

PSM scan report

20120906_balliau_extract_1_A01_urnb-1

Scan index=5737 mh=1240.4975 mz=620.7524 charge=2
rt=1306.13s

[+132.97]?KEMAPDNFR

GRMZM2G059083_P02

+221 YTGLAWLFLGVPTSDTLLYKEELEKMKEMAPDNFRLDFAVSREQTNAAGEKMYIQTRMAE +280

GRMZM2G059083_P03

+247 YTGLAWLFLGVPTSDTLLYKEELEKMKEMAPDNFRLDFAVSREQTNAAGEKMYIQTRMAE +306

GRMZM2G059083_P04

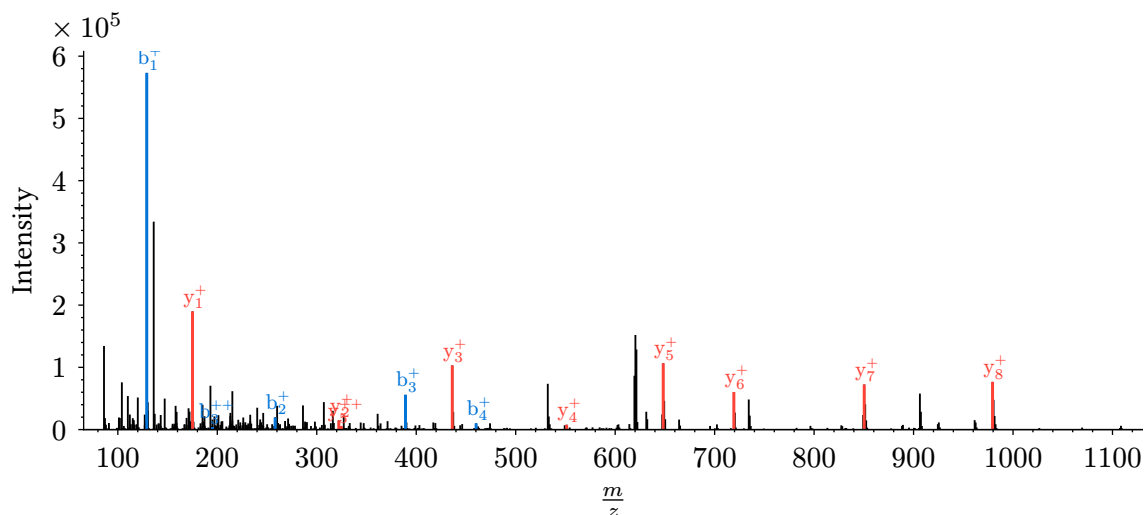
+247 YTGLAWLFLGVPTSDTLLYKEELEKMKEMAPDNFRLDFAVSREQTNAAGEKMYIQTRMAE +306

GRMZM2G059083_P05

+0 MKEMAPDNFRLDFAVSREQTNAAGEKMYIQTRMAEYKEELWELLKKDNTYVYMCGLKGME +59

GRMZM2G059191_P01

+248 FSGLAWLFLGVPTSDSLLYKEELEKMKEMAPDNFRLDFAVSREQTNAAGEKMYIQTRMAE +307



hyperscore 39.98

peptide length 9

peptide mh 1107.52515

charge 2

Δ mz 66.48617

Δ mh 132.97233

Spoms

isotope error 0

bracket [261.07]EMA[P][D]NFR

nam -0.000871082298544934

spc 8

score 42

Spoms post

bracket KEMA[P][D]NFR

nam 132.97232888789586

spc 11

score 55