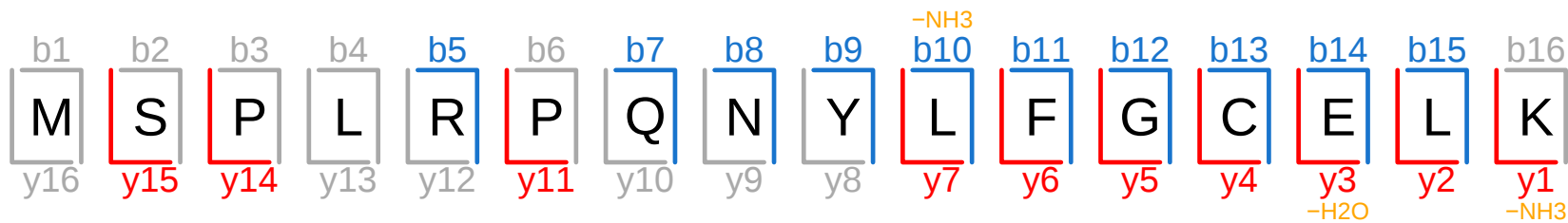


Raw File: Kermit_20140724_KK_HS_B1large

Scan Number: 45019

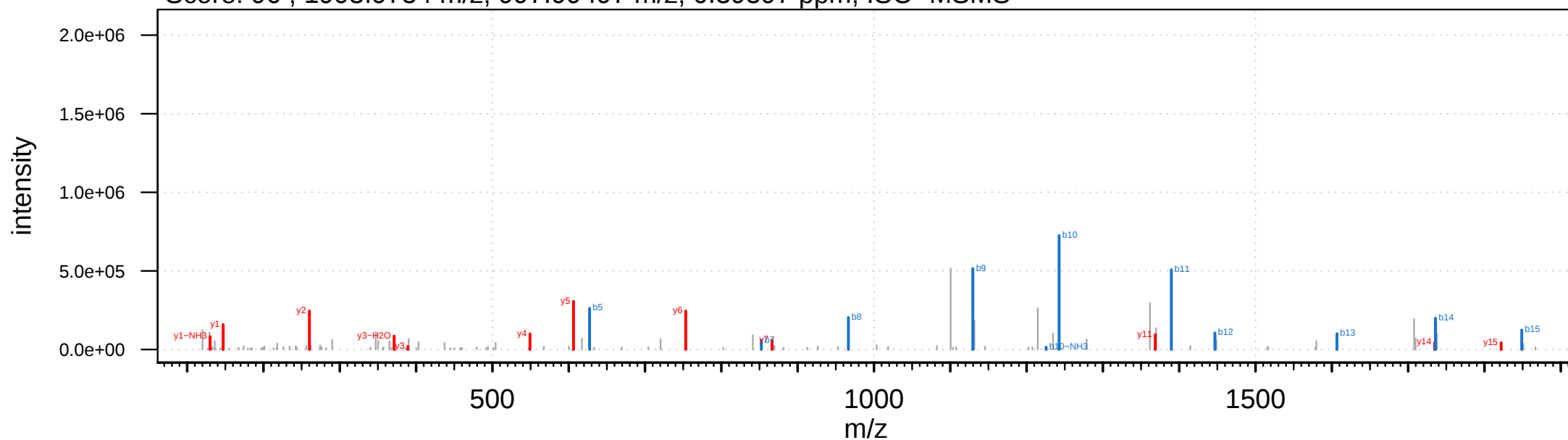
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 96 ; 1993.9754 m/z; 997.99497 m/z; 0.39597 ppm; ISO-MSMS

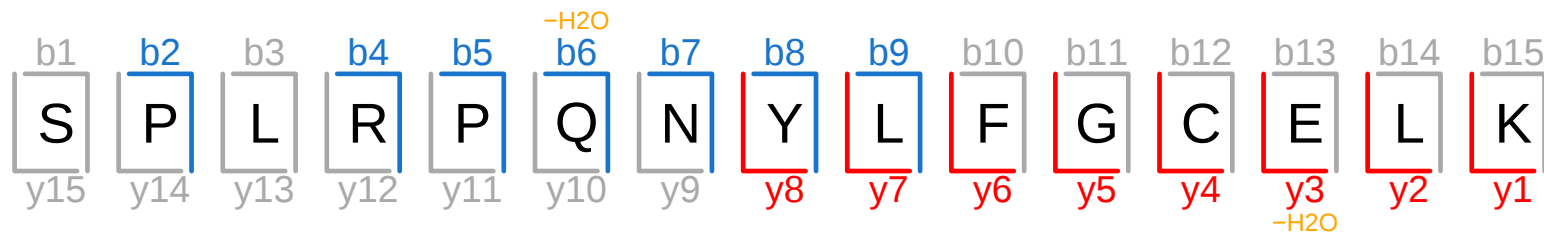


Raw File: Kermit_20140724_KK_HS_B1large

Scan Number: 45048

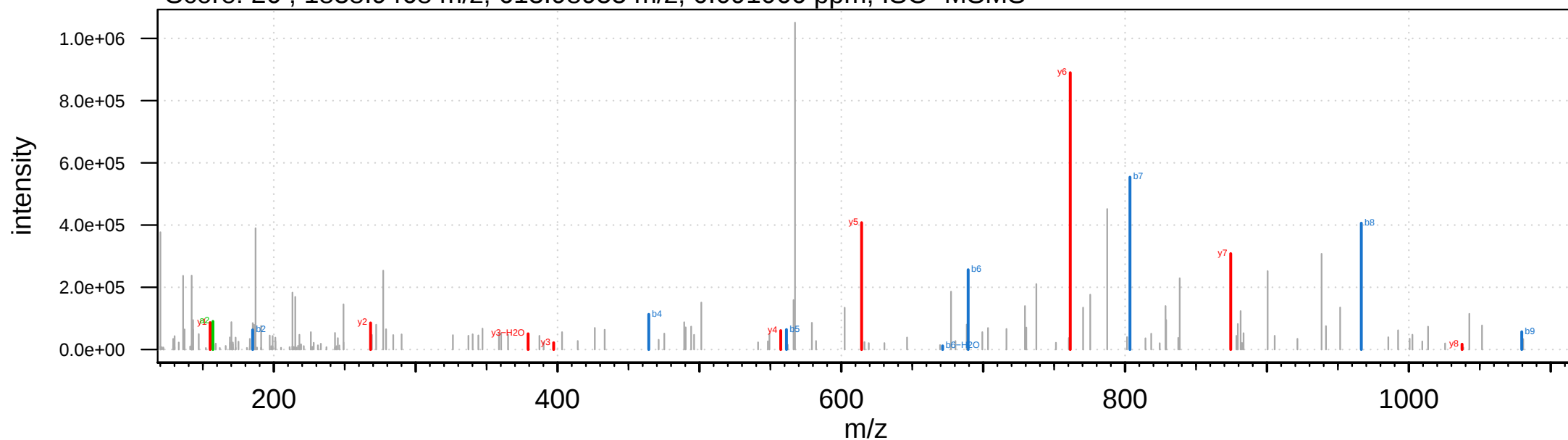
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 26 ; 1838.9468 m/z; 613.98955 m/z; 0.091066 ppm; ISO-MSMS

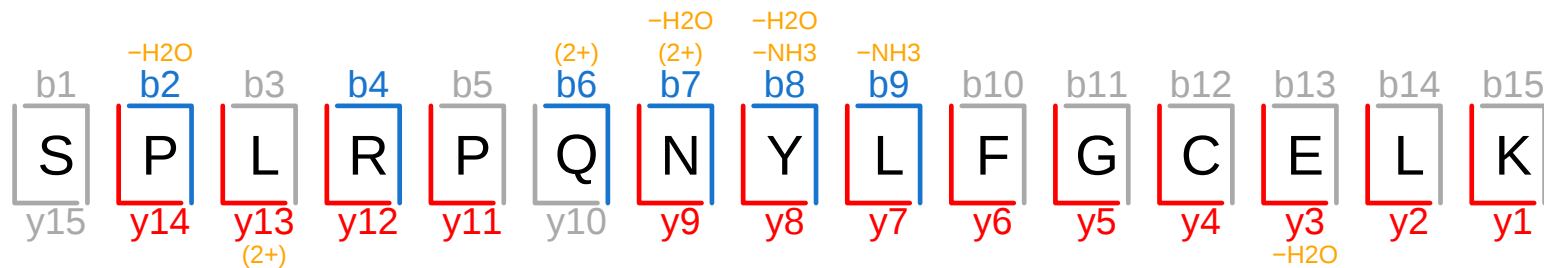


Raw File: Kermit_20140724_KK_HS_B1large

Scan Number: 30986

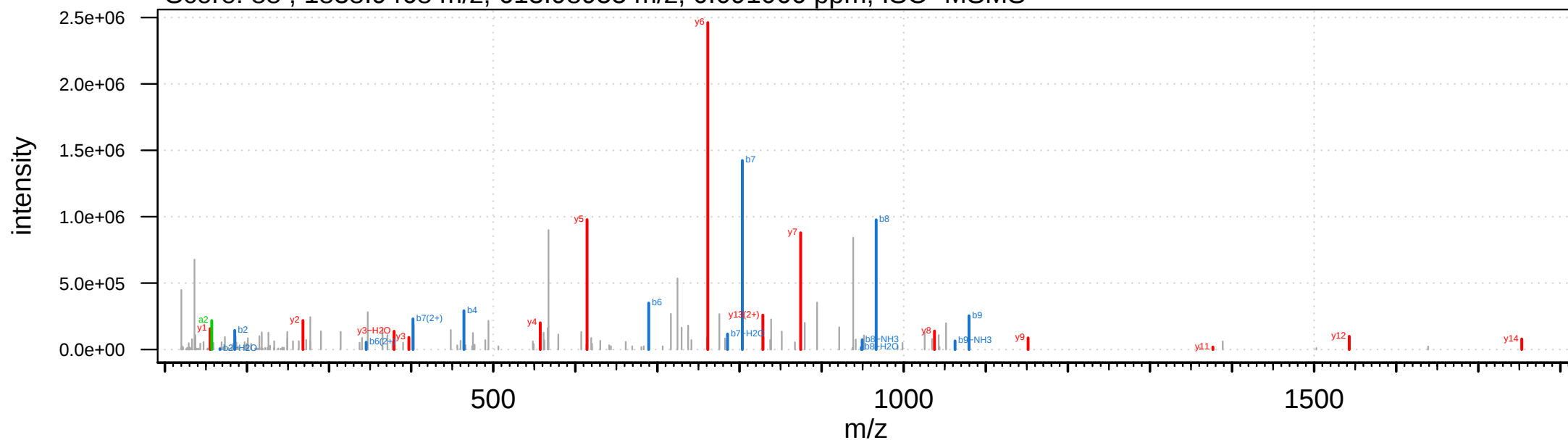
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 88 ; 1838.9468 m/z; 613.98955 m/z; 0.091066 ppm; ISO-MSMS

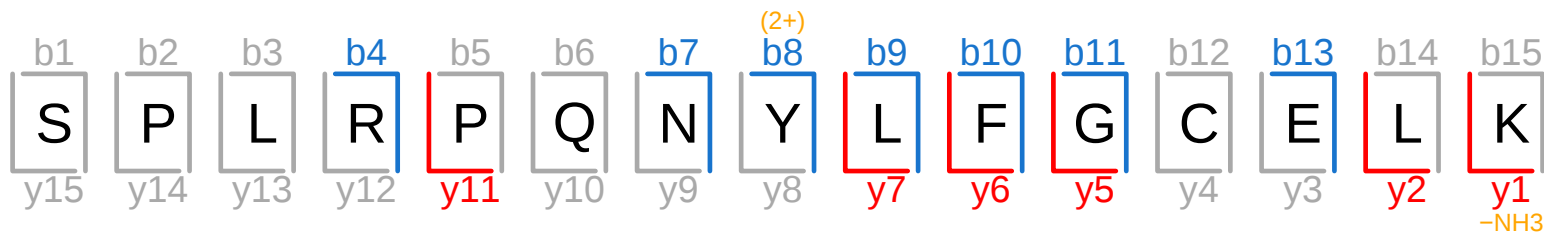


Raw File: Kermit_20140724_KK_HS_B1large

Scan Number: 31258

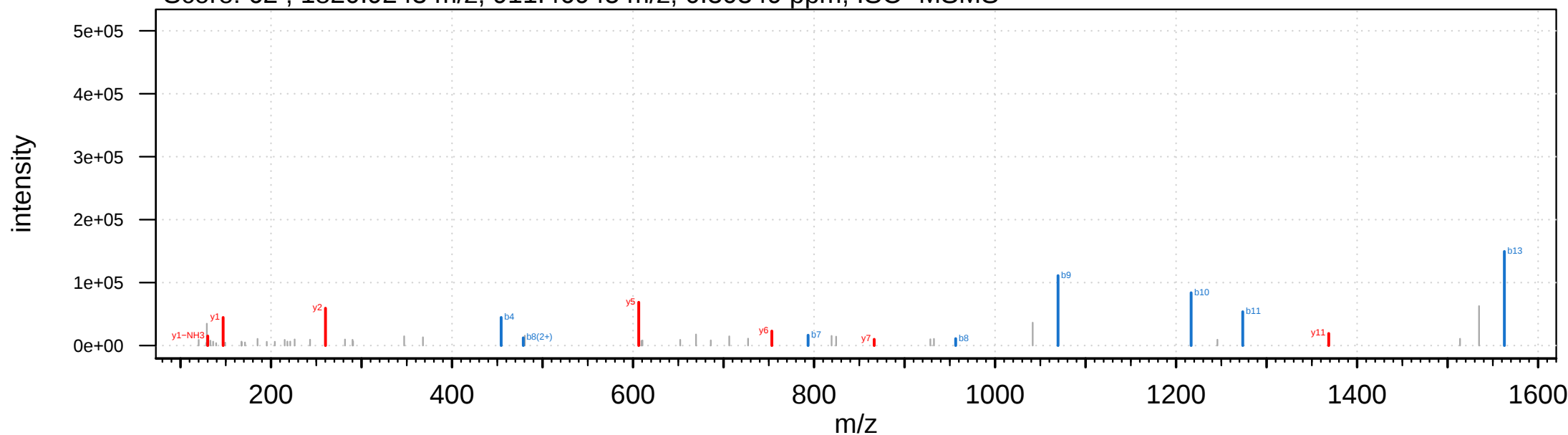
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 62 ; 1820.9243 m/z; 911.46945 m/z; 0.30349 ppm; ISO-MSMS

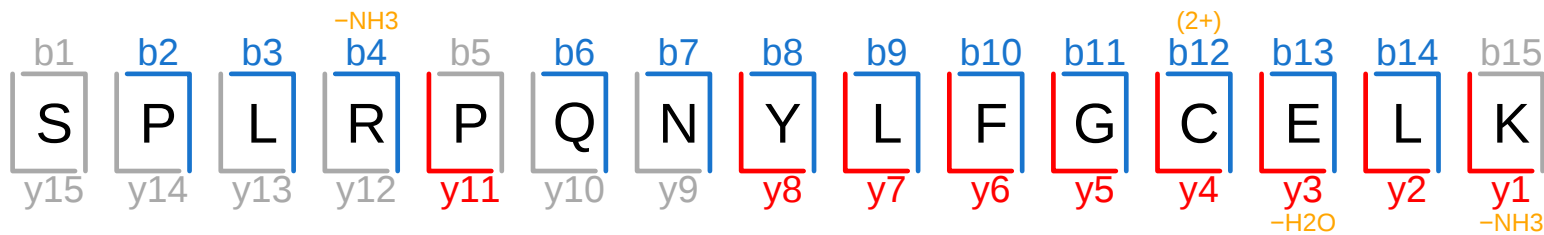


Raw File: Kermit_20140724_KK_HS_B1large

Scan Number: 31007

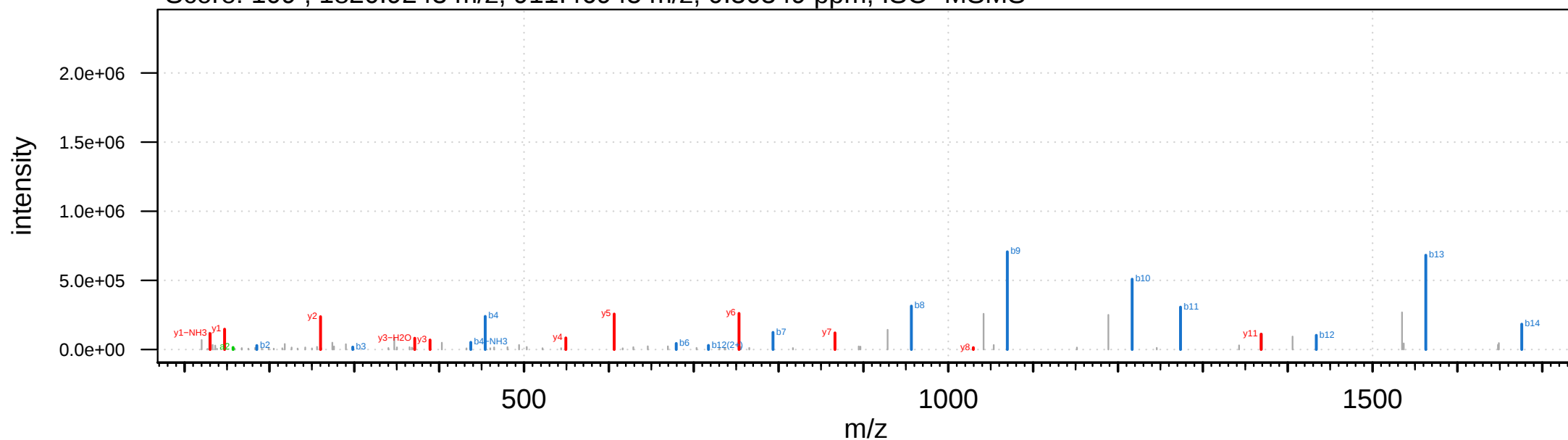
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 109 ; 1820.9243 m/z; 911.46945 m/z; 0.30349 ppm; ISO-MSMS

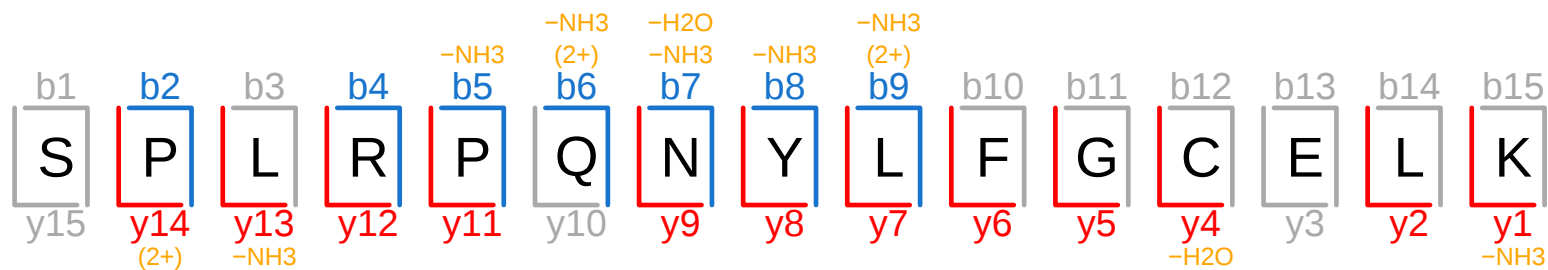


Raw File: Kermit_20140724_KK_HS_B1large

Scan Number: 31280

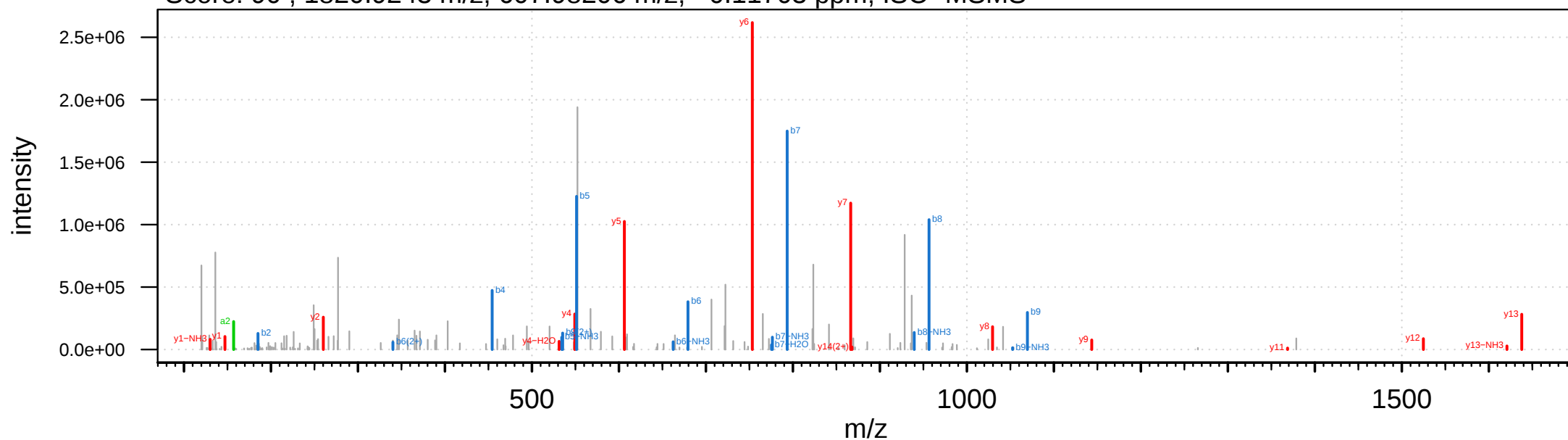
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 90 ; 1820.9243 m/z; 607.98206 m/z; -0.11703 ppm; ISO-MSMS

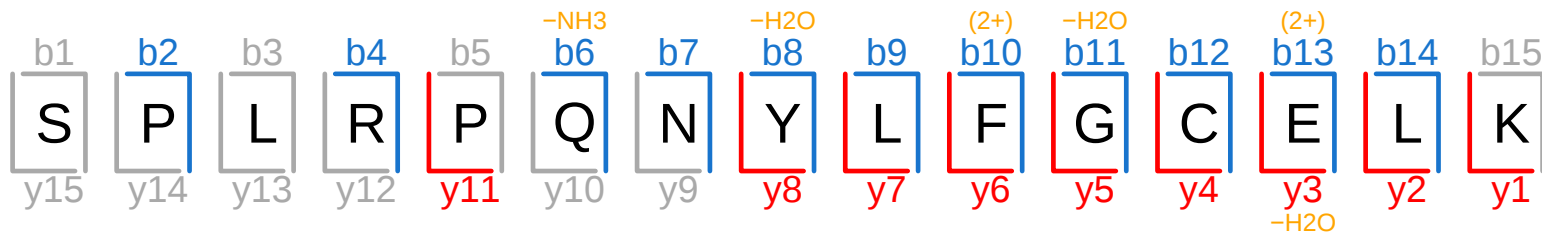


Raw File: Kermit_20140724_KK_HS_B1large

Scan Number: 31096

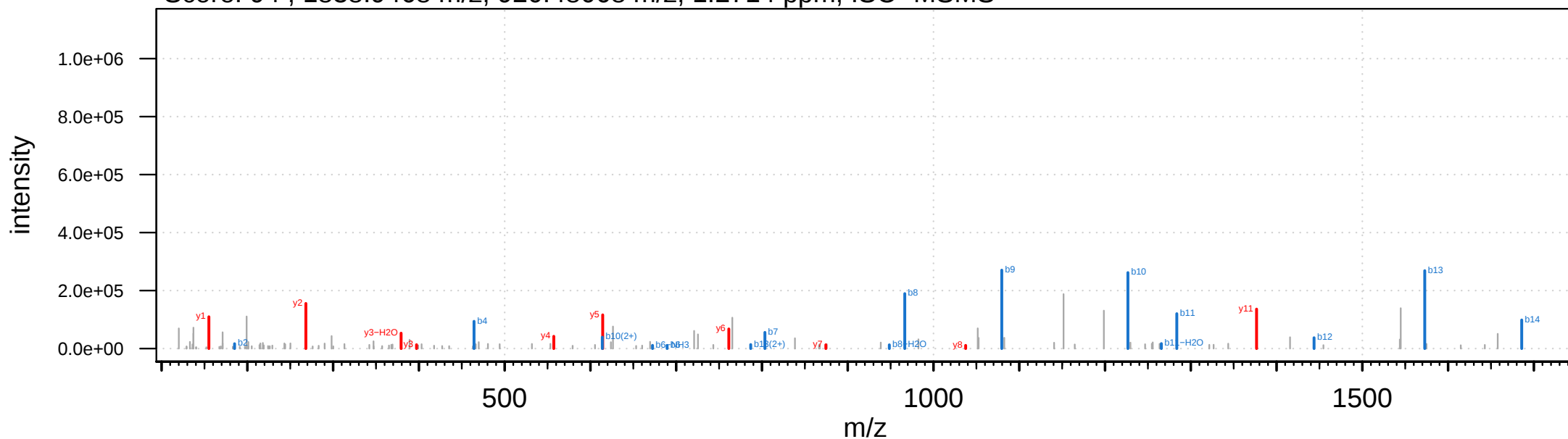
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 94 ; 1838.9468 m/z; 920.48068 m/z; 1.2714 ppm; ISO-MSMS

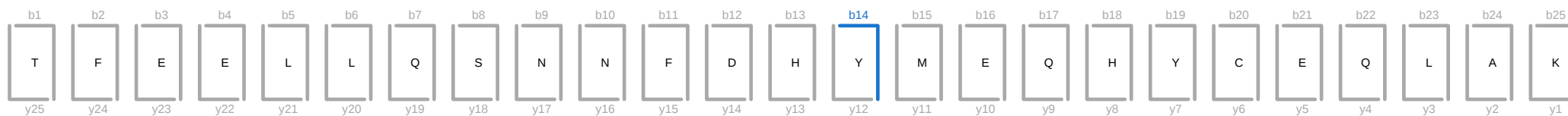


Raw File: Kermit_20140724_KK_HS_B1large

Scan Number: 31270

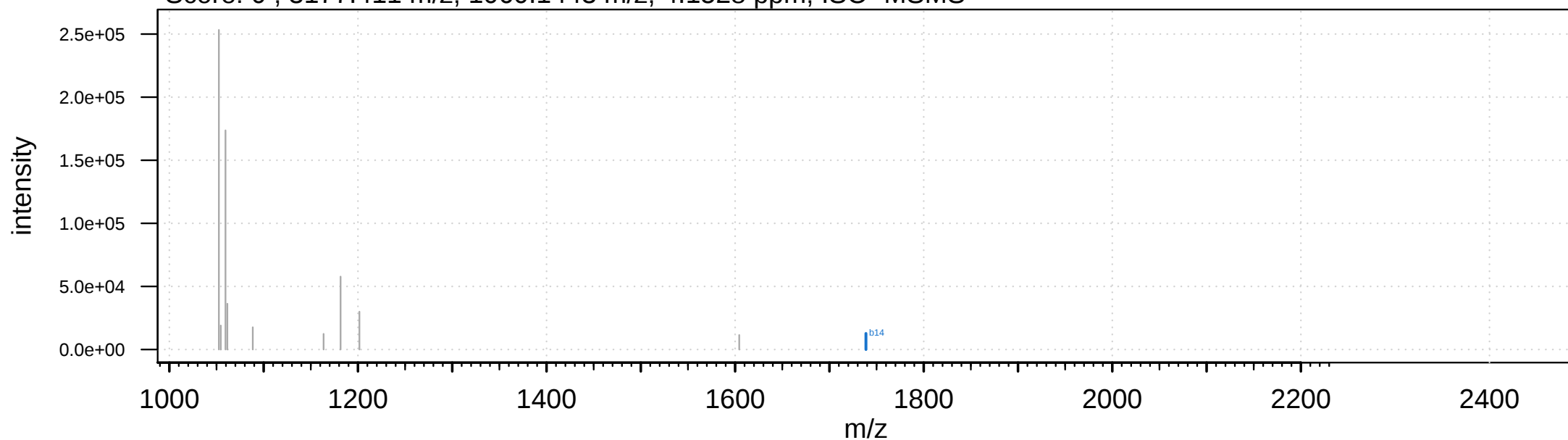
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



TFEELLQSNNFDHYMEQHYCEQLAK

Score: 0 ; 3177.411 m/z; 1060.1443 m/z; 4.1528 ppm; ISO-MSMS

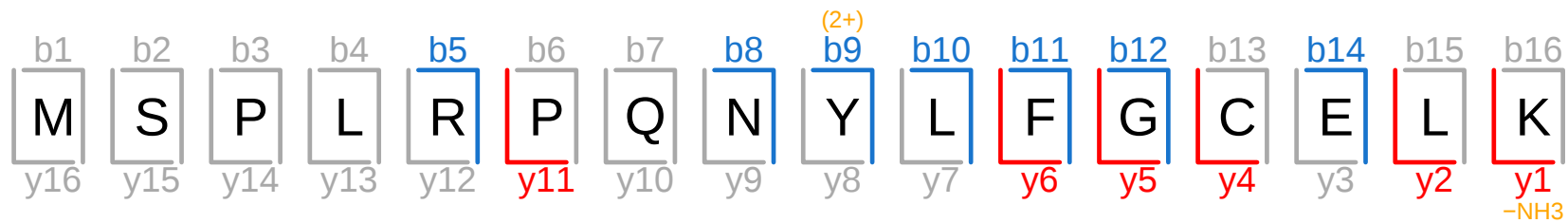


Raw File: Kermit_20140724_KK_HS_B1large

Scan Number: 19842

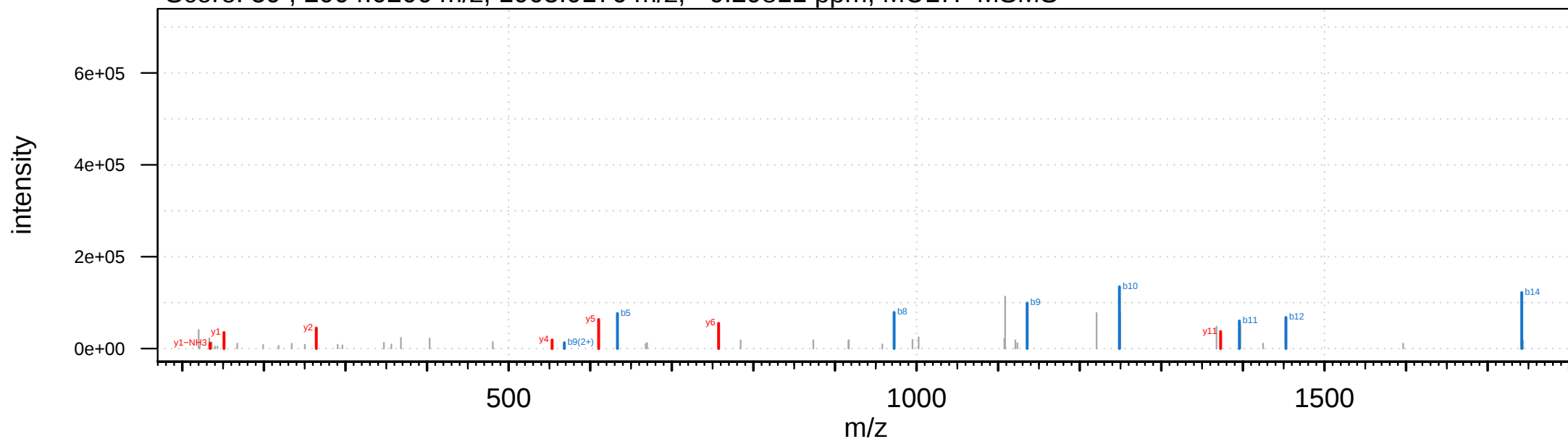
Proteins:

ENST00000608683_chr1:149649167-149701093:-



(ac)MSPLRPQNYLFGCELK

Score: 59 ; 2004.0206 m/z; 1003.0176 m/z; -0.29811 ppm; MULTI-MSMS

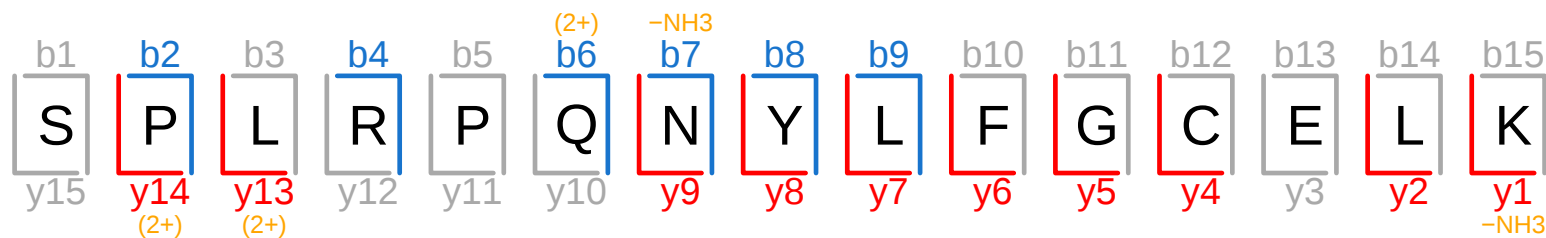


Raw File: Kermit_20140724_KK_HS_A1large

Scan Number: 48480

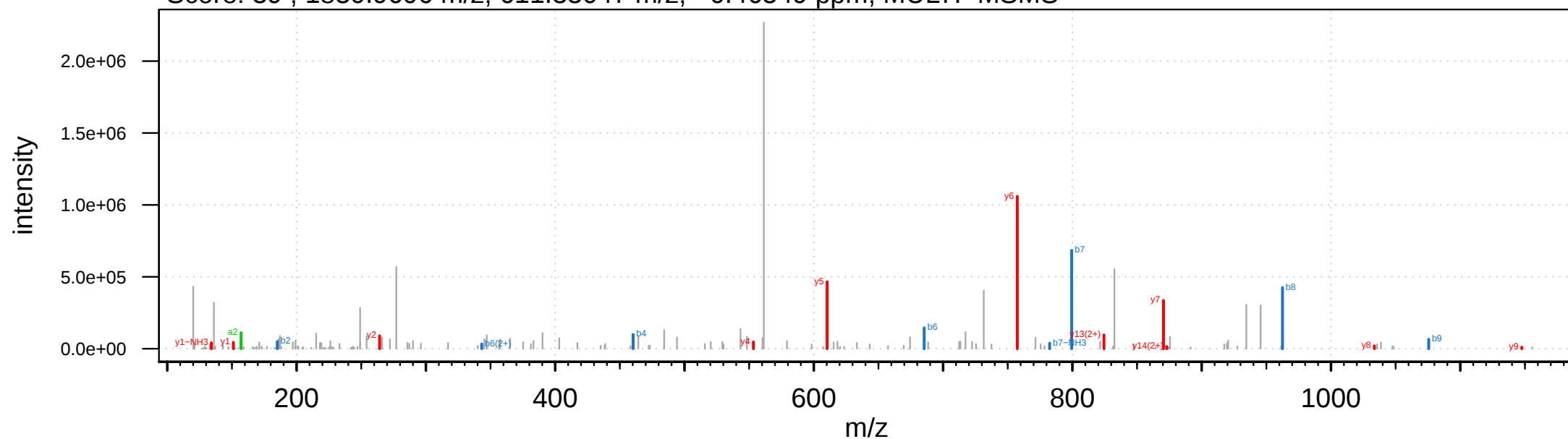
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 39 ; 1830.9696 m/z; 611.33047 m/z; -0.46549 ppm; MULTI-MSMS

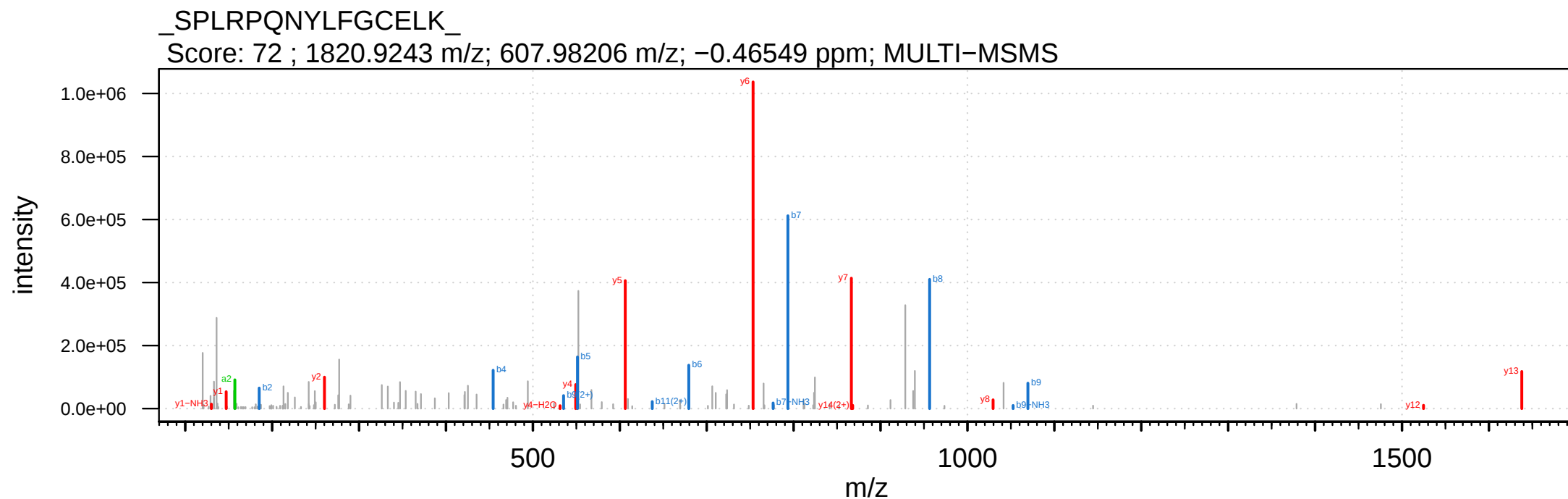
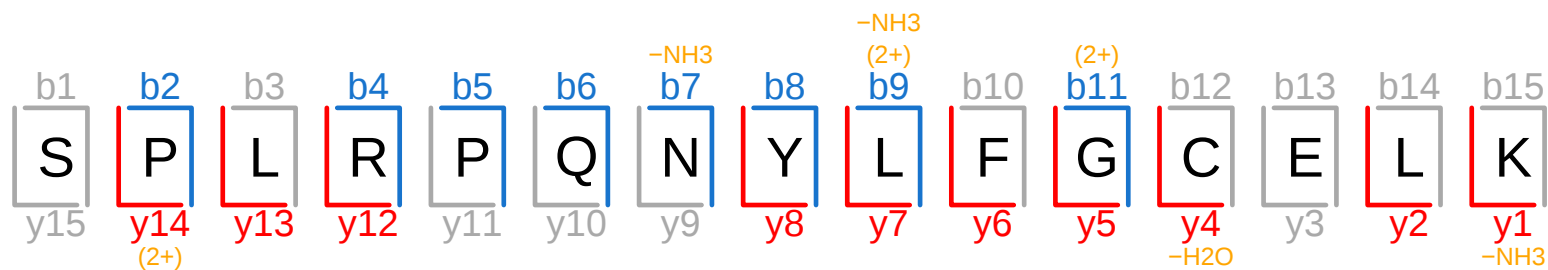


Raw File: Kermit_20140724_KK_HS_A1large

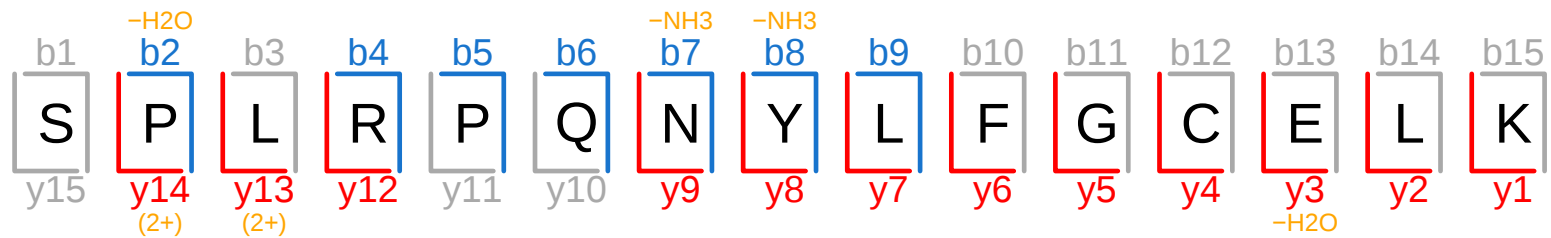
Scan Number: 33272

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

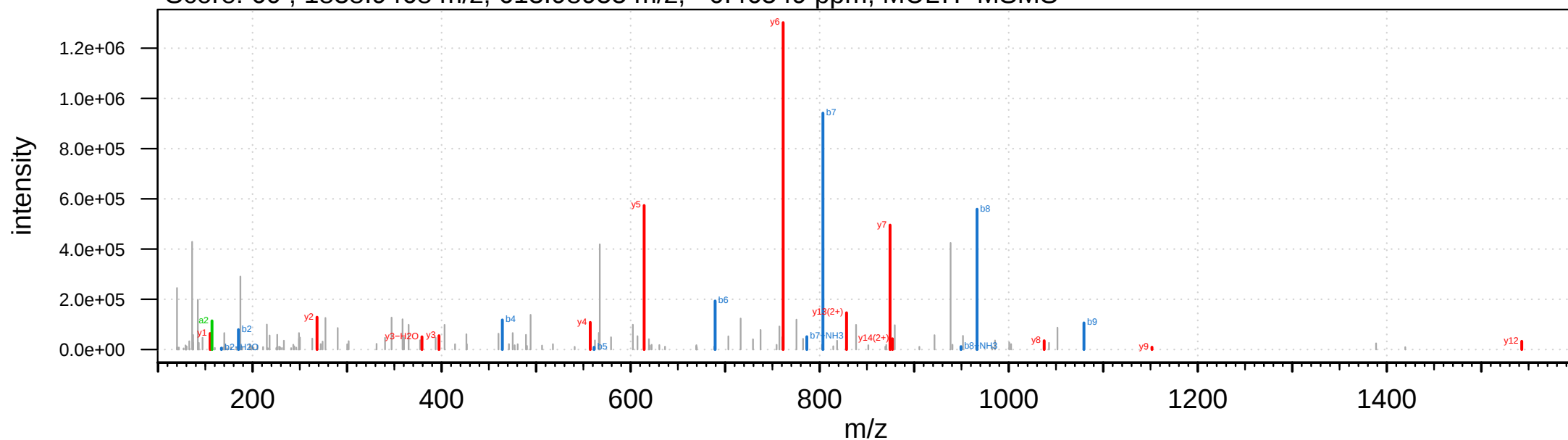


Raw File: Kermit_20140724_KK_HS_A1large
 Scan Number: 33303
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 60 ; 1838.9468 m/z; 613.98955 m/z; -0.46549 ppm; MULTI-MSMS

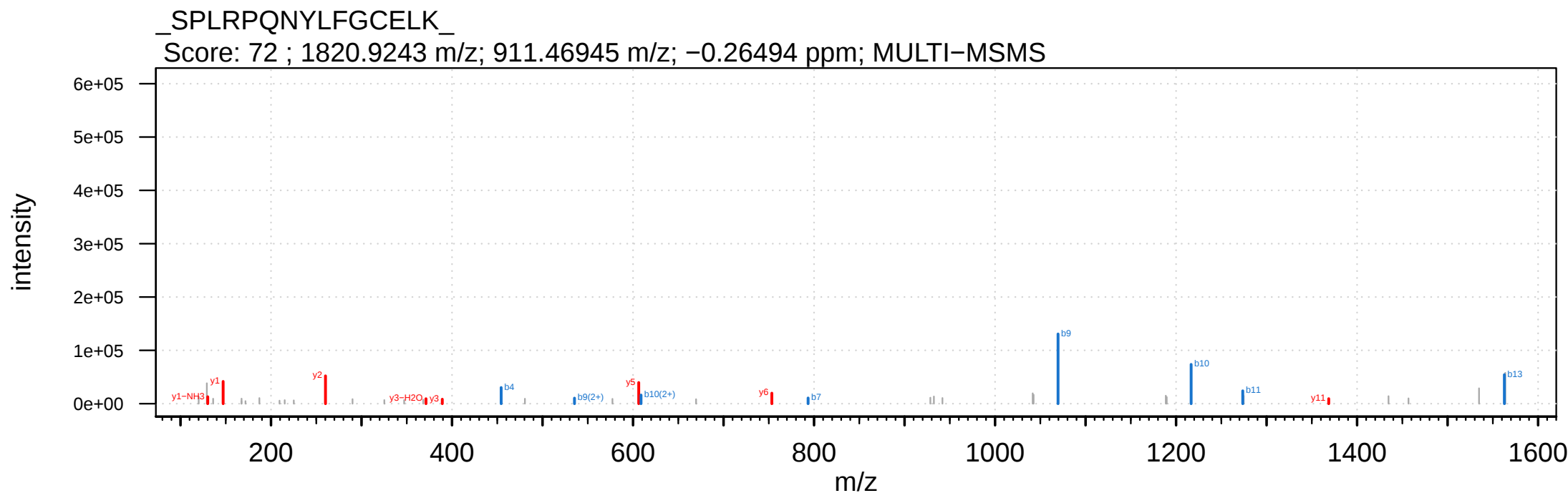
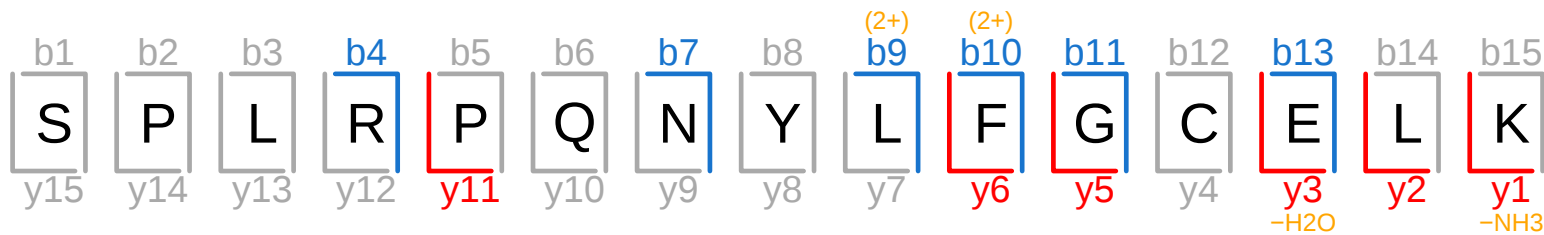


Raw File: Kermit_20140724_KK_HS_A1large

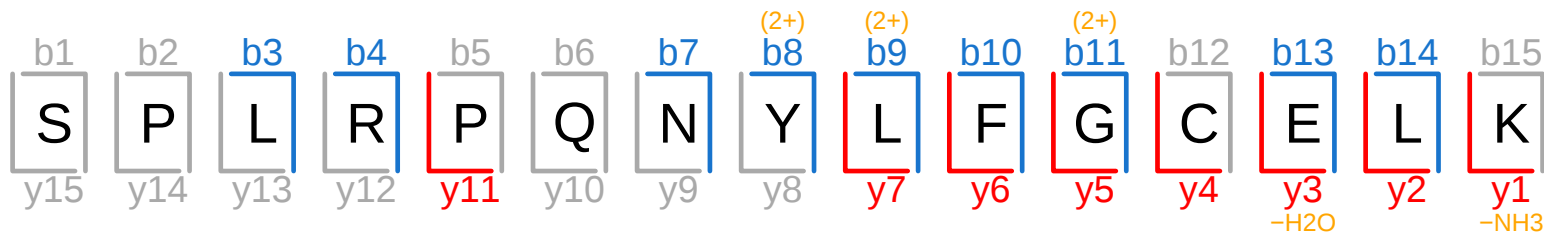
Scan Number: 33314

Proteins:

TCONS_I2_00008829_ch15:92829088-92829258:+

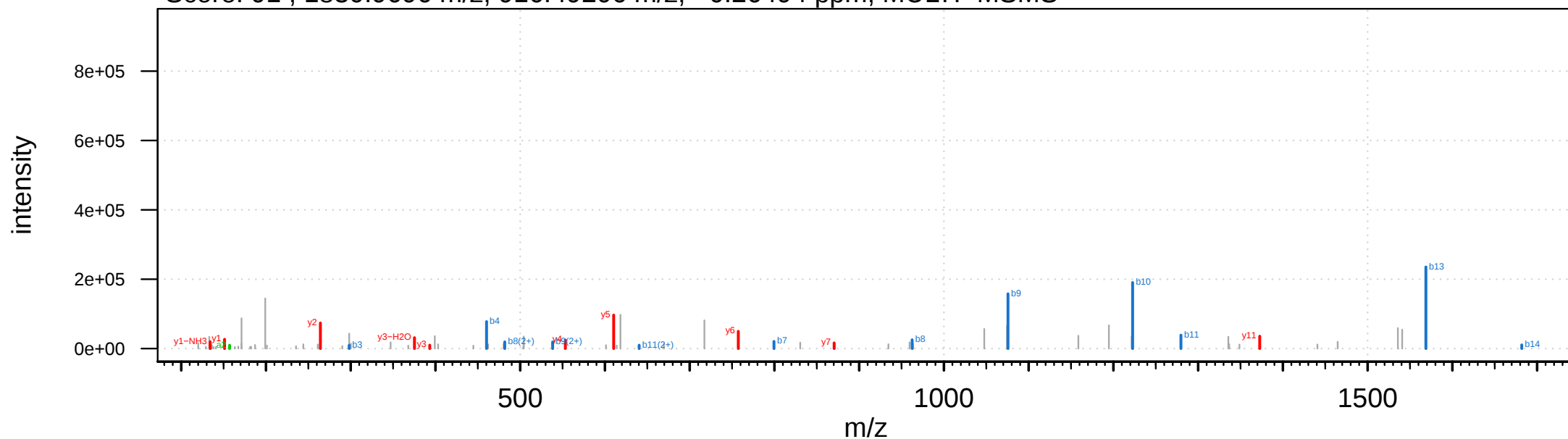


Raw File: Kermit_20140724_KK_HS_A1large
 Scan Number: 33286
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 91 ; 1830.9696 m/z; 916.49206 m/z; -0.26494 ppm; MULTI-MSMS

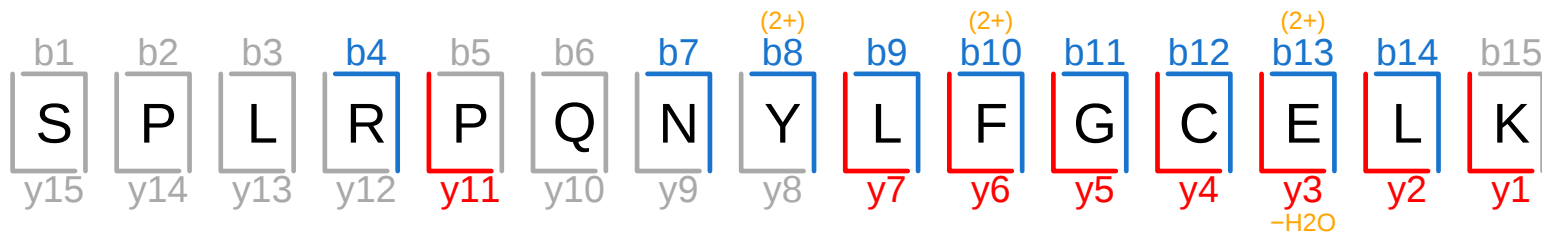


Raw File: Kermit_20140724_KK_HS_A1large

Scan Number: 33316

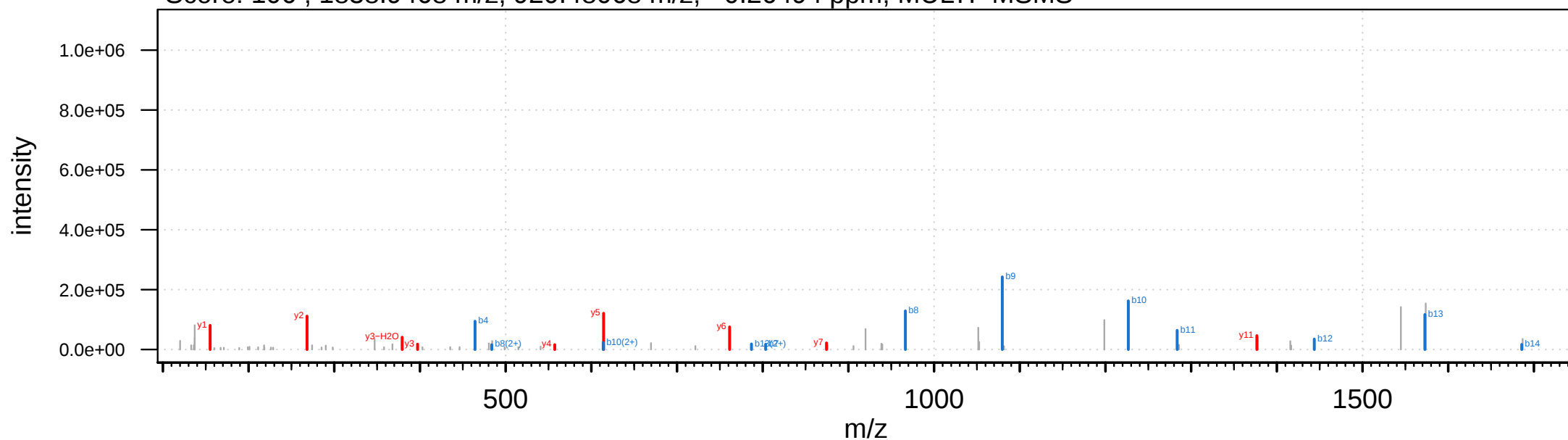
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 106 ; 1838.9468 m/z; 920.48068 m/z; -0.26494 ppm; MULTI-MSMS

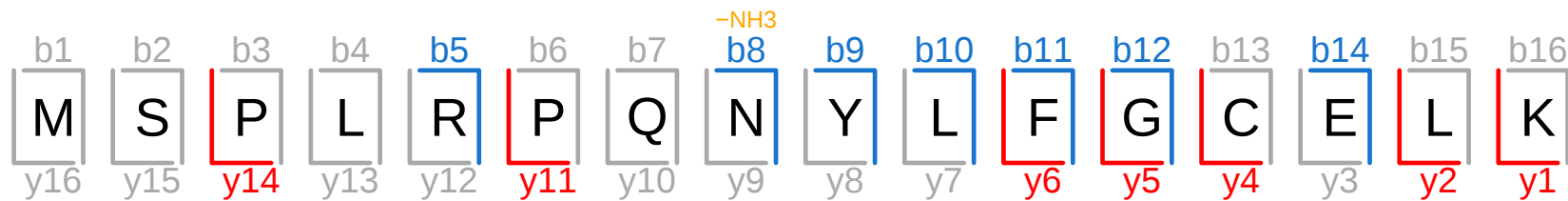


Raw File: Kermit_20140724_KK_HS_A1large

Scan Number: 33392

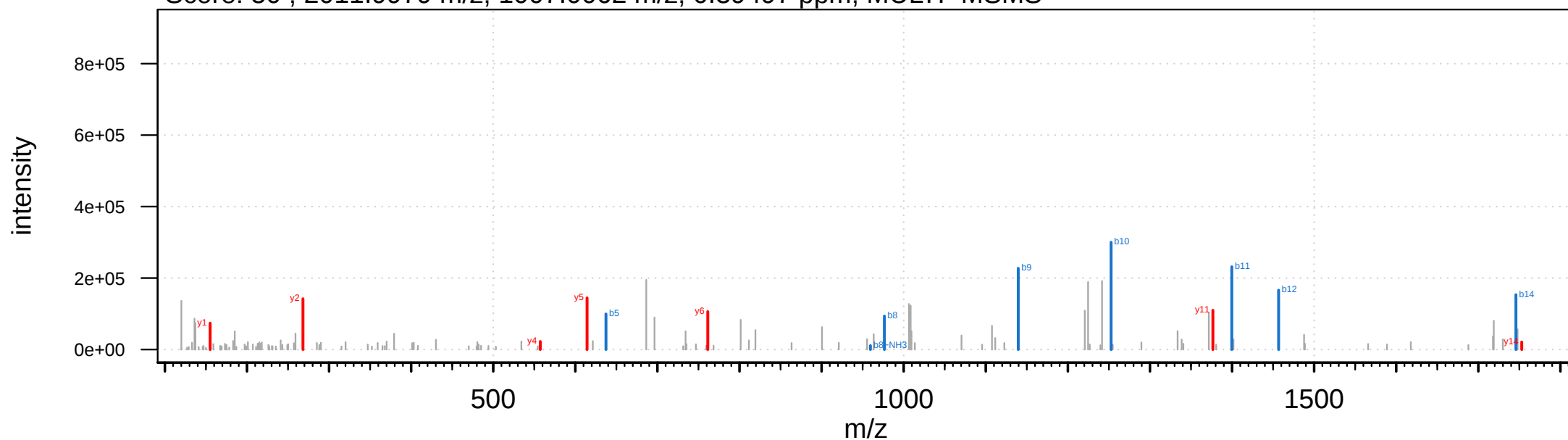
Proteins:

TCONS_I2_00008829_ch15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 50 ; 2011.9979 m/z; 1007.0062 m/z; 0.39497 ppm; MULTI-MSMS

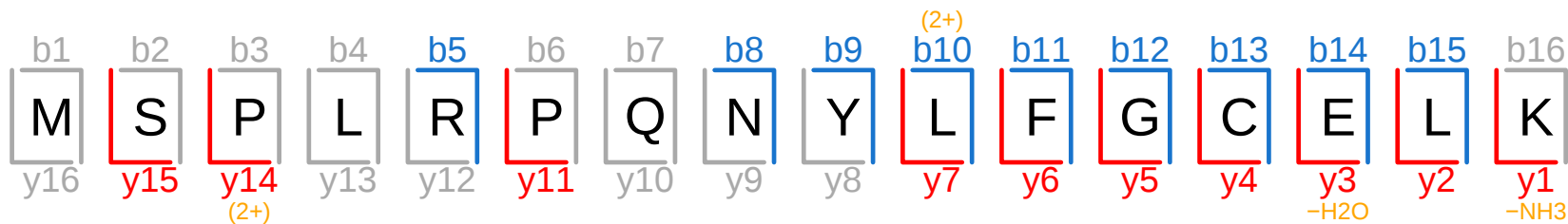


Raw File: Kermit_20140724_KK_HS_large

Scan Number: 47640

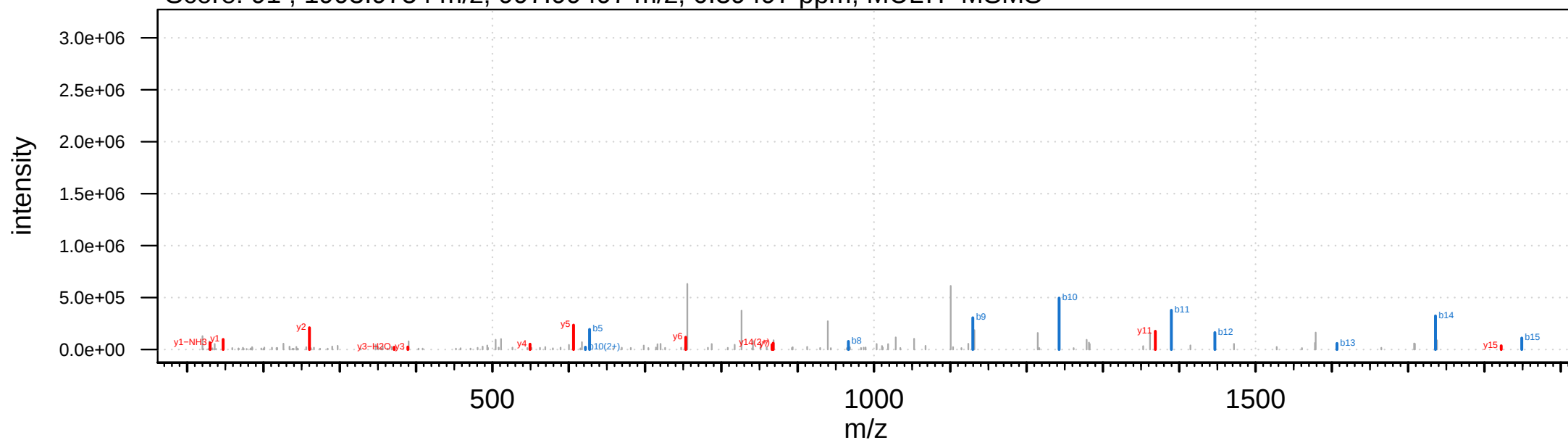
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 91 ; 1993.9754 m/z; 997.99497 m/z; 0.39497 ppm; MULTI-MSMS

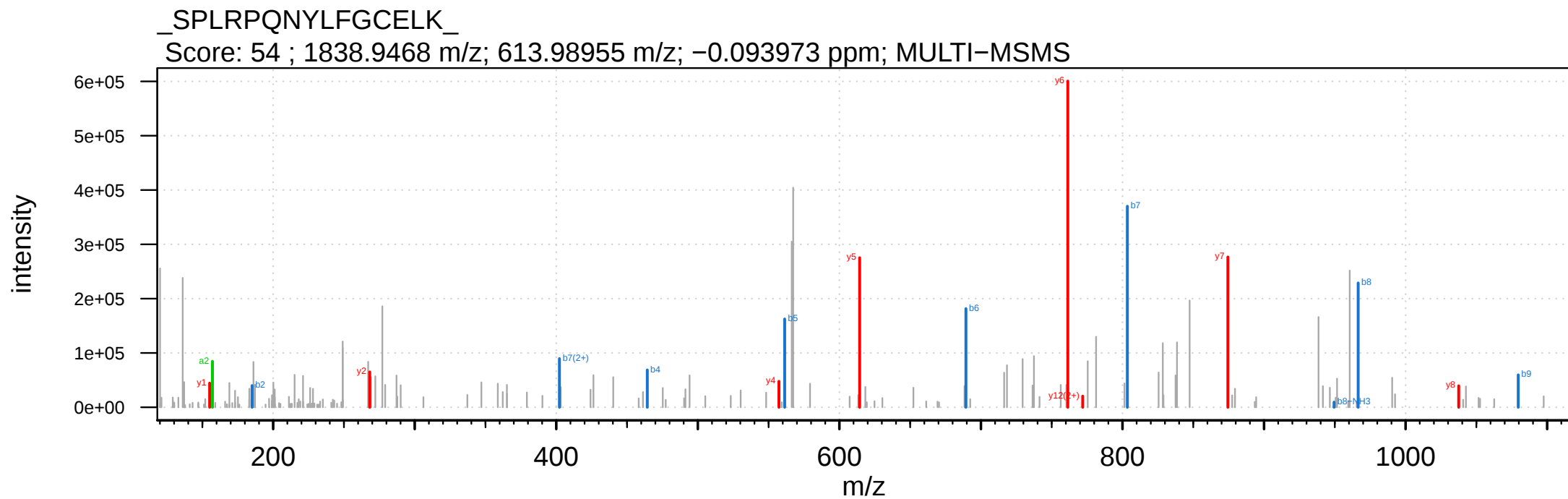
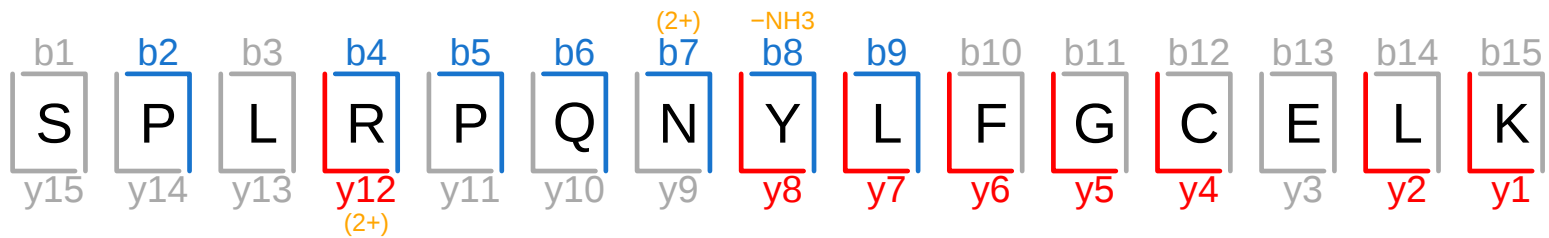


Raw File: Kermit_20140724_KK_HS_large

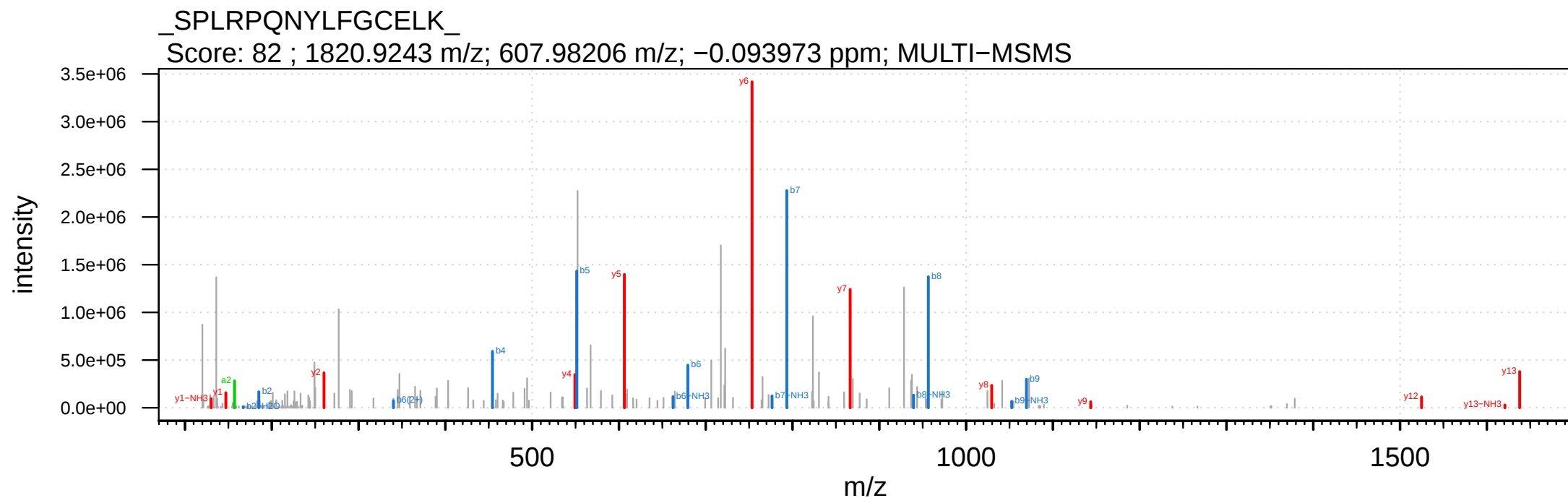
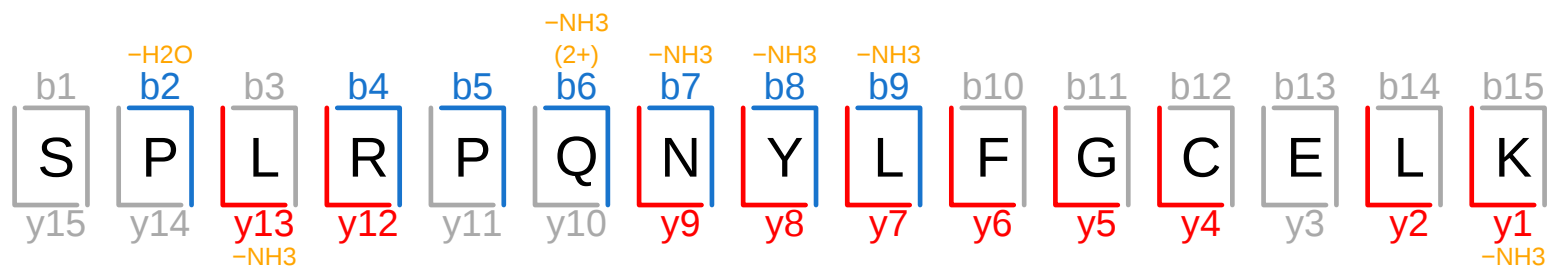
Scan Number: 47656

Proteins:

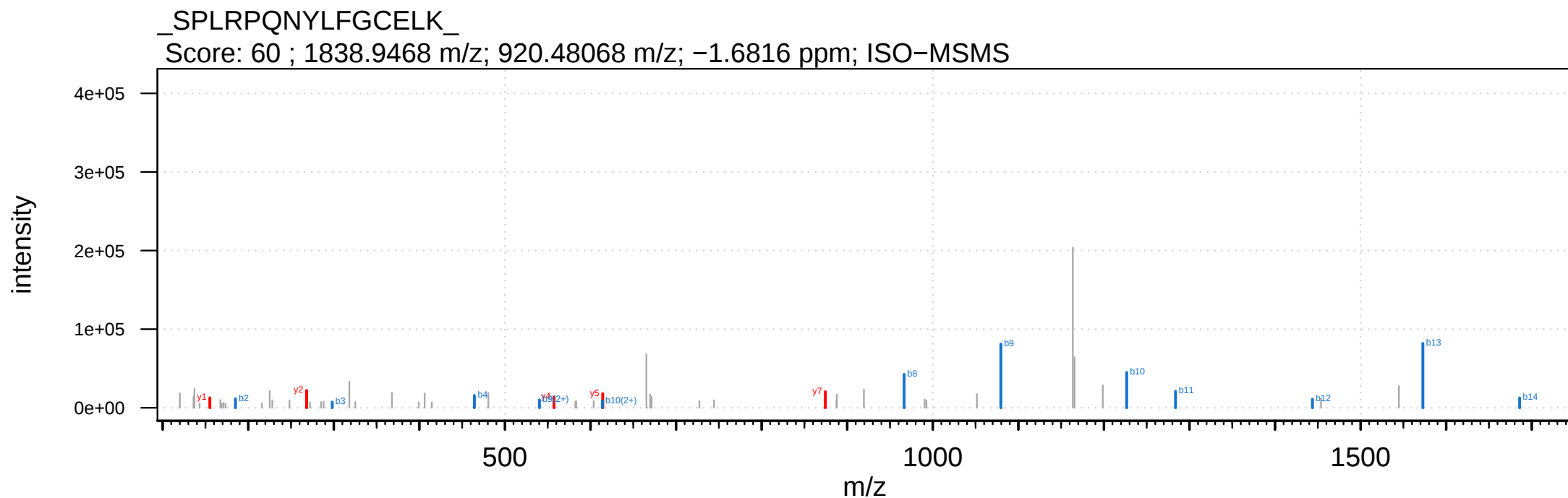
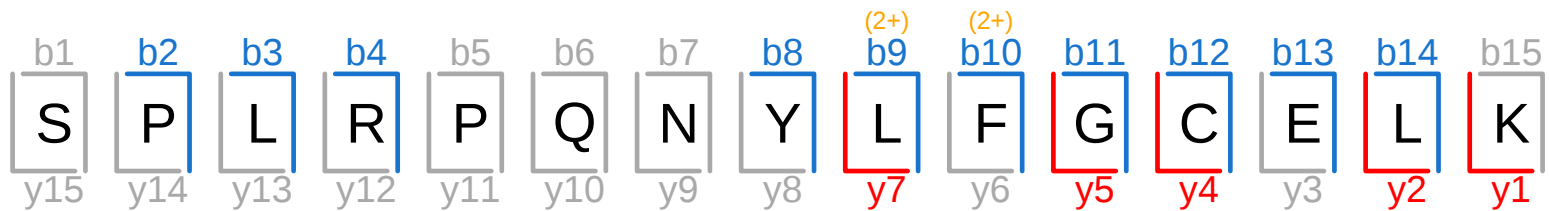
TCONS_I2_00008829_chr15:92829088-92829258:+



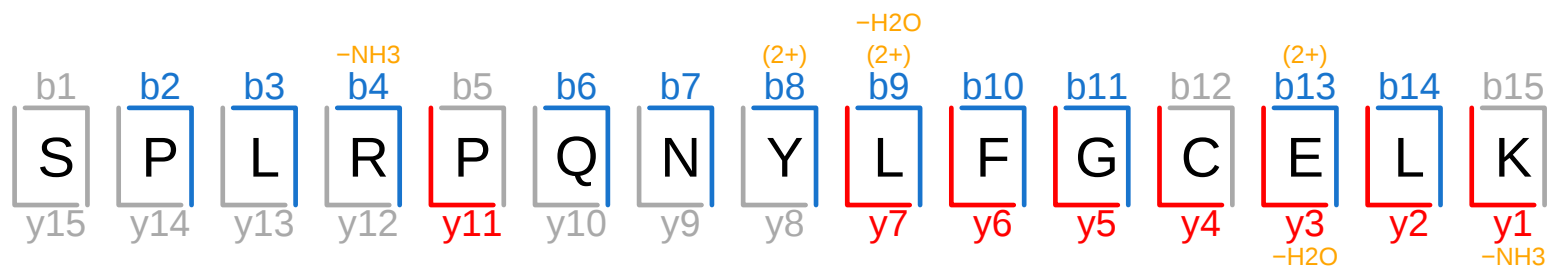
Raw File: Kermit_20140724_KK_HS_large
 Scan Number: 31899
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



Raw File: Kermit_20140724_KK_HS_large
 Scan Number: 32052
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

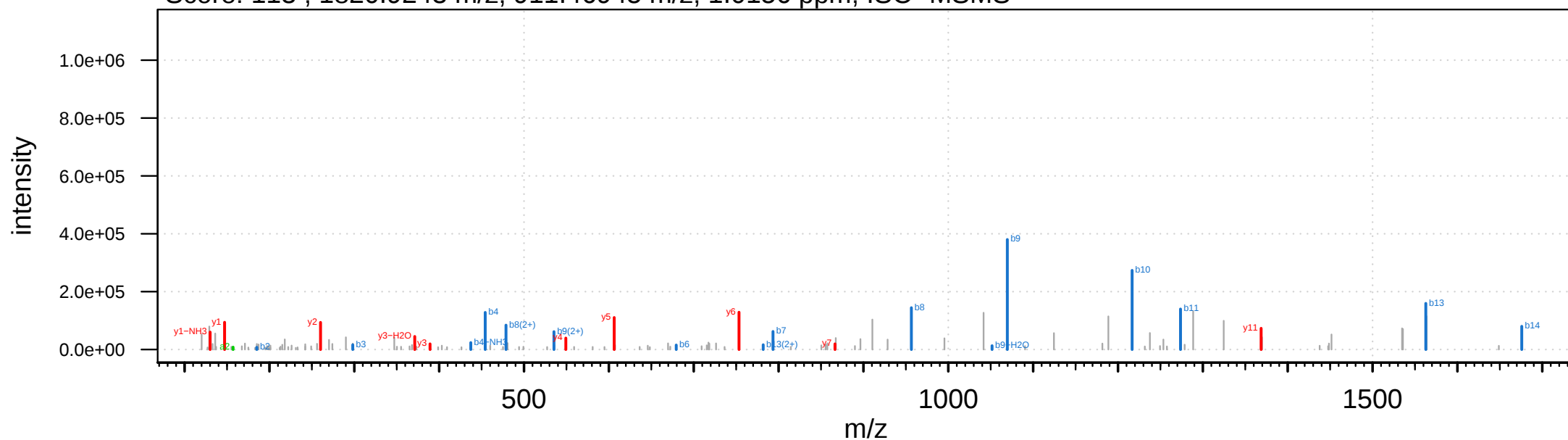


Raw File: Kermit_20140724_KK_HS_large
Scan Number: 31859
Proteins:
TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 113 ; 1820.9243 m/z; 911.46945 m/z; 1.0156 ppm; ISO-MSMS

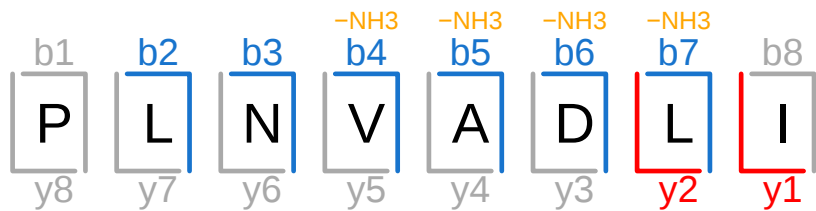


Raw File: Kermit_20140724_KK_HS_large

Scan Number: 31900

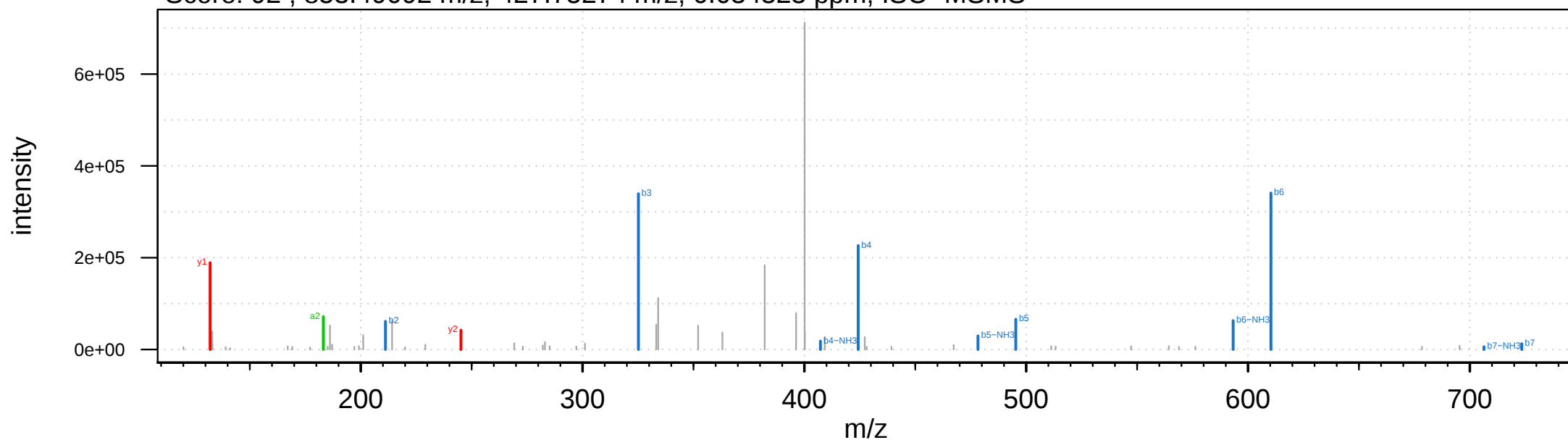
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



PLNVADLI

Score: 92 ; 853.49092 m/z; 427.75274 m/z; 0.034323 ppm; ISO-MSMS

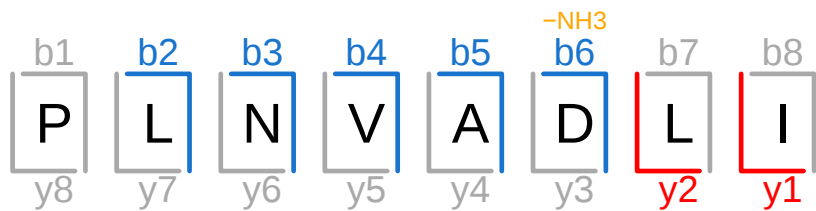


Raw File: Grobi_20140814_HZ_HS_A1S_MicroPeps_TriSilac

Scan Number: 75836

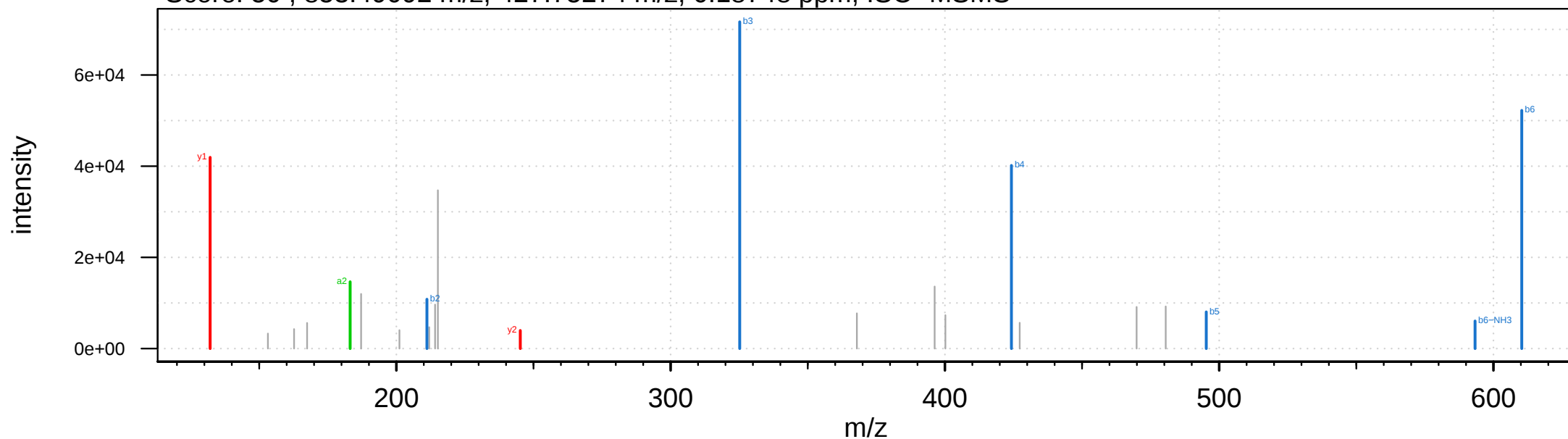
Proteins:

TCONS_I2_00005685_chr12:53548245-53548274:+



PLNVADLI

Score: 50 ; 853.49092 m/z; 427.75274 m/z; 0.18748 ppm; ISO-MSMS

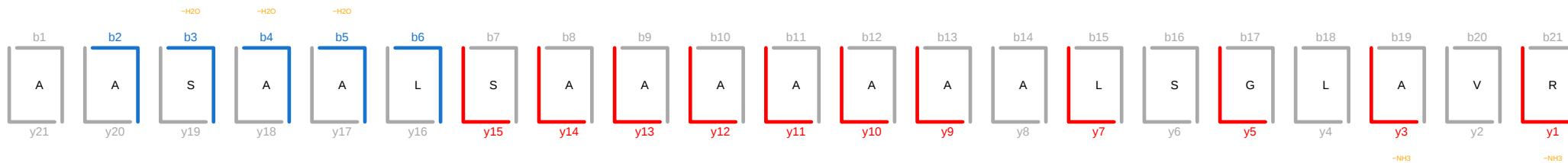


Raw File: Kermit_20140724_KK_HS_B1small

Scan Number: 15593

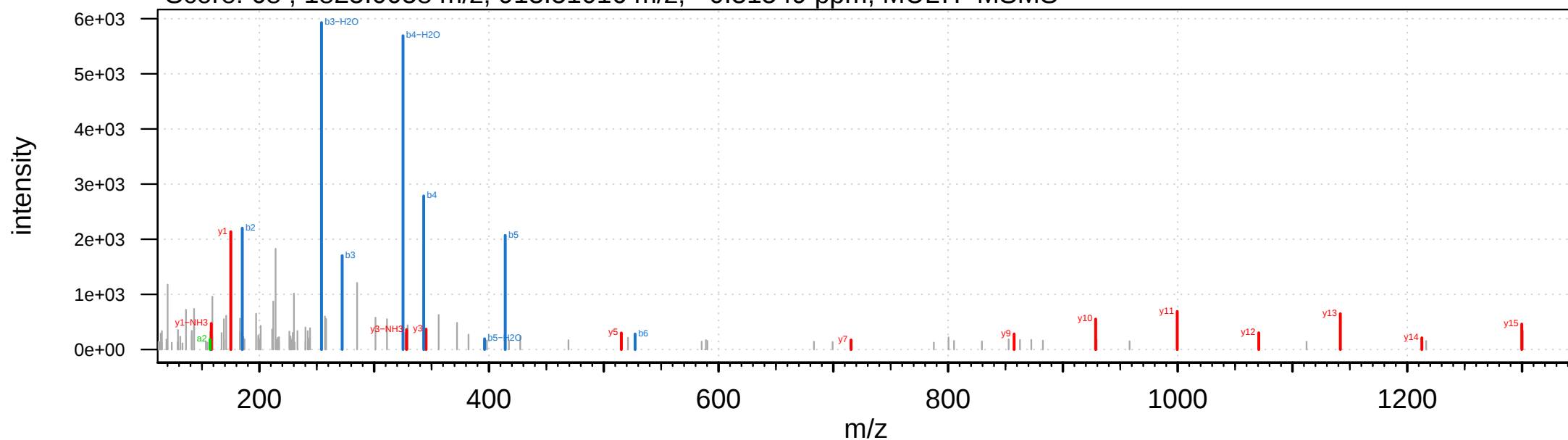
Proteins:

TCONS_I2_00005685_ch12:53548245-53548274:+



(ac)AASAALSAAAAAALSGLAVR

Score: 68 ; 1825.0058 m/z; 913.51016 m/z; -0.31549 ppm; MULTI-MSMS



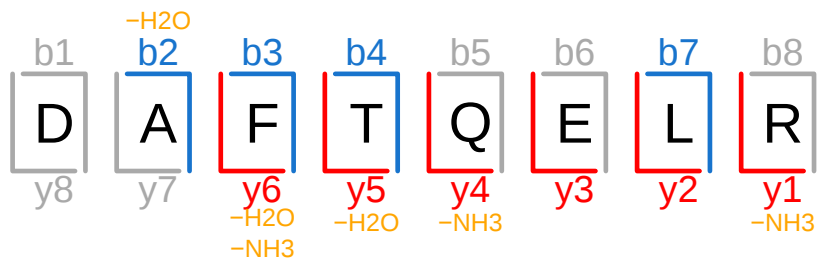
Raw File: 20100719_Velos1_TaGe_SA_LnCap_5

Scan Number: 47630

Proteins:

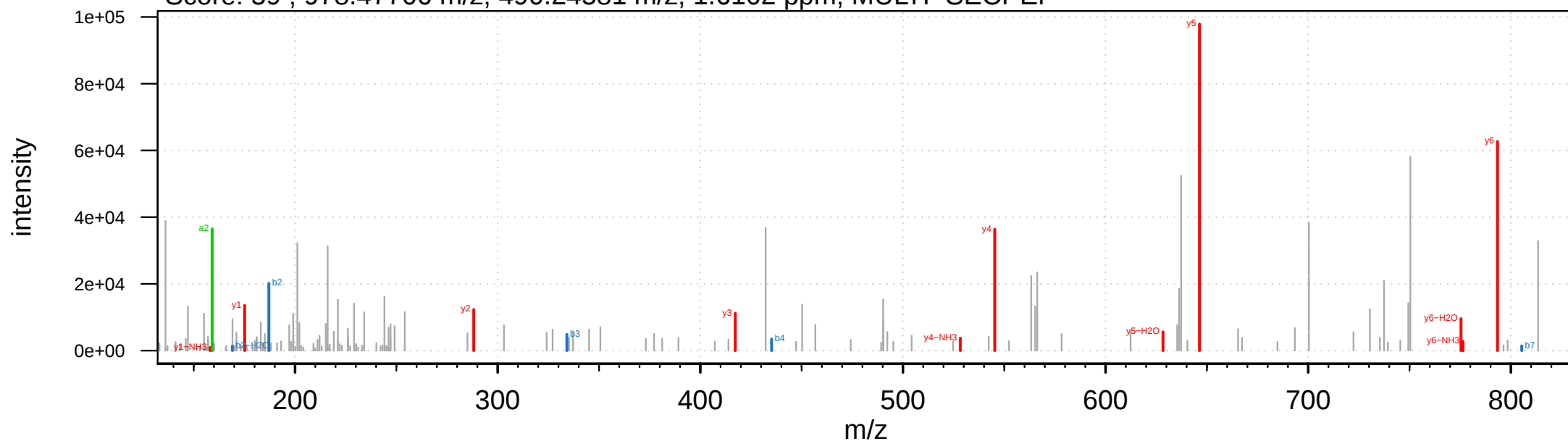
ENST00000433425_chrX:134232427-134232663:-

ENST00000417443_chrX:134556057-134556293:+

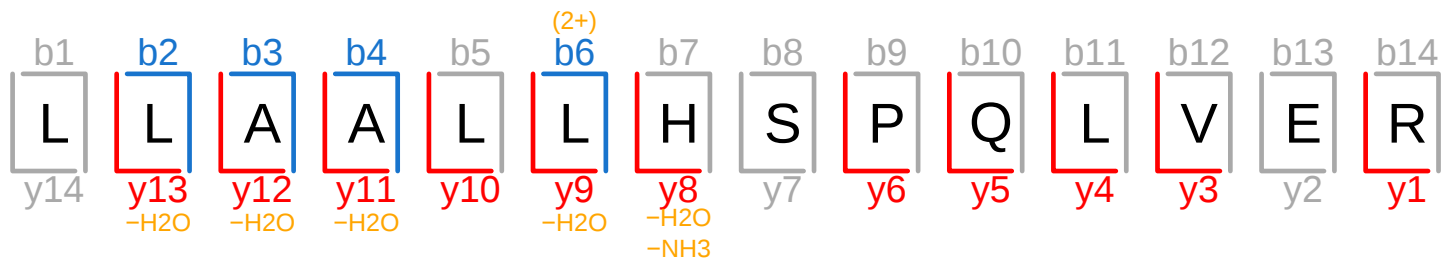


DAFTQELR

Score: 59 ; 978.47706 m/z; 490.24581 m/z; 1.6102 ppm; MULTI-SECPEP

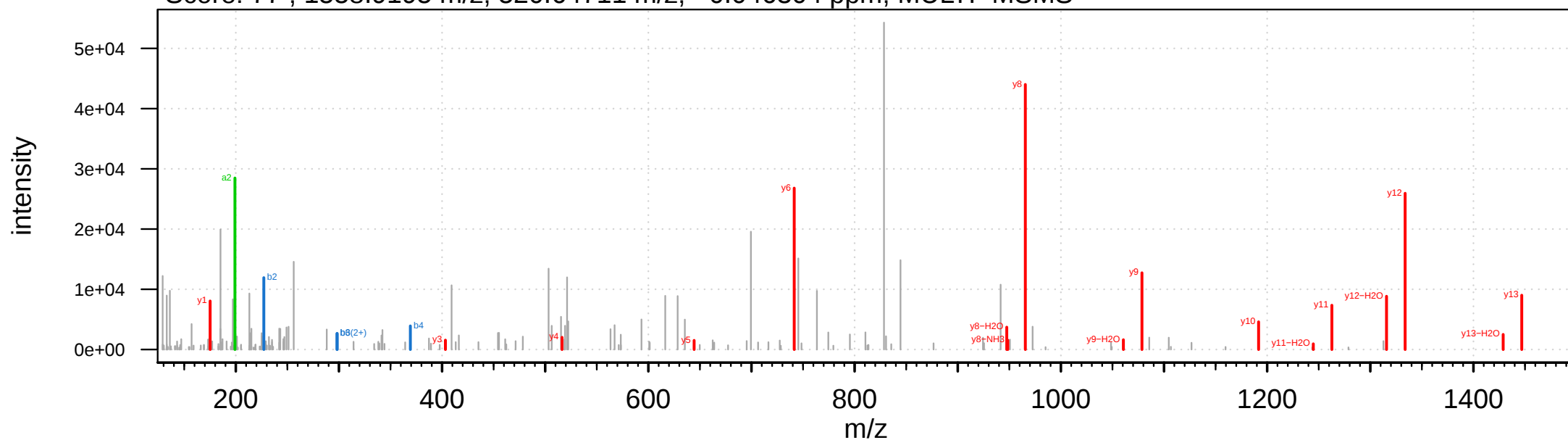


Raw File: 20100611_Velos1_TaGe_SA_Hela_1
 Scan Number: 12547
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+



LLAALLHSPQLVER

Score: 77 ; 1558.9195 m/z; 520.64711 m/z; -0.046304 ppm; MULTI-MSMS

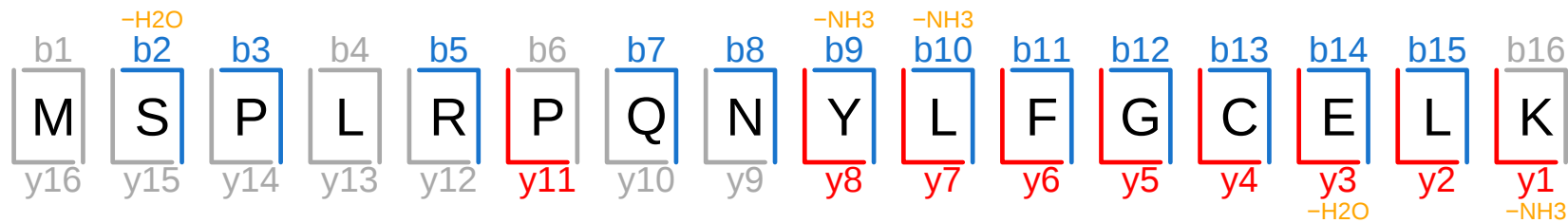


Raw File: 20100611_Velos1_TaGe_SA_Hela_1

Scan Number: 26961

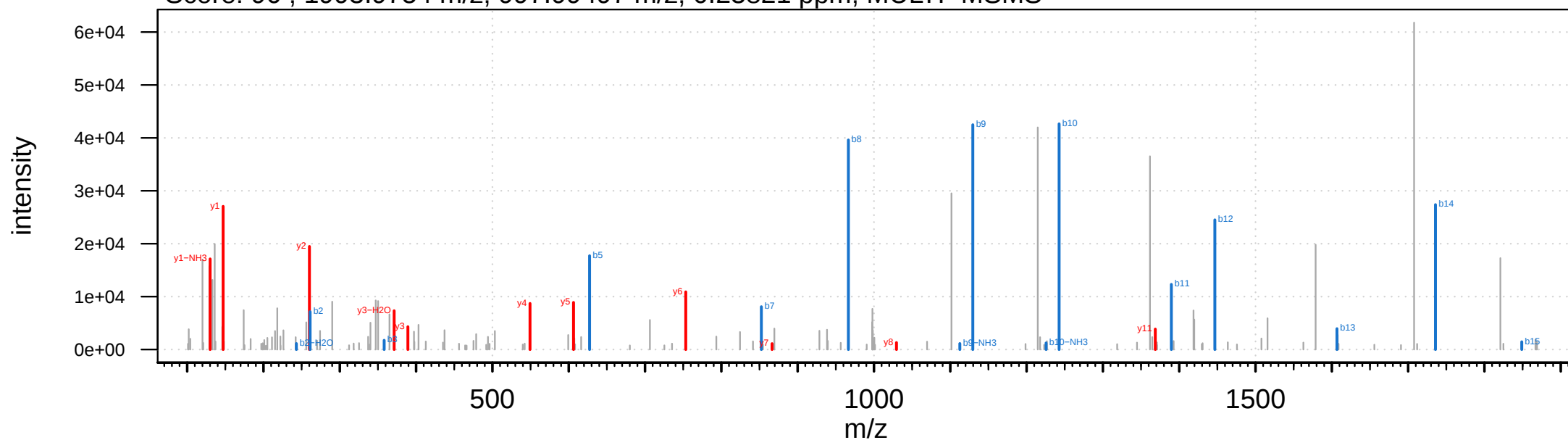
Proteins:

ENST00000602845_chr3:196669588-196669887:+



(ac)MSPLRPQNYLFGCELK

Score: 96 ; 1993.9754 m/z; 997.99497 m/z; 0.23821 ppm; MULTI-MSMS

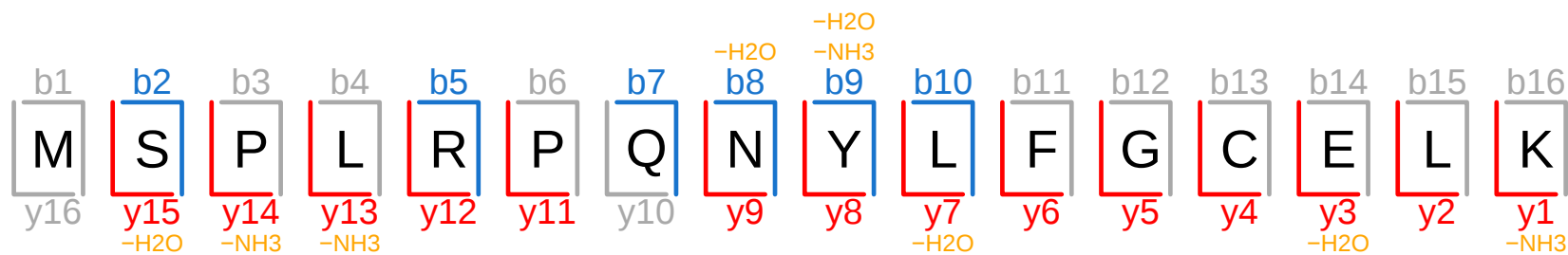


Raw File: 20100611_Velos1_TaGe_SA_Hela_1

Scan Number: 35799

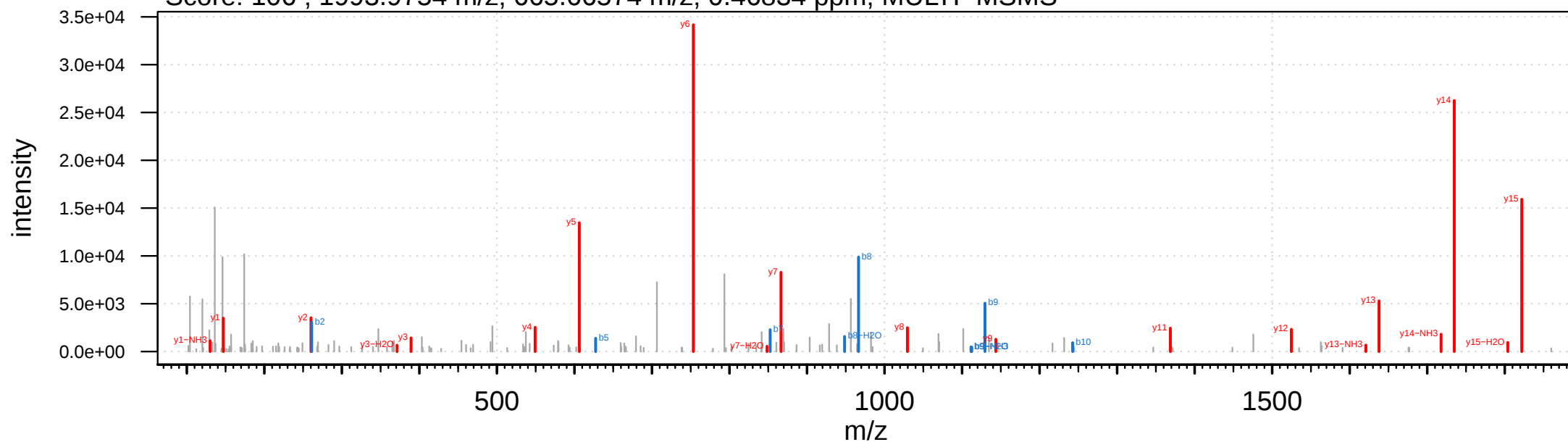
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 106 ; 1993.9754 m/z; 665.66574 m/z; 0.46834 ppm; MULTI-MSMS

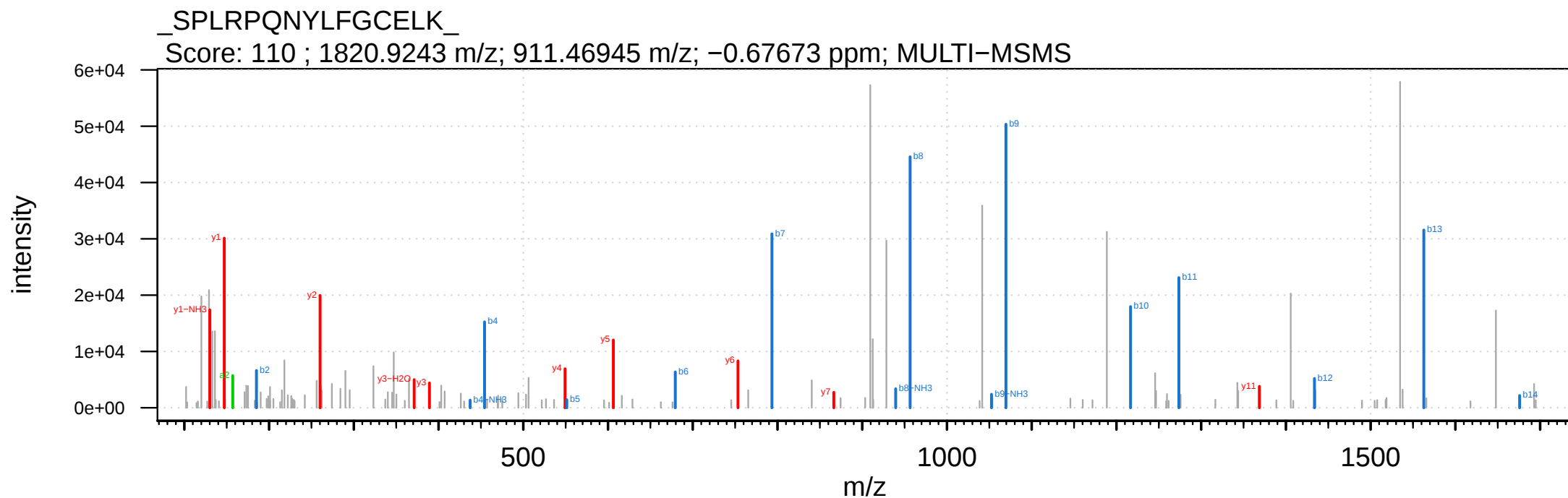
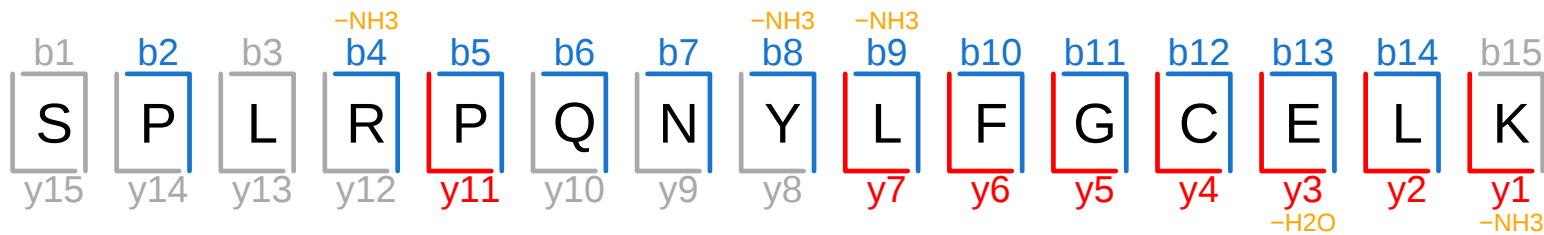


Raw File: 20100611_Velos1-TaGe_SA_Hela_1

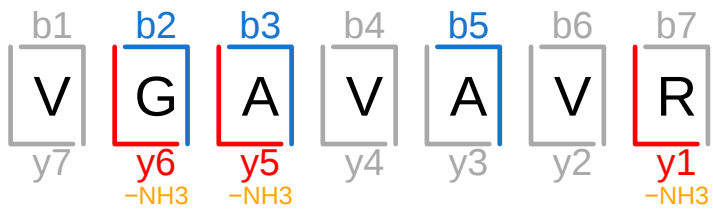
Scan Number: 35813

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

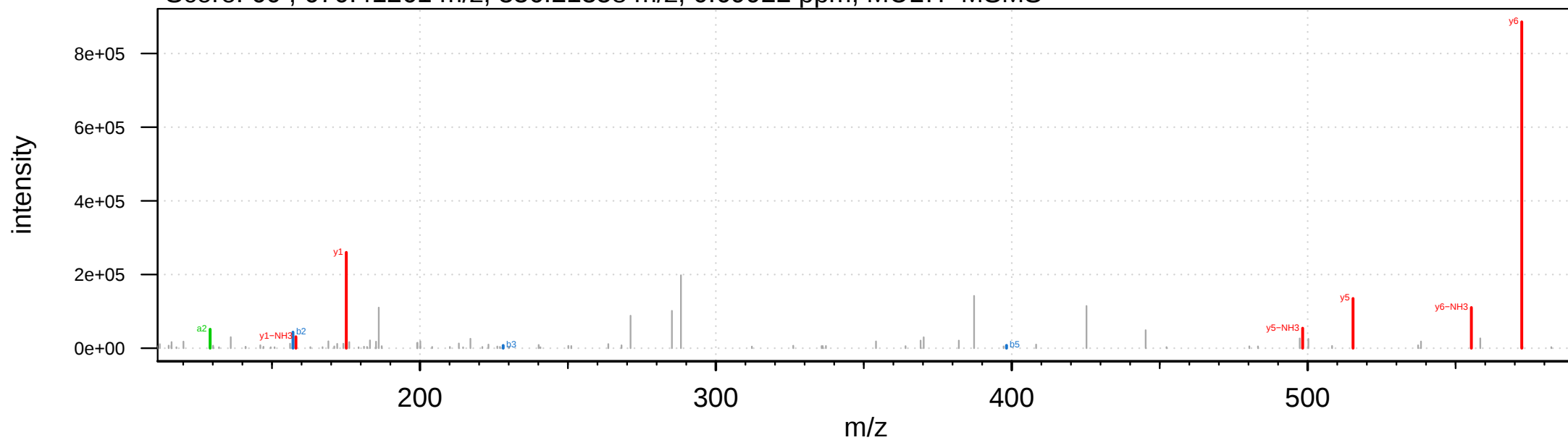


Raw File: 20100611_Velos1_TaGe_SA_Hela_1
 Scan Number: 22310
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



VGAVAVR

Score: 60 ; 670.41261 m/z; 336.21358 m/z; 0.69922 ppm; MULTI-MSMS



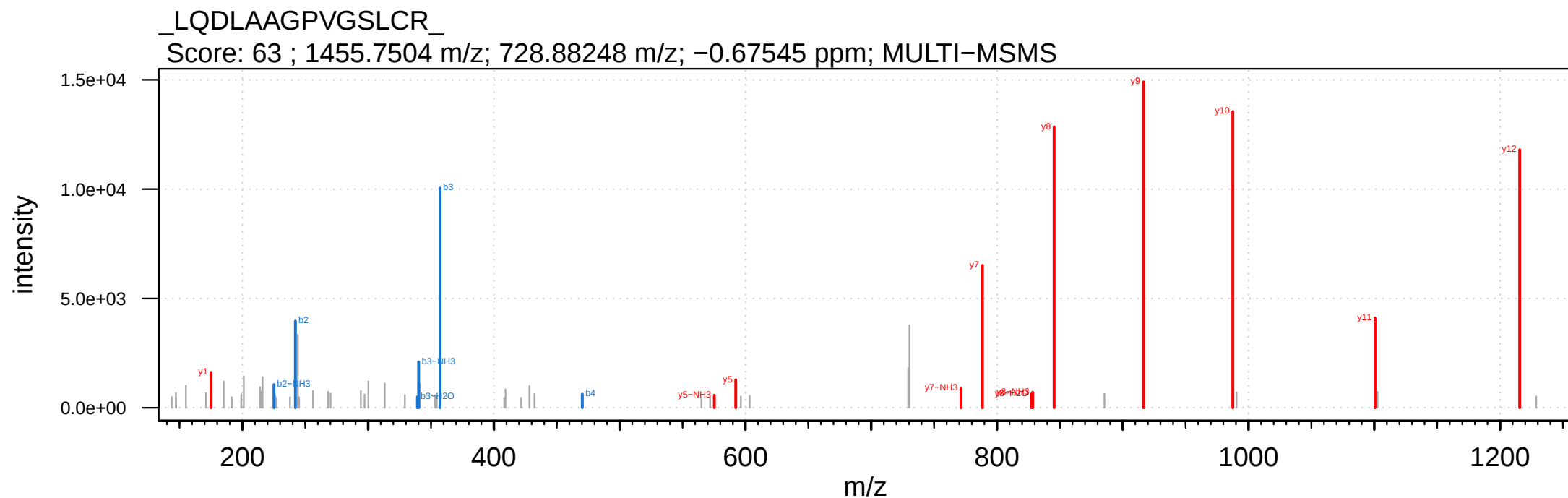
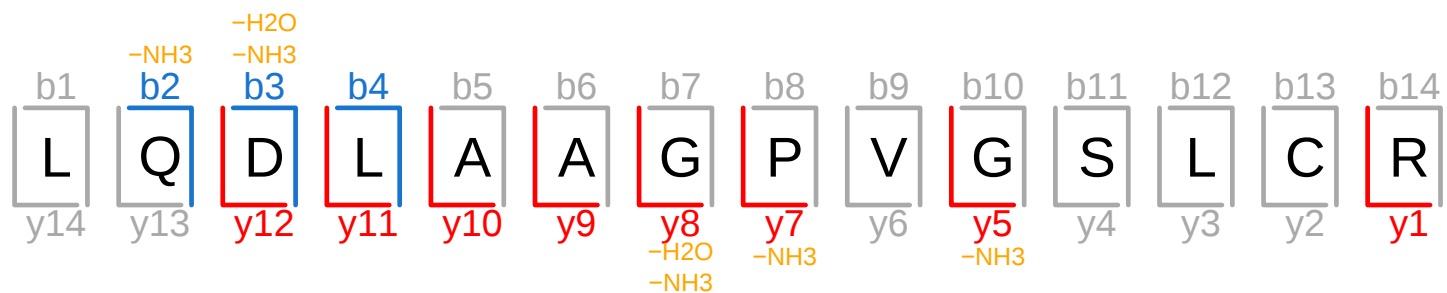
Raw File: 20100611_Velos1_TaGe_SA_Hela_1

Scan Number: 2321

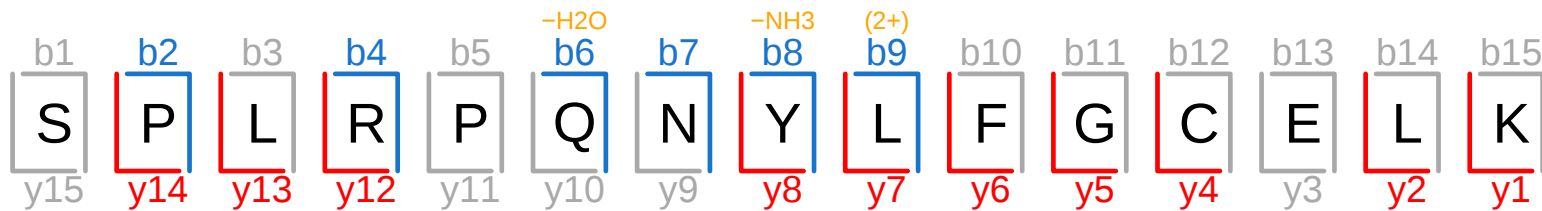
Proteins:

TCONS_I2_00030545_chrX:79544539-79546436:-

Raw File: 20100616_Velos1_TaGe_SA_MCF7_1
Scan Number: 13313
Proteins:
ENST00000602845_chr3:196669588-196669887:+

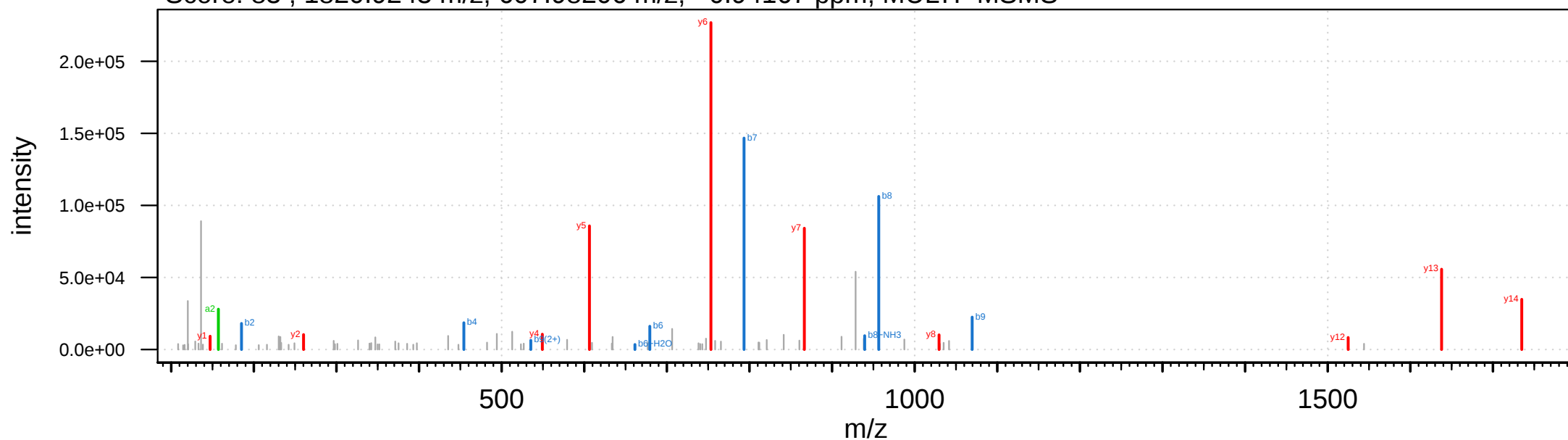


Raw File: 20100616_Velos1_TaGe_SA_MCF7_1
Scan Number: 21903
Proteins:
ENST00000602845_chr3:196669588-196669887:+



SPLRPQNYLFGCELK

Score: 83 ; 1820.9243 m/z; 607.98206 m/z; -0.94167 ppm; MULTI-MSMS

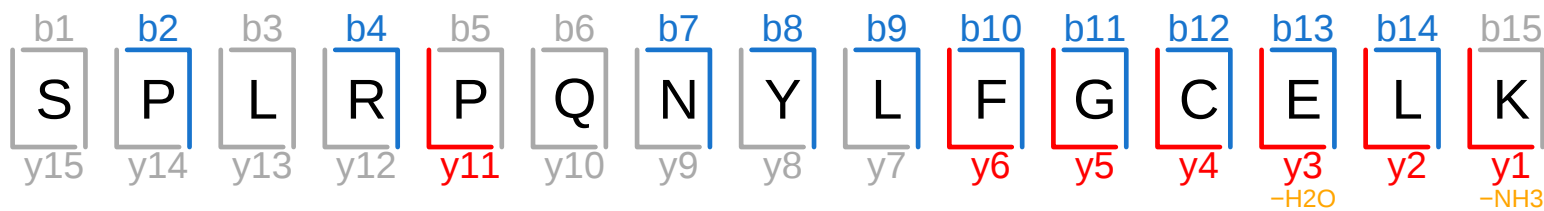


Raw File: 20100616_Velos1_TaGe_SA_MCF7_1

Scan Number: 23529

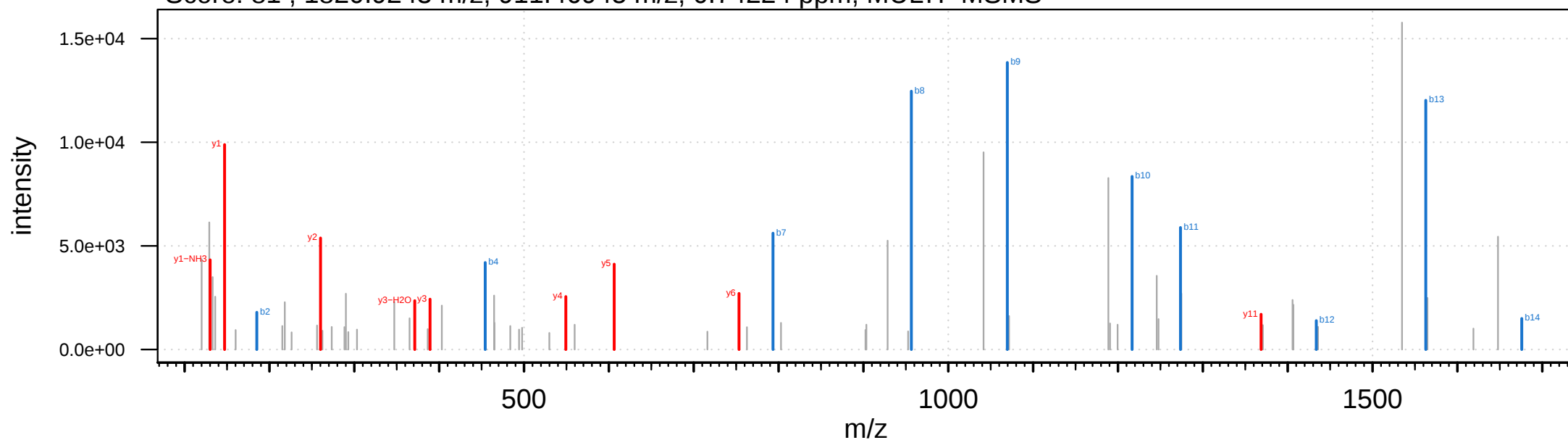
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 81 ; 1820.9243 m/z; 911.46945 m/z; 0.74224 ppm; MULTI-MSMS

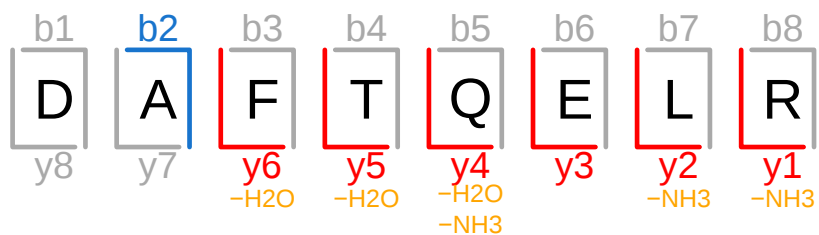


Raw File: 20100616_Velos1_TaGe_SA_MCF7_1

Scan Number: 23544

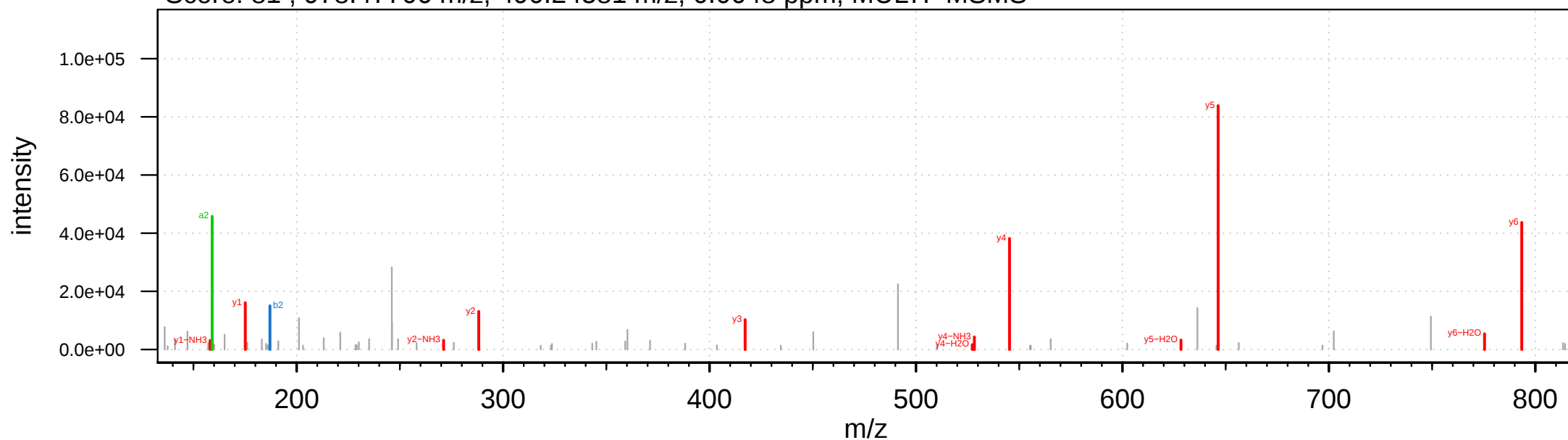
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



DAFTQELR

Score: 81 ; 978.47706 m/z; 490.24581 m/z; 0.0648 ppm; MULTI-MSMS

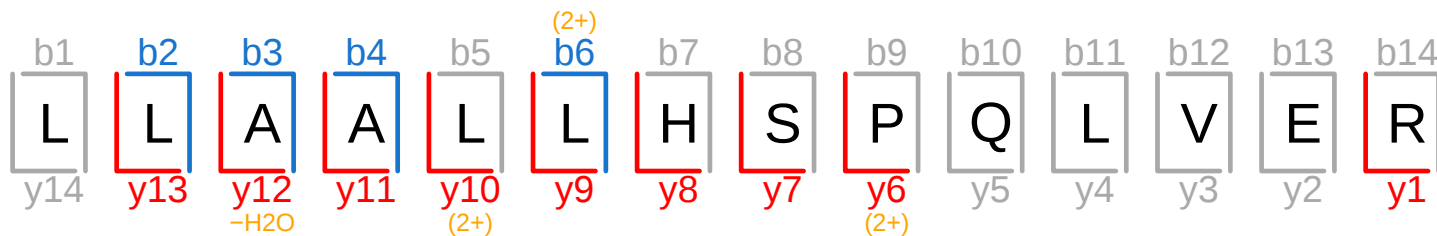


Raw File: 20100616_Velos1_TaGe_SA_MCF7_2

Scan Number: 12441

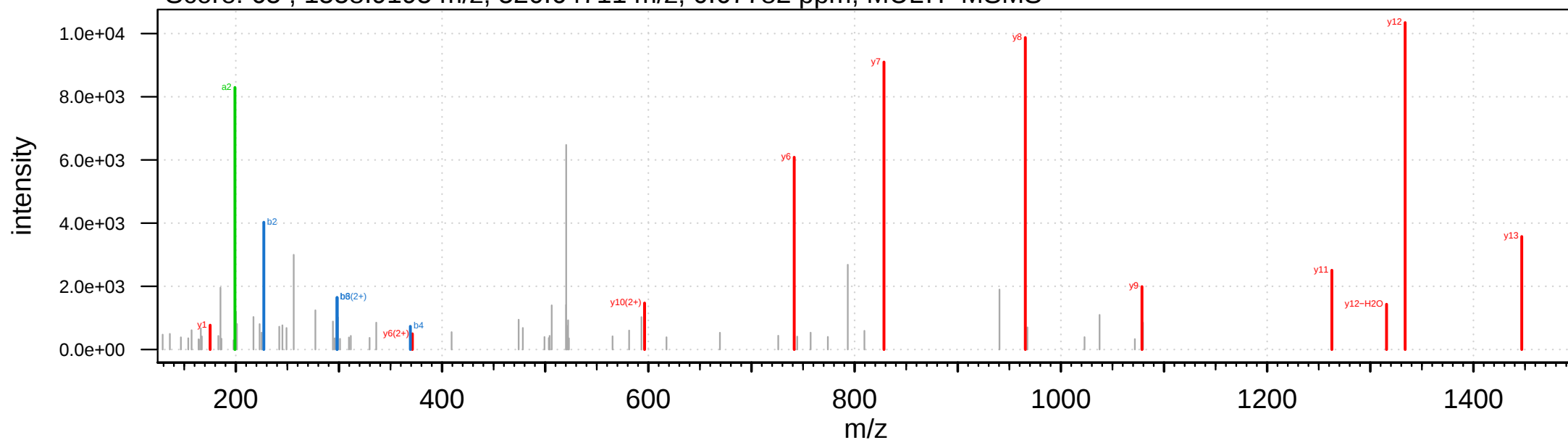
Proteins:

ENST00000602845_chr3:196669588-196669887:+



LLAALLHSPQLVER

Score: 65 ; 1558.9195 m/z; 520.64711 m/z; 0.67782 ppm; MULTI-MSMS

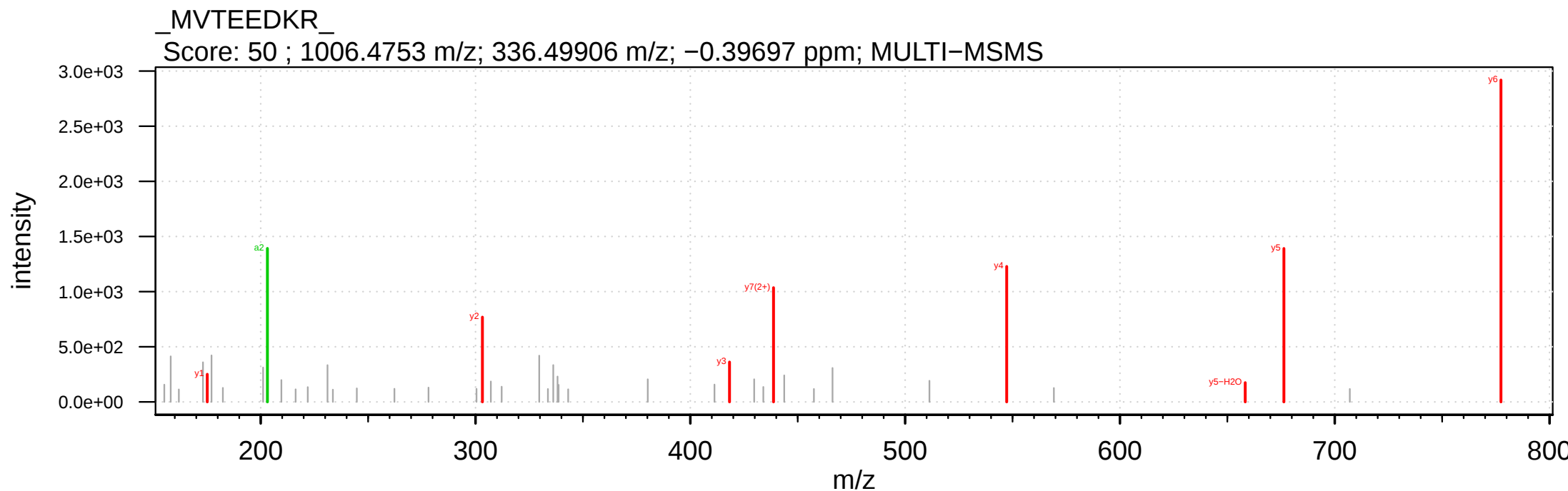
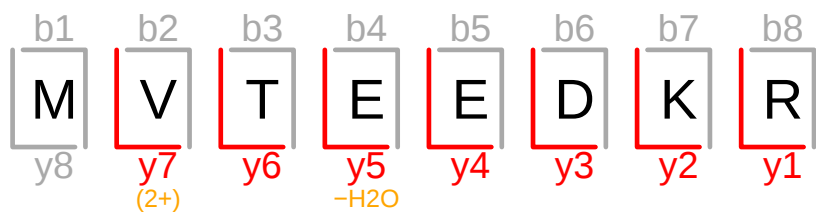


Raw File: 20100616_Velos1_TaGe_SA_MCF7_2

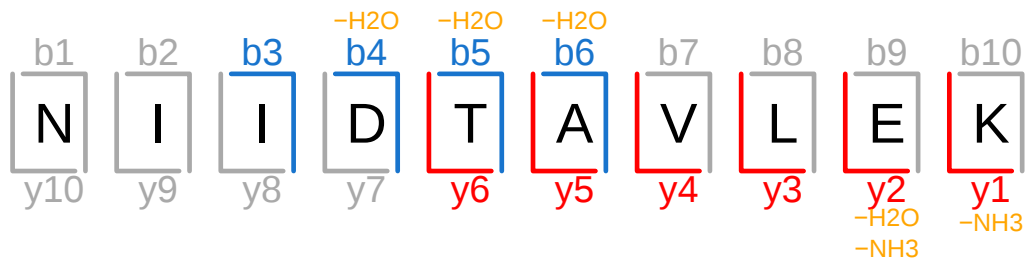
Scan Number: 24215

Proteins:

ENST00000602845_chr3:196669588-196669887:+

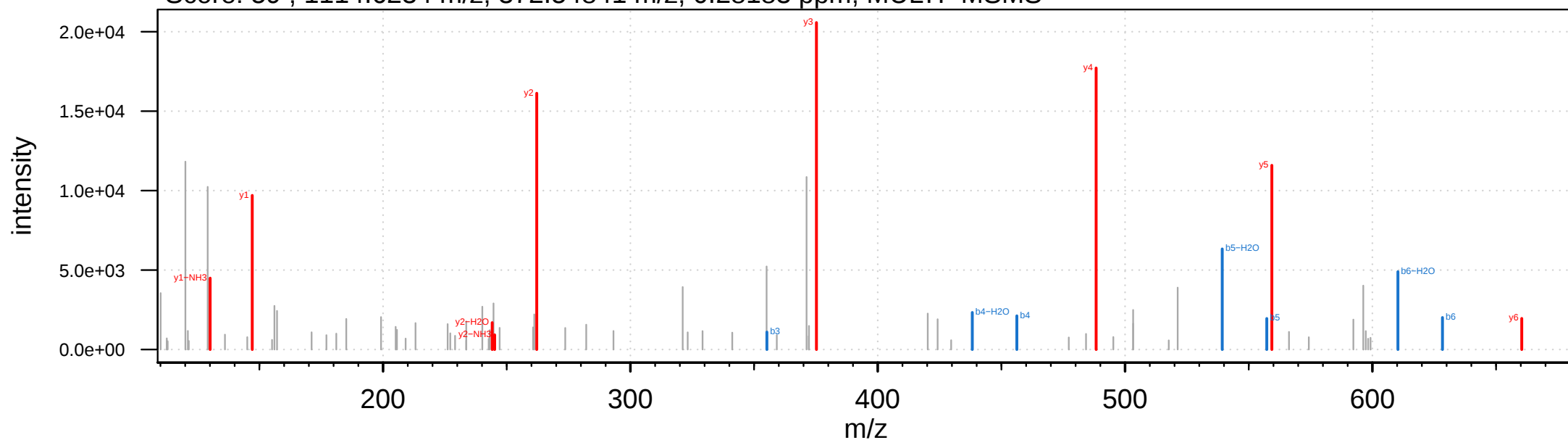


Raw File: 20100616_Velos1_TaGe_SA_MCF7_2
Scan Number: 1495
Proteins:
TCONS_I2_00006768_chr13:29174091-29174123:+

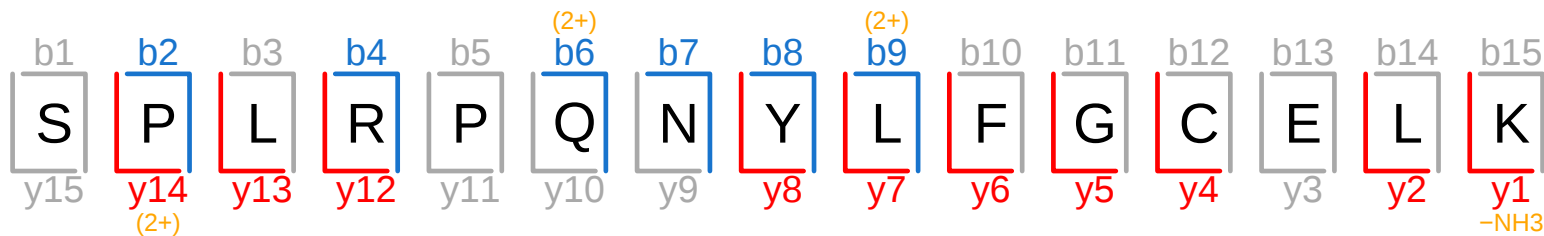


NIIDTAVLEK

Score: 59 ; 1114.6234 m/z; 372.54841 m/z; 0.28183 ppm; MULTI-MSMS

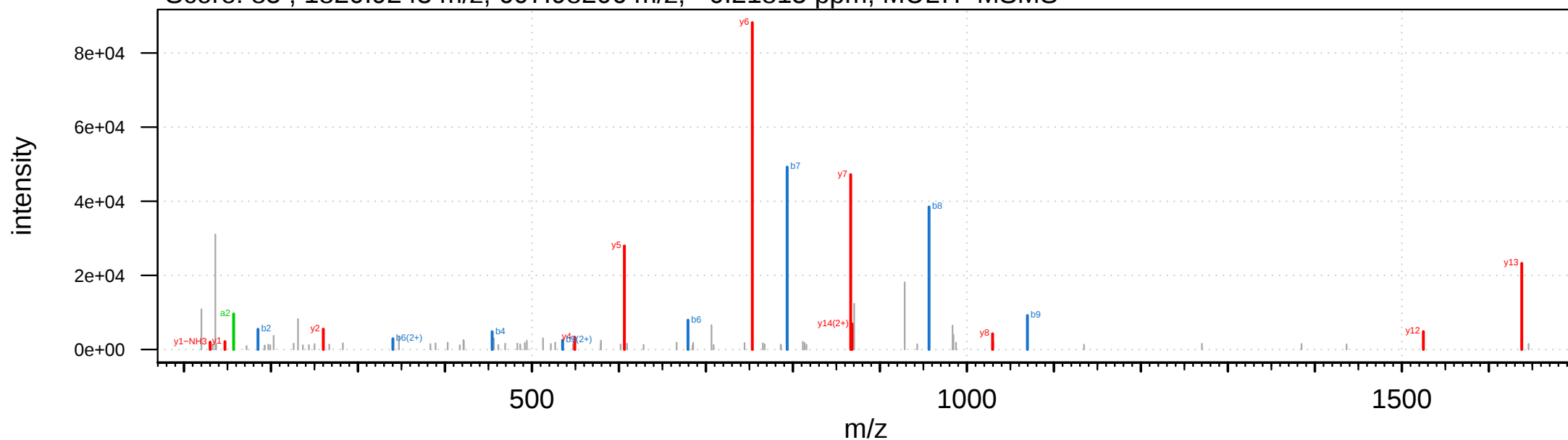


Raw File: 20100616_Velos1_TaGe_SA_MCF7_2
 Scan Number: 7680
 Proteins:
 ENST00000497138_chr20:56806826-56807846:-



SPLRPQNYLFGCELK

Score: 85 ; 1820.9243 m/z; 607.98206 m/z; -0.21815 ppm; MULTI-MSMS

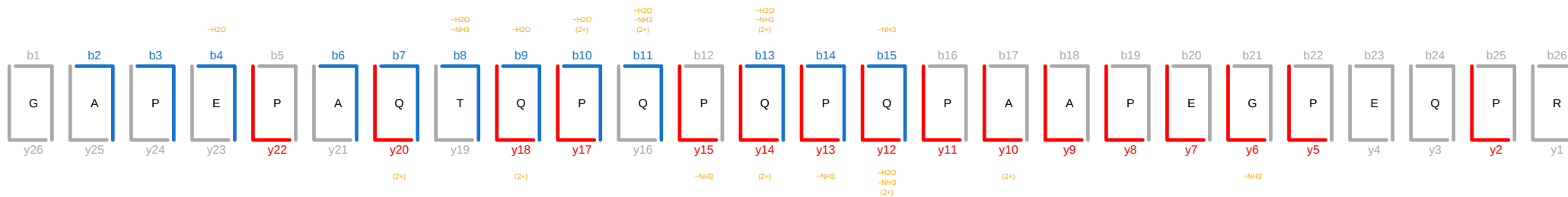


Raw File: 20100616_Velos1_TaGe_SA_MCF7_2

Scan Number: 21019

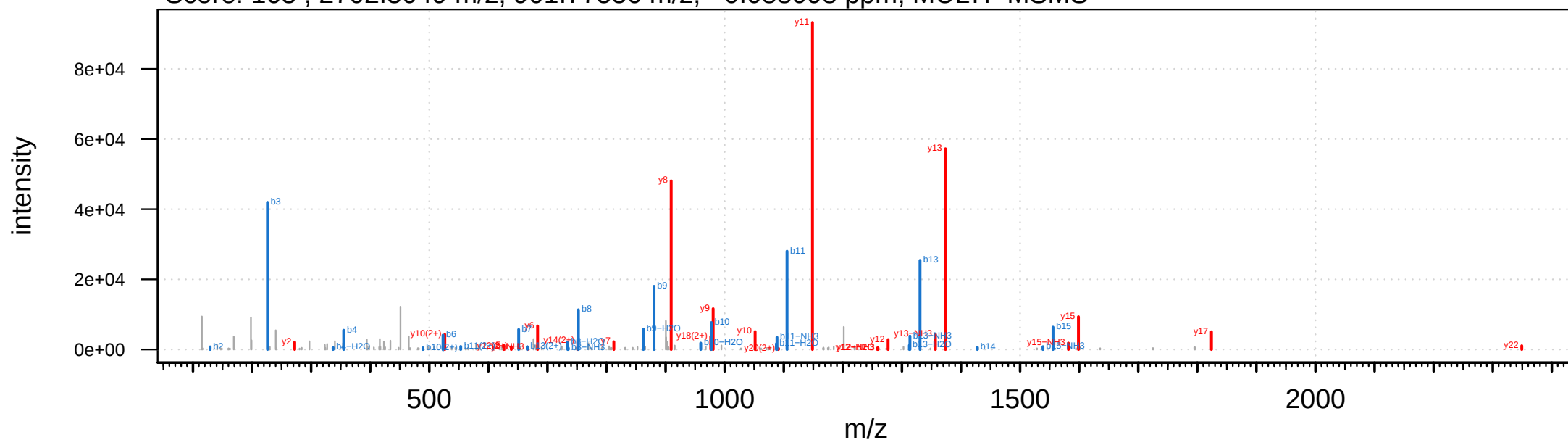
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

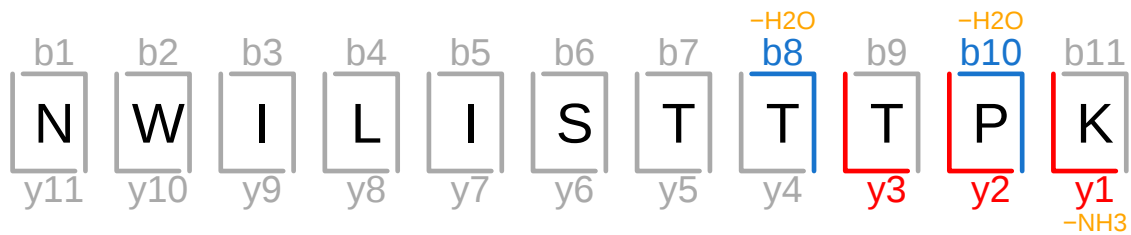


GAPEPAQTQPQPQPQPAAPGPEQPR

Score: 163 ; 2702.3049 m/z; 901.77556 m/z; -0.088098 ppm; MULTI-MSMS

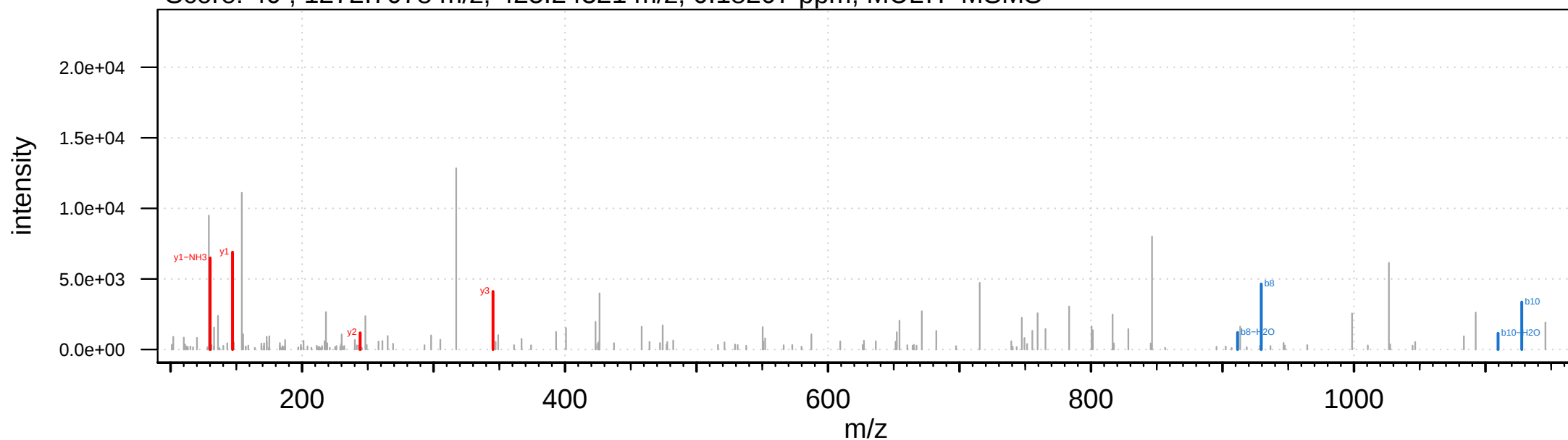


Raw File: 20101215_Velos1_TaGe_SA_A549_01
 Scan Number: 11582
 Proteins:
 ENST00000424358_chr20:33865517-33865732:-

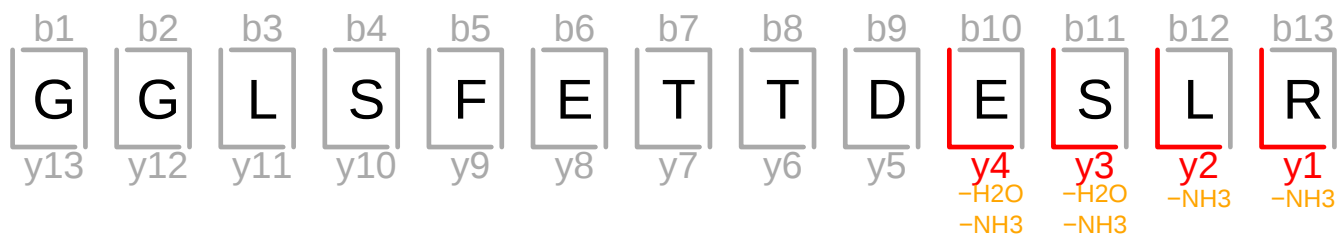


NWILISTTTPK

Score: 40 ; 1272.7078 m/z; 425.24321 m/z; 0.18207 ppm; MULTI-MSMS

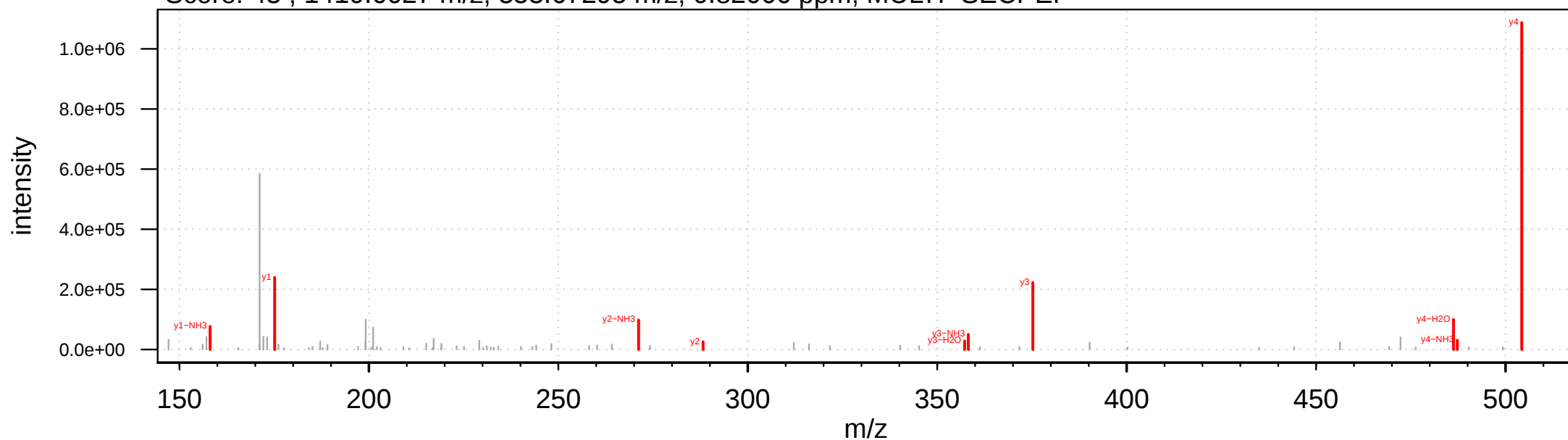


Raw File: 20101215_Velos1_TaGe_SA_A549_01
 Scan Number: 23826
 Proteins:
 ENST00000430431_chr10:61496818-61496958:-



GGLSFETTDESLR

Score: 43 ; 1410.6627 m/z; 353.67295 m/z; 0.82066 ppm; MULTI-SECPEP

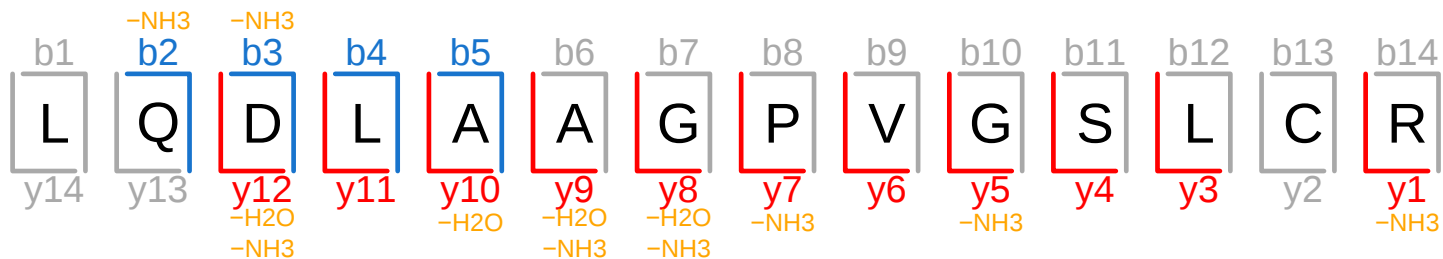


Raw File: 20101210_Velos1_AnWe_SA_MCF7_1

Scan Number: 6288

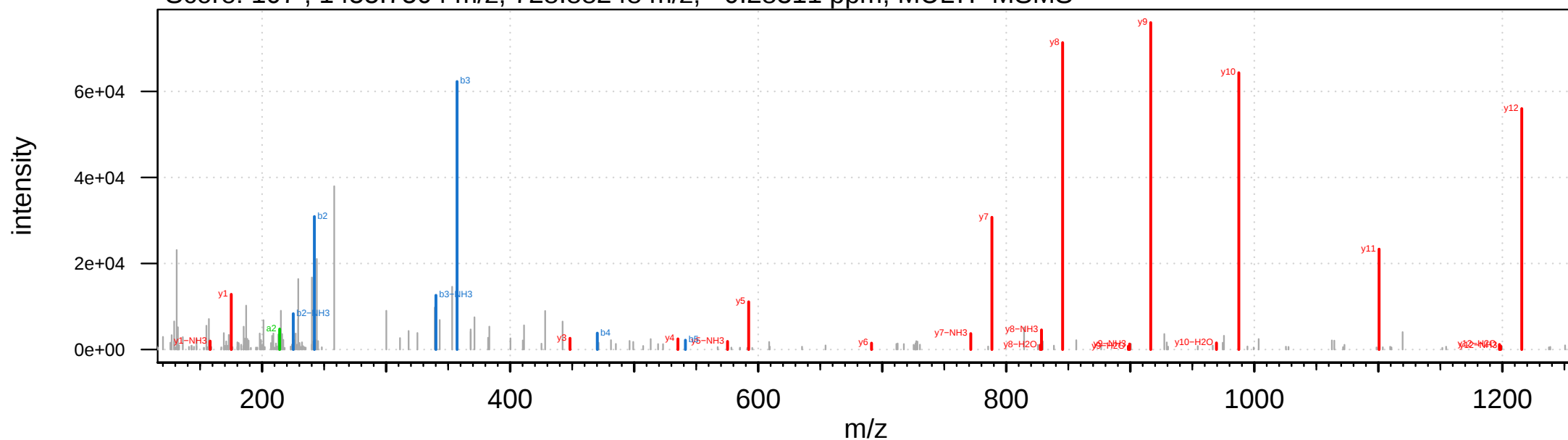
Proteins:

ENST00000511590_chr4:113802003-113803254:-

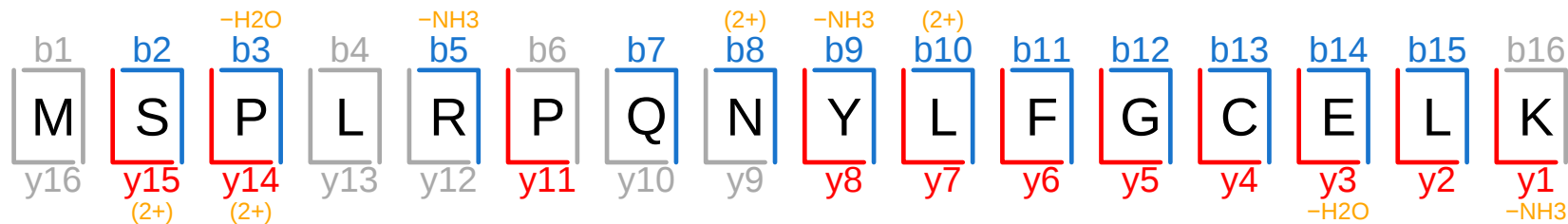


LQDLAAGPVGSLCR

Score: 107 ; 1455.7504 m/z; 728.88248 m/z; -0.28311 ppm; MULTI-MSMS

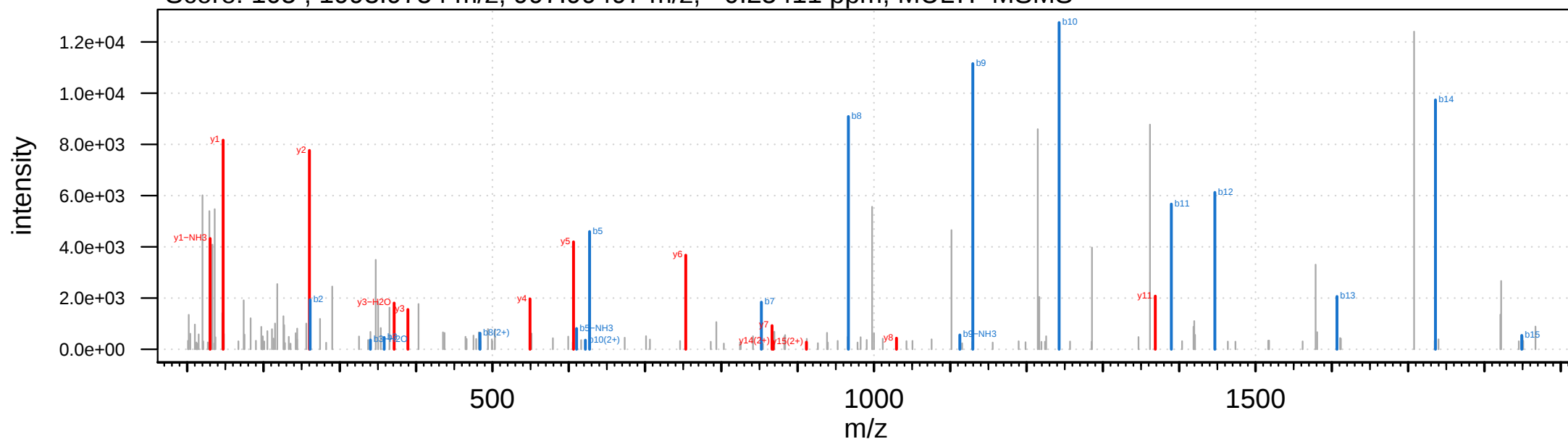


Raw File: 20101210_Velos1_AnWe_SA_MCF7_1
 Scan Number: 21471
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+



(ac)MSPLRPQNYLFGCELK

Score: 105 ; 1993.9754 m/z; 997.99497 m/z; -0.25411 ppm; MULTI-MSMS

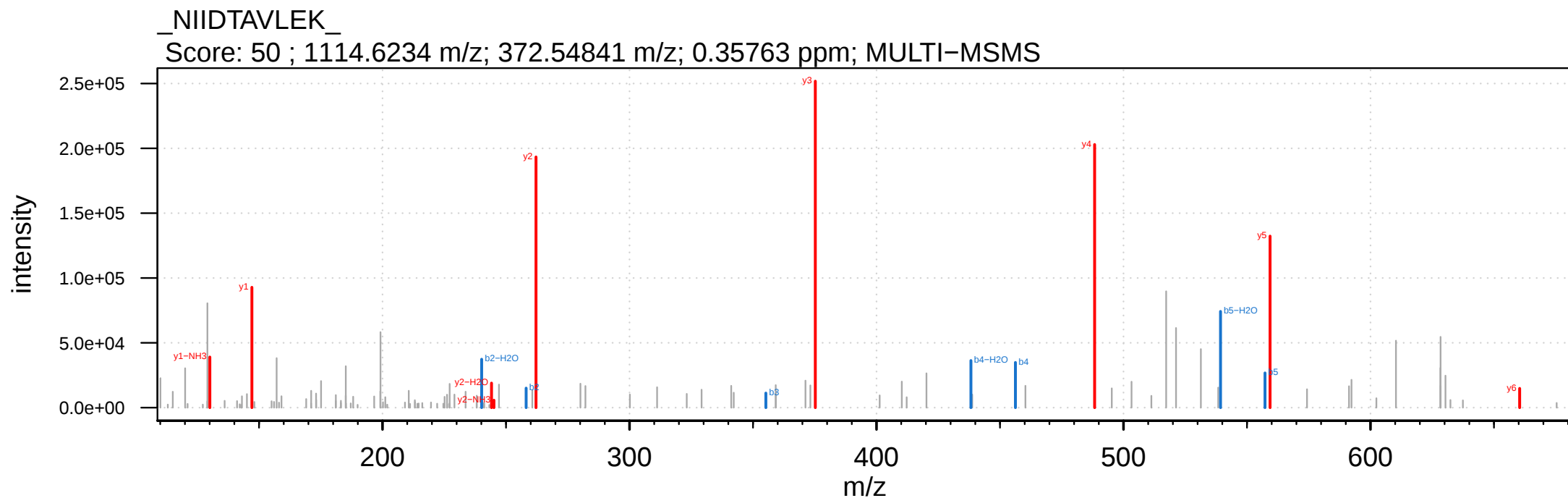
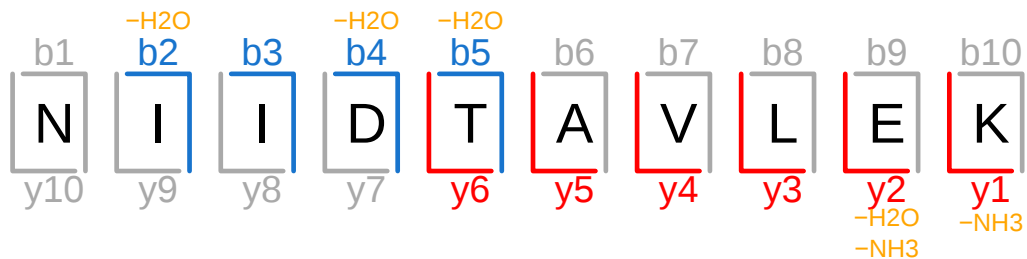


Raw File: 20101210_Velos1_AnWe_SA_MCF7_1

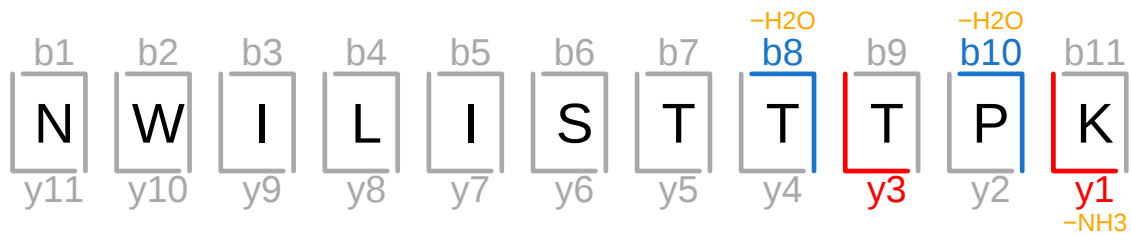
Scan Number: 36201

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

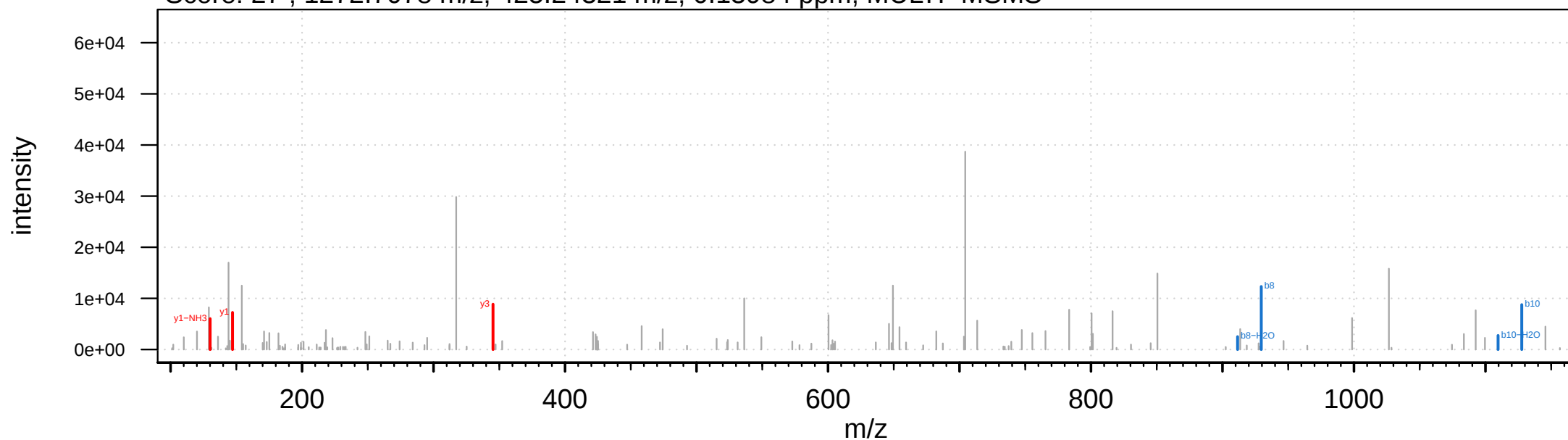


Raw File: 20101210_Velos1_AnWe_SA_MCF7_1
Scan Number: 9268
Proteins:
ENST00000497138_chr20:56806826-56807846:-



NWILISTTTPK

Score: 27 ; 1272.7078 m/z; 425.24321 m/z; 0.13984 ppm; MULTI-MSMS

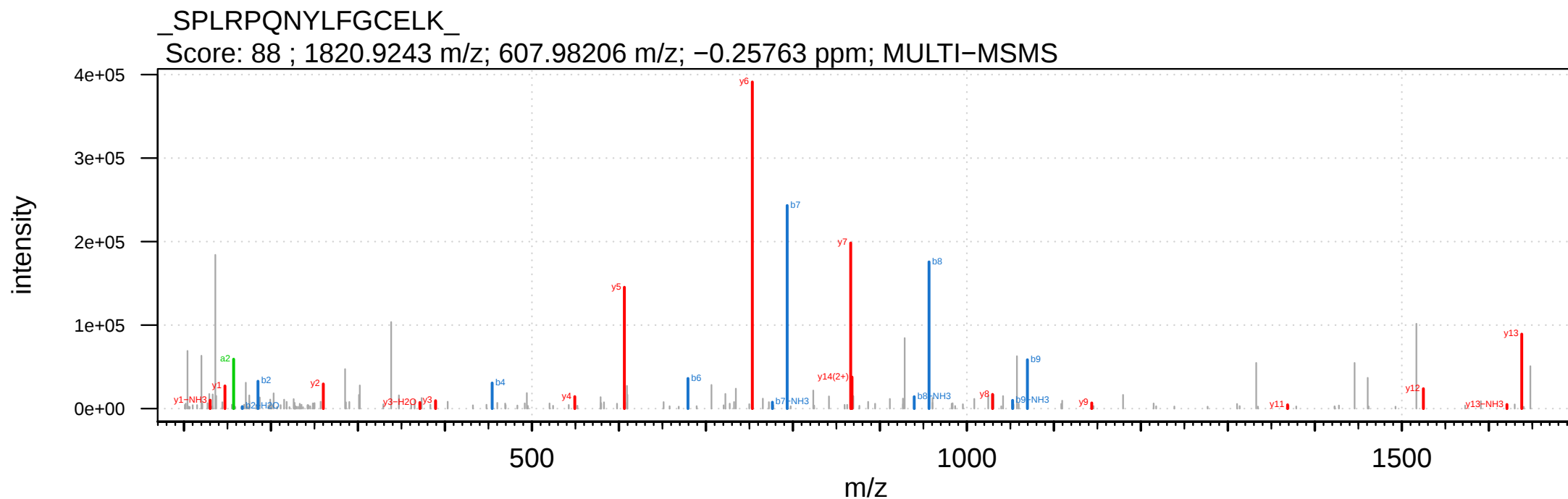
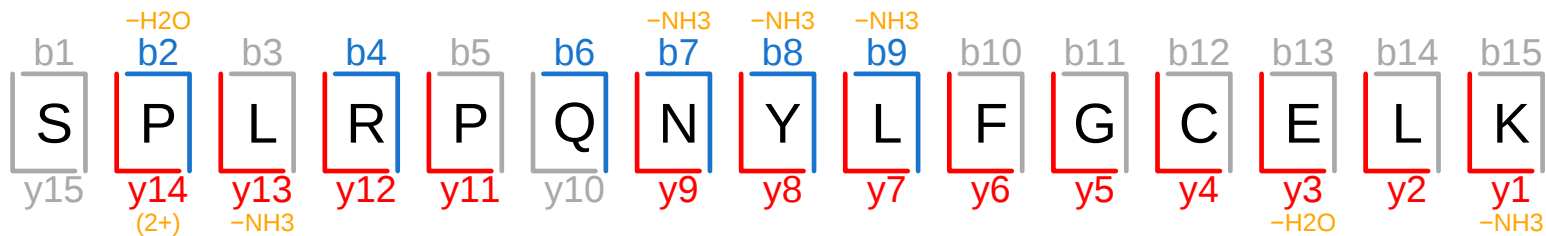


Raw File: 20101210_Velos1_AnWe_SA_MCF7_1

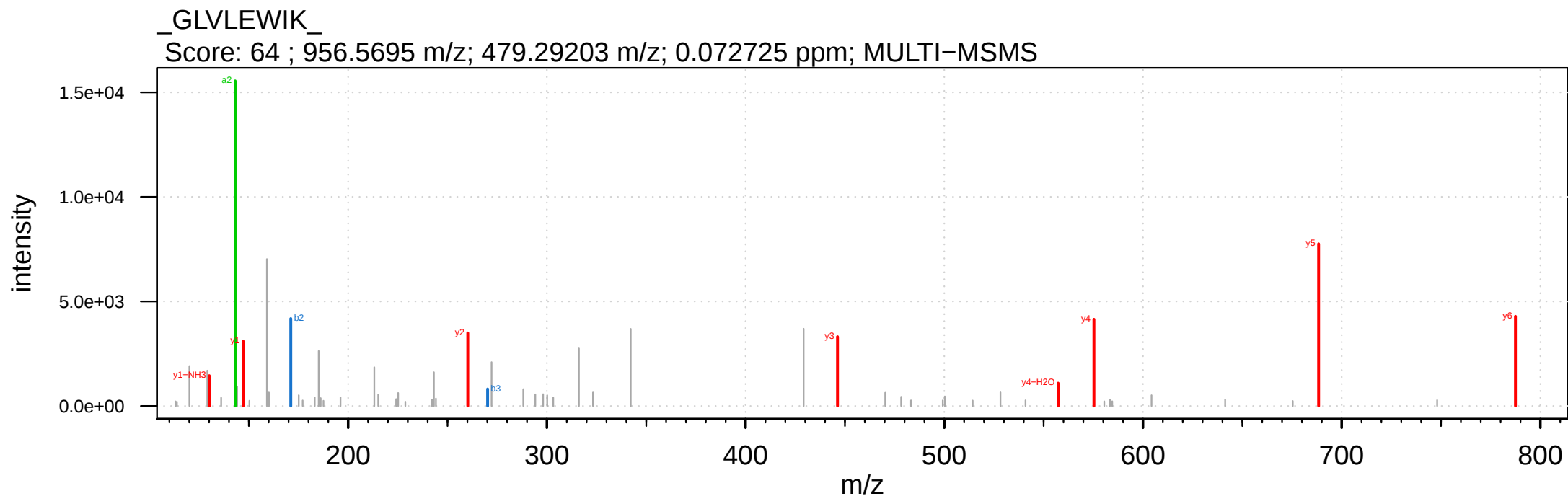
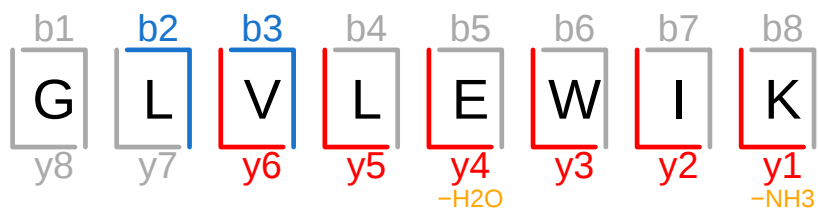
Scan Number: 26493

Proteins:

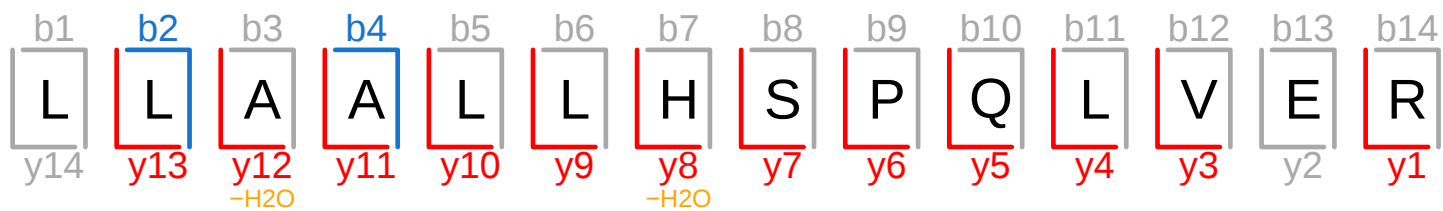
ENST00000430431_chr10:61496818-61496958:-



Raw File: 20101210_Velos1_AnWe_SA_MCF7_1
 Scan Number: 23299
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

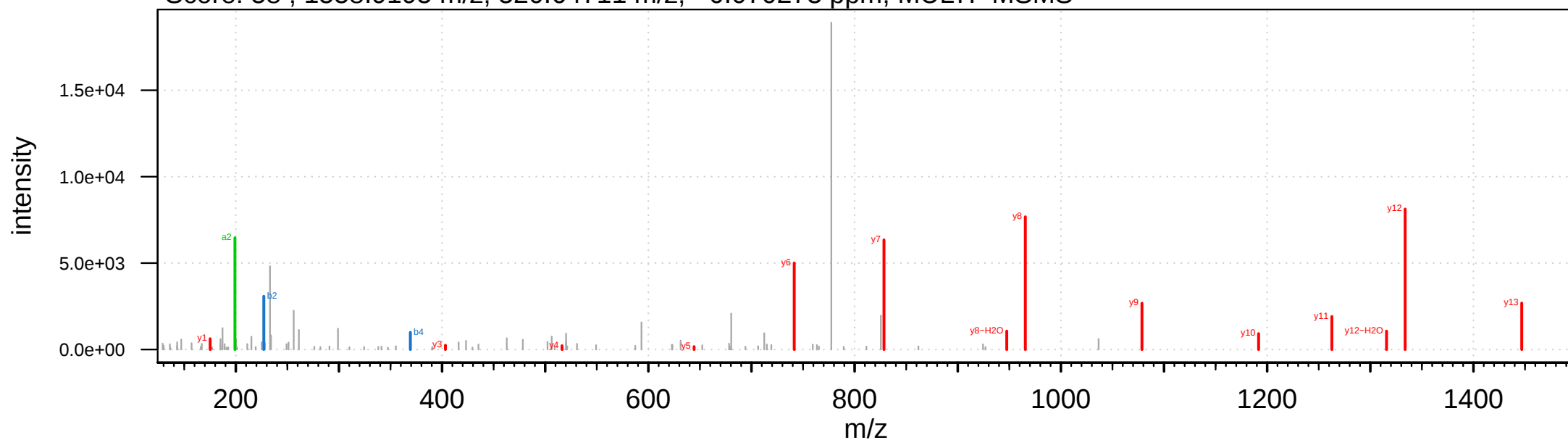


Raw File: 20100618_Velos1_TaGe_SA_LanCap_2
 Scan Number: 25327
 Proteins:
 TCONS_I2_00001296_chr1:79520703-79520992:-



LLAALLHSPQLVER

Score: 58 ; 1558.9195 m/z; 520.64711 m/z; -0.070273 ppm; MULTI-MSMS

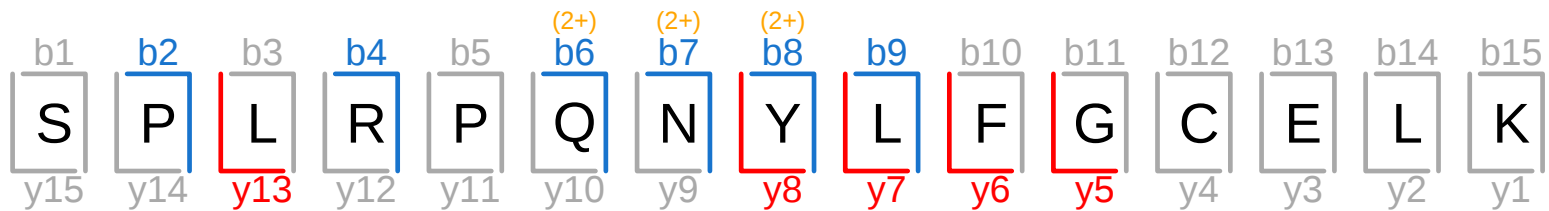


Raw File: 20100618_Velos1_TaGe_SA_LanCap_2

Scan Number: 23130

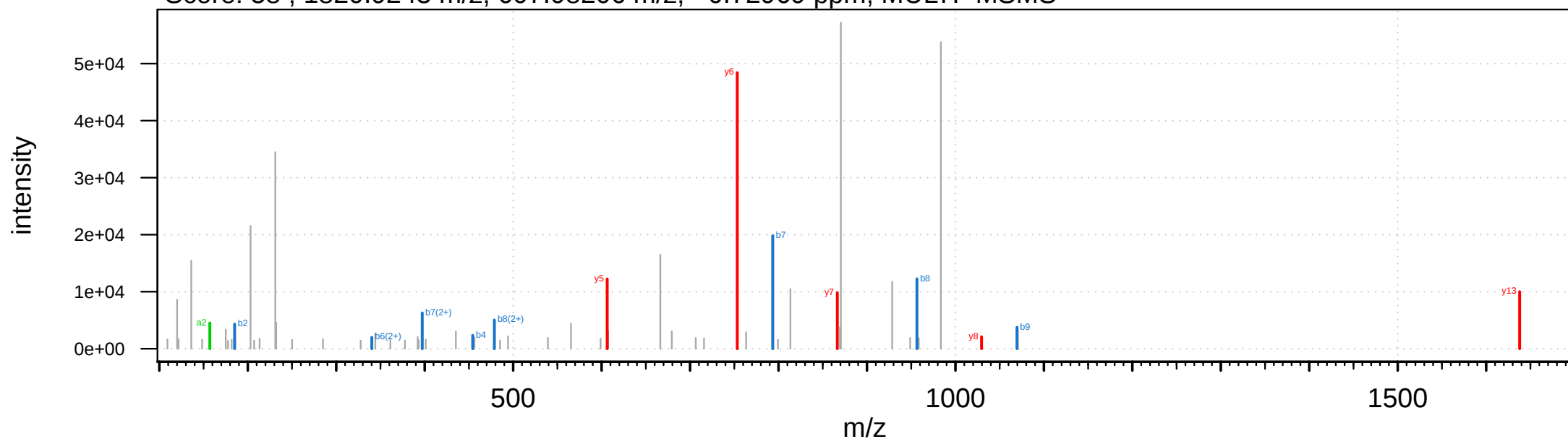
Proteins:

ENST00000602845_chr3:196669588-196669887:+



SPLRPQNYLFGCELK

Score: 58 ; 1820.9243 m/z; 607.98206 m/z; -0.72969 ppm; MULTI-MSMS

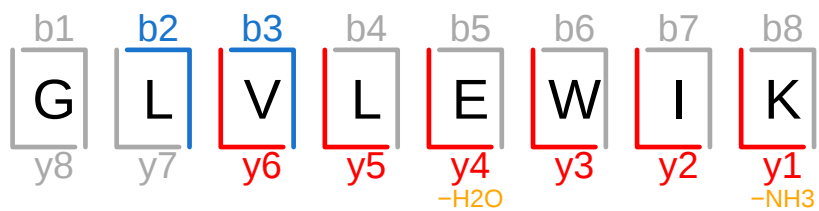


Raw File: 20100618_Velos1_TaGe_SA_LanCap_2

Scan Number: 19882

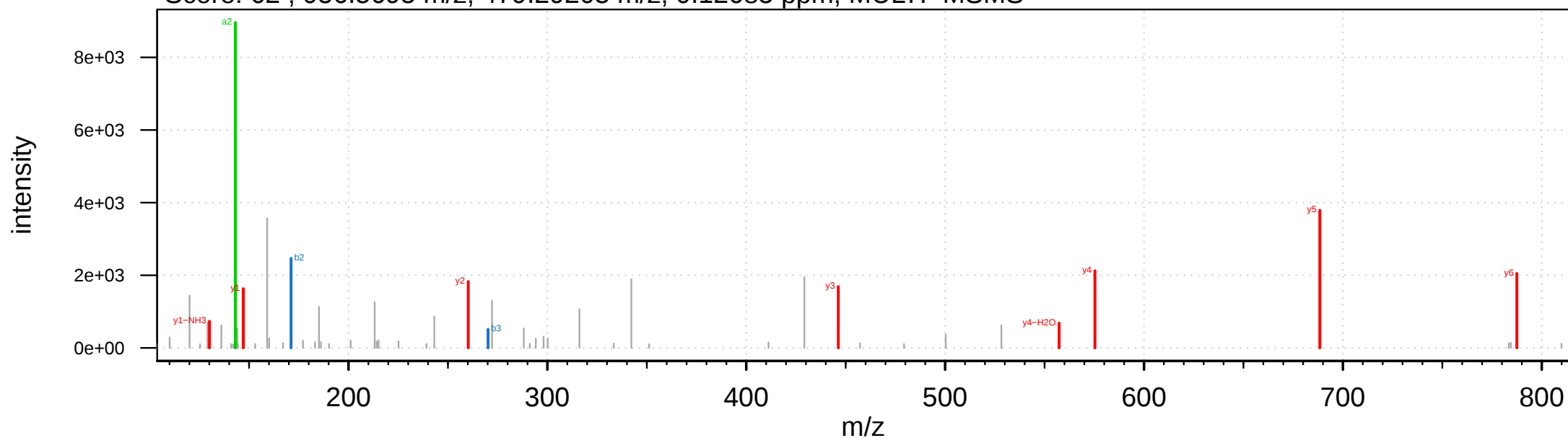
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



GLVLEWIK

Score: 62 ; 956.5695 m/z; 479.29203 m/z; 0.12685 ppm; MULTI-MSMS

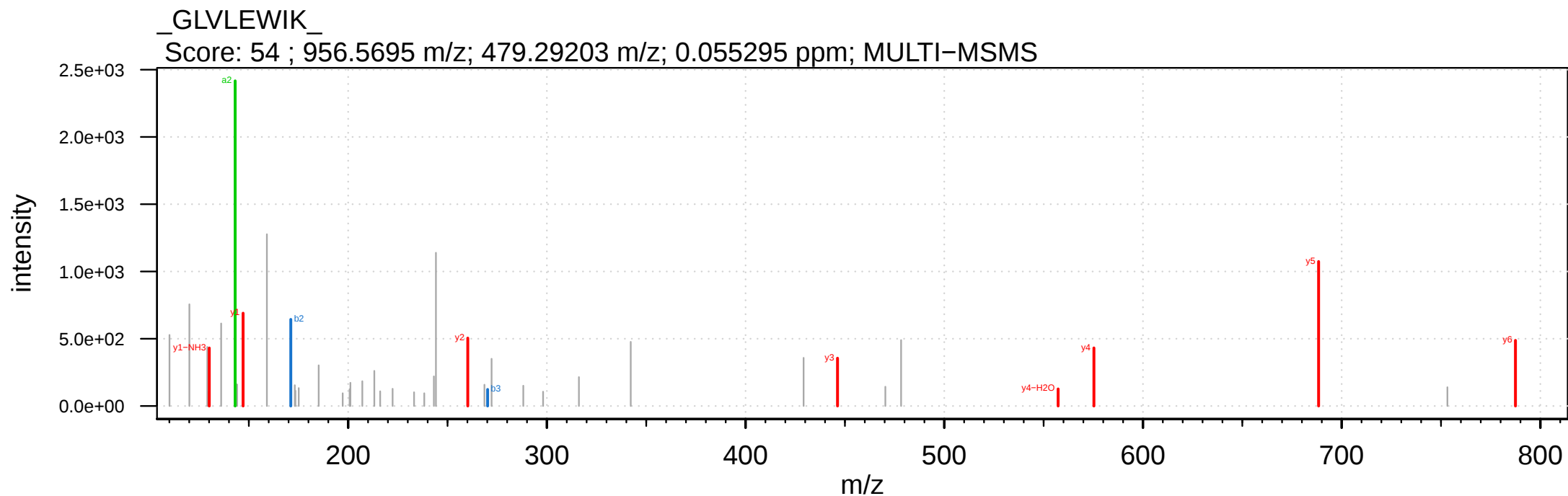
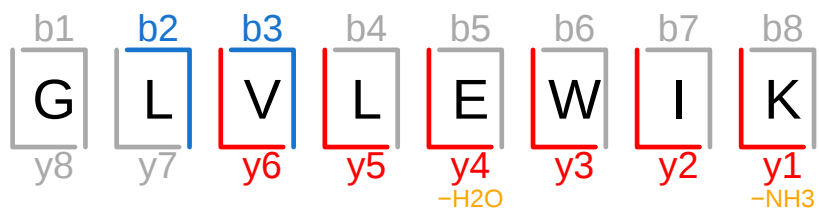


Raw File: 20100618_Velos1_TaGe_SA_LanCap_3

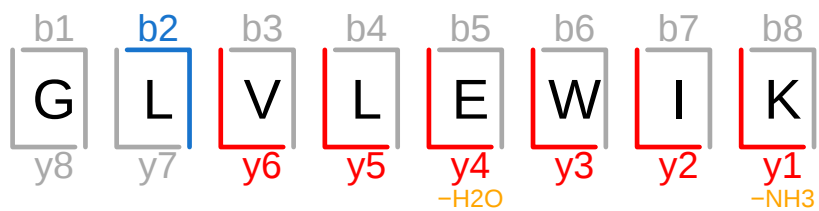
Scan Number: 23438

Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-

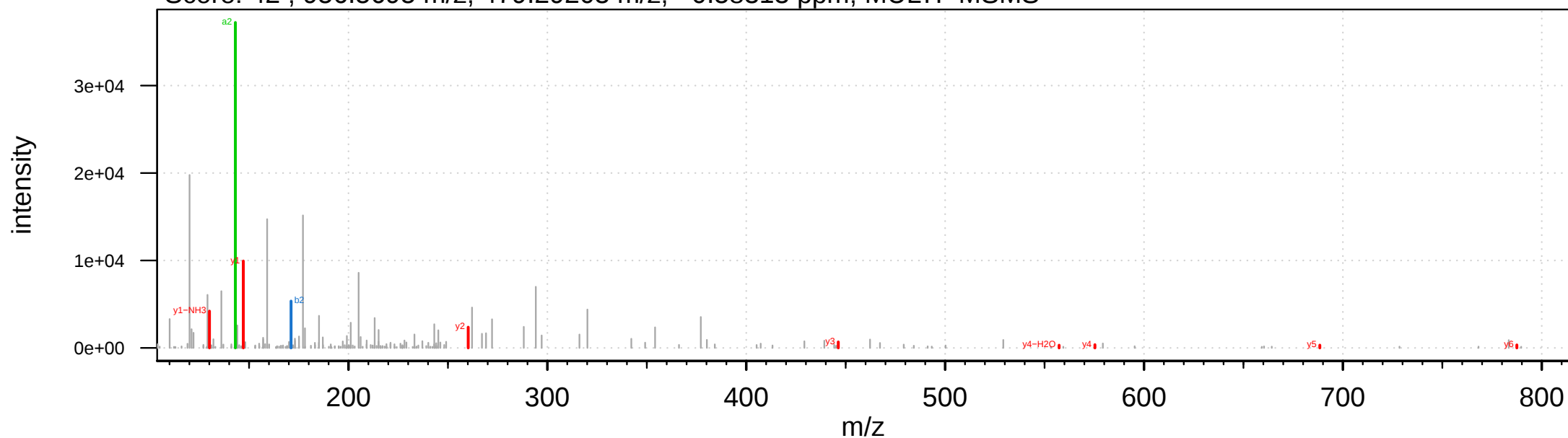


Raw File: 20100618_Velos1_TaGe_SA_LanCap_4
Scan Number: 20191
Proteins:
TCONS_I2_00001296_chr1:79520703-79520992:-



GLVLEWIK

Score: 42 ; 956.5695 m/z; 479.29203 m/z; -0.58313 ppm; MULTI-MSMS

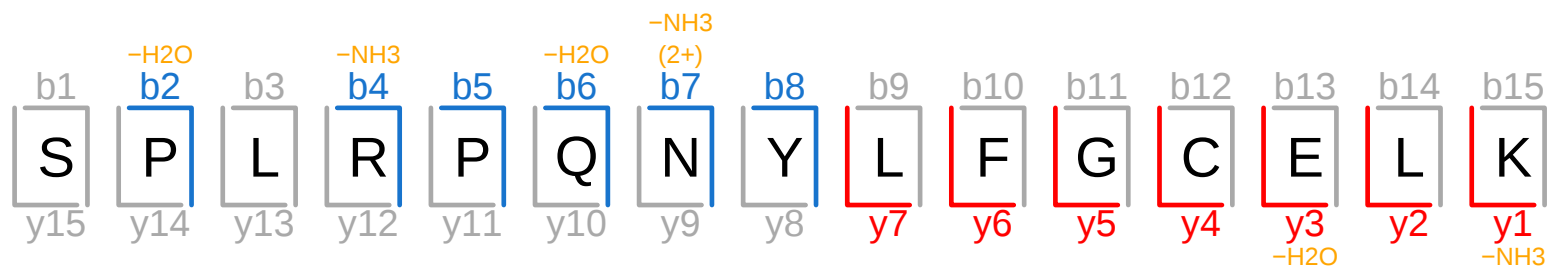


Raw File: 20100726_Velos1_TaGe_SA_HepG2_2

Scan Number: 22614

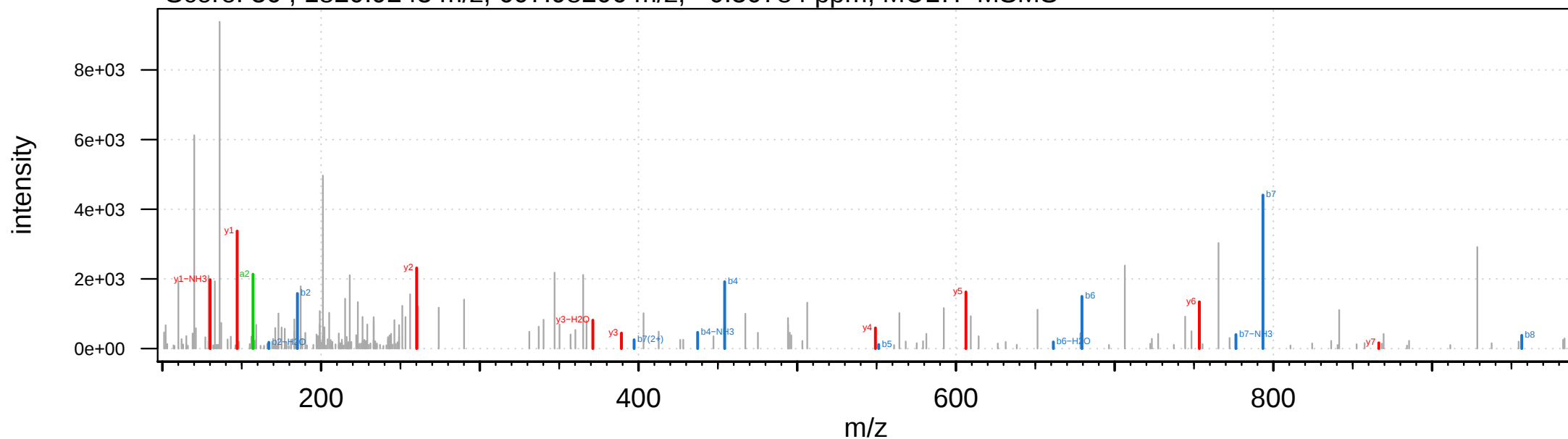
Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-



SPLRPQNYLFGCELK

Score: 36 ; 1820.9243 m/z; 607.98206 m/z; -0.30784 ppm; MULTI-MSMS

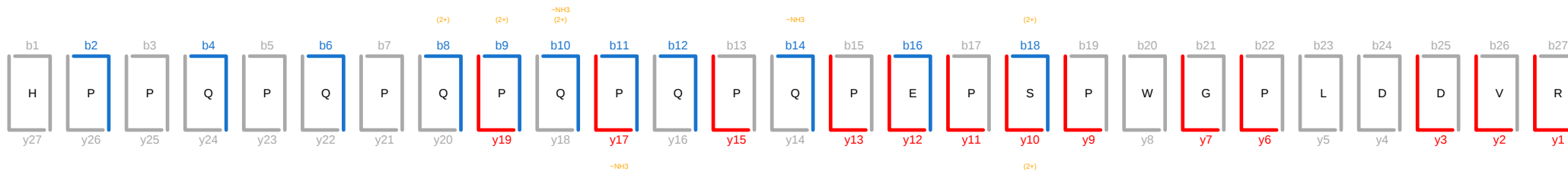


Raw File: 20100726_Velos1_TaGe_SA_HepG2_2

Scan Number: 16923

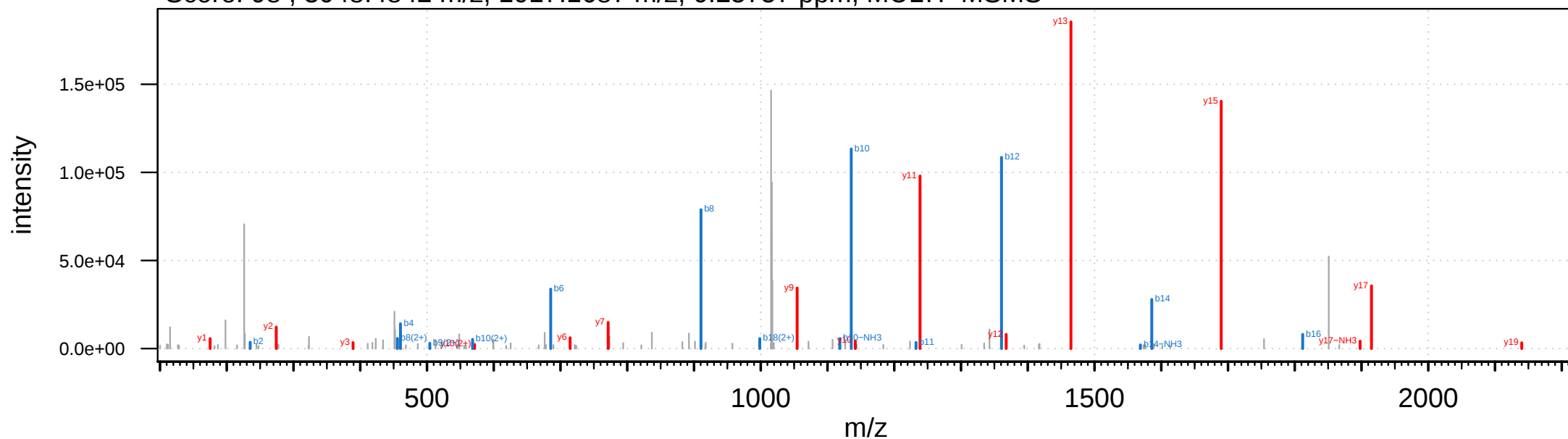
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



HPPQPQPQPQPQPPEPSPWGPLDDVR

Score: 98 ; 3048.4842 m/z; 1017.1687 m/z; 0.25757 ppm; MULTI-MSMS

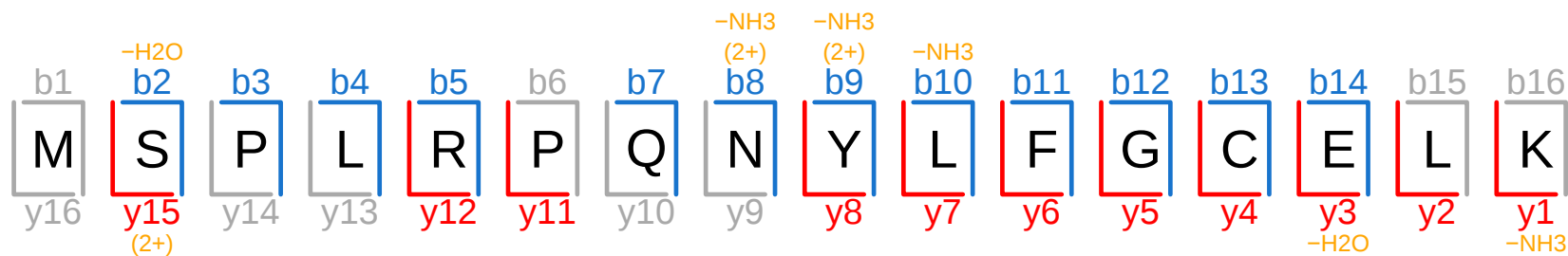


Raw File: 20100609_Velos1_TaGe_SA_GAMG_1

Scan Number: 22989

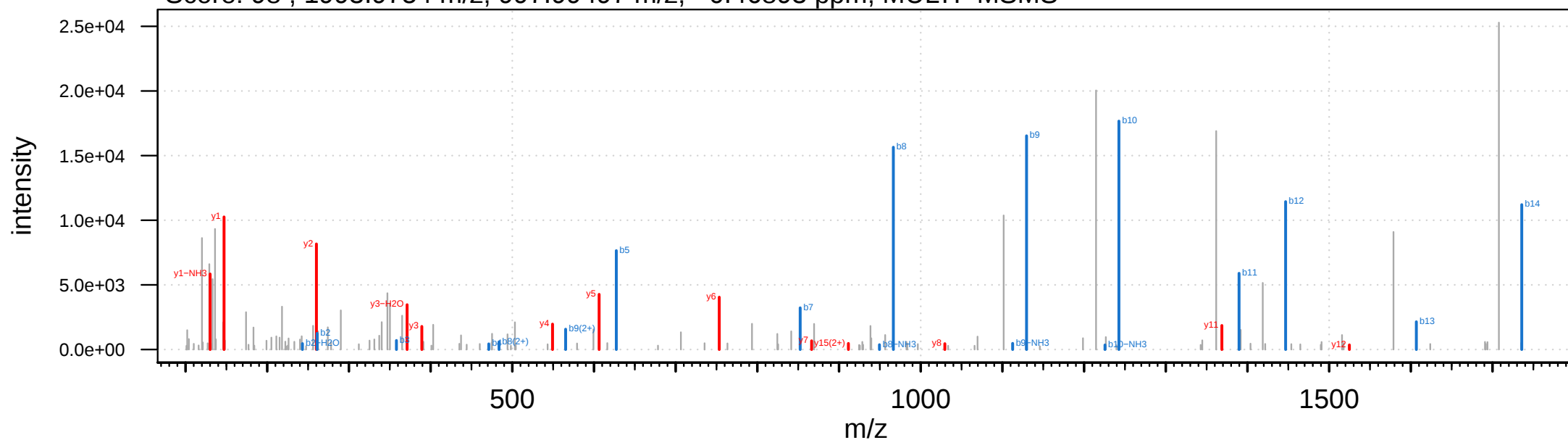
Proteins:

ENST00000424358_chr20:33865517-33865732:-



(ac)MSPLRPQNYLFGCELK

Score: 98 ; 1993.9754 m/z; 997.99497 m/z; -0.46893 ppm; MULTI-MSMS

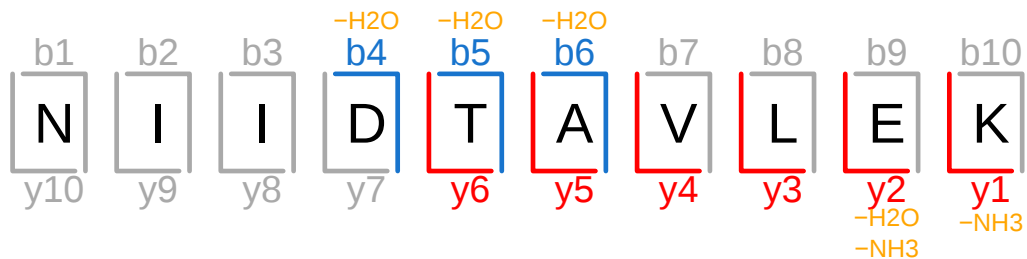


Raw File: 20100609_Velos1_TaGe_SA_GAMG_1

Scan Number: 36079

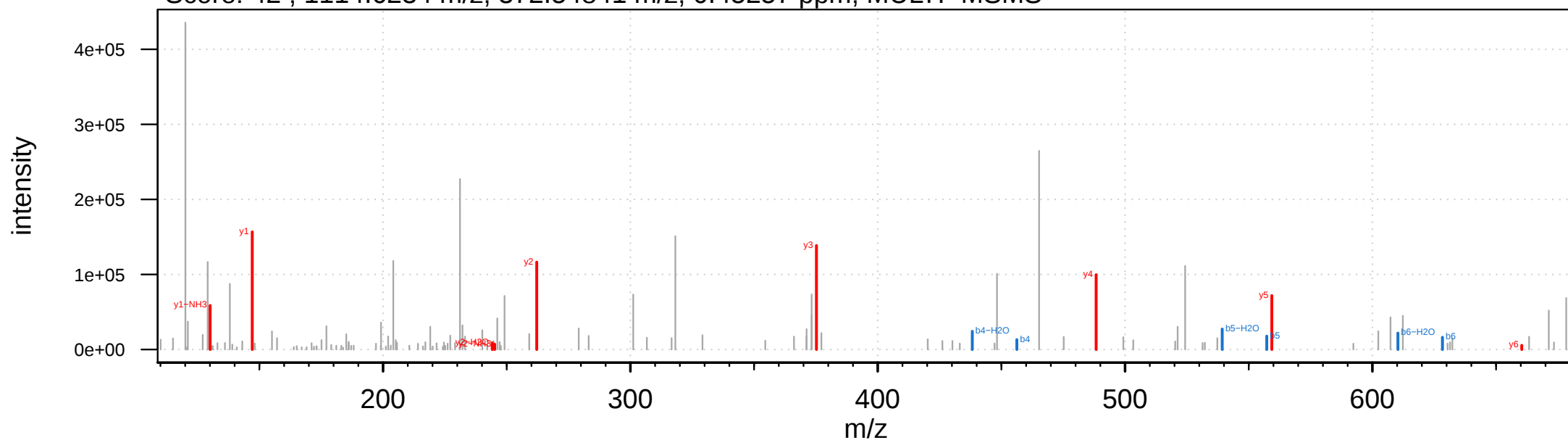
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



NIIDTAVLEK

Score: 42 ; 1114.6234 m/z; 372.54841 m/z; 0.45257 ppm; MULTI-MSMS

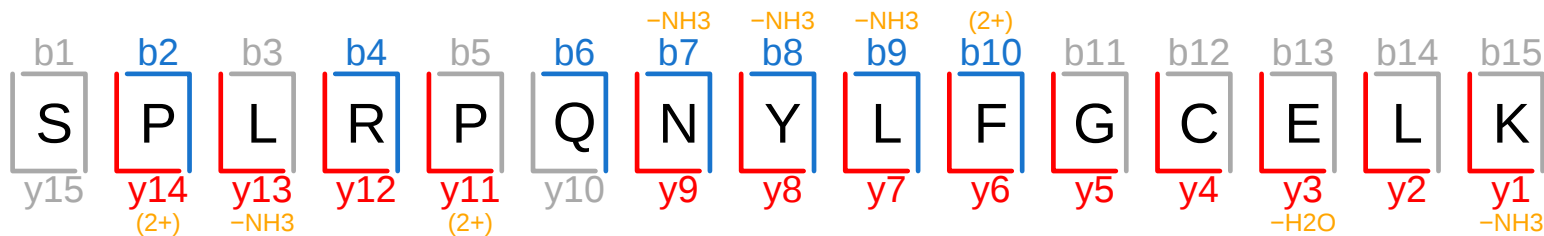


Raw File: 20100609_Velos1_TaGe_SA_GAMG_1

Scan Number: 4709

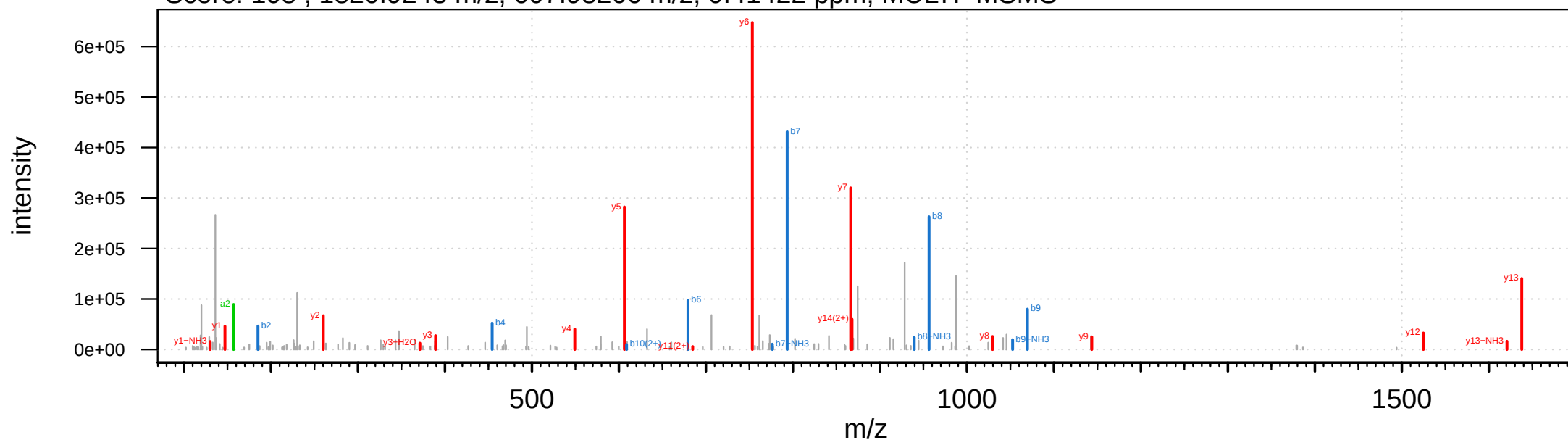
Proteins:

ENST00000497138_chr20:56806826-56807846:-



SPLRPQNYLFGCELK

Score: 108 ; 1820.9243 m/z; 607.98206 m/z; 0.41422 ppm; MULTI-MSMS

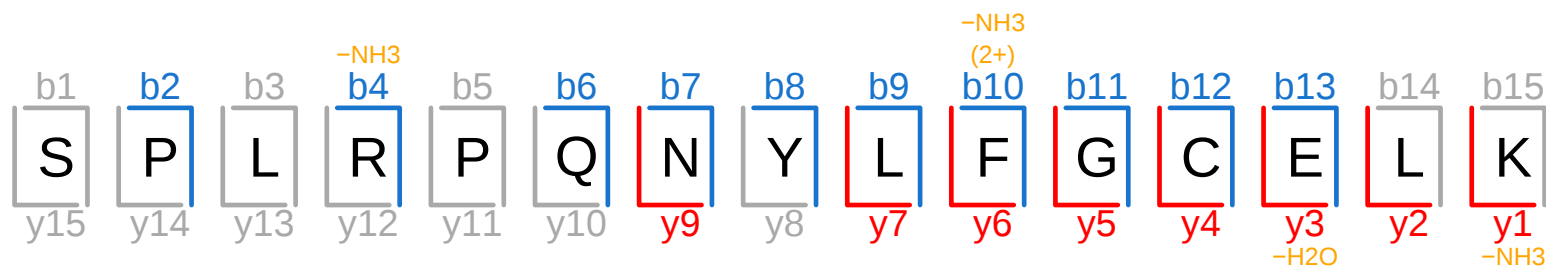


Raw File: 20100609_Velos1_TaGe_SA_GAMG_1

Scan Number: 20632

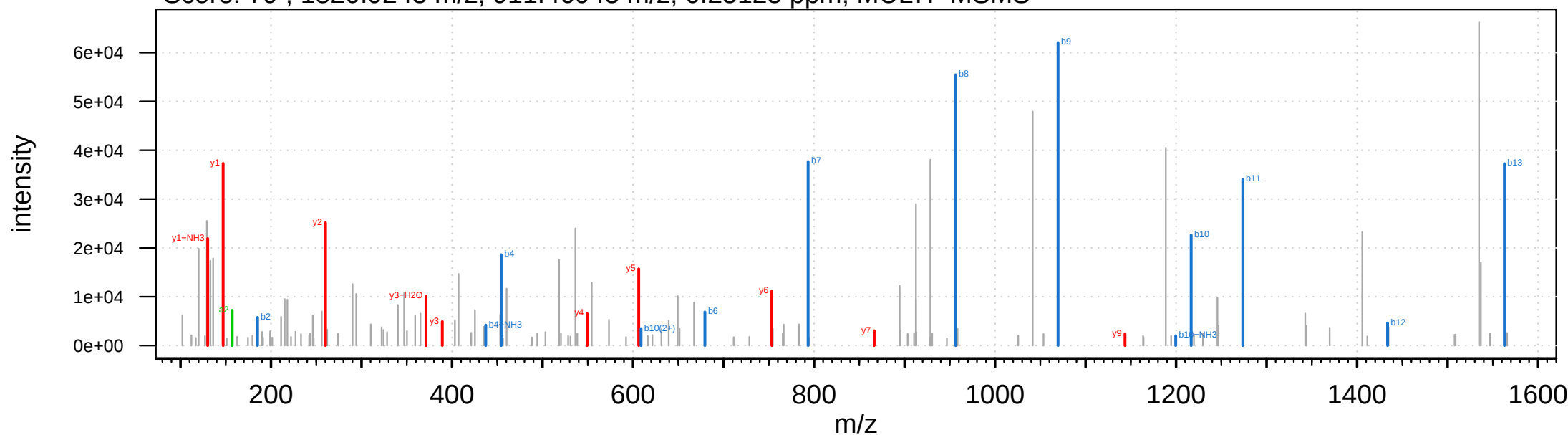
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 79 ; 1820.9243 m/z; 911.46945 m/z; 0.25125 ppm; MULTI-MSMS

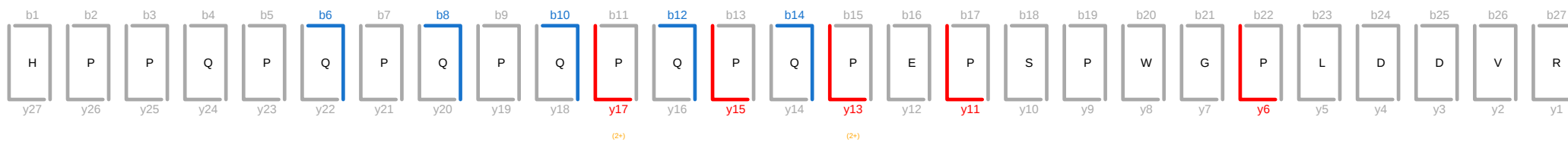


Raw File: 20100609_Velos1_TaGe_SA_GAMG_1

Scan Number: 20645

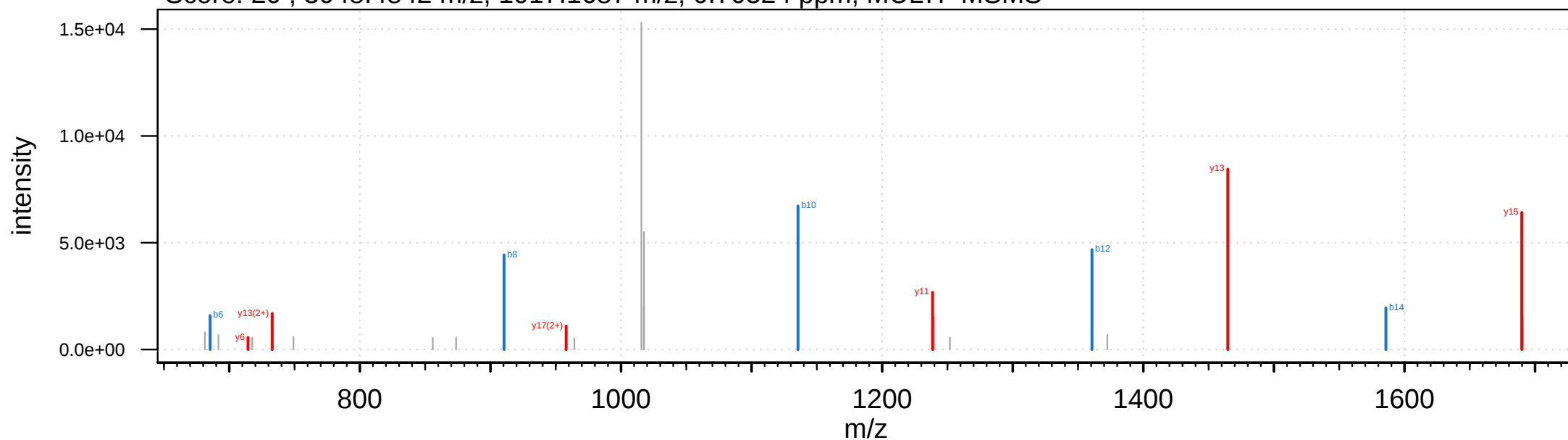
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



HPPQPQPQPQPQPPEPSPWGPLDDVR

Score: 20 ; 3048.4842 m/z; 1017.1687 m/z; 0.70524 ppm; MULTI-MSMS

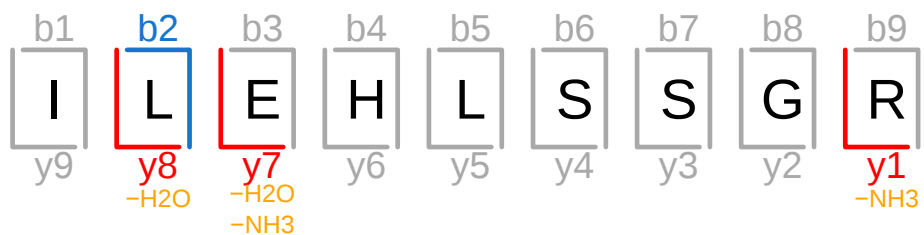


Raw File: 20100618_Velos1_TaGe_SA_LanCap_1

Scan Number: 24132

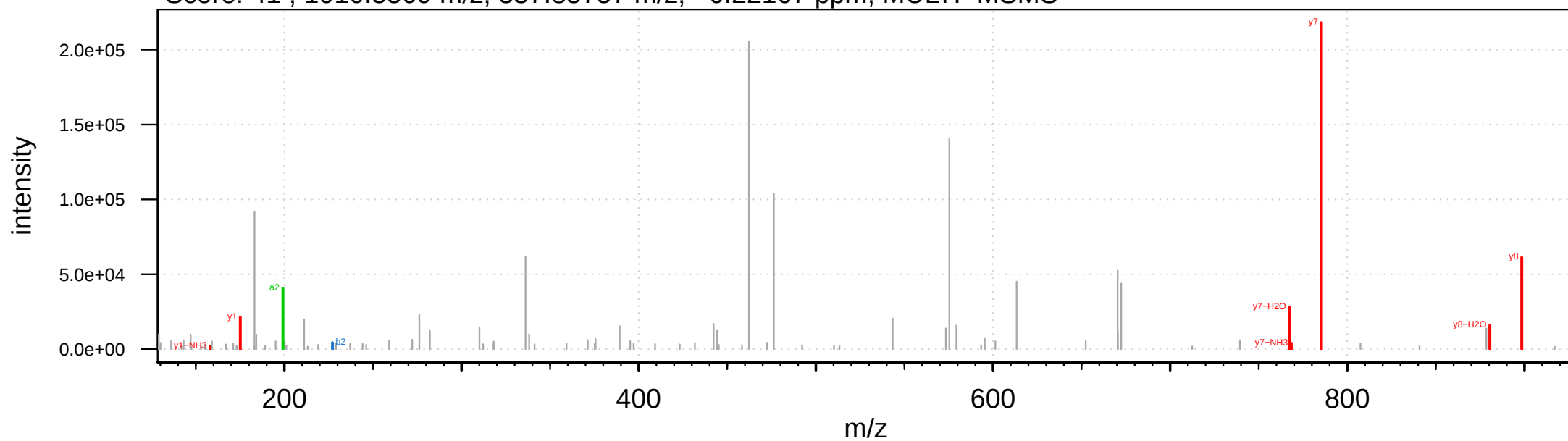
Proteins:

ENST00000424358_chr20:33865517-33865732:-



ILEHLSSGR

Score: 41 ; 1010.5509 m/z; 337.85757 m/z; -0.22167 ppm; MULTI-MSMS

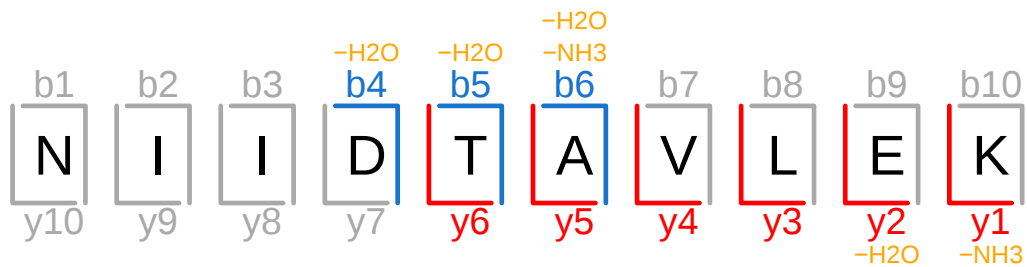


Raw File: 20100618_Velos1_TaGe_SA_LanCap_1

Scan Number: 6018

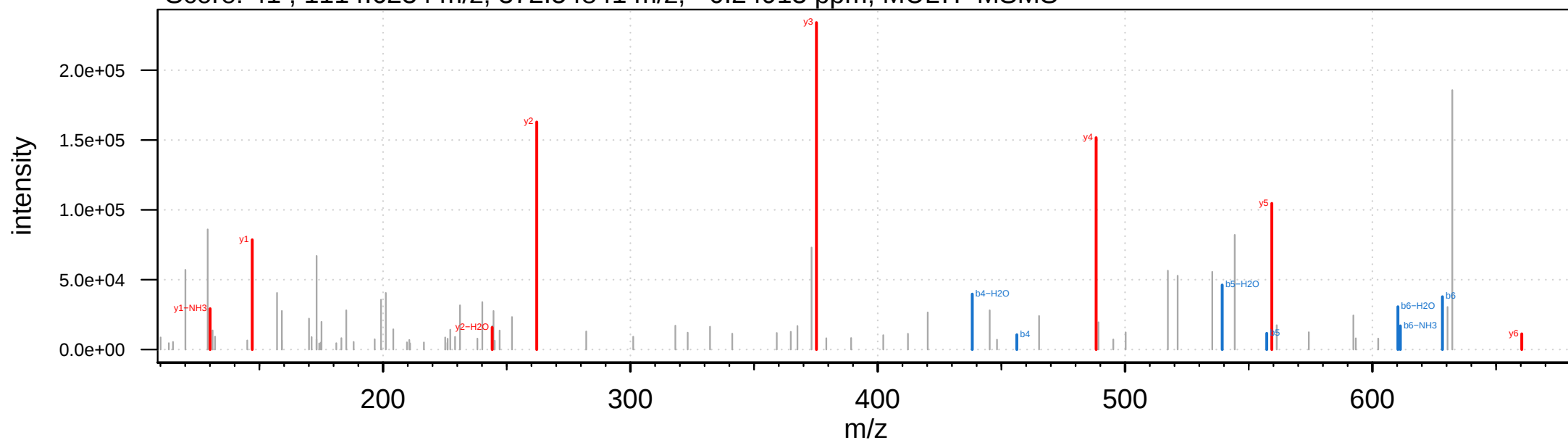
Proteins:

ENST00000452079_chr1:3663088-3663306:-



NIIDTAVLEK

Score: 41 ; 1114.6234 m/z; 372.54841 m/z; -0.24913 ppm; MULTI-MSMS

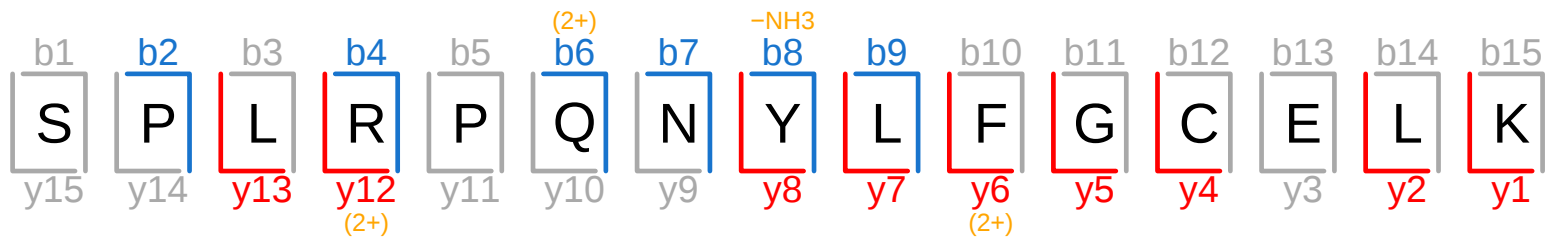


Raw File: 20100618_Velos1_TaGe_SA_LanCap_1

Scan Number: 5368

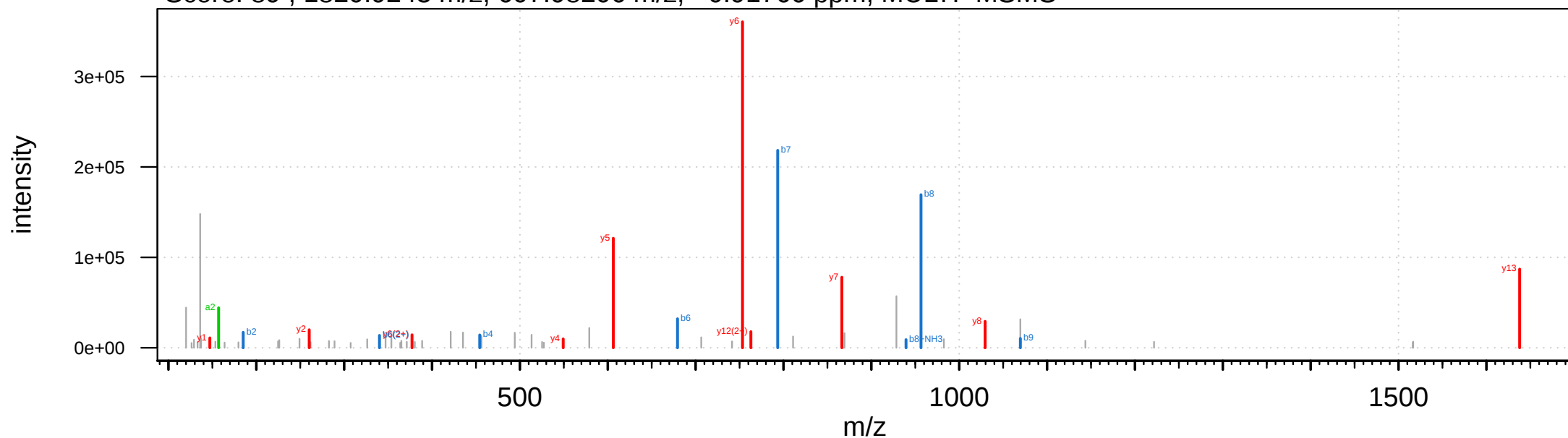
Proteins:

ENST00000497138_chr20:56806826-56807846:-

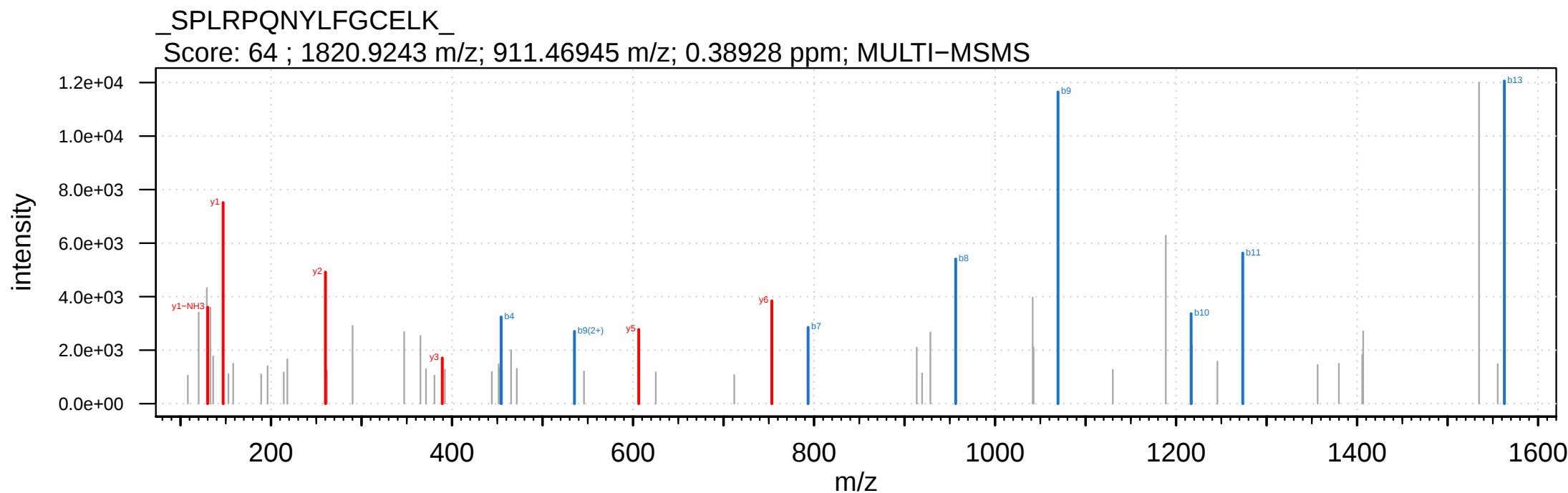
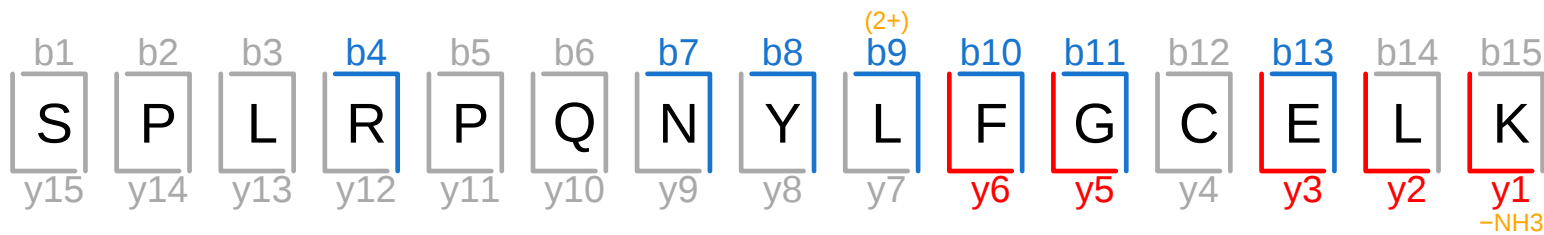


SPLRPQNYLFGCELK

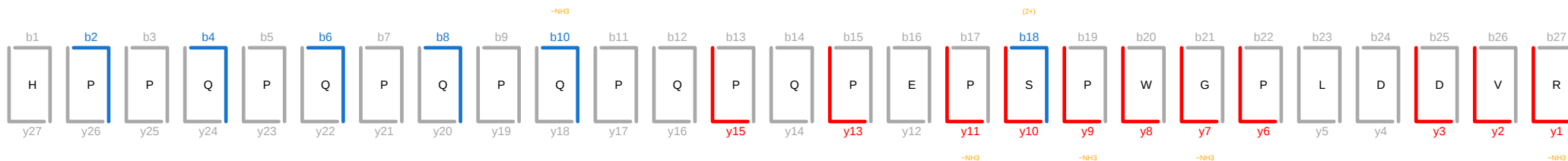
Score: 89 ; 1820.9243 m/z; 607.98206 m/z; -0.91766 ppm; MULTI-MSMS



Raw File: 20100618_Velos1_TaGe_SA_LanCap_1
 Scan Number: 22913
 Proteins:
 TCONS_I2_00008829_ch15:92829088-92829258:+

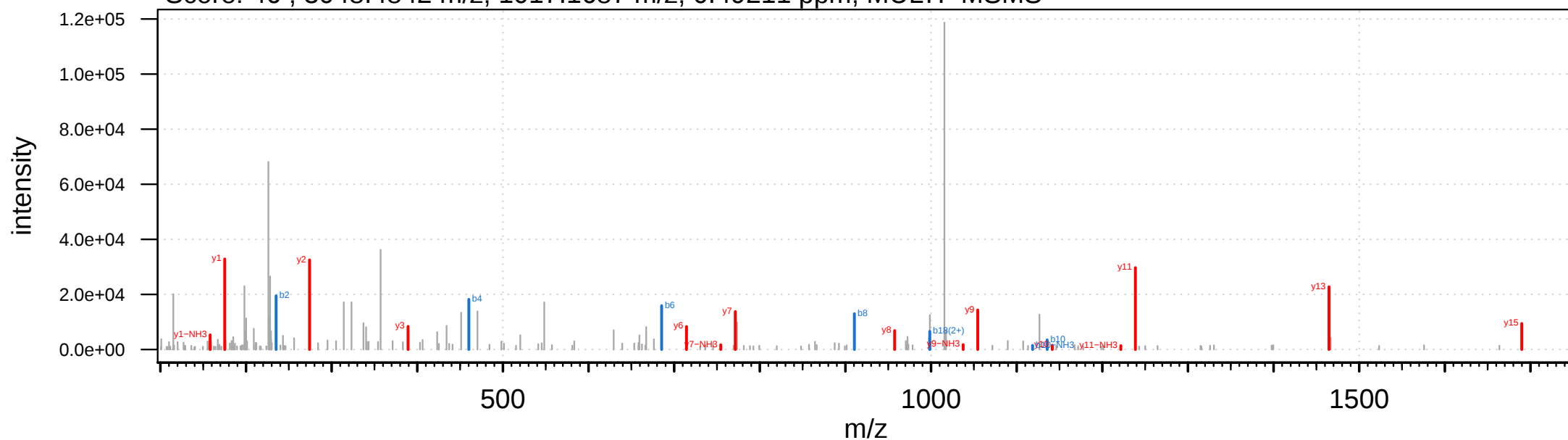


Raw File: 20100618_Velos1_TaGe_SA_LanCap_1
 Scan Number: 22917
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

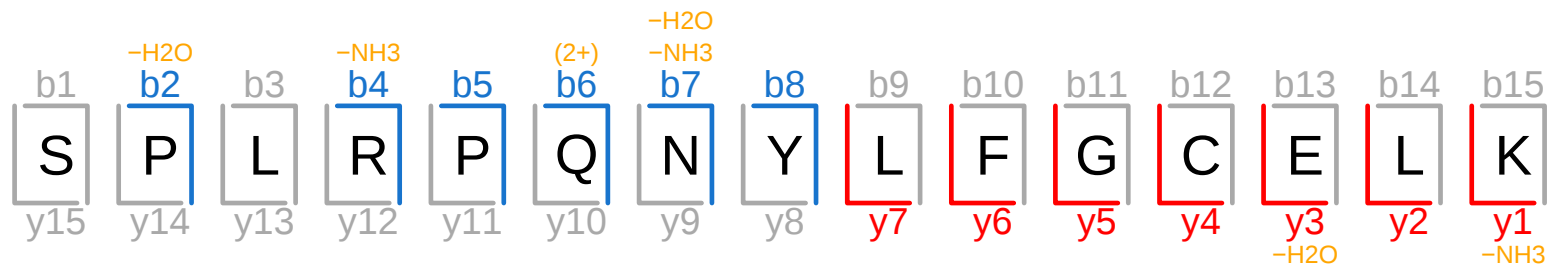


HPPQPQPQPQPQPPEPSPWGPLDDVR

Score: 46 ; 3048.4842 m/z; 1017.1687 m/z; 0.49211 ppm; MULTI-MSMS

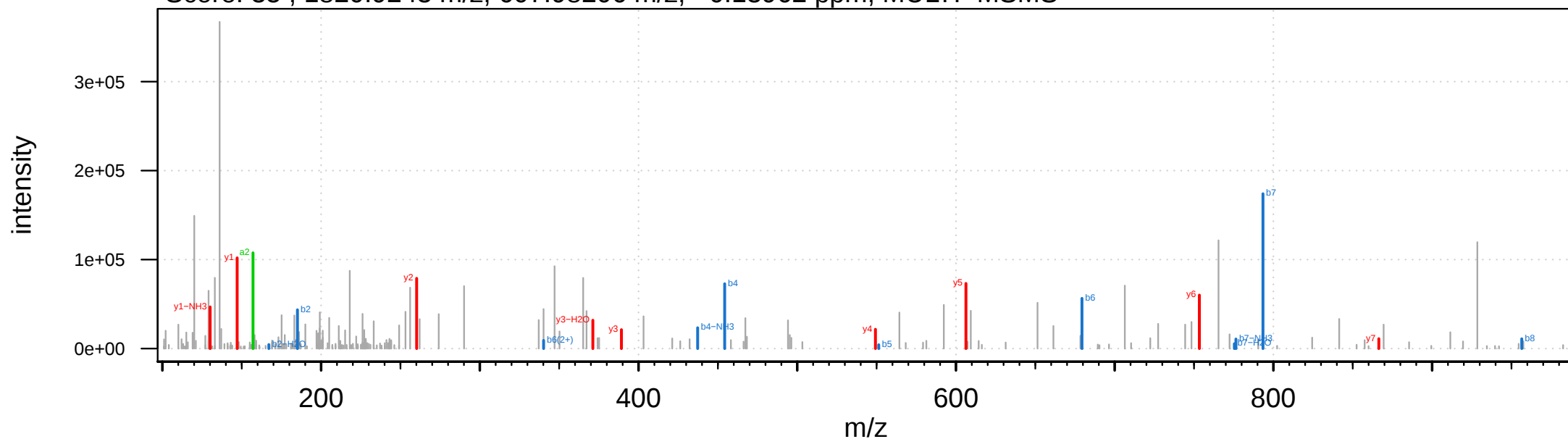


Raw File: 20100719_Velos1_TaGe_SA_LnCap_1
Scan Number: 18918
Proteins:
ENST00000424358_chr20:33865517-33865732:-

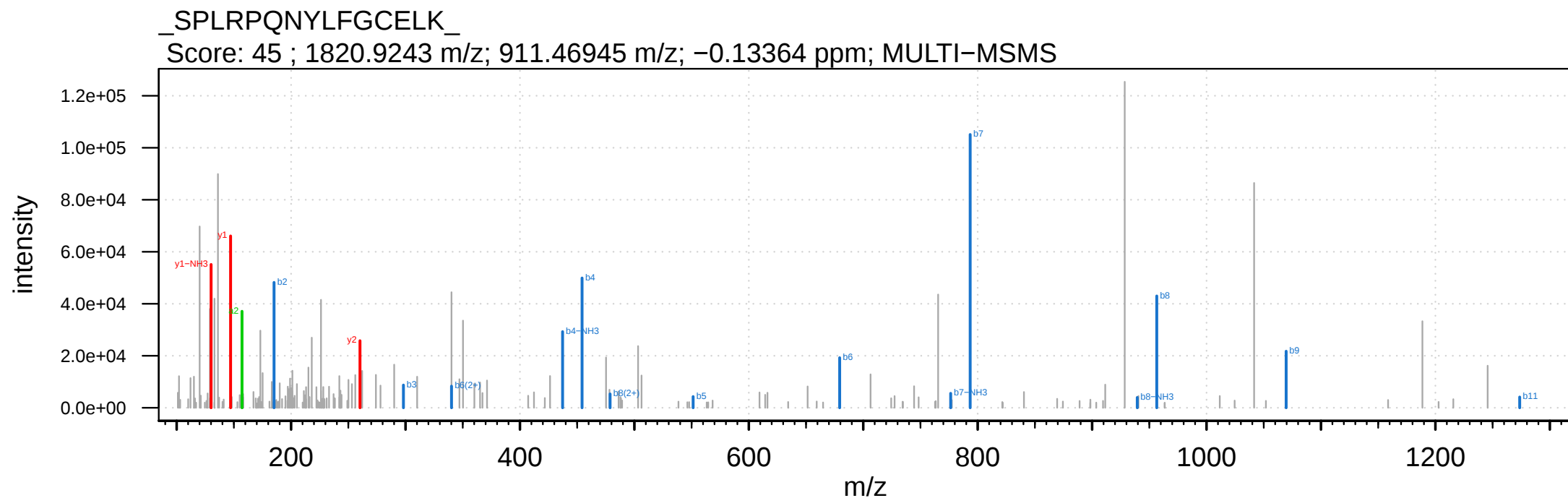
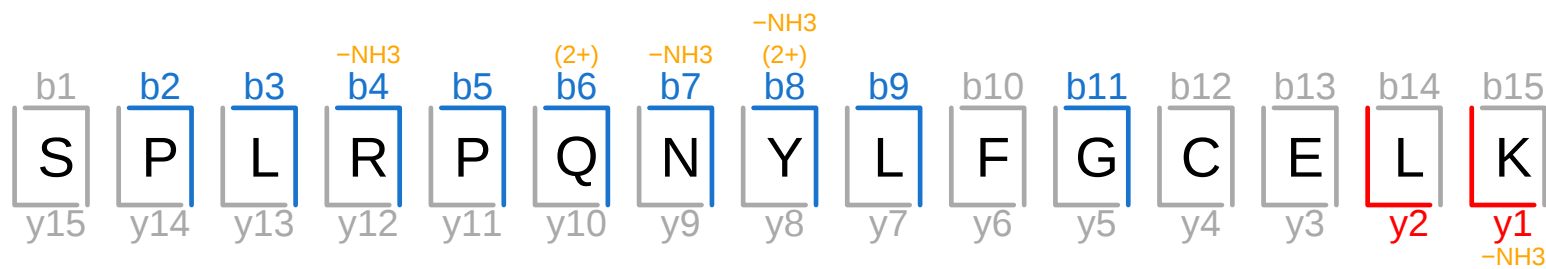


SPLRPQNYLFGCELK

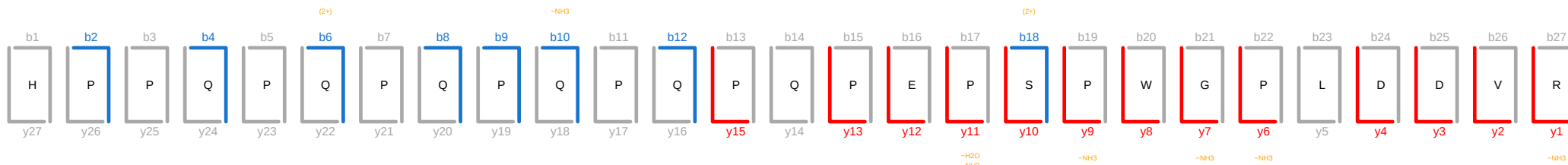
Score: 35 ; 1820.9243 m/z; 607.98206 m/z; -0.13962 ppm; MULTI-MSMS



Raw File: 20100719_Velos1_TaGe_SA_LnCap_1
 Scan Number: 18162
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

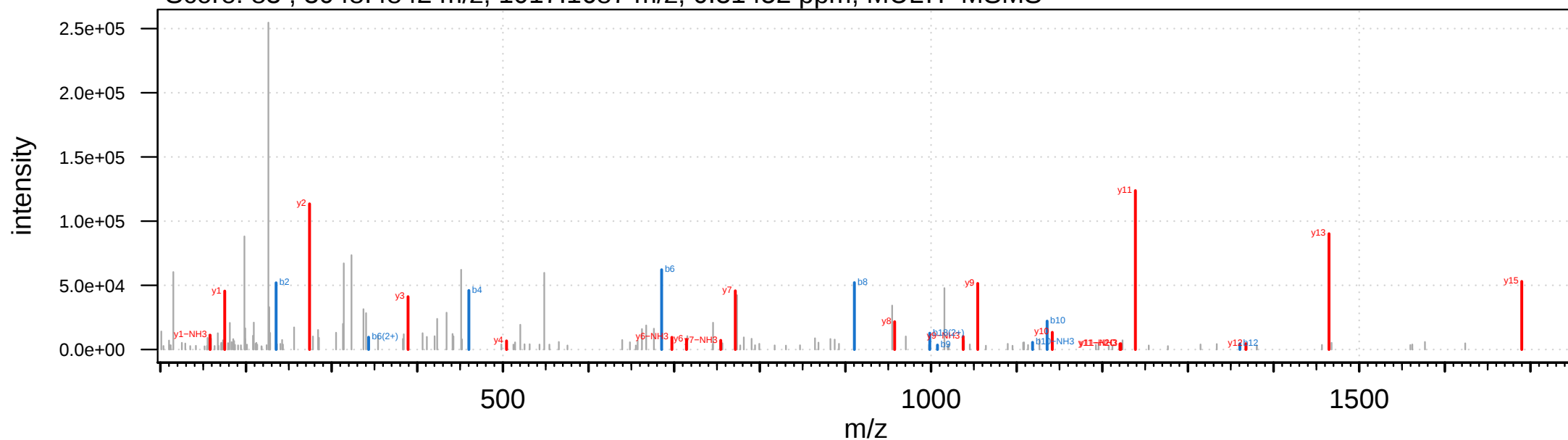


Raw File: 20100719_Velos1_TaGe_SA_LnCap_1
 Scan Number: 18173
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



HPPQPQPQPQPQPPEPSPWGPLDDVR

Score: 83 ; 3048.4842 m/z; 1017.1687 m/z; 0.31452 ppm; MULTI-MSMS

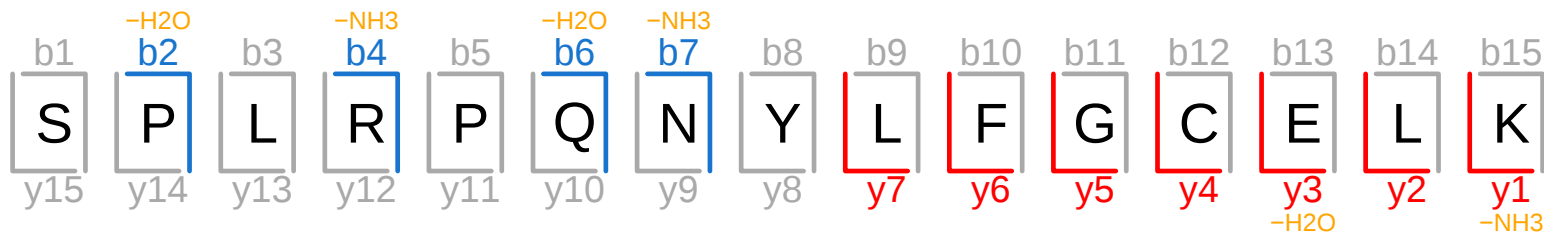


Raw File: 20100723_Velos1_TaGe_SA_Gamg_1

Scan Number: 19123

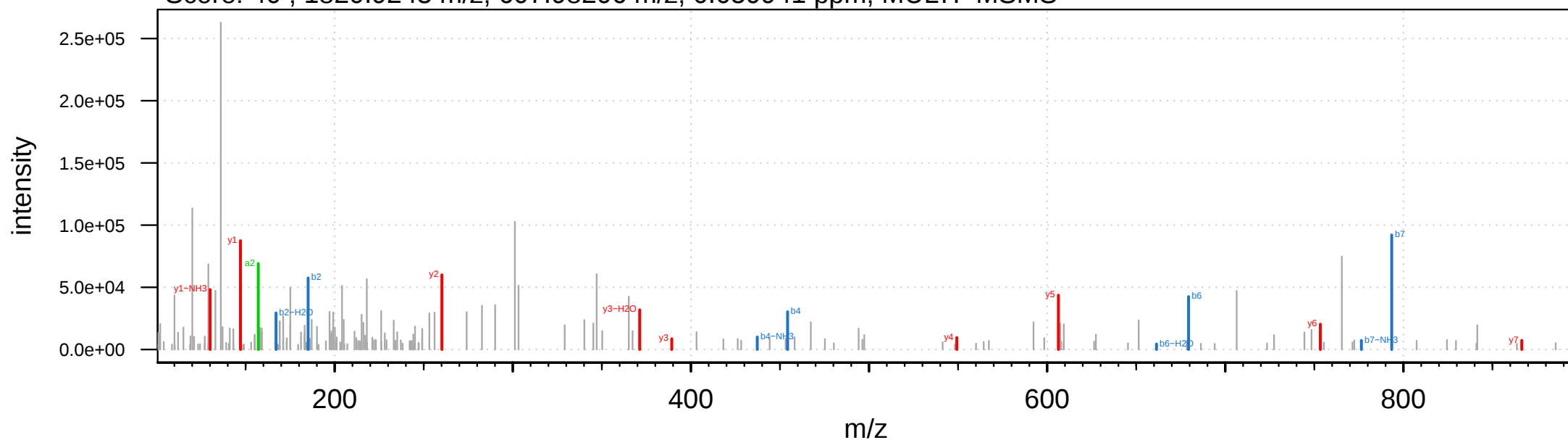
Proteins:

ENST00000424358_chr20:33865517-33865732:-



SPLRPQNYLFGCELK

Score: 40 ; 1820.9243 m/z; 607.98206 m/z; 0.059941 ppm; MULTI-MSMS

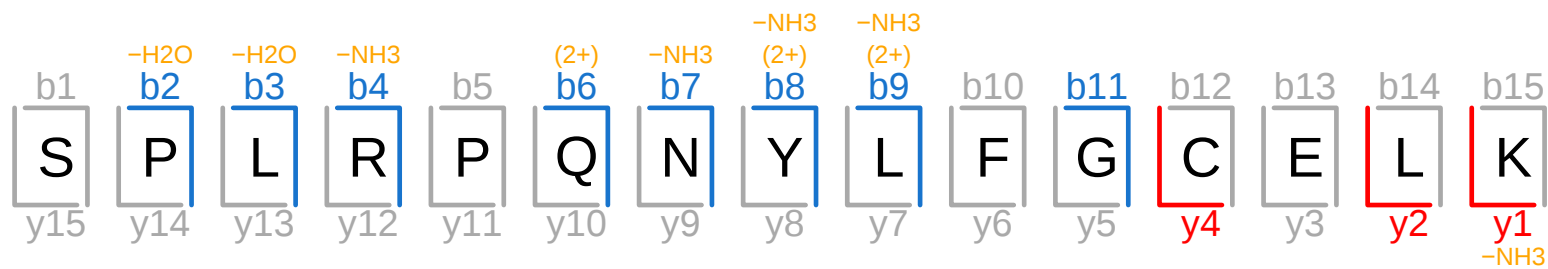


Raw File: 20100723_Velos1_TaGe_SA_Gamg_1

Scan Number: 18354

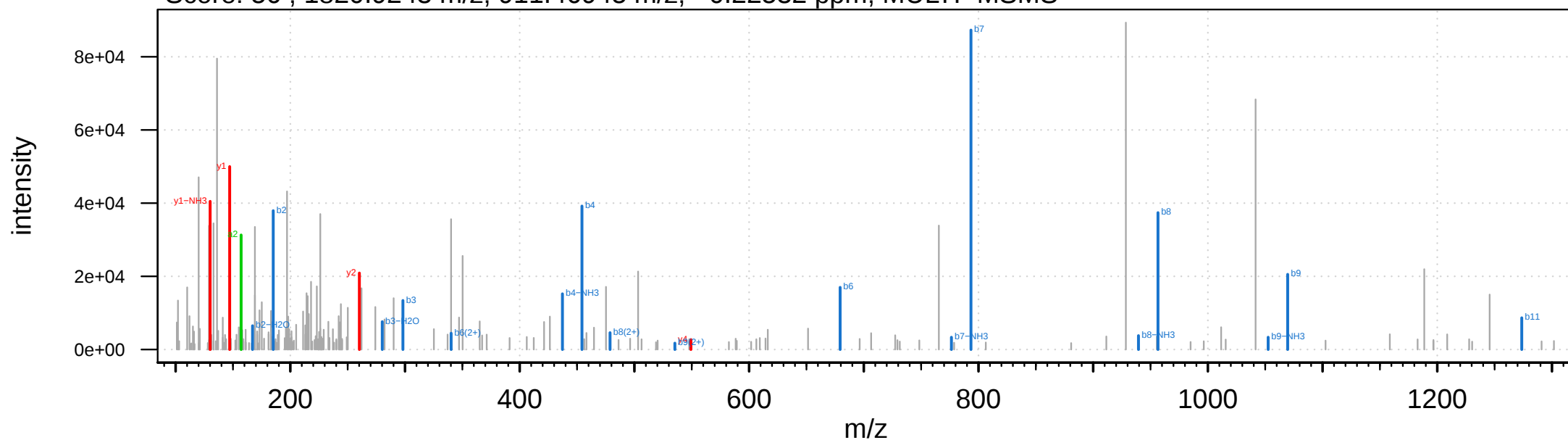
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 56 ; 1820.9243 m/z; 911.46945 m/z; -0.22532 ppm; MULTI-MSMS

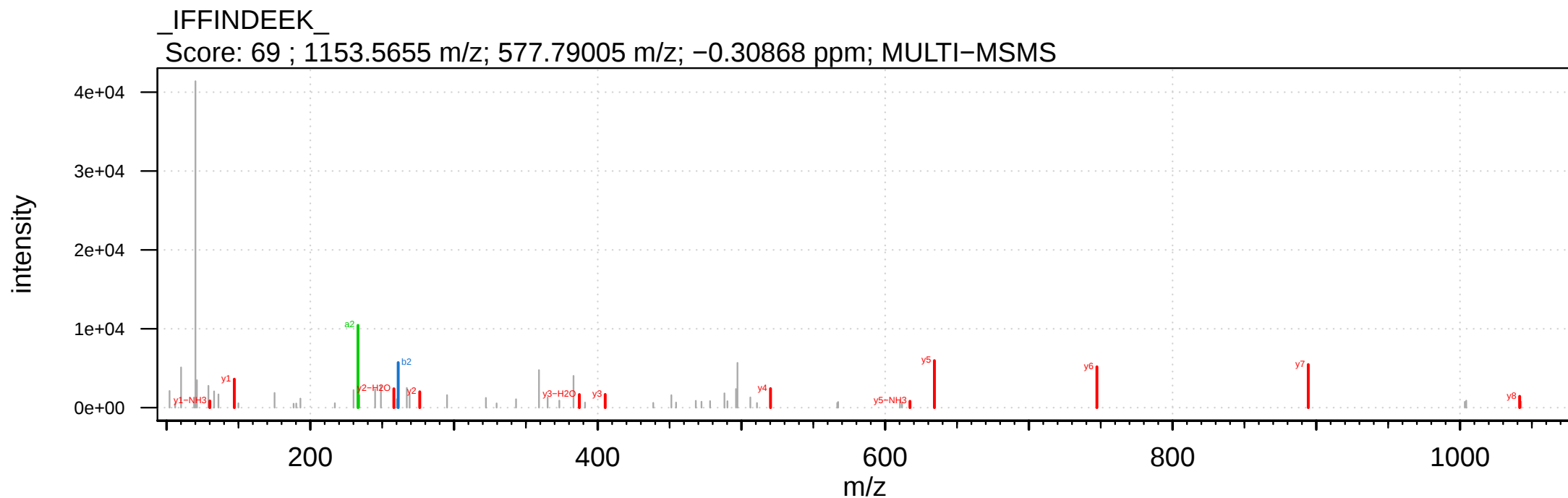
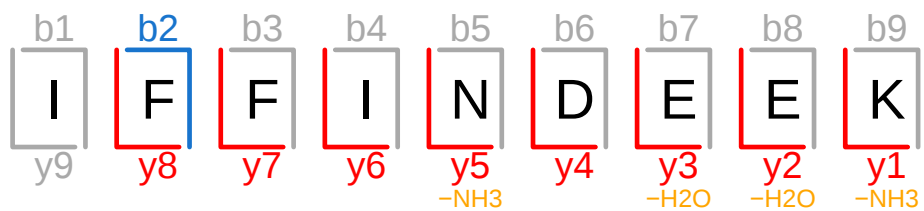


Raw File: 20100723_Velos1_TaGe_SA_Gamg_1

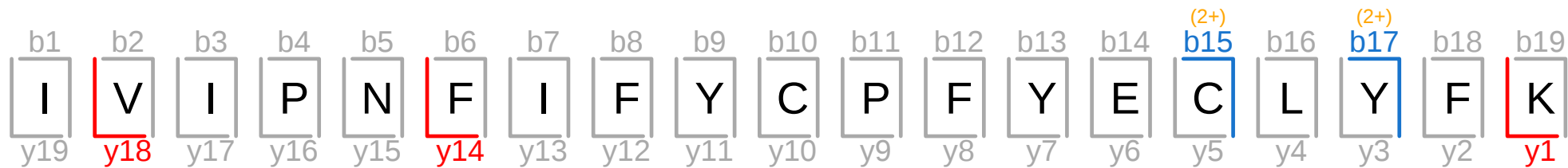
Scan Number: 18363

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

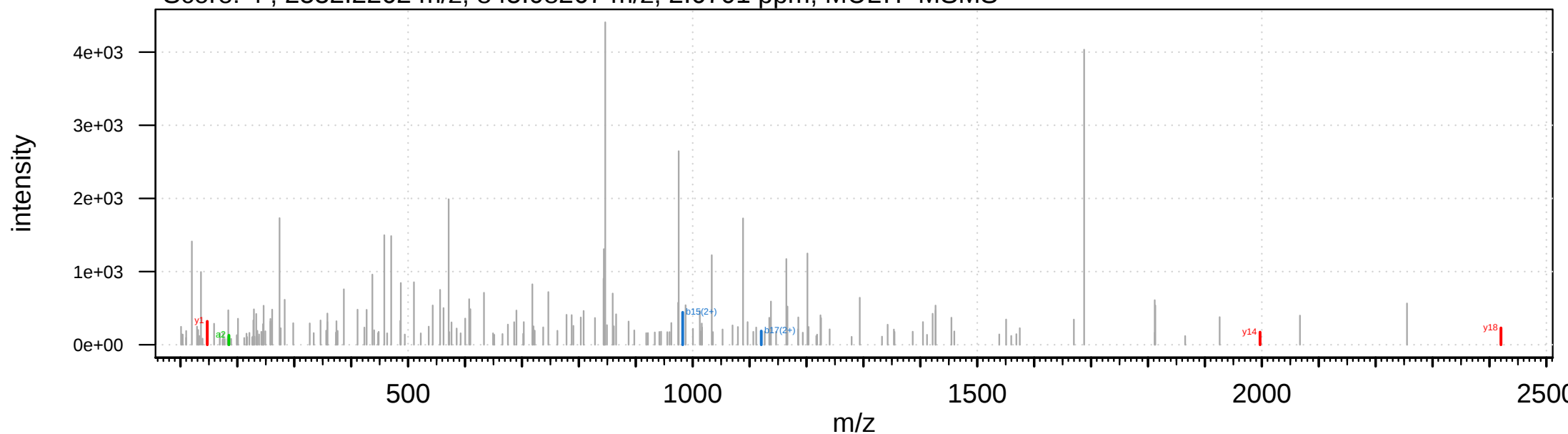


Raw File: 20100604_Velos1_TaGe_SA_A549_5
Scan Number: 11134
Proteins:
TCONS_I2_00001718_chr1:204336923-204346793:-



IVIPNFIFYCPFYECLYFK

Score: 4 ; 2532.2262 m/z; 845.08267 m/z; 2.6701 ppm; MULTI-MSMS

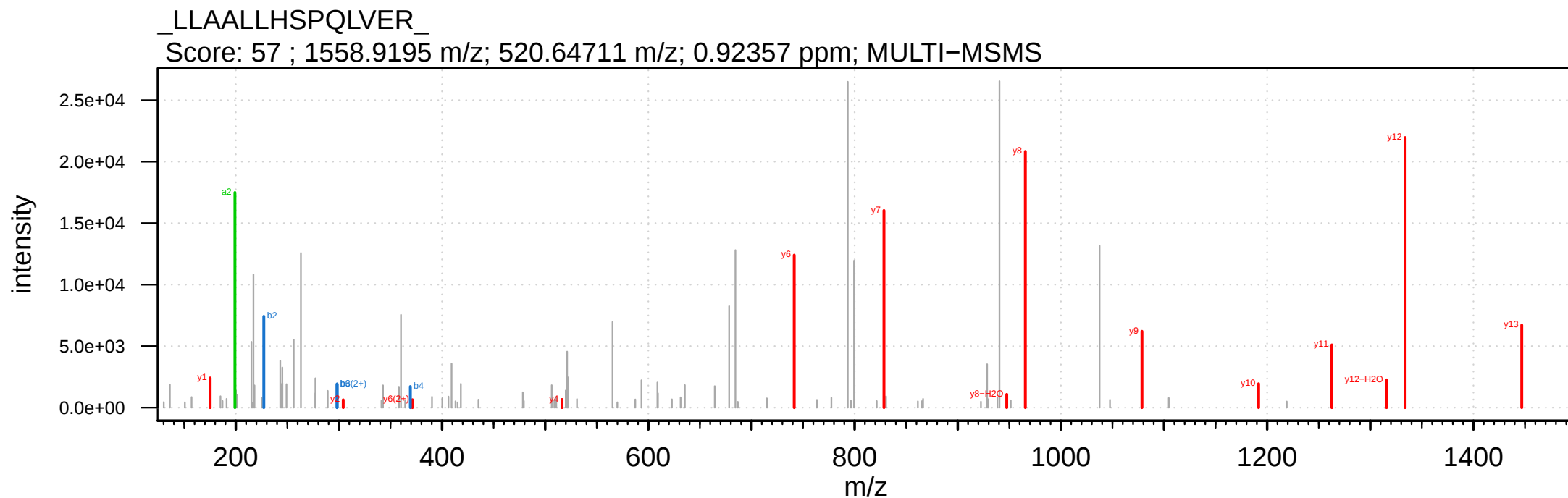
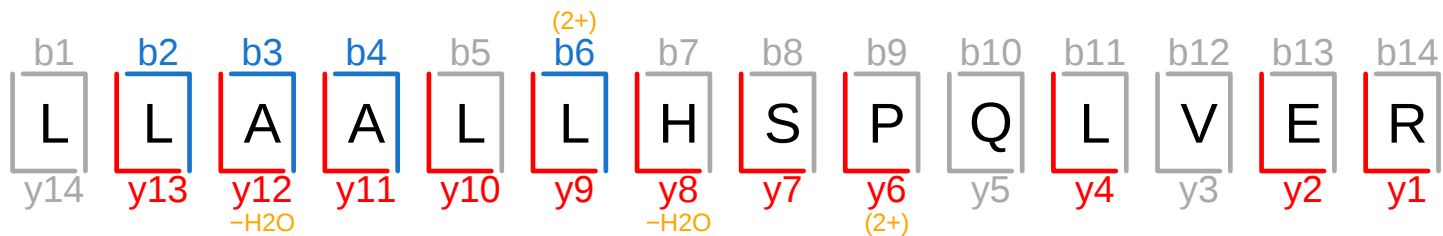


Raw File: 20101227_Velos1_TaGe_SA_GAMG6

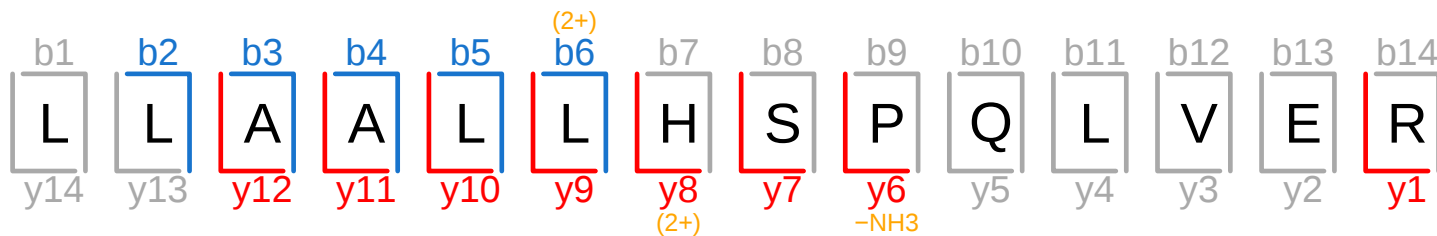
Scan Number: 35131

Proteins:

ENST00000291592_chr21:45878569-45878637:+

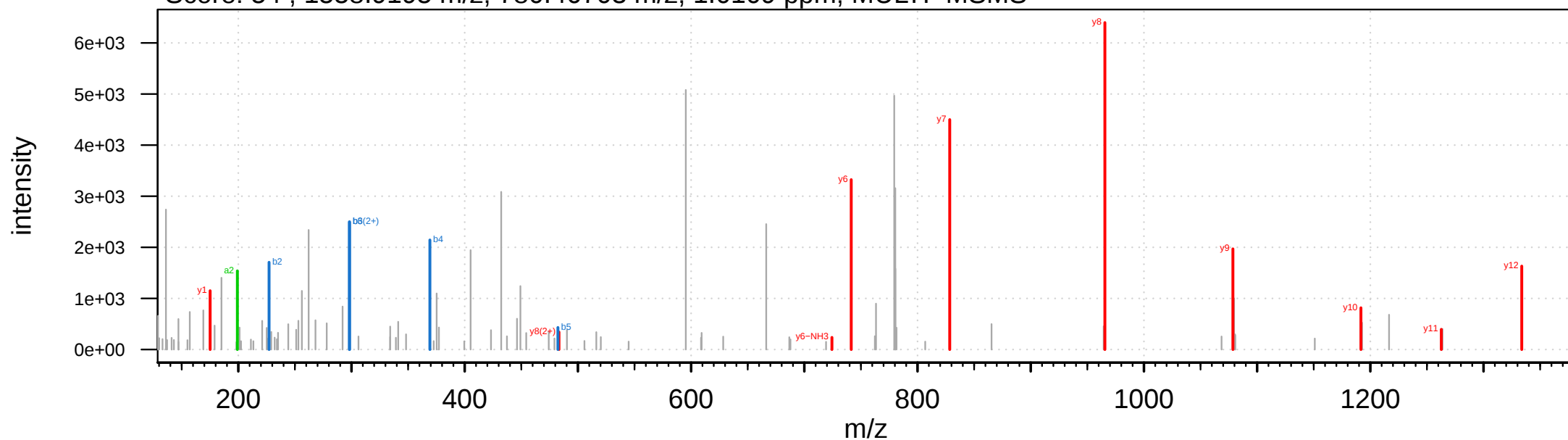


Raw File: 20100616_Velos1_TaGe_SA_MCF7_3
Scan Number: 23325
Proteins:
ENST00000602845_chr3:196669588-196669887:+



LLAALLHSPQLVER

Score: 54 ; 1558.9195 m/z; 780.46703 m/z; 1.0109 ppm; MULTI-MSMS

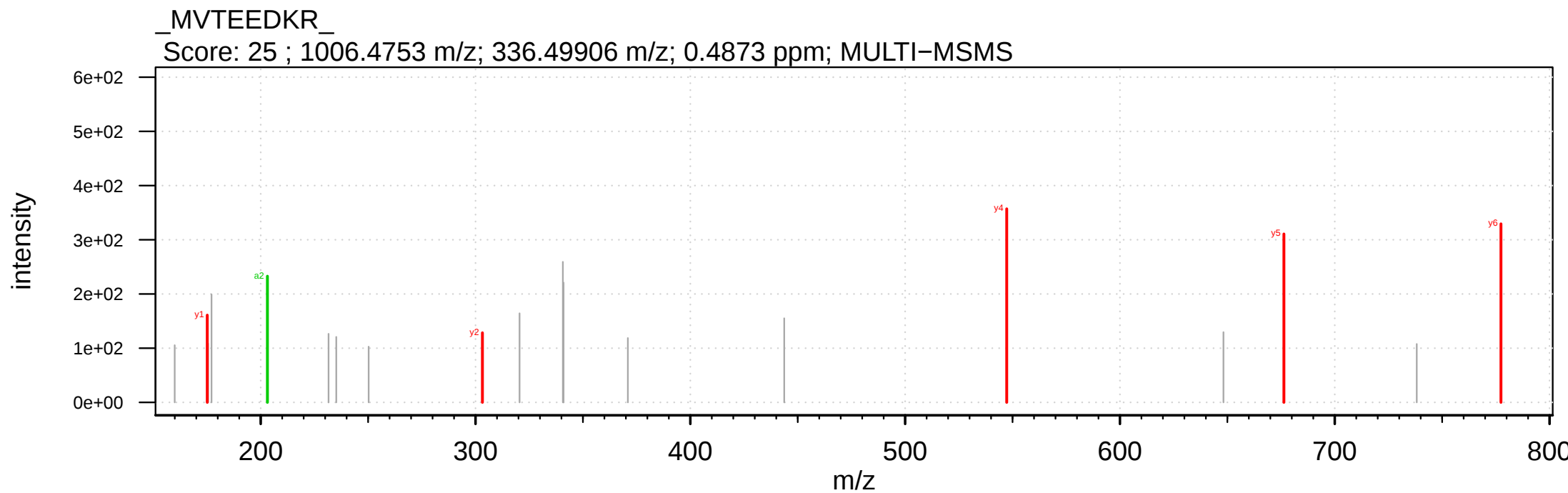
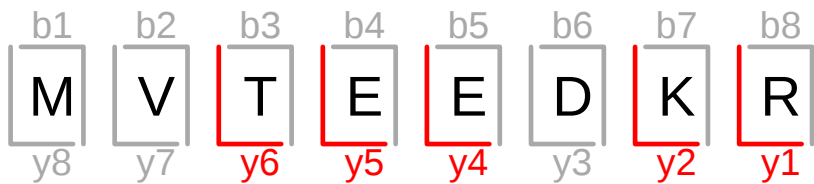


Raw File: 20100616_Velos1_TaGe_SA_MCF7_3

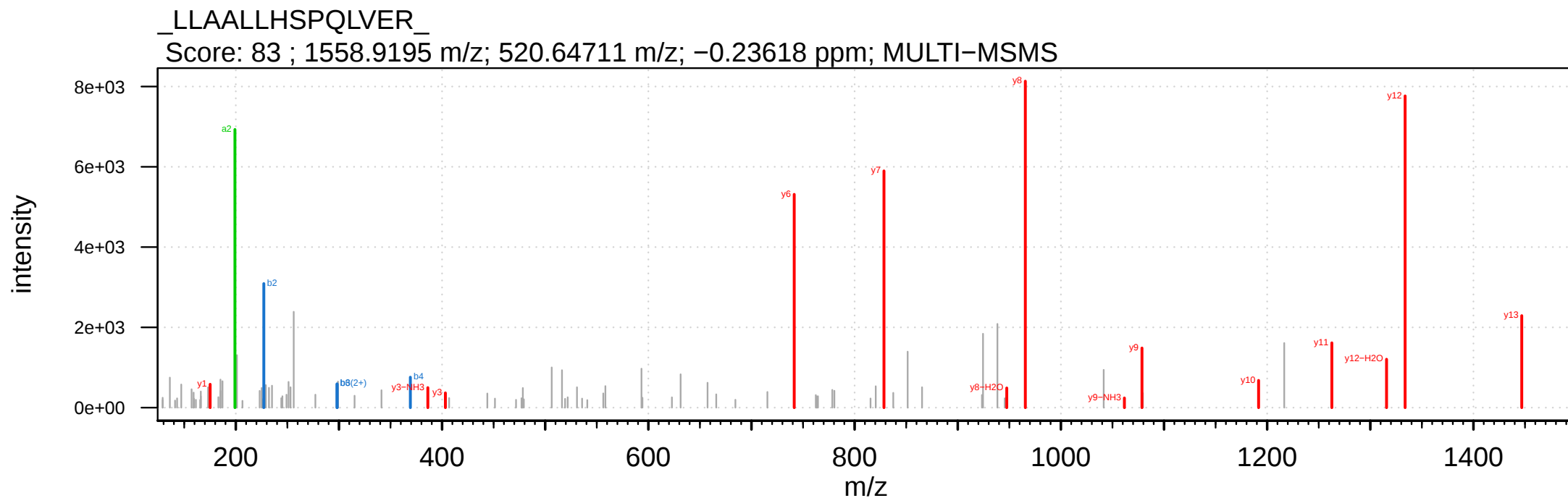
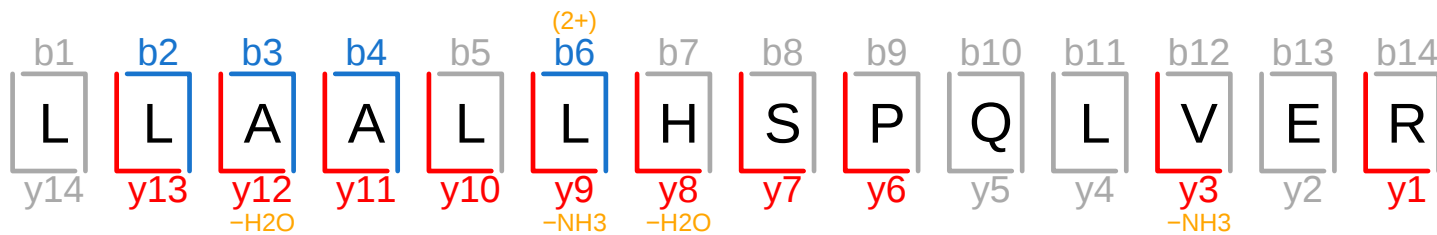
Scan Number: 23362

Proteins:

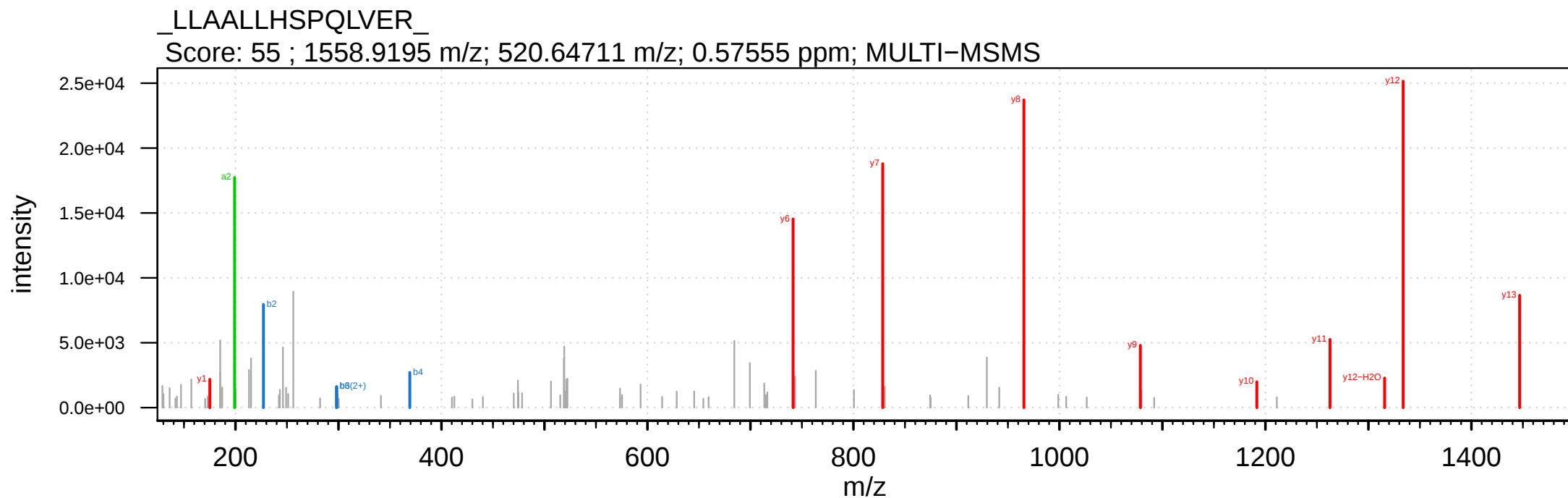
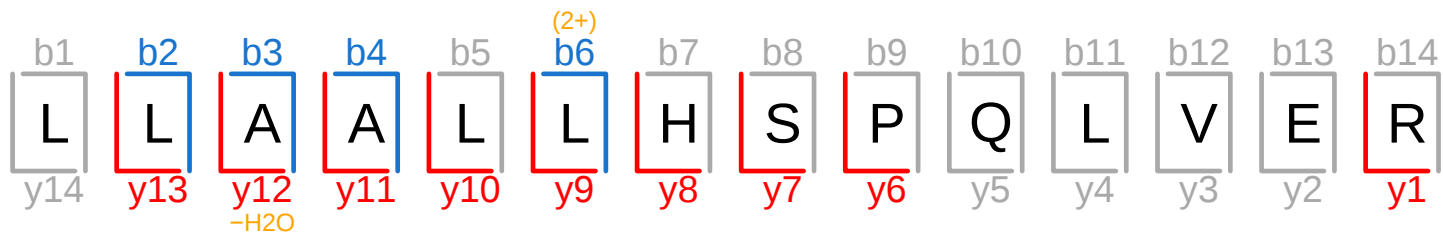
ENST00000602845_chr3:196669588-196669887:+



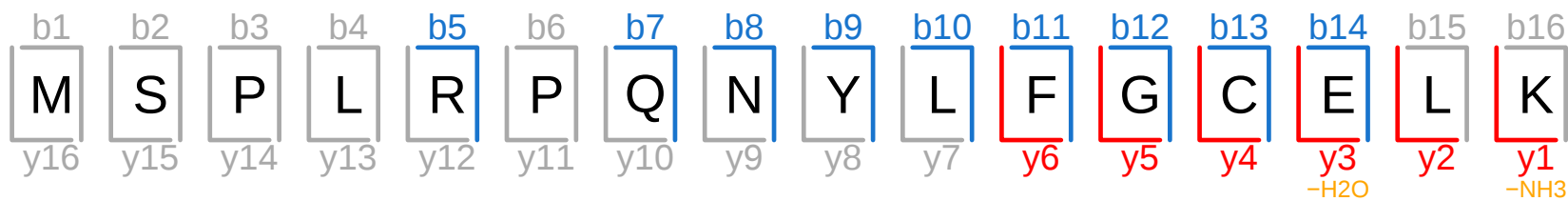
Raw File: 20100616_Velos1_TaGe_SA_MCF7_3
Scan Number: 1594
Proteins:
TCONS_I2_00006768_chr13:29174091-29174123:+



Raw File: 20100616_Velos1_TaGe_SA_MCF7_4
 Scan Number: 19997
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+

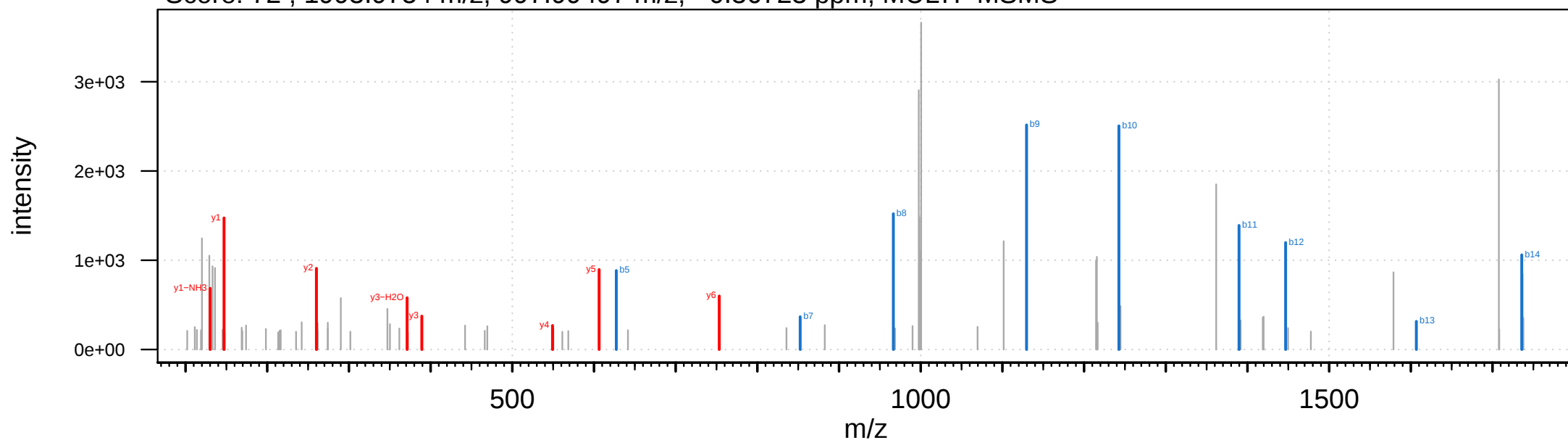


Raw File: 20100616_Velos1_TaGe_SA_RKO_1
Scan Number: 27484
Proteins:
ENST00000602845_chr3:196669588-196669887:+



(ac)MSPLRPQNYLFGCELK

Score: 72 ; 1993.9754 m/z; 997.99497 m/z; -0.36723 ppm; MULTI-MSMS

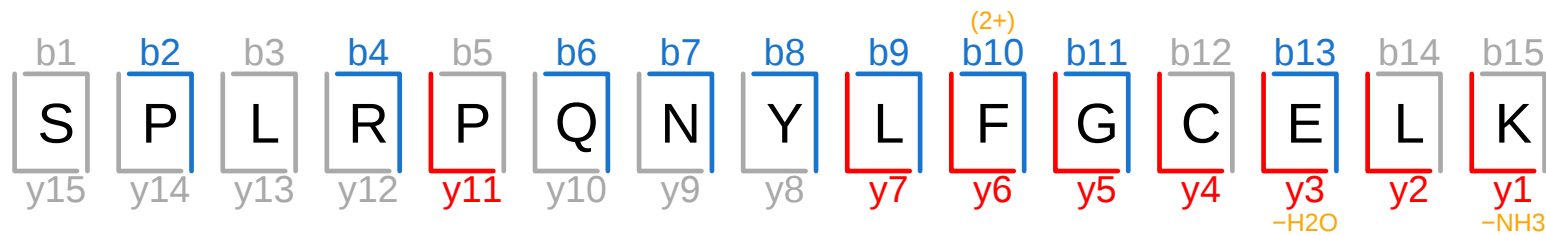


Raw File: 20100616_Velos1_TaGe_SA_RKO_1

Scan Number: 36901

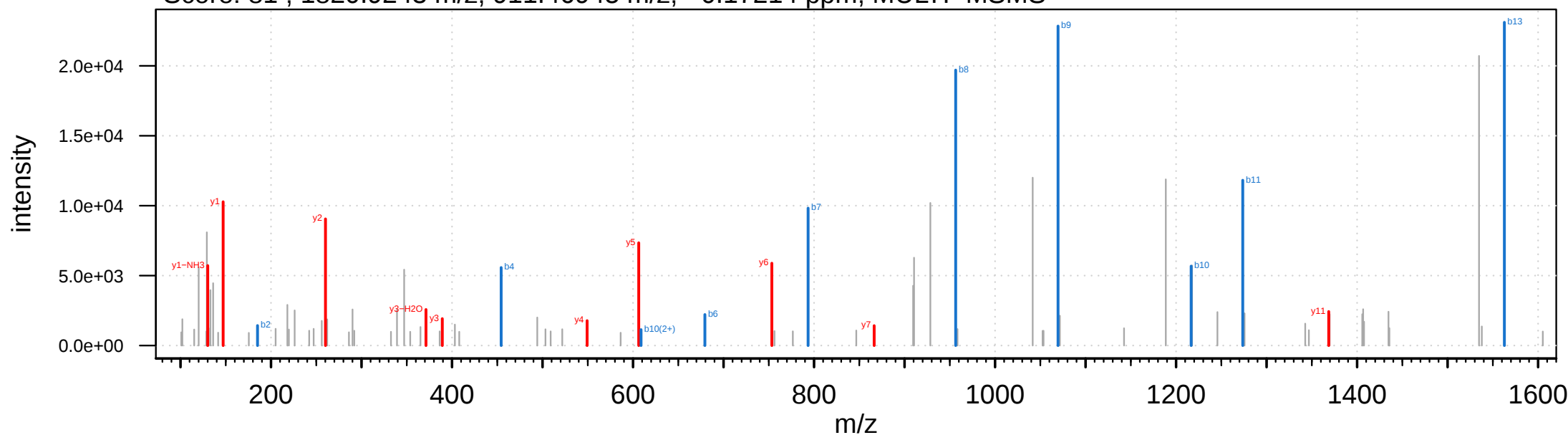
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 81 ; 1820.9243 m/z; 911.46945 m/z; -0.17214 ppm; MULTI-MSMS

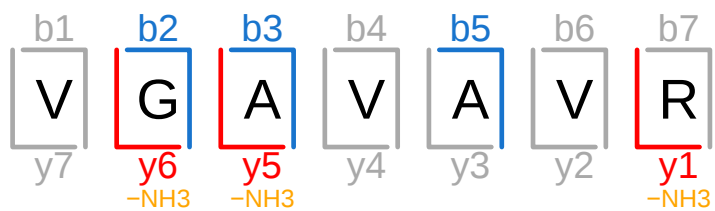


Raw File: 20100616_Velos1_TaGe_SA_RKO_1

Scan Number: 22671

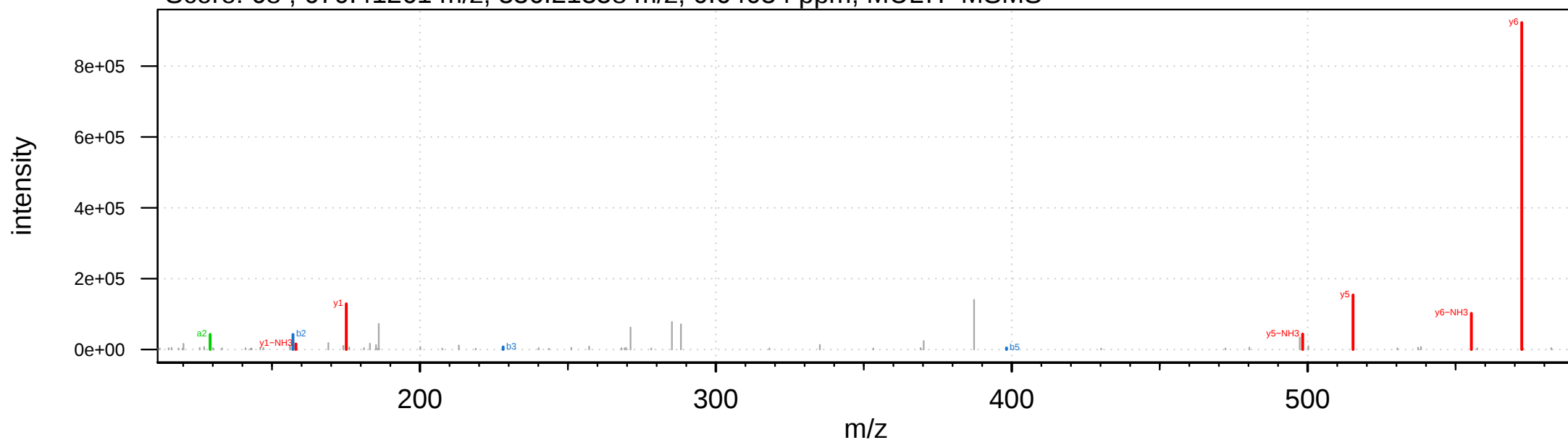
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

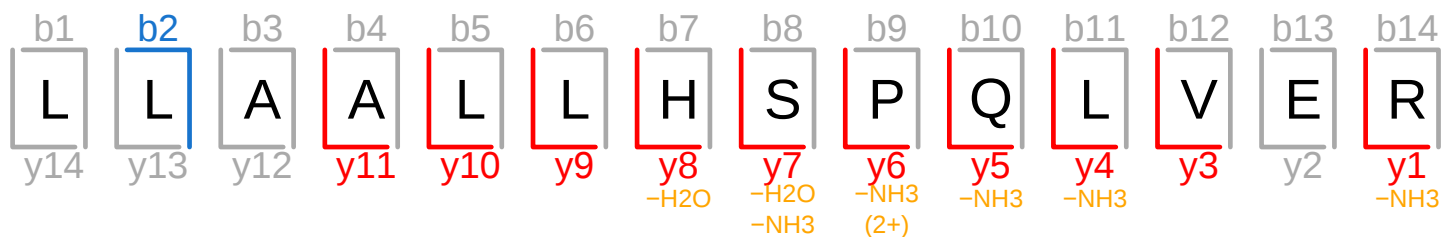


VGAVAVR

Score: 68 ; 670.41261 m/z; 336.21358 m/z; 0.64034 ppm; MULTI-MSMS

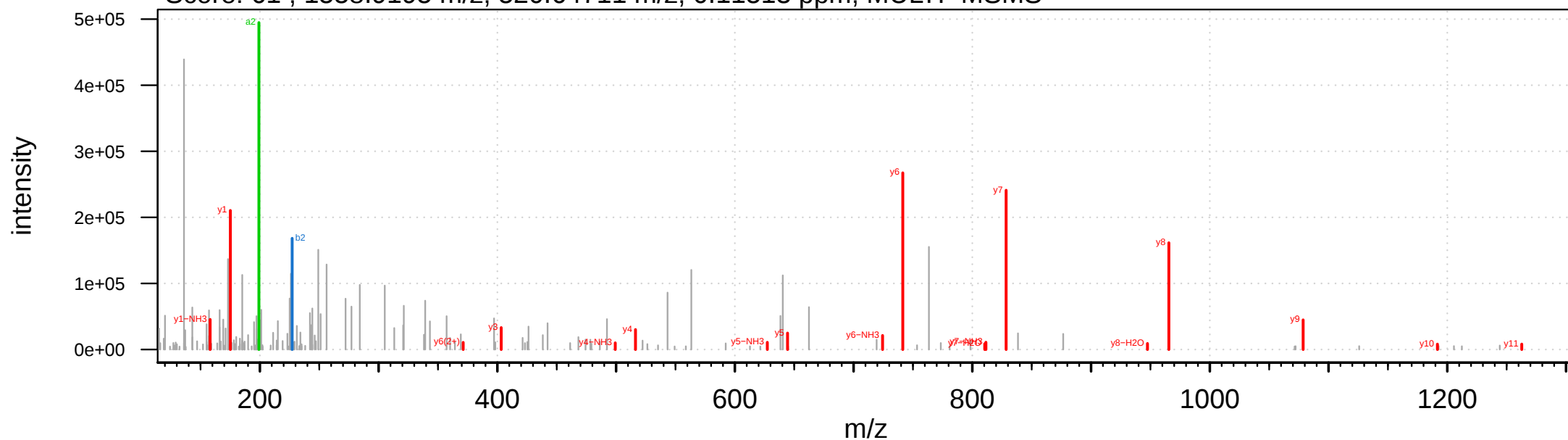


Raw File: 20100616_Velos1_TaGe_SA_RKO_1
 Scan Number: 2969
 Proteins:
 TCONS_I2_00030545_chrX:79544539-79546436:-

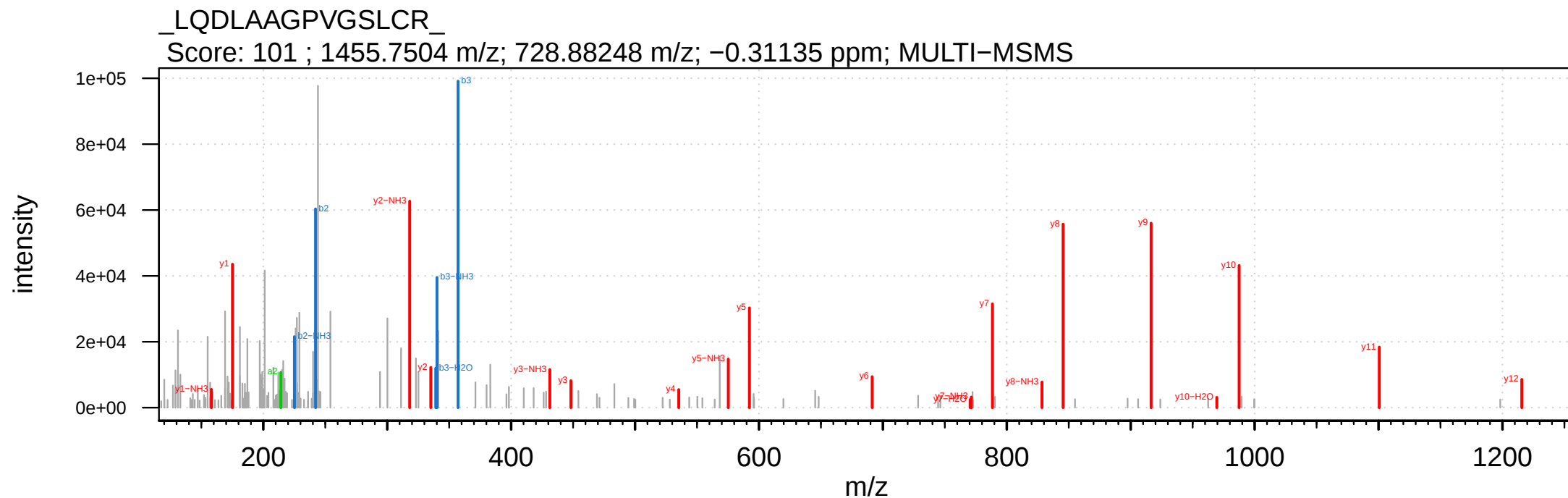
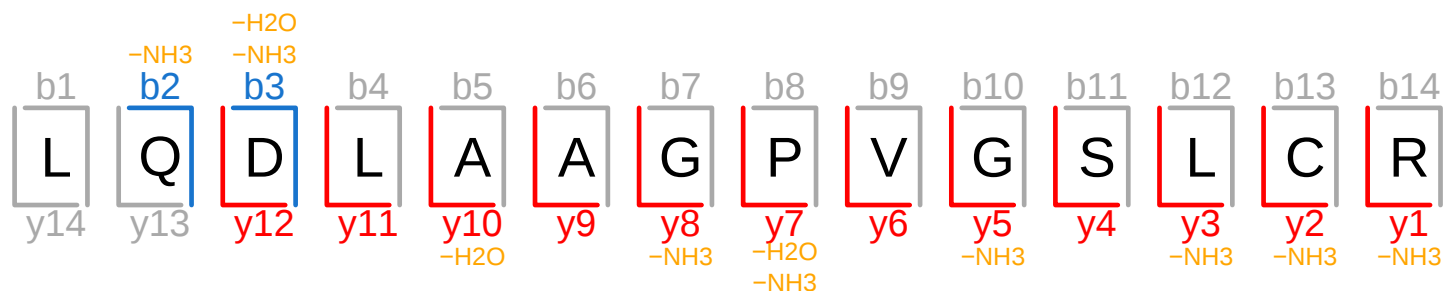


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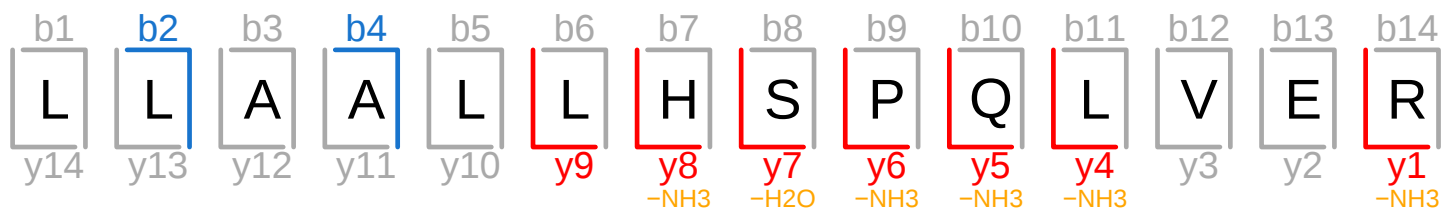
Score: 61 ; 1558.9195 m/z; 520.64711 m/z; 0.11513 ppm; MULTI-MSMS



Raw File: 20100719_Velos1_TaGe_SA_MCF7_01
 Scan Number: 26418
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+

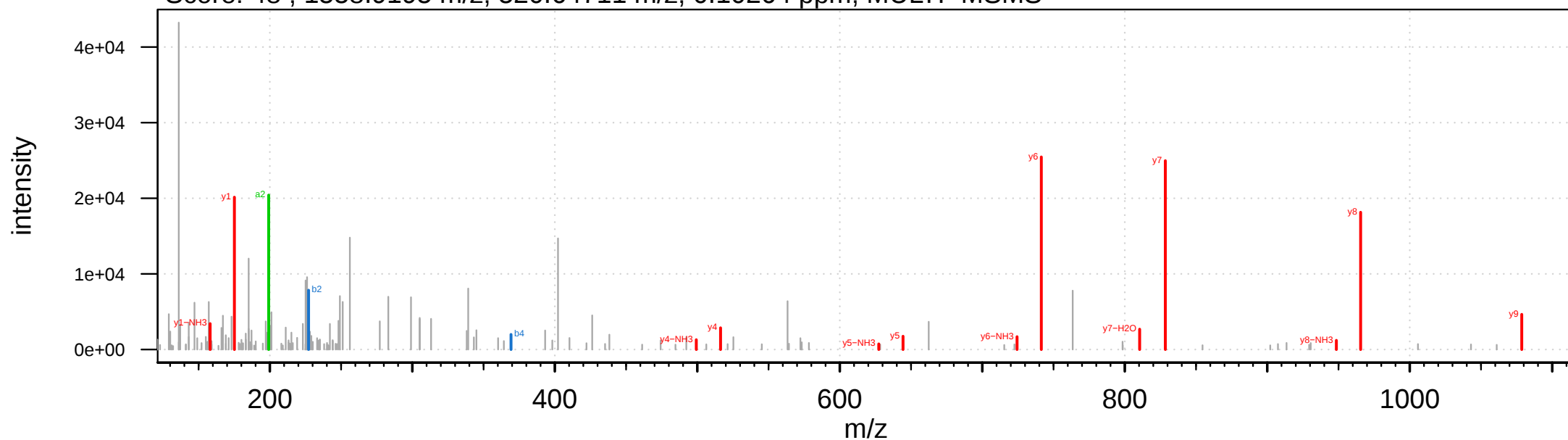


Raw File: 20100719_Velos1_TaGe_SA_MCF7_01
Scan Number: 22824
Proteins:
ENST00000602845_chr3:196669588-196669887:+

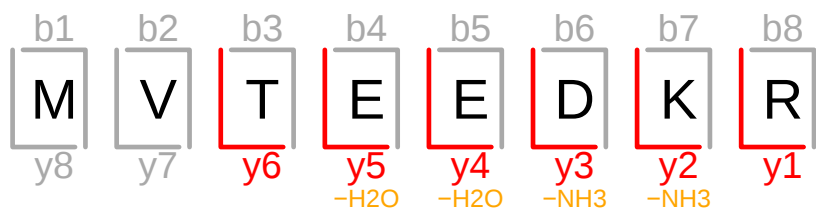


LLAALLHSPQLVER

Score: 48 ; 1558.9195 m/z; 520.64711 m/z; 0.10264 ppm; MULTI-MSMS

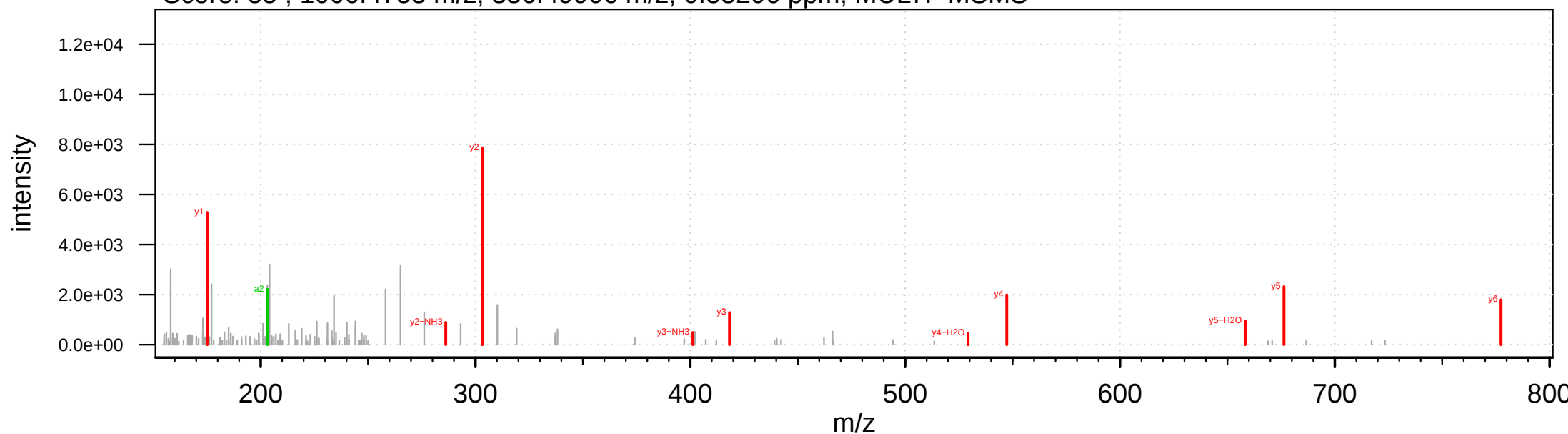


Raw File: 20100719_Velos1_TaGe_SA_MCF7_02
 Scan Number: 23454
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+



MVTEEDKR

Score: 55 ; 1006.4753 m/z; 336.49906 m/z; 0.53206 ppm; MULTI-MSMS

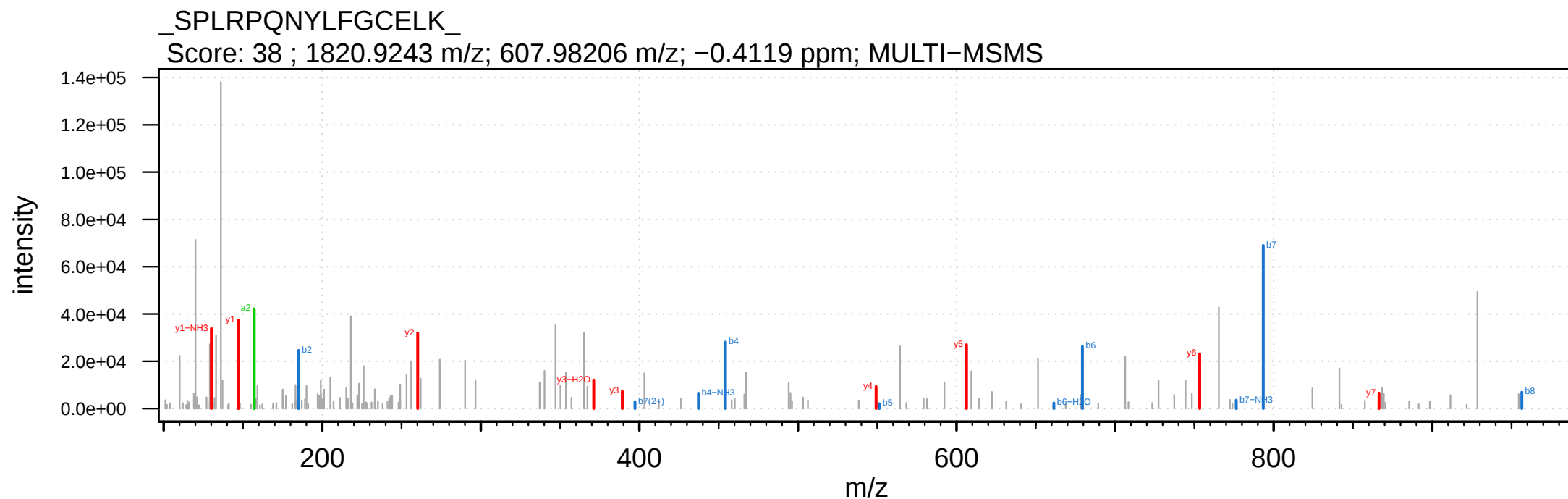
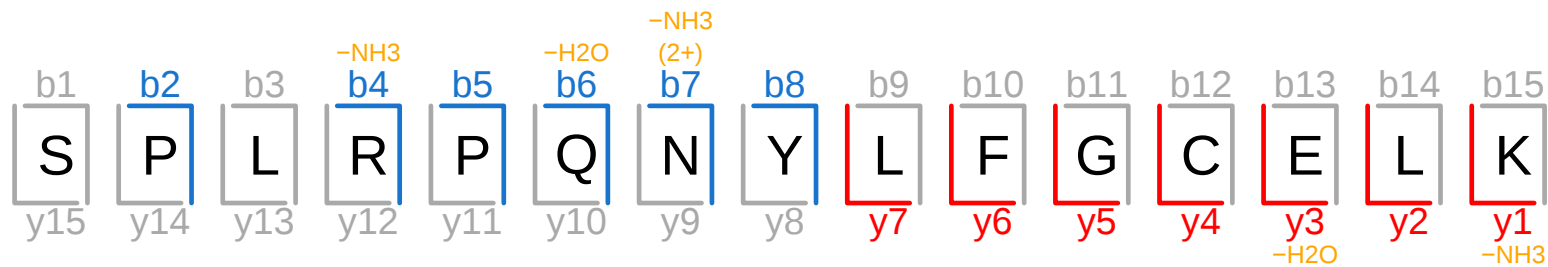


Raw File: 20100719_Velos1_TaGe_SA_MCF7_02

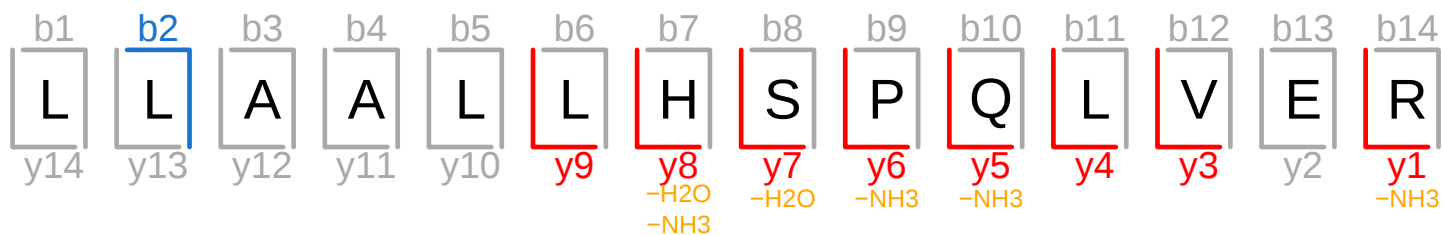
Scan Number: 4195

Proteins:

TCONS_I2_00006768_chr13:29174091-29174123:+

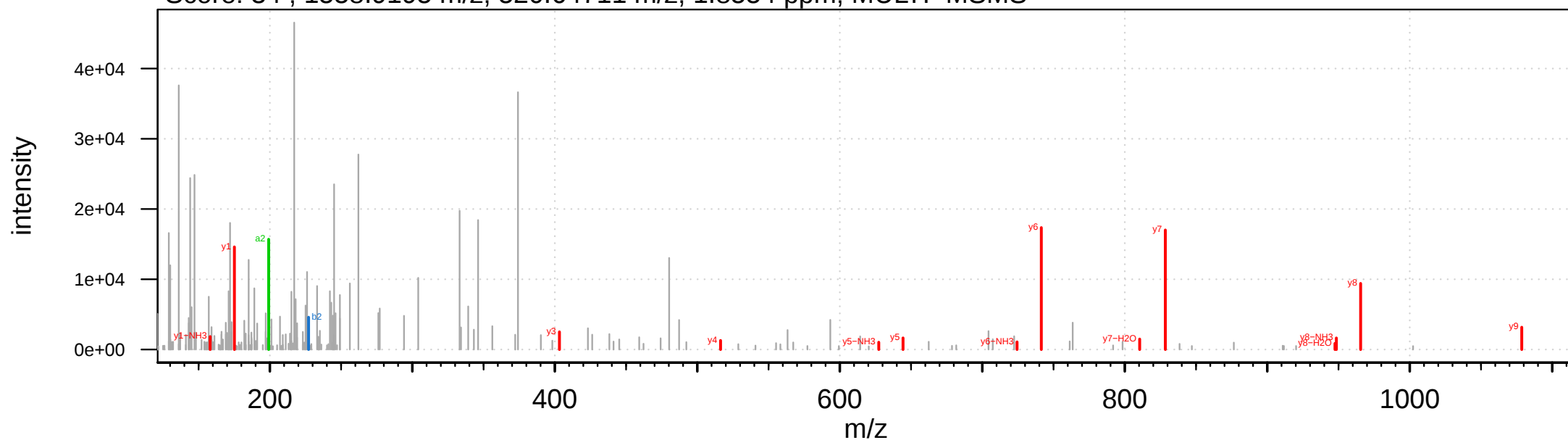


Raw File: 20100719_Velos1_TaGe_SA_MCF7_02
 Scan Number: 19997
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

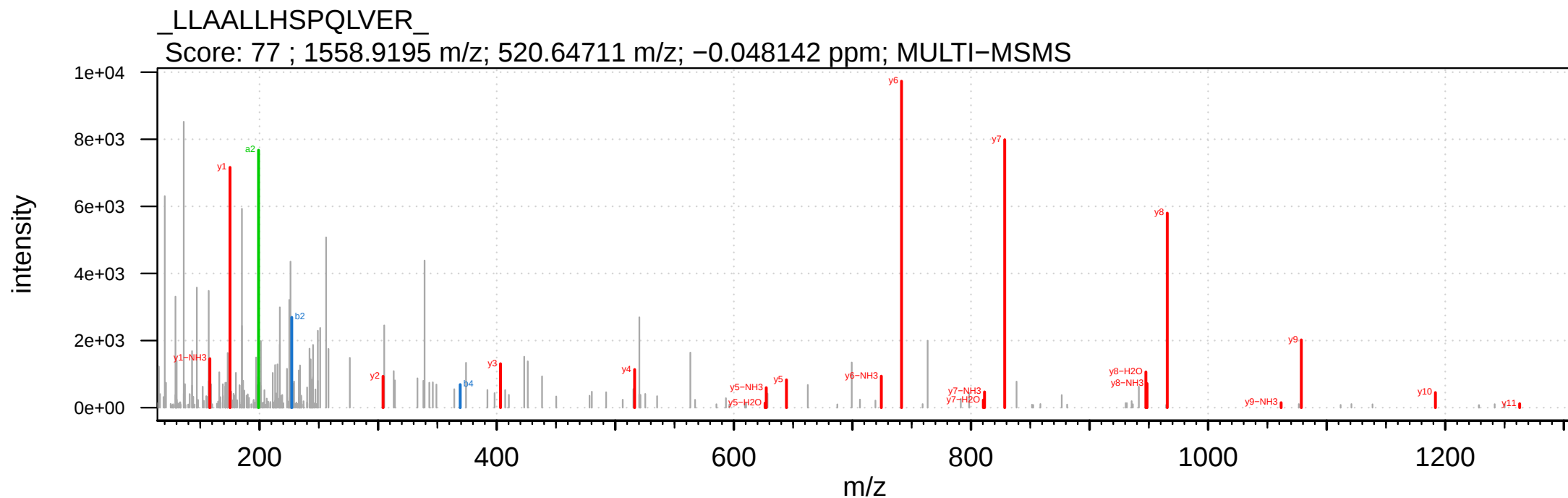
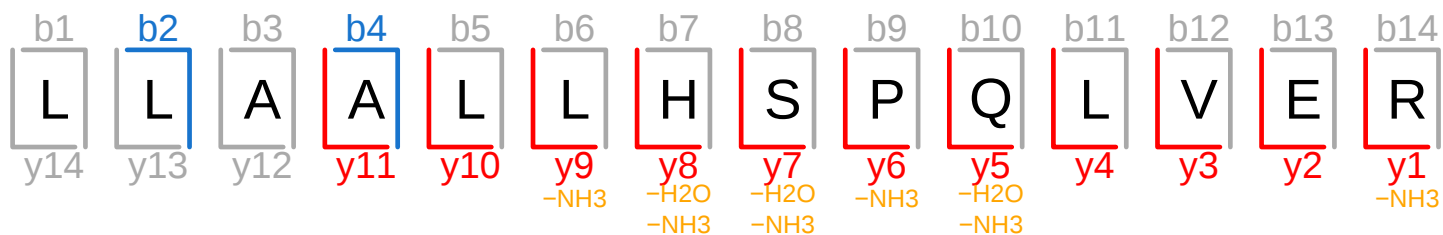


LLAALLHSPQLVER

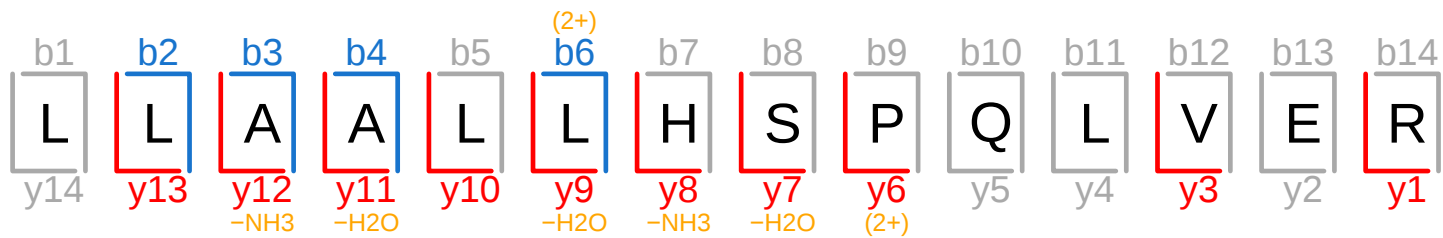
Score: 34 ; 1558.9195 m/z; 520.64711 m/z; 1.8554 ppm; MULTI-MSMS



Raw File: 20100719_Velos1_TaGe_SA_MCF7_03
 Scan Number: 22752
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+

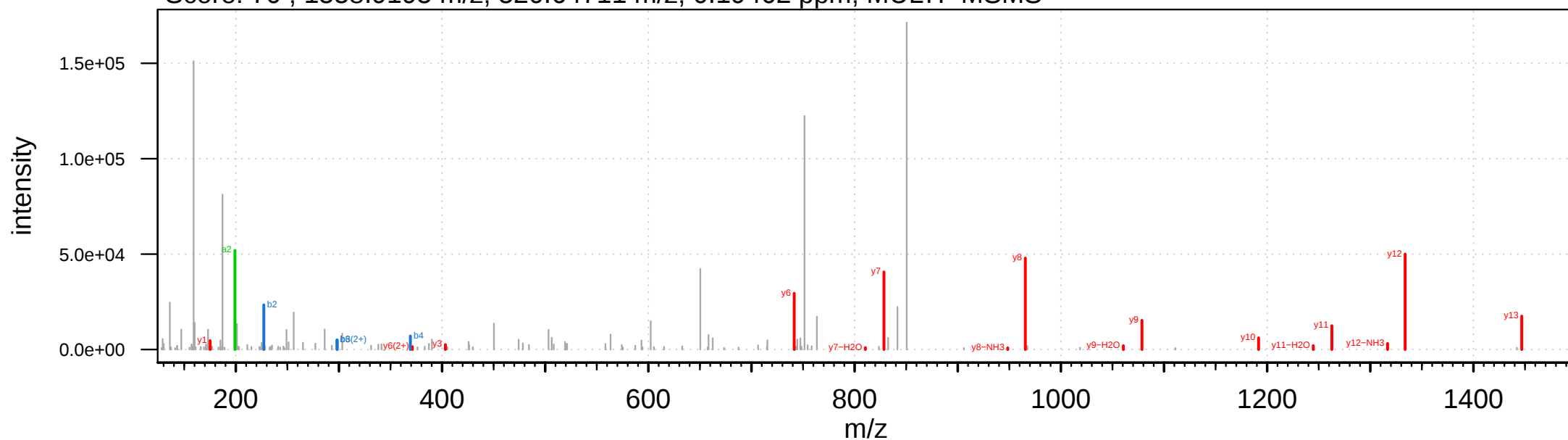


Raw File: 20100726_Velos1_TaGe_SA_HeLa_2
 Scan Number: 21111
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+

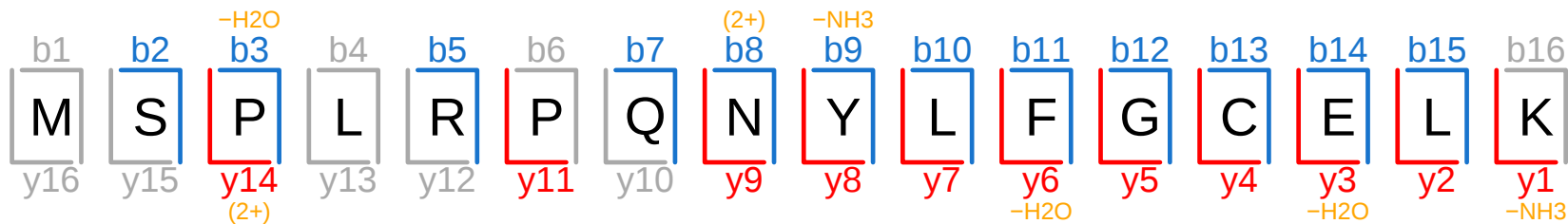


LLAALLHSPQLVER

Score: 70 ; 1558.9195 m/z; 520.64711 m/z; 0.19462 ppm; MULTI-MSMS

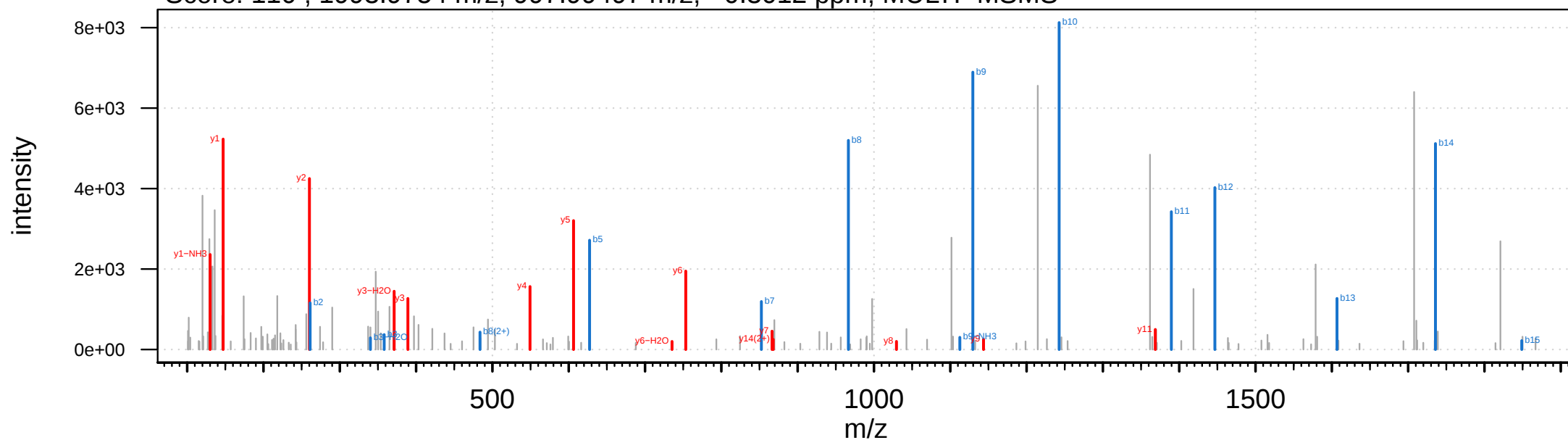


Raw File: 20101210_Velos1_AnWe_SA_MCF7_2
 Scan Number: 22820
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+



(ac)MSPLRPQNYLFGCELK

Score: 110 ; 1993.9754 m/z; 997.99497 m/z; -0.3012 ppm; MULTI-MSMS



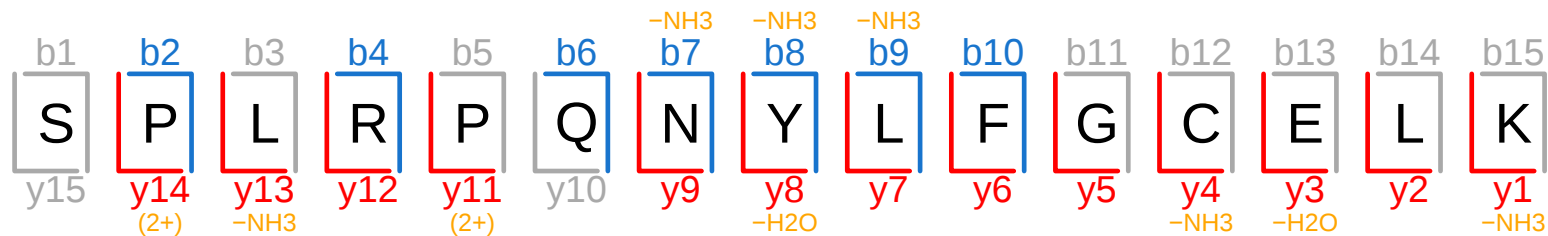
Raw File: 20101210_Velos1_AnWe_SA_MCF7_2

Scan Number: 30652

Proteins:

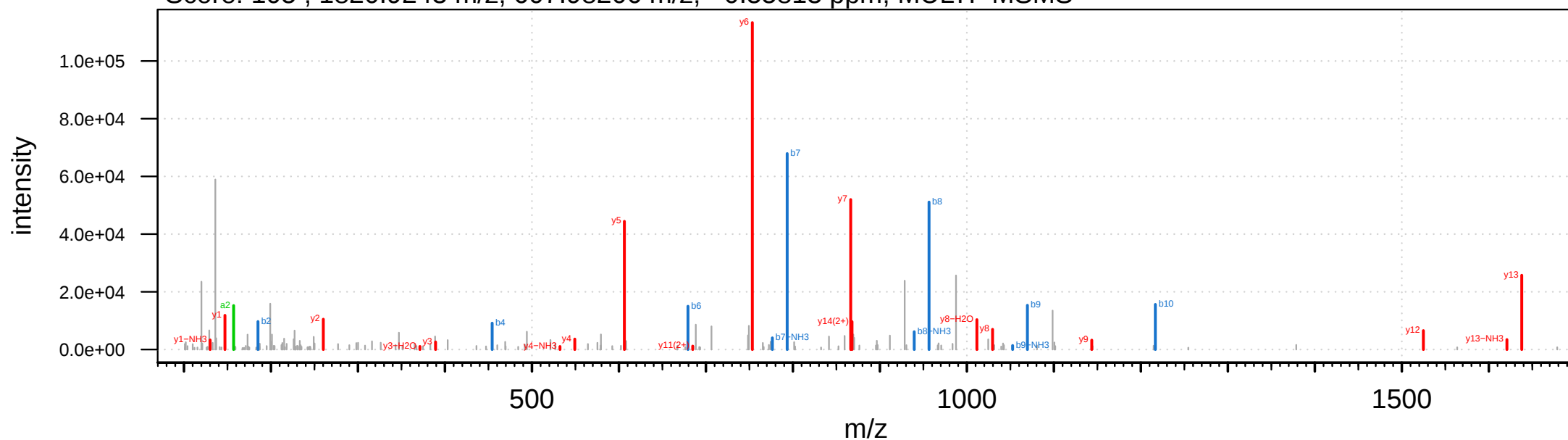
TCONS_I2_00008829_chr15:92829088-92829258:+

Raw File: 20101210_Velos1_AnWe_SA_MCF7_2
Scan Number: 22814
Proteins:
TCONS_ID 00008829 chr15:92829088-92829258:+

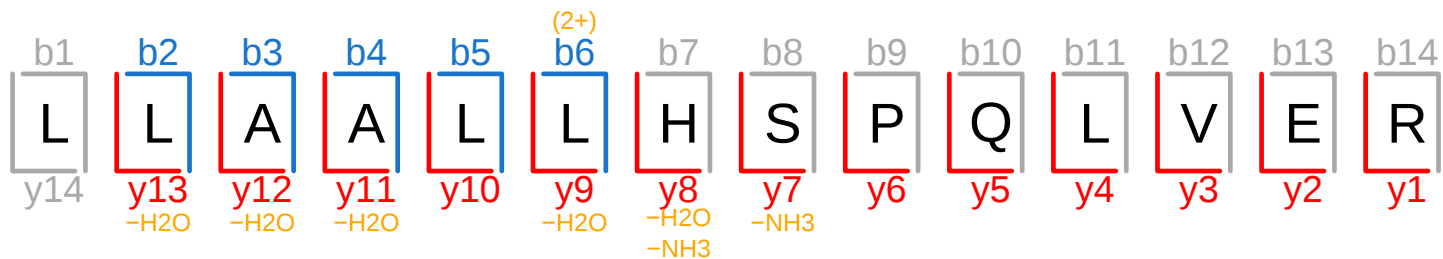


SPLRPQNYLFGCELK

Score: 105 ; 1820.9243 m/z; 607.98206 m/z; -0.55813 ppm; MULTI-MSMS

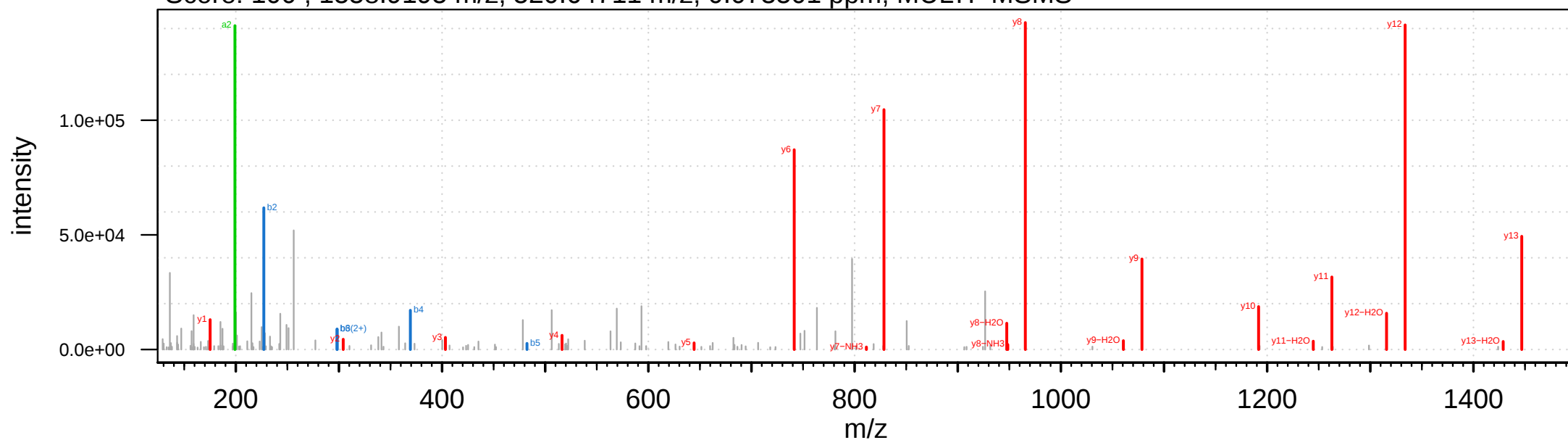


Raw File: 20101210_Velos1_AnWe_SA_MCF7_2
 Scan Number: 20040
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

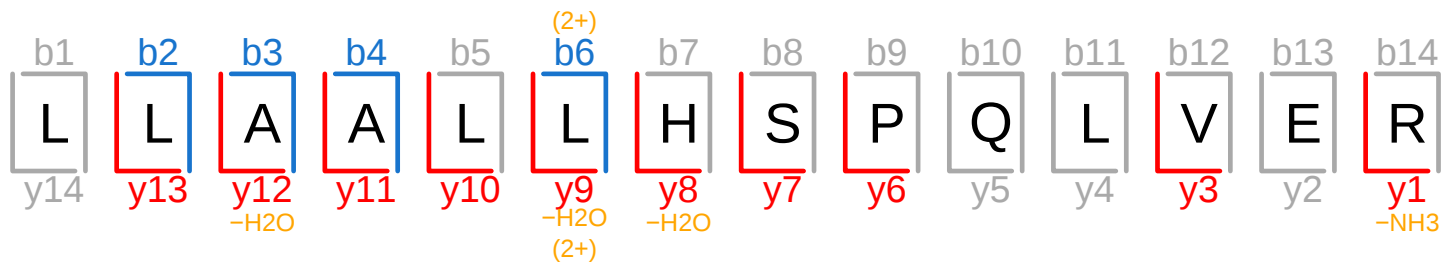


LLAALLHSPQLVER

Score: 100 ; 1558.9195 m/z; 520.64711 m/z; 0.073501 ppm; MULTI-MSMS

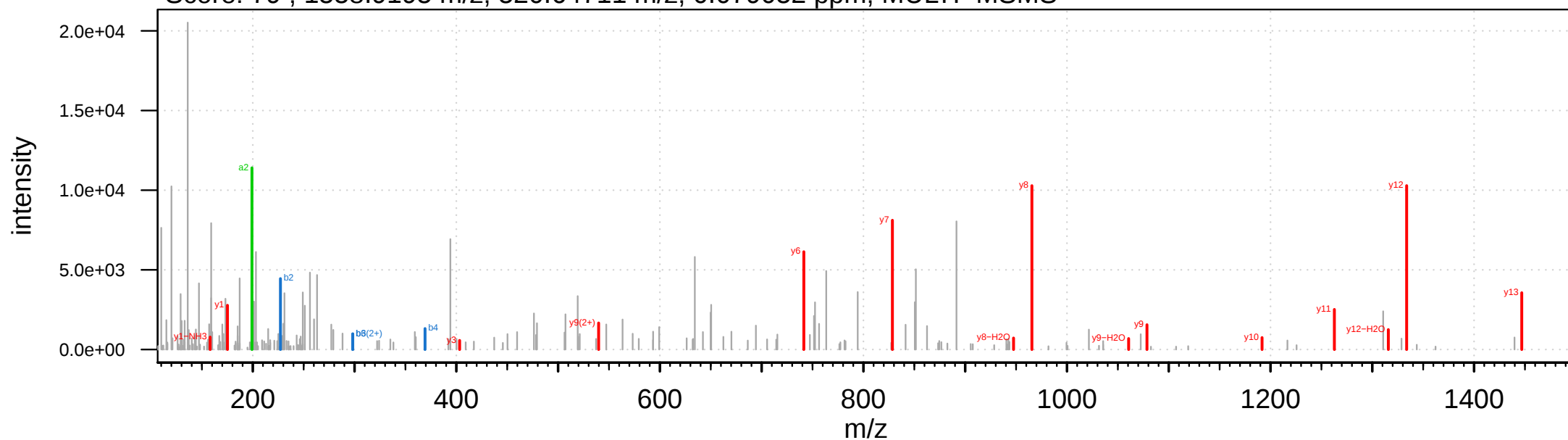


Raw File: 20101210_Velos1_AnWe_SA_MCF7_3
 Scan Number: 22064
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+

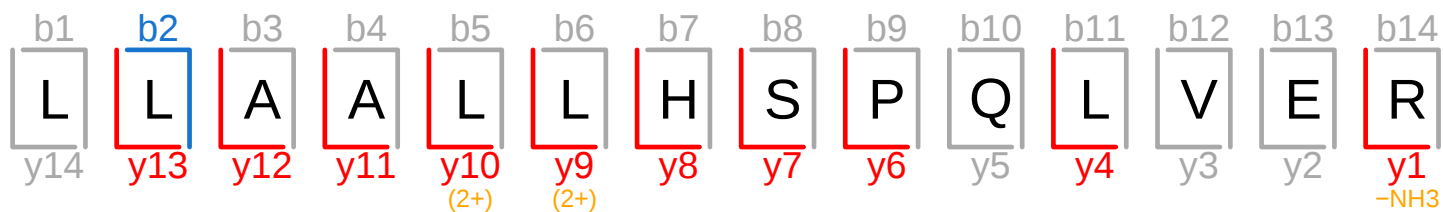


LLAALLHSPQLVER

Score: 79 ; 1558.9195 m/z; 520.64711 m/z; 0.079052 ppm; MULTI-MSMS

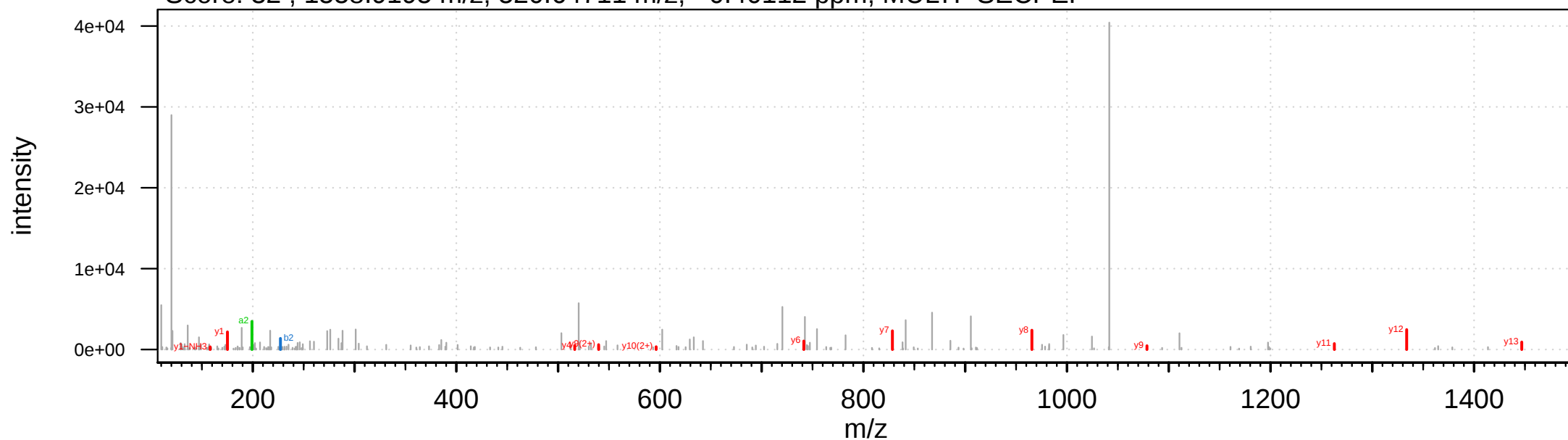


Raw File: 20101210_Velos1_AnWe_SA_MCF7_4
 Scan Number: 20013
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+



LLAALLHSPQLVER

Score: 32 ; 1558.9195 m/z; 520.64711 m/z; -0.40112 ppm; MULTI-SECPEP

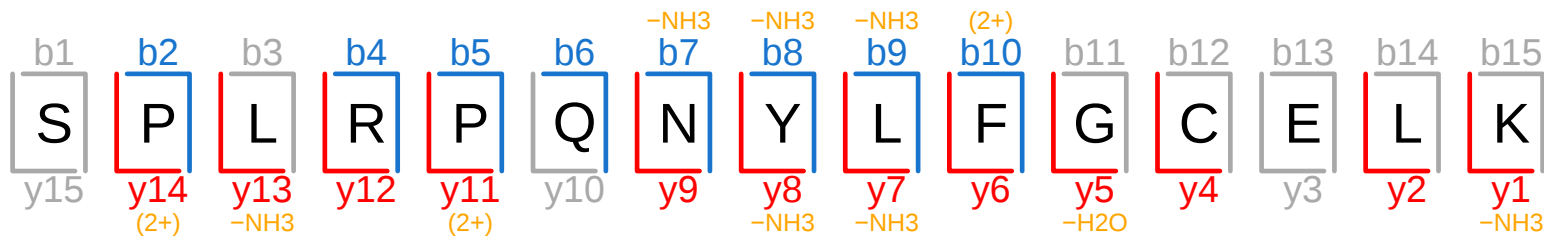


Raw File: 20101224_Velos1-TaGe_SA-HepG2_3

Scan Number: 24144

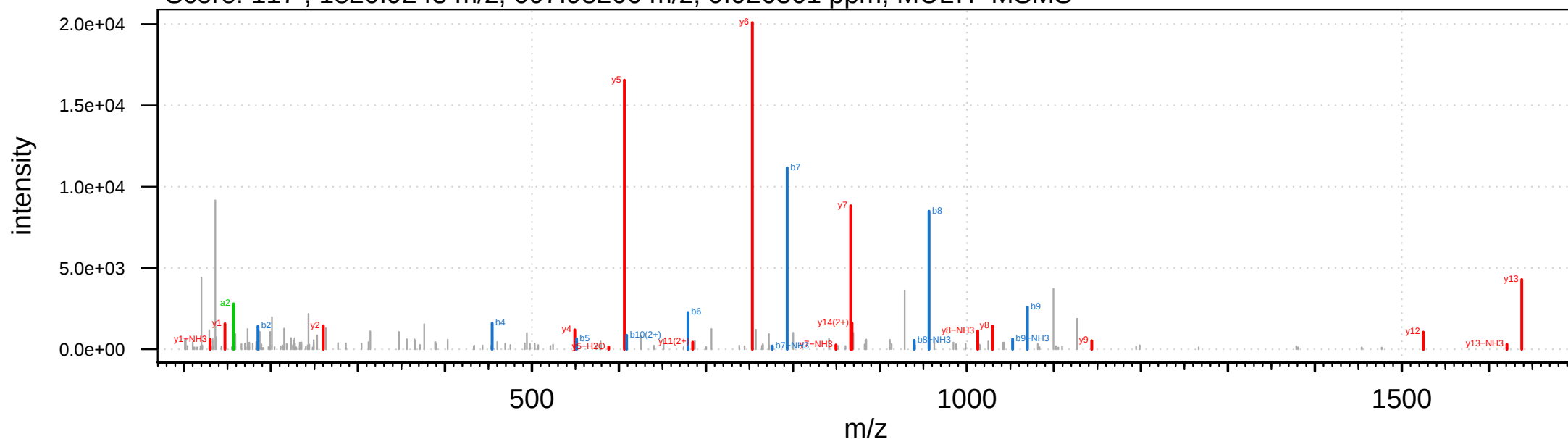
Proteins:

ENST00000602845_chr3:196669588-196669887:+



SPLRPQNYLFGCELK

Score: 117 ; 1820.9243 m/z; 607.98206 m/z; 0.026361 ppm; MULTI-MSMS

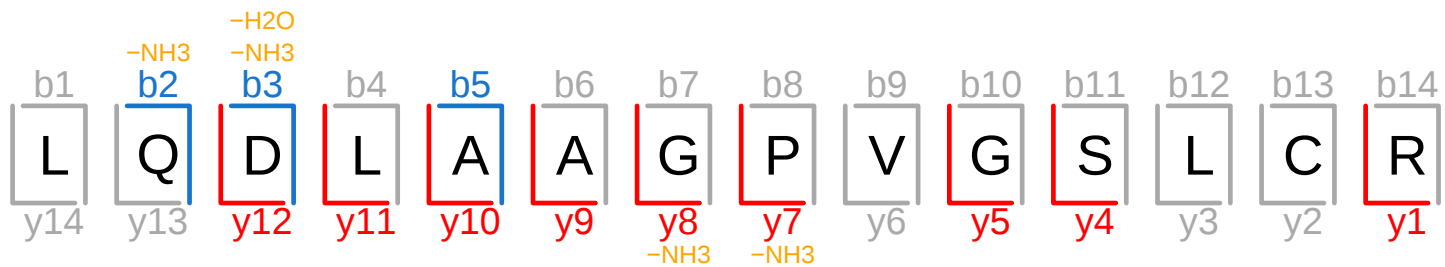


Raw File: 20101224_Velos1_TaGe_SA_HepG2_3

Scan Number: 20547

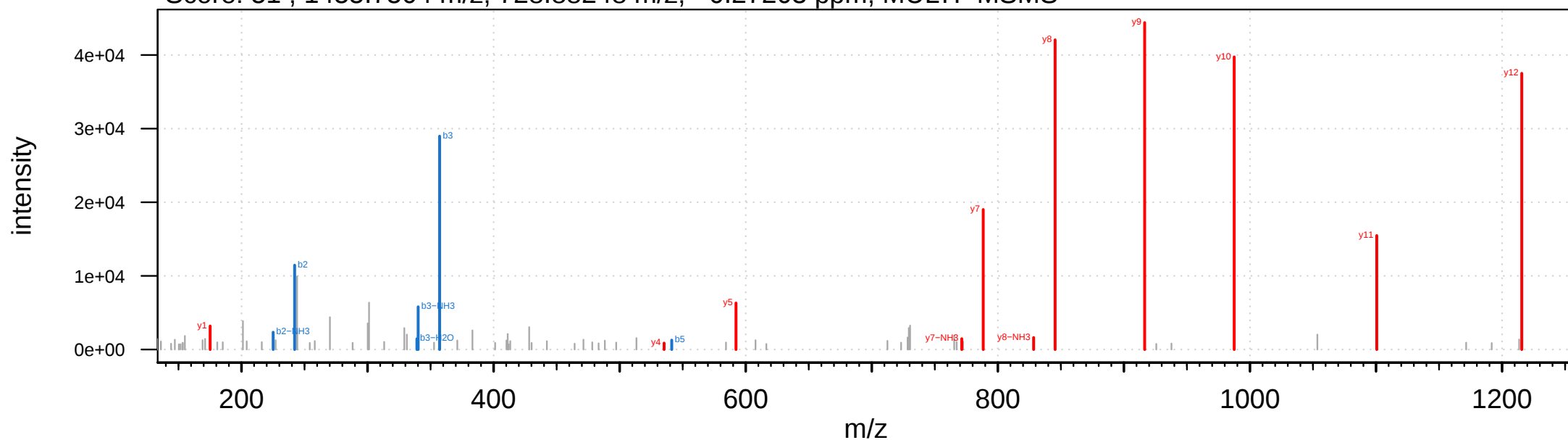
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

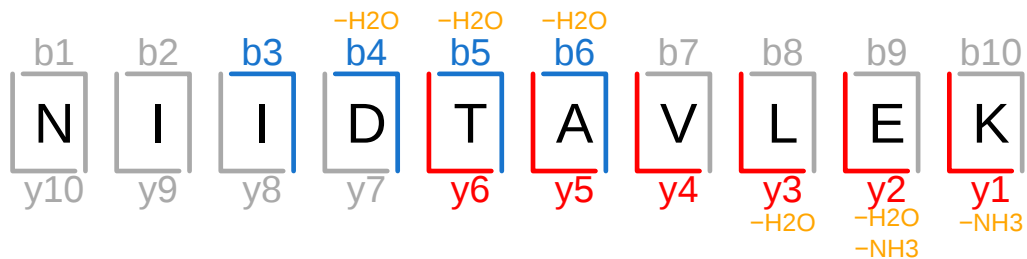


LQDLAAGPVGSLCR

Score: 51 ; 1455.7504 m/z; 728.88248 m/z; -0.27203 ppm; MULTI-MSMS

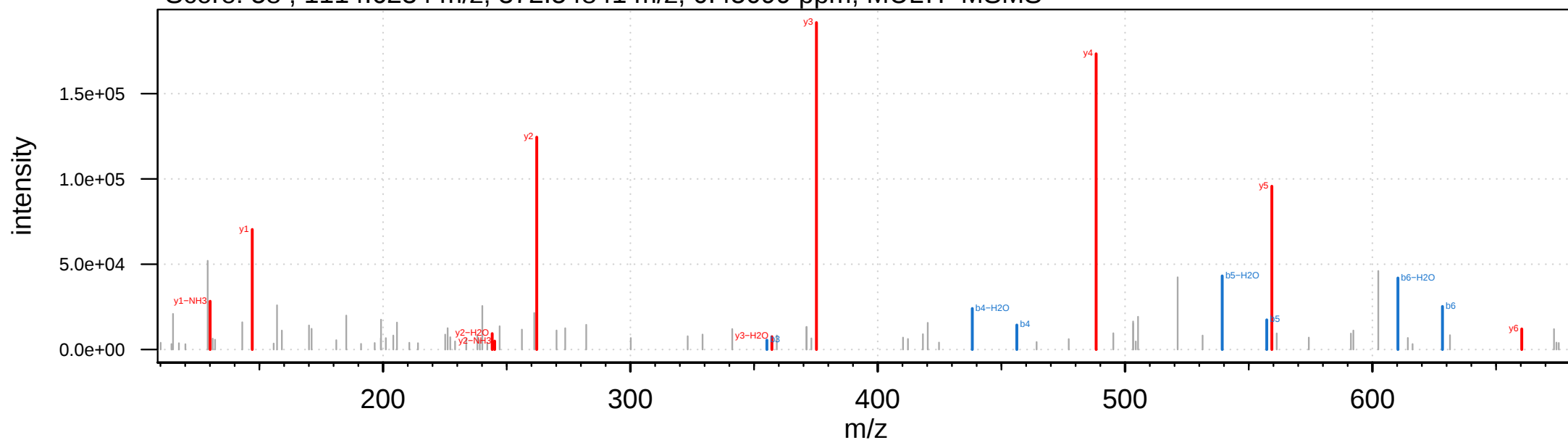


Raw File: 20100618_Velos1_TaGe_SA_U2OS_1
 Scan Number: 21570
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+



NIIDTAVLEK

Score: 58 ; 1114.6234 m/z; 372.54841 m/z; 0.45099 ppm; MULTI-MSMS

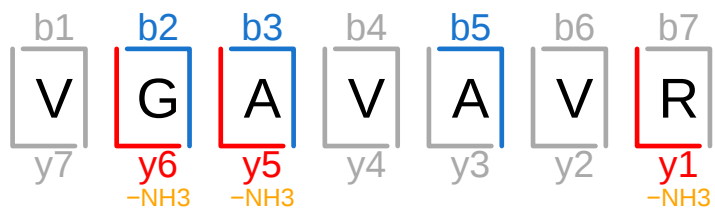


Raw File: 20100618_Velos1_TaGe_SA_U2OS_1

Scan Number: 5637

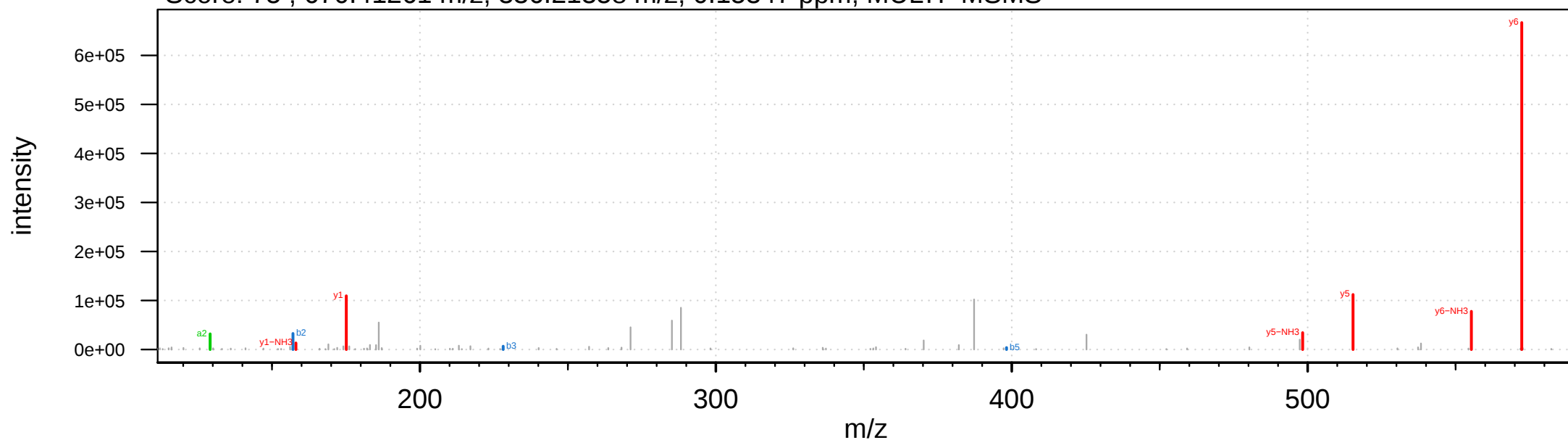
Proteins:

ENST00000497138_chr20:56806826-56807846:-

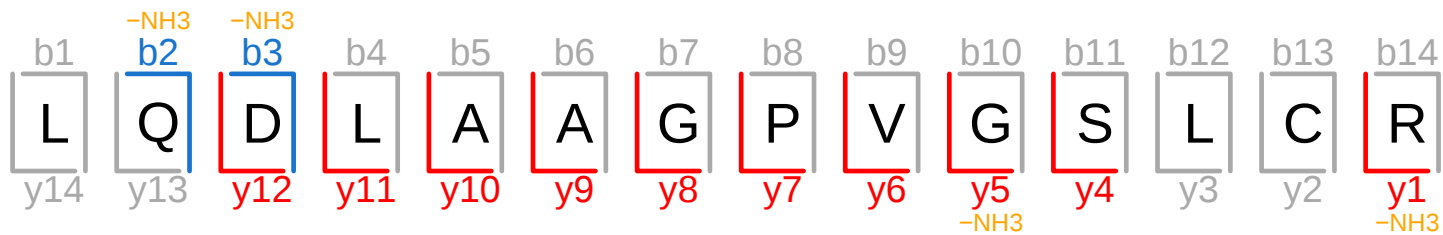


VGAVAVR

Score: 75 ; 670.41261 m/z; 336.21358 m/z; 0.15347 ppm; MULTI-MSMS

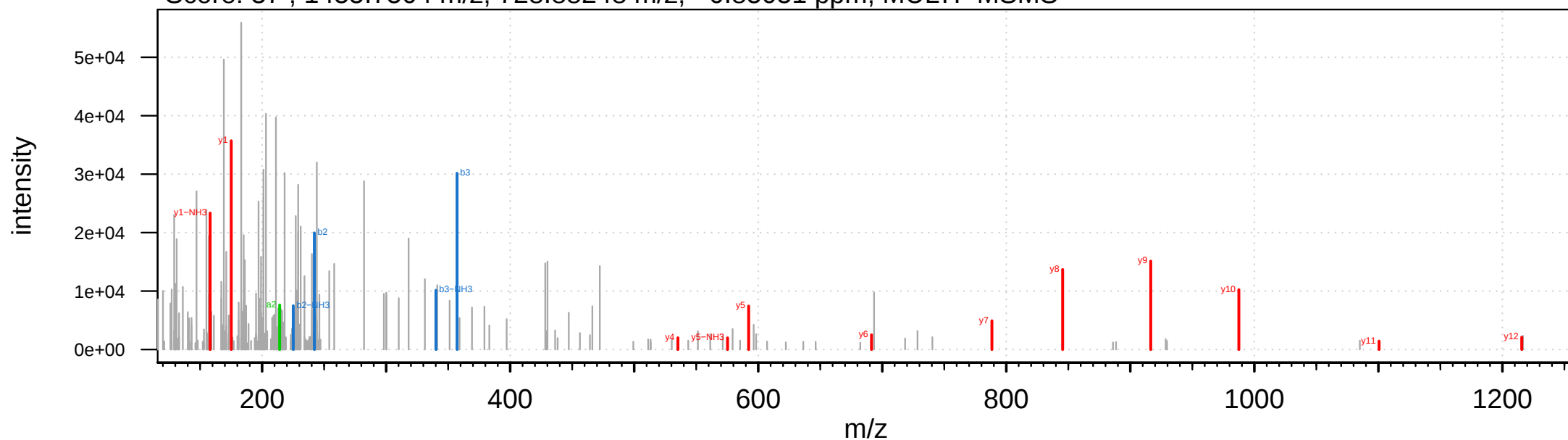


Raw File: 20100618_Velos1_TaGe_SA_U2OS_1
 Scan Number: 2825
 Proteins:
 TCONS_I2_00030545_chrX:79544539-79546436:-



LQDLAAGPVGSLCR

Score: 37 ; 1455.7504 m/z; 728.88248 m/z; -0.85051 ppm; MULTI-MSMS

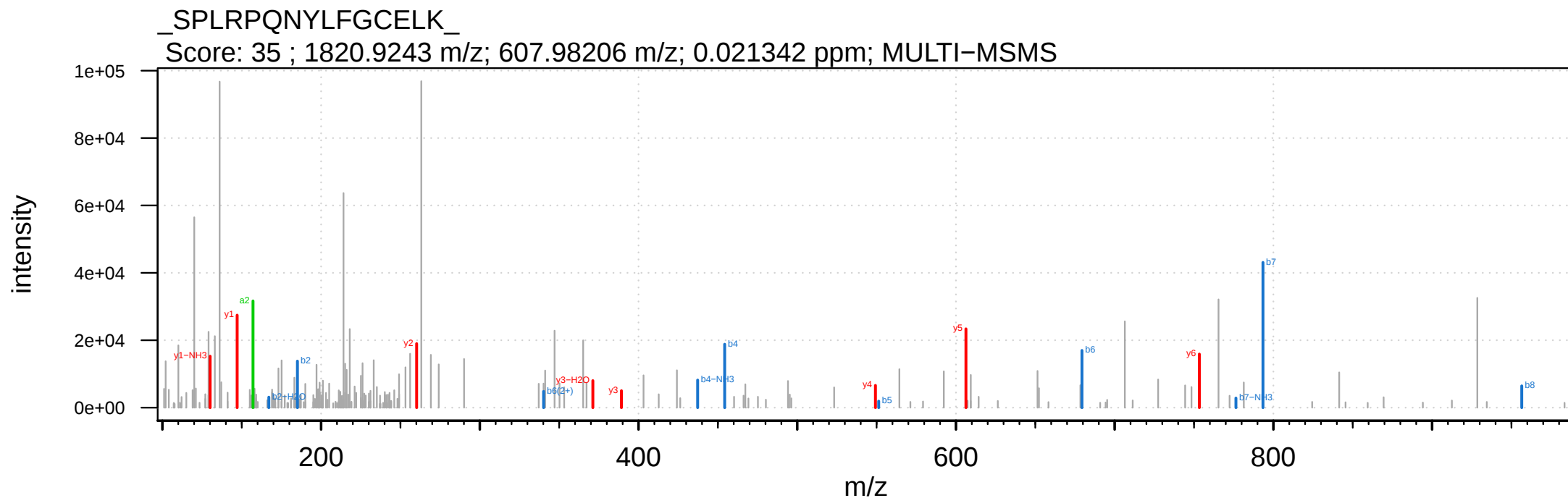
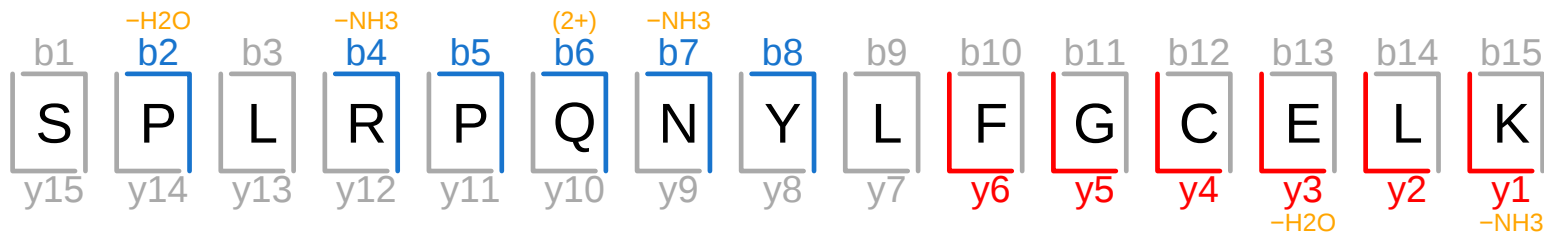


Raw File: 20100721_Velos1_TaGe_SA_U2OS_1

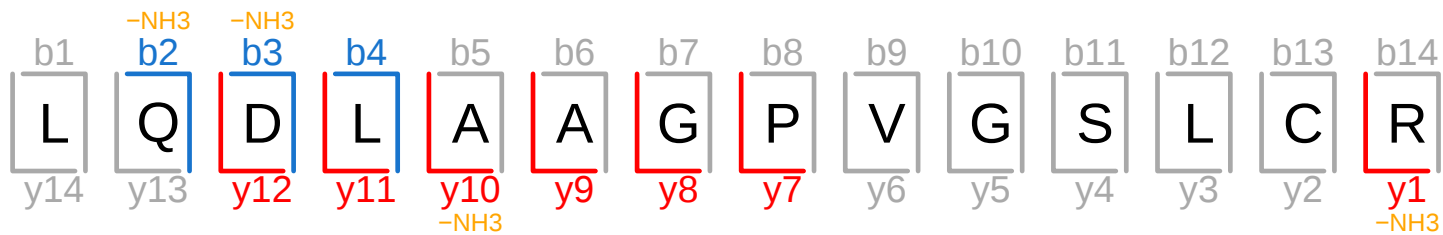
Scan Number: 17691

Proteins:

ENST00000602845_chr3:196669588-196669887:+

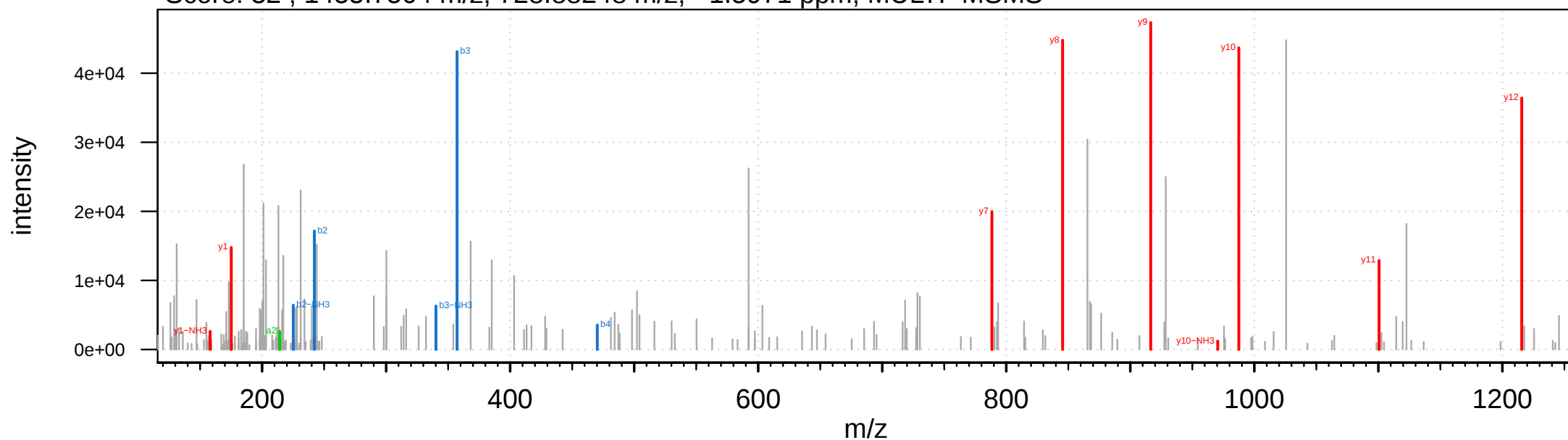


Raw File: 20100721_Velos1_TaGe_SA_U2OS_1
 Scan Number: 19584
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

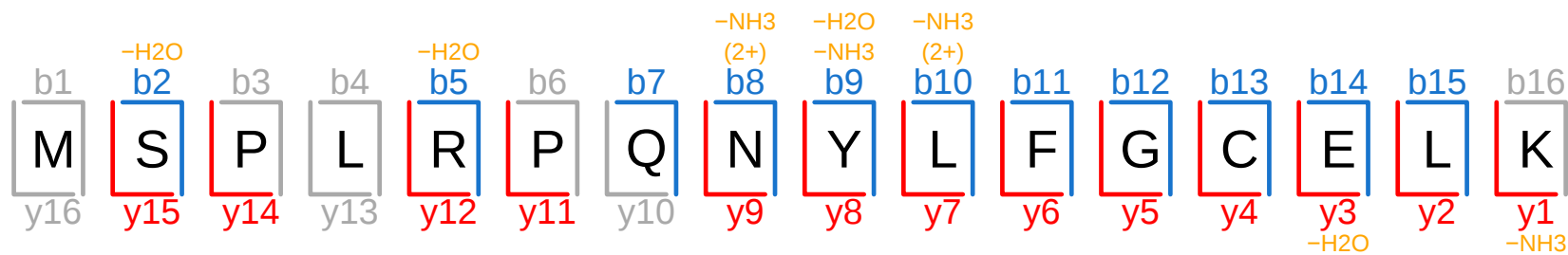


LQDLAAGPVGSLCR

Score: 32 ; 1455.7504 m/z; 728.88248 m/z; -1.5971 ppm; MULTI-MSMS

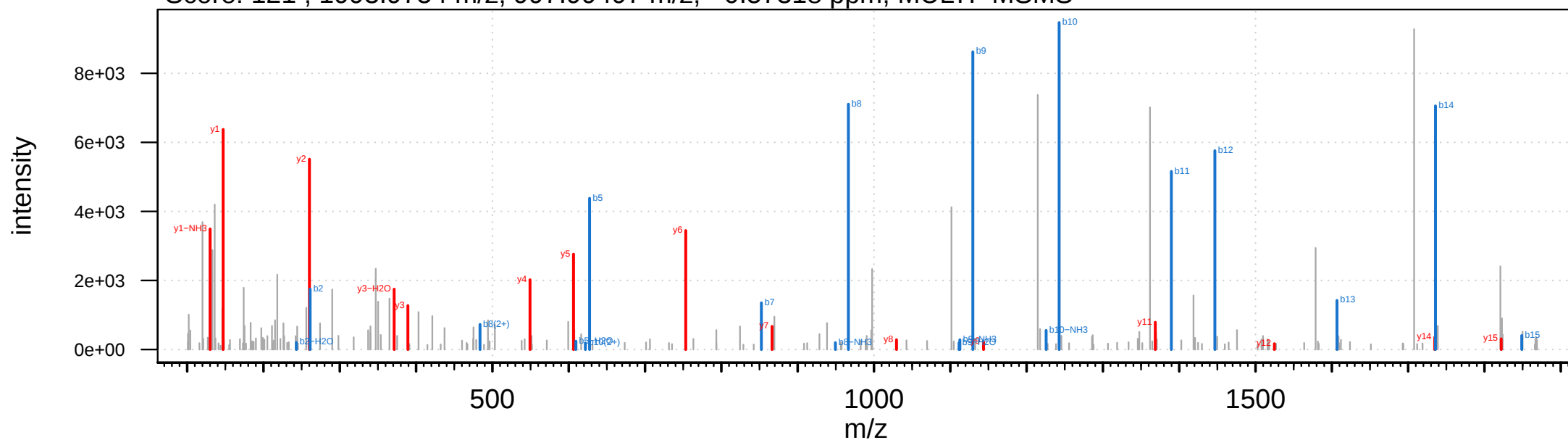


Raw File: 20101210_Velos1_AnWe_SA_LnCap_1
 Scan Number: 20166
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+



(ac)MSPLRPQNYLFGCELK

Score: 121 ; 1993.9754 m/z; 997.99497 m/z; -0.57318 ppm; MULTI-MSMS

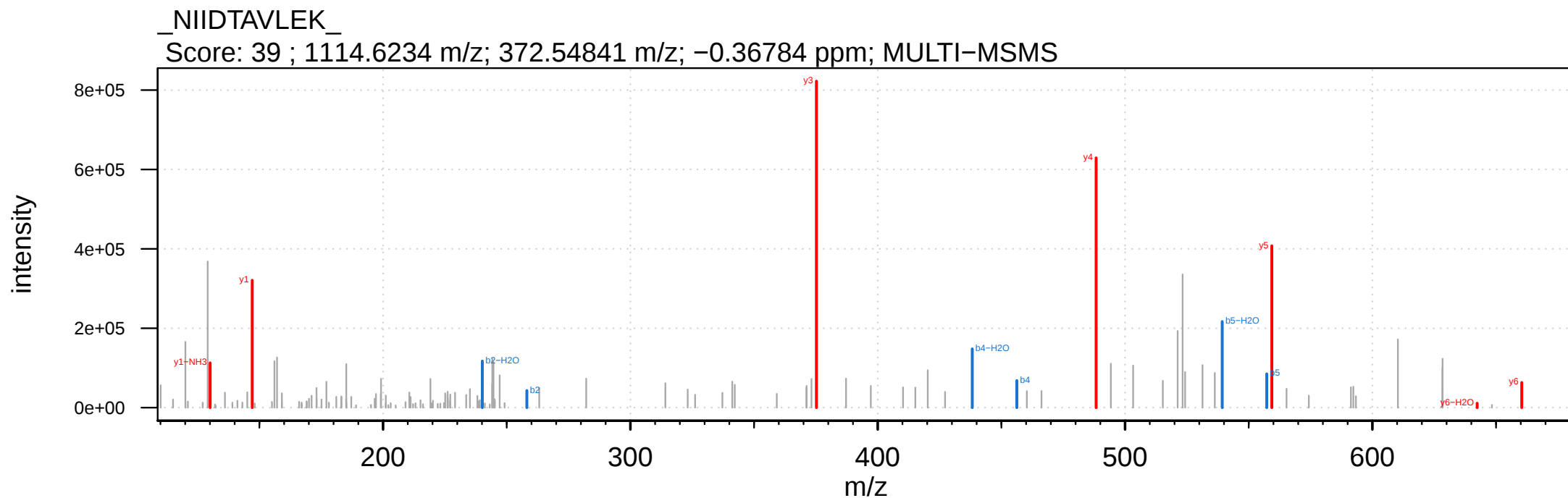
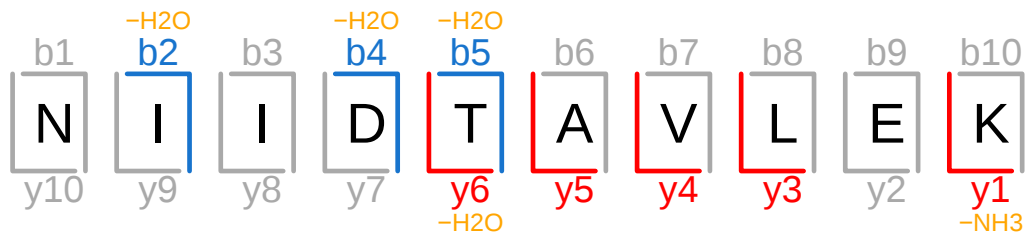


Raw File: 20101210_Velos1_AnWe_SA_LnCap_1

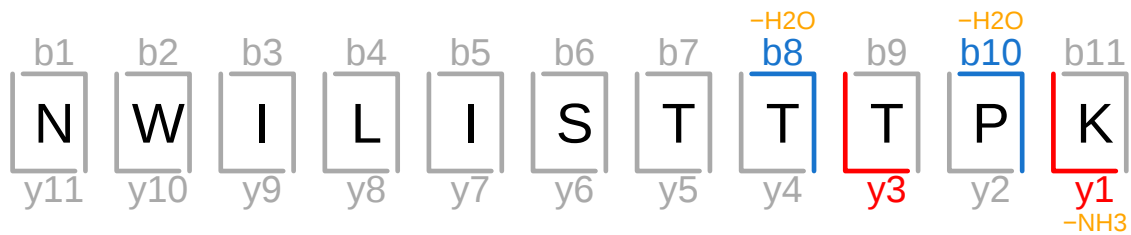
Scan Number: 36029

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

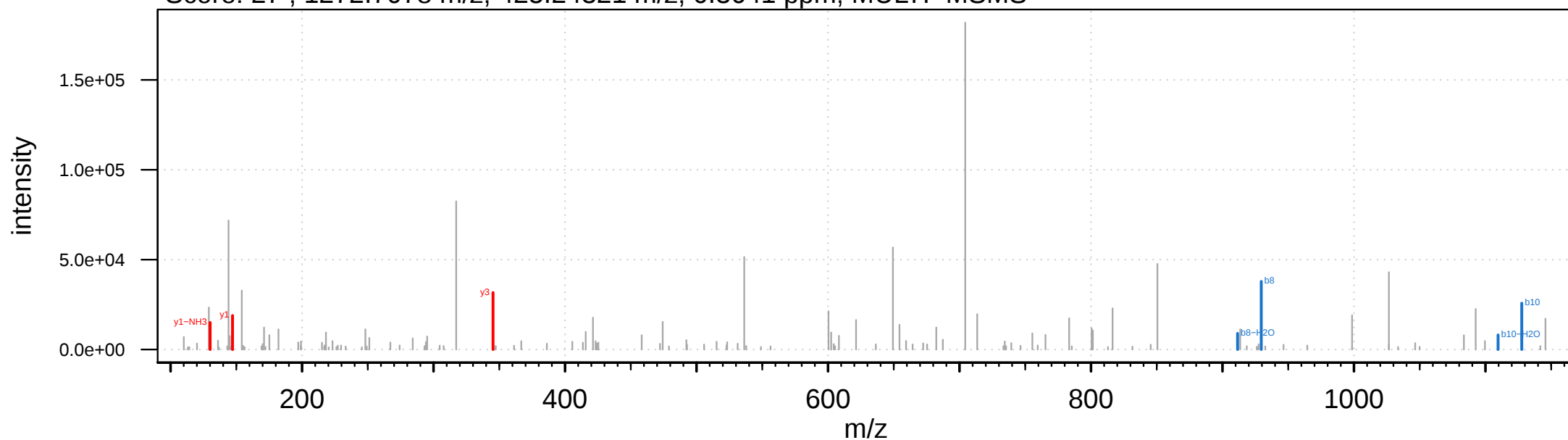


Raw File: 20101210_Velos1_AnWe_SA_LnCap_1
Scan Number: 5308
Proteins:
ENST00000497138_chr20:56806826-56807846:-



NWILISTTTPK

Score: 27 ; 1272.7078 m/z; 425.24321 m/z; 0.3641 ppm; MULTI-MSMS

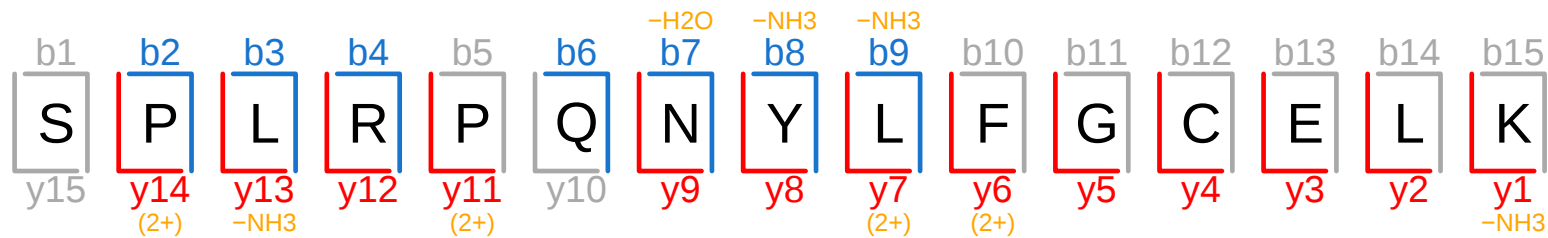


Raw File: 20101210_Velos1_AnWe_SA_LnCap_1

Scan Number: 25645

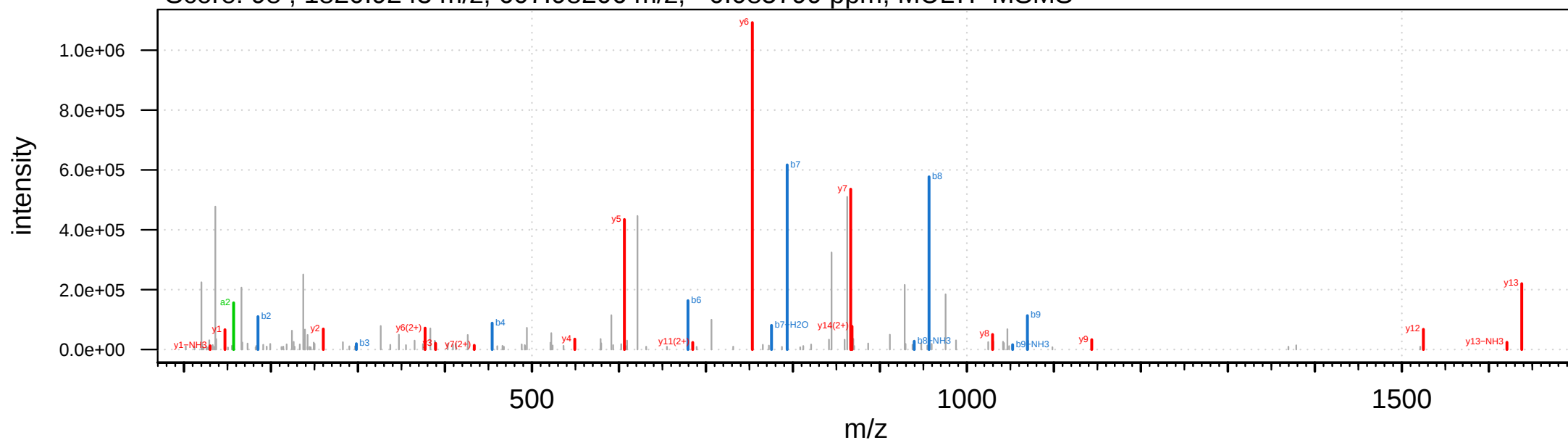
Proteins:

ENST00000430431_chr10:61496818-61496958:-

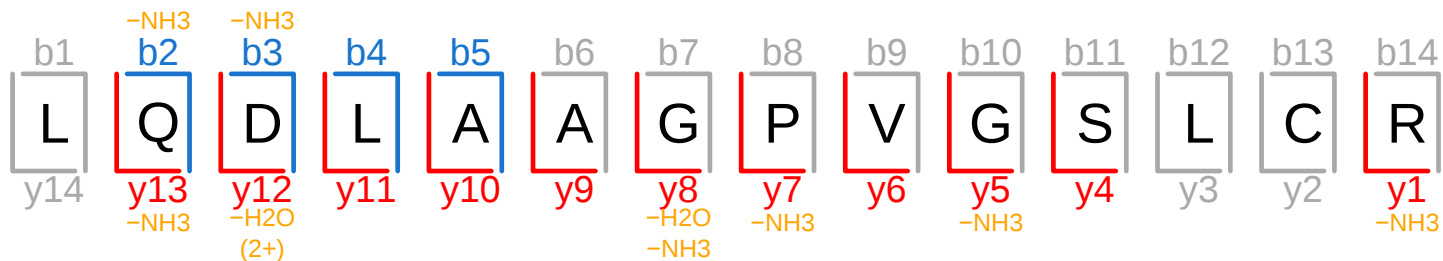


SPLRPQNYLFGCELK

Score: 98 ; 1820.9243 m/z; 607.98206 m/z; -0.085799 ppm; MULTI-MSMS

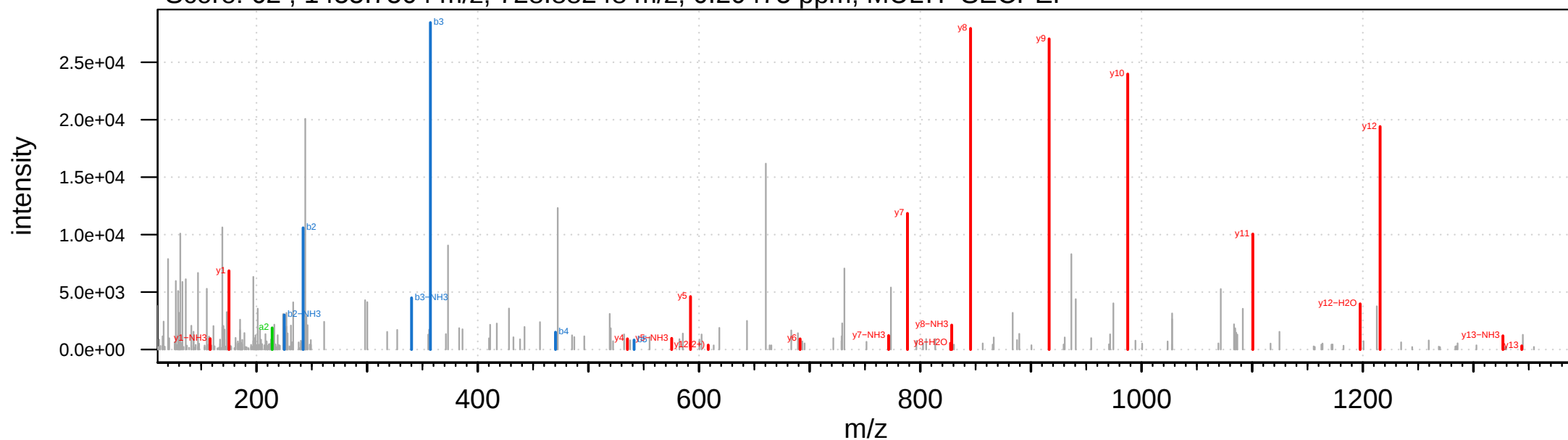


Raw File: 20101210_Velos1_AnWe_SA_LnCap_1
 Scan Number: 22329
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

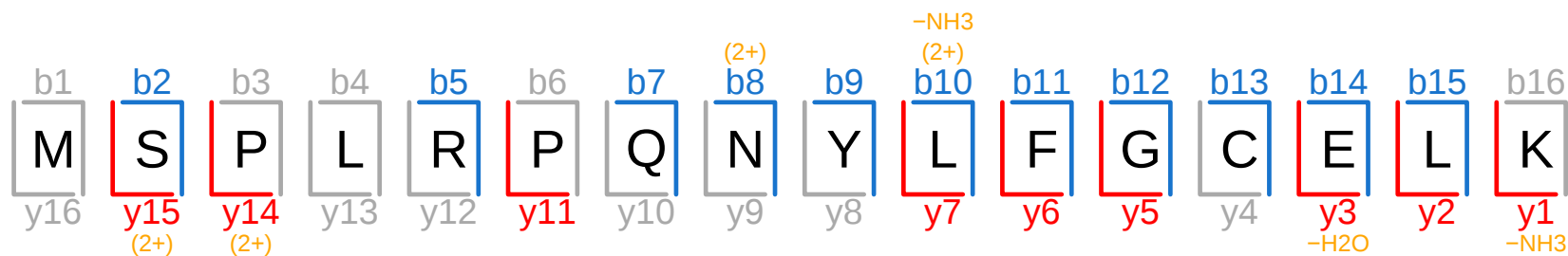


LQDLAAGPVGSLCR

Score: 62 ; 1455.7504 m/z; 728.88248 m/z; 0.20475 ppm; MULTI-SECPEP

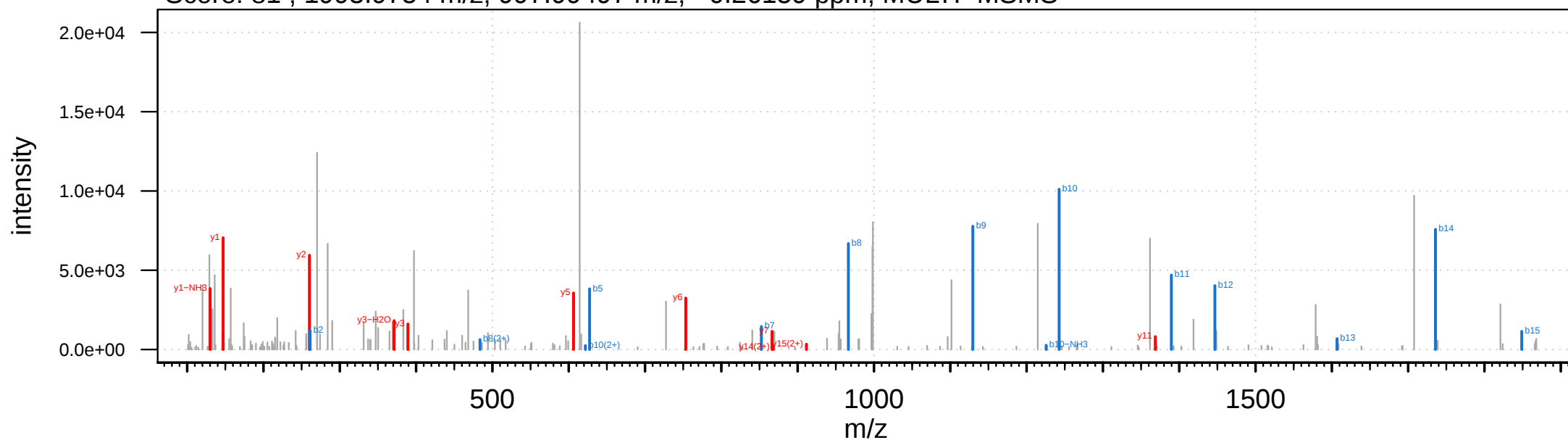


Raw File: 20101224_Velos1_TaGe_SA_HeLa_01
 Scan Number: 20424
 Proteins:
 ENST00000602845_chr3:196669588-196669887:+



(ac)MSPLRPQNYLFGCELK

Score: 81 ; 1993.9754 m/z; 997.99497 m/z; -0.26159 ppm; MULTI-MSMS

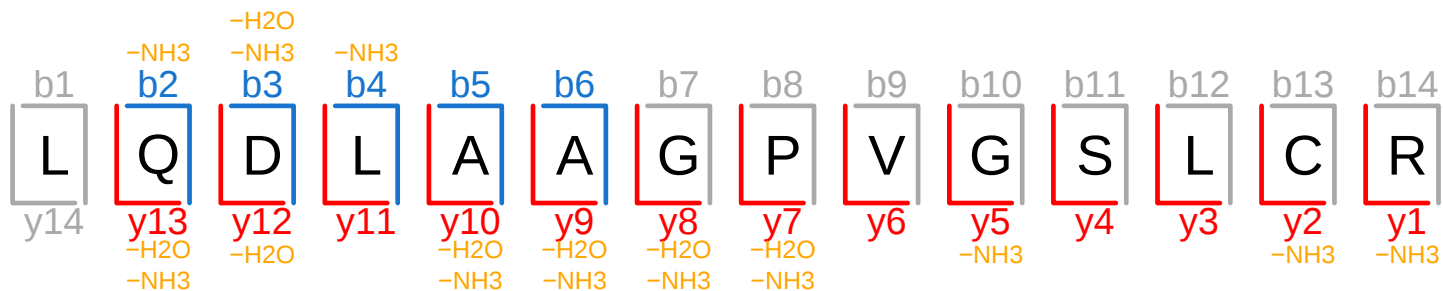


Raw File: 20101224_Velos1_TaGe_SA_HeLa_01

Scan Number: 37071

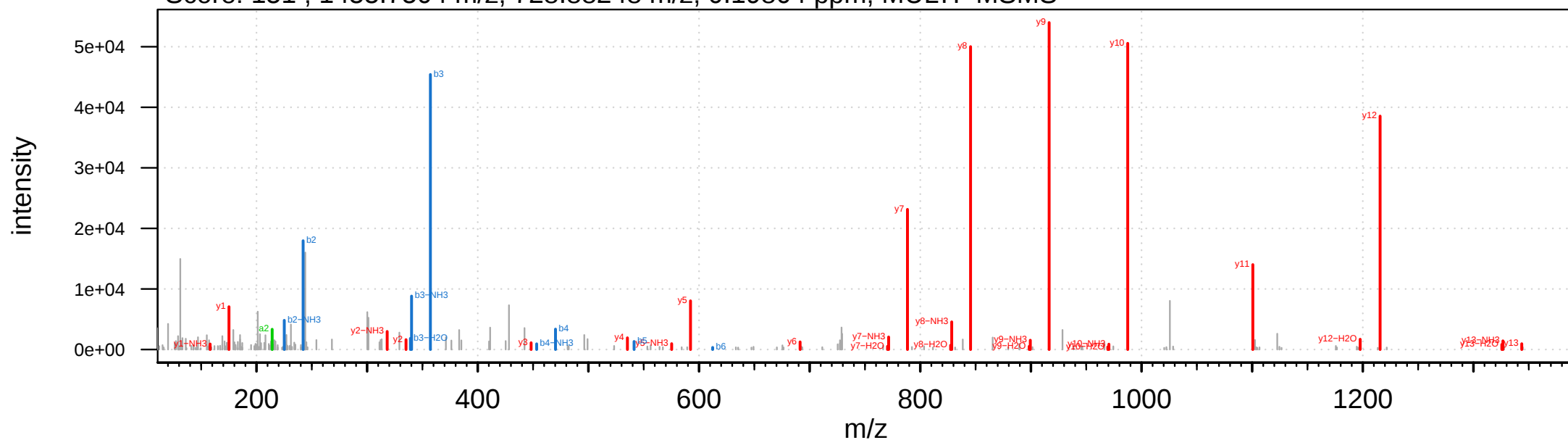
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



LQDLAAGPVGSLCR

Score: 151 ; 1455.7504 m/z; 728.88248 m/z; 0.19804 ppm; MULTI-MSMS

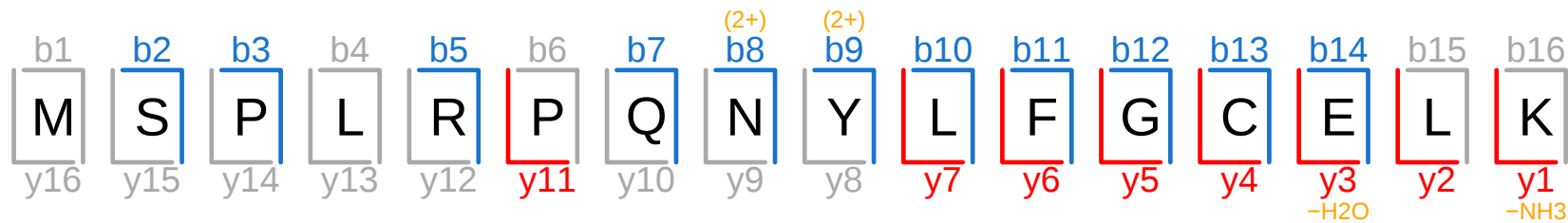


Raw File: 20101224_Velos1-TaGe_SA-HepG2_1

Scan Number: 20233

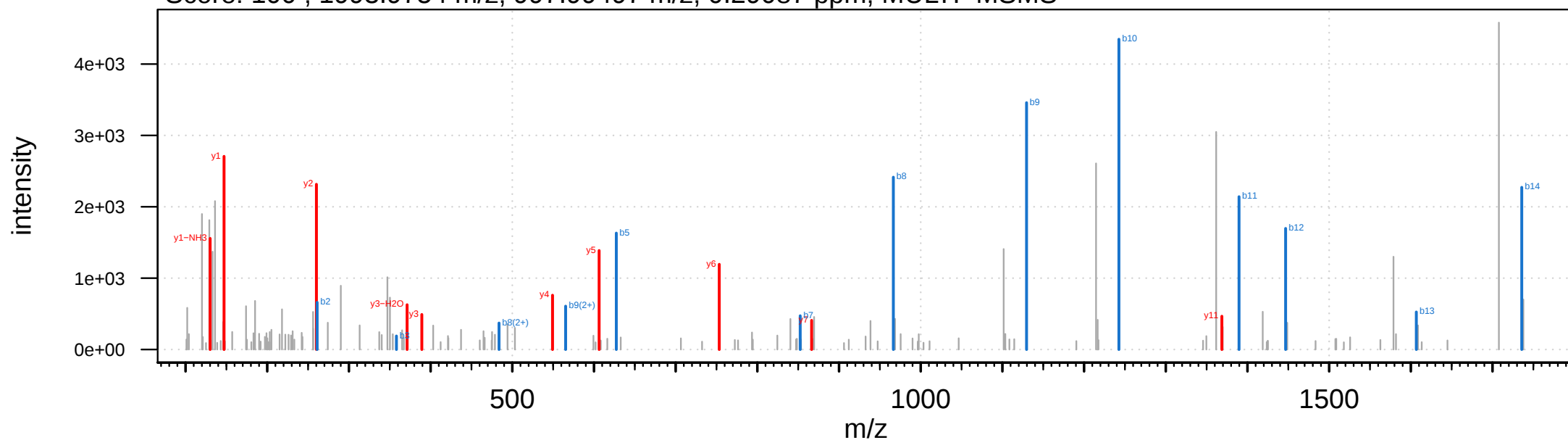
Proteins:

ENST00000602845_chr3:196669588-196669887:+



(ac)MSPLRPQNYLFGCELK

Score: 100 ; 1993.9754 m/z; 997.99497 m/z; 0.29087 ppm; MULTI-MSMS

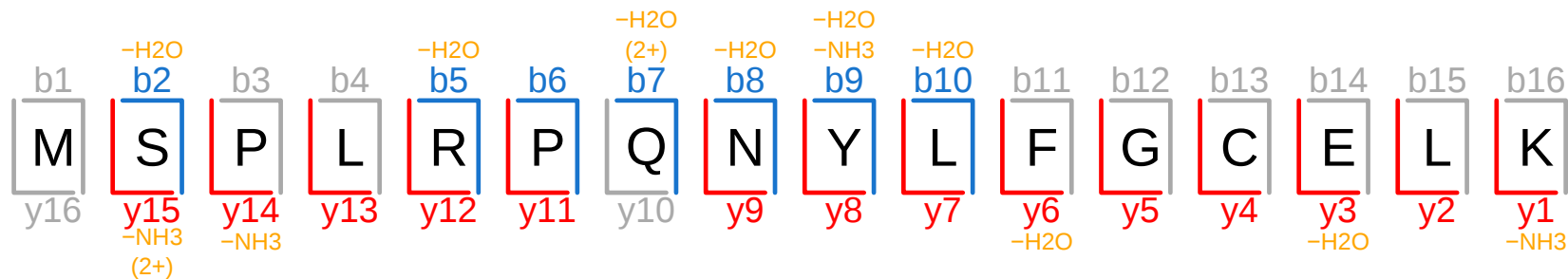


Raw File: 20101224_Velos1_TaGe_SA_HepG2_1

Scan Number: 37022

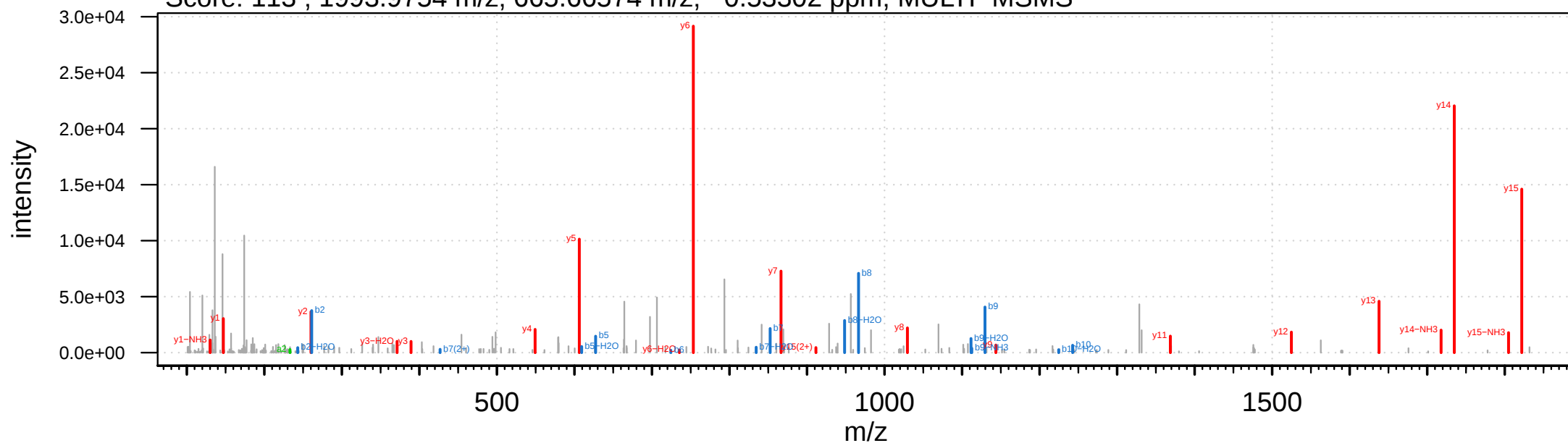
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 113 ; 1993.9754 m/z; 665.66574 m/z; -0.53302 ppm; MULTI-MSMS

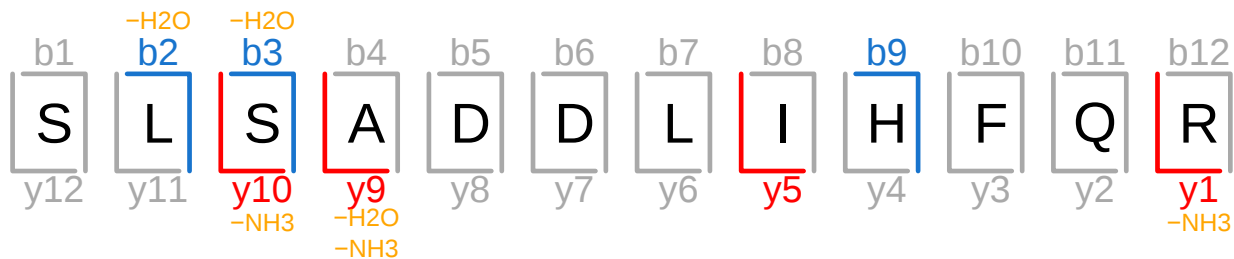


Raw File: 20101224_Velos1_TaGe_SA_HepG2_1

Scan Number: 37032

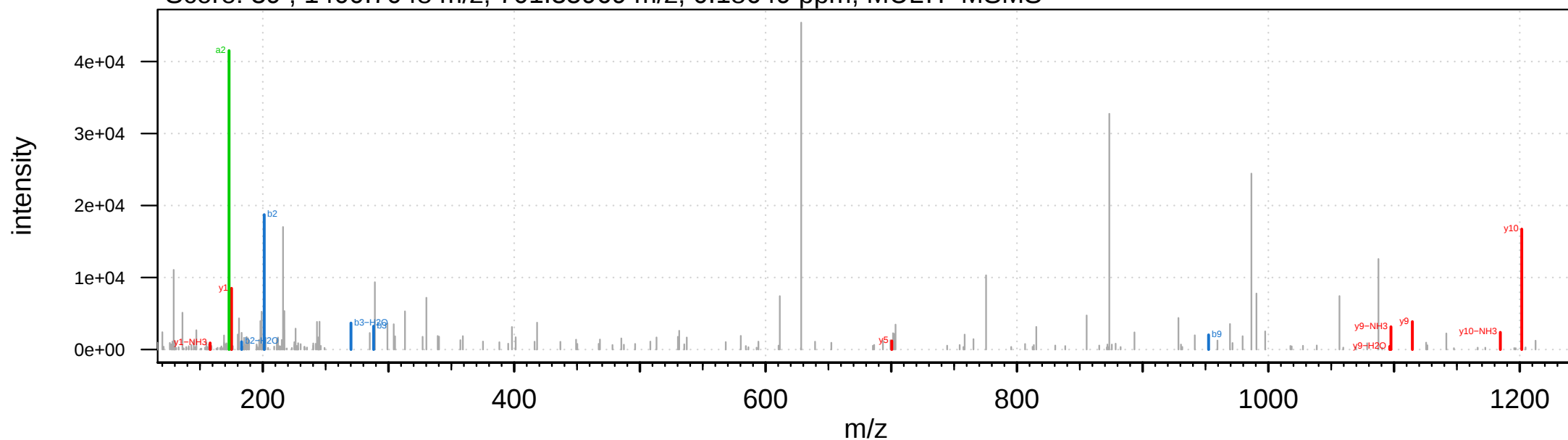
Proteins:

TCONS_I2_00008829_ch15:92829088-92829258:+



SLSADDLIHFQR

Score: 39 ; 1400.7048 m/z; 701.35969 m/z; 0.18649 ppm; MULTI-MSMS

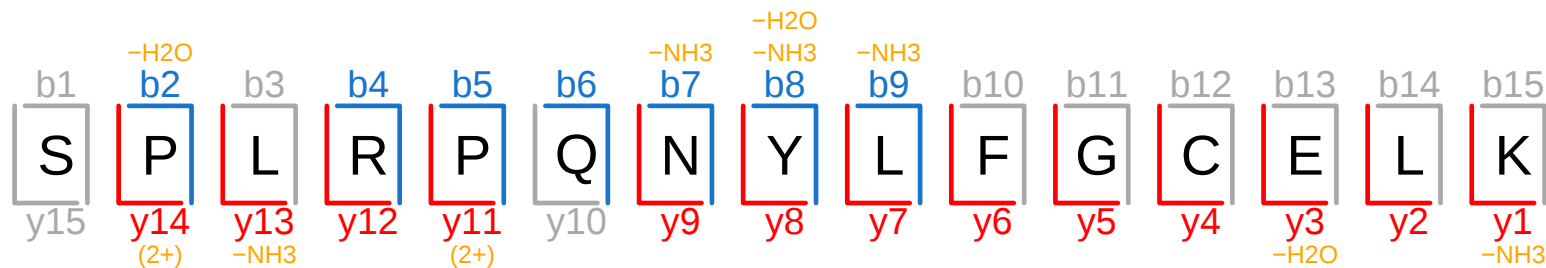


Raw File: 20101224_Velos1-TaGe_SA-HepG2_1

Scan Number: 17037

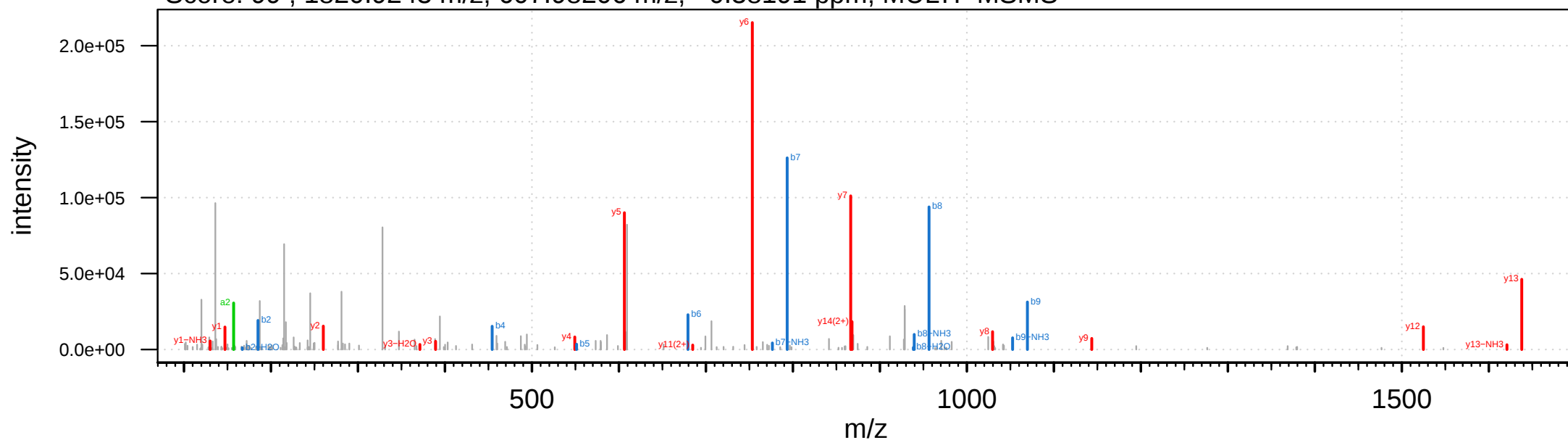
Proteins:

TCONS_I2_00002988_chr10:38499673-38501345:+



SPLRPQNYLFGCELK

Score: 99 ; 1820.9243 m/z; 607.98206 m/z; -0.38191 ppm; MULTI-MSMS

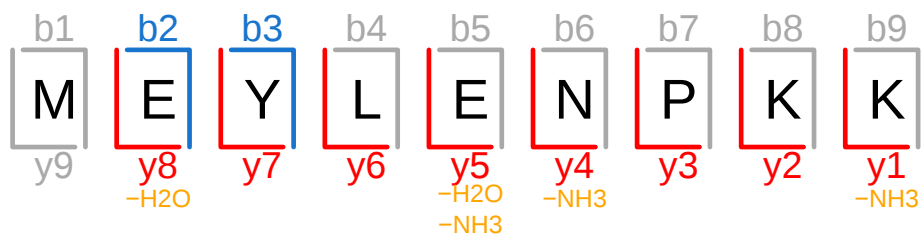


Raw File: 20101224_Velos1_TaGe_SA_HepG2_1

Scan Number: 23847

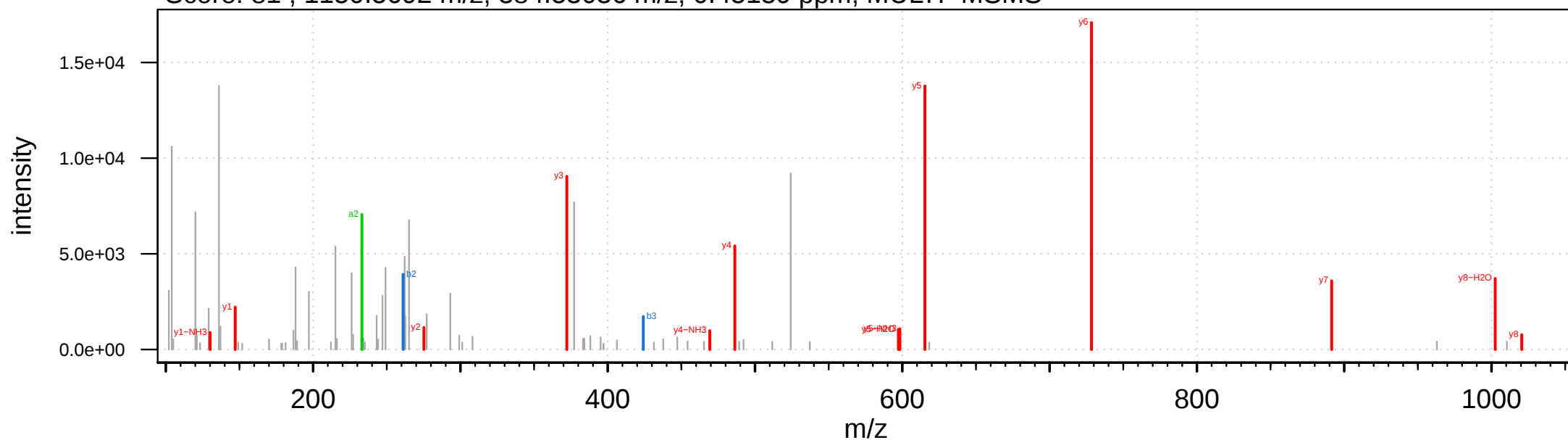
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



MEYLENPKK

Score: 81 ; 1150.5692 m/z; 384.53036 m/z; 0.45139 ppm; MULTI-MSMS

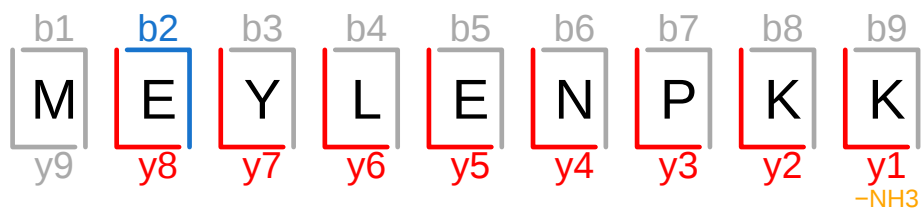


Raw File: 20100618_Velos1_TaGe_SA_U2OS_6

Scan Number: 5833

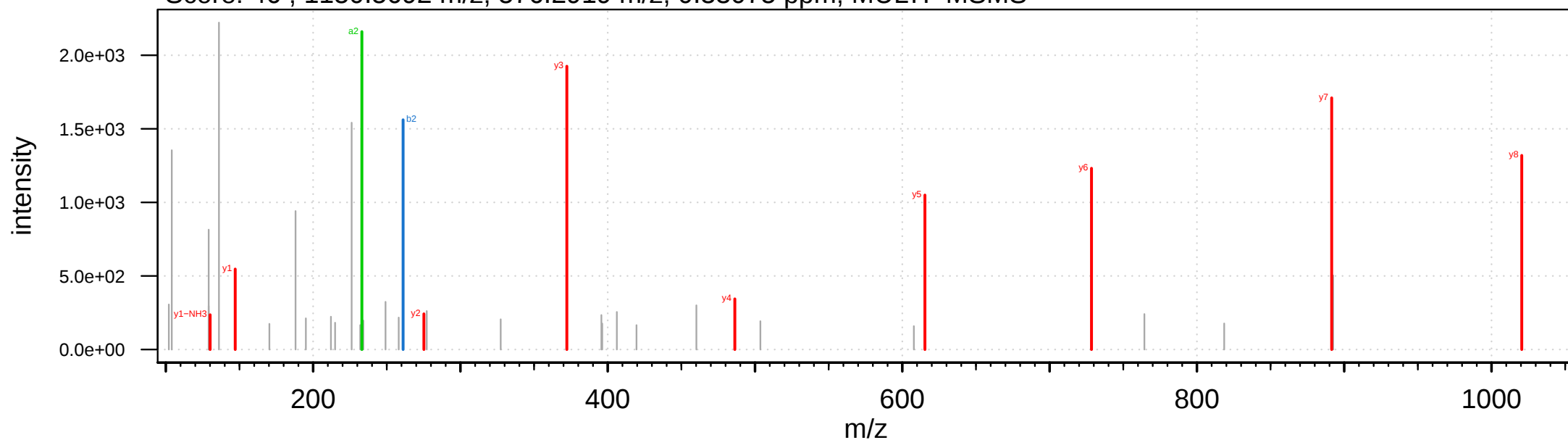
Proteins:

ENST00000518958_chr8:120630400-120630519:-



MEYLENPKK

Score: 46 ; 1150.5692 m/z; 576.2919 m/z; 0.33075 ppm; MULTI-MSMS

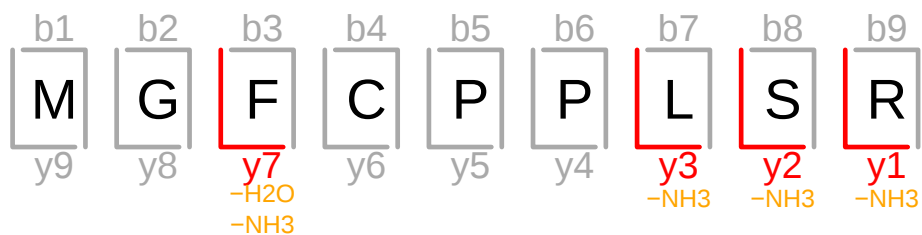


Raw File: 20100618_Velos1_TaGe_SA_U2OS_6

Scan Number: 5845

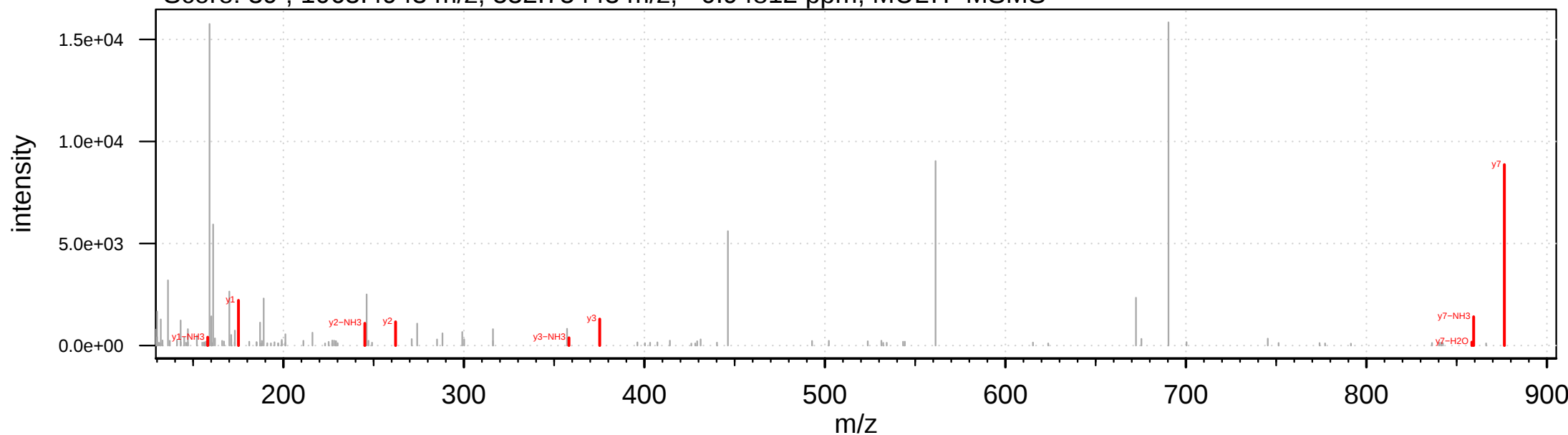
Proteins:

ENST00000518958_chr8:120630400-120630519:-



MGFCPPLSR

Score: 39 ; 1063.4943 m/z; 532.75443 m/z; -0.94812 ppm; MULTI-MSMS

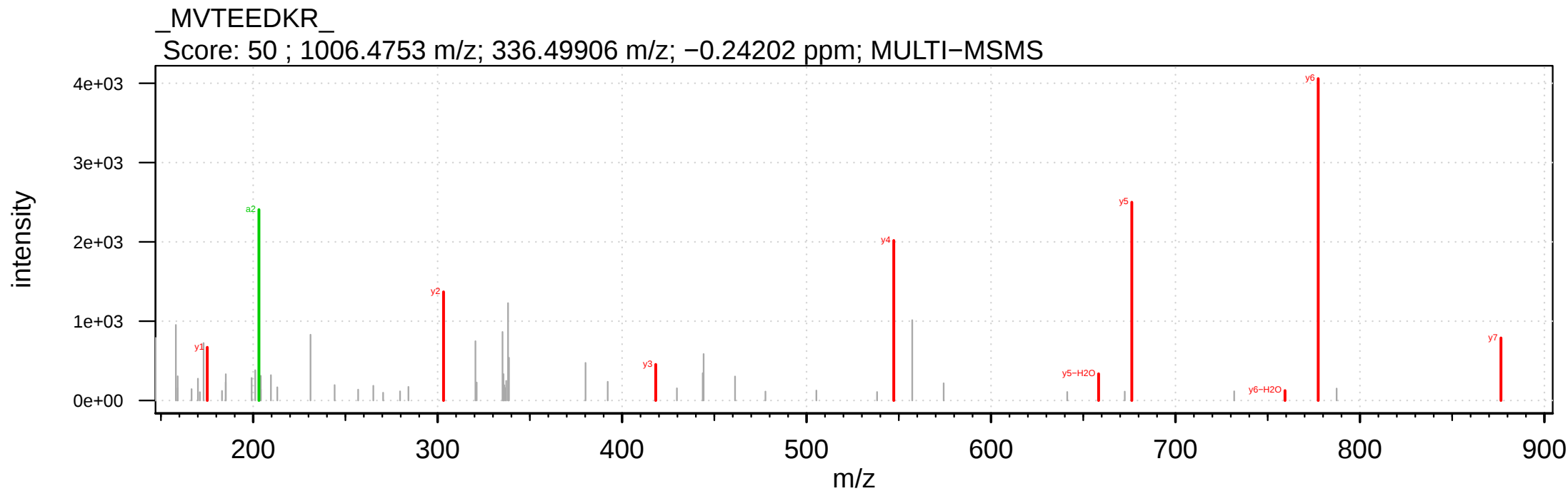
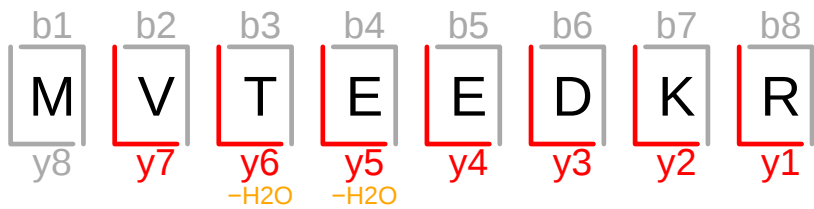


Raw File: 20101224_Velos1_TaGe_SA_HepG2_4

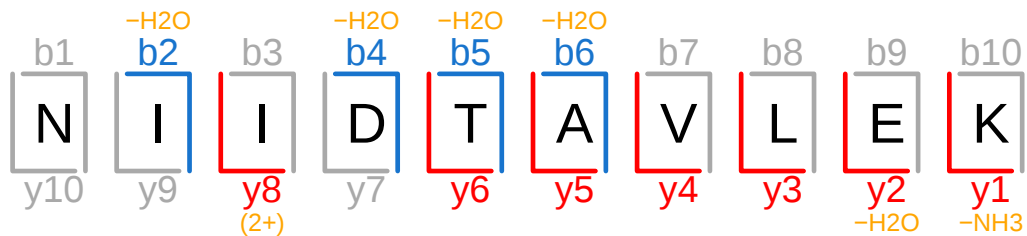
Scan Number: 13185

Proteins:

ENST00000381213_chr5:56218114-56218146:-

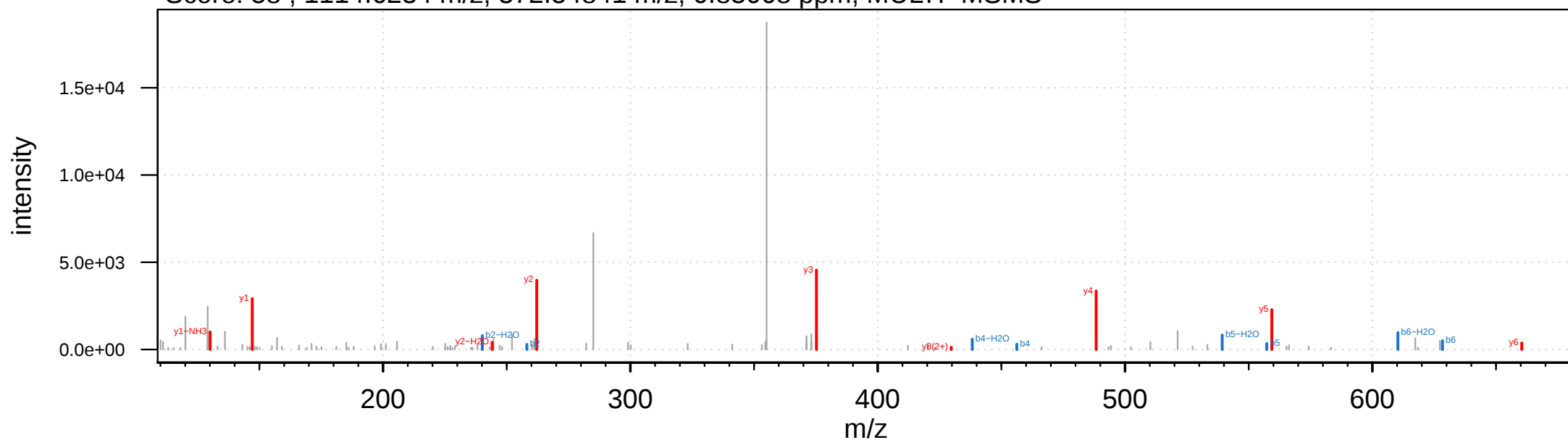


Raw File: 20101224_Velos1_TaGe_SA_HepG2_4
Scan Number: 929
Proteins:
TCONS_I2_00006768_chr13:29174091-29174123:+



NIIDTAVLEK

Score: 58 ; 1114.6234 m/z; 372.54841 m/z; 0.83068 ppm; MULTI-MSMS

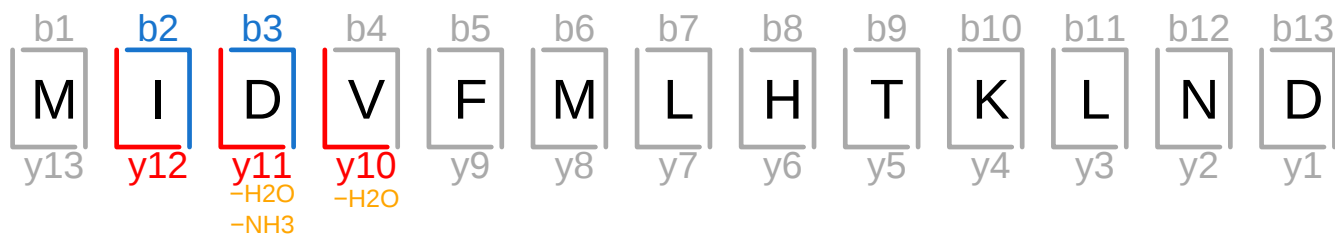


Raw File: 20101224_Velos1_TaGe_SA_HepG2_4

Scan Number: 6466

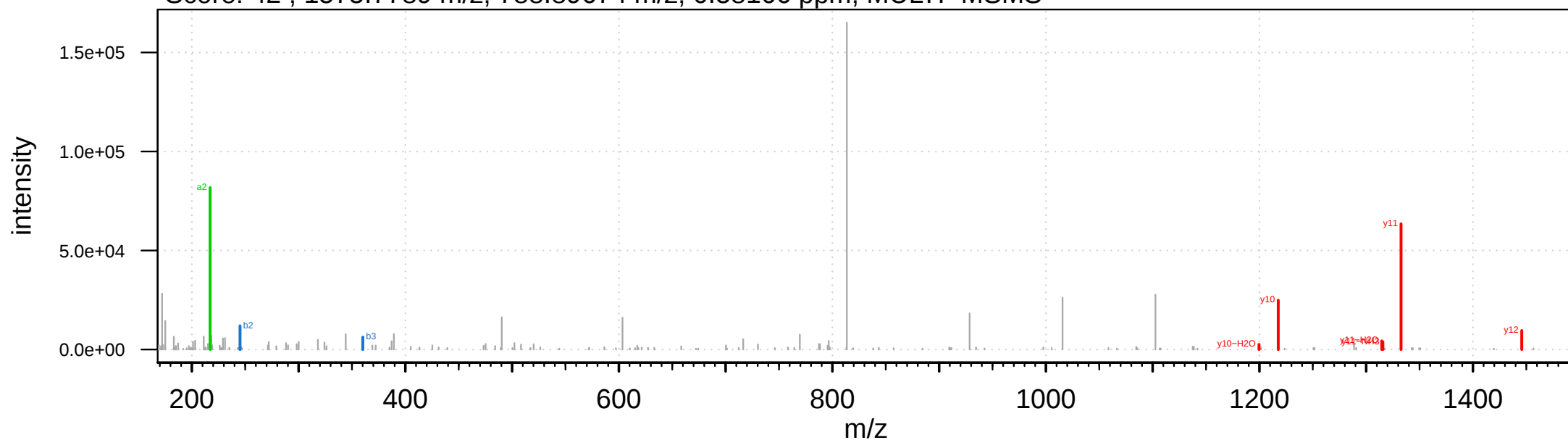
Proteins:

ENST00000497138_chr20:56806826-56807846:-



MIDVFMLHTKLND

Score: 42 ; 1575.7789 m/z; 788.89674 m/z; 0.58166 ppm; MULTI-MSMS

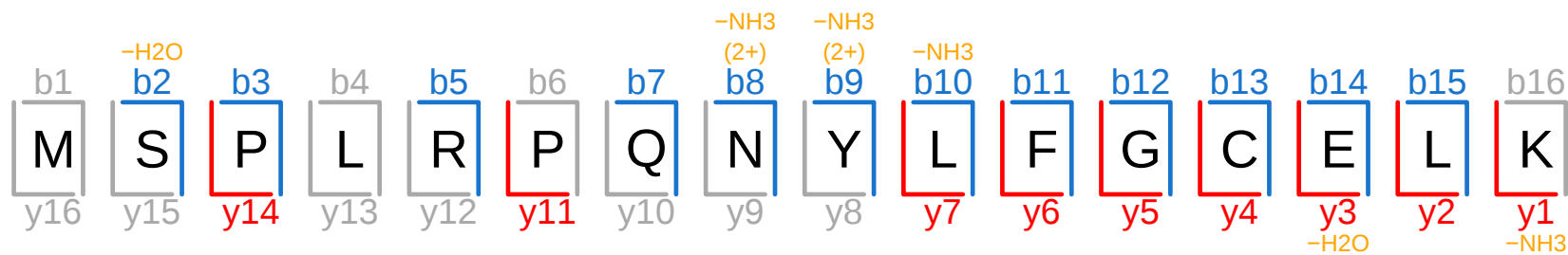


Raw File: 20100609_Velos1_TaGe_SA_293_1

Scan Number: 30776

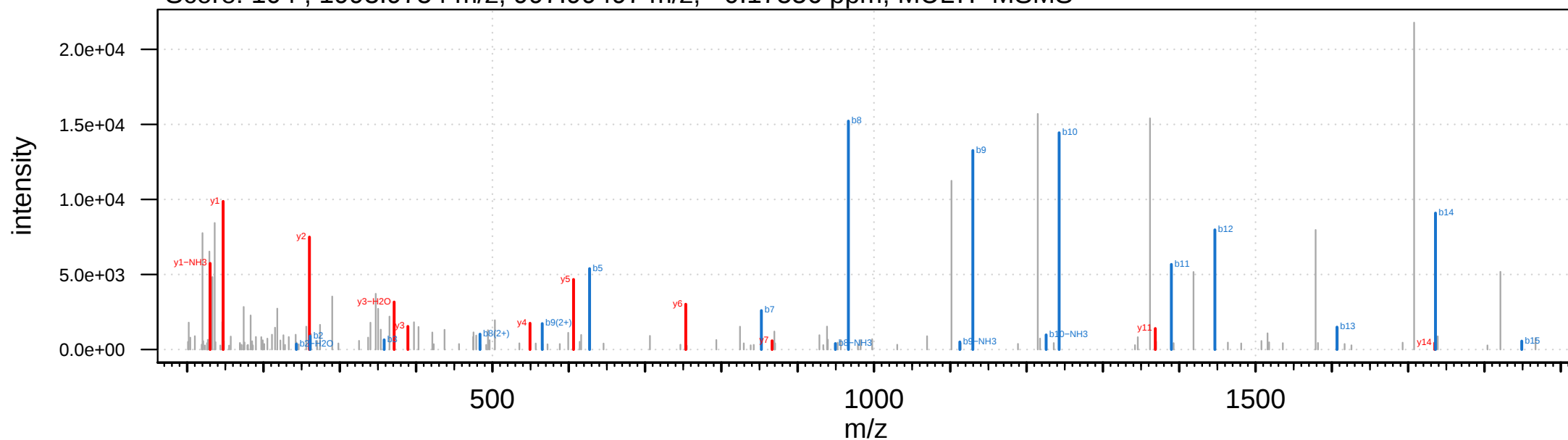
Proteins:

ENST00000335670_chr15:60788697-60788738:-



(ac)MSPLRPQNYLFGCELK

Score: 104 ; 1993.9754 m/z; 997.99497 m/z; -0.17556 ppm; MULTI-MSMS

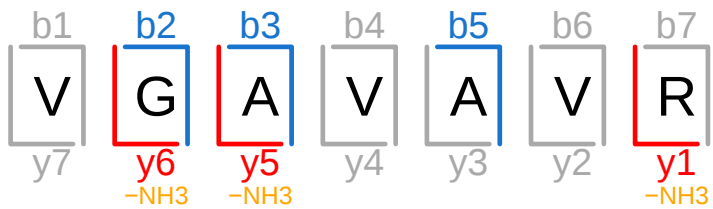


Raw File: 20100609_Velos1_TaGe_SA_293_1

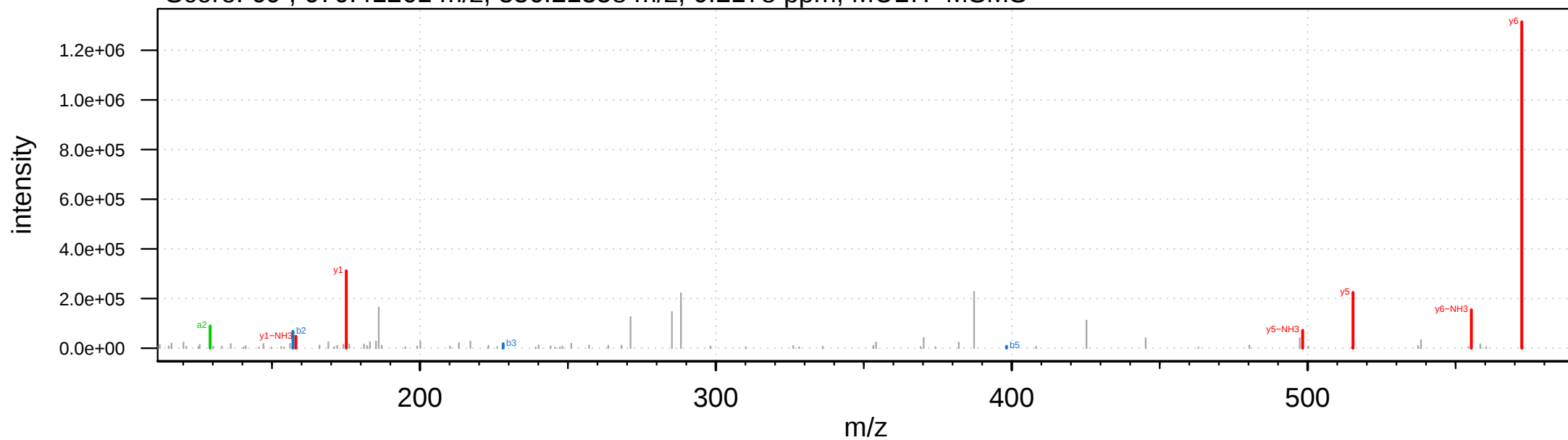
Scan Number: 36580

Proteins:

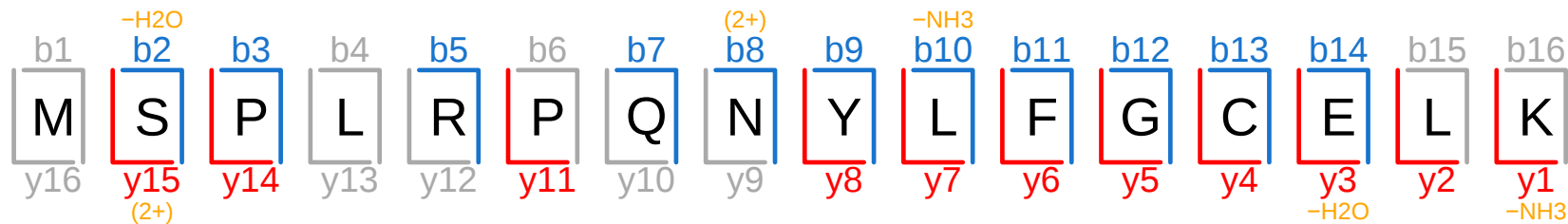
TCONS_I2_00008829_chr15:92829088-92829258:+



VGAVAVR
 Score: 69 ; 670.41261 m/z; 336.21358 m/z; 0.2178 ppm; MULTI-MSMS

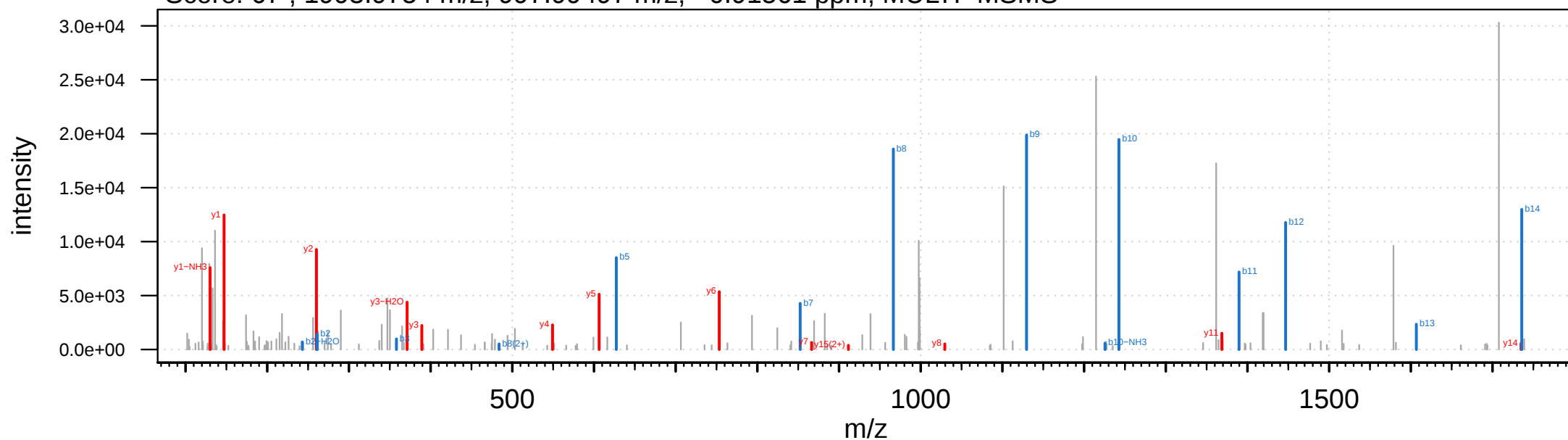


Raw File: 20100609_Velos1_TaGe_SA_293_1
 Scan Number: 2254
 Proteins:
 TCONS_I2_00030545_chrX:79544539-79546436:-



(ac)MSPLRPQNYLFGCELK

Score: 67 ; 1993.9754 m/z; 997.99497 m/z; -0.91561 ppm; MULTI-MSMS

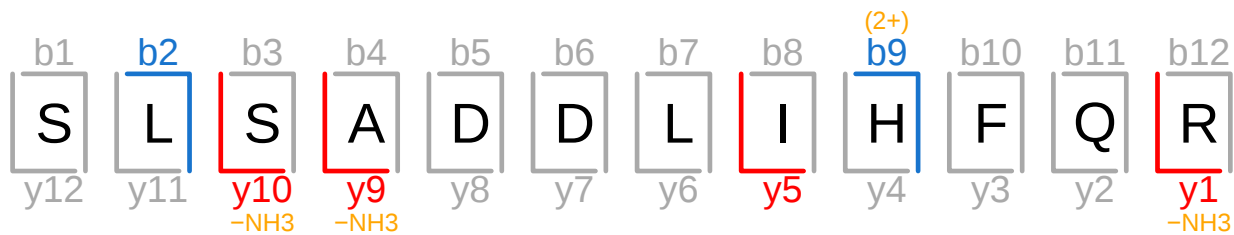


Raw File: 20100604_Velos1_TaGe_SA_A549_1

Scan Number: 34597

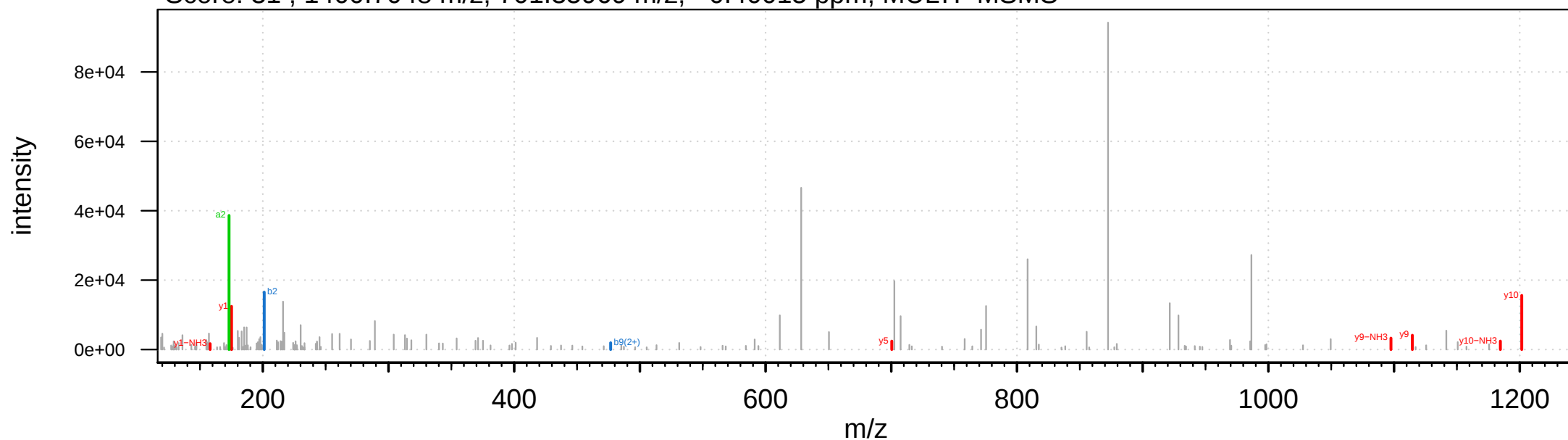
Proteins:

TCONS_I2_00008829_ch15:92829088-92829258:+



SLSADDLIHFQR

Score: 31 ; 1400.7048 m/z; 701.35969 m/z; -0.40015 ppm; MULTI-MSMS

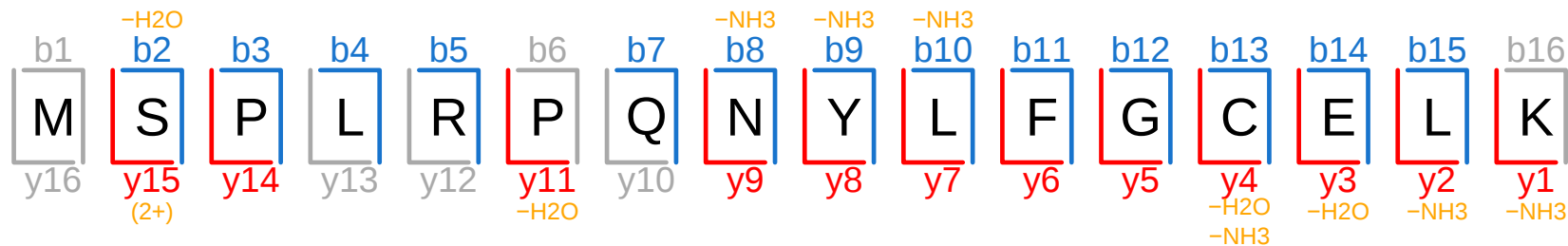


Raw File: 20100604_Velos1_TaGe_SA_A549_1

Scan Number: 16586

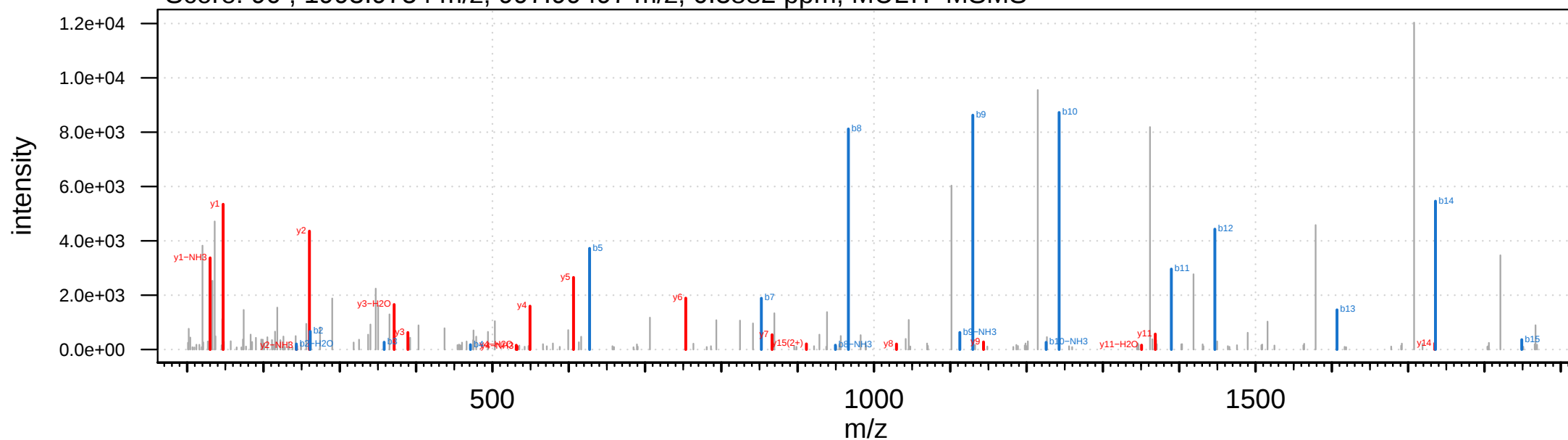
Proteins:

TCONS_I2_00002988_chr10:38499673-38501345:+



(ac)MSPLRPQNYLFGCELK

Score: 90 ; 1993.9754 m/z; 997.99497 m/z; 0.3882 ppm; MULTI-MSMS

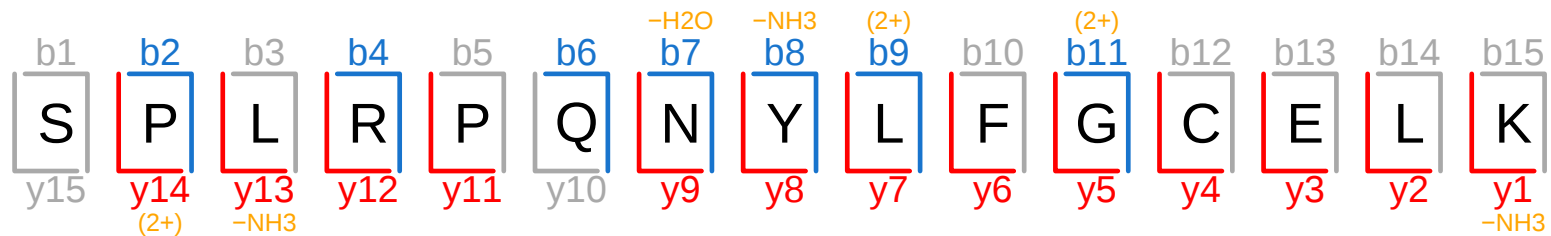


Raw File: 20100611_Velos1_TaGe_SA_Hela_2

Scan Number: 25725

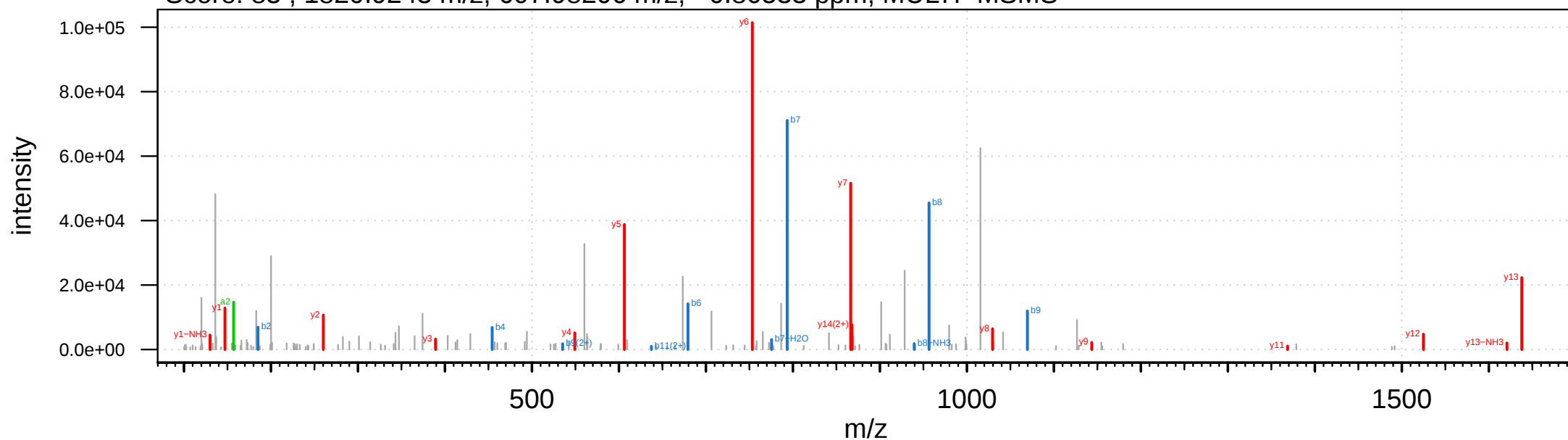
Proteins:

TCONS_I2_00008829_ch15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 83 ; 1820.9243 m/z; 607.98206 m/z; -0.80533 ppm; MULTI-MSMS

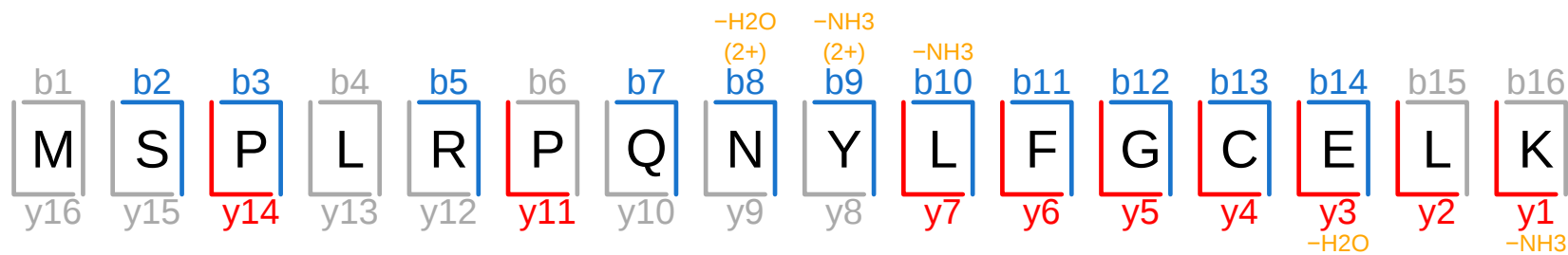


Raw File: 20100611_Velos1_TaGe_SA_Hela_2

Scan Number: 16815

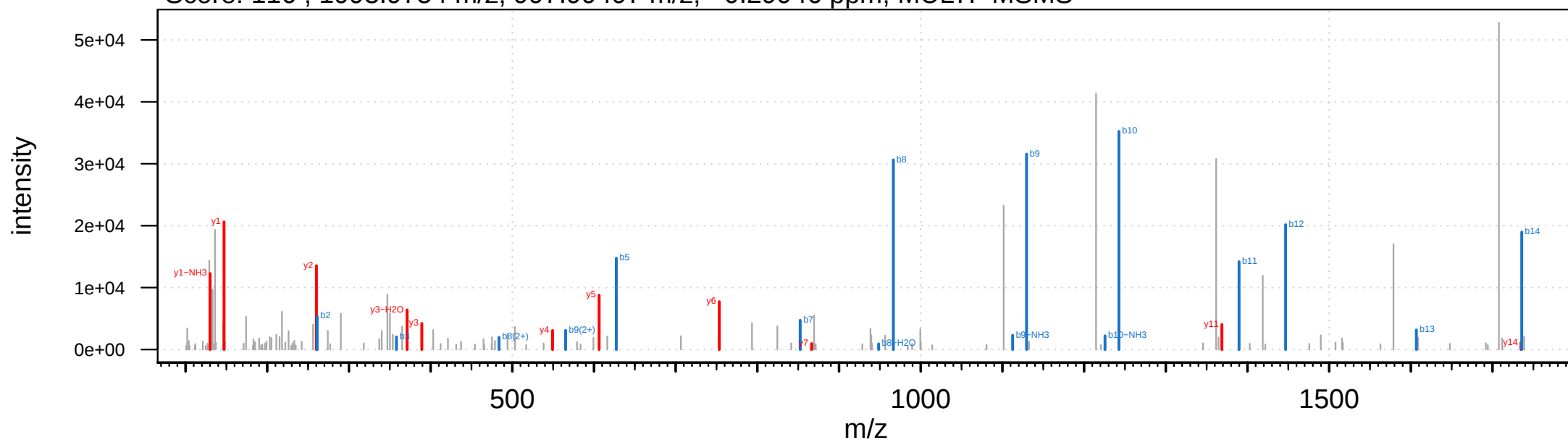
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 116 ; 1993.9754 m/z; 997.99497 m/z; -0.29946 ppm; MULTI-MSMS

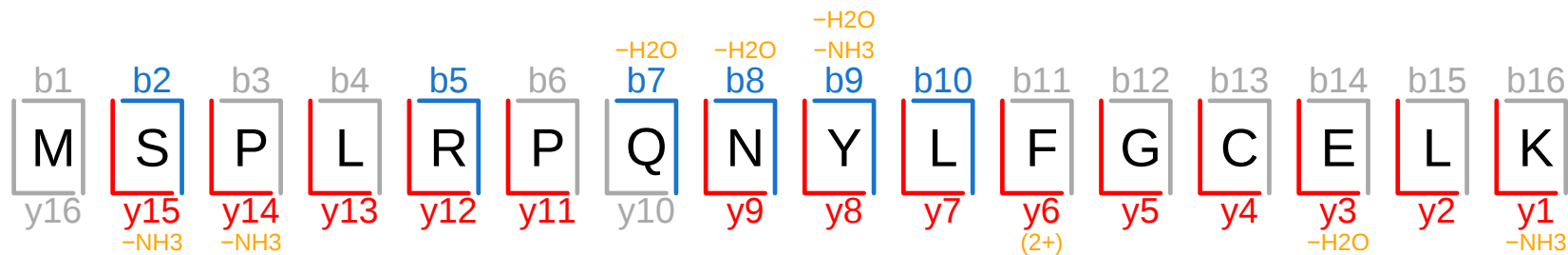


Raw File: 20100611_Velos1_TaGe_SA_HepG2_1

Scan Number: 36267

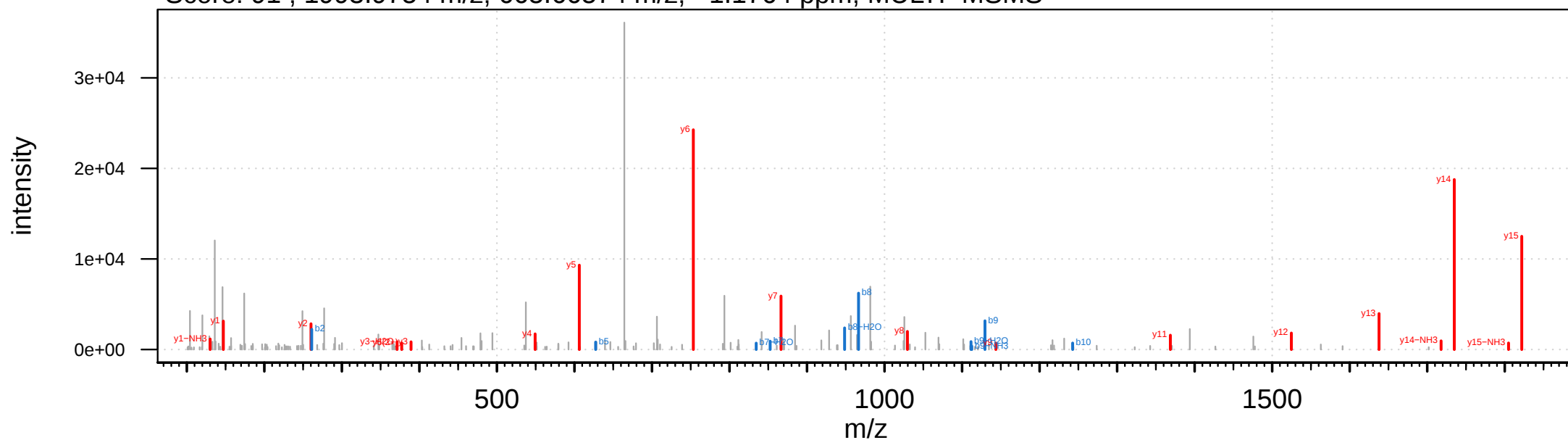
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 91 ; 1993.9754 m/z; 665.66574 m/z; -1.1764 ppm; MULTI-MSMS

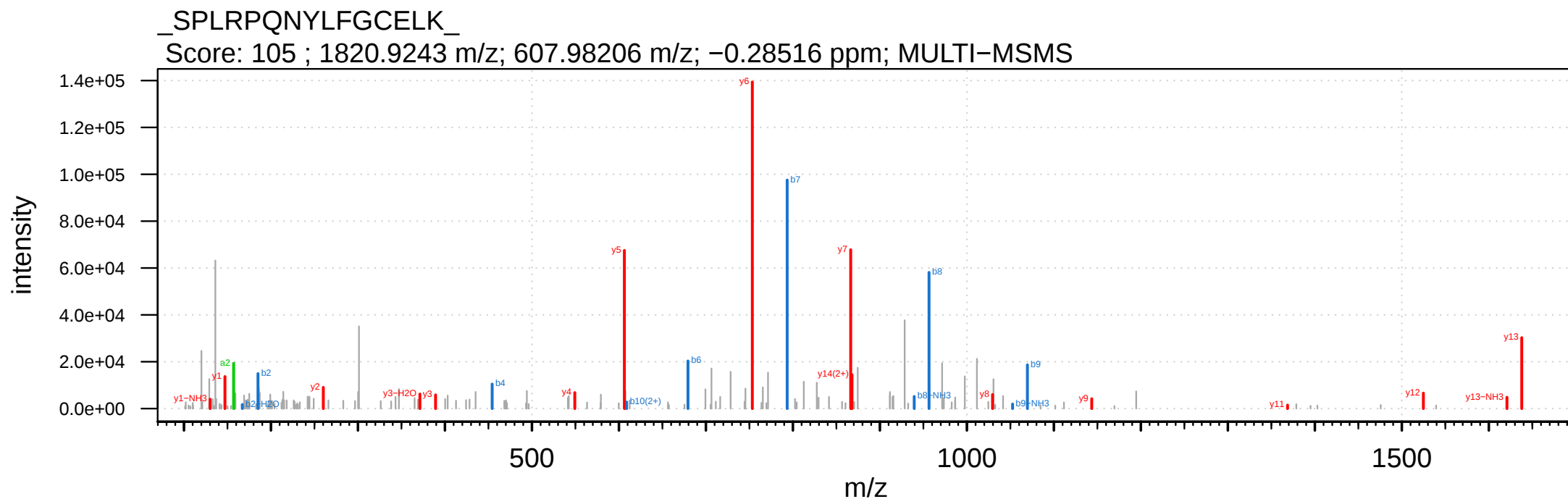
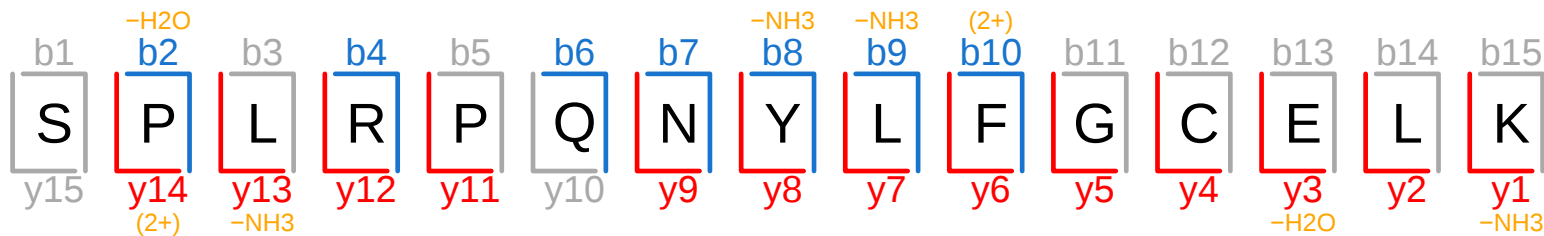


Raw File: 20100611_Velos1_TaGe_SA_HepG2_1

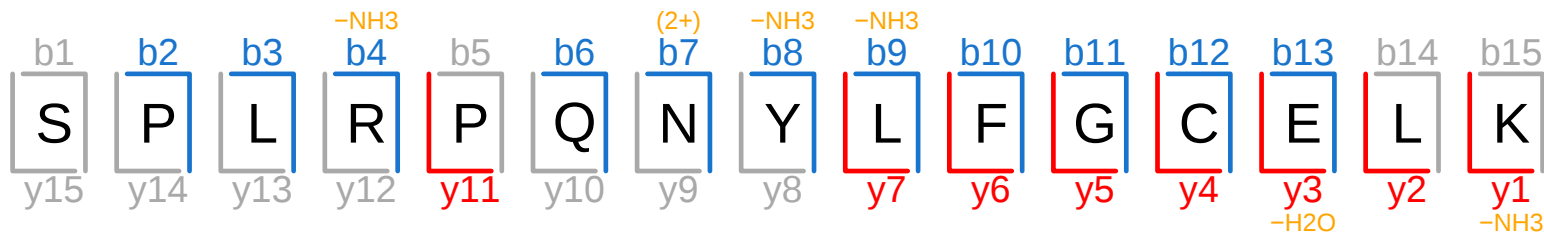
Scan Number: 36292

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

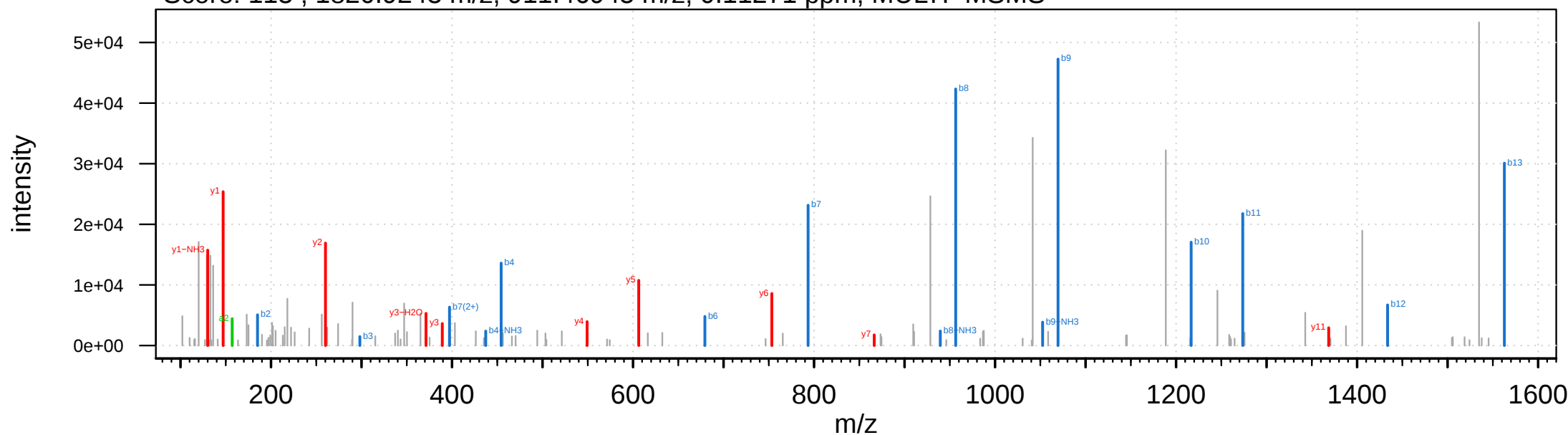


Raw File: 20100611_Velos1_TaGe_SA_HepG2_1
 Scan Number: 21108
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 113 ; 1820.9243 m/z; 911.46945 m/z; 0.11271 ppm; MULTI-MSMS

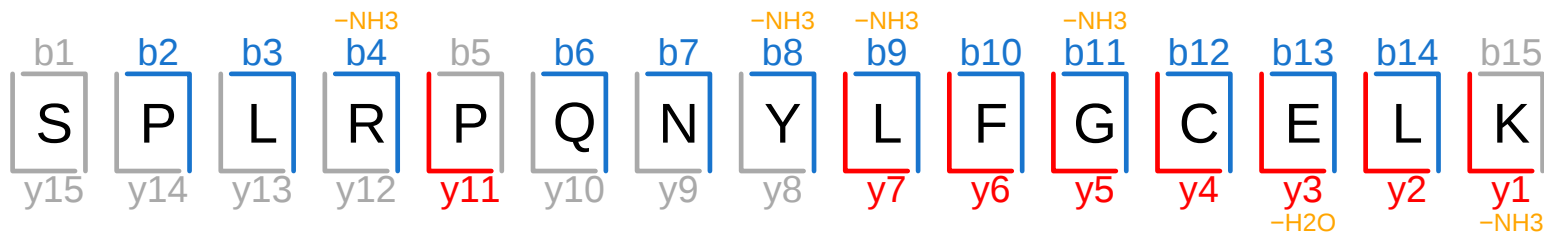


Raw File: 20100611_Velos1_TaGe_SA_HepG2_1

Scan Number: 21134

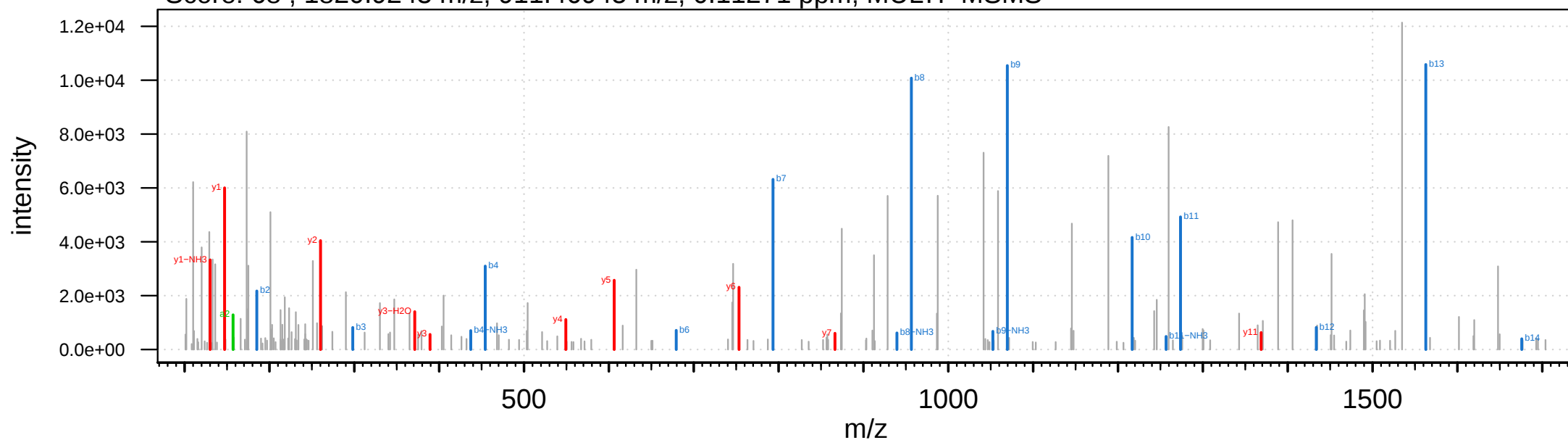
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 68 ; 1820.9243 m/z; 911.46945 m/z; 0.11271 ppm; MULTI-MSMS

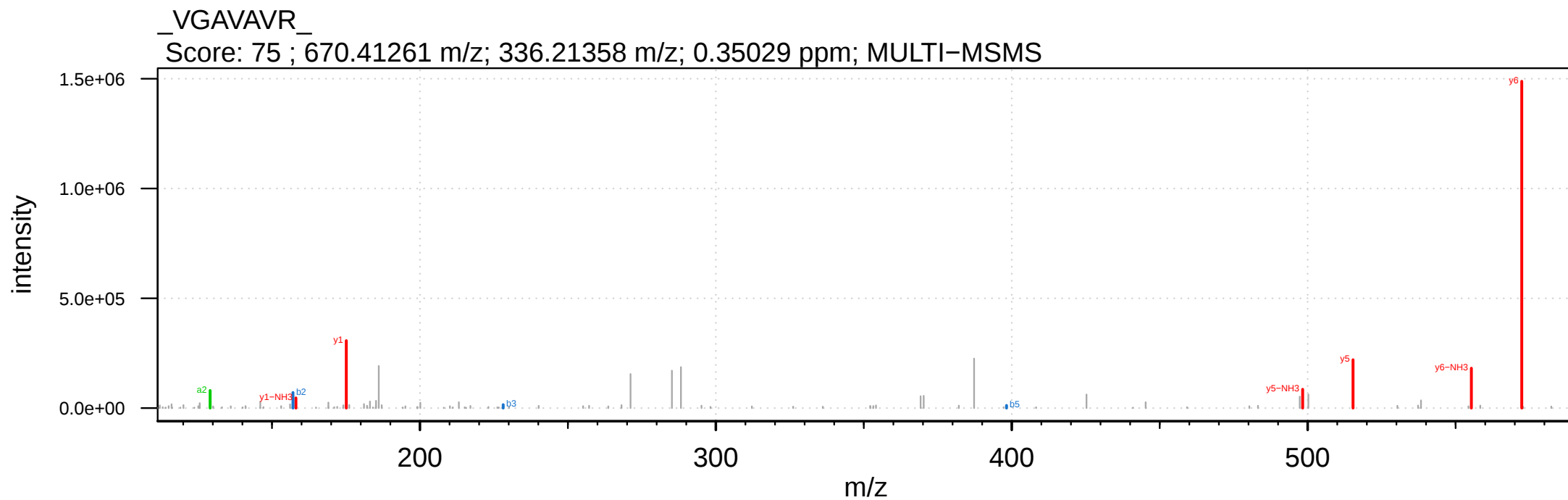
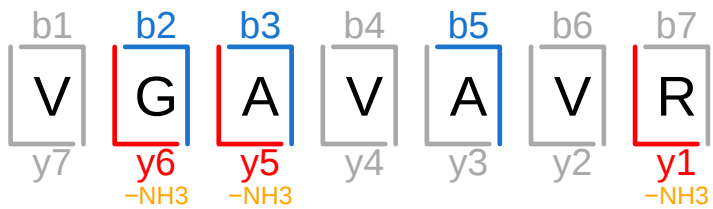


Raw File: 20100611_Velos1_TaGe_SA_HepG2_1

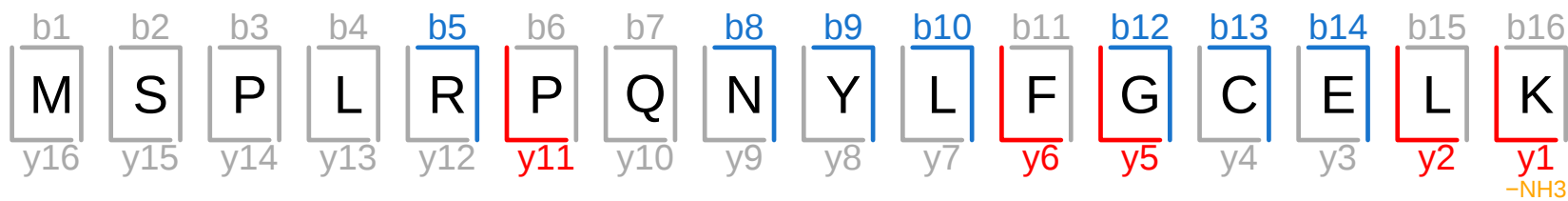
Scan Number: 21265

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

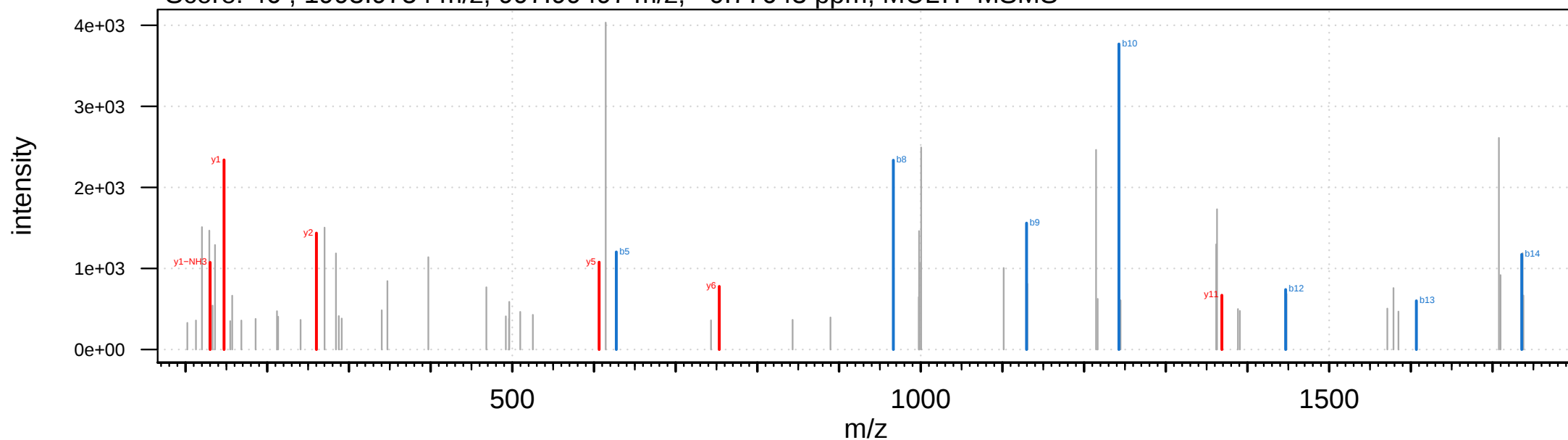


Raw File: 20100611_Velos1_TaGe_SA_HepG2_1
Scan Number: 2142
Proteins:
TCONS_I2_00030545_chrX:79544539-79546436:-



(ac)MSPLRPQNYLFGCELK

Score: 46 ; 1993.9754 m/z; 997.99497 m/z; -0.77643 ppm; MULTI-MSMS

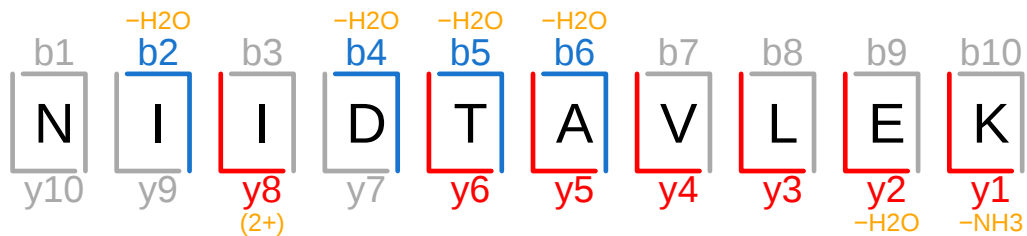


Raw File: 20100614_Velos1_TaGe_SA_Jurkat_1

Scan Number: 36266

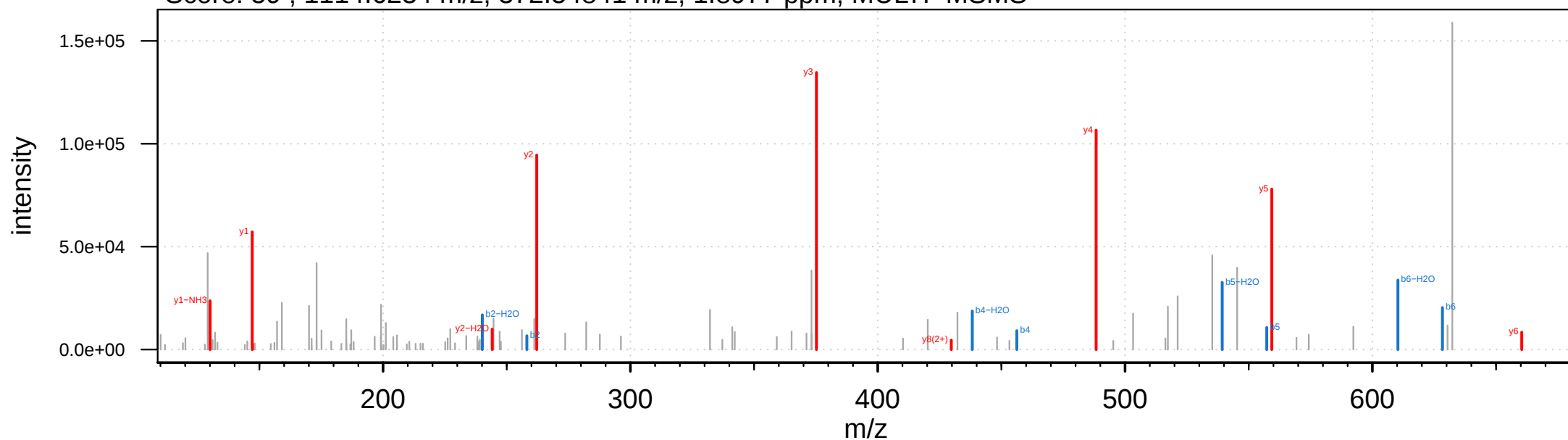
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



NIIDTAVLEK

Score: 59 ; 1114.6234 m/z; 372.54841 m/z; 1.8977 ppm; MULTI-MSMS

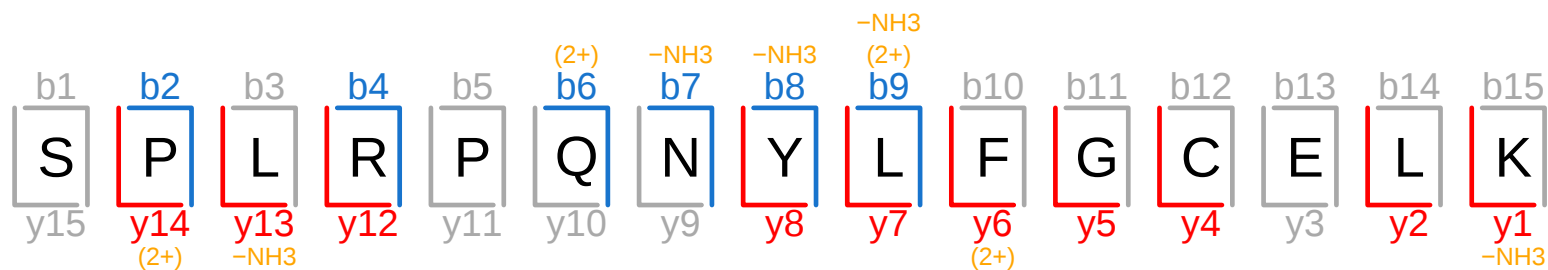


Raw File: 20100614_Velos1_TaGe_SA_Jurkat_1

Scan Number: 5508

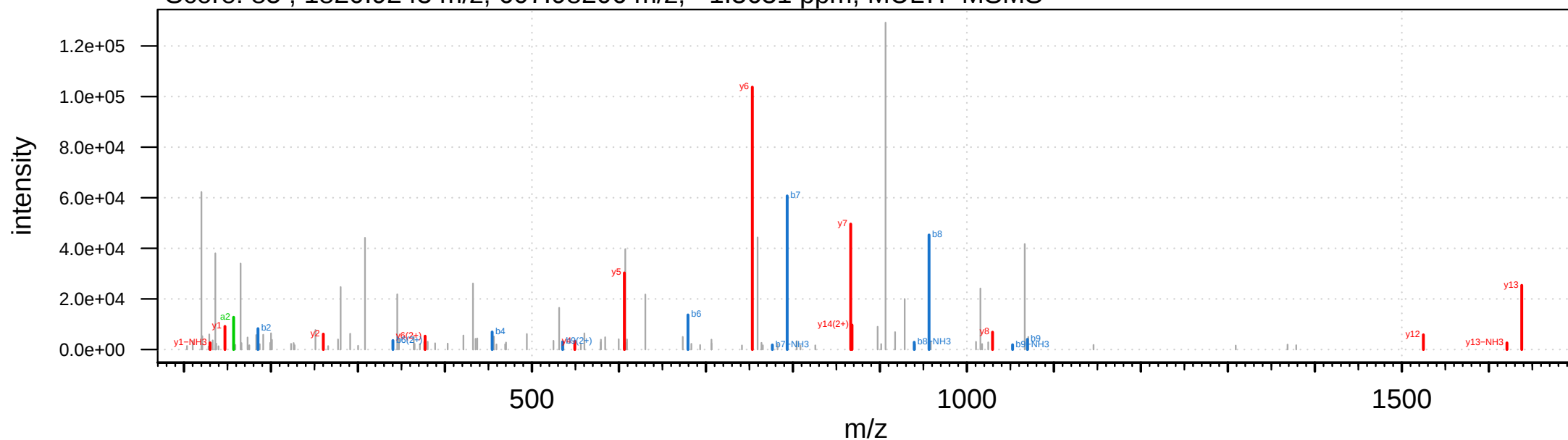
Proteins:

ENST00000497138_chr20:56806826-56807846:-



SPLRPQNYLFGCELK

Score: 85 ; 1820.9243 m/z; 607.98206 m/z; -1.3631 ppm; MULTI-MSMS

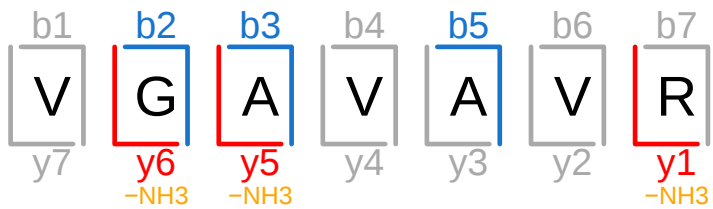


Raw File: 20100614_Velos1_TaGe_SA_Jurkat_1

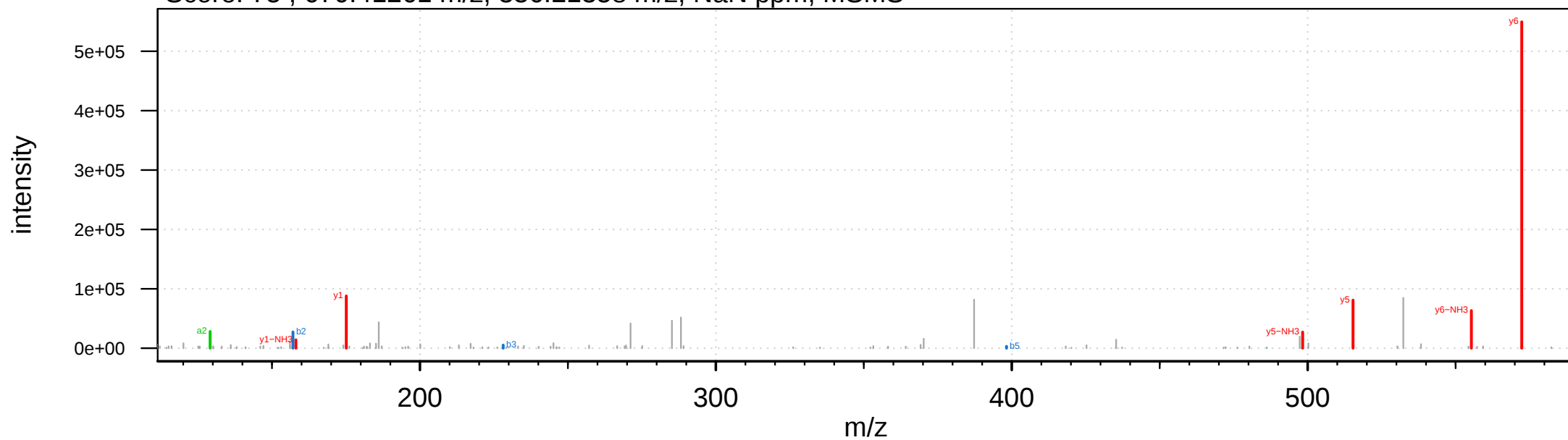
Scan Number: 22522

Proteins:

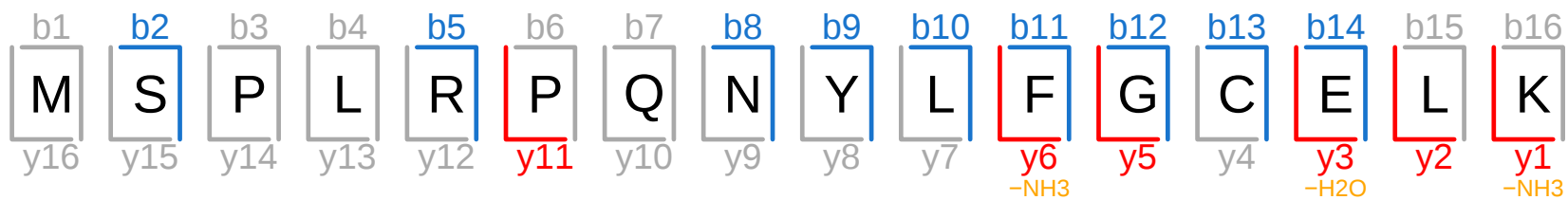
TCONS_I2_00008829_chr15:92829088-92829258:+



VGAVAVR
 Score: 75 ; 670.41261 m/z; 336.21358 m/z; NaN ppm; MSMS

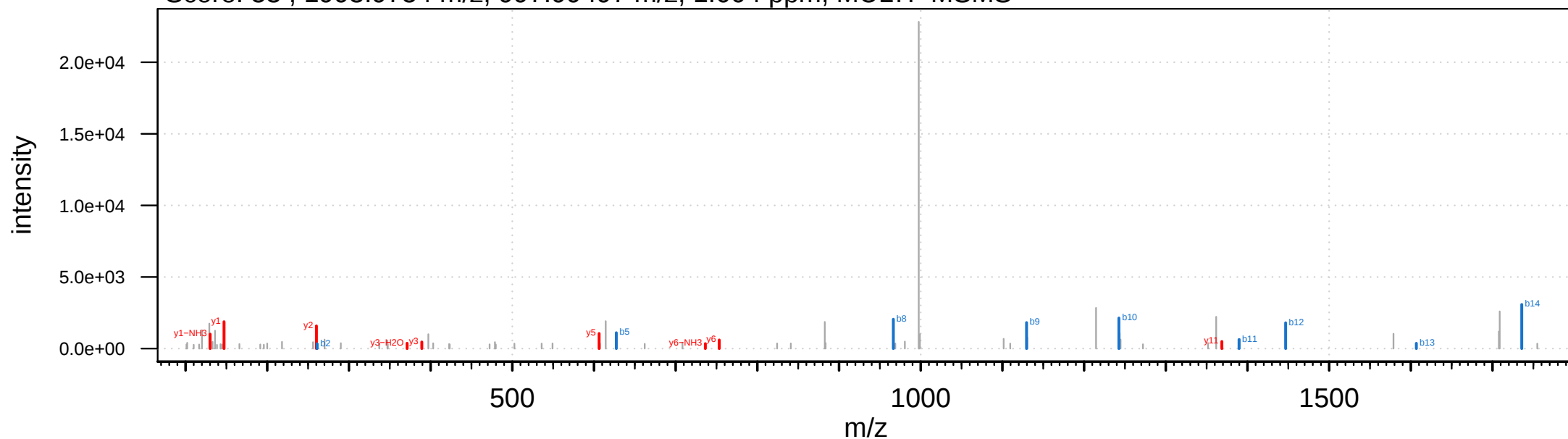


Raw File: 20100614_Velos1_TaGe_SA_Jurkat_1
 Scan Number: 2948
 Proteins:
 TCONS_I2_00030545_chrX:79544539-79546436:-



(ac)MSPLRPQNYLFGCELK

Score: 53 ; 1993.9754 m/z; 997.99497 m/z; 1.004 ppm; MULTI-MSMS

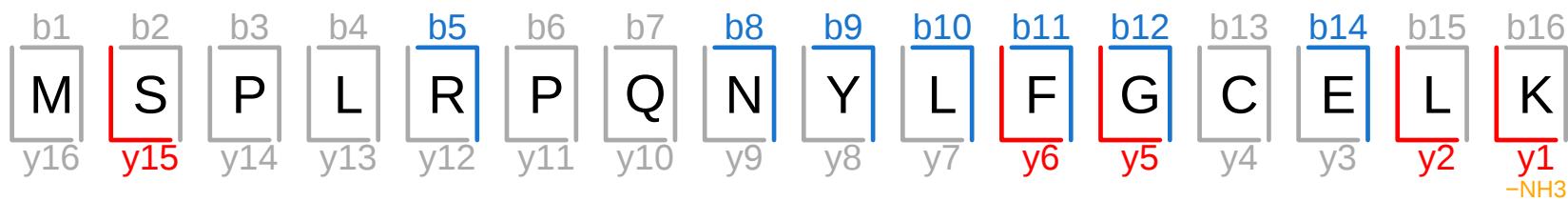


Raw File: 20100614_Velos1_TaGe_SA_K562_1

Scan Number: 37185

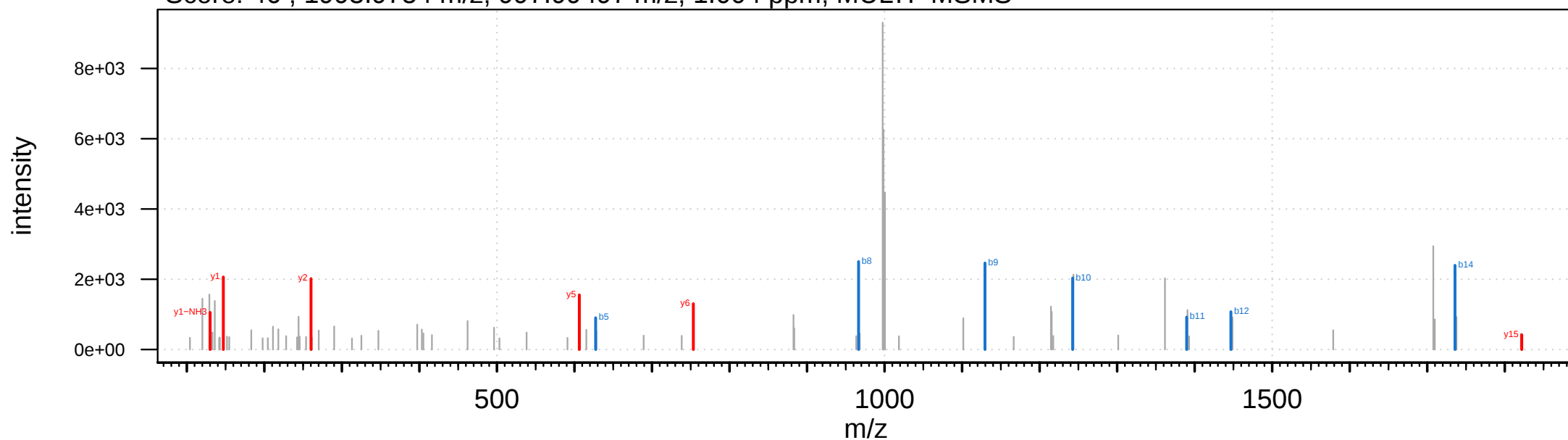
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 46 ; 1993.9754 m/z; 997.99497 m/z; 1.004 ppm; MULTI-MSMS

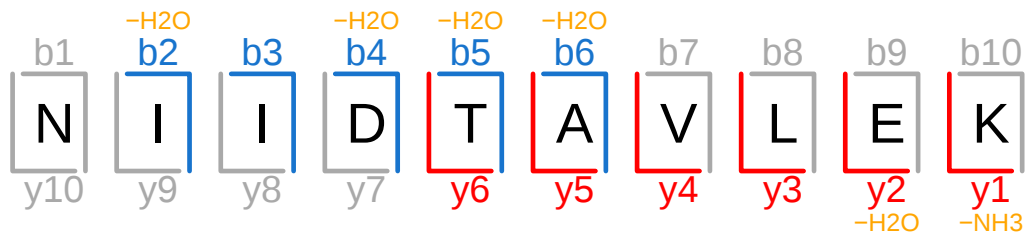


Raw File: 20100614_Velos1_TaGe_SA_K562_1

Scan Number: 37223

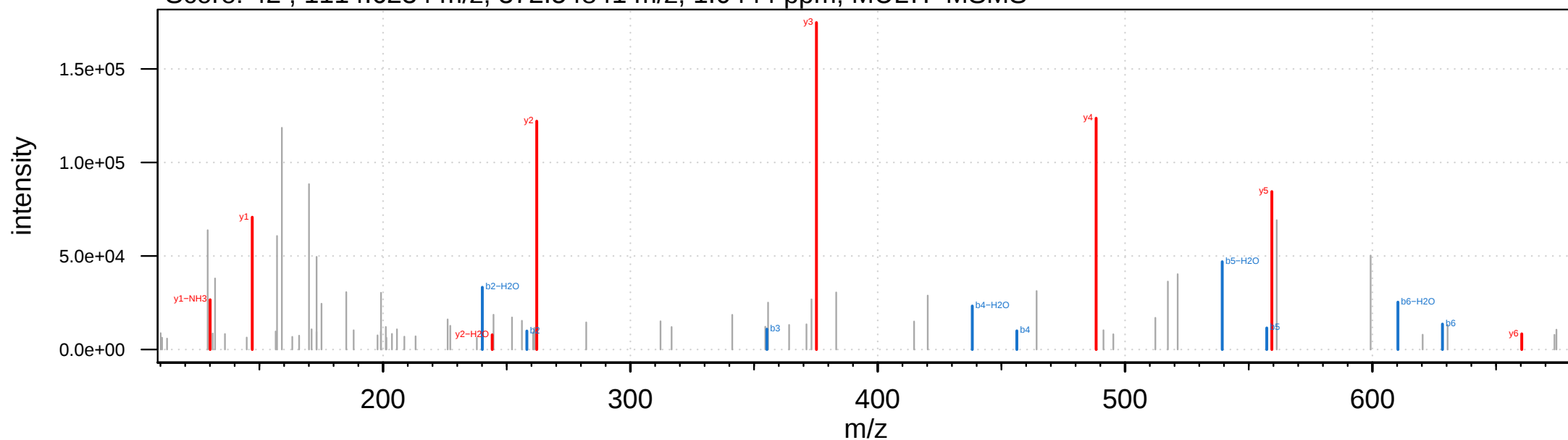
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

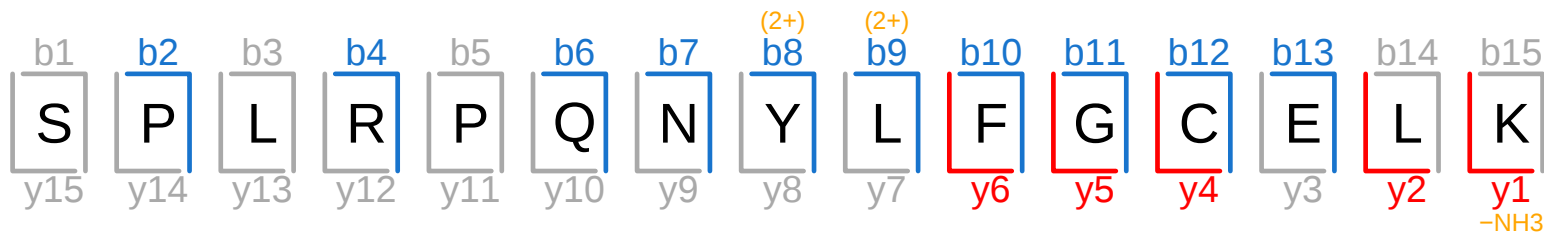


NIIDTAVLEK

Score: 42 ; 1114.6234 m/z; 372.54841 m/z; 1.0444 ppm; MULTI-MSMS

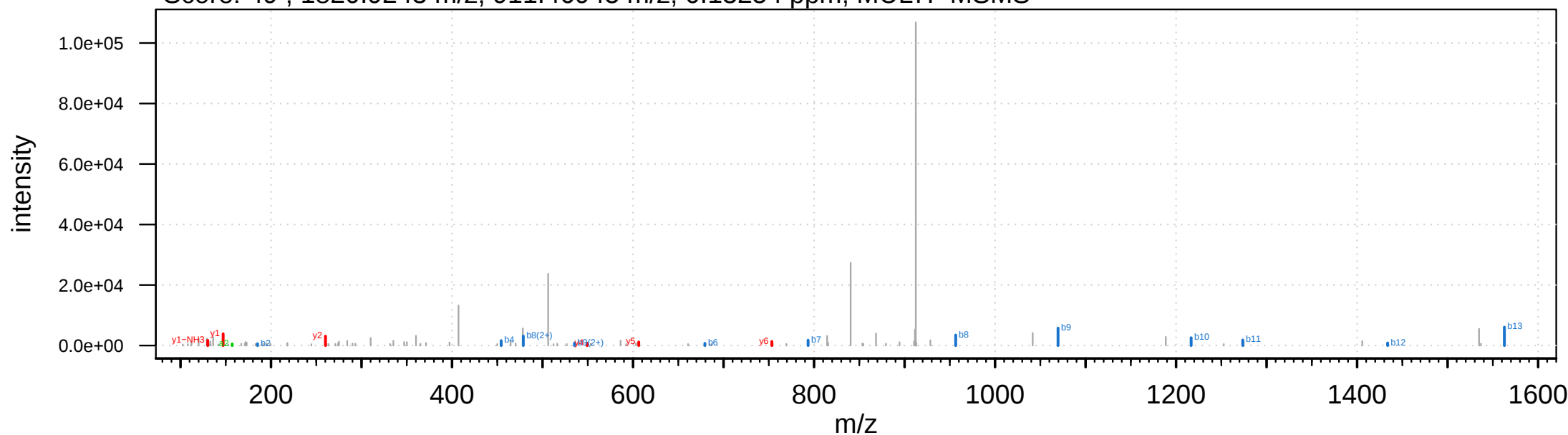


Raw File: 20100614_Velos1_TaGe_SA_K562_1
 Scan Number: 5533
 Proteins:
 ENST00000497138_chr20:56806826-56807846:-



SPLRPQNYLFGCELK

Score: 49 ; 1820.9243 m/z; 911.46945 m/z; 0.13254 ppm; MULTI-MSMS

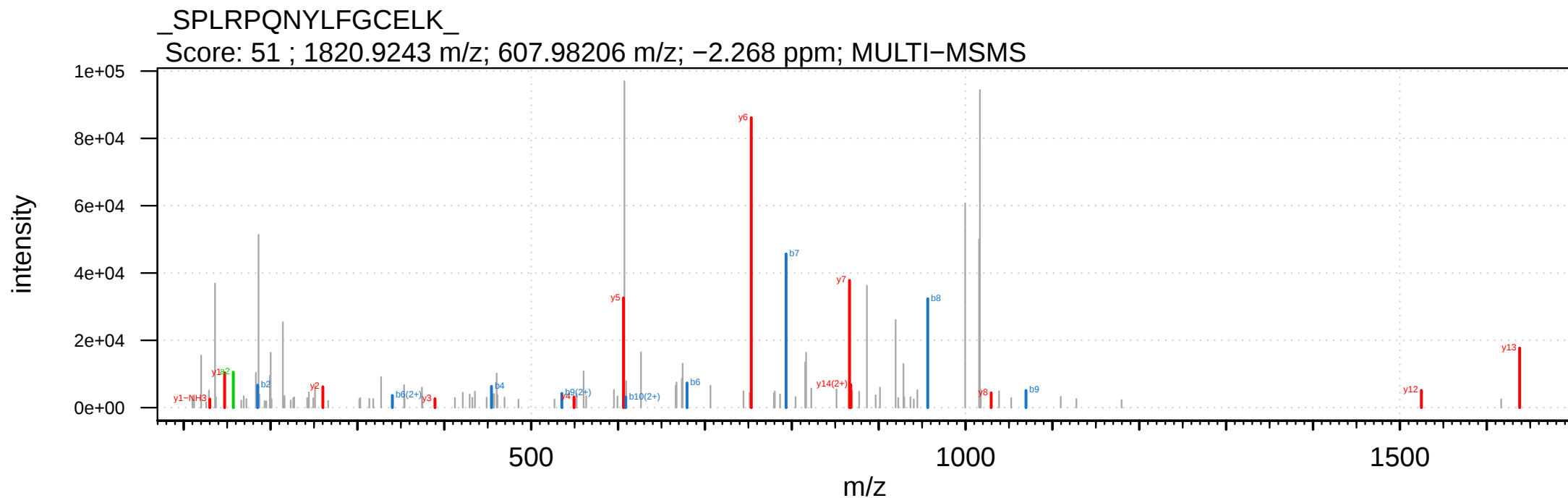
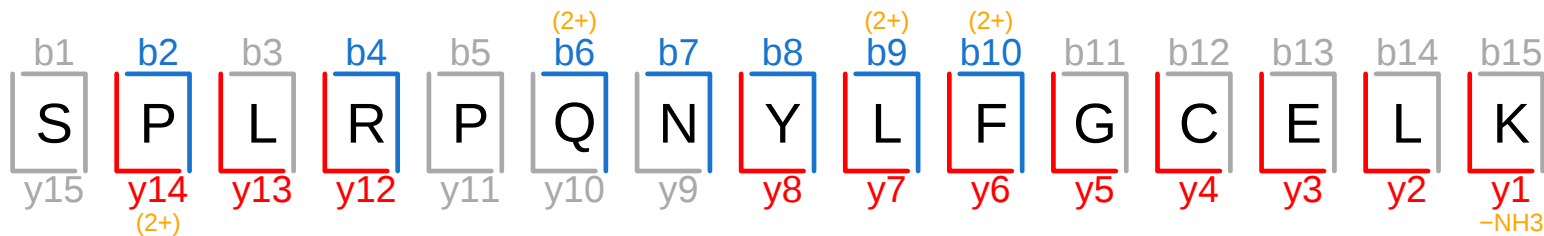


Raw File: 20100614_Velos1_TaGe_SA_K562_1

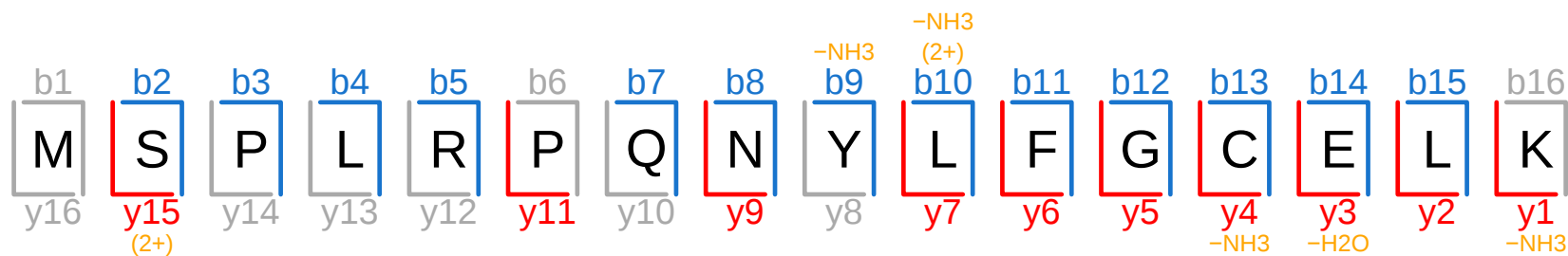
Scan Number: 22646

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

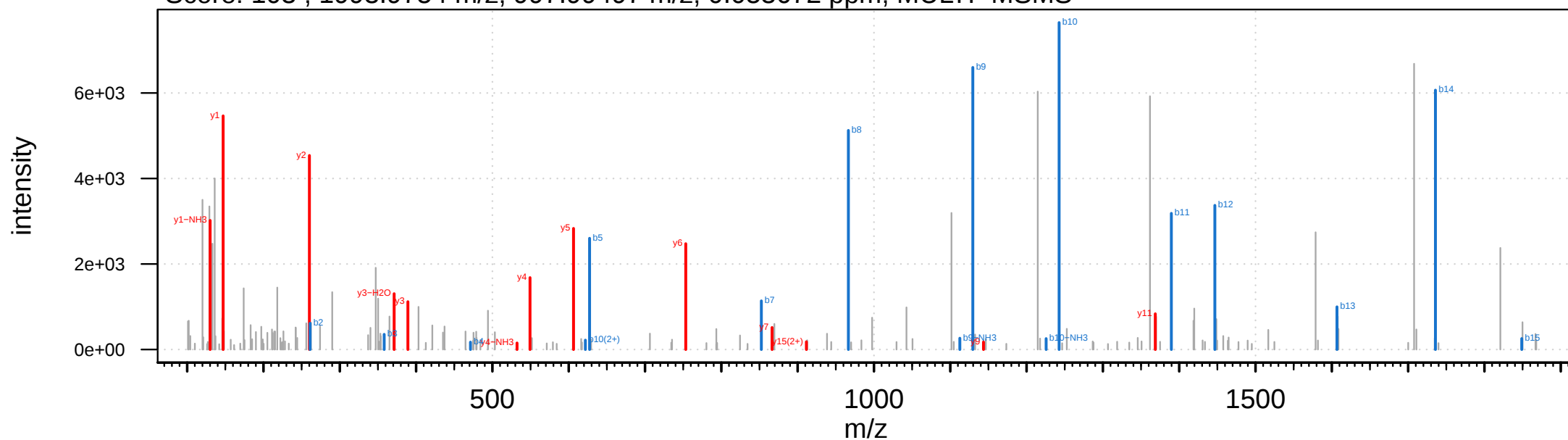


Raw File: 20100614_Velos1_TaGe_SA_K562_1
 Scan Number: 22656
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 103 ; 1993.9754 m/z; 997.99497 m/z; 0.053672 ppm; MULTI-MSMS

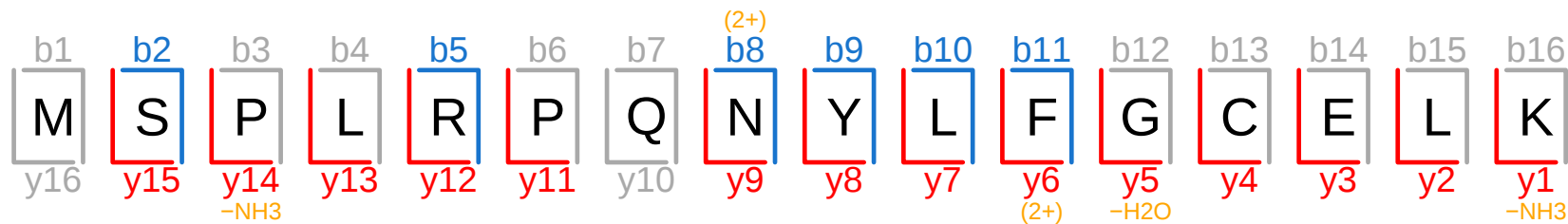


Raw File: 20101210_Velos1_AnWe_SA_LnCap_2

Scan Number: 32137

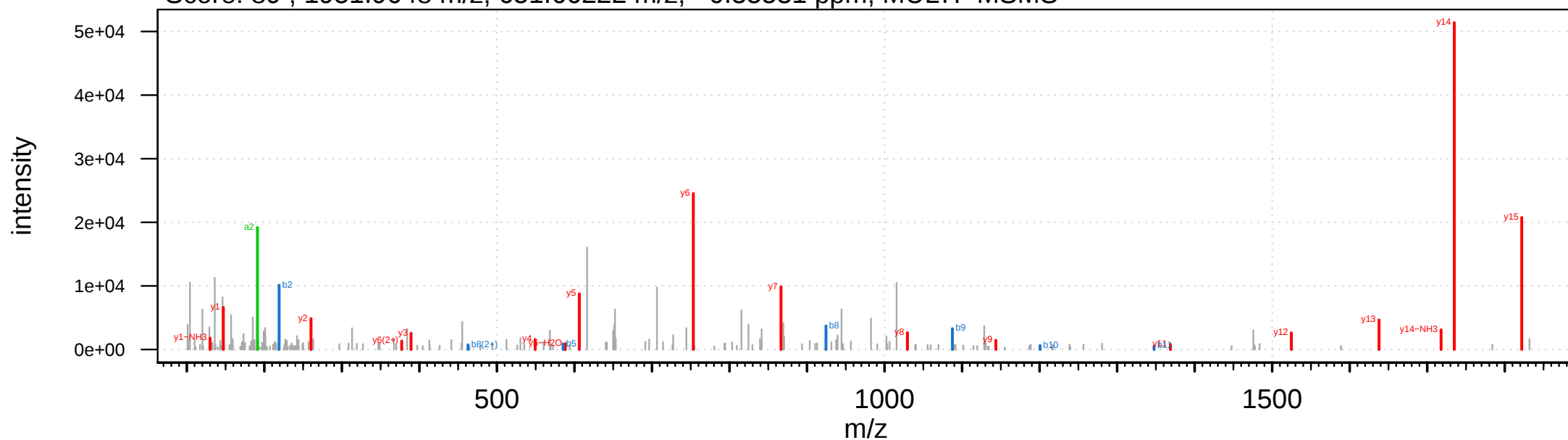
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



MSPLRPQNYLFGCELK

Score: 89 ; 1951.9648 m/z; 651.66222 m/z; -0.35531 ppm; MULTI-MSMS

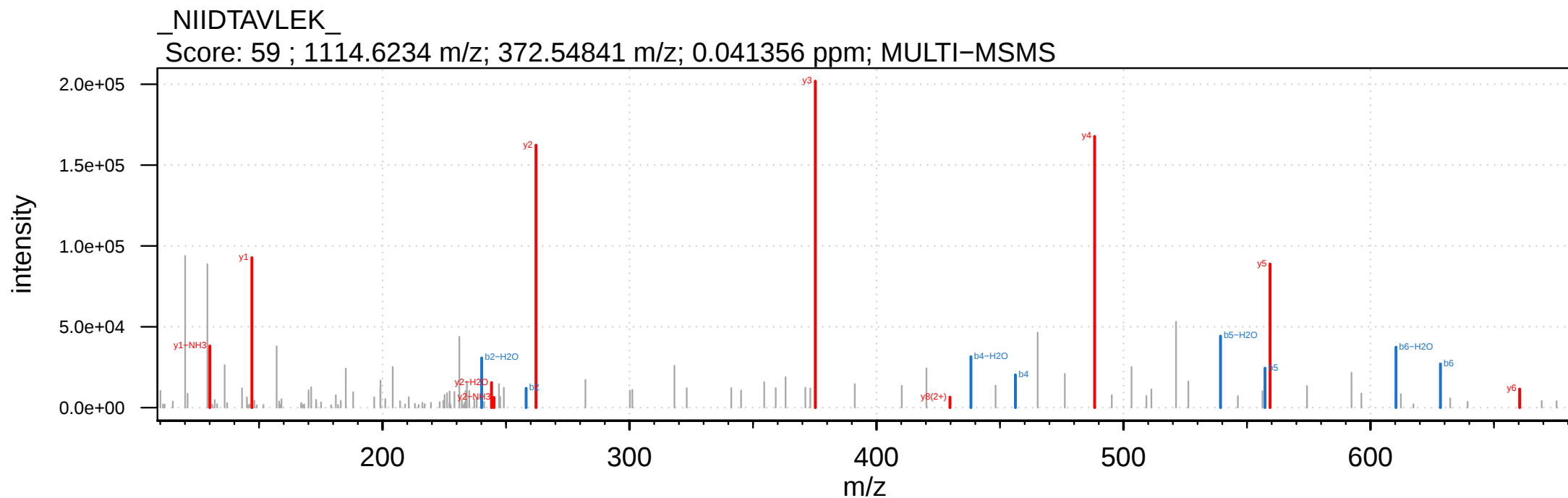
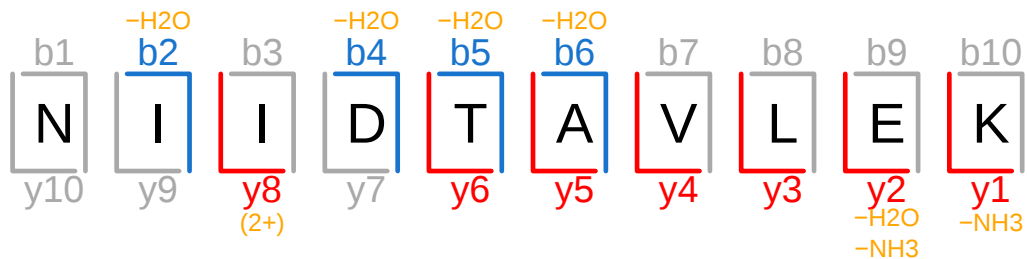


Raw File: 20101210_Velos1_AnWe_SA_LnCap_2

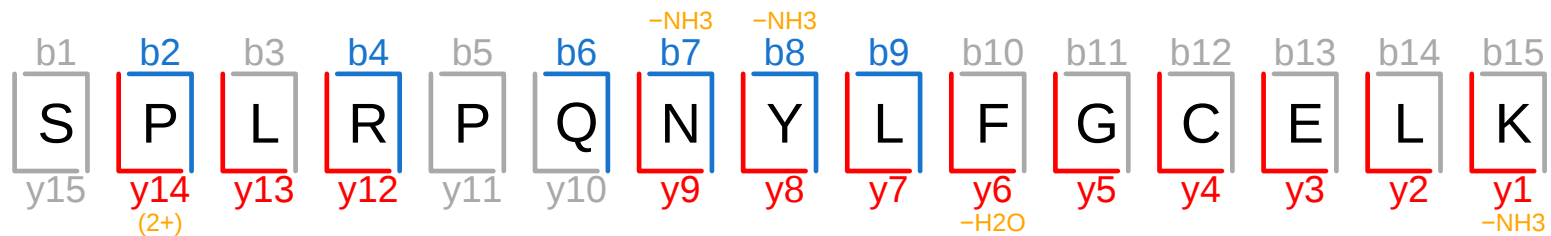
Scan Number: 23919

Proteins:

TCONS_I2_00008829_ch15:92829088-92829258:+

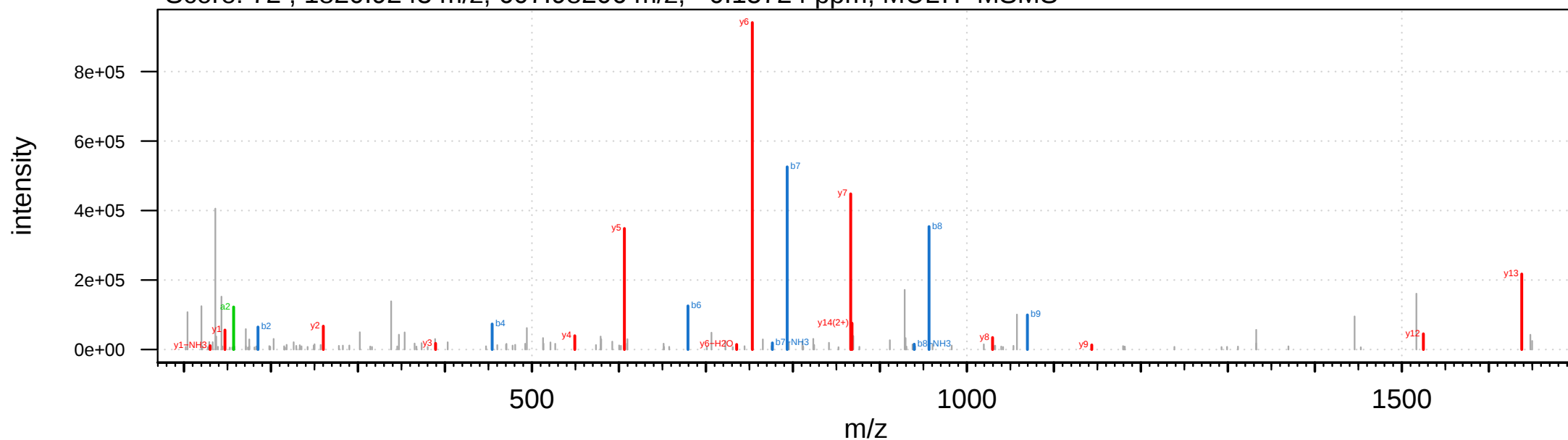


Raw File: 20101210_Velos1_AnWe_SA_LnCap_2
Scan Number: 9707
Proteins:
ENST00000497138_chr20:56806826-56807846:-

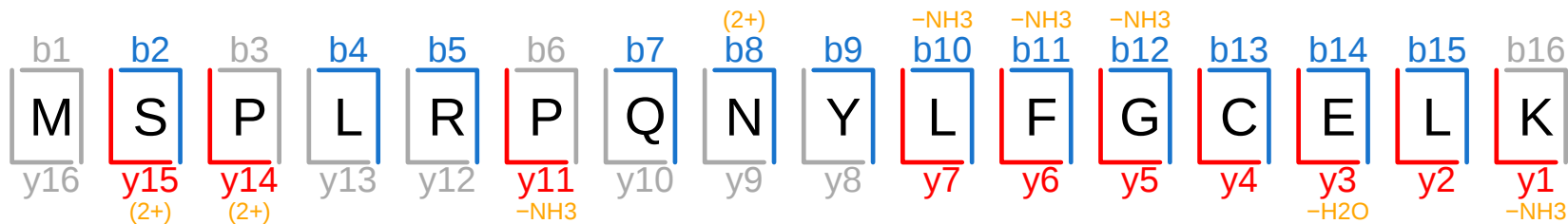


SPLRPQNYLFGCELK

Score: 72 ; 1820.9243 m/z; 607.98206 m/z; -0.15724 ppm; MULTI-MSMS

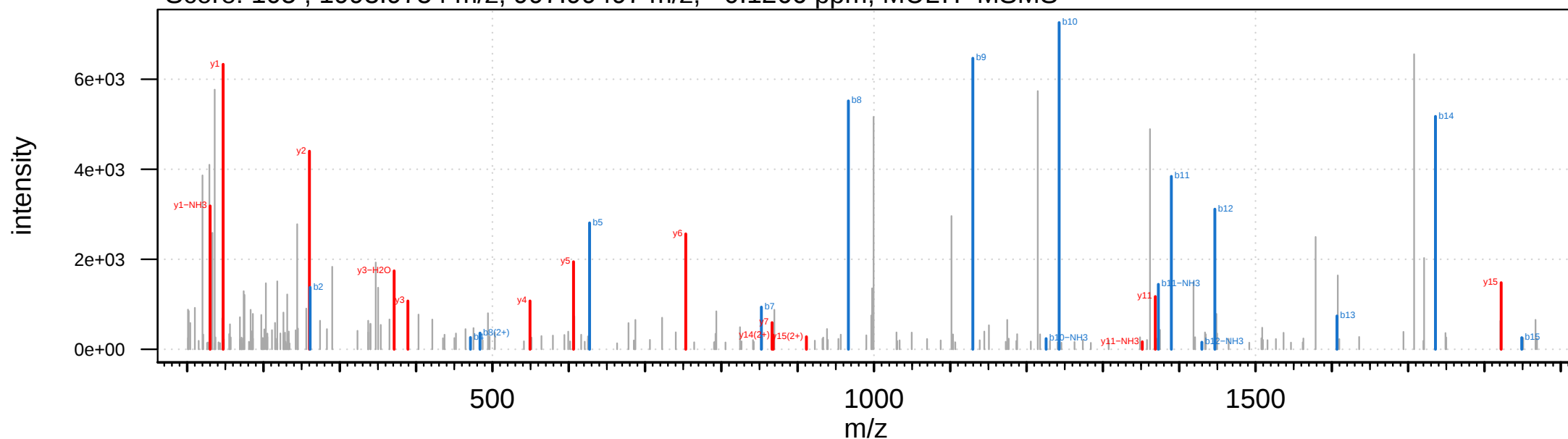


Raw File: 20101210_Velos1_AnWe_SA_LnCap_2
 Scan Number: 21025
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 105 ; 1993.9754 m/z; 997.99497 m/z; -0.1266 ppm; MULTI-MSMS

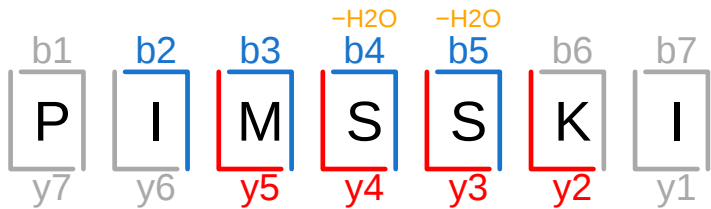


Raw File: 20101210_Velos1_AnWe_SA_U2OS_1

Scan Number: 38163

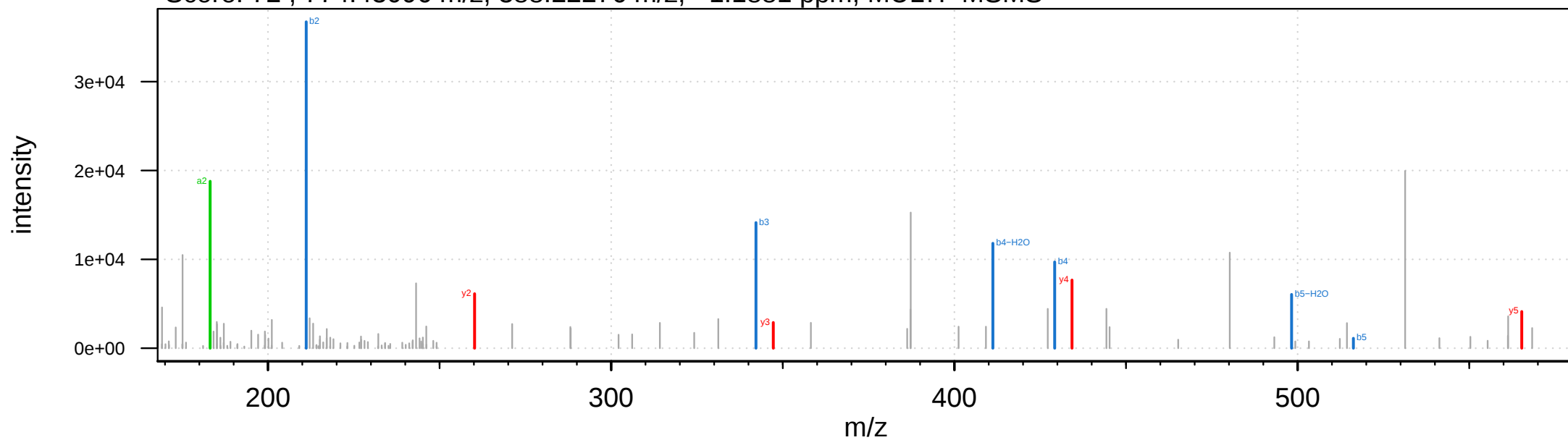
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

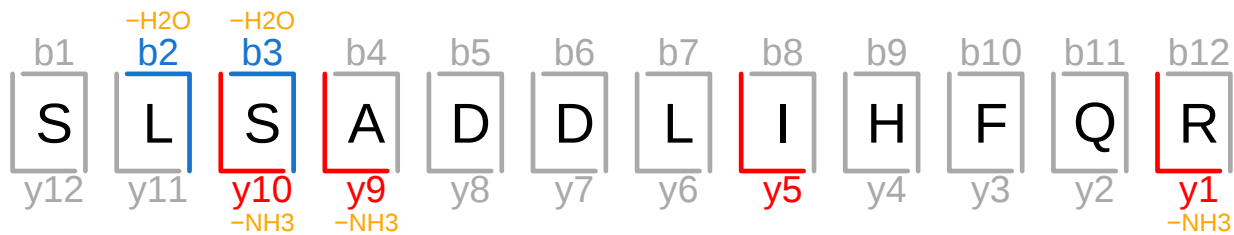


PIMSSKI

Score: 72 ; 774.43096 m/z; 388.22276 m/z; -1.1881 ppm; MULTI-MSMS

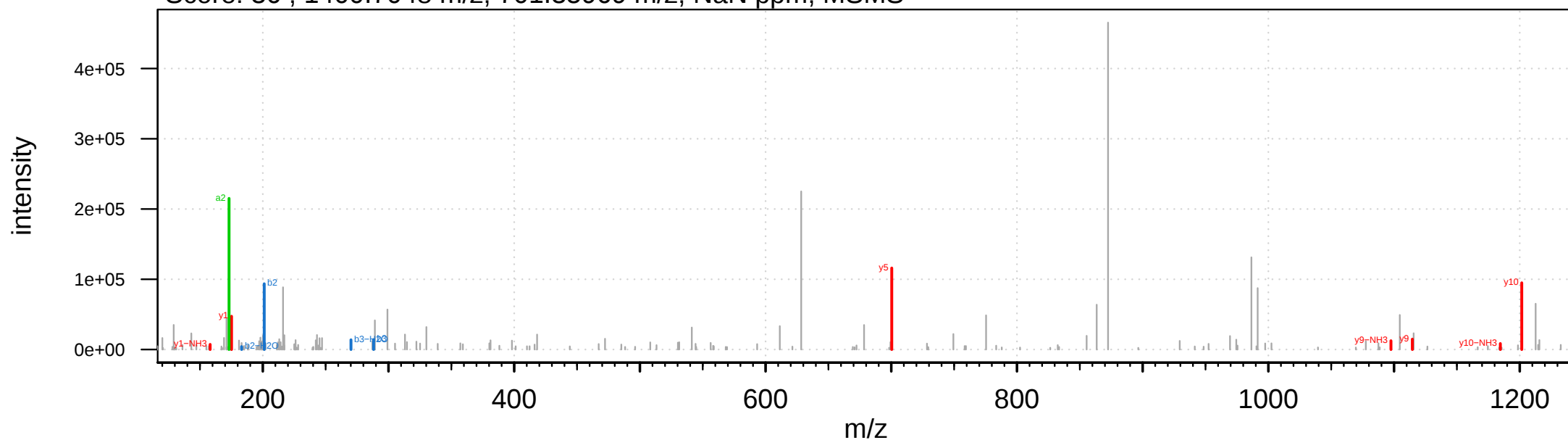


Raw File: 20101210_Velos1_AnWe_SA_U2OS_1
 Scan Number: 22625
 Proteins:
 ENST00000490272_chr19:52826295-52826555:+



SLSADDLIHFQR

Score: 36 ; 1400.7048 m/z; 701.35969 m/z; NaN ppm; MSMS

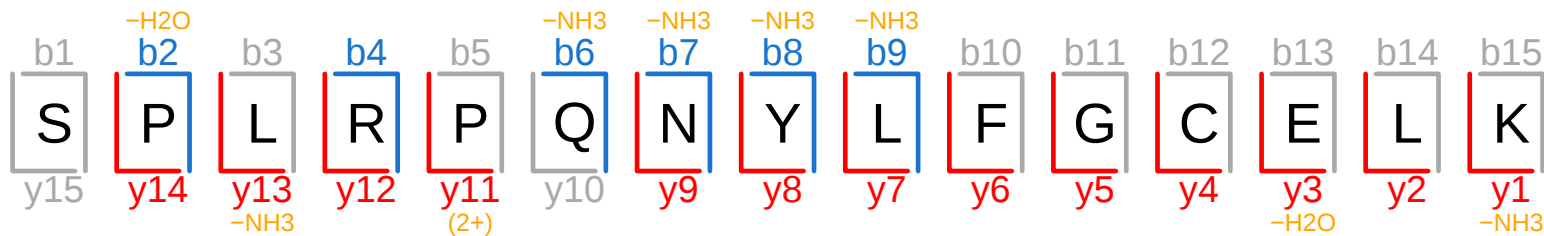


Raw File: 20101210_Velos1_AnWe_SA_U2OS_1

Scan Number: 19671

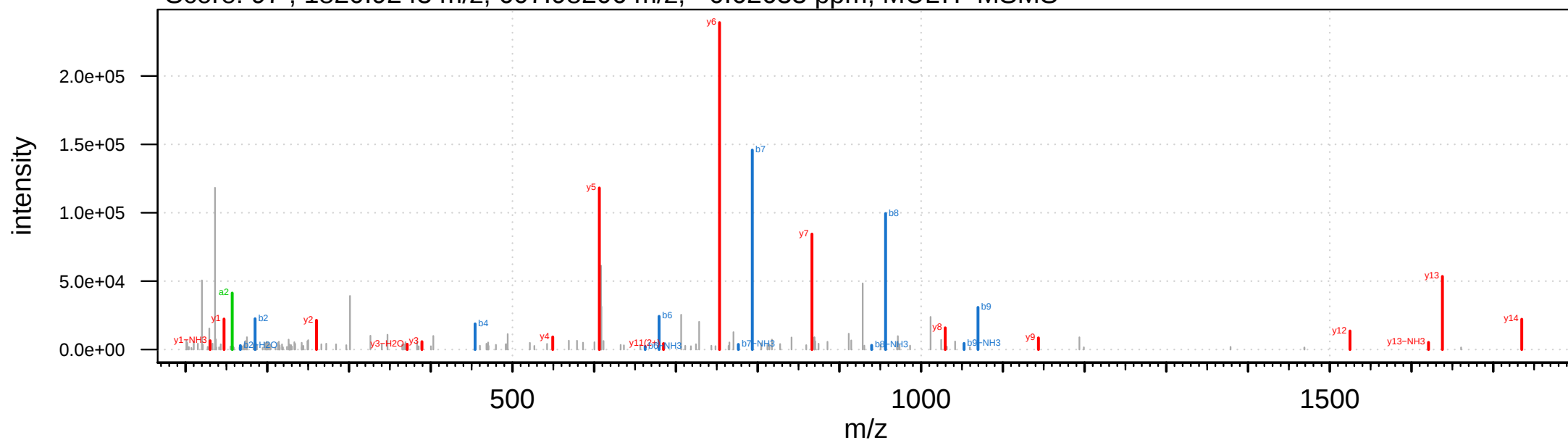
Proteins:

TCONS_I2_00002988_chr10:38499673-38501345:+

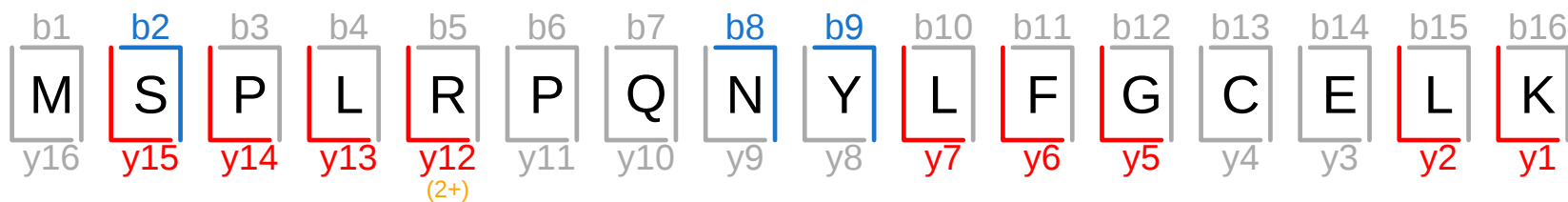


SPLRPQNYLFGCELK

Score: 97 ; 1820.9243 m/z; 607.98206 m/z; -0.62633 ppm; MULTI-MSMS

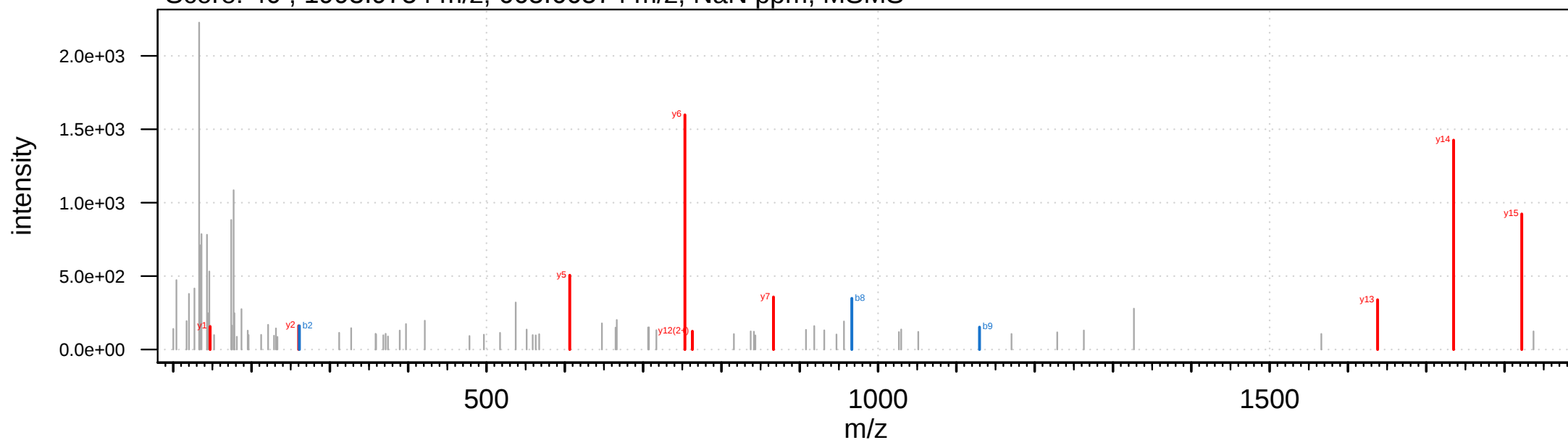


Raw File: 20101210_Velos1_AnWe_SA_U2OS_1
 Scan Number: 23828
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 40 ; 1993.9754 m/z; 665.66574 m/z; NaN ppm; MSMS

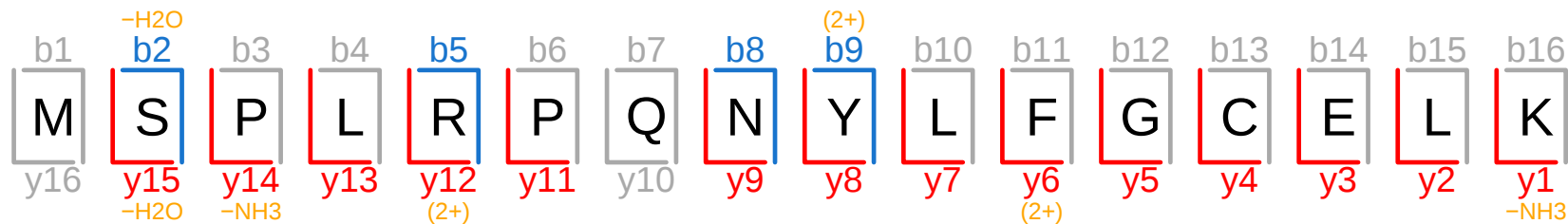


Raw File: 20101222_Velos1-TaGe_SA_K562_01

Scan Number: 22570

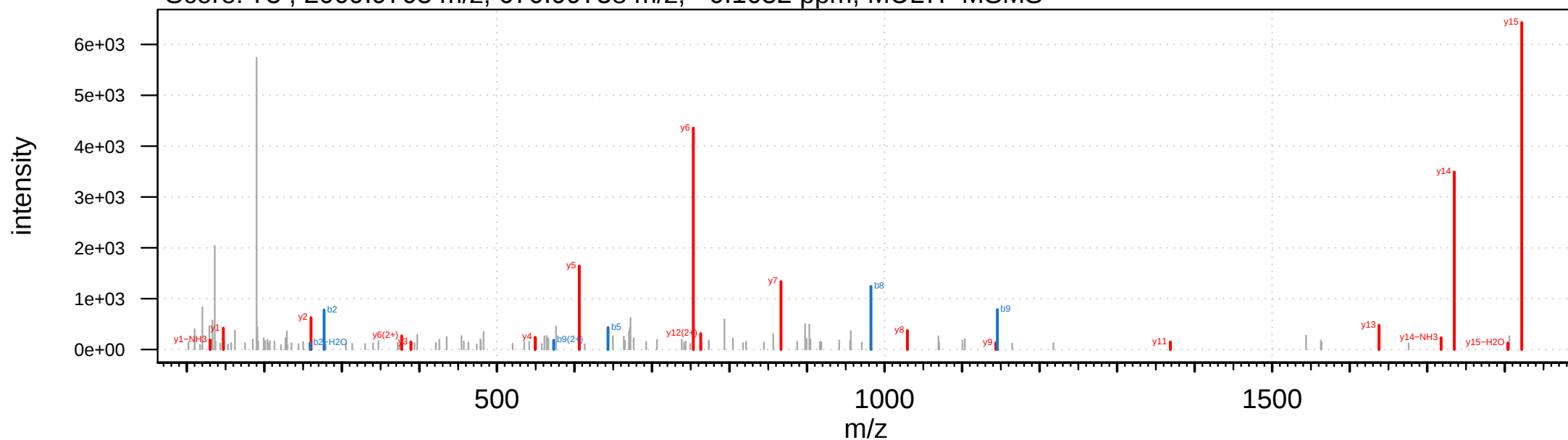
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)M(ox)SPLRPQNYLFGCELK

Score: 73 ; 2009.9703 m/z; 670.99738 m/z; -0.1032 ppm; MULTI-MSMS

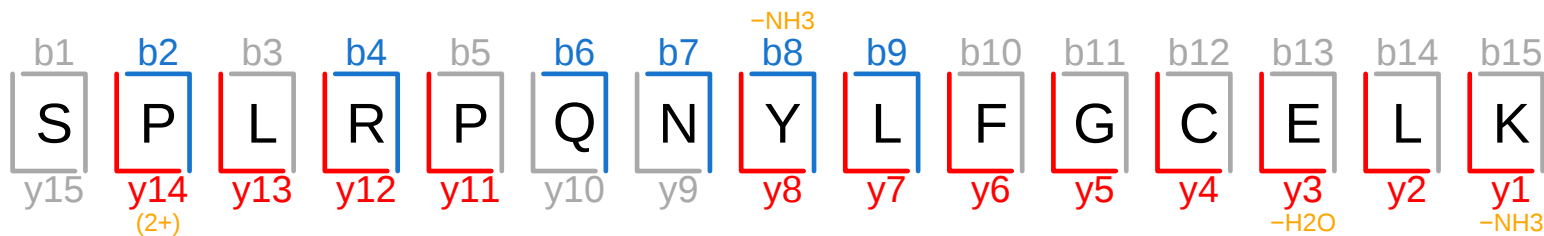


Raw File: 20101222_Velos1_TaGe_SA_K562_01

Scan Number: 19383

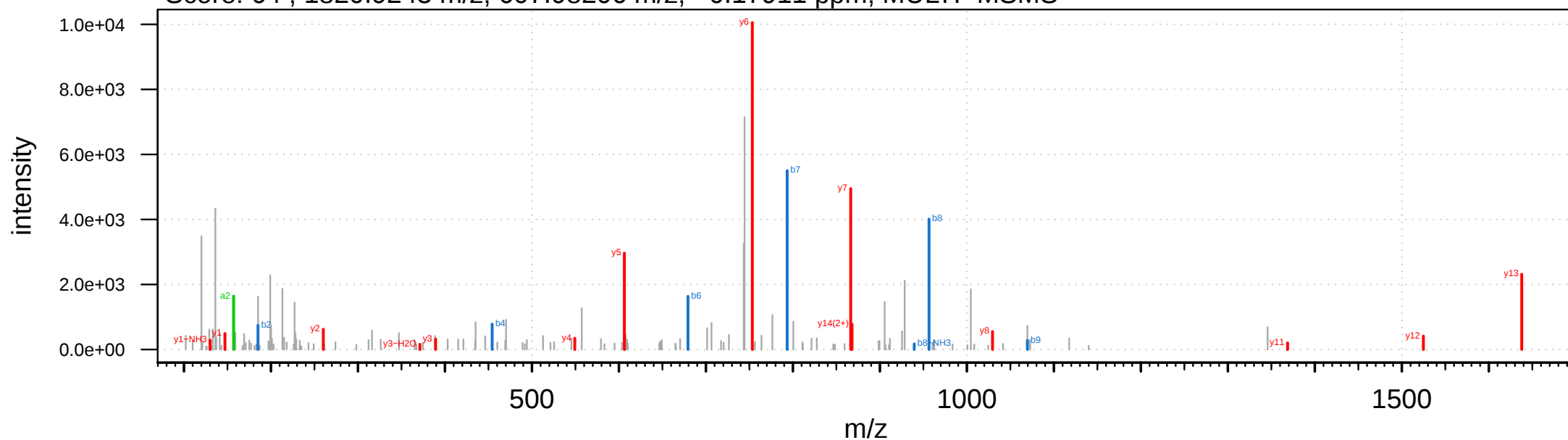
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 64 ; 1820.9243 m/z; 607.98206 m/z; -0.17911 ppm; MULTI-MSMS

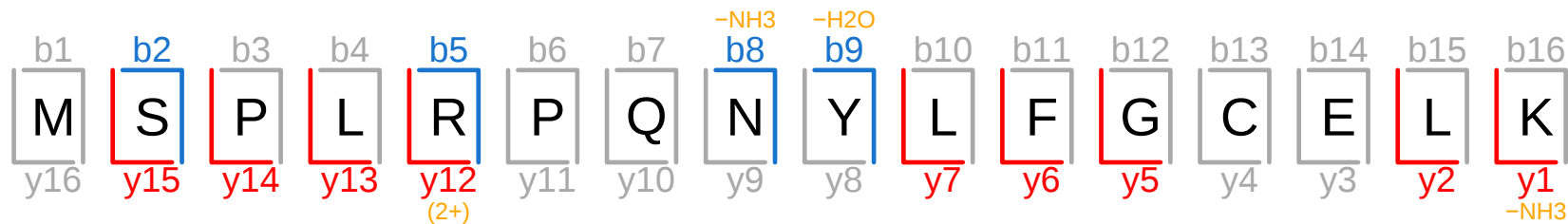


Raw File: 20101222_Velos1_TaGe_SA_K562_01

Scan Number: 14066

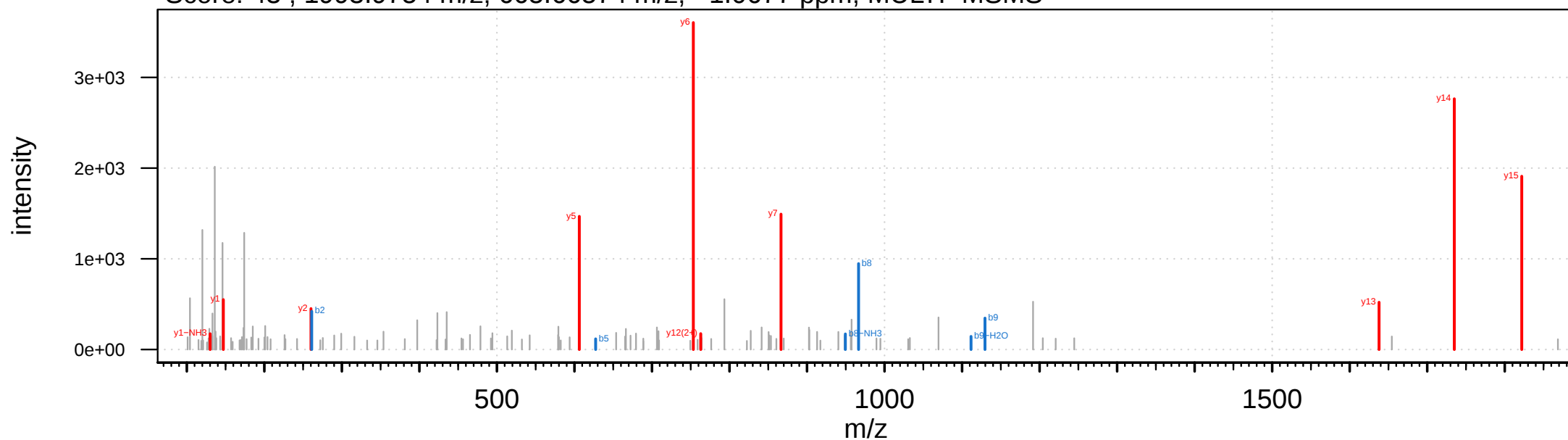
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 43 ; 1993.9754 m/z; 665.66574 m/z; -1.0677 ppm; MULTI-MSMS

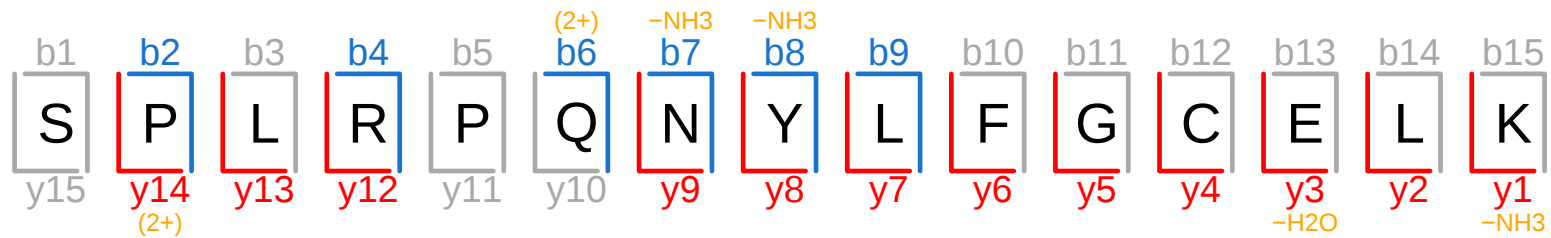


Raw File: 20101222_Velos1_TaGe_SA_K562_02

Scan Number: 20965

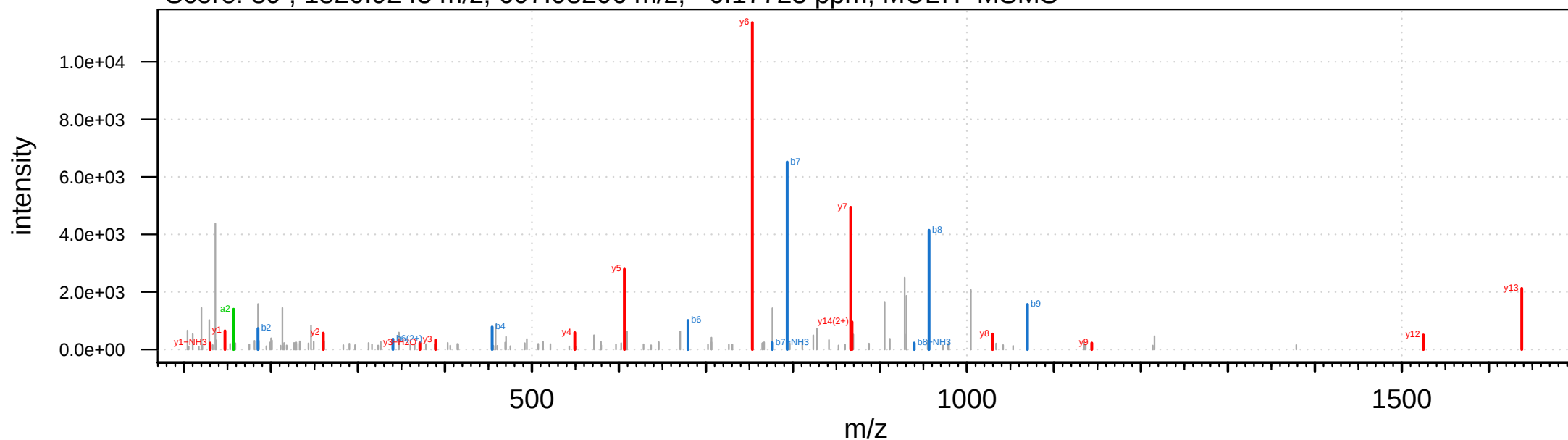
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 89 ; 1820.9243 m/z; 607.98206 m/z; -0.17723 ppm; MULTI-MSMS

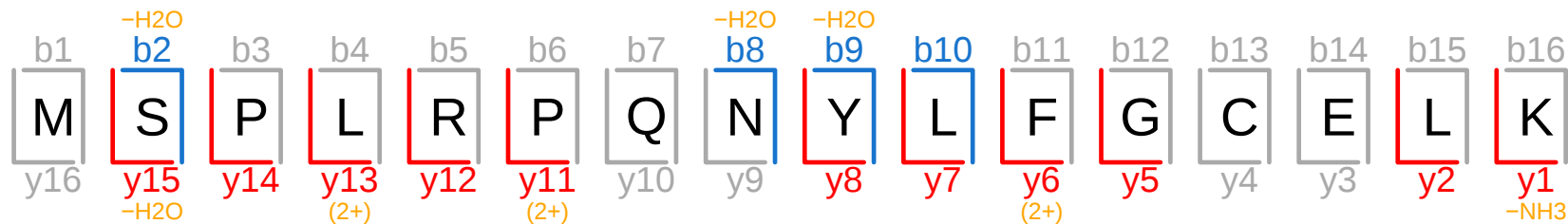


Raw File: 20101222_Velos1_TaGe_SA_K562_02

Scan Number: 12418

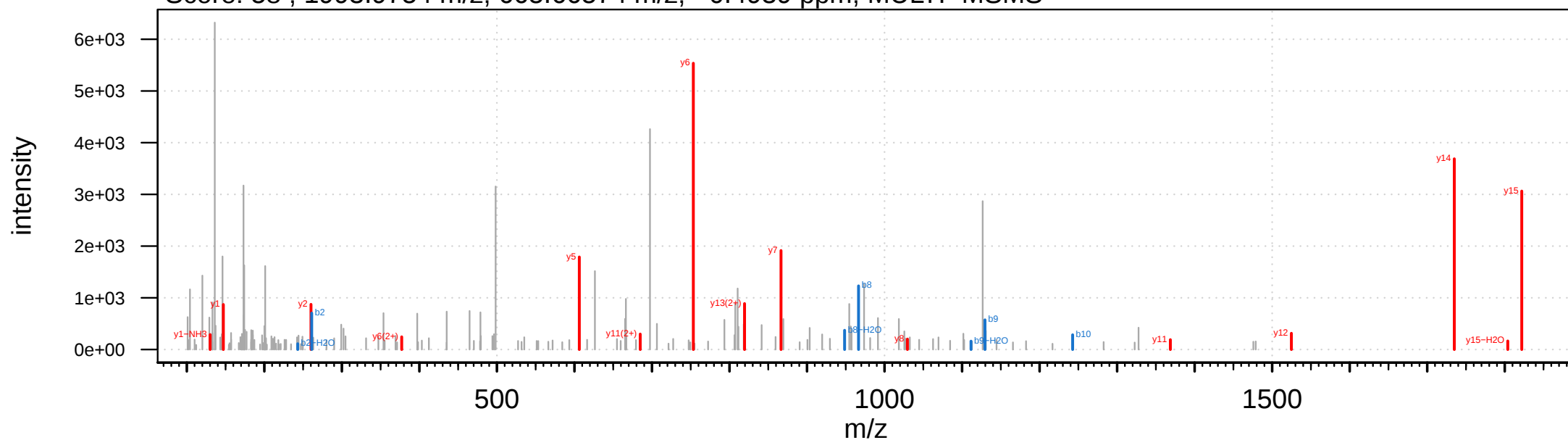
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 58 ; 1993.9754 m/z; 665.66574 m/z; -0.4939 ppm; MULTI-MSMS

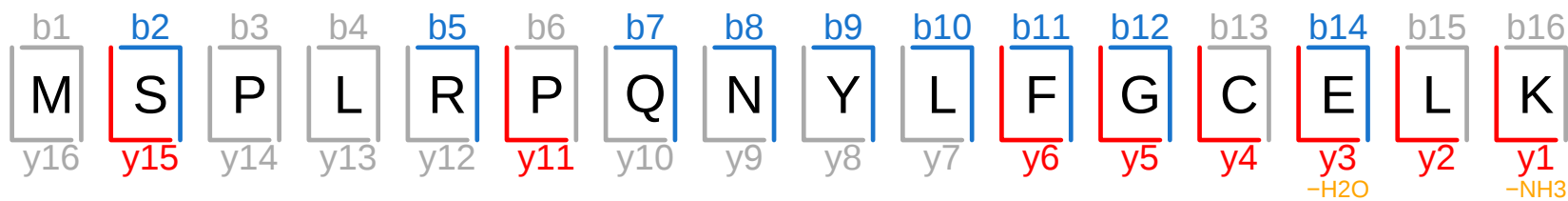


Raw File: 20101224_Velos1-TaGe_SA-HepG2_2

Scan Number: 33012

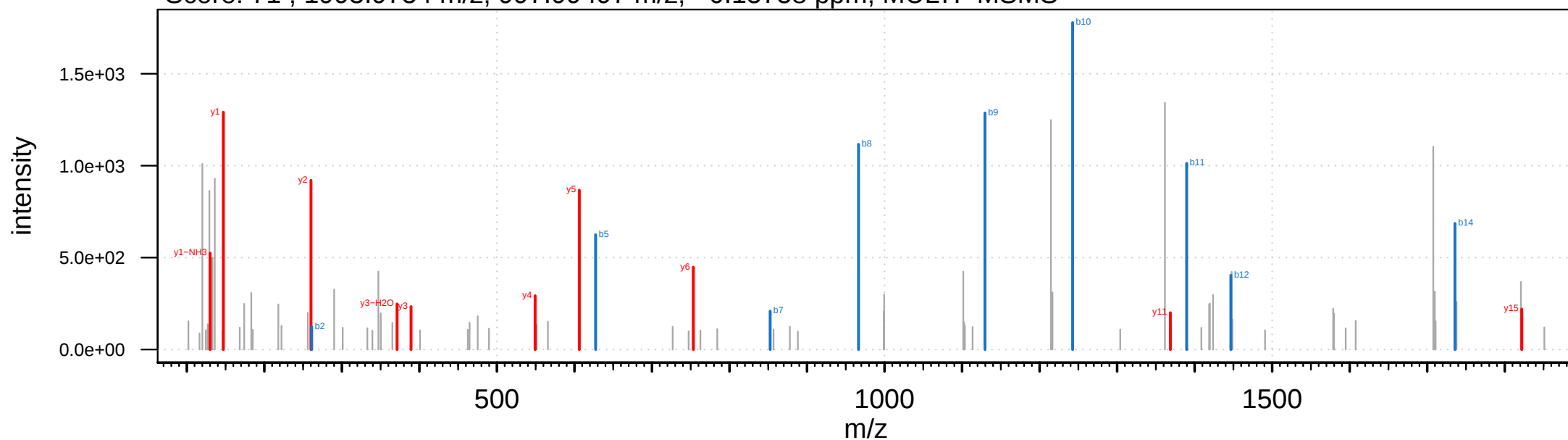
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 71 ; 1993.9754 m/z; 997.99497 m/z; -0.15758 ppm; MULTI-MSMS

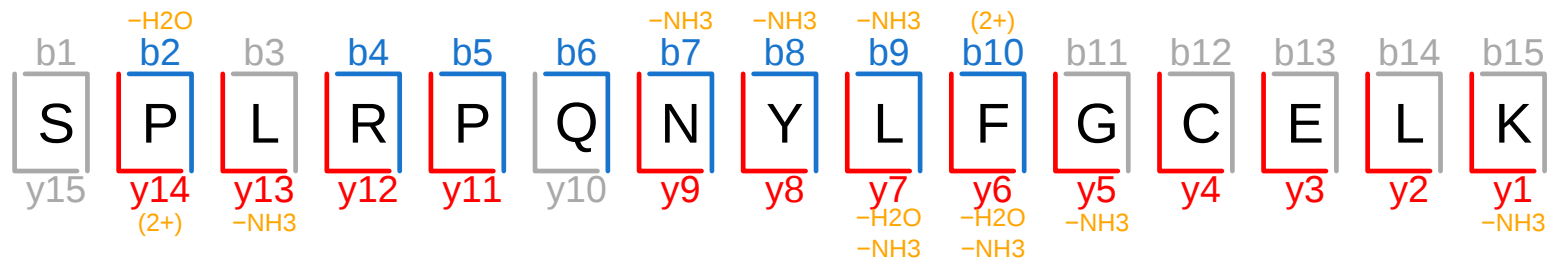


Raw File: 20101224_Velos1-TaGe_SA-HepG2_2

Scan Number: 33013

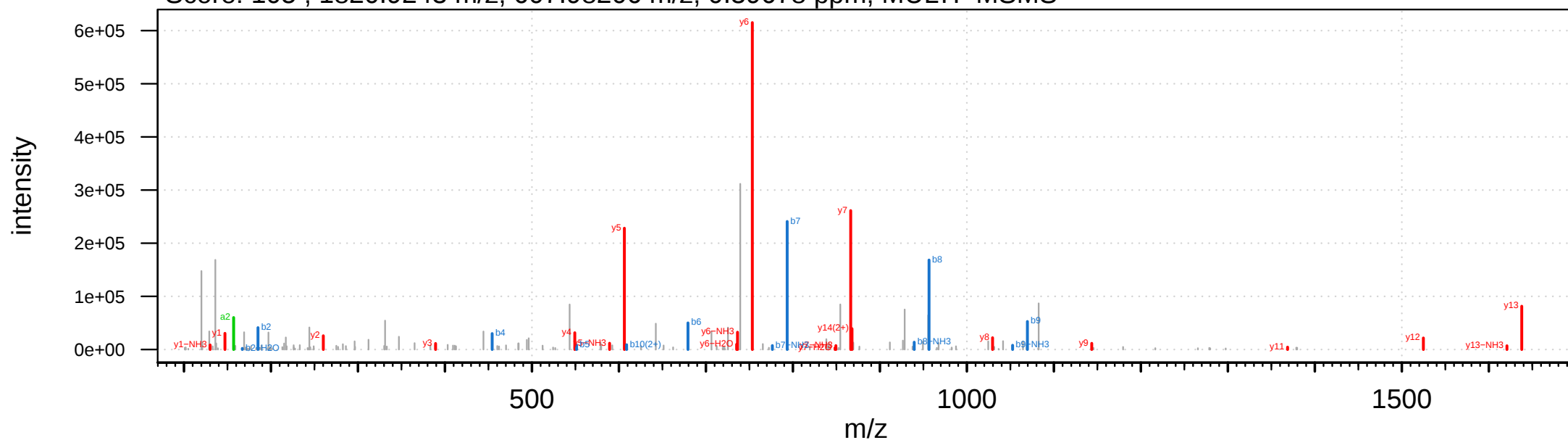
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 105 ; 1820.9243 m/z; 607.98206 m/z; 0.39678 ppm; MULTI-MSMS

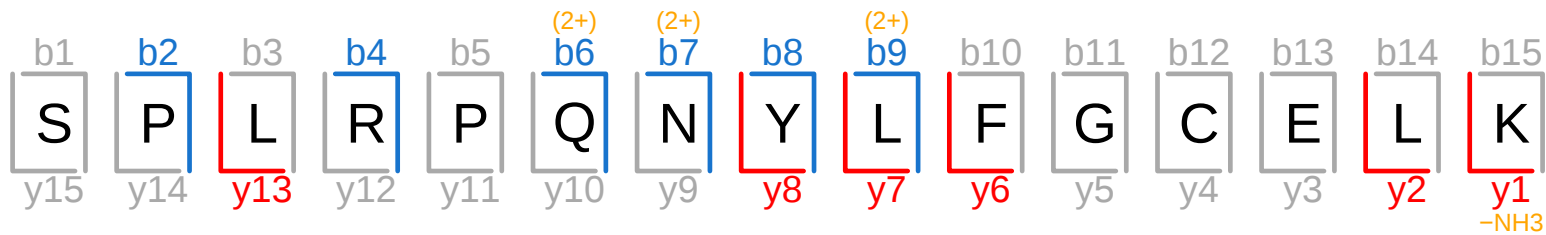


Raw File: 20101224_Velos1_TaGe_SA_HepG2_2

Scan Number: 22244

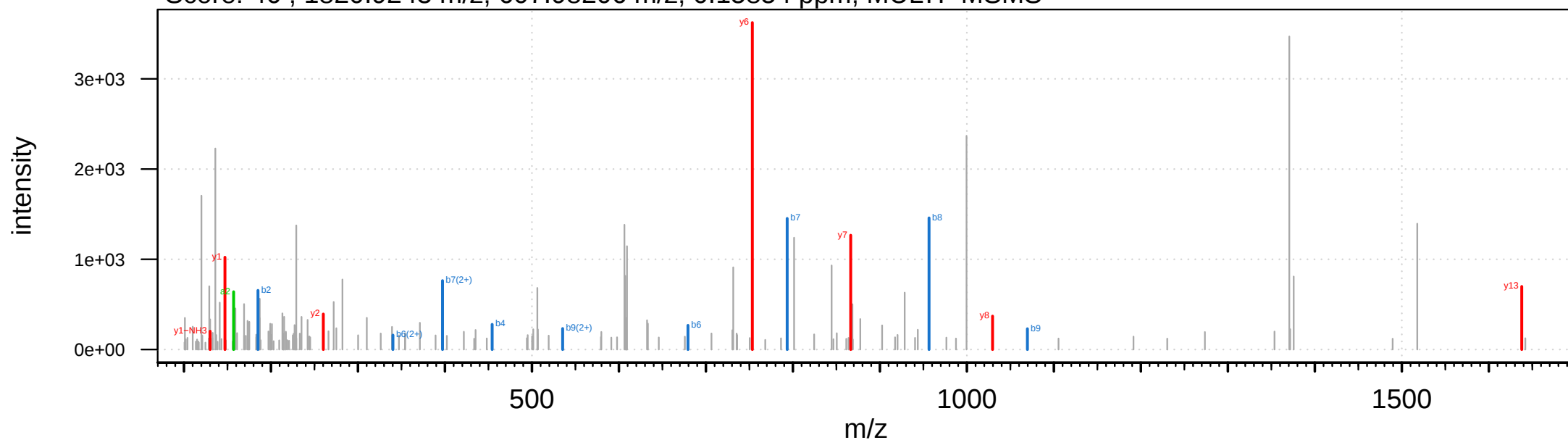
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 46 ; 1820.9243 m/z; 607.98206 m/z; 0.15854 ppm; MULTI-MSMS

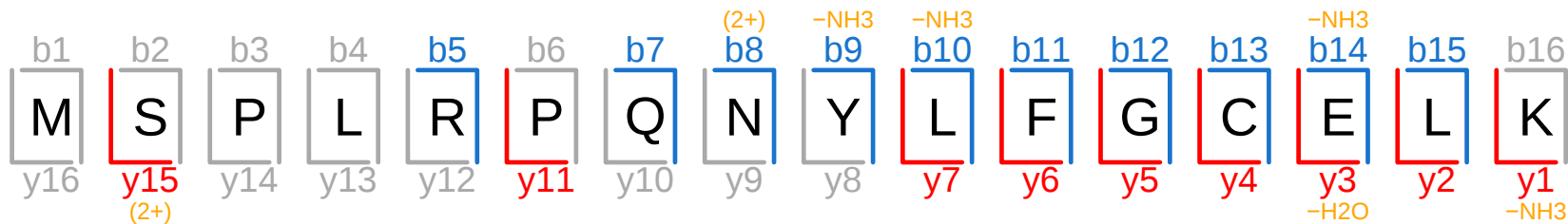


Raw File: 20101224_Velos1-TaGe_SA-HepG2_2

Scan Number: 33935

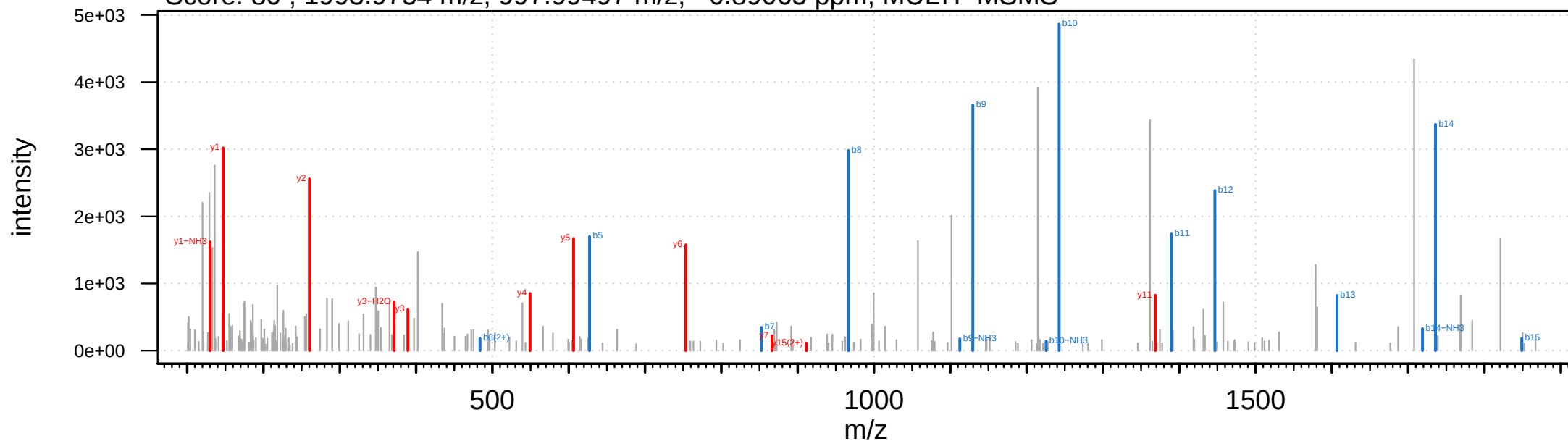
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 80 ; 1993.9754 m/z; 997.99497 m/z; -0.89065 ppm; MULTI-MSMS

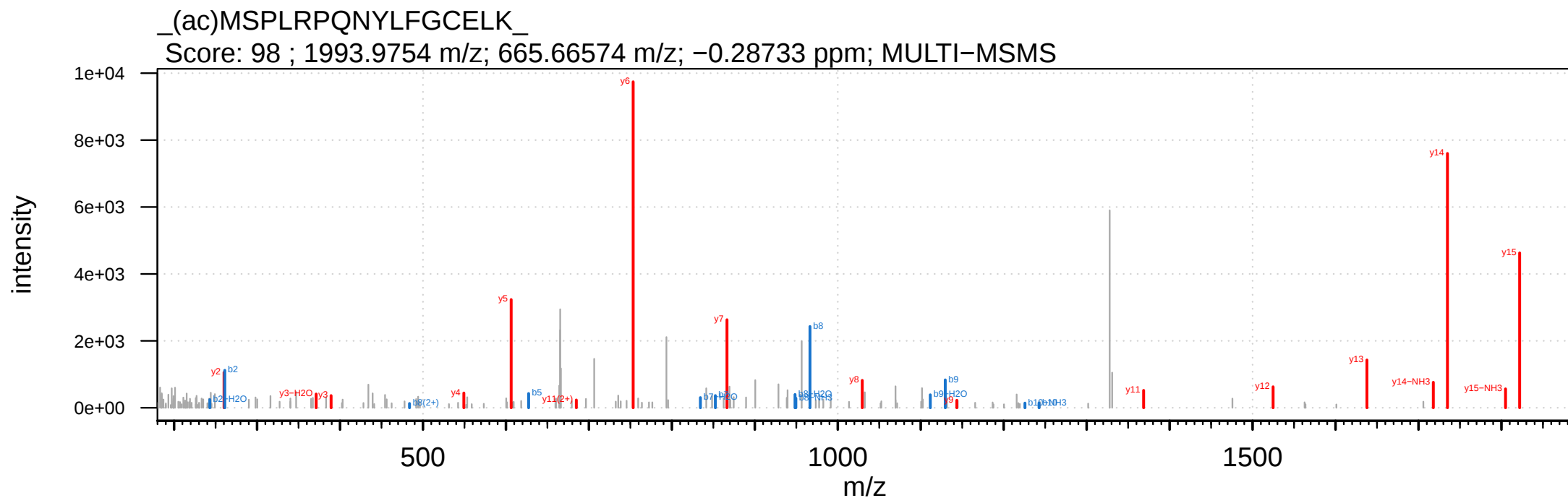
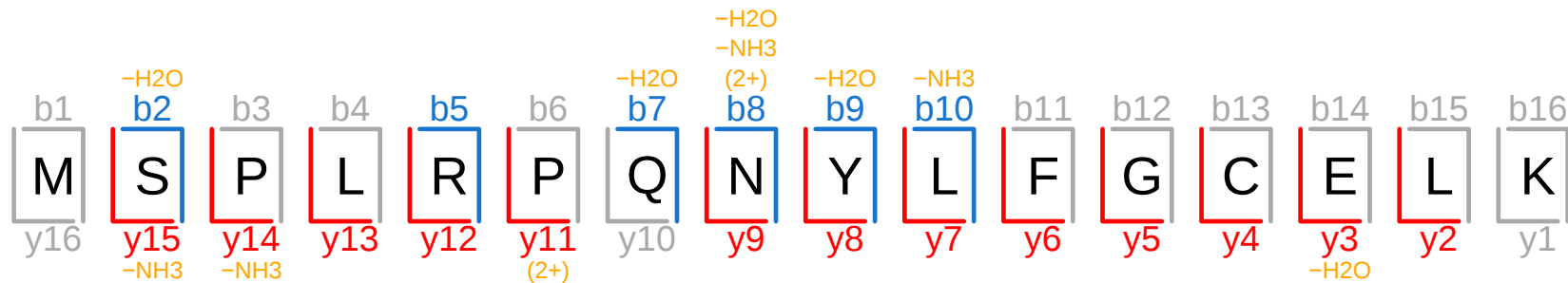


Raw File: 20101227_Velos1_TaGe_SA_GAMG_101230100451

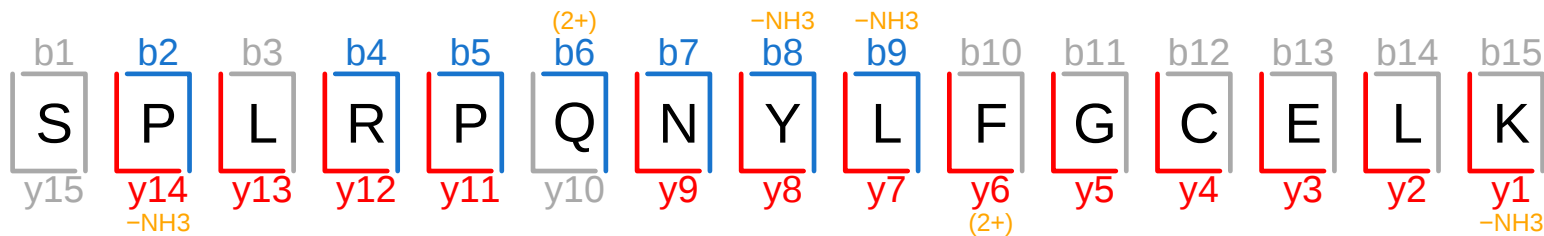
Scan Number: 38847

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

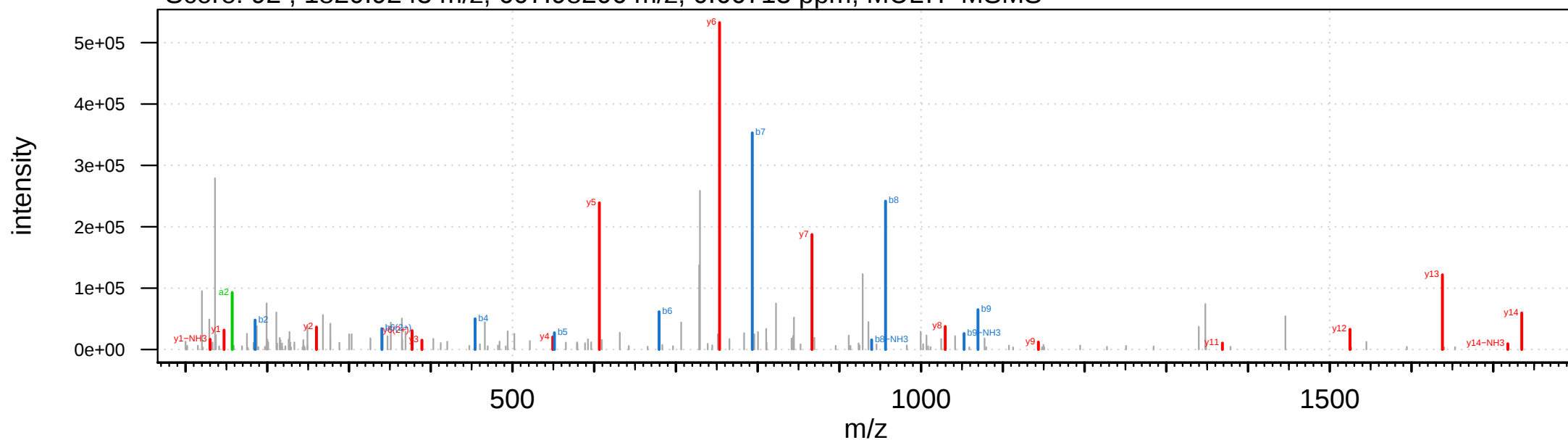


Raw File: 20101227_Velos1_TaGe_SA_GAMG_101230100451
 Scan Number: 38898
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 92 ; 1820.9243 m/z; 607.98206 m/z; 0.66713 ppm; MULTI-MSMS

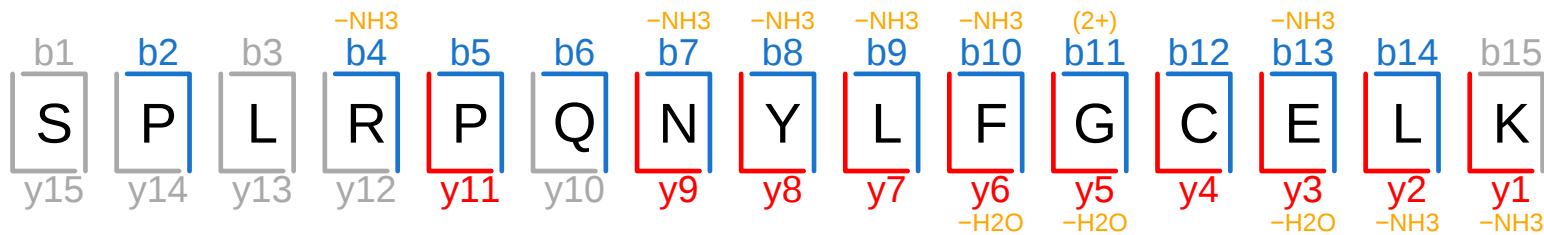


Raw File: 20101227_Velos1_TaGe_SA_GAMG_101230100451

Scan Number: 25043

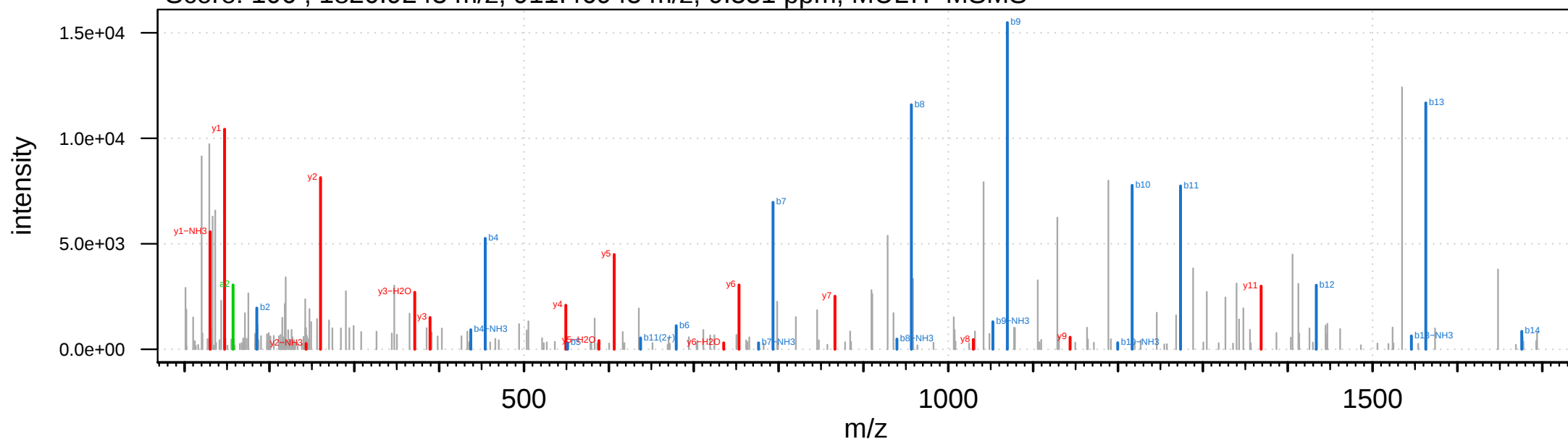
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 106 ; 1820.9243 m/z; 911.46945 m/z; 0.351 ppm; MULTI-MSMS

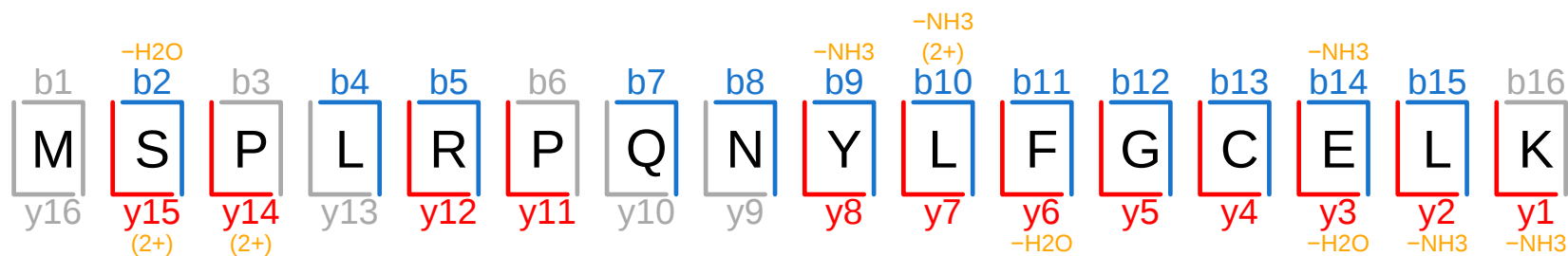


Raw File: 20101227_Velos1_TaGe_SA_GAMG_101230100451

Scan Number: 25047

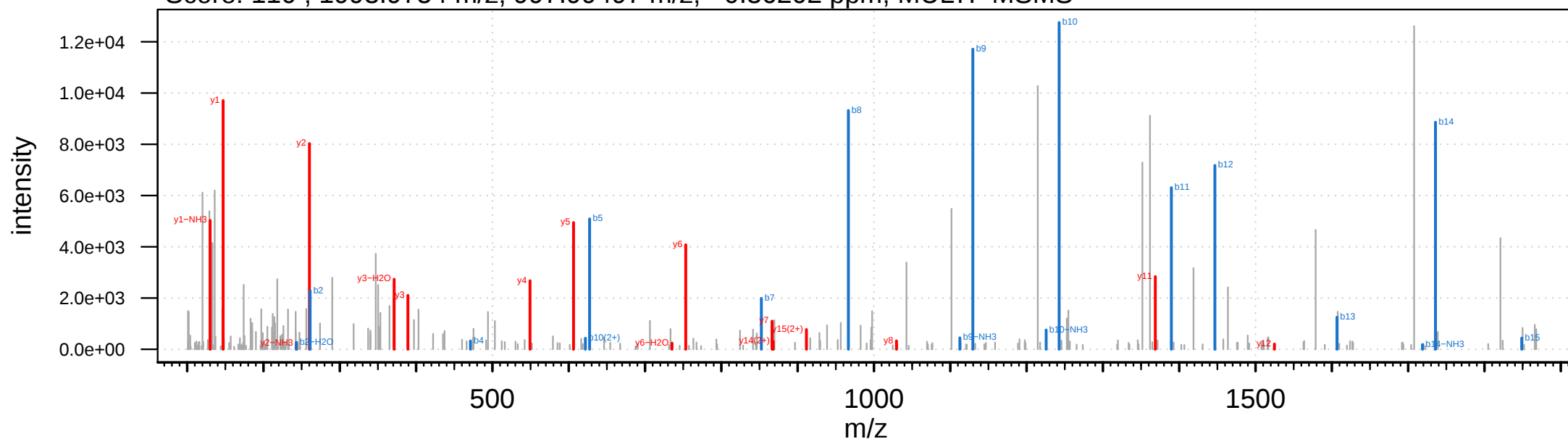
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 110 ; 1993.9754 m/z; 997.99497 m/z; -0.36202 ppm; MULTI-MSMS

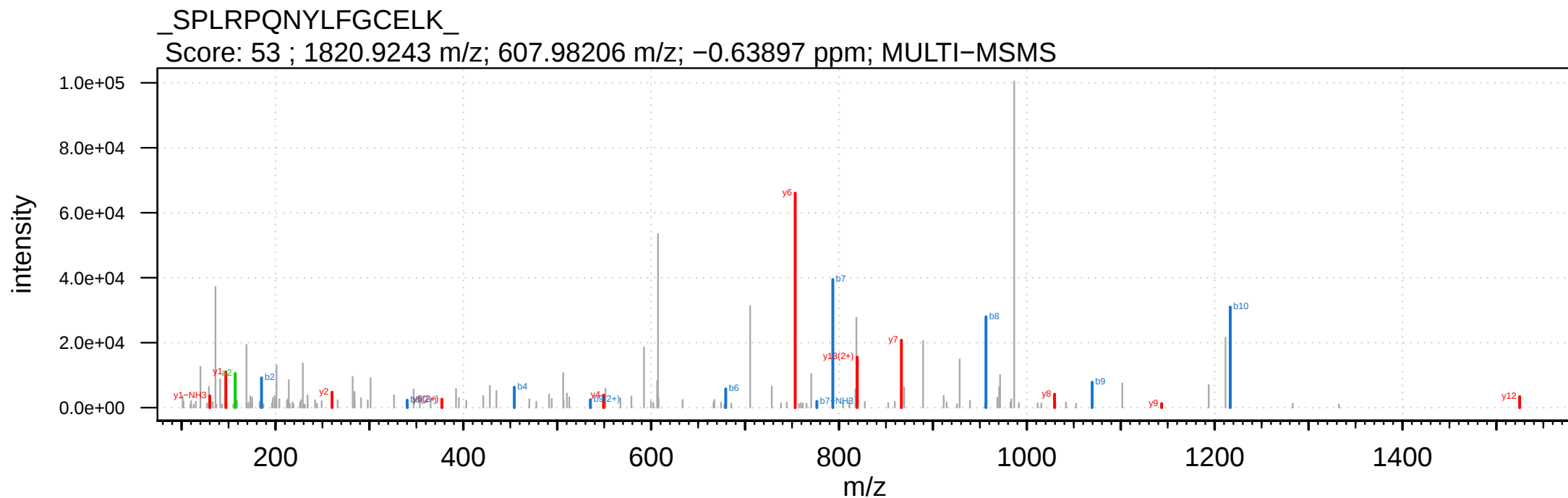
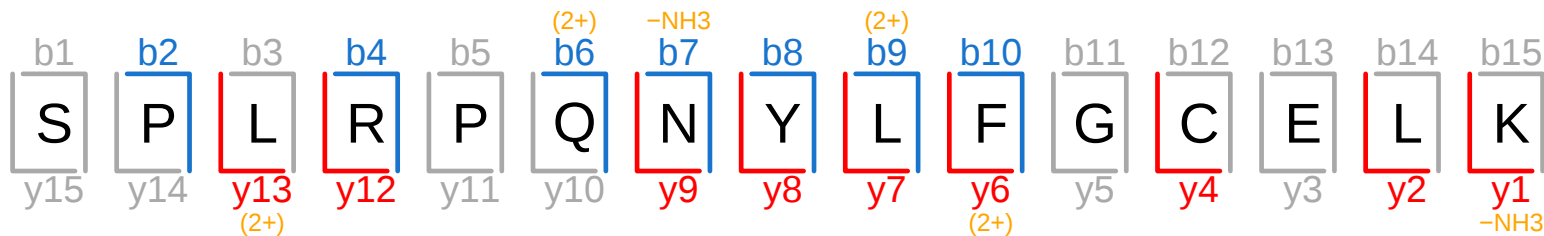


Raw File: 20101227_Velos1_TaGe_SA_HEK293_01

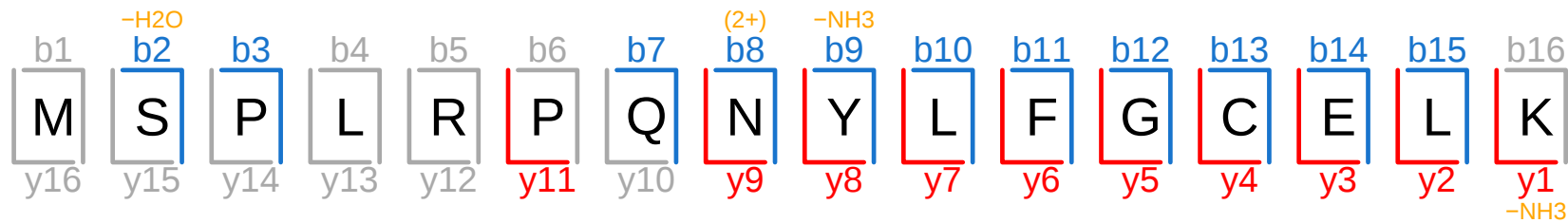
Scan Number: 36838

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

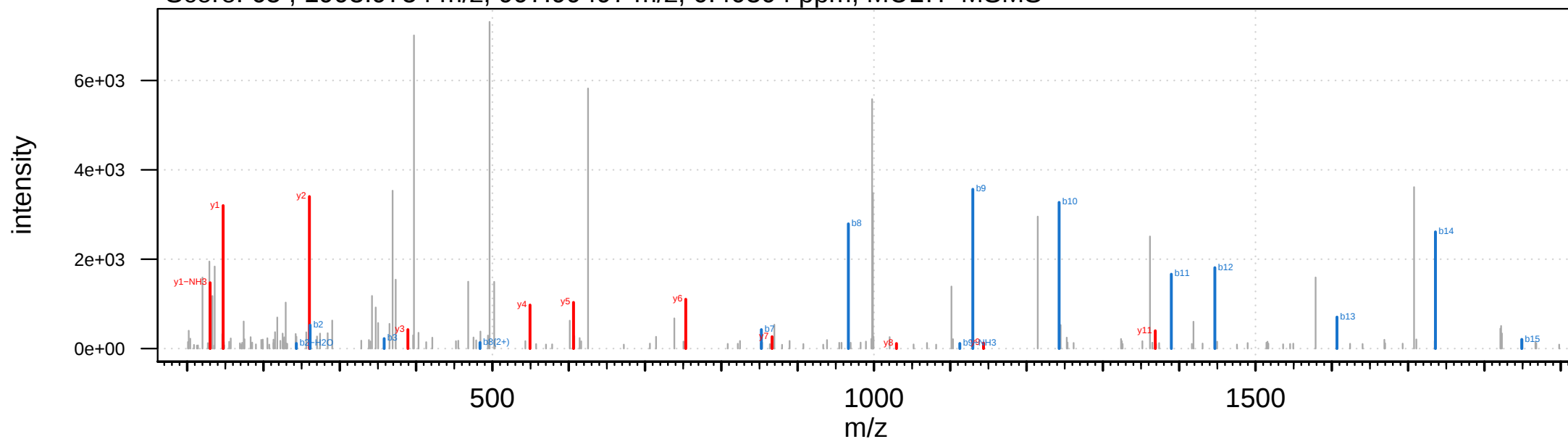


Raw File: 20101227_Velos1_TaGe_SA_HEK293_01
 Scan Number: 21139
 Proteins:
 TCONS_I2_00008829_ch15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 65 ; 1993.9754 m/z; 997.99497 m/z; 0.40304 ppm; MULTI-MSMS

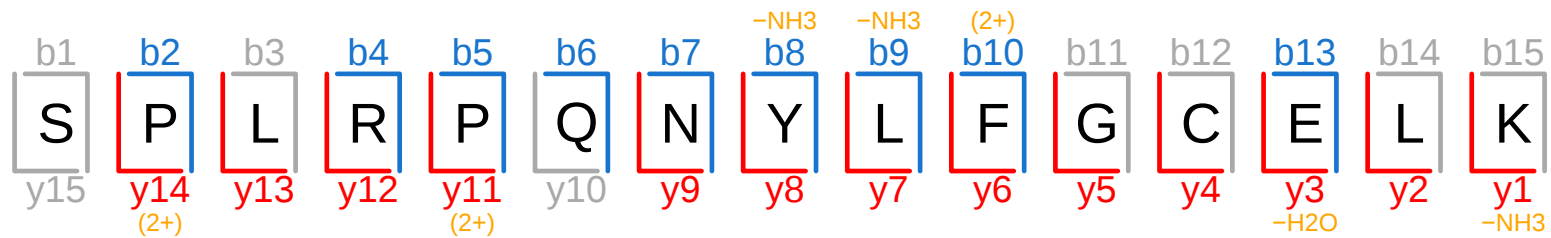


Raw File: 20101230_Velos1_TaGe_SA_Jurkat1

Scan Number: 39660

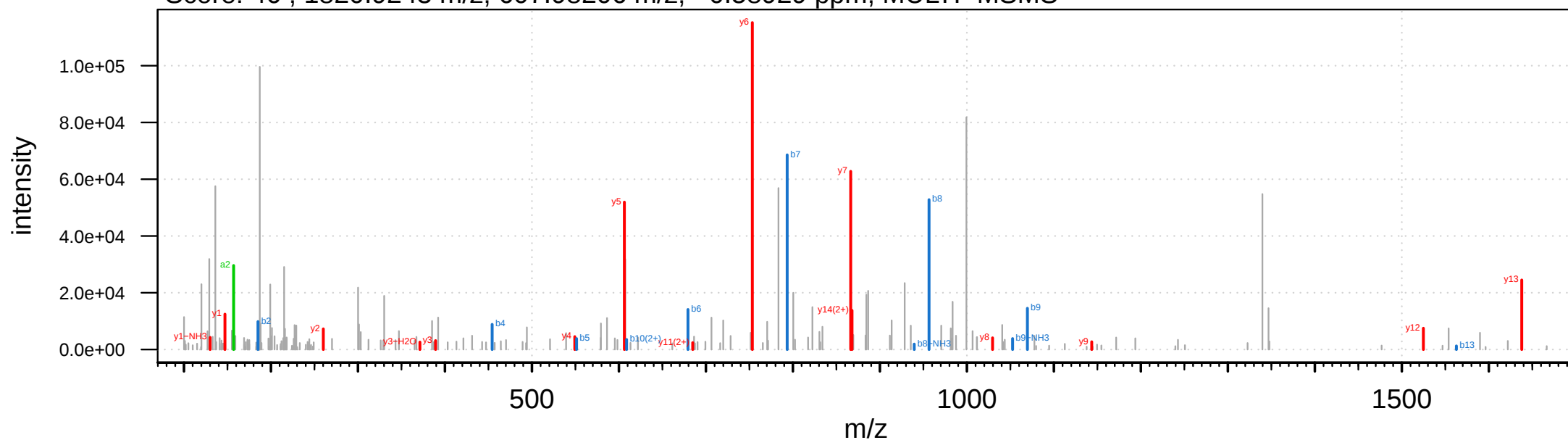
Proteins:

TCONS_I2_00008829_ch15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 46 ; 1820.9243 m/z; 607.98206 m/z; -0.58929 ppm; MULTI-MSMS

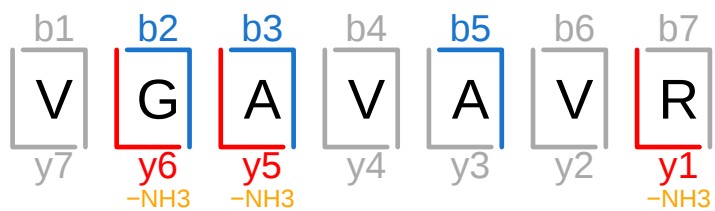


Raw File: 20101230_Velos1_TaGe_SA_Jurkat1

Scan Number: 24939

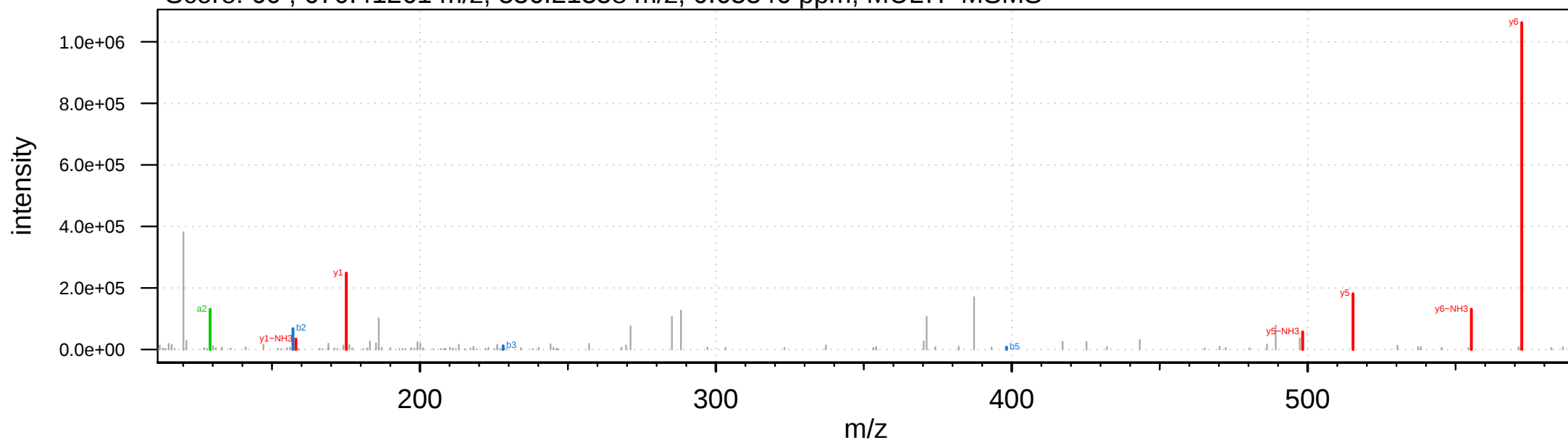
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

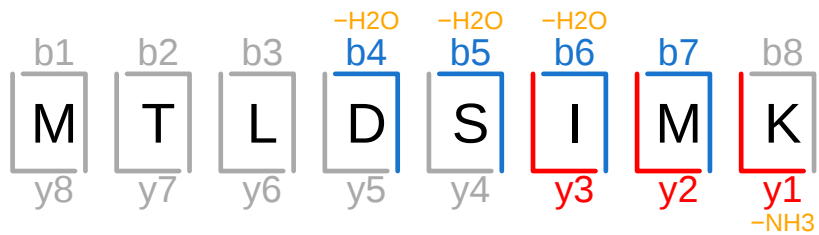


VGAVAVR

Score: 60 ; 670.41261 m/z; 336.21358 m/z; 0.63346 ppm; MULTI-MSMS

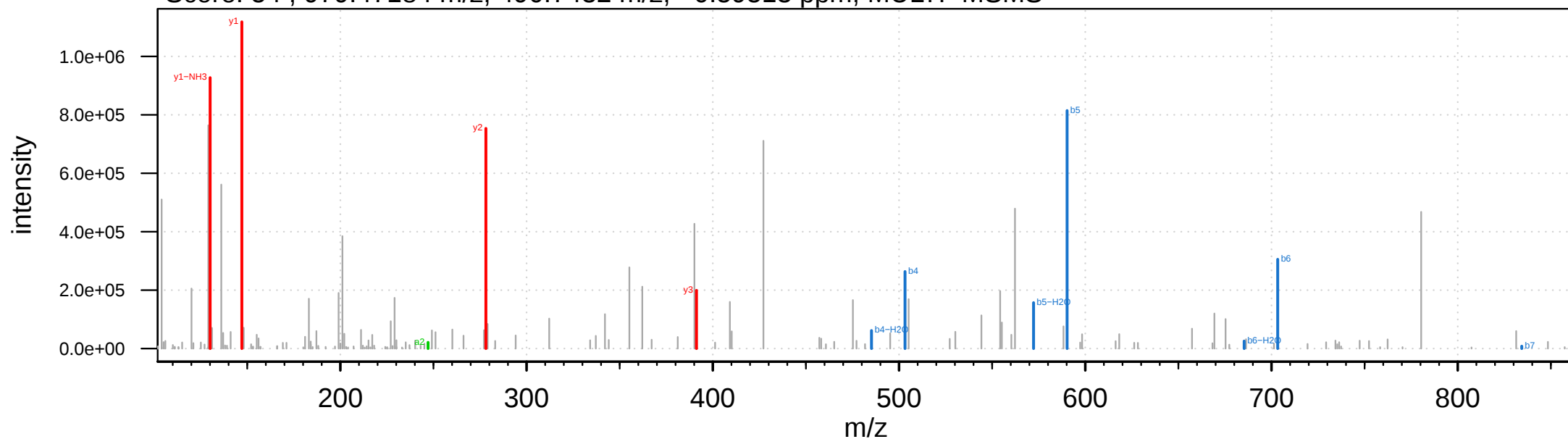


Raw File: 20101230_Velos1_TaGe_SA_Jurkat1
 Scan Number: 3081
 Proteins:
 TCONS_I2_00030545_chrX:79544539-79546436:-



(ac)MTLDSIMK

Score: 54 ; 979.47184 m/z; 490.7432 m/z; -0.59513 ppm; MULTI-MSMS

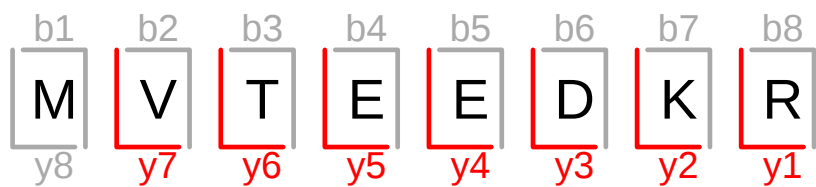


Raw File: 20101210_Velos1_AnWe_SA_U2OS_5

Scan Number: 17472

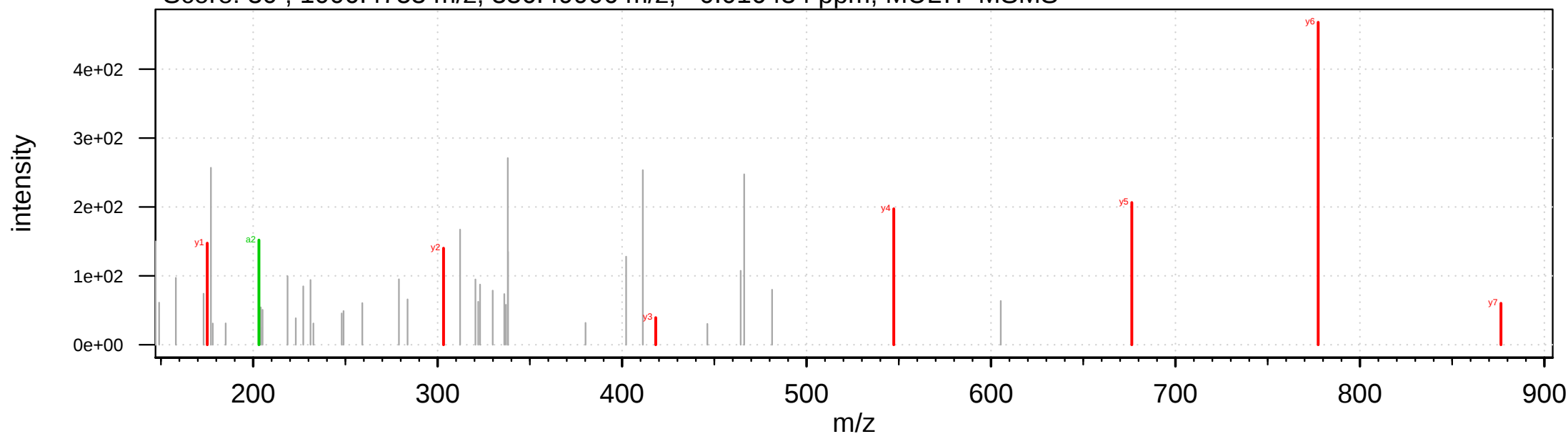
Proteins:

TCONS_I2_00001211_chr1:47348822-47348905:-



MVTEEDKR

Score: 30 ; 1006.4753 m/z; 336.49906 m/z; -0.016434 ppm; MULTI-MSMS

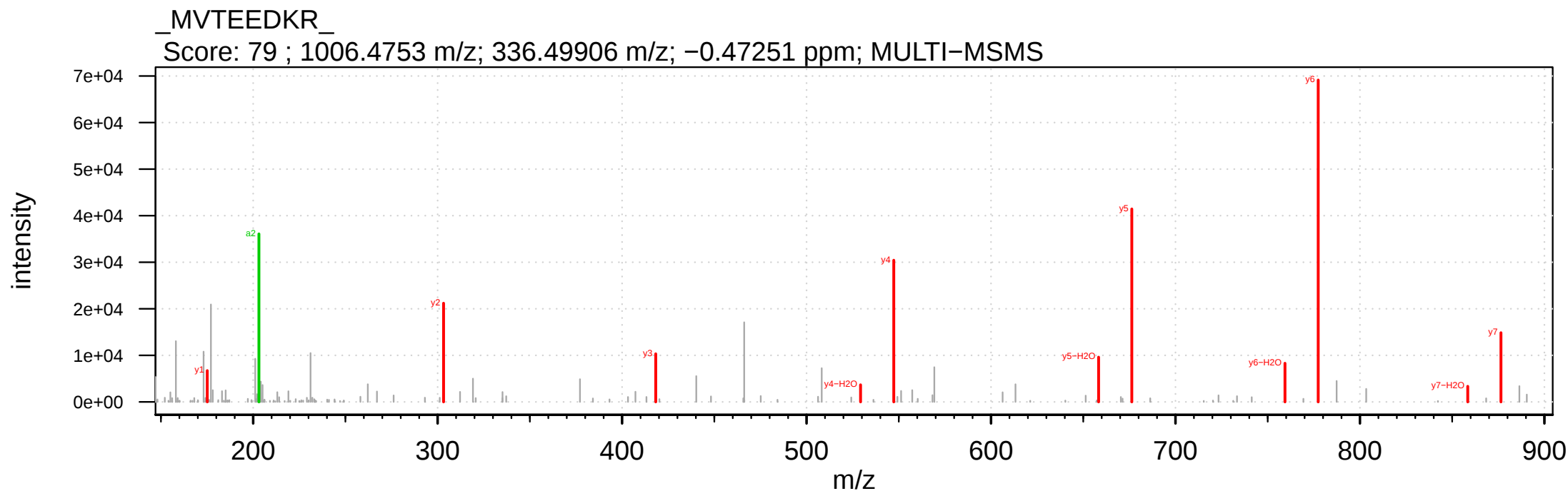
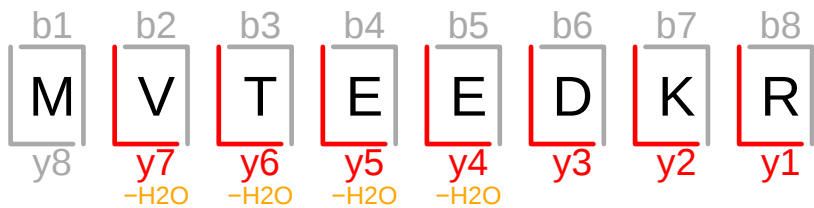


Raw File: 20100611_Velos1_TaGe_SA_Hela_3

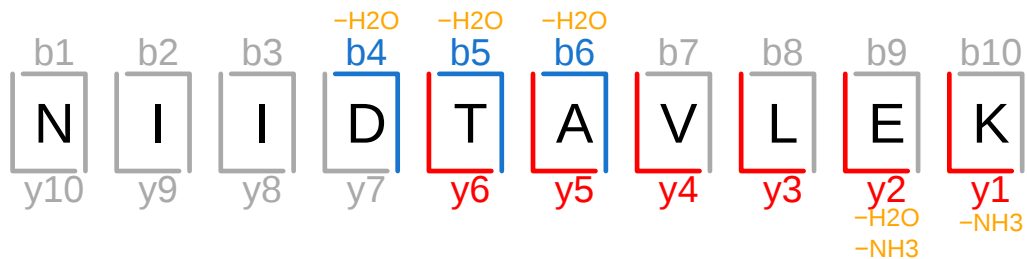
Scan Number: 1127

Proteins:

TCONS_I2_00006768_chr13:29174091-29174123:+

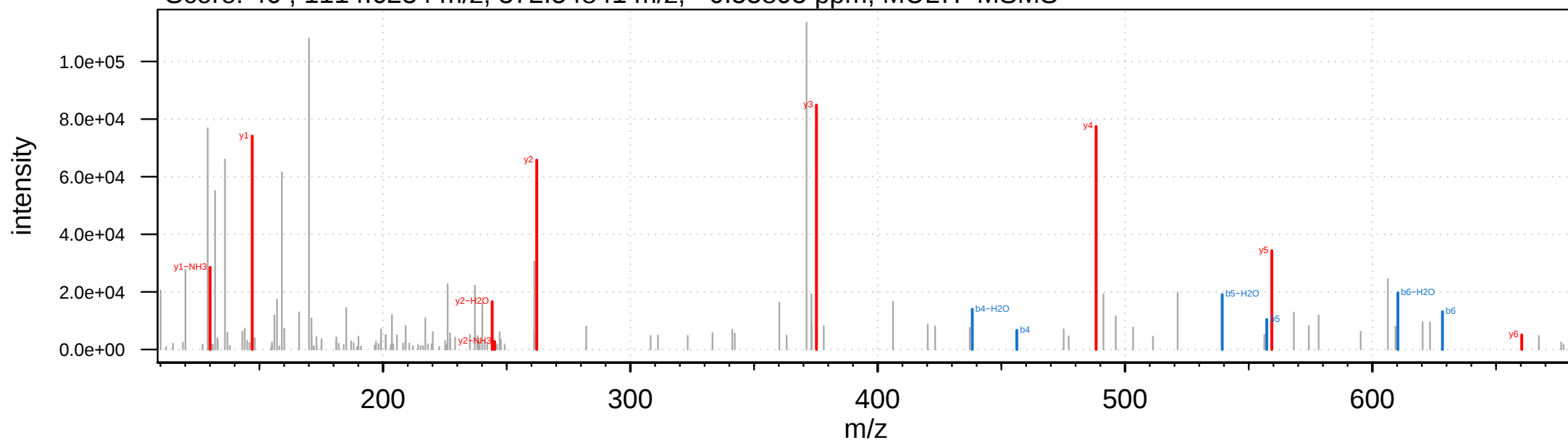


Raw File: 20101210_Velos1_AnWe_SA_U2OS_3
Scan Number: 1563
Proteins:
TCONS_I2_00006768_chr13:29174091-29174123:+



NIIDTAVLEK

Score: 46 ; 1114.6234 m/z; 372.54841 m/z; -0.55895 ppm; MULTI-MSMS

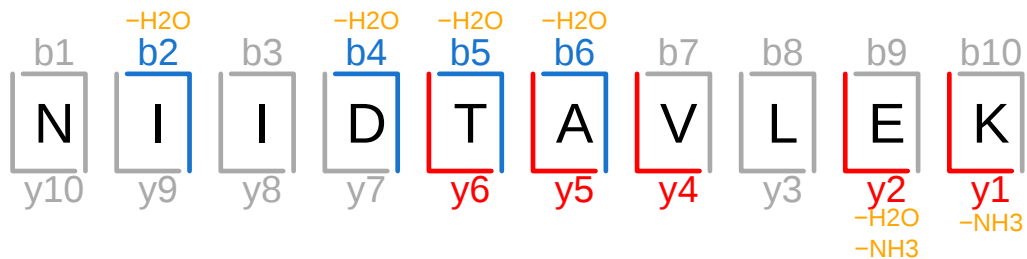


Raw File: 20101210_Velos1_AnWe_SA_U2OS_3

Scan Number: 8671

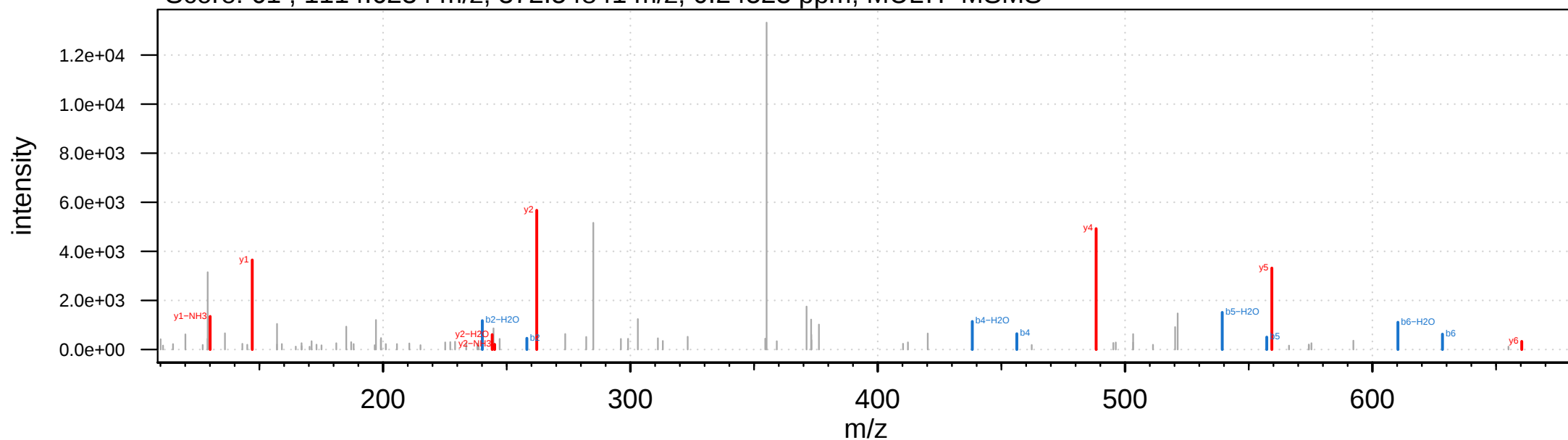
Proteins:

ENST00000497138_chr20:56806826-56807846:-



NIIDTAVLEK

Score: 61 ; 1114.6234 m/z; 372.54841 m/z; 0.24523 ppm; MULTI-MSMS

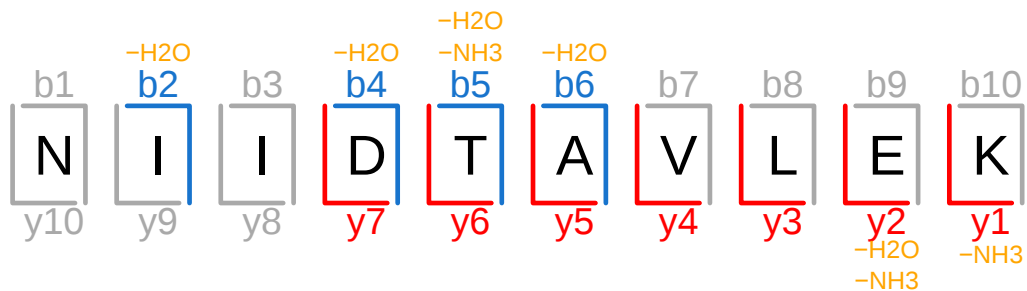


Raw File: 20100609_Velos1_TaGe_SA_293_5

Scan Number: 4694

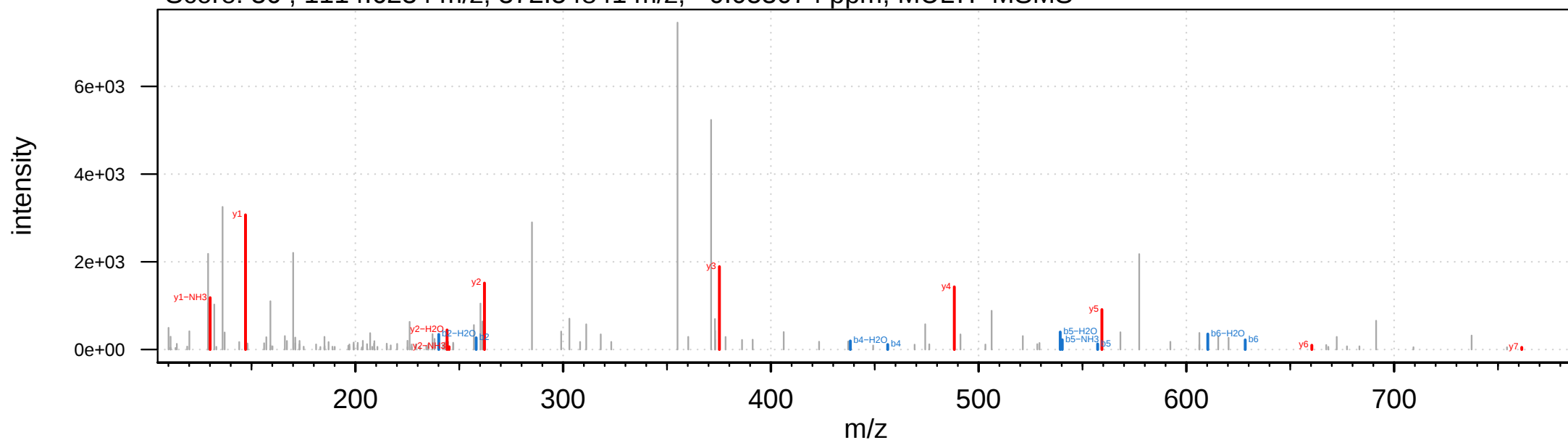
Proteins:

ENST00000497138_chr20:56806826-56807846:-



NIIDTAVLEK

Score: 36 ; 1114.6234 m/z; 372.54841 m/z; -0.053074 ppm; MULTI-MSMS

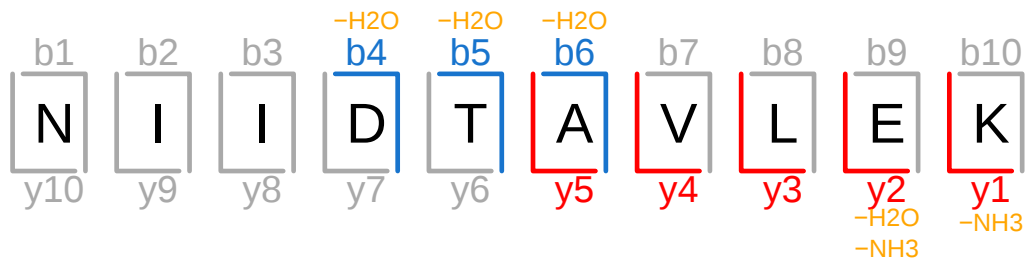


Raw File: 20100609_Velos1_TaGe_SA_GAMG_5

Scan Number: 4529

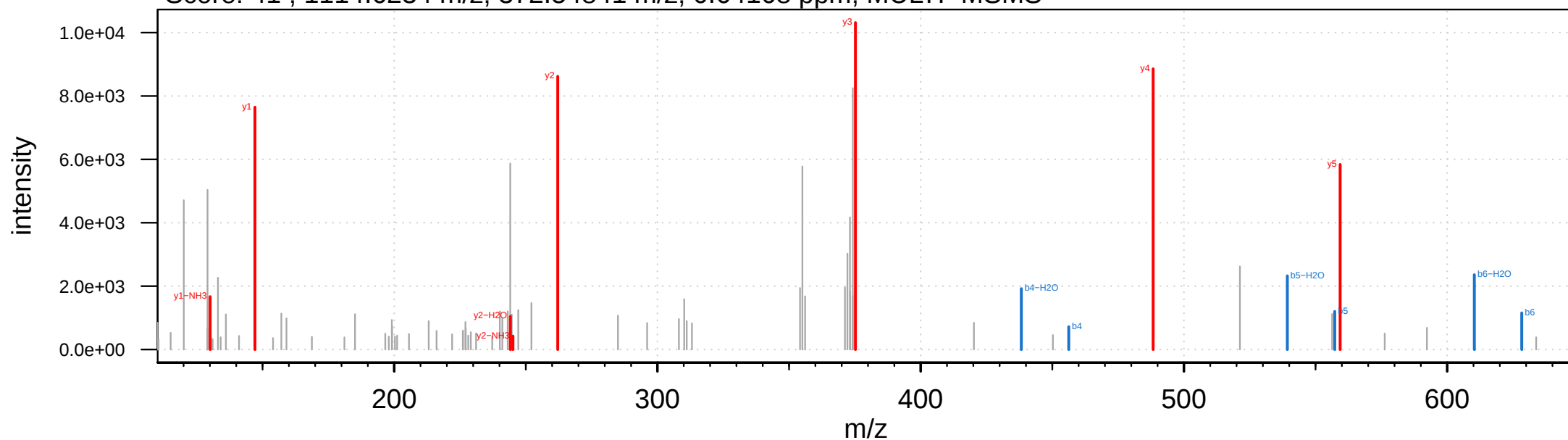
Proteins:

ENST00000497138_chr20:56806826-56807846:-



NIIDTAVLEK

Score: 41 ; 1114.6234 m/z; 372.54841 m/z; 0.64168 ppm; MULTI-MSMS

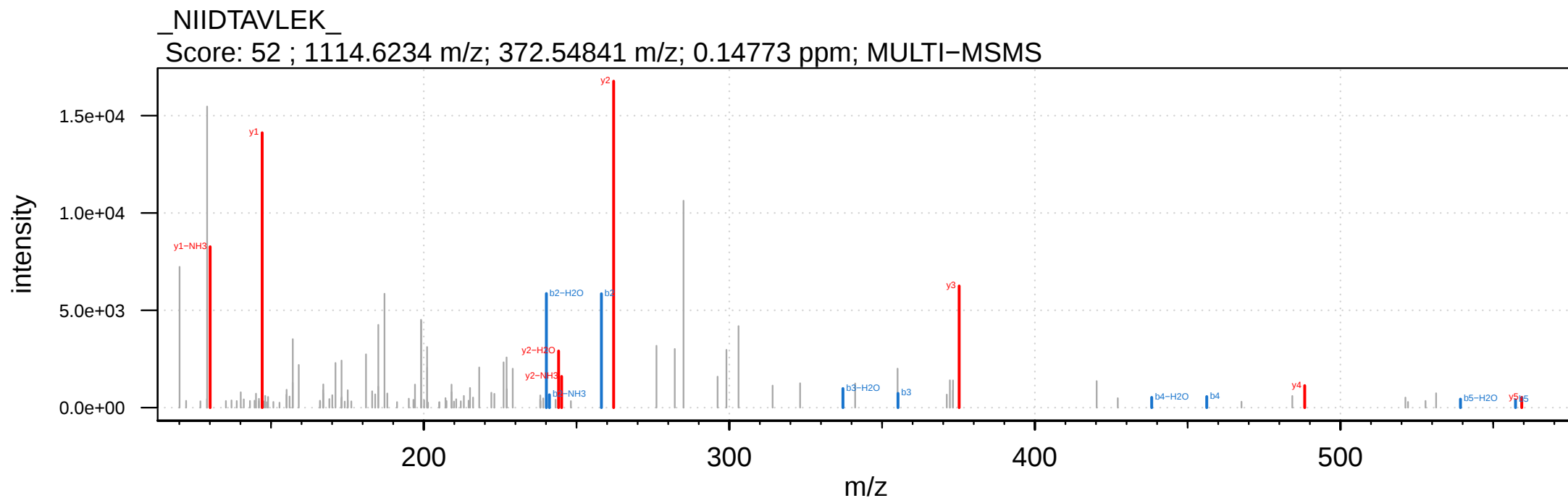
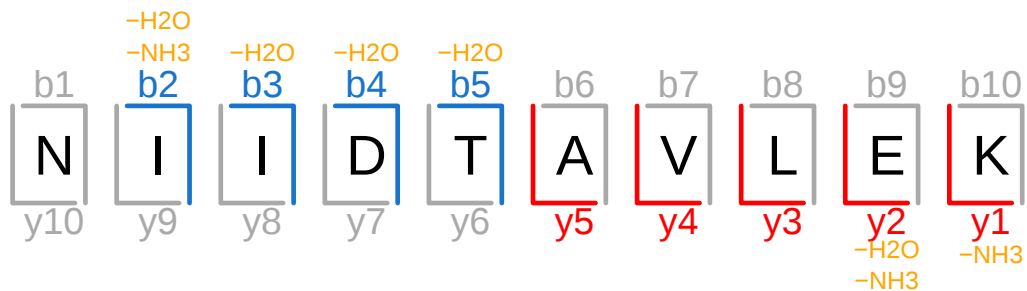


Raw File: 20100616_Velos1_TaGe_SA_RKO_3

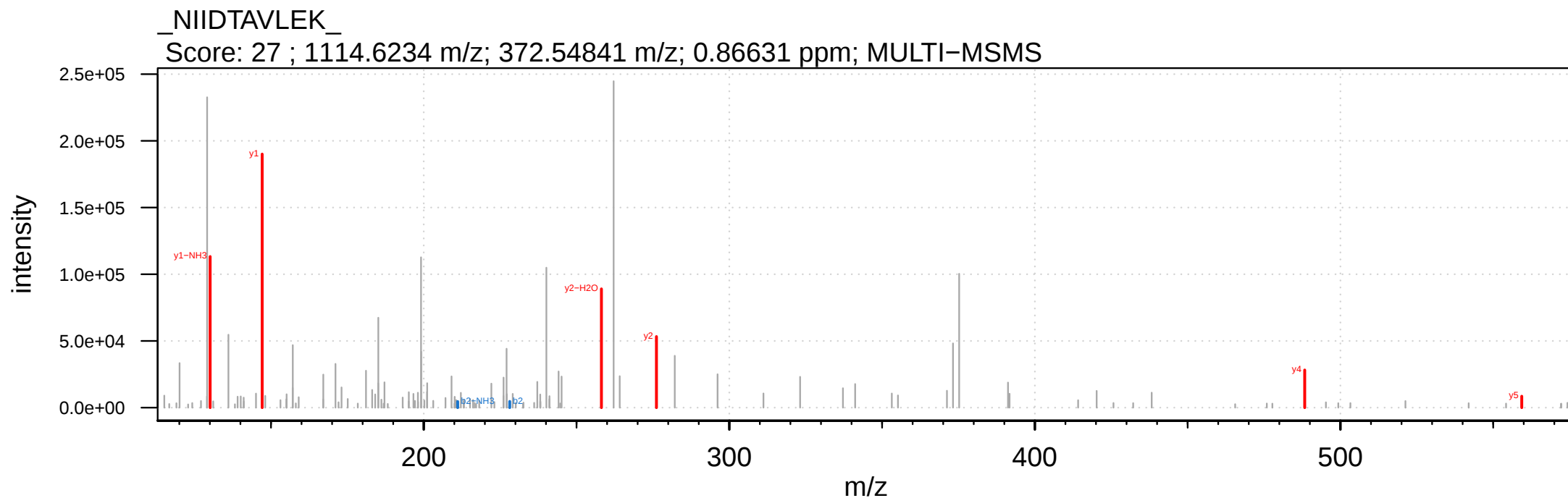
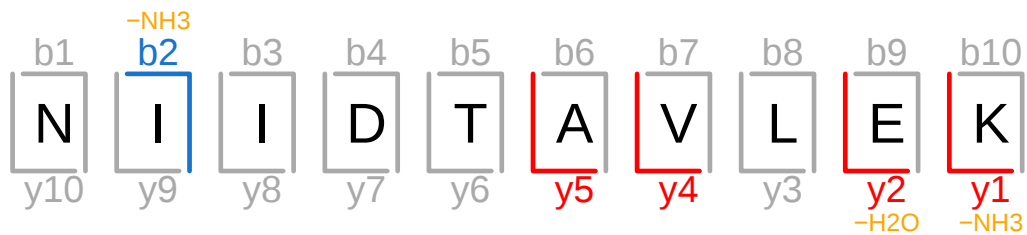
Scan Number: 7681

Proteins:

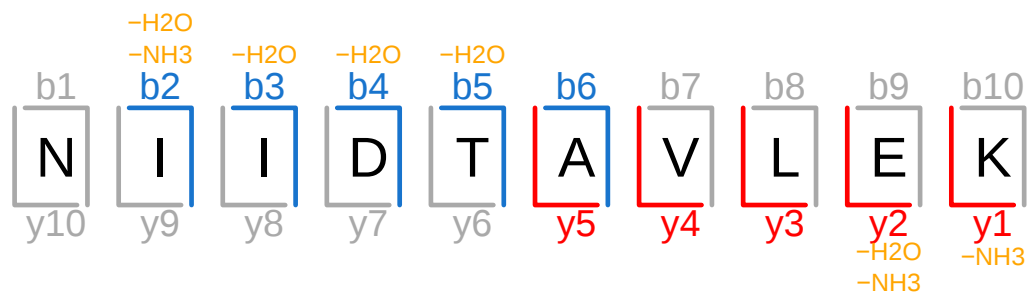
ENST00000497138_chr20:56806826-56807846:-



Raw File: 20100721_Velos1_TaGe_SA_A549_04
 Scan Number: 7276
 Proteins:
 ENST00000497138_chr20:56806826-56807846:-

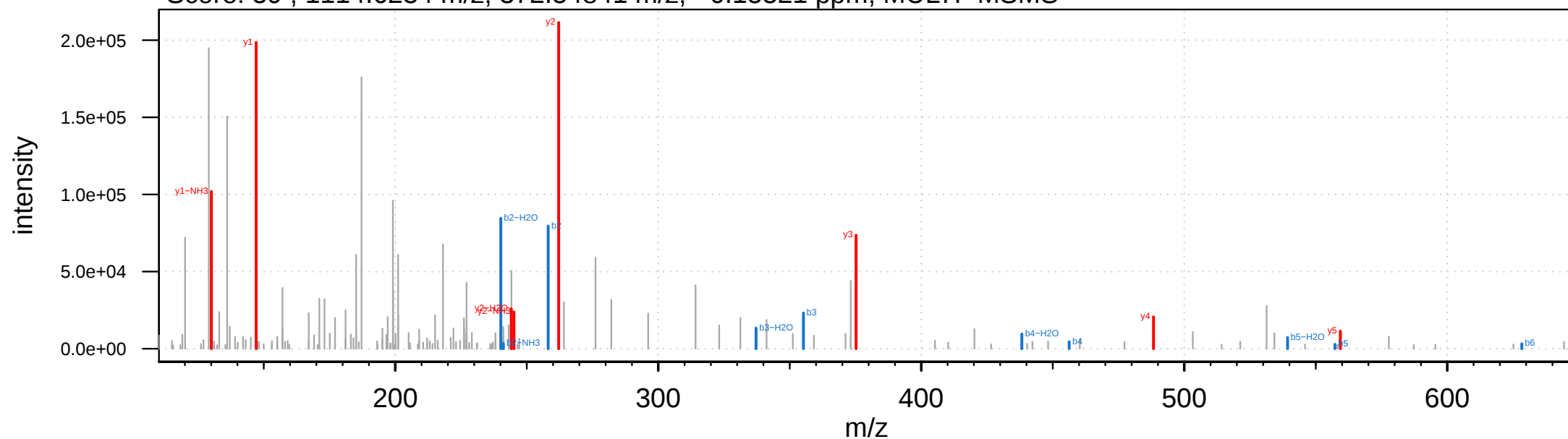


Raw File: 20100723_Velos1_TaGe_SA_Hek293_02
Scan Number: 7568
Proteins:
ENST00000497138_chr20:56806826-56807846:-



NIIDTAVLEK

Score: 59 ; 1114.6234 m/z; 372.54841 m/z; -0.15321 ppm; MULTI-MSMS

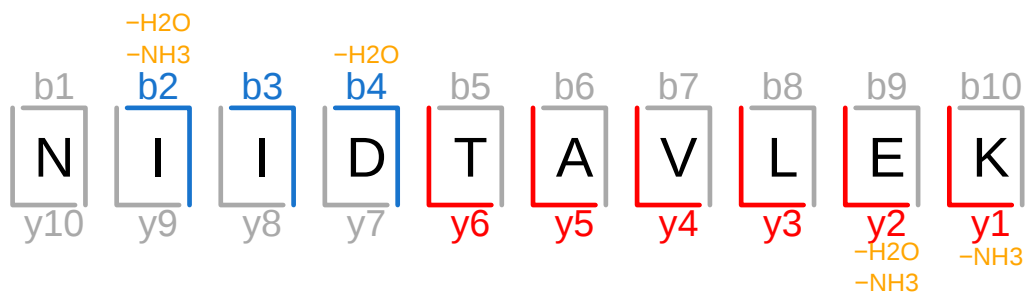


Raw File: 20100723_Velos1_TaGe_SA_Hek293_03

Scan Number: 7807

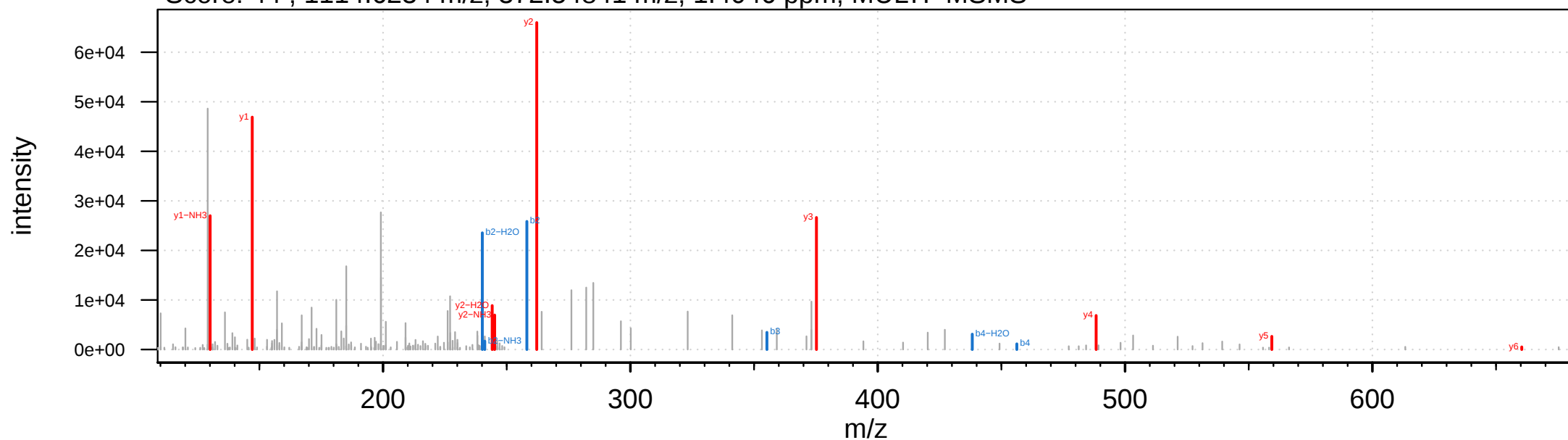
Proteins:

ENST00000497138_chr20:56806826-56807846:-



NIIDTAVLEK

Score: 44 ; 1114.6234 m/z; 372.54841 m/z; 1.4046 ppm; MULTI-MSMS

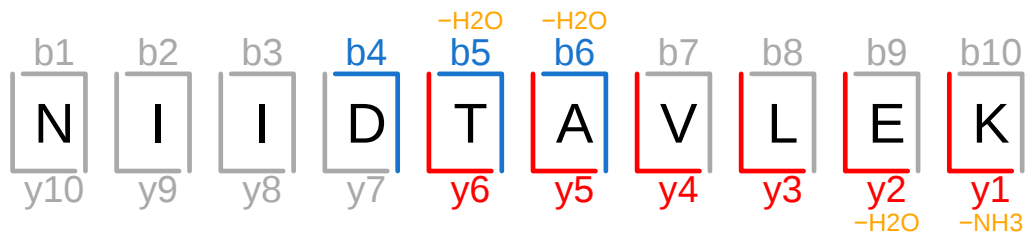


Raw File: 20100723_Velos1_TaGe_SA_Hek293_04

Scan Number: 7858

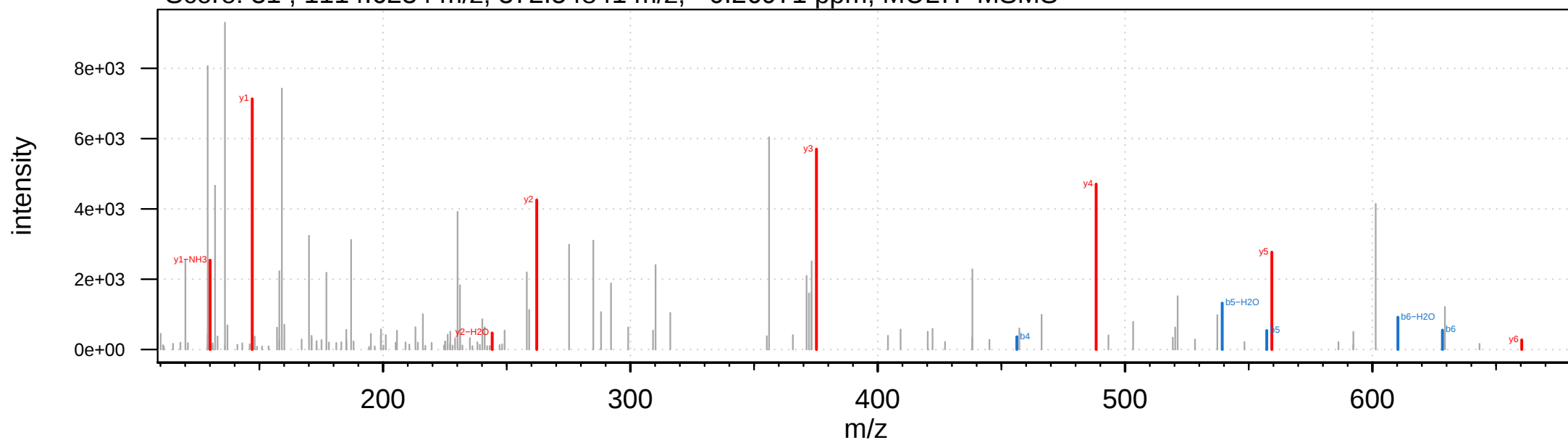
Proteins:

ENST00000497138_chr20:56806826-56807846:-

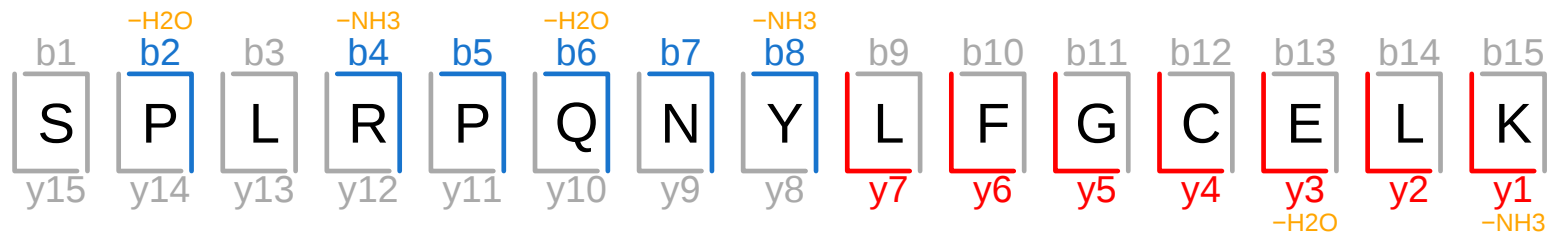


NIIDTAVLEK

Score: 31 ; 1114.6234 m/z; 372.54841 m/z; -0.26971 ppm; MULTI-MSMS

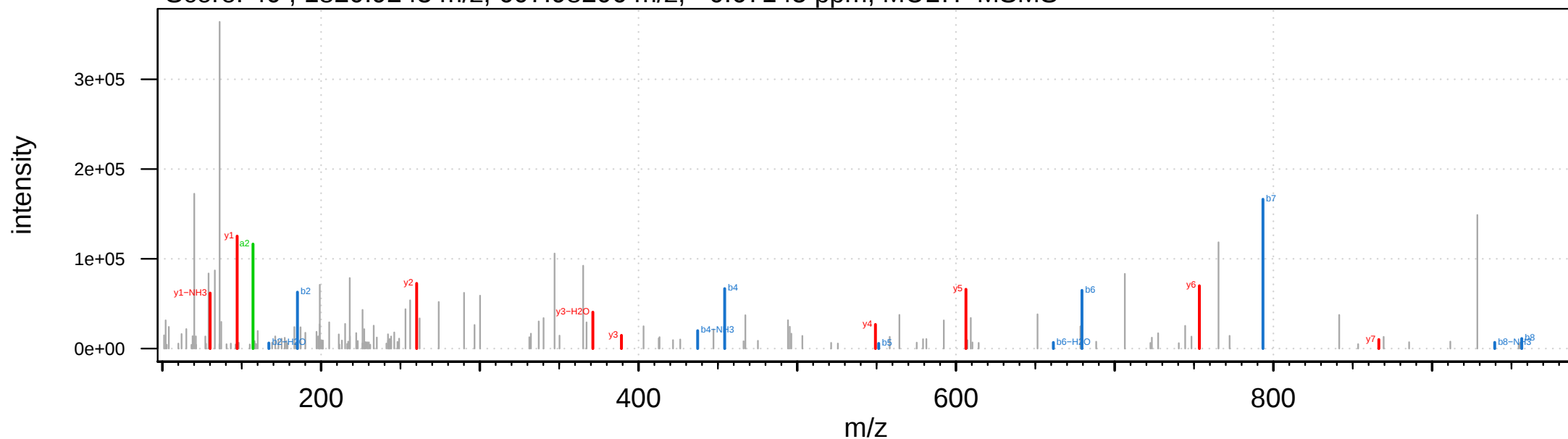


Raw File: 20101230_Velos1_TaGe_SA_Jurkat4
 Scan Number: 6642
 Proteins:
 ENST00000497138_chr20:56806826-56807846:-

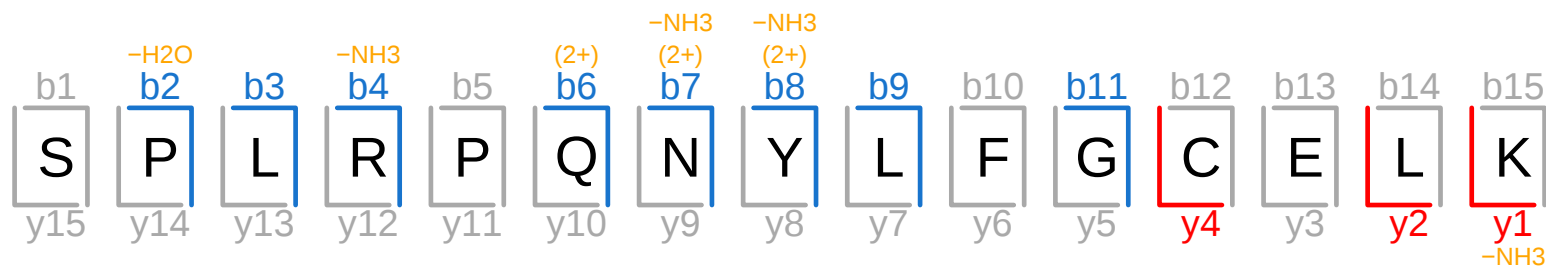


SPLRPQNYLFGCELK

Score: 40 ; 1820.9243 m/z; 607.98206 m/z; -0.67145 ppm; MULTI-MSMS

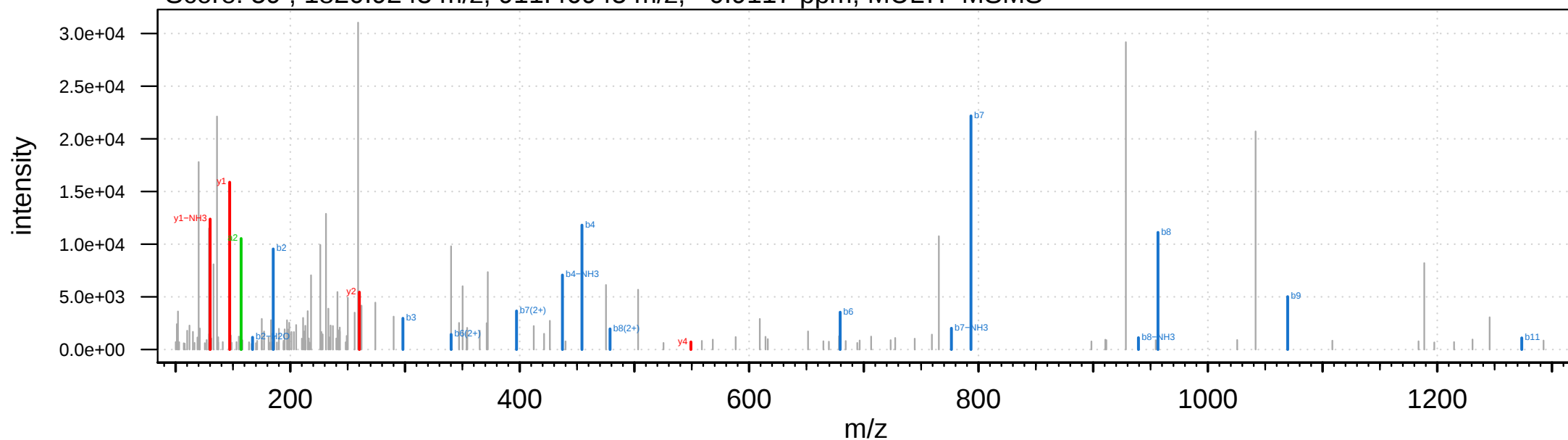


Raw File: 20100719_Velos1_TaGe_SA_LnCap_2
 Scan Number: 20183
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

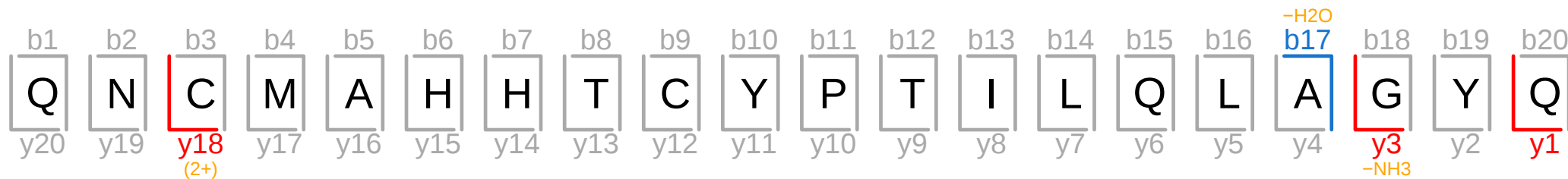


SPLRPQNYLFGCELK

Score: 59 ; 1820.9243 m/z; 911.46945 m/z; -0.9117 ppm; MULTI-MSMS

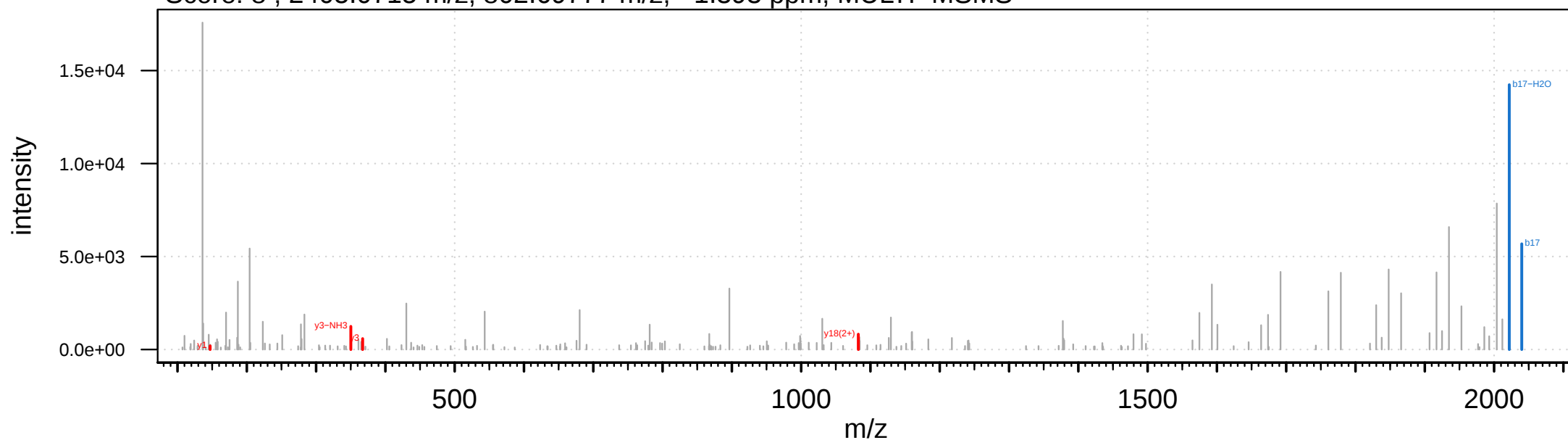


Raw File: 20100719_Velos1_TaGe_SA_LnCap_2
 Scan Number: 20212
 Proteins:
 TCONS_I2_00008829_ch15:92829088-92829258:+



QNCMAHHTCYPTILQLAGYQ

Score: 8 ; 2405.0715 m/z; 802.69777 m/z; -1.393 ppm; MULTI-MSMS

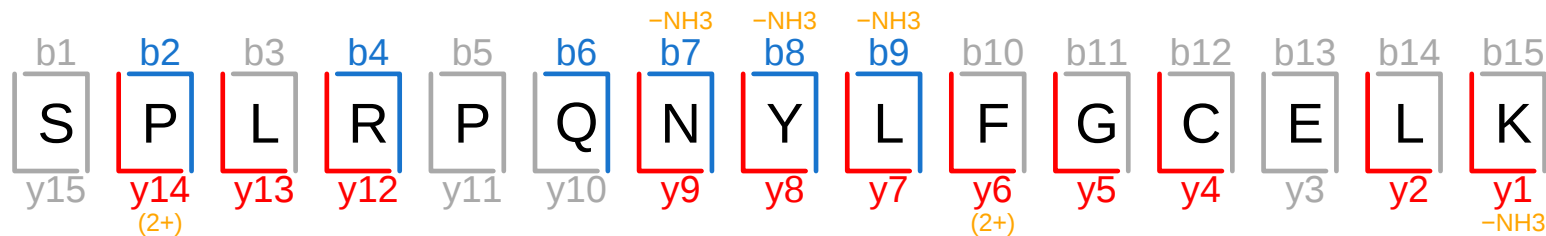


Raw File: 20100604_Velos1_TaGe_SA_A549_3

Scan Number: 12478

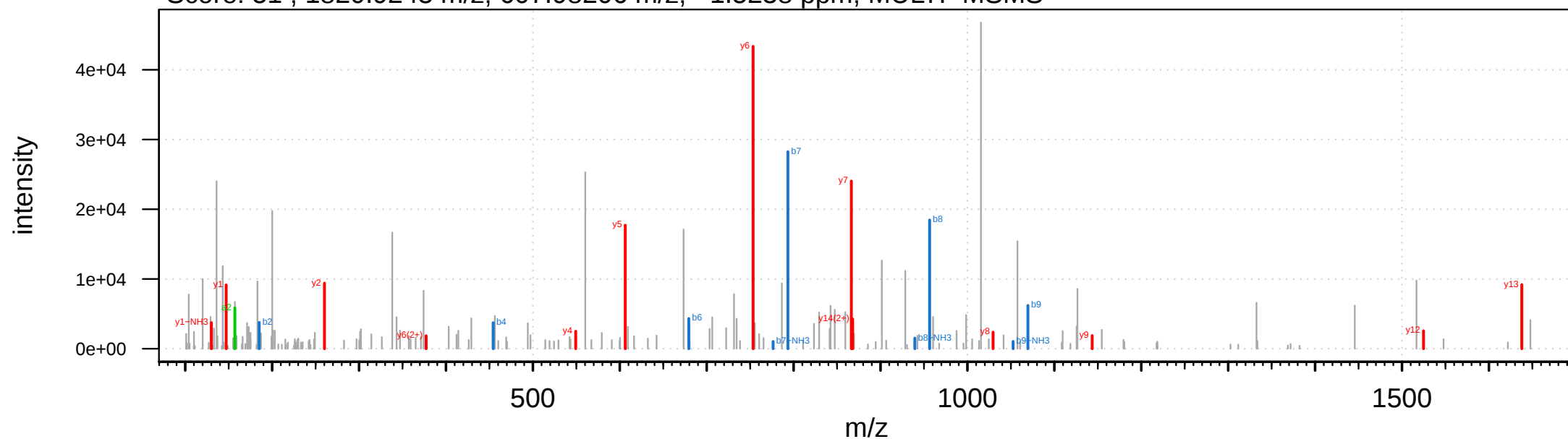
Proteins:

TCONS_I2_00030931_chrY:13309531-13309647:-



SPLRPQNYLFGCELK

Score: 51 ; 1820.9243 m/z; 607.98206 m/z; -1.5238 ppm; MULTI-MSMS

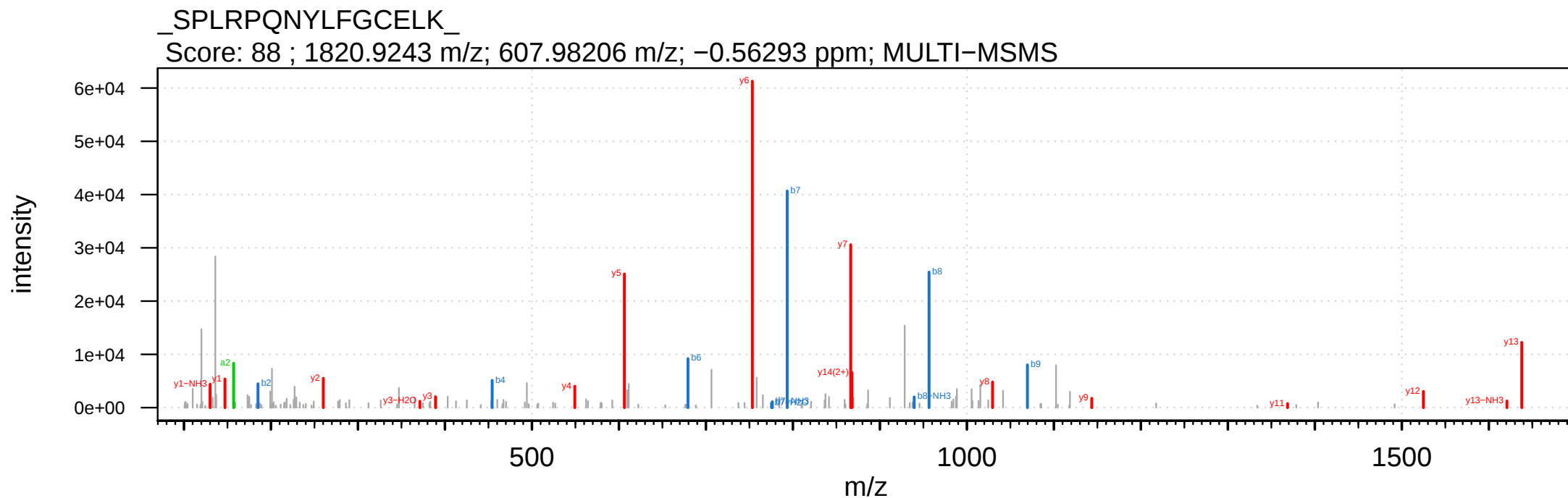
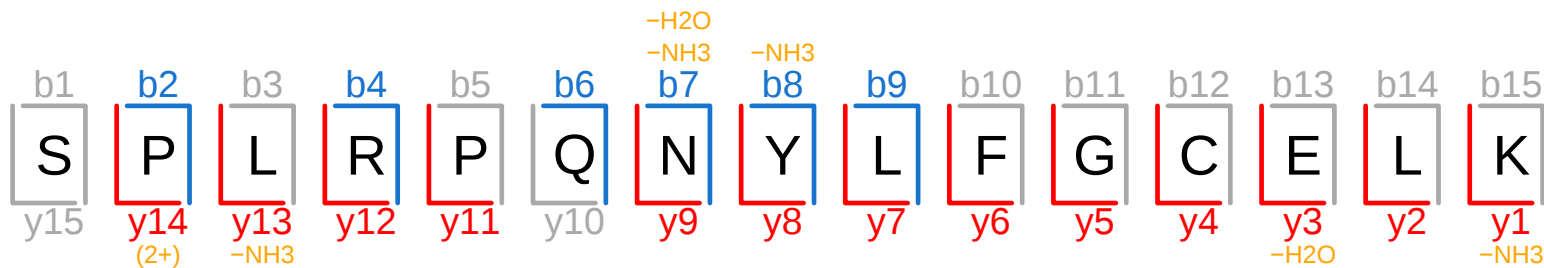


Raw File: 20100604_Velos1_TaGe_SA_A549_2

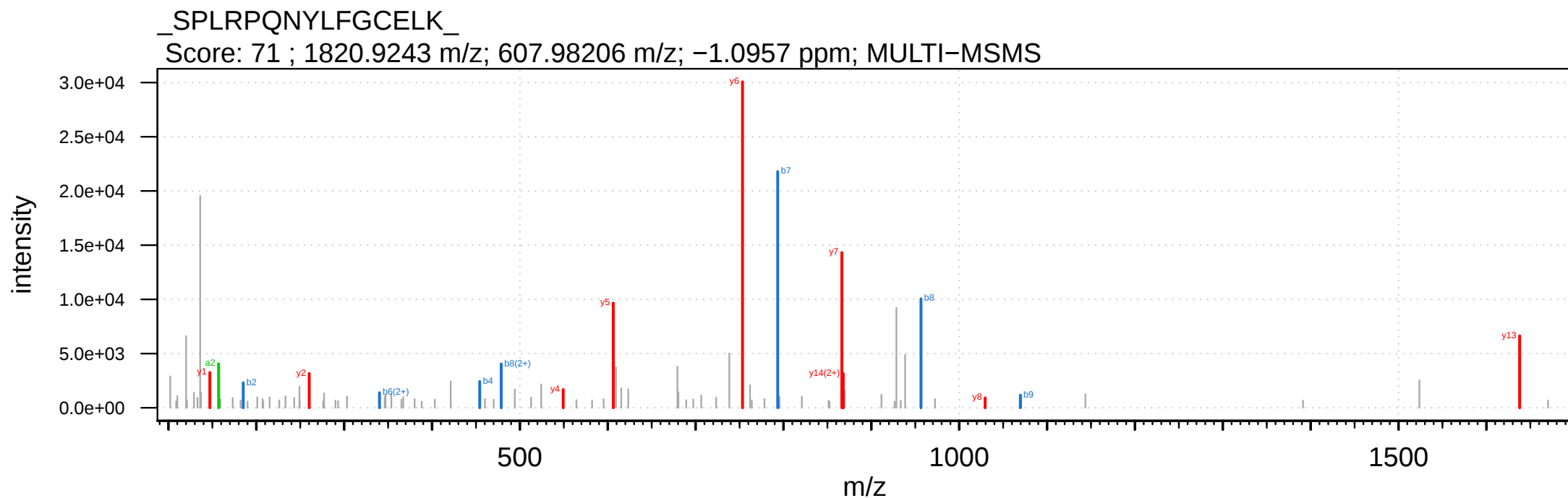
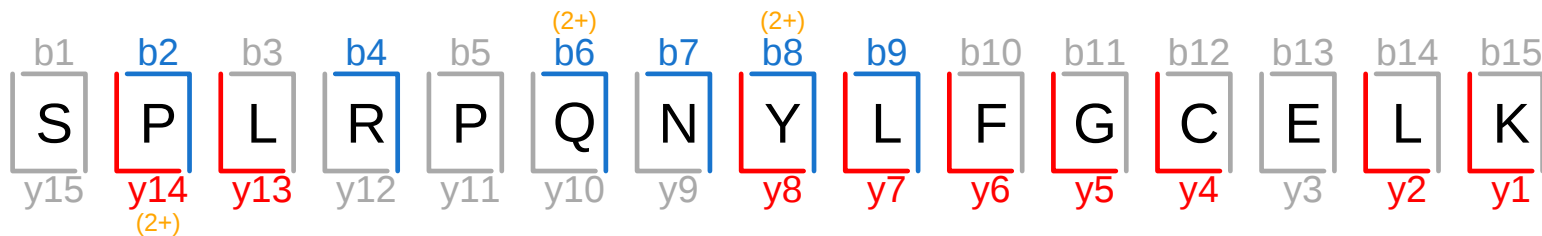
Scan Number: 15896

Proteins:

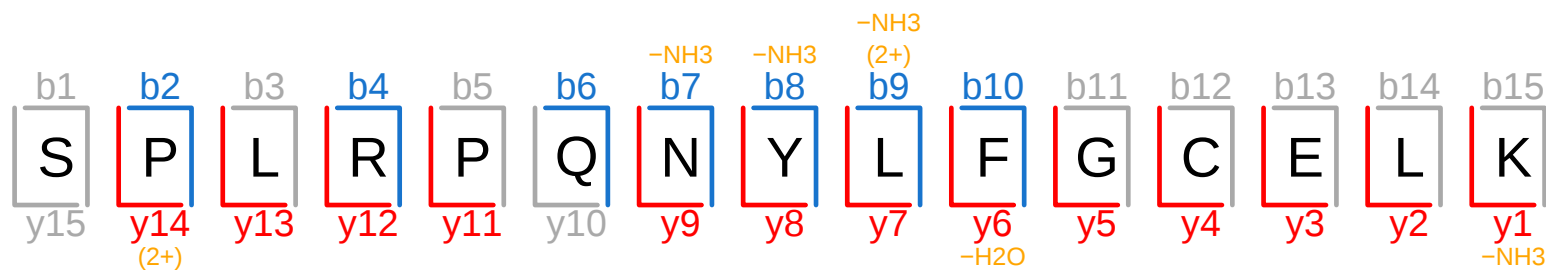
TCONS_I2_00008829_chr15:92829088-92829258:+



Raw File: 20100609_Velos1_TaGe_SA_GAMG_2
 Scan Number: 16160
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

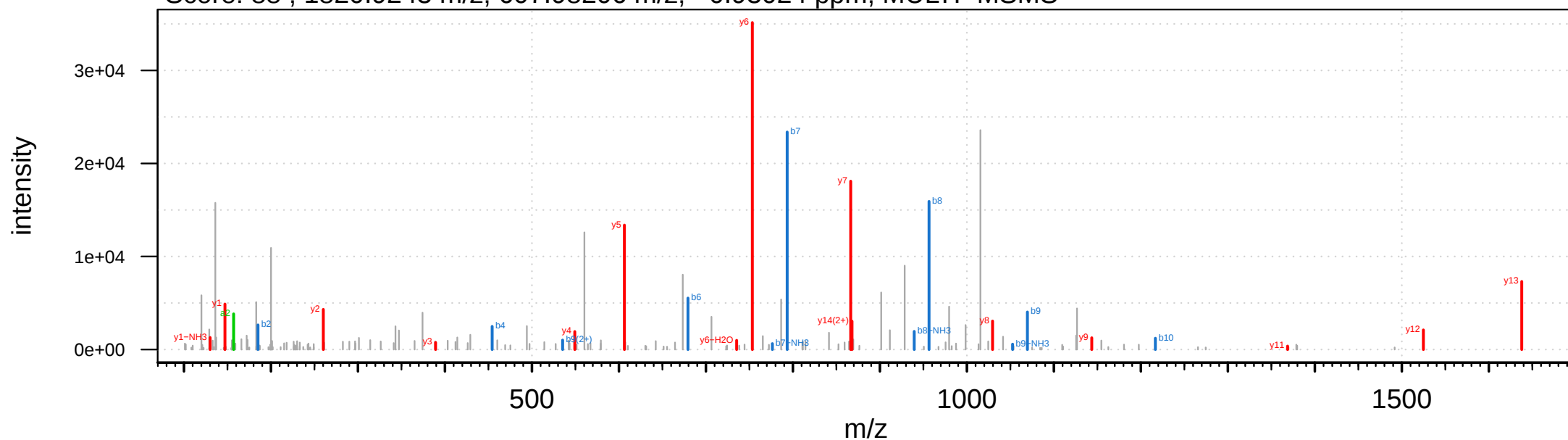


Raw File: 20100611_Velos1_TaGe_SA_Hela_6
 Scan Number: 13703
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 88 ; 1820.9243 m/z; 607.98206 m/z; -0.93024 ppm; MULTI-MSMS

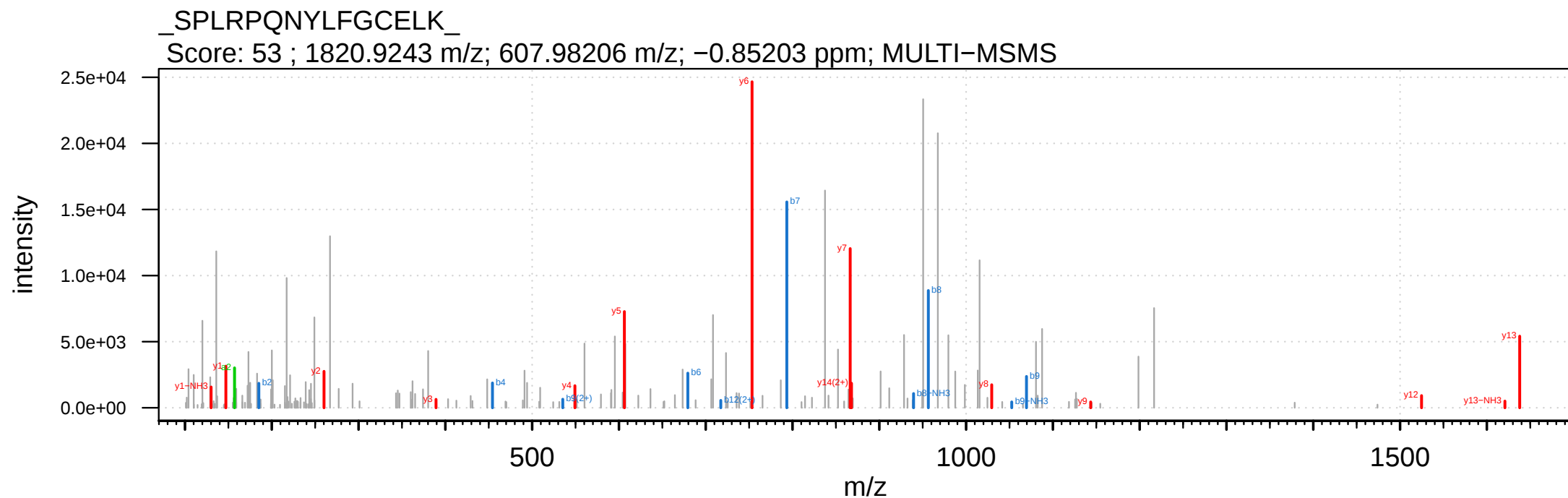
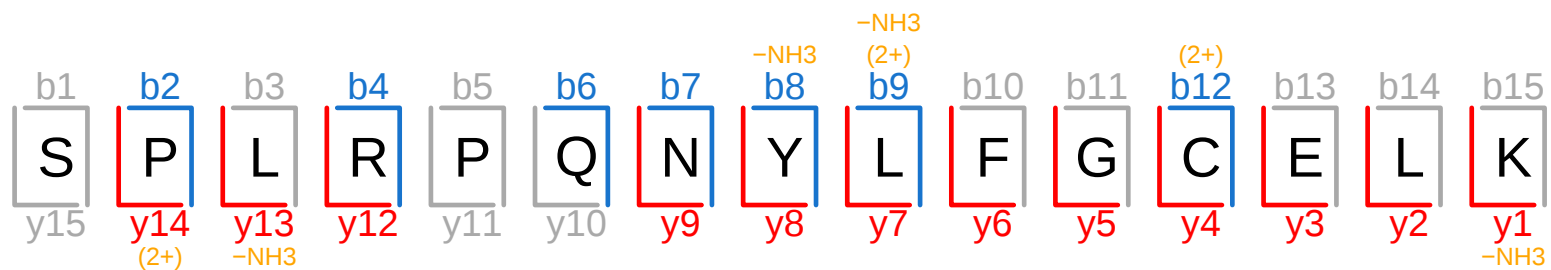


Raw File: 20100611_Velos1_TaGe_SA_HepG2_2

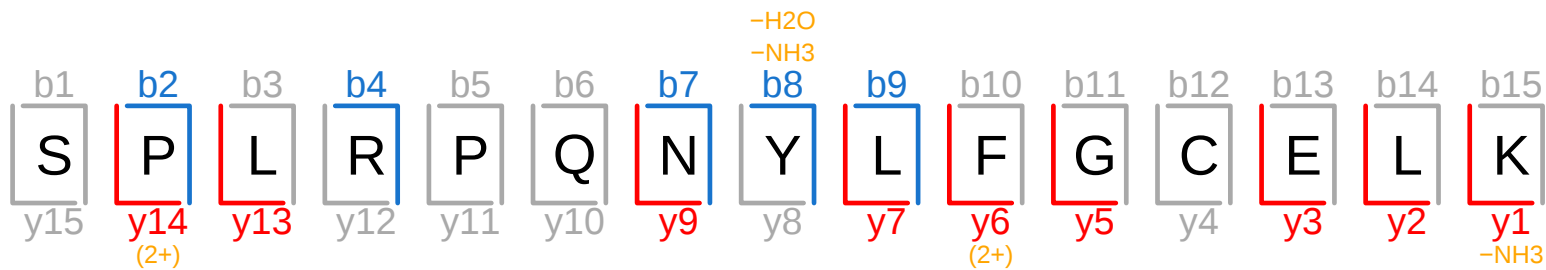
Scan Number: 17273

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

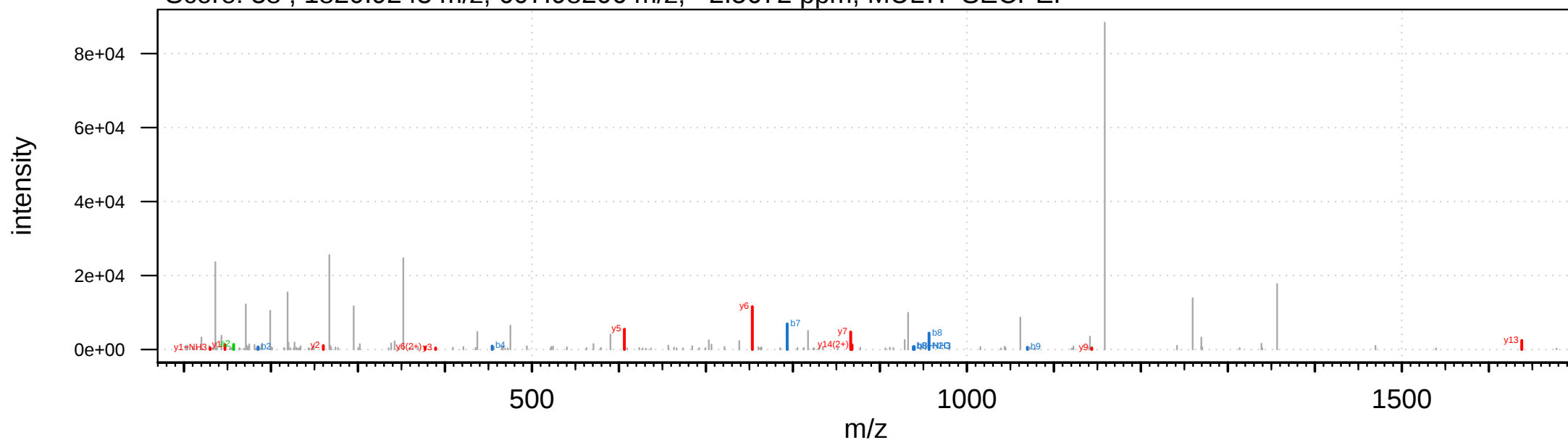


Raw File: 20100611_Velos1_TaGe_SA_HepG2_3
 Scan Number: 16508
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 38 ; 1820.9243 m/z; 607.98206 m/z; -2.5672 ppm; MULTI-SECPEP

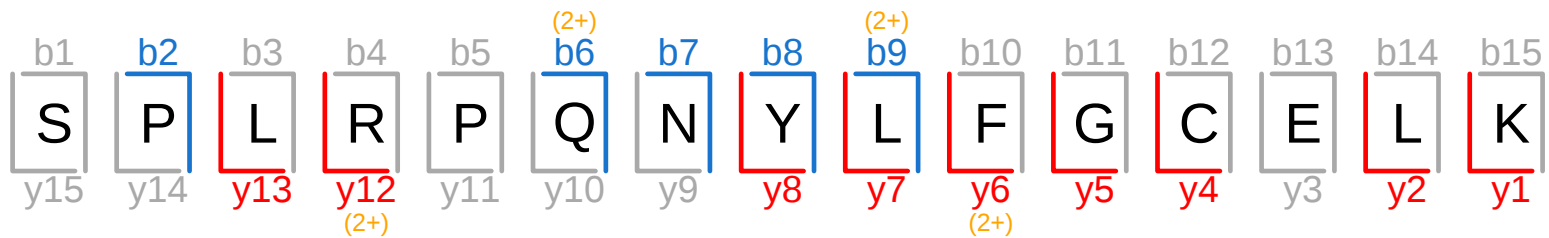


Raw File: 20100611_Velos1_TaGe_SA_HepG2_6

Scan Number: 11162

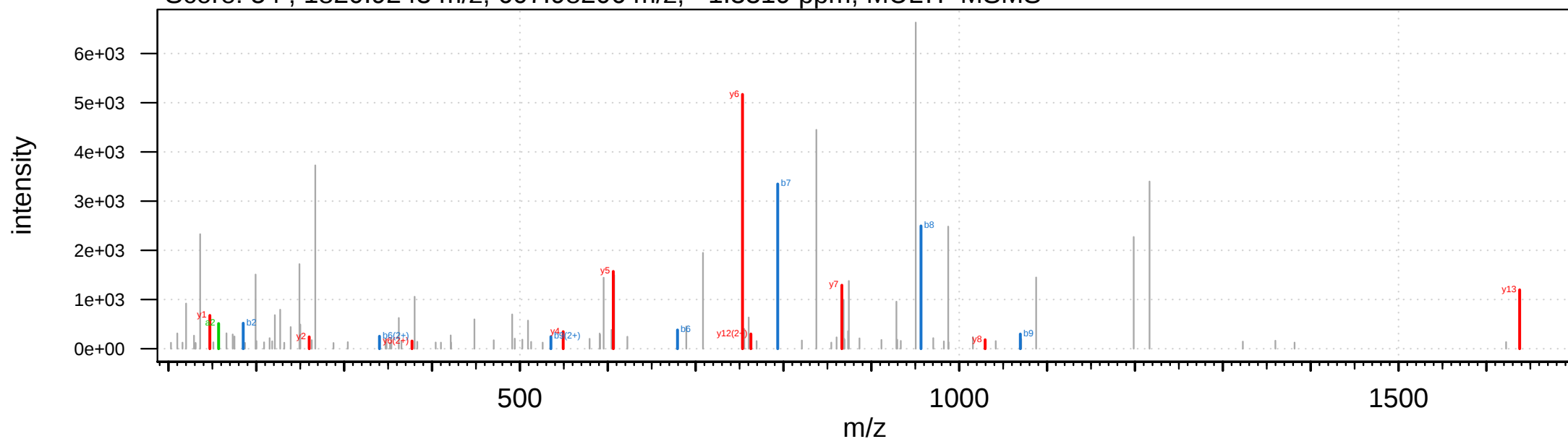
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 54 ; 1820.9243 m/z; 607.98206 m/z; -1.3319 ppm; MULTI-MSMS

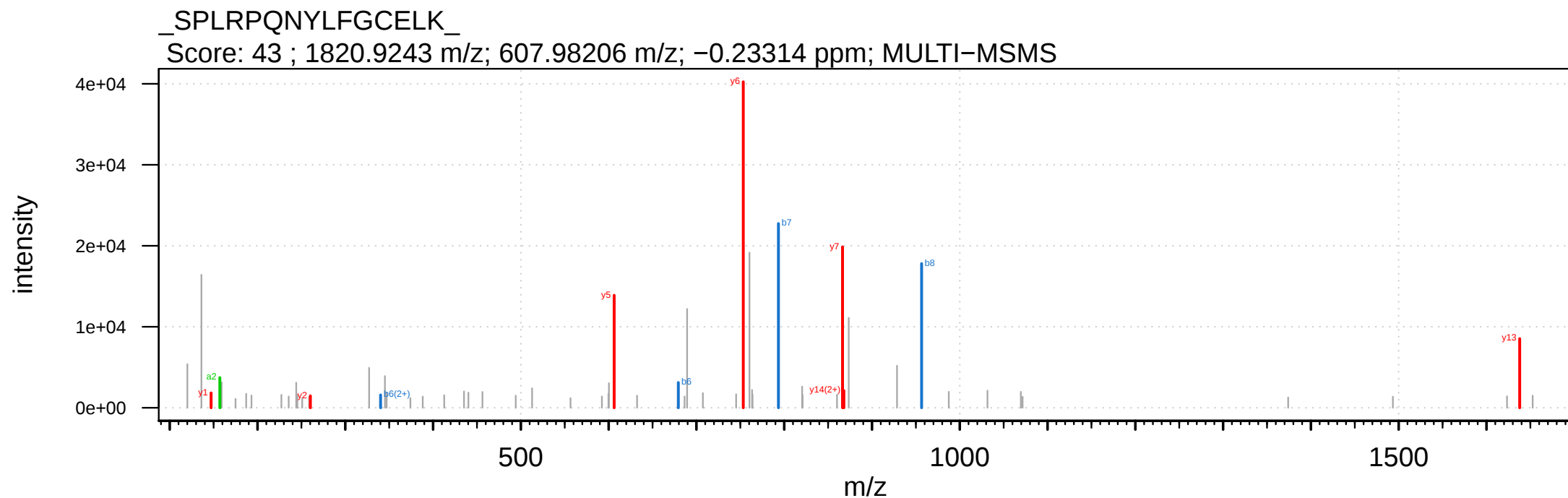
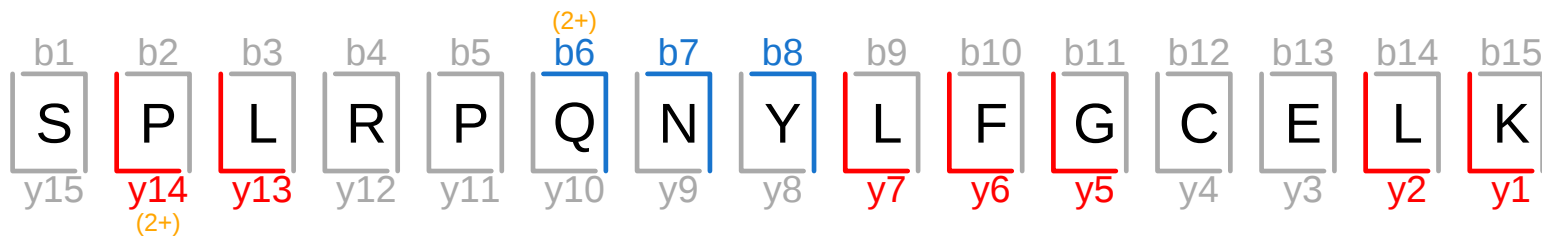


Raw File: 20100614_Velos1-TaGe_SA_Jurkat_2

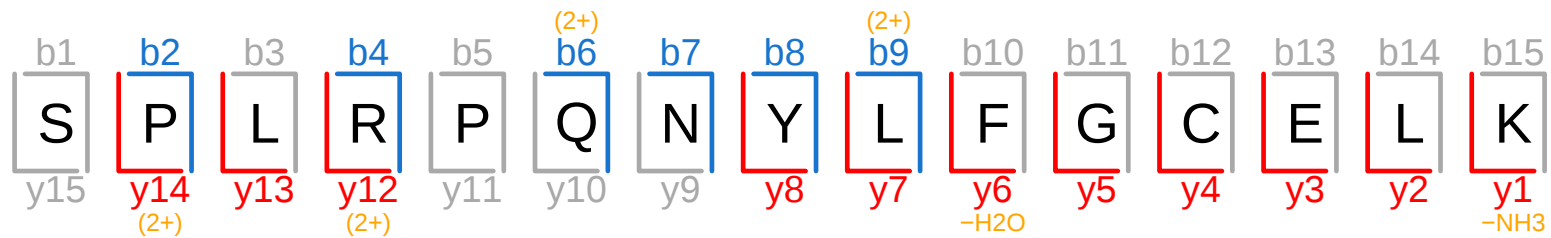
Scan Number: 18378

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

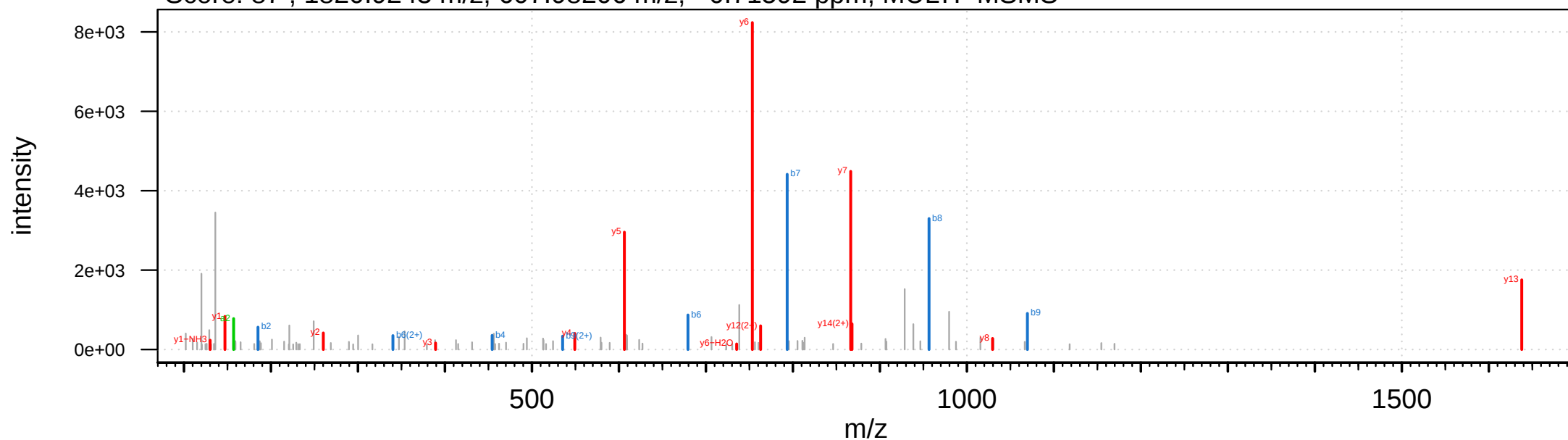


Raw File: 20100616_Velos1_TaGe_SA_RKO_2
Scan Number: 19011
Proteins:
TCONS_I2_00008829_ch15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 87 ; 1820.9243 m/z; 607.98206 m/z; -0.71592 ppm; MULTI-MSMS

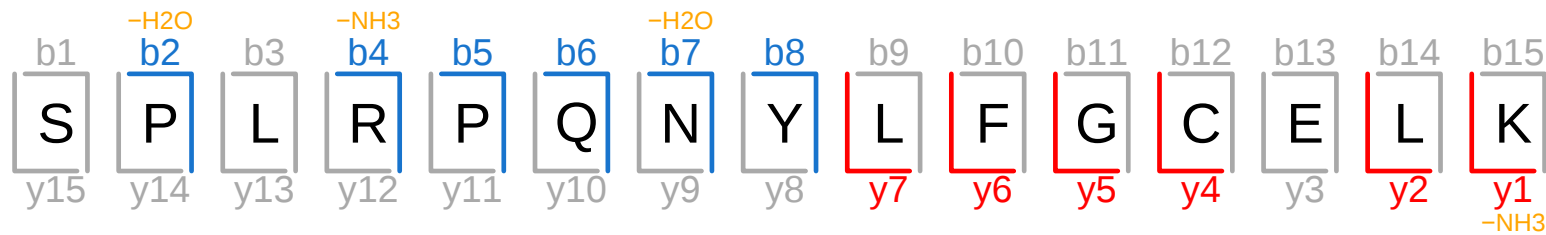


Raw File: 20100616_Velos1_TaGe_SA_RKO_6

Scan Number: 15140

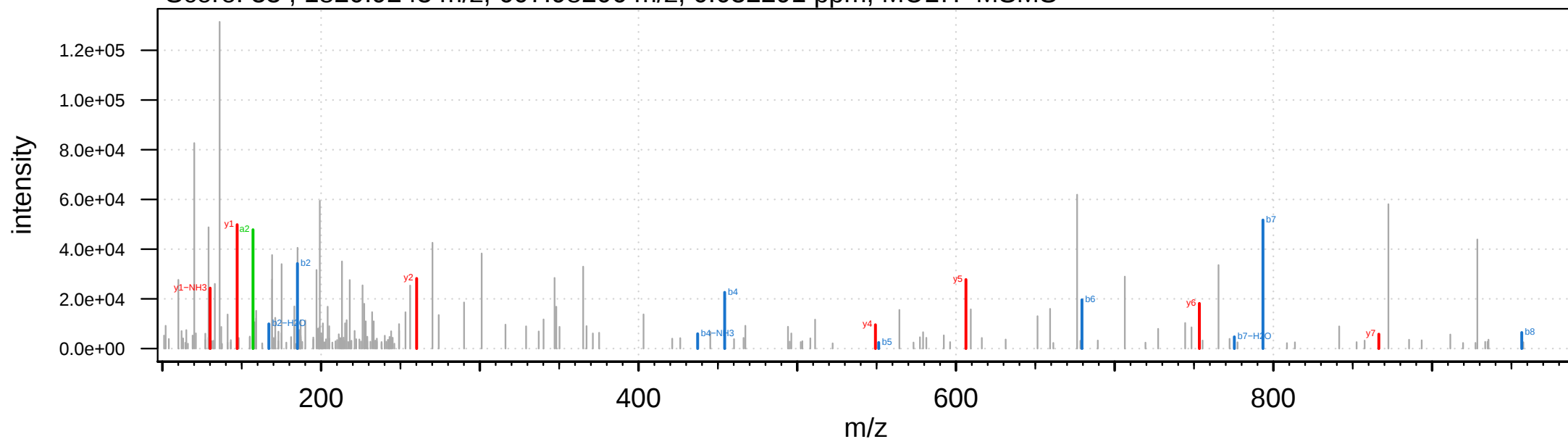
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 33 ; 1820.9243 m/z; 607.98206 m/z; 0.032291 ppm; MULTI-MSMS

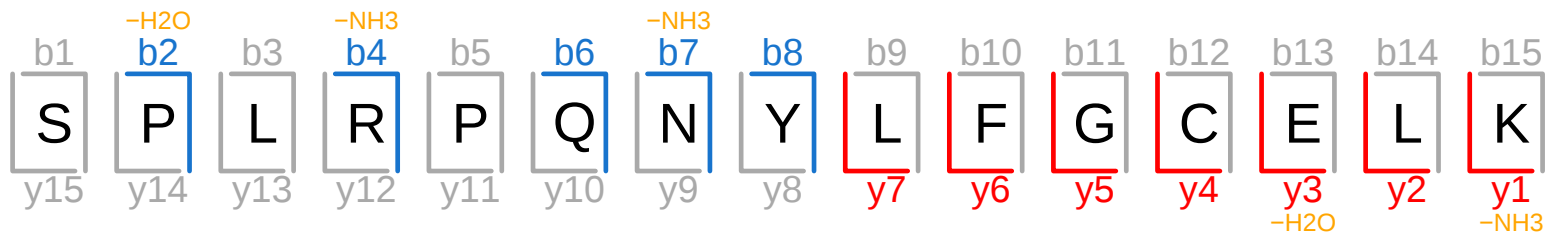


Raw File: 20100721_Velos1_TaGe_SA_A549_01

Scan Number: 18501

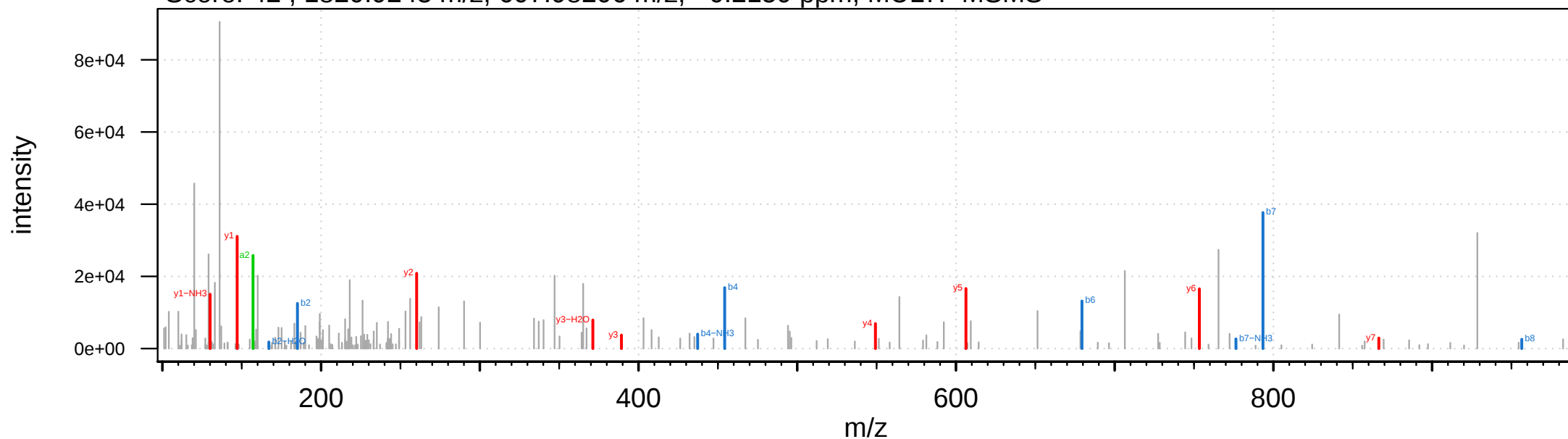
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 42 ; 1820.9243 m/z; 607.98206 m/z; -0.2159 ppm; MULTI-MSMS

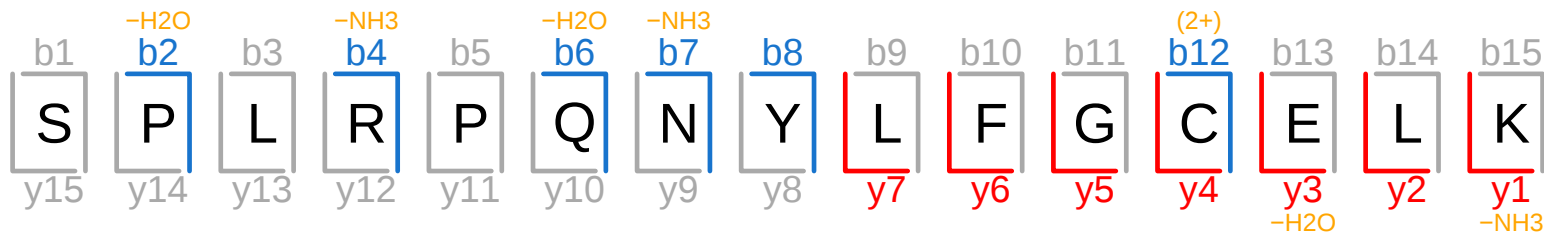


Raw File: 20100721_Velos1_TaGe_SA_A549_02

Scan Number: 19876

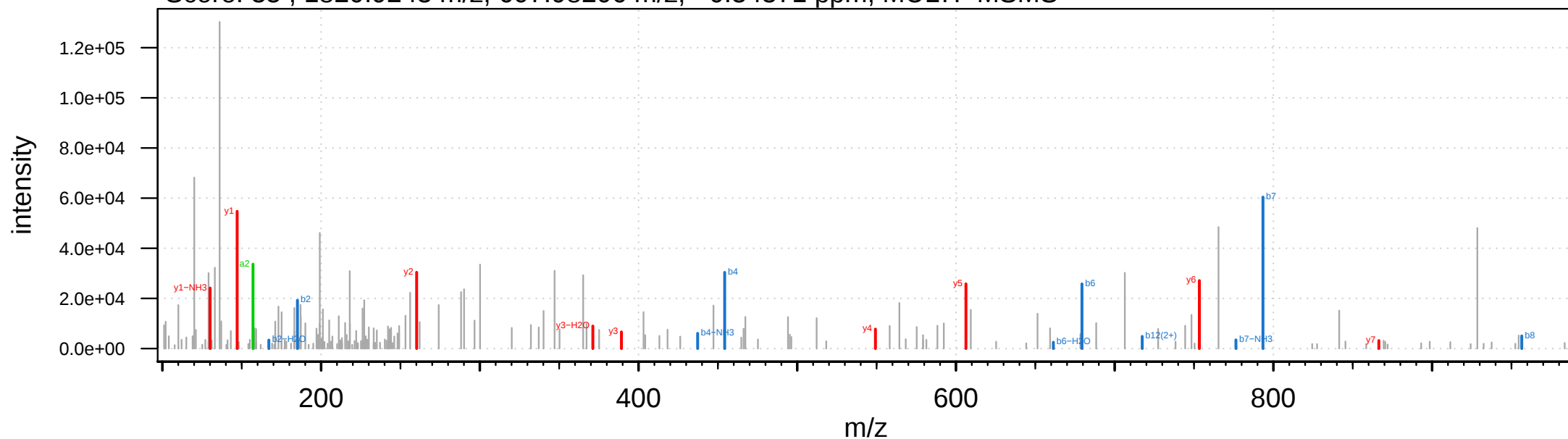
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 35 ; 1820.9243 m/z; 607.98206 m/z; -0.34371 ppm; MULTI-MSMS

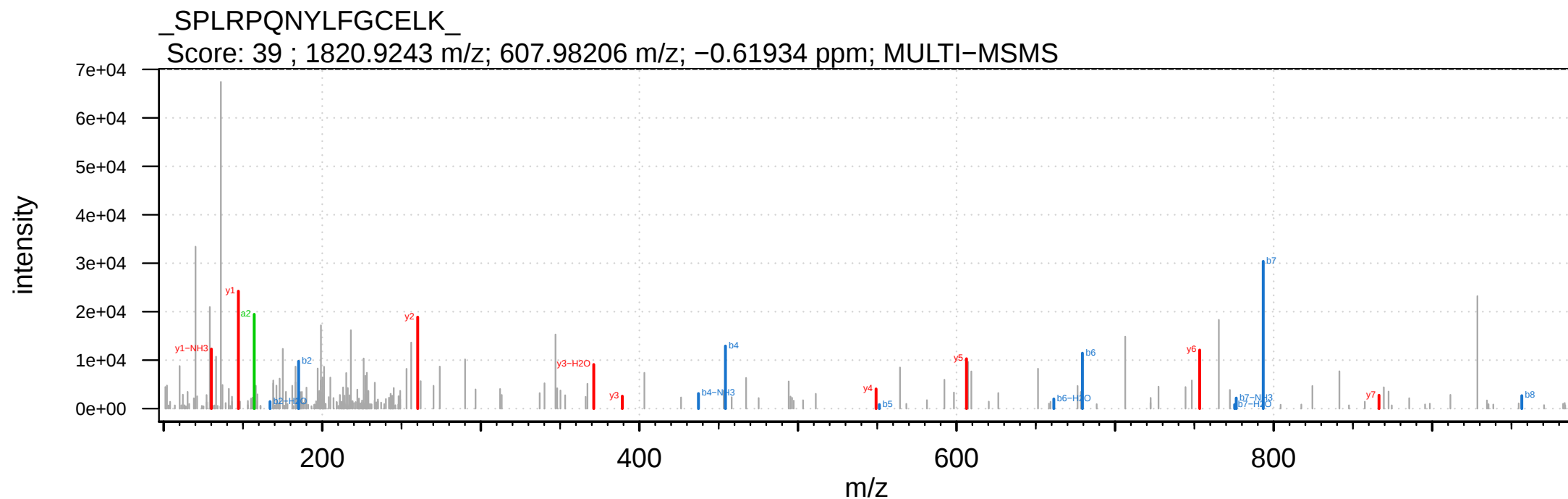
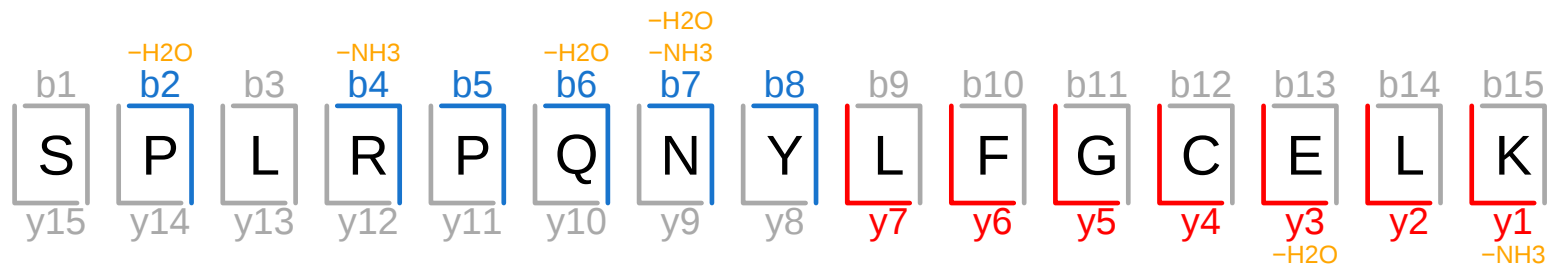


Raw File: 20100723_Velos1_TaGe_SA_Gamg_2

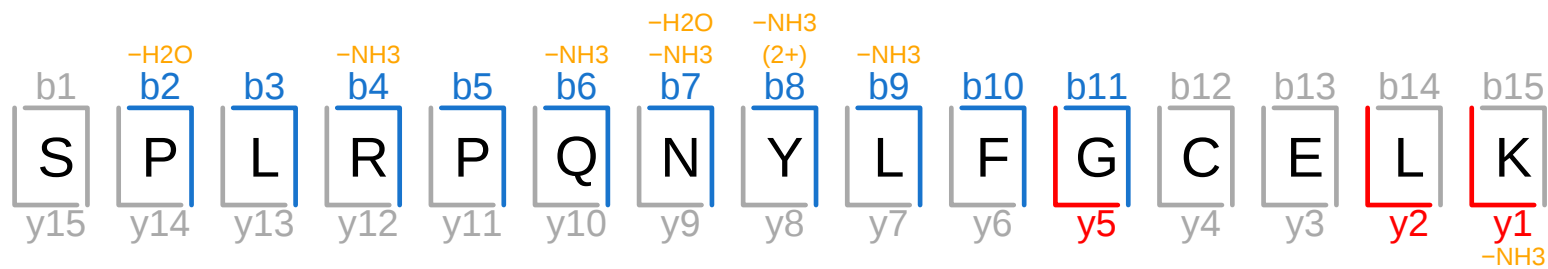
Scan Number: 20646

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

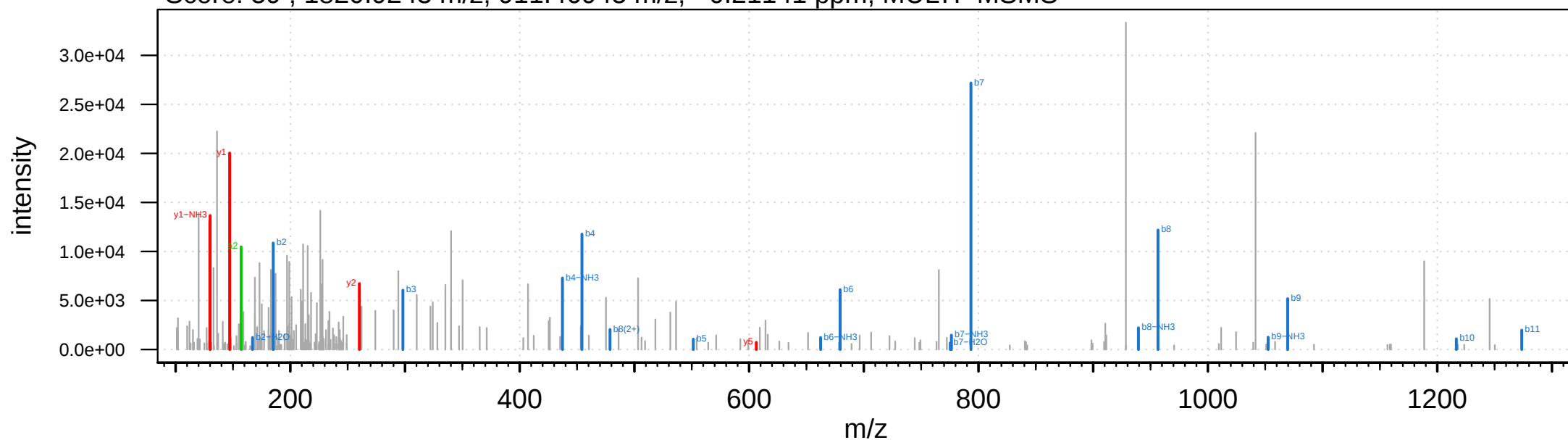


Raw File: 20100726_Velos1_TaGe_SA_HeLa_1
 Scan Number: 17509
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 59 ; 1820.9243 m/z; 911.46945 m/z; -0.21141 ppm; MULTI-MSMS

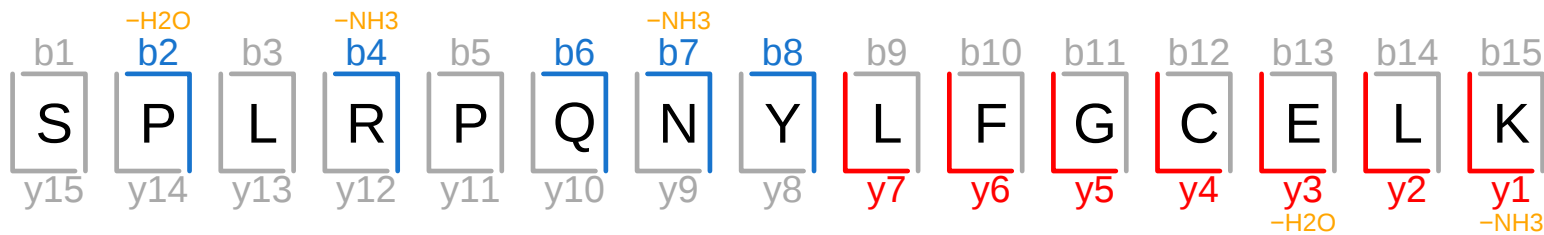


Raw File: 20100726_Velos1_TaGe_SA_HeLa_1

Scan Number: 17529

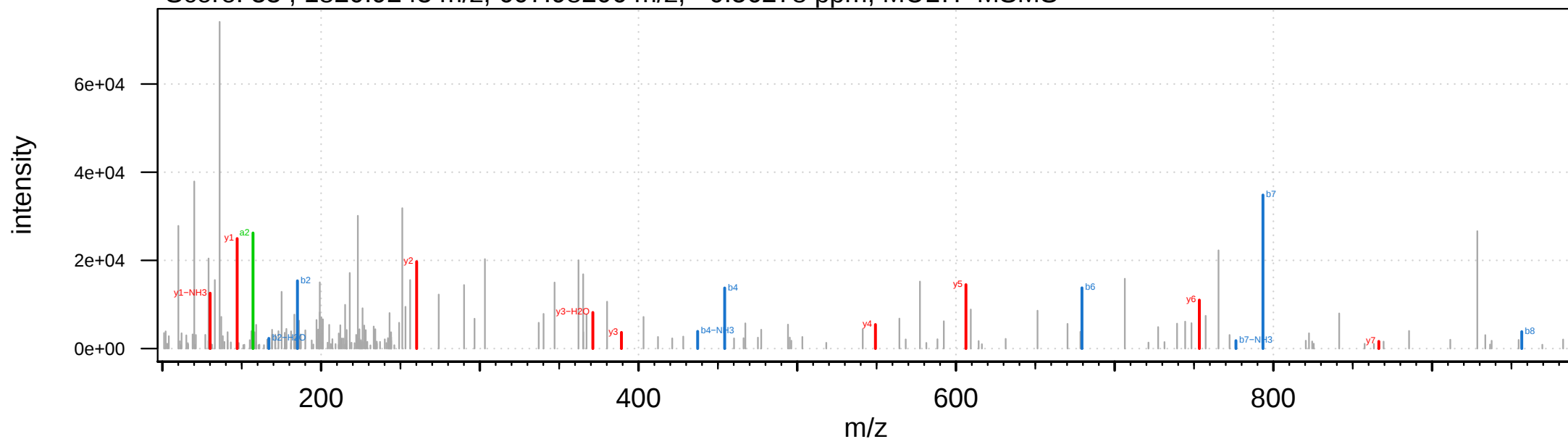
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 33 ; 1820.9243 m/z; 607.98206 m/z; -0.56278 ppm; MULTI-MSMS

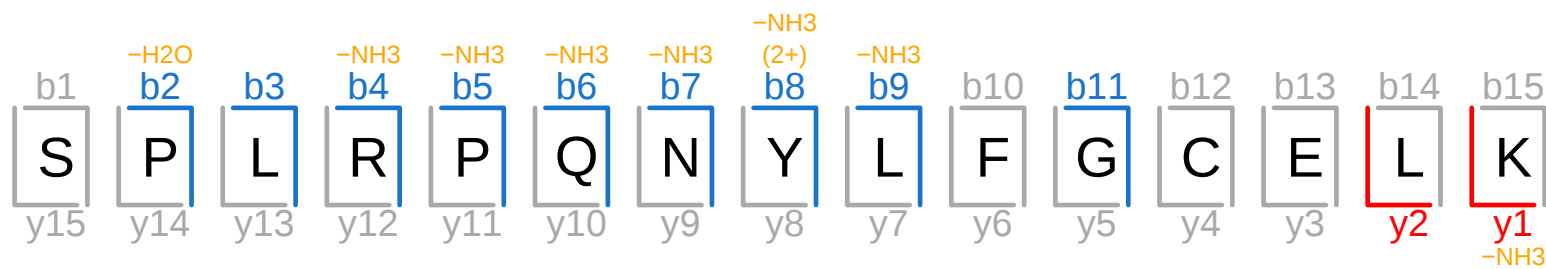


Raw File: 20100726_Velos1_TaGe_SA_HepG2_1

Scan Number: 17794

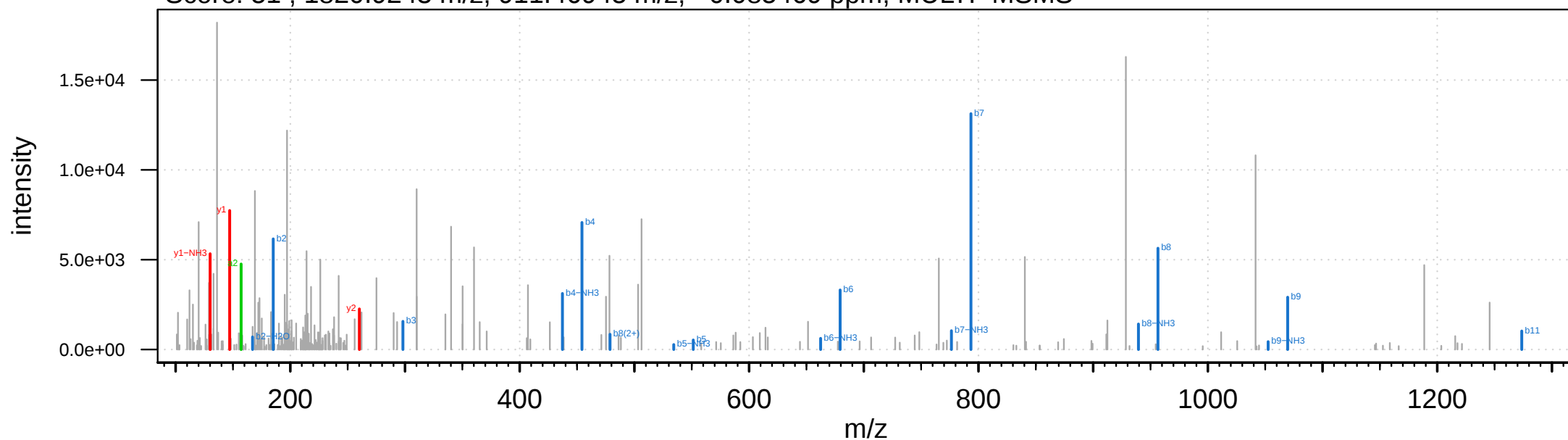
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 51 ; 1820.9243 m/z; 911.46945 m/z; -0.085469 ppm; MULTI-MSMS

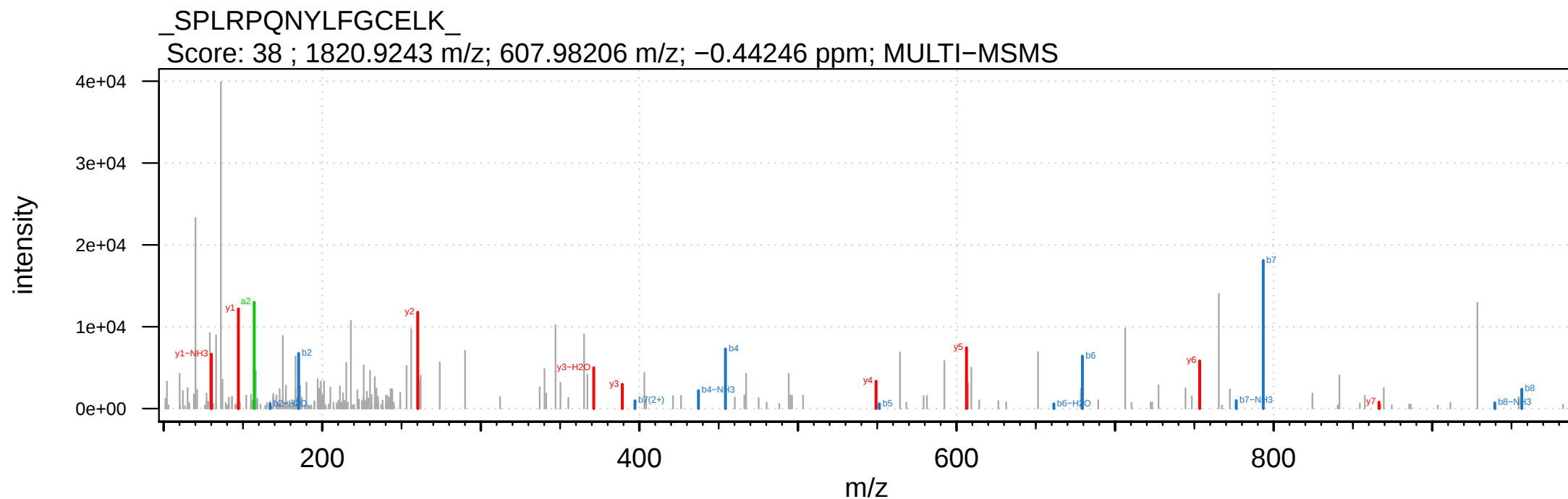
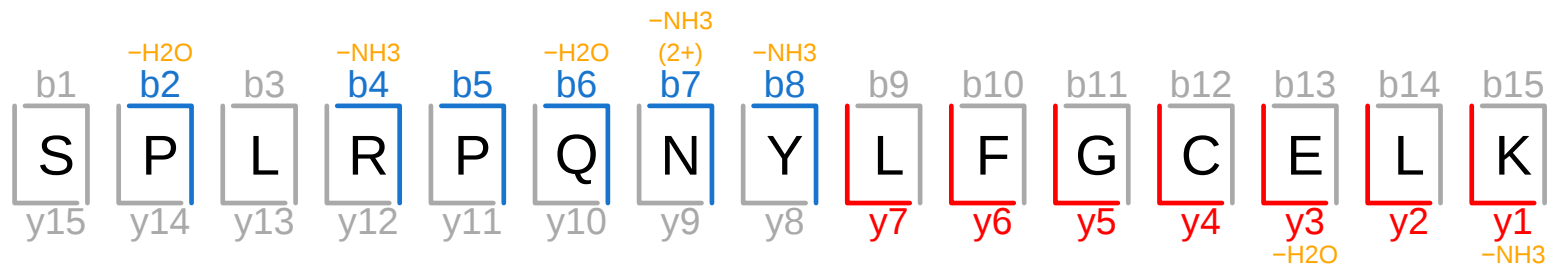


Raw File: 20100726_Velos1_TaGe_SA_HepG2_1

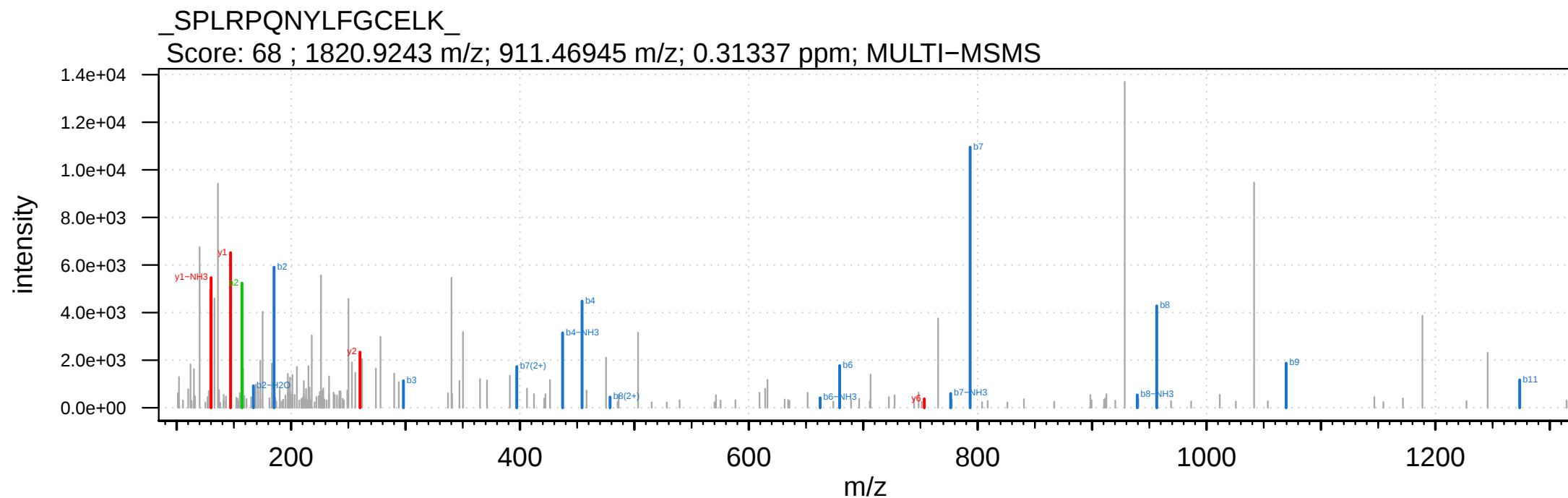
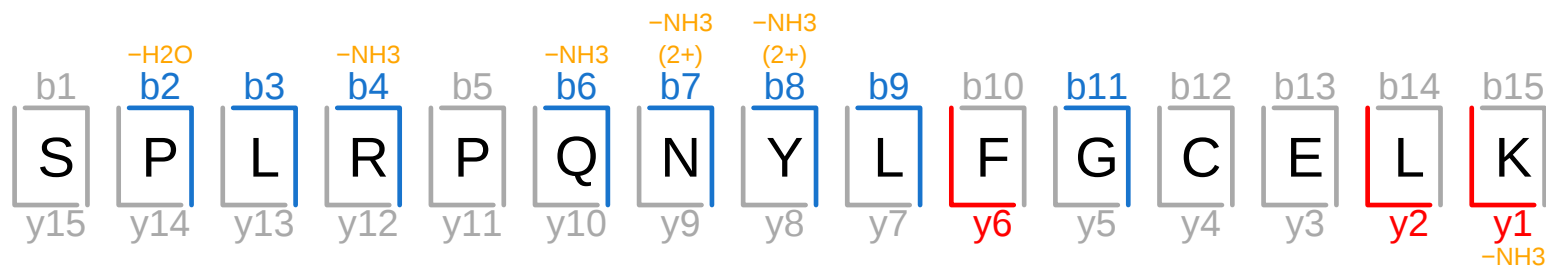
Scan Number: 17800

Proteins:

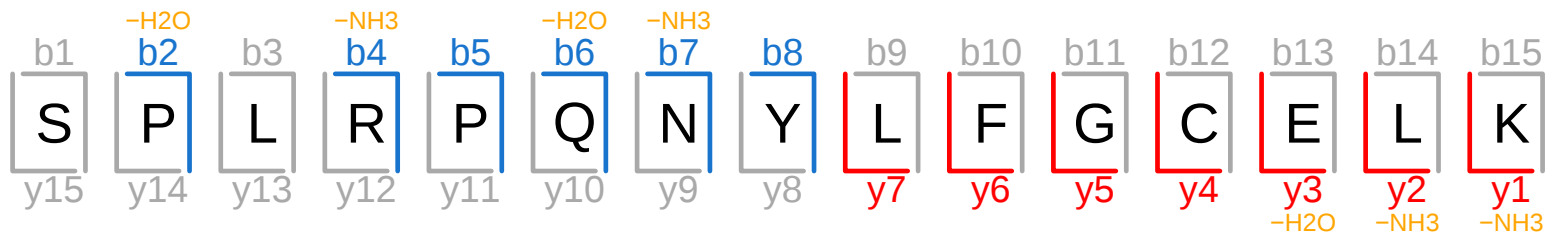
TCONS_I2_00008829_chr15:92829088-92829258:+



Raw File: 20100801_Velos1_TaGe_SA_RKO_01
 Scan Number: 15215
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

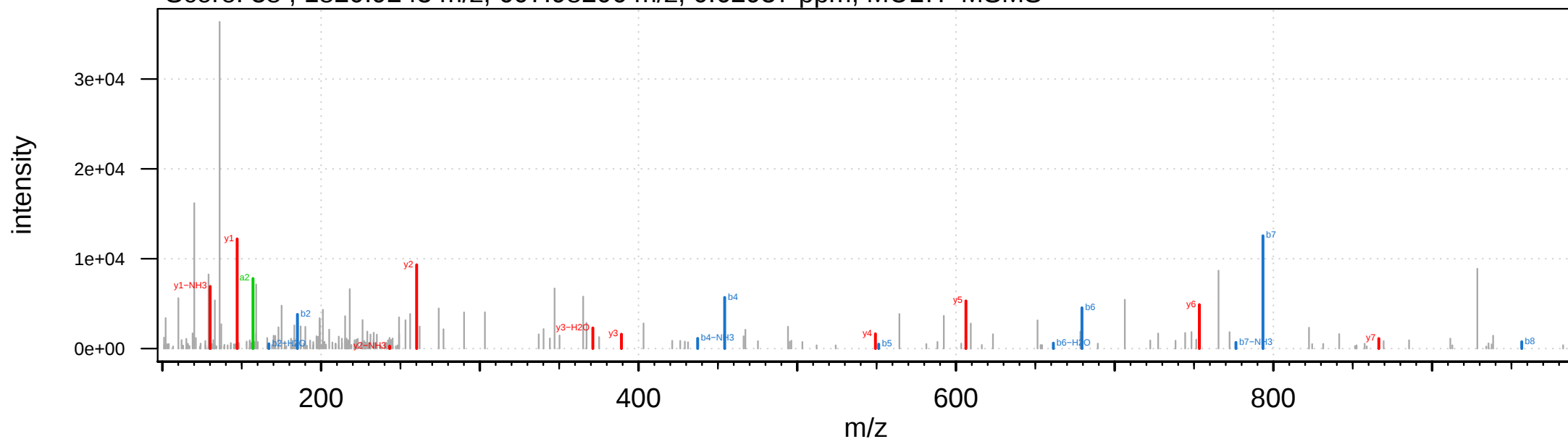


Raw File: 20100801_Velos1_TaGe_SA_RKO_01
Scan Number: 15226
Proteins:
TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 38 ; 1820.9243 m/z; 607.98206 m/z; 0.62937 ppm; MULTI-MSMS

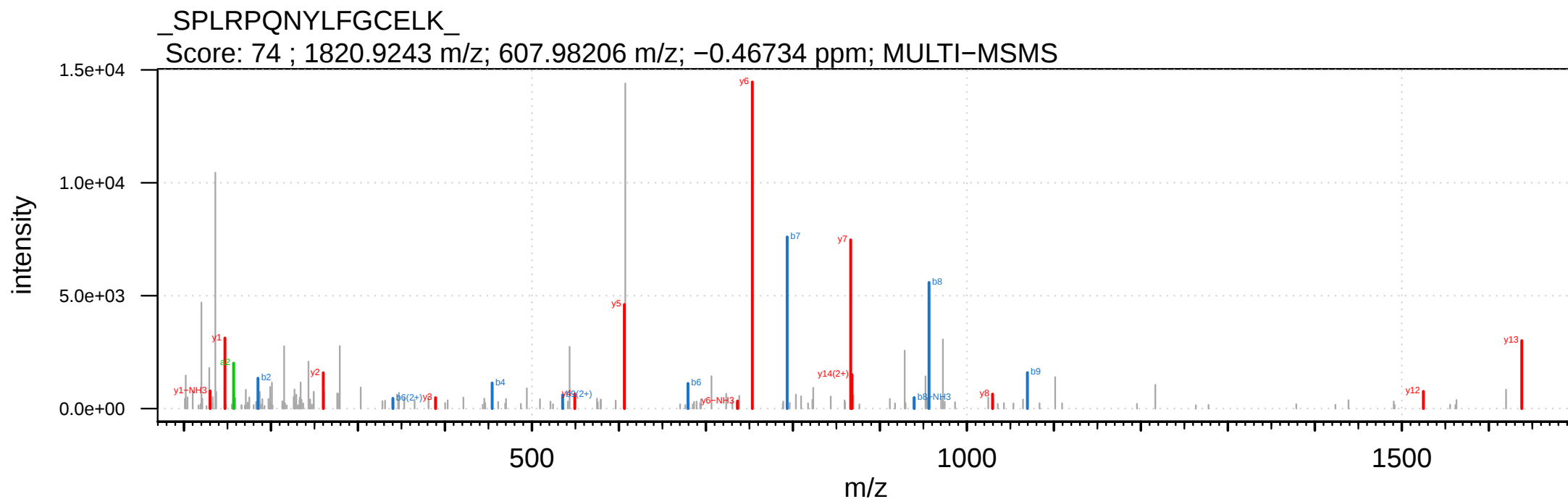
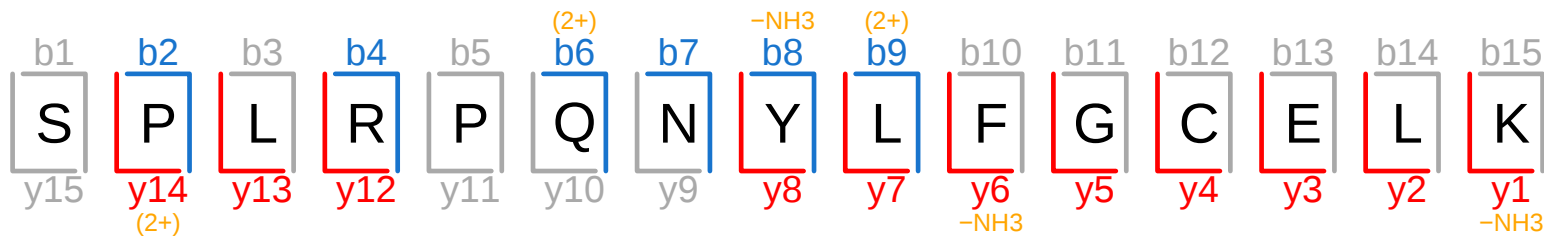


Raw File: 20100805_Velos1_TaGe_SA_RKO_06

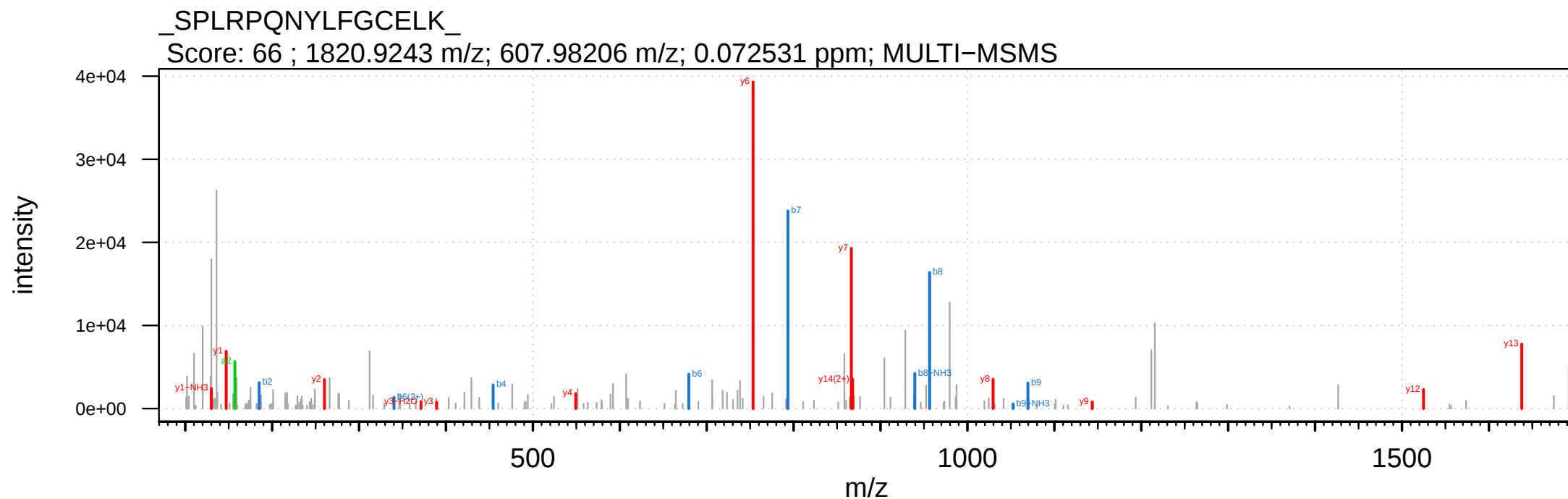
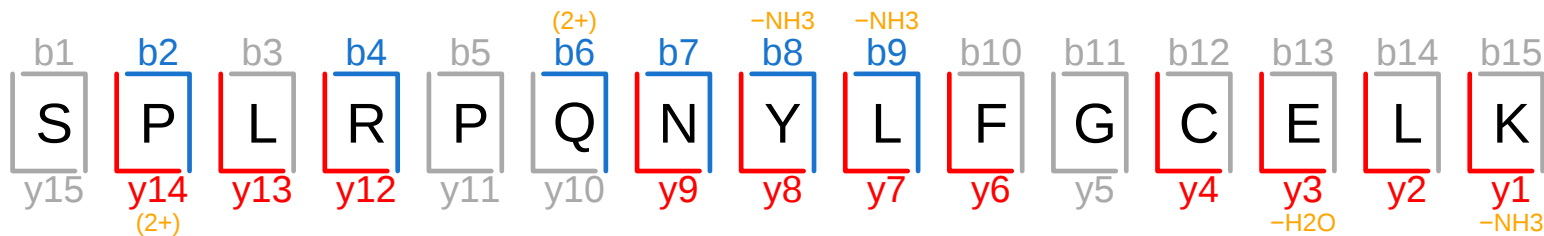
Scan Number: 15209

Proteins:

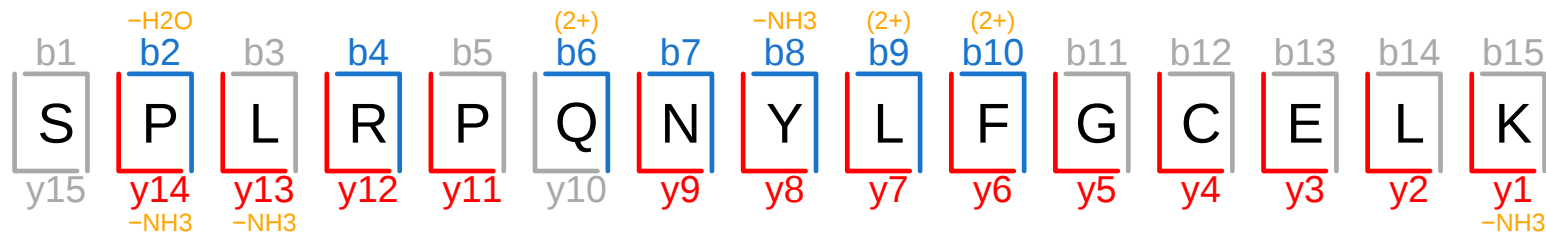
TCONS_I2_00008829_chr15:92829088-92829258:+



Raw File: 20101210_Velos1_AnWe_SA_LnCap_6
 Scan Number: 15445
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

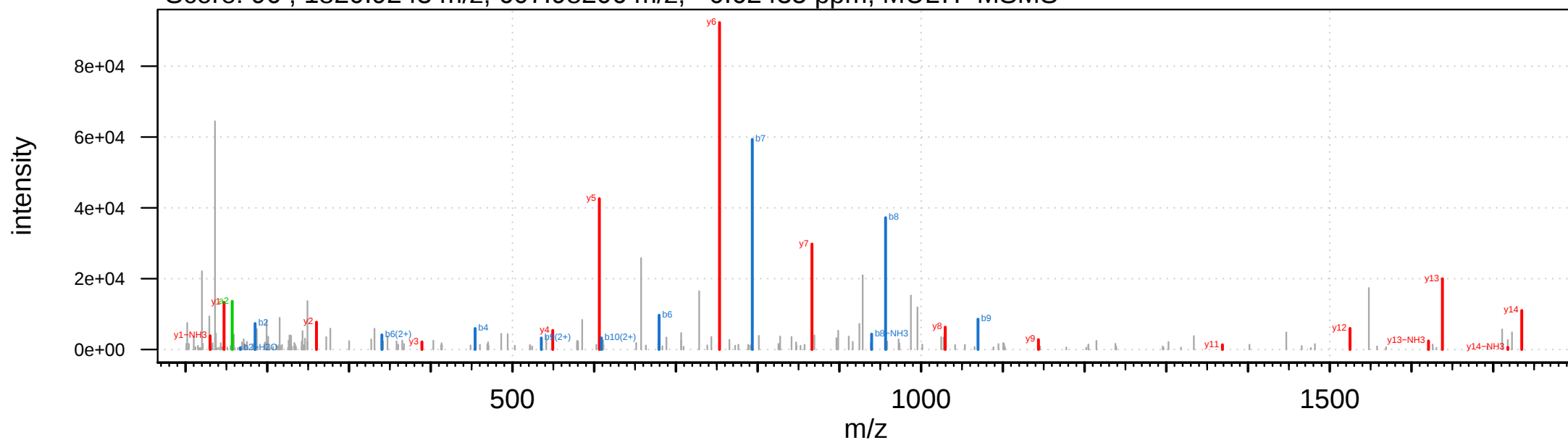


Raw File: 20101210_Velos1_AnWe_SA_MCF7_6
Scan Number: 16375
Proteins:
TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 96 ; 1820.9243 m/z; 607.98206 m/z; -0.62435 ppm; MULTI-MSMS

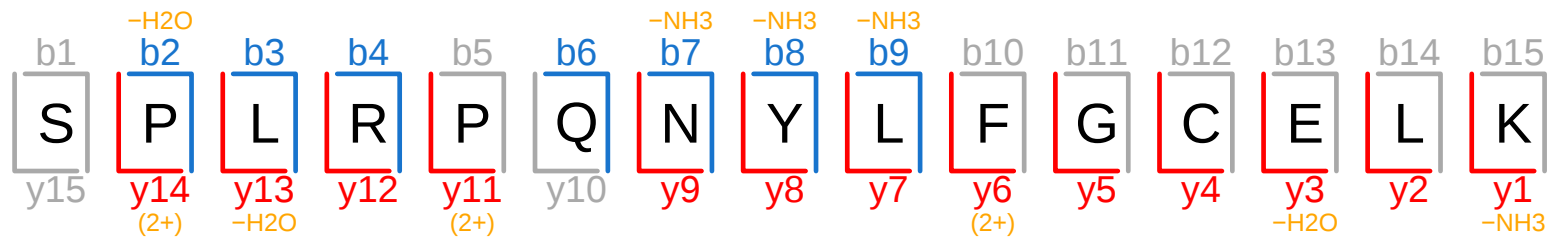


Raw File: 20101210_Velos1_AnWe_SA_U2OS_6

Scan Number: 17394

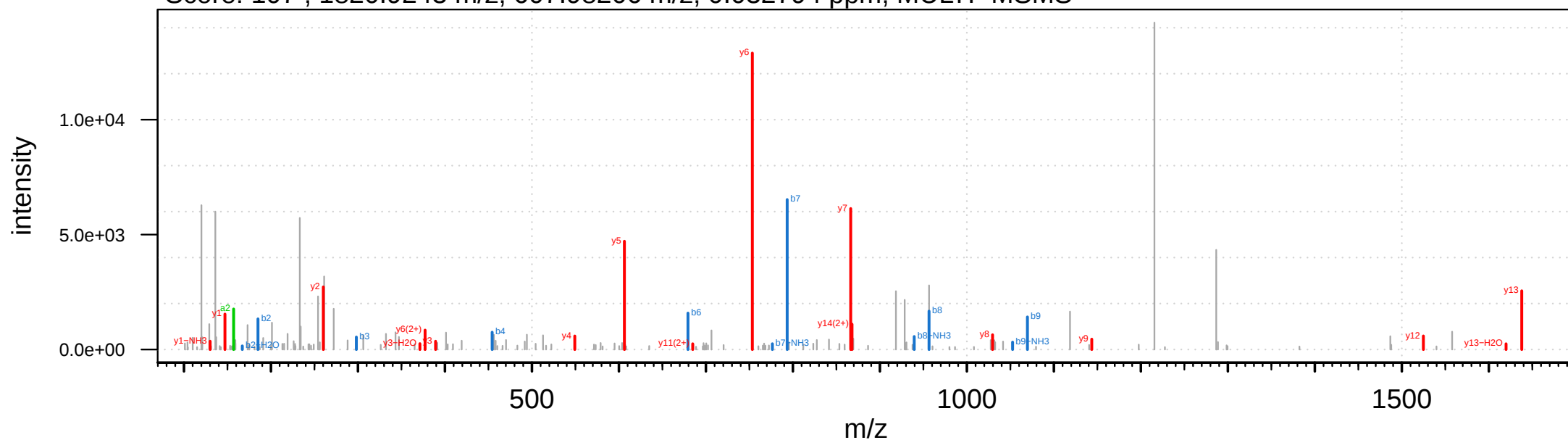
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 107 ; 1820.9243 m/z; 607.98206 m/z; 0.032794 ppm; MULTI-MSMS

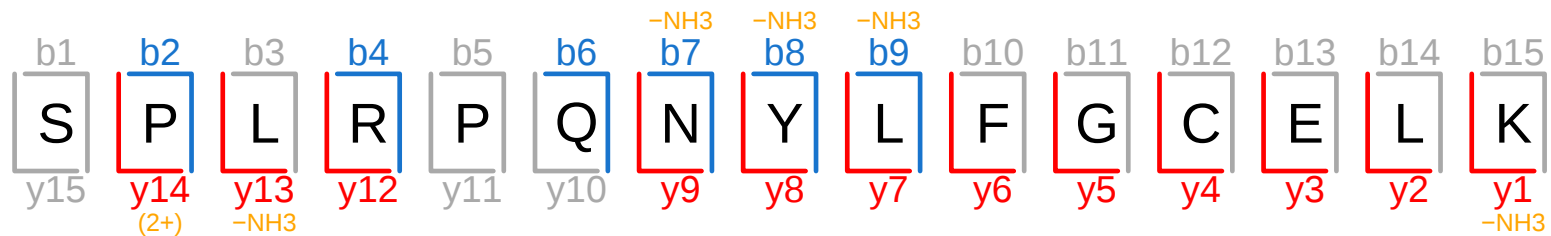


Raw File: 20101222_Velos1_TaGe_SA_K562_03

Scan Number: 12933

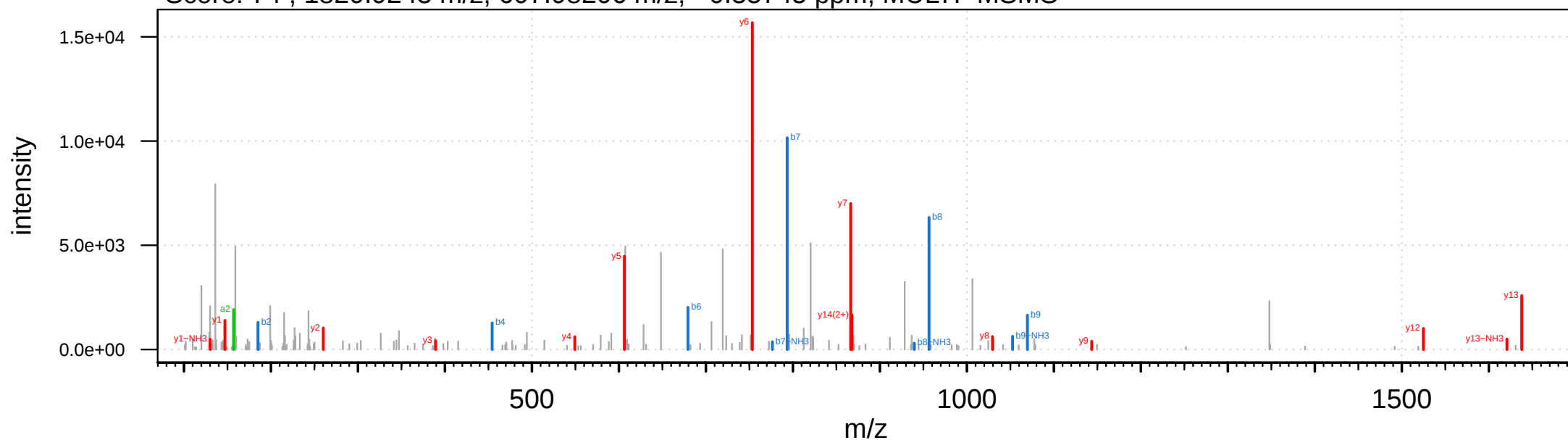
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 74 ; 1820.9243 m/z; 607.98206 m/z; -0.35745 ppm; MULTI-MSMS

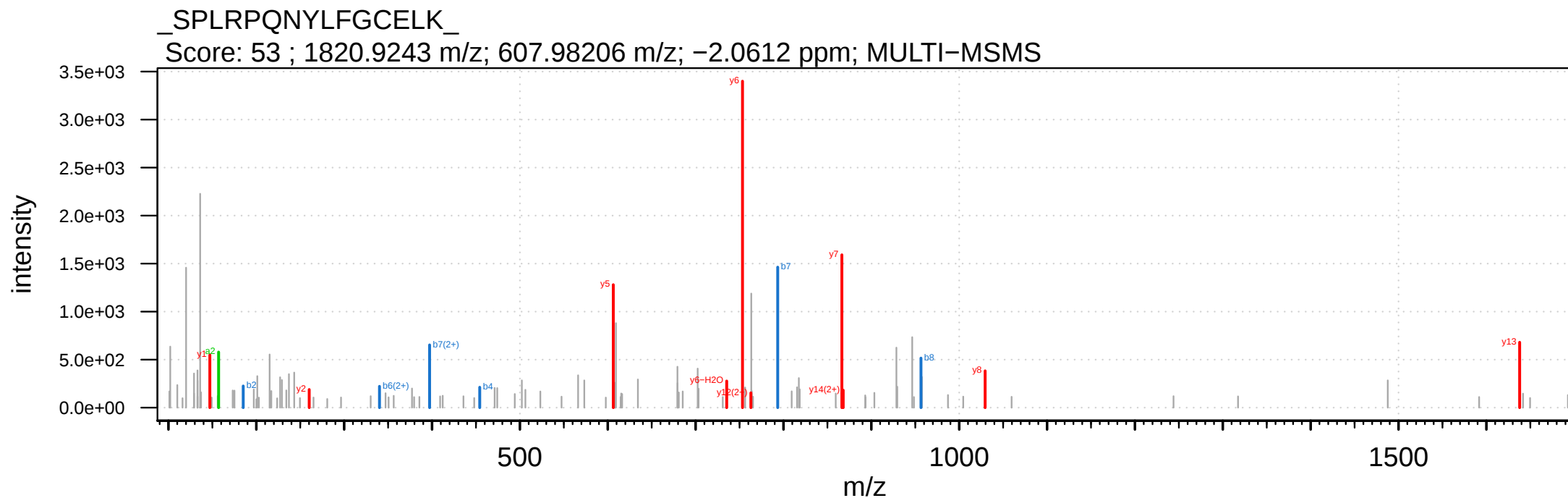
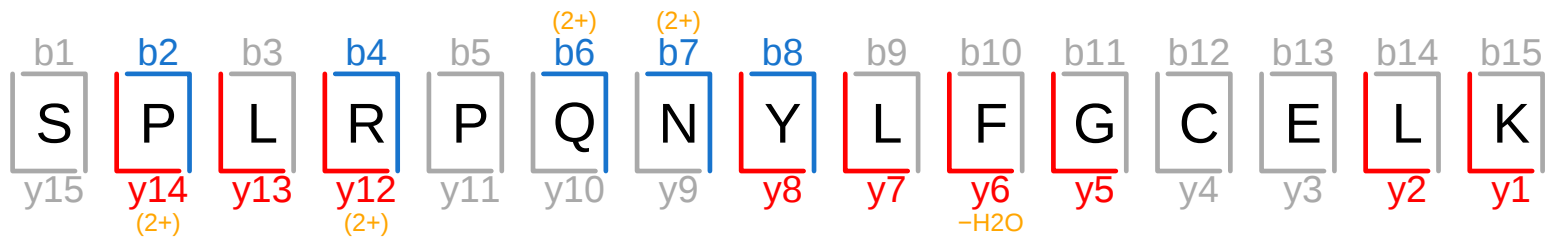


Raw File: 20101222_Velos1_TaGe_SA_K562_05

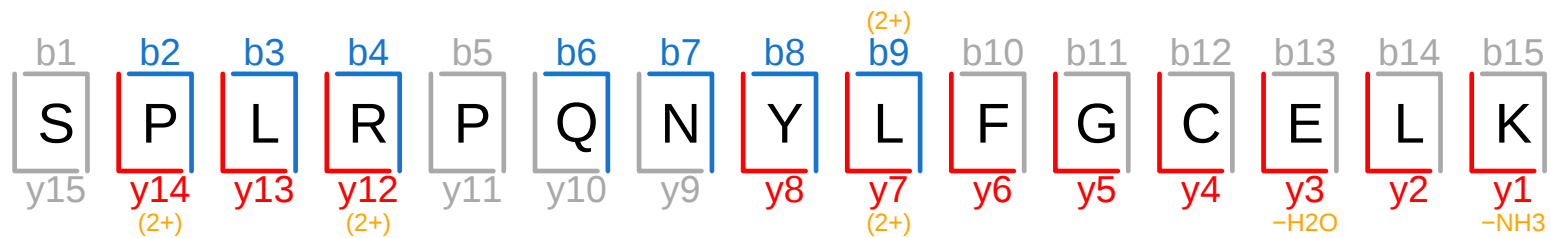
Scan Number: 12446

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

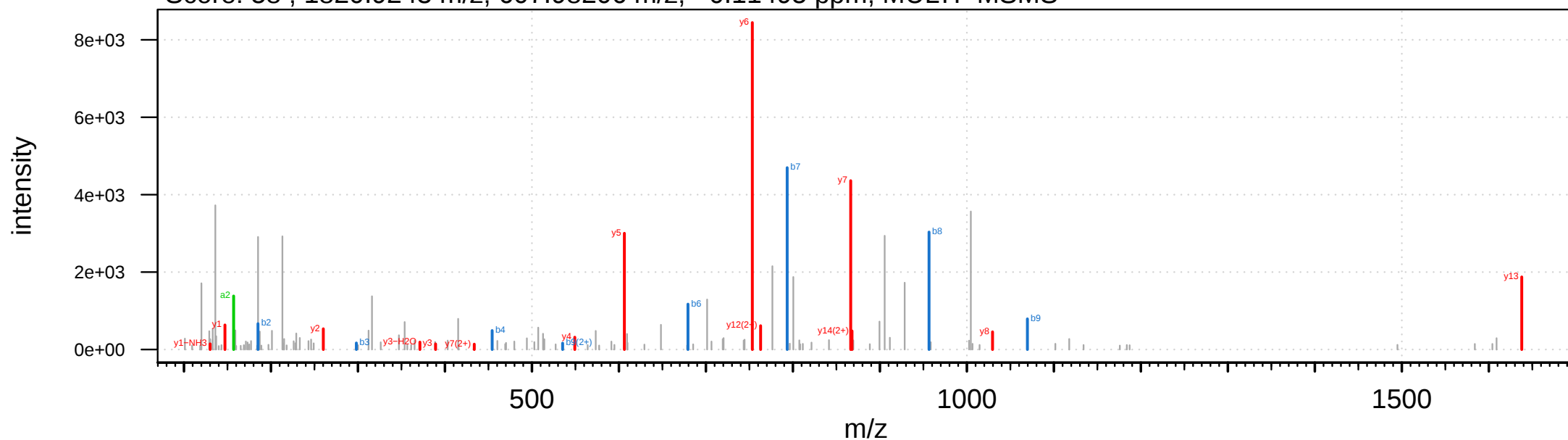


Raw File: 20101222_Velos1_TaGe_SA_K562_06
 Scan Number: 11938
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 58 ; 1820.9243 m/z; 607.98206 m/z; -0.11495 ppm; MULTI-MSMS

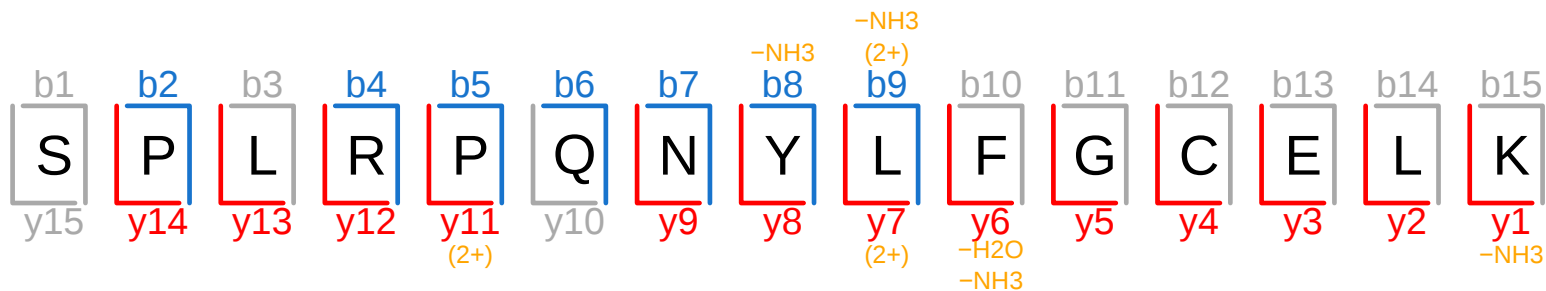


Raw File: 20101223_Velos1_TaGe_SA_RKO_01

Scan Number: 14825

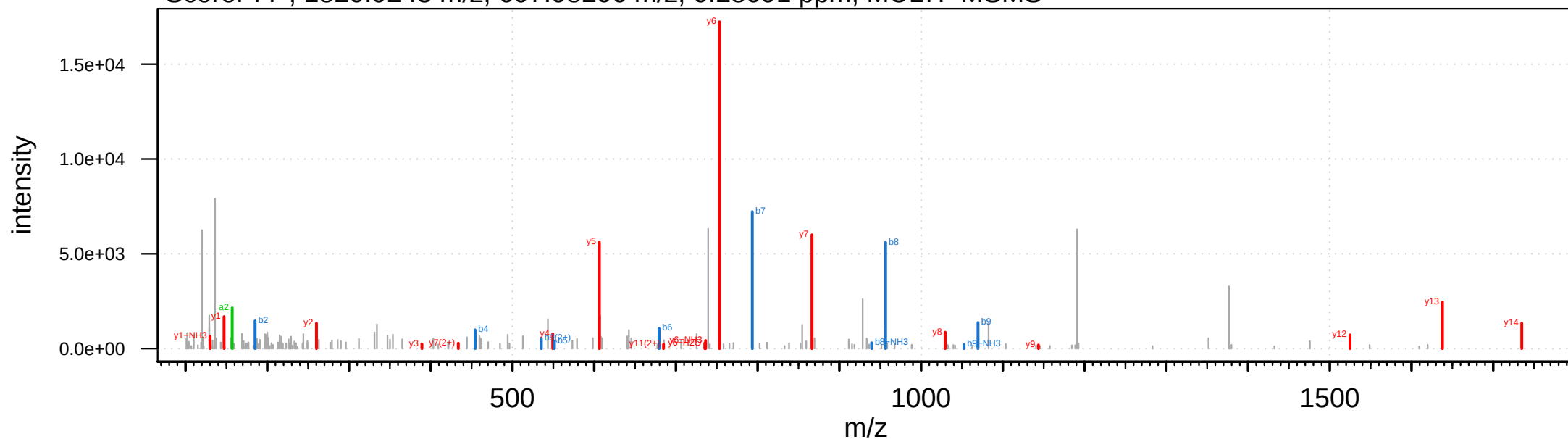
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

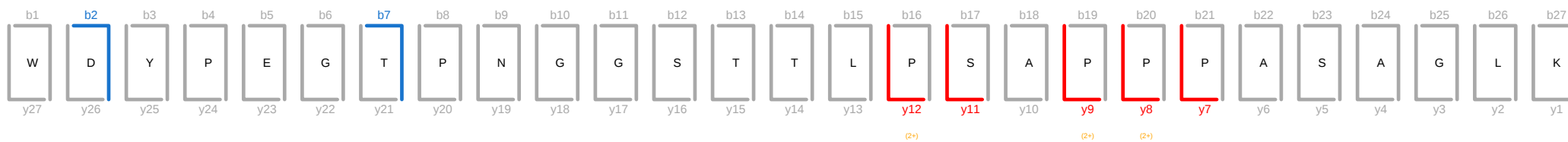


SPLRPQNYLFGCELK

Score: 77 ; 1820.9243 m/z; 607.98206 m/z; 0.28091 ppm; MULTI-MSMS

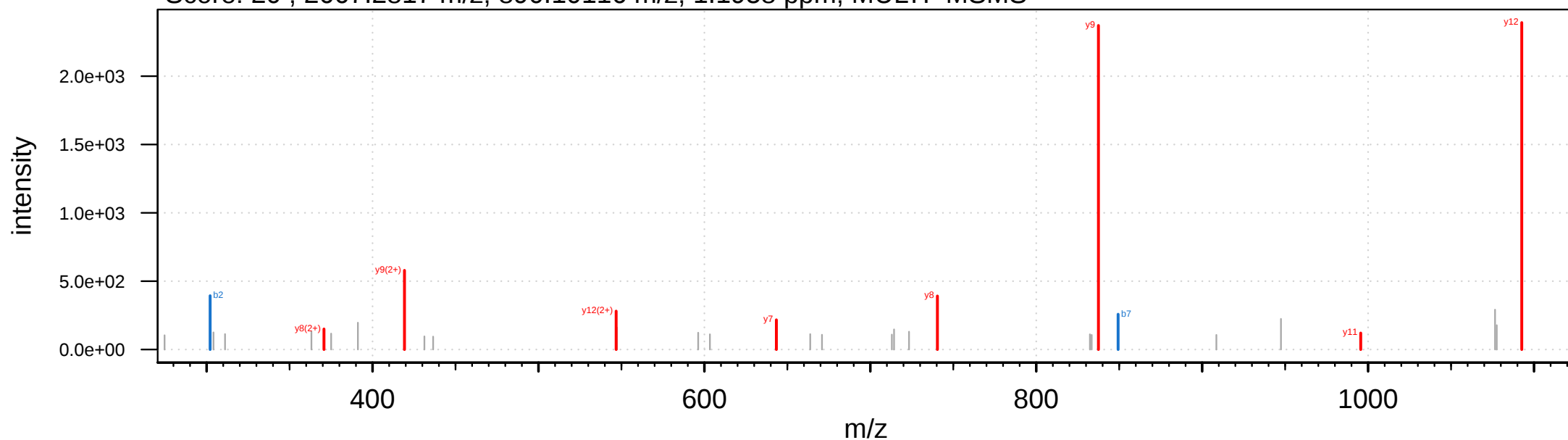


Raw File: 20101224_Velos1_TaGe_SA_HeLa_04
 Scan Number: 19194
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



WDYPEGTPNGGSTTLPSAPPPASAGLK

Score: 20 ; 2667.2817 m/z; 890.10116 m/z; 1.1958 ppm; MULTI-MSMS

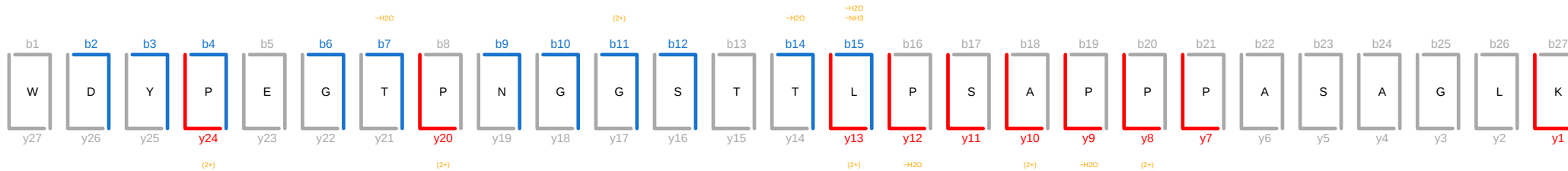


Raw File: 20101215_Velos1_TaGe_SA_A549_04

Scan Number: 19320

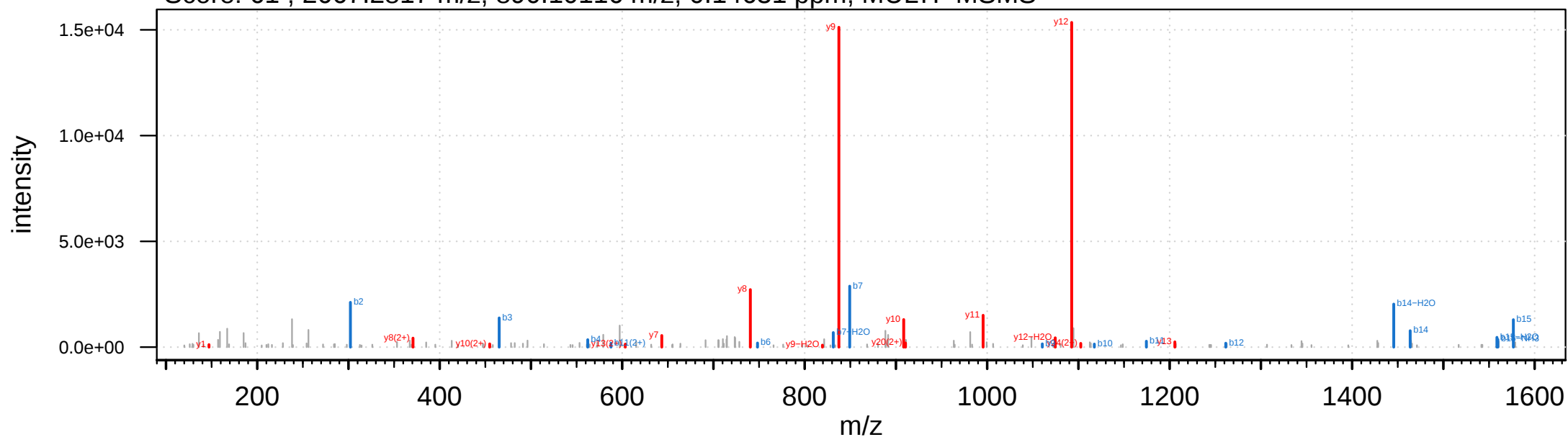
Proteins:

ENST00000374922_chrX:56755787-56755993:+

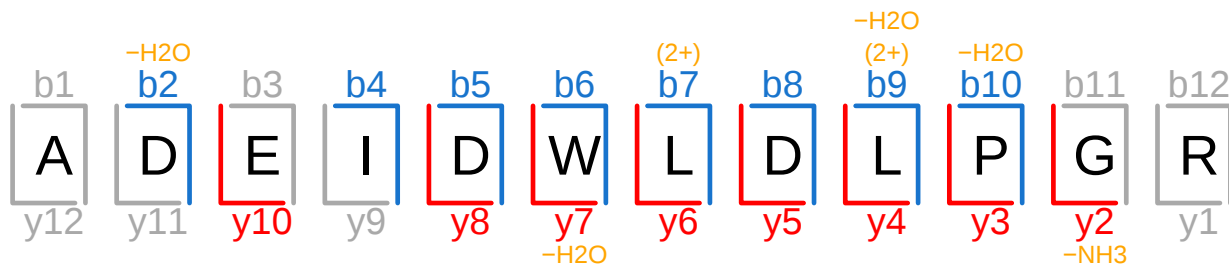


WDYPEGTPNGGSTTLPSAPPPASAGLK

Score: 61 ; 2667.2817 m/z; 890.10116 m/z; 0.14631 ppm; MULTI-MSMS

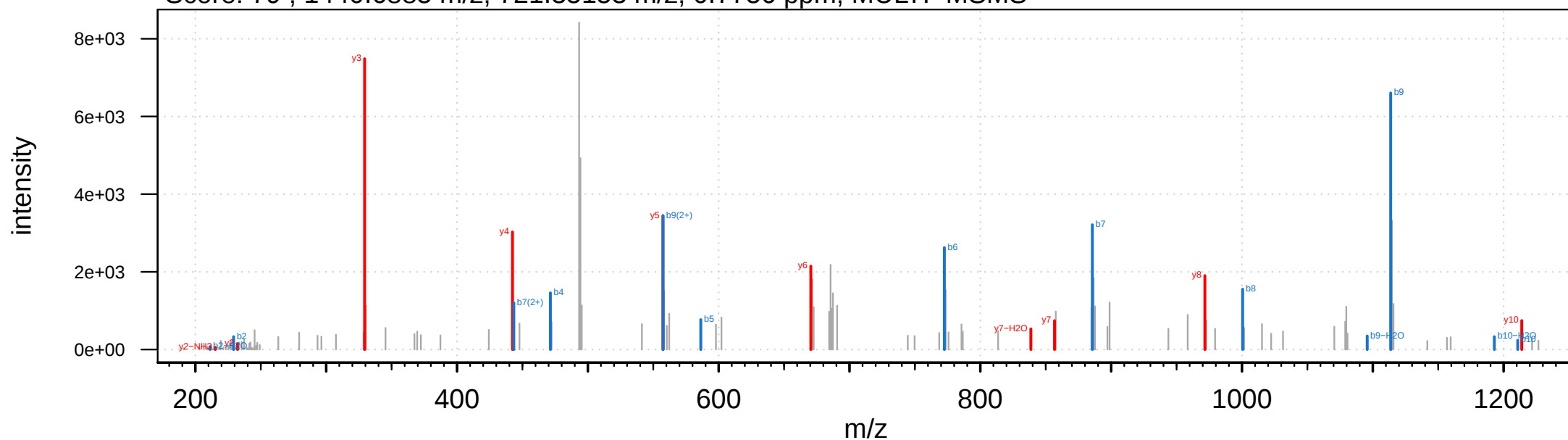


Raw File: 20101230_Velos1_TaGe_SA_Jurkat3
 Scan Number: 22281
 Proteins:
 ENST00000374922_chrX:56755787-56755993:+



(ac)ADEIDWLDLPGR

Score: 79 ; 1440.6885 m/z; 721.35153 m/z; 0.7756 ppm; MULTI-MSMS

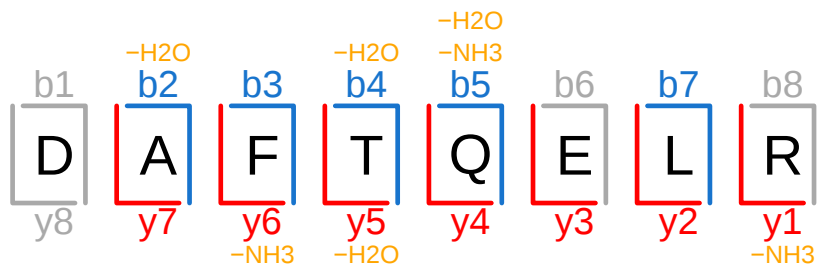


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F3_01

Scan Number: 31694

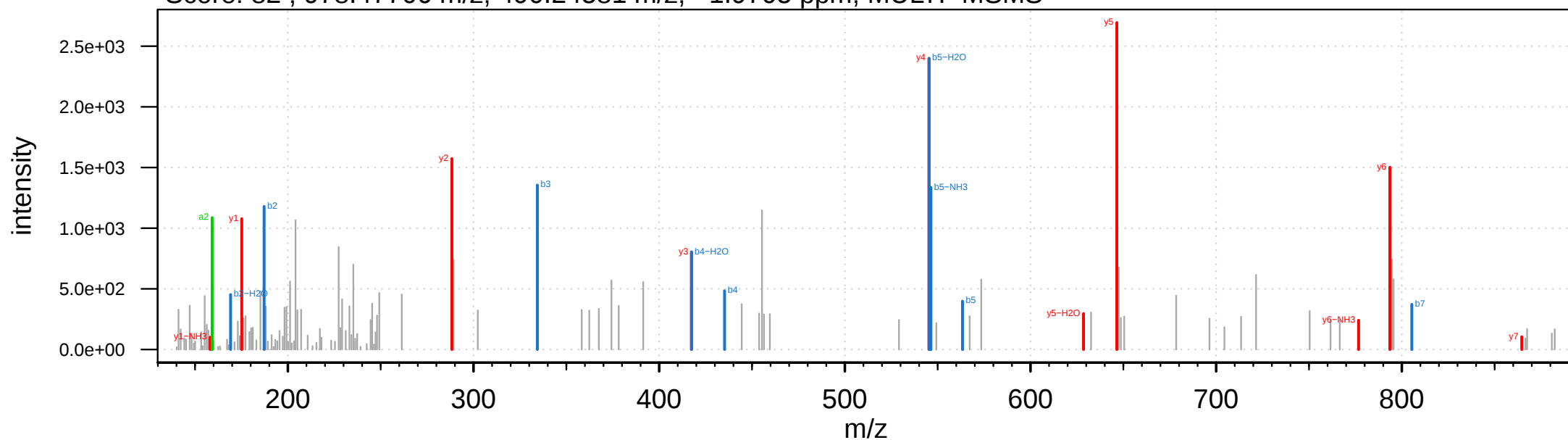
Proteins:

TCONS_I2_00001718_chr1:204336923-204346793:-



DAFTQELR

Score: 82 ; 978.47706 m/z; 490.24581 m/z; -1.9705 ppm; MULTI-MSMS



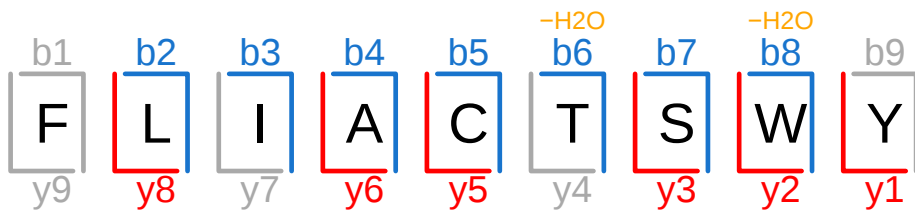
Raw File: Toni_20111125_FB_MCF7_F4_2

Scan Number: 30437

Proteins:

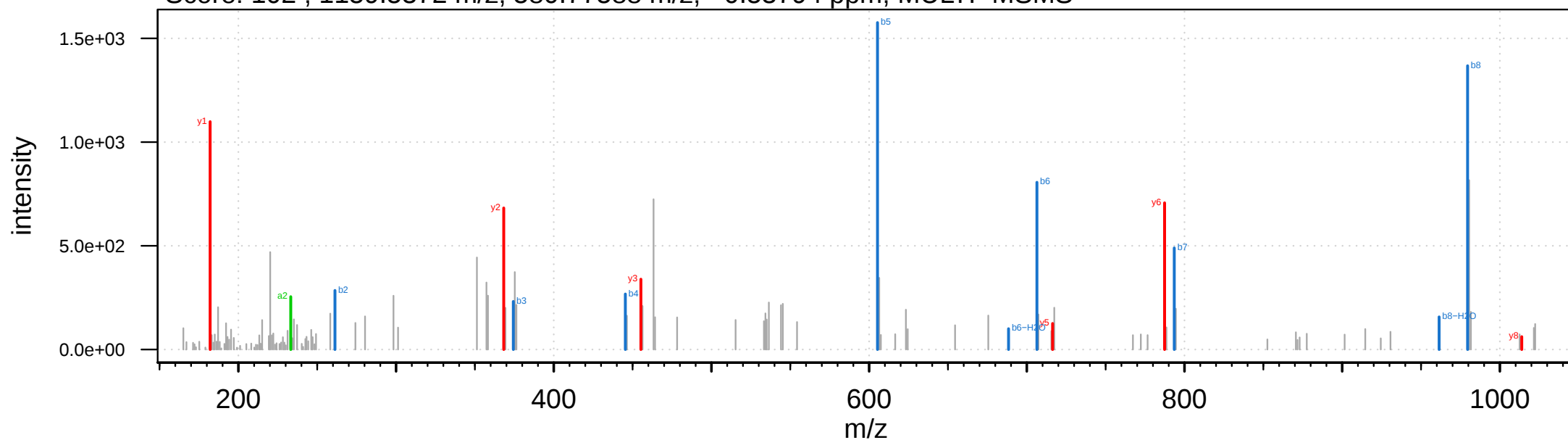
Q69YL0

ENST00000602845_chr3:196669588-196669887:+



FLIACTSWY

Score: 102 ; 1159.5372 m/z; 580.77588 m/z; -0.53794 ppm; MULTI-MSMS

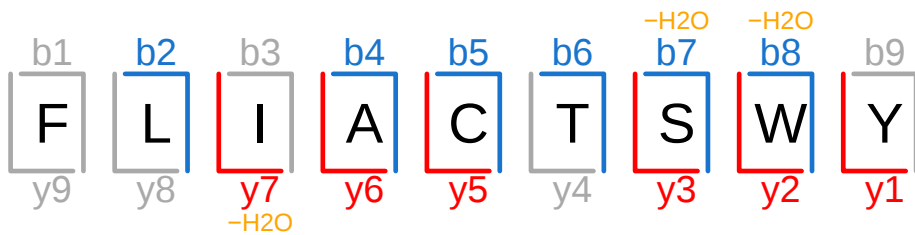


Raw File: Toni_20111114_FB_MDA-MB-231_F6_2

Scan Number: 53029

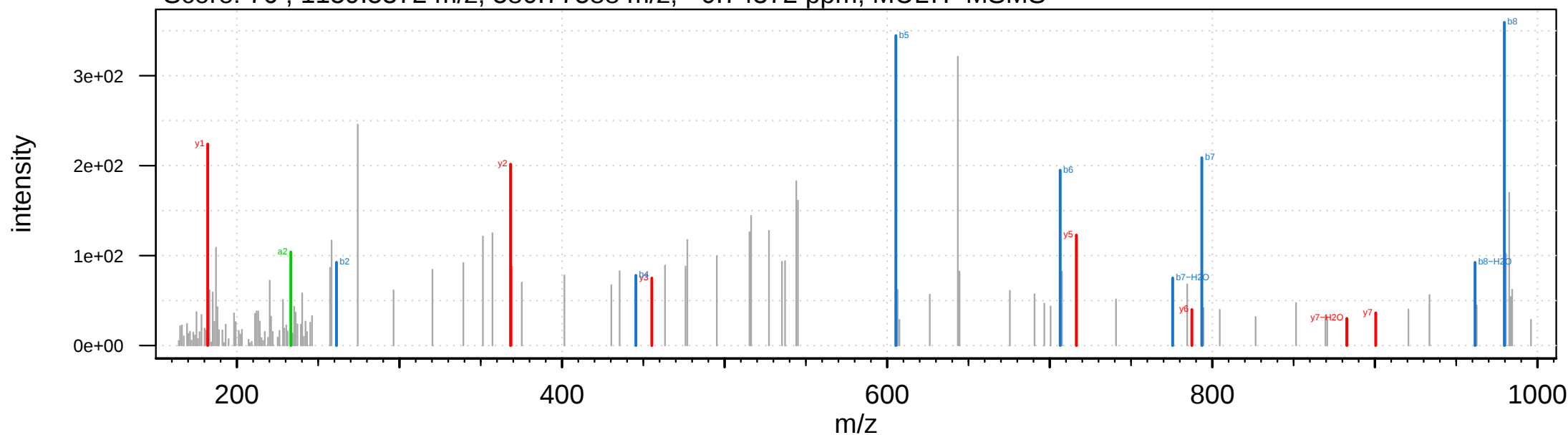
Proteins:

ENST00000424358_chr20:33865517-33865732:-



FLIACTSWY

Score: 76 ; 1159.5372 m/z; 580.77588 m/z; -0.74372 ppm; MULTI-MSMS

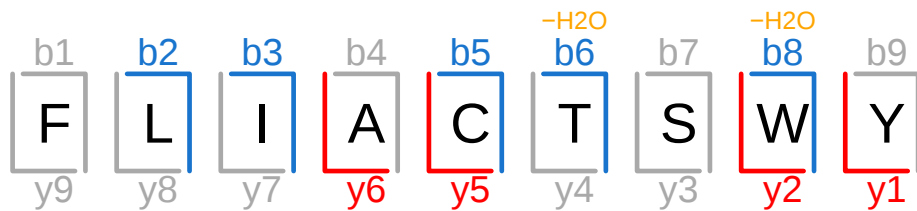


Raw File: Toni_20111114_FB_MDA-MB-231_F7_2

Scan Number: 51019

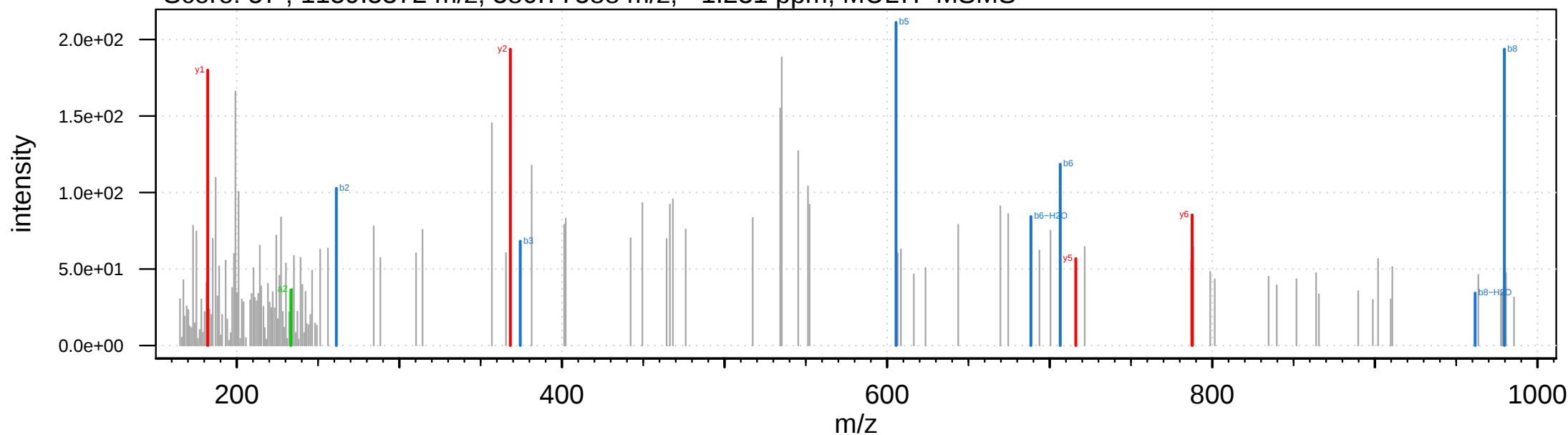
Proteins:

ENST00000424358_chr20:33865517-33865732:-



FLIACTSWY

Score: 57 ; 1159.5372 m/z; 580.77588 m/z; -1.231 ppm; MULTI-MSMS

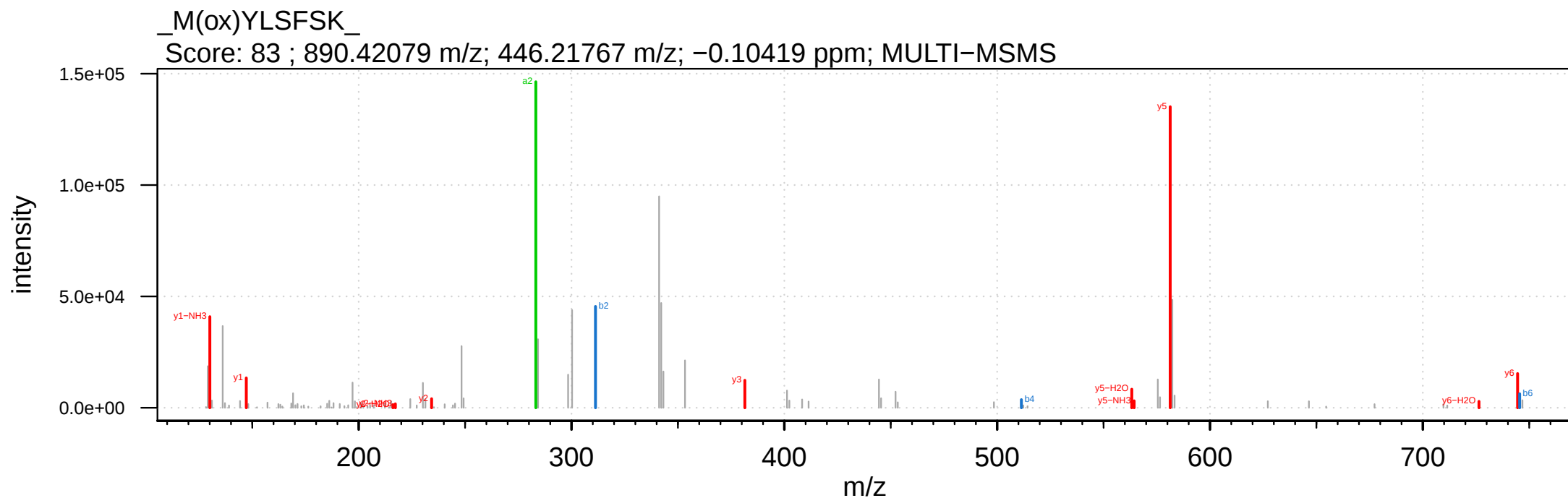
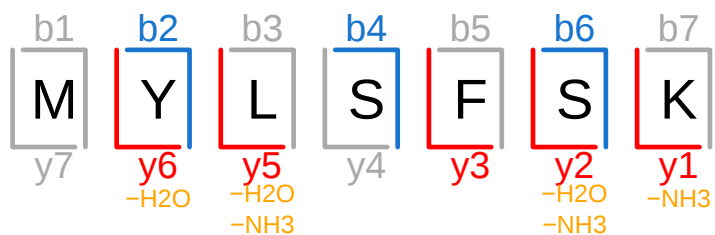


Raw File: Toni_20111125_FB_MCF7_F7_1

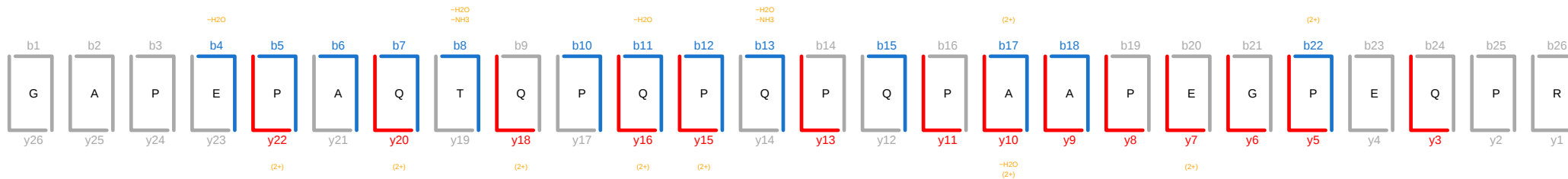
Scan Number: 56501

Proteins:

ENST00000424358_chr20:33865517-33865732:-

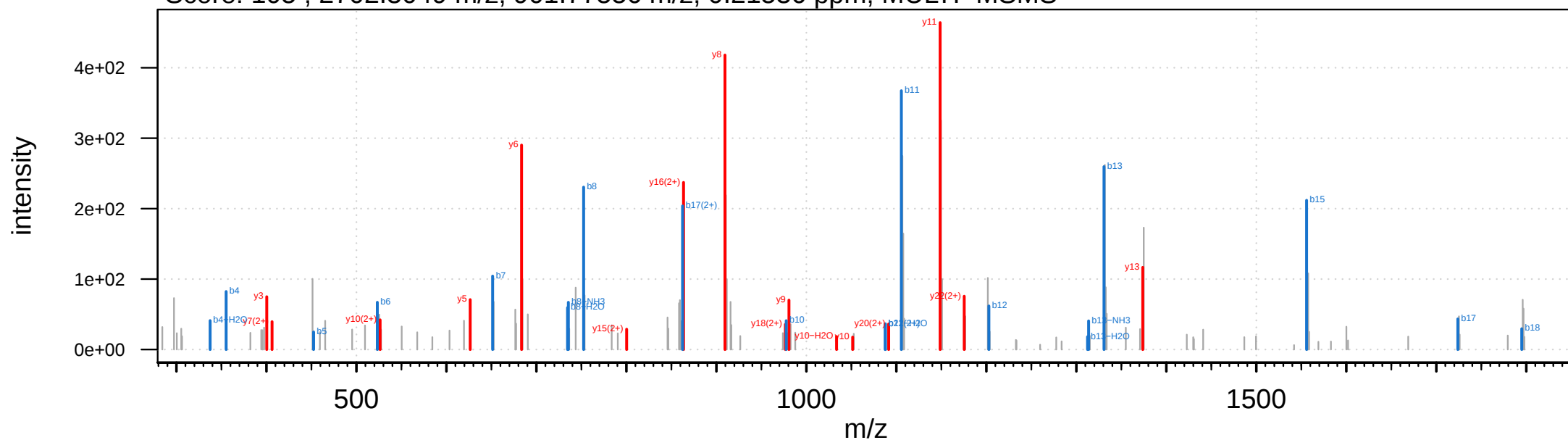


Raw File: Toni_20111125_FB_MCF7_F7_1
 Scan Number: 26760
 Proteins:
 ENST00000392145_chr6:160874572-160874622:+



GAPEPAQTQPQPQPQPAAPEGPEQPR

Score: 105 ; 2702.3049 m/z; 901.77556 m/z; 0.21536 ppm; MULTI-MSMS

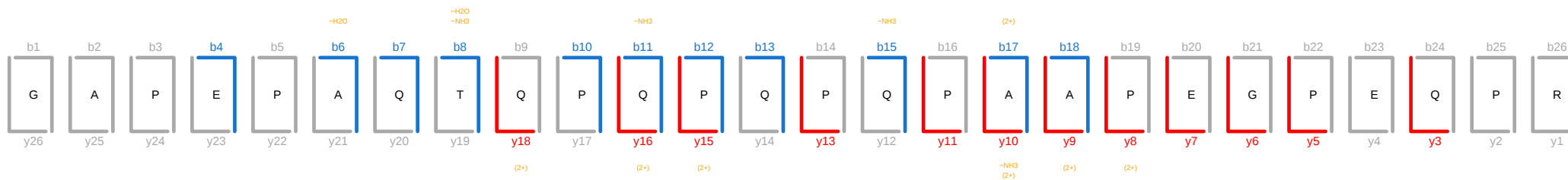


Raw File: Toni_20111114_FB_MDA-MB-231_F5_1

Scan Number: 22679

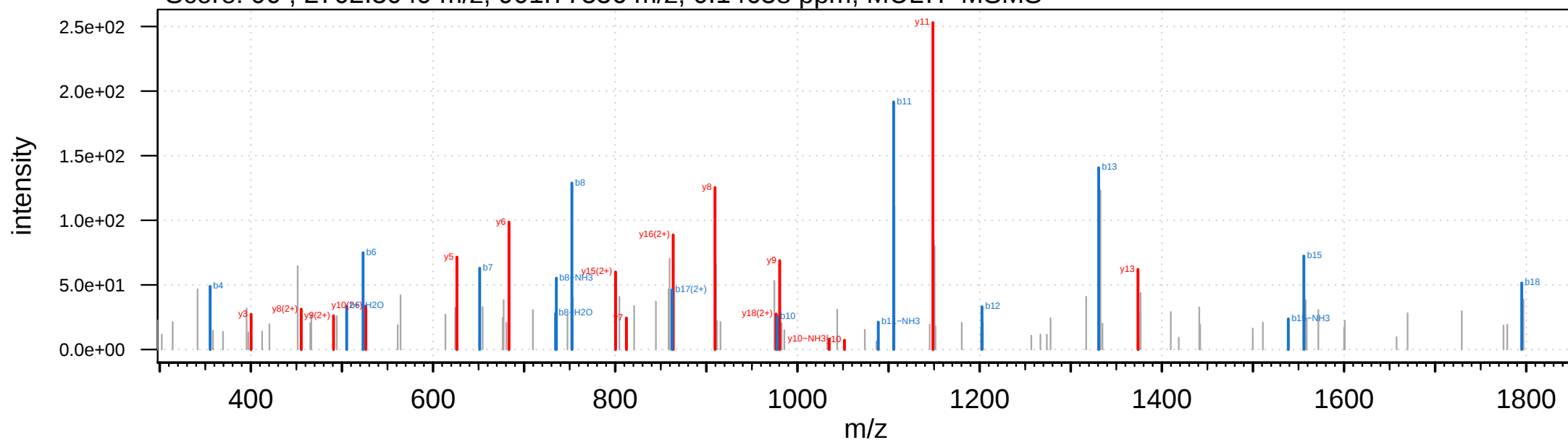
Proteins:

ENST00000424358_chr20:33865517-33865732:-



GAPEPAQTQPQPQPQPAAPEGPEQPR

Score: 90 ; 2702.3049 m/z; 901.77556 m/z; 0.14638 ppm; MULTI-MSMS

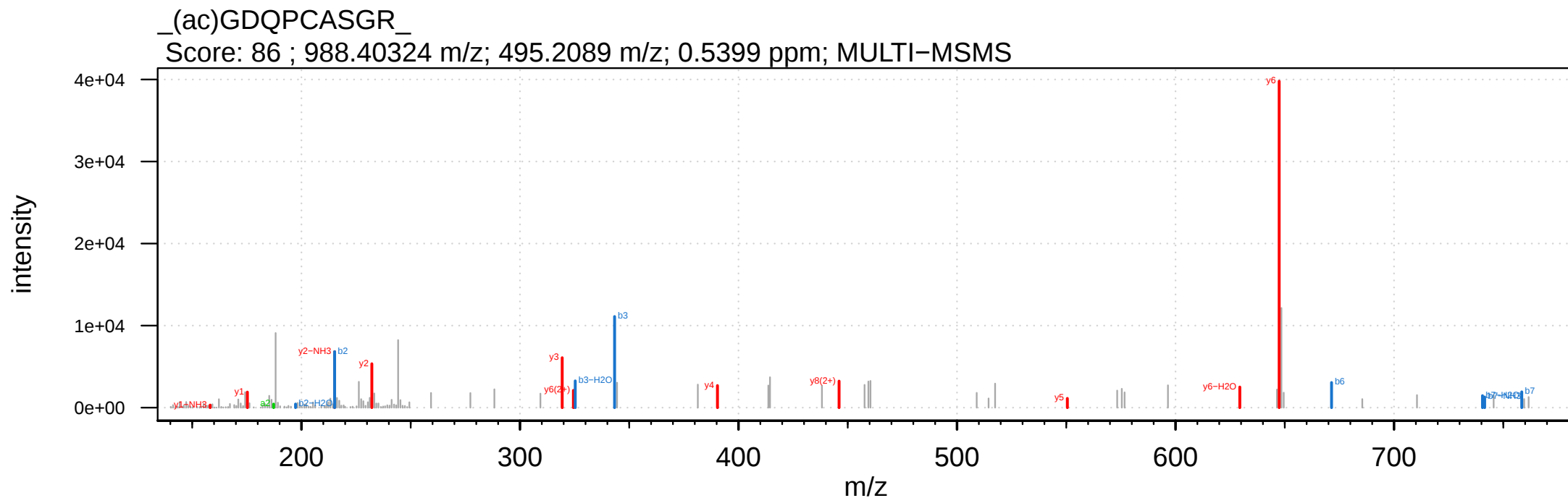
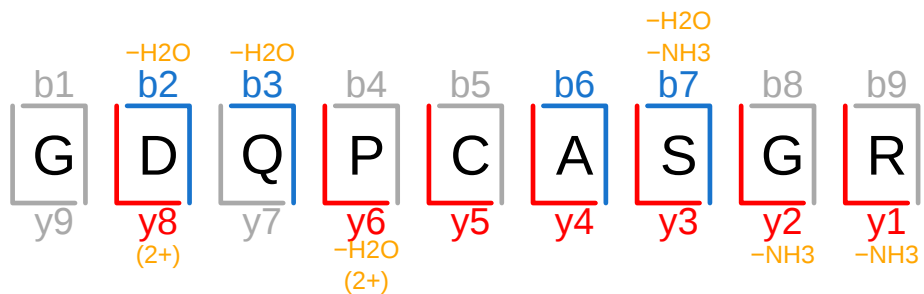


Raw File: Toni_20111114_FB_MDA-MB-231_F5_2

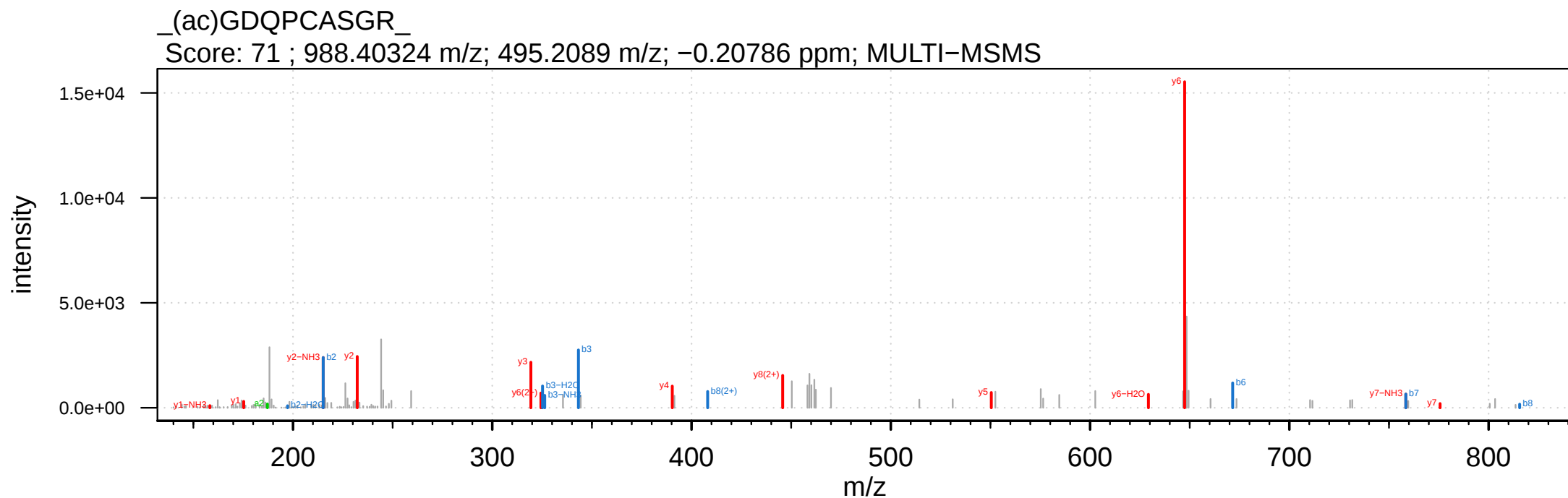
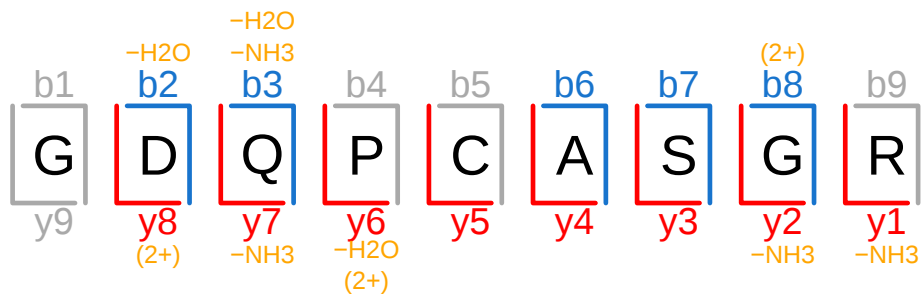
Scan Number: 22692

Proteins:

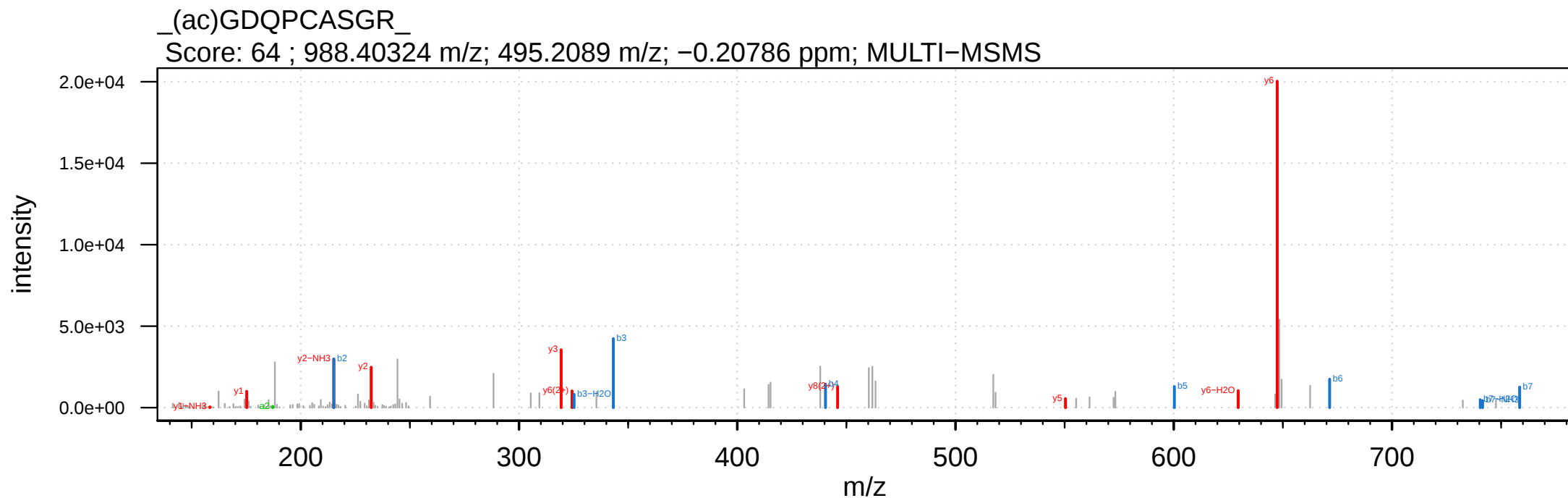
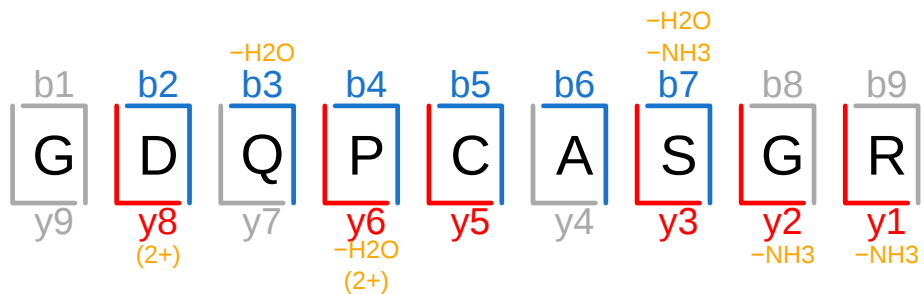
ENST00000424358_chr20:33865517-33865732:-



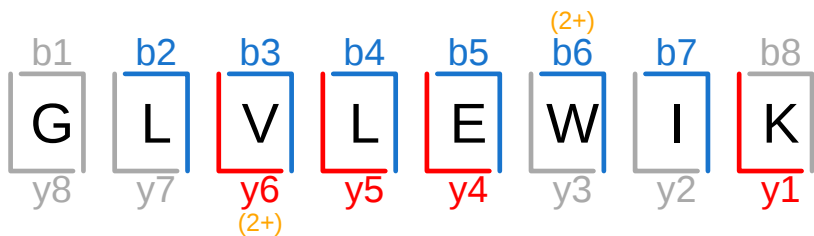
Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F4_01
 Scan Number: 6359
 Proteins:
 ENST00000374922_chrX:56755787-56755993:+



Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F4_02
Scan Number: 4047
Proteins:
ENST00000374922_chrX:56755787-56755993:+

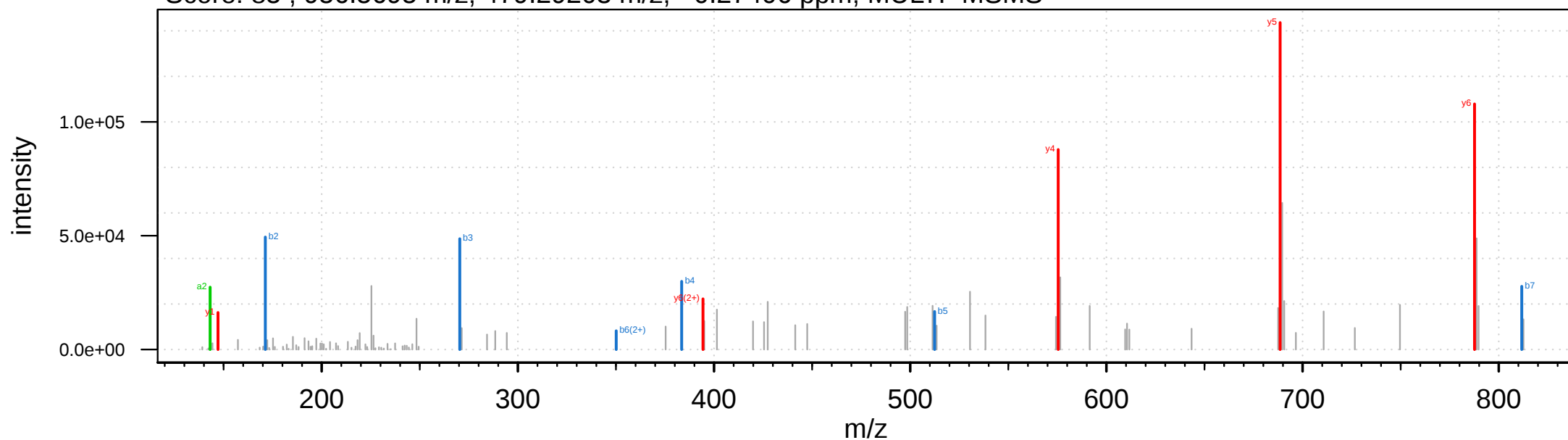


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F4_02
 Scan Number: 4154
 Proteins:
 ENST00000374922_chrX:56755787-56755993:+



GLVLEWIK

Score: 85 ; 956.5695 m/z; 479.29203 m/z; -0.27406 ppm; MULTI-MSMS

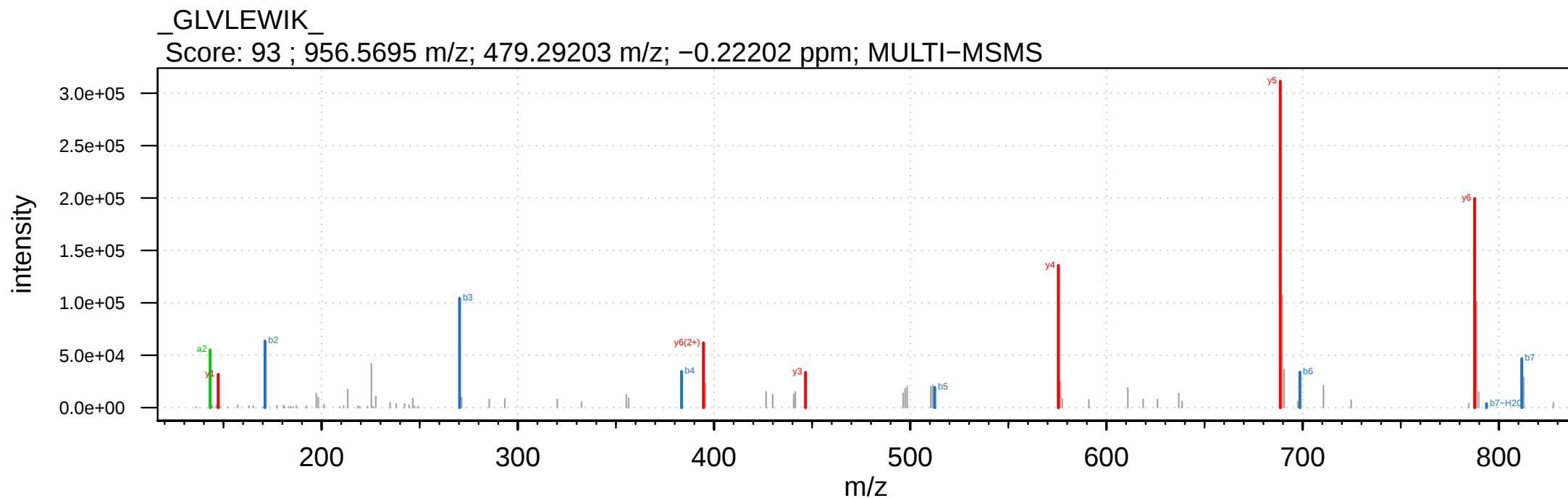
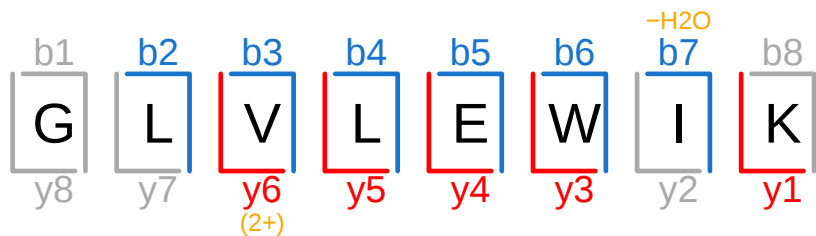


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F6_01

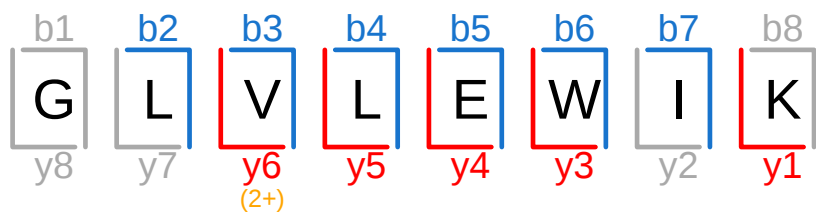
Scan Number: 36545

Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-

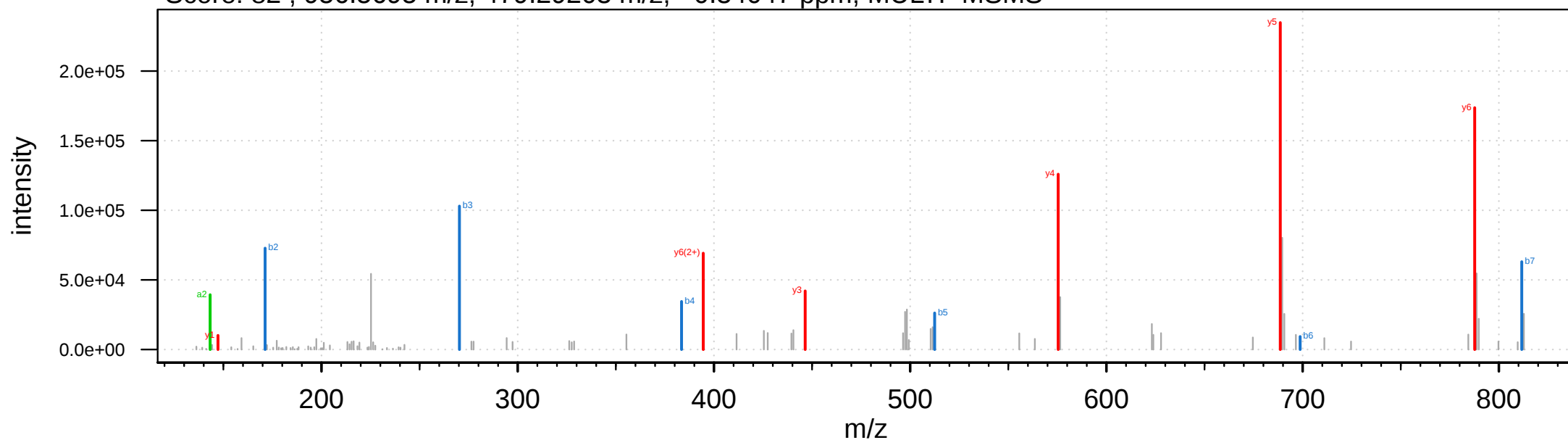


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F7_01
Scan Number: 35694
Proteins:
TCONS_I2_00001296_chr1:79520703-79520992:-



GLVLEWIK

Score: 82 ; 956.5695 m/z; 479.29203 m/z; -0.34047 ppm; MULTI-MSMS

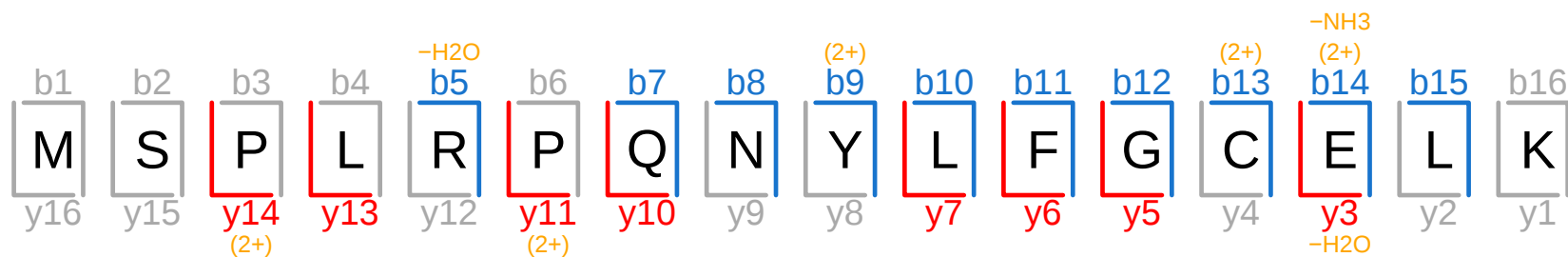


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F7_02

Scan Number: 35780

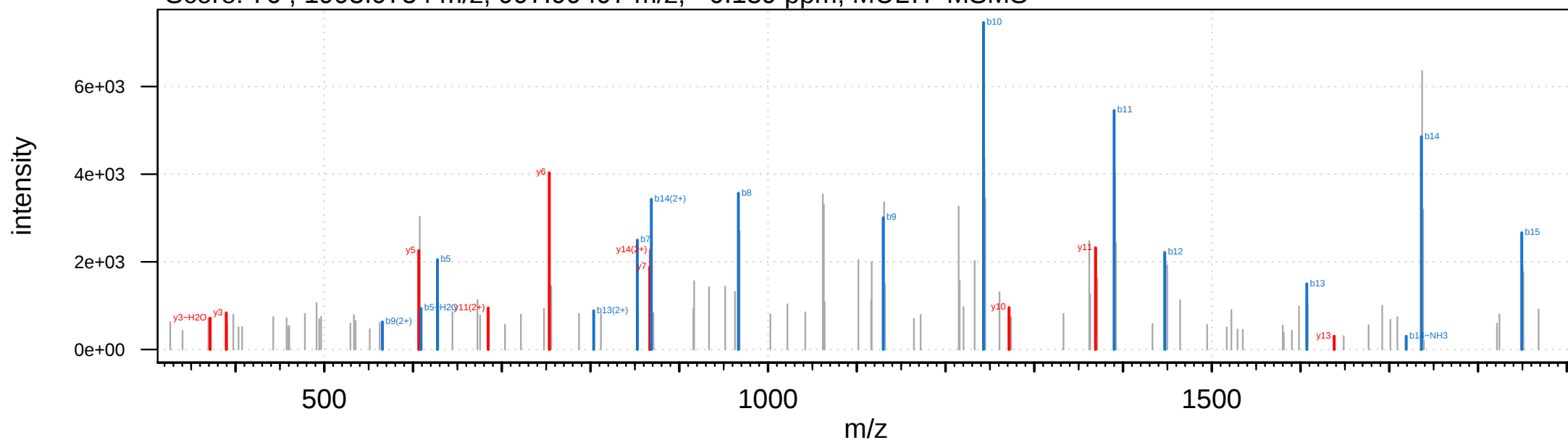
Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-



(ac)MSPLRPQNYLFGCELK

Score: 76 ; 1993.9754 m/z; 997.99497 m/z; -0.139 ppm; MULTI-MSMS

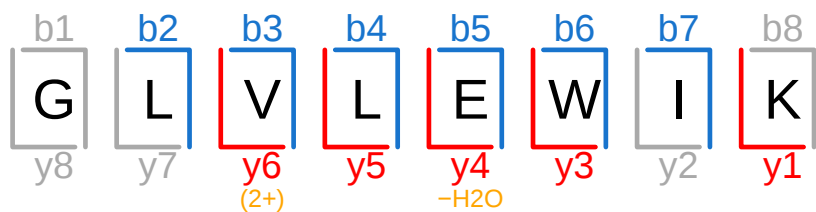


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F7_02

Scan Number: 37820

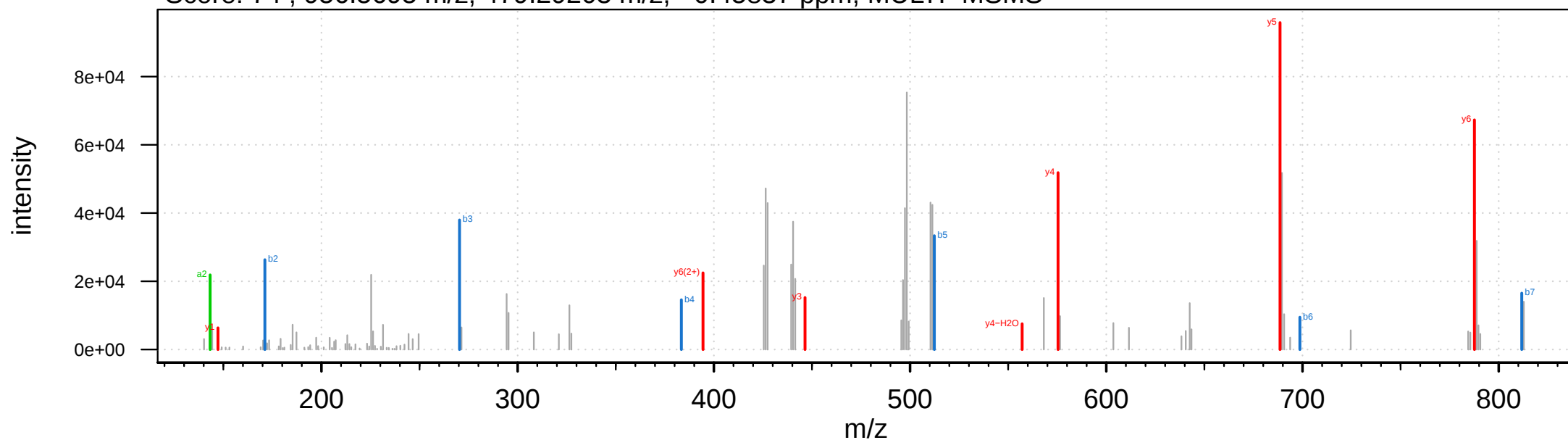
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



GLVLEWIK

Score: 74 ; 956.5695 m/z; 479.29203 m/z; -0.45837 ppm; MULTI-MSMS

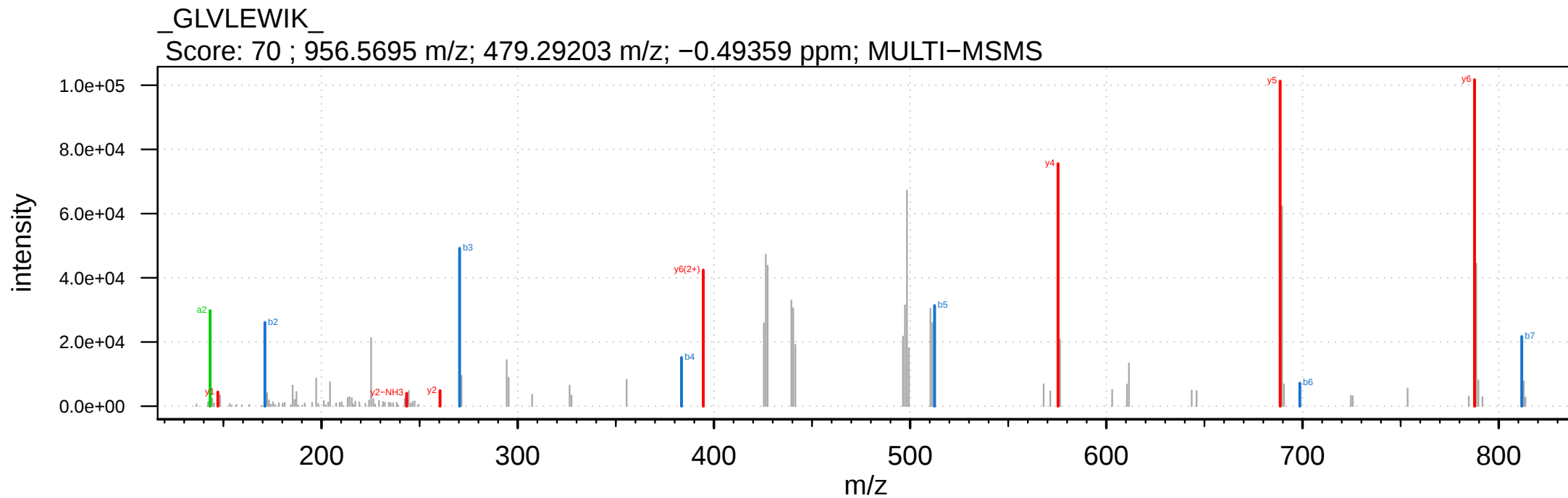
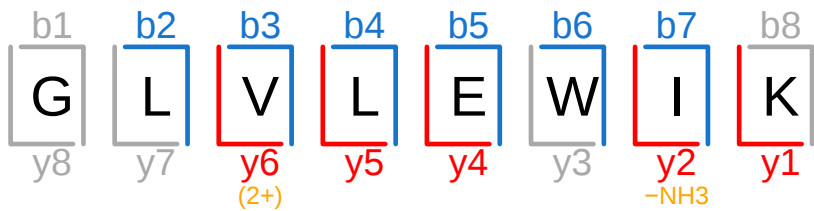


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F8_01

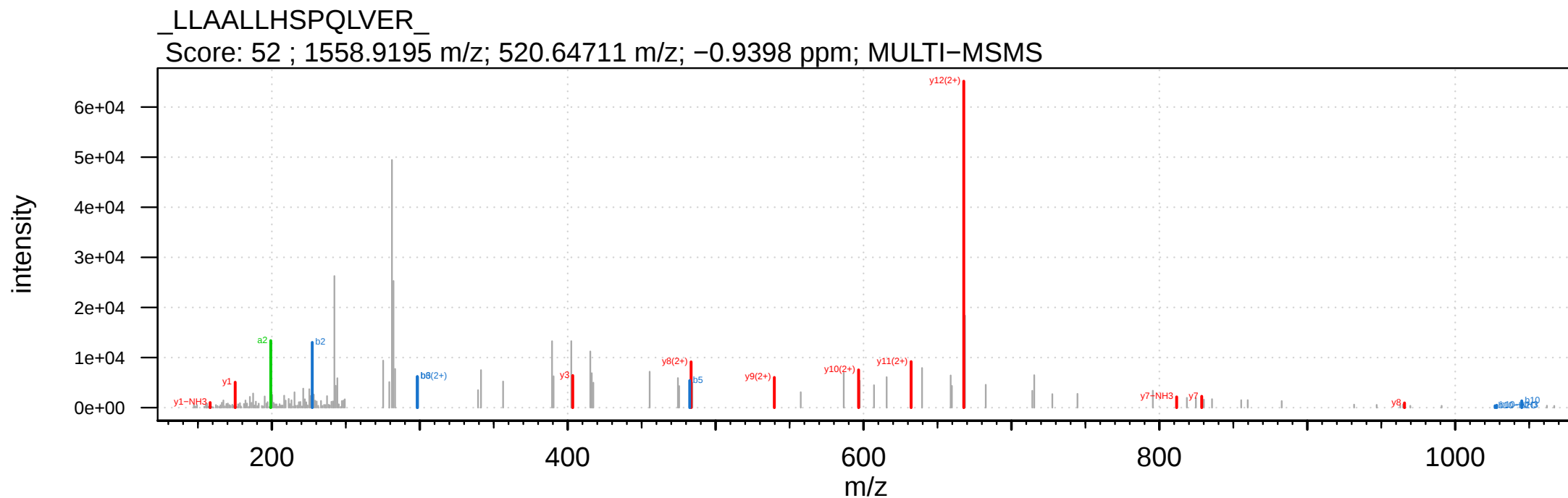
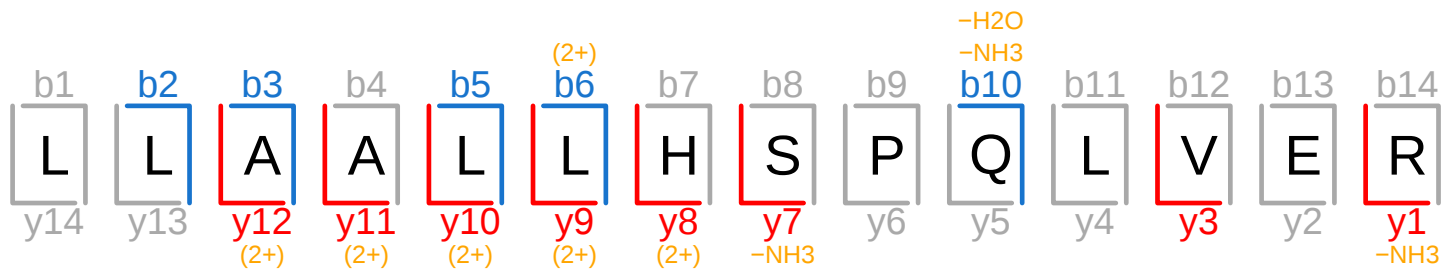
Scan Number: 33753

Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-



Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F8_02
Scan Number: 33679
Proteins:
TCONS_I2_00001296_chr1:79520703-79520992:-



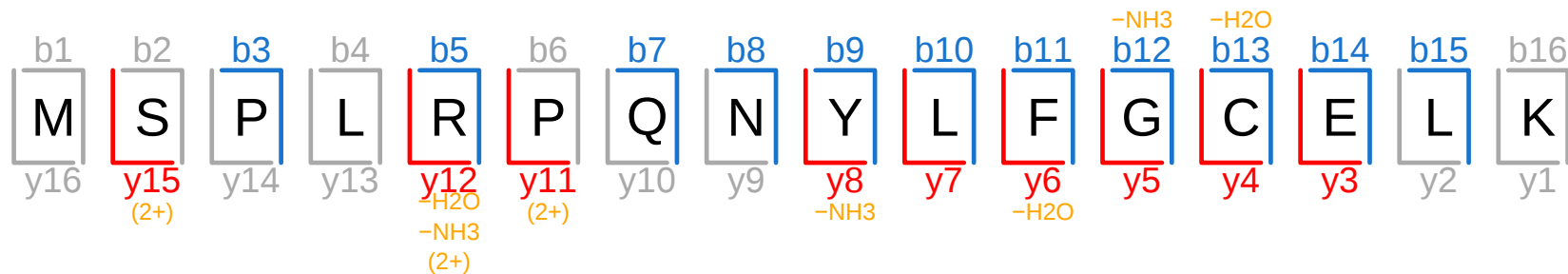
Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F8_02

Scan Number: 30072

Proteins:

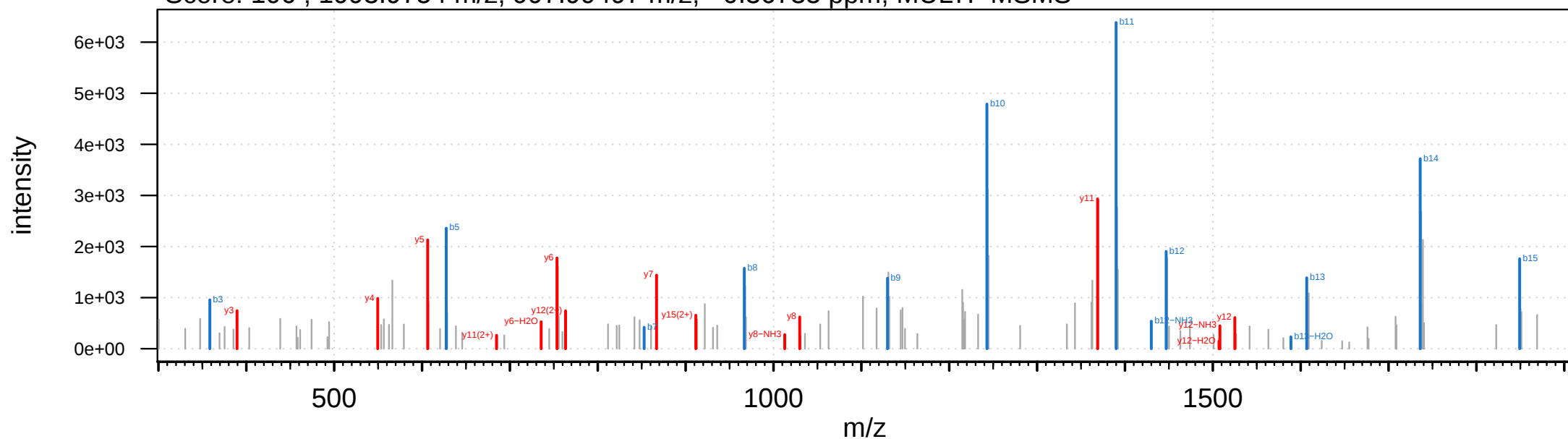
Q69YL0

ENST00000602845_chr3:196669588-196669887:+



(ac)MSPLRPQNYLFGCELK

Score: 106 ; 1993.9754 m/z; 997.99497 m/z; -0.36735 ppm; MULTI-MSMS

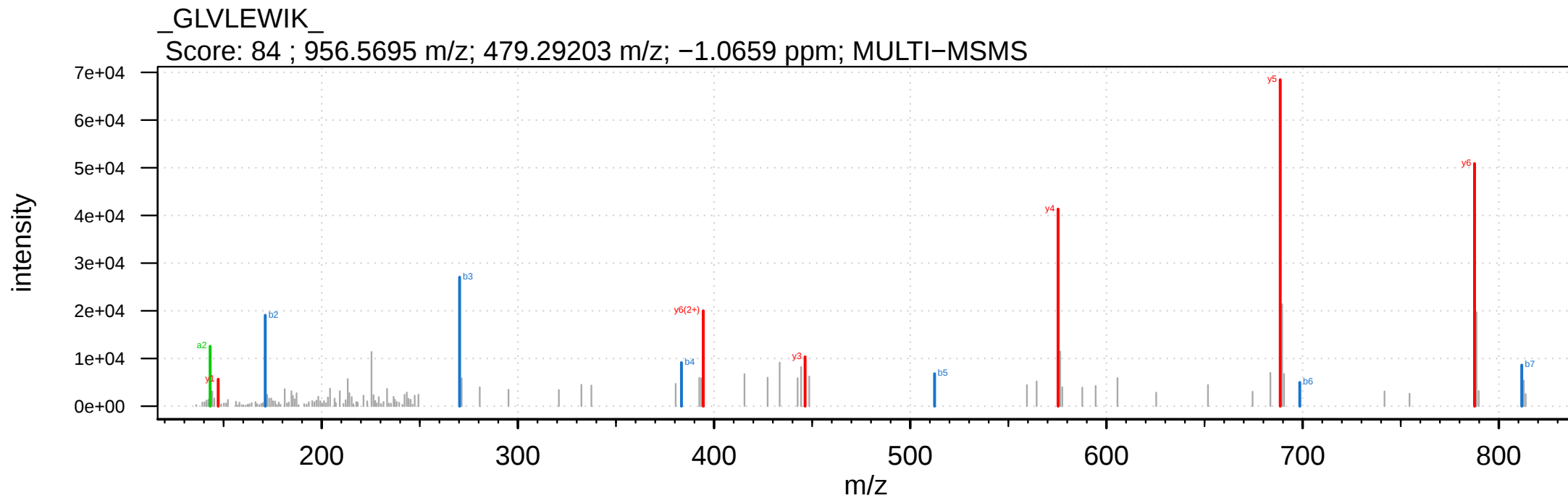
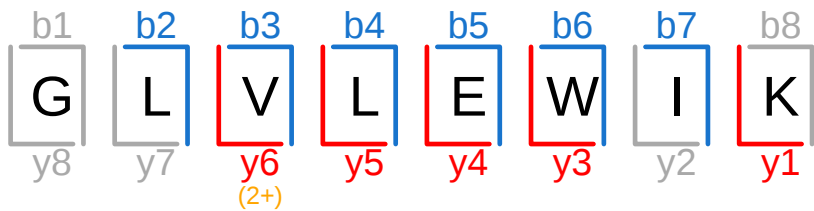


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F8_02

Scan Number: 35611

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

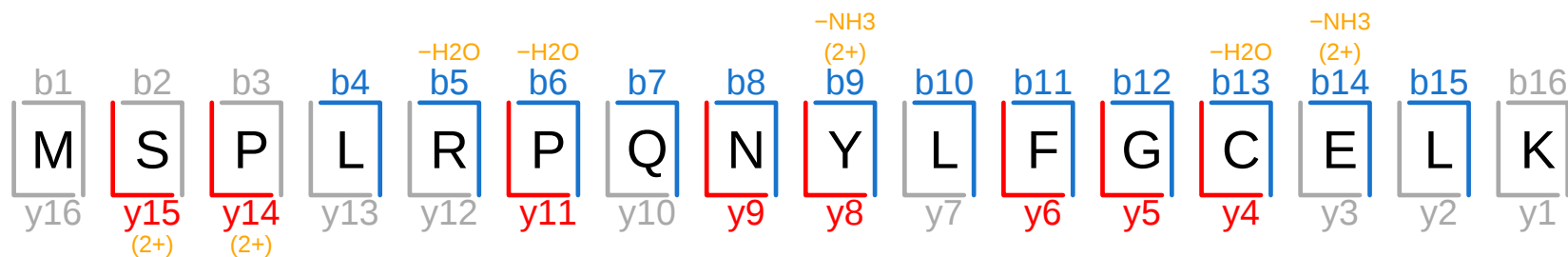


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F9_01

Scan Number: 33104

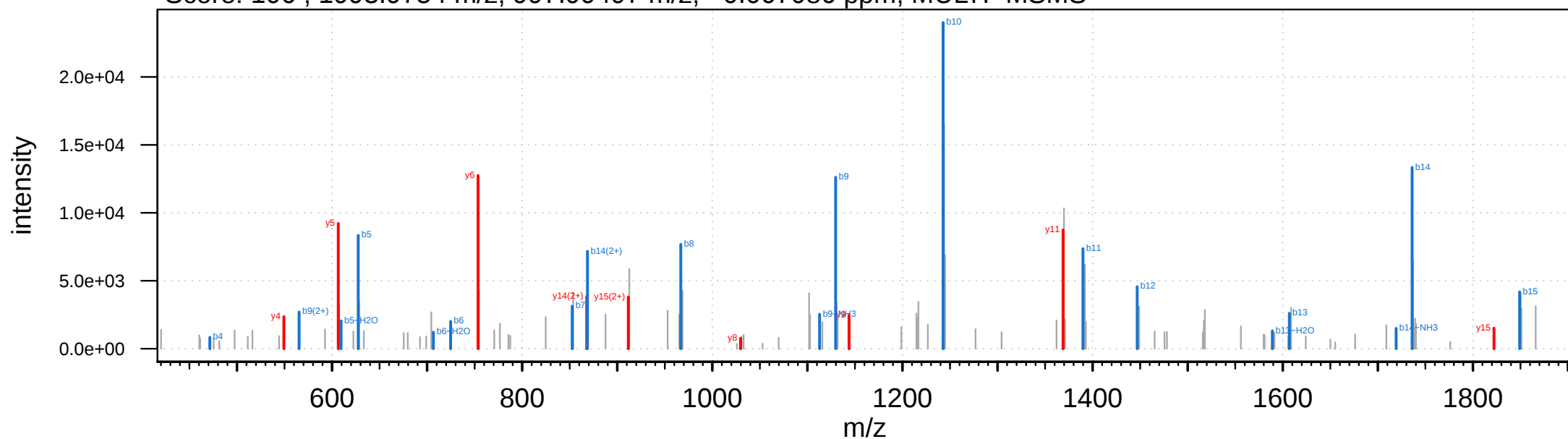
Proteins:

TCONS_I2_00001296_chr1:79520703-79520992:-



(ac)MSPLRPQNYLFGCELK

Score: 106 ; 1993.9754 m/z; 997.99497 m/z; -0.067086 ppm; MULTI-MSMS

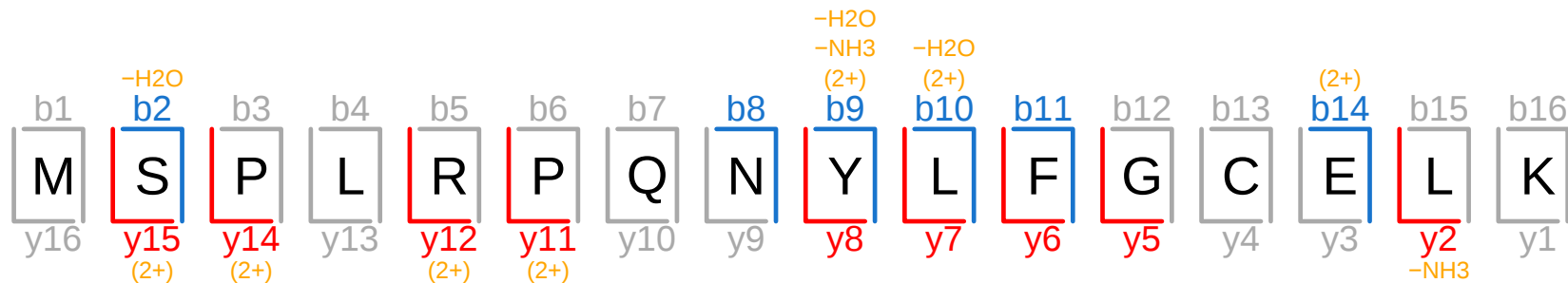


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F9_01

Scan Number: 35025

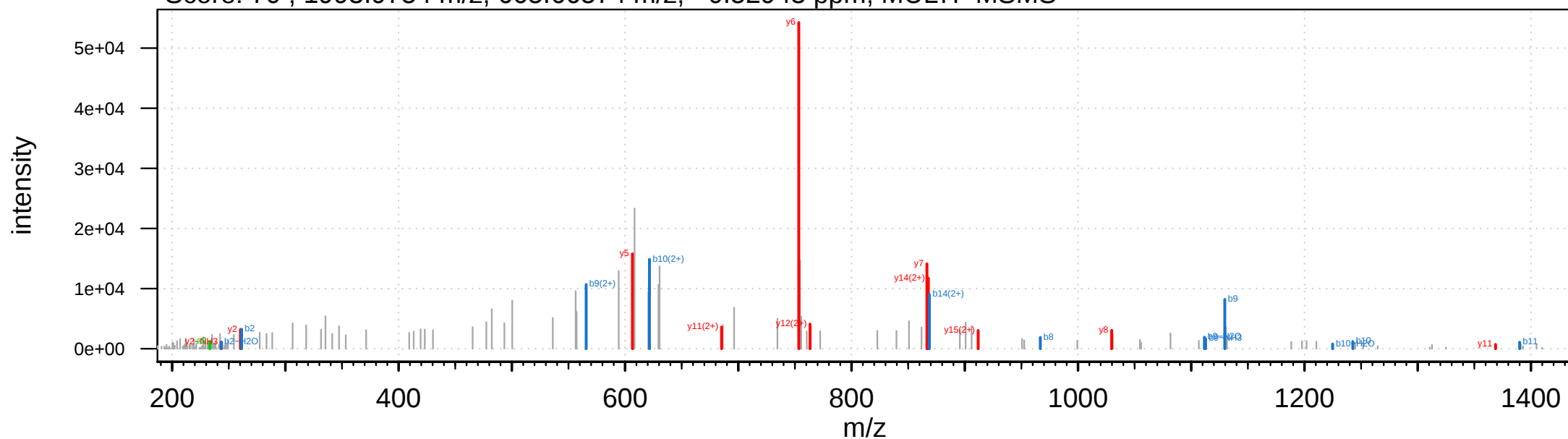
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 70 ; 1993.9754 m/z; 665.66574 m/z; -0.52943 ppm; MULTI-MSMS

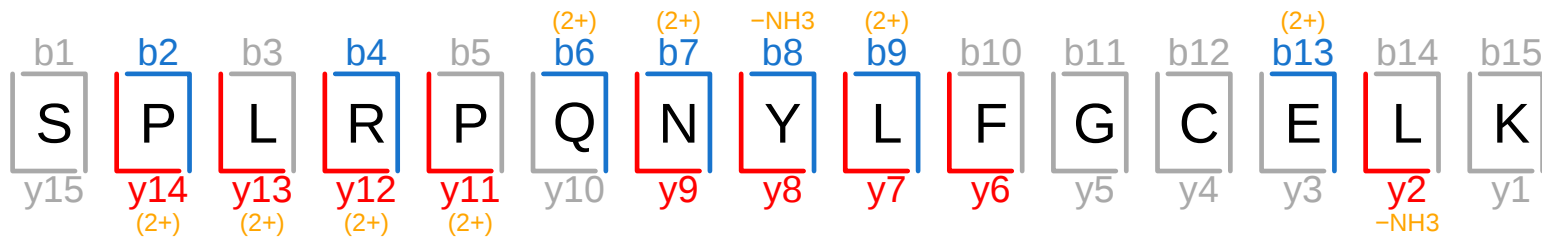


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F9_01

Scan Number: 35027

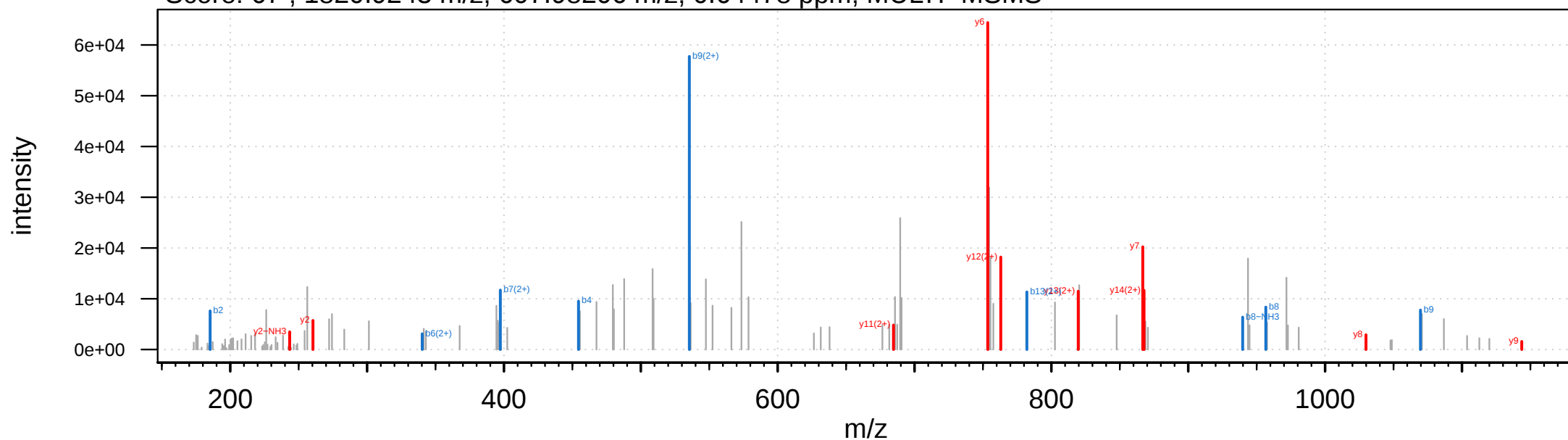
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 67 ; 1820.9243 m/z; 607.98206 m/z; 0.64478 ppm; MULTI-MSMS

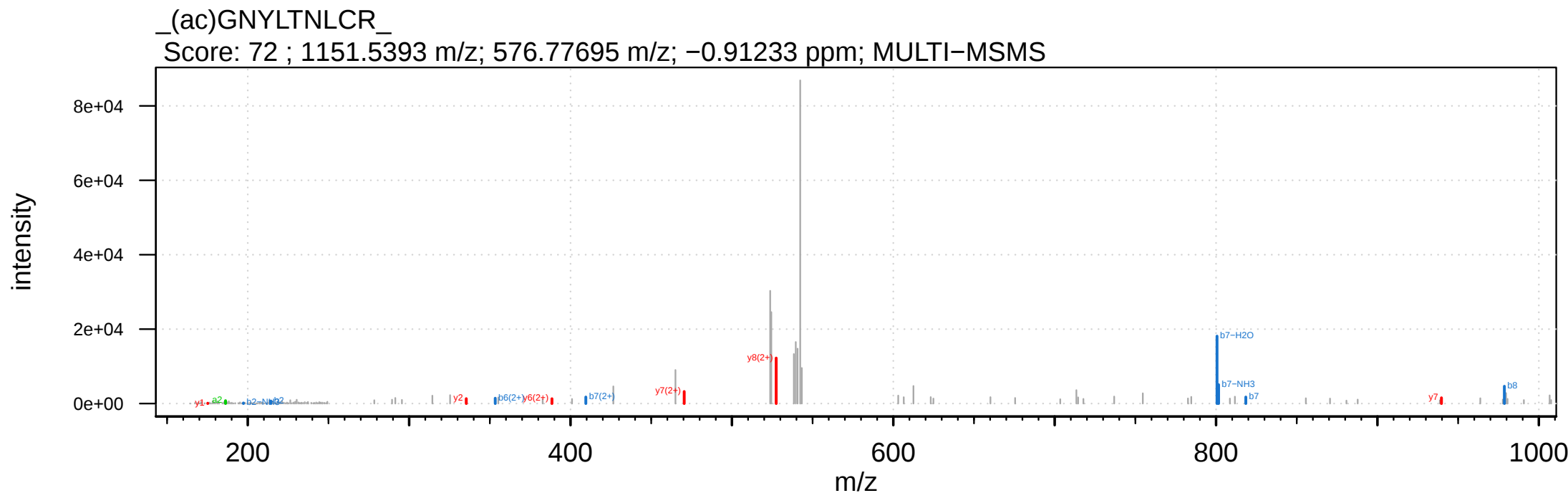
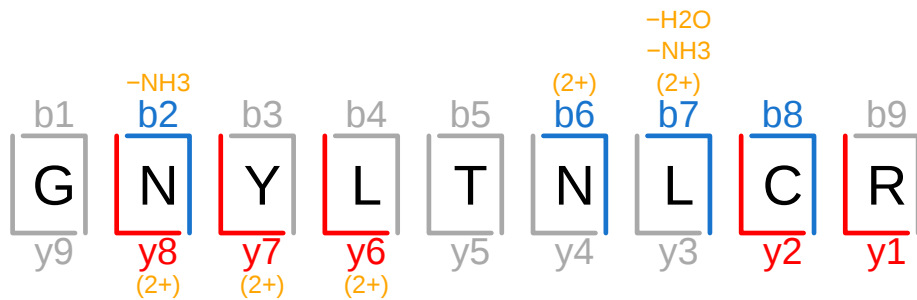


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F9_01

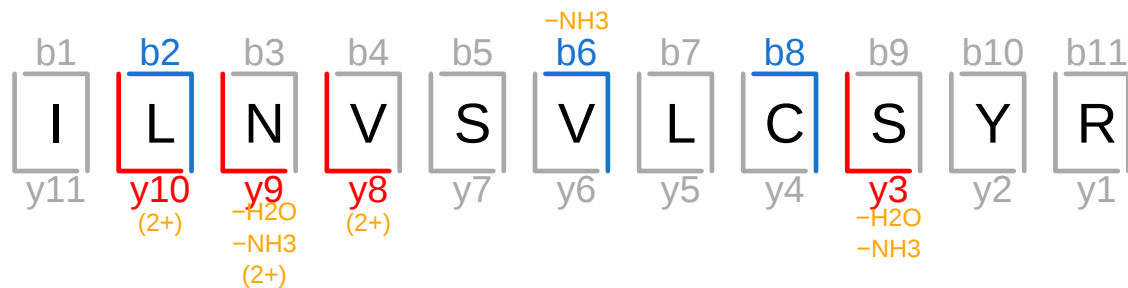
Scan Number: 28225

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

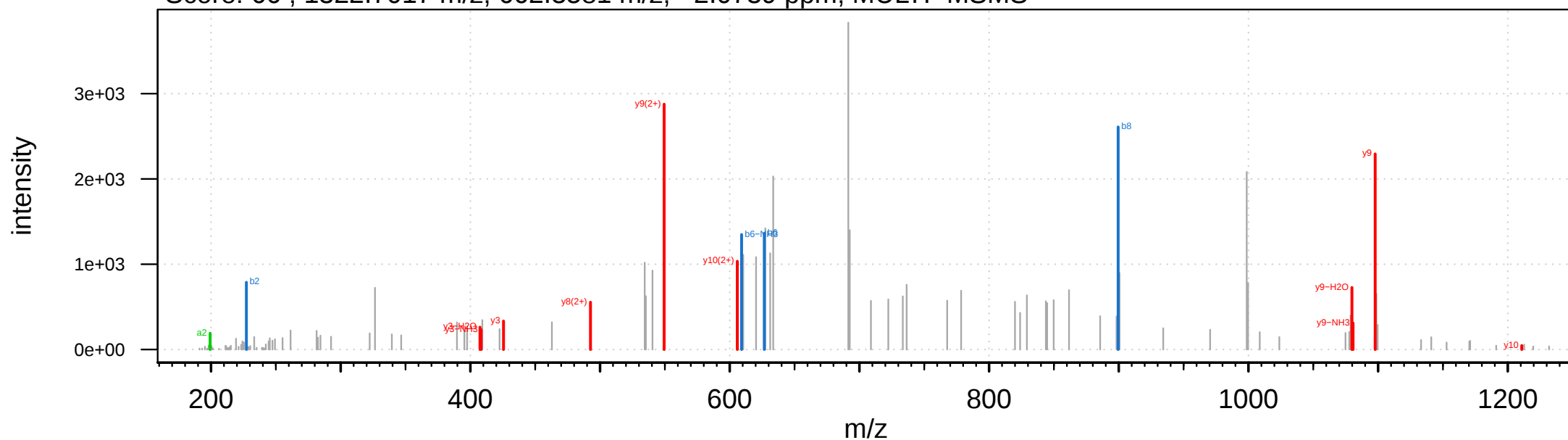


Raw File: Toni_20110506_FB_25_roto_WI38_F7_2
 Scan Number: 8295
 Proteins:
 TCONS_I2_00030467_chrX:38343975-38344091:-



ILNVSVLCSYR

Score: 66 ; 1322.7017 m/z; 662.3581 m/z; -2.0739 ppm; MULTI-MSMS

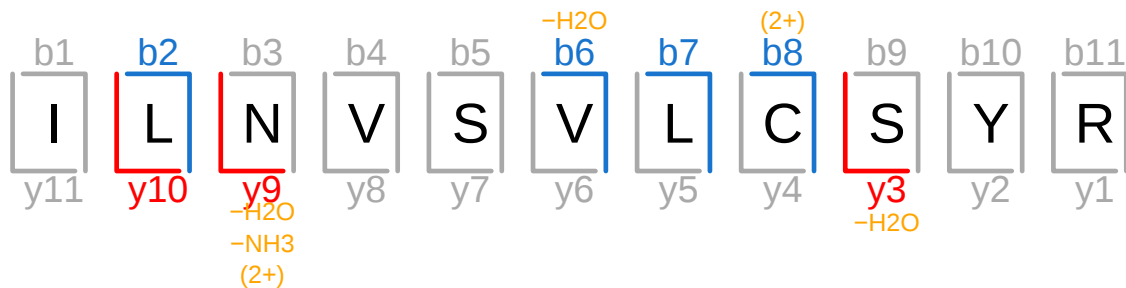


Raw File: Toni_20110506_FB_25_roto_WI38_F7_2

Scan Number: 30370

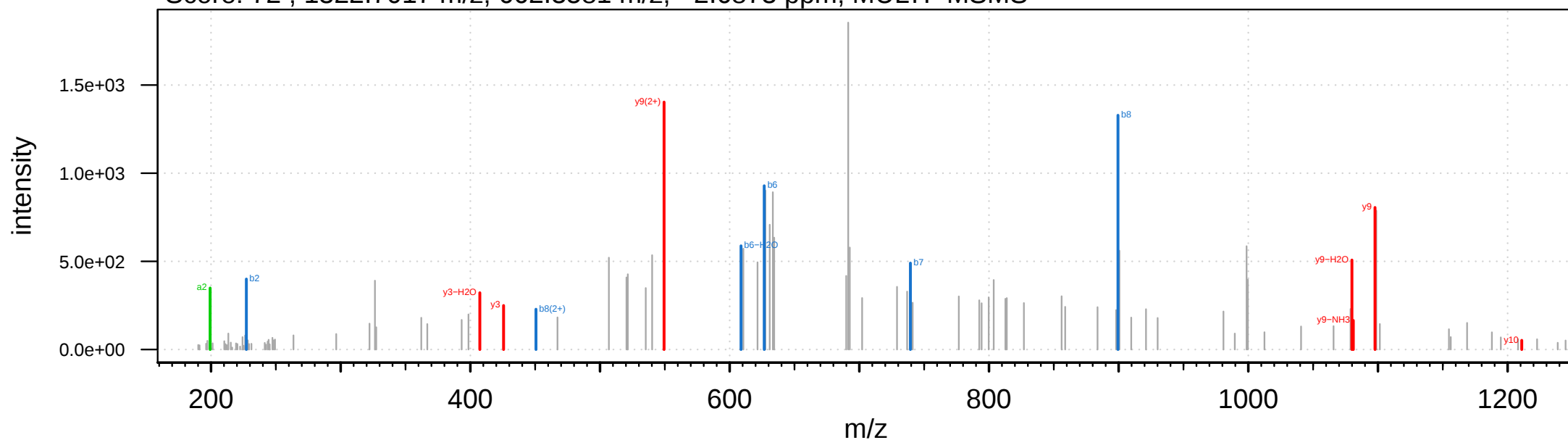
Proteins:

ENST00000297293_chr7:97835505-97835555:+



ILNVSVLCSYR

Score: 72 ; 1322.7017 m/z; 662.3581 m/z; -2.6873 ppm; MULTI-MSMS

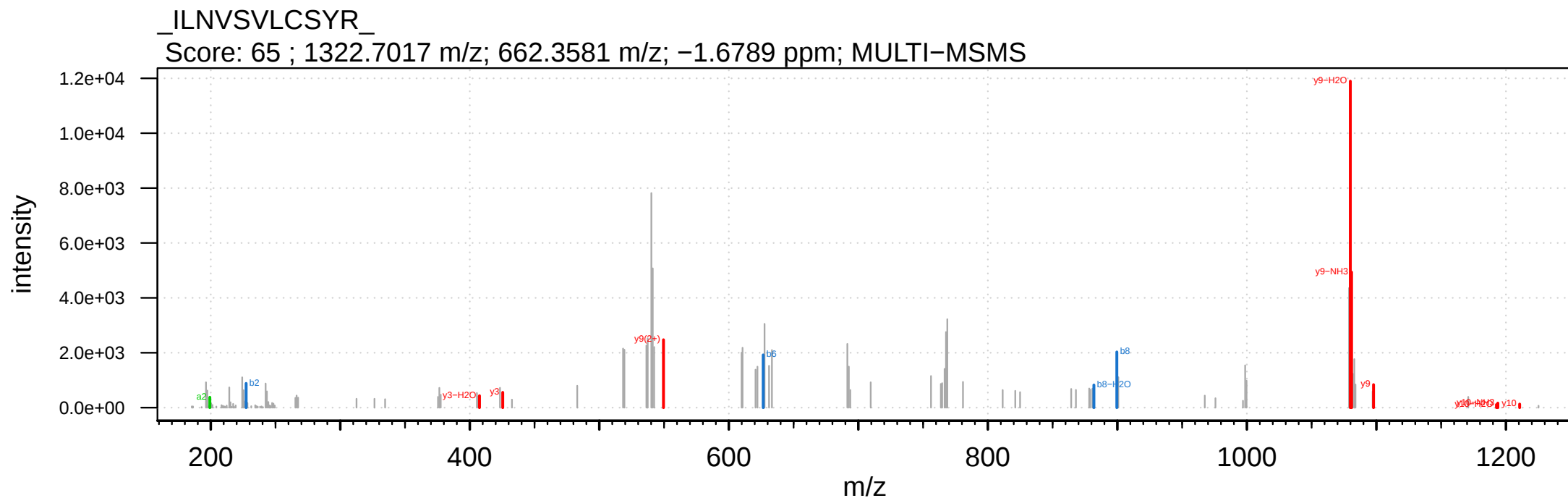
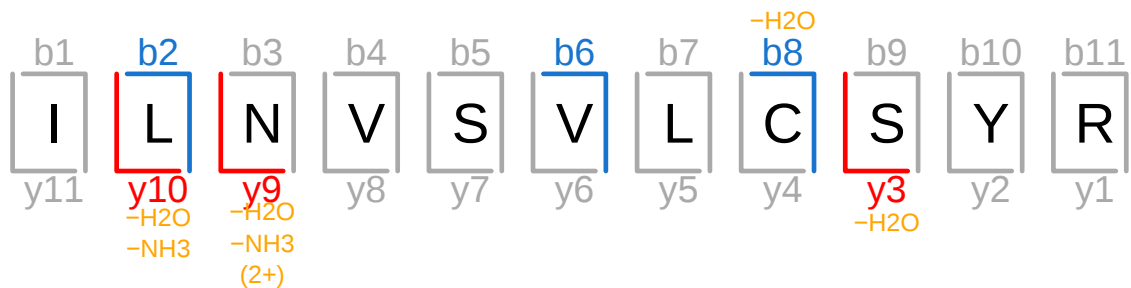


Raw File: Toni_20110506_FB_25_roto_WI38_F6_2

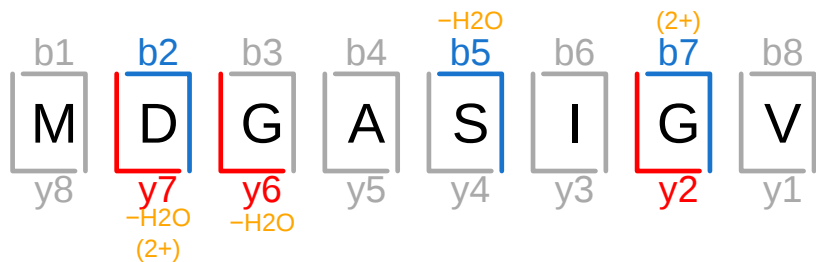
Scan Number: 29288

Proteins:

ENST00000297293_chr7:97835505-97835555:+

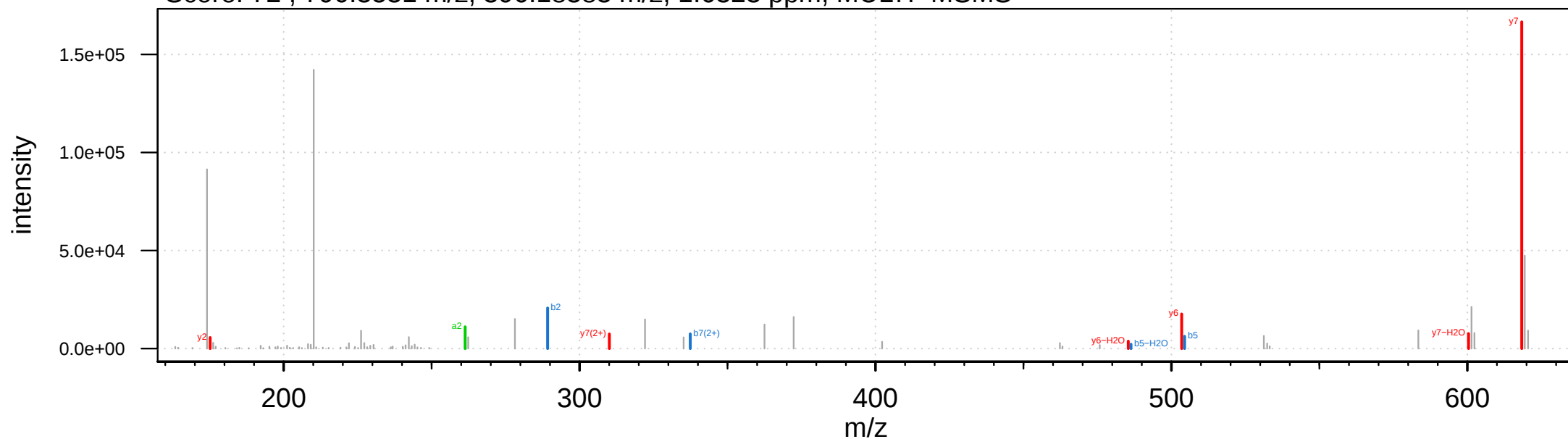


Raw File: Toni_20110506_FB_25_roto_WI38_F9_2
Scan Number: 29385
Proteins:
ENST00000297293_chr7:97835505-97835555:+



(ac)MDGASIGV

Score: 72 ; 790.3531 m/z; 396.18383 m/z; 1.6323 ppm; MULTI-MSMS

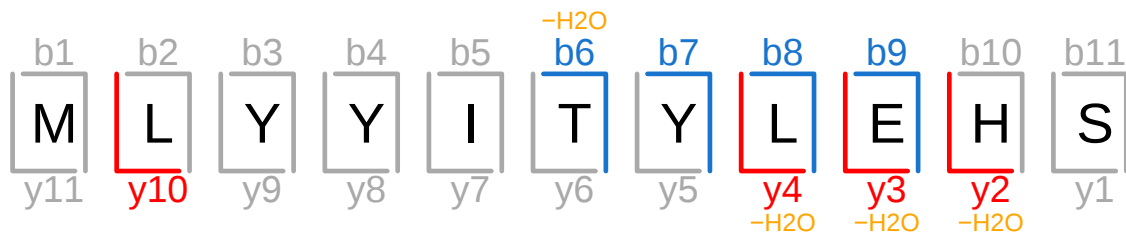


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F3_02

Scan Number: 17796

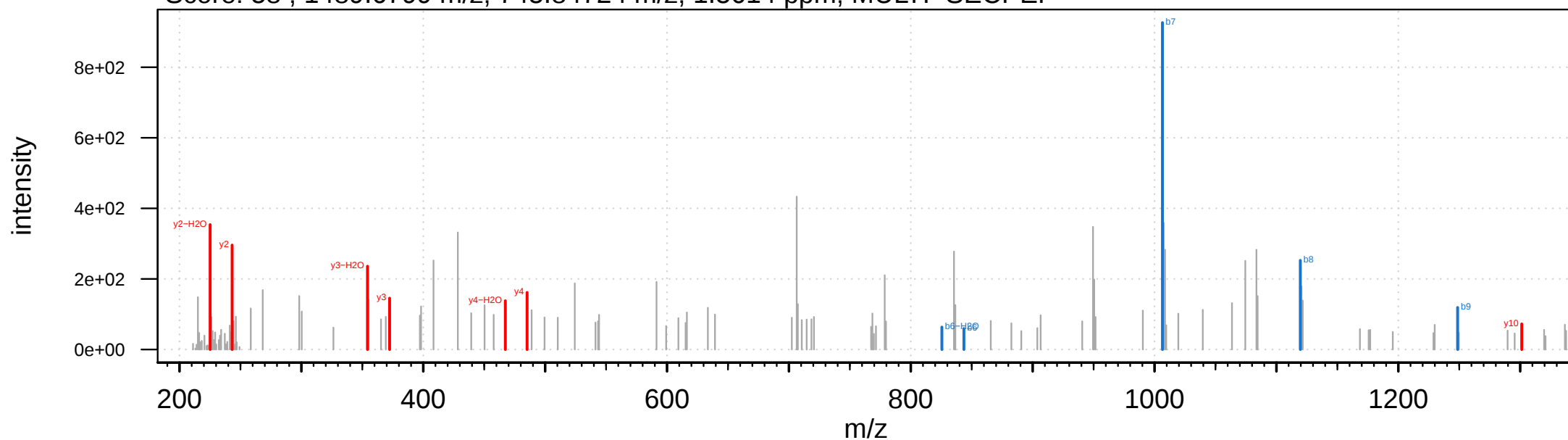
Proteins:

ENST00000529489_chr1:236681646-236681672:+



(ac)M(ox)LYYITYLEHS

Score: 58 ; 1489.6799 m/z; 745.84724 m/z; 1.5614 ppm; MULTI-SECPEP

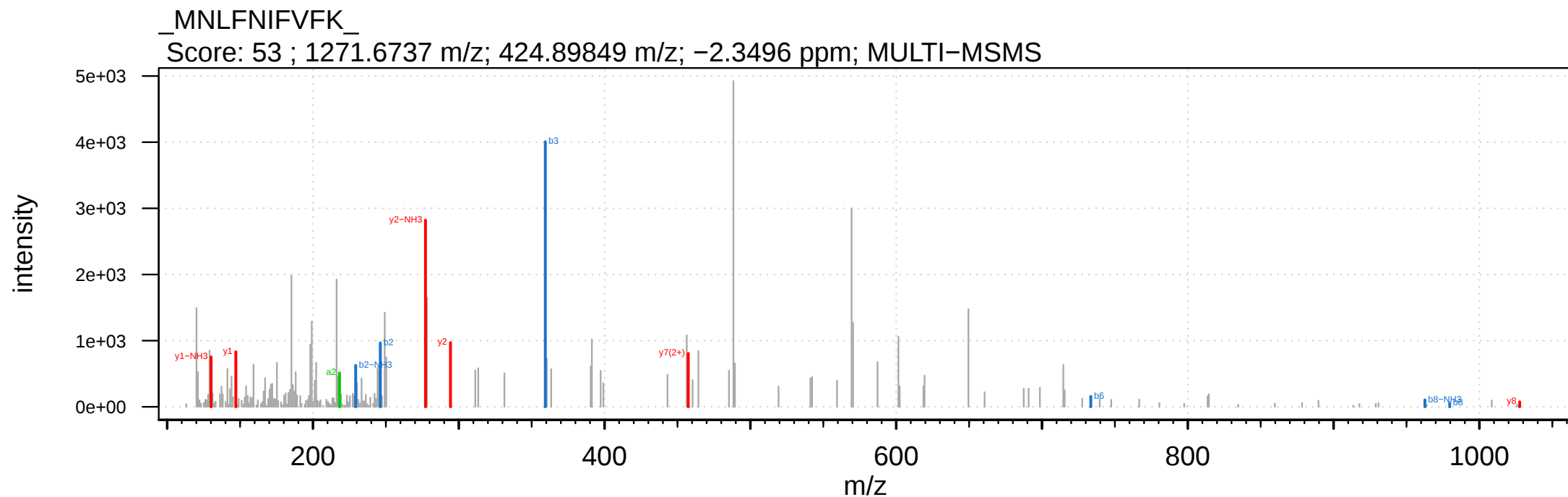
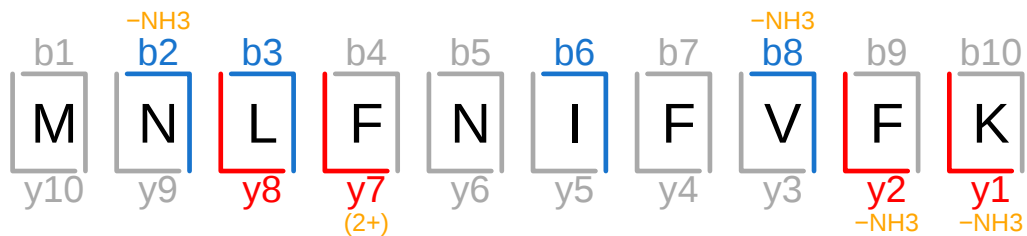


Raw File: Toni_20111125_FB_MCF7_F4_1

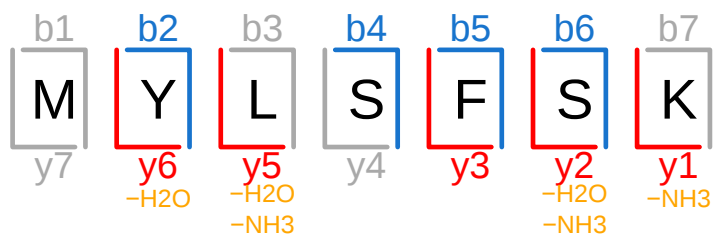
Scan Number: 39180

Proteins:

ENST00000317881_chrX:118605057-118605092:+

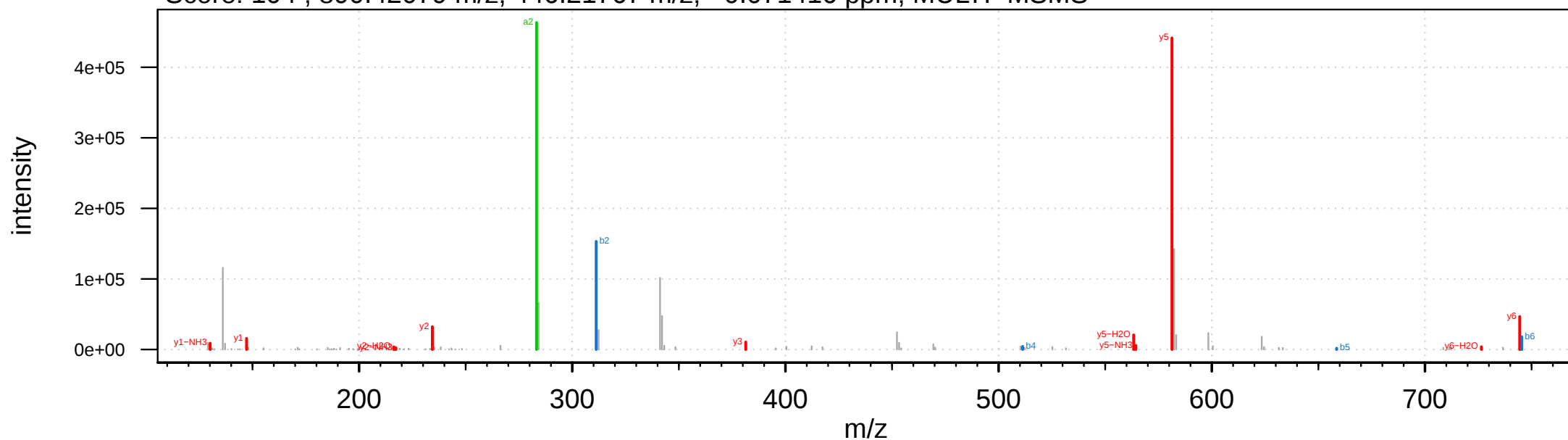


Raw File: Toni_20111125_FB_MCF7_F6_2
Scan Number: 33622
Proteins:
ENST00000274364_chr5:76003185-76003223:+



M(ox)YLSFSK

Score: 104 ; 890.42079 m/z; 446.21767 m/z; -0.071416 ppm; MULTI-MSMS

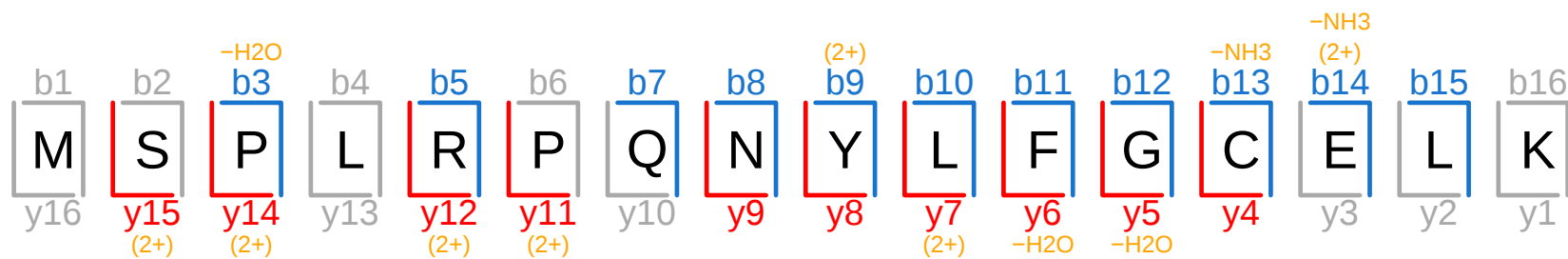


Raw File: Toni_20111125_FB_MCF7_F6_2

Scan Number: 27380

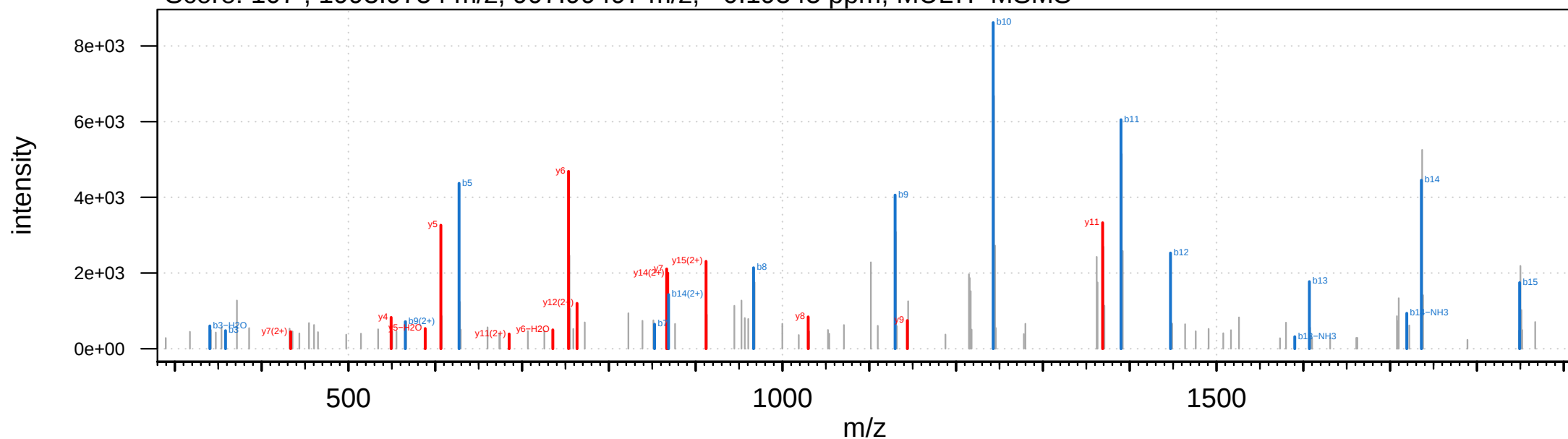
Proteins:

ENST00000392145_chr6:160874572-160874622:+



(ac)MSPLRPQNYLFGCELEK

Score: 107 ; 1993.9754 m/z; 997.99497 m/z; -0.19545 ppm; MULTI-MSMS

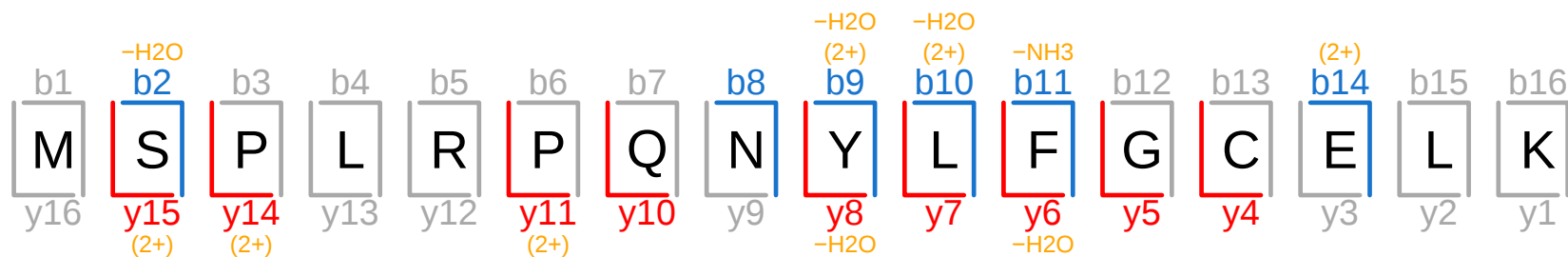


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F9_02

Scan Number: 35058

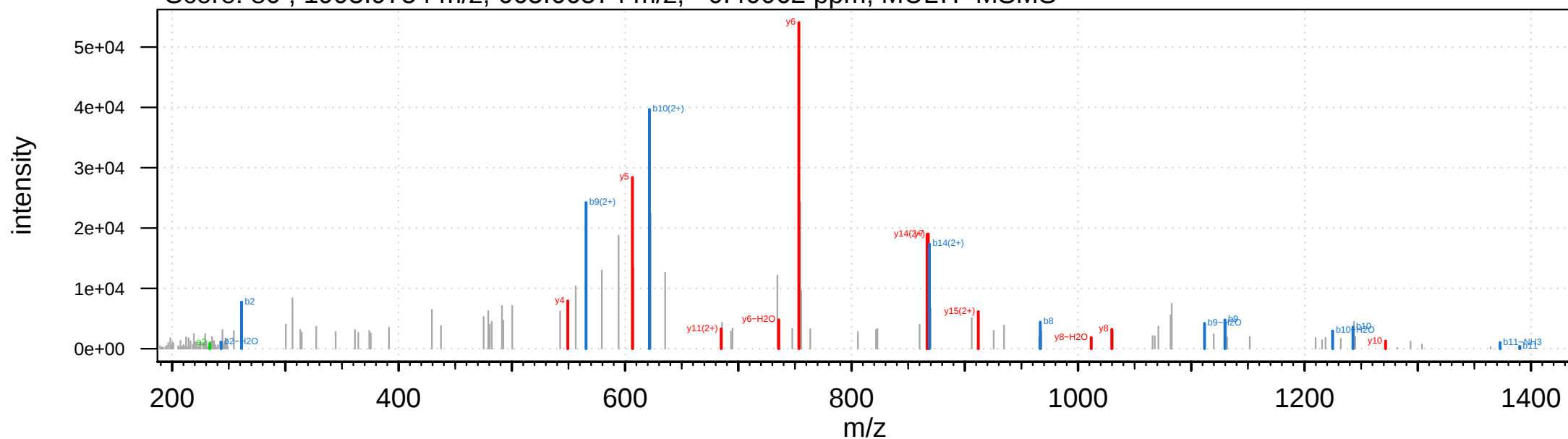
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 80 ; 1993.9754 m/z; 665.66574 m/z; -0.40062 ppm; MULTI-MSMS

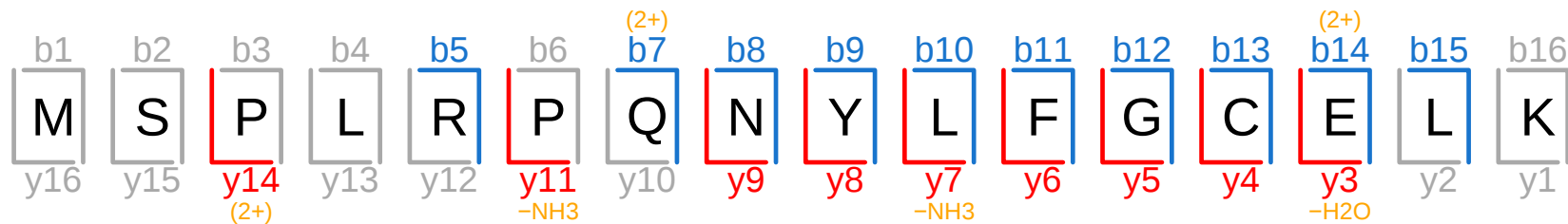


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F9_02

Scan Number: 35089

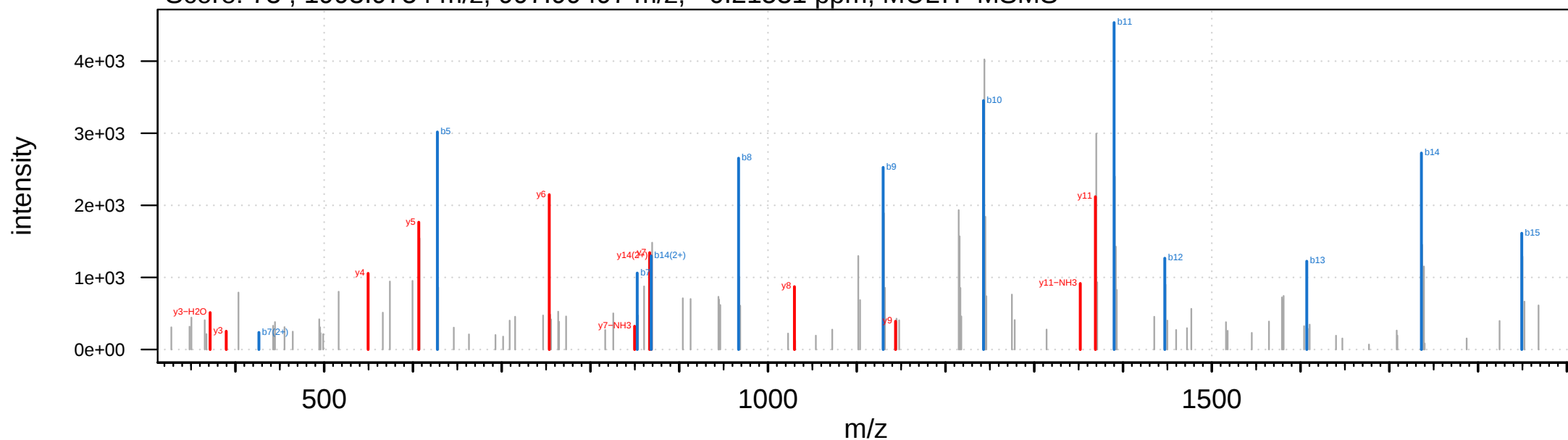
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELEK

Score: 73 ; 1993.9754 m/z; 997.99497 m/z; -0.21531 ppm; MULTI-MSMS

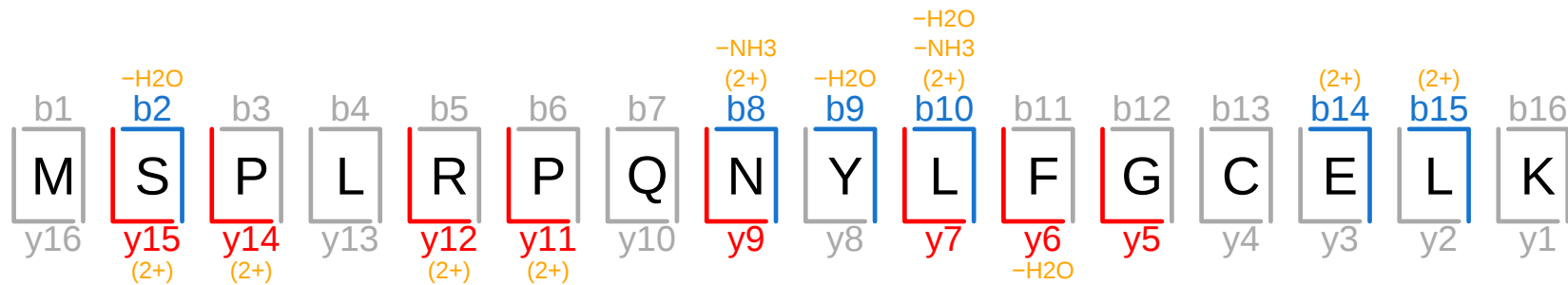


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F10_01

Scan Number: 34770

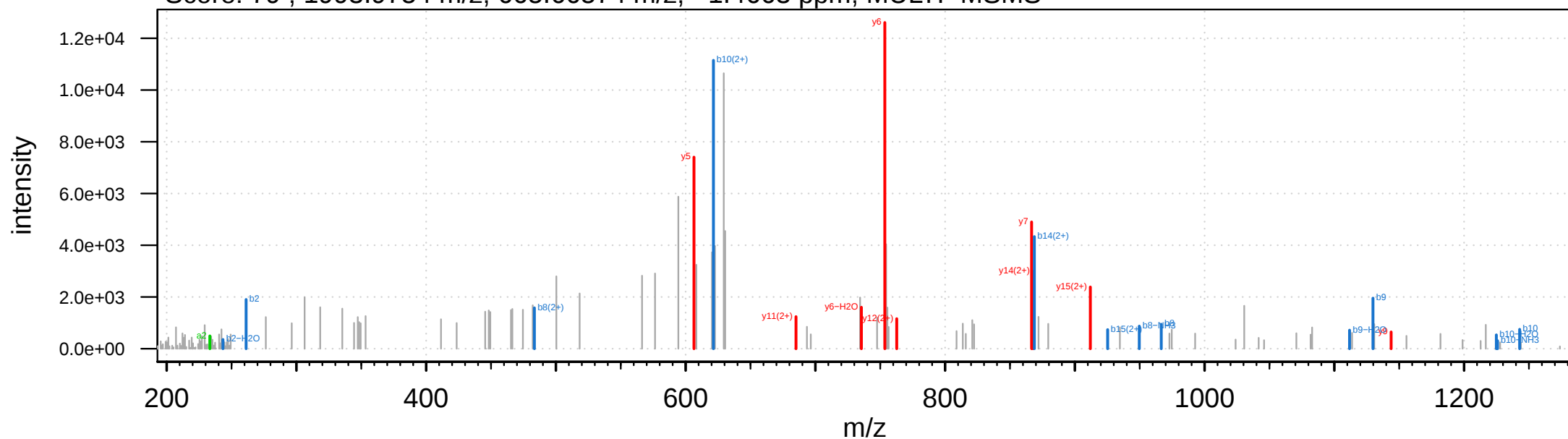
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 79 ; 1993.9754 m/z; 665.66574 m/z; -1.4663 ppm; MULTI-MSMS

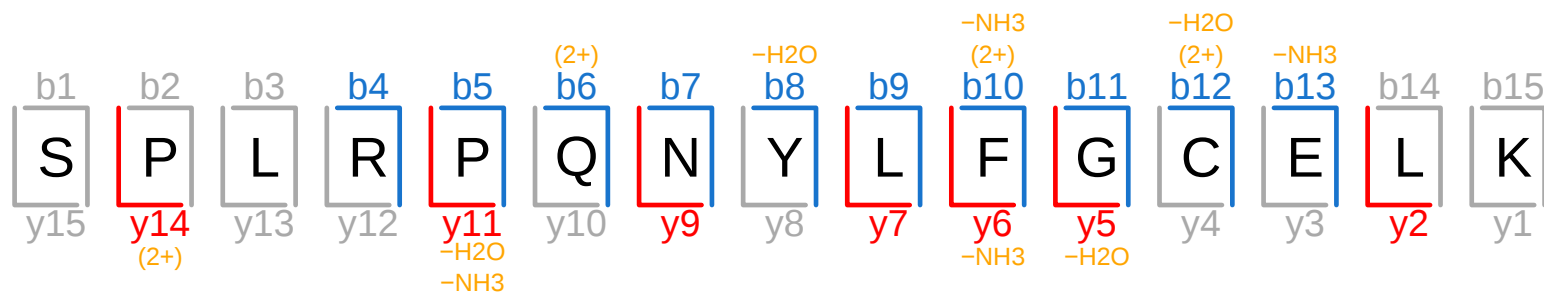


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F10_01

Scan Number: 34780

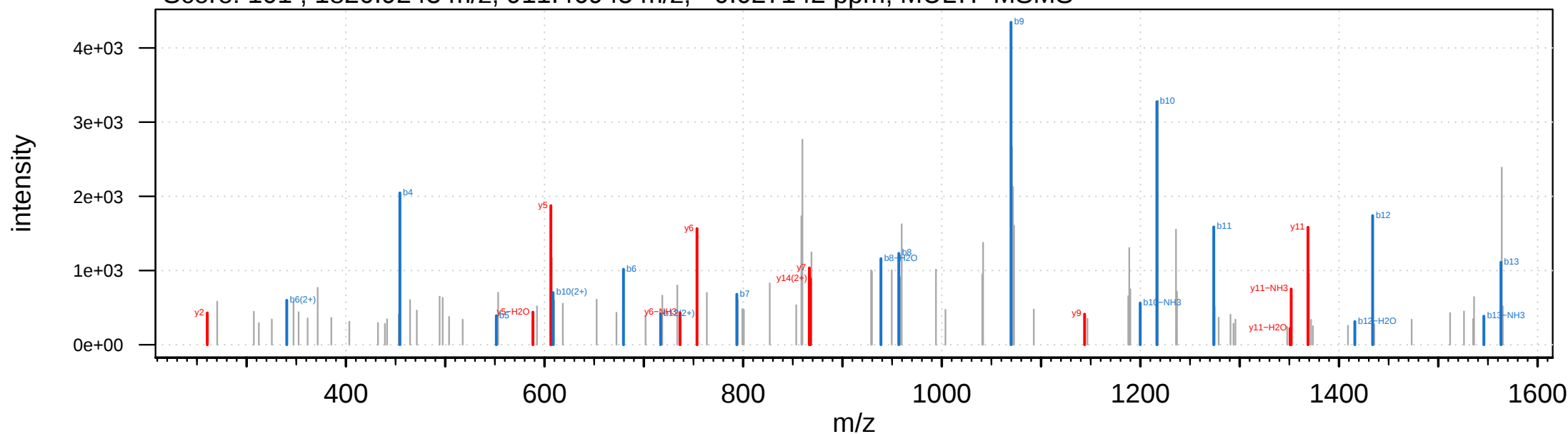
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 101 ; 1820.9243 m/z; 911.46945 m/z; -0.027142 ppm; MULTI-MSMS

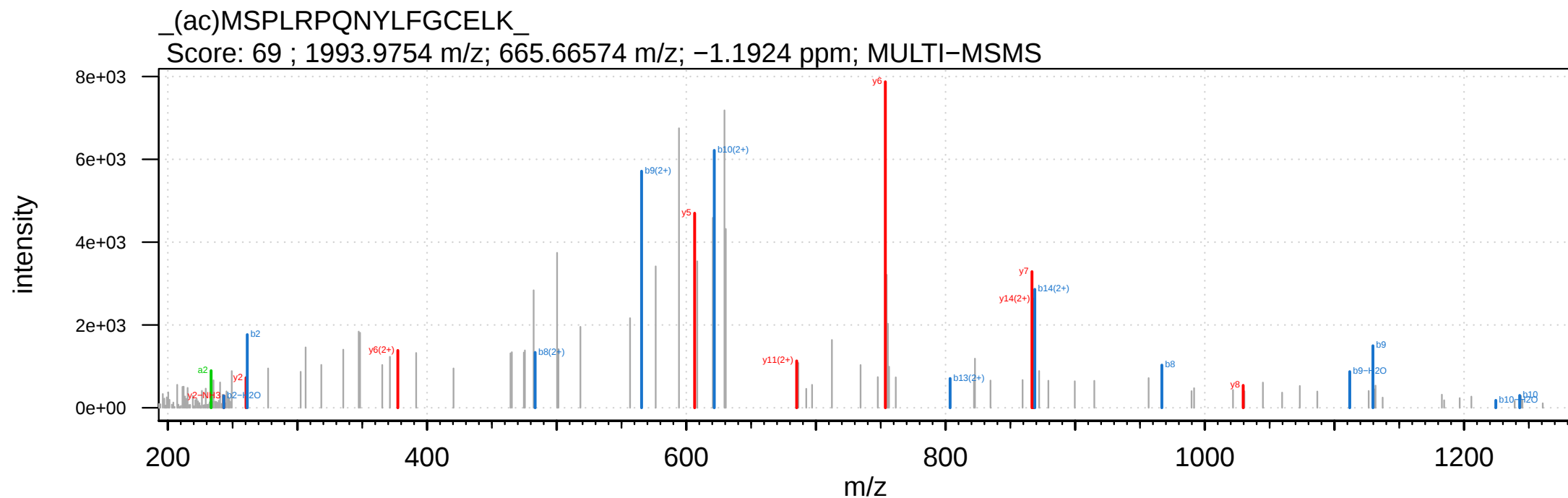
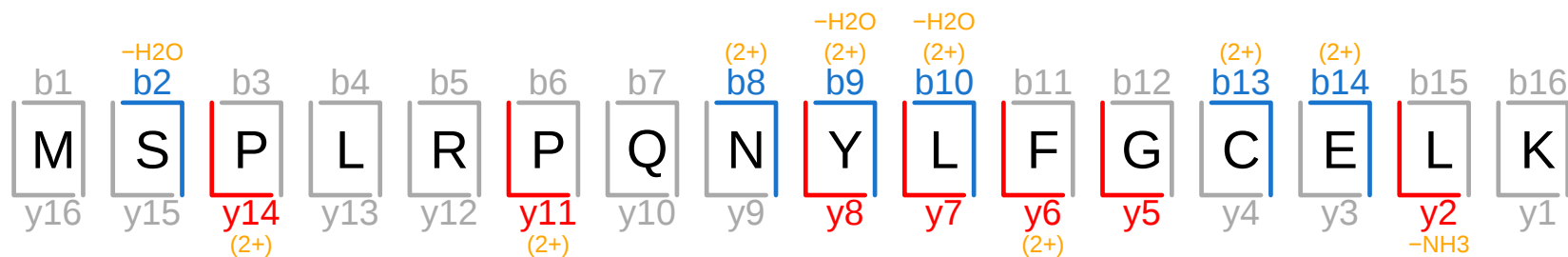


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F10_01

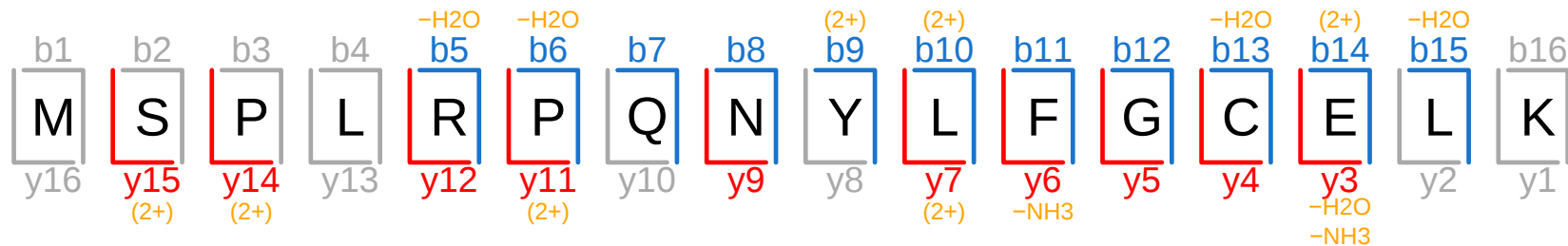
Scan Number: 28299

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+

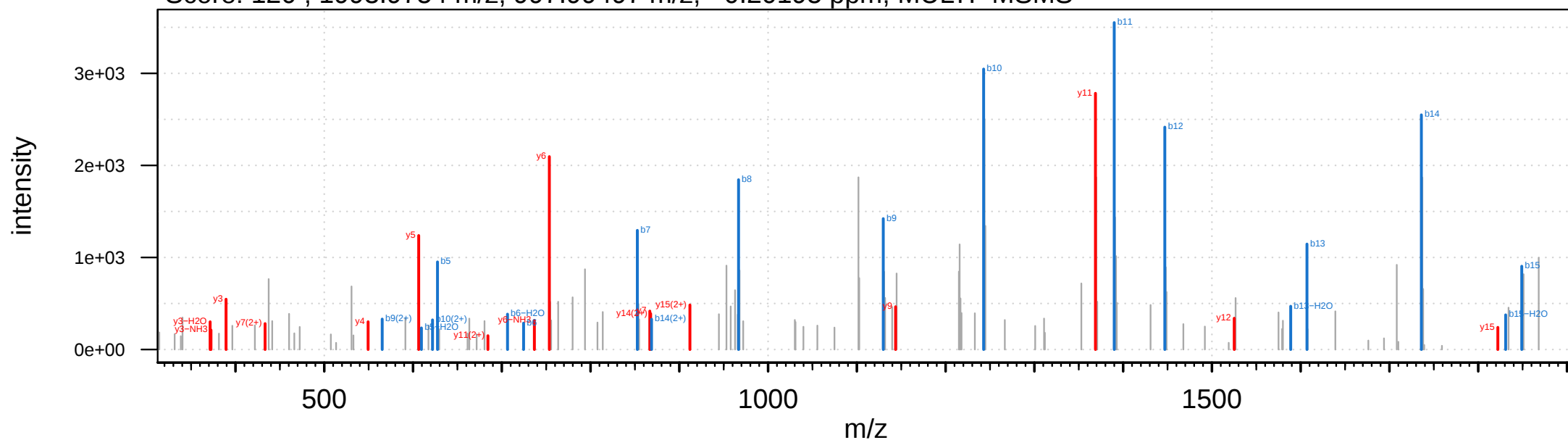


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F10_02
 Scan Number: 34946
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)MSPLRPQNYLFGCELK

Score: 120 ; 1993.9754 m/z; 997.99497 m/z; -0.20195 ppm; MULTI-MSMS

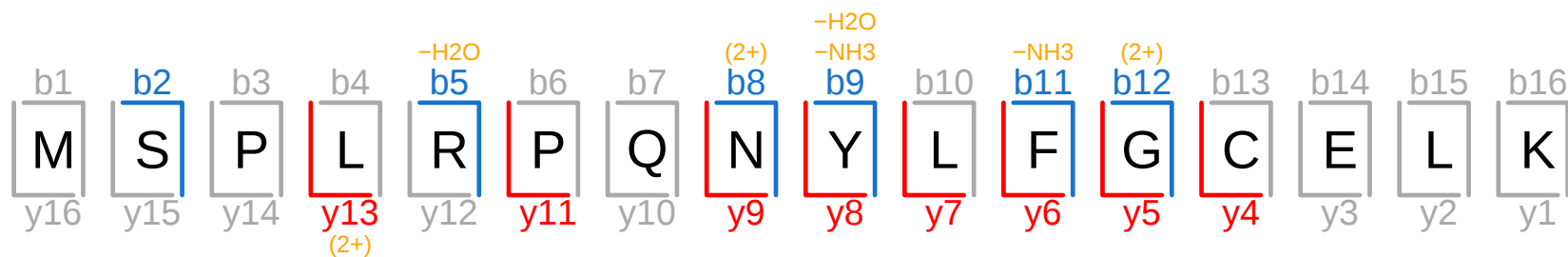


Raw File: Toni_20110714_FB_HepG2_2dot5_uROTO_F10_02

Scan Number: 34948

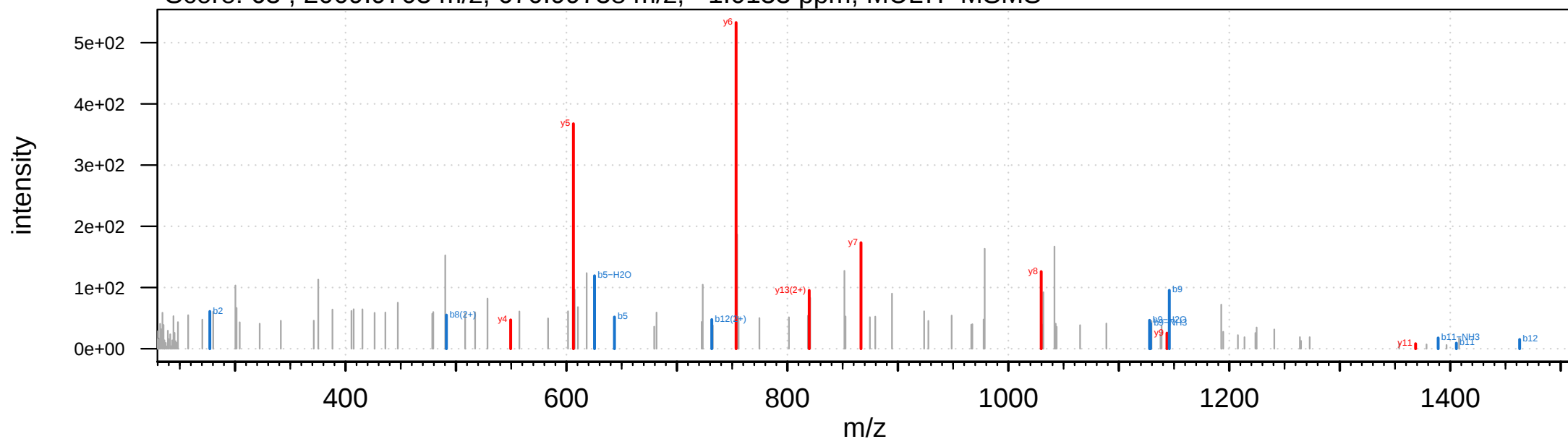
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



(ac)M(ox)SPLRPQNYLFGCELK

Score: 63 ; 2009.9703 m/z; 670.99738 m/z; -1.0133 ppm; MULTI-MSMS

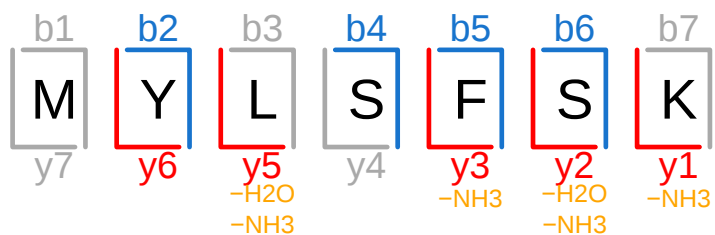


Raw File: Toni_20111125_FB_MCF7_F9_2

Scan Number: 42305

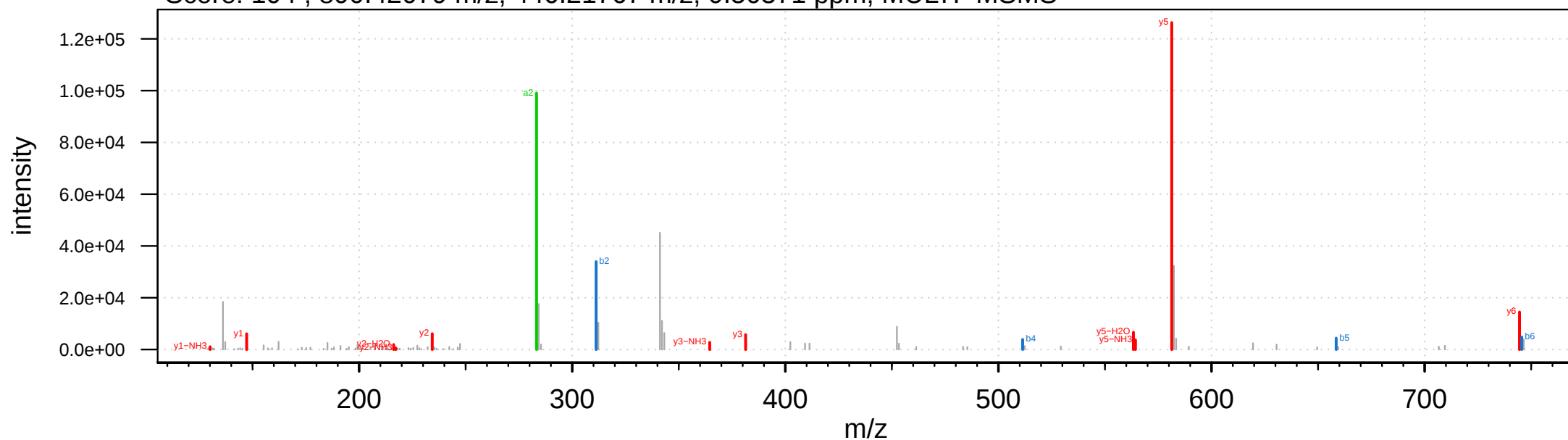
Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



M(ox)YLSFSK

Score: 104 ; 890.42079 m/z; 446.21767 m/z; 0.50371 ppm; MULTI-MSMS

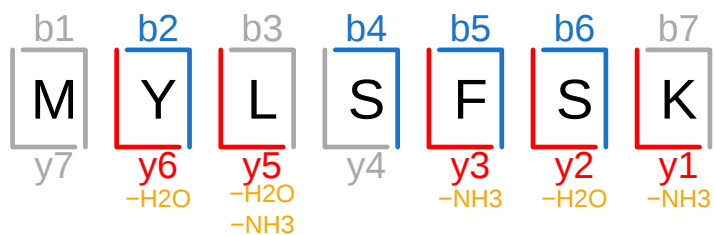


Raw File: Toni_20111114_FB_MDA-MB-231_F7_1

Scan Number: 23407

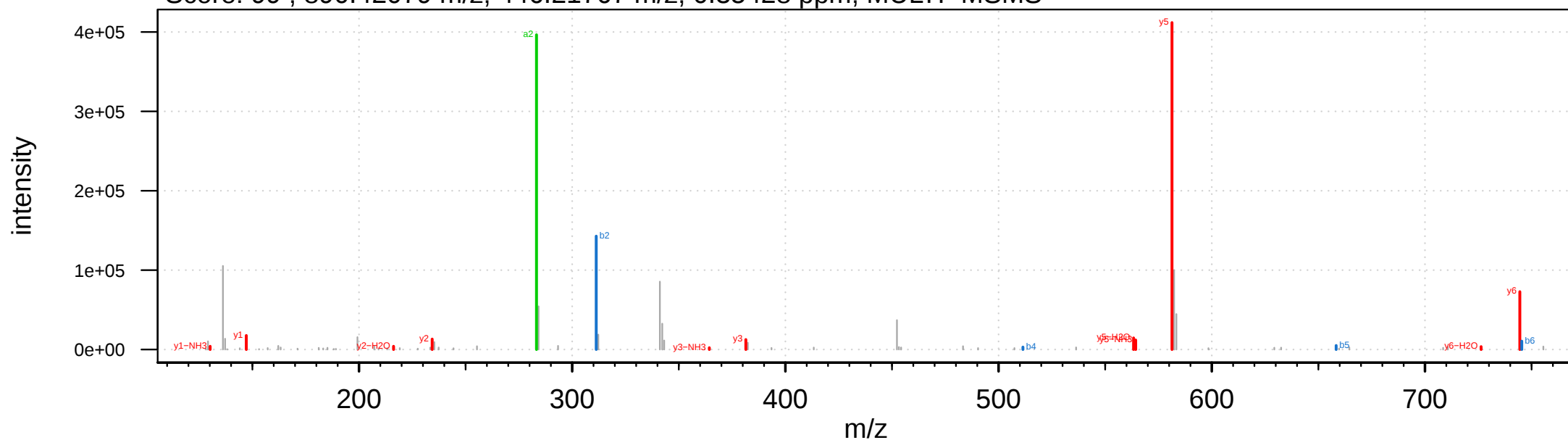
Proteins:

ENST00000392145_chr6:160874572-160874622:+



M(ox)YLSFSK

Score: 99 ; 890.42079 m/z; 446.21767 m/z; 0.35428 ppm; MULTI-MSMS

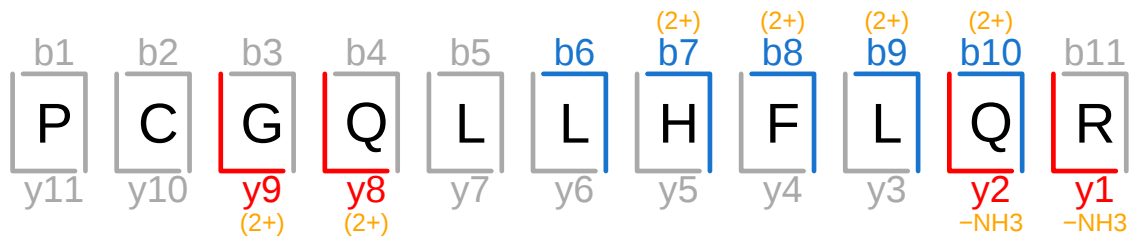


Raw File: Toni_20111125_FB_MCF7_F7_2

Scan Number: 27408

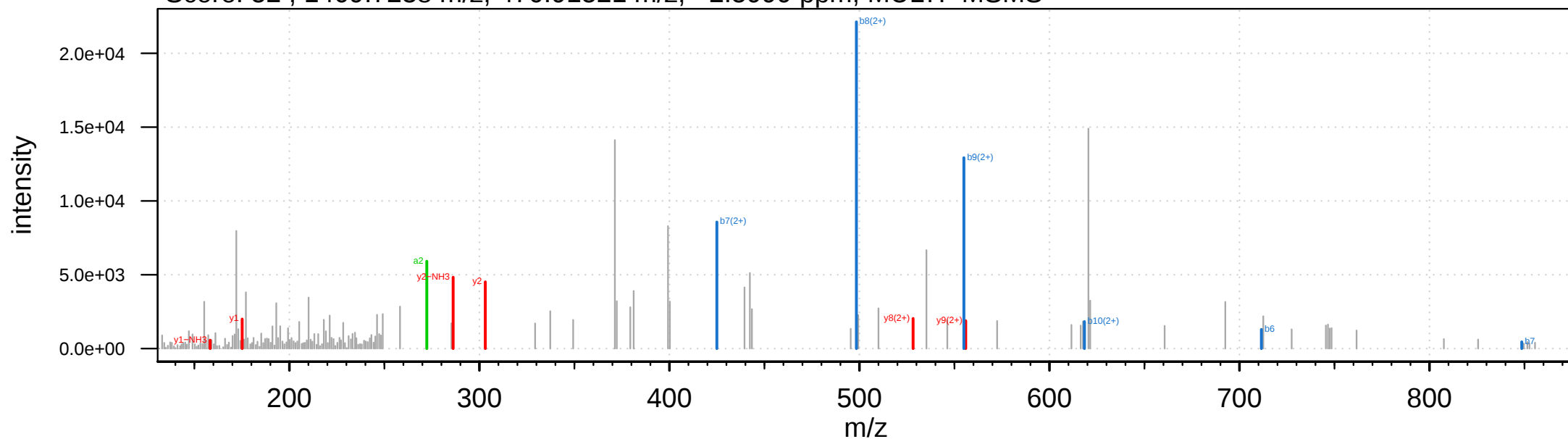
Proteins:

ENST00000392145_chr6:160874572-160874622:+



(ac)PCGQLLHFLQR

Score: 52 ; 1409.7238 m/z; 470.91521 m/z; -2.5999 ppm; MULTI-MSMS

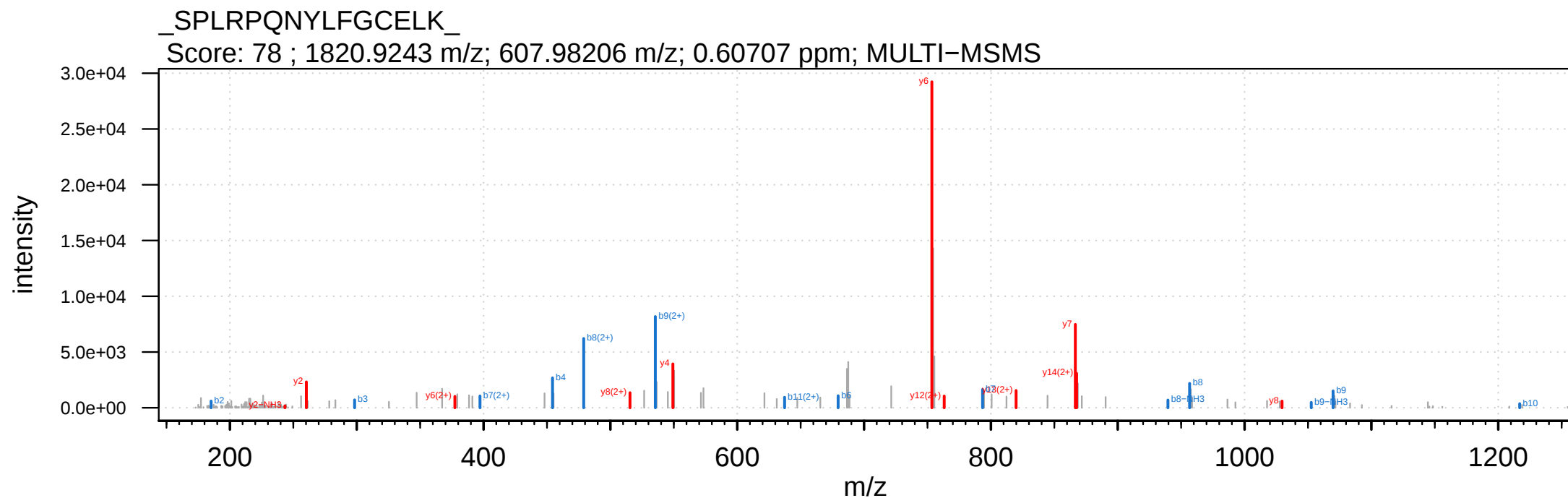
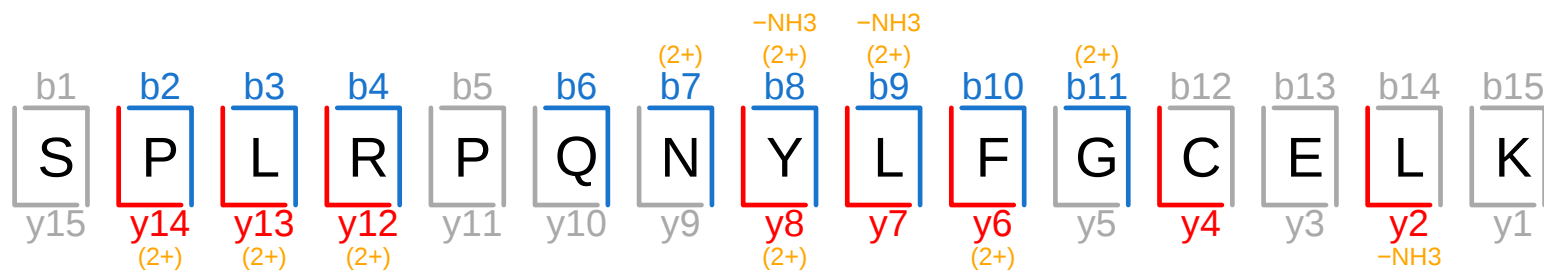


Raw File: Toni_20111109_FB_10A_F8_2

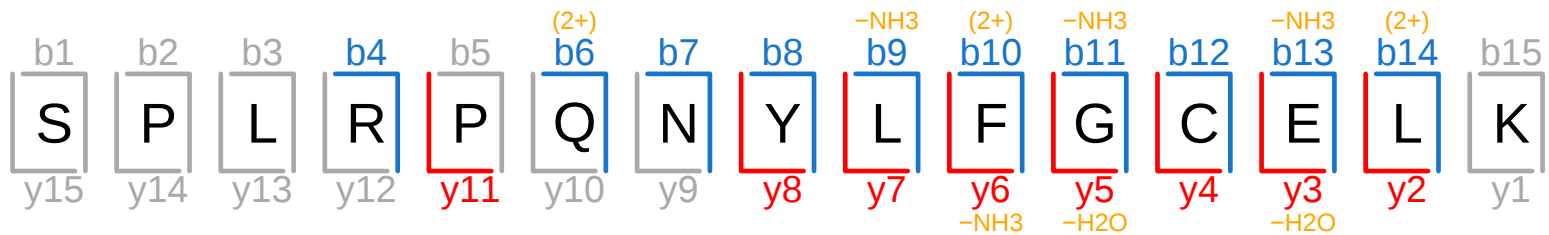
Scan Number: 42791

Proteins:

ENST00000605331_chr2:98317634-98318539:+

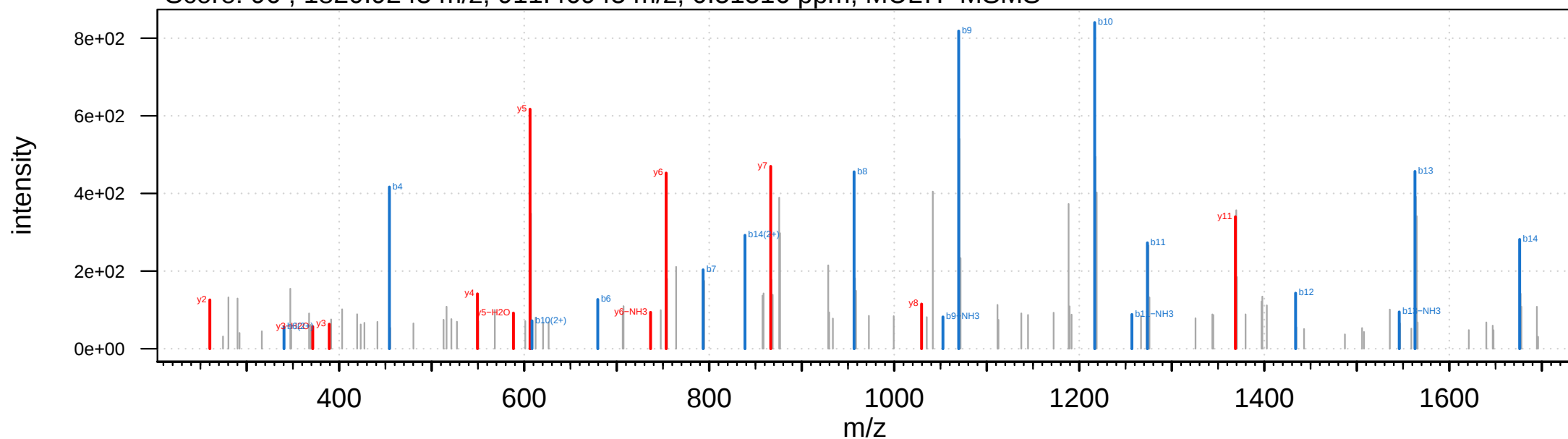


Raw File: Toni_20111109_FB_10A_F9_1
 Scan Number: 37732
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 96 ; 1820.9243 m/z; 911.46945 m/z; 0.51516 ppm; MULTI-MSMS

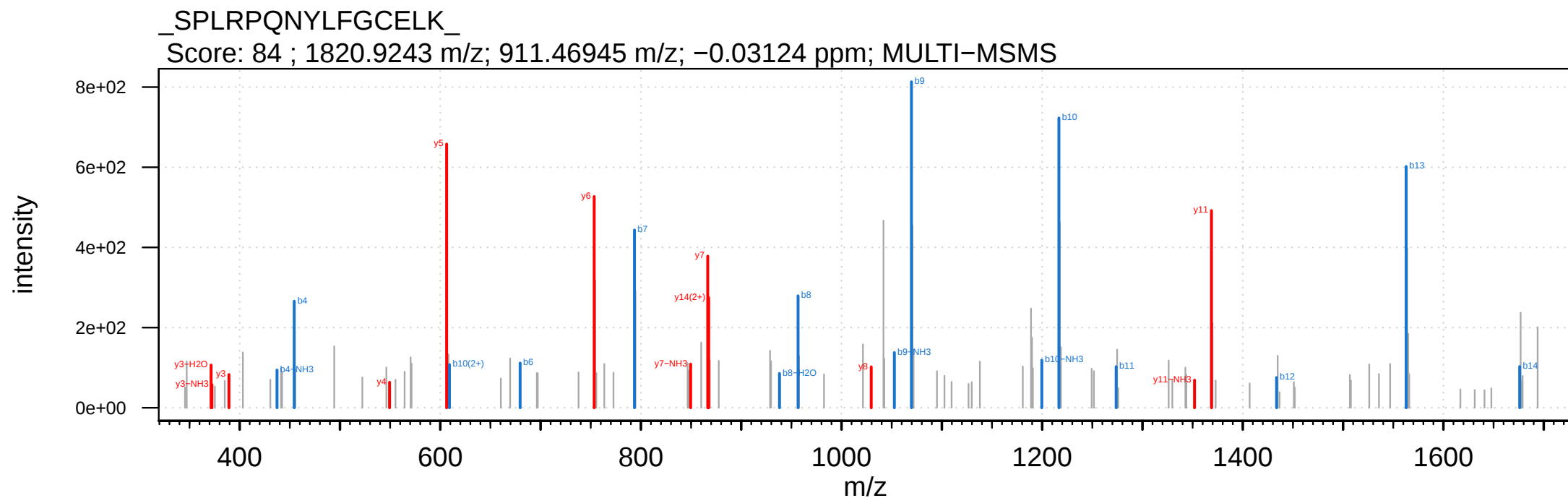
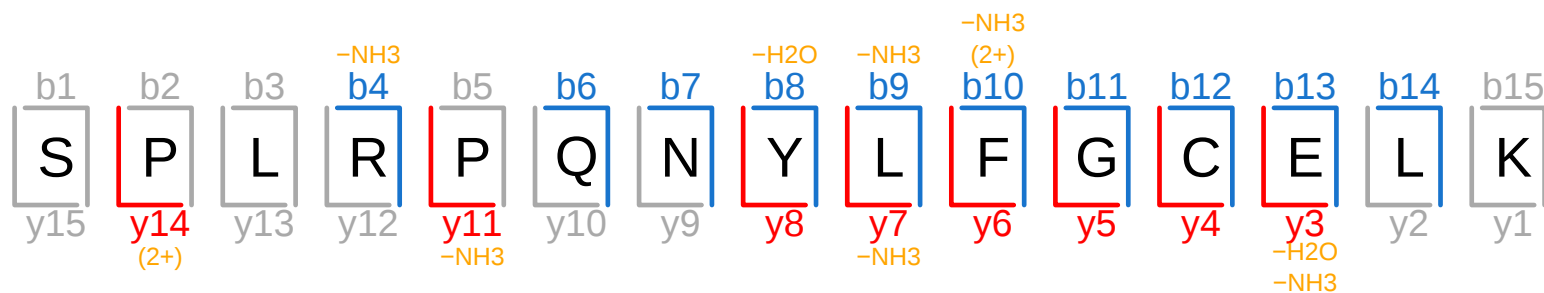


Raw File: Toni_20111109_FB_10A_F9_1

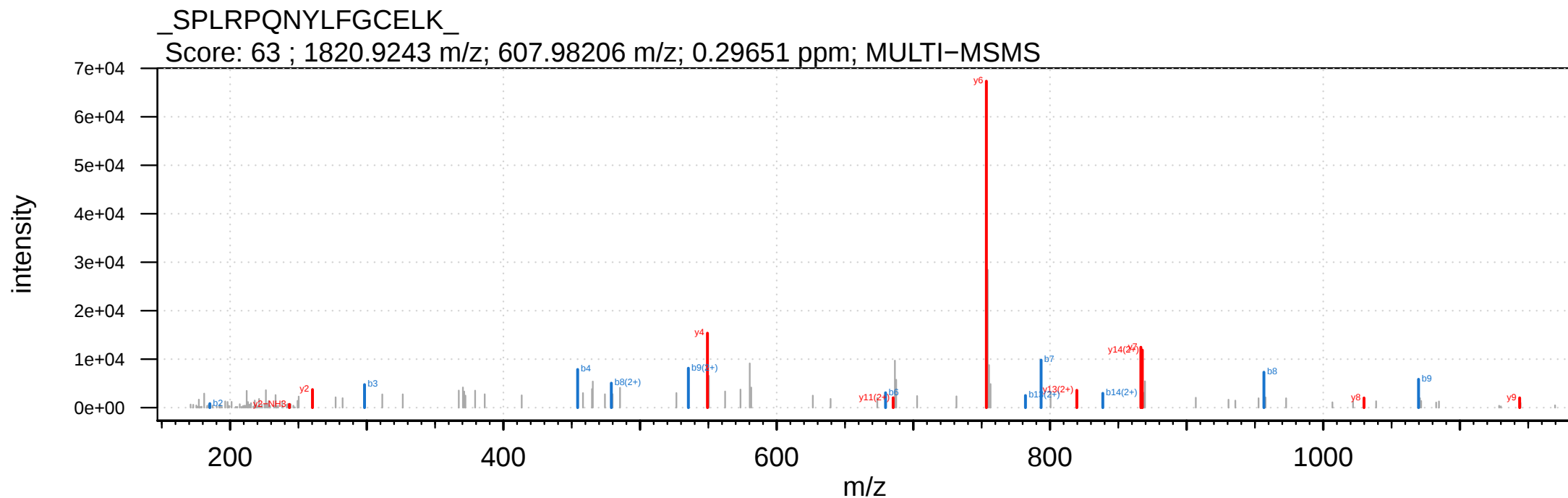
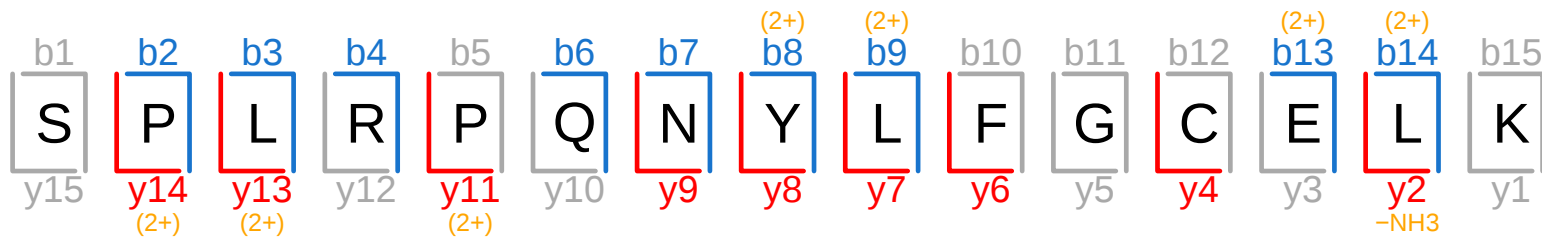
Scan Number: 37764

Proteins:

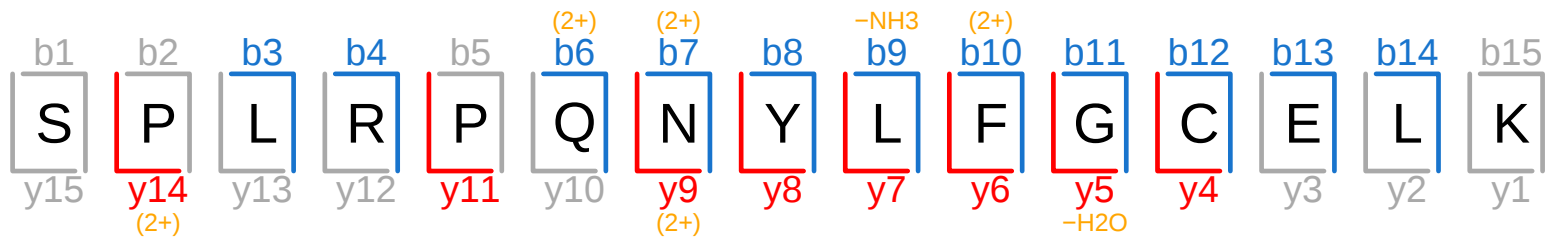
TCONS_I2_00008829_chr15:92829088-92829258:+



Raw File: Toni_20111109_FB_10A_F9_2
 Scan Number: 37890
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+

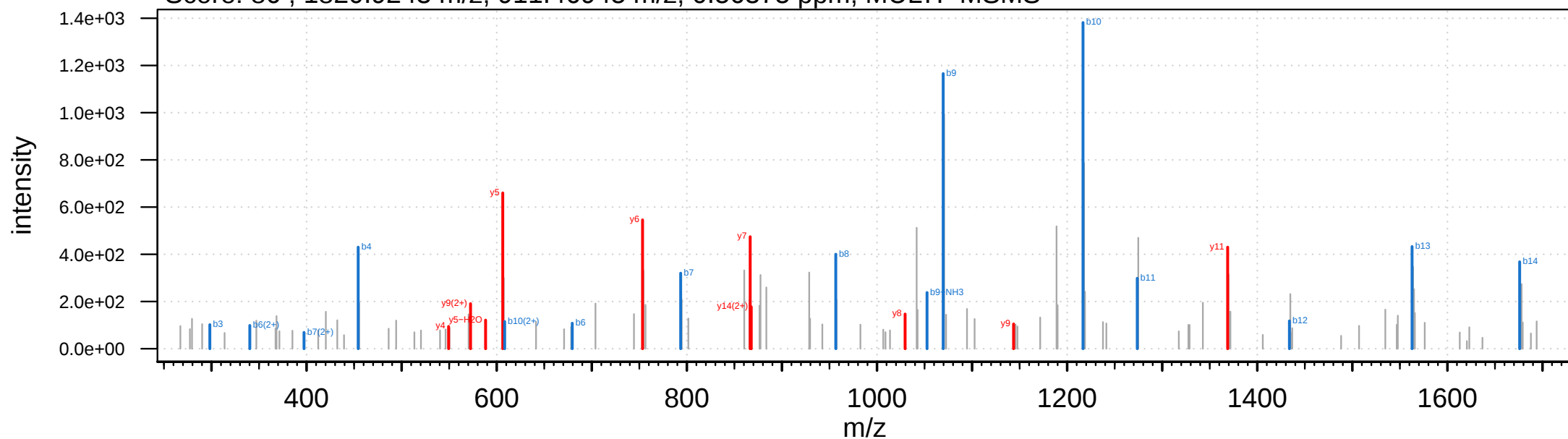


Raw File: Toni_20111109_FB_10A_F10_1
 Scan Number: 38286
 Proteins:
 TCONS_I2_00008829_chr15:92829088-92829258:+



SPLRPQNYLFGCELK

Score: 86 ; 1820.9243 m/z; 911.46945 m/z; 0.56375 ppm; MULTI-MSMS

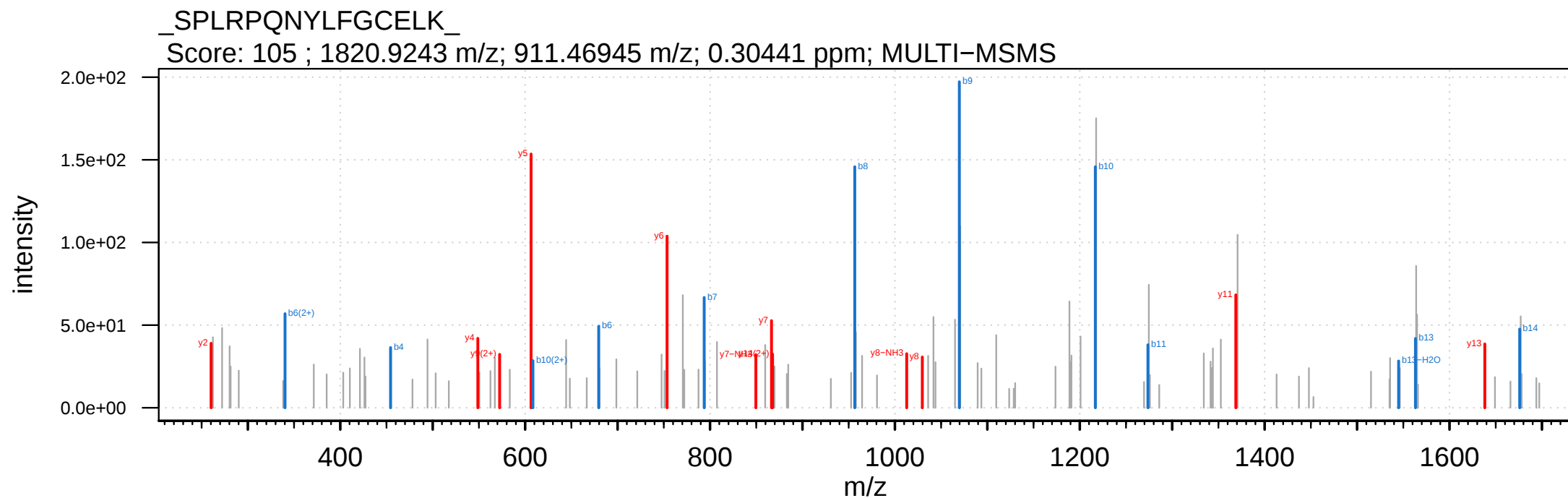
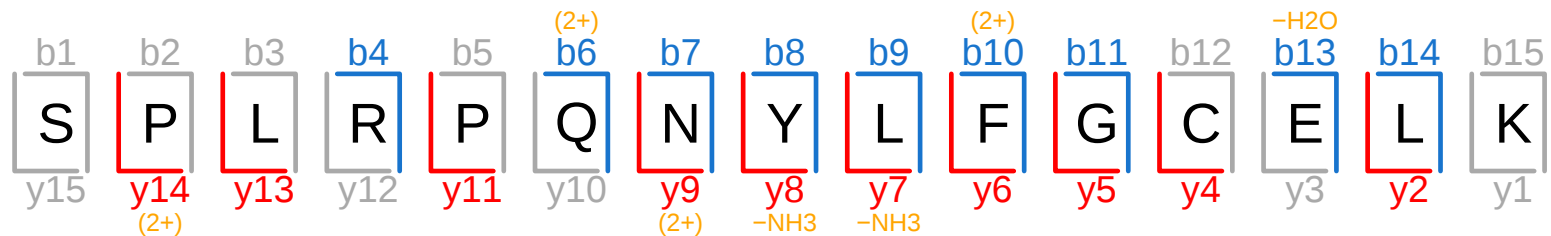


Raw File: Toni_20111109_FB_10A_F10_1

Scan Number: 38299

Proteins:

TCONS_I2_00008829_chr15:92829088-92829258:+



Raw File: Toni_20111114_FB_MDA-MB-231_F10_2
Scan Number: 41088
Proteins:
TCONS_I2_00008829_chr15:92829088-92829258:+