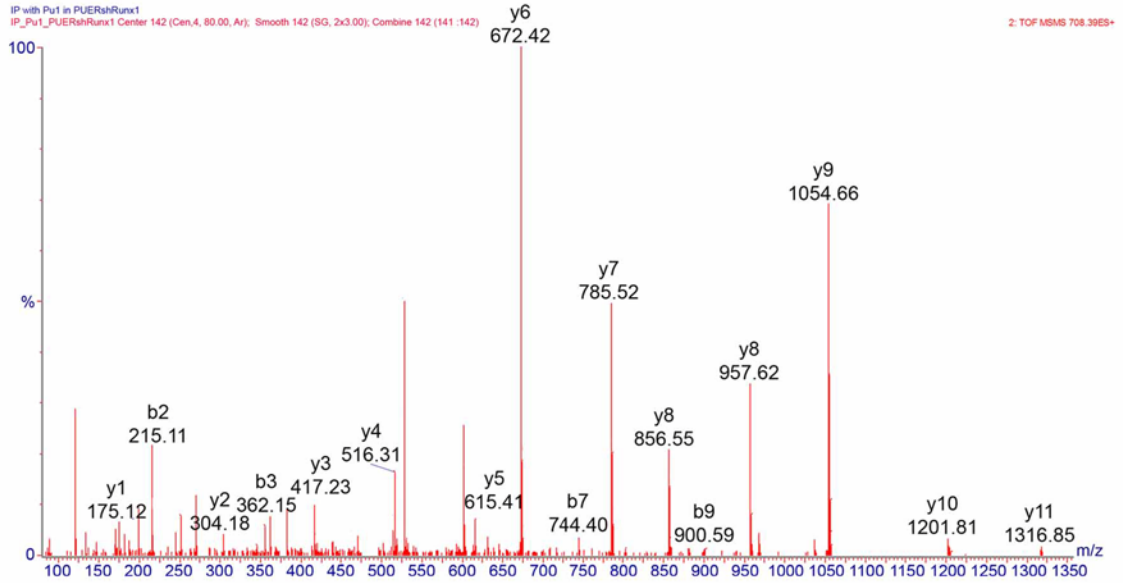


SUPPLEMENTARY MATERIAL (RUNX1 regulates PU.1 corepressor/coactivator exchange)

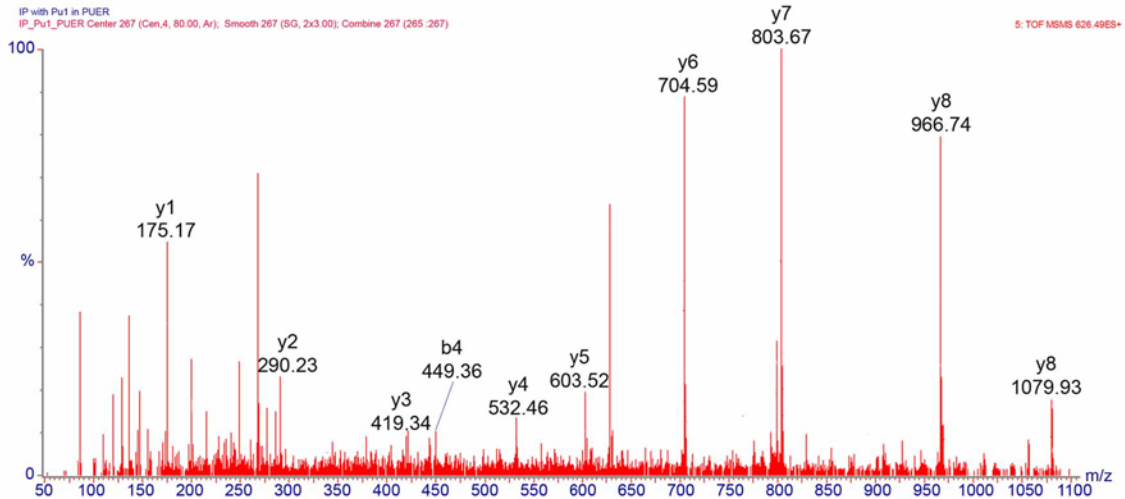
Contents:

Figure S1: Tandem mass spectrometry (MS/MS) spectrum of single peptide identification. Respective b and y'' ions are indicated.

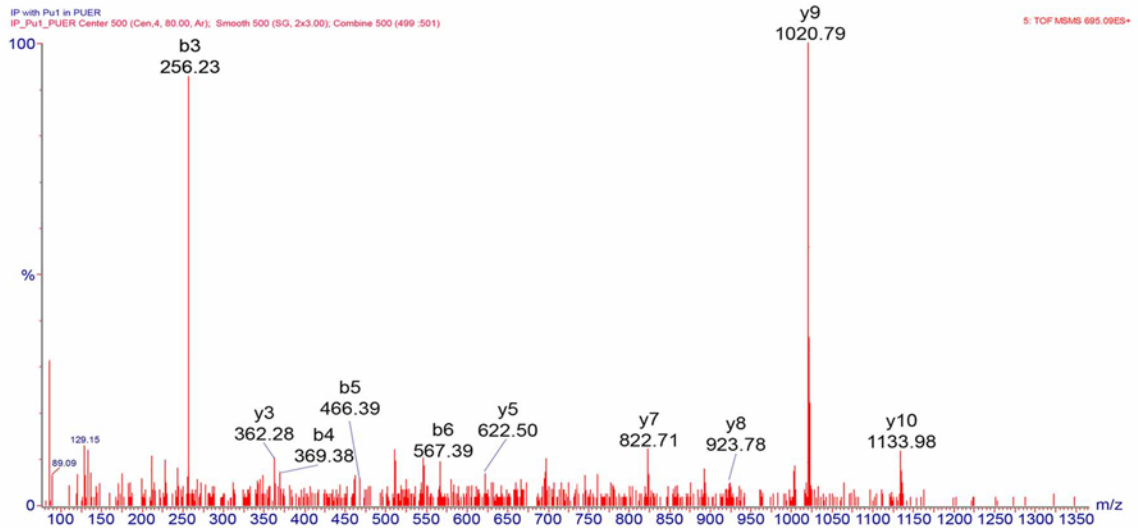
ACL6A IP exp3 PUERshRUNX1 VDFPTAIGVVLER, m/z 708.39 (2+)



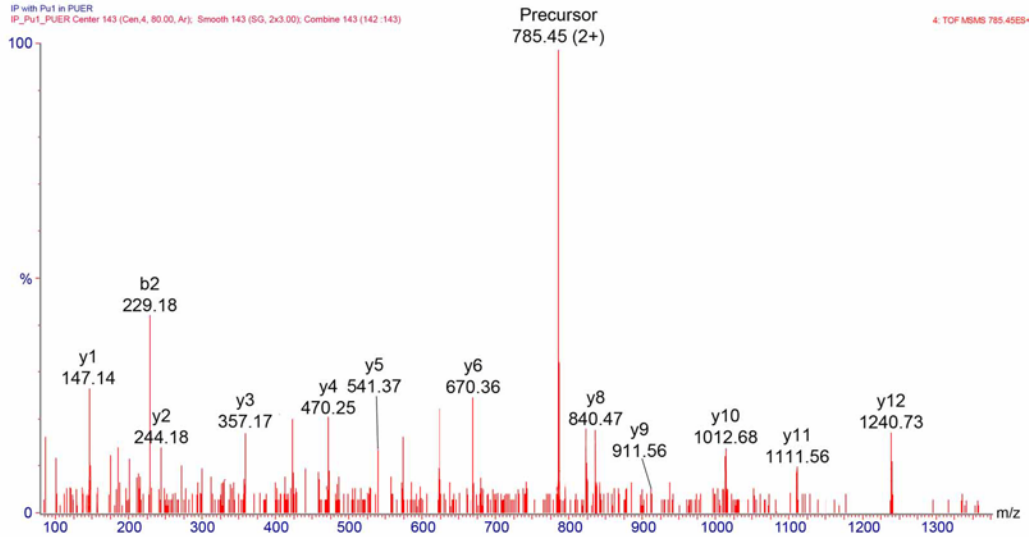
ANM1 IP exp1 PUER ATLYVTAIEDR, m/z 626.49 (2+)



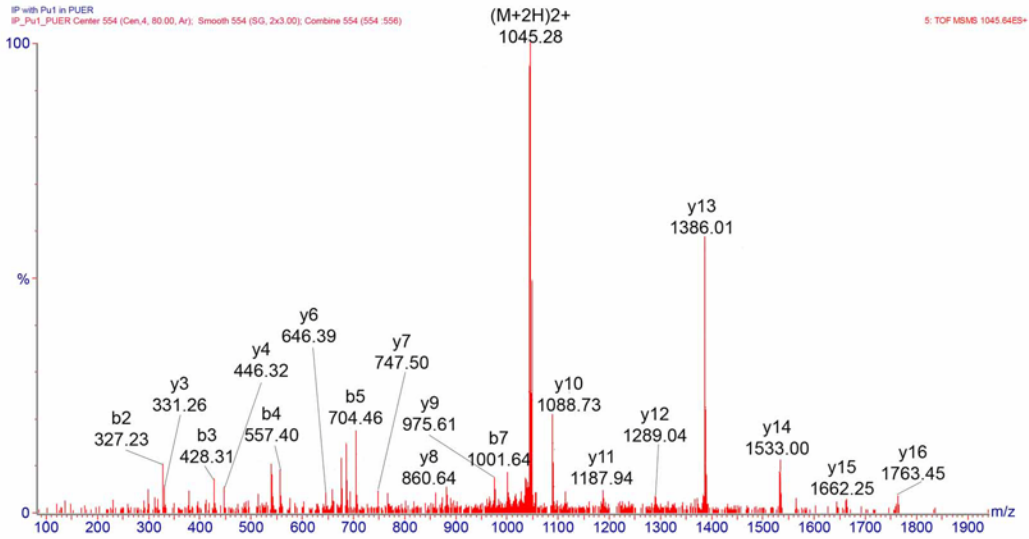
ANM5 IP exp1 PUER AAILPTSIFLTNK, m/z 695.09 (2+)



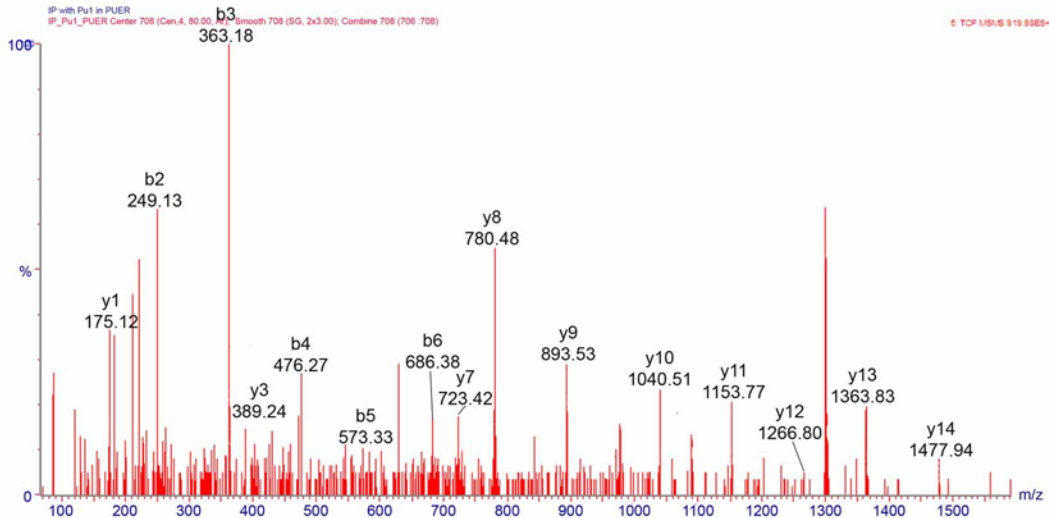
EP400 IP exp3 PUER DITEVTAVAEAILPK, m/z 785.45 (2+)



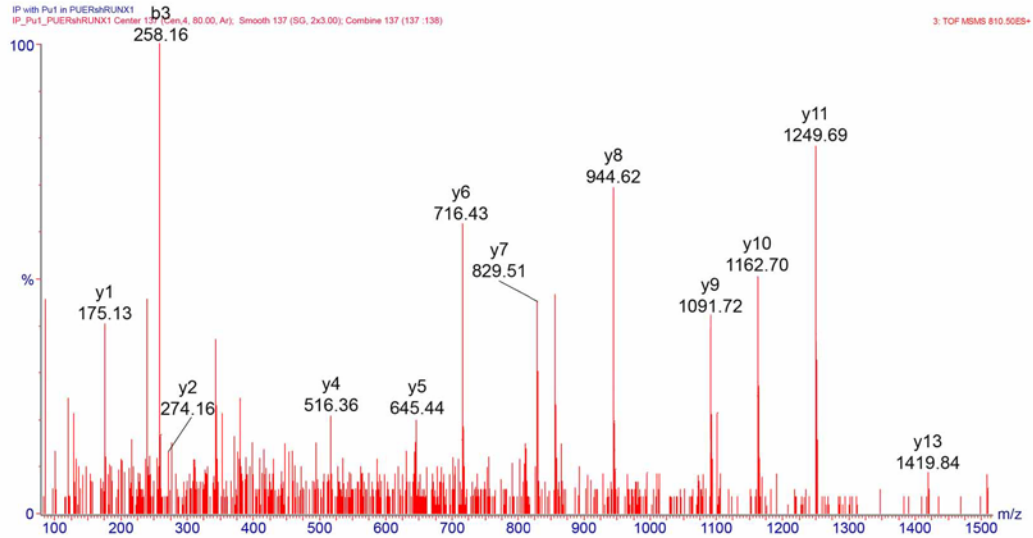
HAT1 IP exp1 PUER YYTEFPTVLDITAEDPSK, m/z 1045.28 (2+)



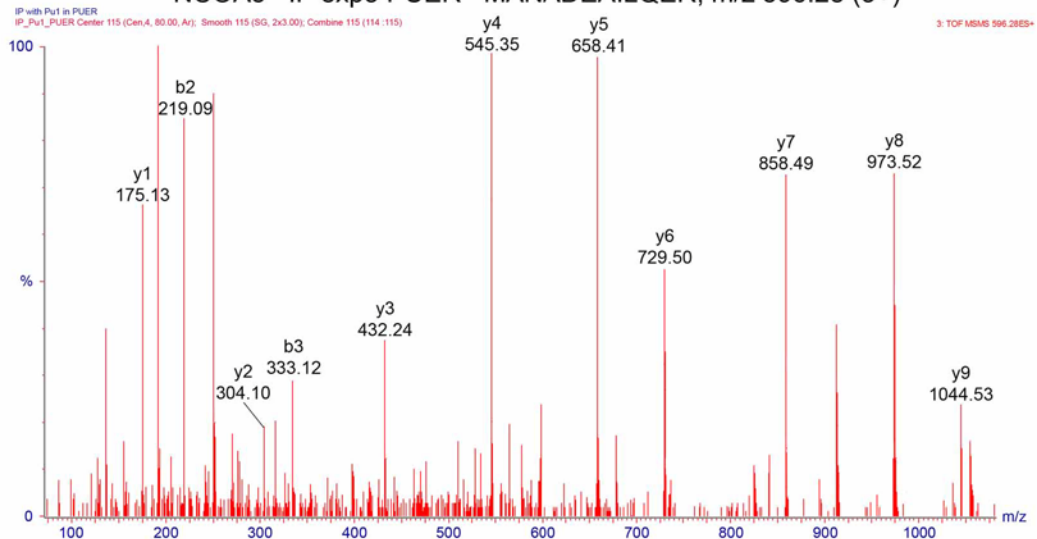
HDAC2 IP exp1 PUER TFNLPLLM(ox)LGGGGYTIR, m/z 919.98



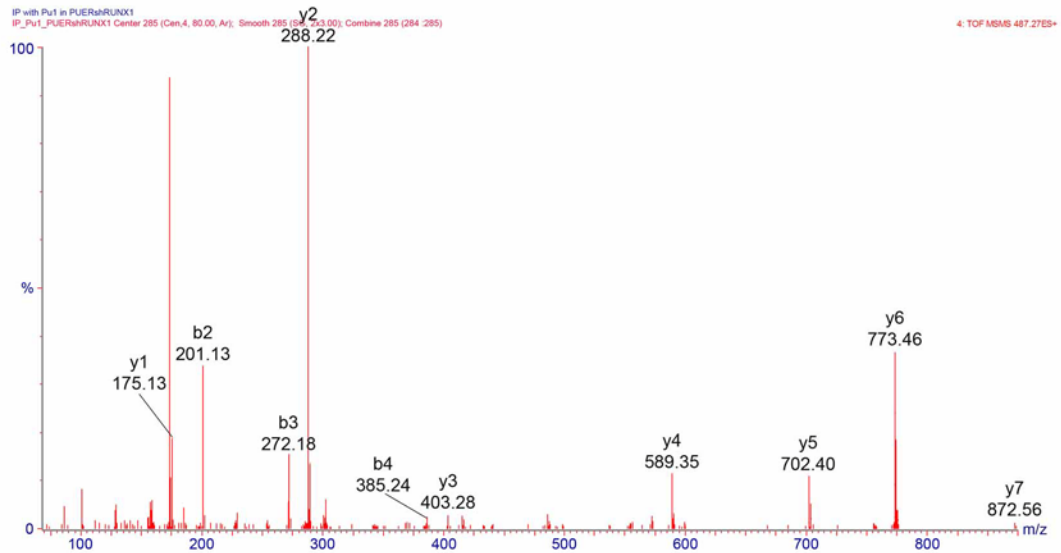
MBD3 IP exp3 PUERshRUNX1 LSGLSAFDIAEELVR, 810.50 (2+)



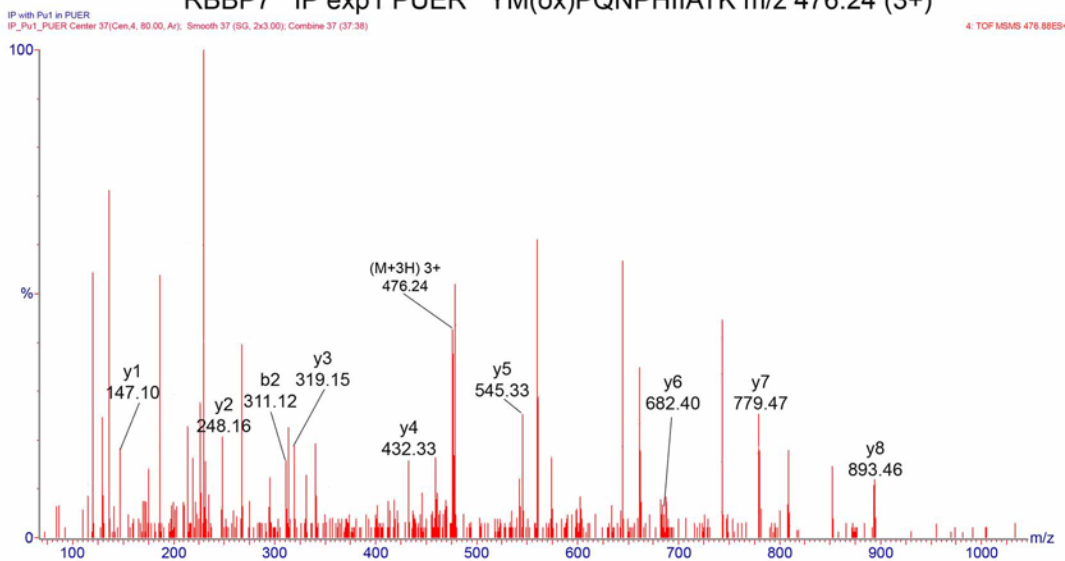
NCOA5 IP exp3 PUER MANADEAILQER, m/z 596.28 (3+)



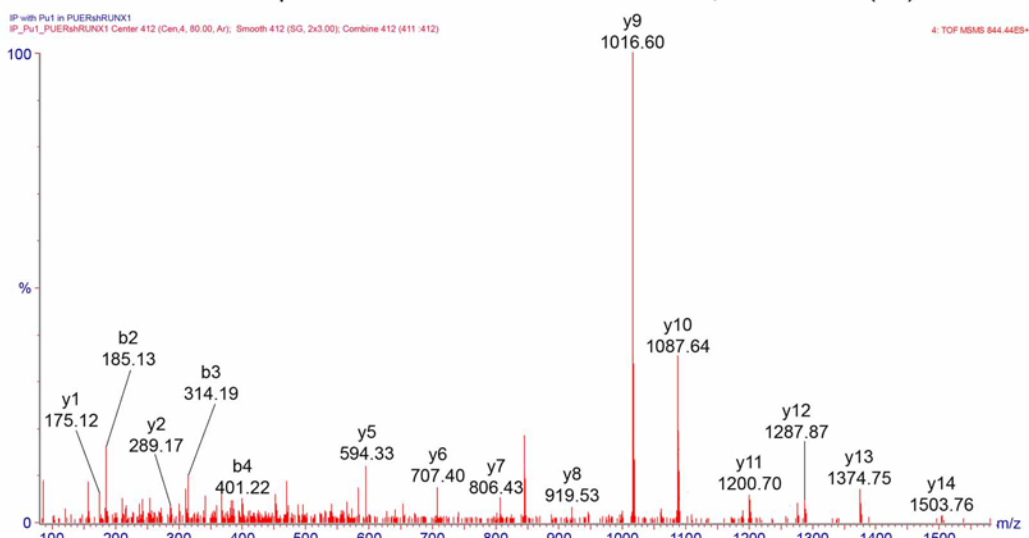
RBBP4 IP exp3 PUERshRUNX1 TVALWDLR, 487.27 (2+)



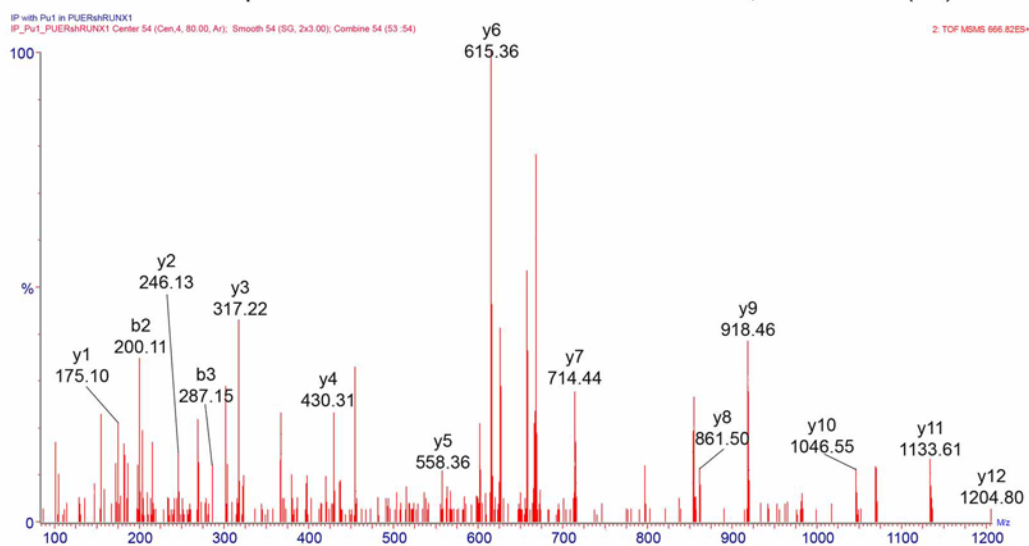
RBBP7 IP exp1 PUER YM(ox)PQNPHIATK m/z 476.24 (3+)



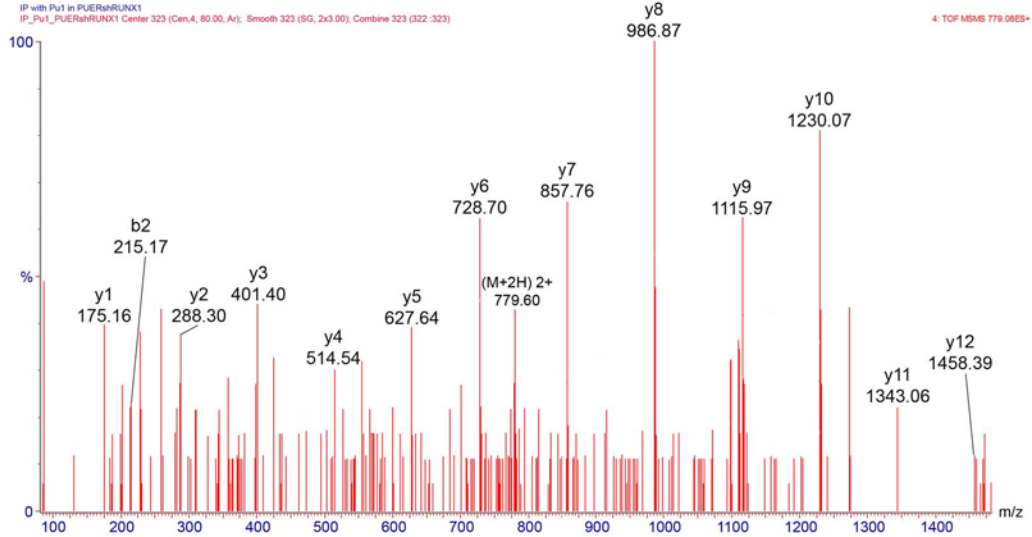
RUVB1 IP exp3 PUERshR ALESSIPIVIFASNR, m/z 844.44 (2+)



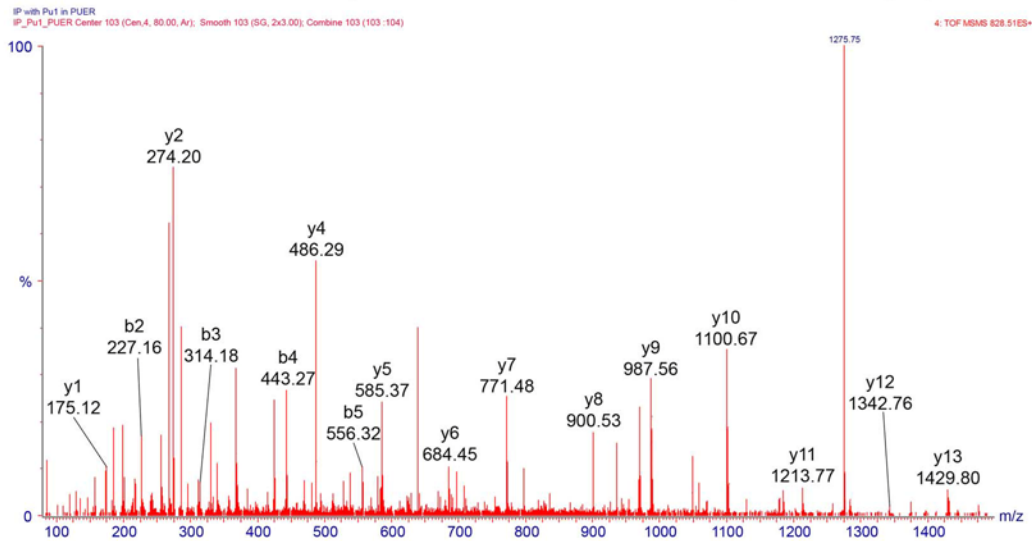
RUVB2 IP exp3 PUERshRUNX1 QASQGMVGQLAAR, m/z 666.82 (3+)



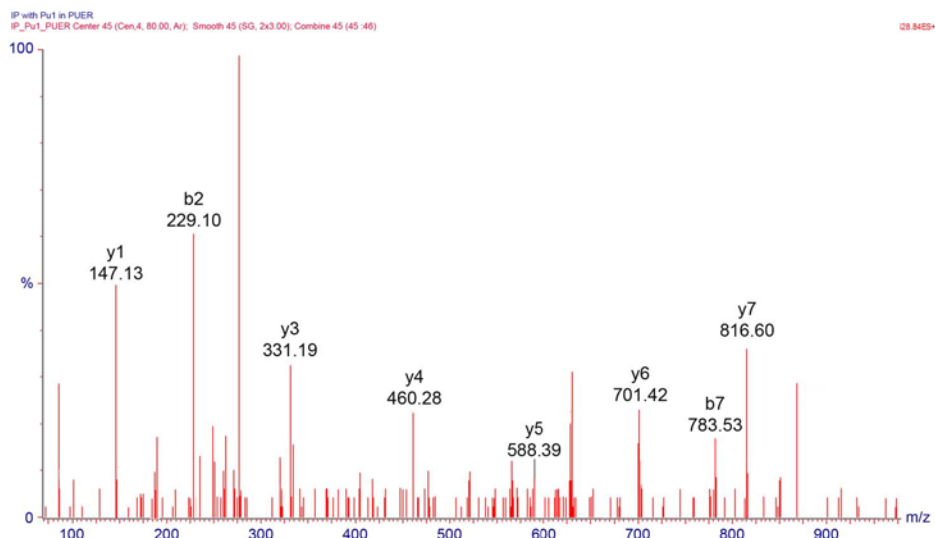
SMARCA2 IP exp3 PUER VDLNEEETILIIR, m/z 799.08 (2+)



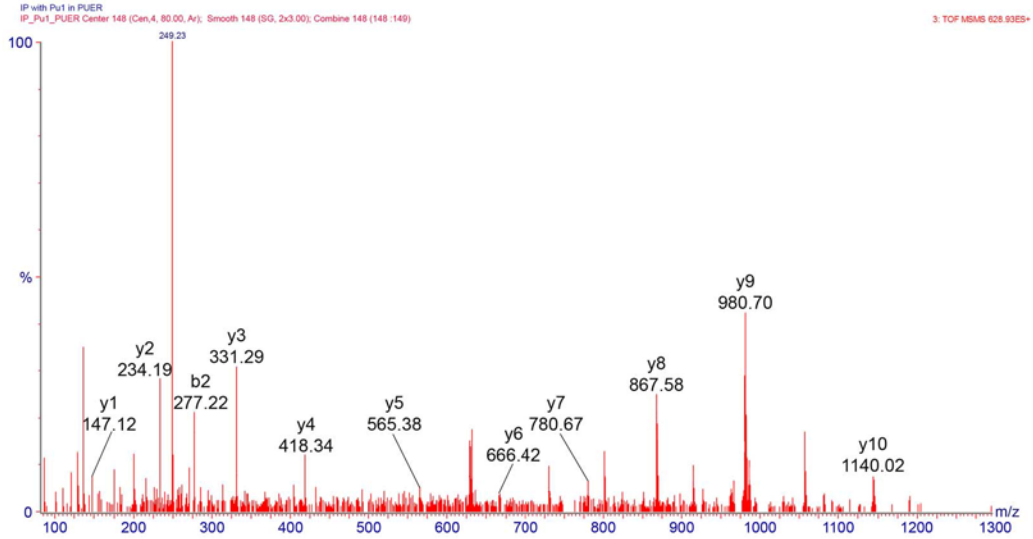
SMARCE1 IP exp1 PUER LISELSESVVPDVR, m/z 828.51 (2+)



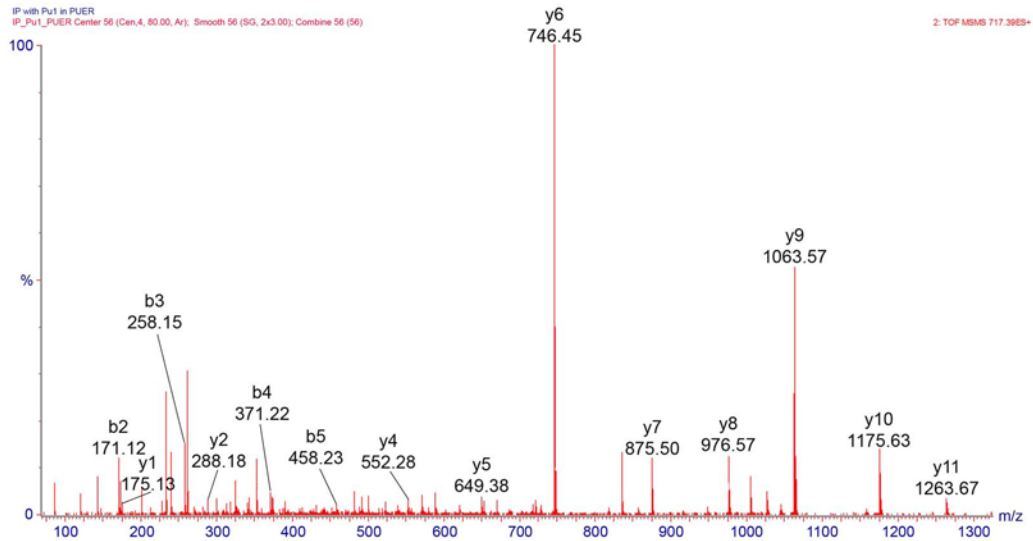
SMARCD1 IP exp1 PUER LDIQEALK, m/z 465.37 (2+)



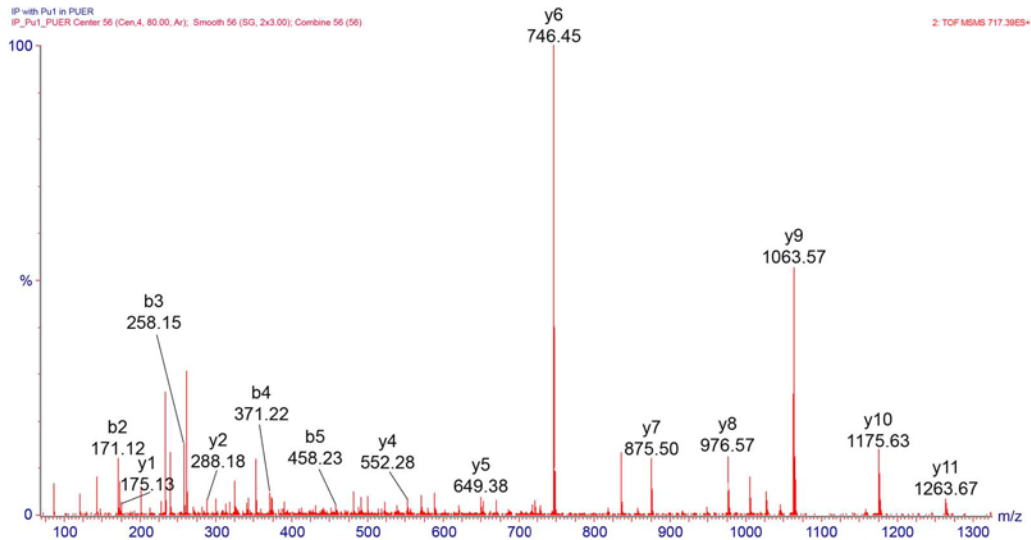
SMARCD2 IP exp1 PUER IYISNTFSPSK, m/z 628.93 (2+)



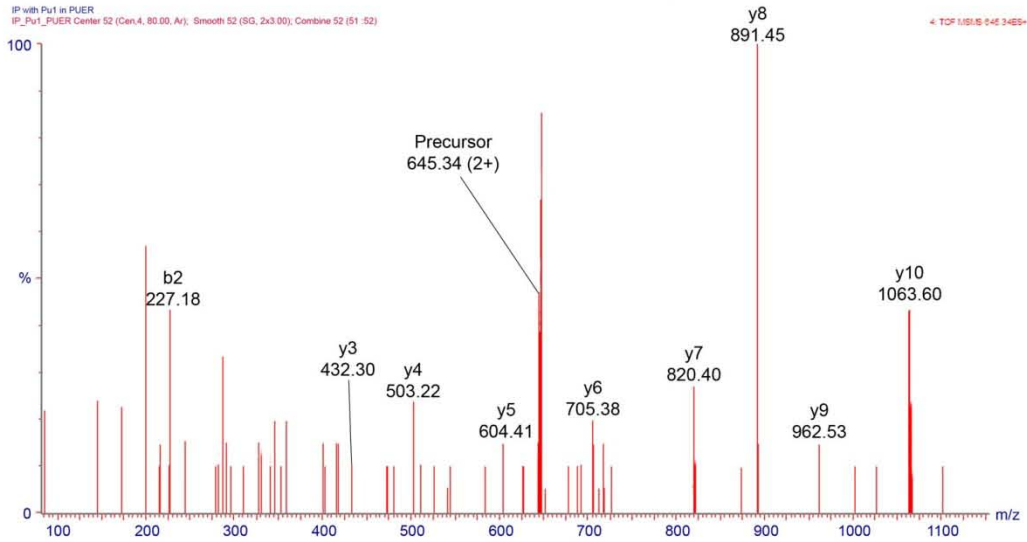
SNF5 IP exp3 PUER AVSISTEPPTYLR, m/z 717.39 (2+)



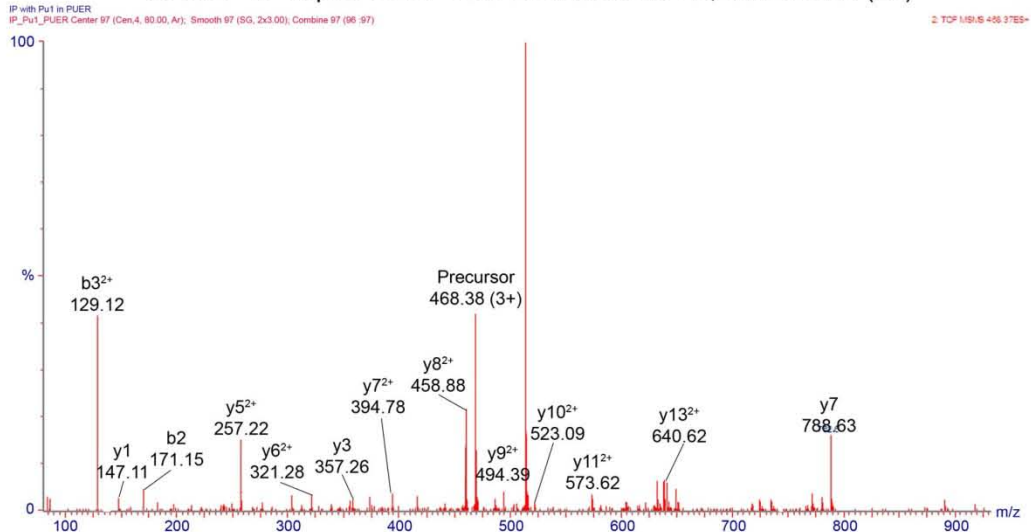
SNF5 IP exp3 PUER AVSISTEPPTYLR, m/z 717.39 (2+)



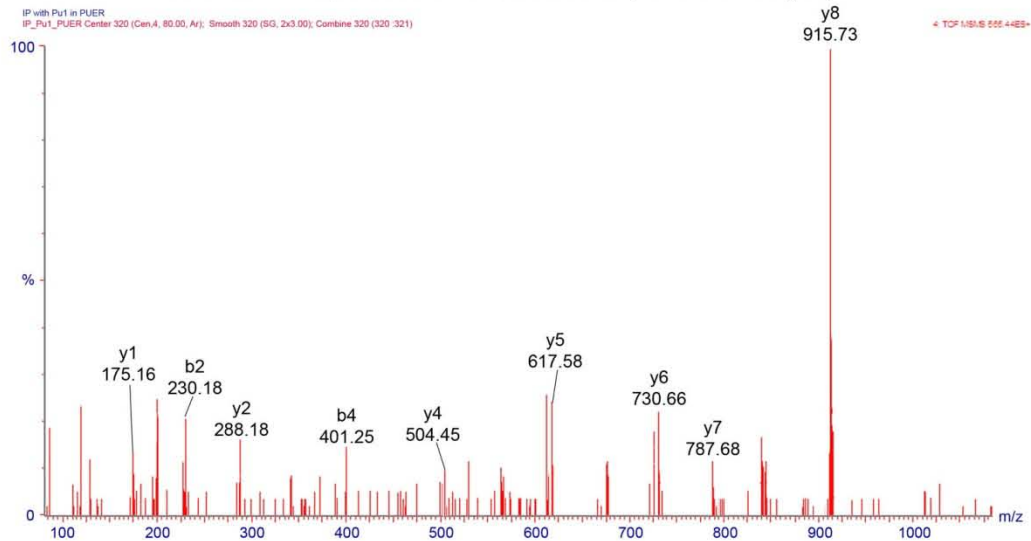
DMAP1 IP exp3 PUER LITAADTTAEQR, m/z 645.34 (2+)



SATB1 IP exp3 PUER LGSTGAKMQGVPLK, m/z 468.35 (2+)



SATB2 IP exp3 PUER TQGLLSEILR, m/z 461.34 (2+)



CARM1 IP exp1 PUER QPVVDTFDIR, m/z 595.45 (2+)

