

Experimental Spectrum Colour Code:

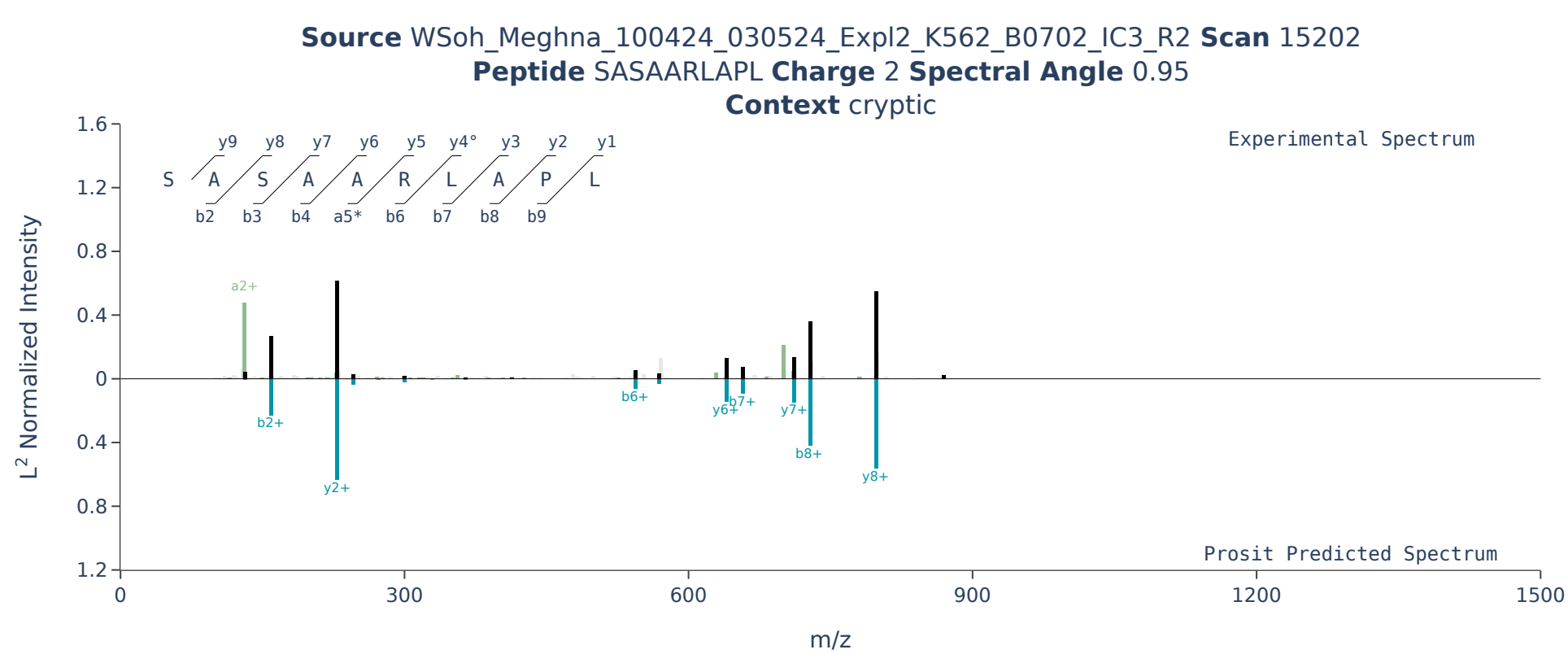
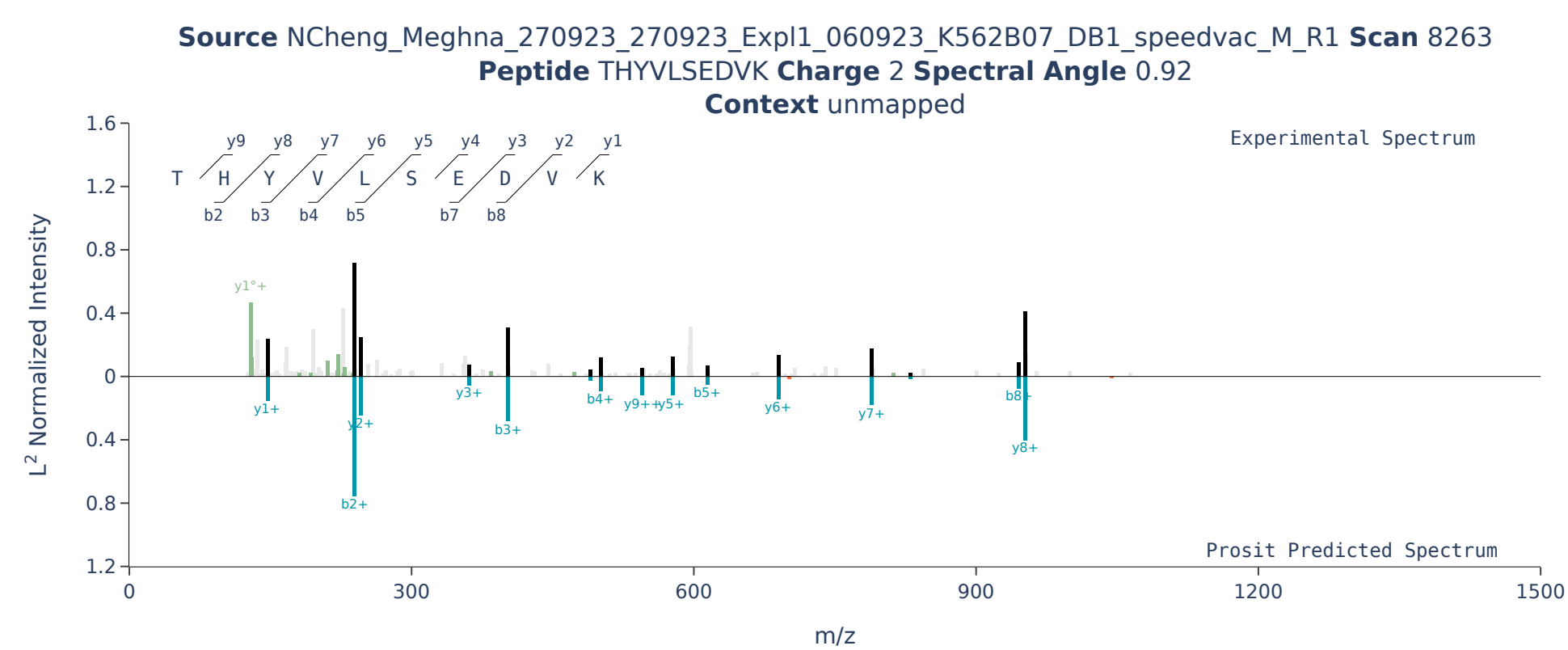
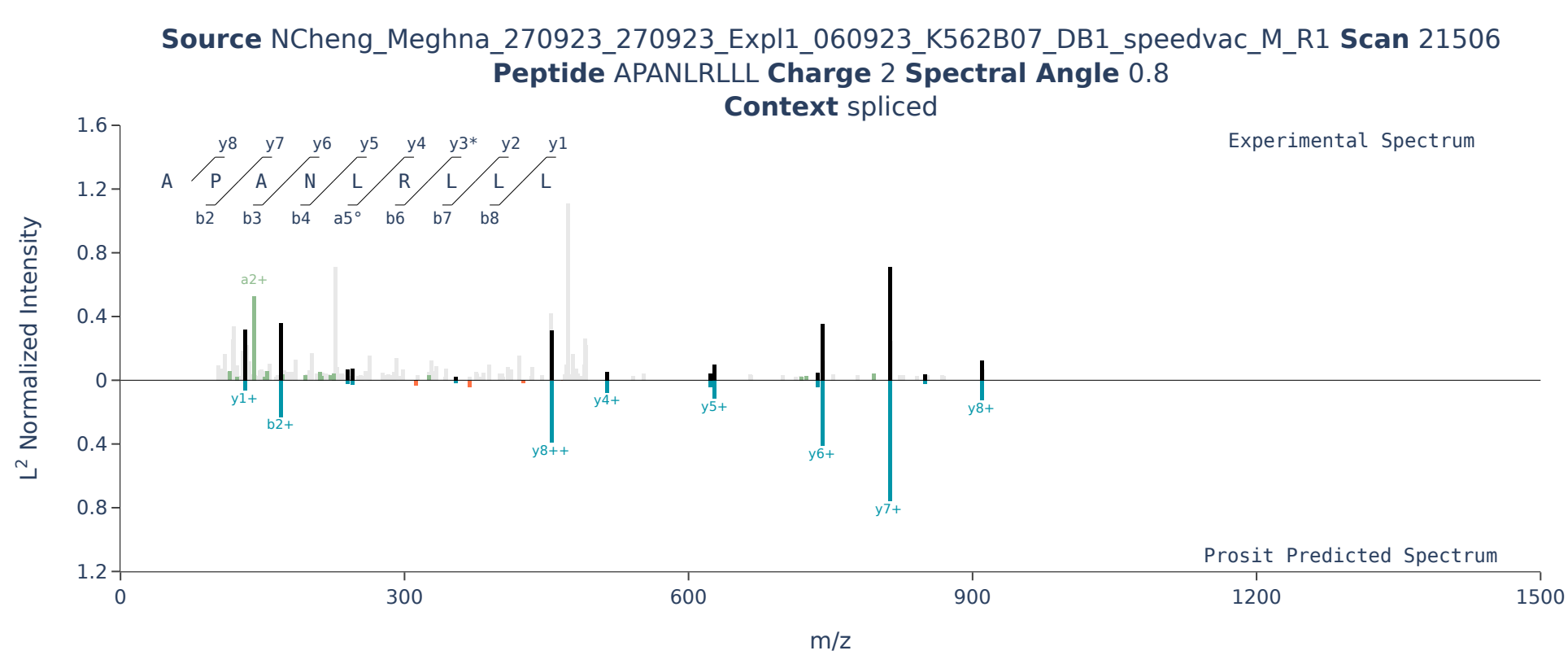
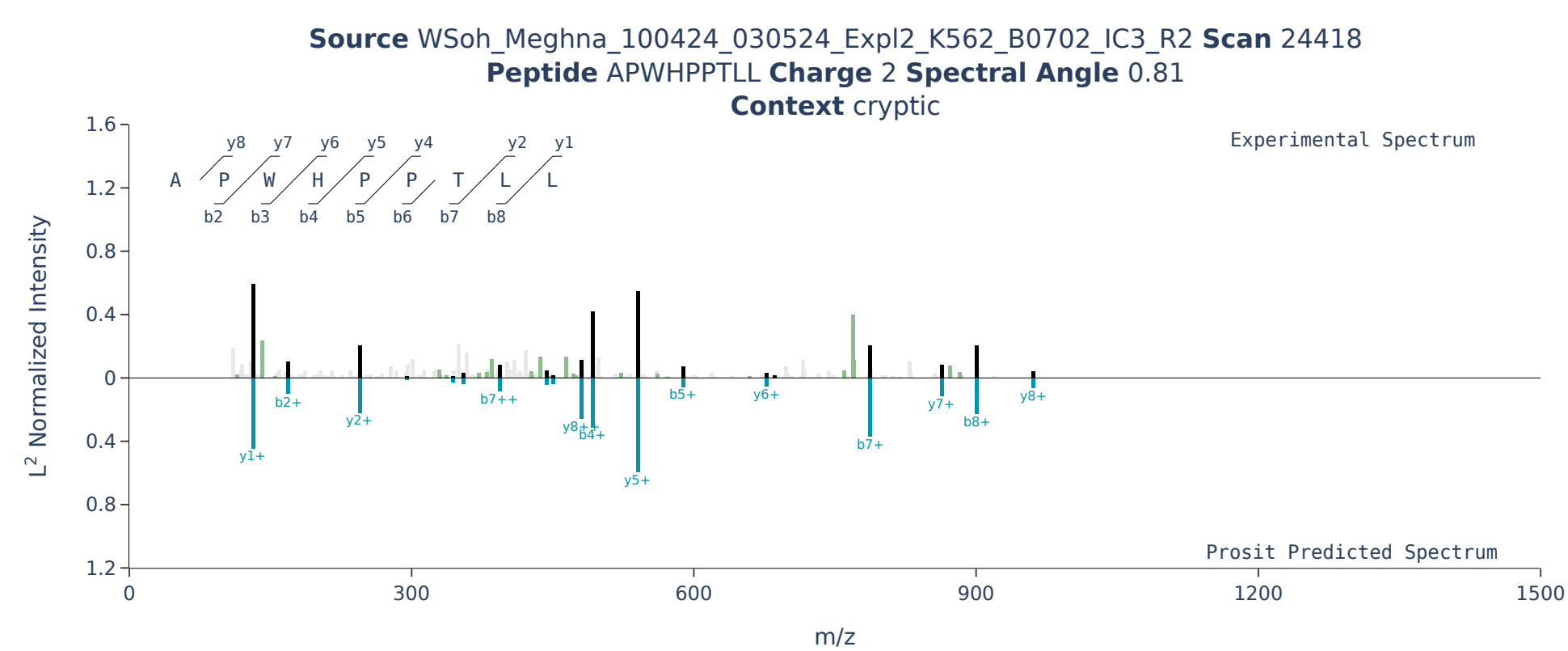
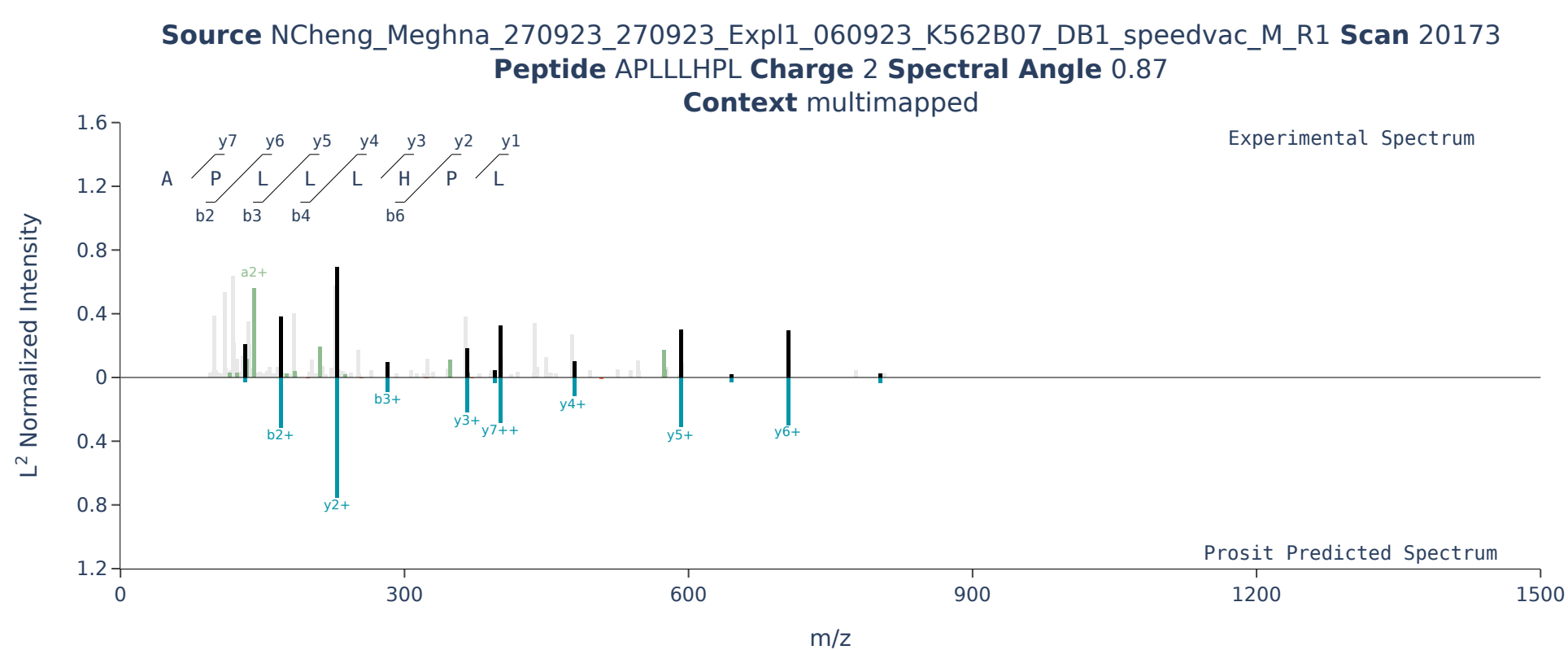
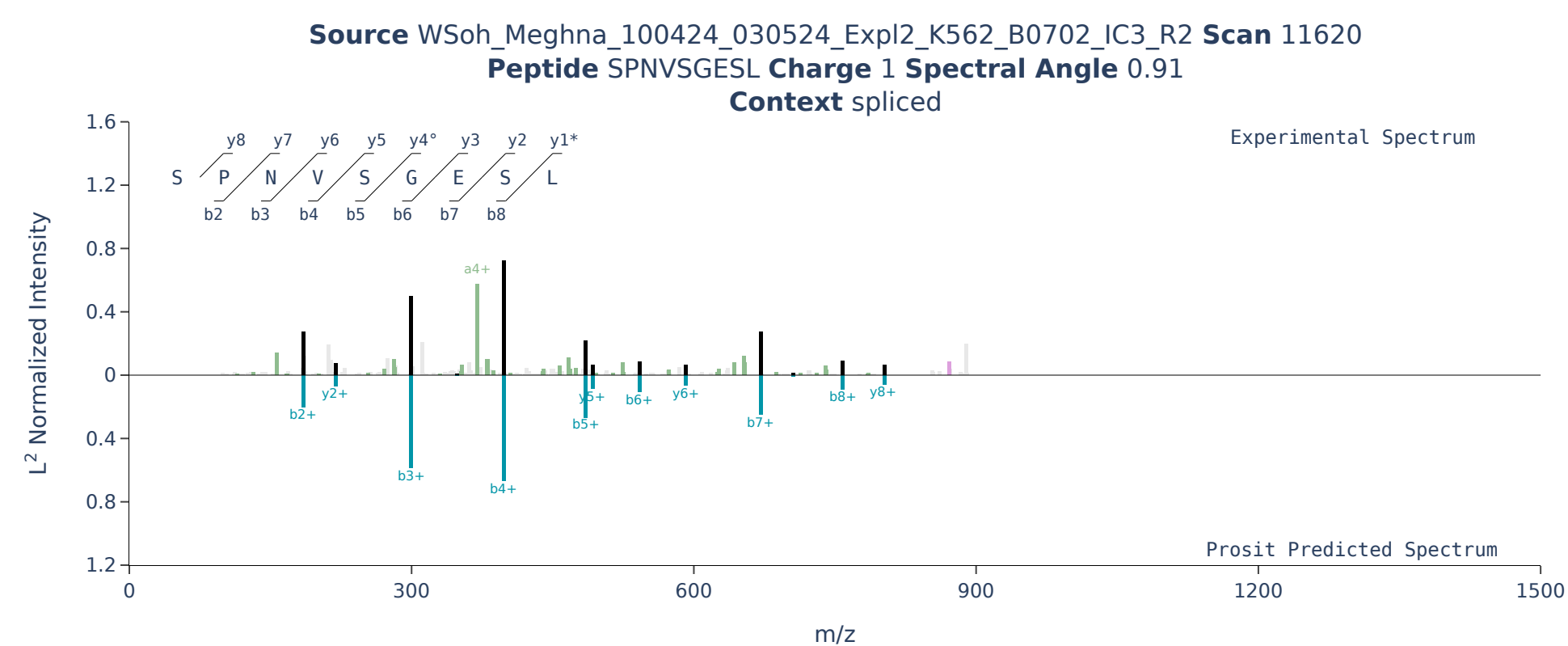
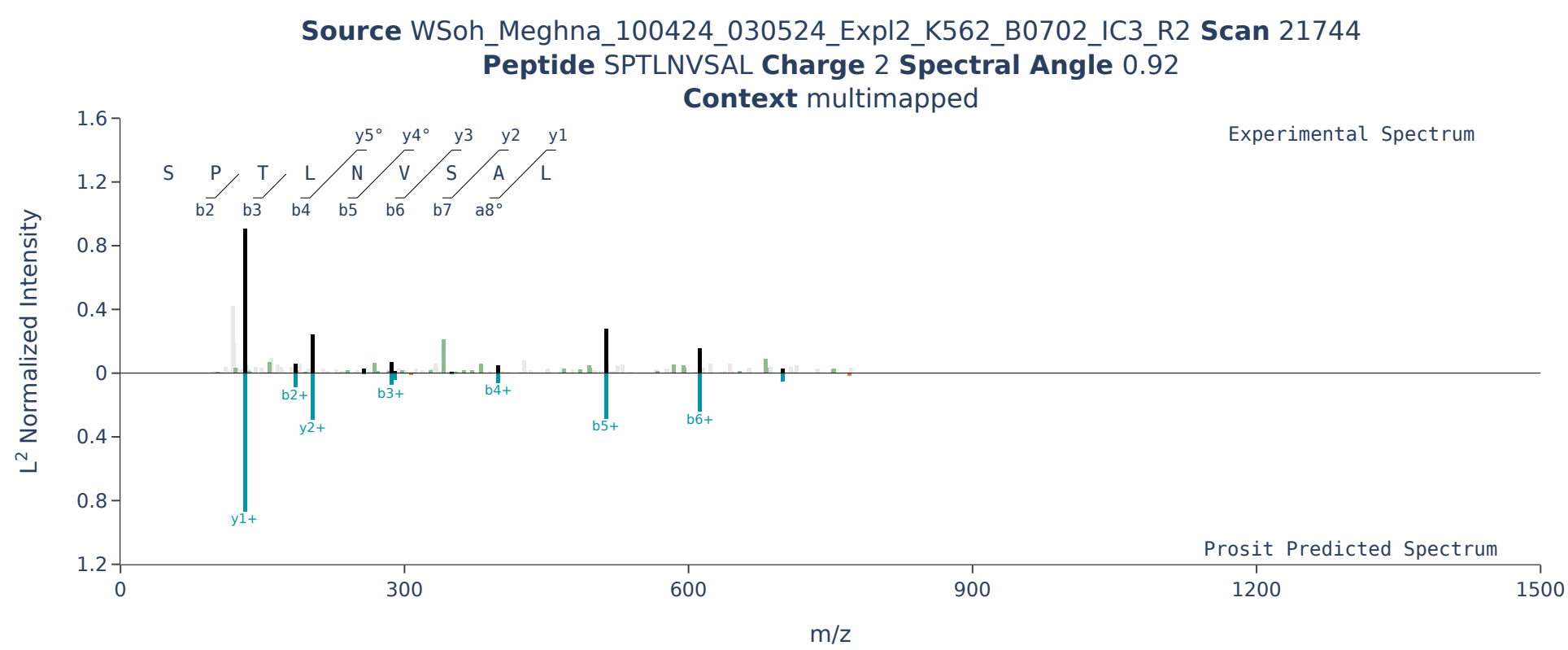
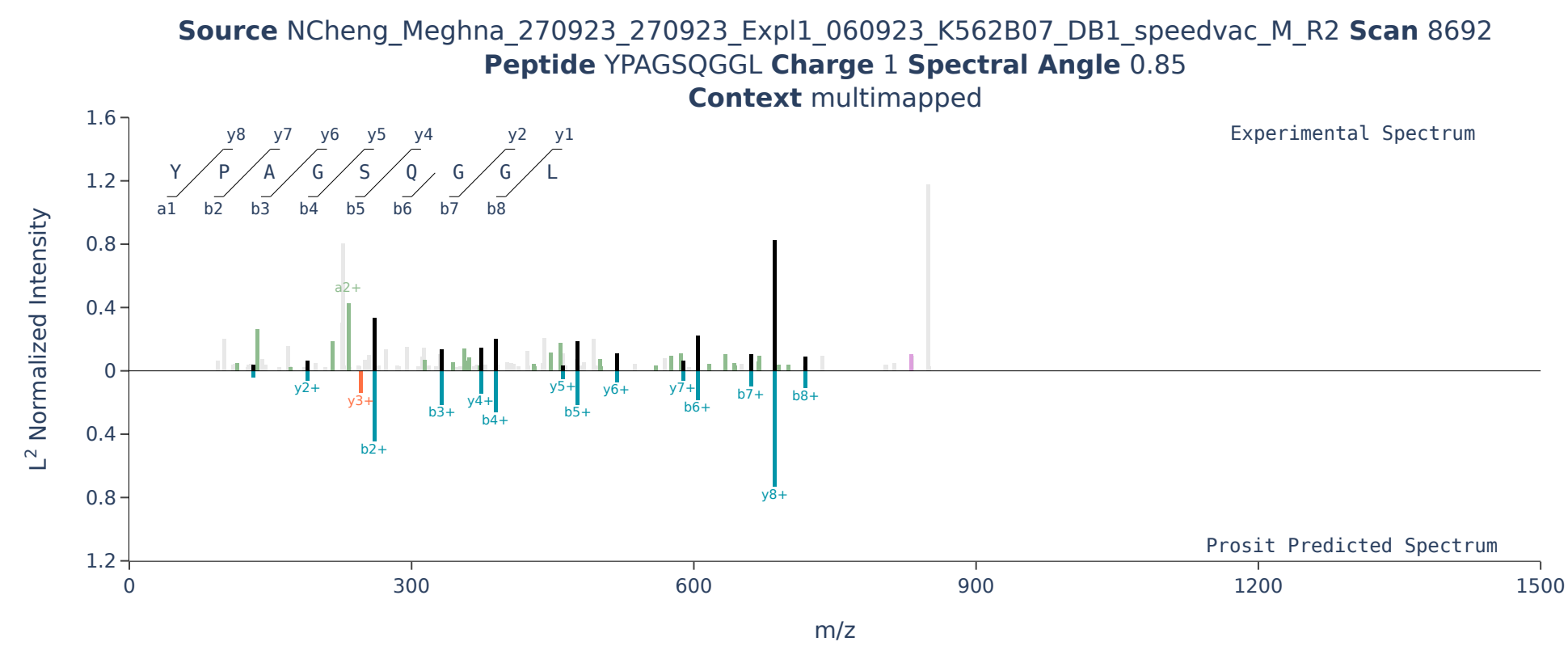
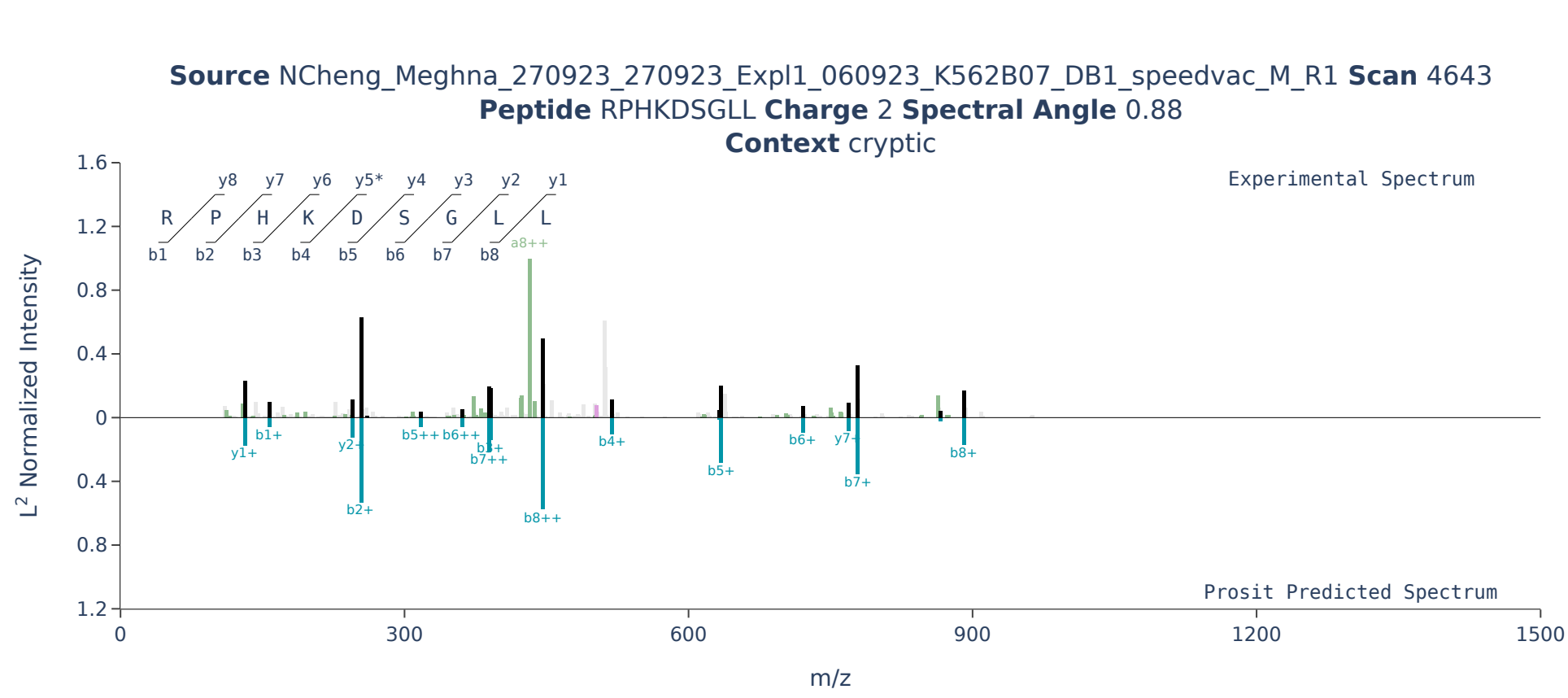
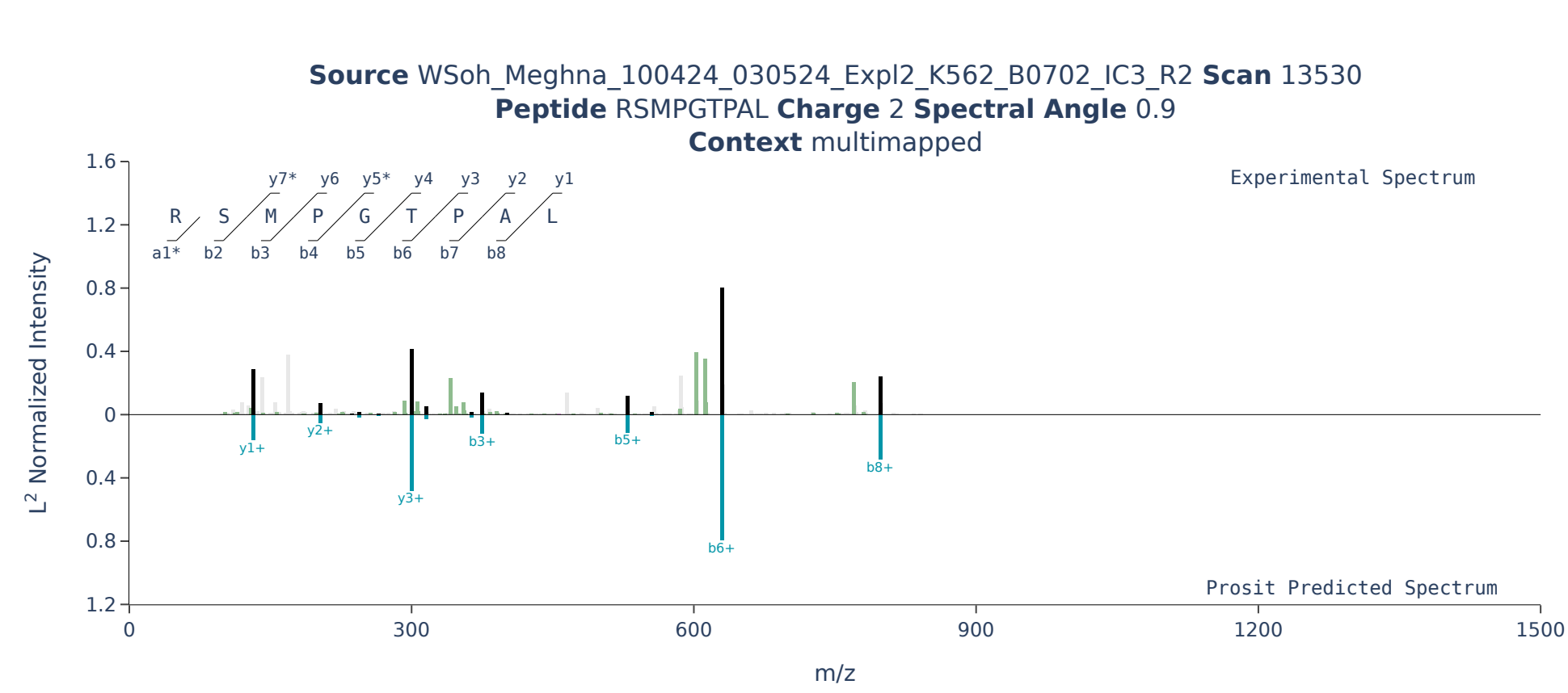
- Experimental peak matched to a Prosit predicted peak.
- Possible ion unknown to Prosit.
- Precursor matched peak.
- Experimental peak not matched to any potential ion.

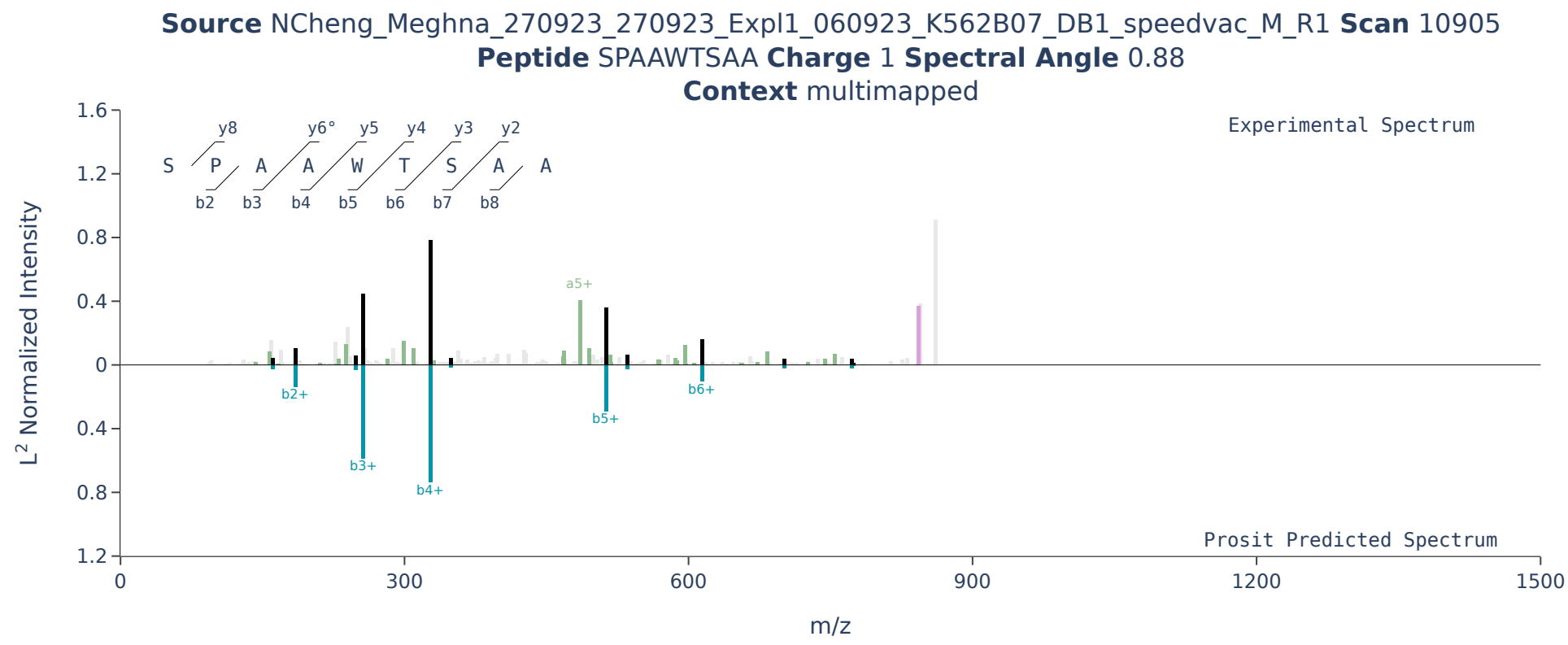
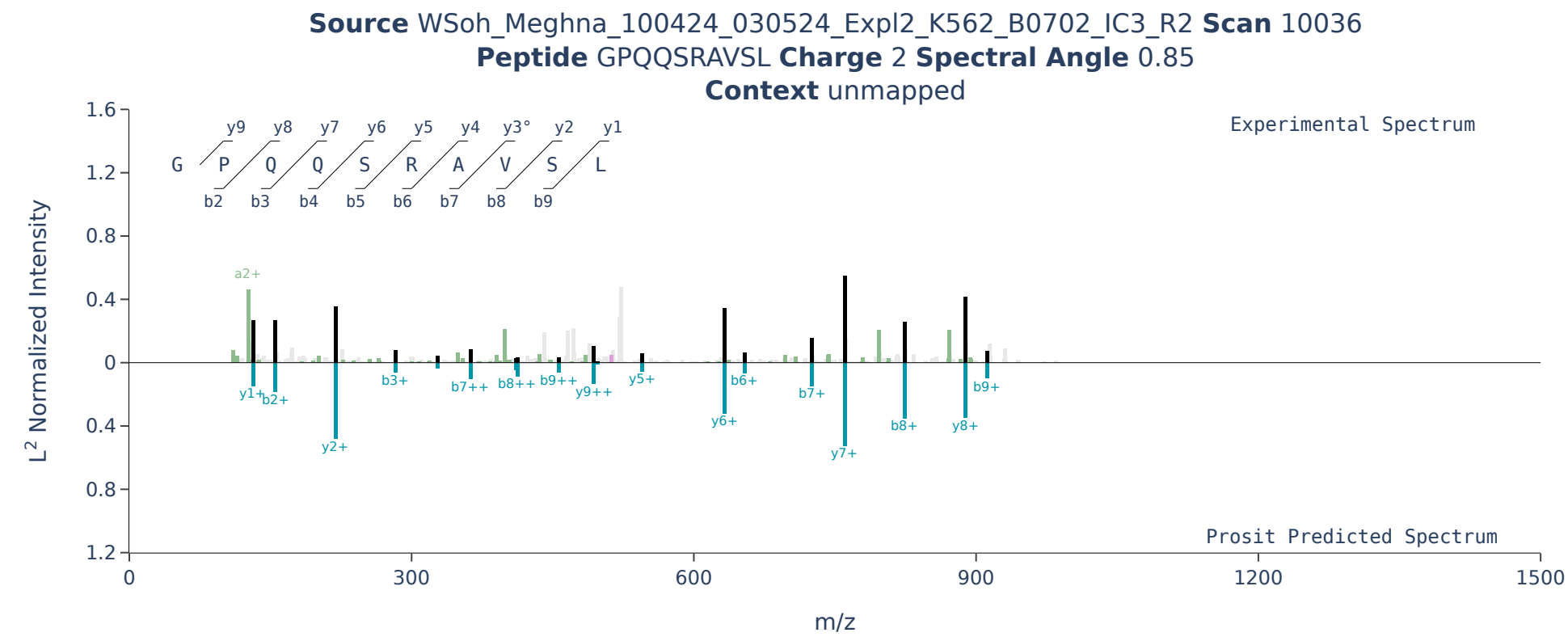
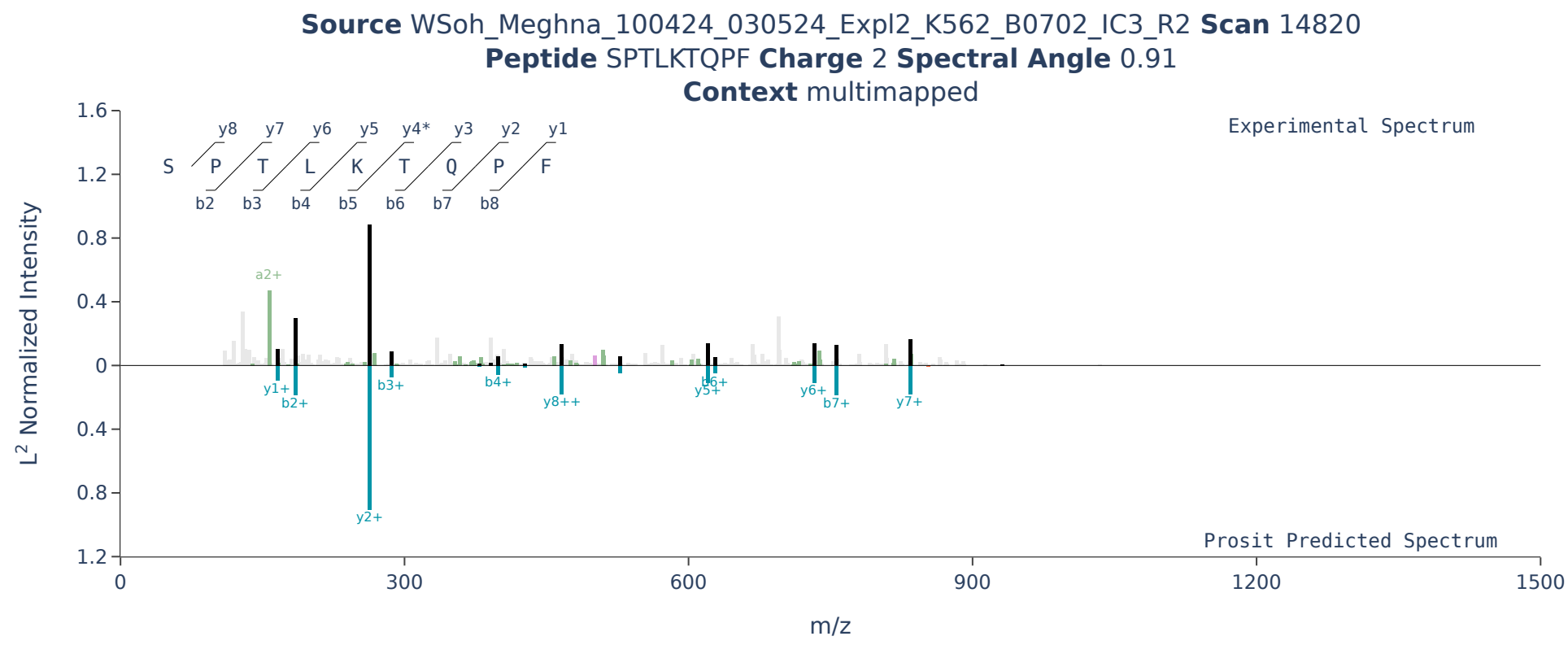
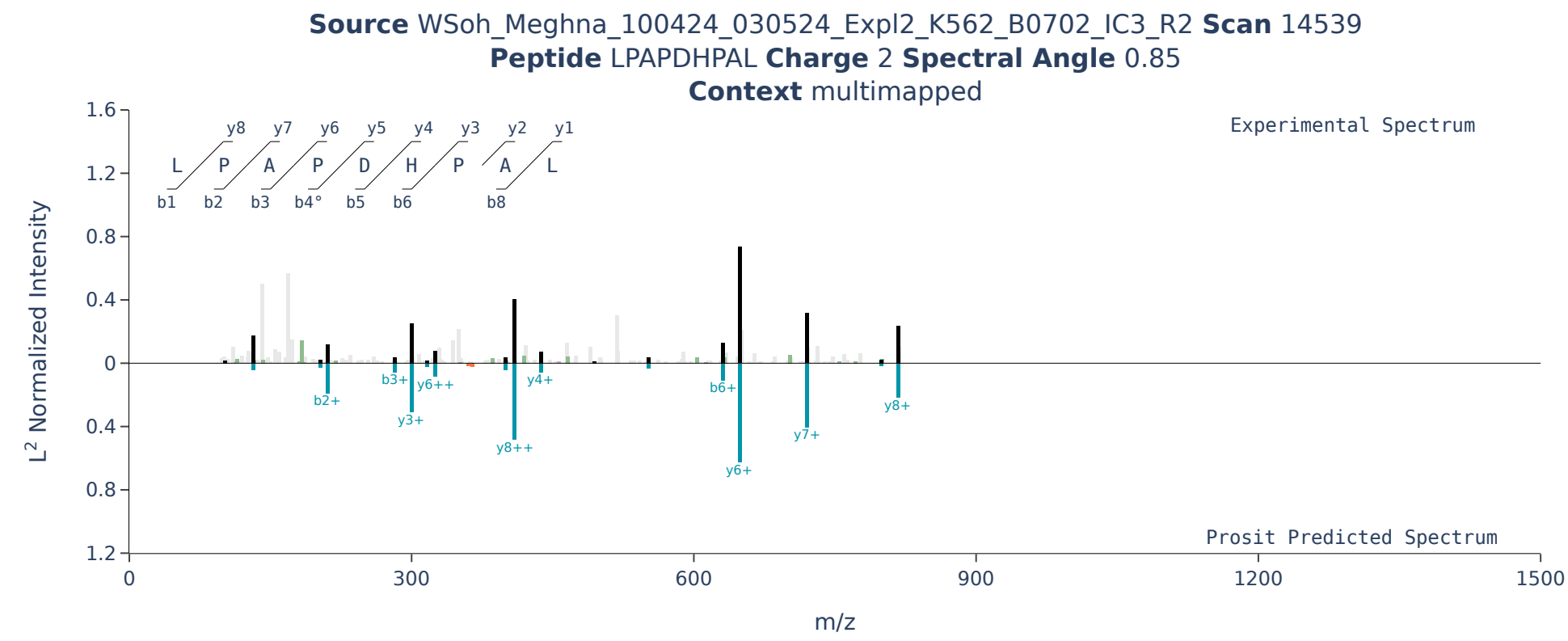
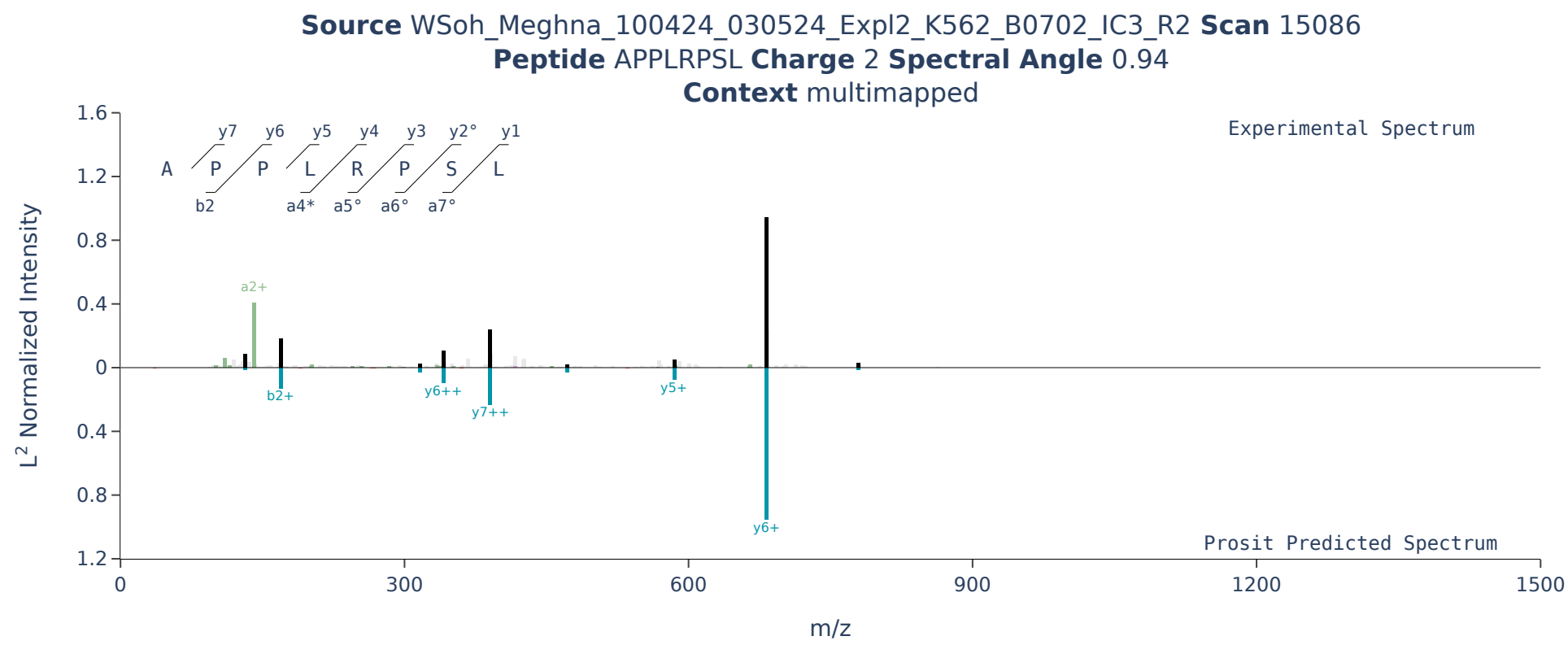
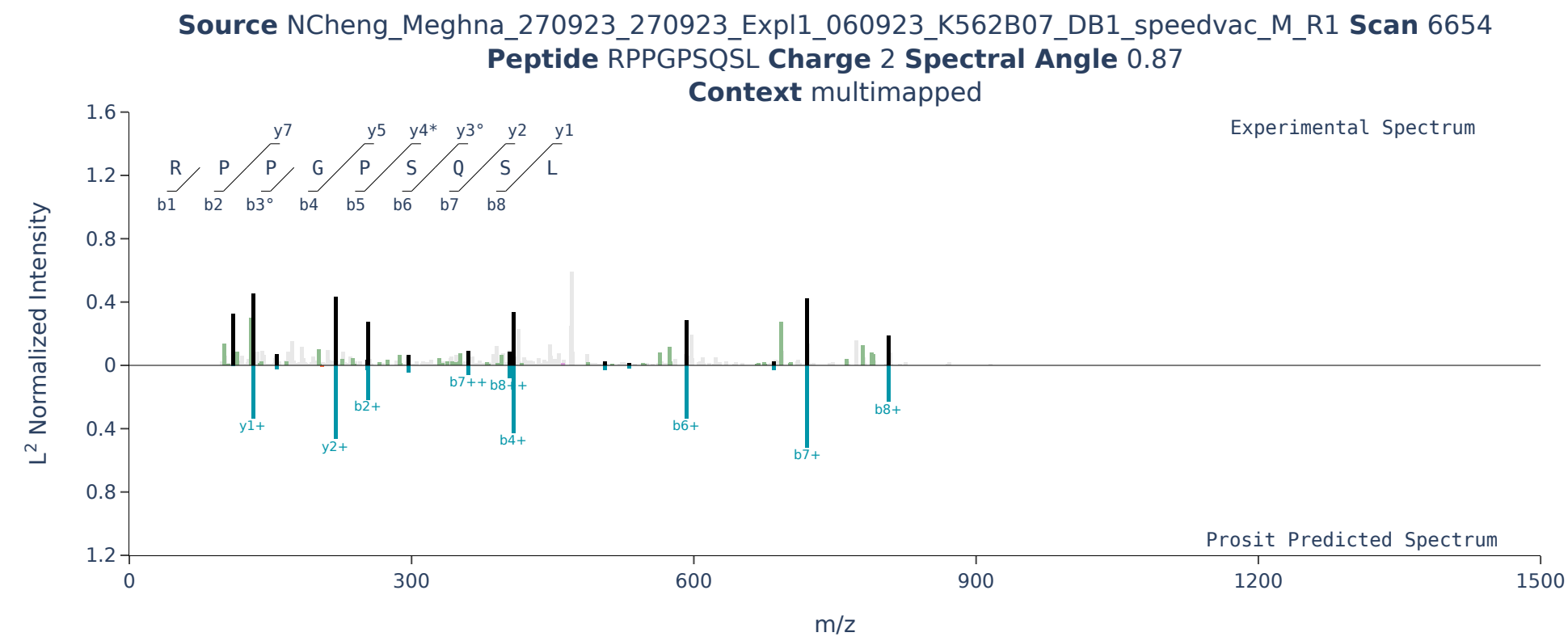
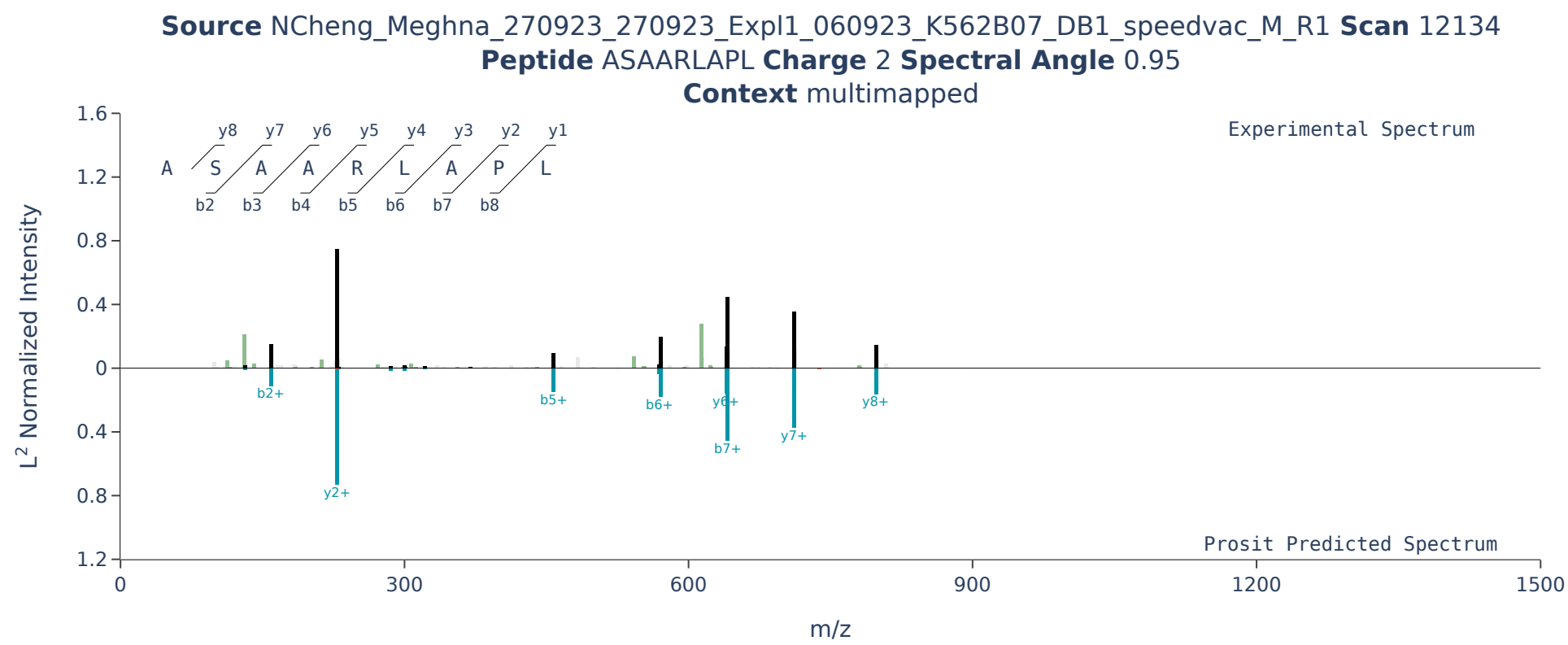
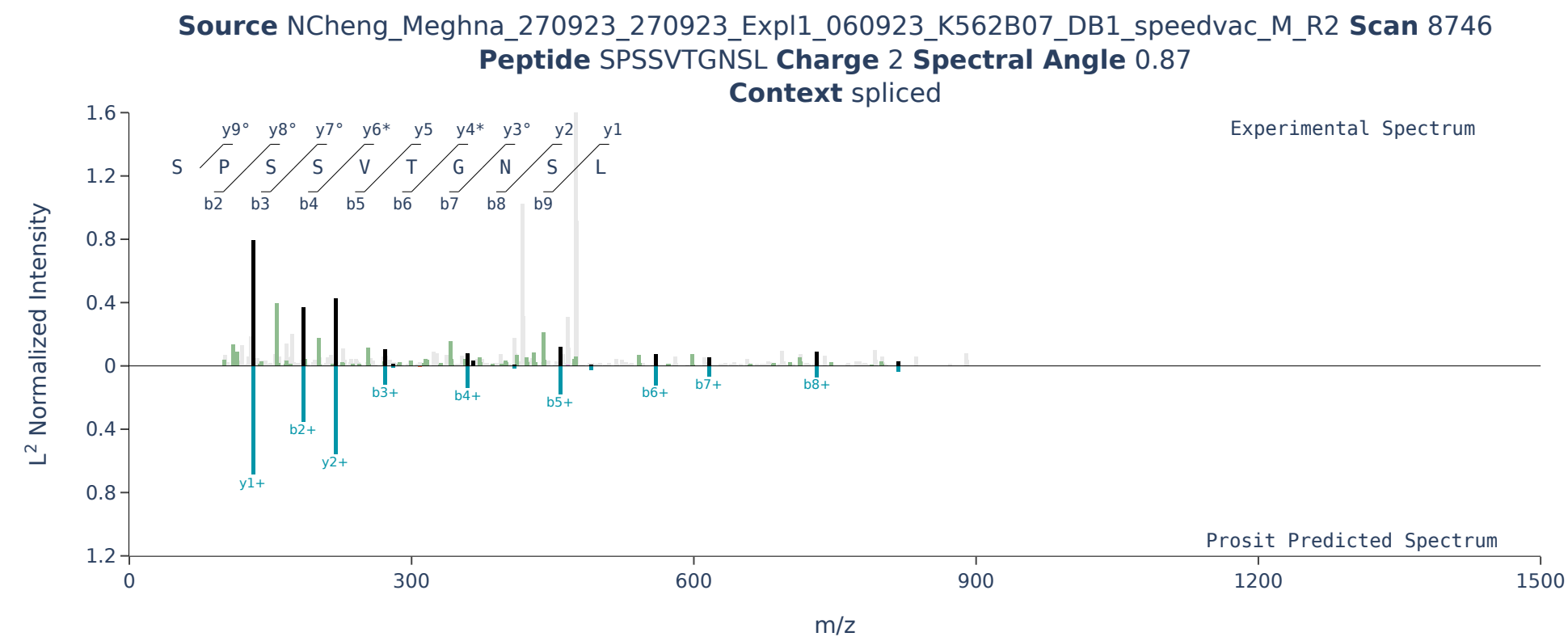
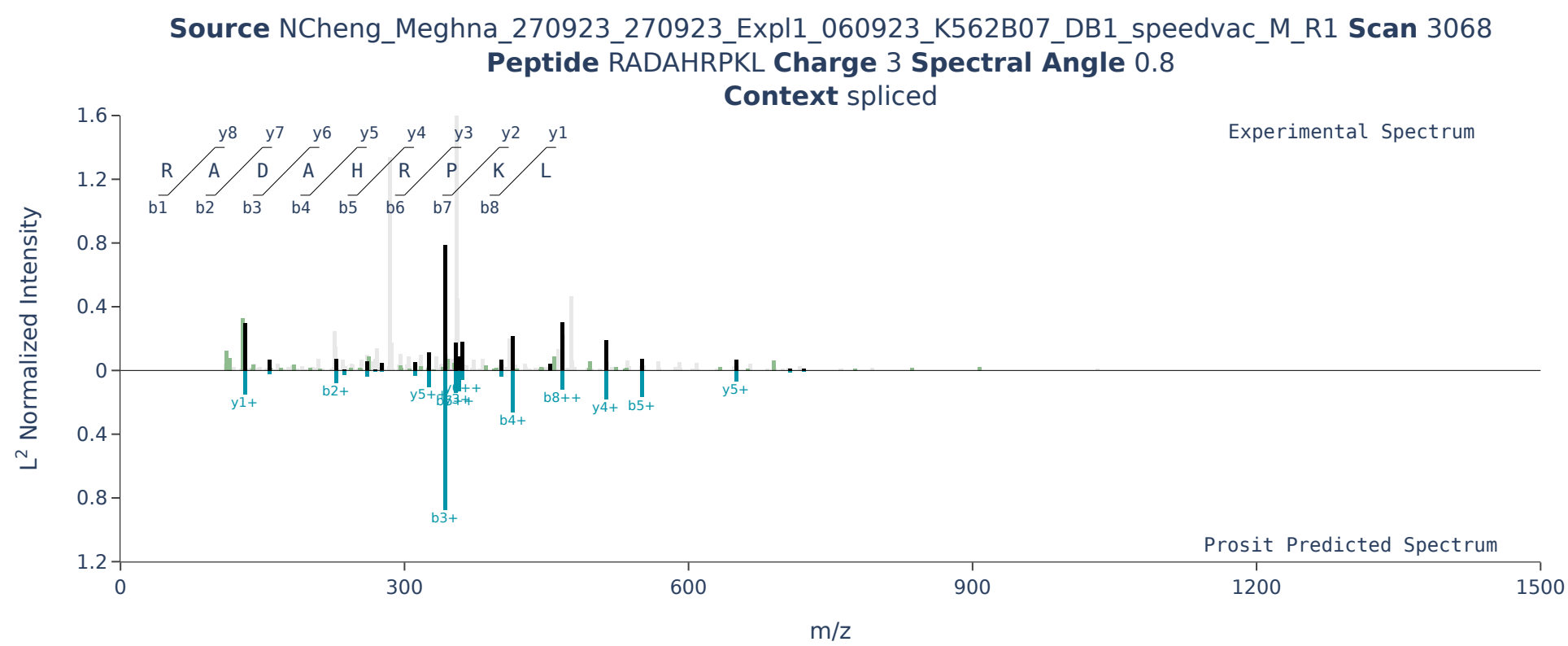
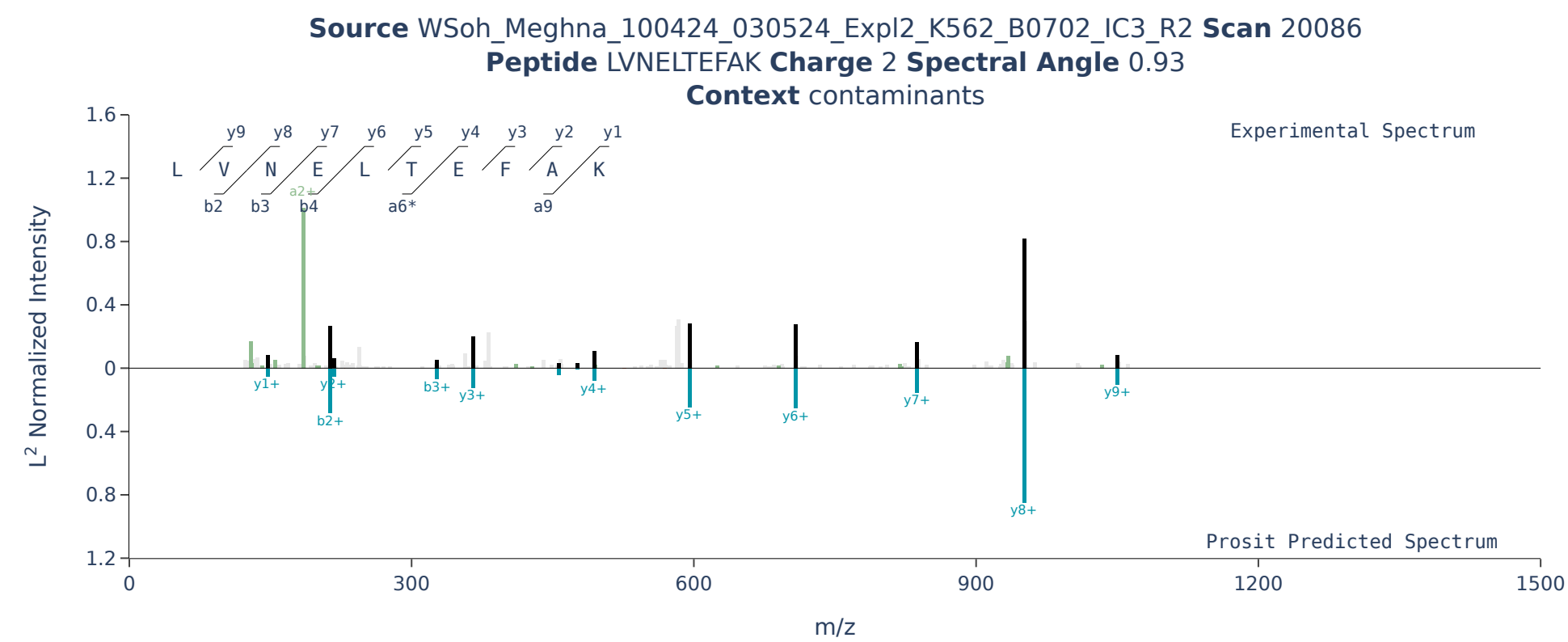
Prosit Spectrum Colour Code:

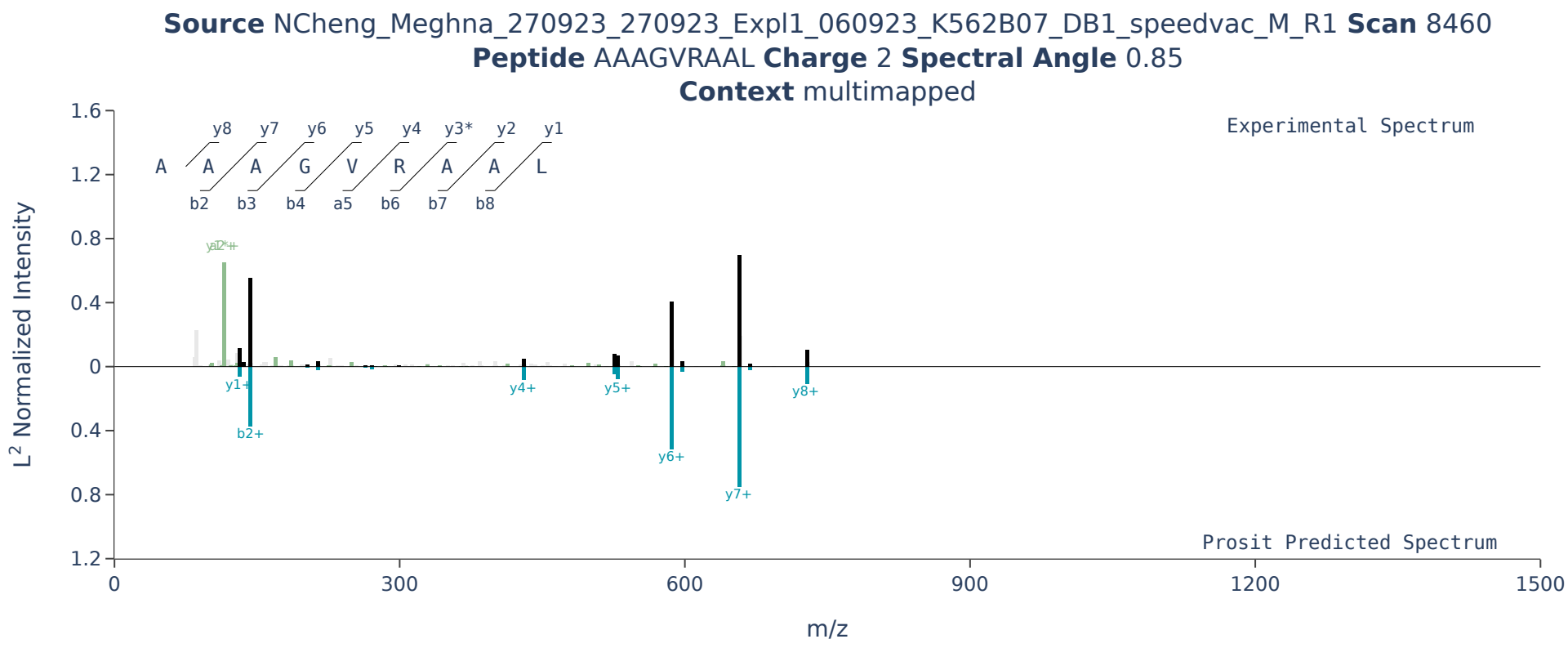
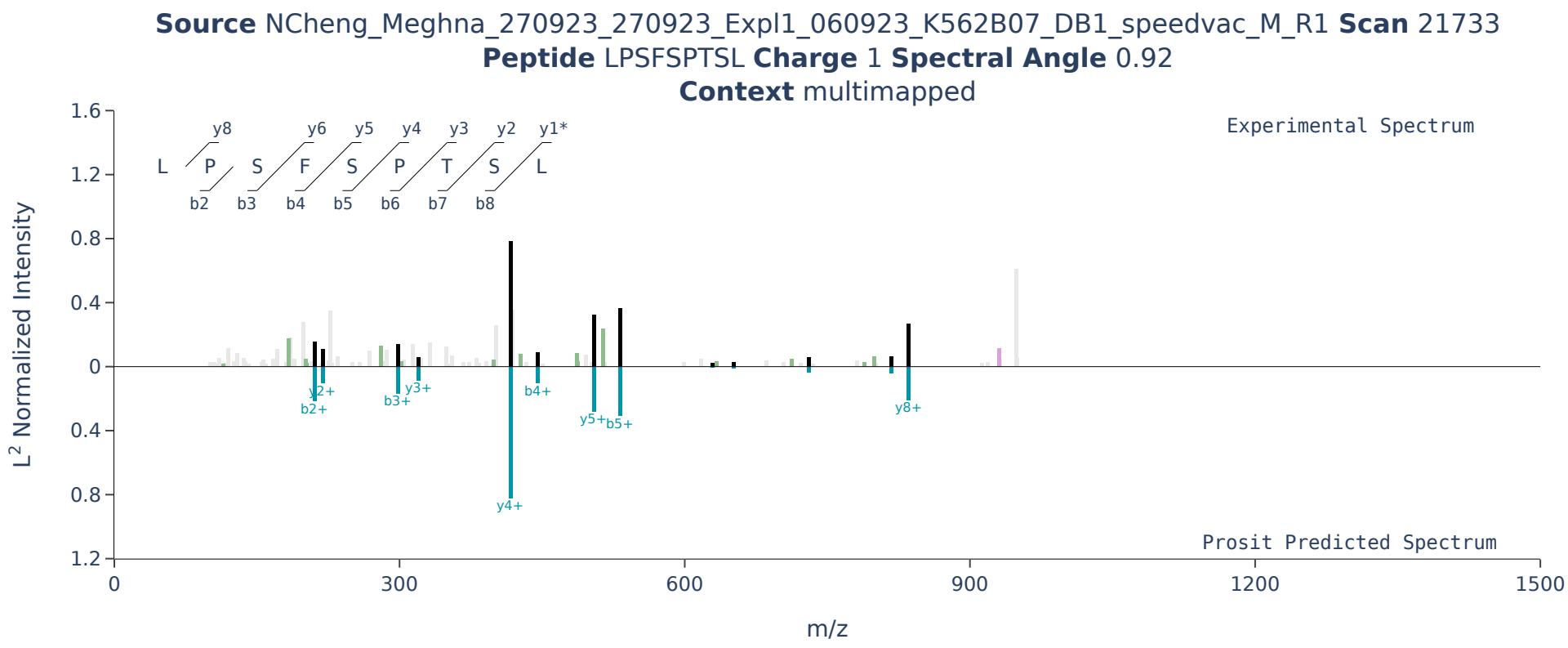
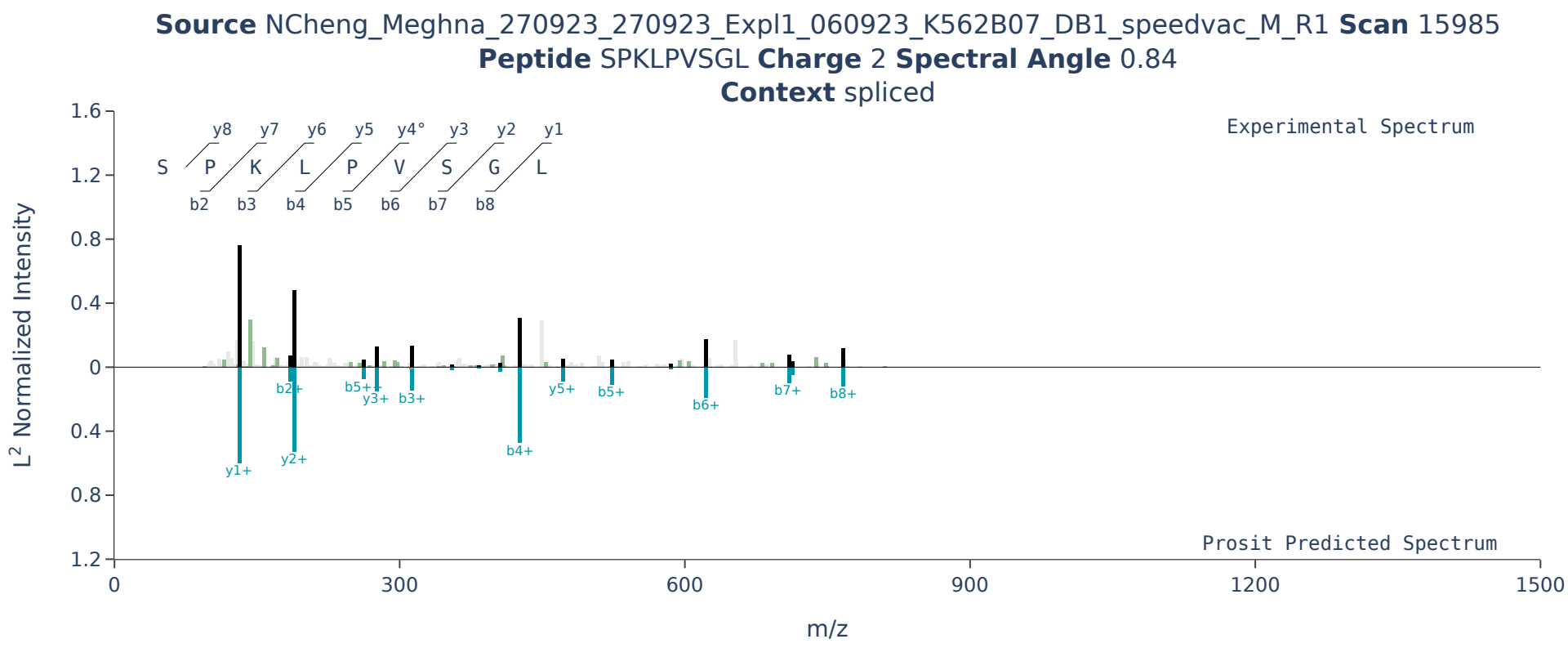
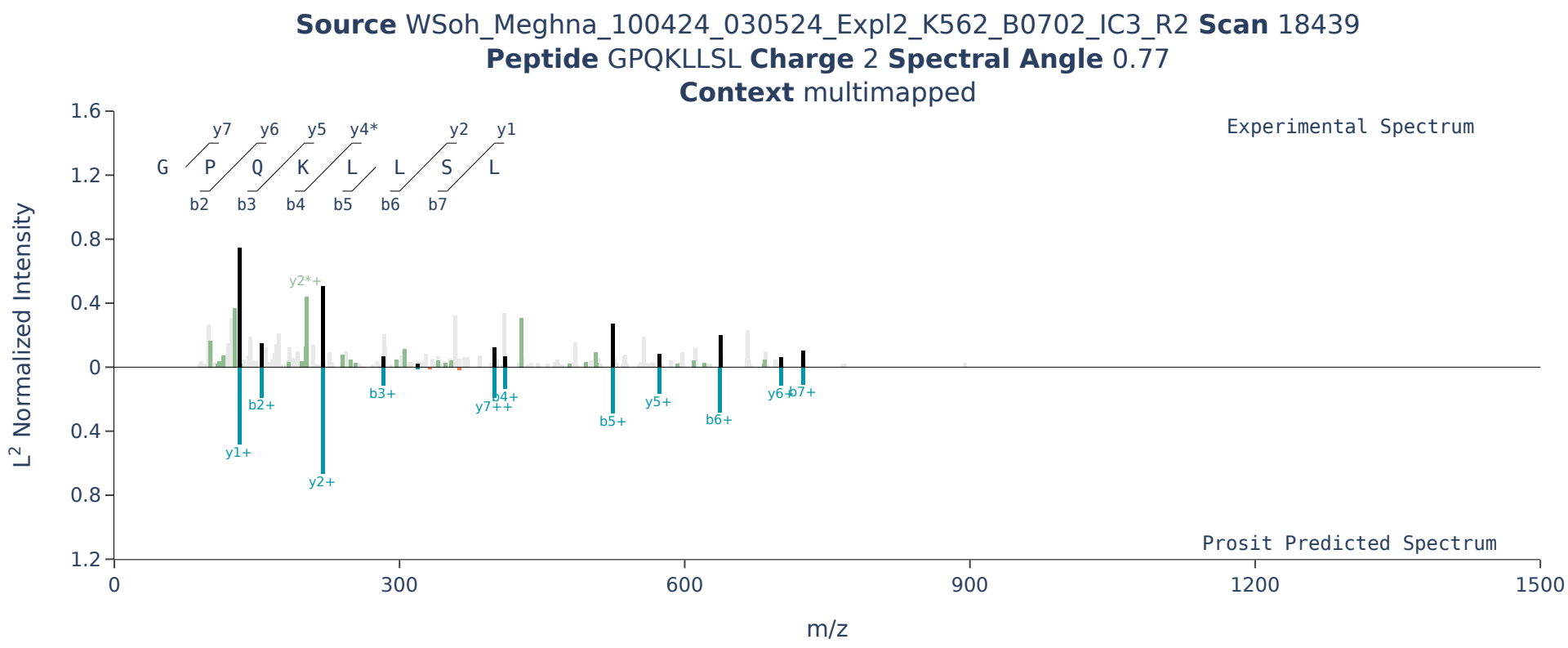
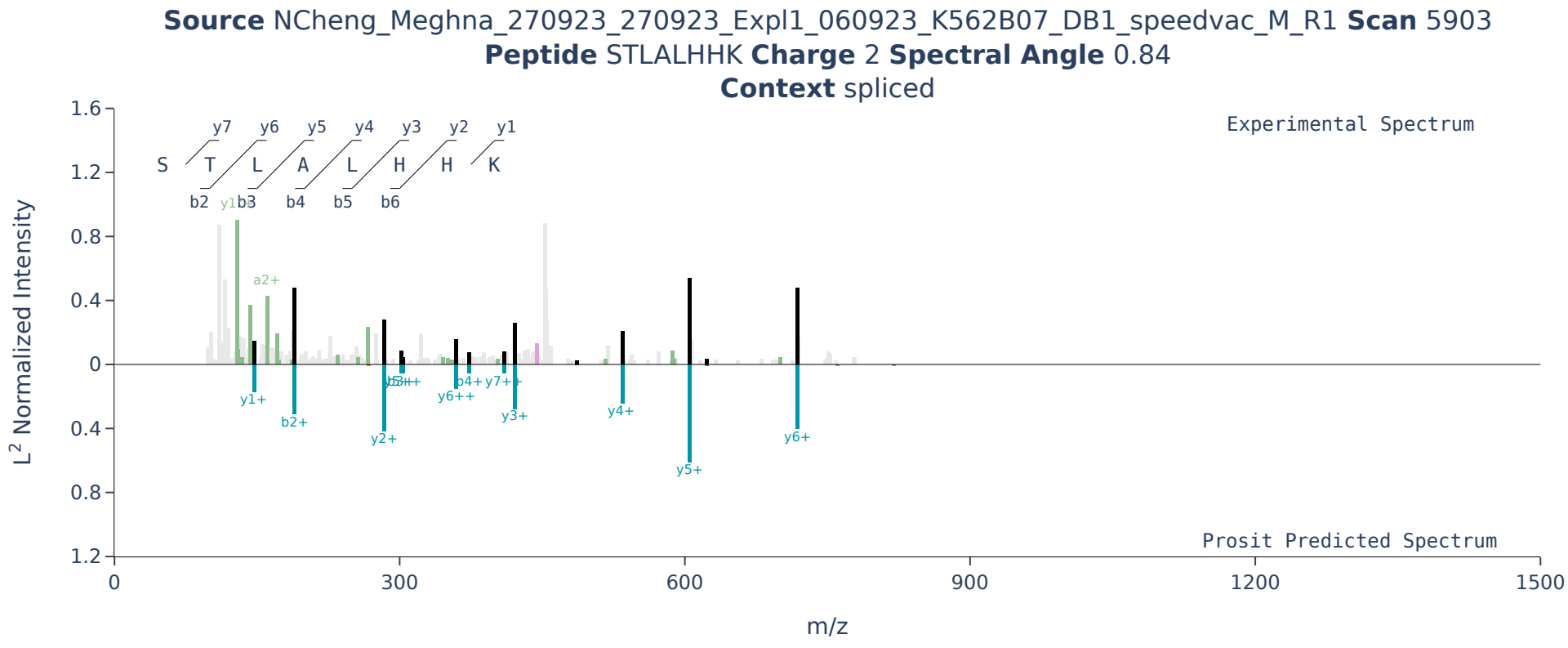
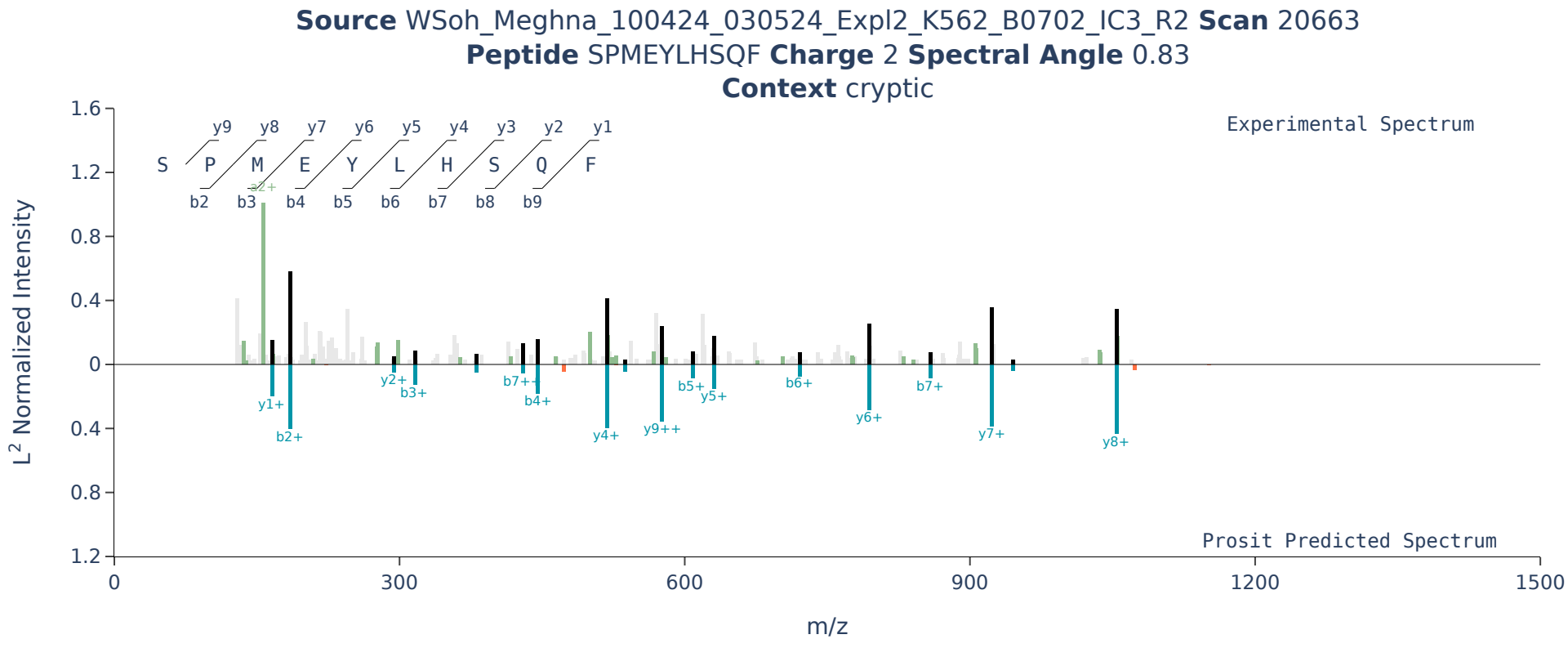
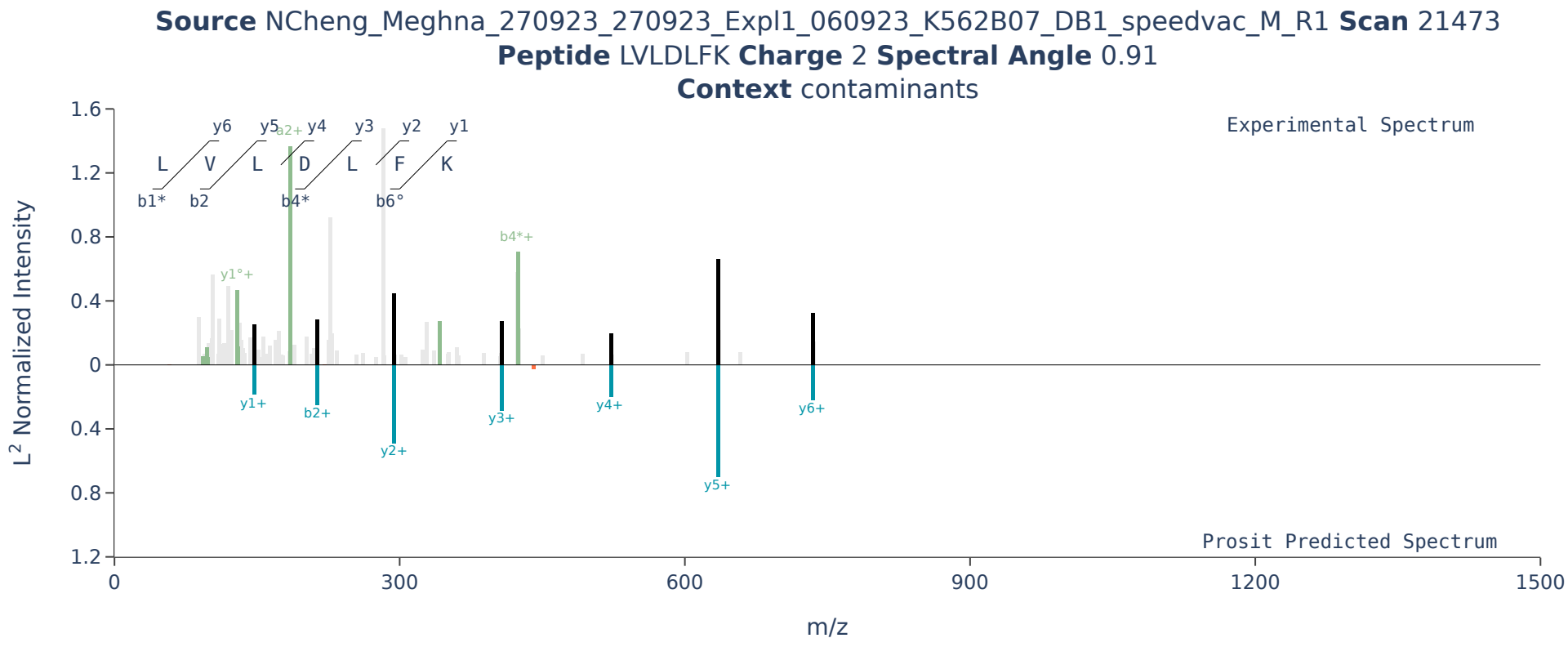
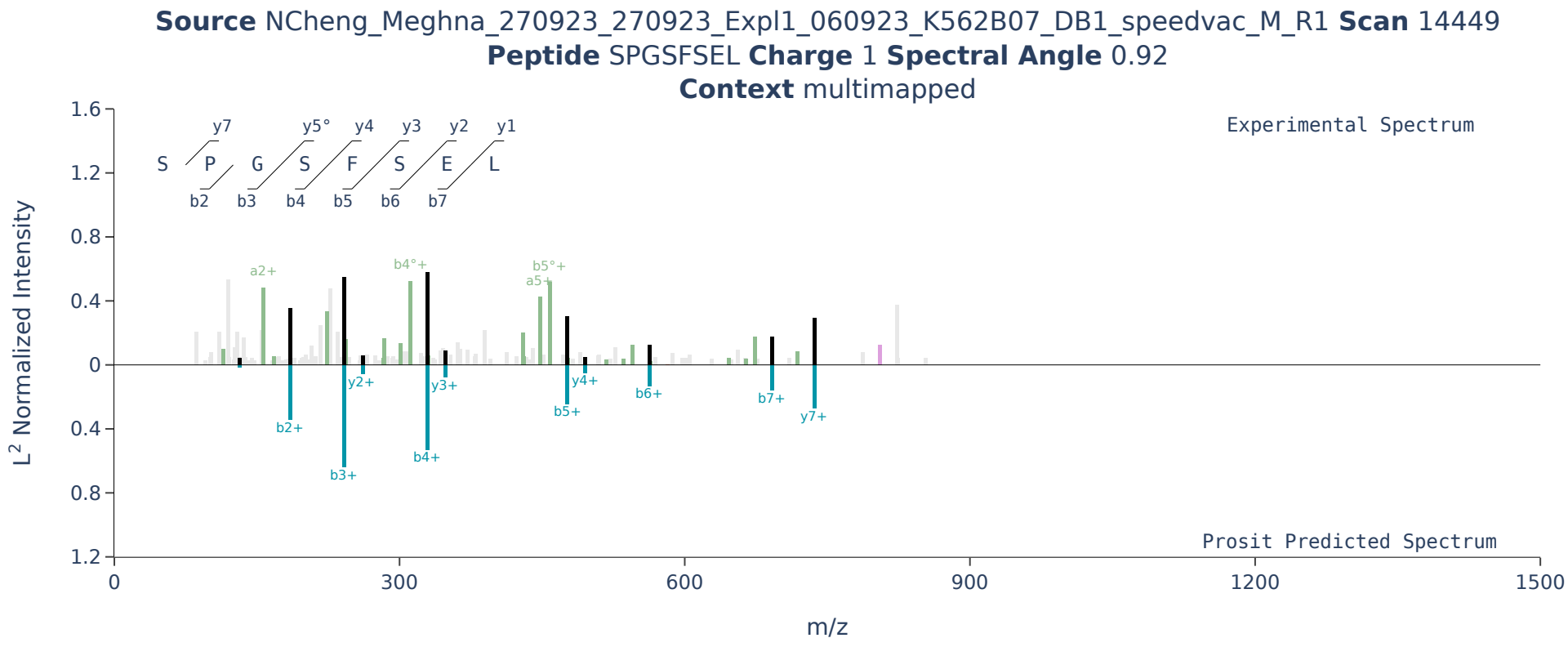
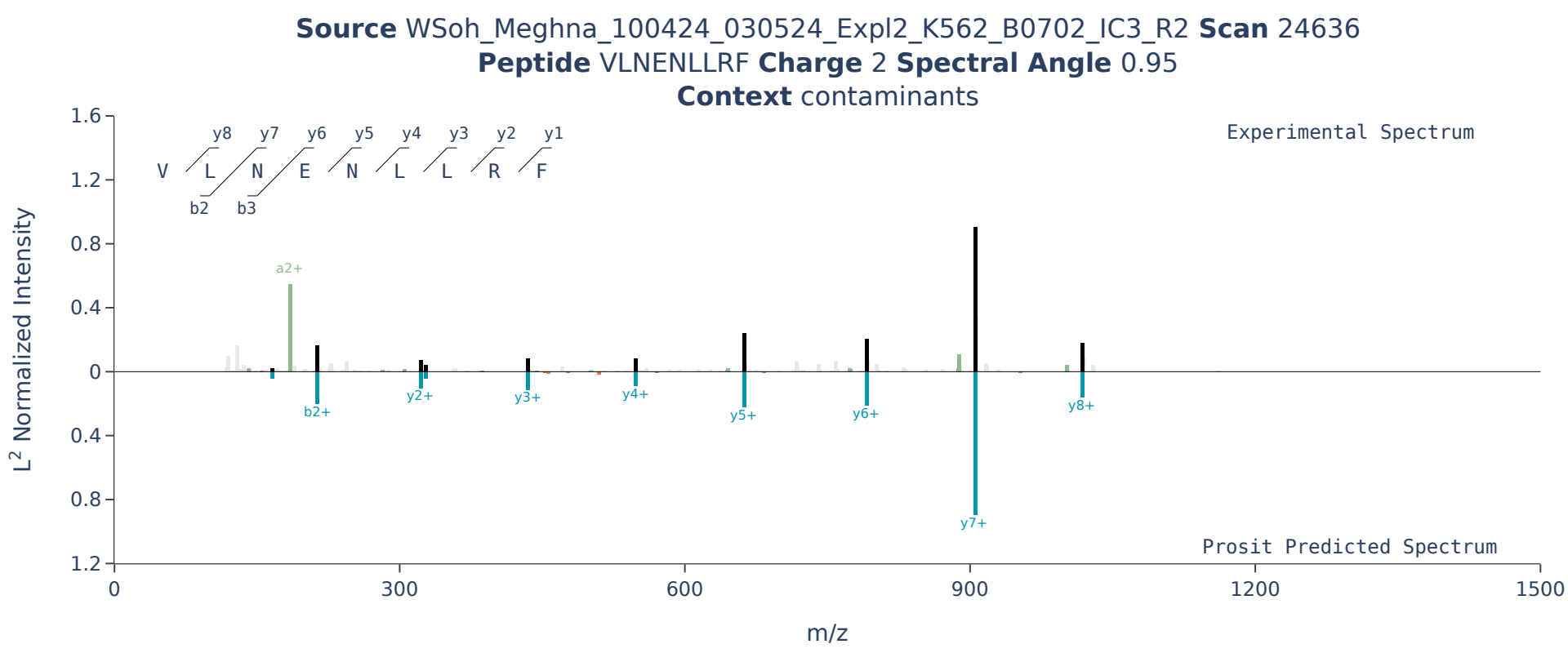
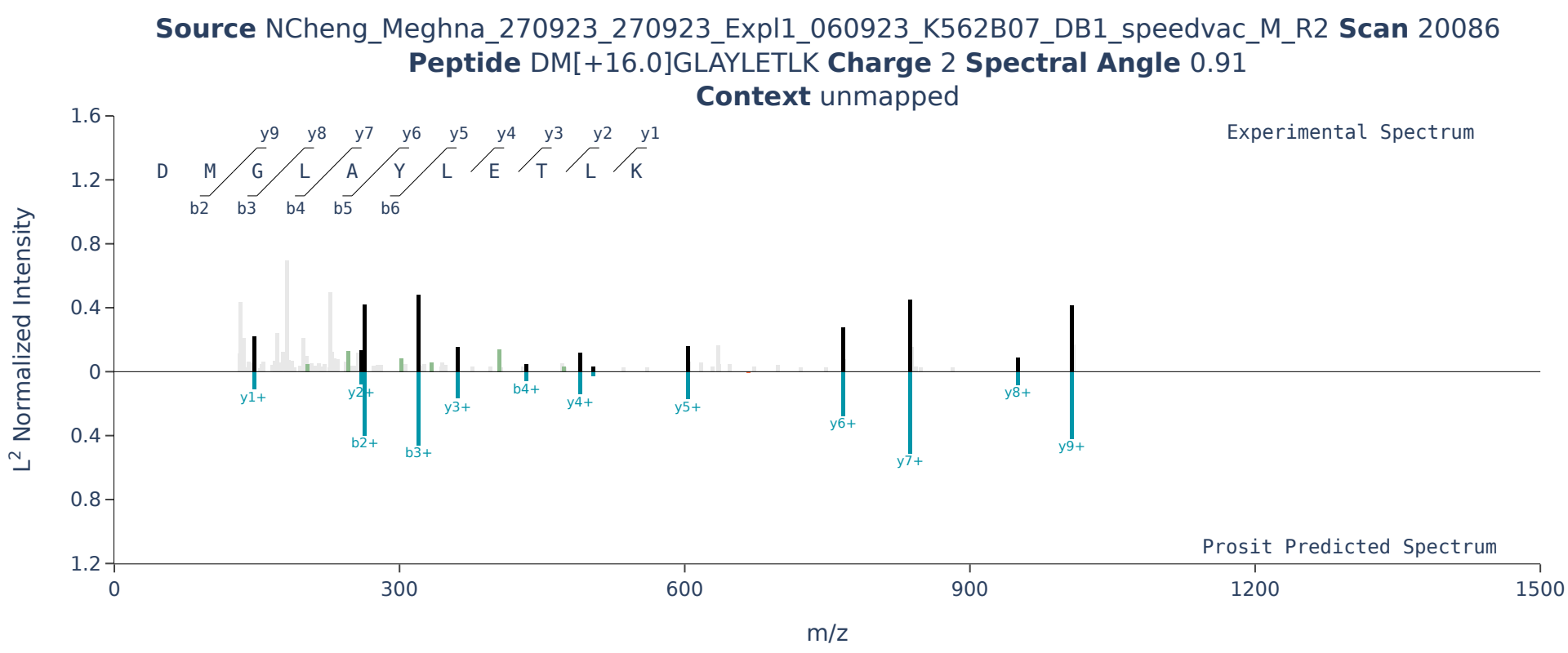
- Prosit predicted peak matched to experimental spectrum.
- Prosit predicted peak not matched to experimental spectrum.

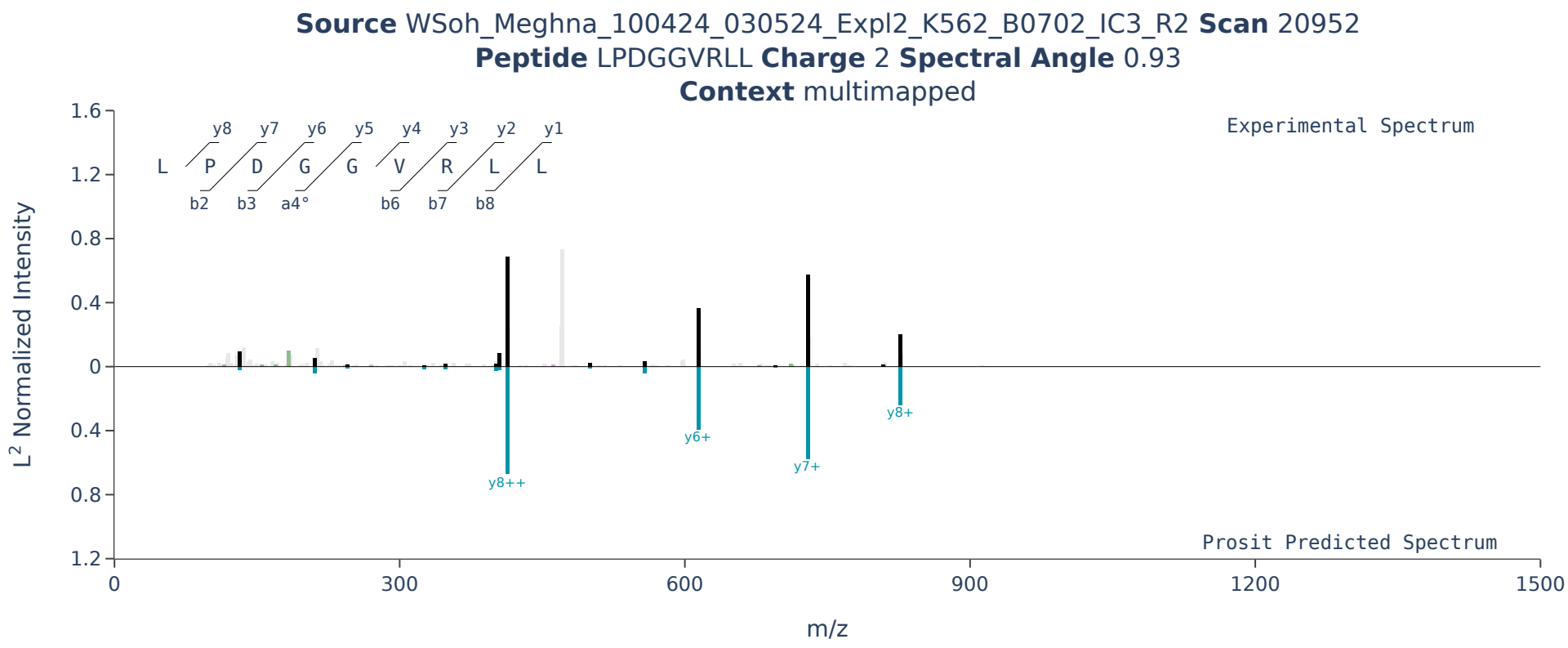
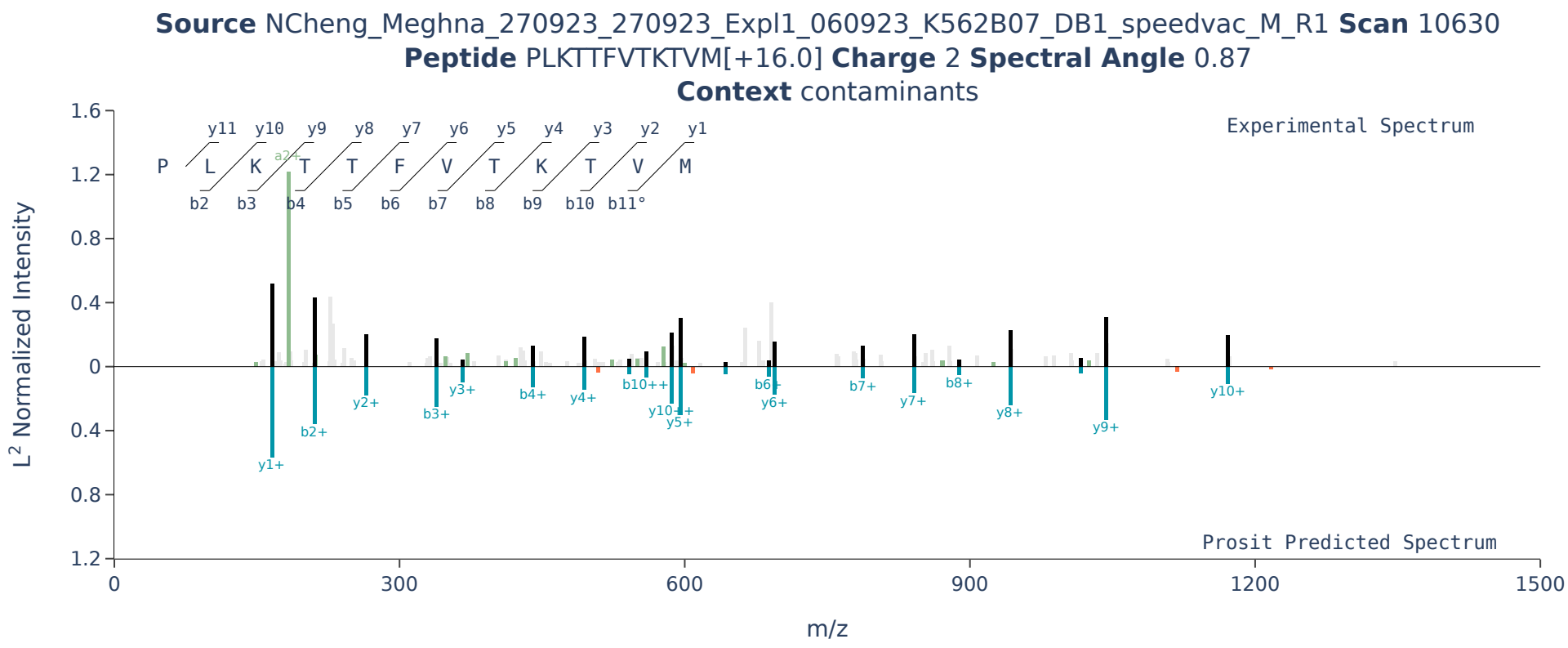
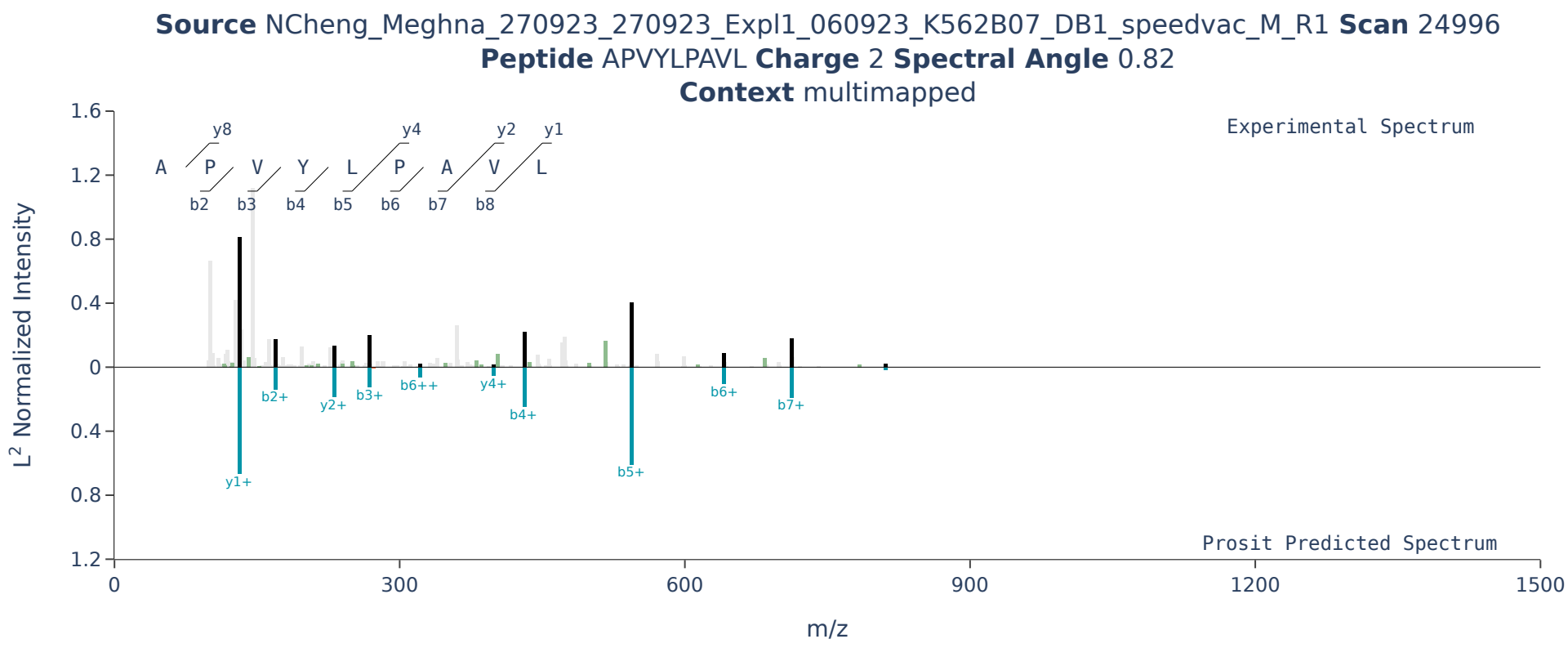
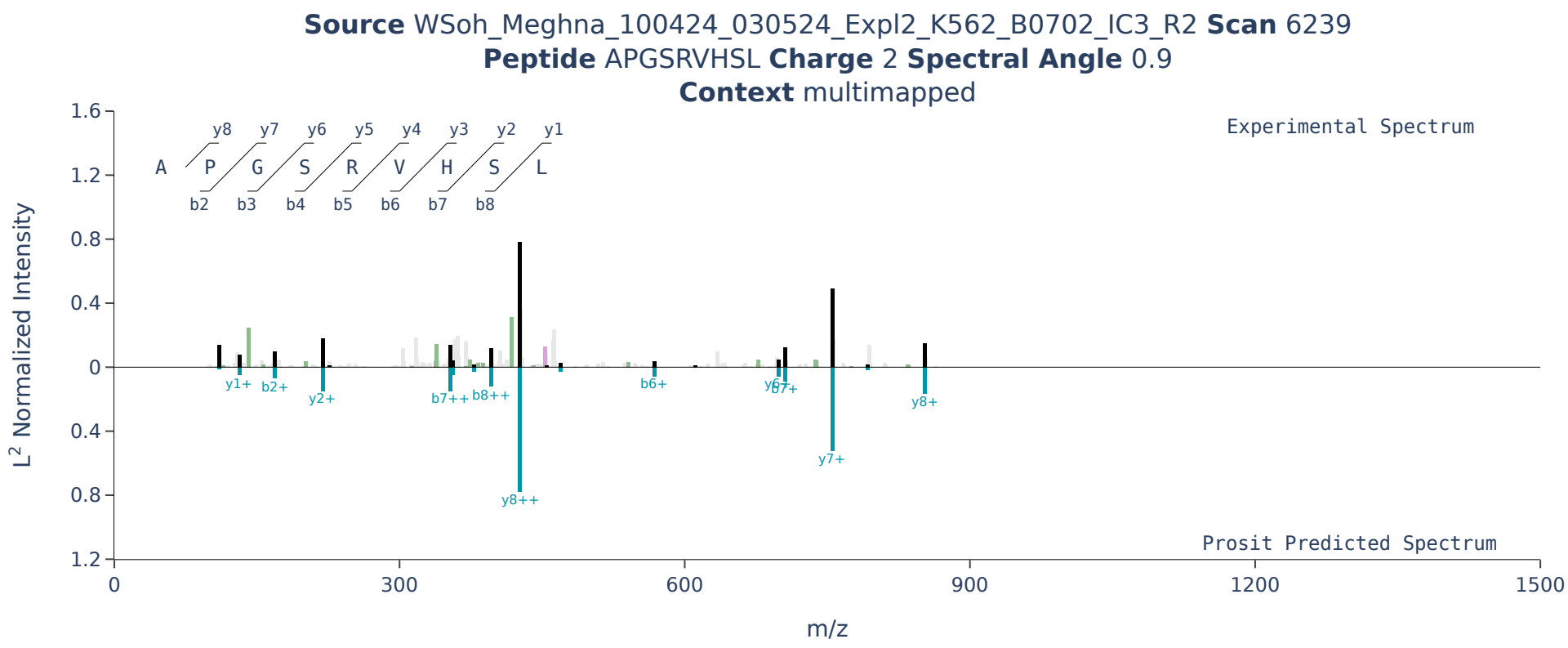
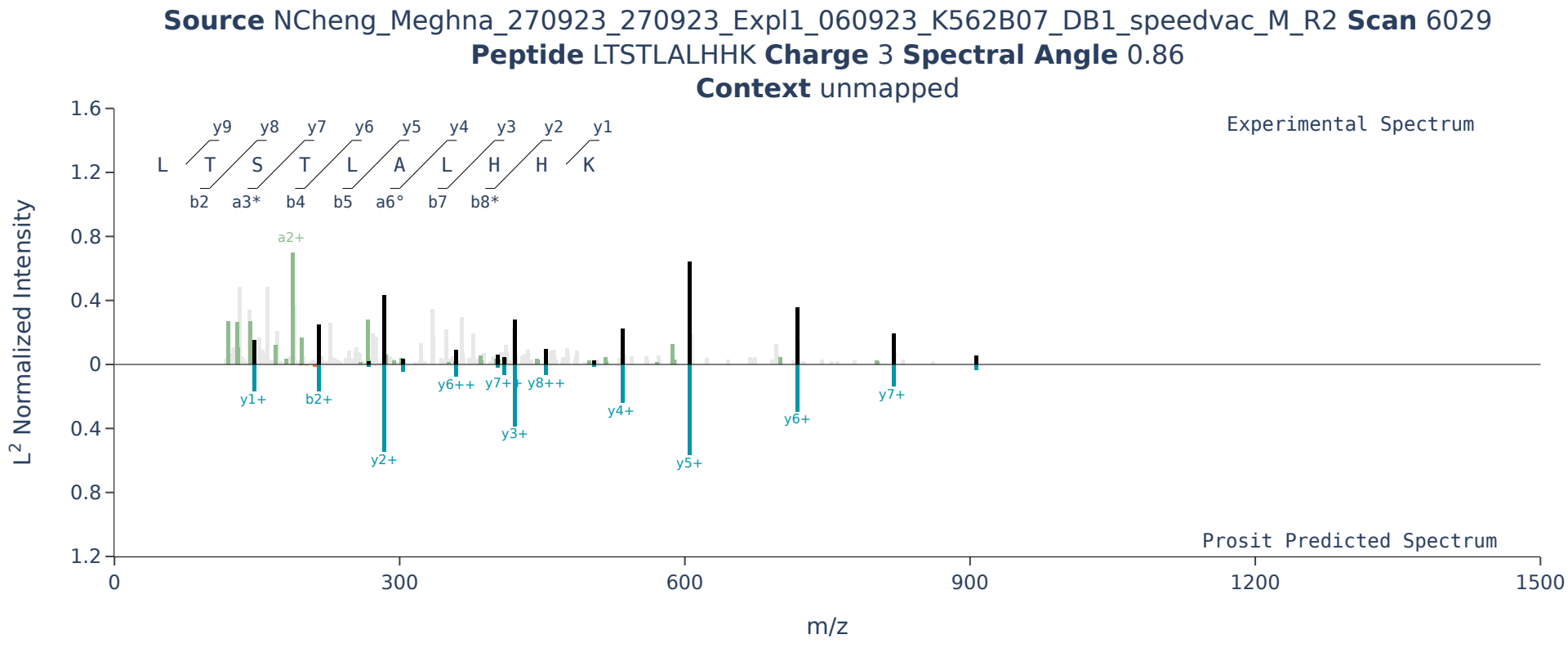
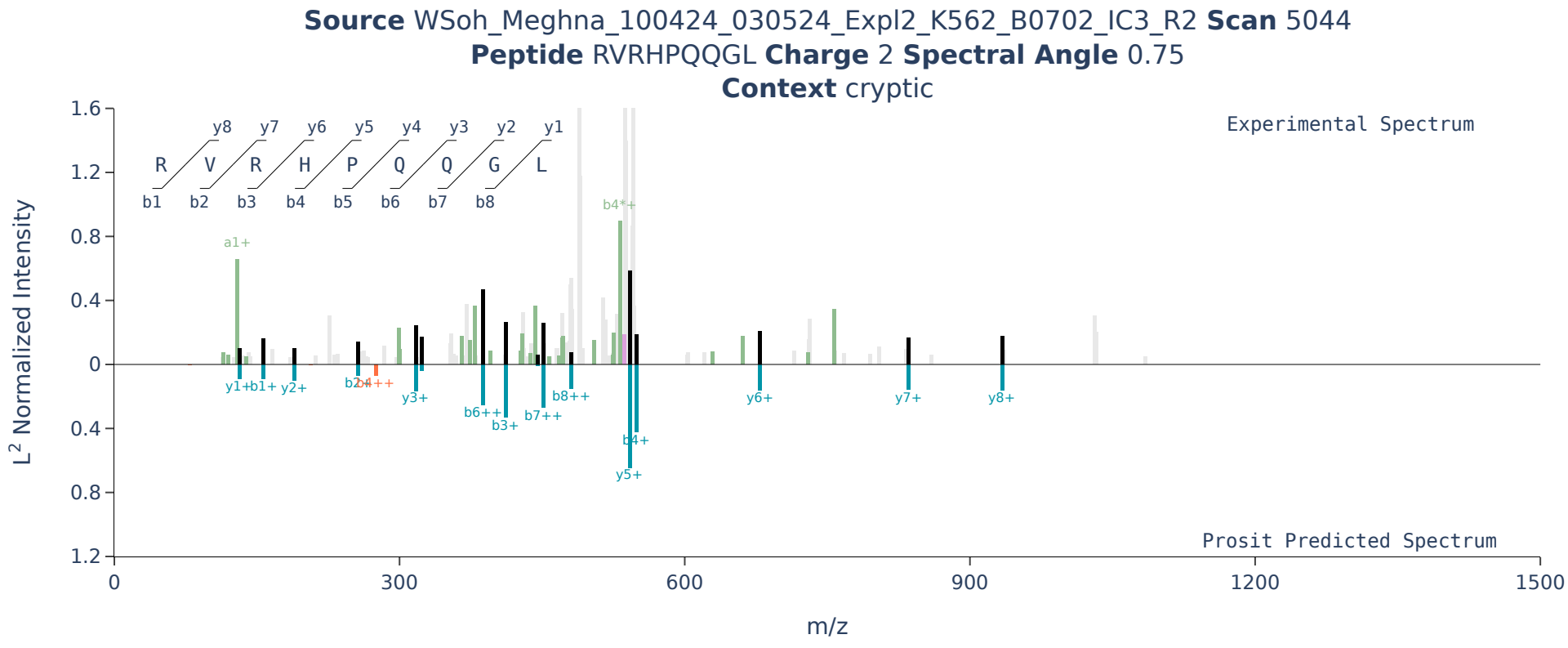
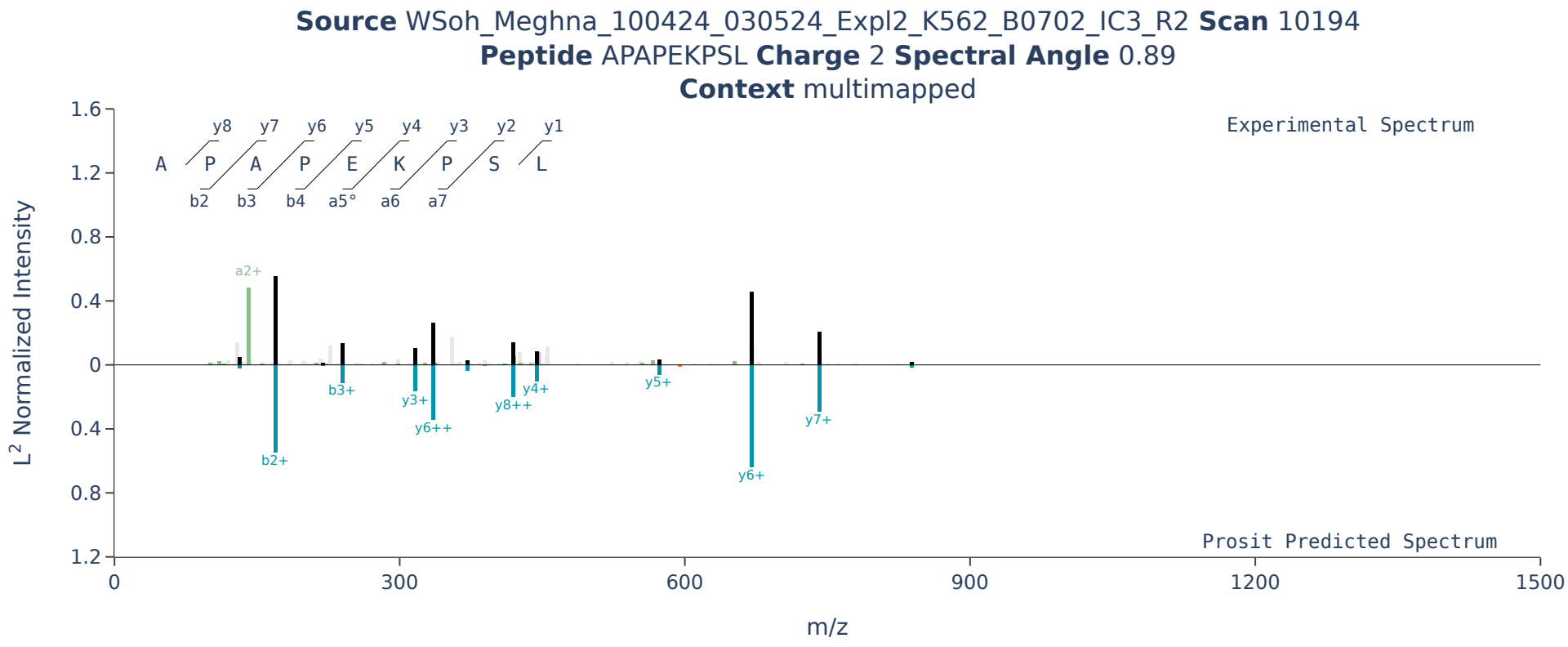
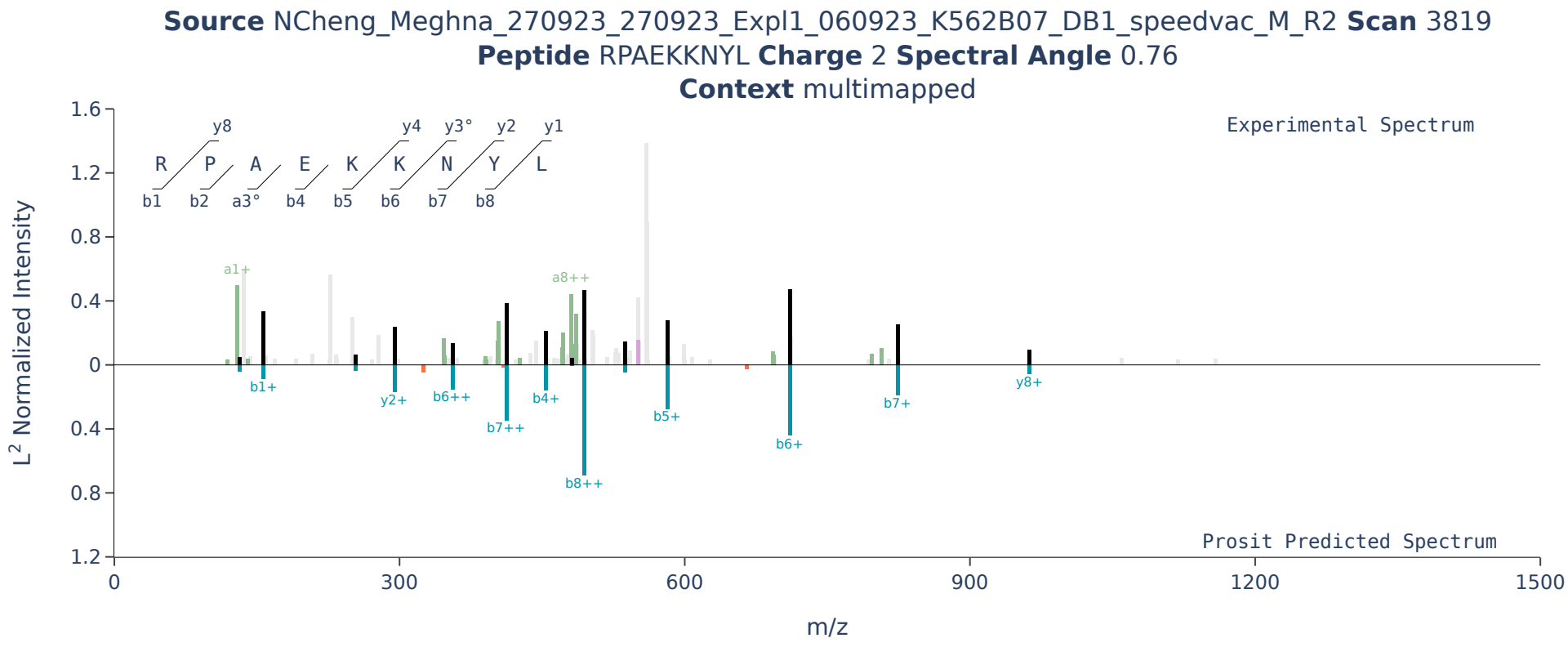
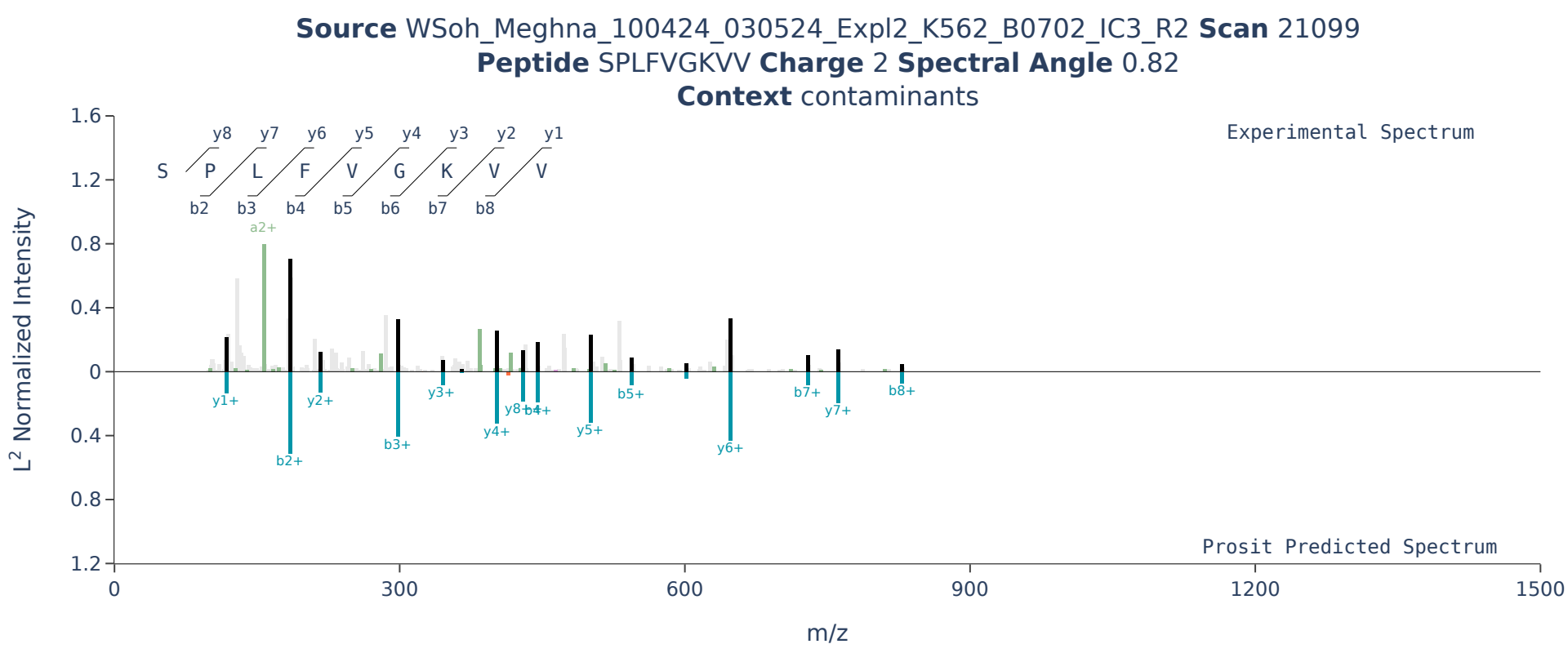
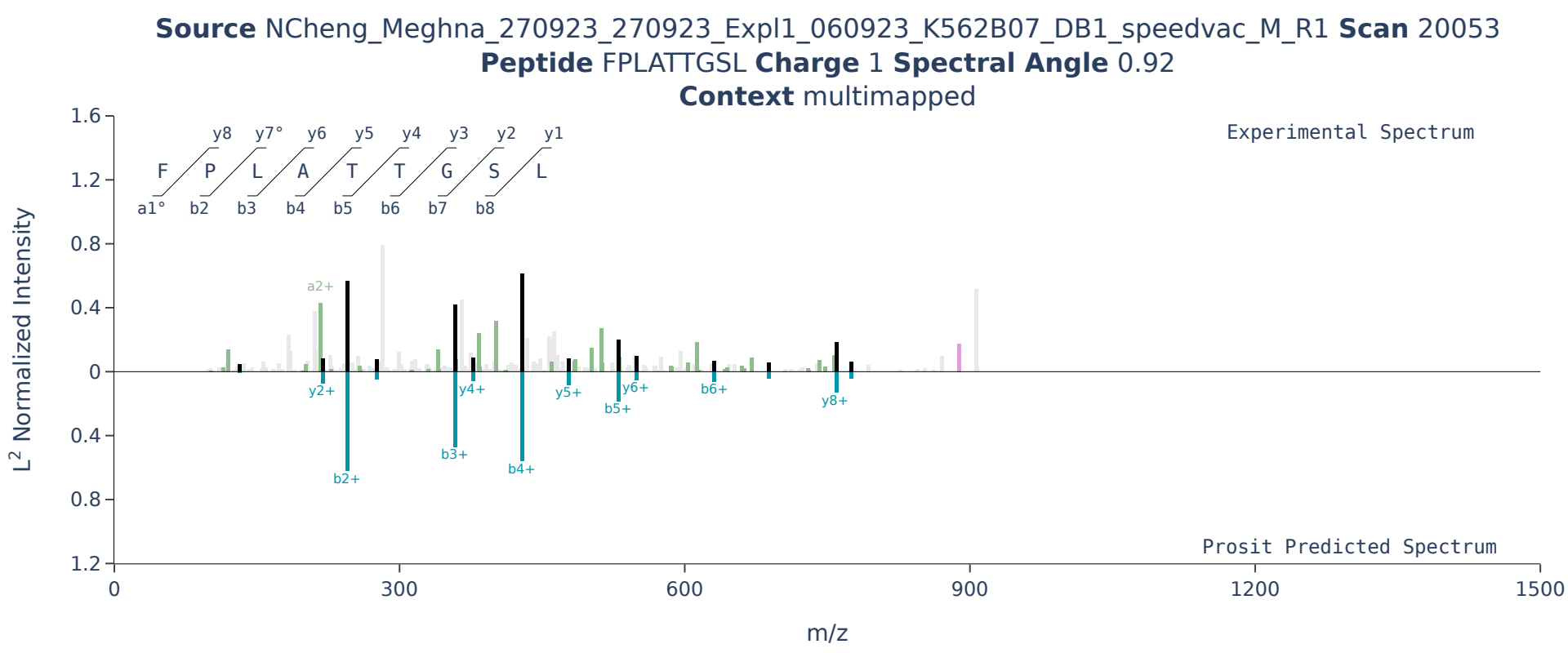
Additional Notes:

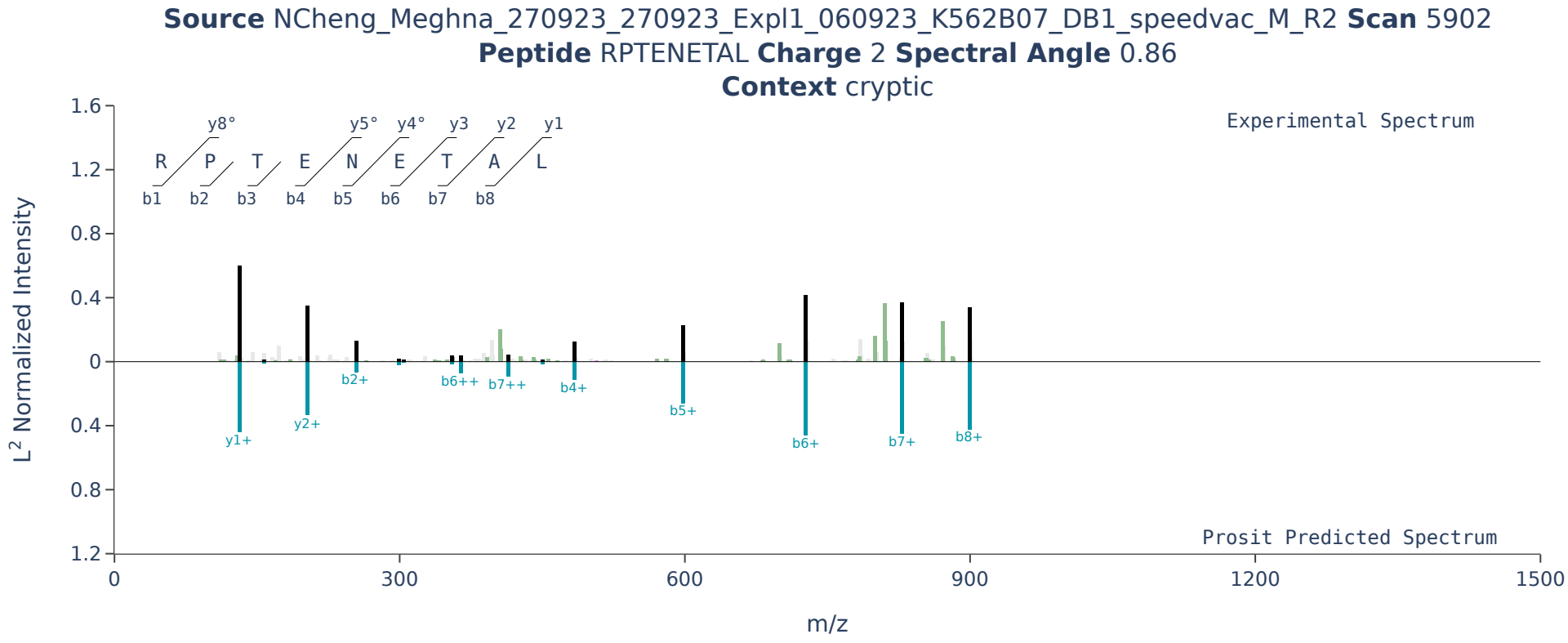
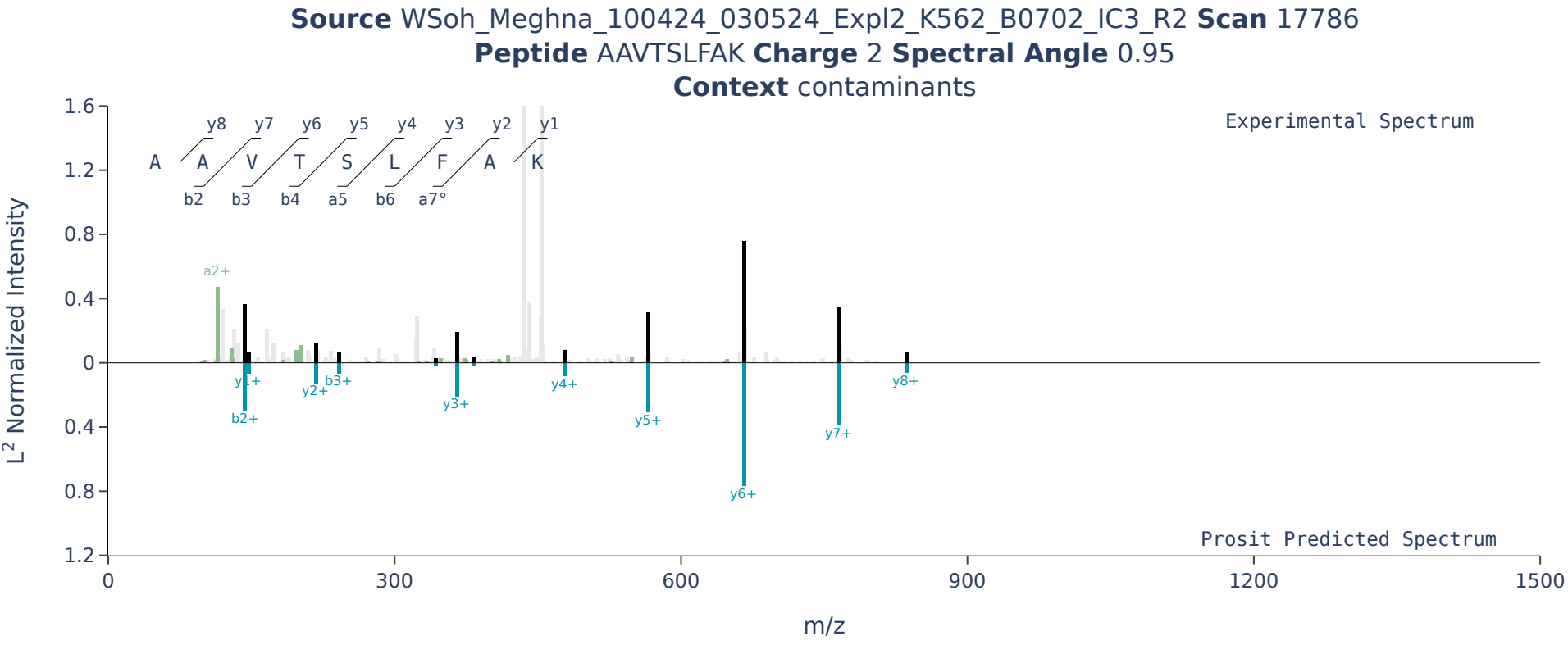
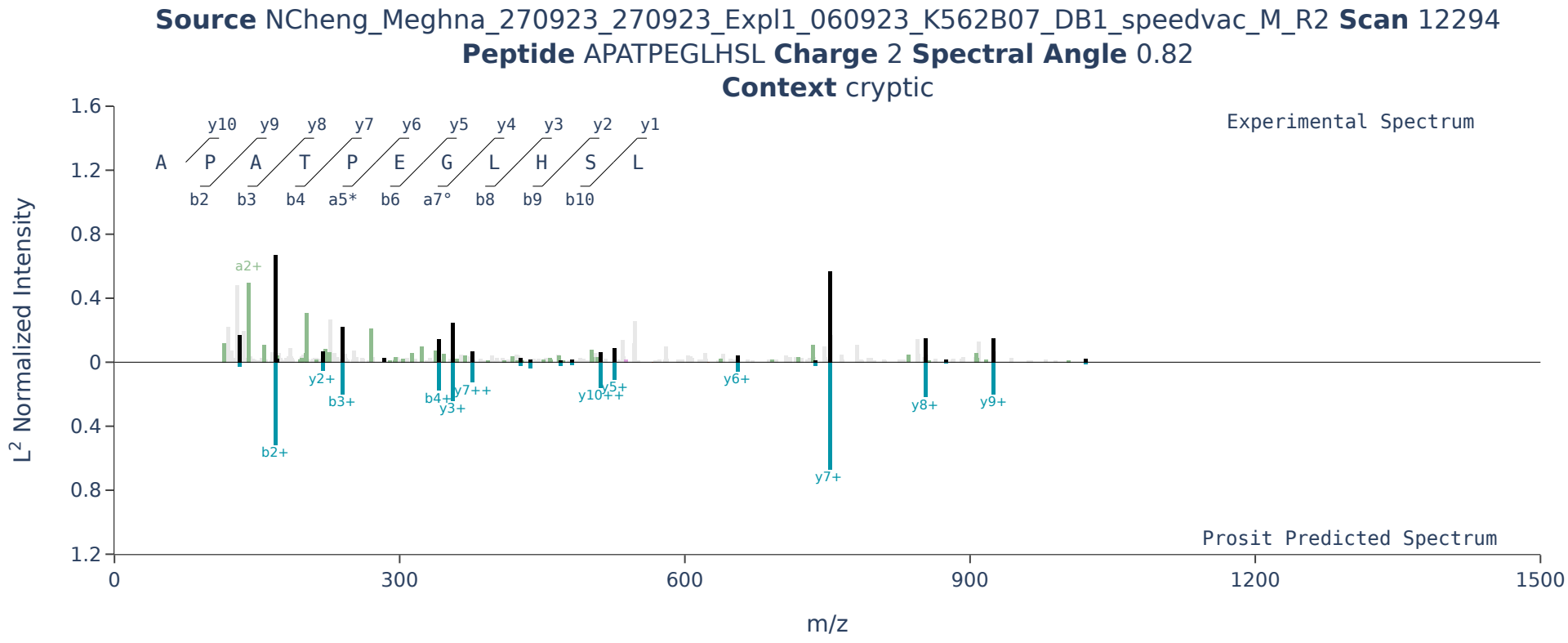
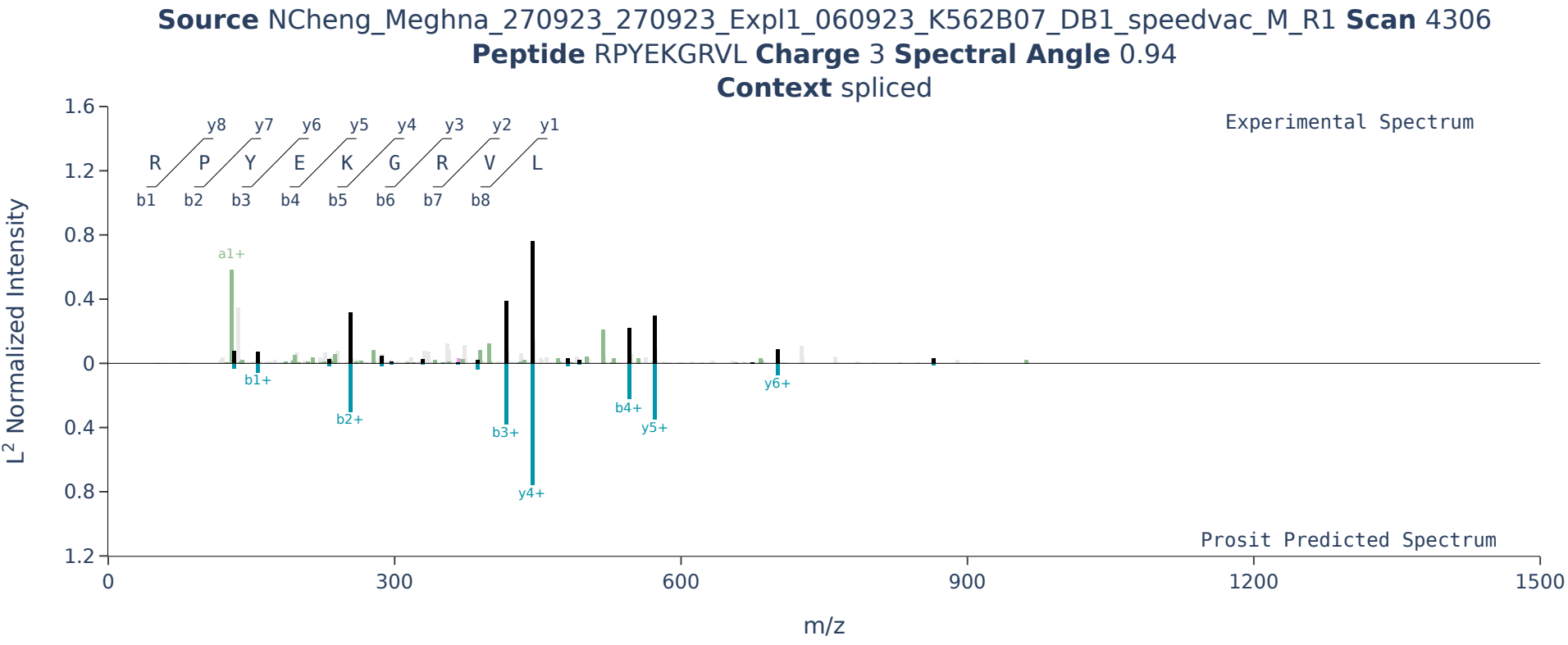
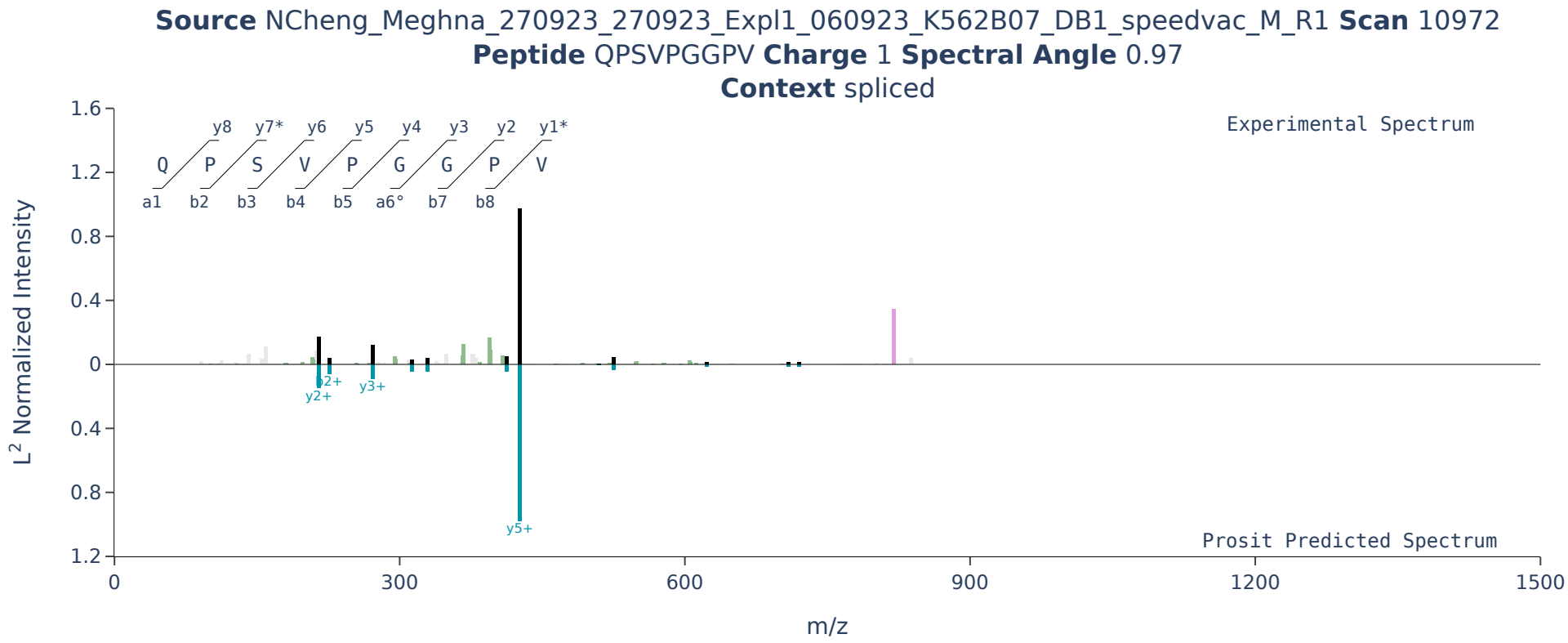
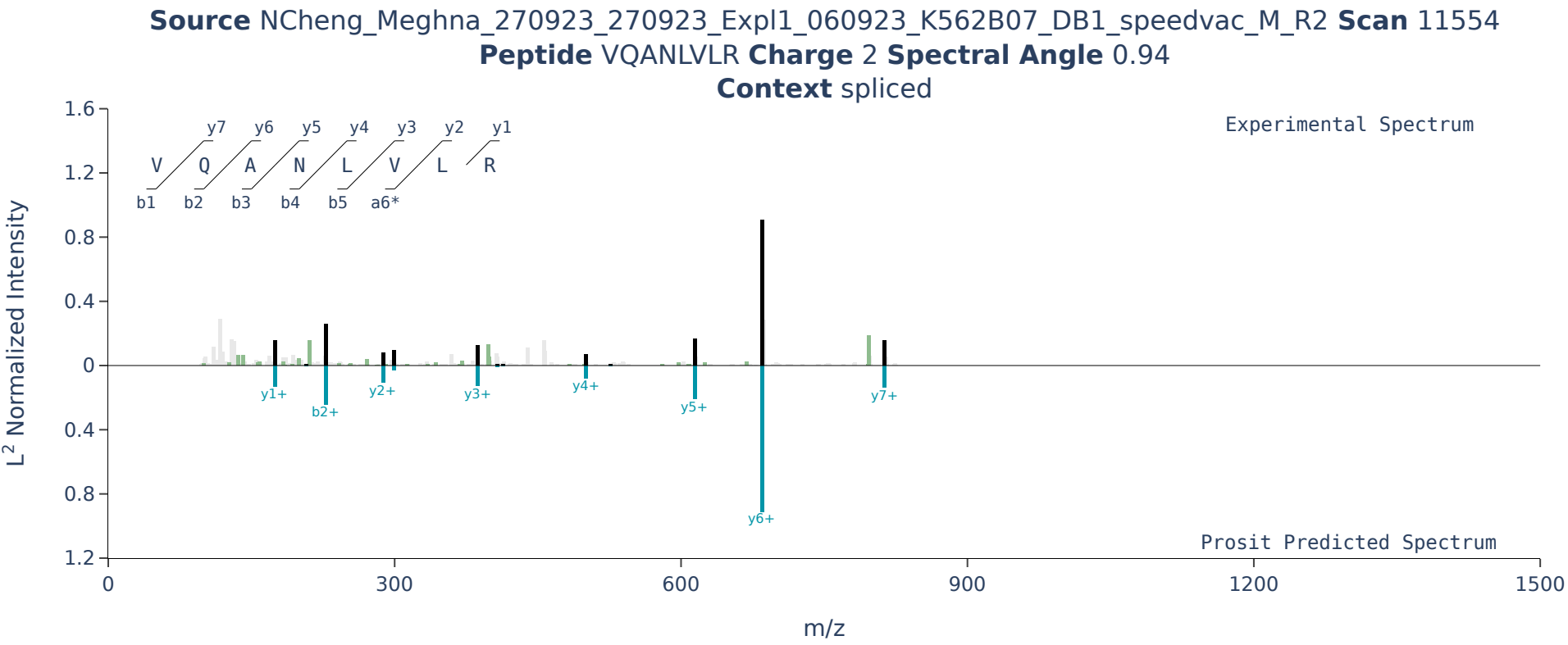
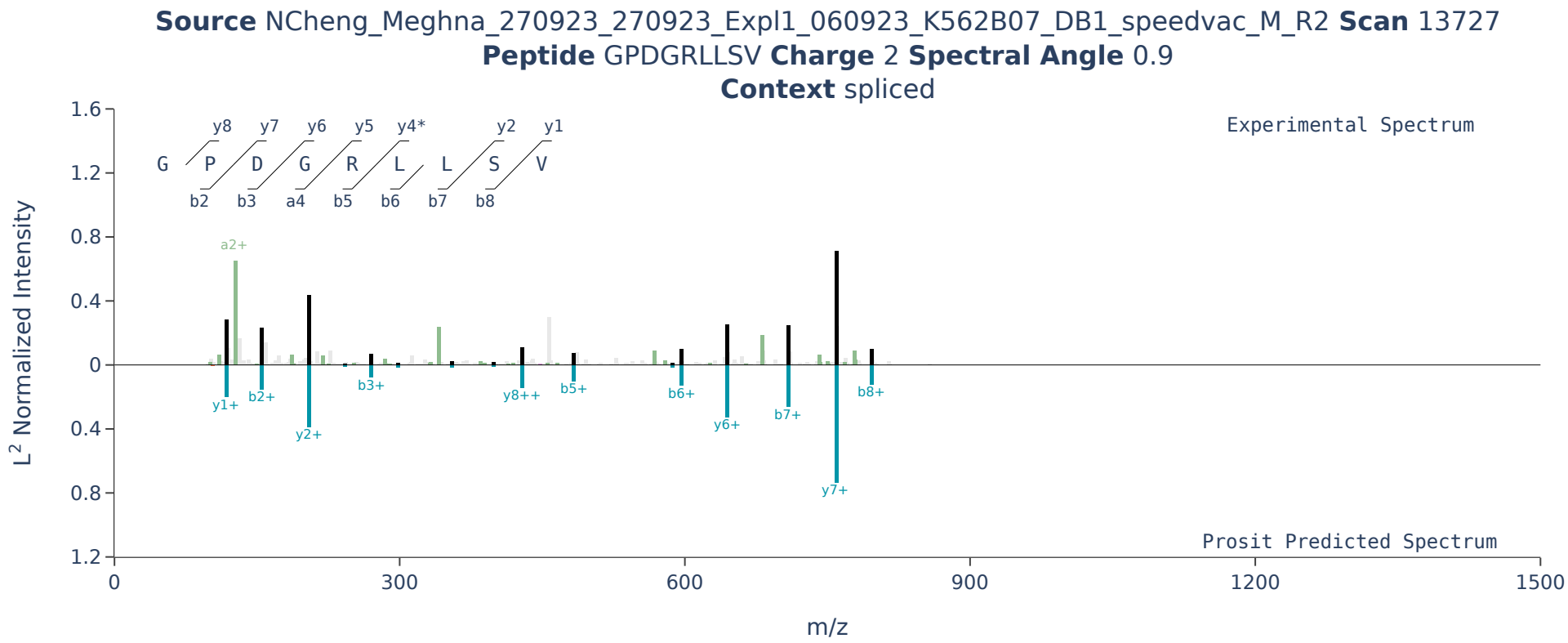
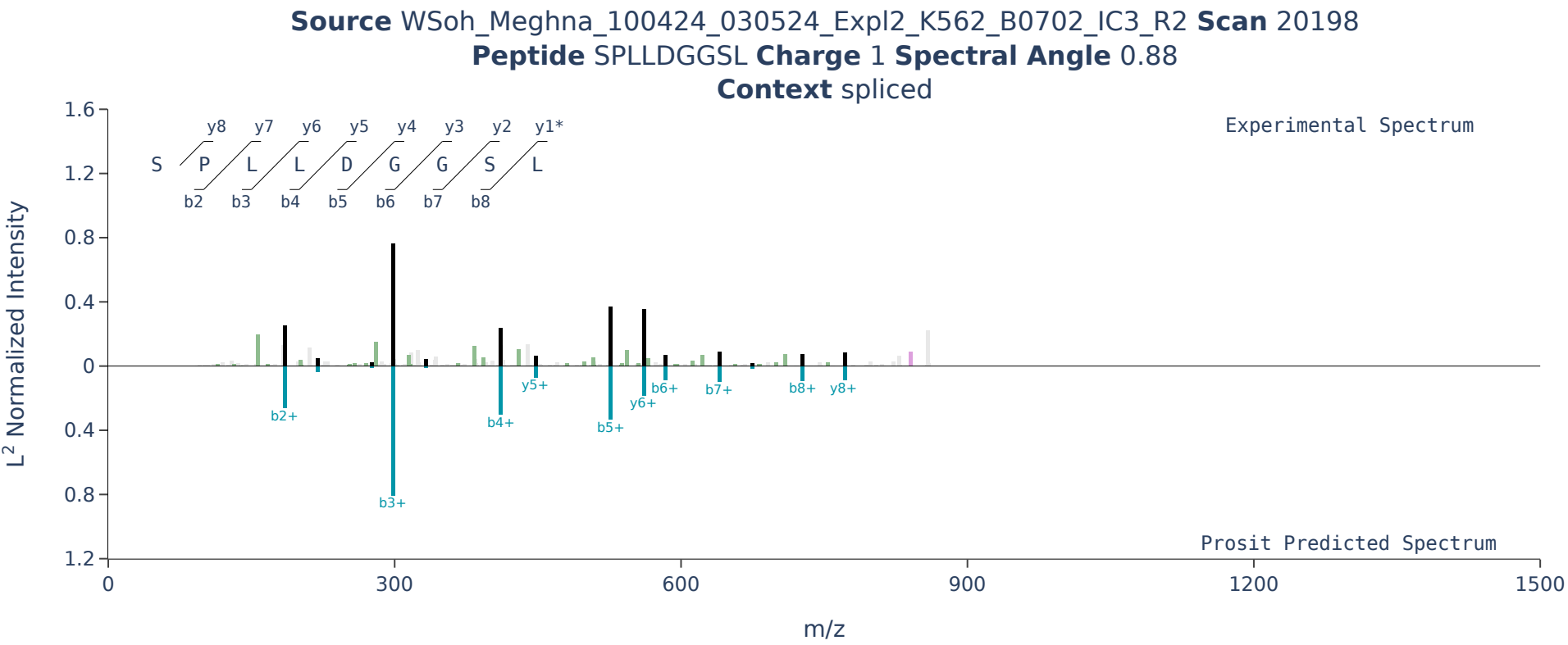
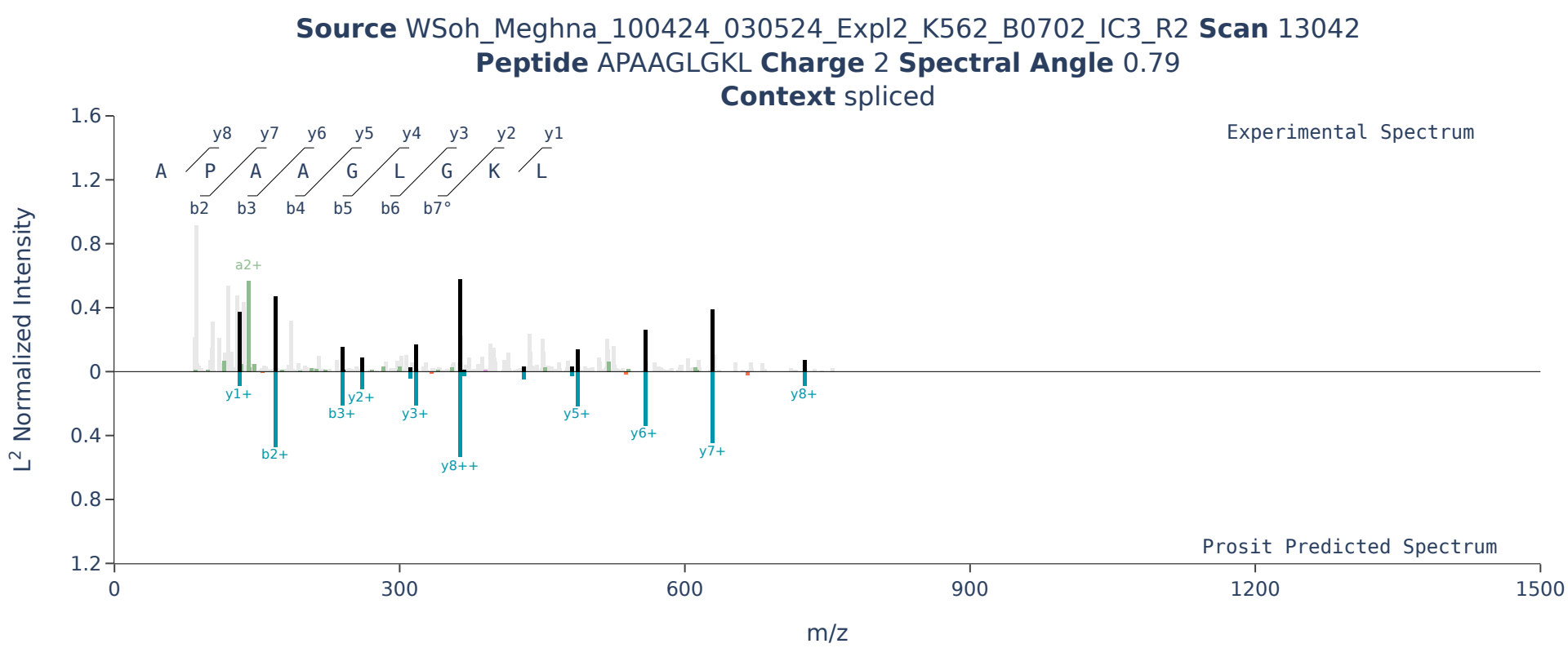
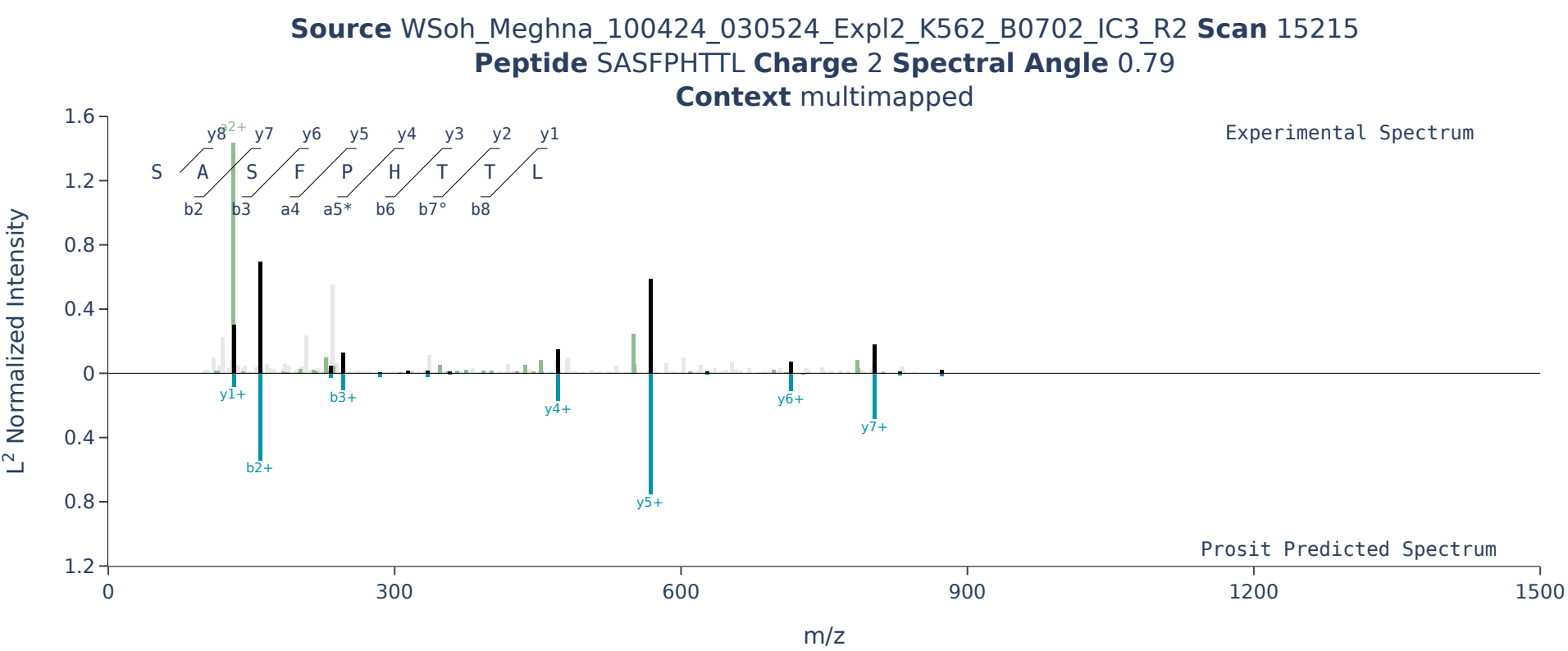
- ° indicates an ion with loss of H₂O.
- * indicates an ion with loss of NH₃.

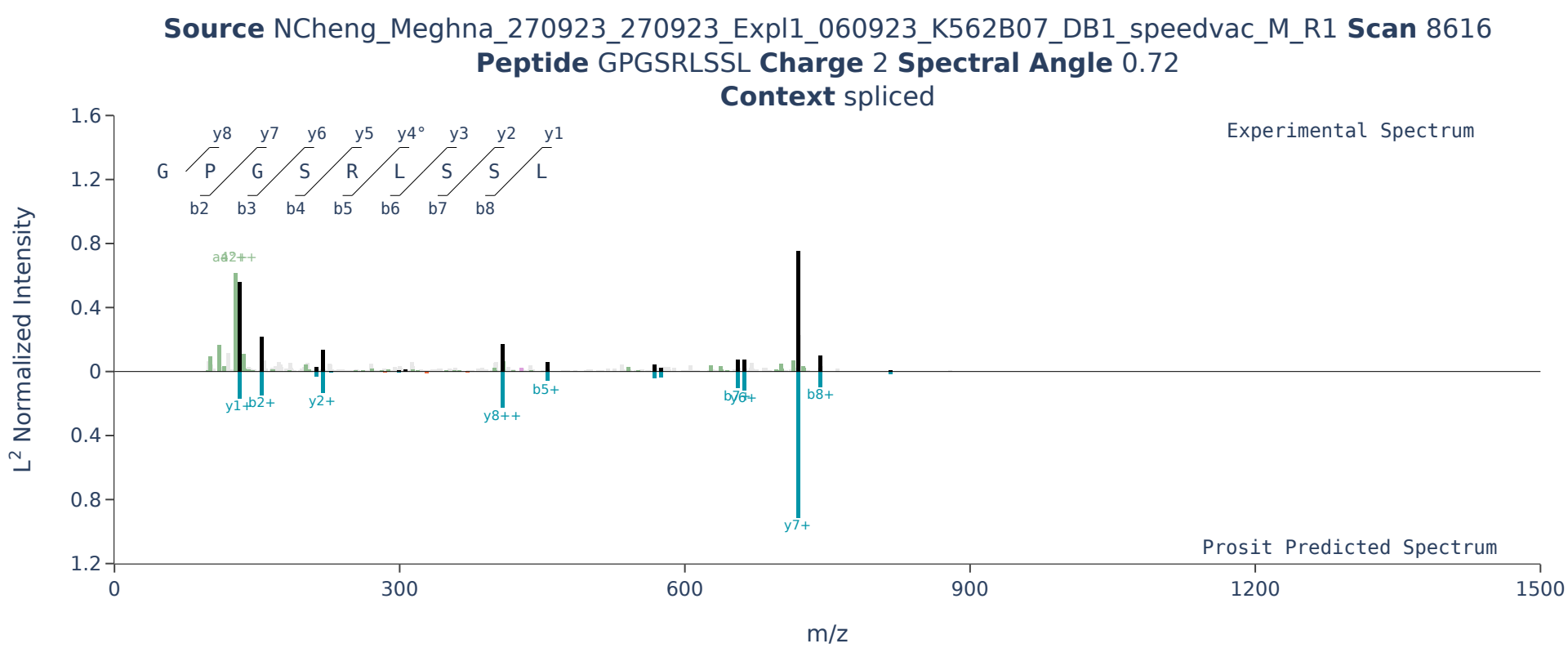
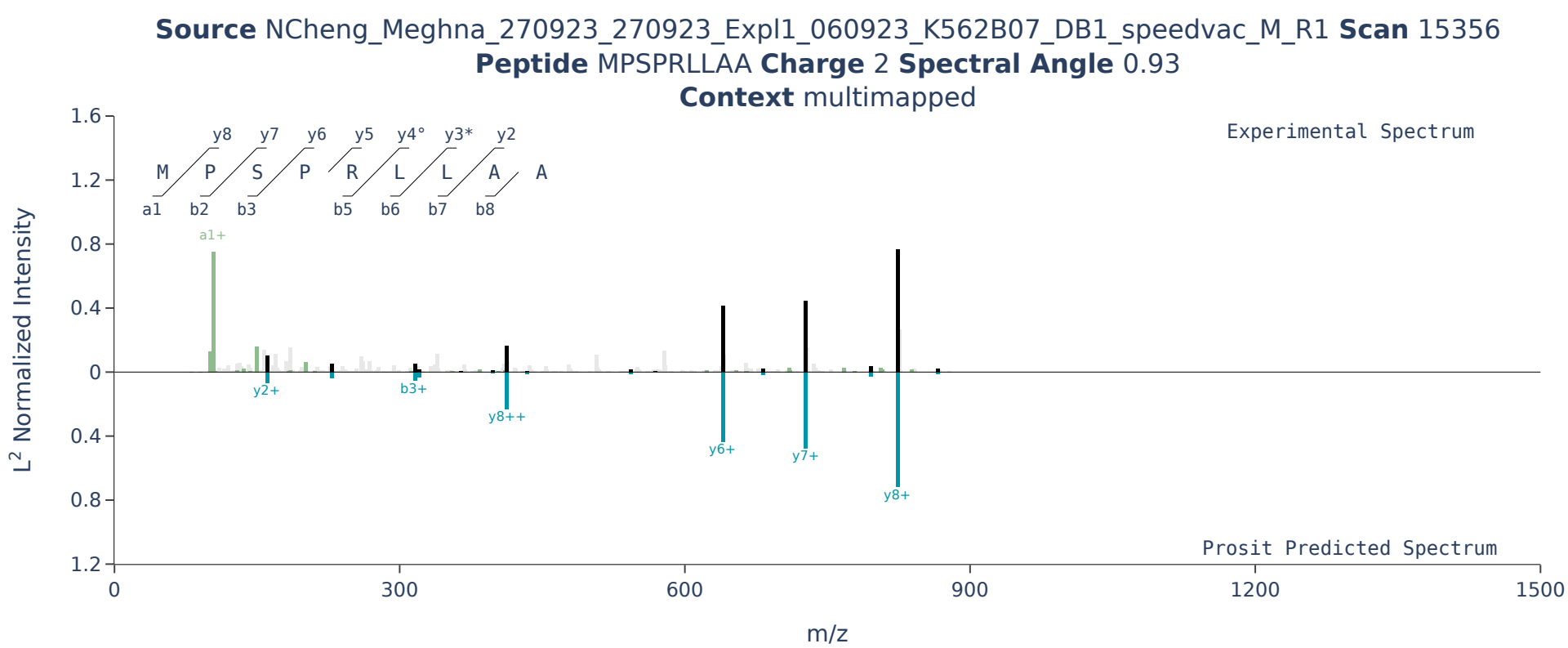
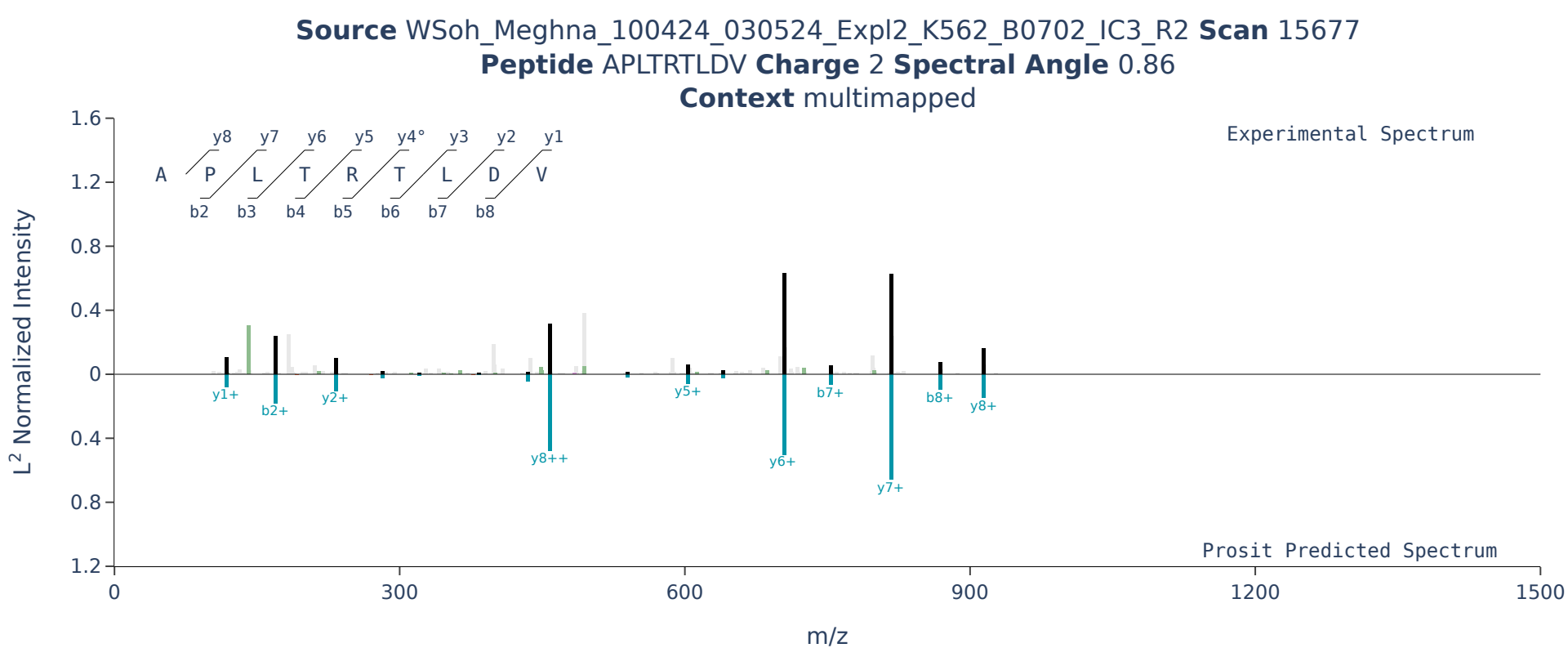
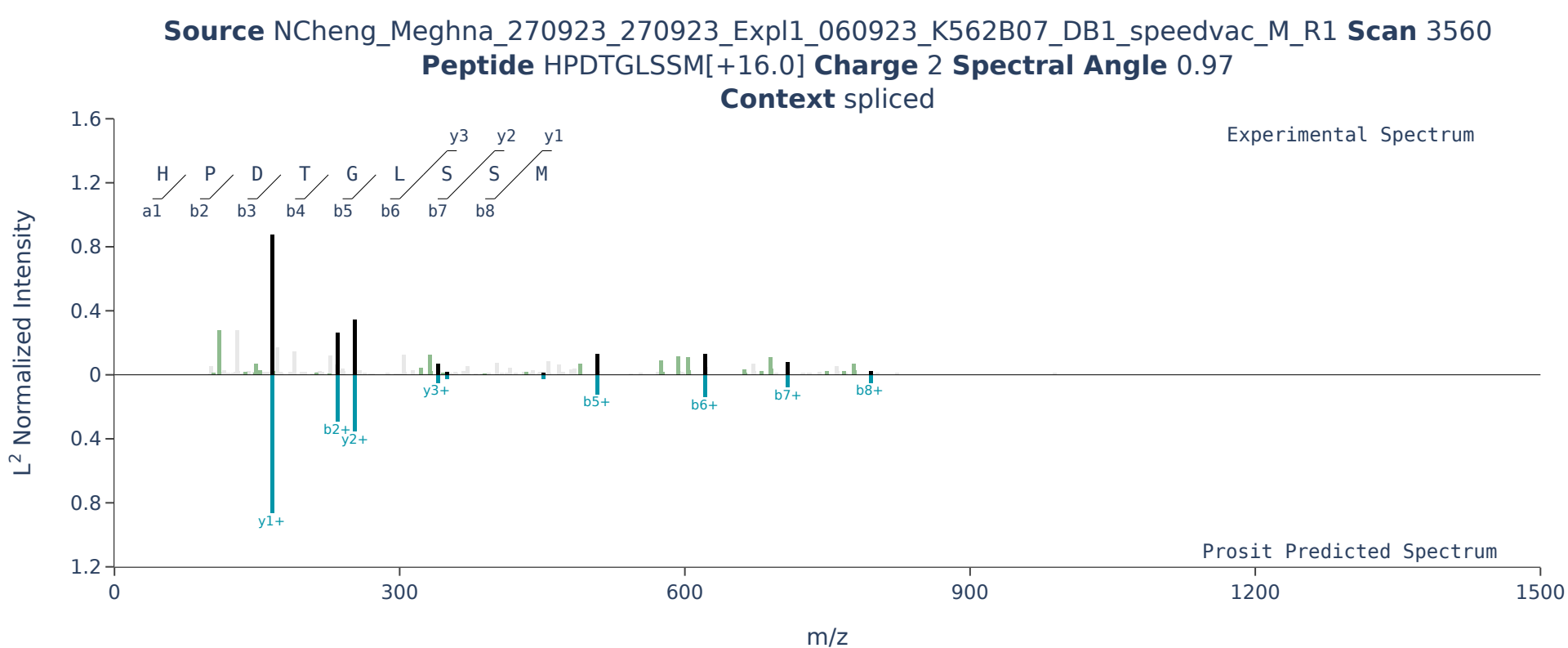
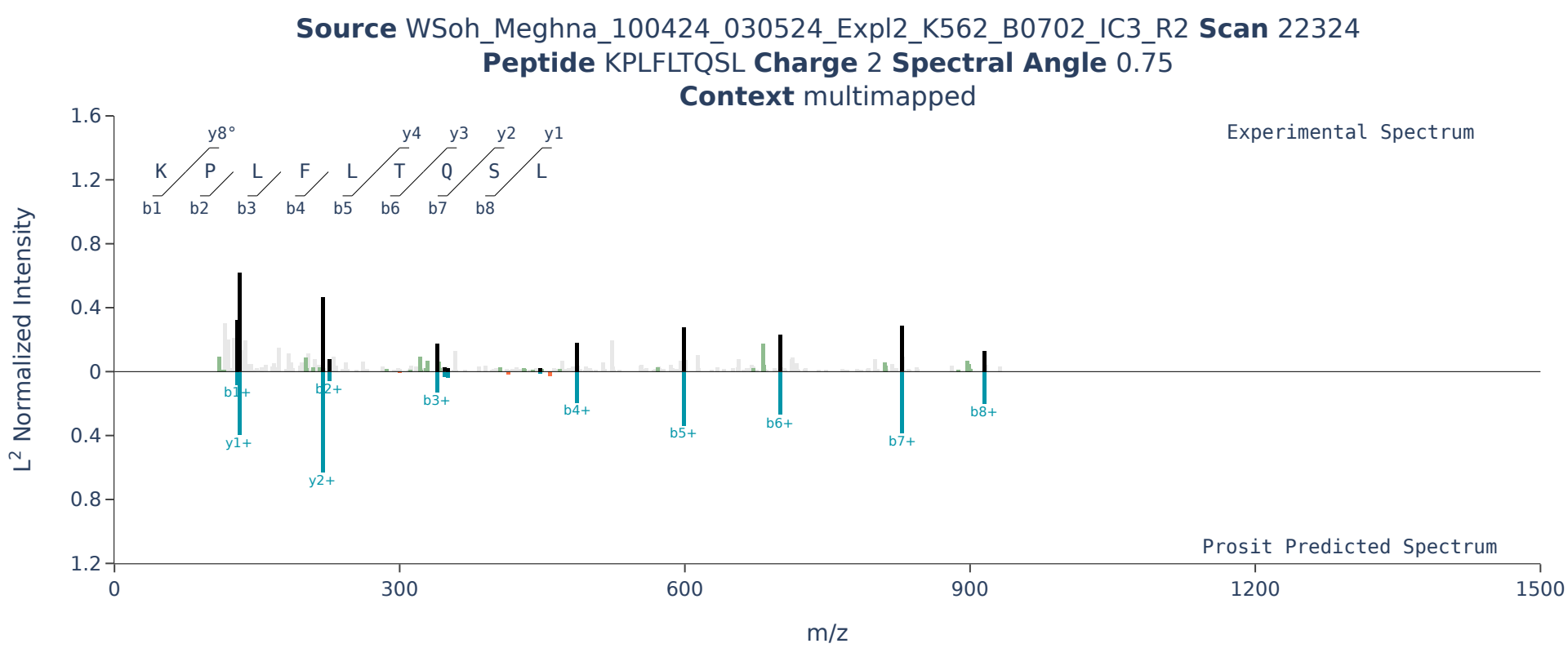


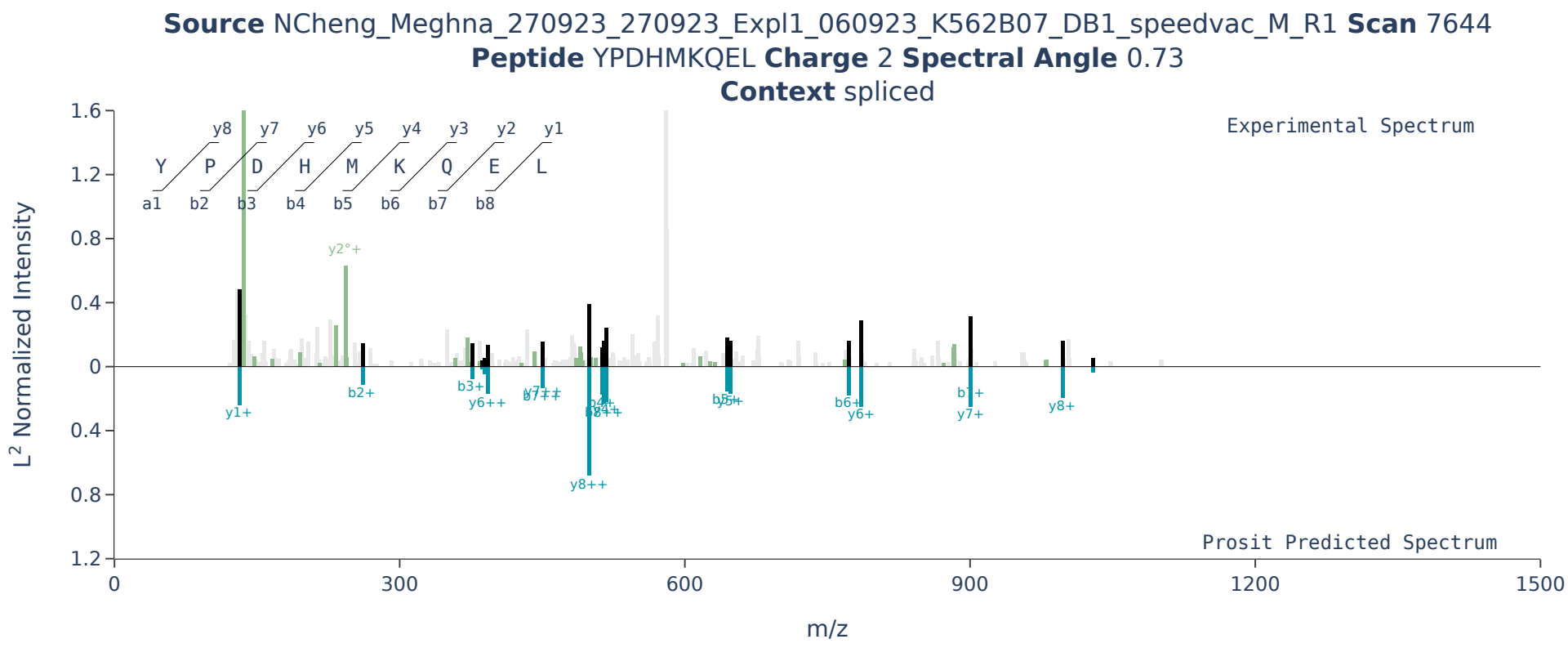
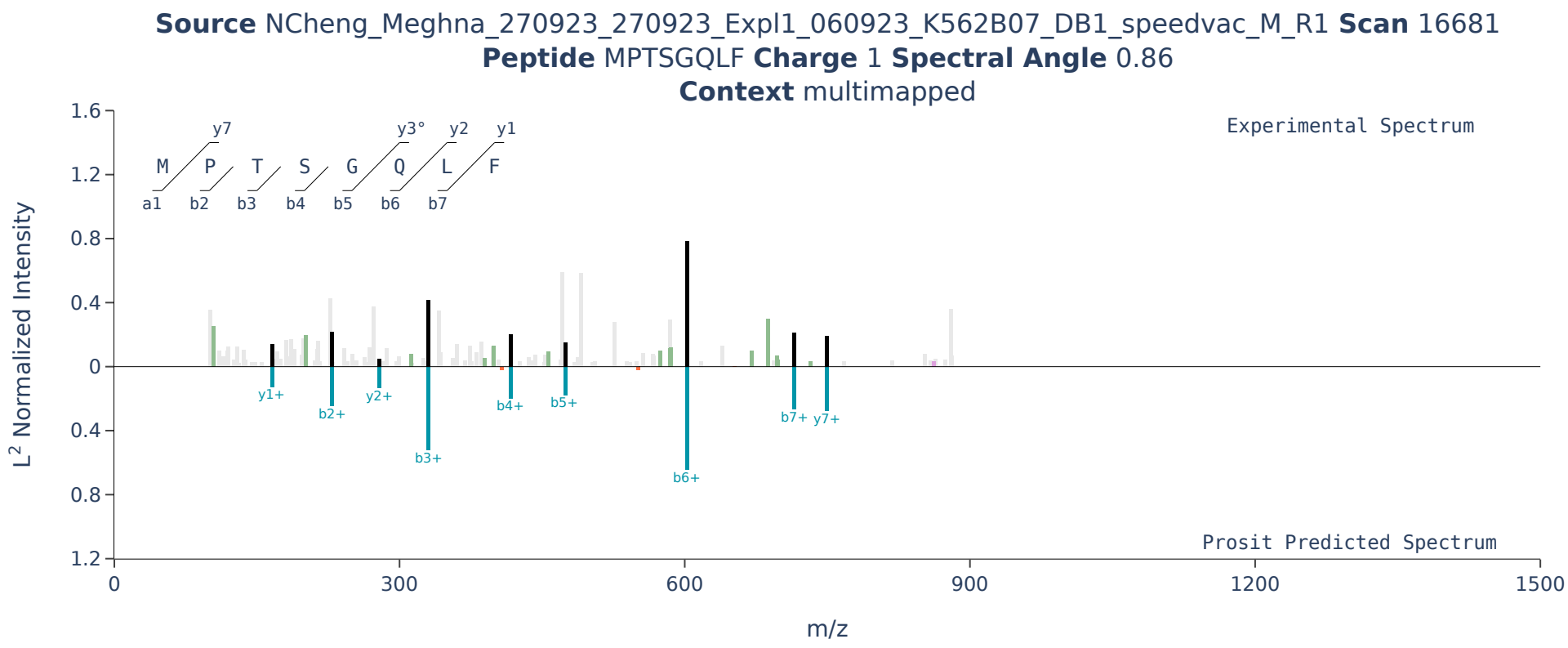
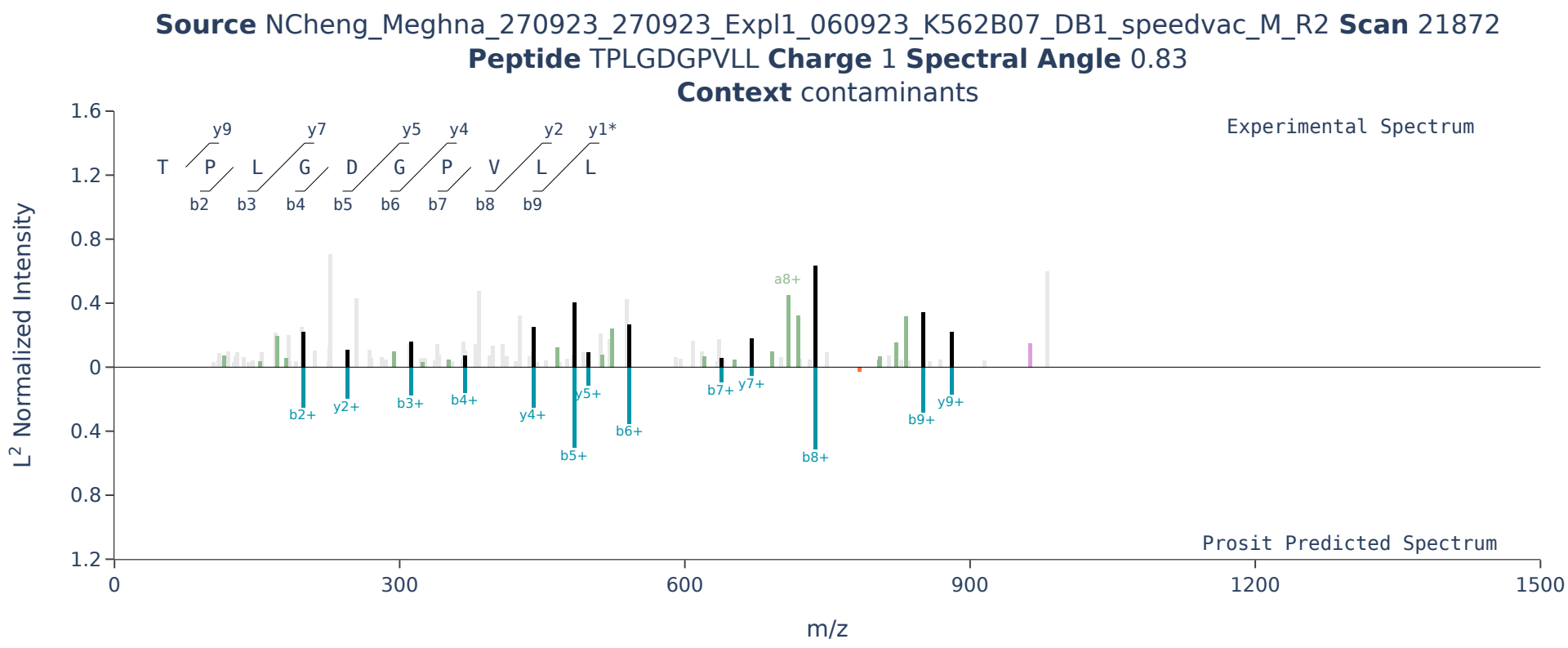
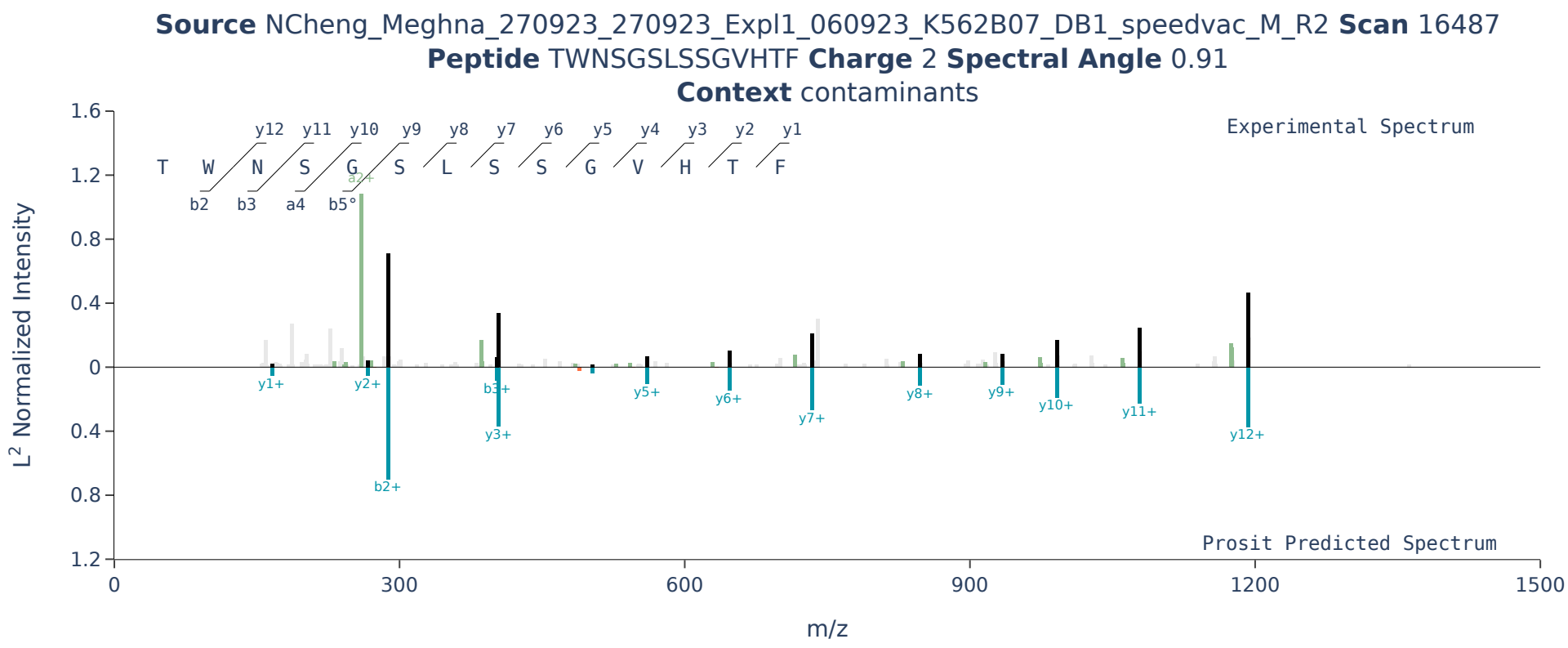
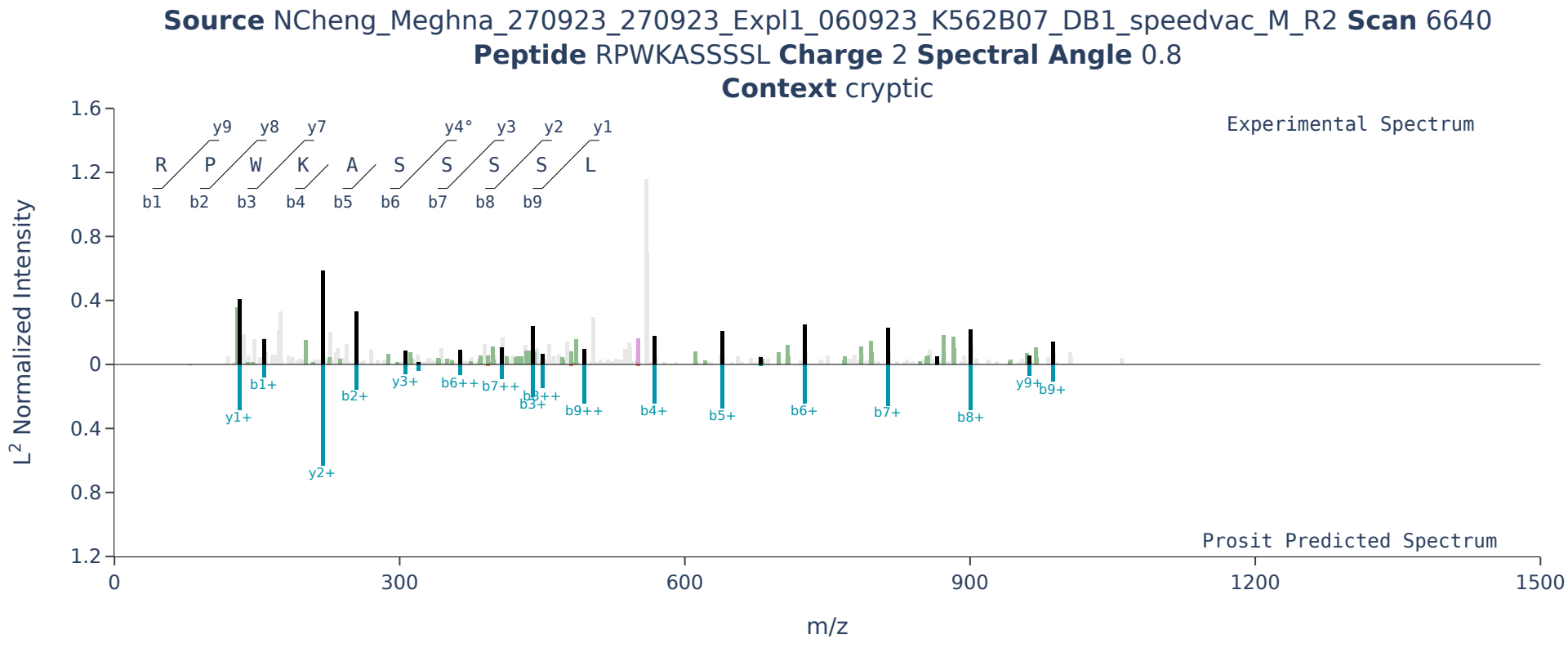
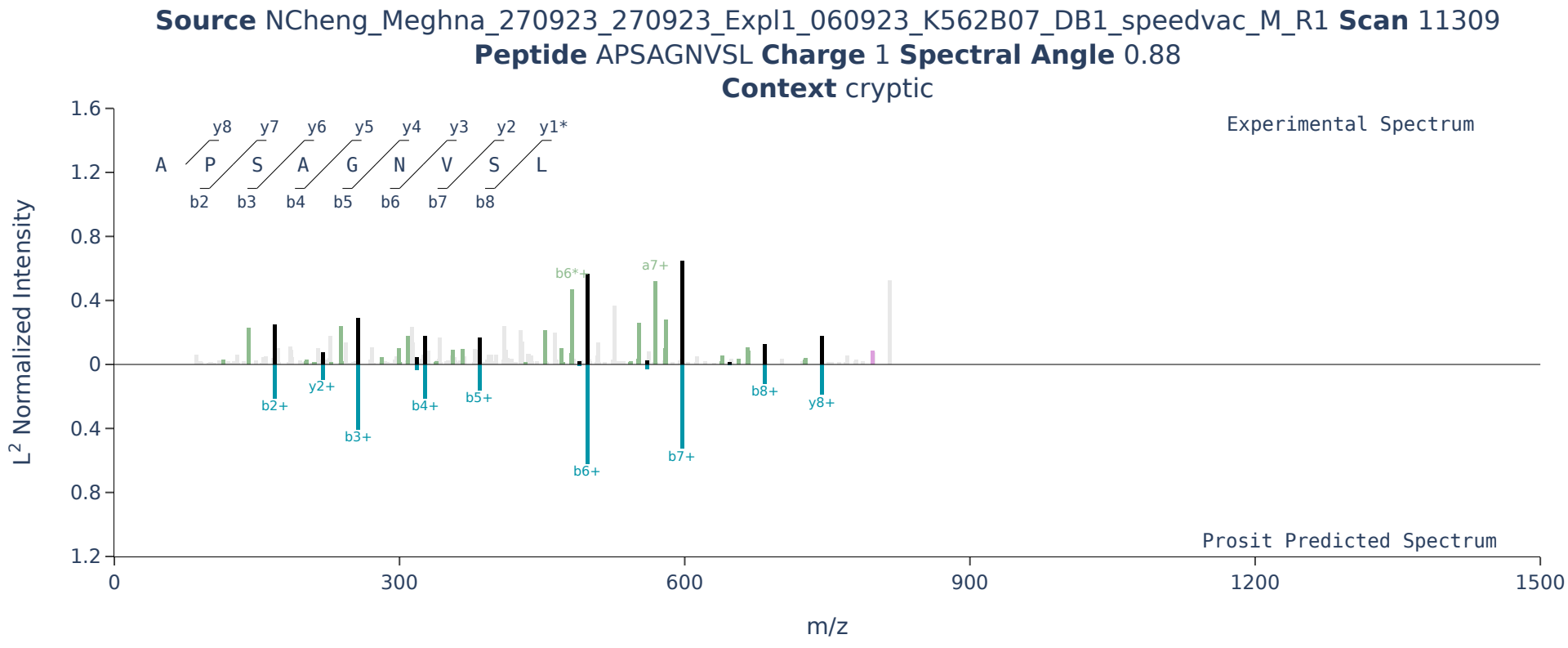
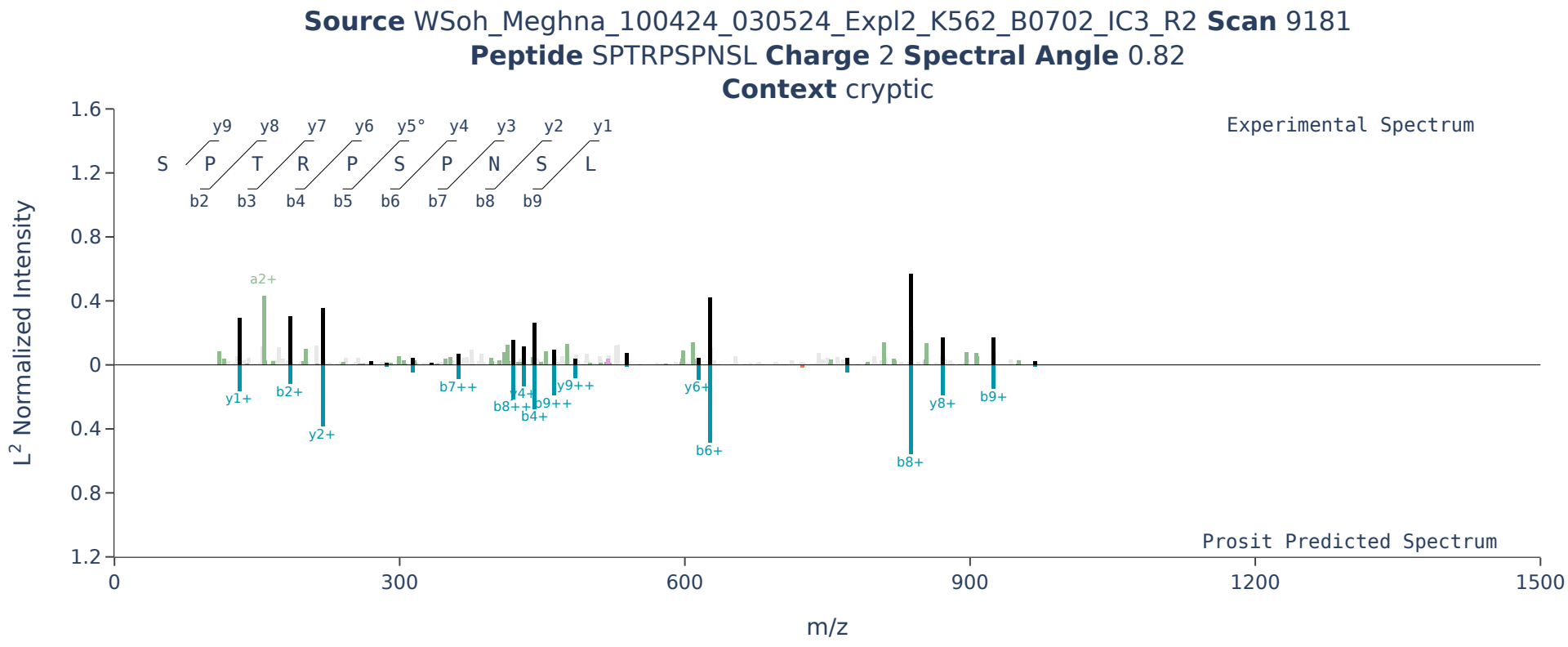
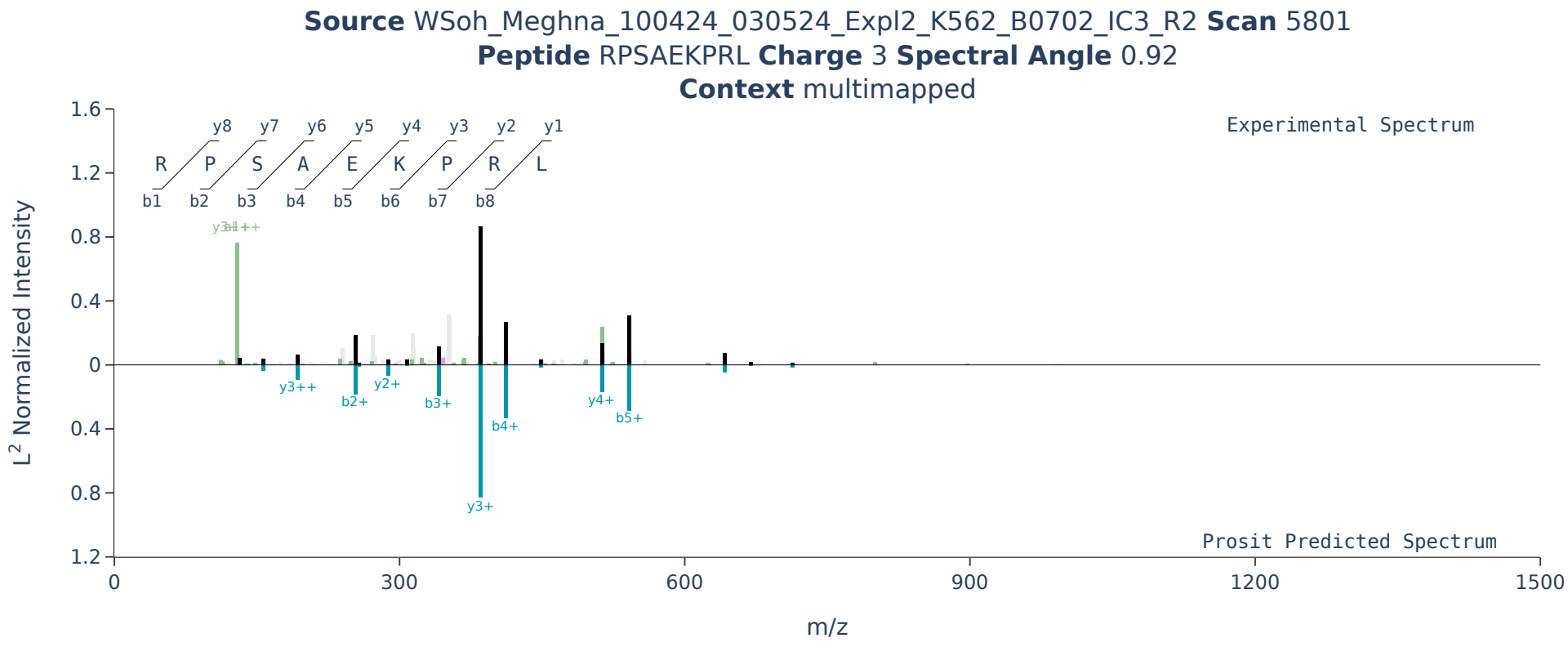
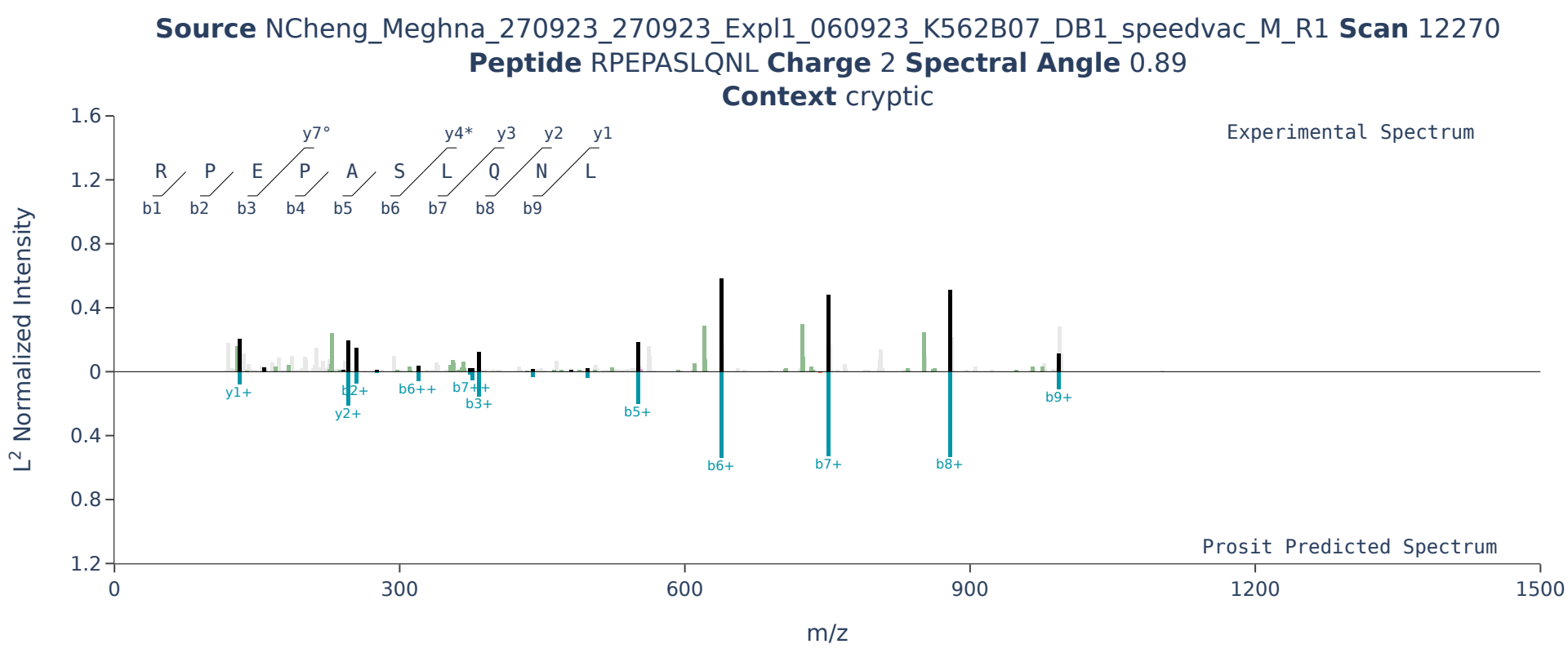
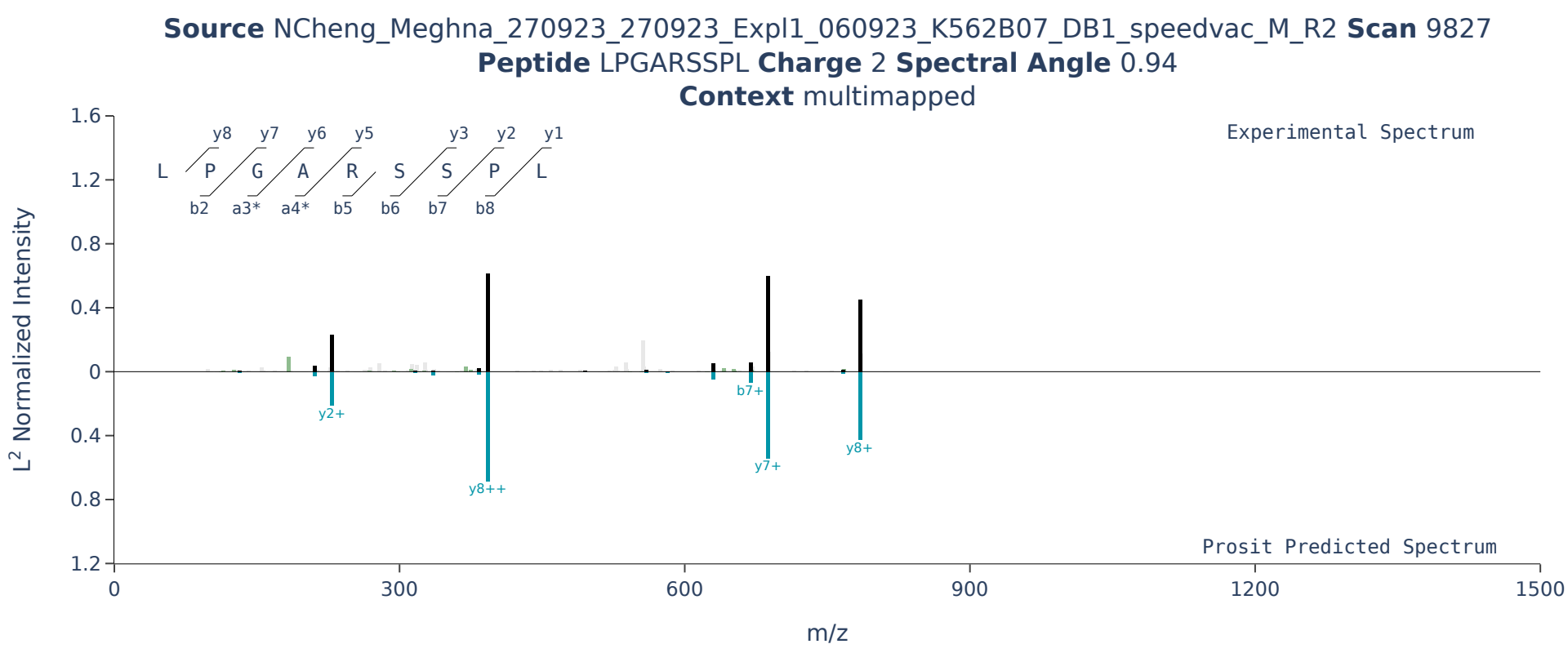




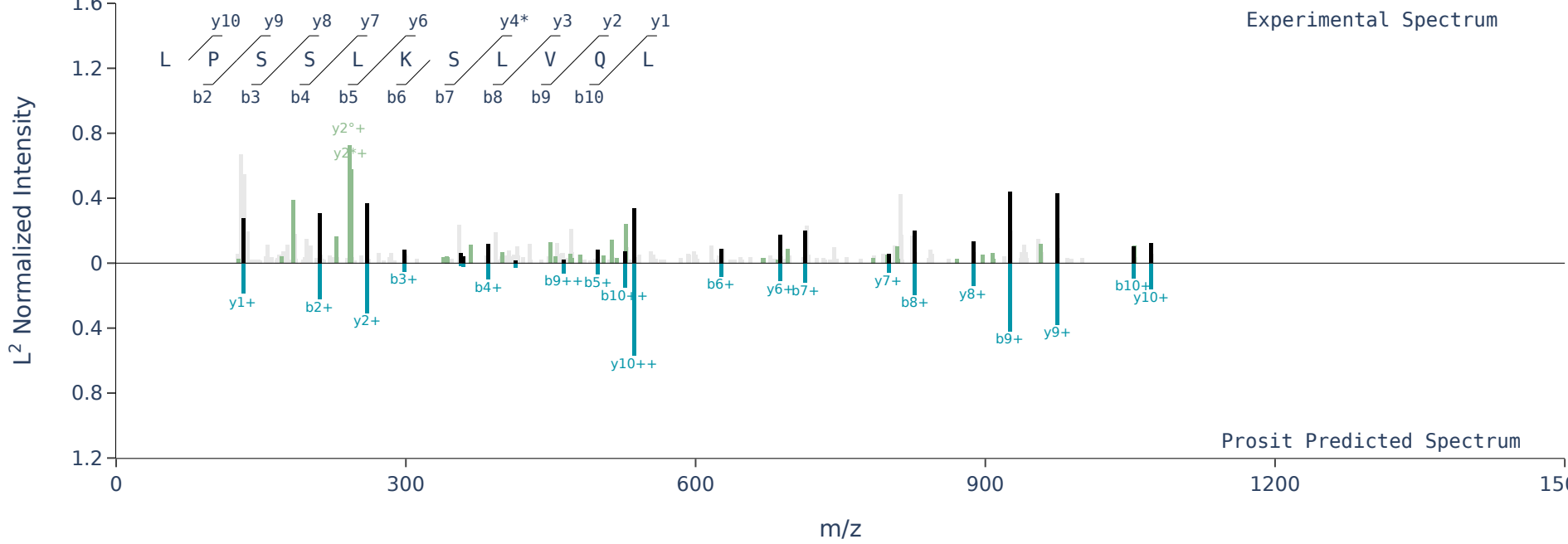




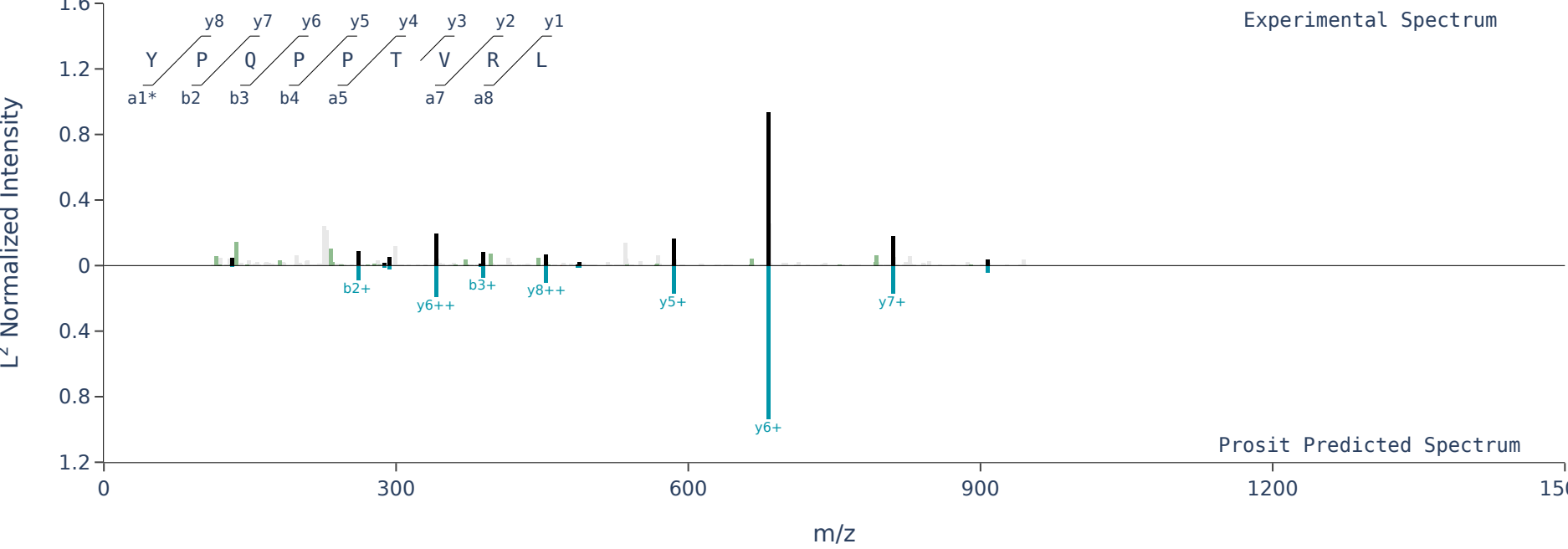




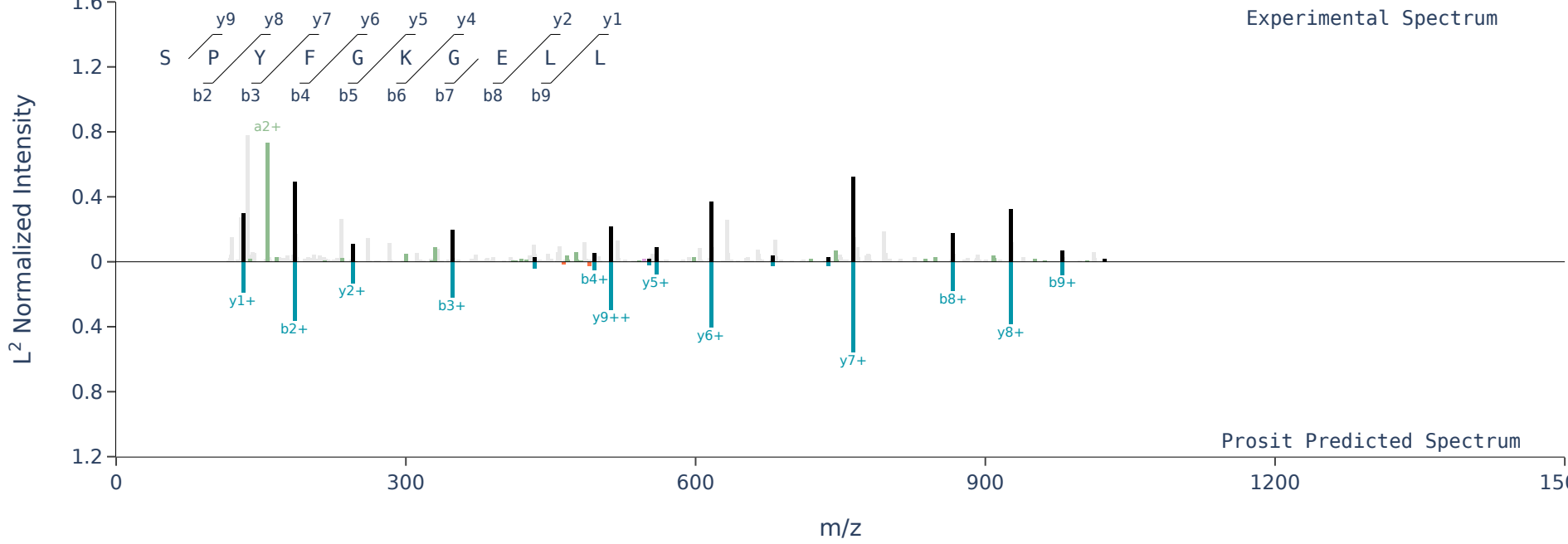
Peptide LPSSLKSLVQL **Charge** 2 **Spectral Angle** 0.8
Context cryptic



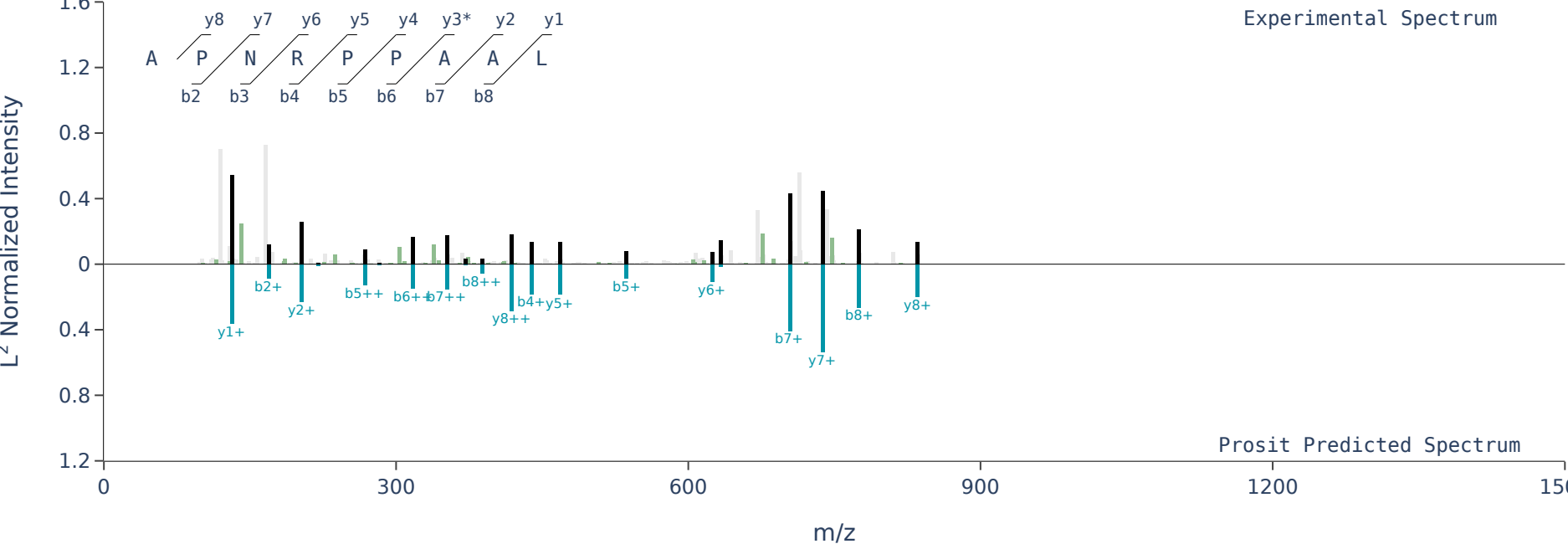
Peptide YPQPPTVRL **Charge** 2 **Spectral Angle** 0.96
Context multimapped



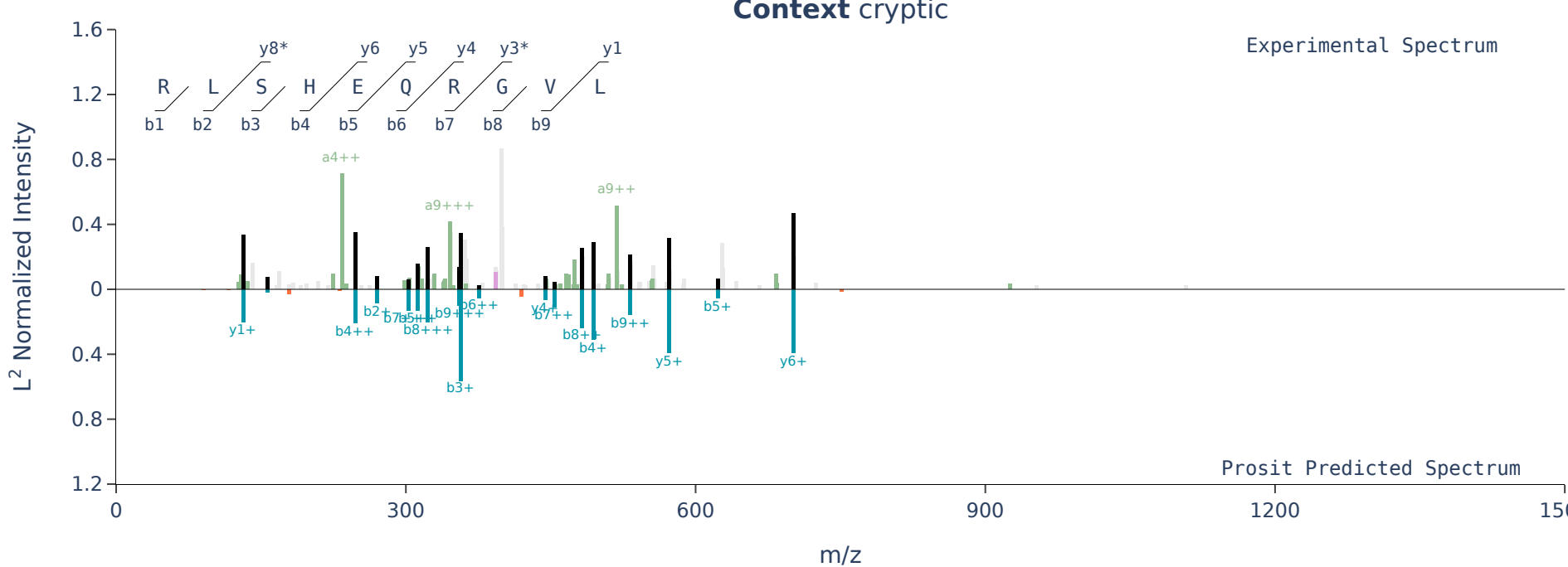
Peptide SPYFGKGELL **Charge** 2 **Spectral Angle** 0.87
Context unmapped



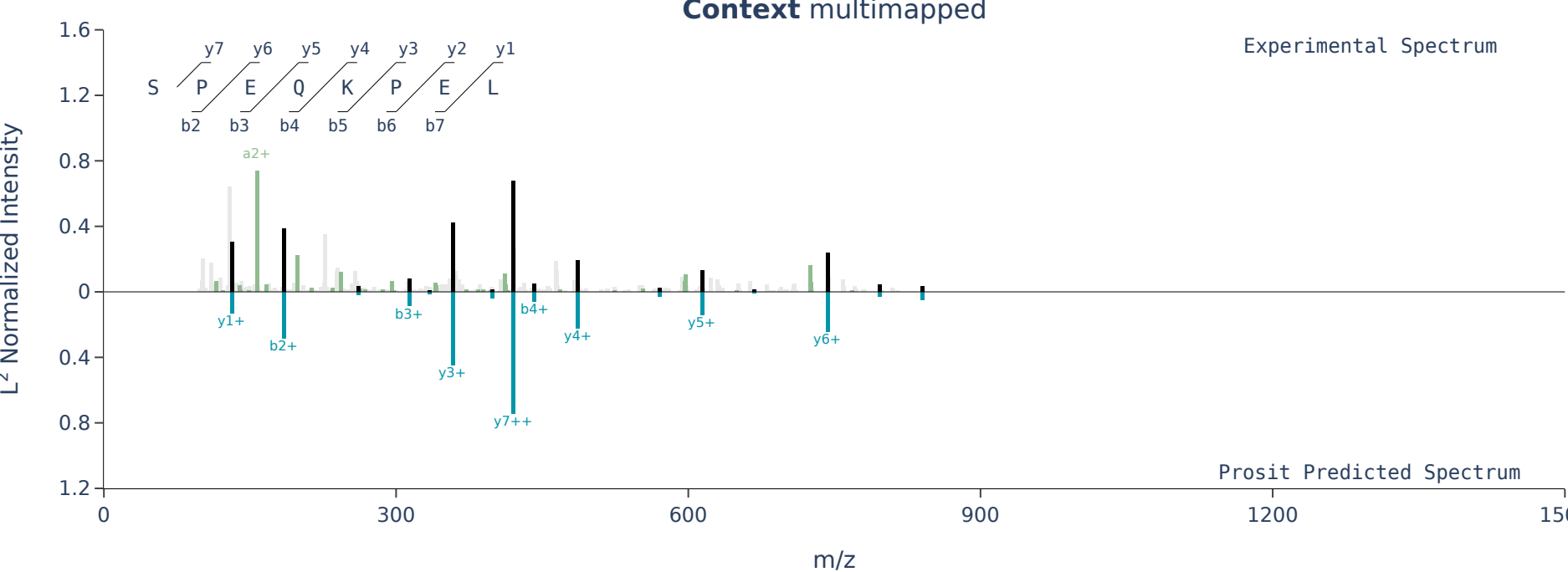
Peptide APNRPPAAL **Charge** 2 **Spectral Angle** 0.81
Context multimapped



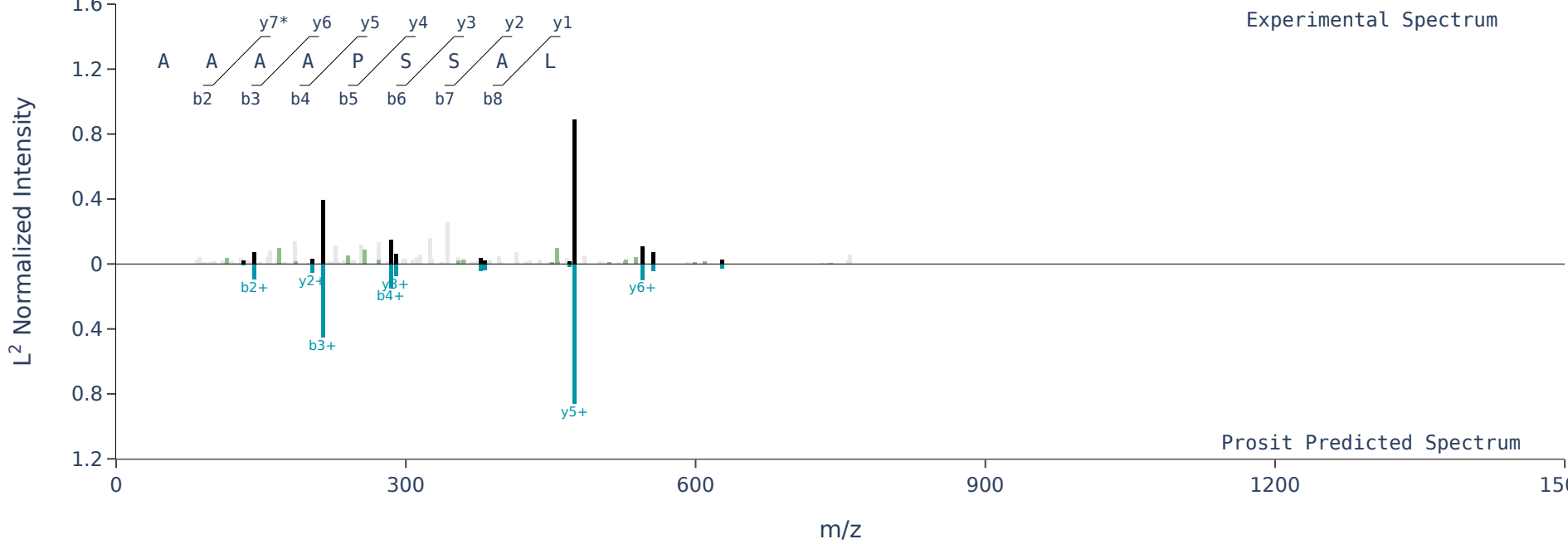
Peptide RLSHEQRGVL Charge 3 Spectral Angle 0.78



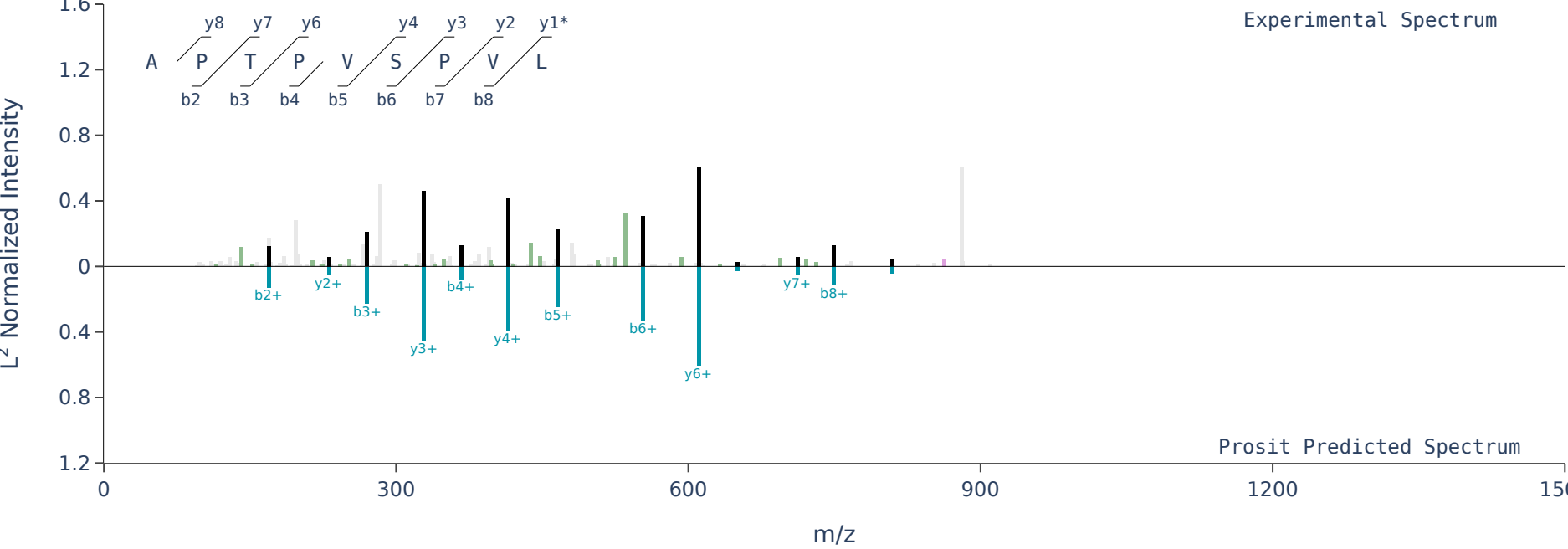
Peptide SPEQKPEL Charge 2 Spectral Angle 0.86
Context multimapped



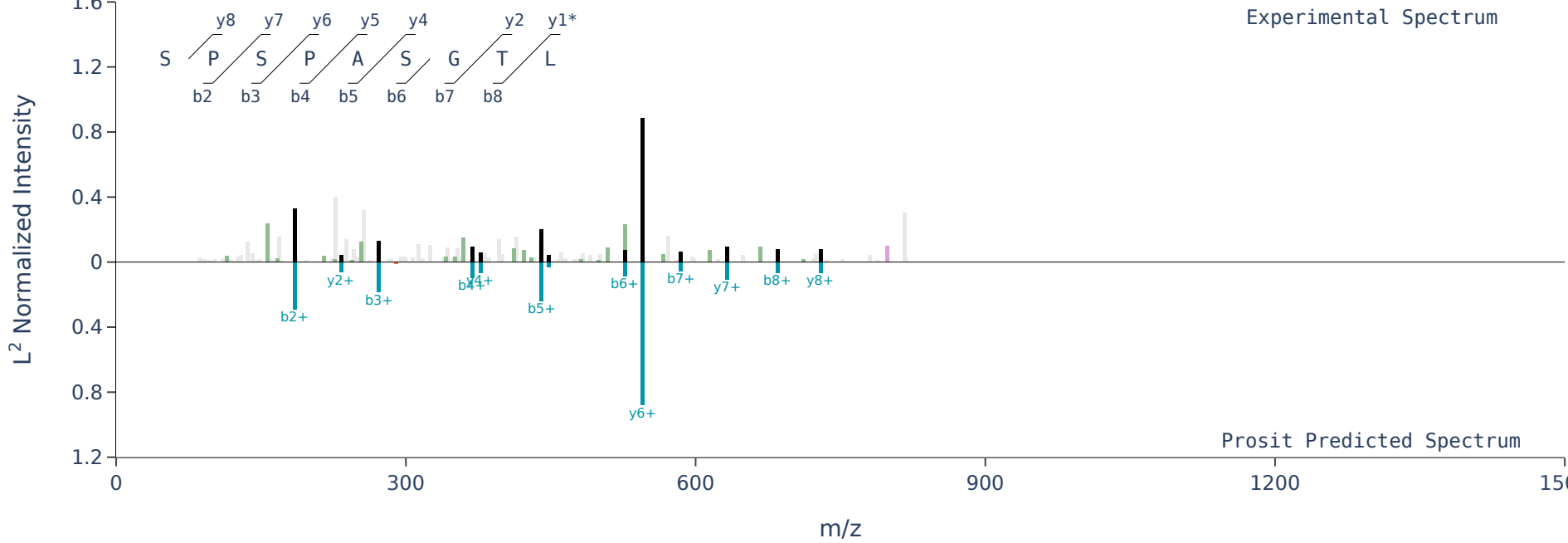
Peptide AAAAPSSAL Charge 1 Spectral Angle 0.95
Context multimapped



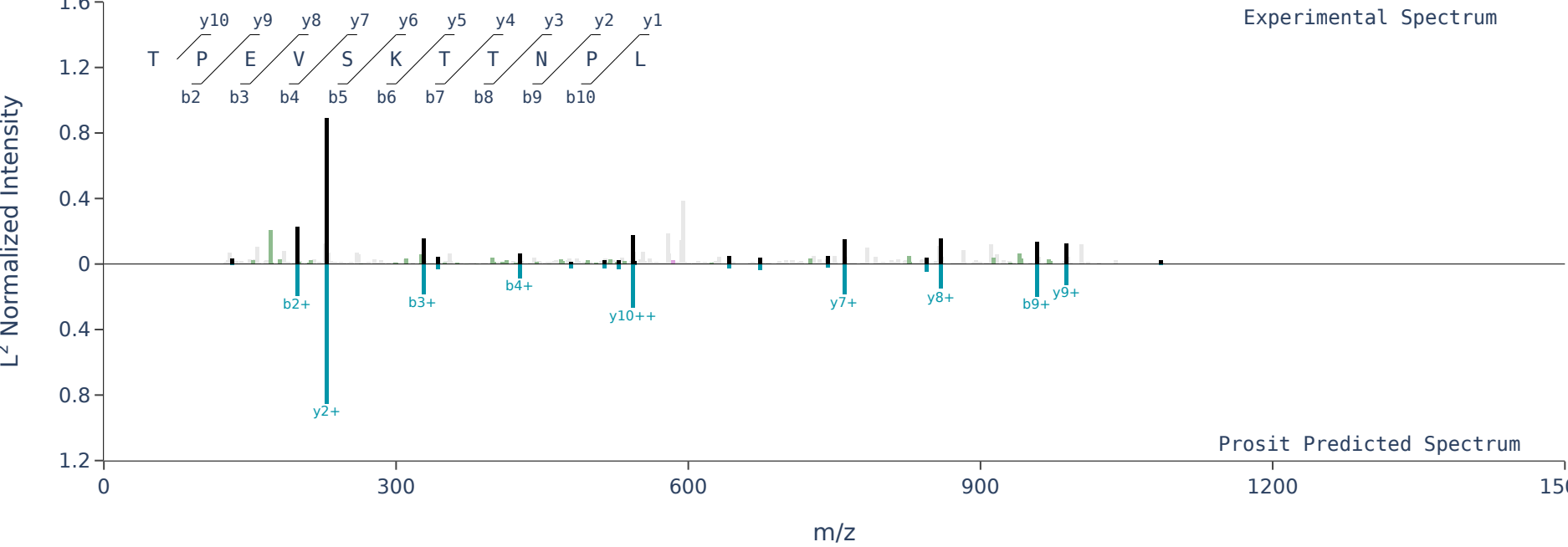
Peptide APTVSPVL **Charge** 1 **Spectral Angle** 0.95
Context multimapped

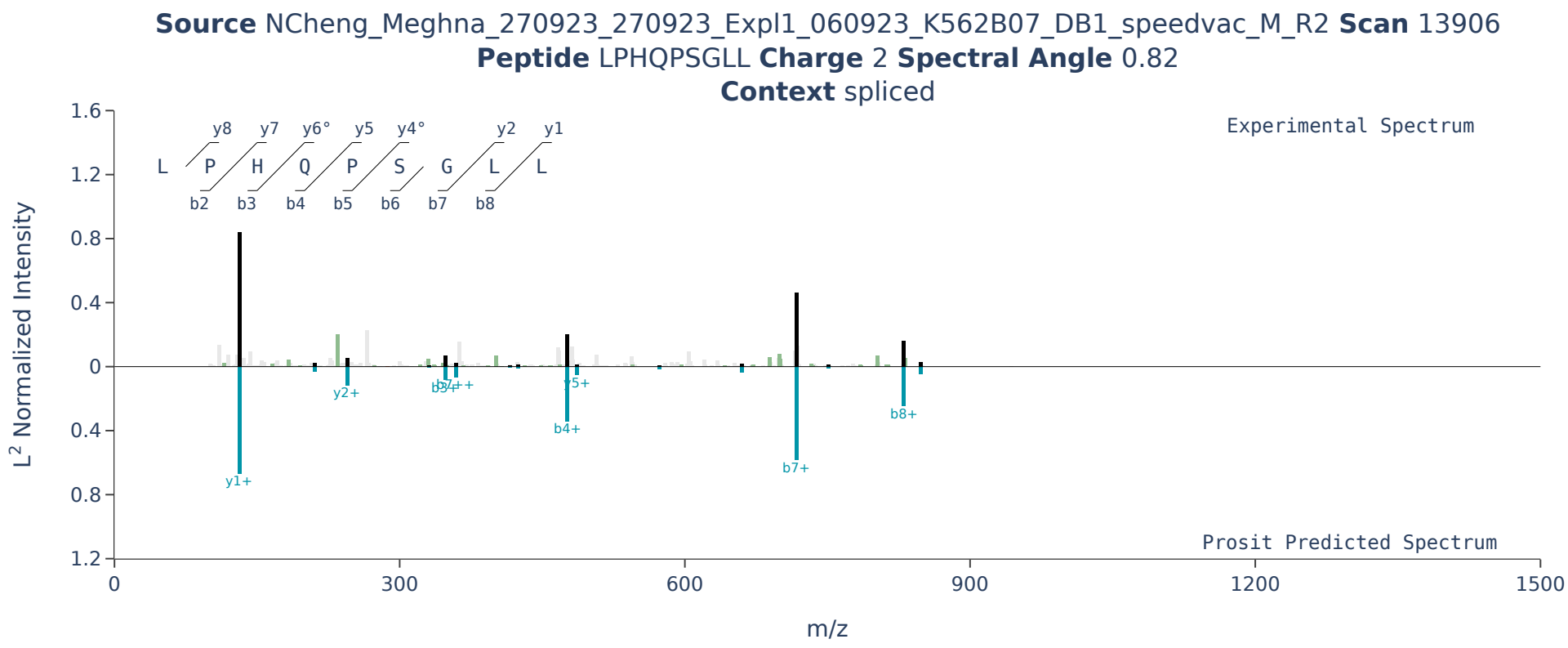
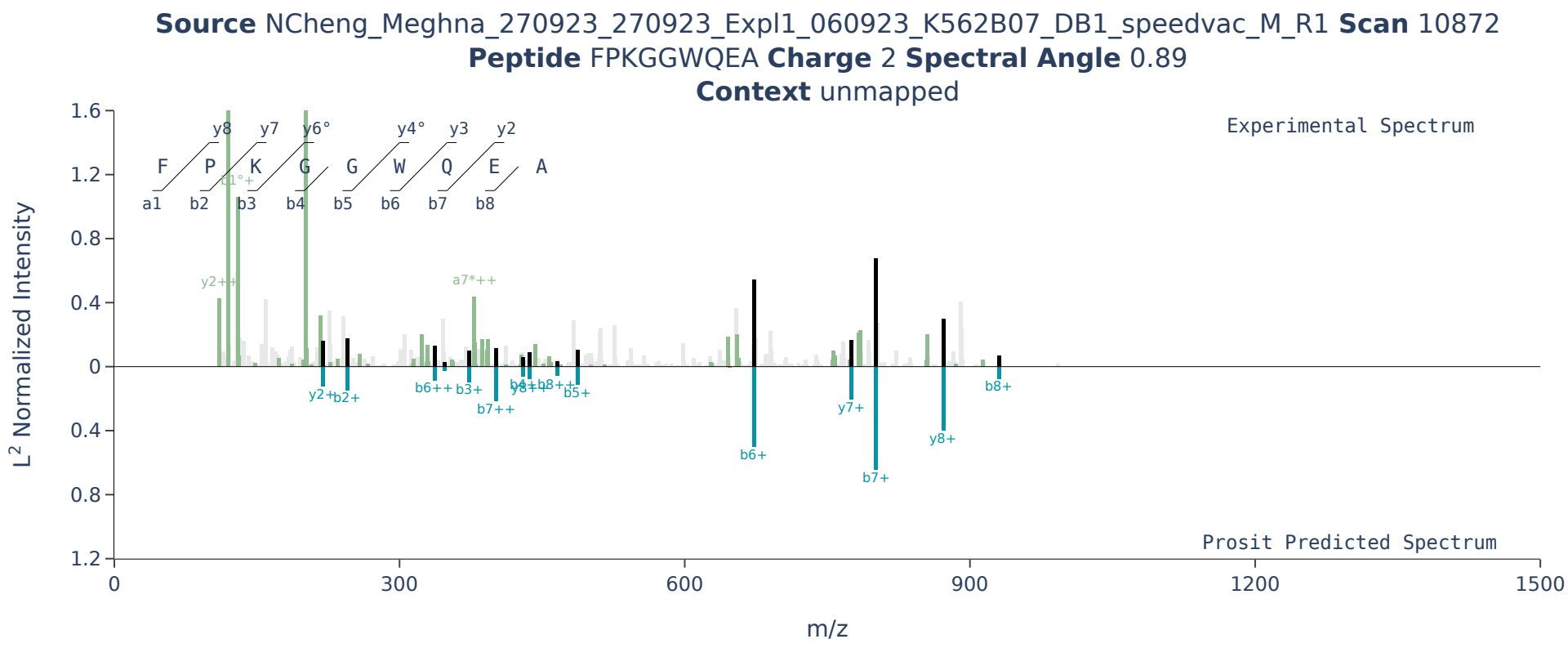
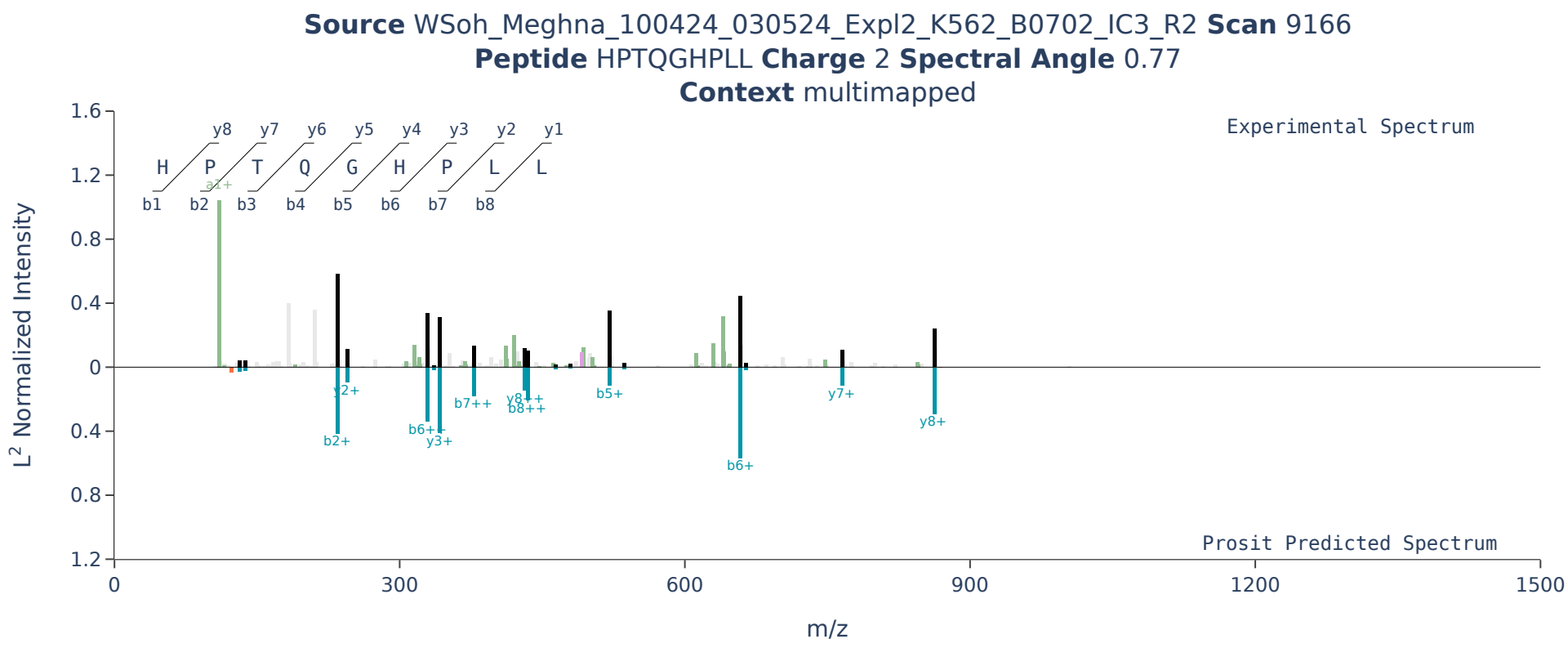
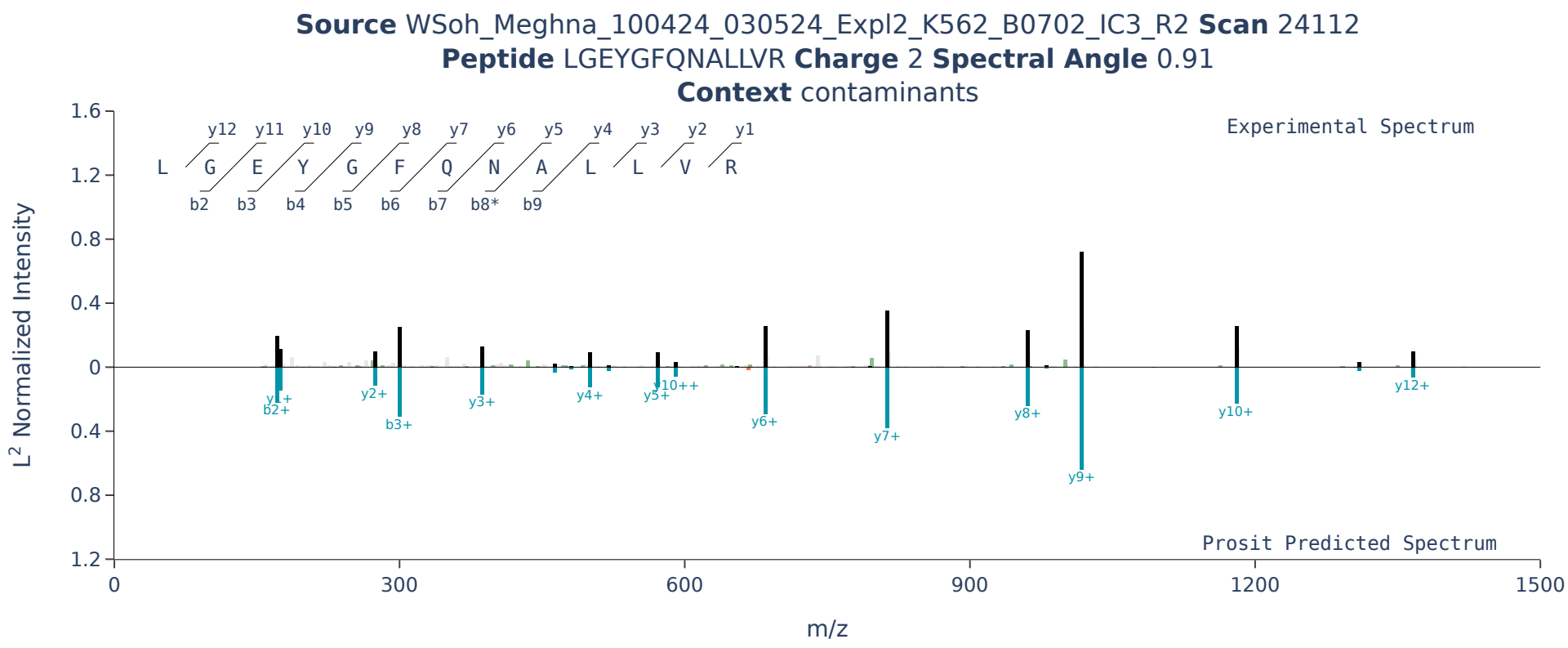
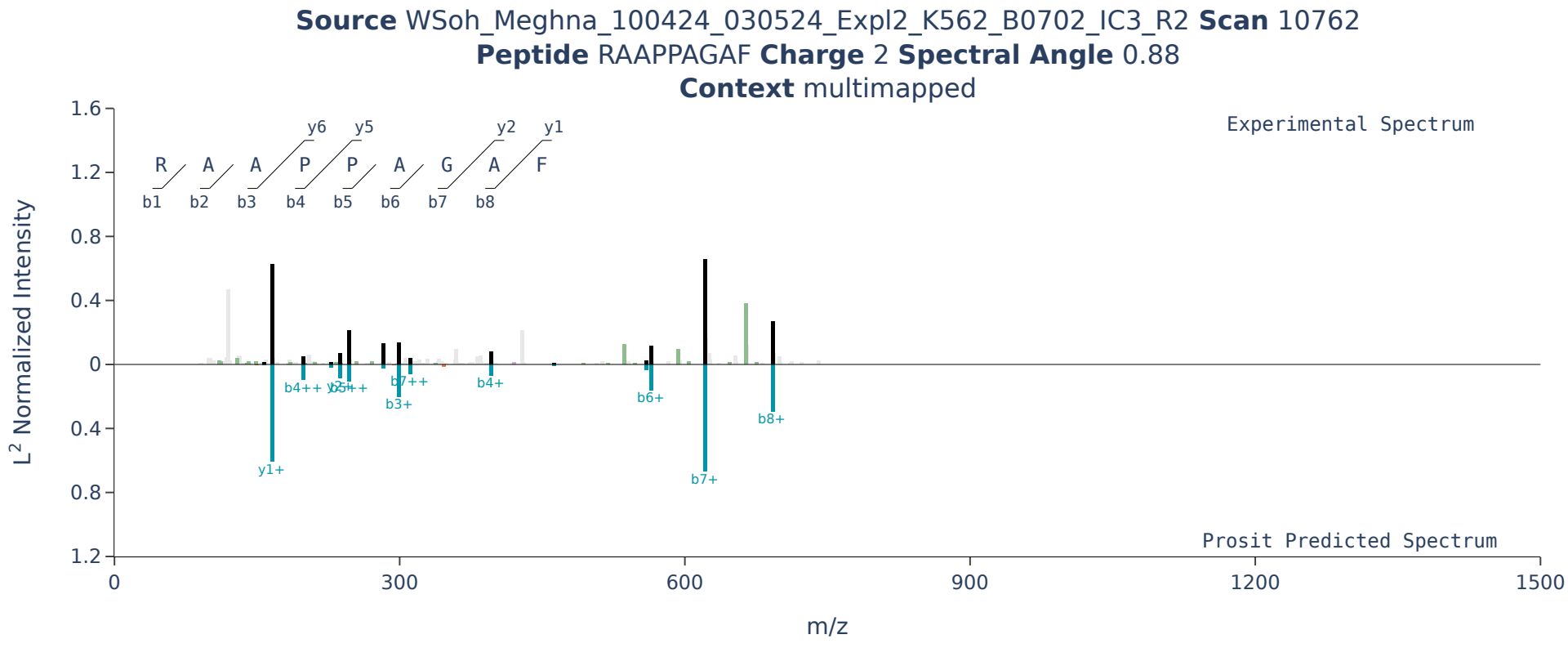
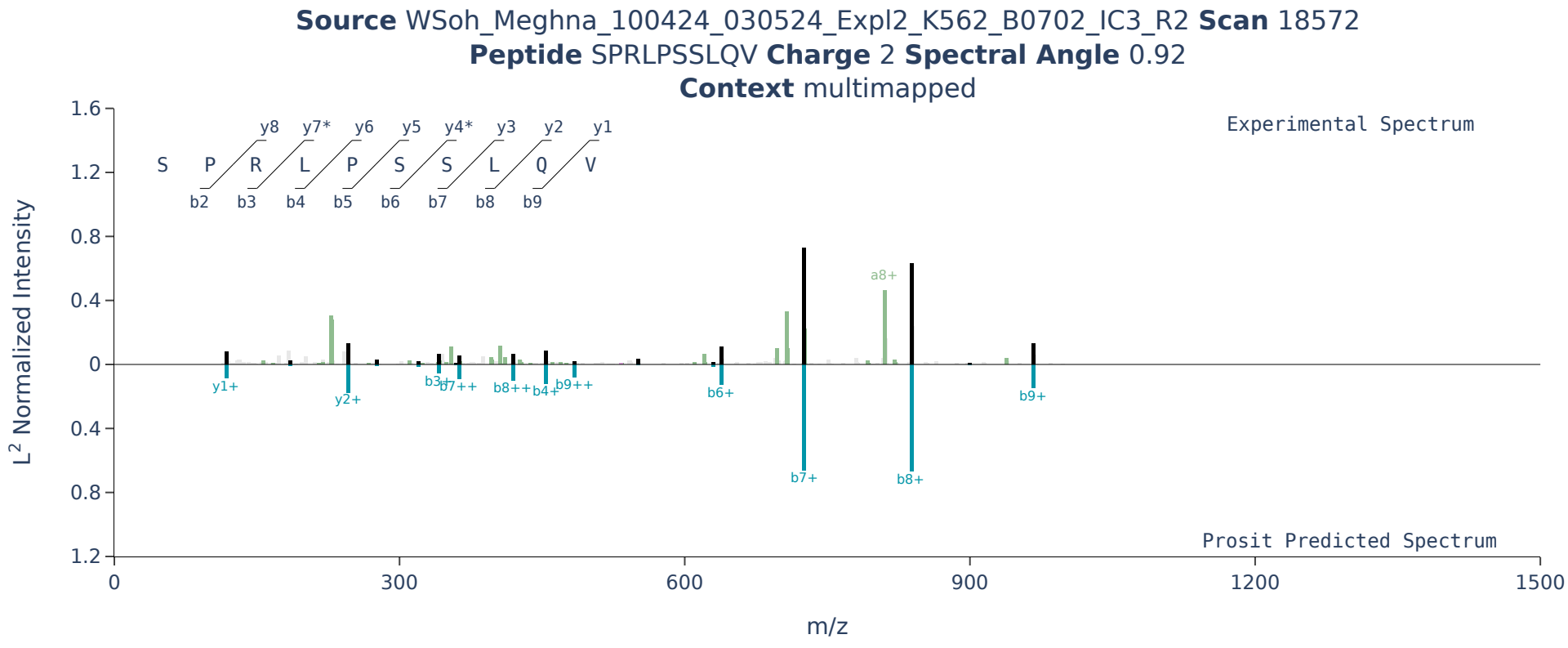
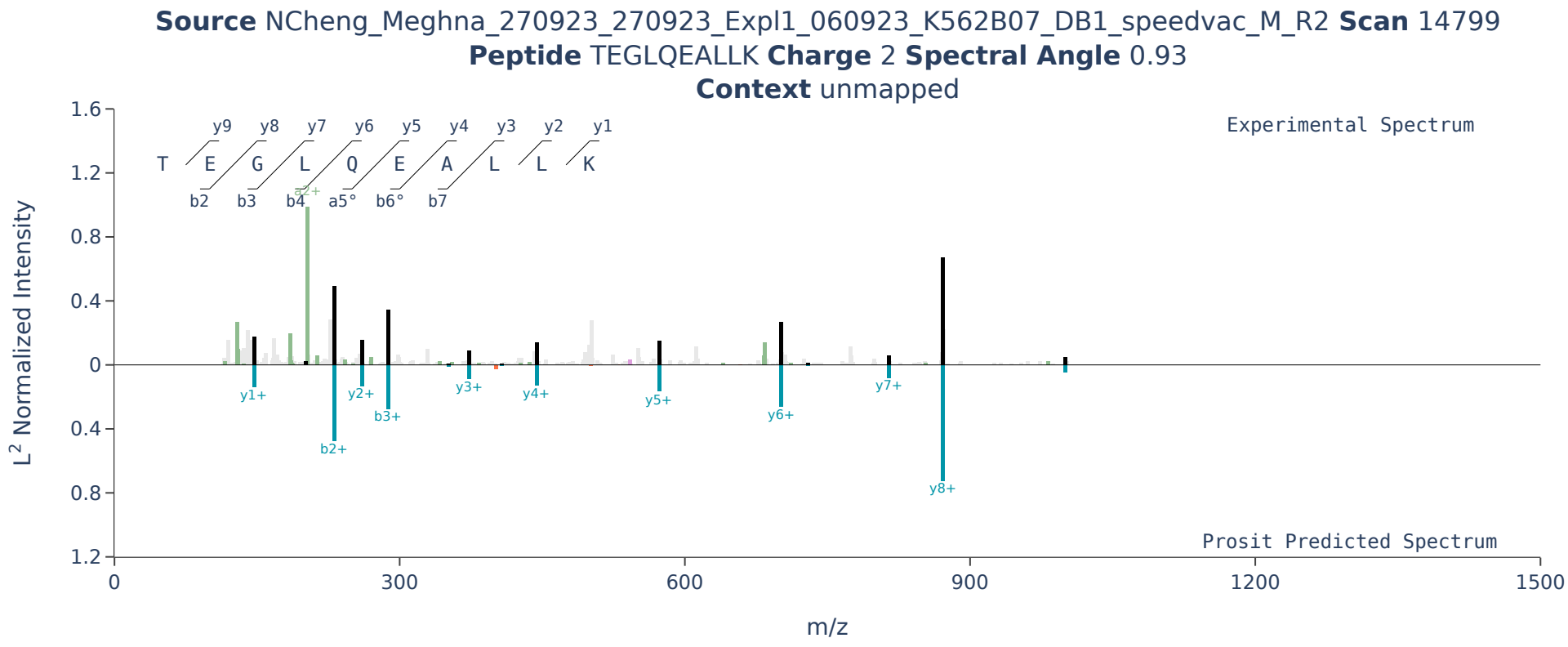
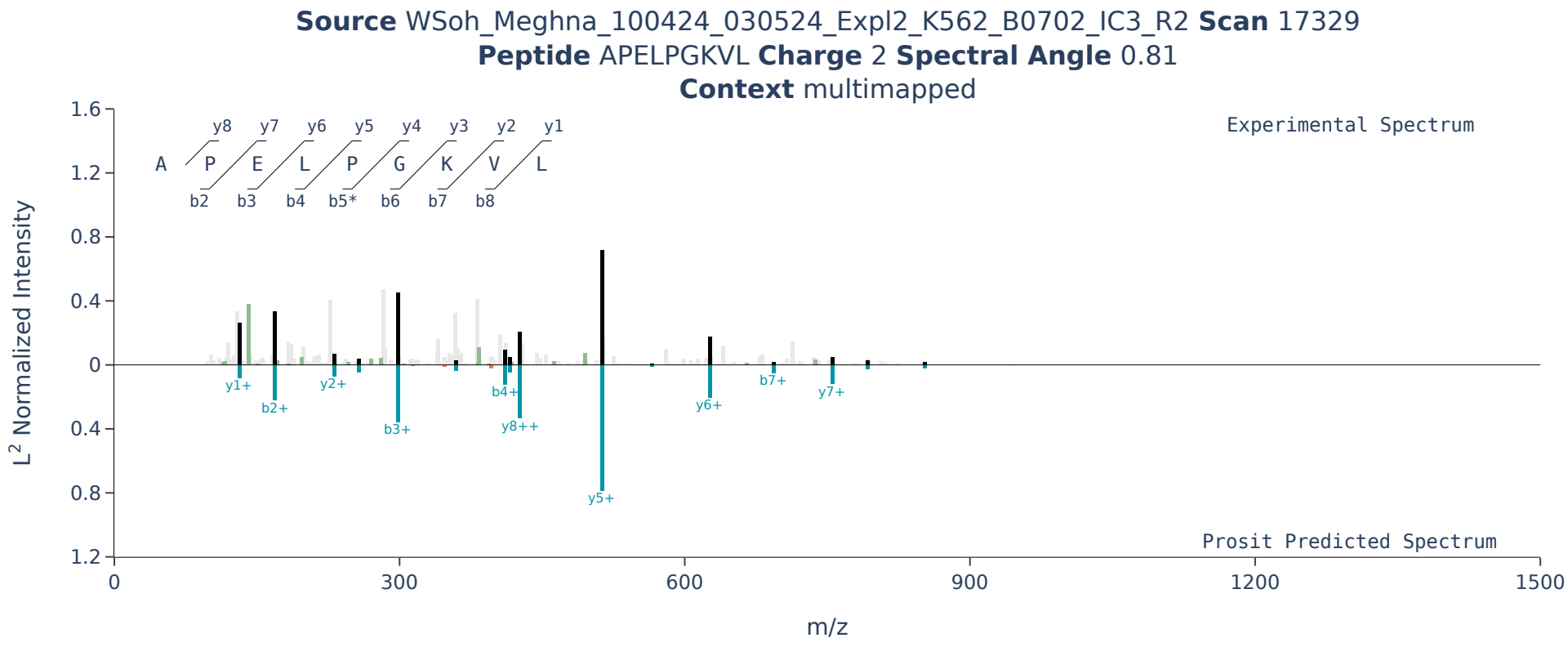
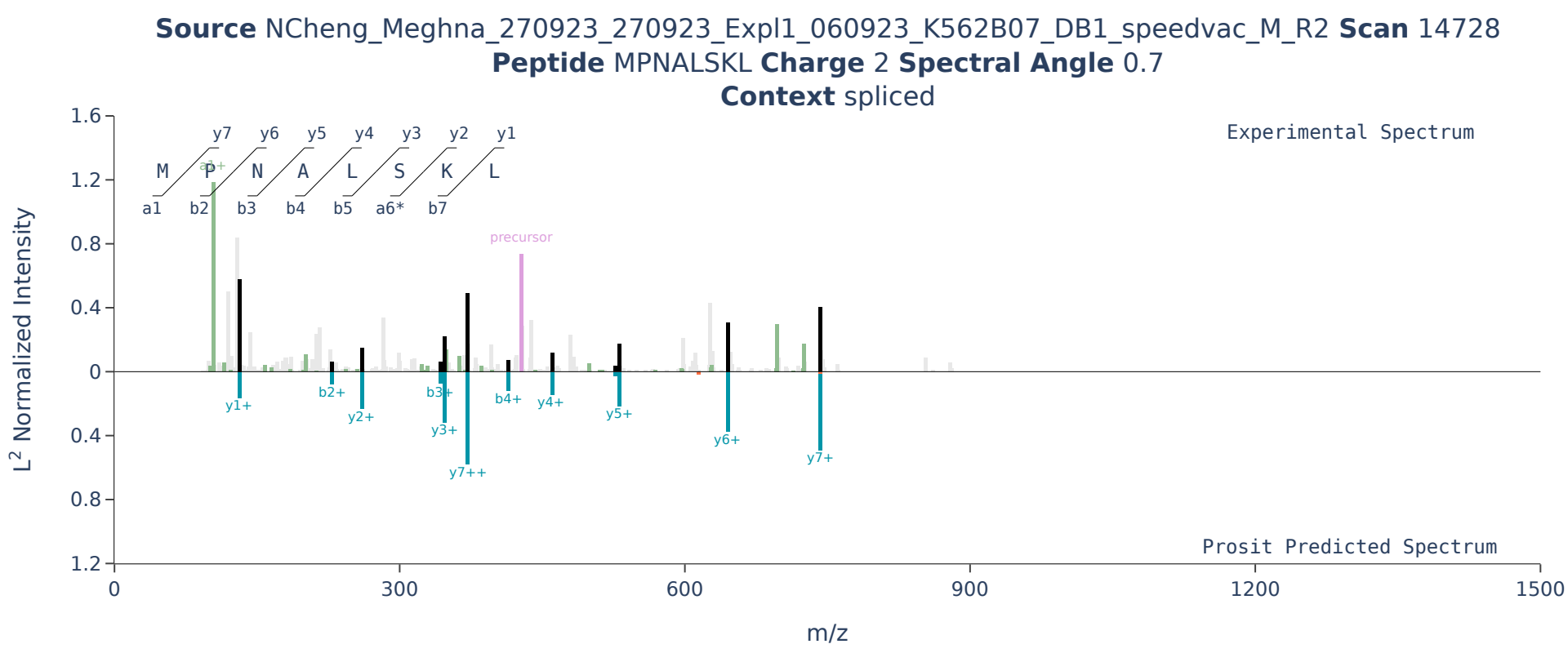
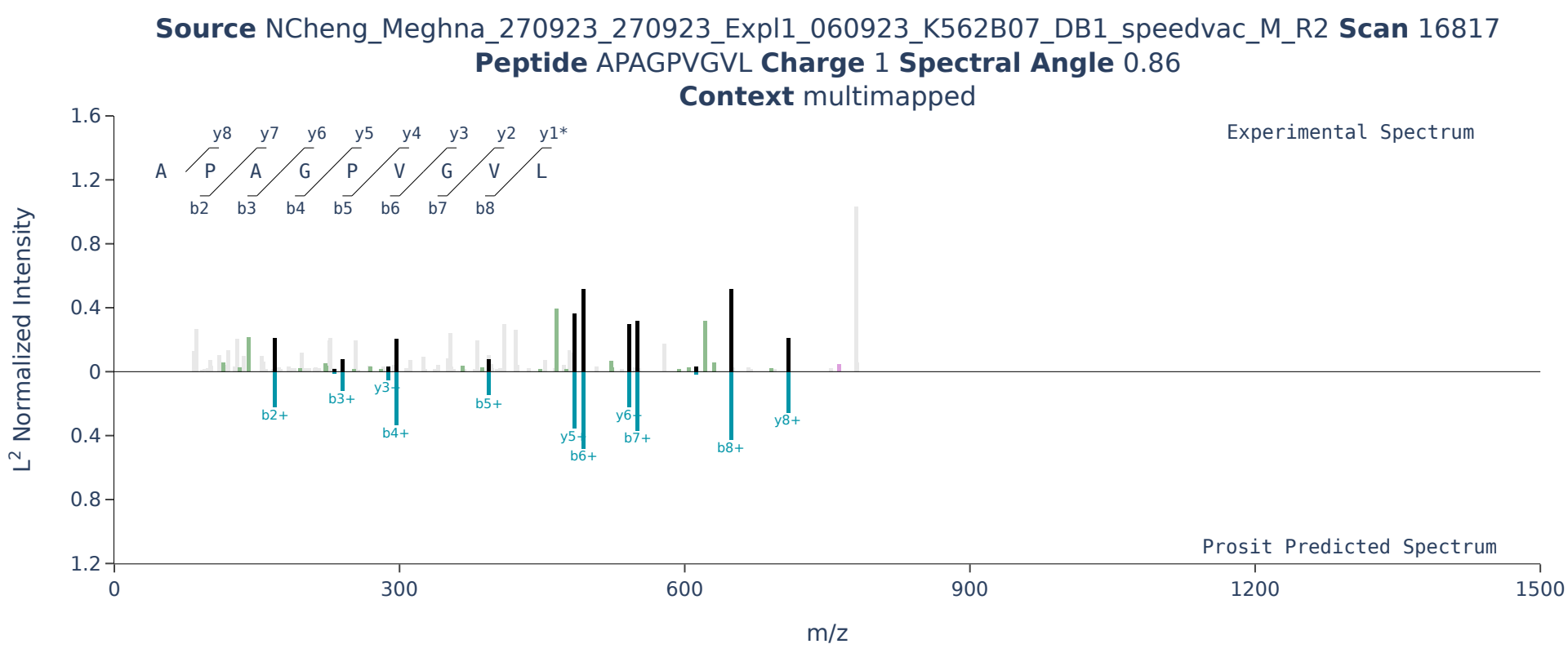


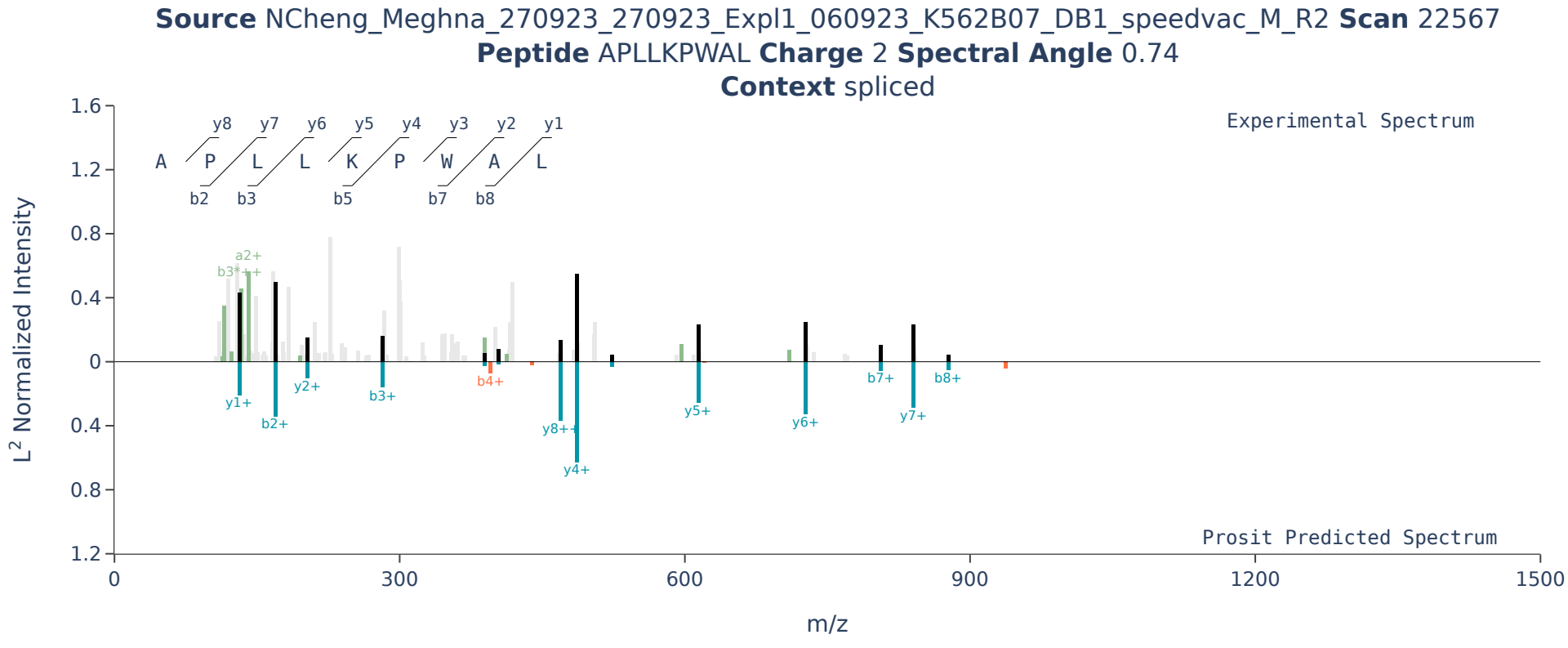
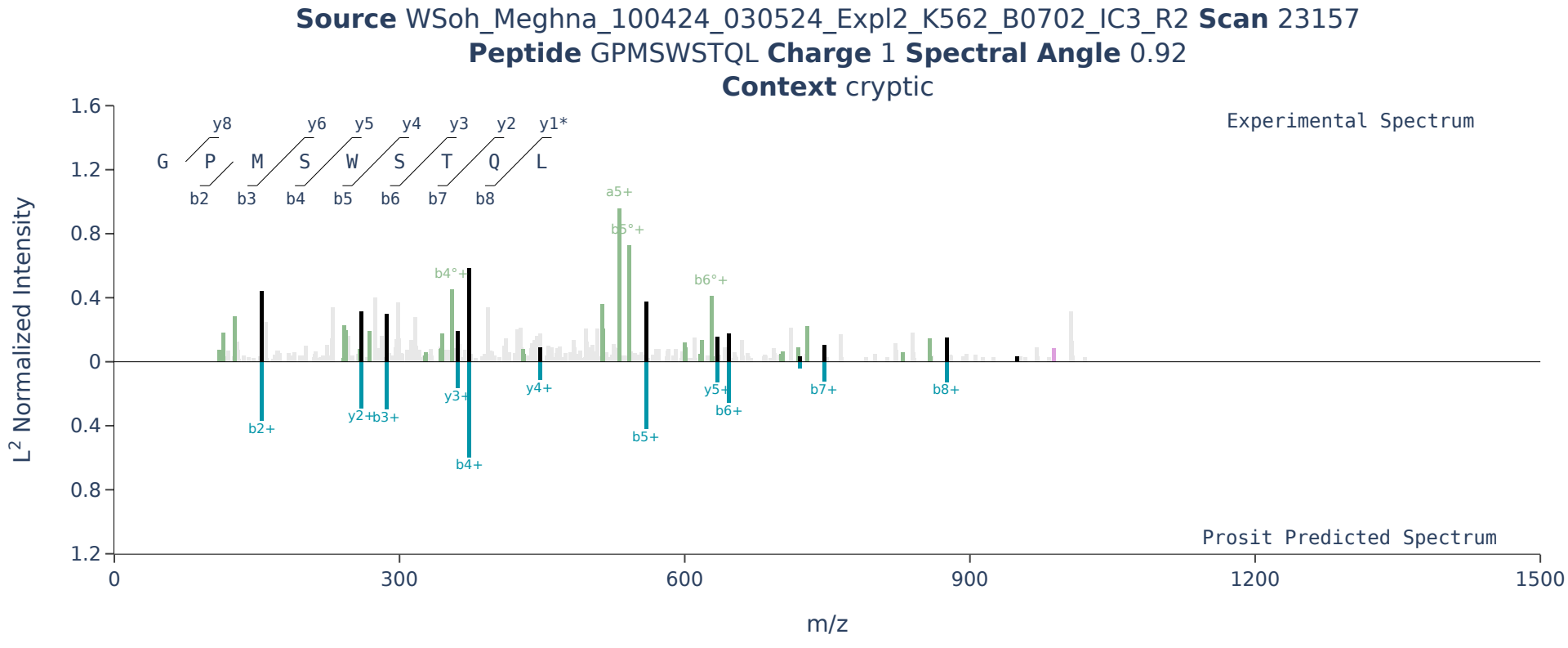
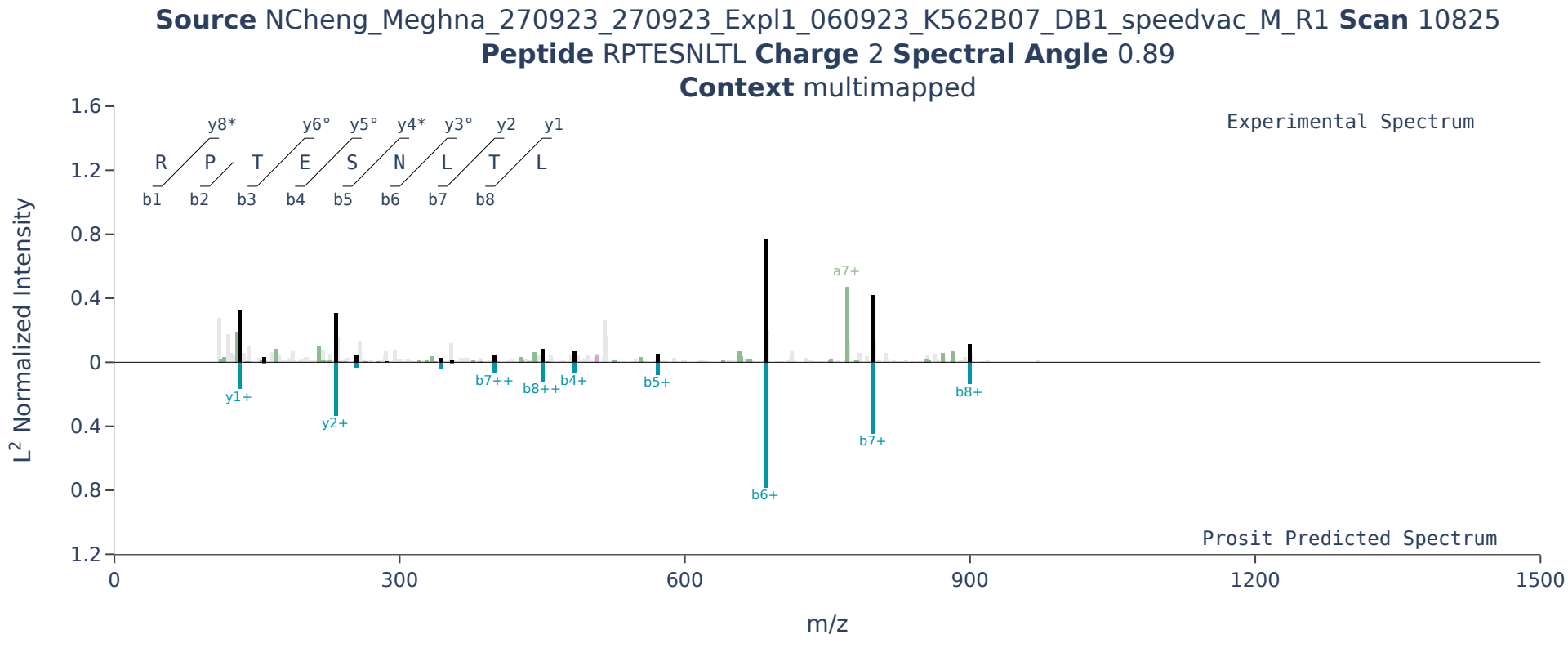
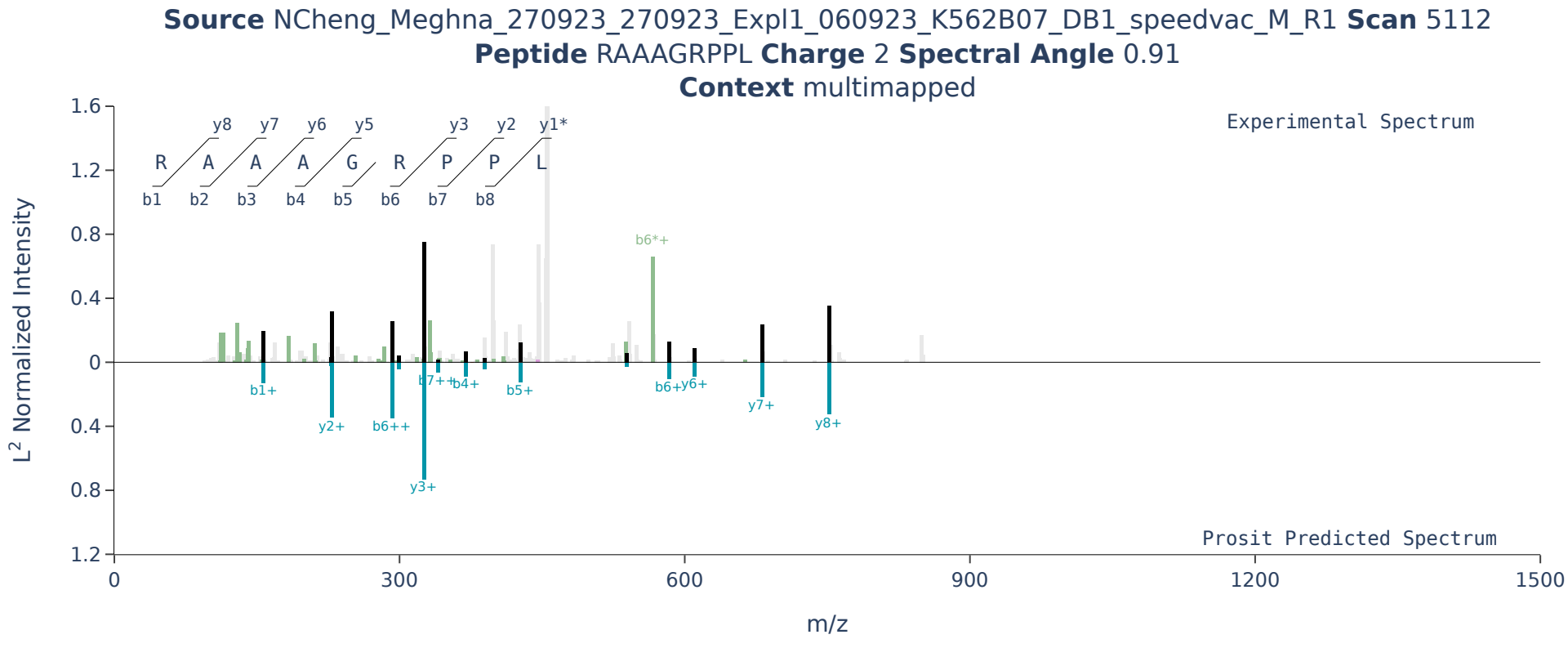
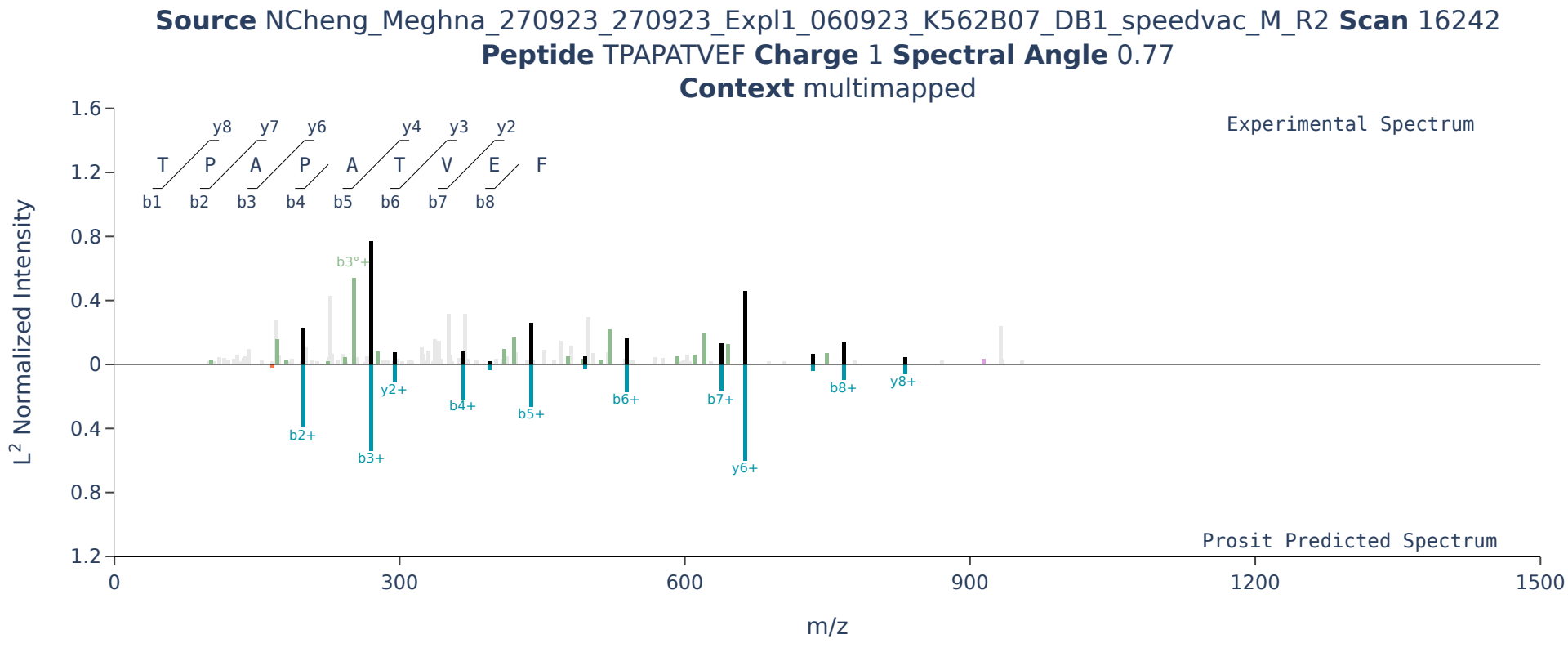
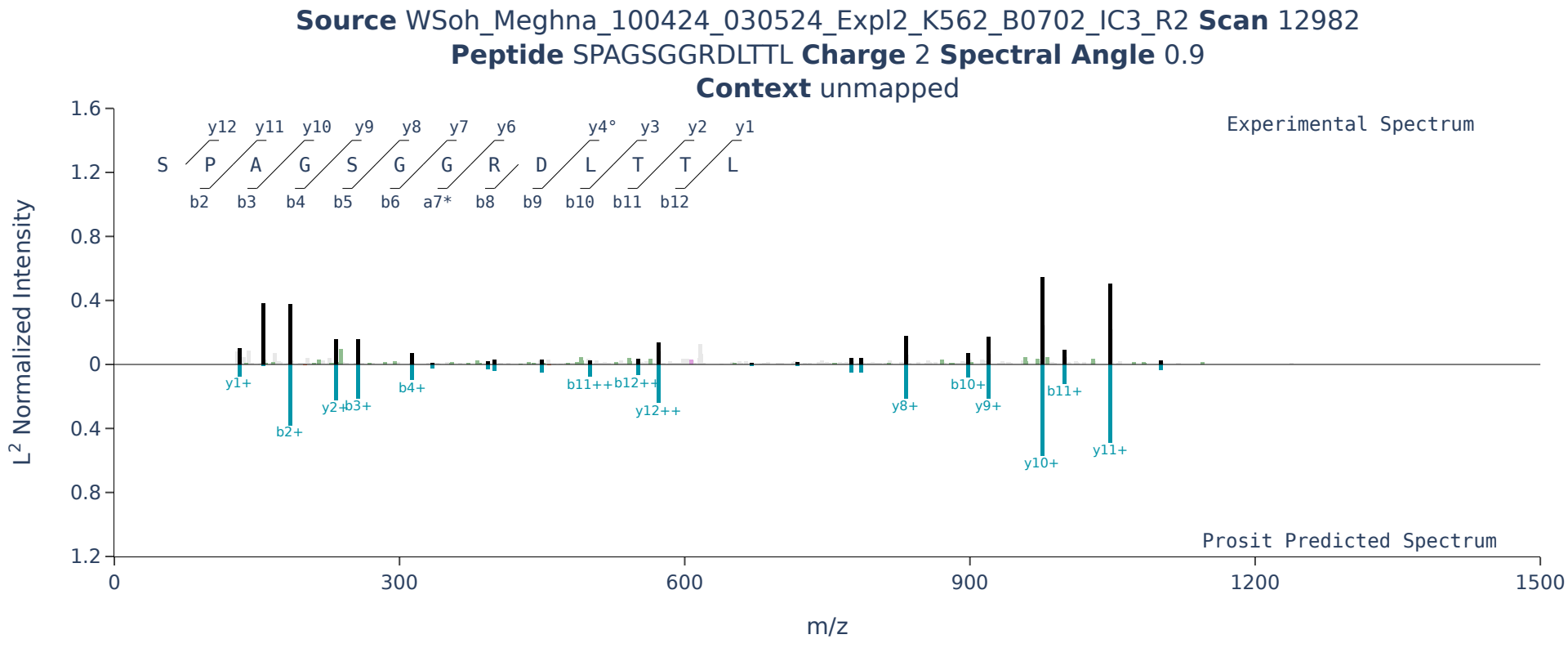
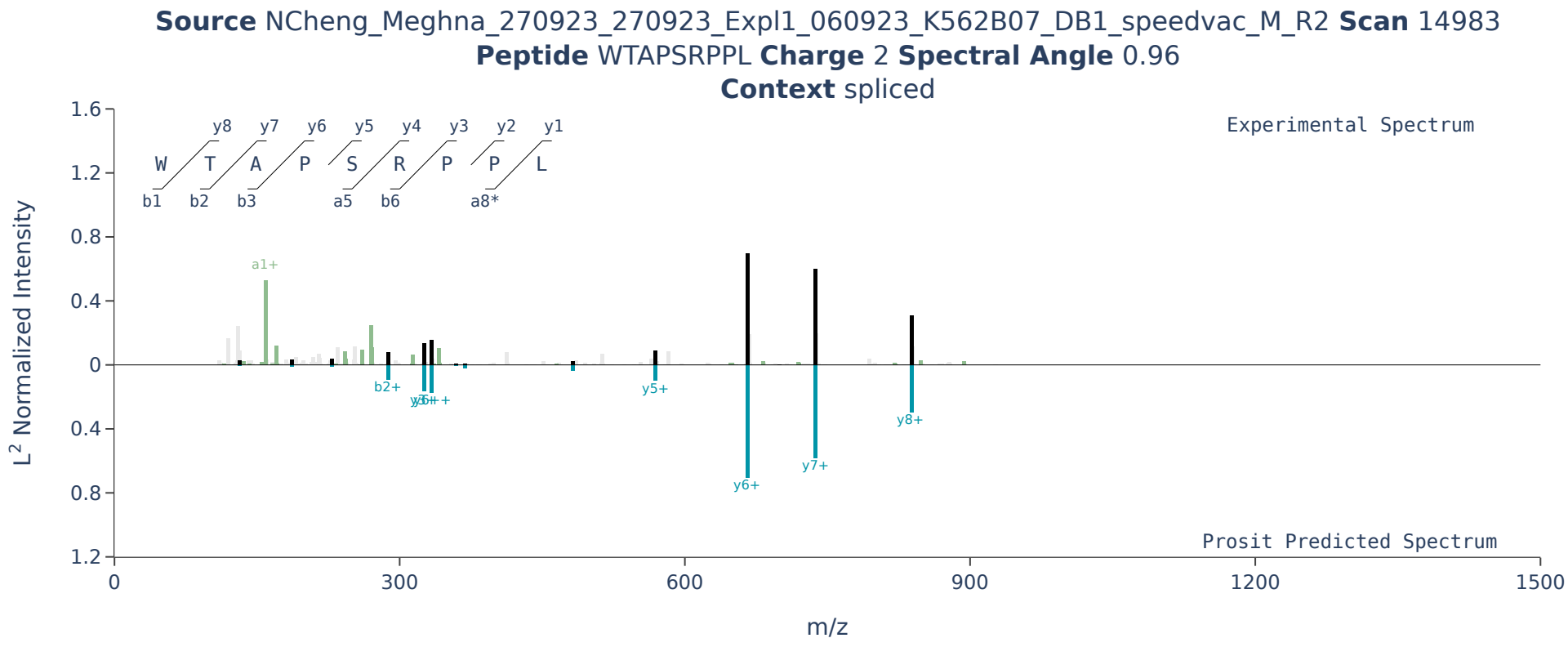
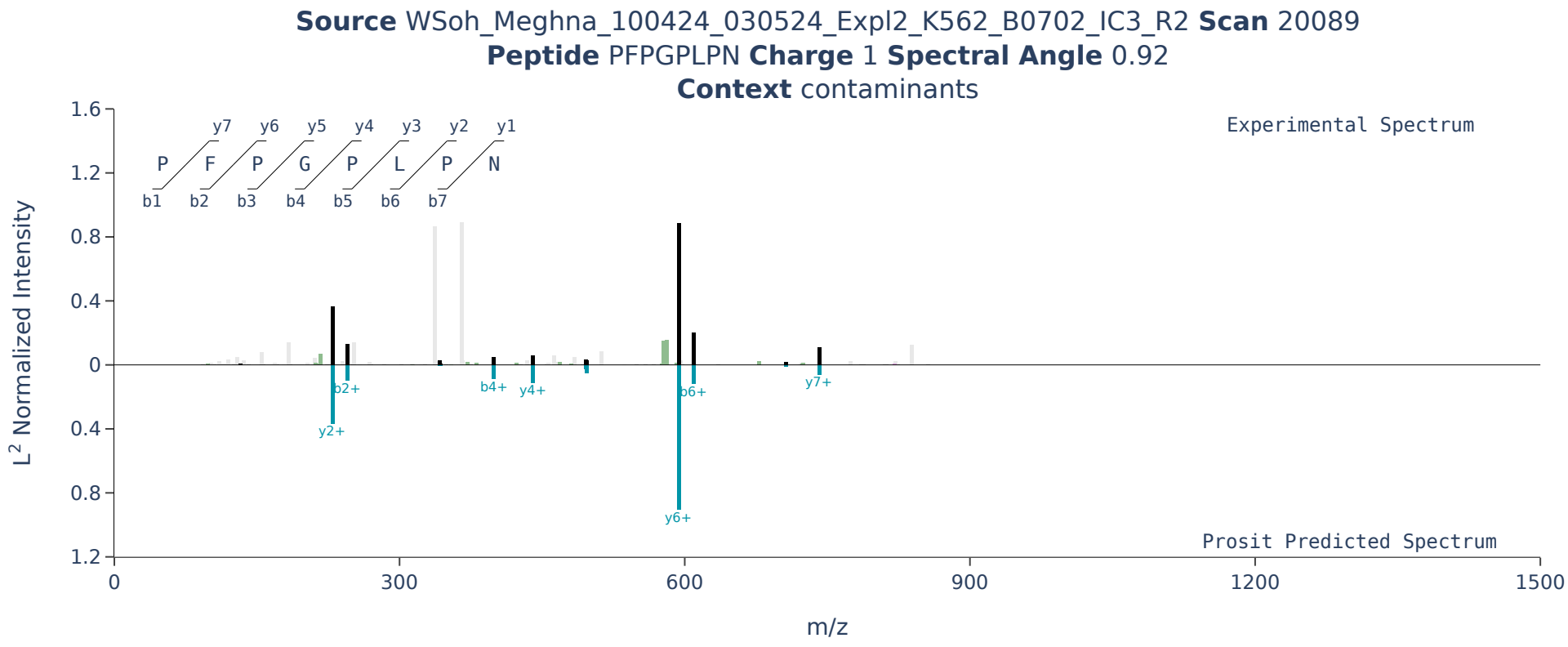
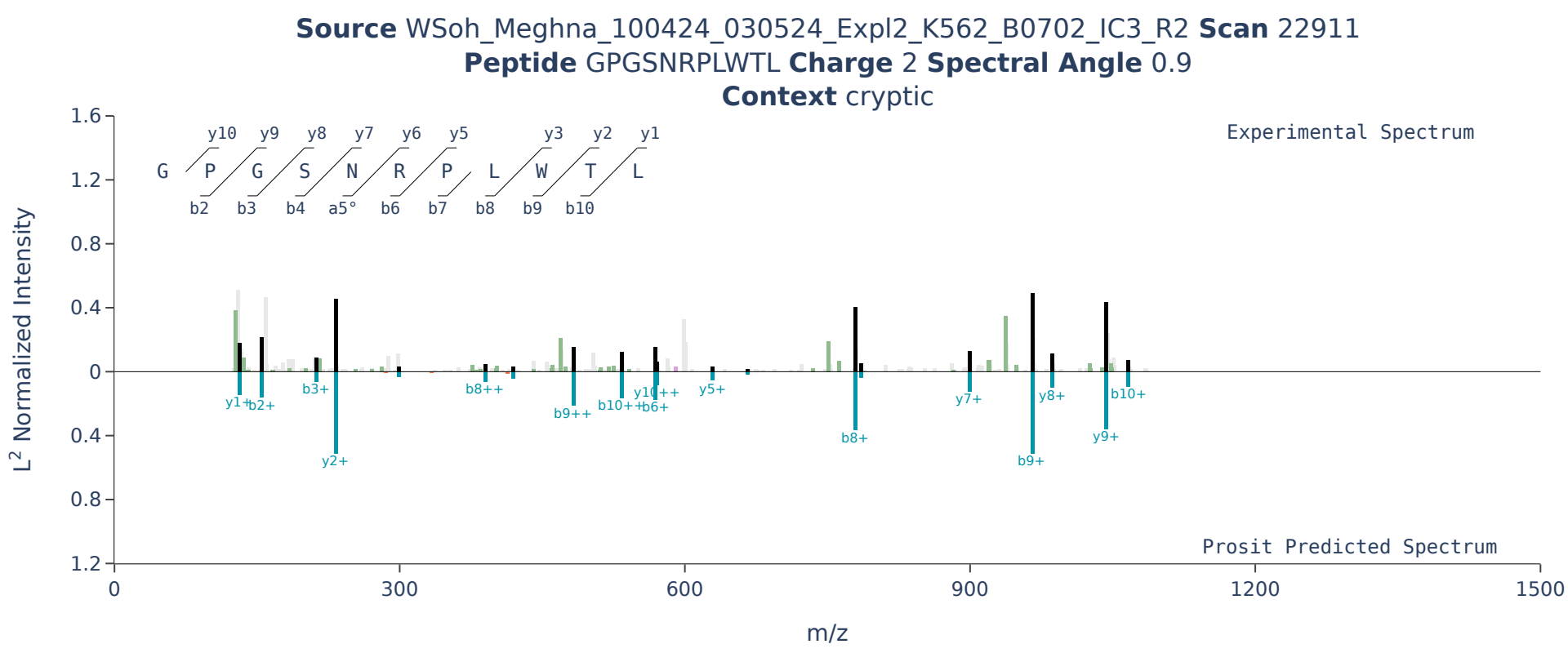
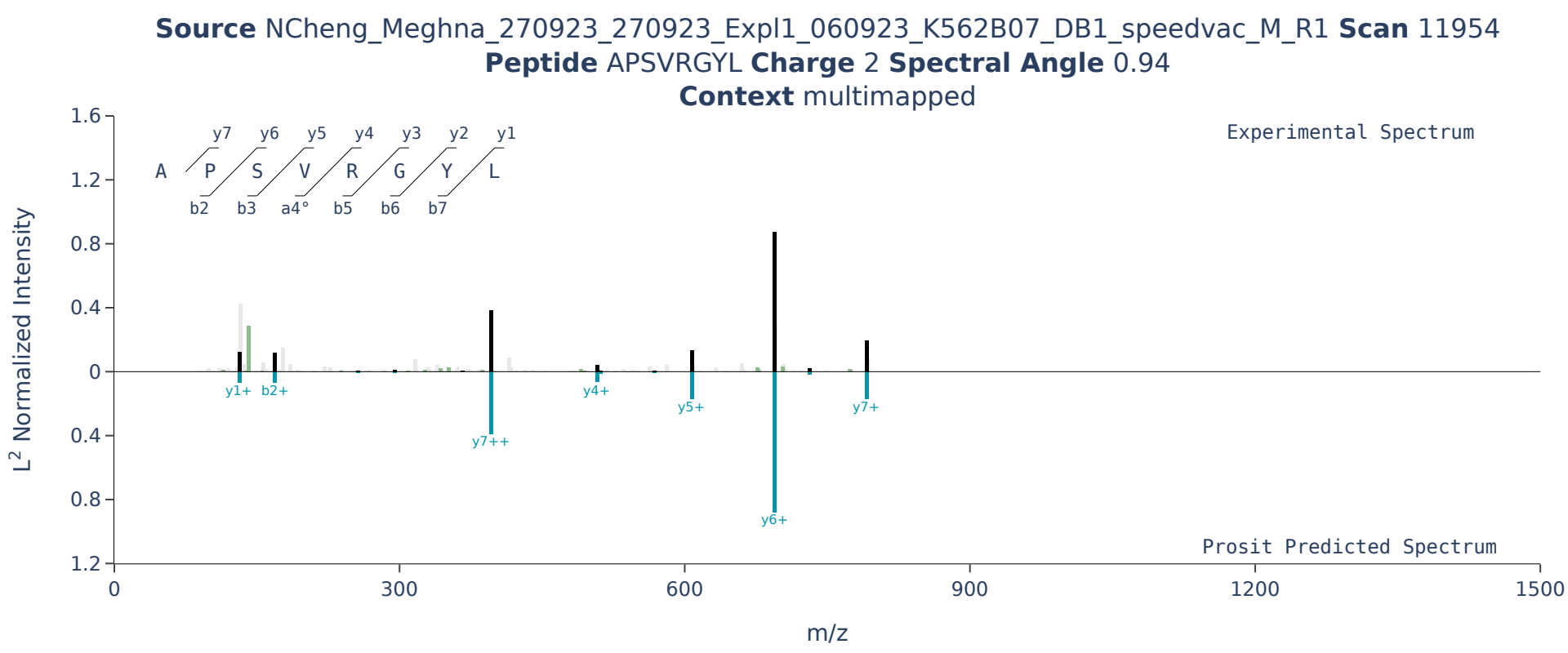
Peptide SPSPASGTL Charge 1 Spectral Angle 0.94
Context multimapped

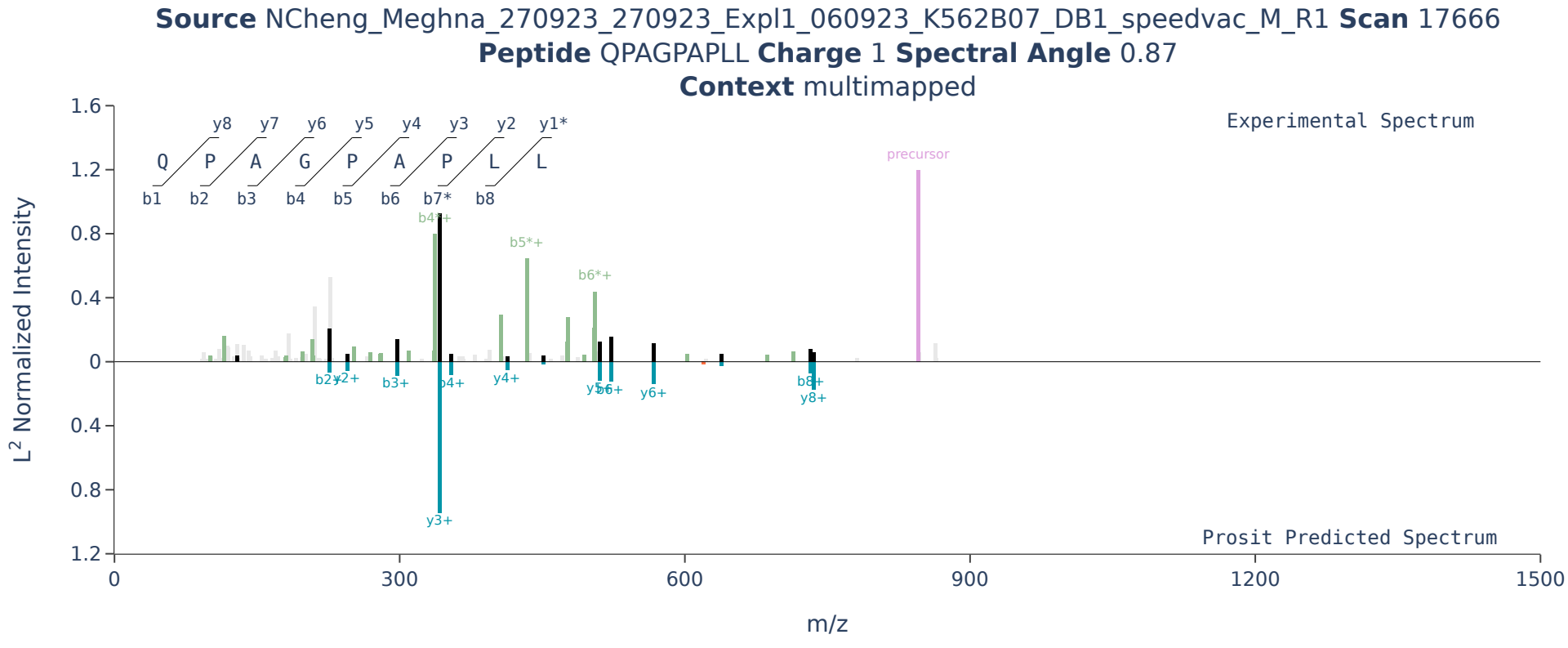
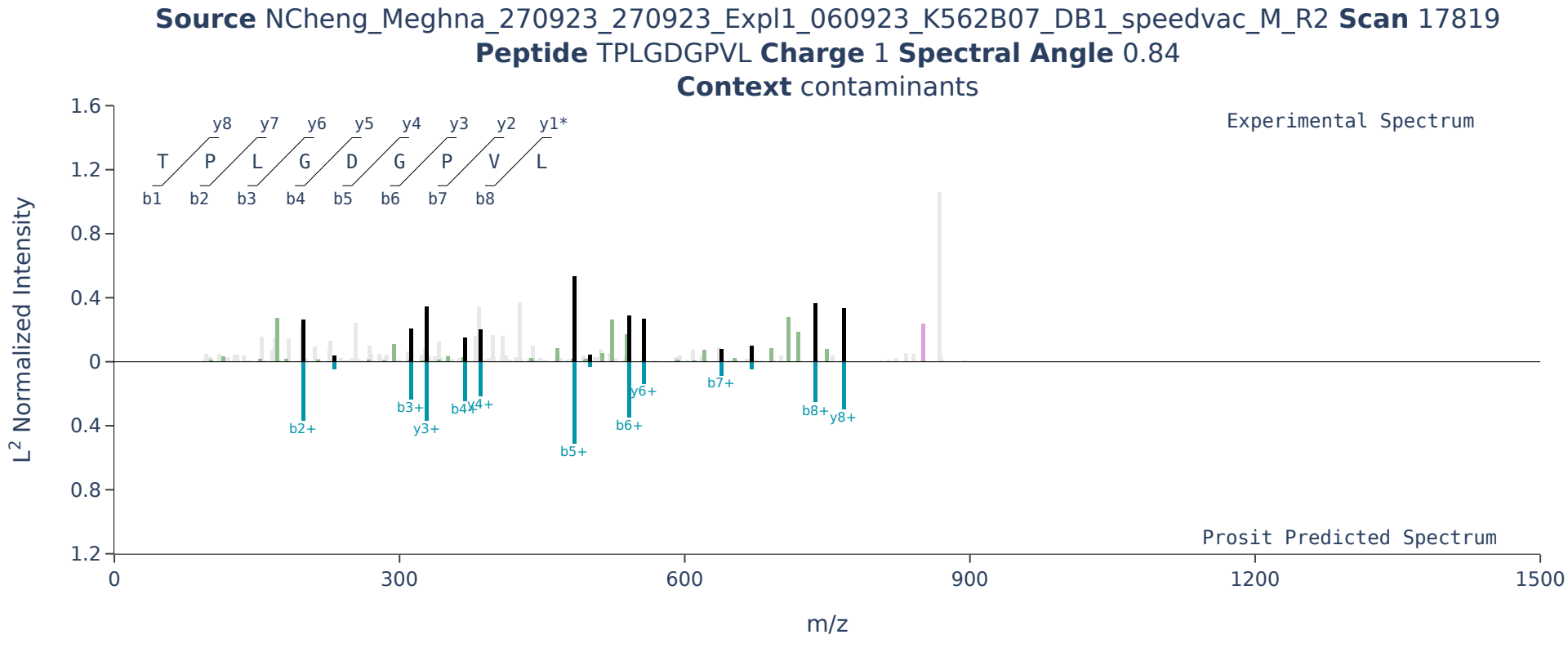
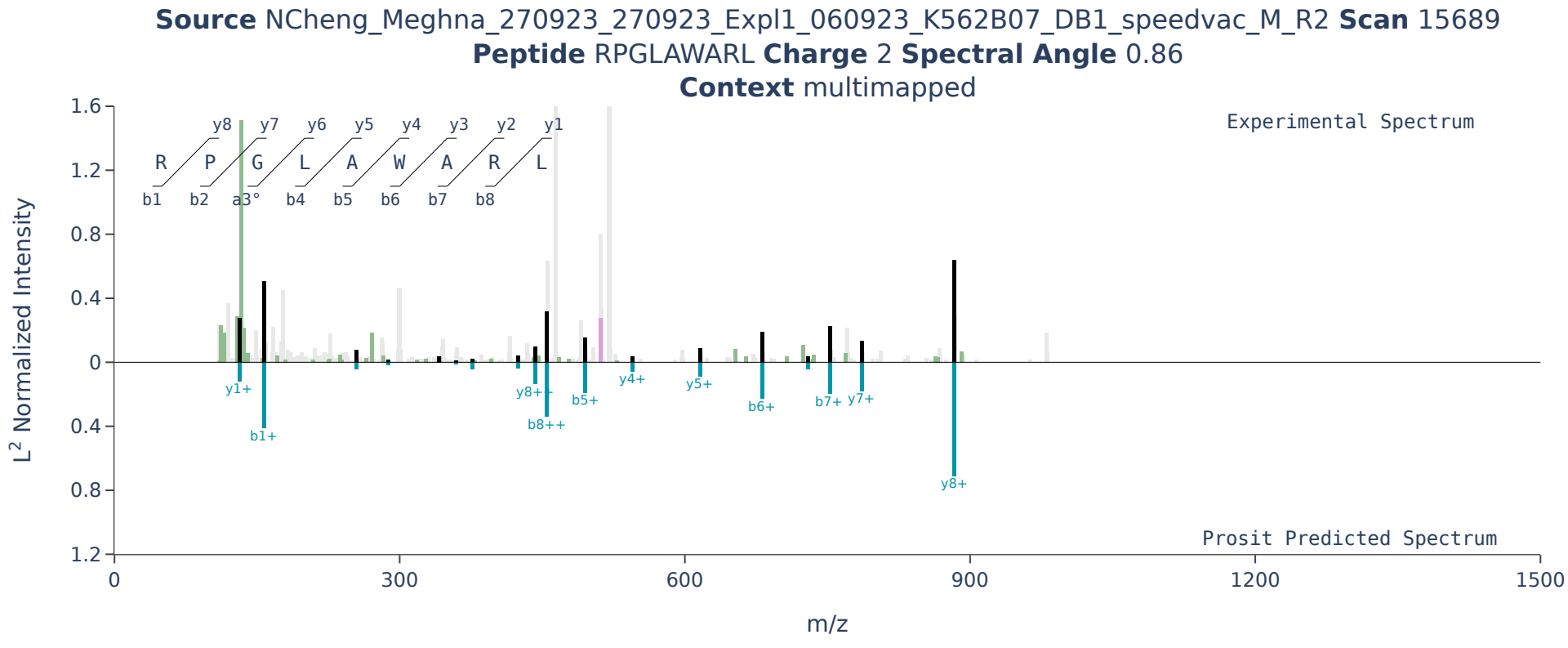
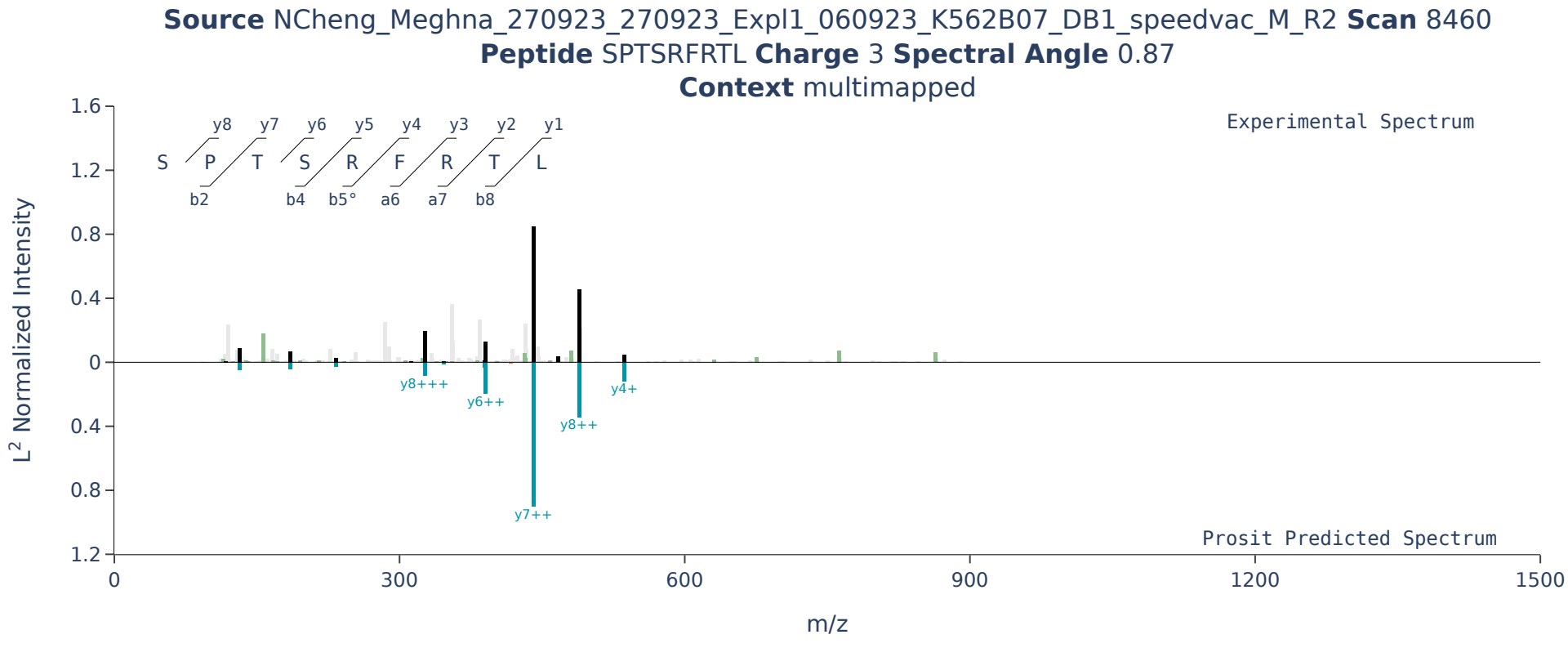
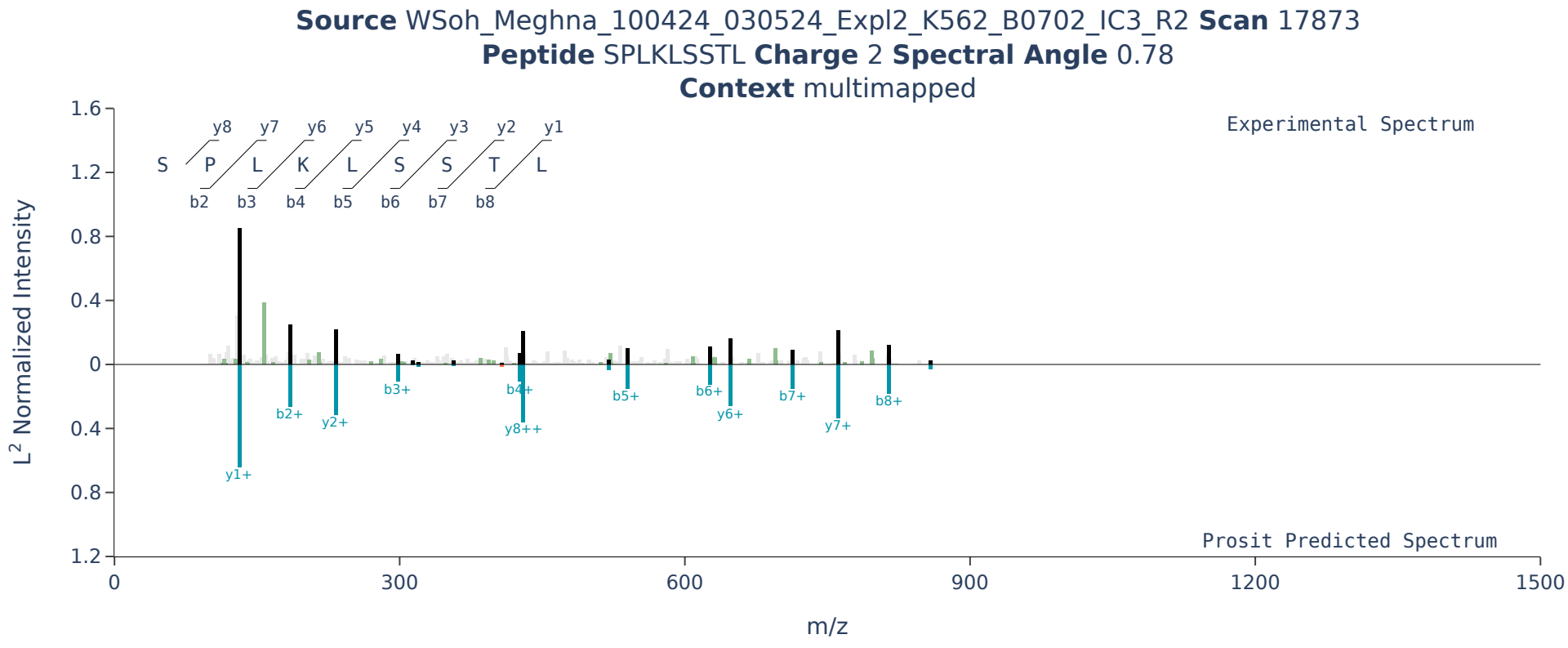
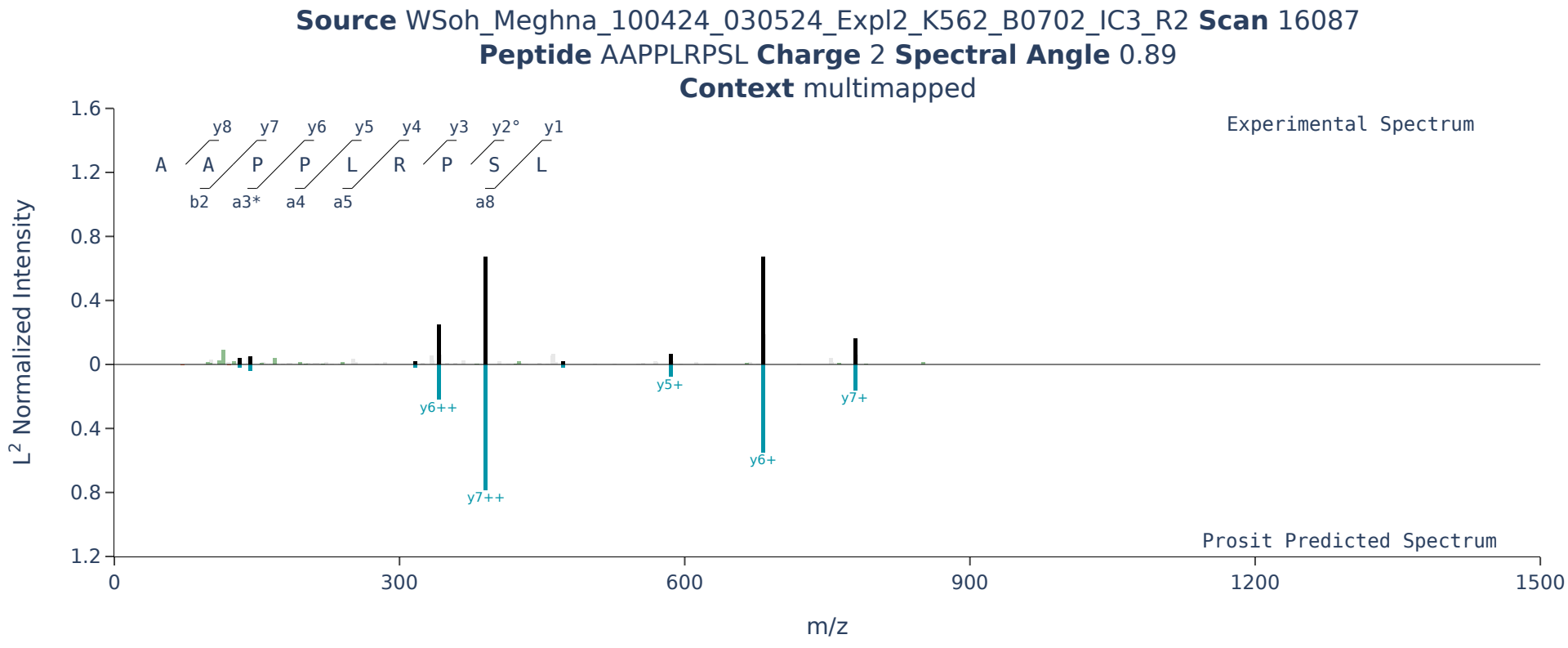
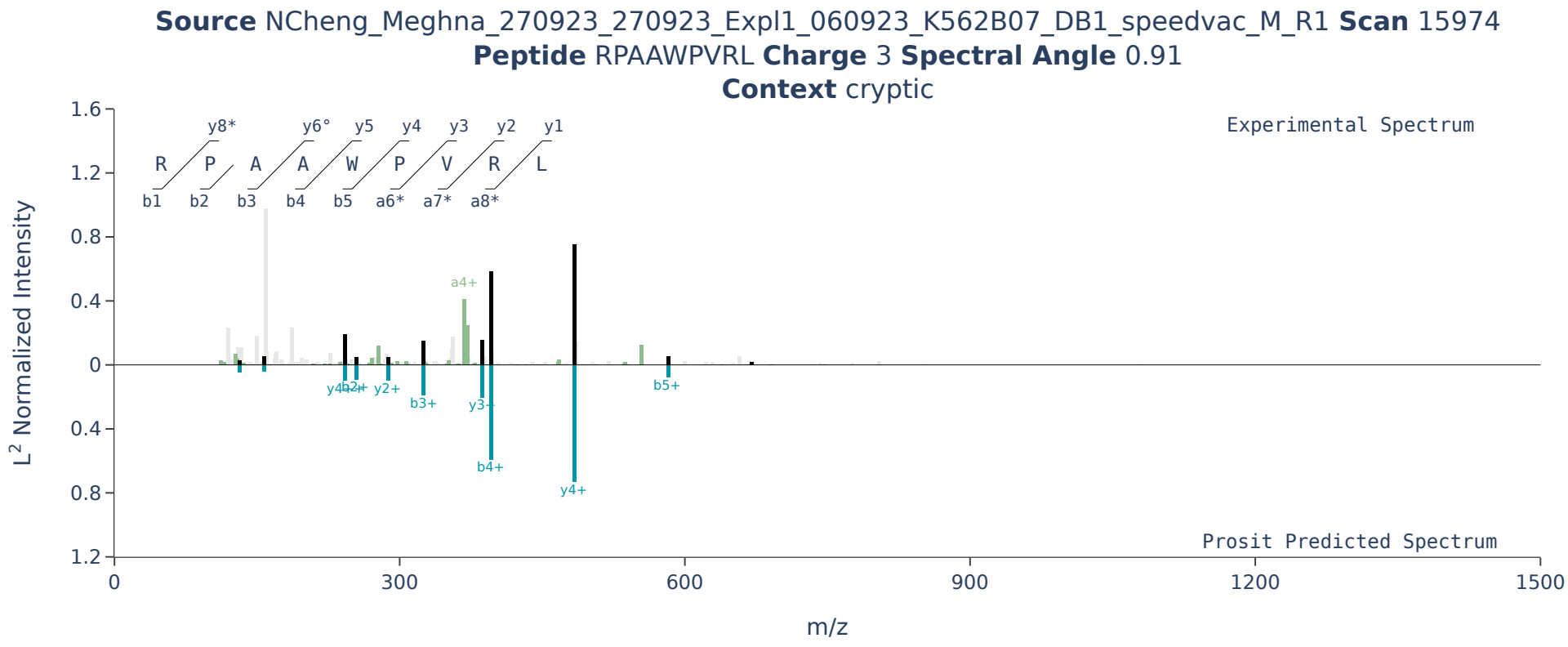
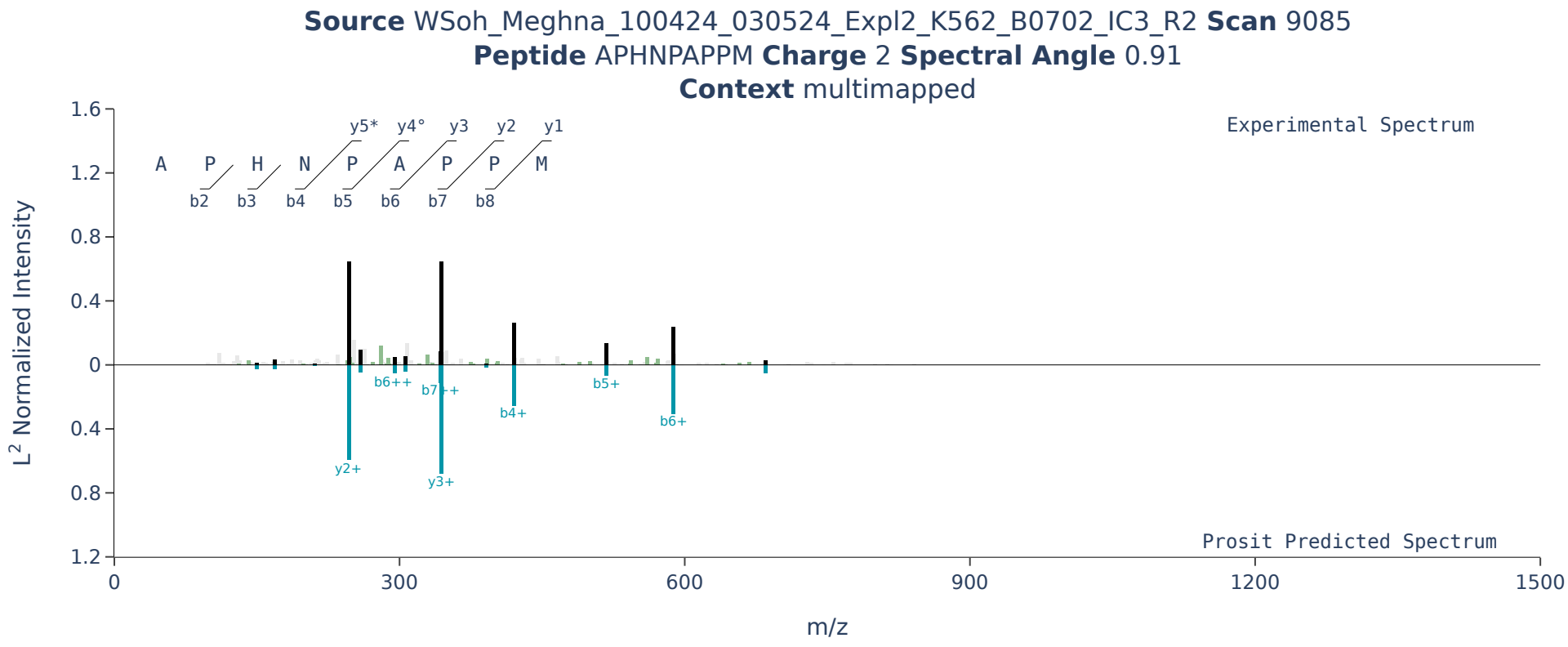
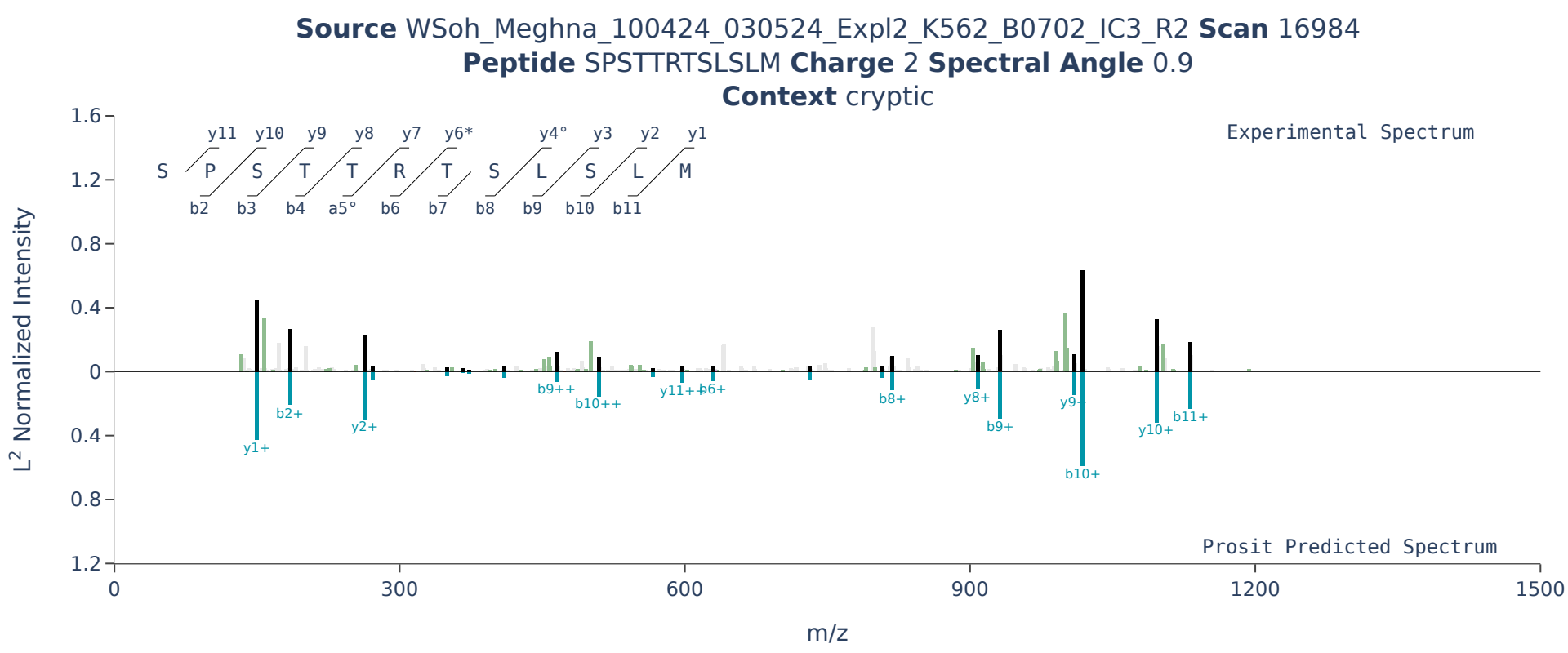
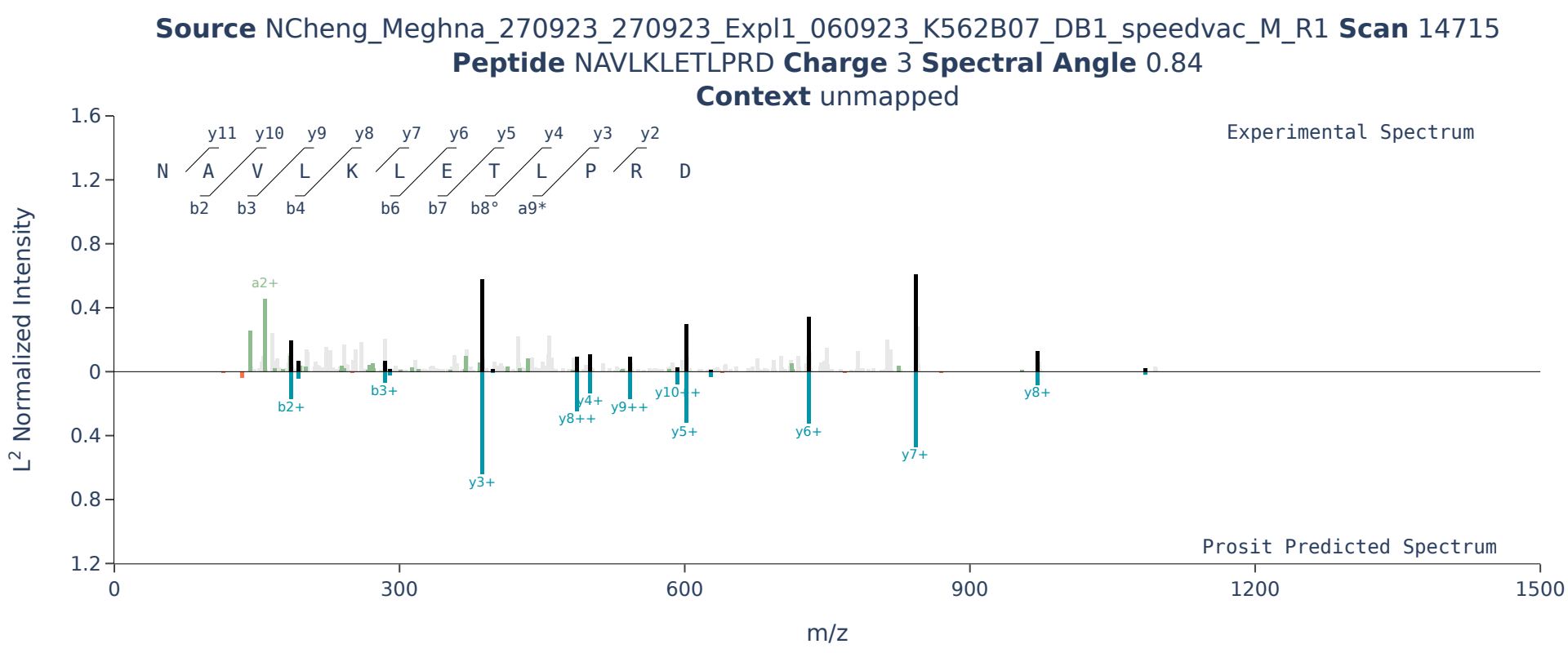


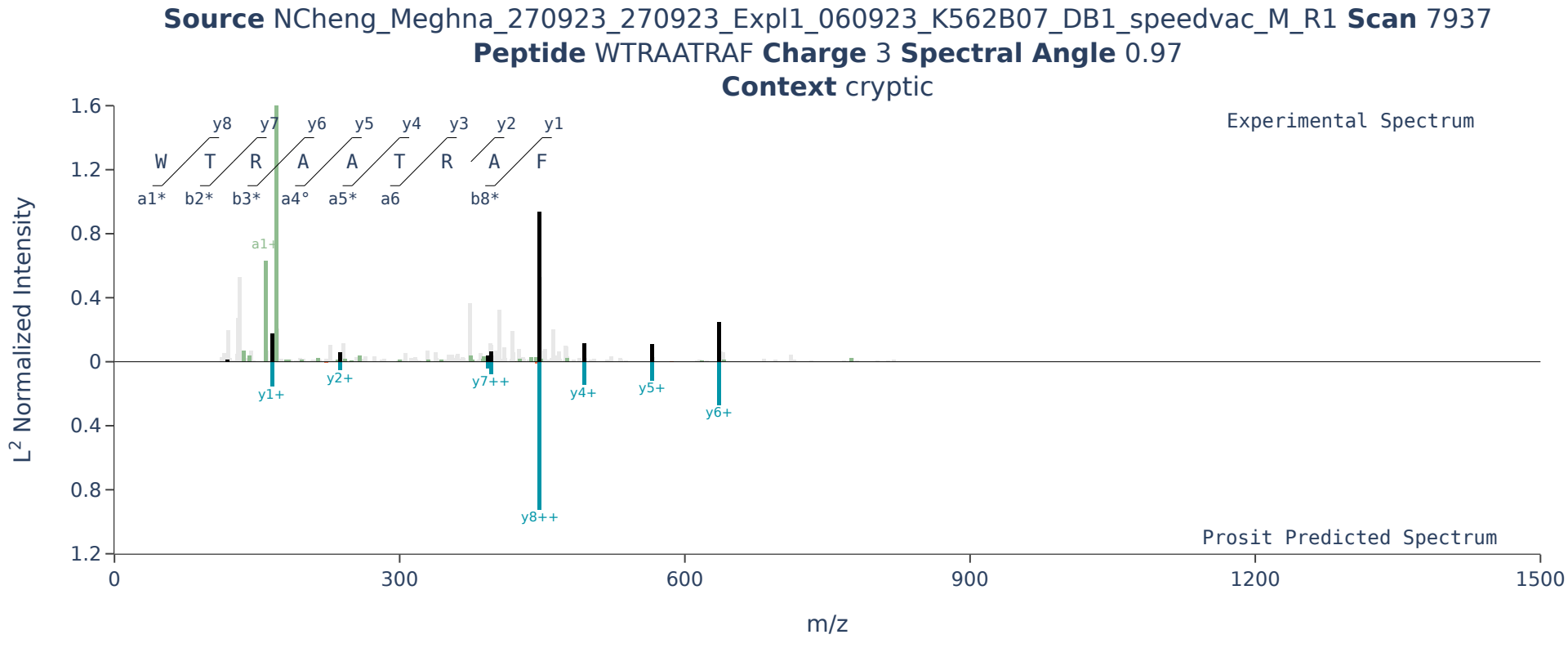
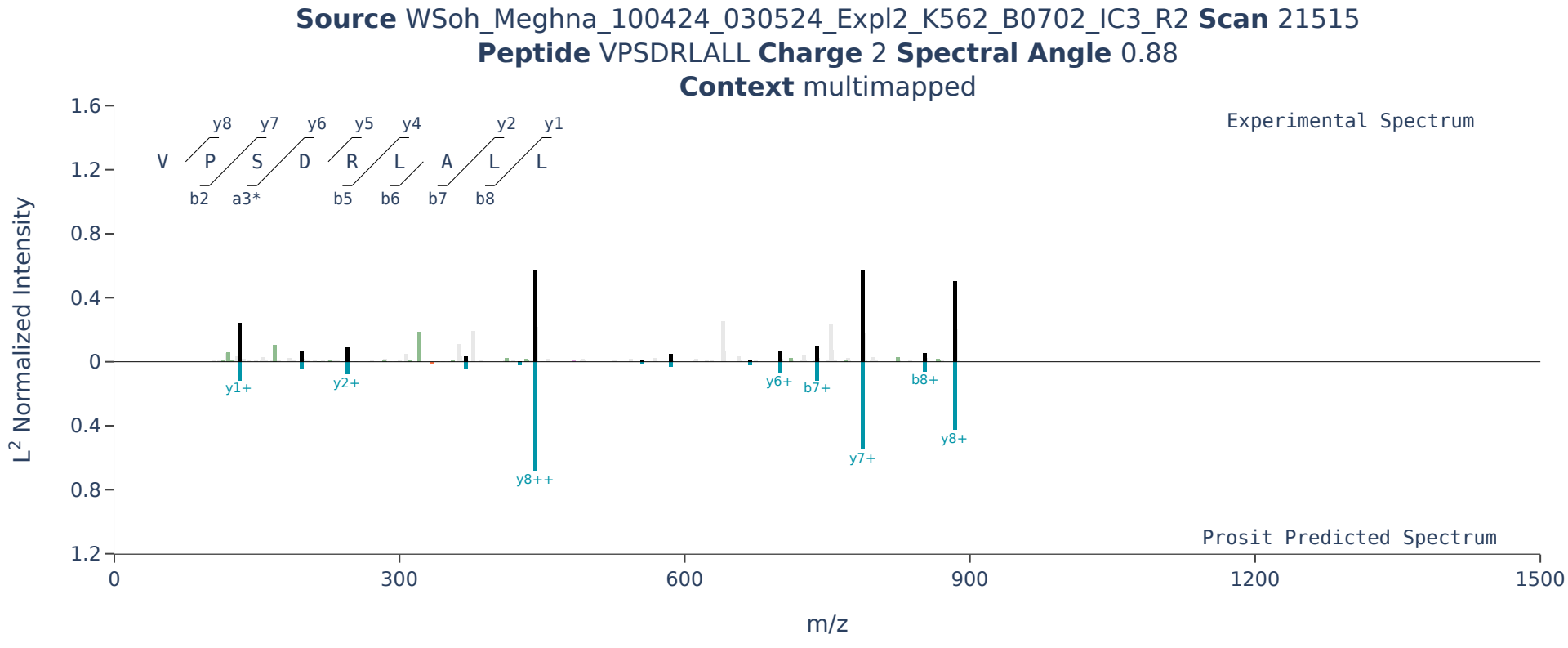
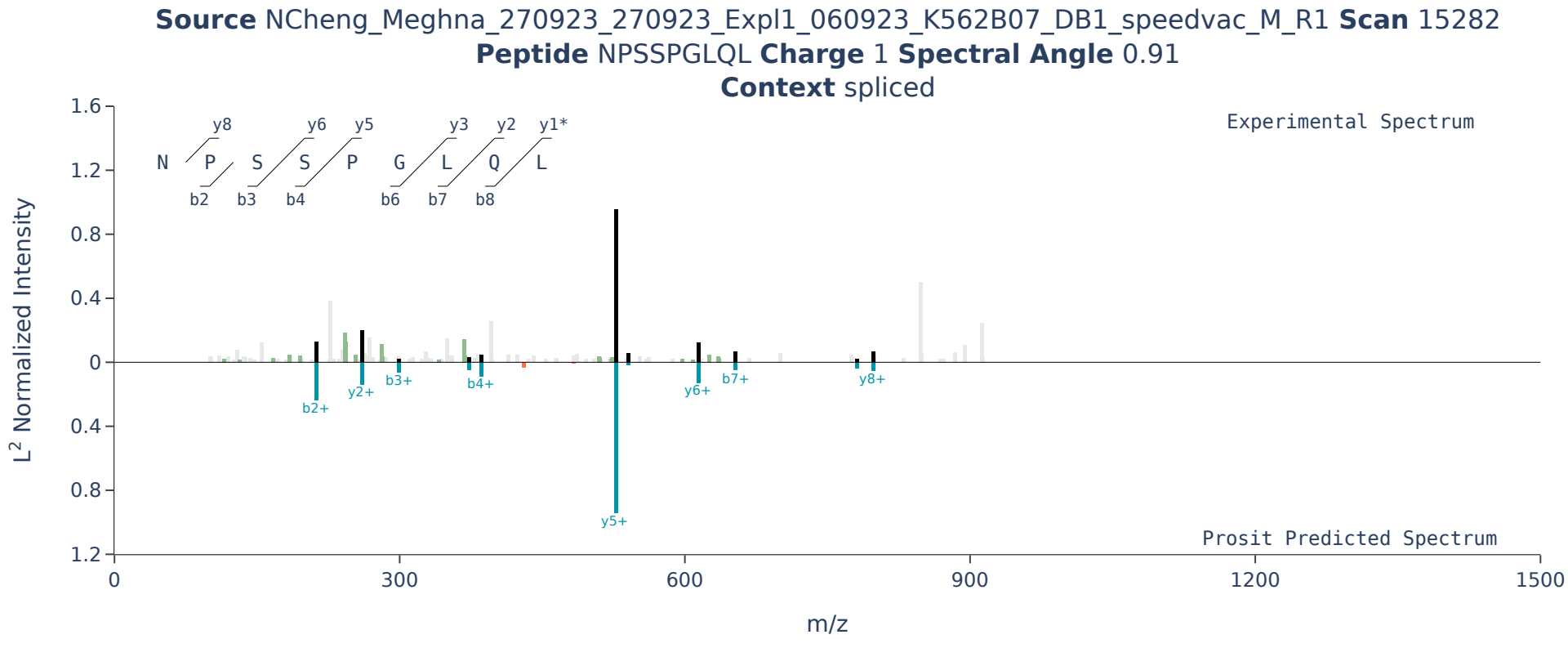
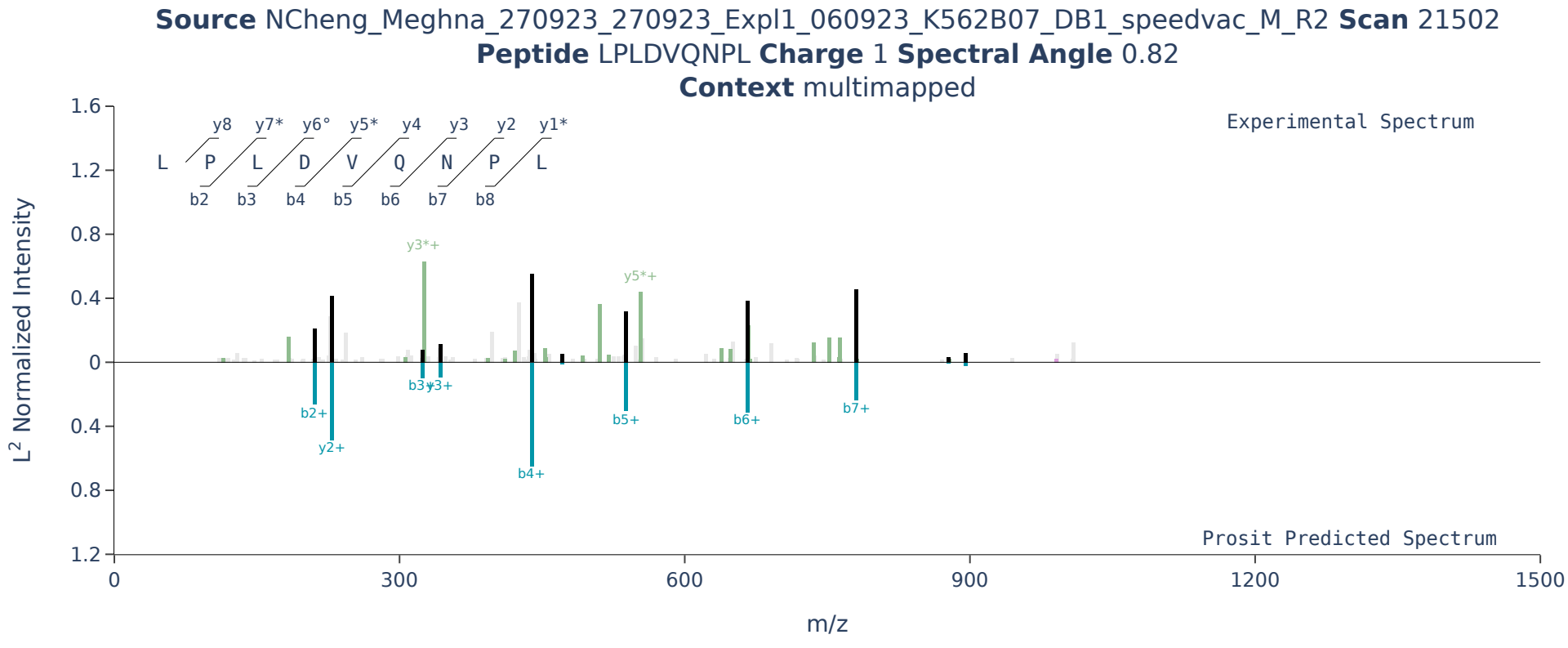
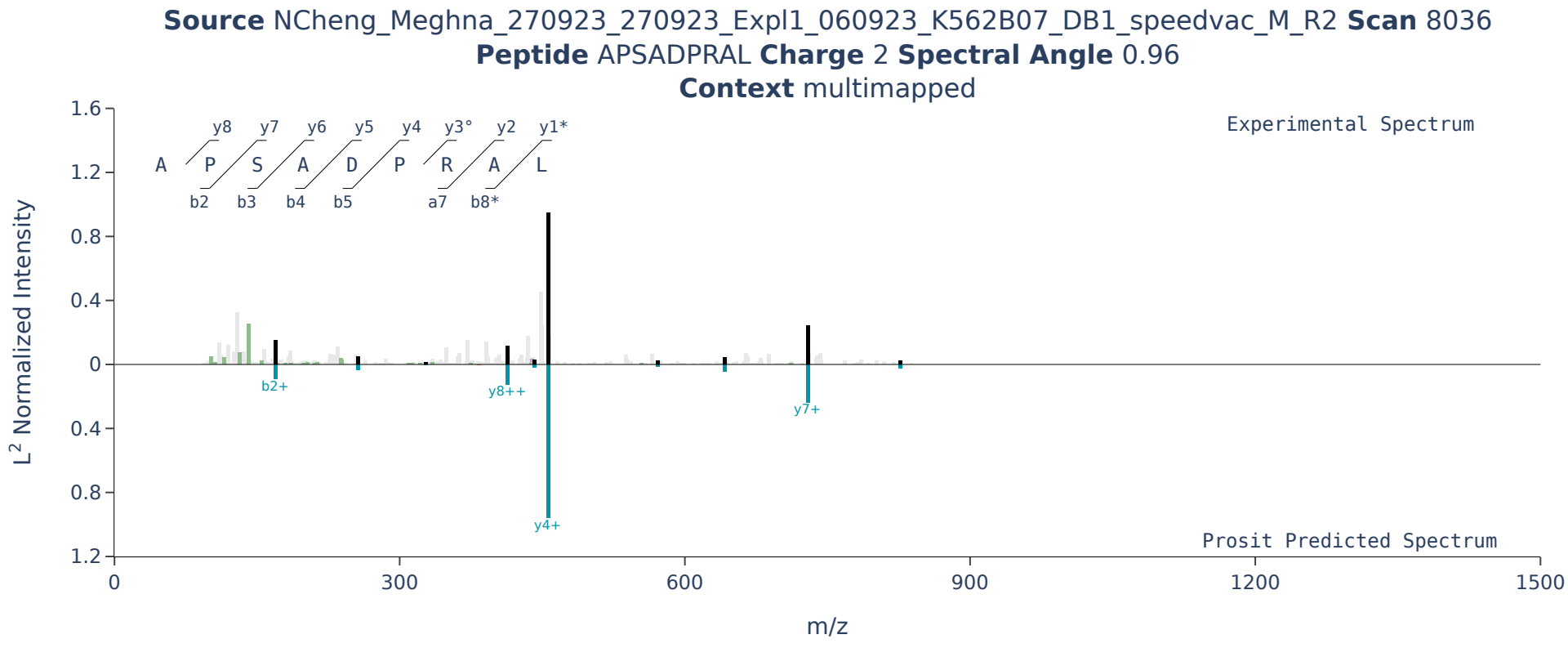
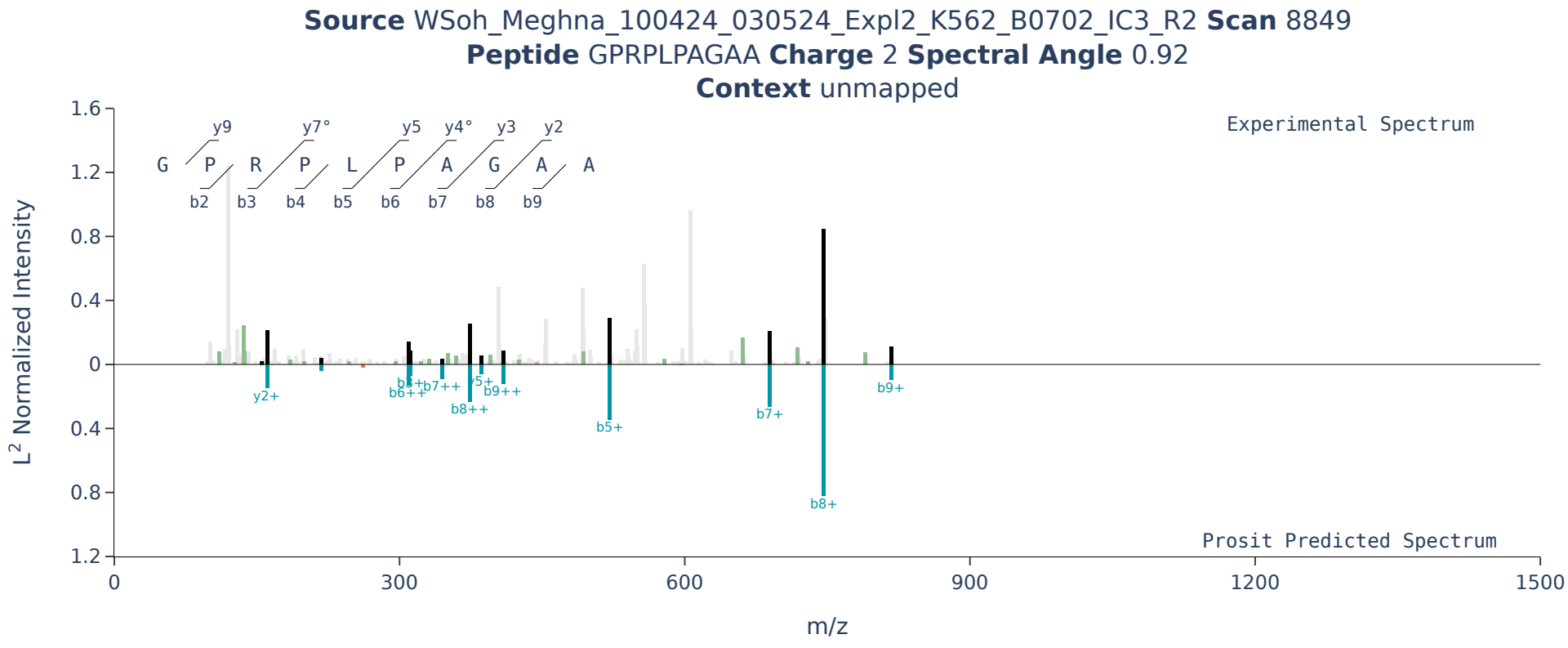
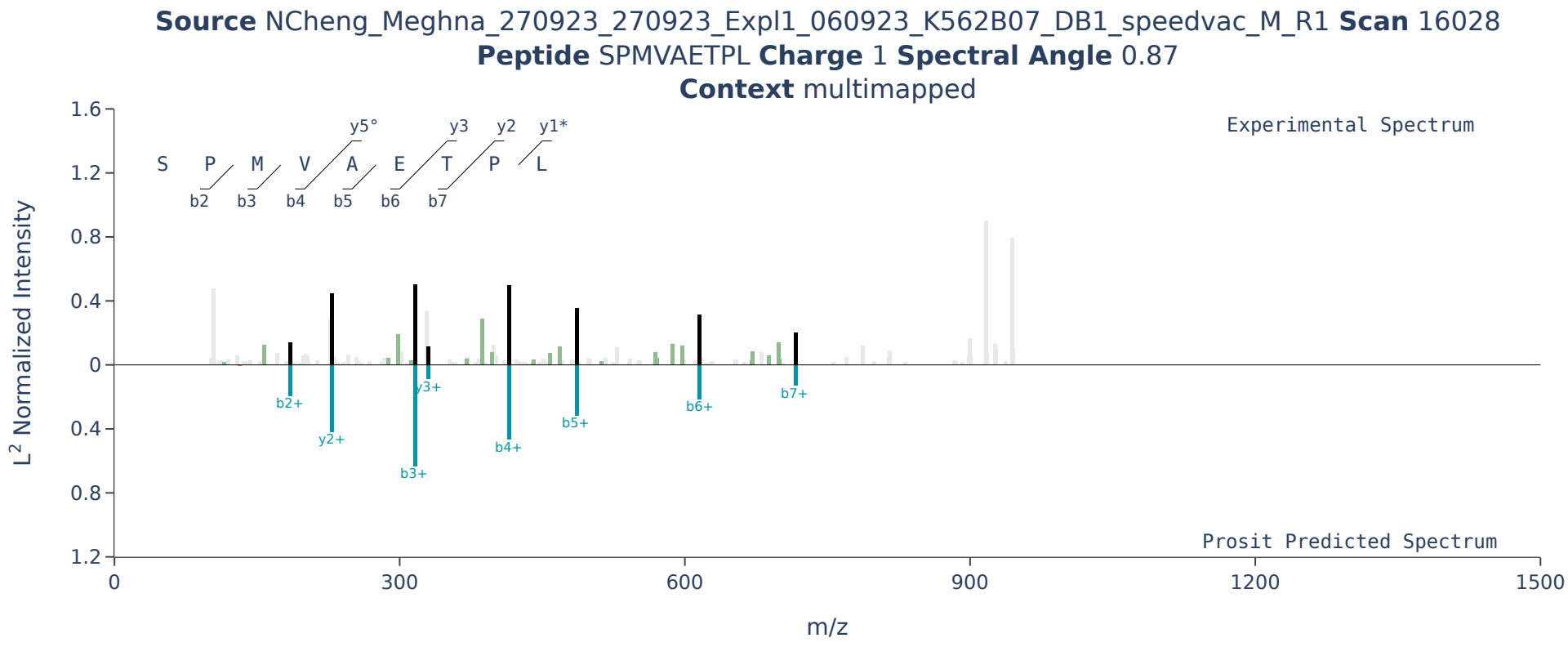
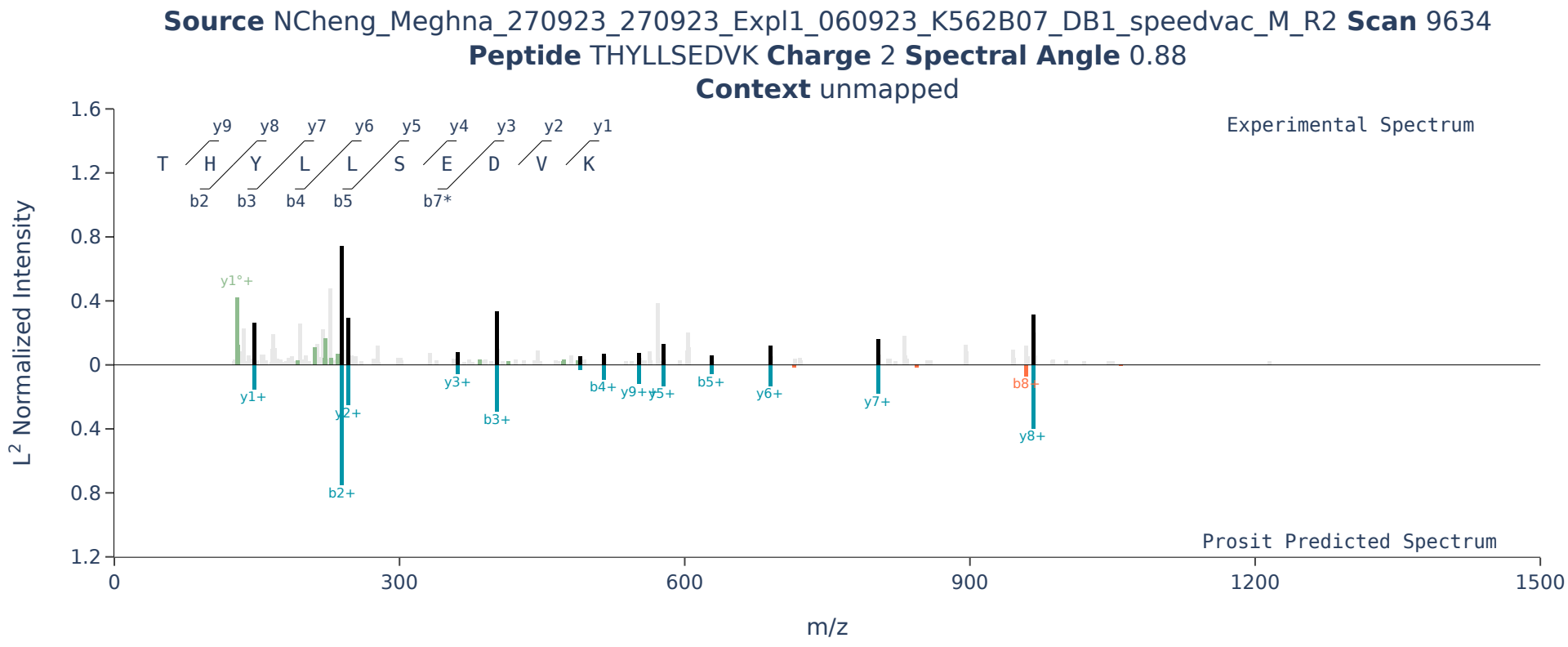
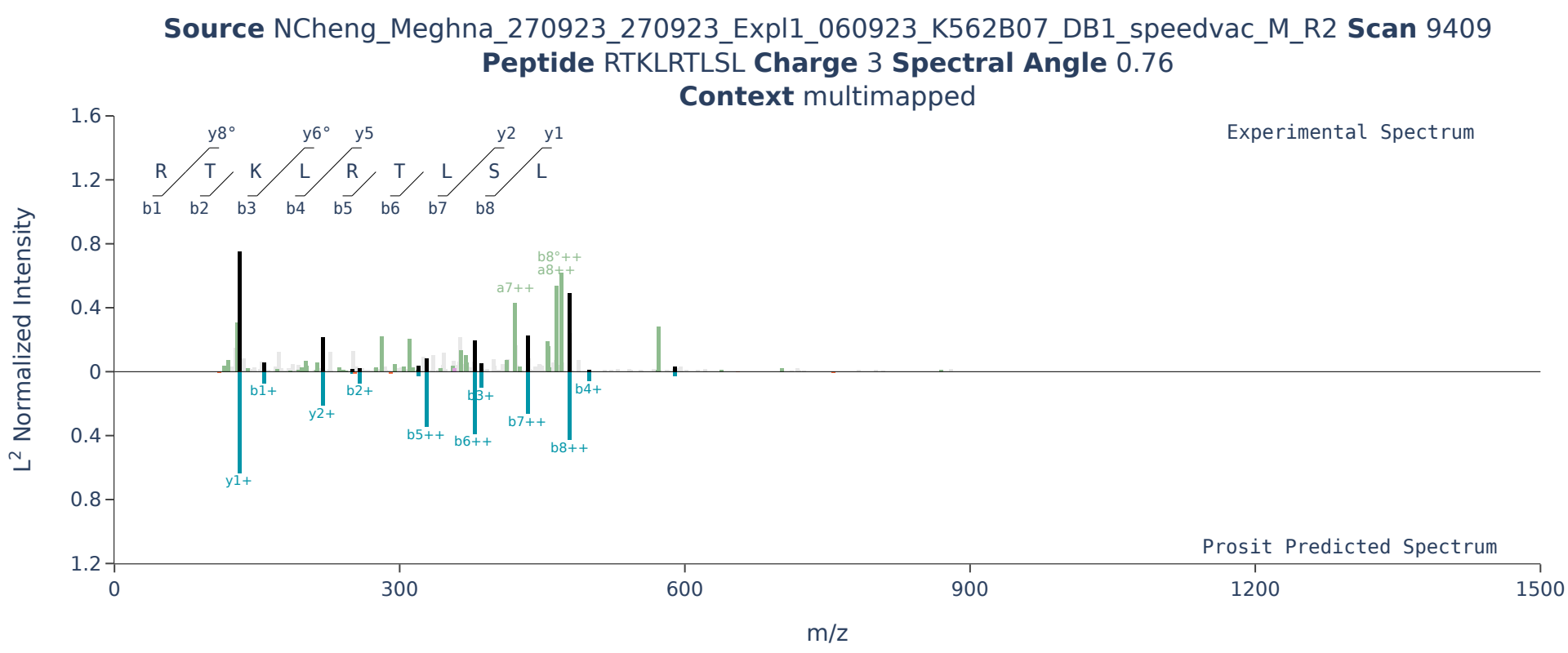
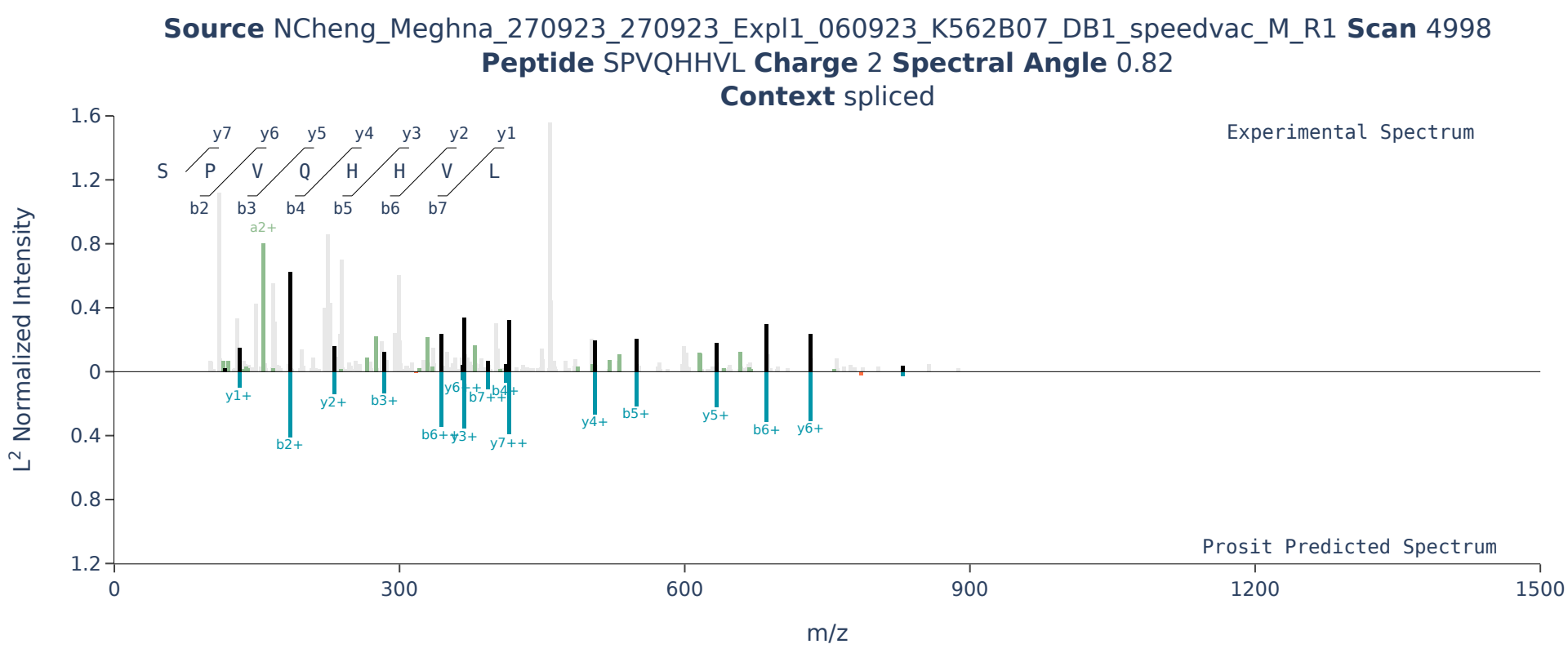
Peptide TPEVSKTTNPL **Charge** 2 **Spectral Angle** 0.91
Context cryptic

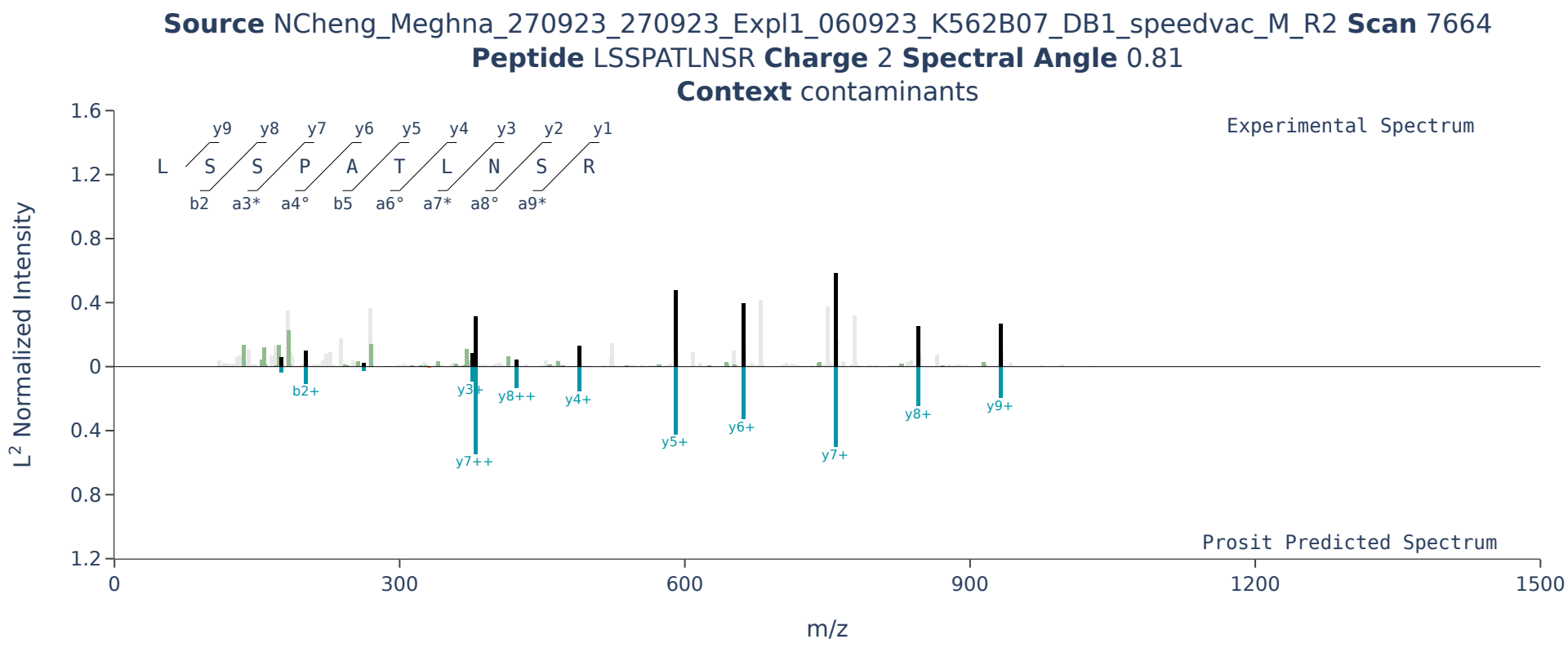
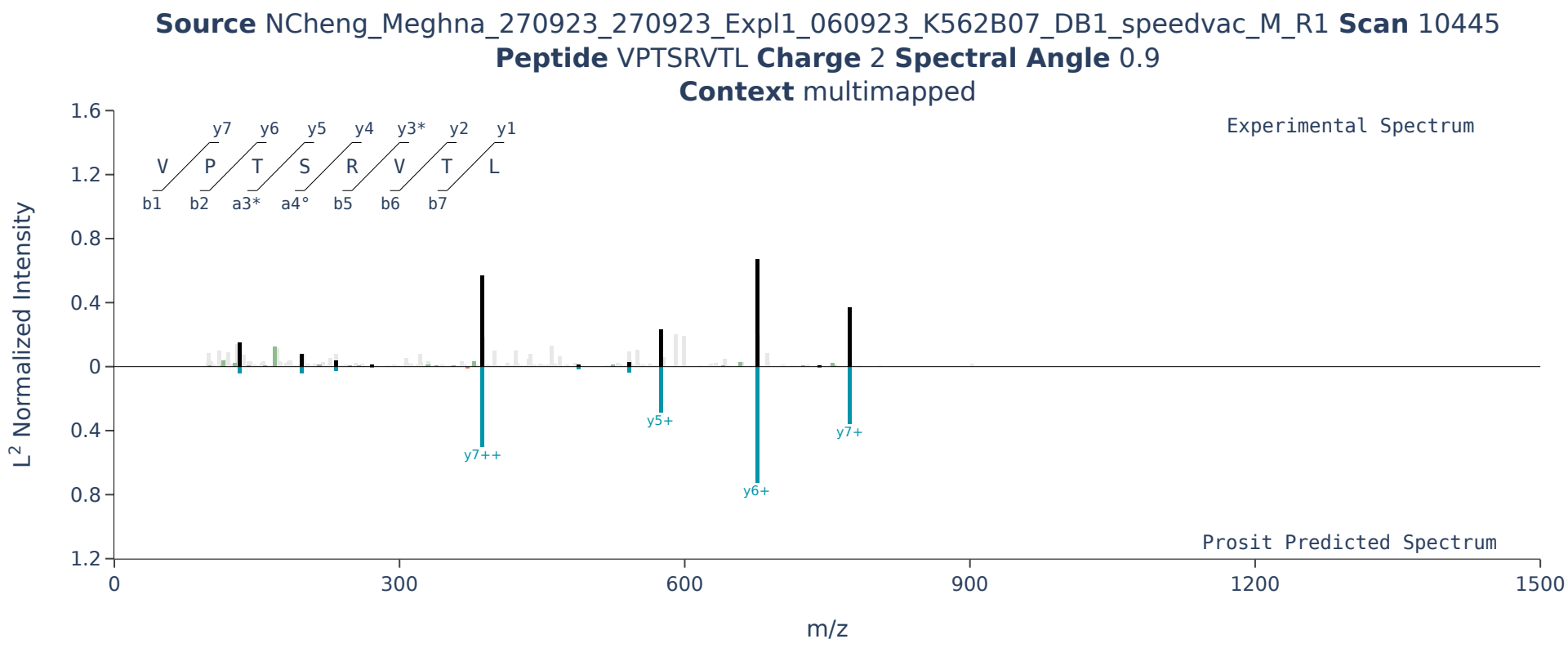
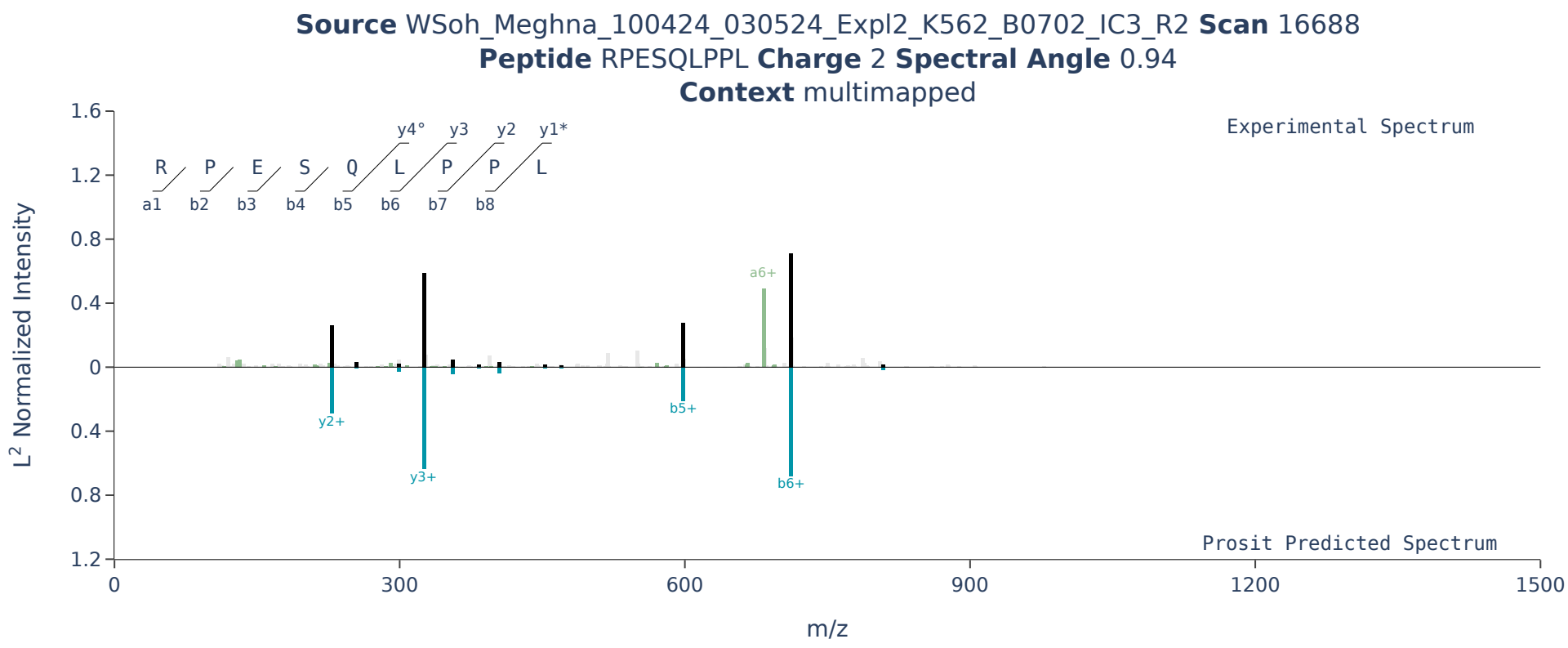
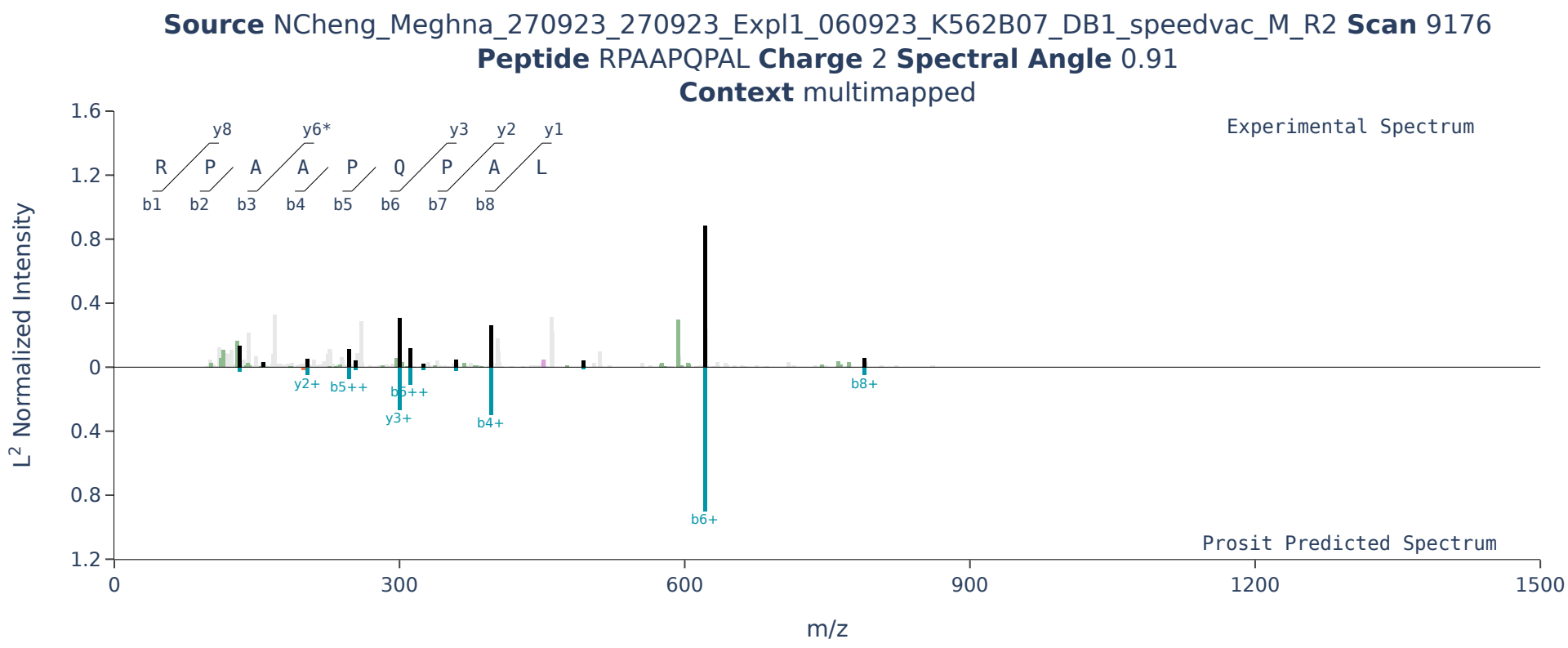
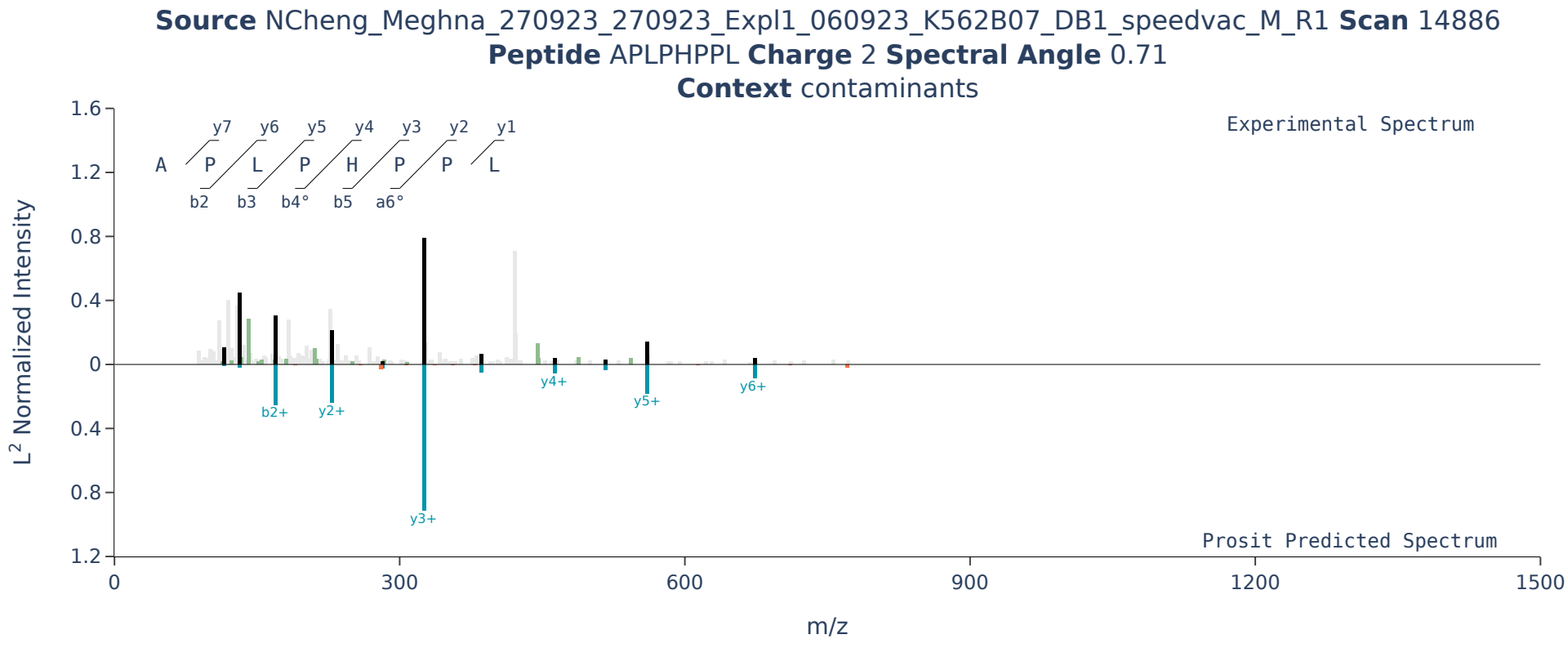
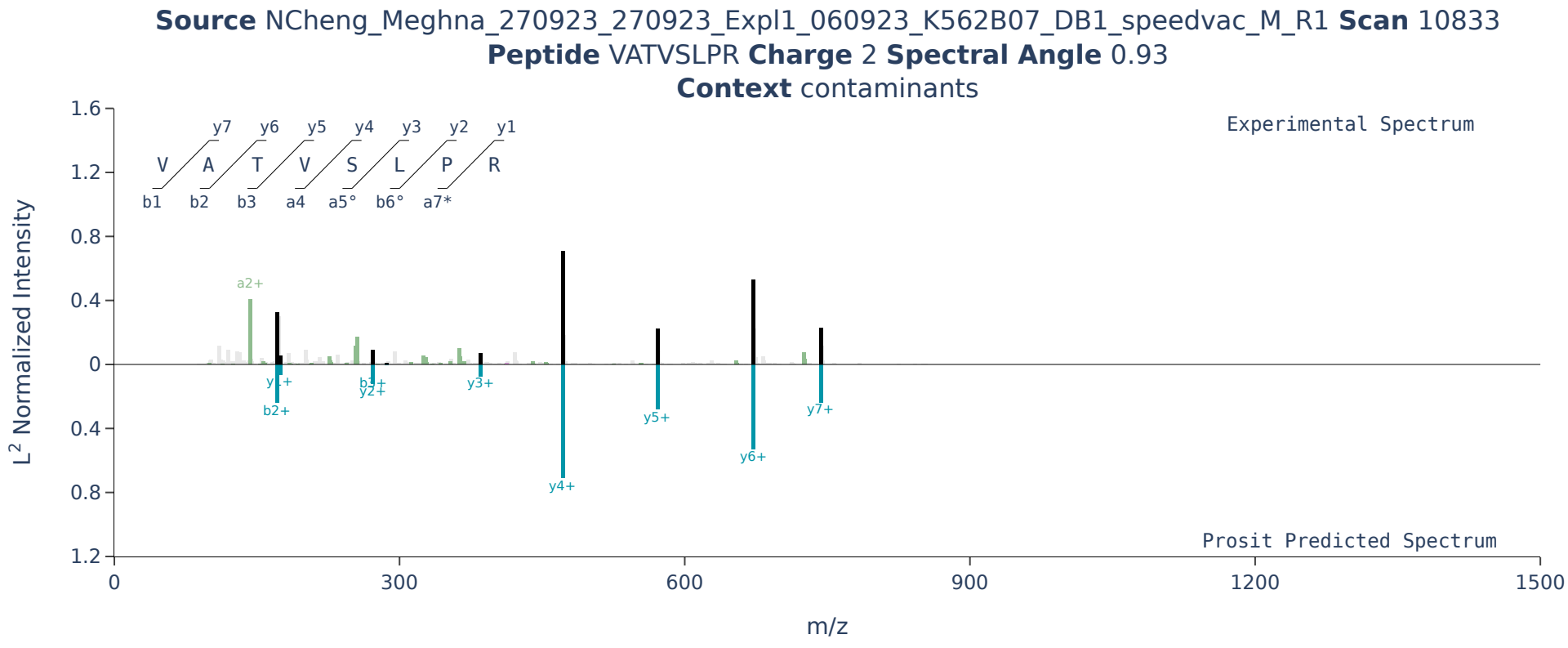
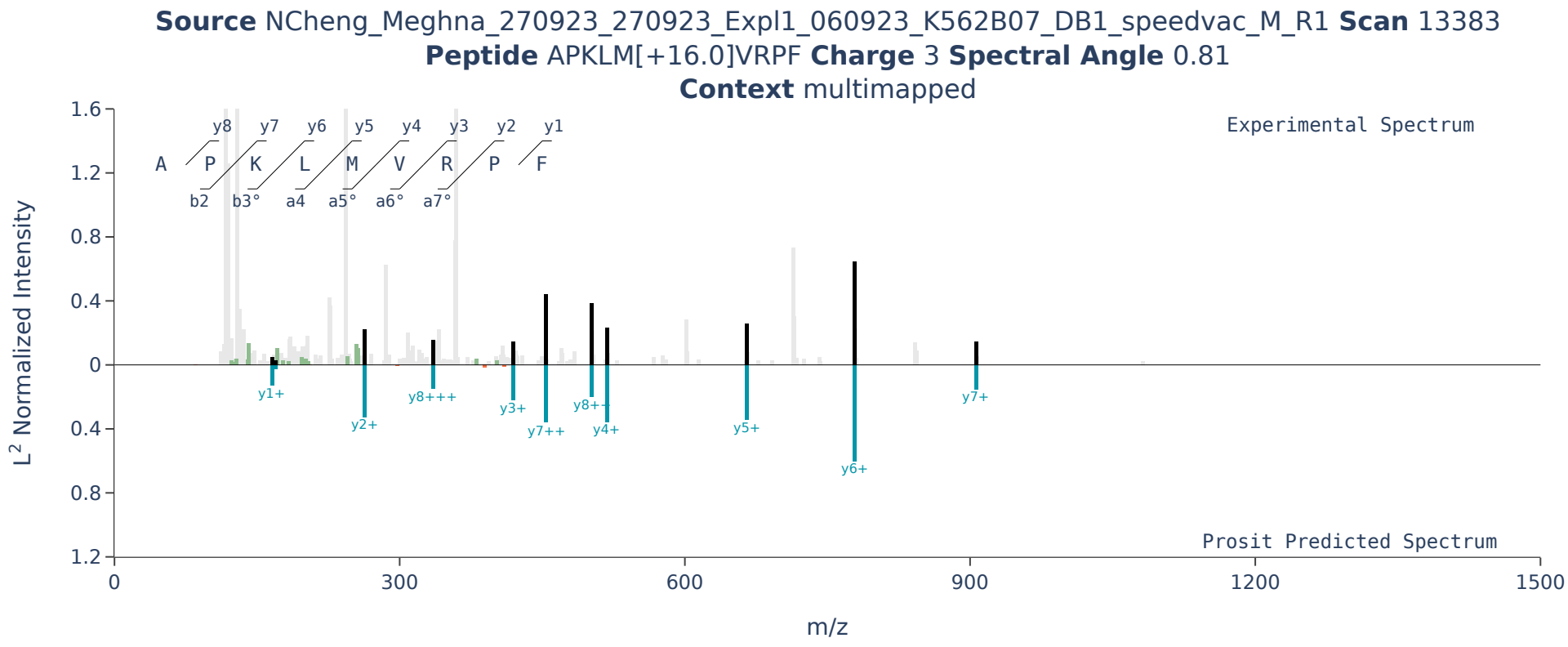
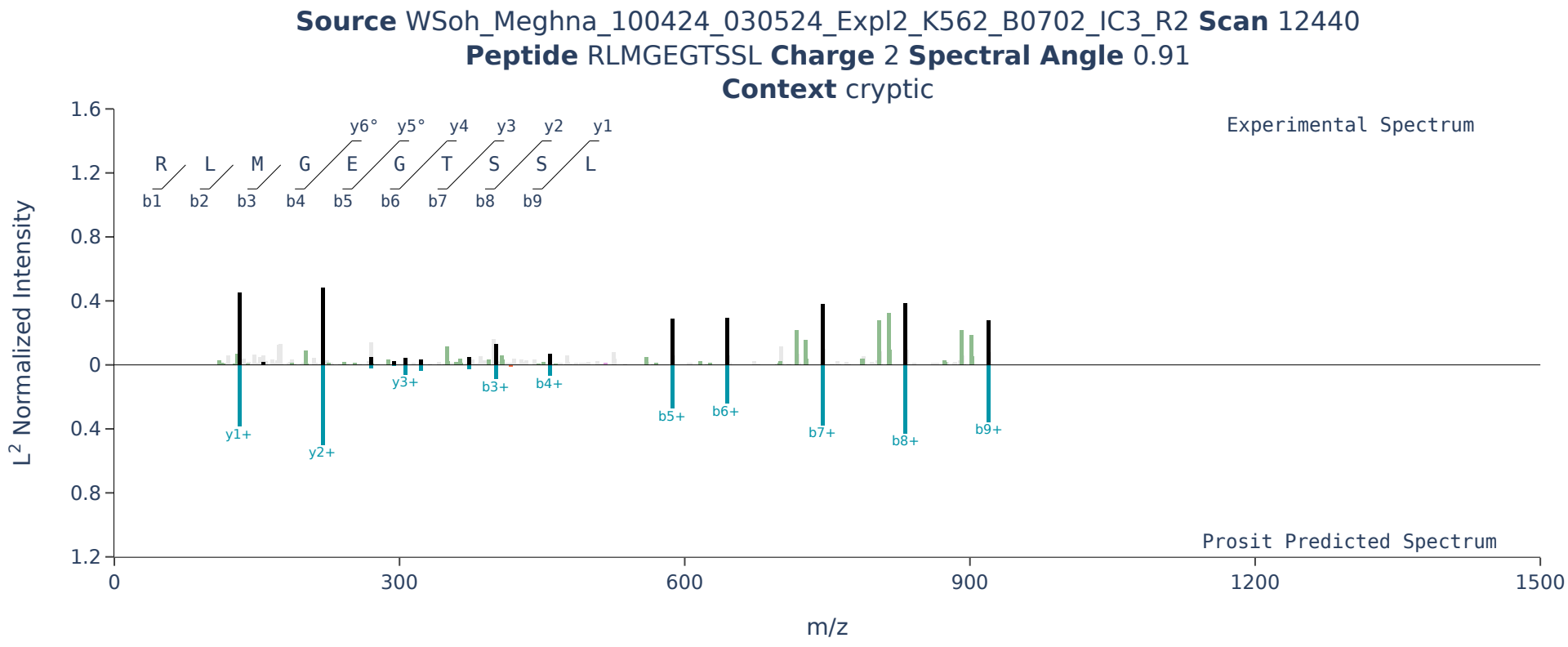
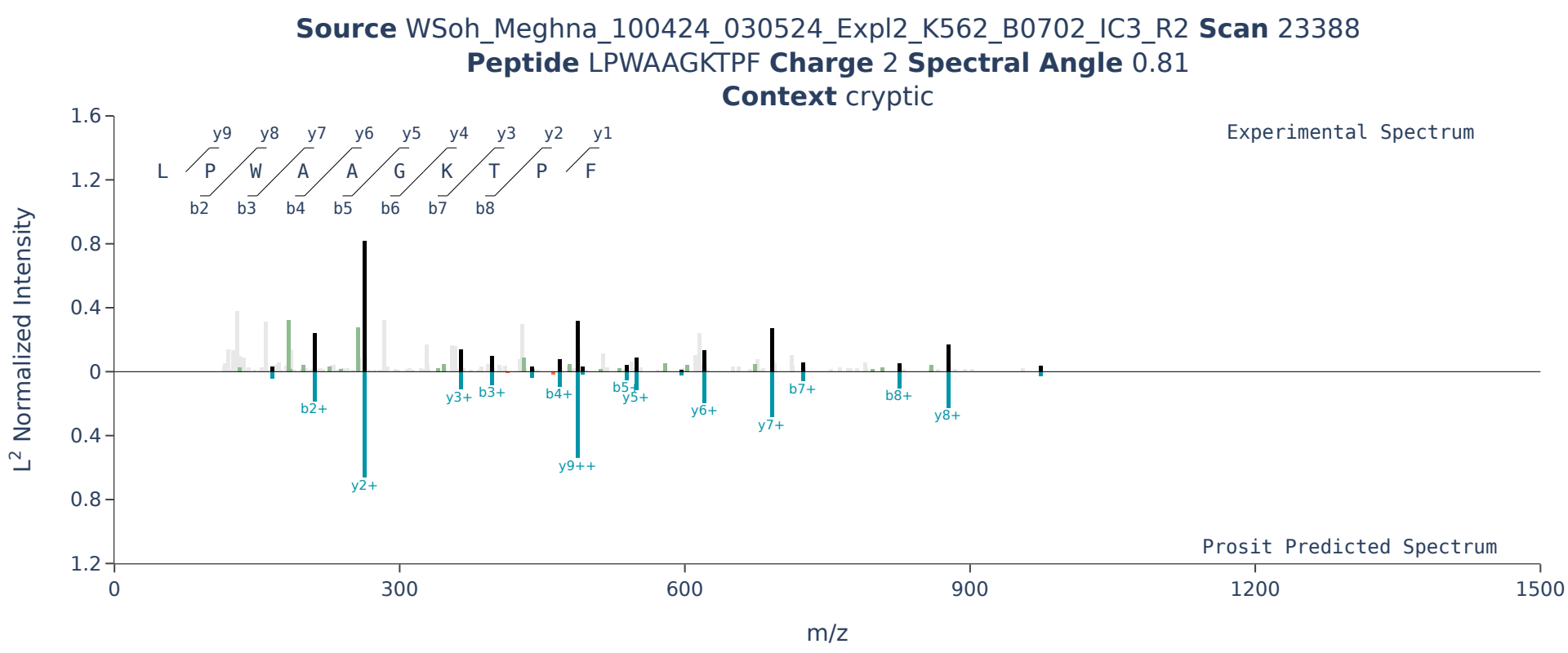
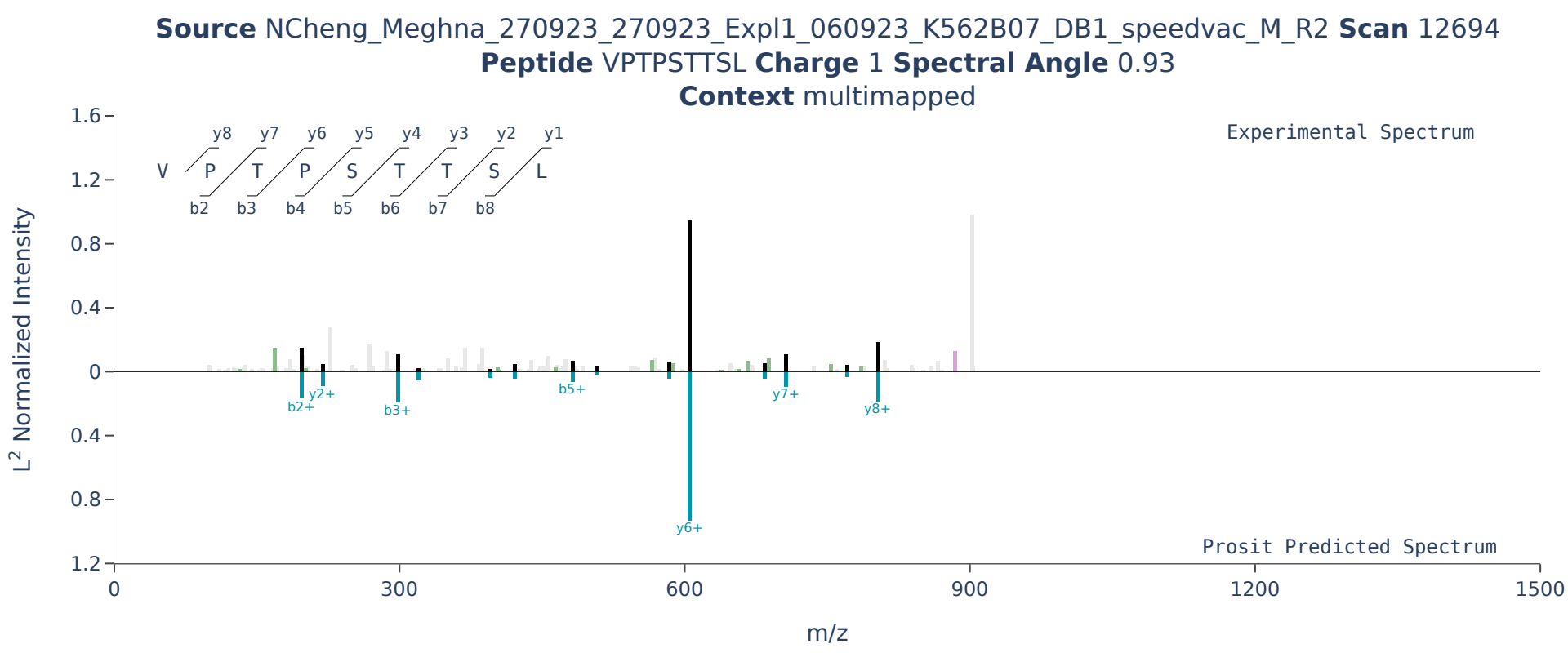


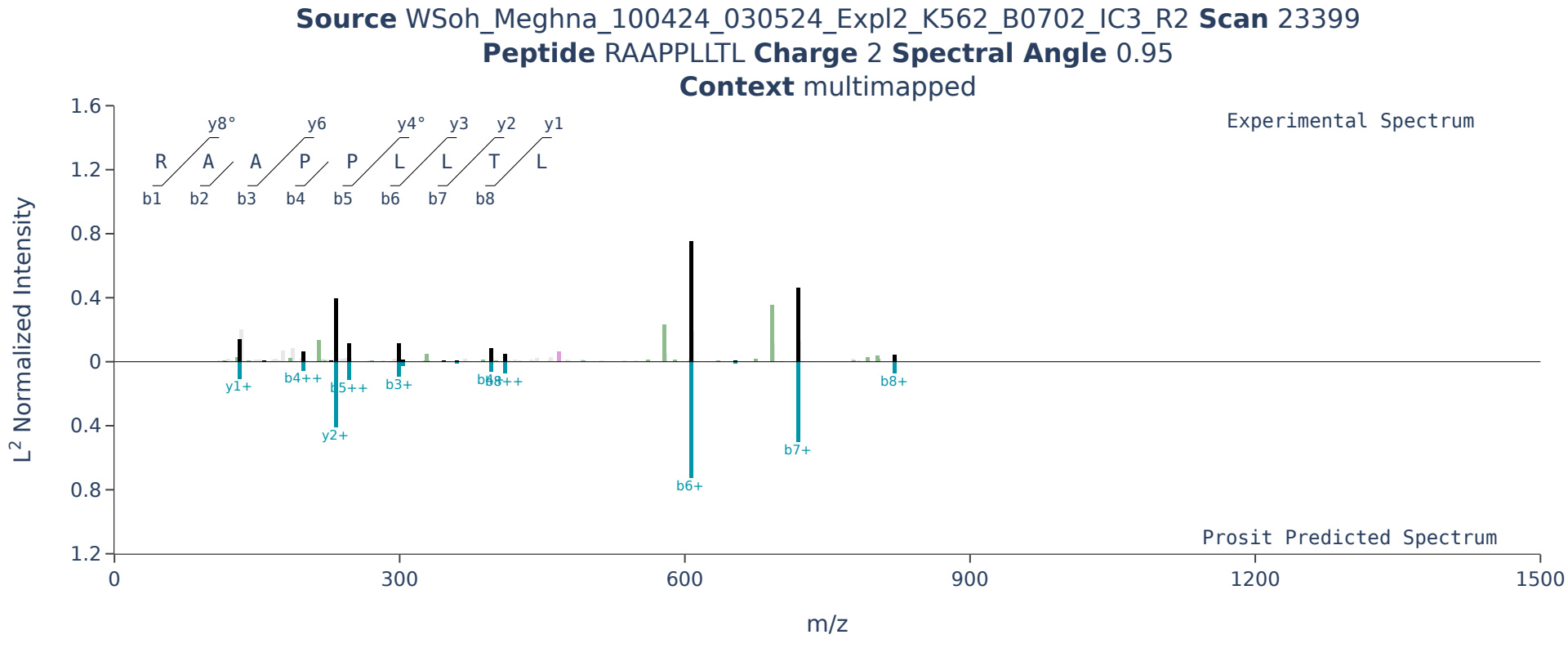
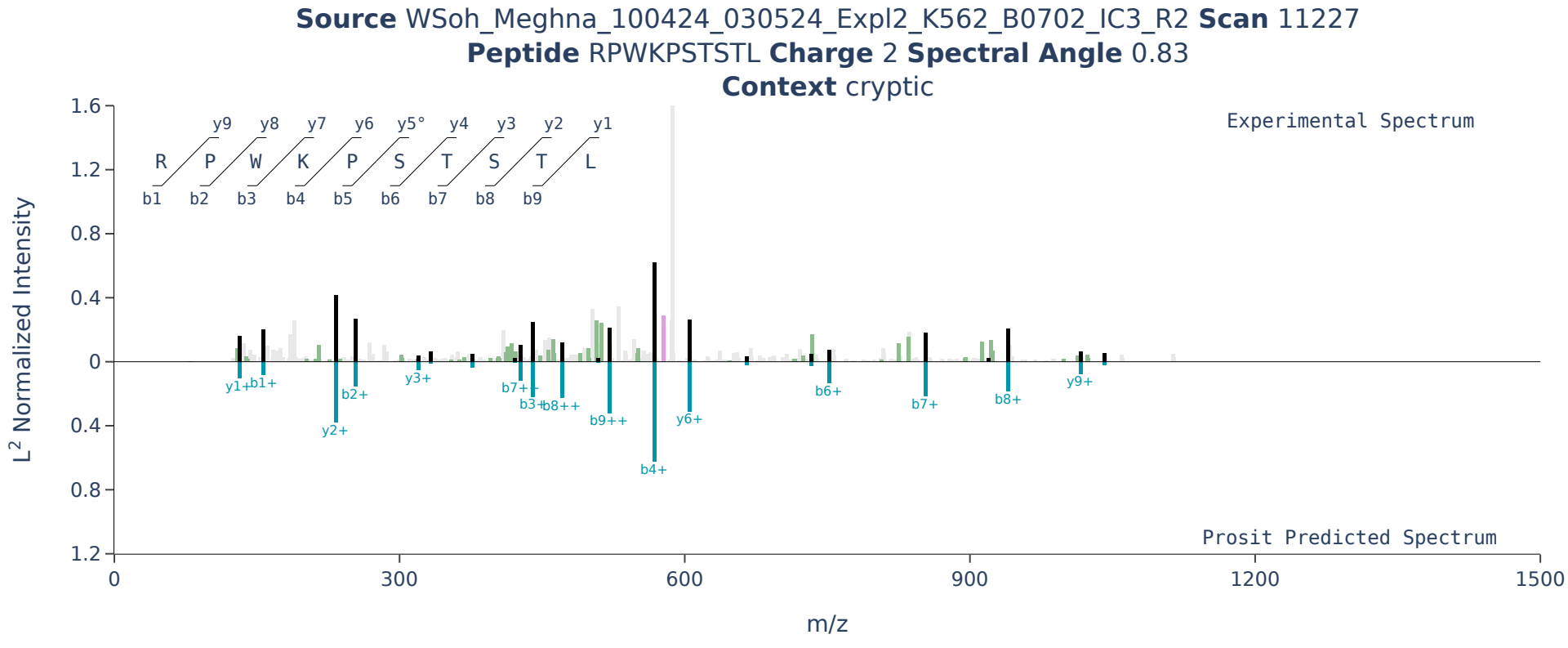
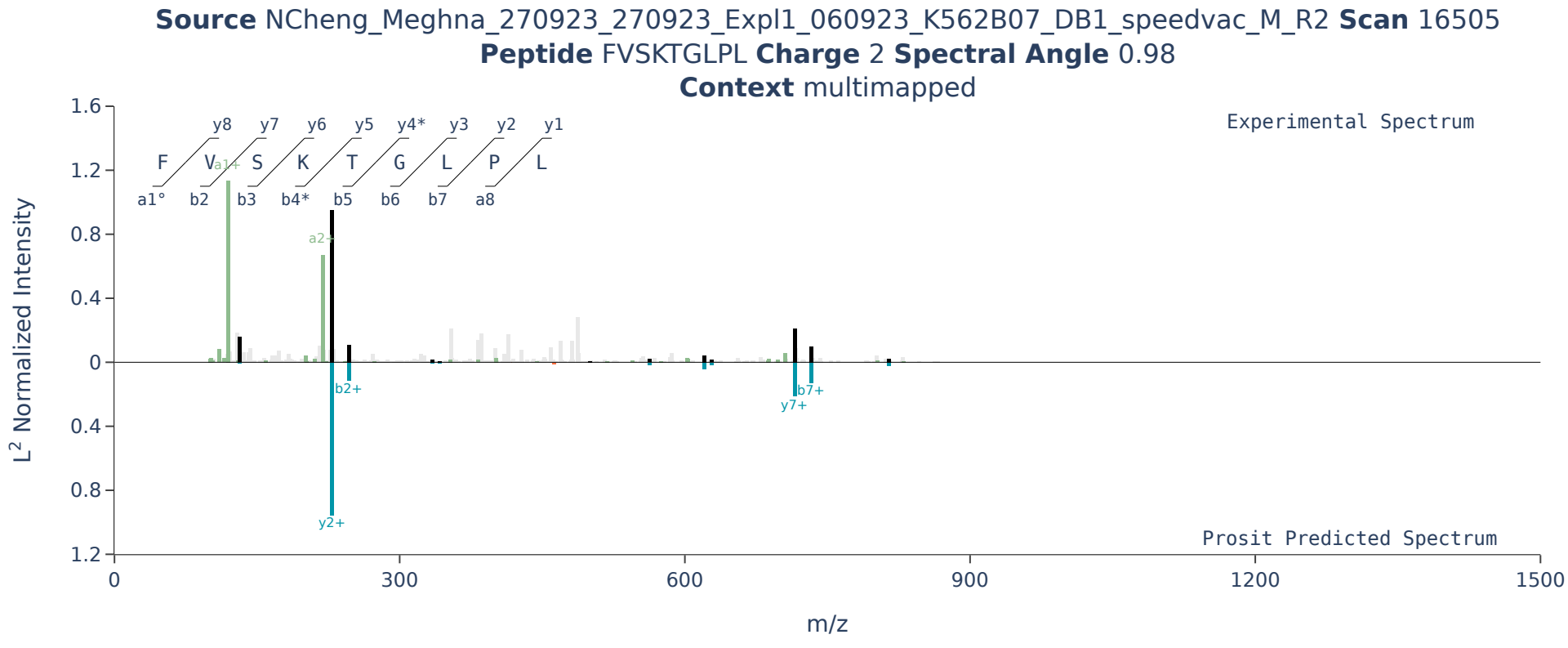
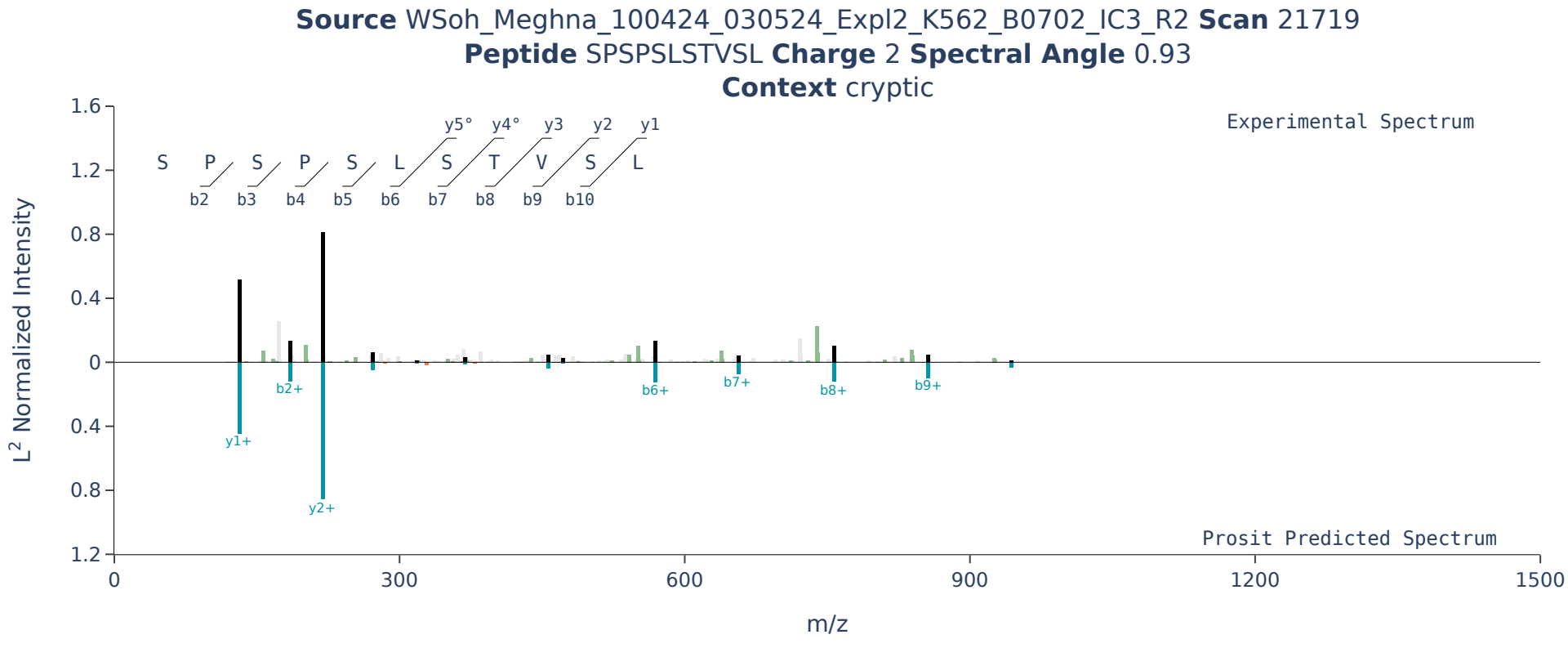
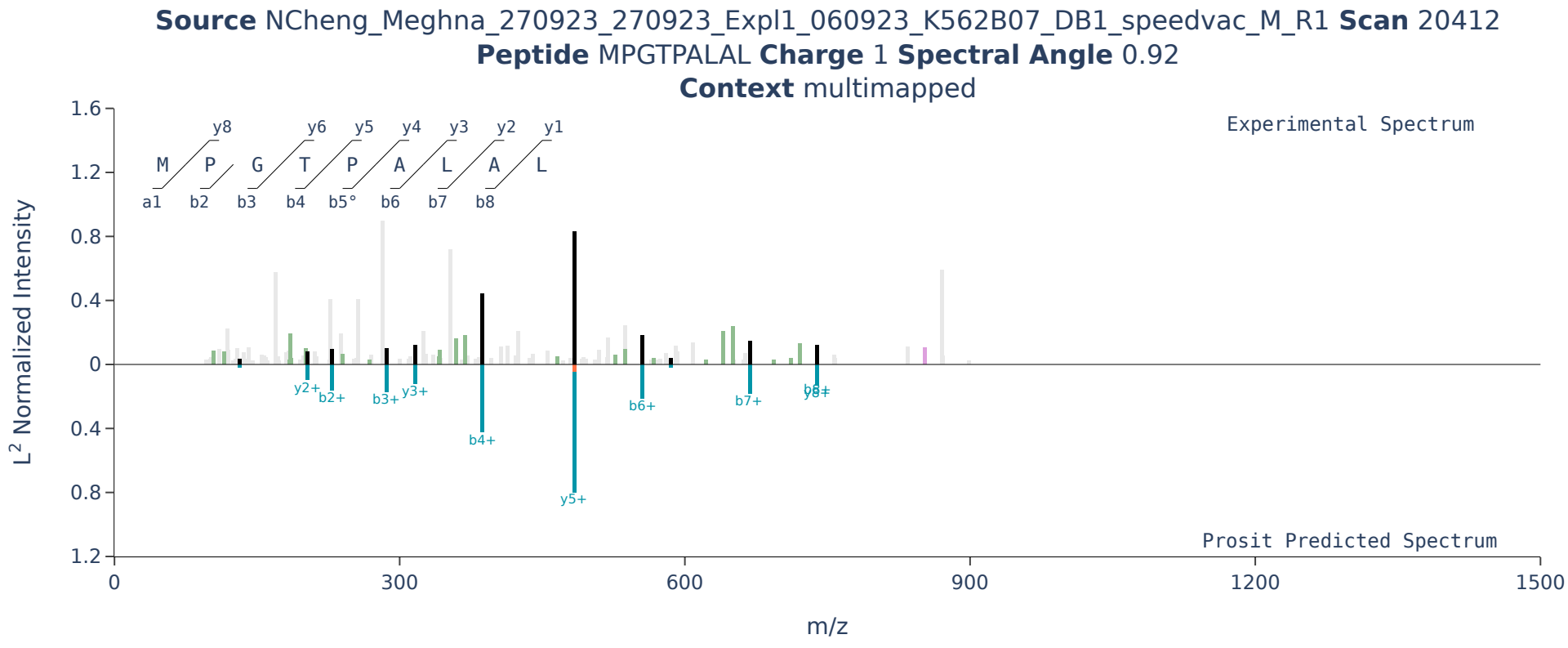
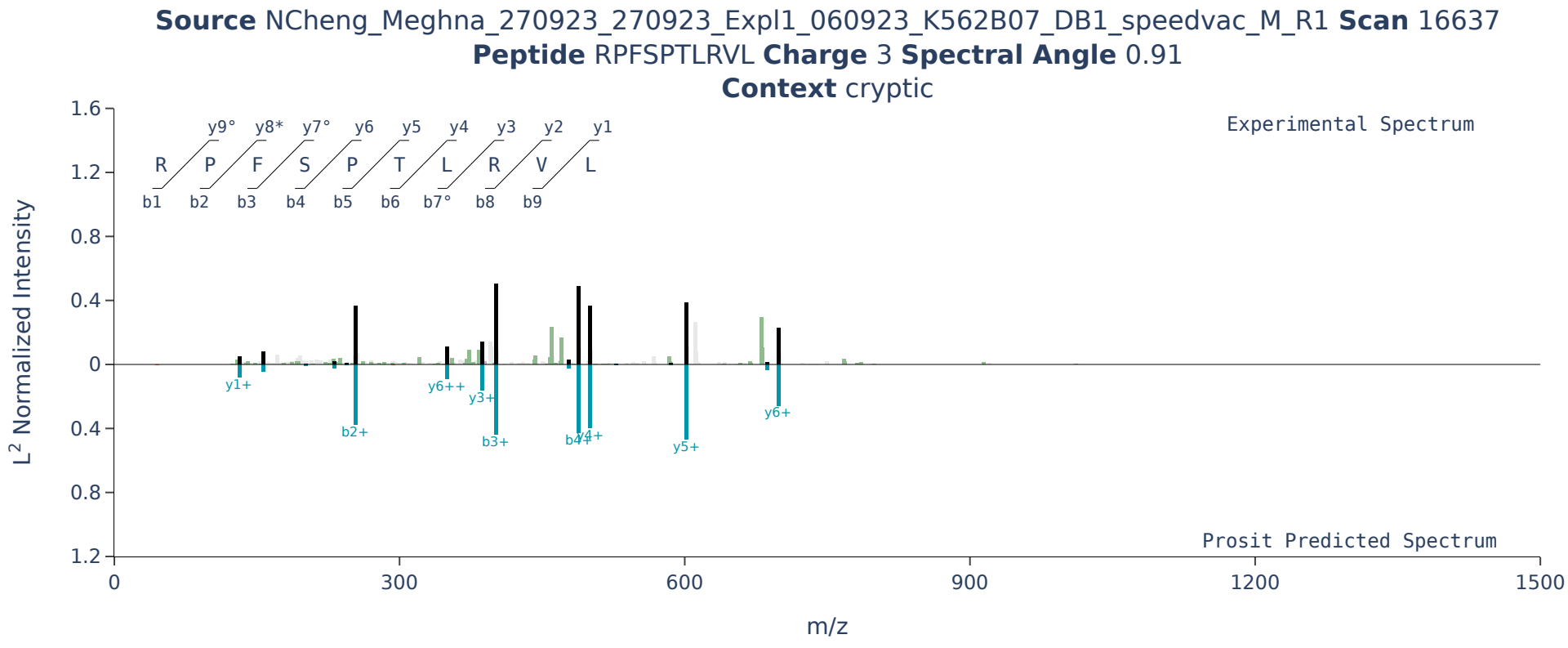
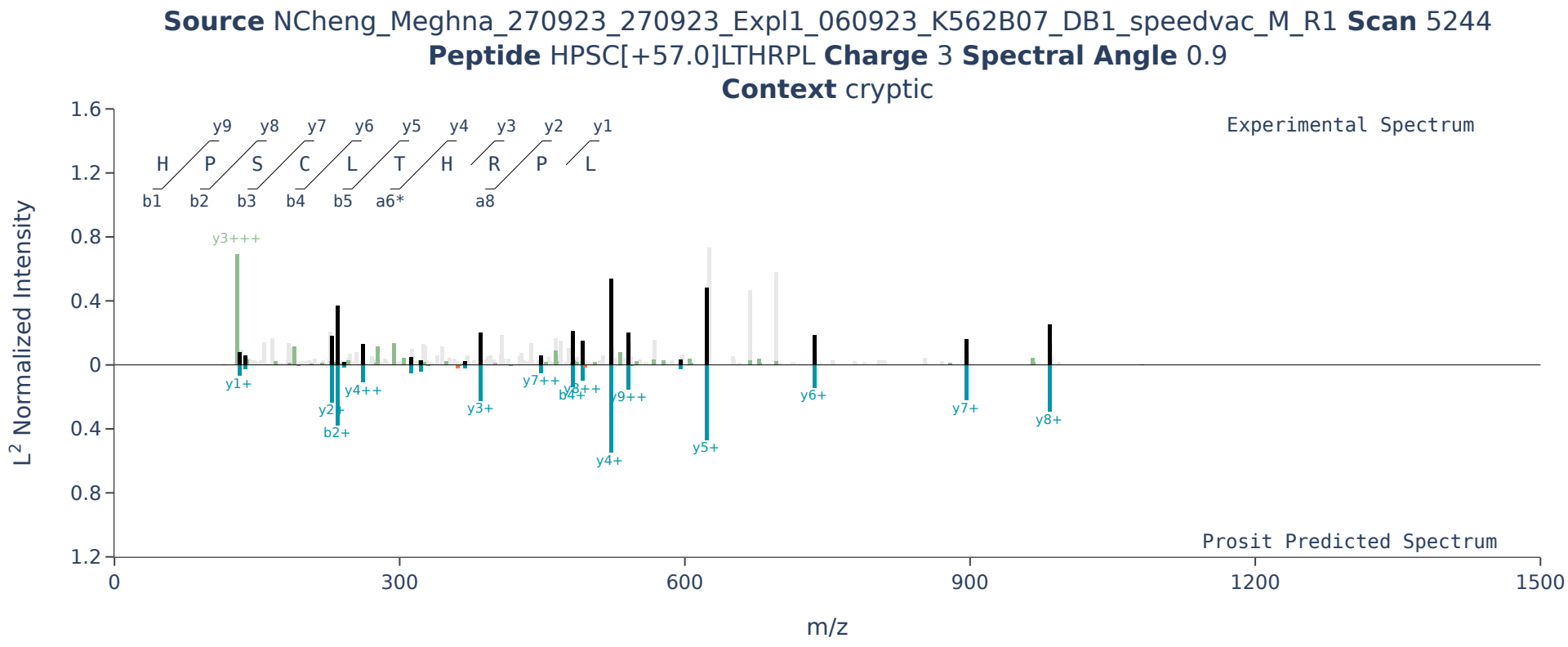
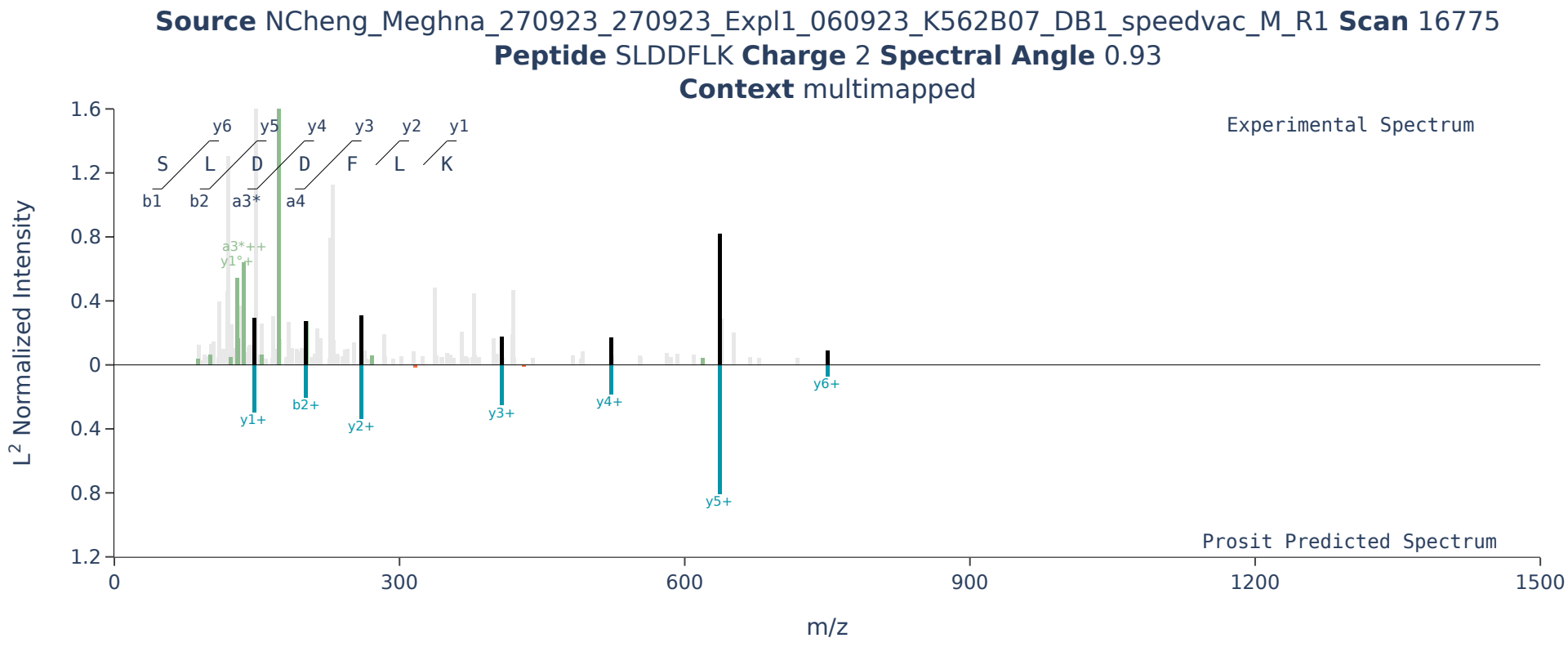
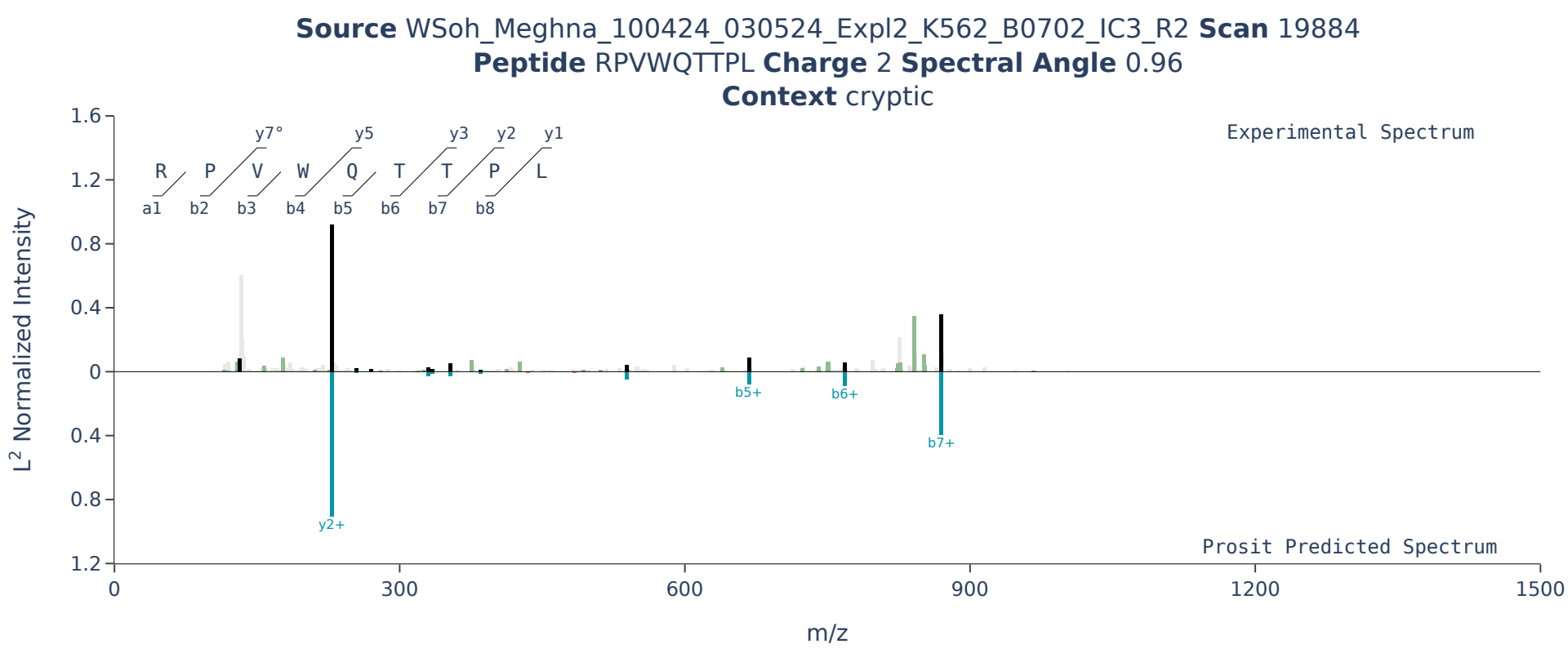
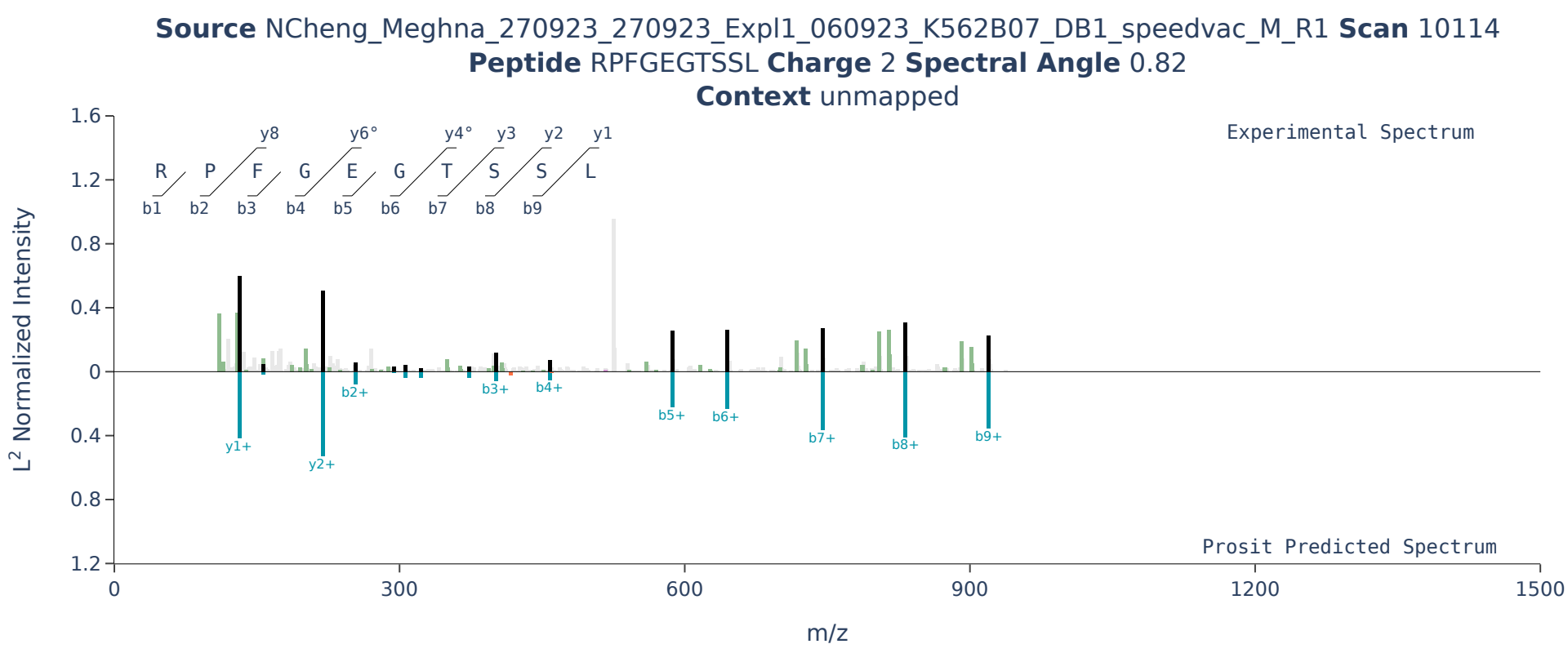


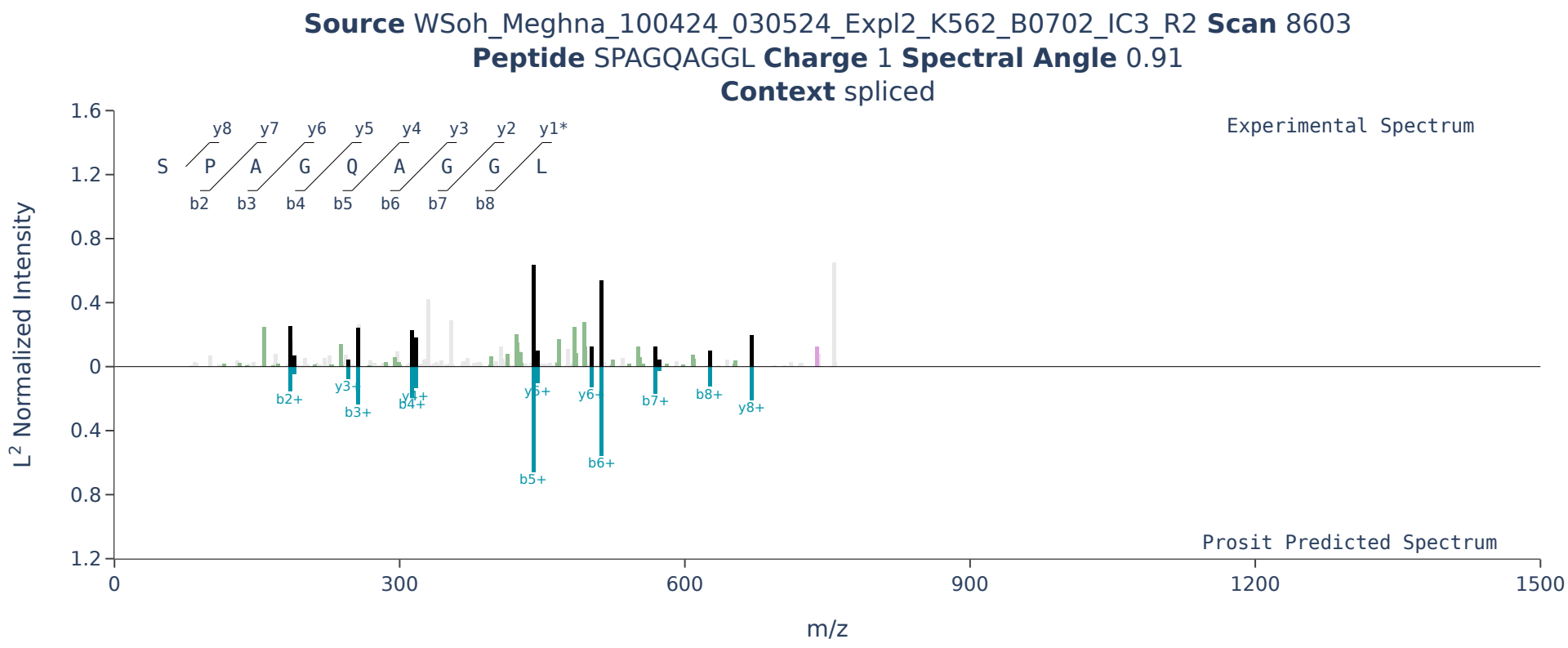
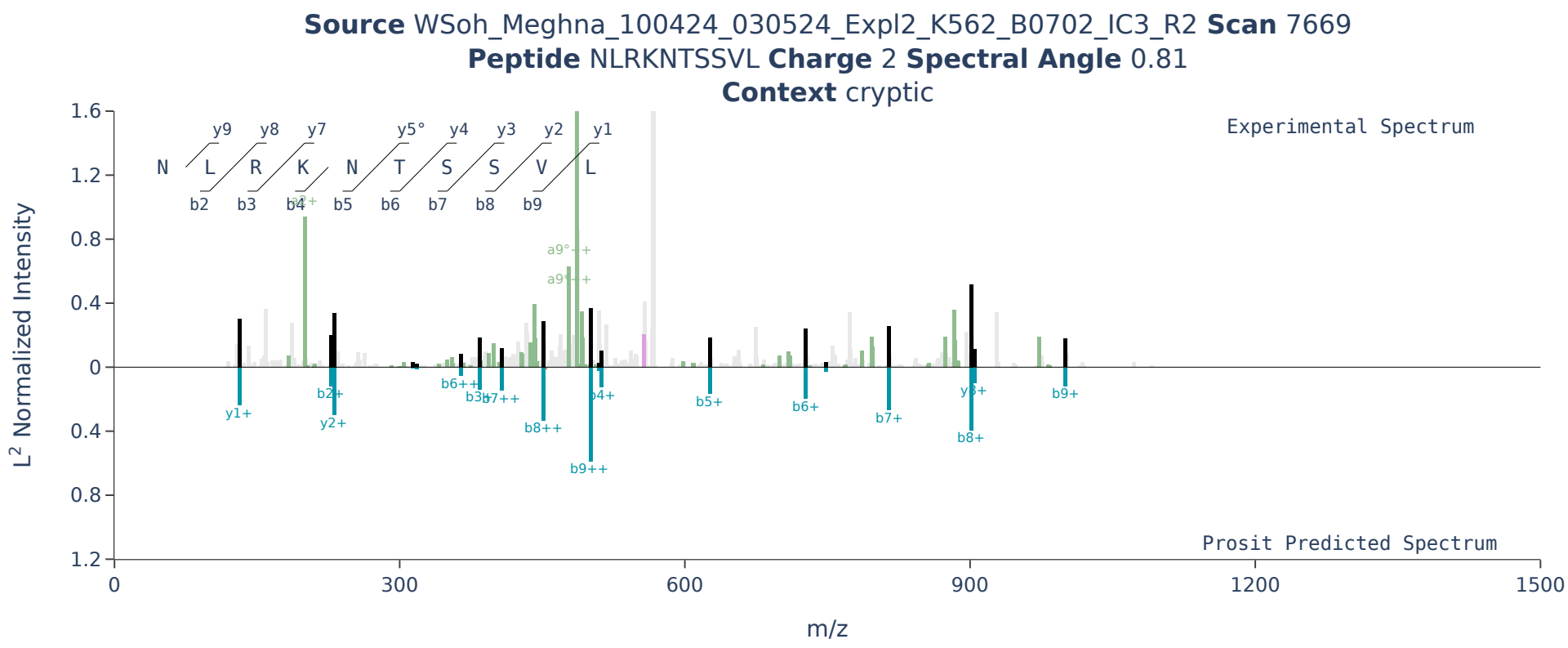
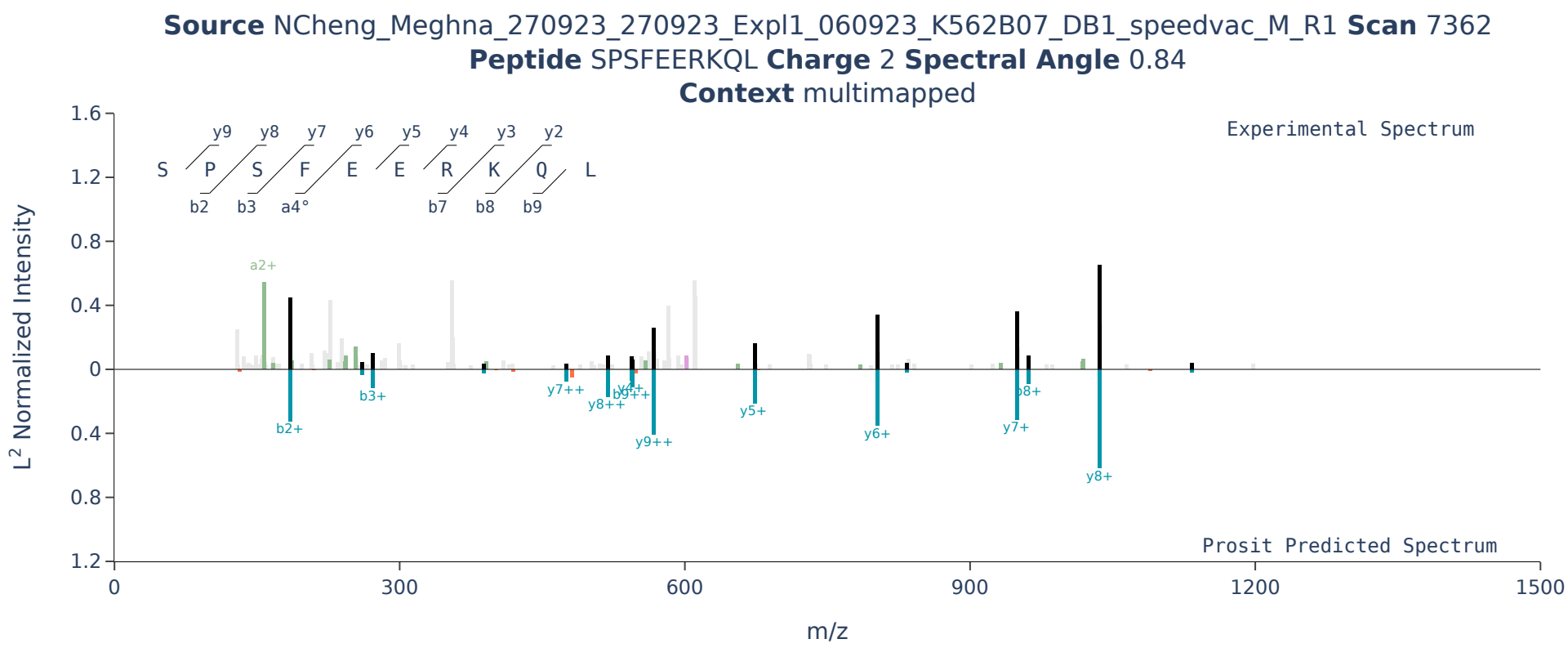
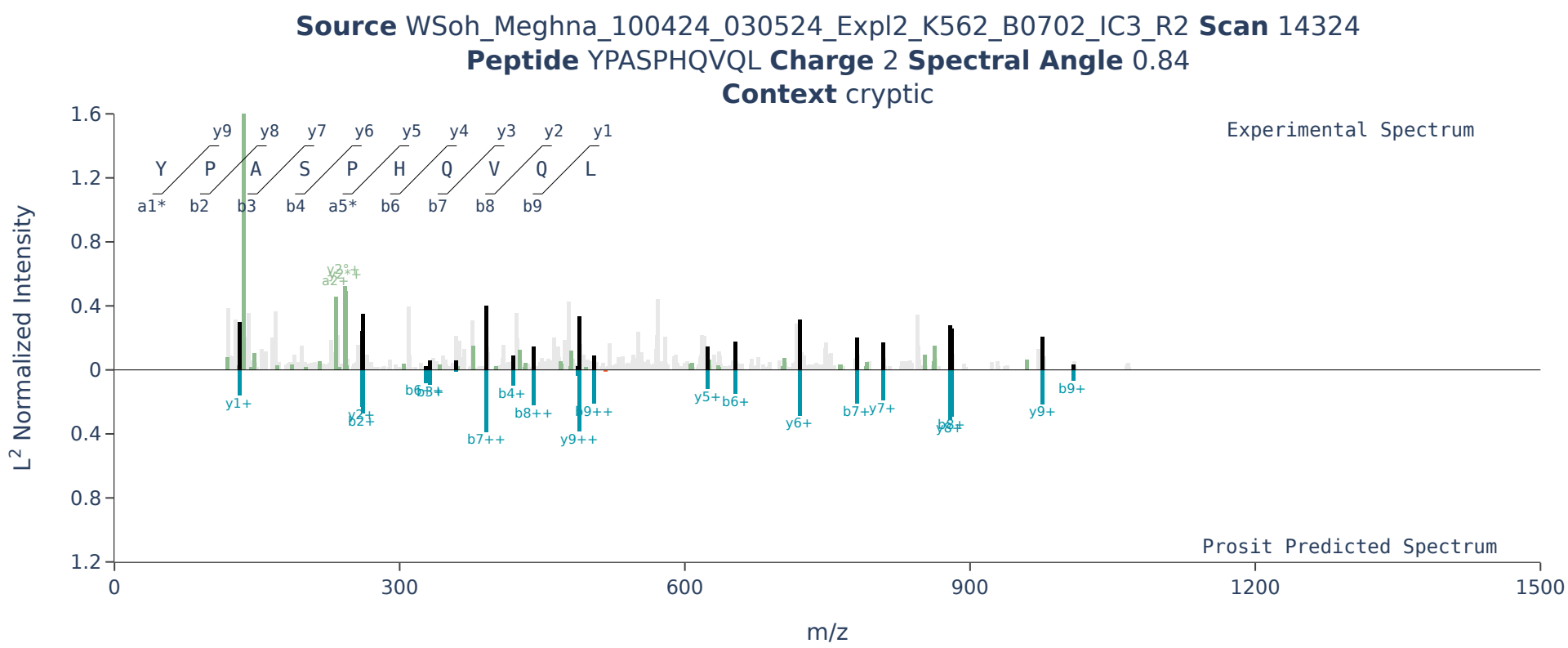
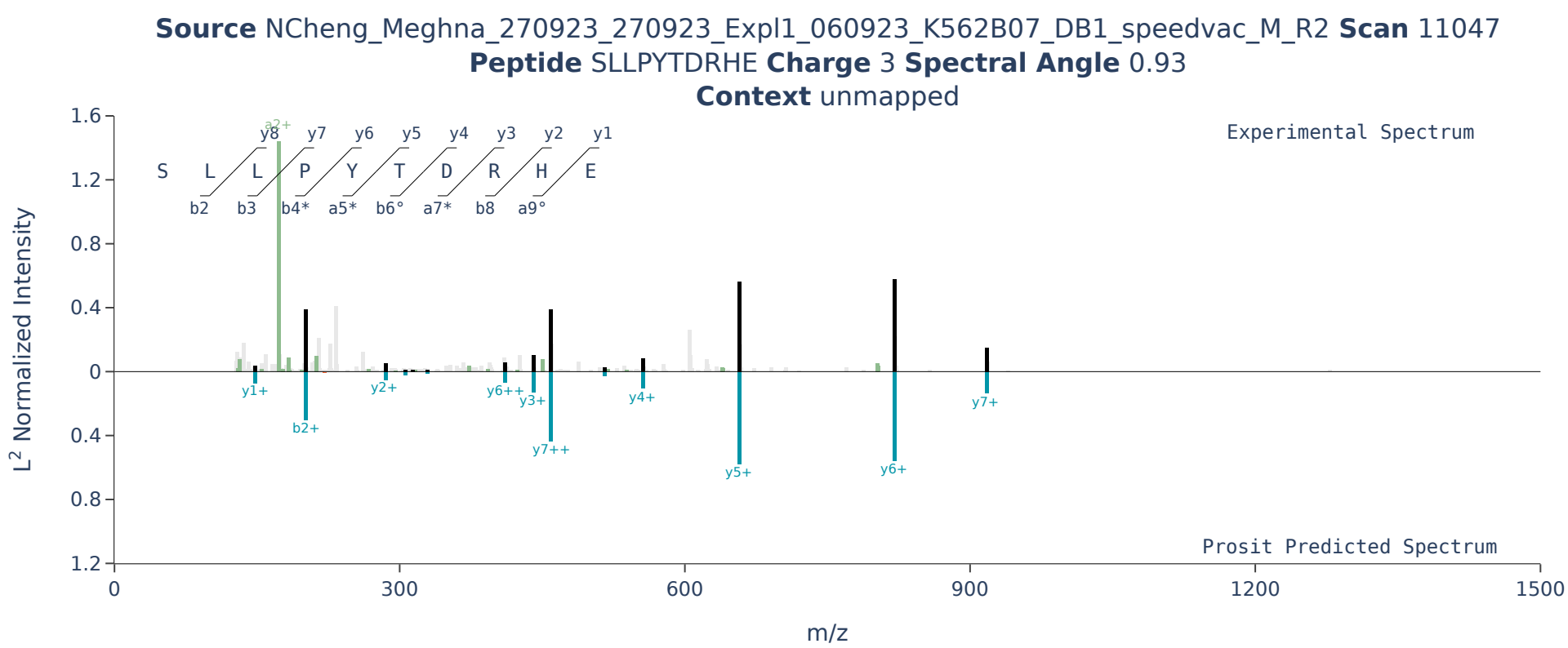


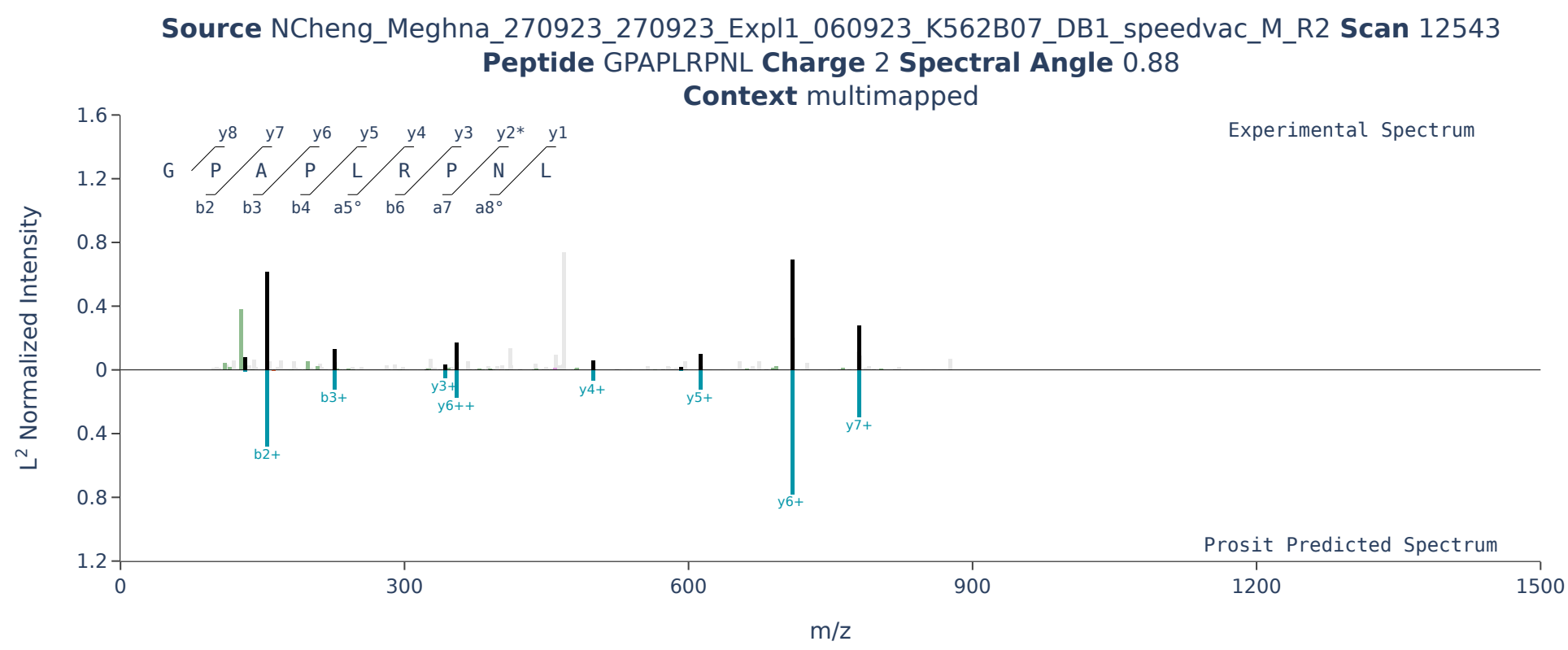
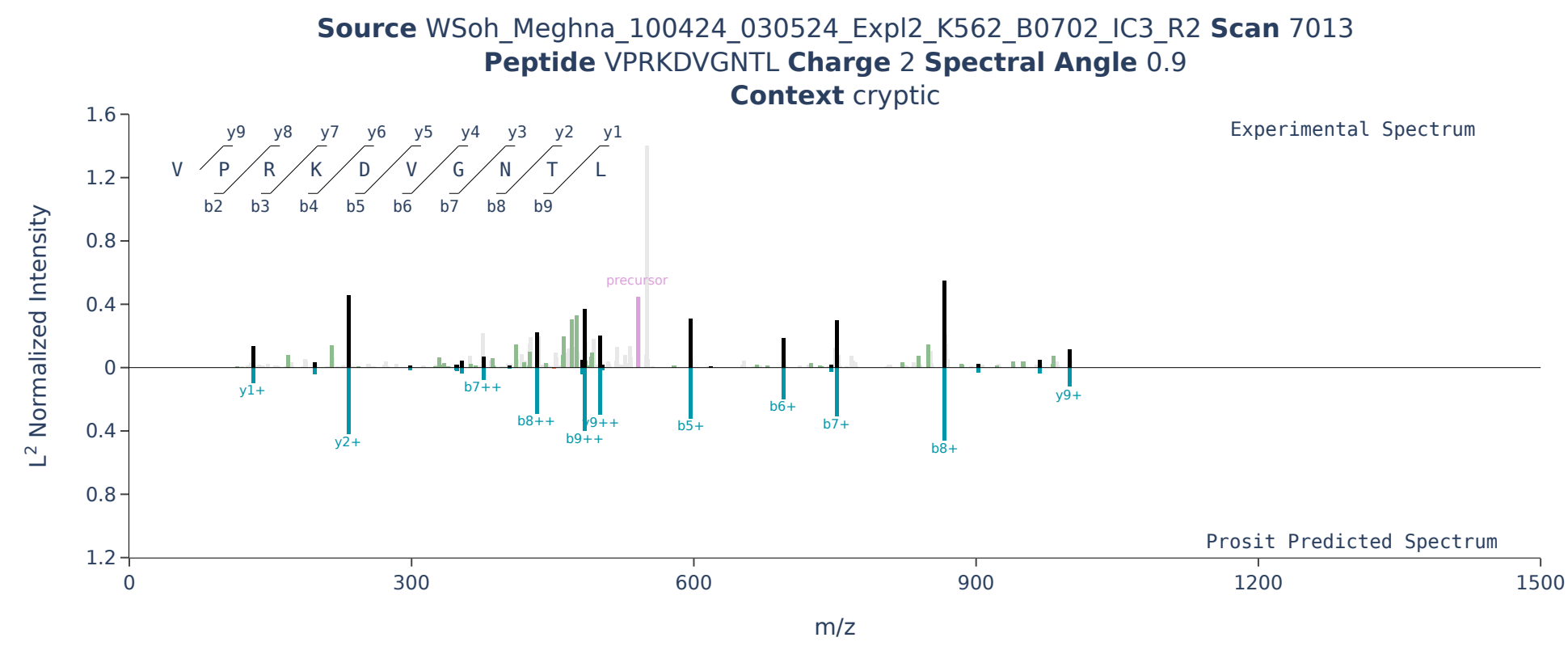
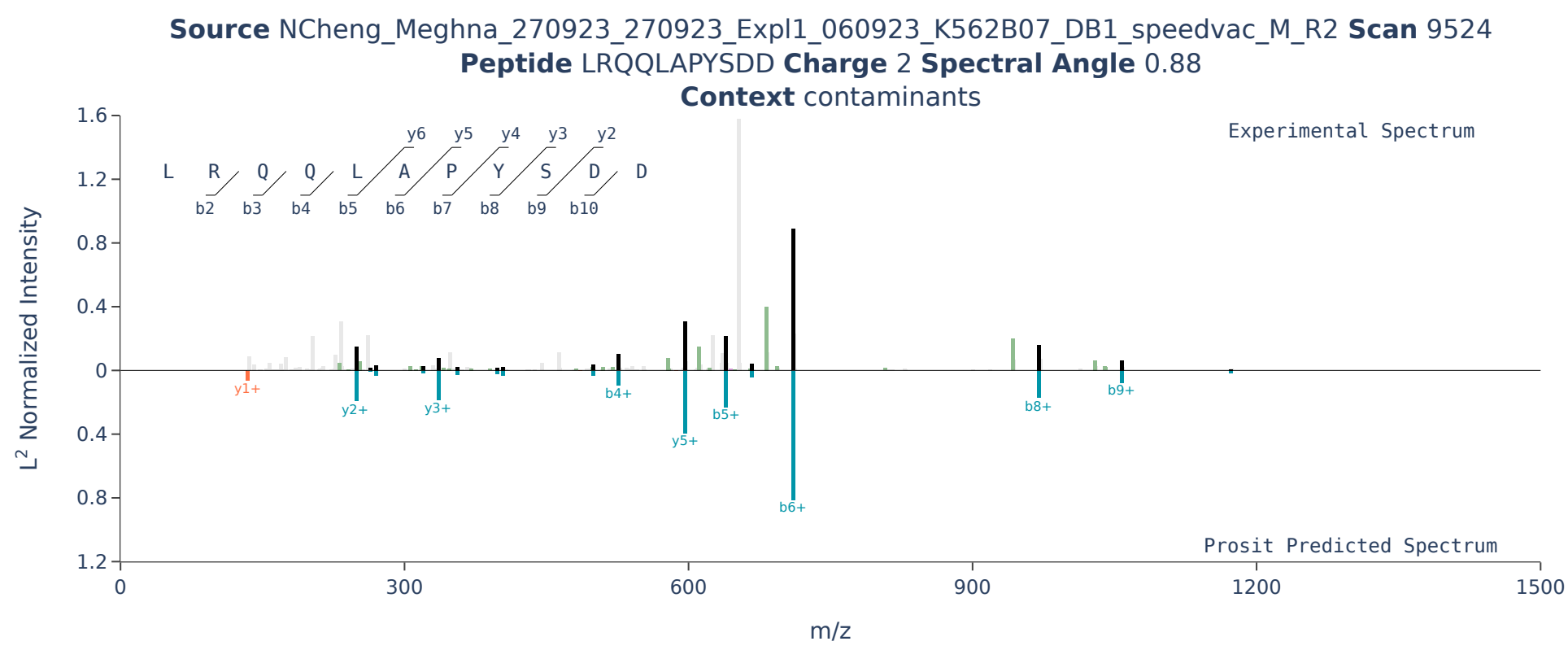
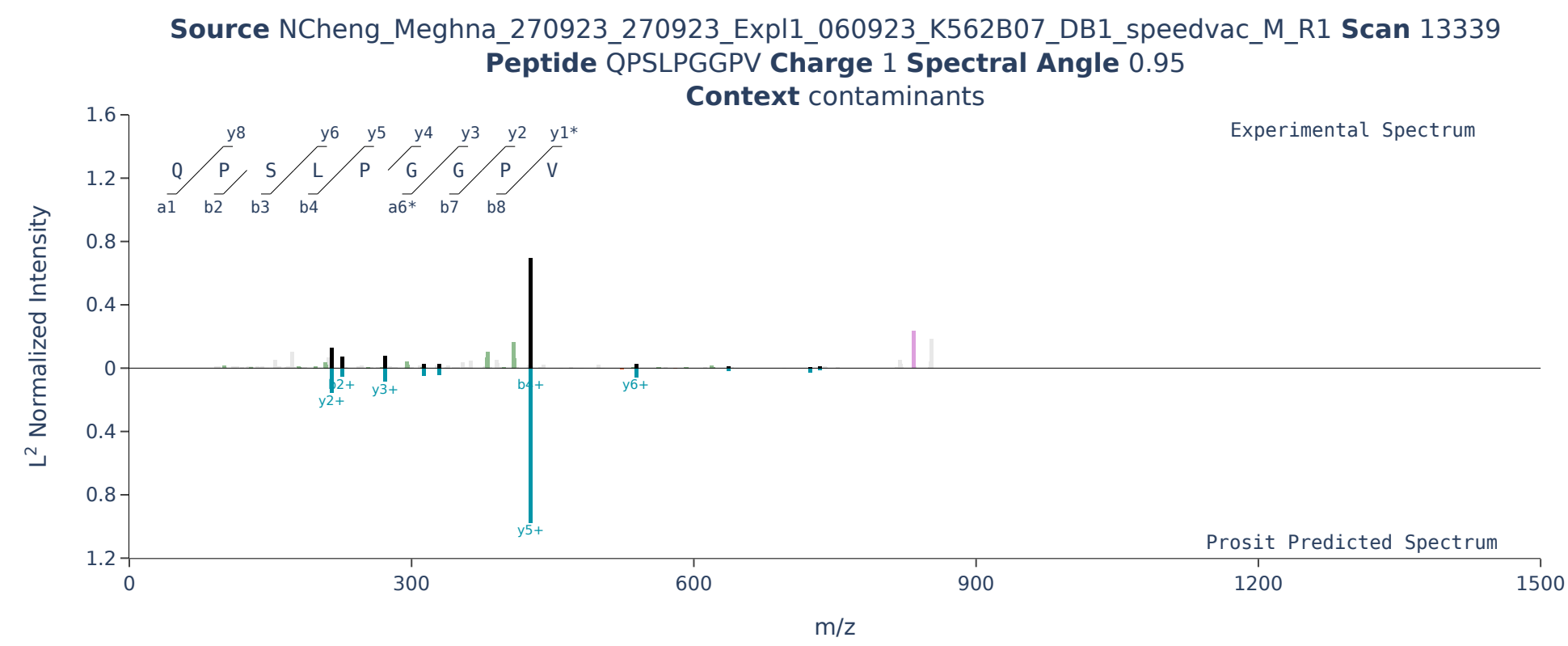
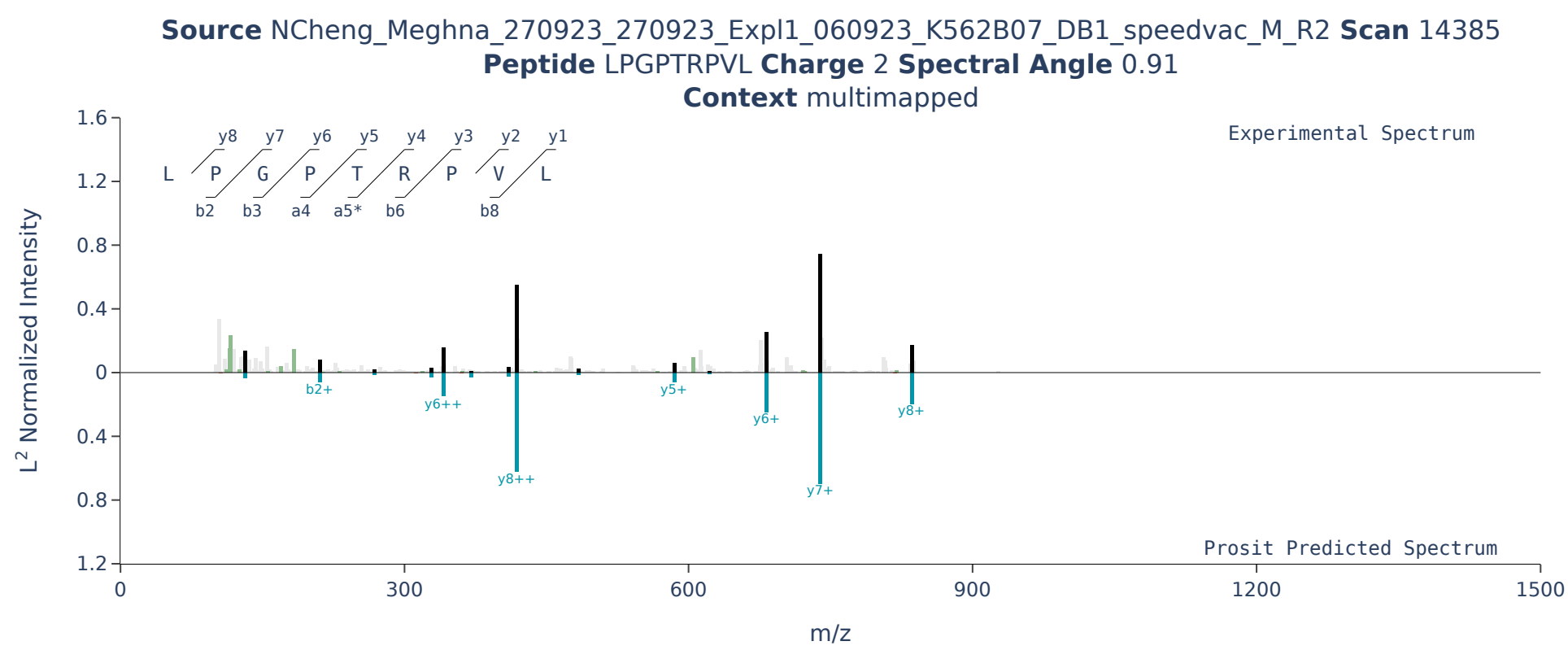
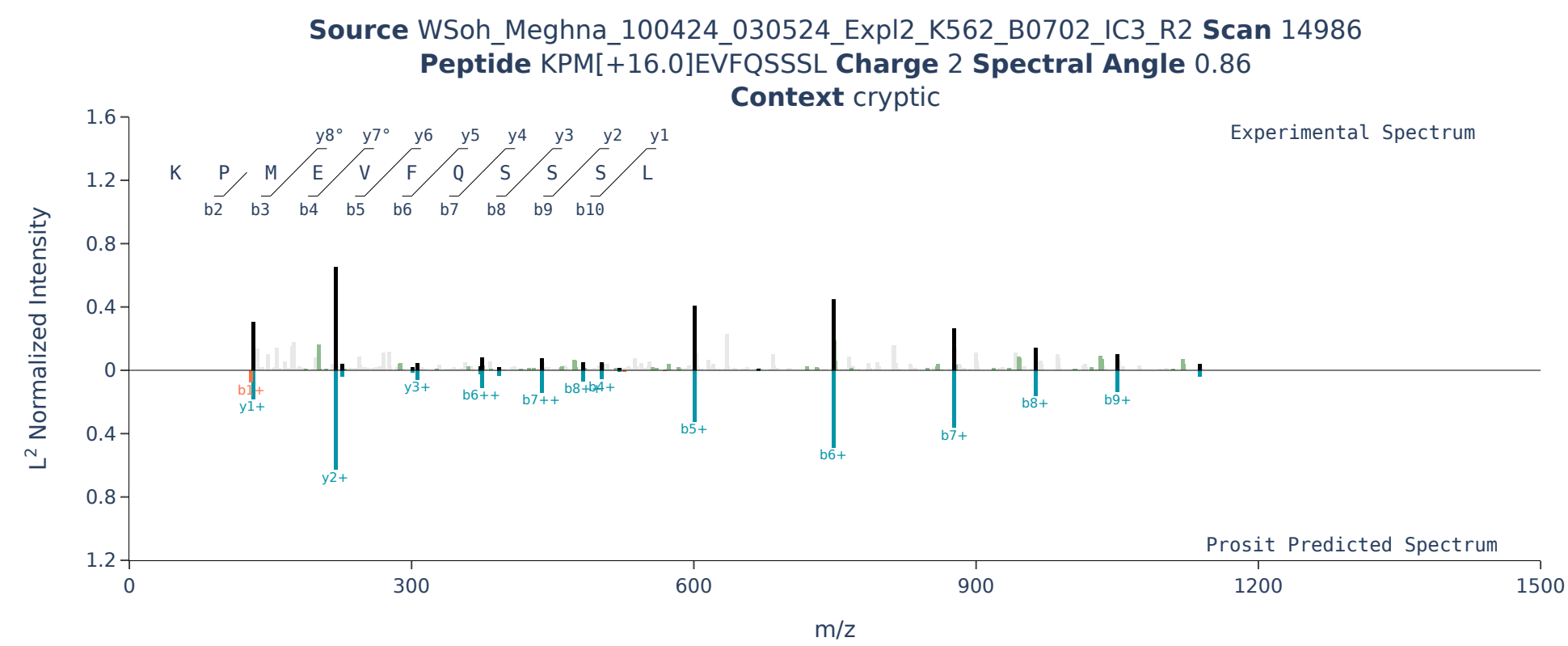
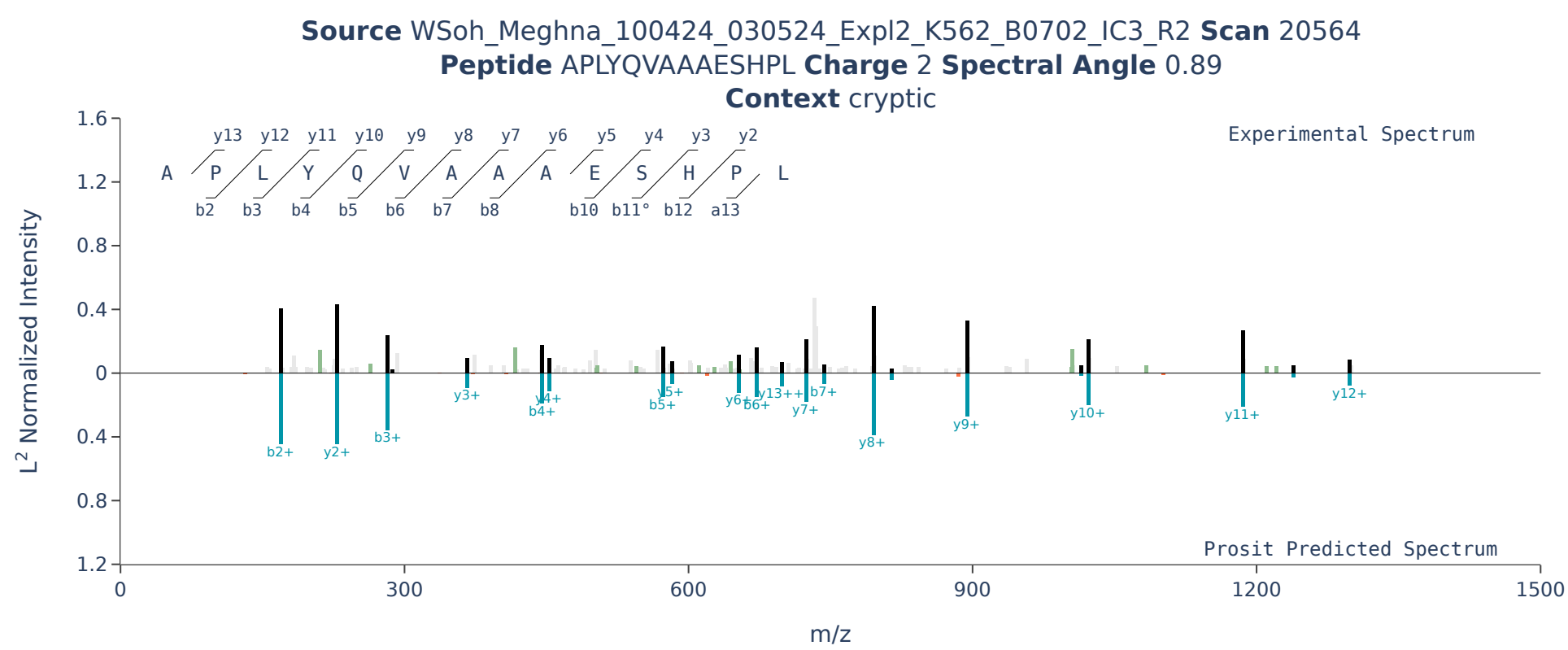
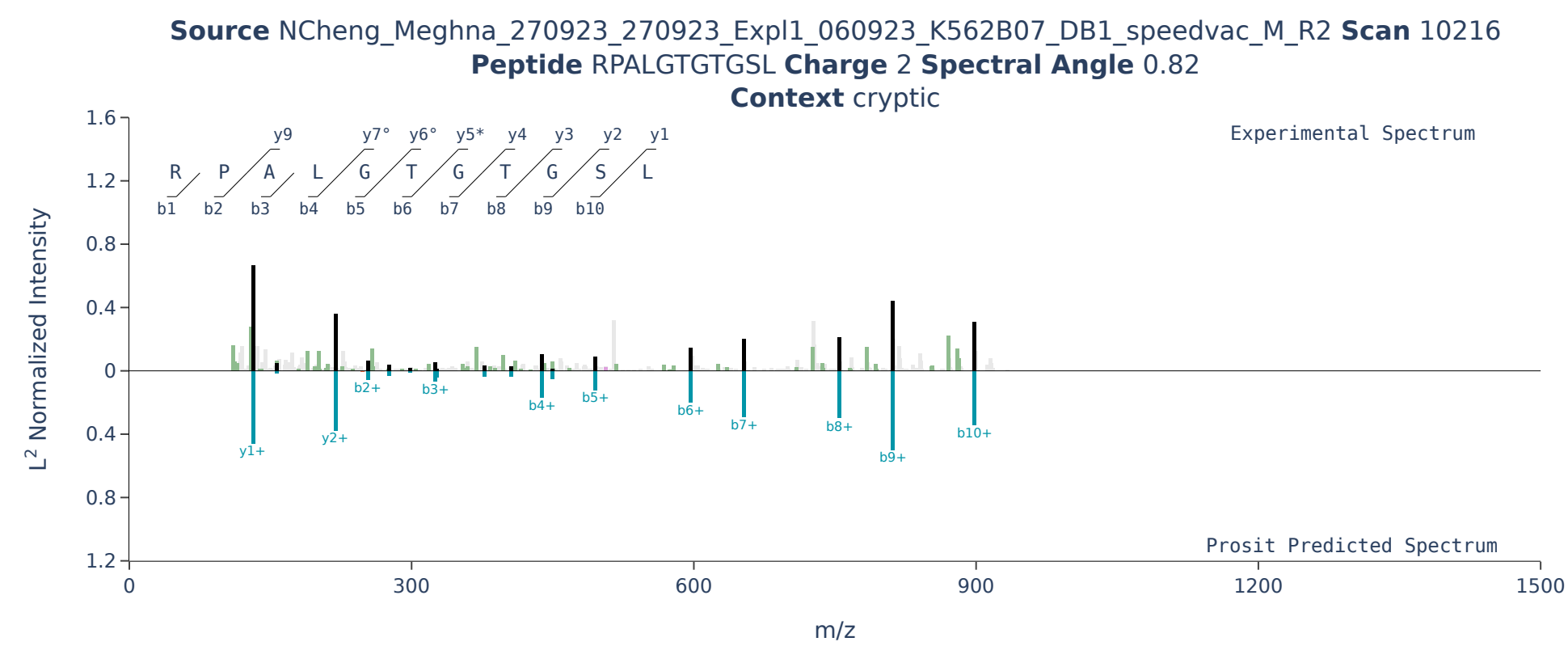
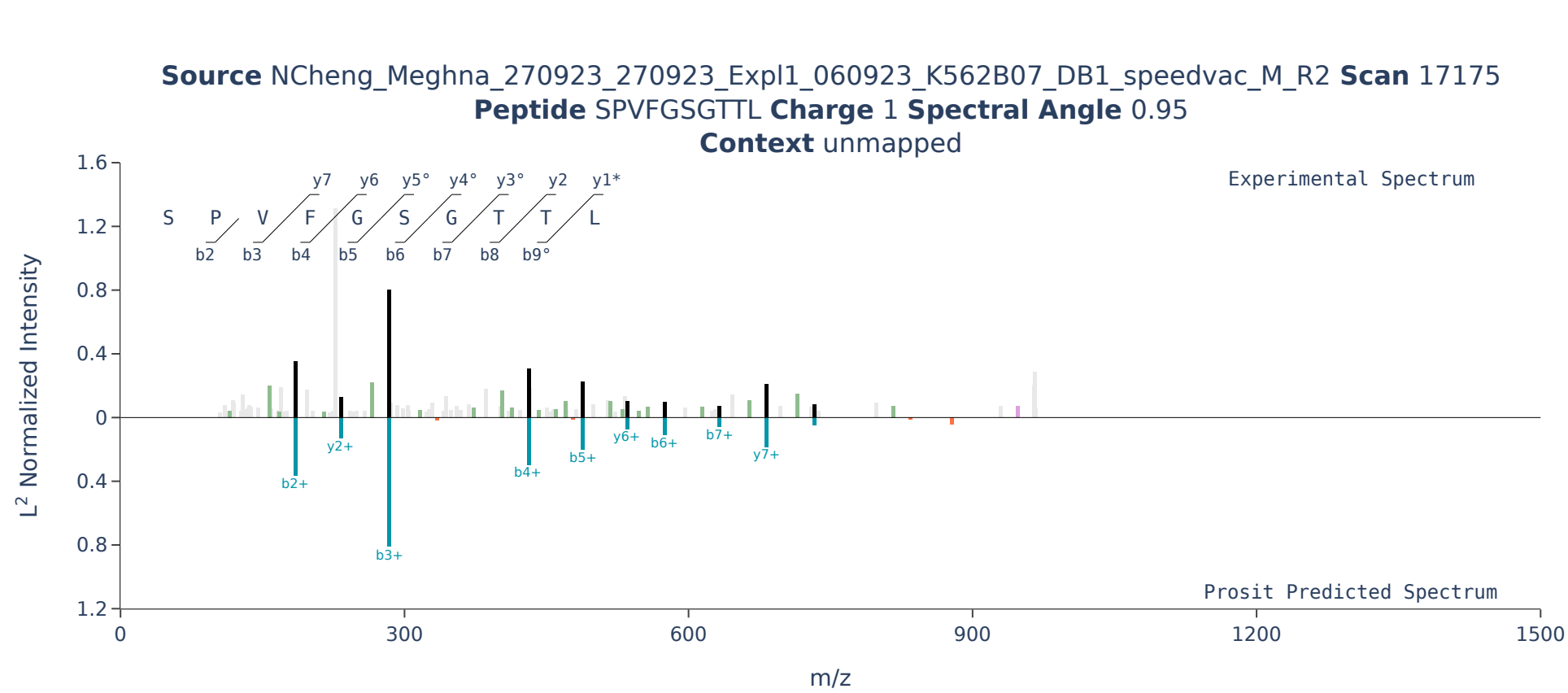
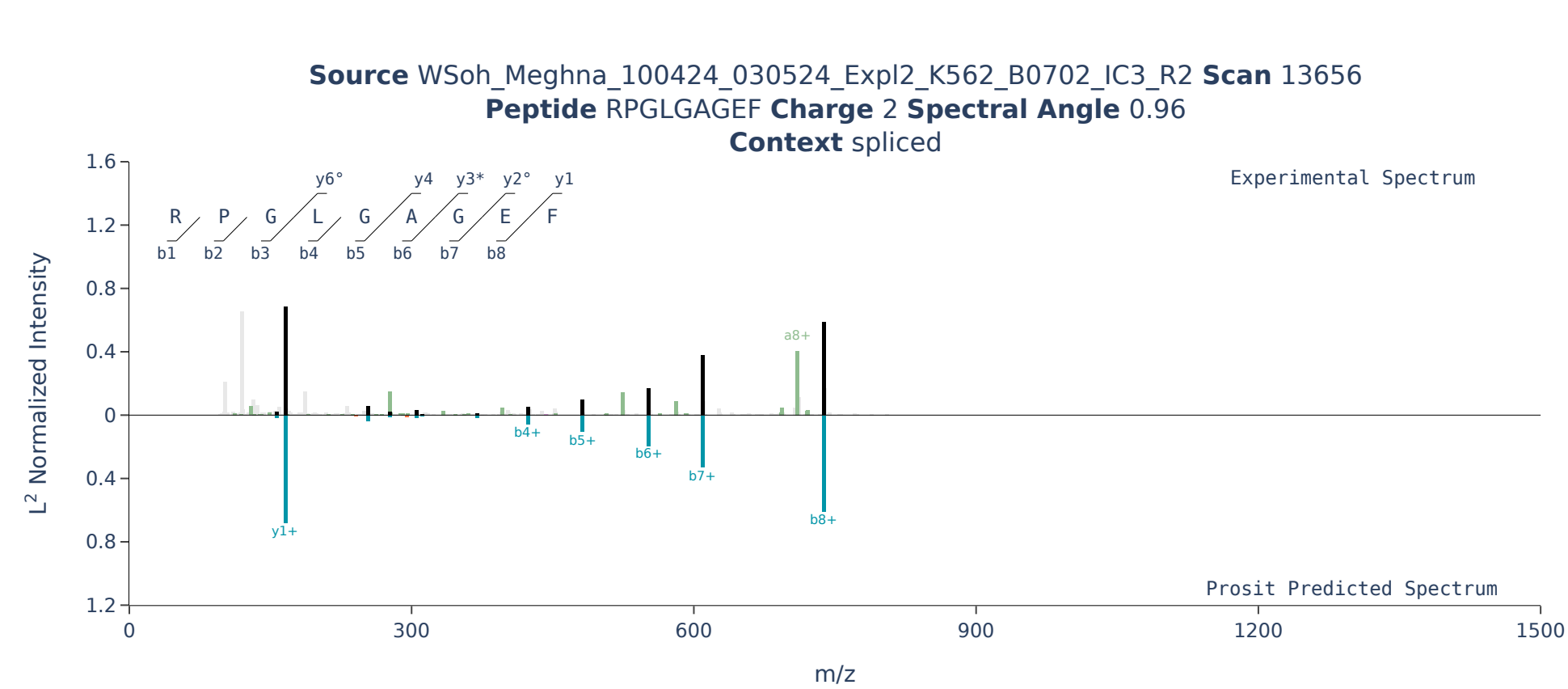




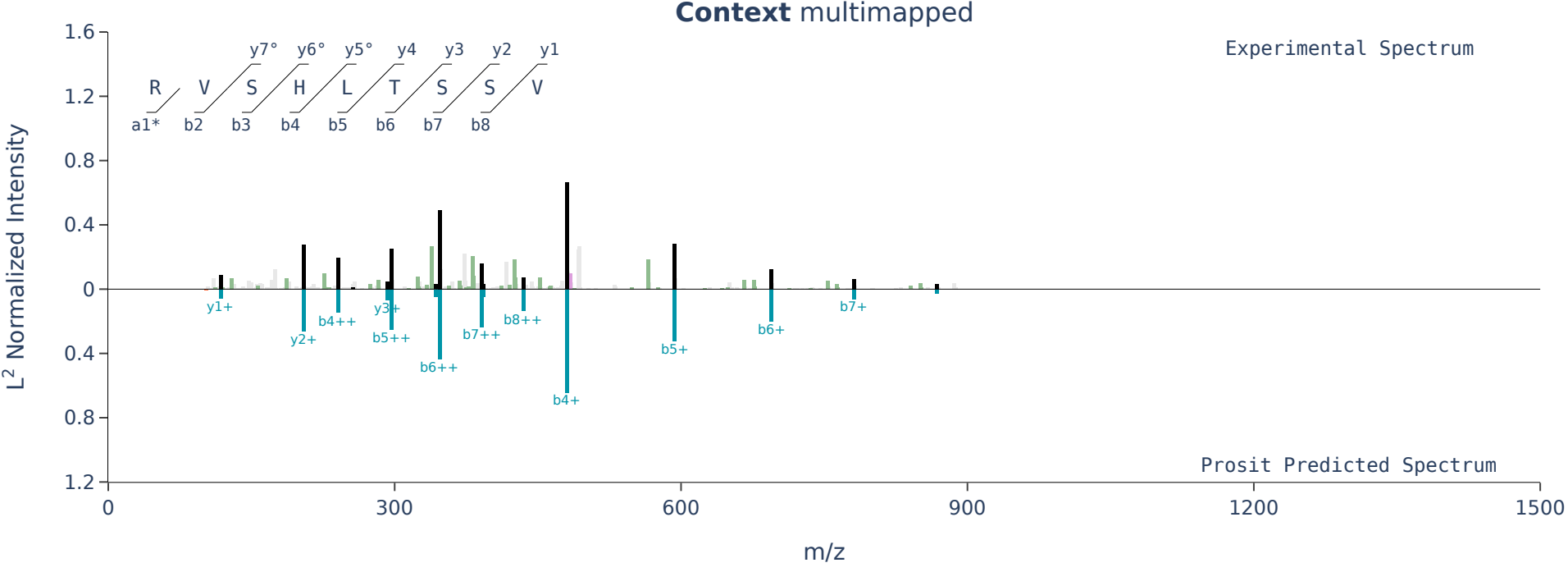




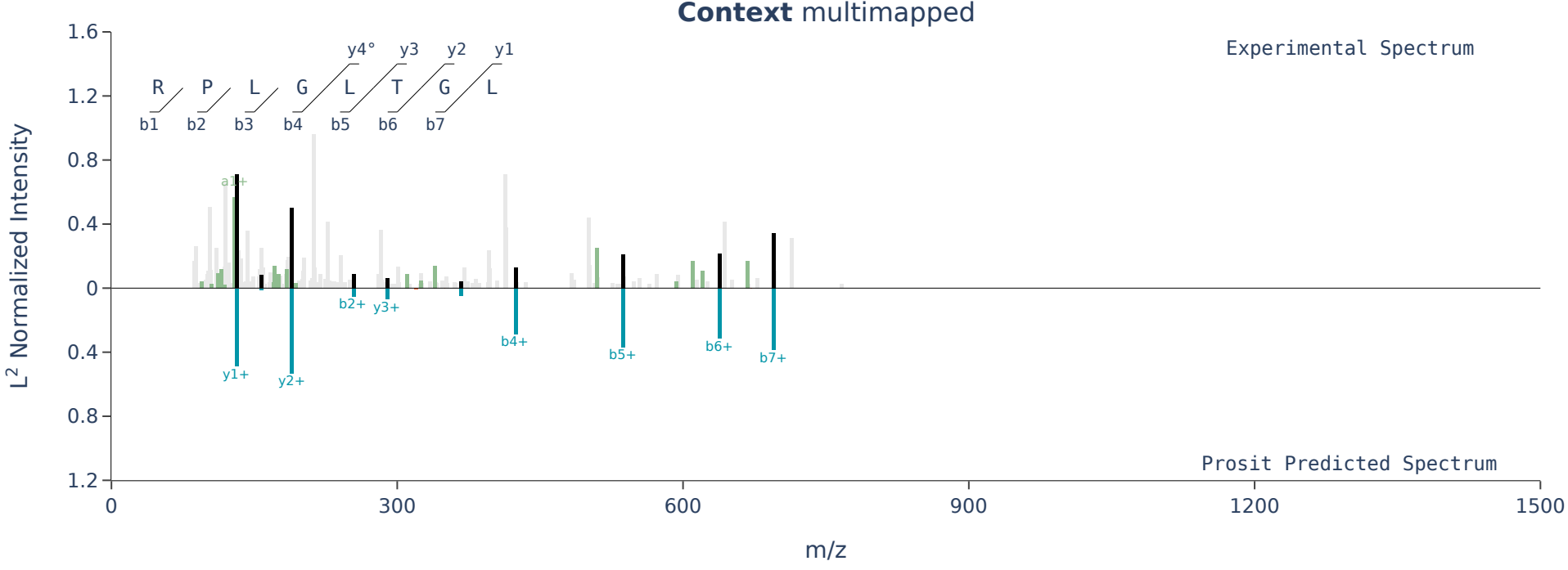




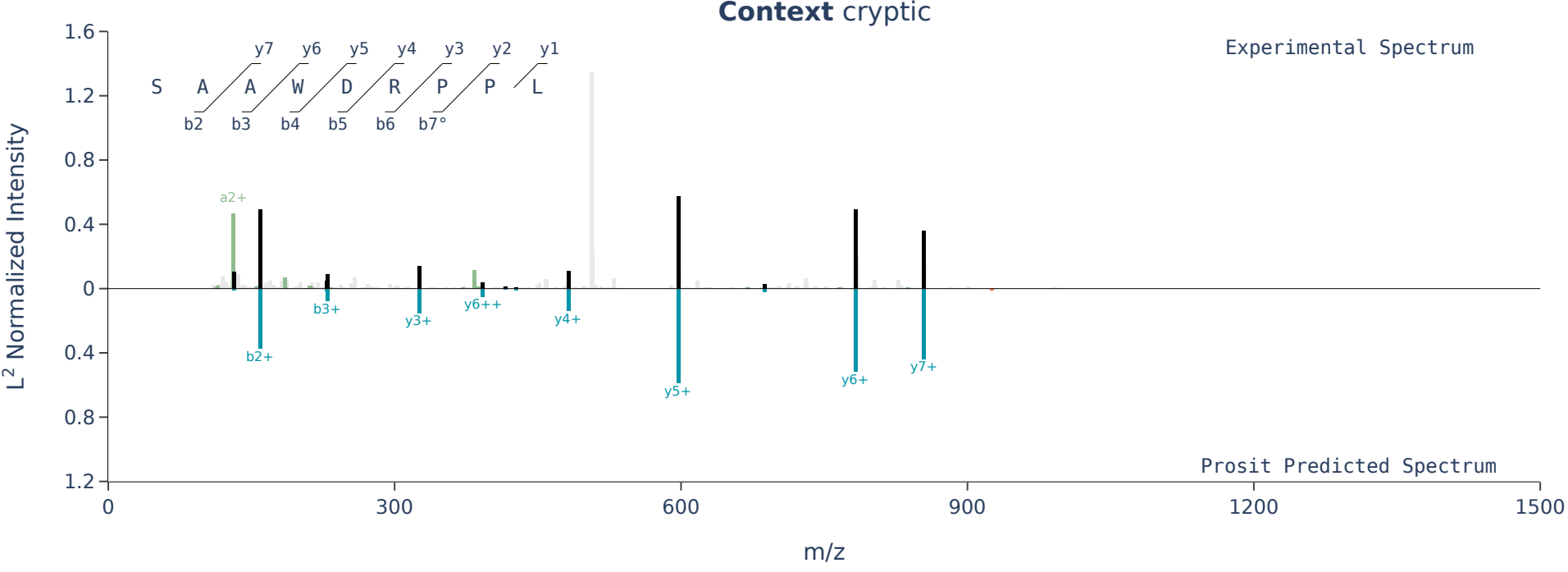
Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 6685
Peptide RVSHLTSSV Charge 2 Spectral Angle 0.9



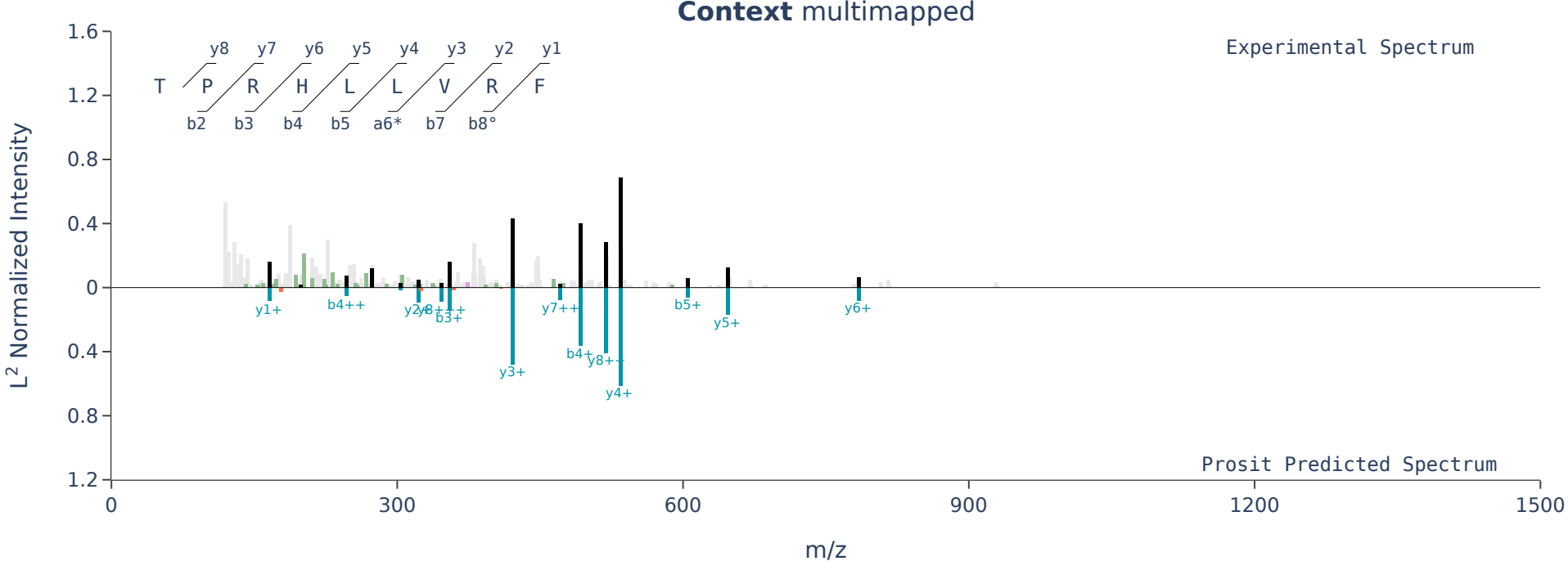
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Peptide RPLGLTGL Charge 2 Spectral Angle 0.78



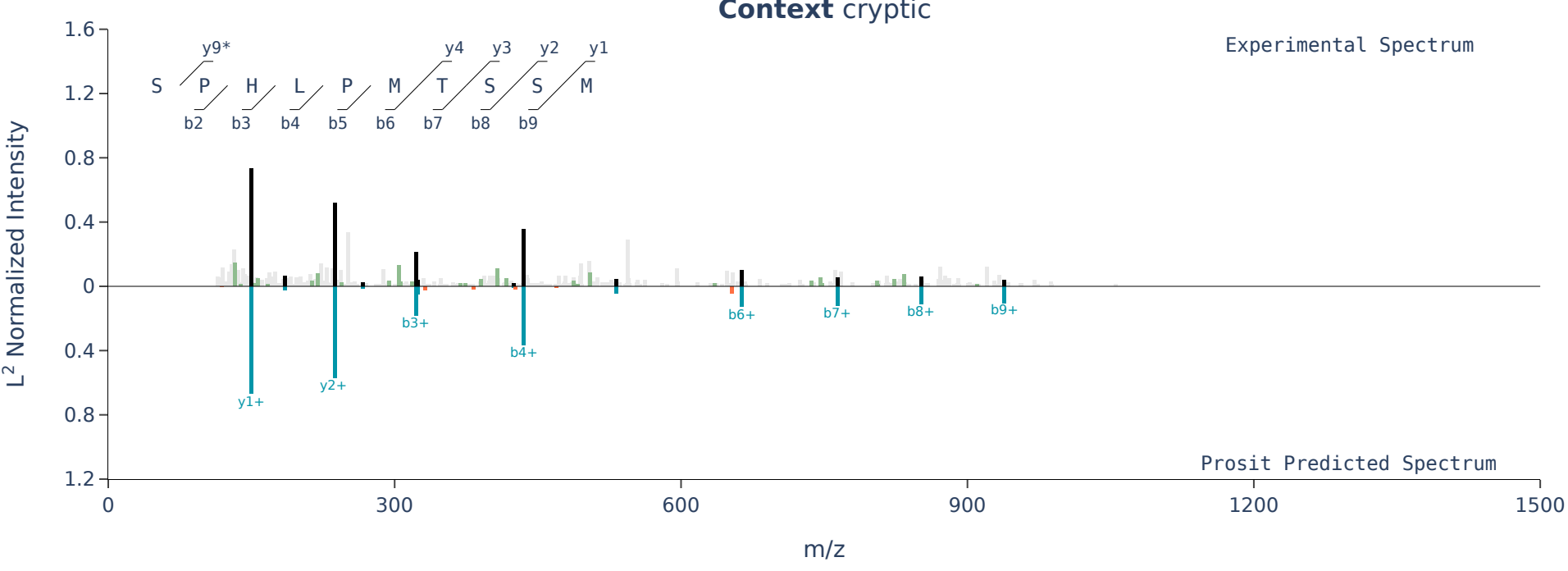
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Peptide SAAWDRPPL Charge 2 Spectral Angle 0.9



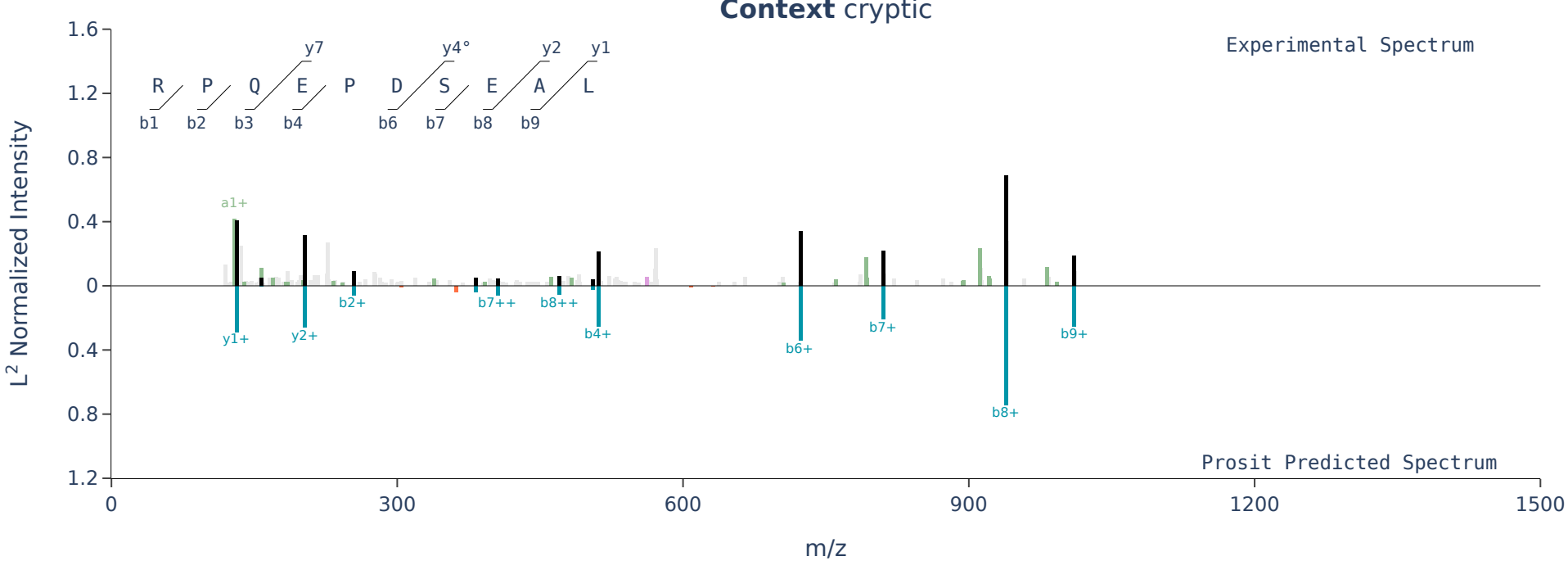
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Peptide TPRHLLVRF Charge 3 Spectral Angle 0.87



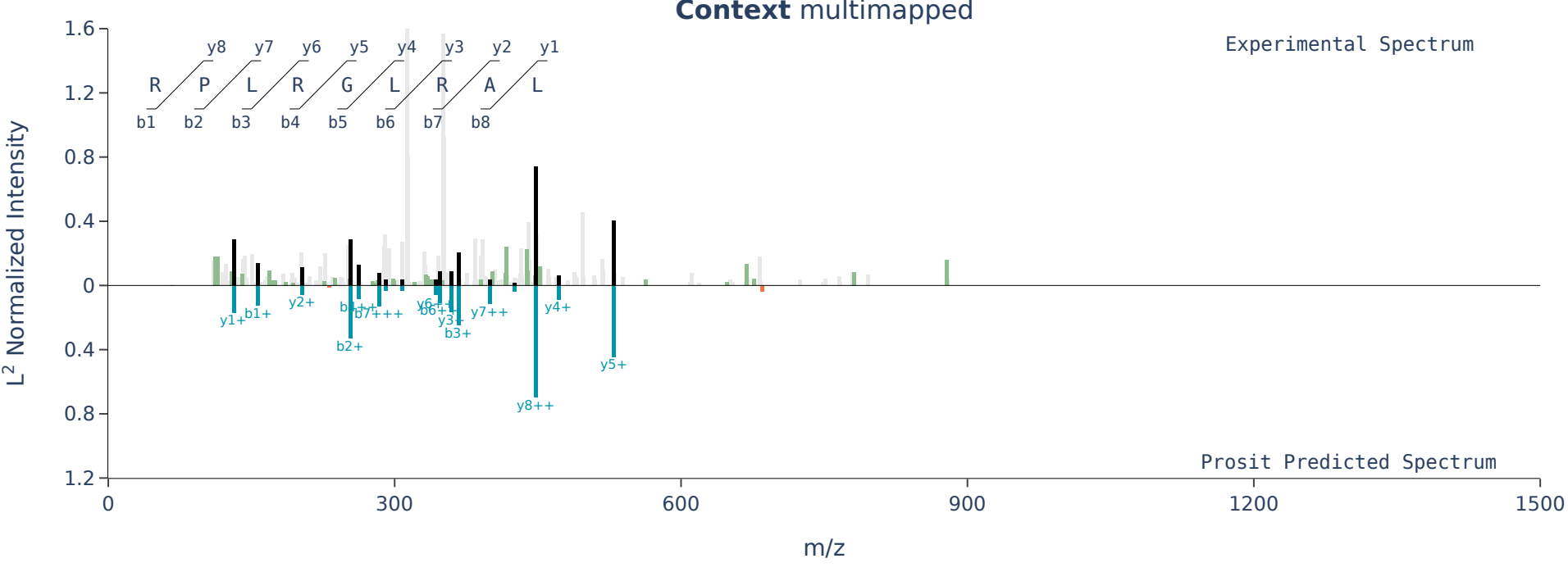
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Peptide SPHLPMTSSM Charge 2 Spectral Angle 0.9



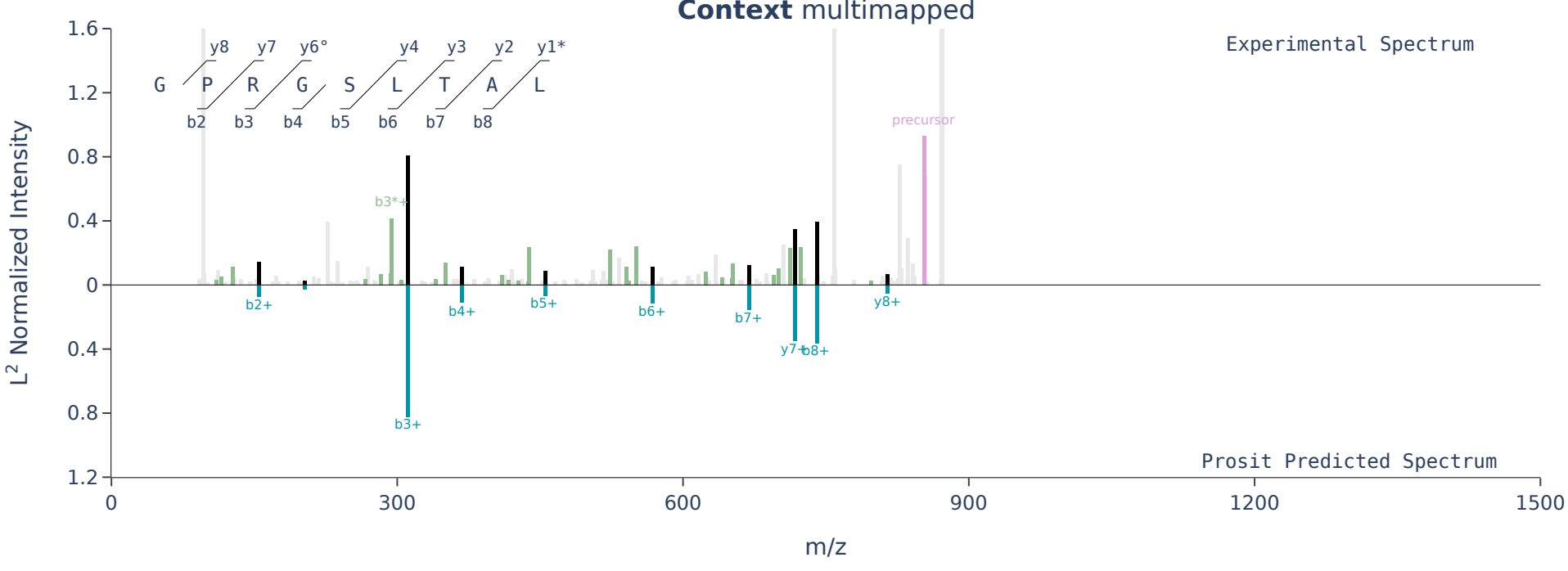
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Peptide RPQEPDSEAL Charge 2 Spectral Angle 0.89



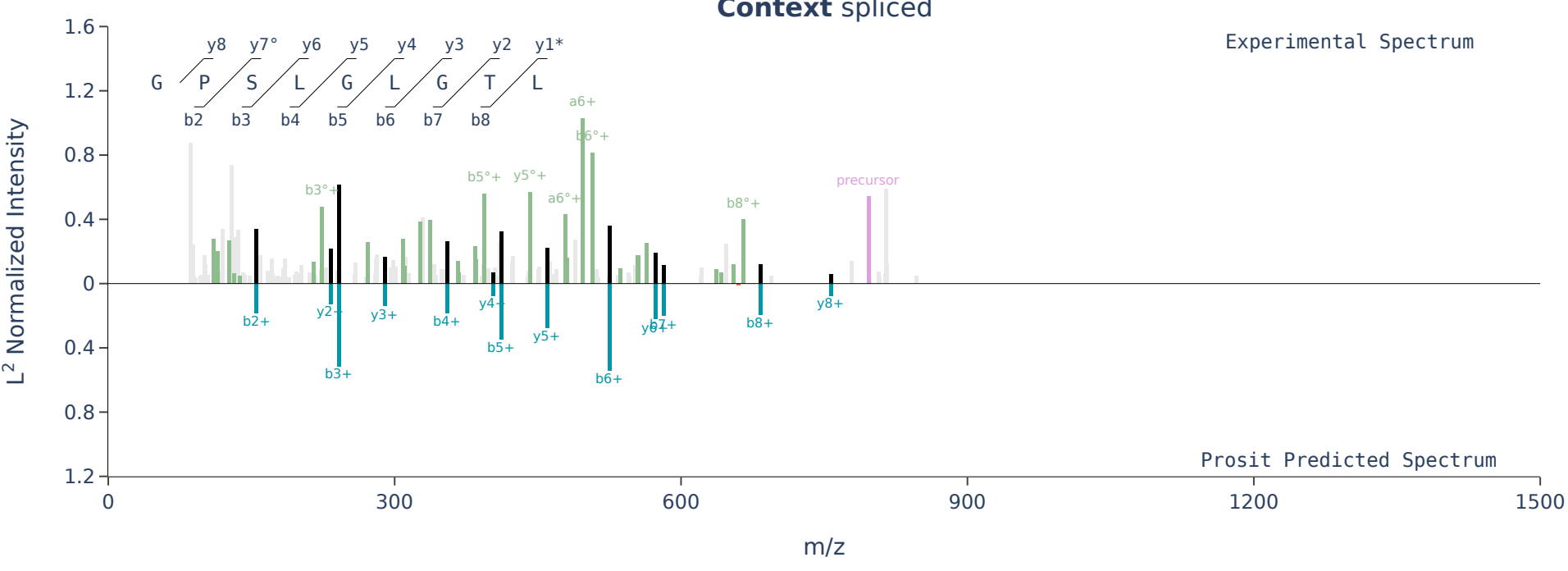
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Peptide RPLRLRAL Charge 3 Spectral Angle 0.86



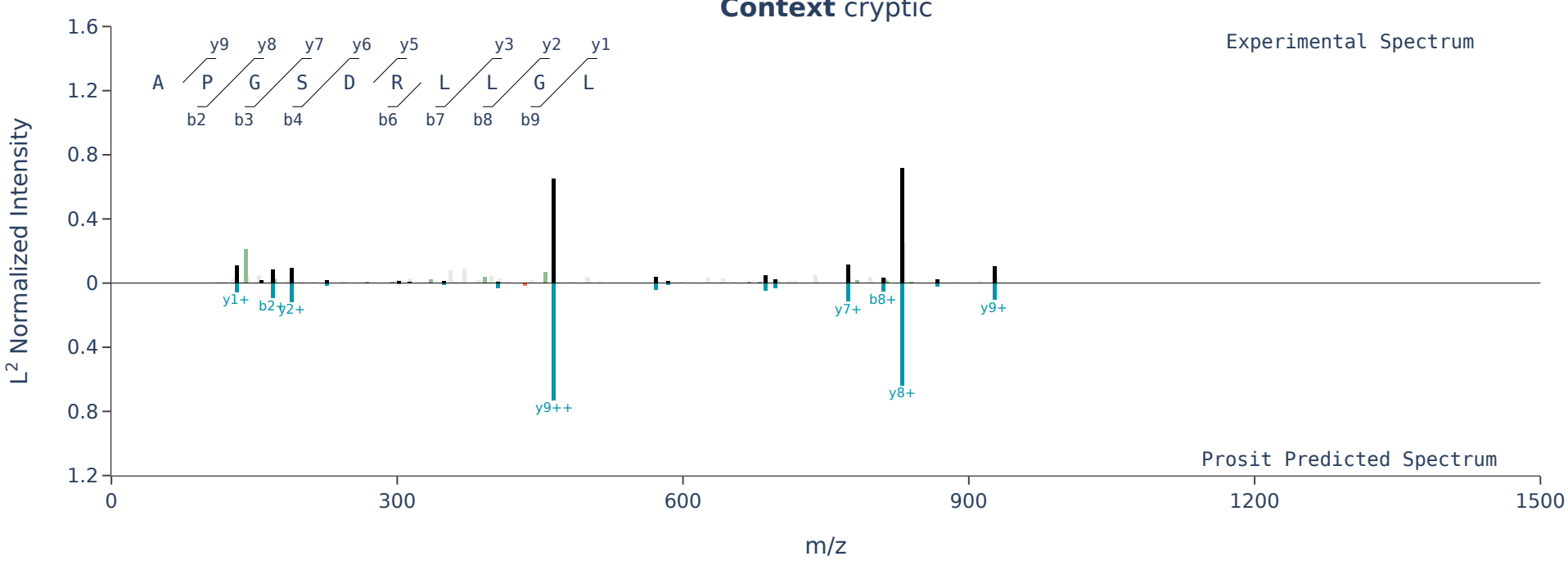
Source NCheng_Meghna_270923_270923_ExpI1_060923_K562B07_DB1_speedvac_M_R1 Scan 10690
Peptide GPRGSLTA Charge 1 Spectral Angle 0.94

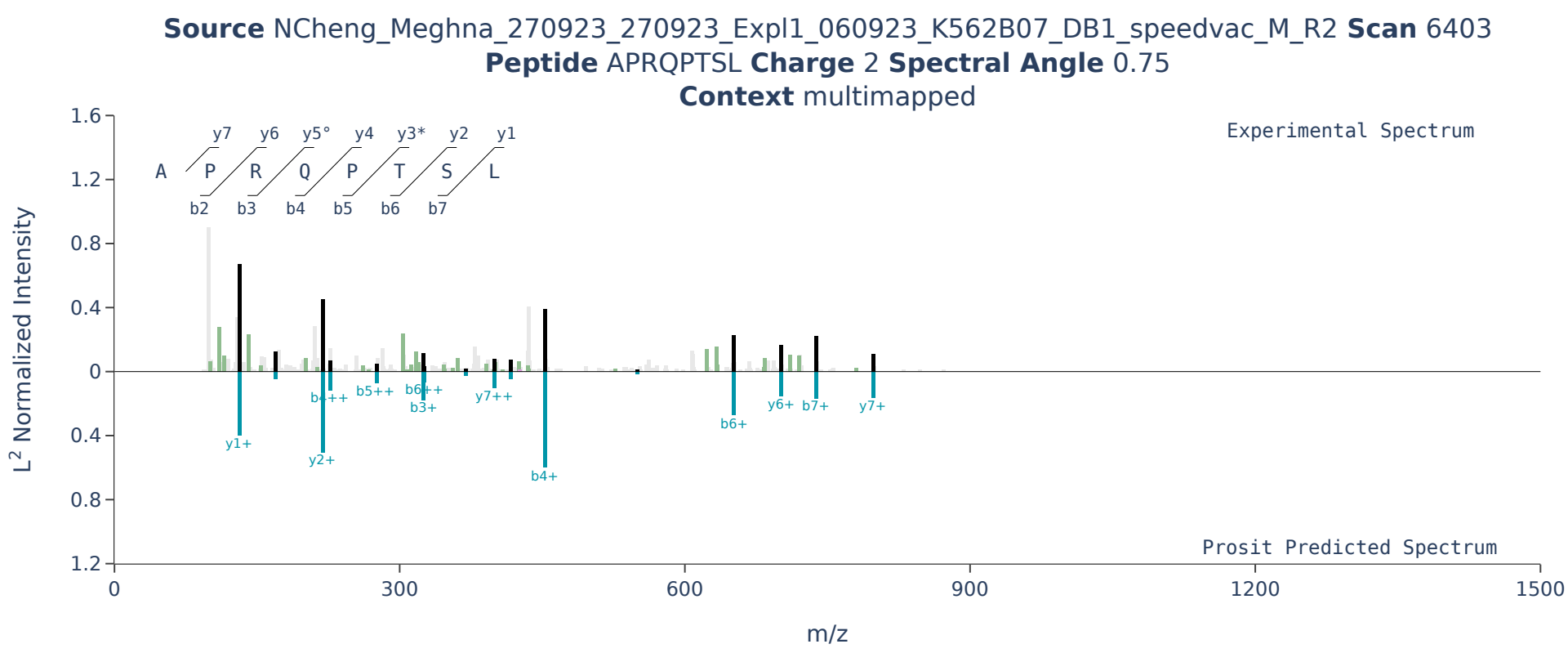
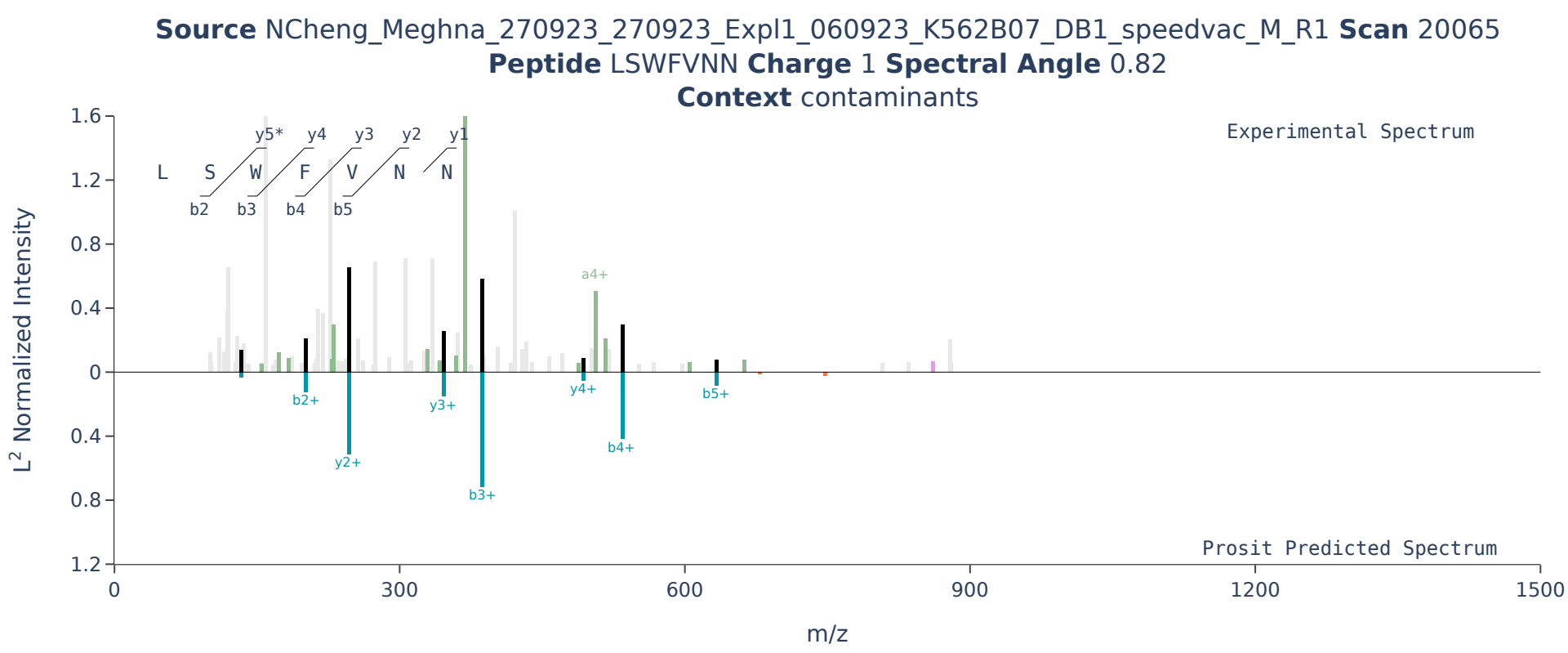
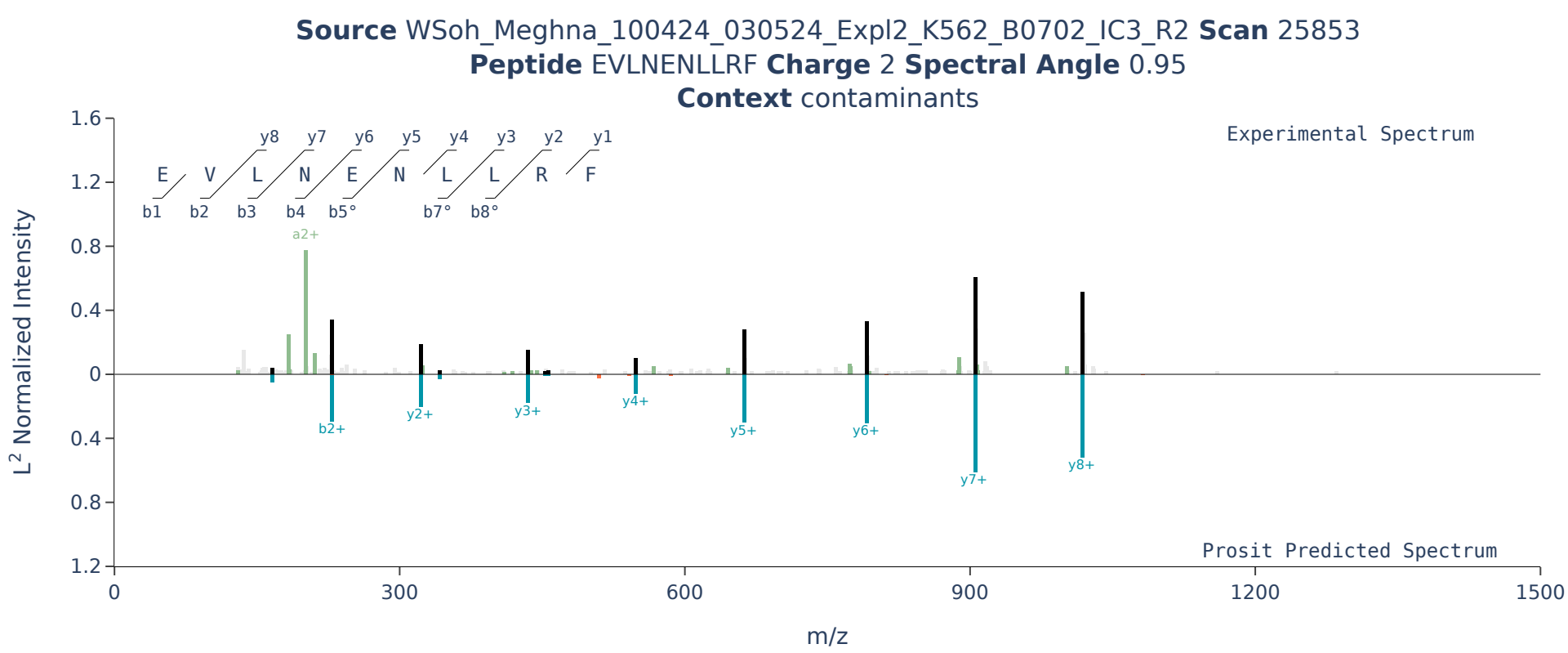
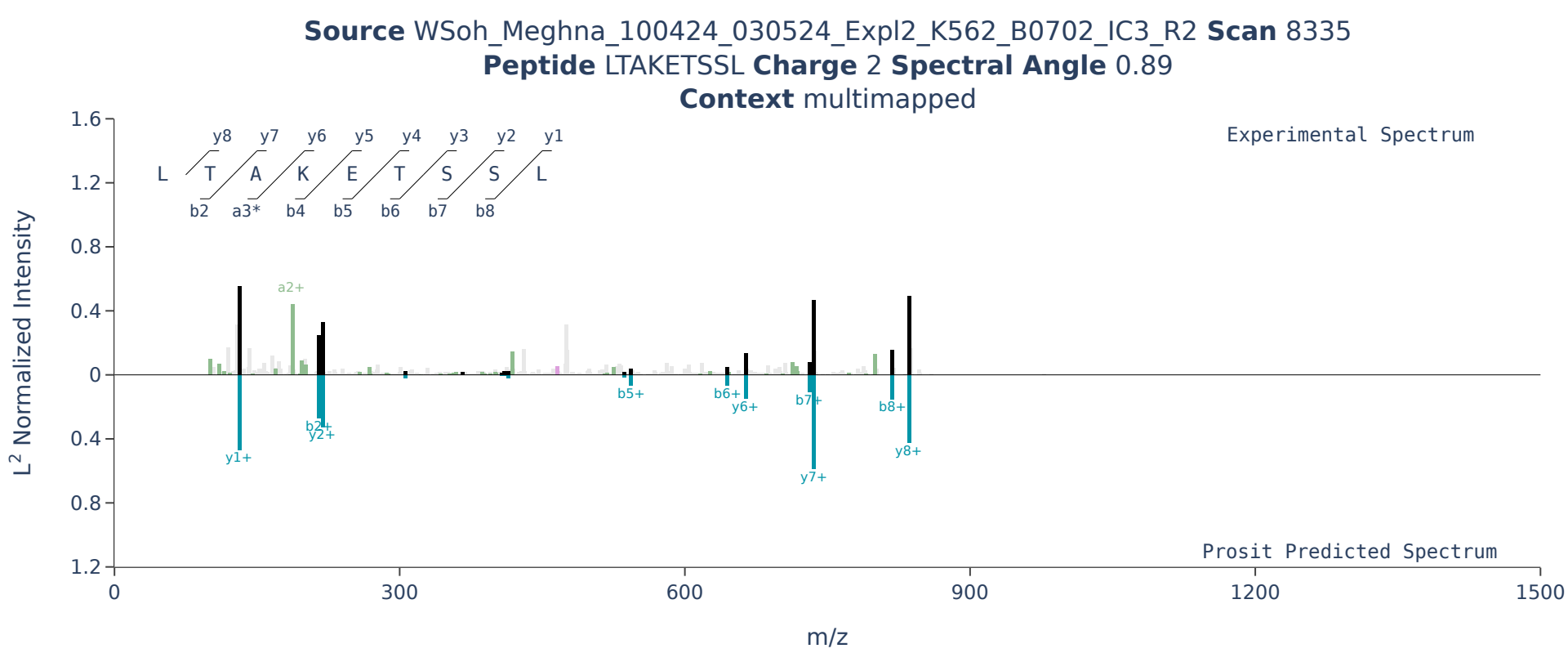
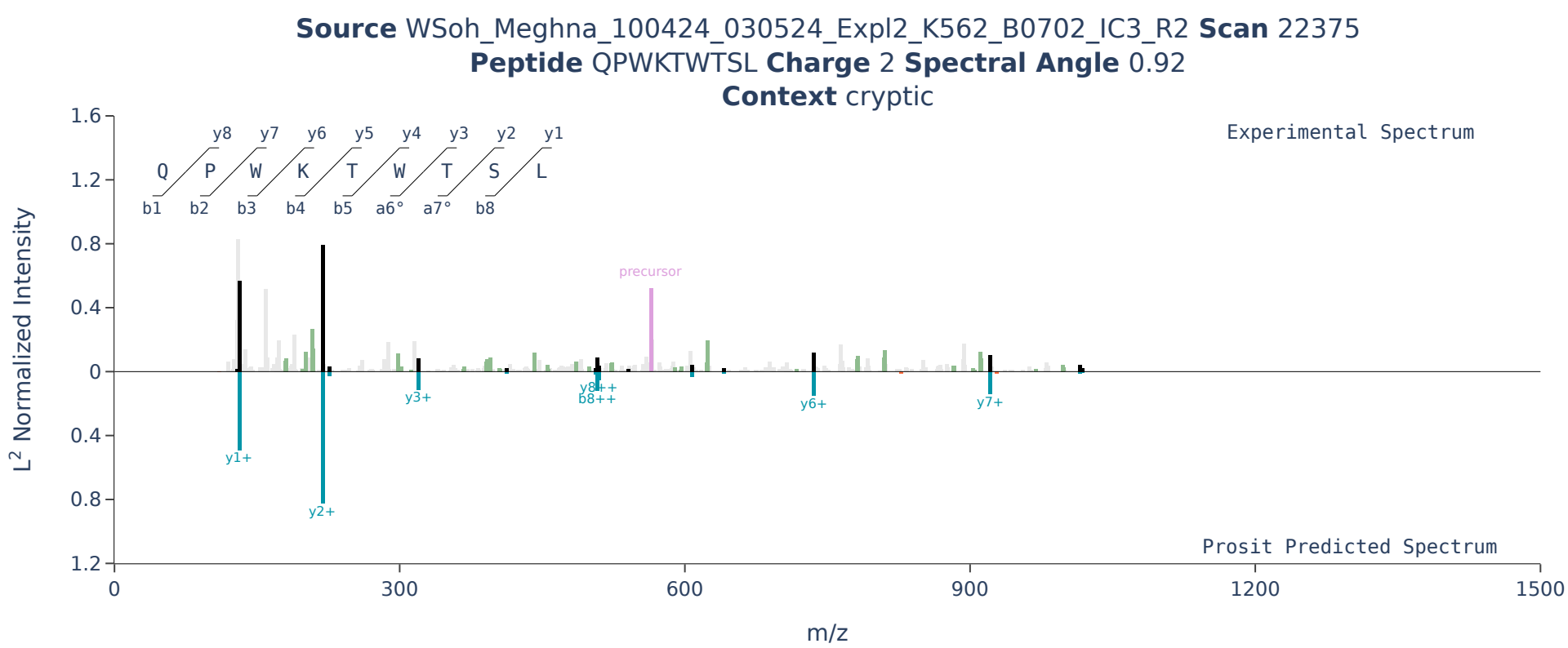


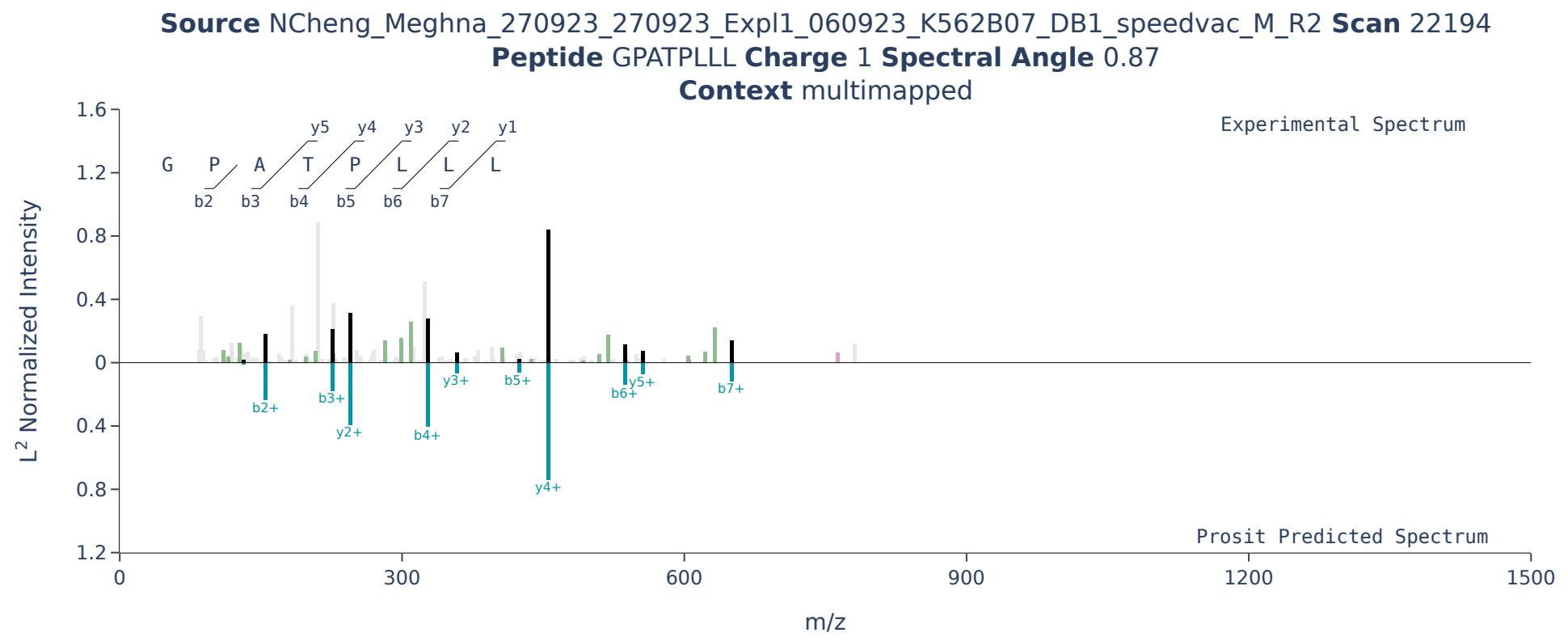
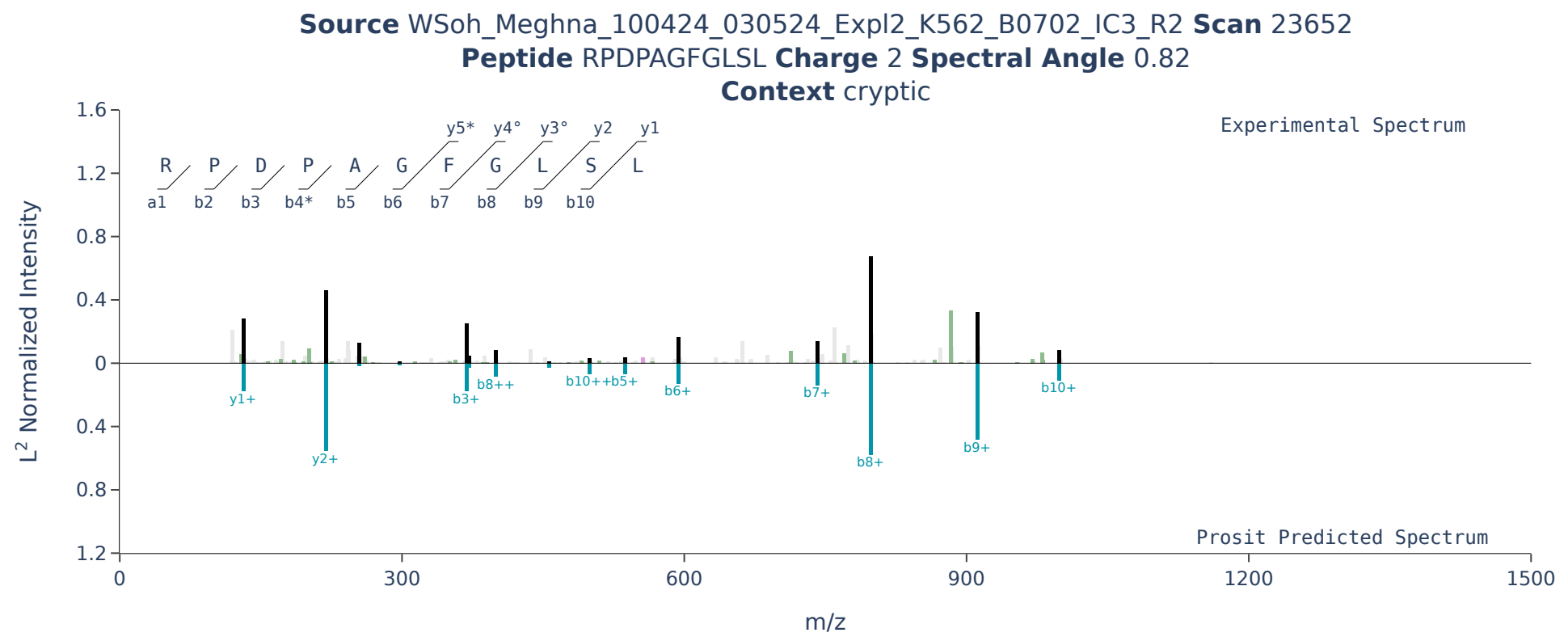
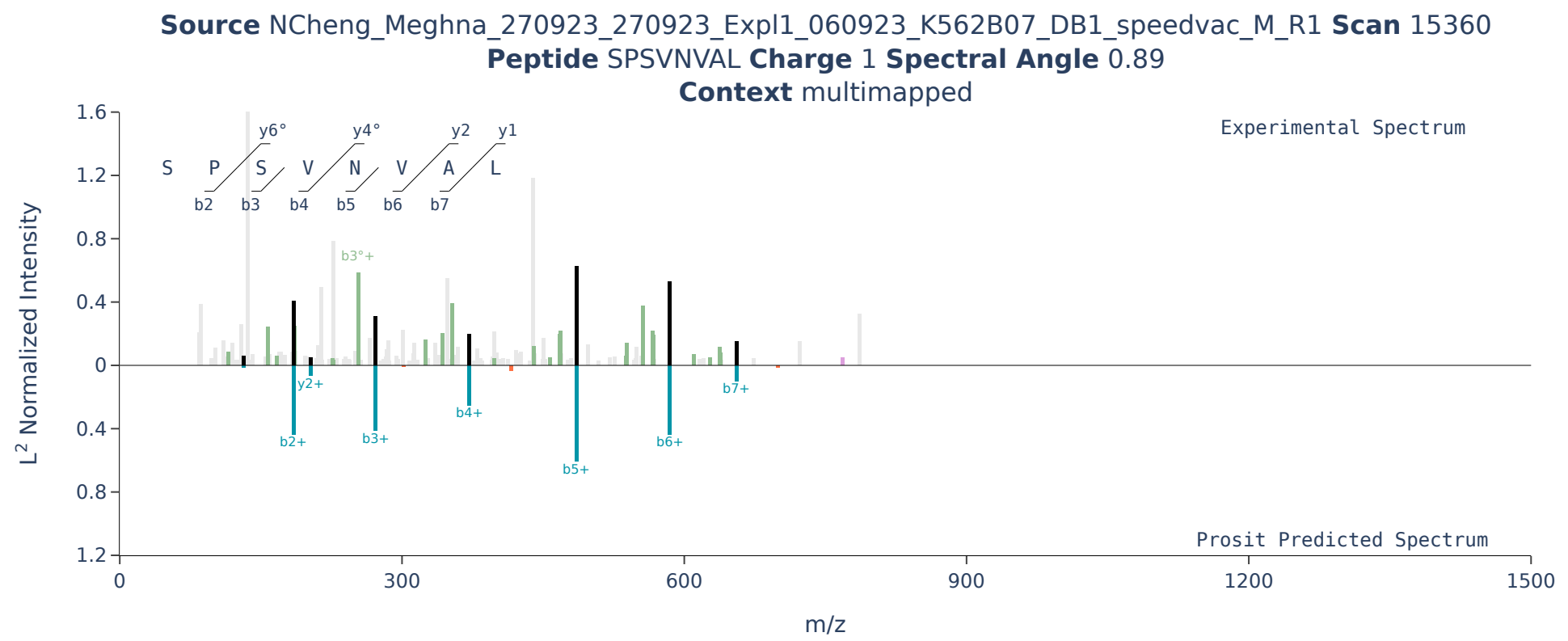
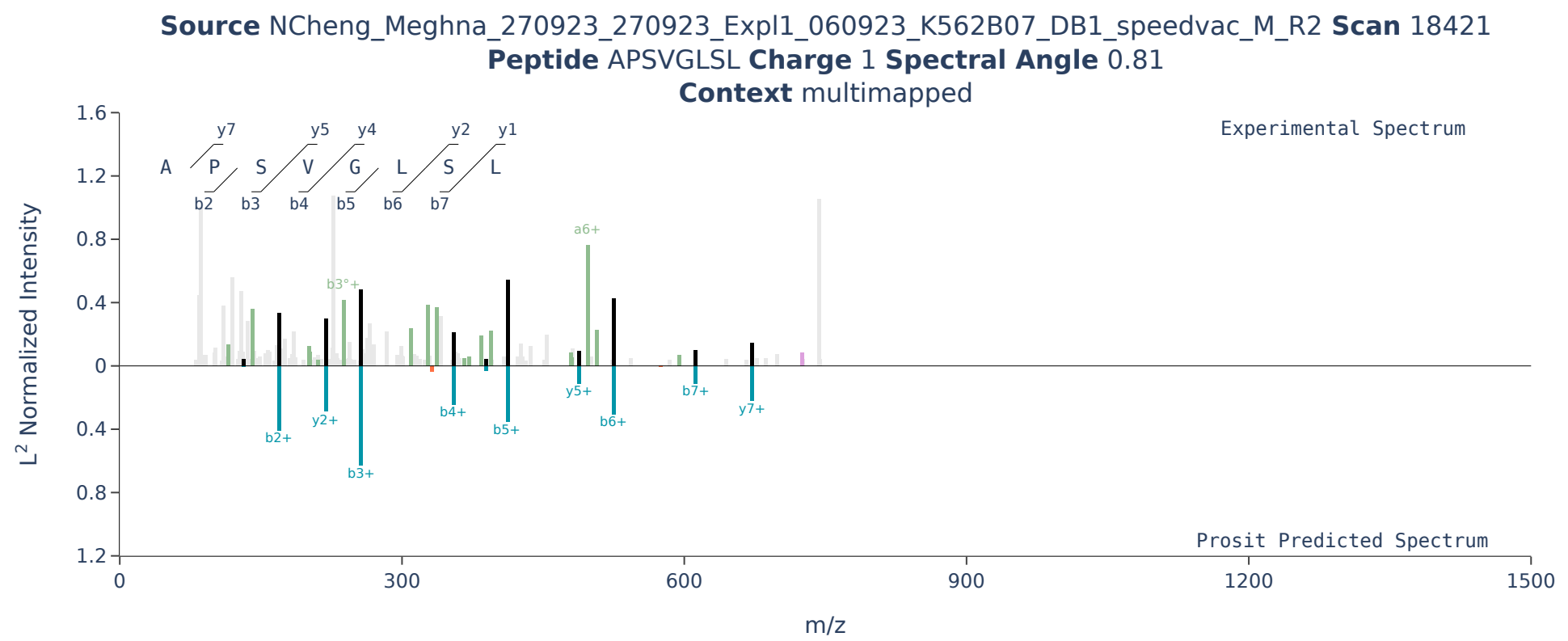
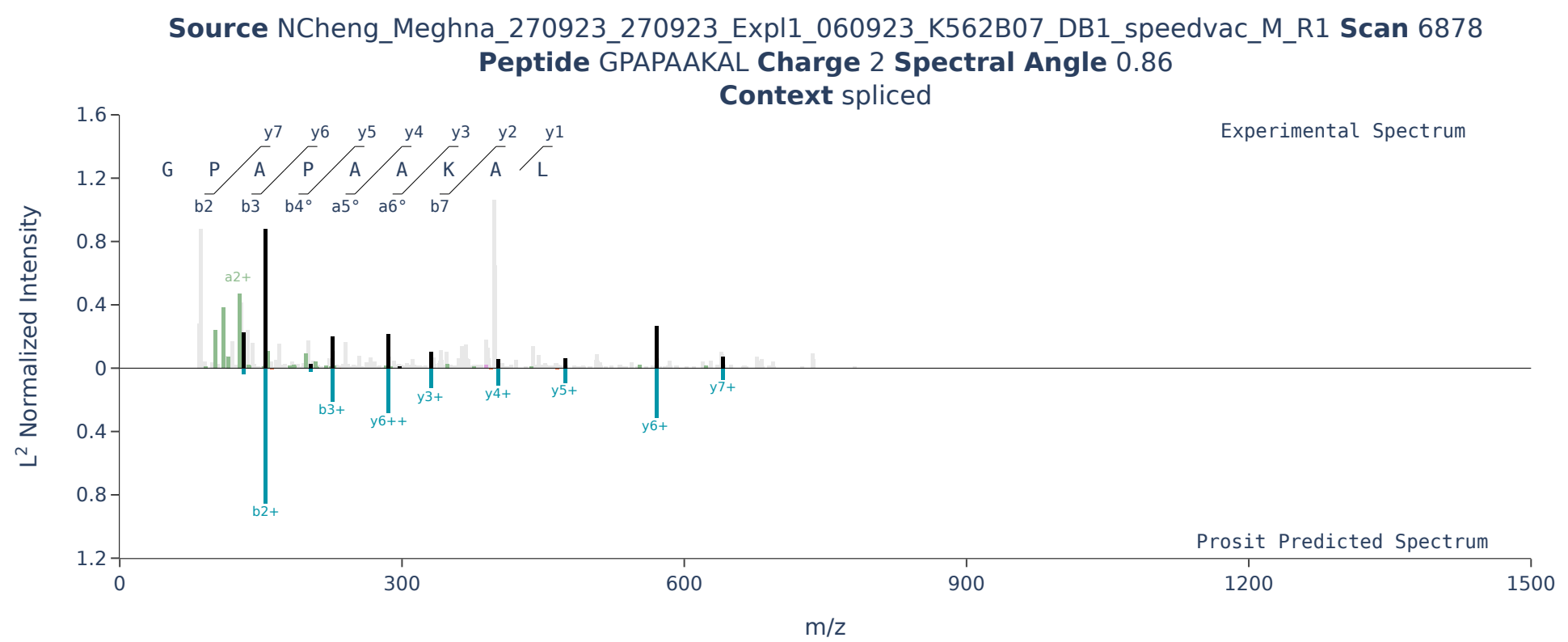
Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 24070
Peptide GPSLGLGTL Charge 1 Spectral Angle 0.8

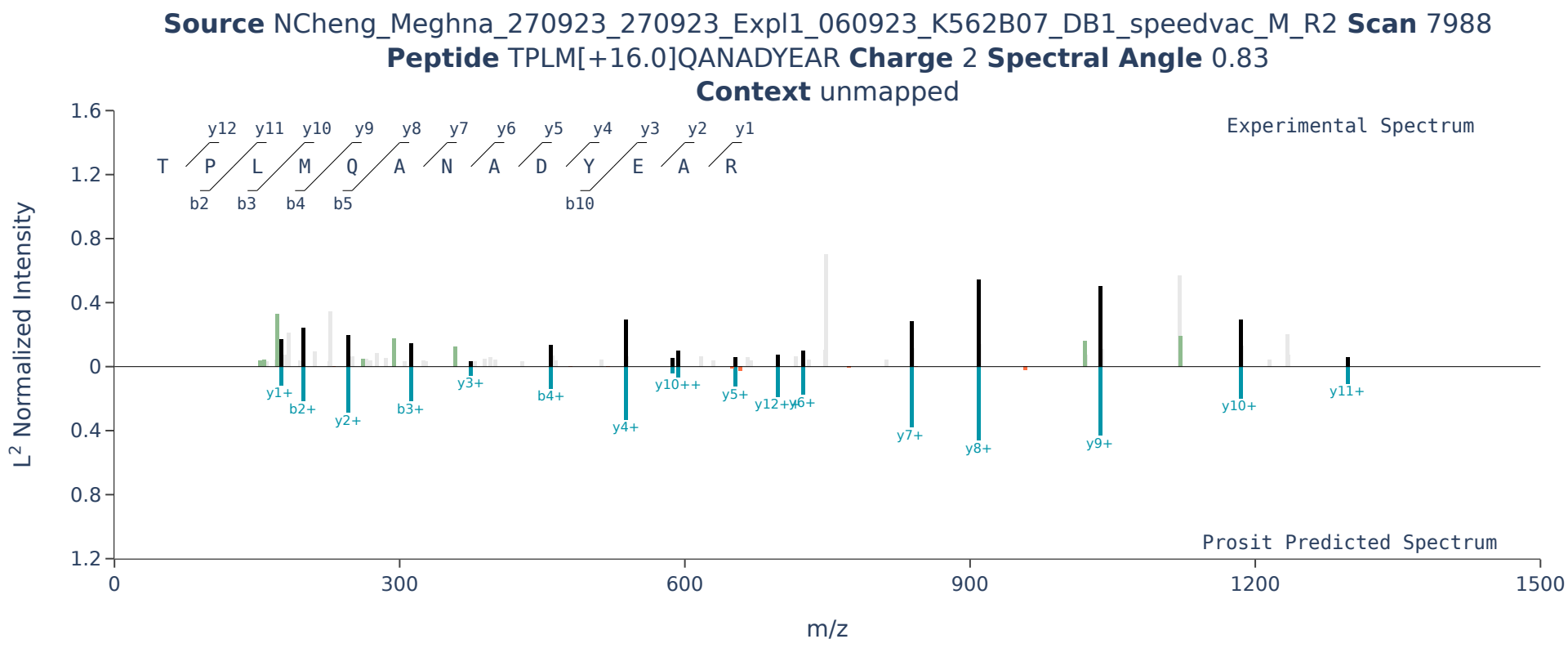
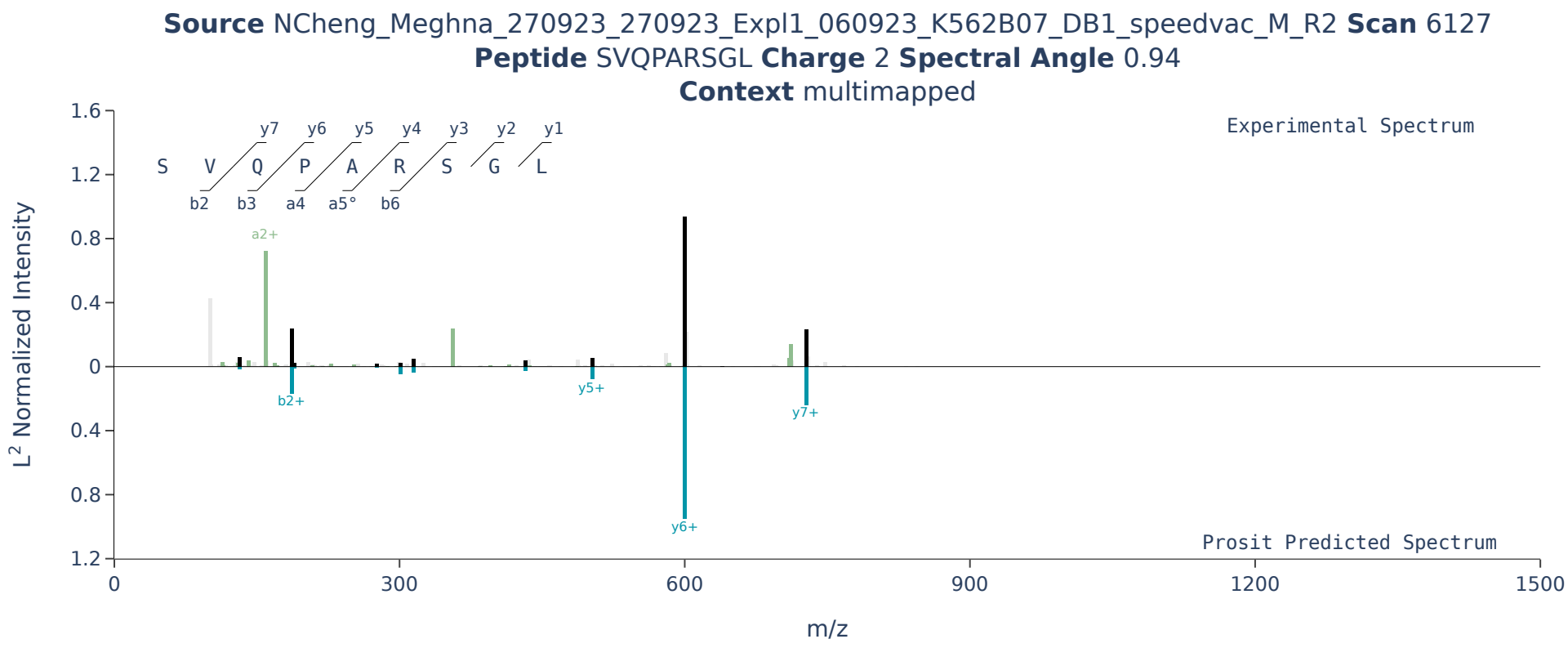
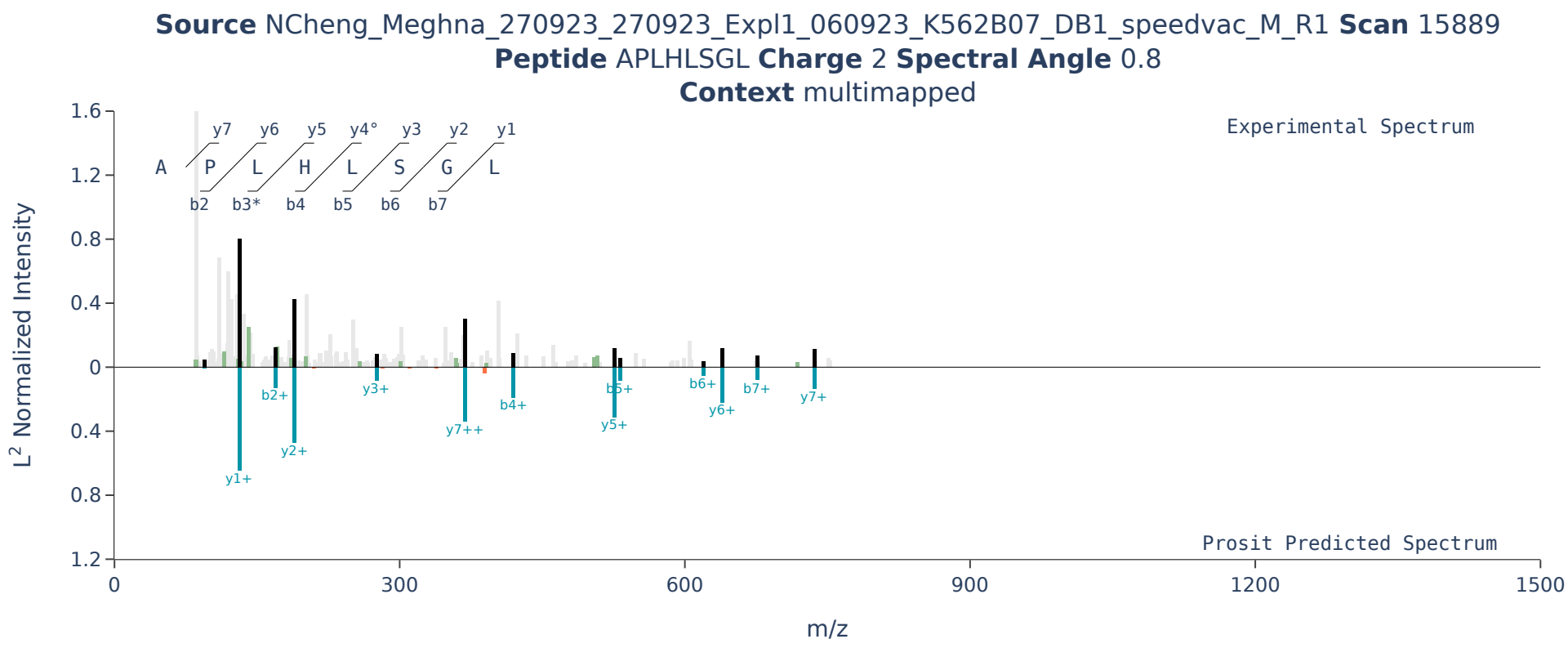
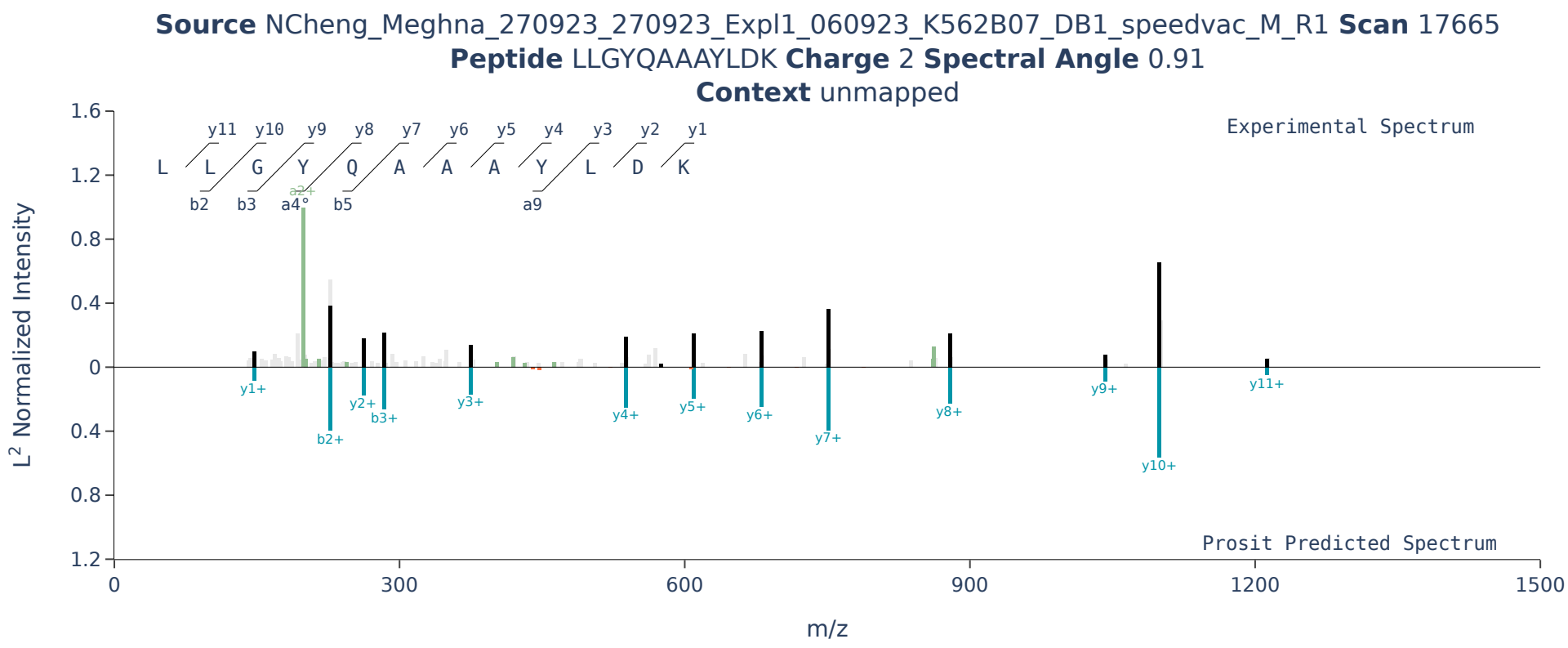
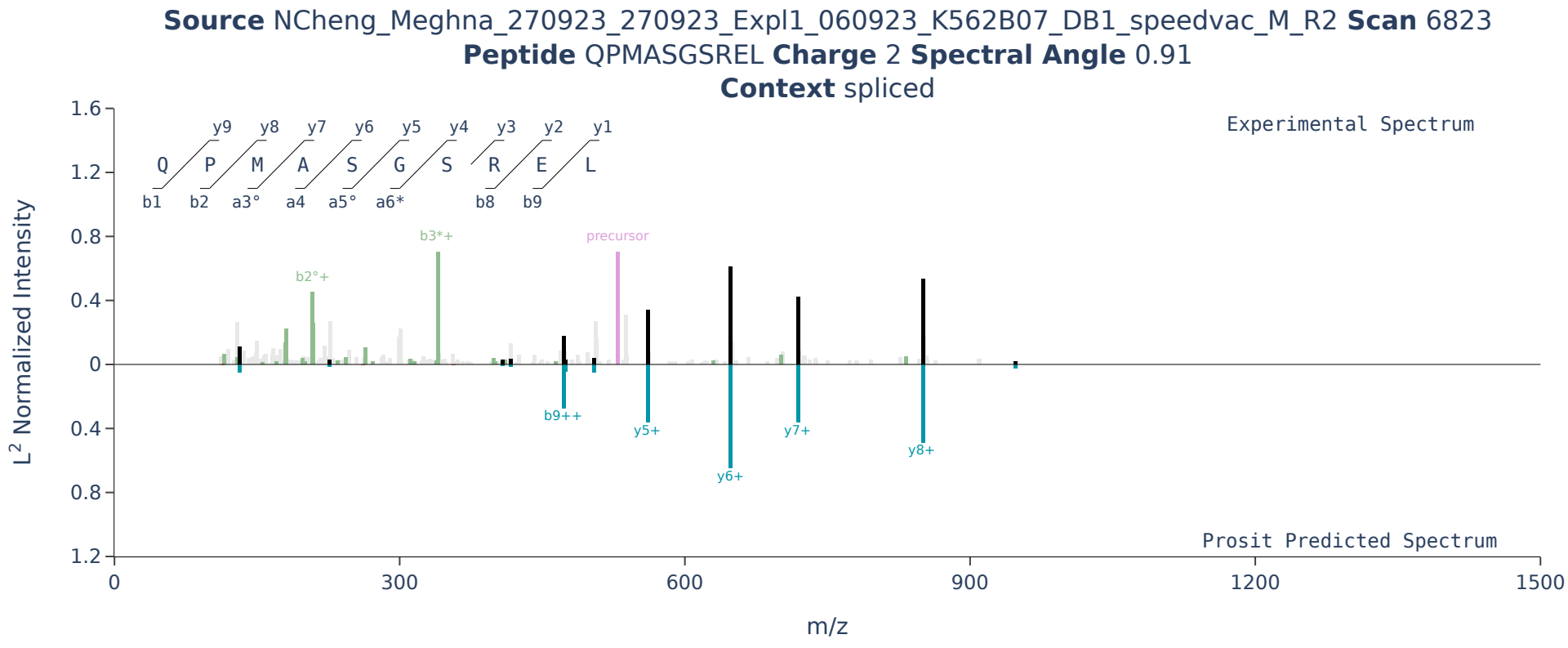
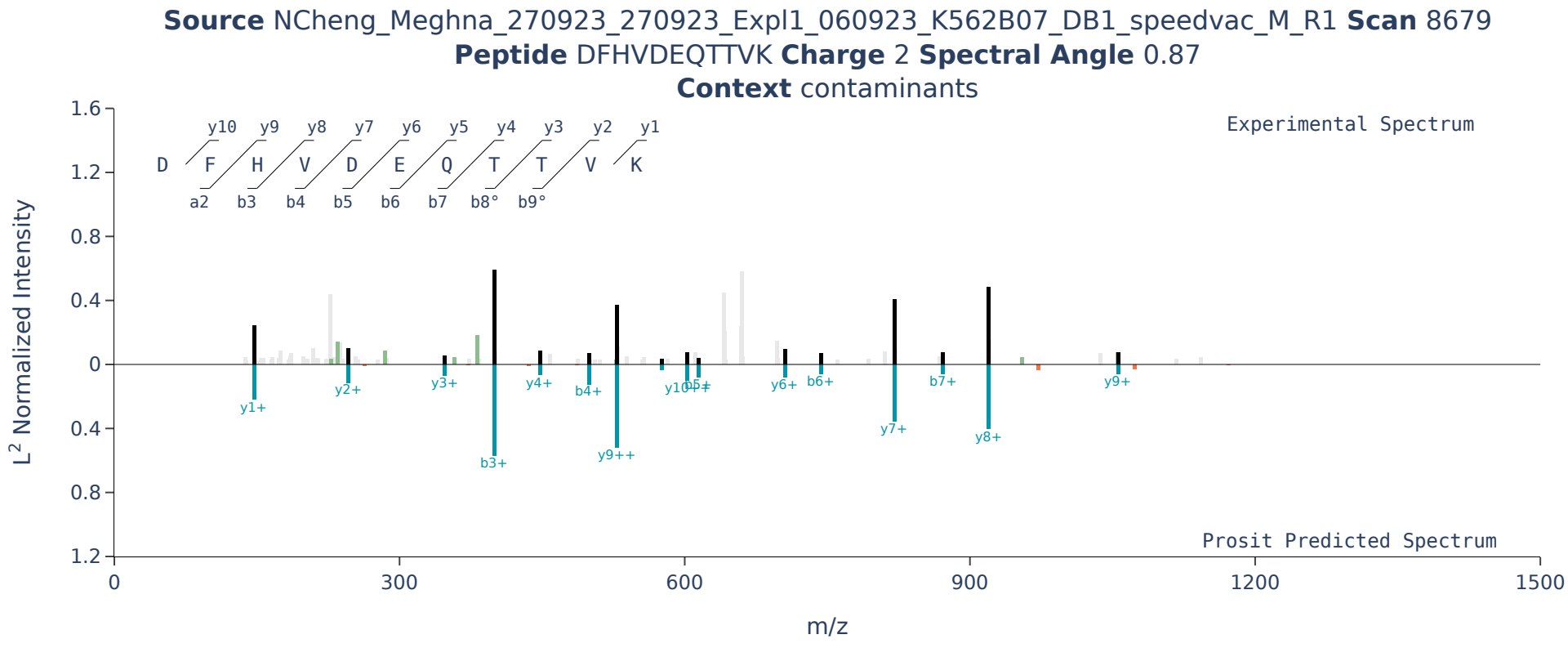
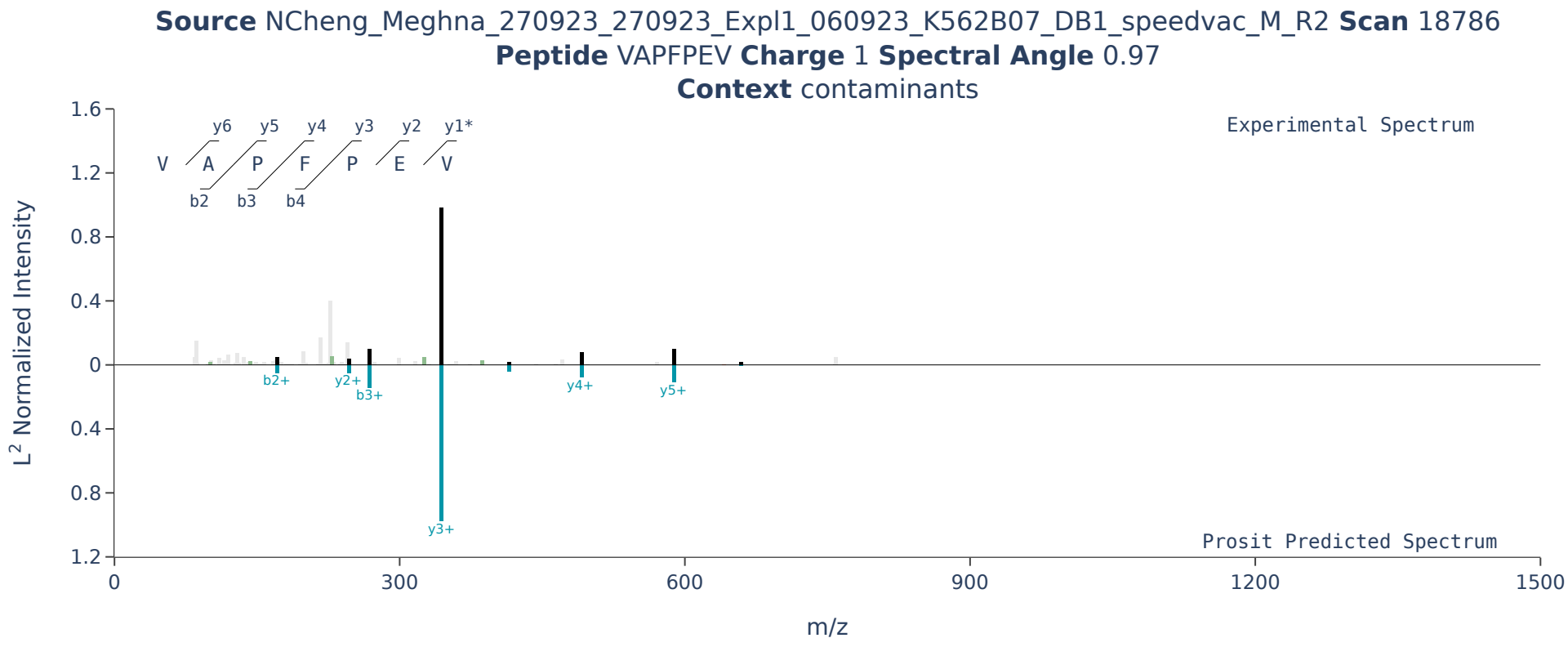
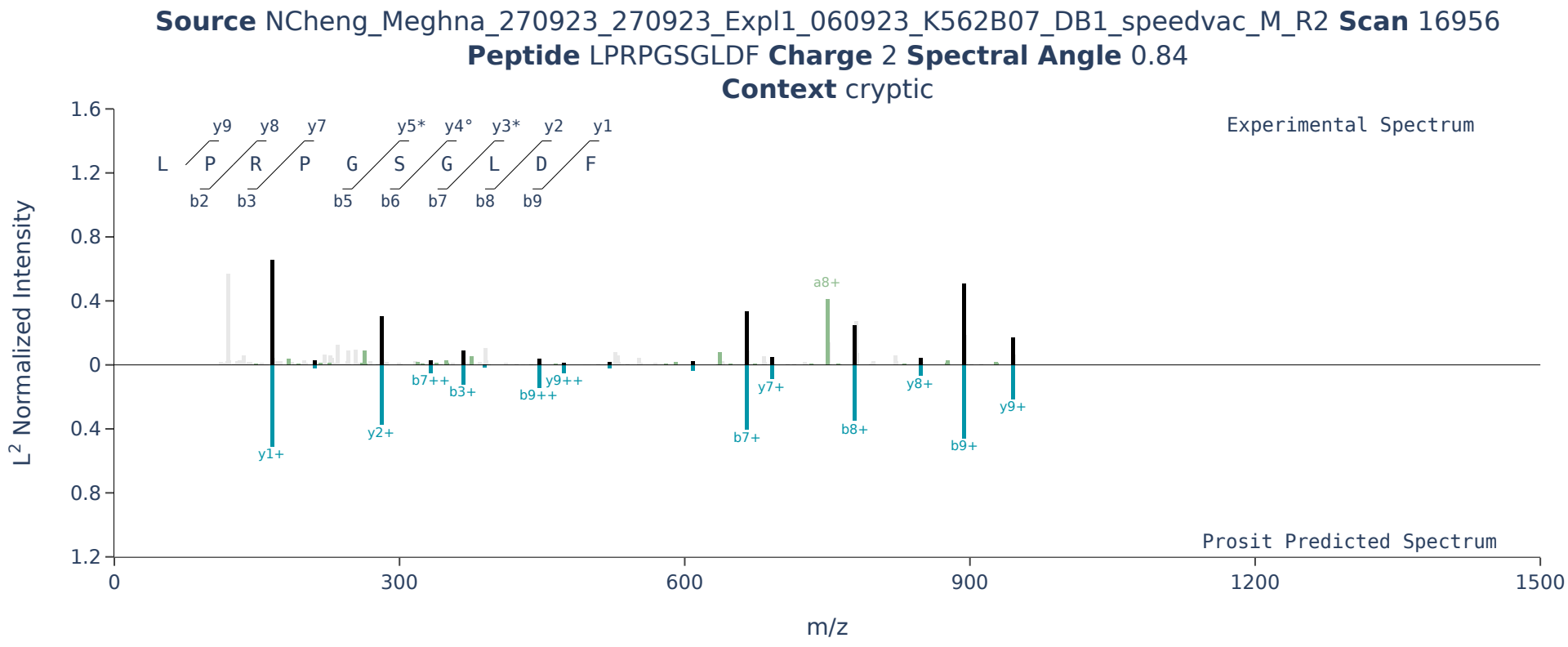
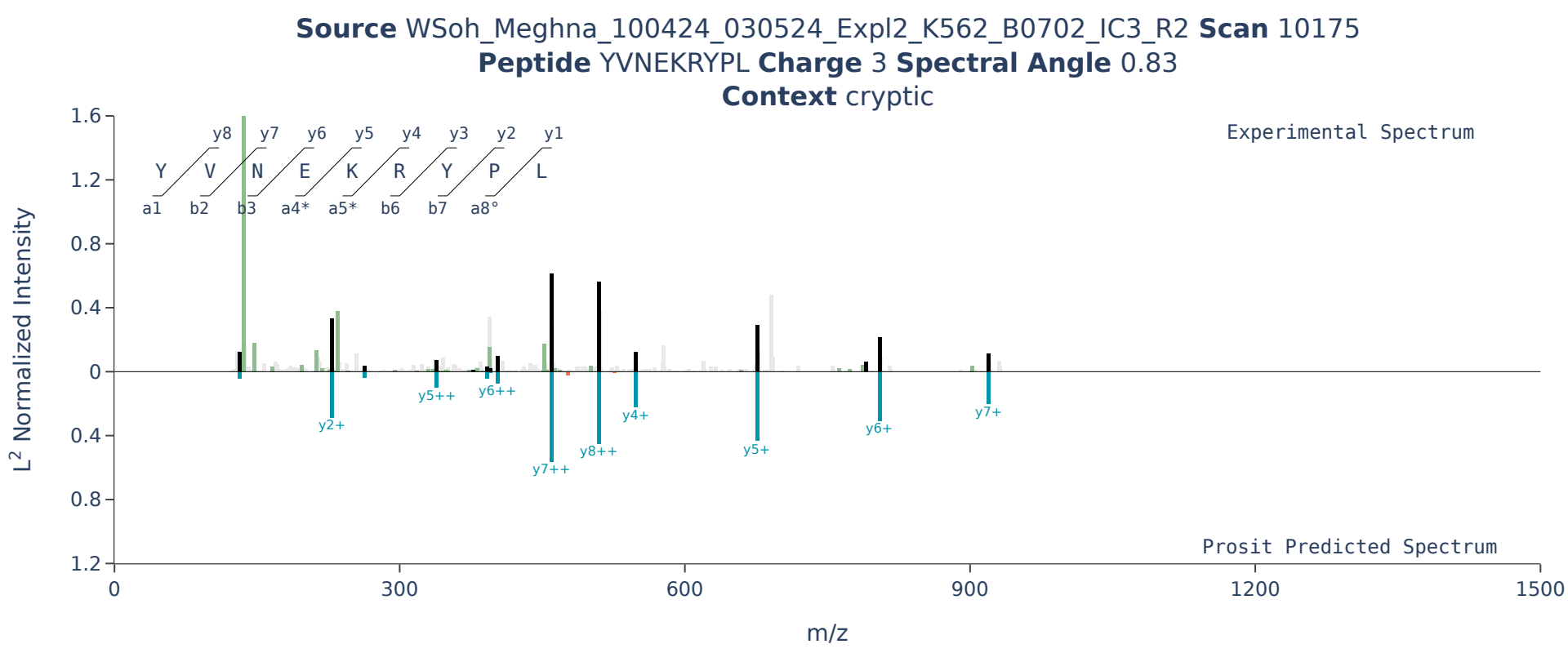
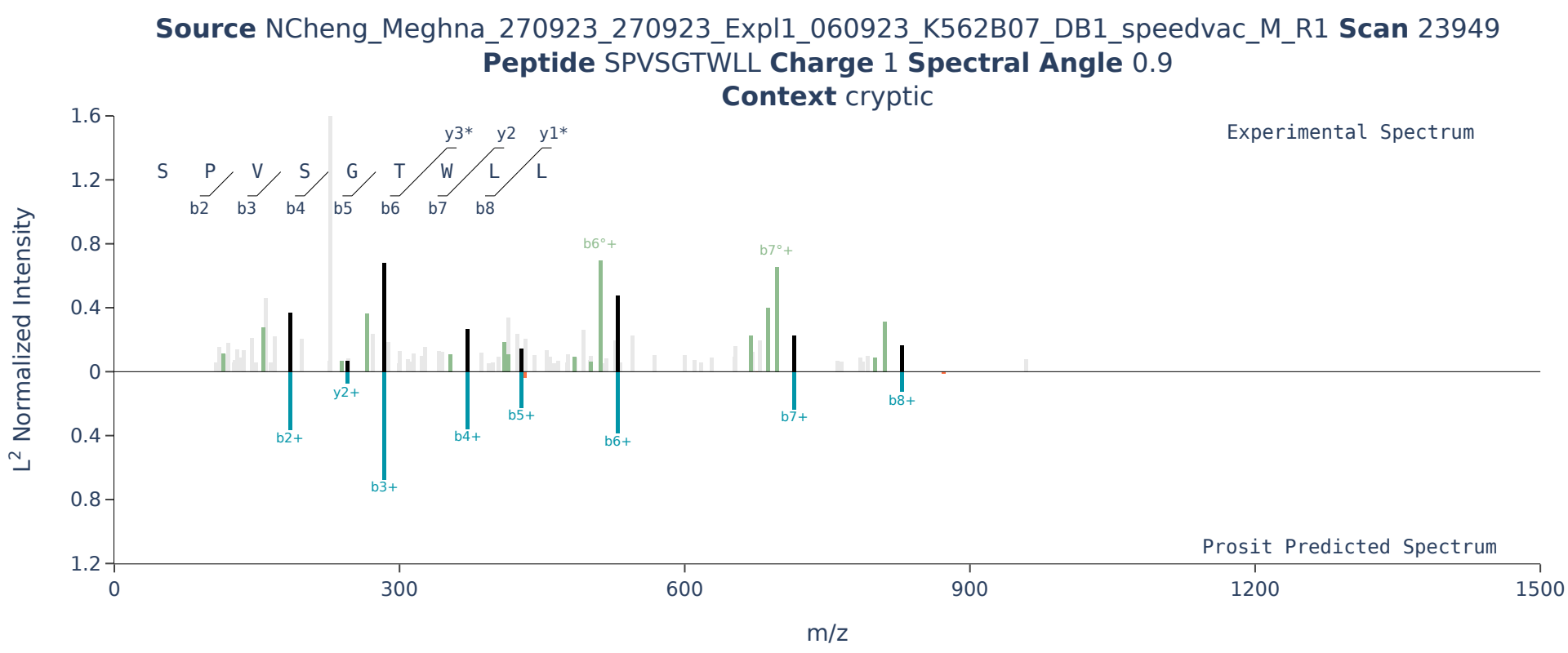


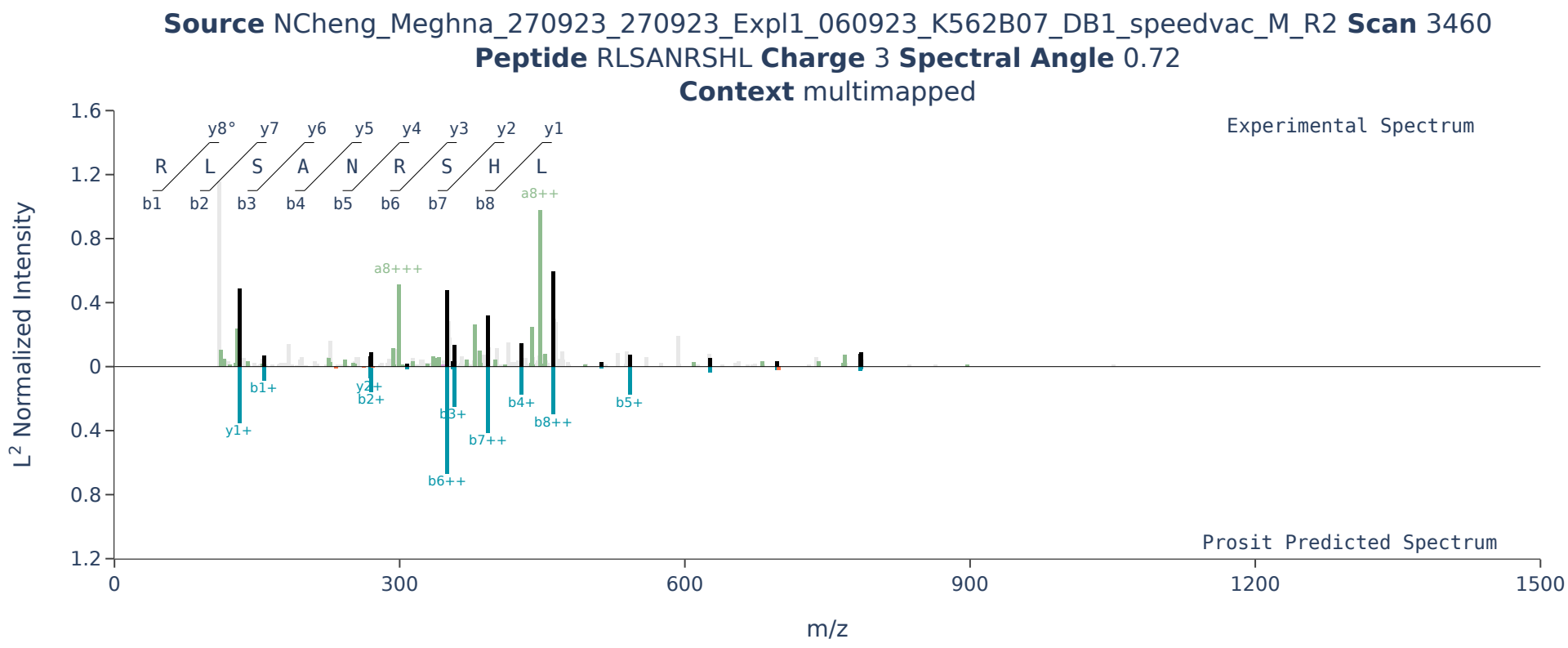
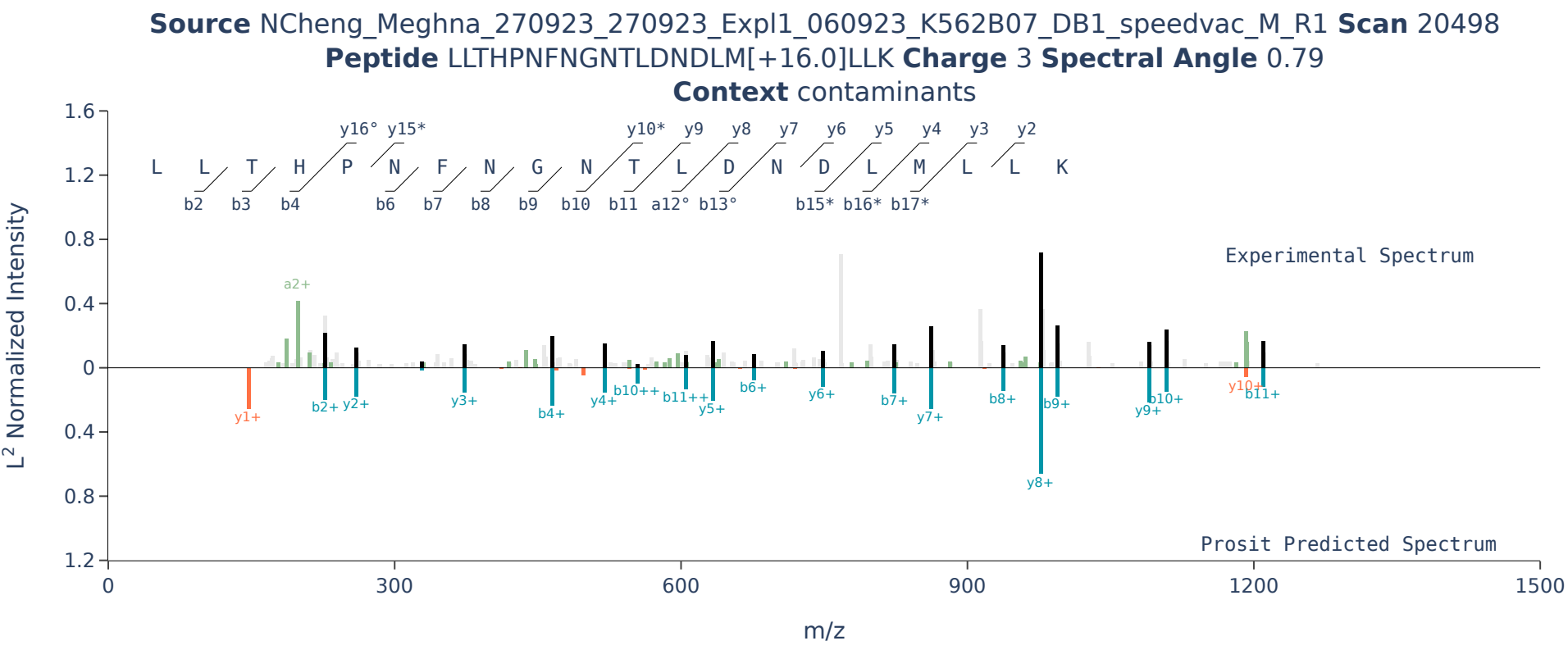
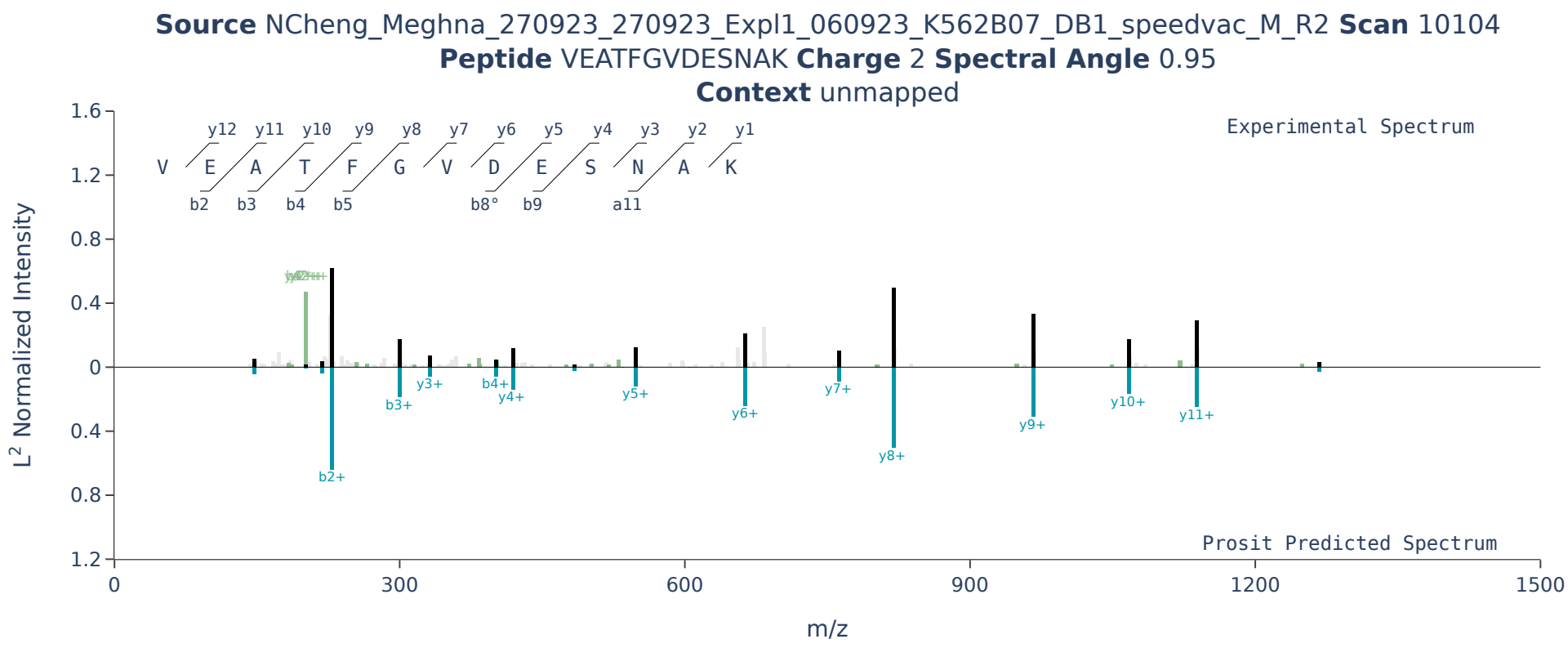
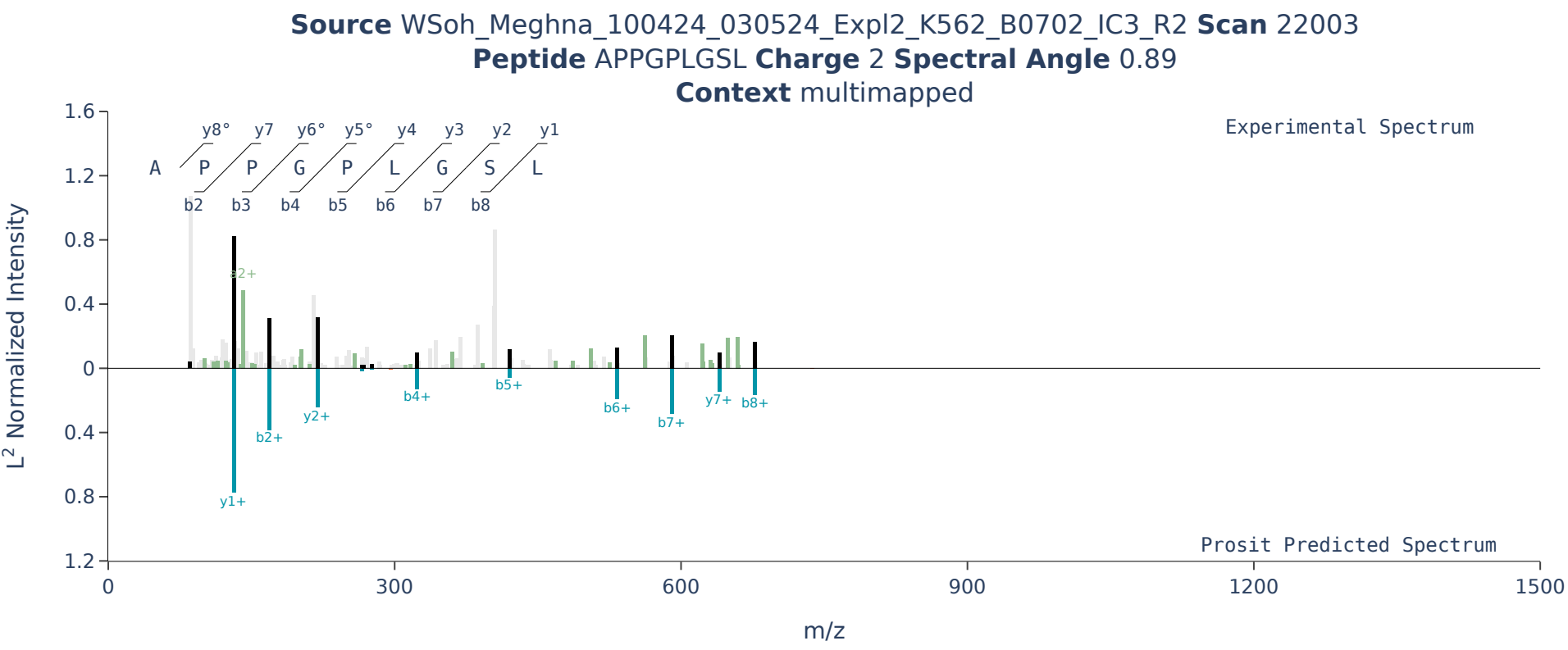
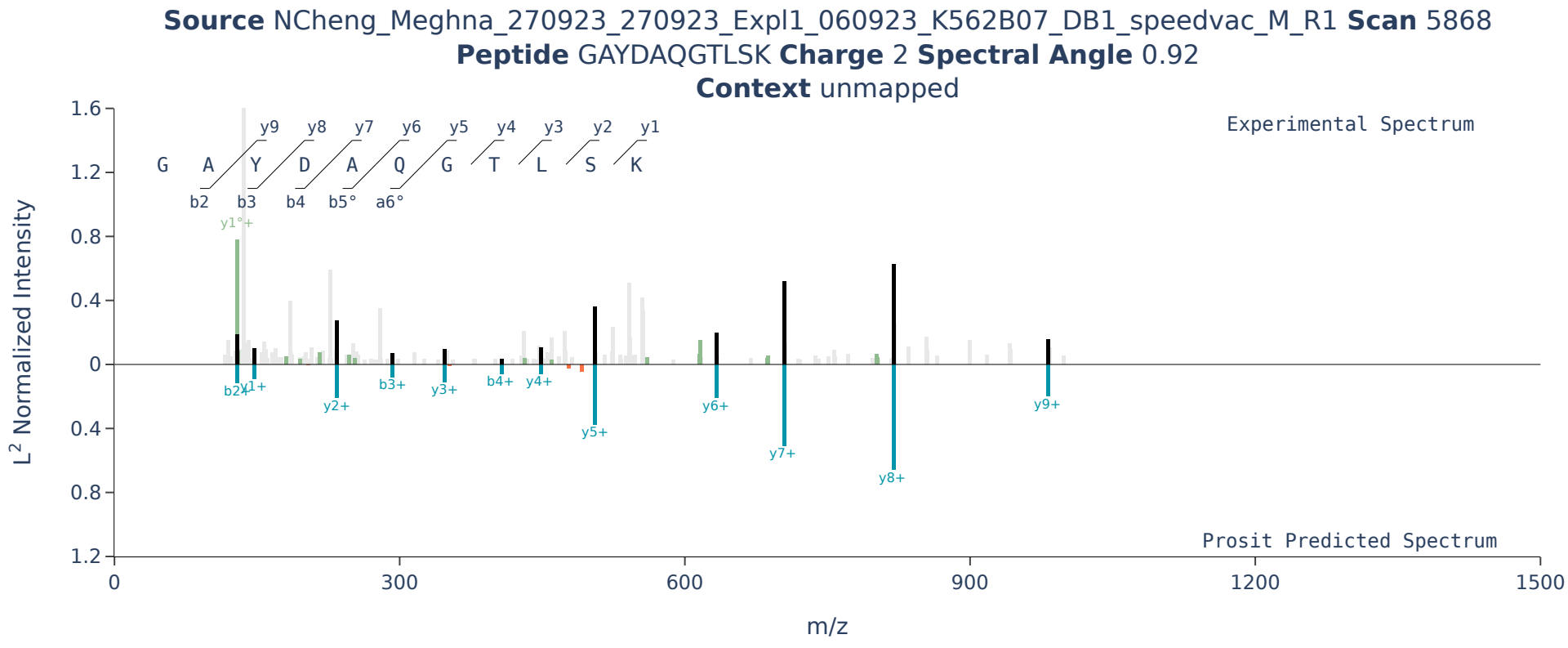
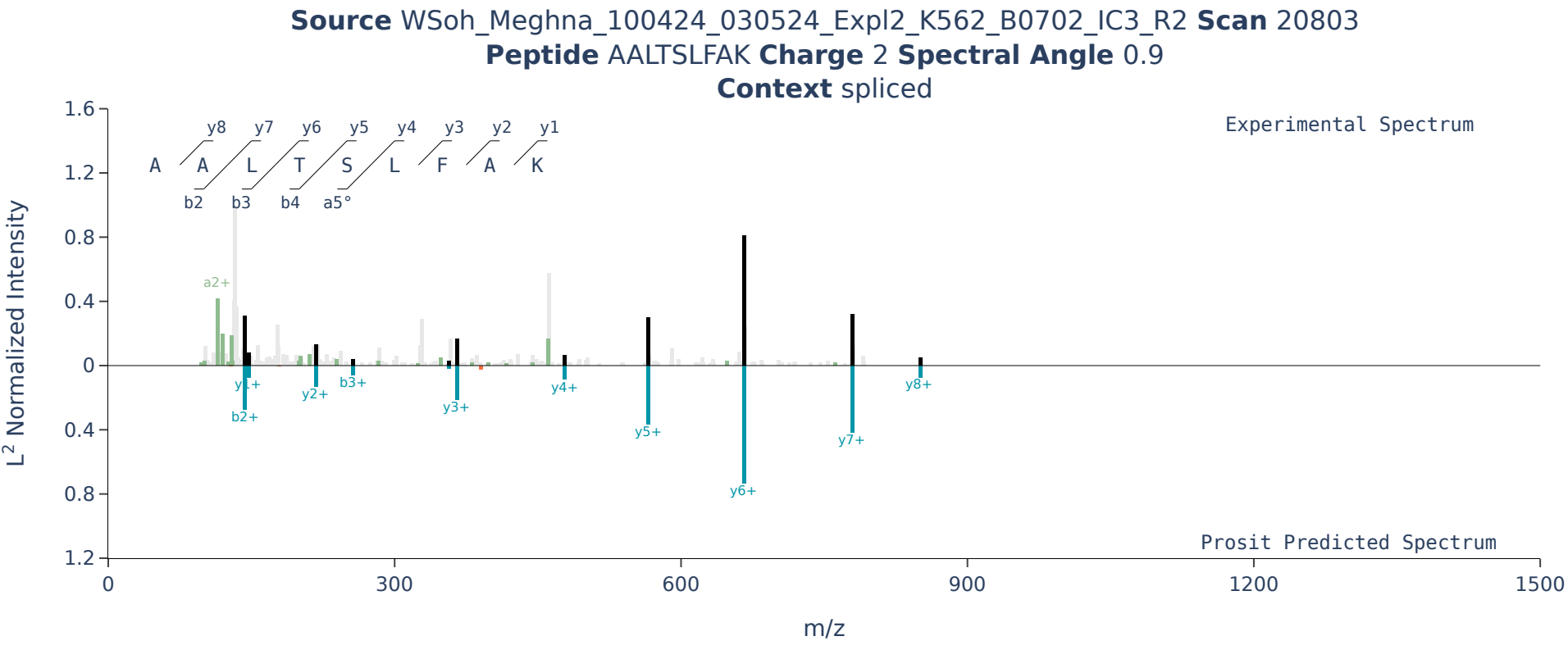
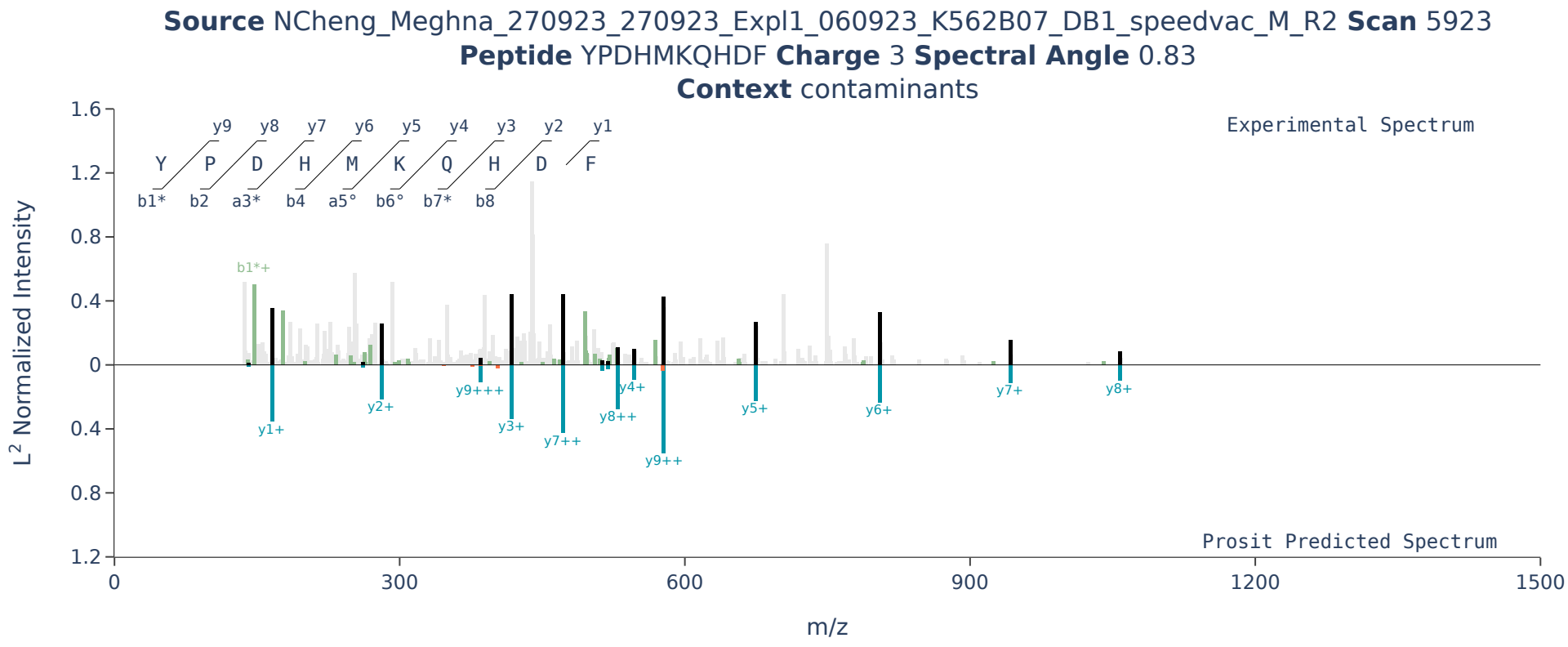
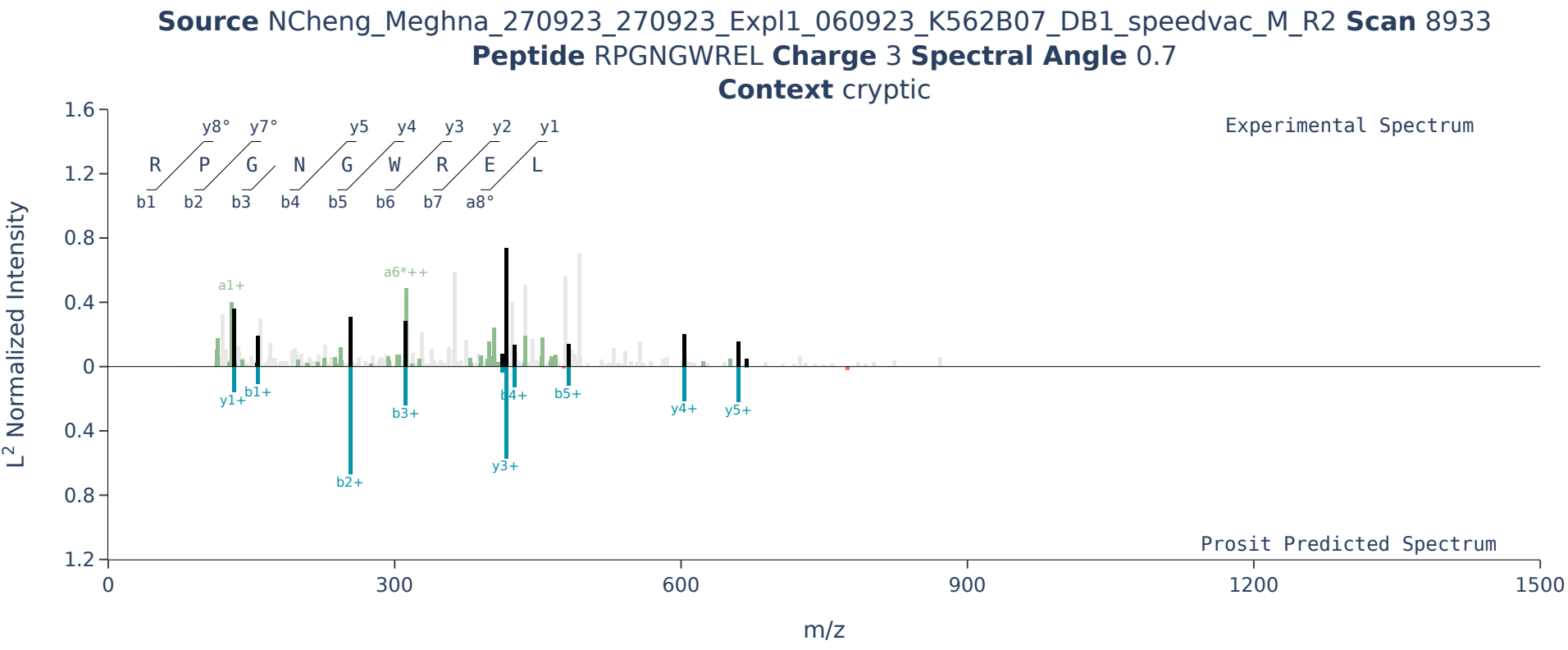
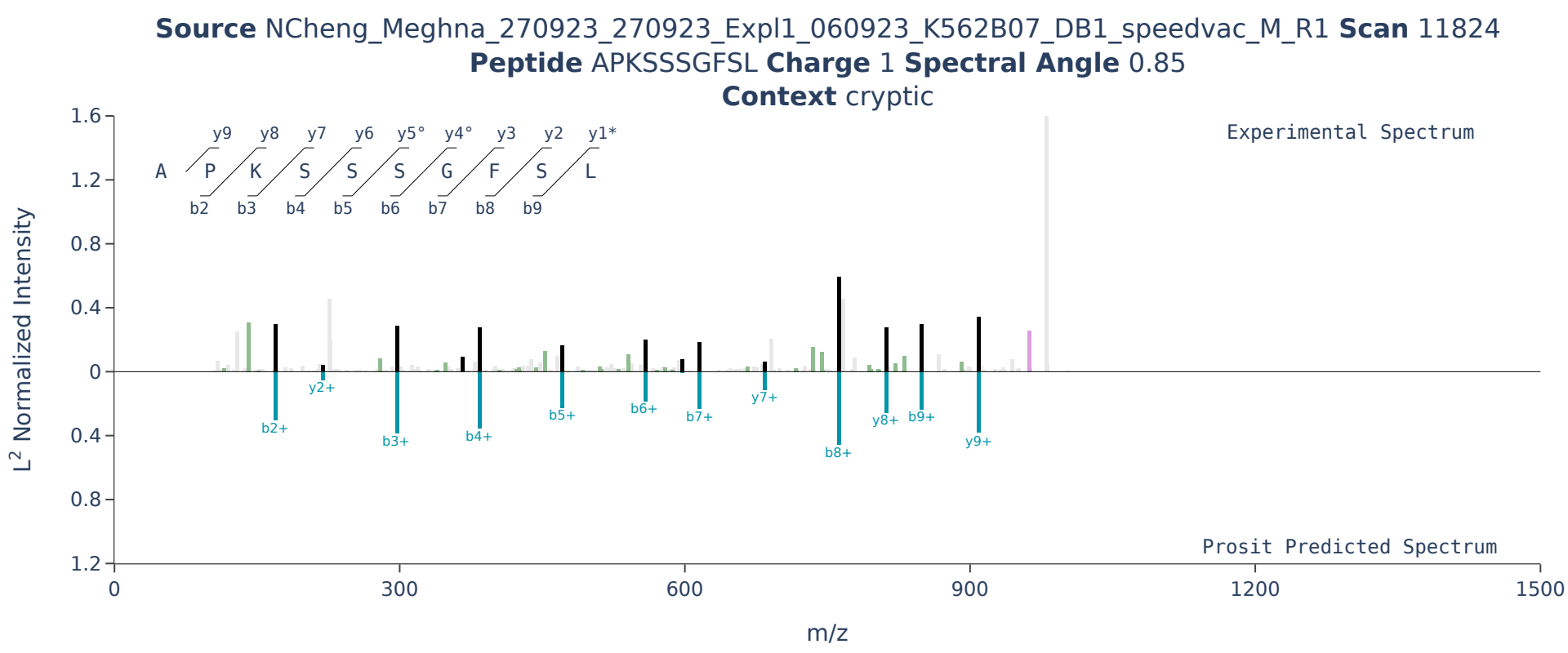
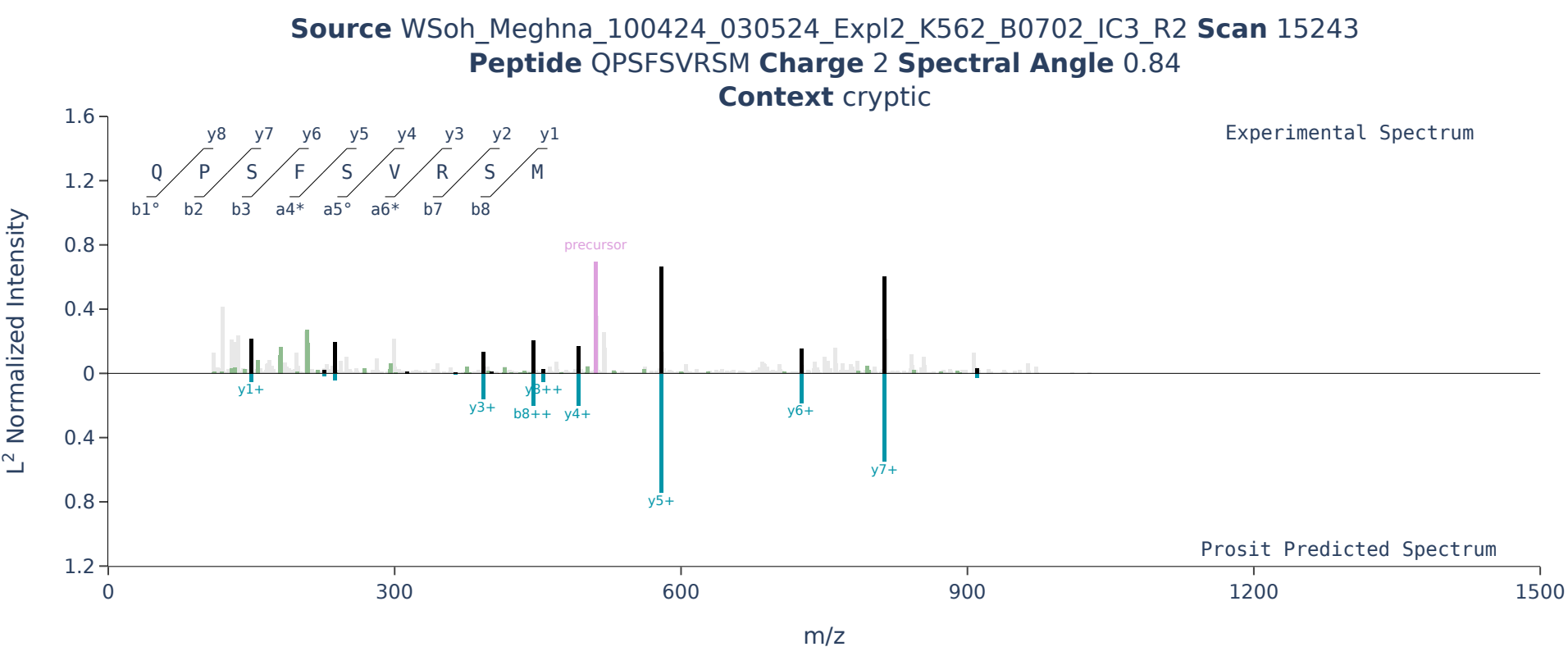
Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 20008
Peptide APGSDRLGL Charge 2 Spectral Angle 0.92

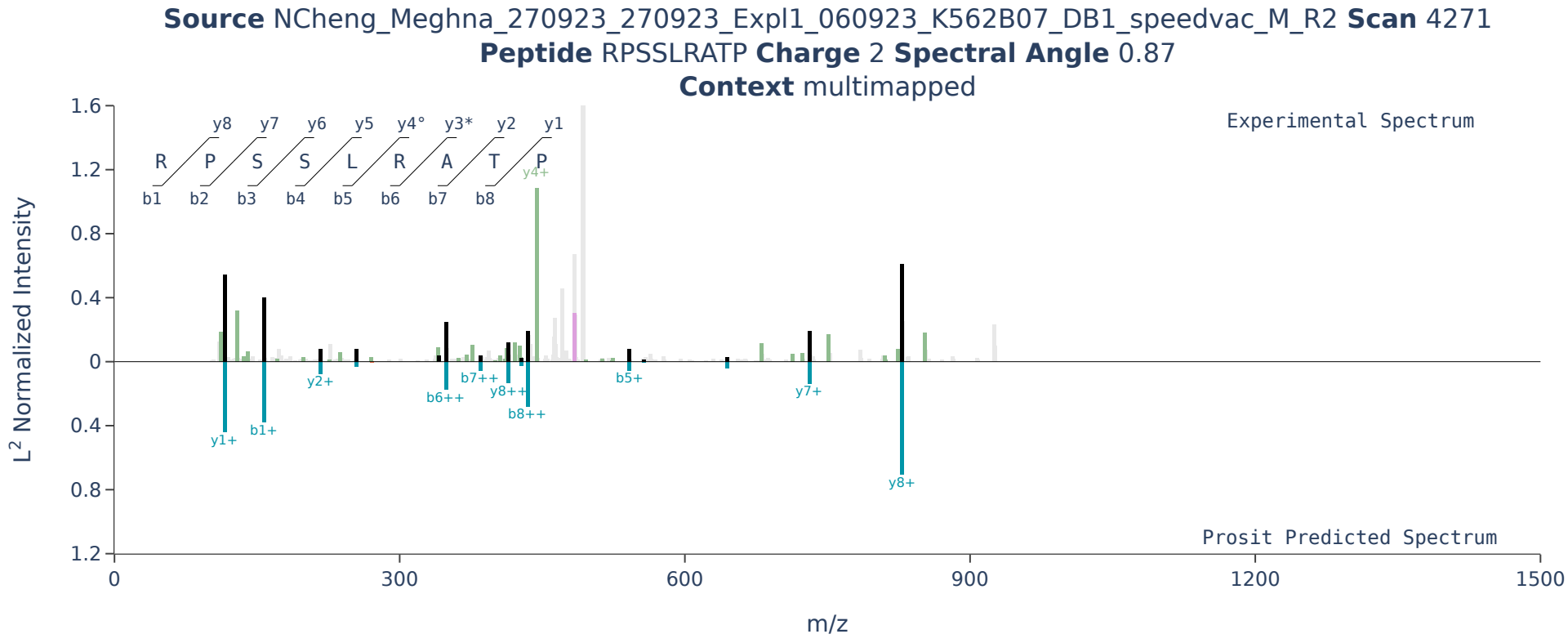
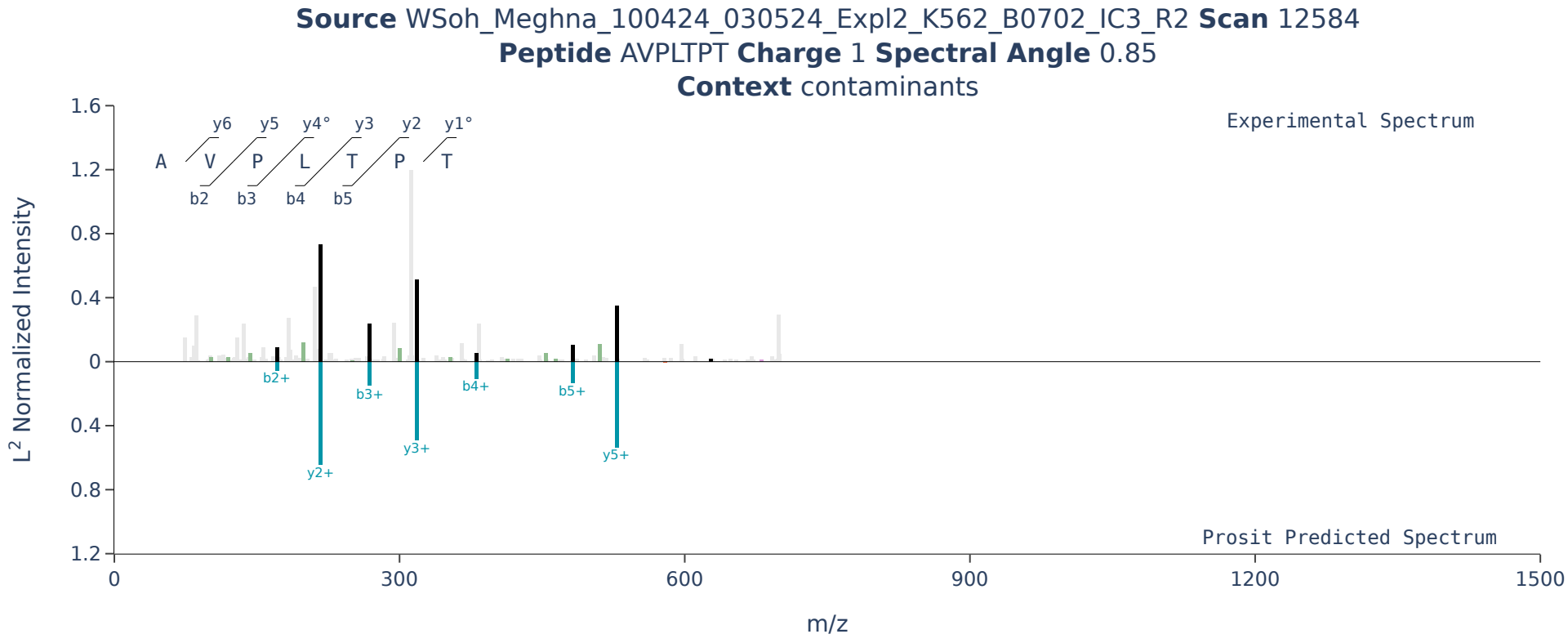
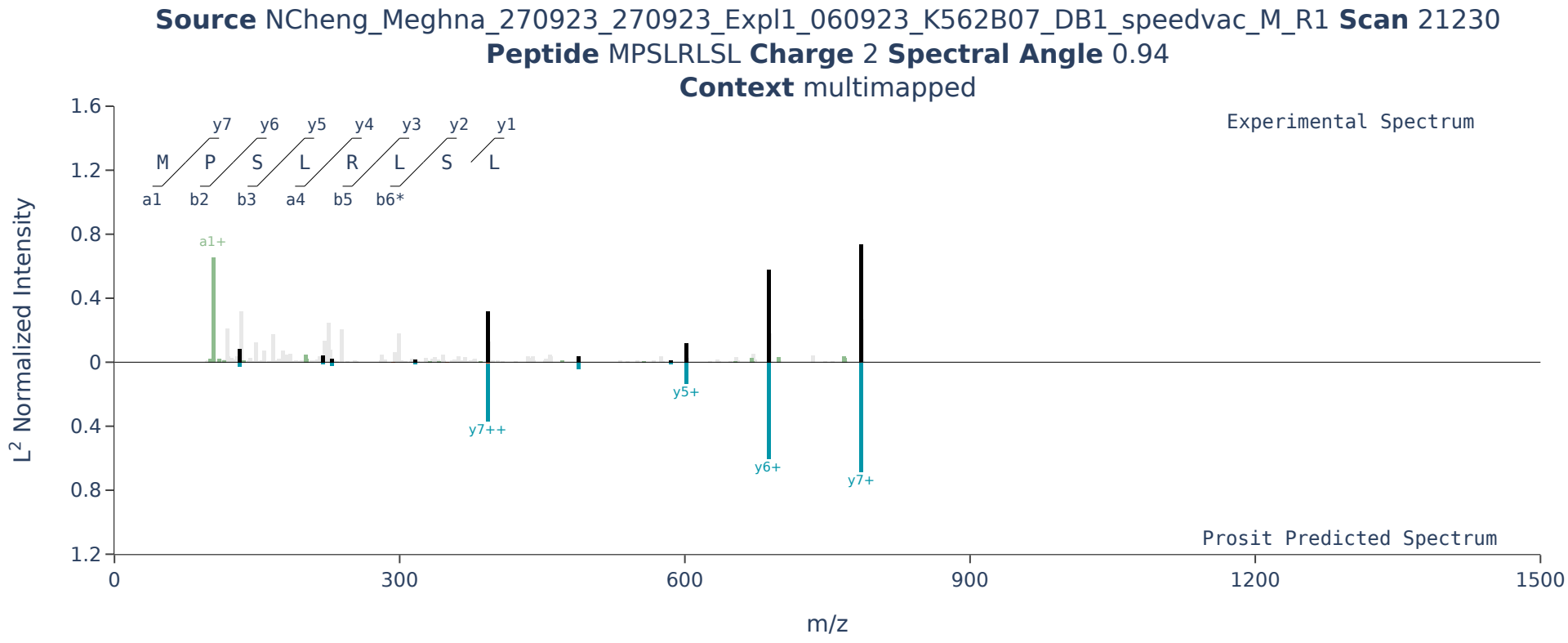
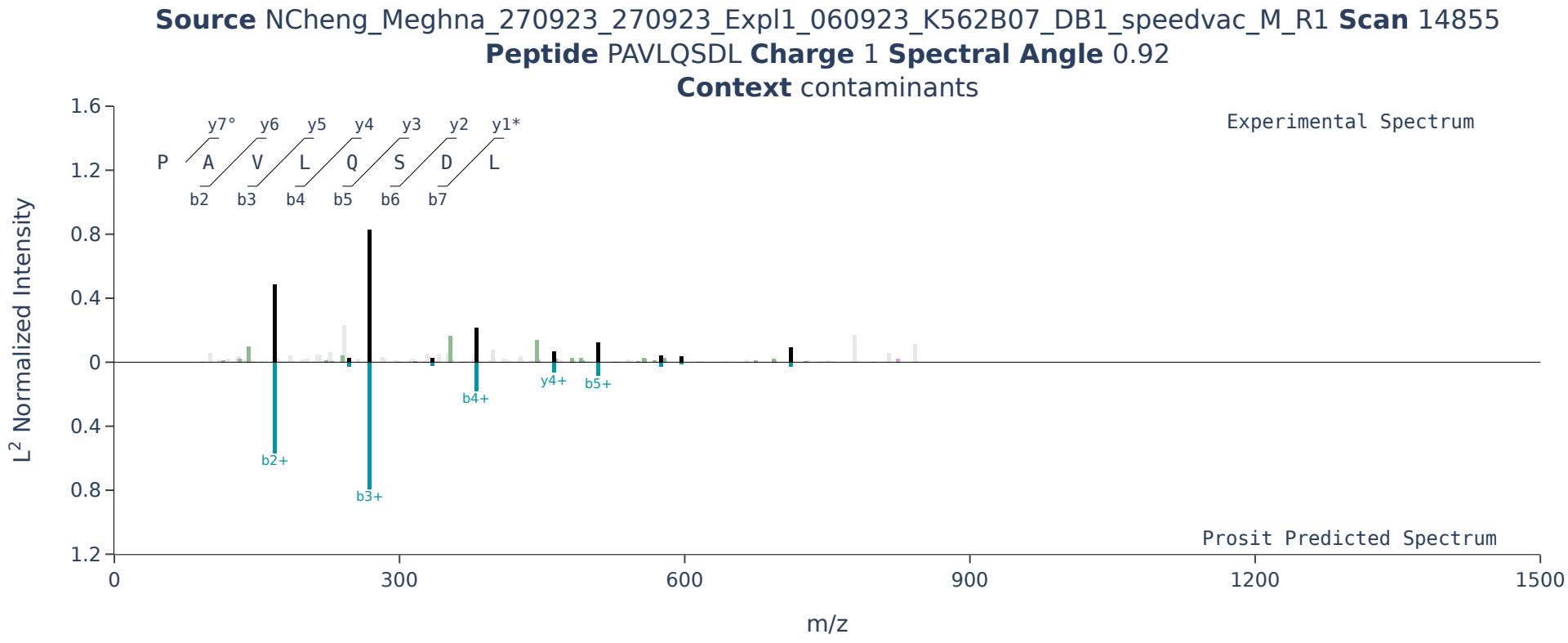
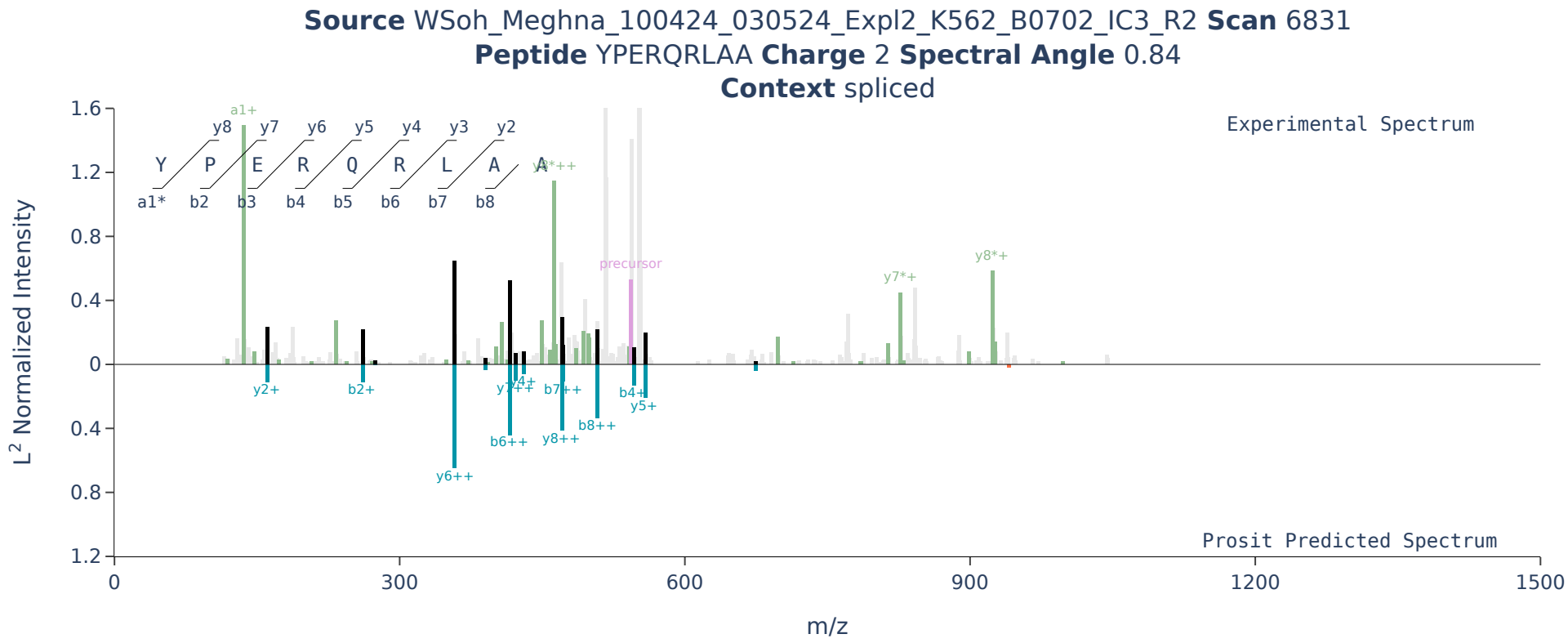
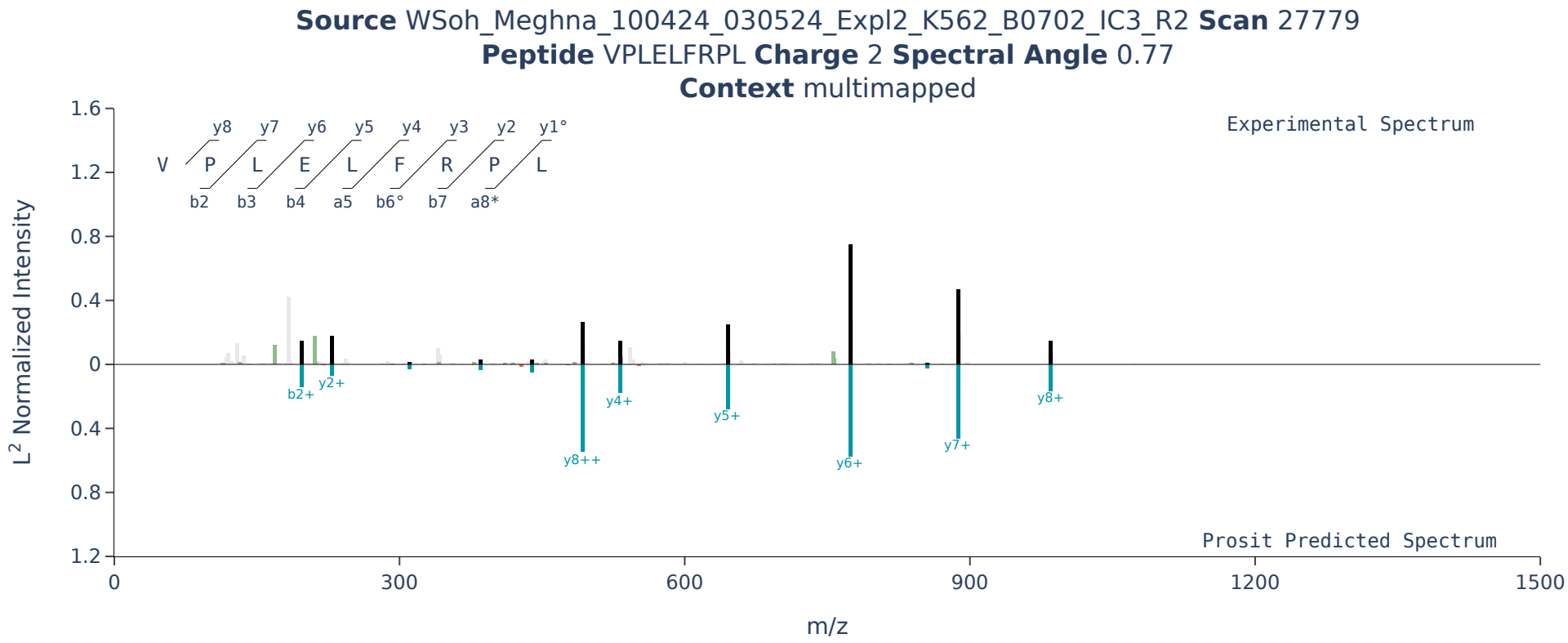
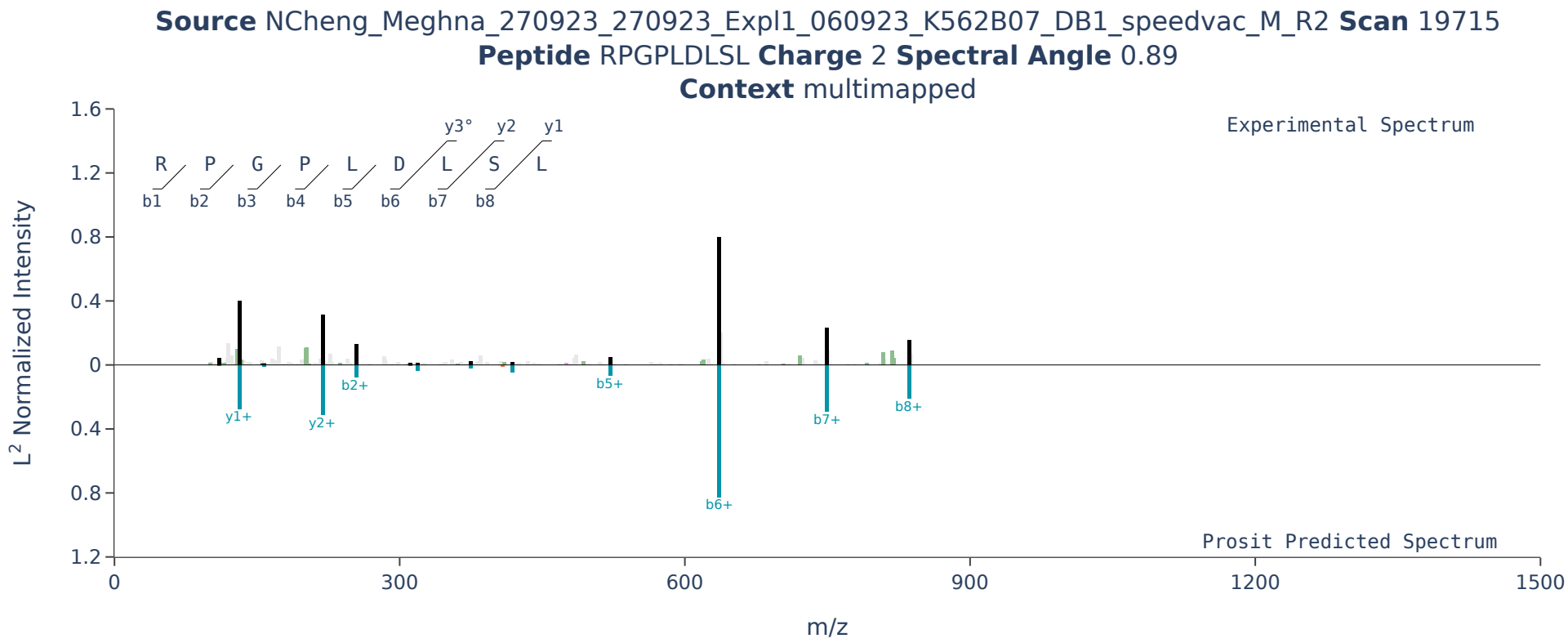
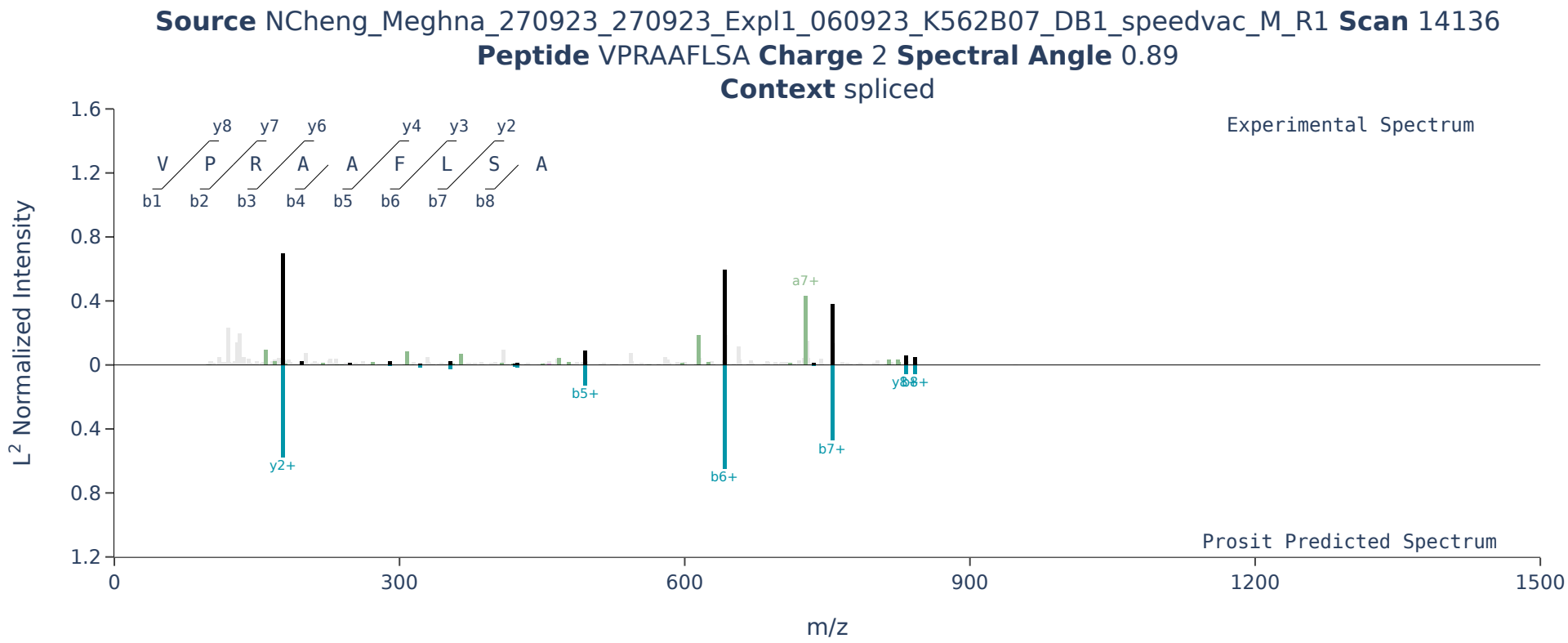
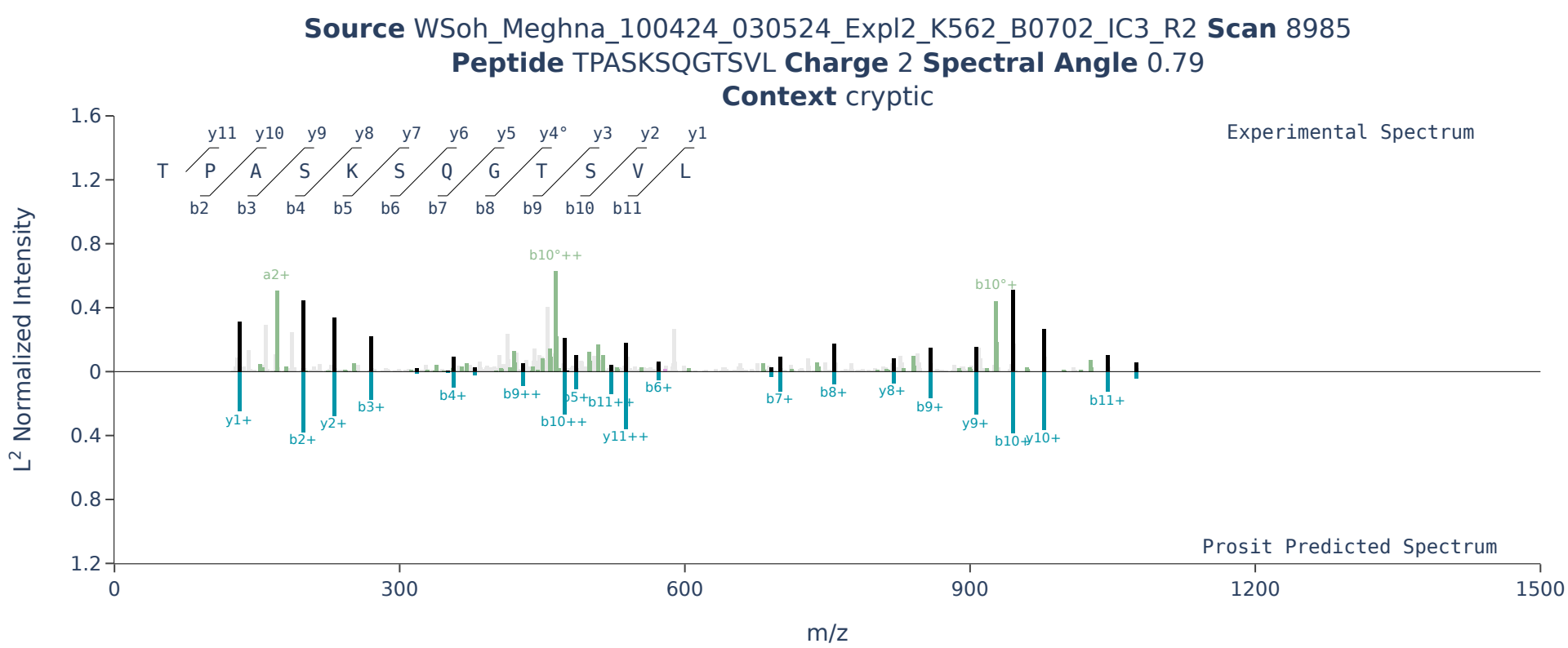
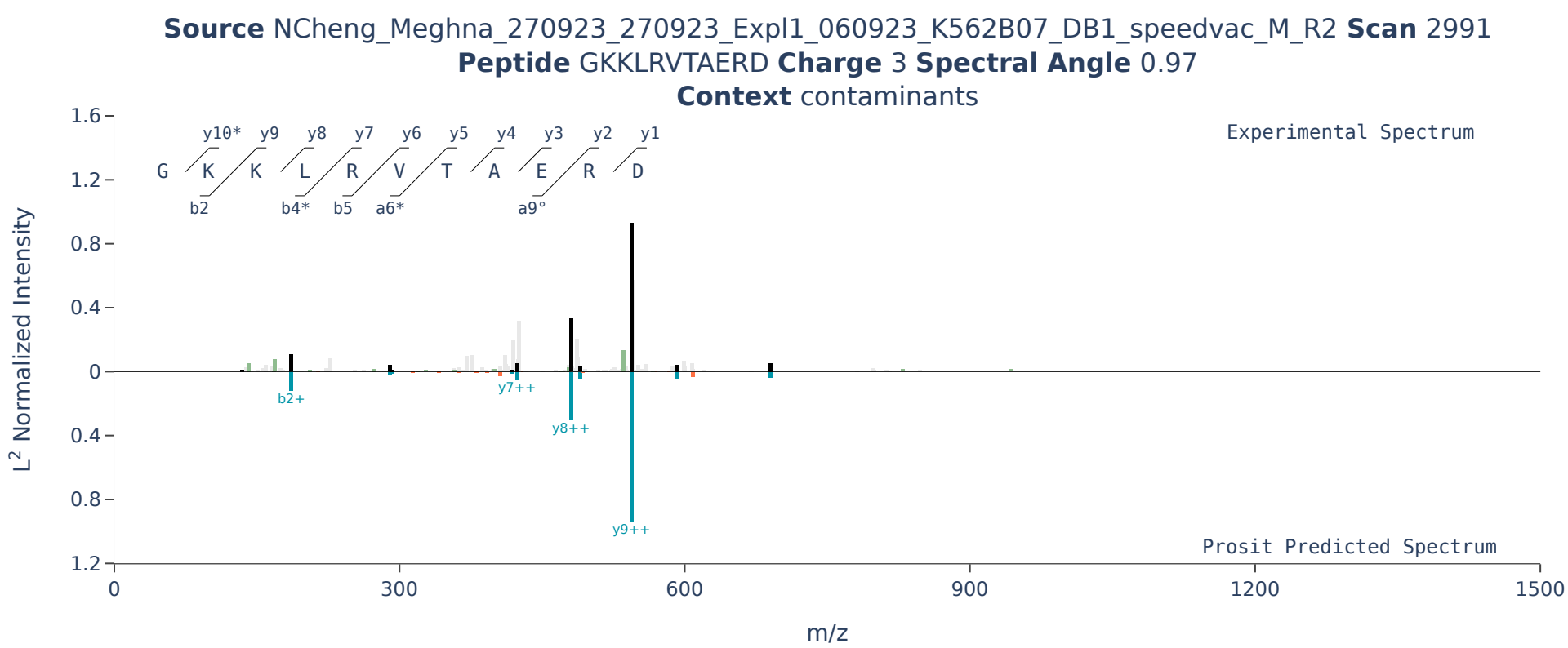


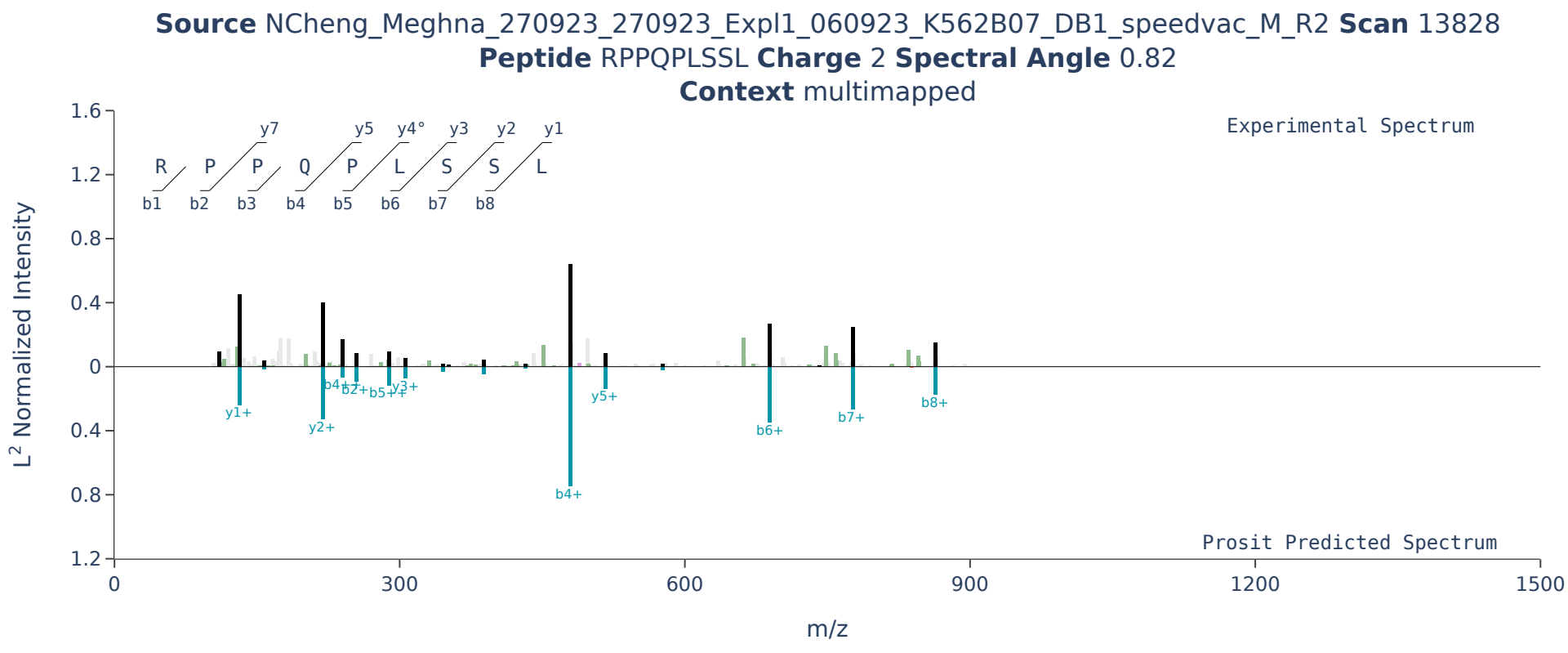
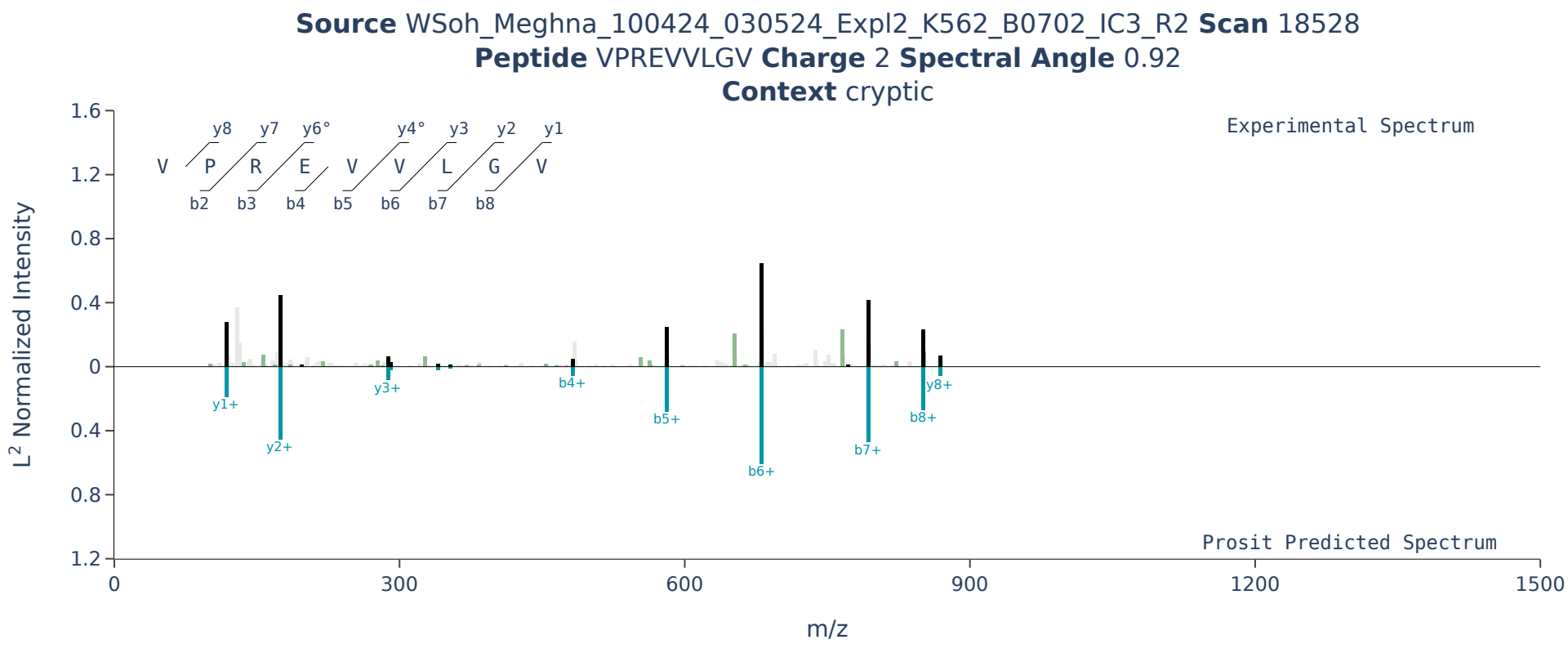
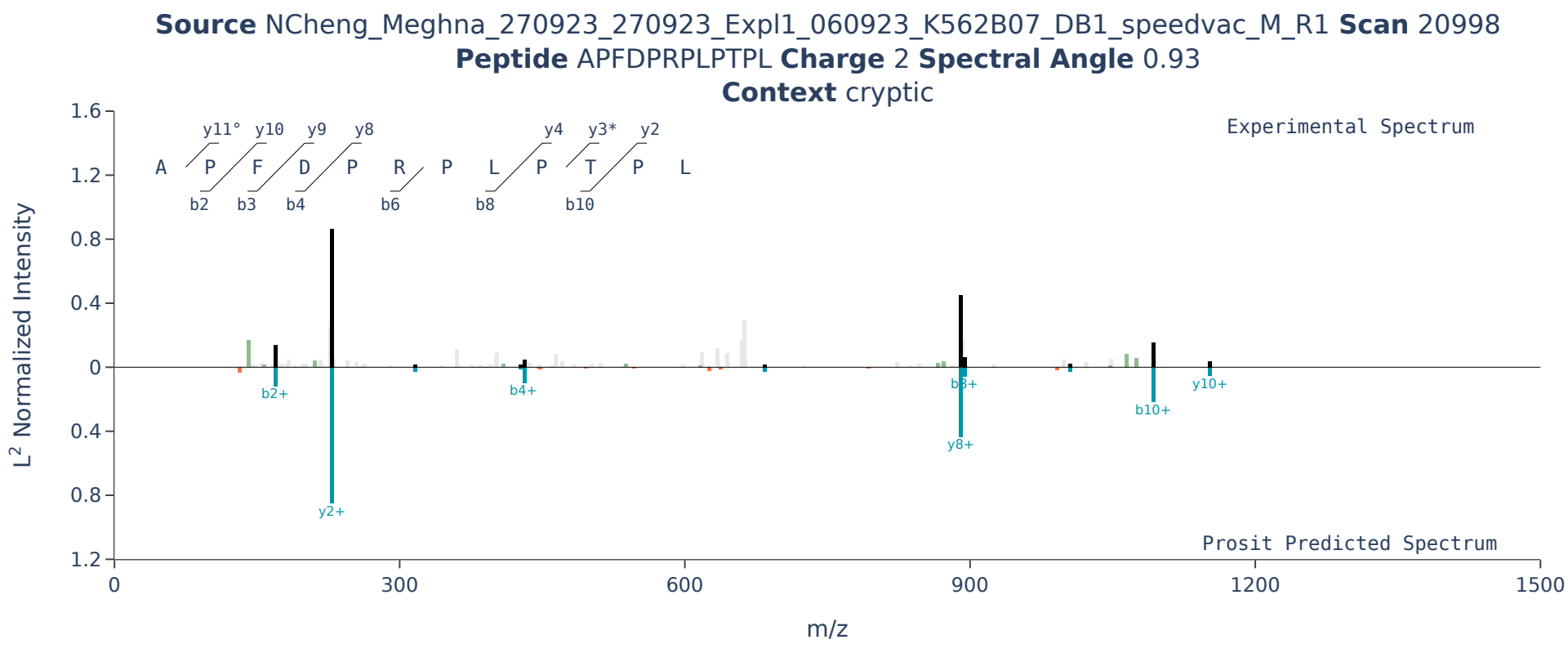
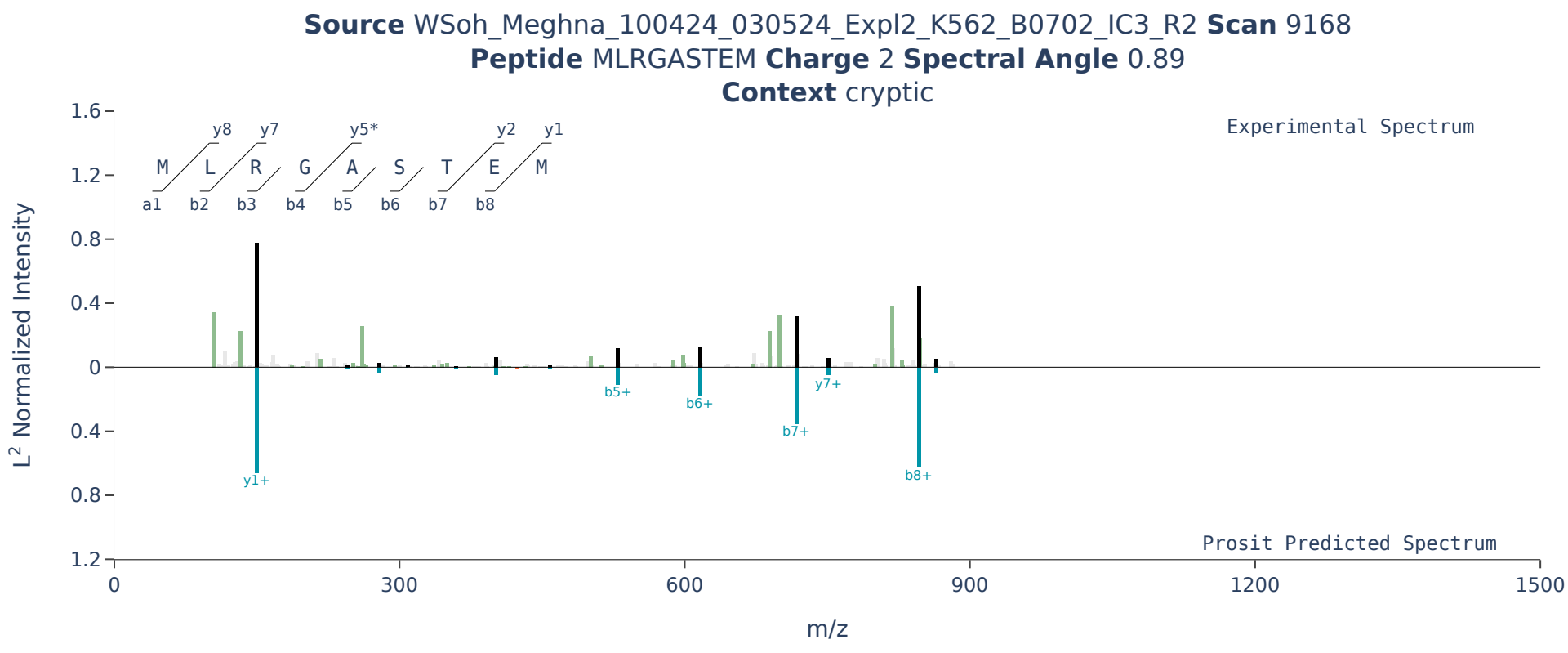
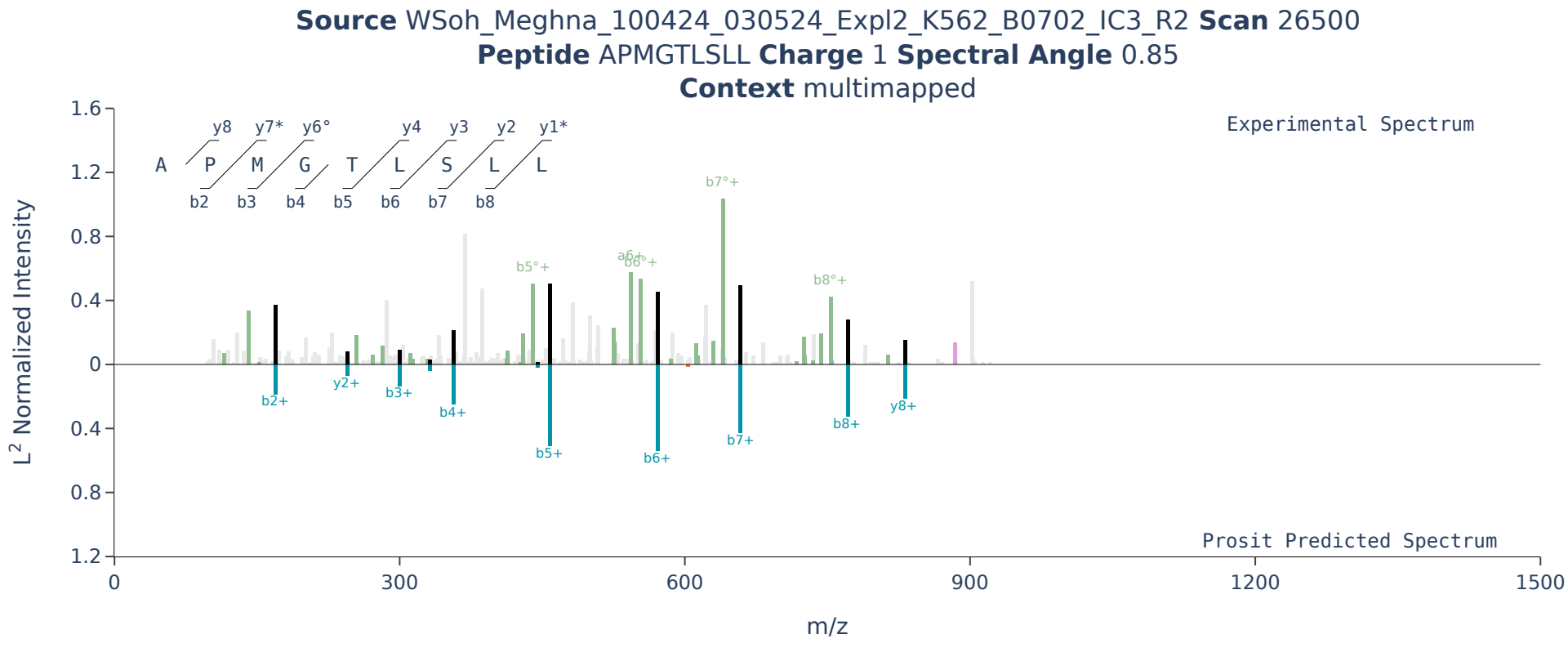
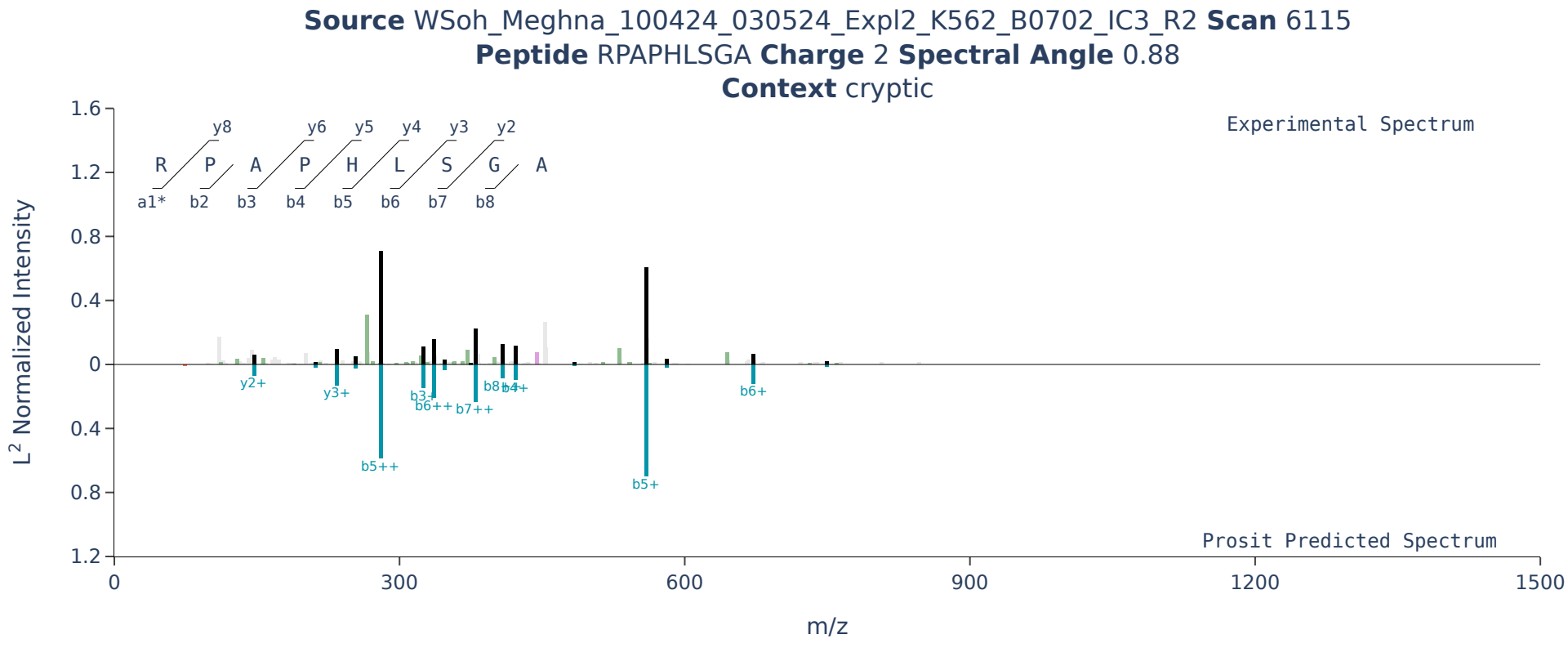
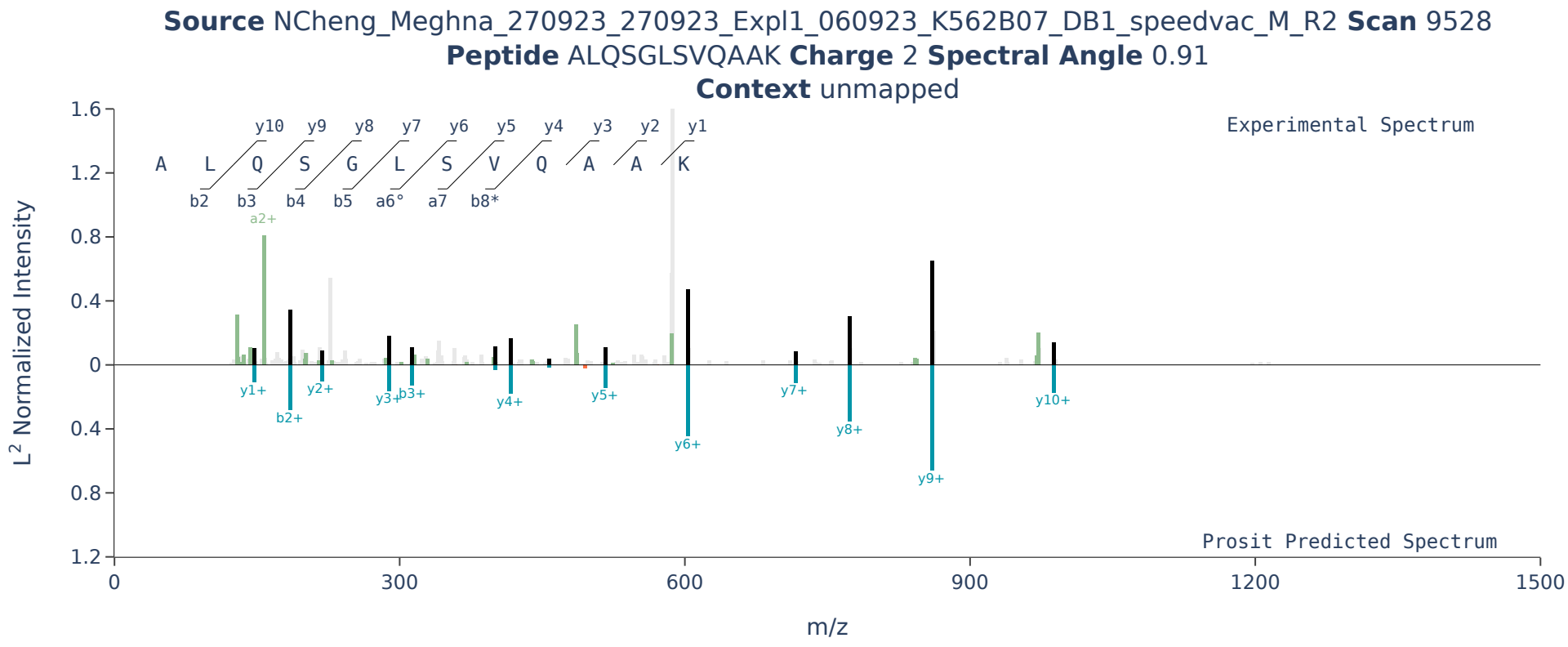
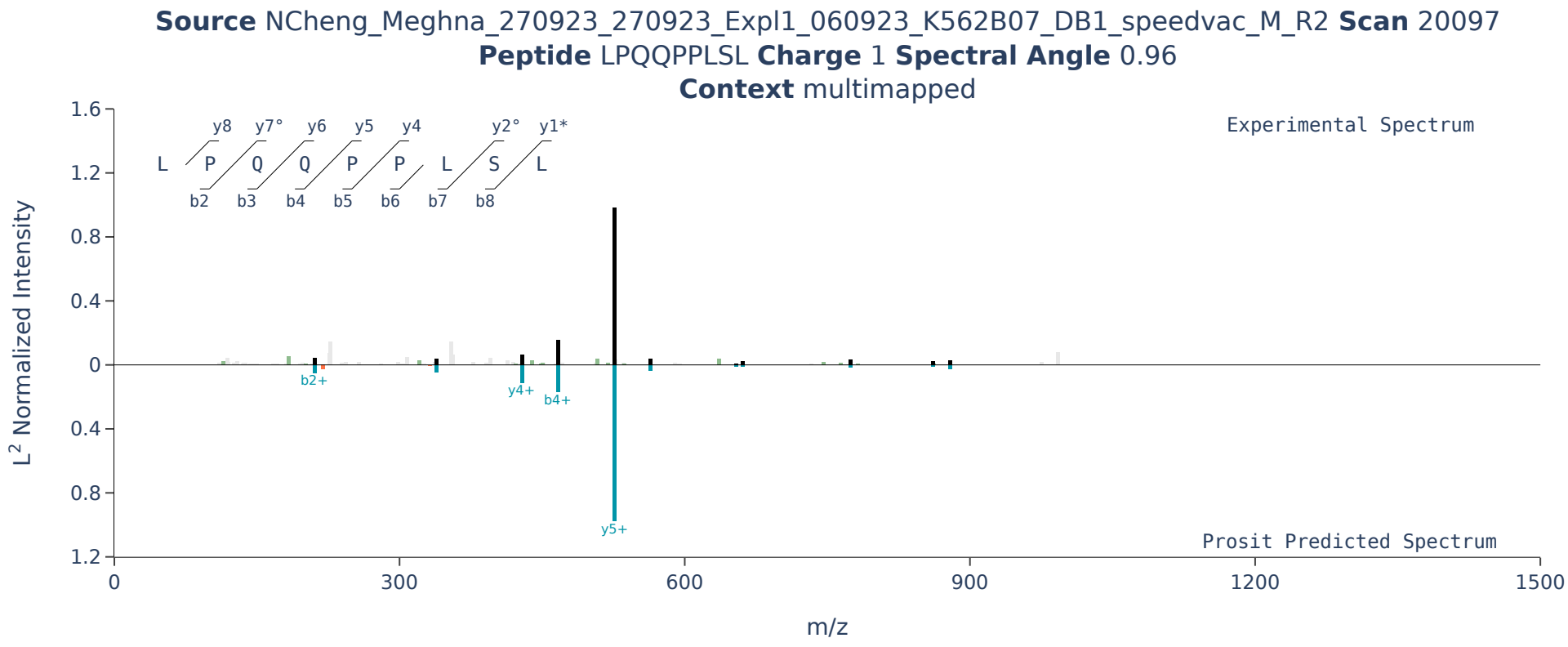
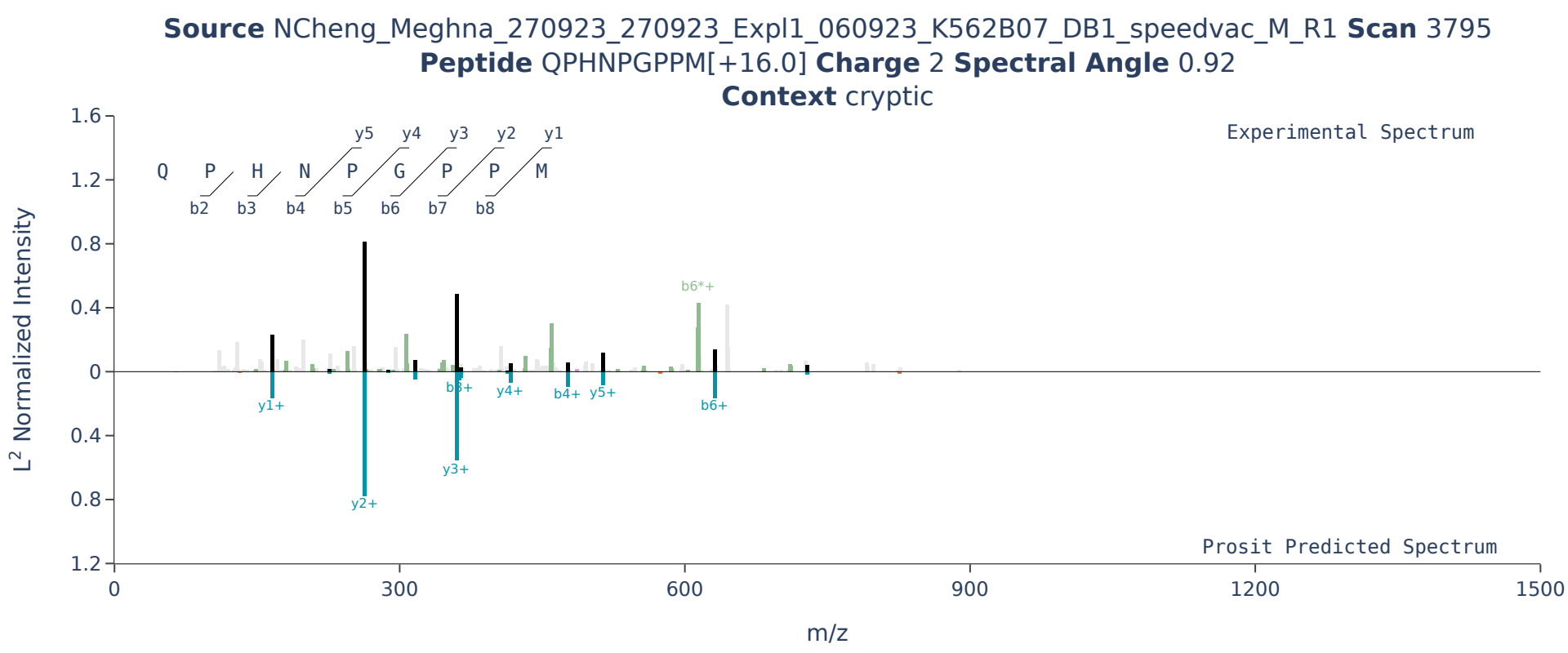
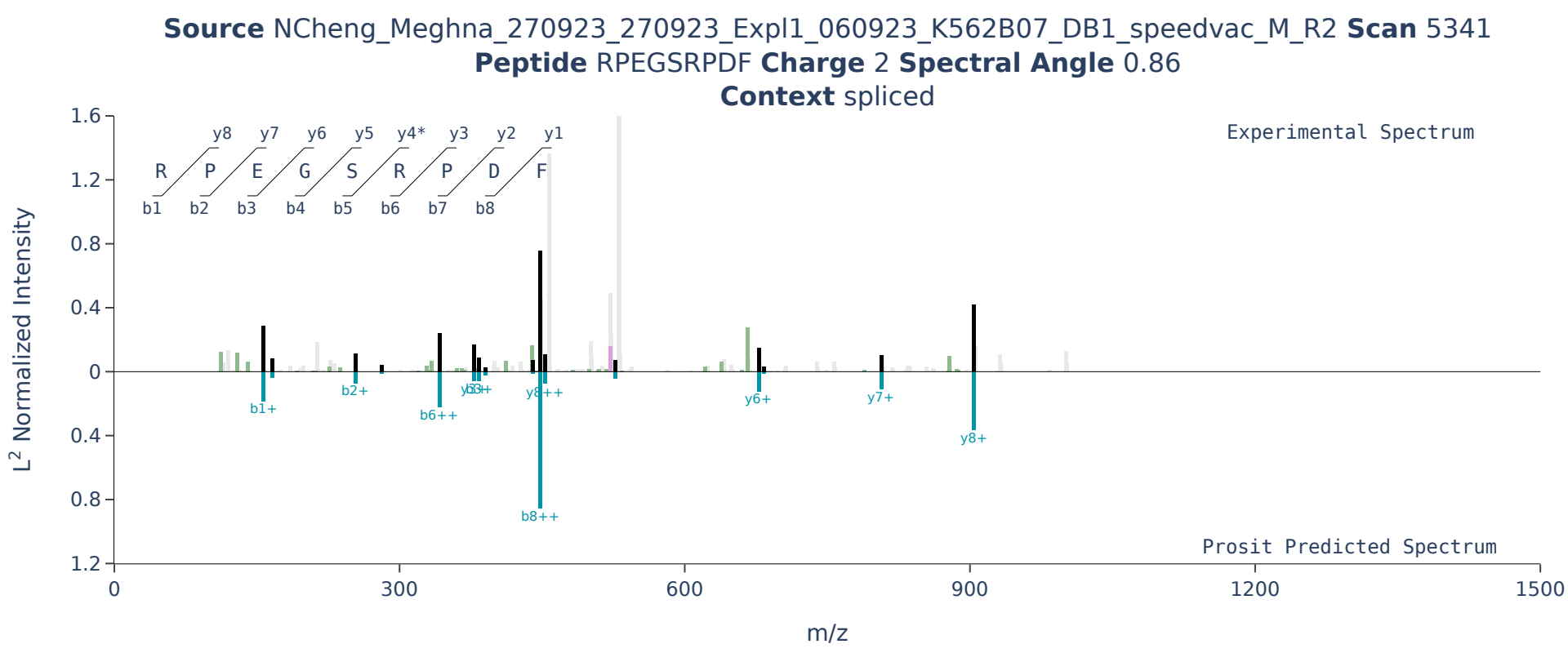






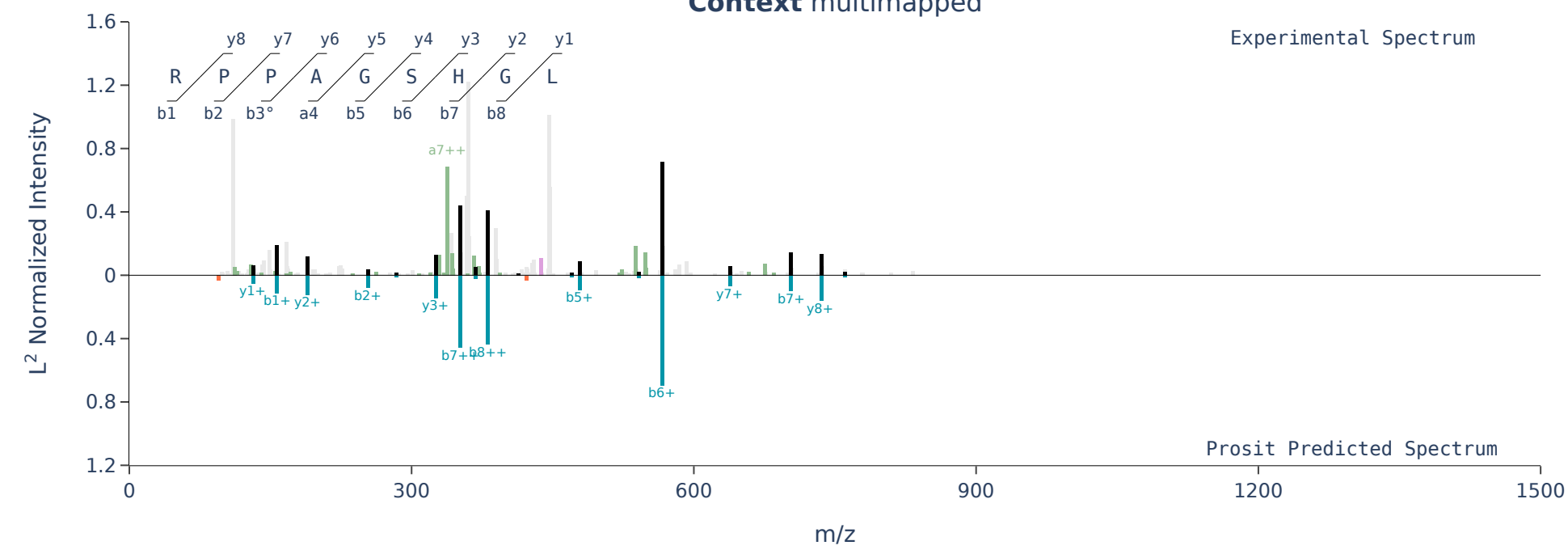






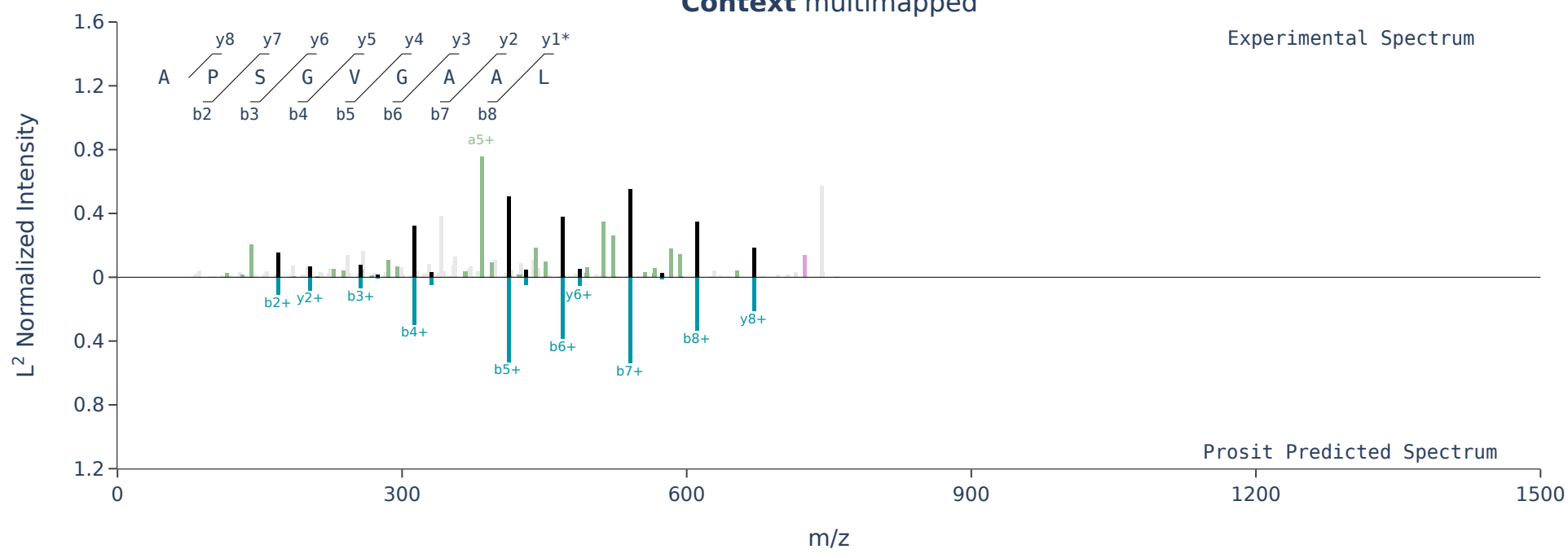
Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 5629
Peptide RPPAGSHGL Charge 2 Spectral Angle 0.92

Context multimapped



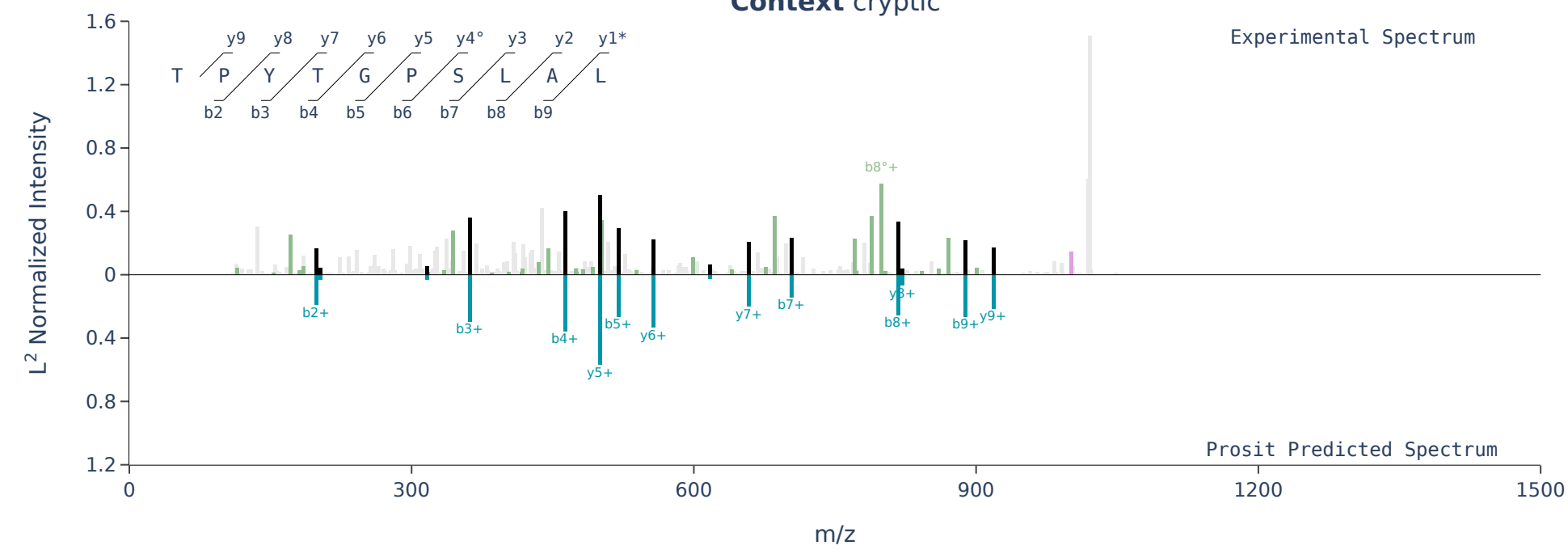
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Peptide APSGVGAAL Charge 1 Spectral Angle 0.95

Context multimapped



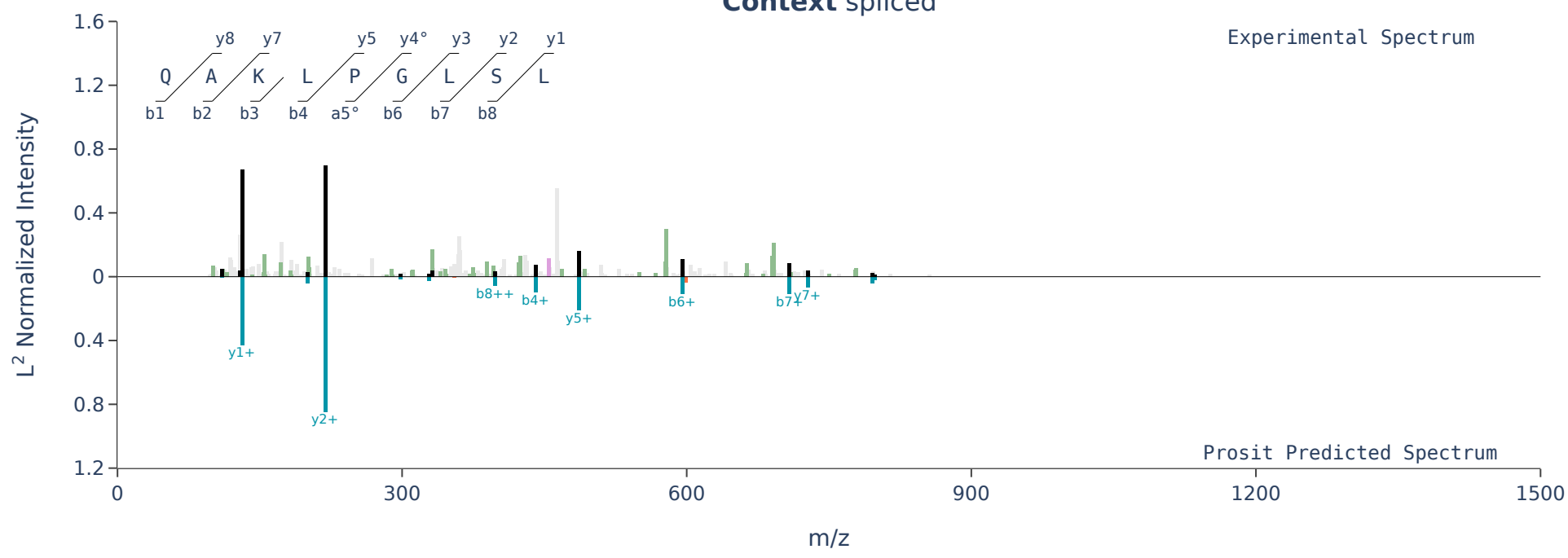
Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 23081
Peptide TPYTGPSLAL Charge 1 Spectral Angle 0.86

Context cryptic



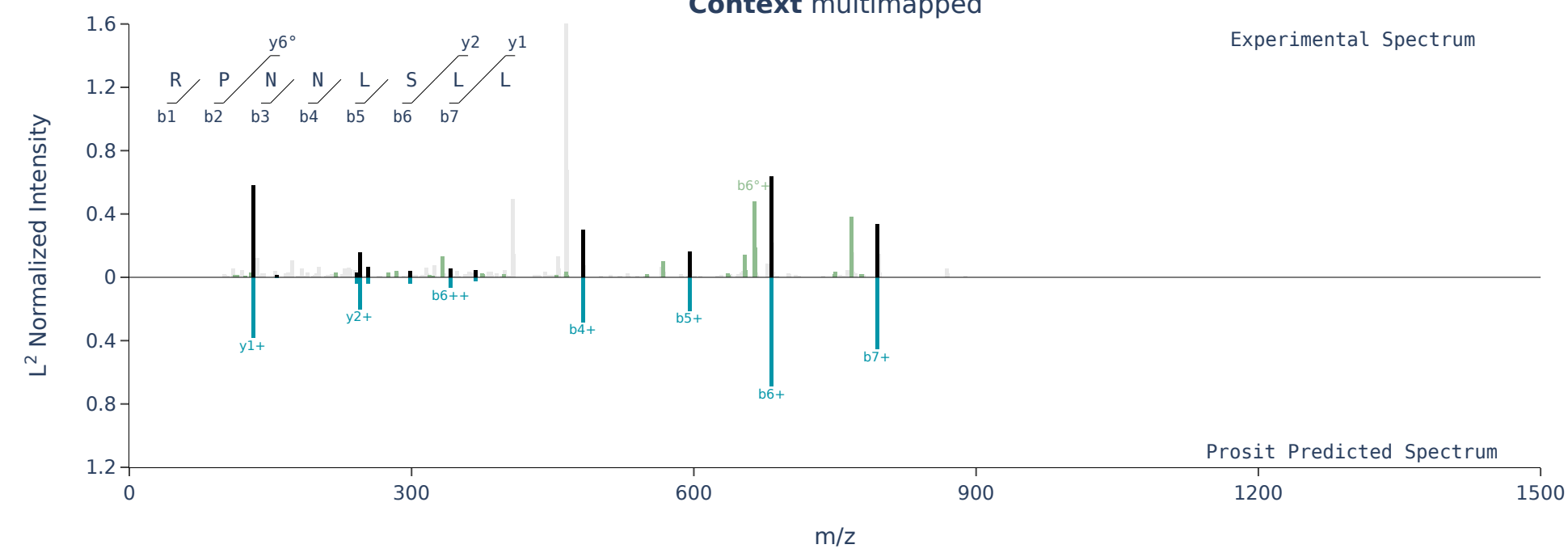
Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 21405
Peptide QAKLPGLSL Charge 2 Spectral Angle 0.81

Context spliced



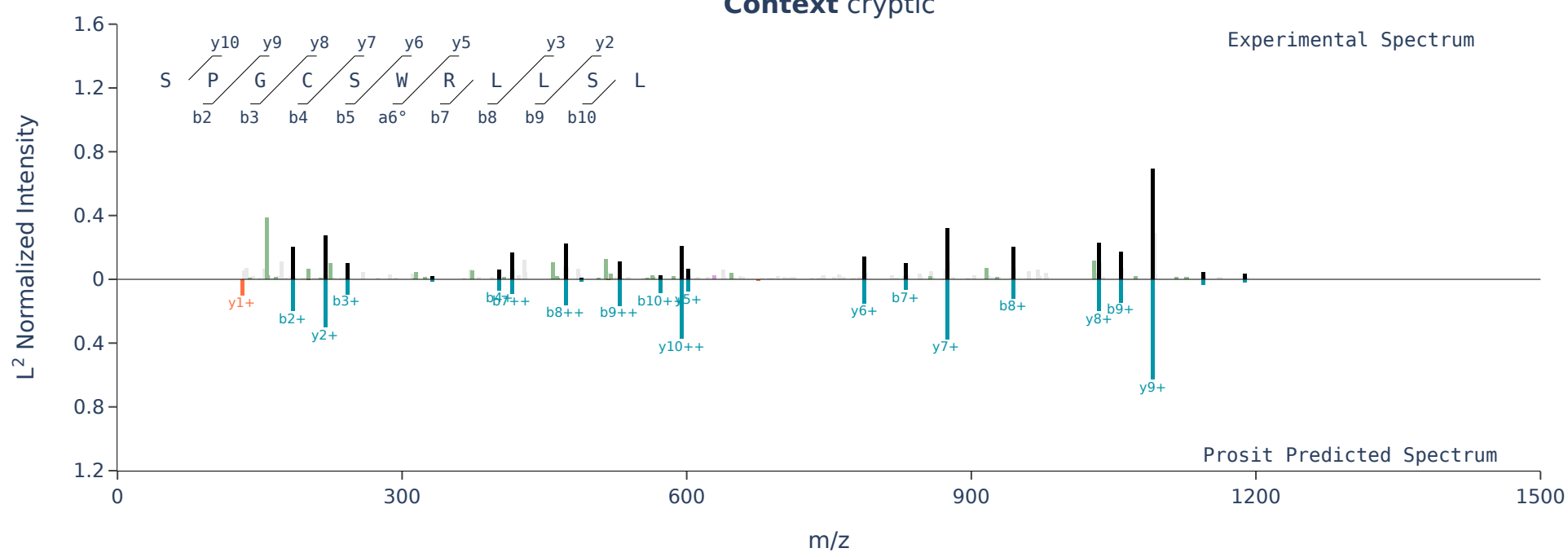
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Peptide RPNNLSLL Charge 2 Spectral Angle 0.84

Context multimapped



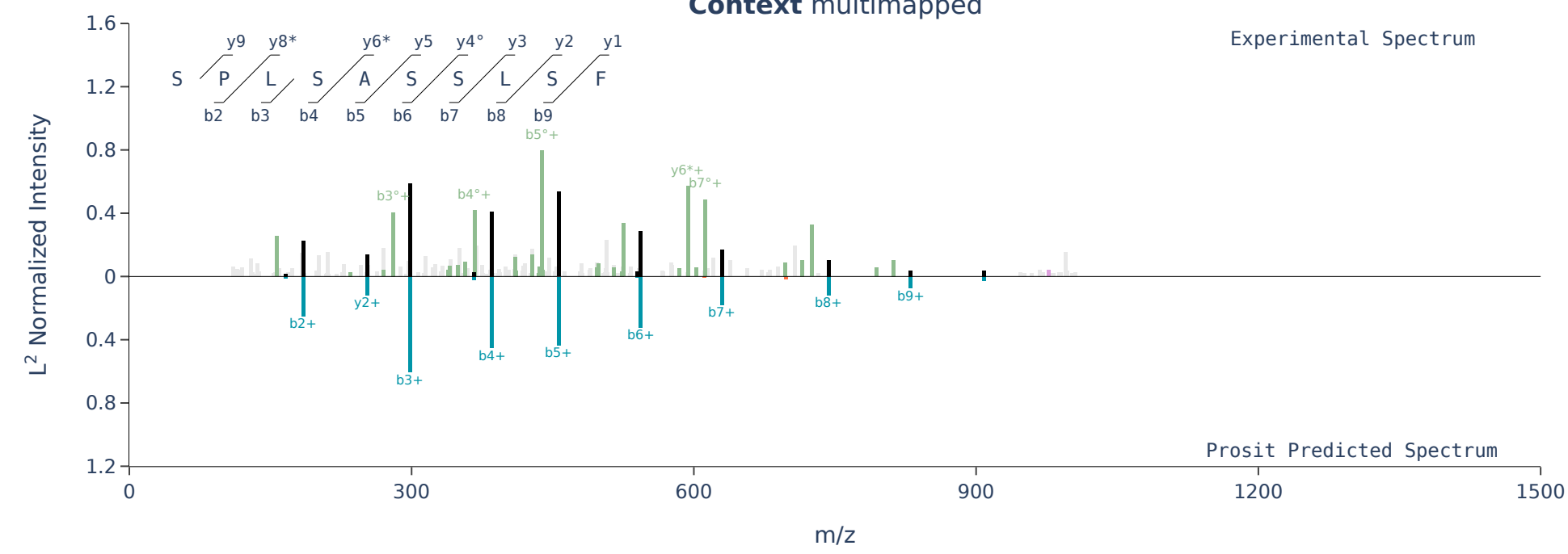
Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 25763
Peptide SPGC[+57.0]SWRLLSL Charge 2 Spectral Angle 0.83

Context cryptic



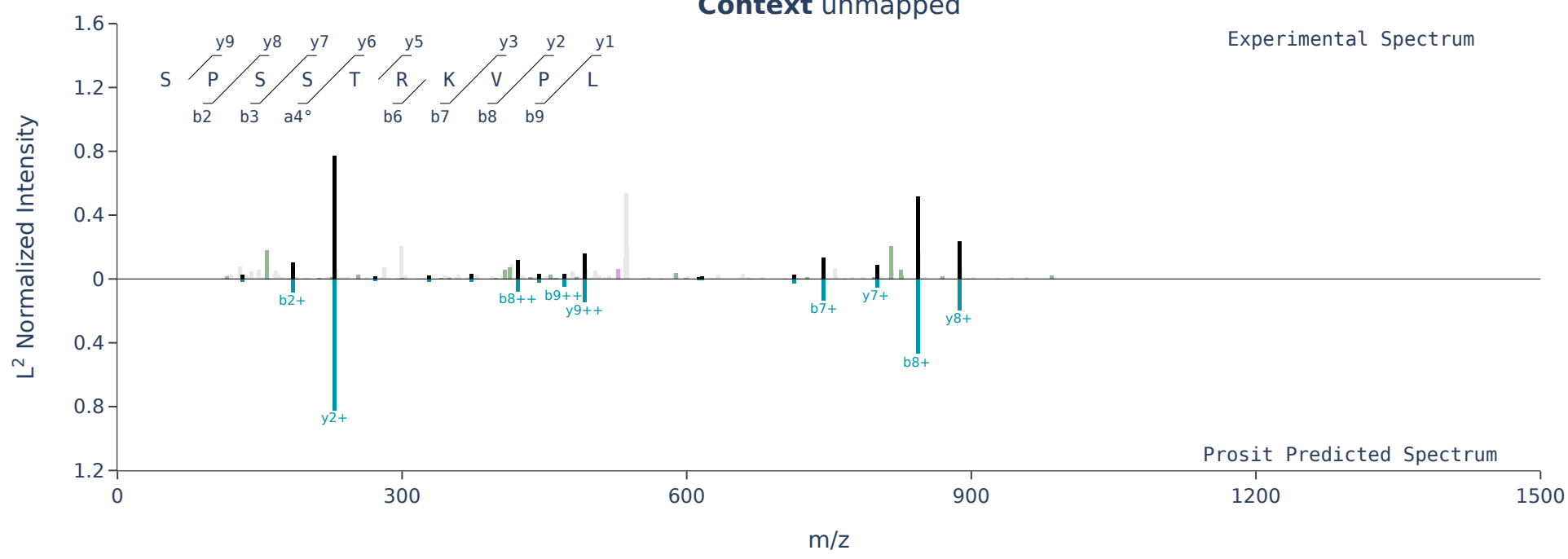
Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 23852
Peptide SPLSASSLSF Charge 1 Spectral Angle 0.92

Context multimapped



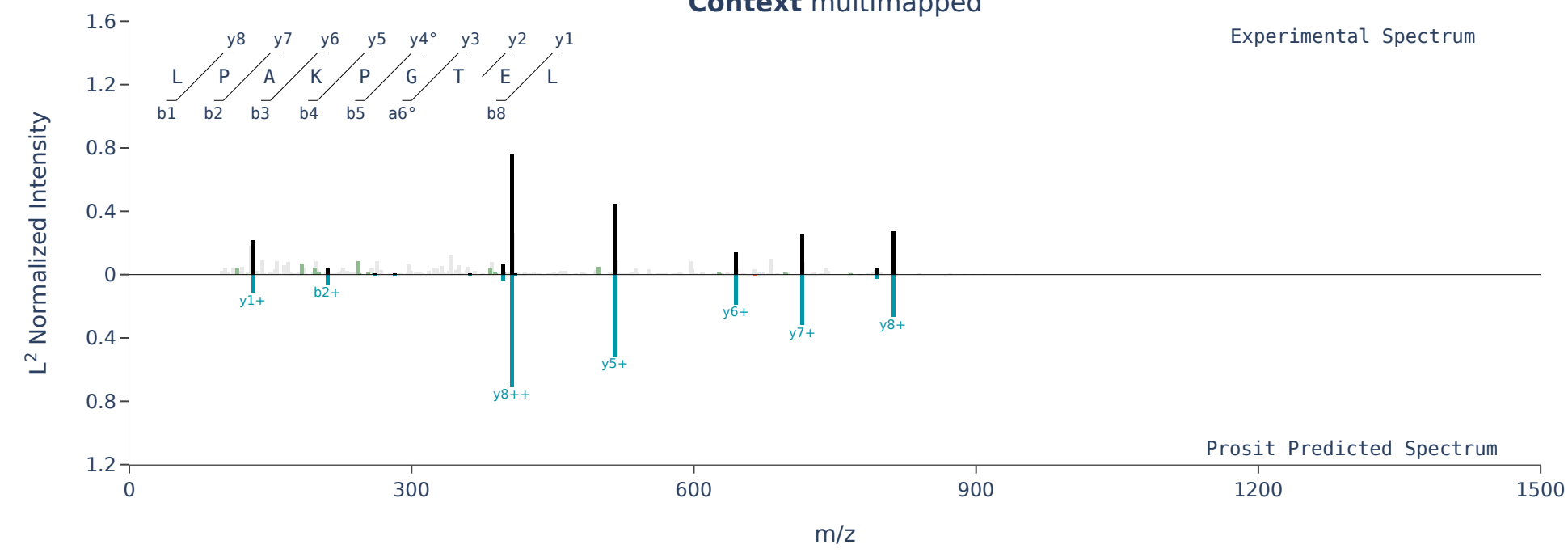
Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 7997
Peptide SPSSTRKVPL Charge 2 Spectral Angle 0.94

Context unmapped



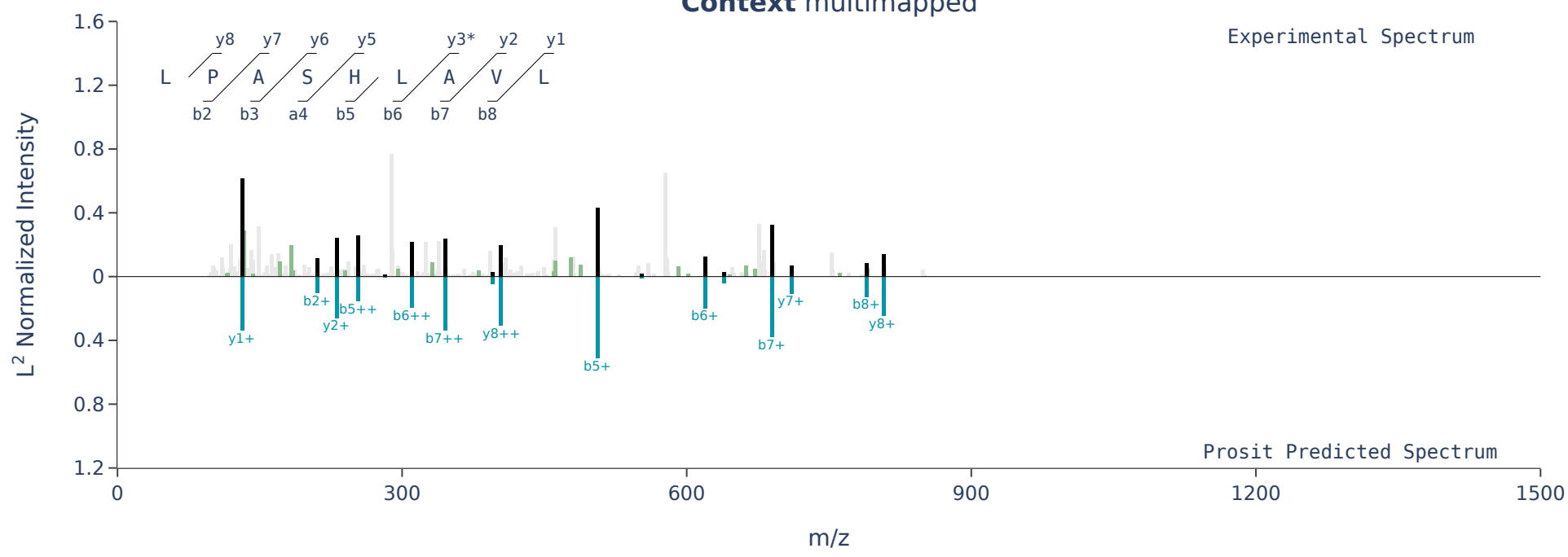
Source NCheng_Meghna_270923_270923_ExpI1_060923_K562B07_DB1_speedvac_M_R2 Scan 8800
Peptide LPAKPGTEL Charge 2 Spectral Angle 0.9

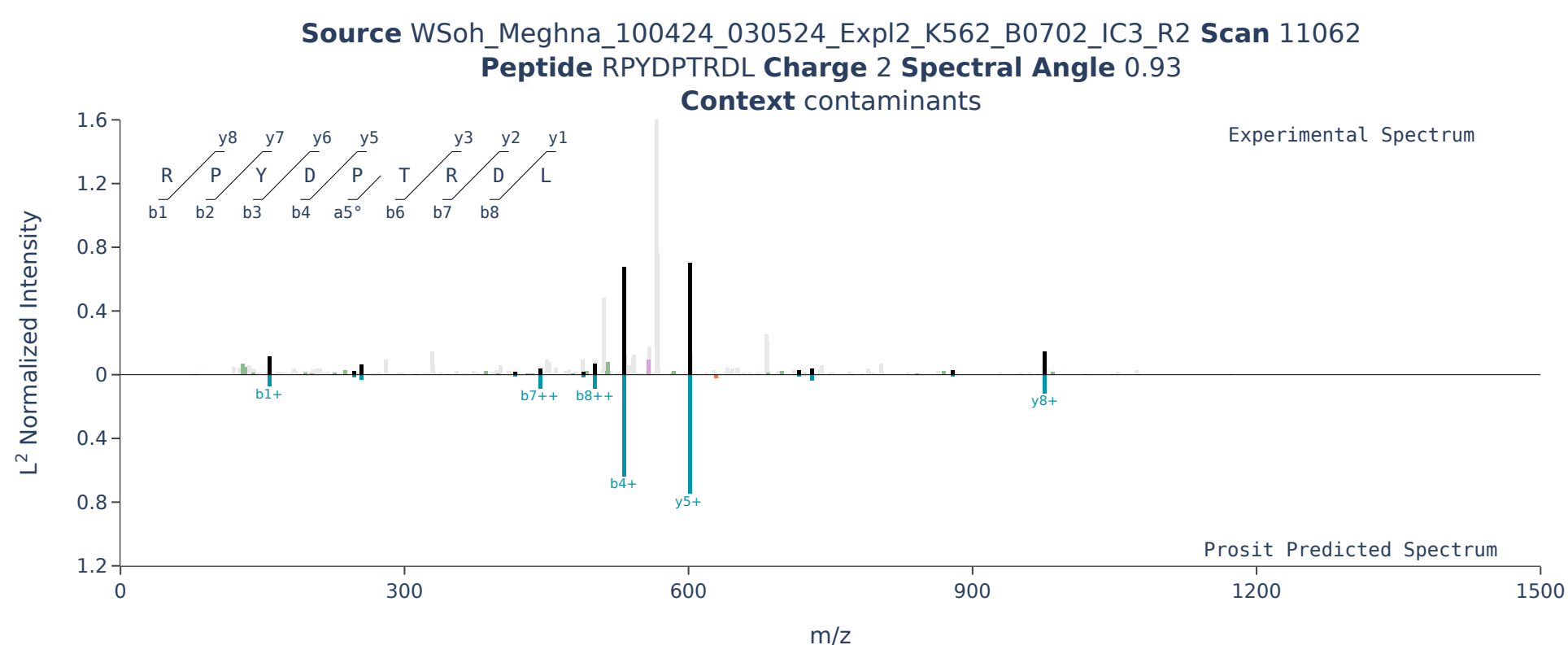
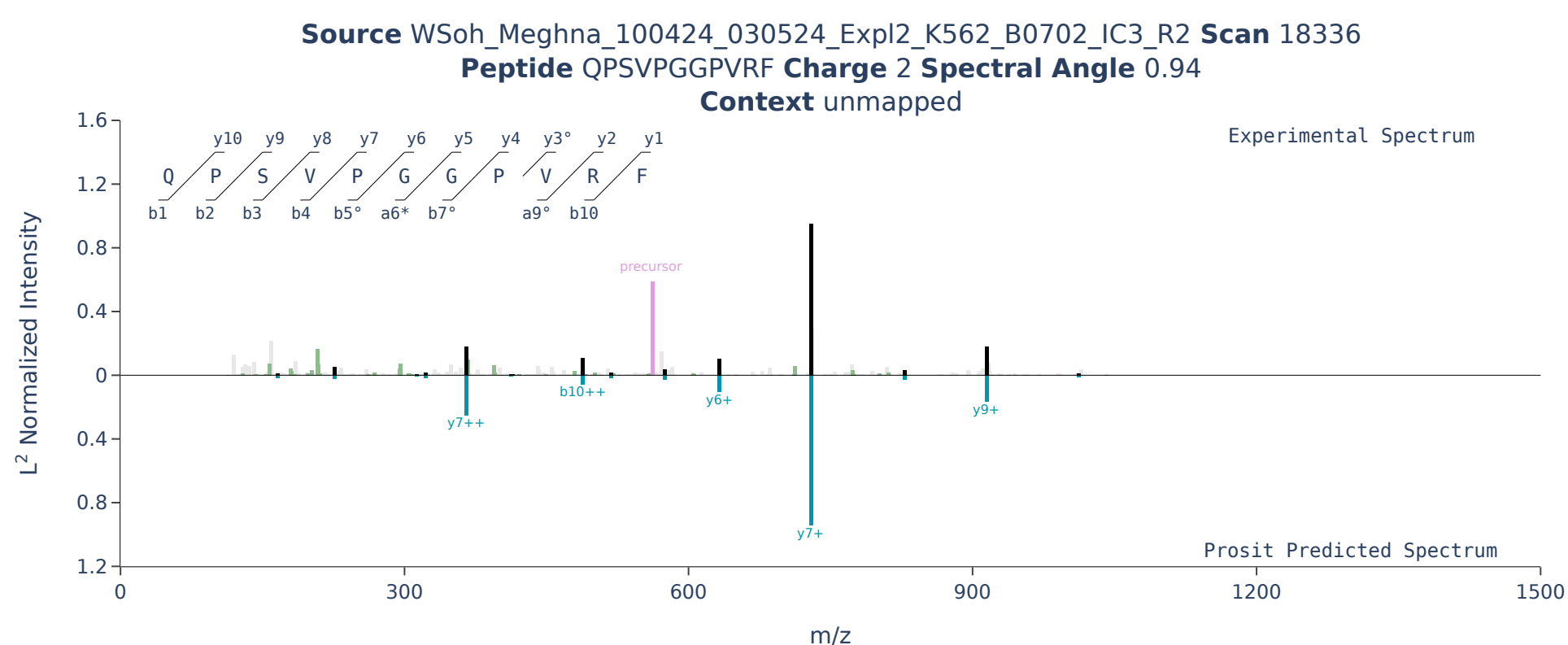
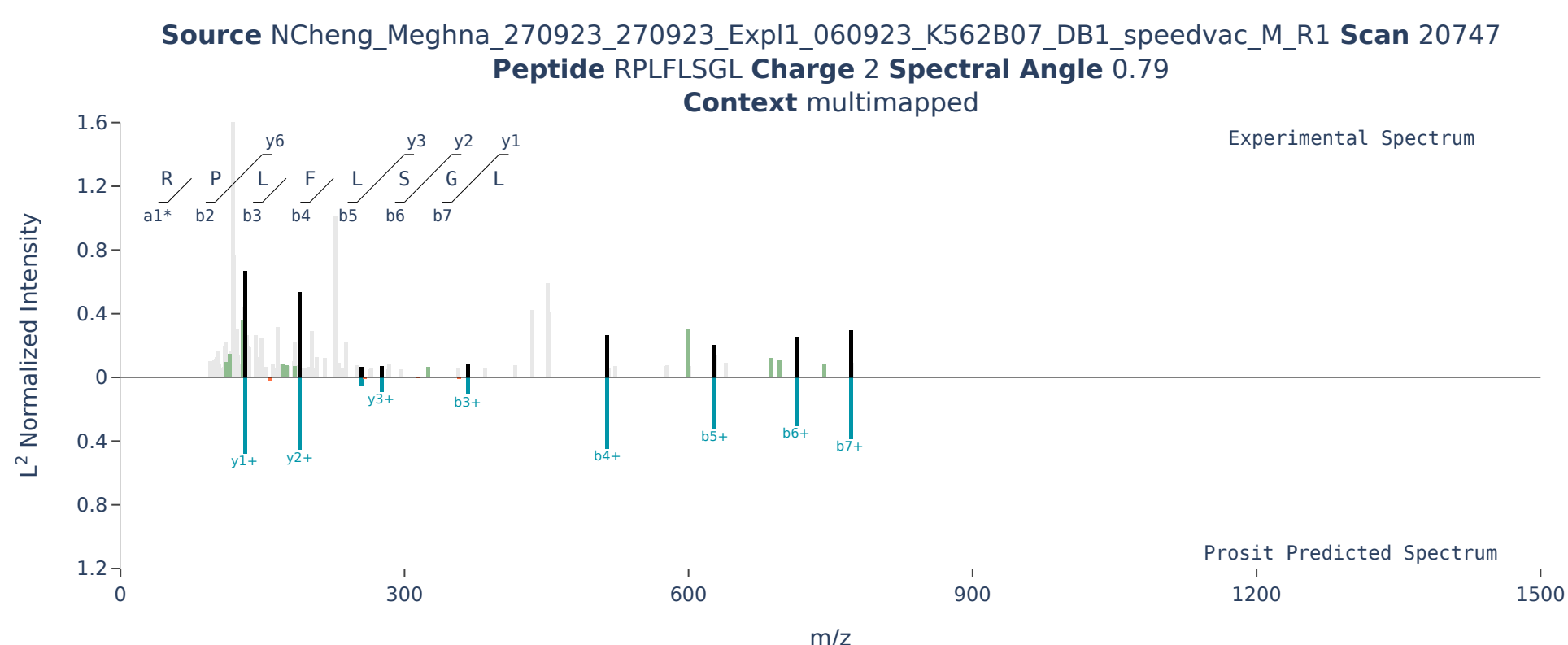
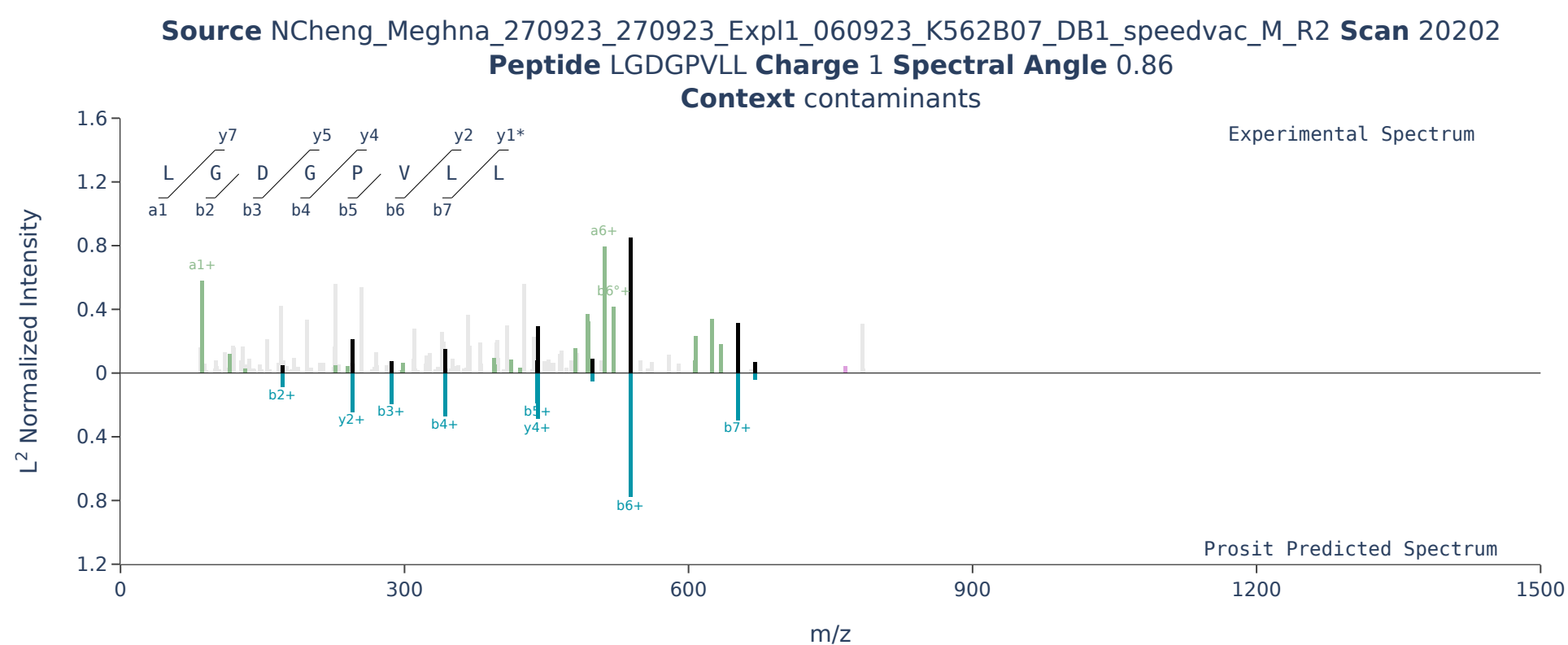
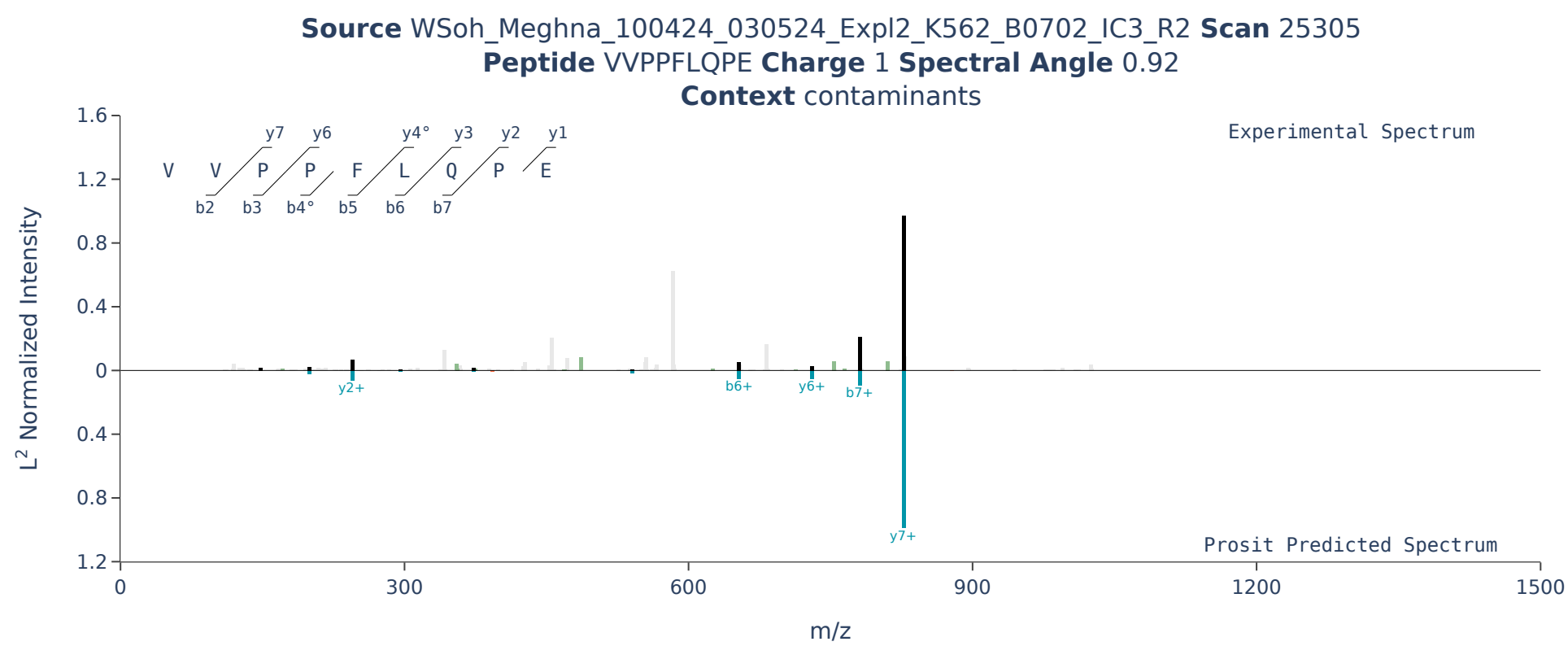
Context multimapped

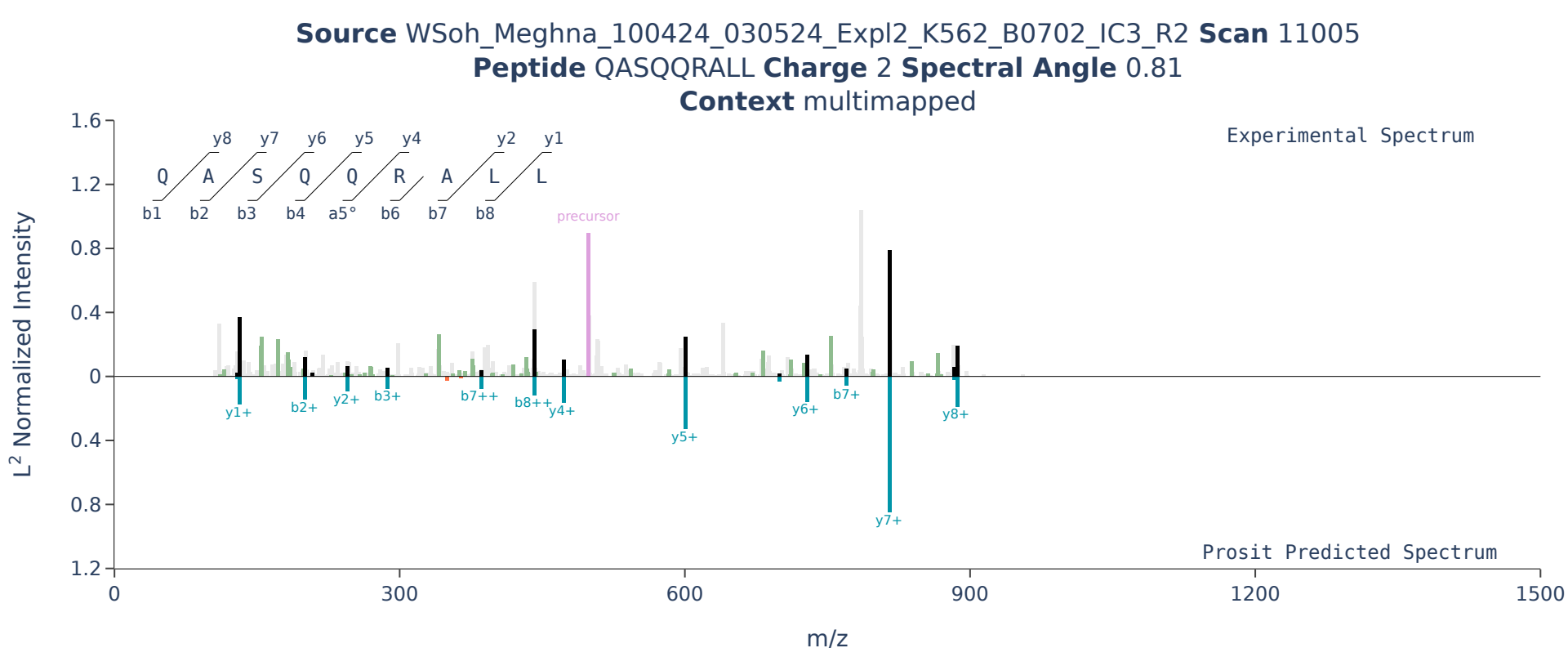
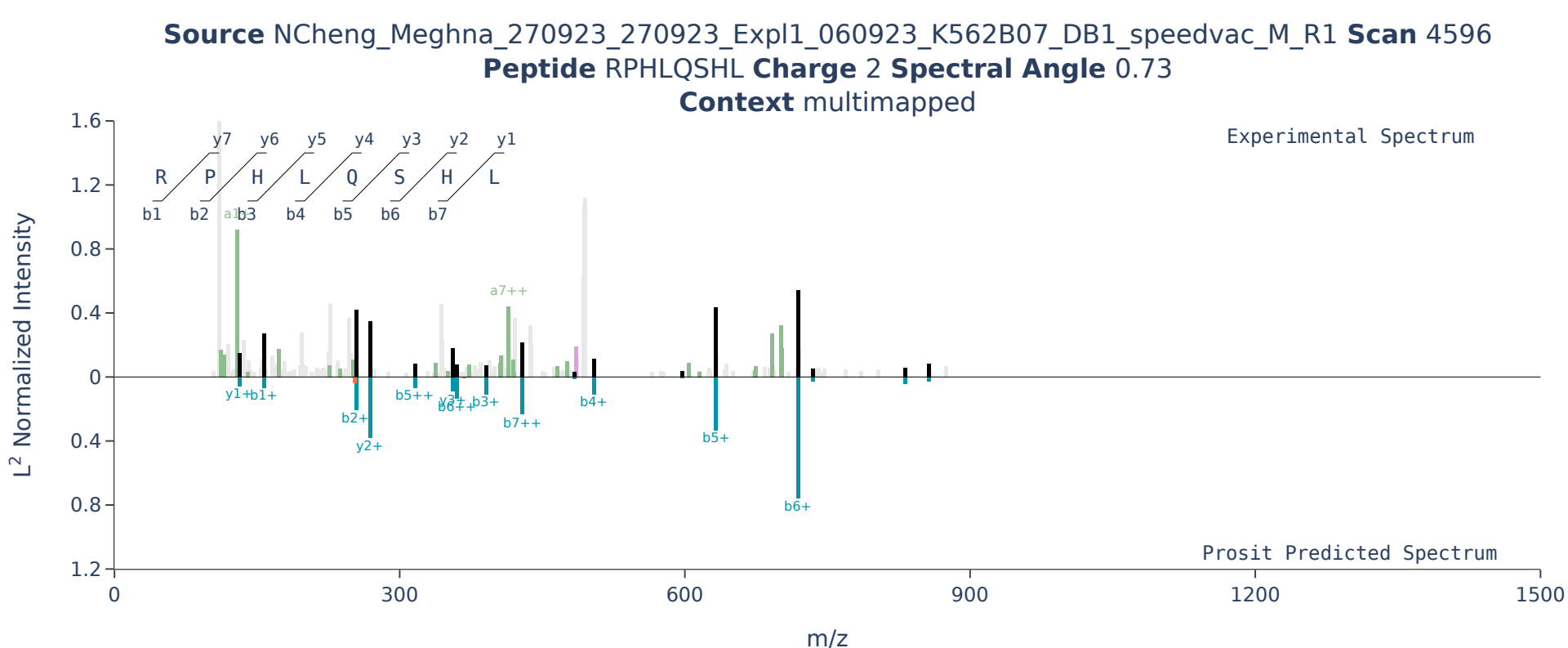
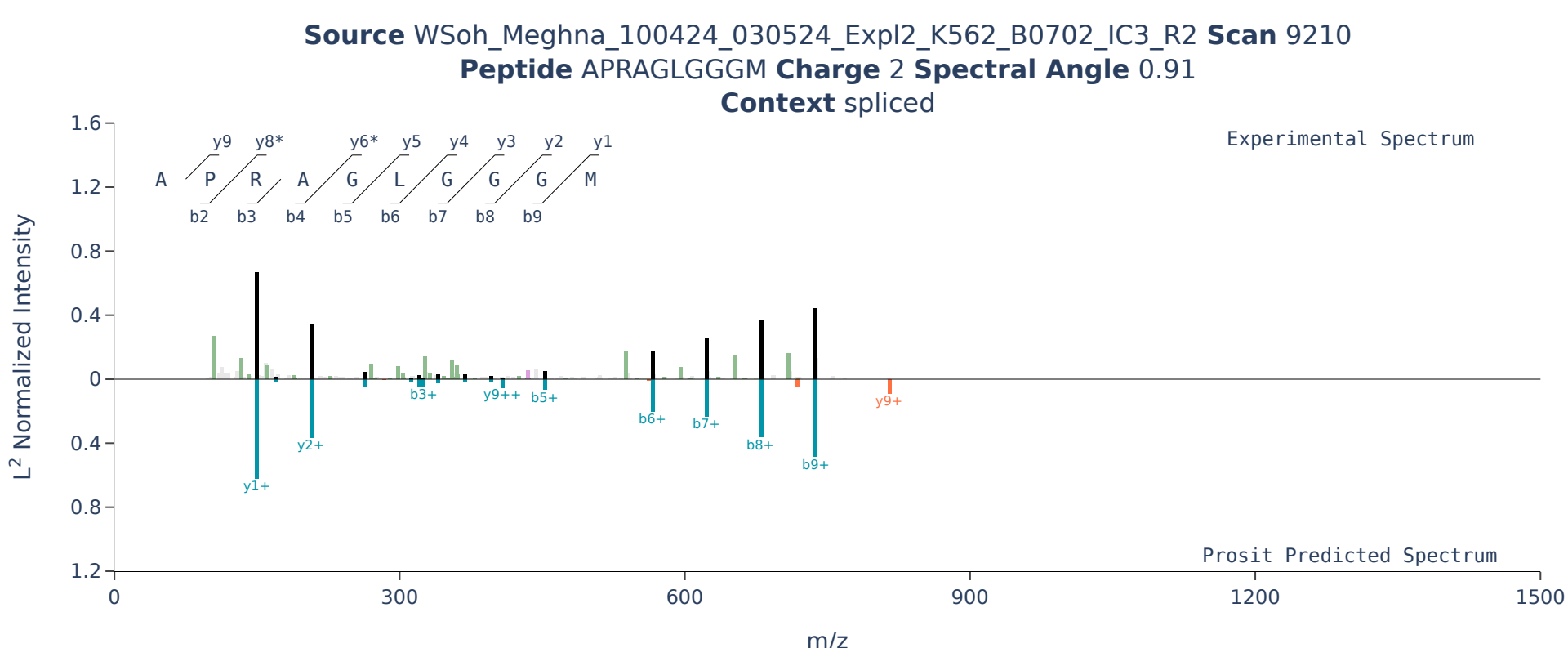
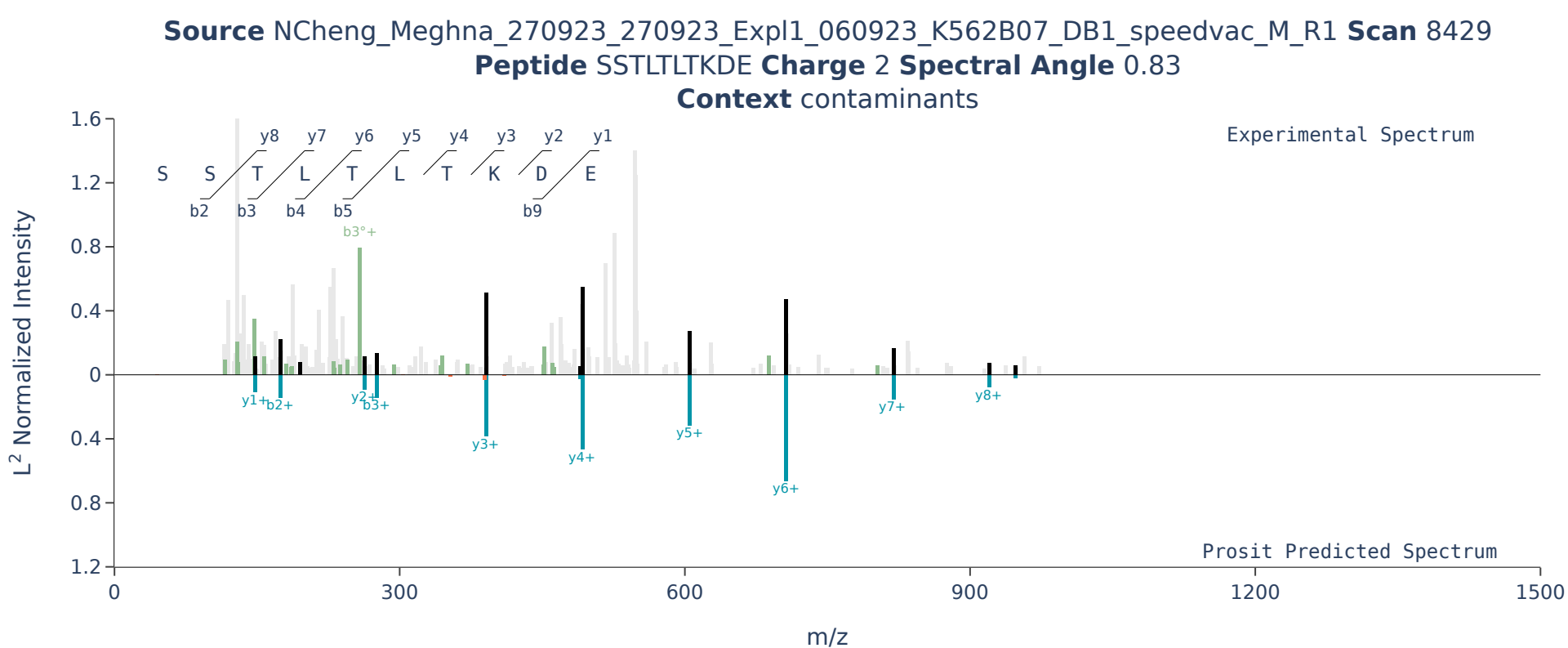
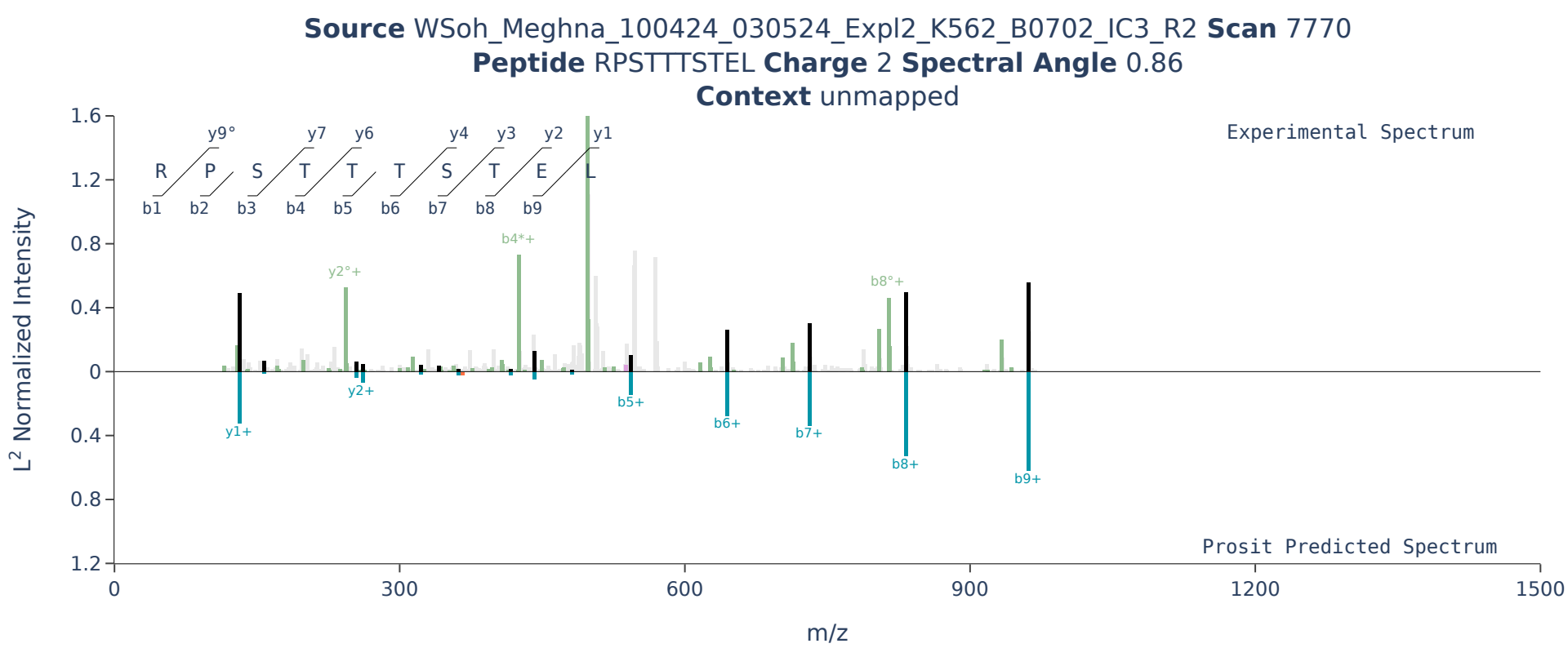


Source WSoh_Meghna_100424_030524_ExpI2_K562_B0702_IC3_R2 Scan 20031
Peptide LPASHLAVL Charge 2 Spectral Angle 0.76

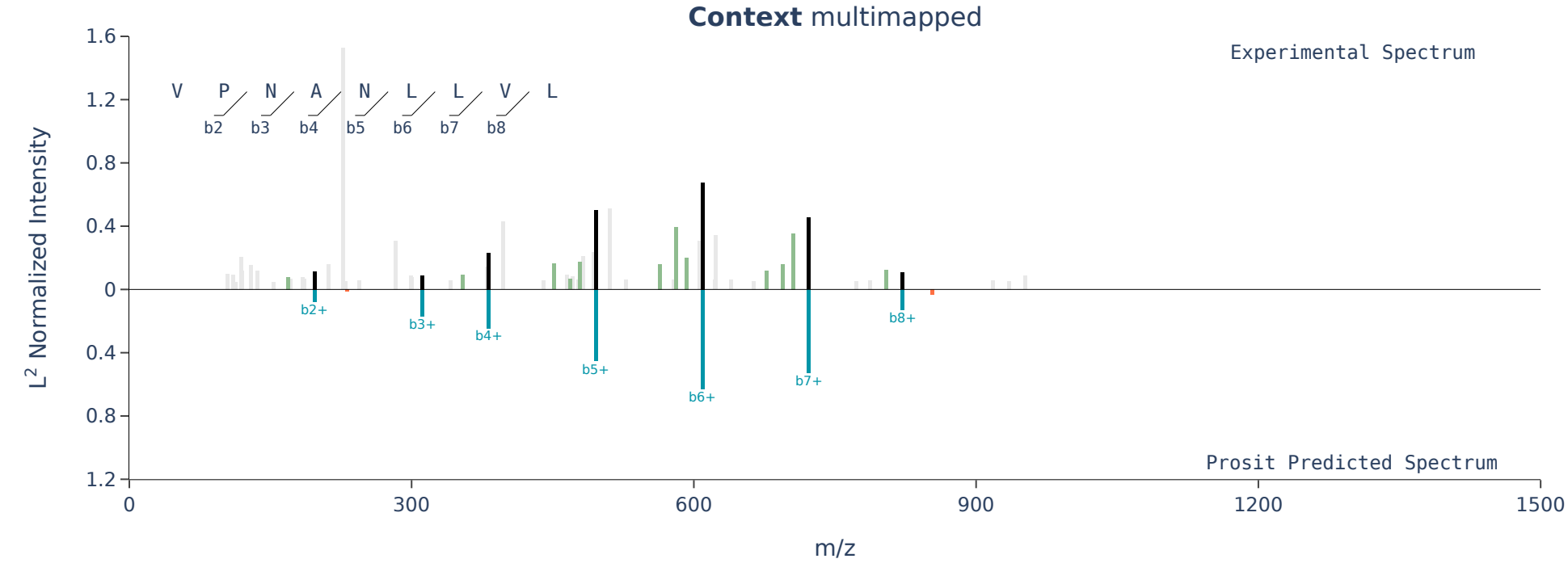
Context multimapped



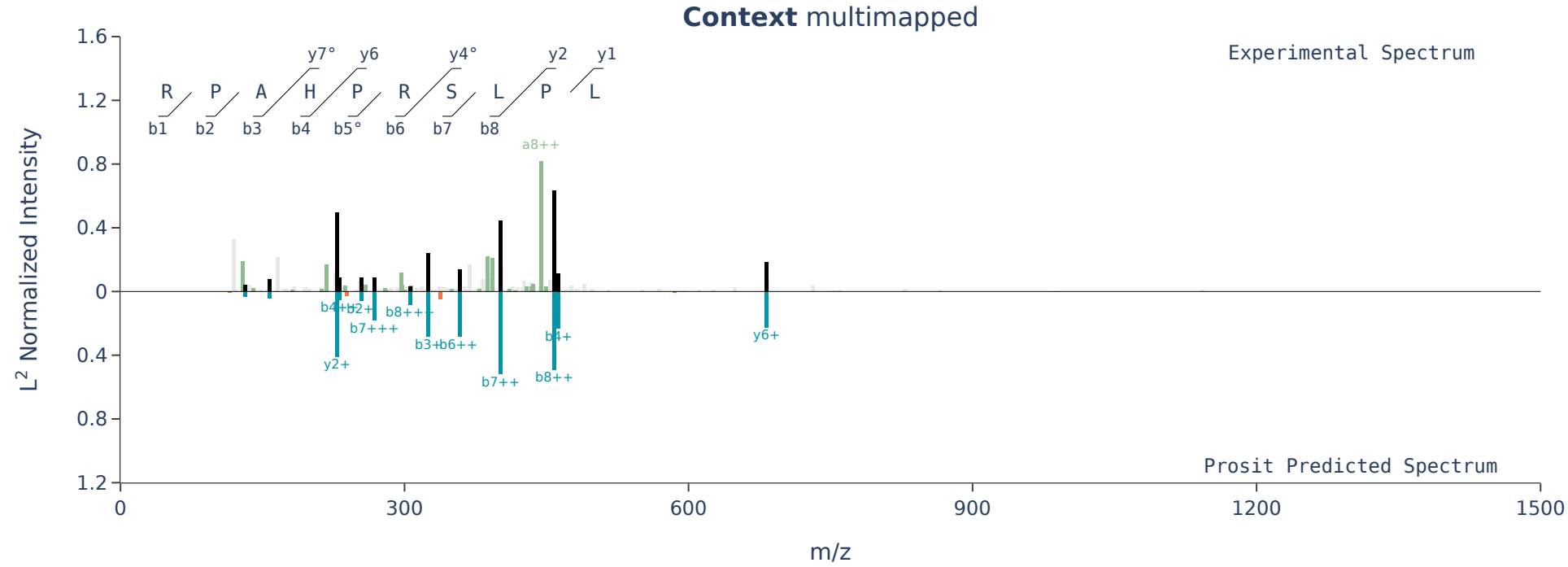




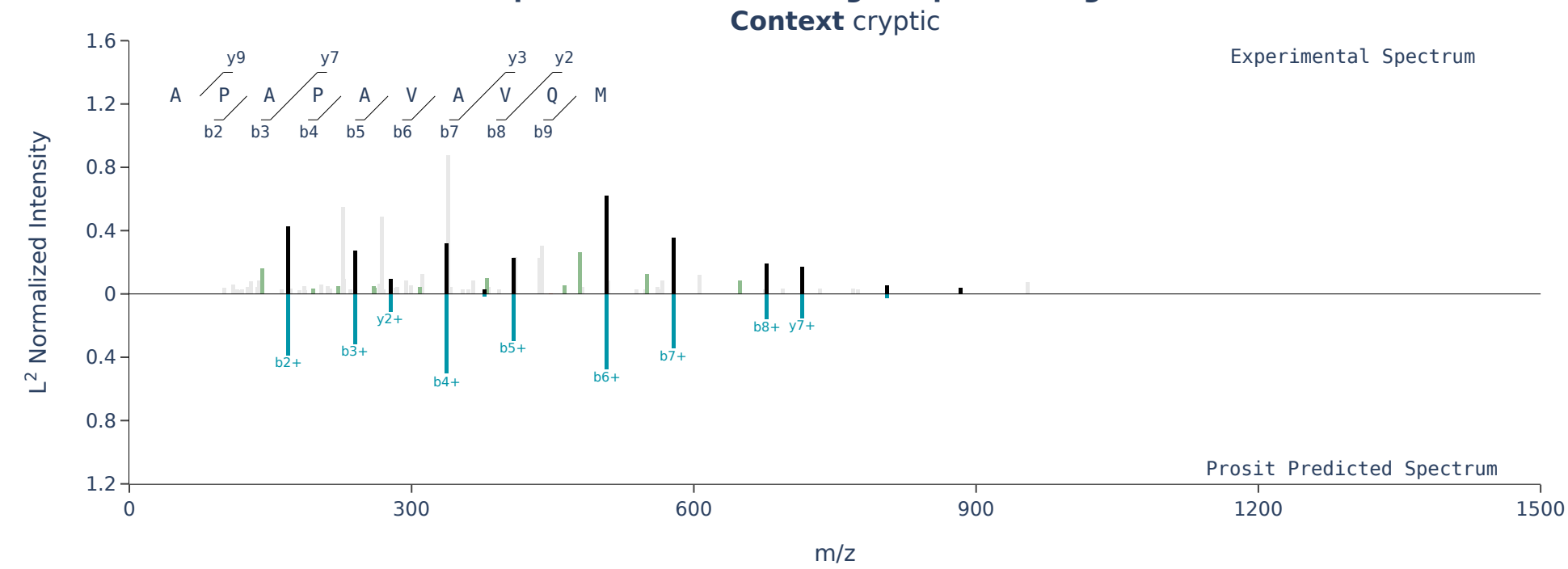
Source NCheng_Meghna_270923_270923_Exp11_060923_K562B07_DB1_speedvac_M_R2 **Scan** 23398
Peptide VPANANLLVL **Charge** 1 **Spectral Angle** 0.91



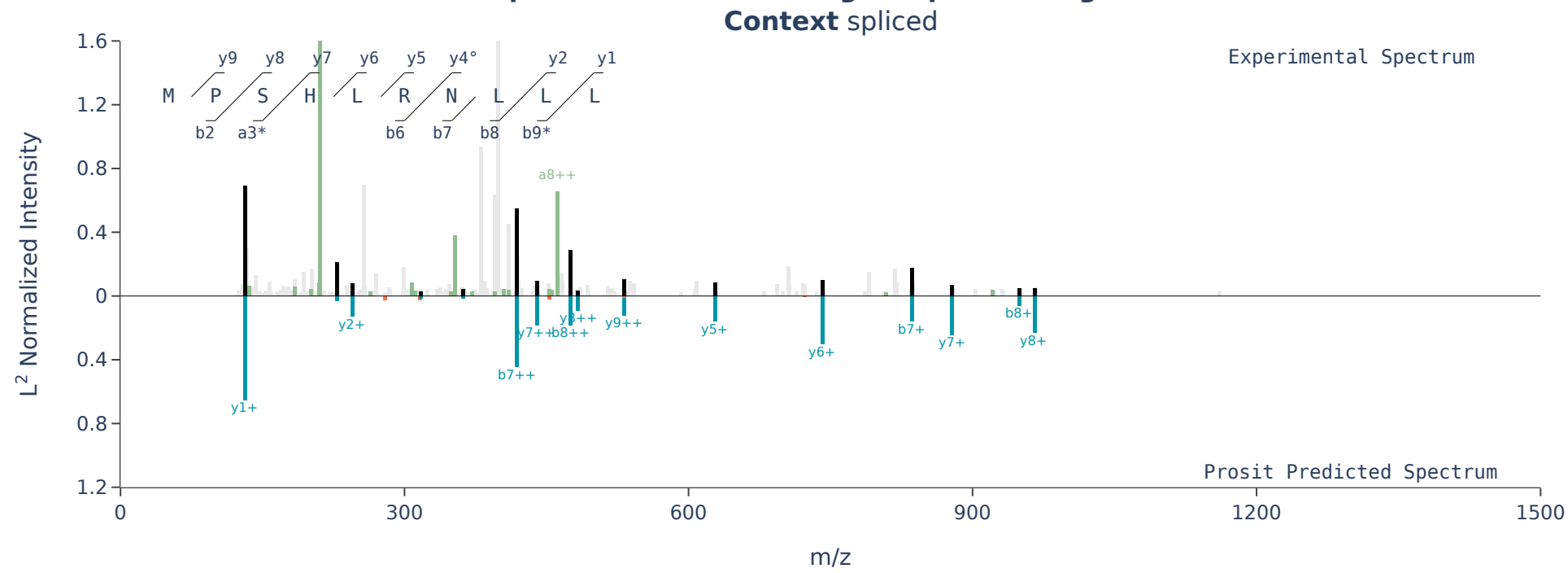
Source NCheng_Meghna_270923_270923_Exp1_060923_K562B07_DB1_speedvac_M_R1 **Scan** 7061
Peptide RPAHPRSLPL **Charge** 3 **Spectral Angle** 0.81



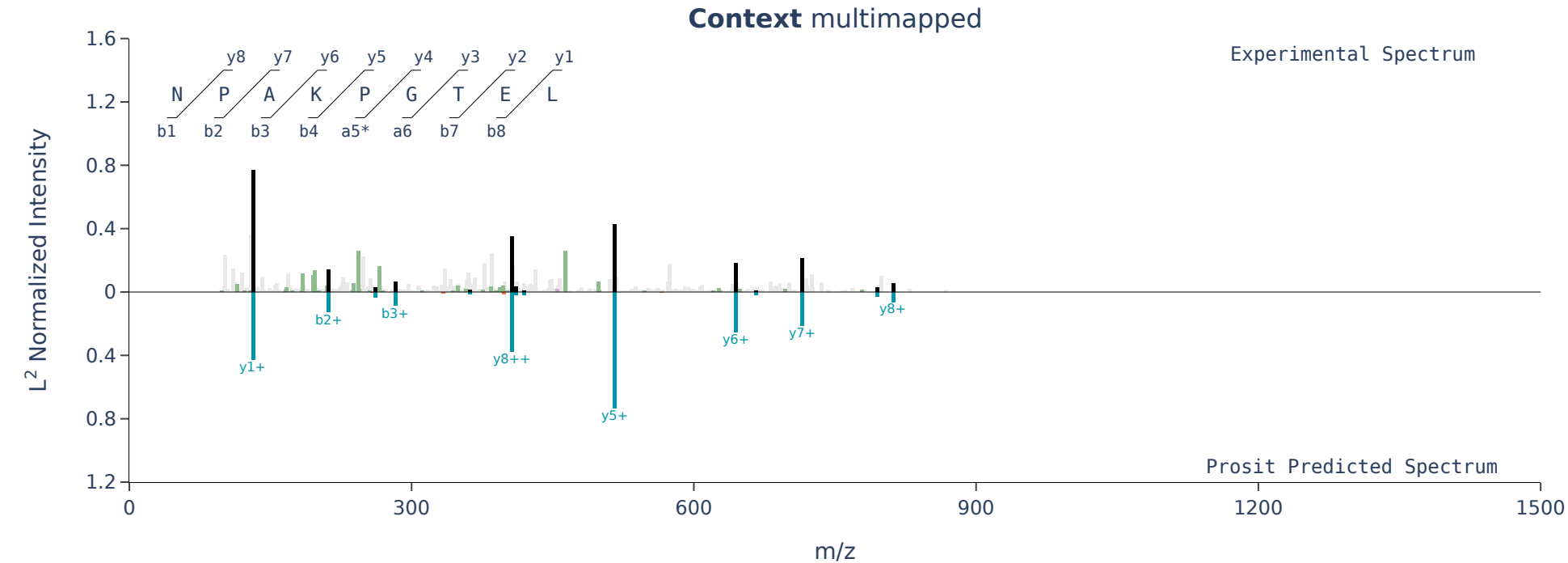
Source NCheng_Meghna_270923_270923_Exp11_060923_K562B07_DB1_speedvac_M_R2 **Scan** 15464
Peptide APAPAVAVQM **Charge** 1 **Spectral Angle** 0.84



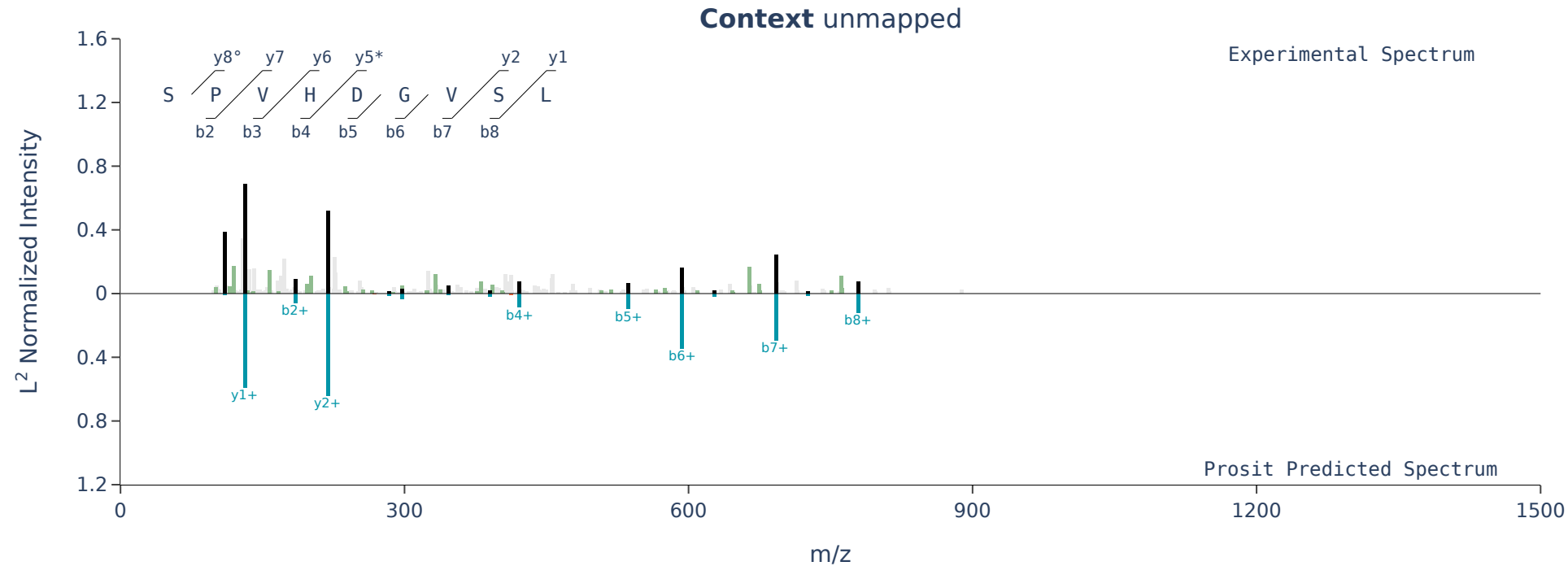
Source WSoh_Meghna_100424_030524_Exp12_K562_B0702_IC3_R2 **Scan** 19893
Peptide MPSHLRNLLL **Charge** 3 **Spectral Angle** 0.73



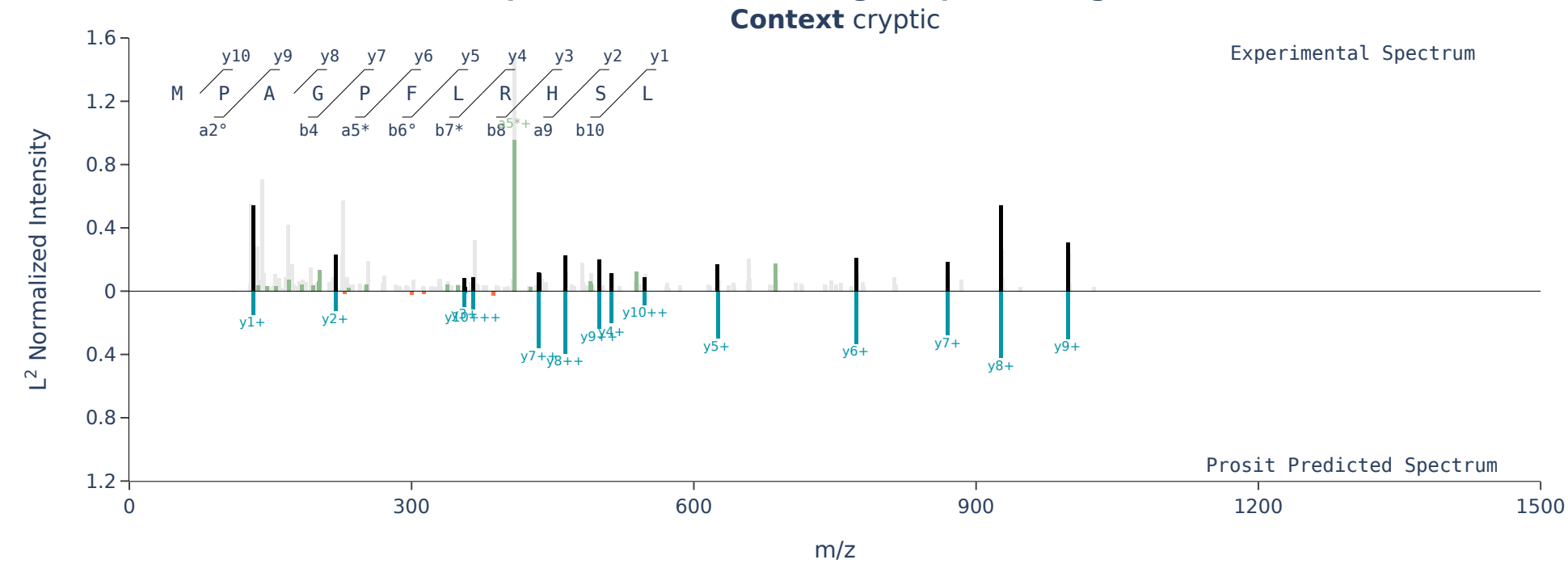
Source NCheng_Meghna_270923_270923_Exp1_060923_K562B07_DB1_speedvac_M_R2 **Scan** 5942
Peptide NPAKPGTEL **Charge** 2 **Spectral Angle** 0.7



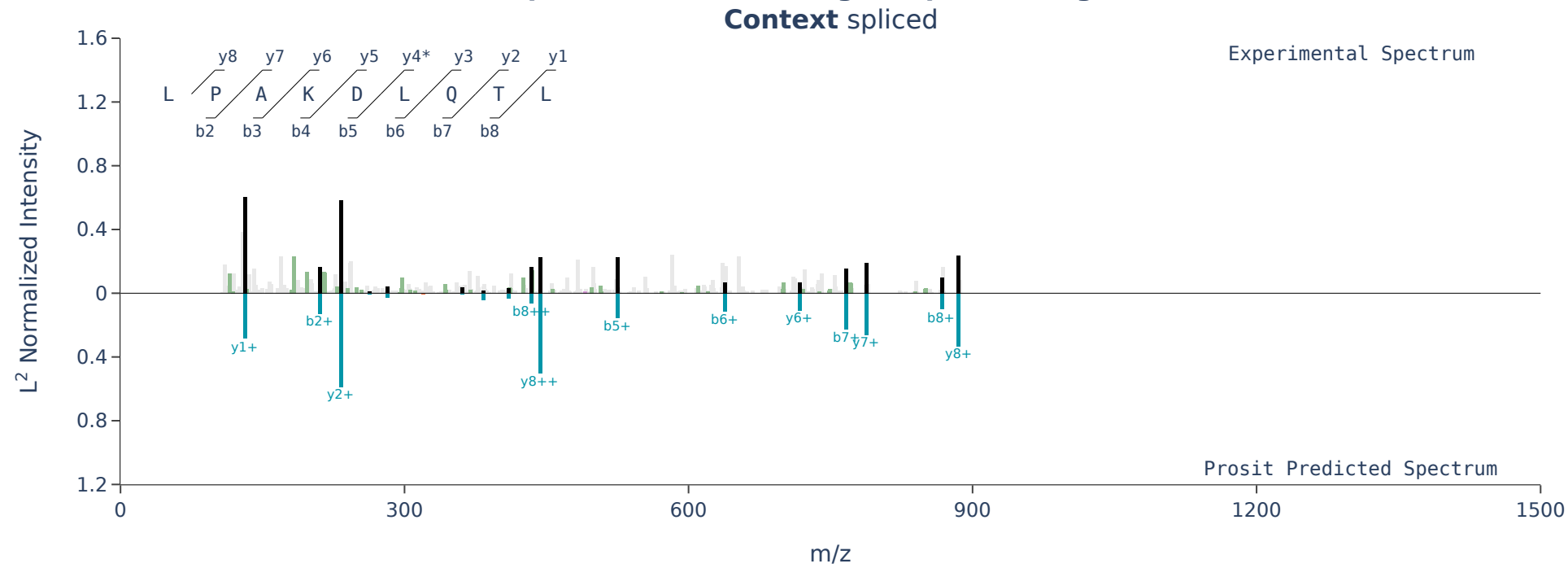
Source NCheng_Meghna_270923_270923_Exp1_060923_K562B07_DB1_speedvac_M_R1 **Scan** 9956
Peptide SPVHDGVSL **Charge** 2 **Spectral Angle** 0.84



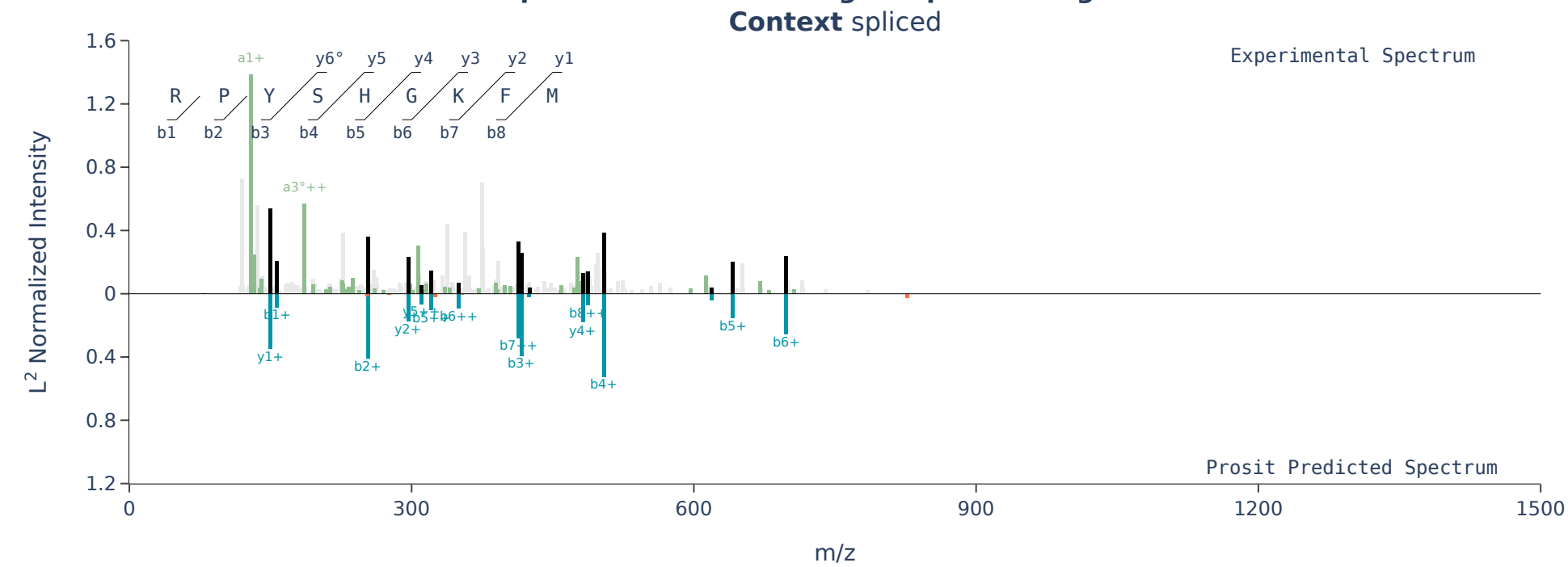
Source NCheng_Meghna_270923_270923_ExpI1_060923_K562B07_DB1_speedvac_M_R2 **Scan** 15364
Peptide MPAGPFLRHSL **Charge** 3 **Spectral Angle** 0.63



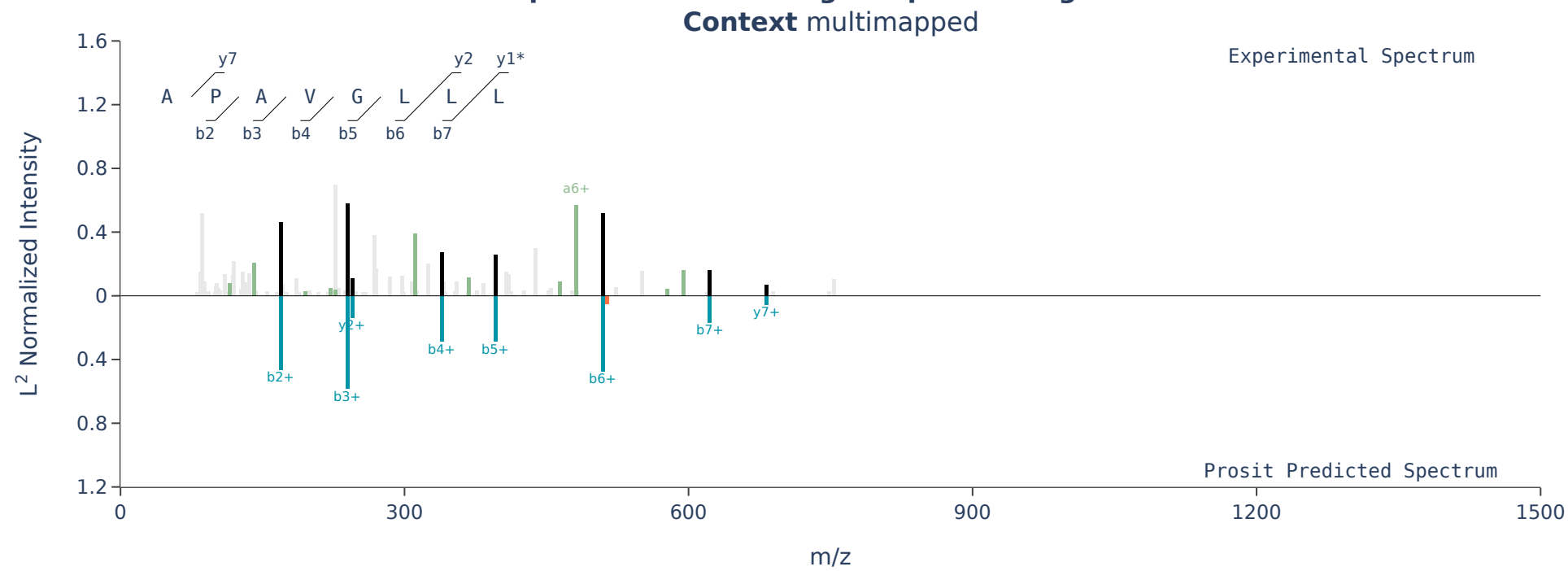
Source NCheng_Meghna_270923_270923_Exp11_060923_K562B07_DB1_speedvac_M_R1 **Scan** 11946
Peptide LPAKDLQTL **Charge** 2 **Spectral Angle** 0.7

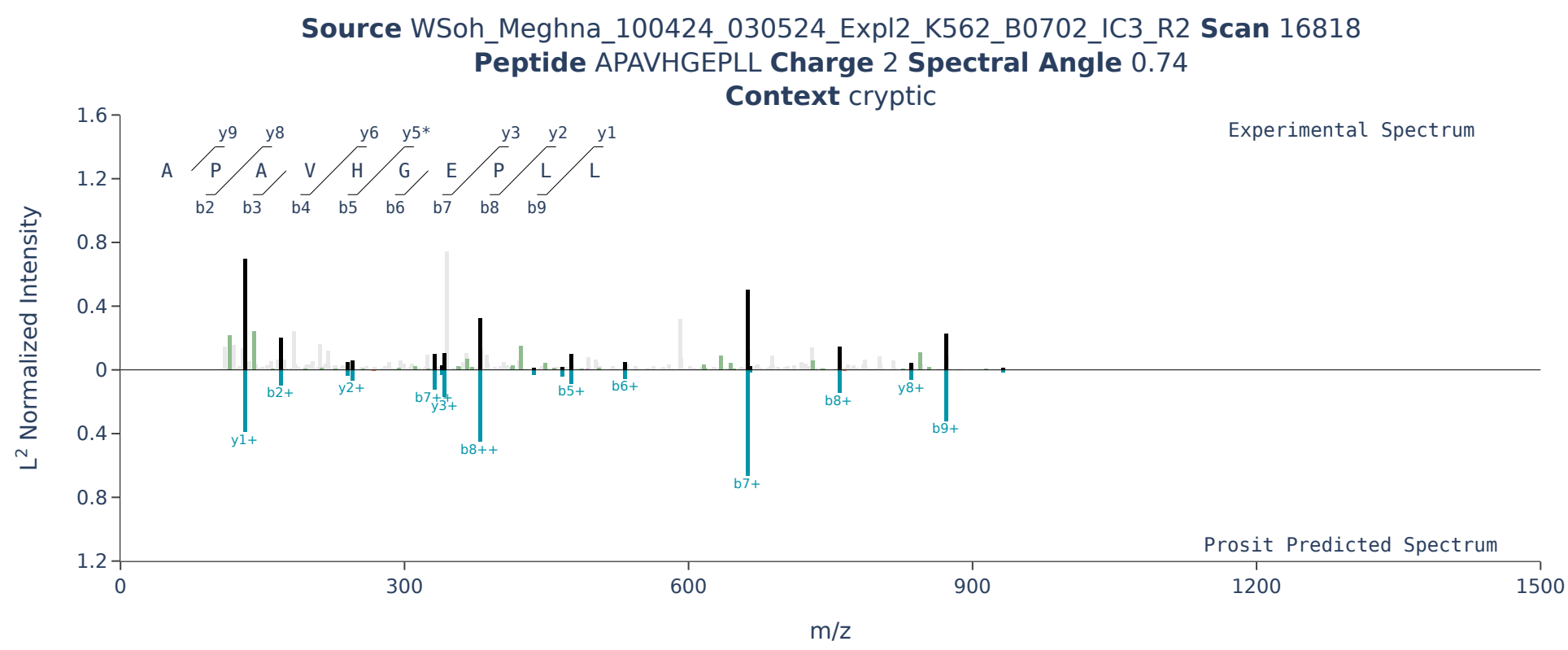
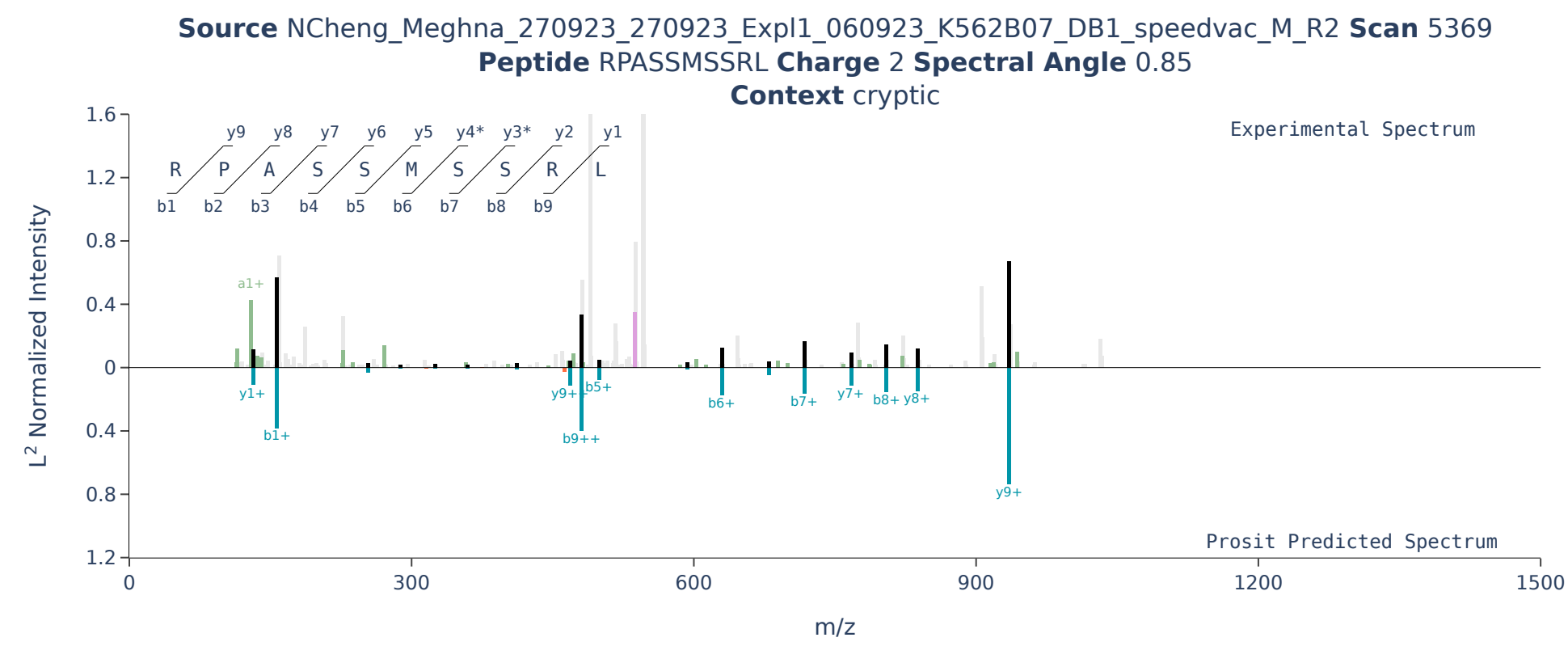
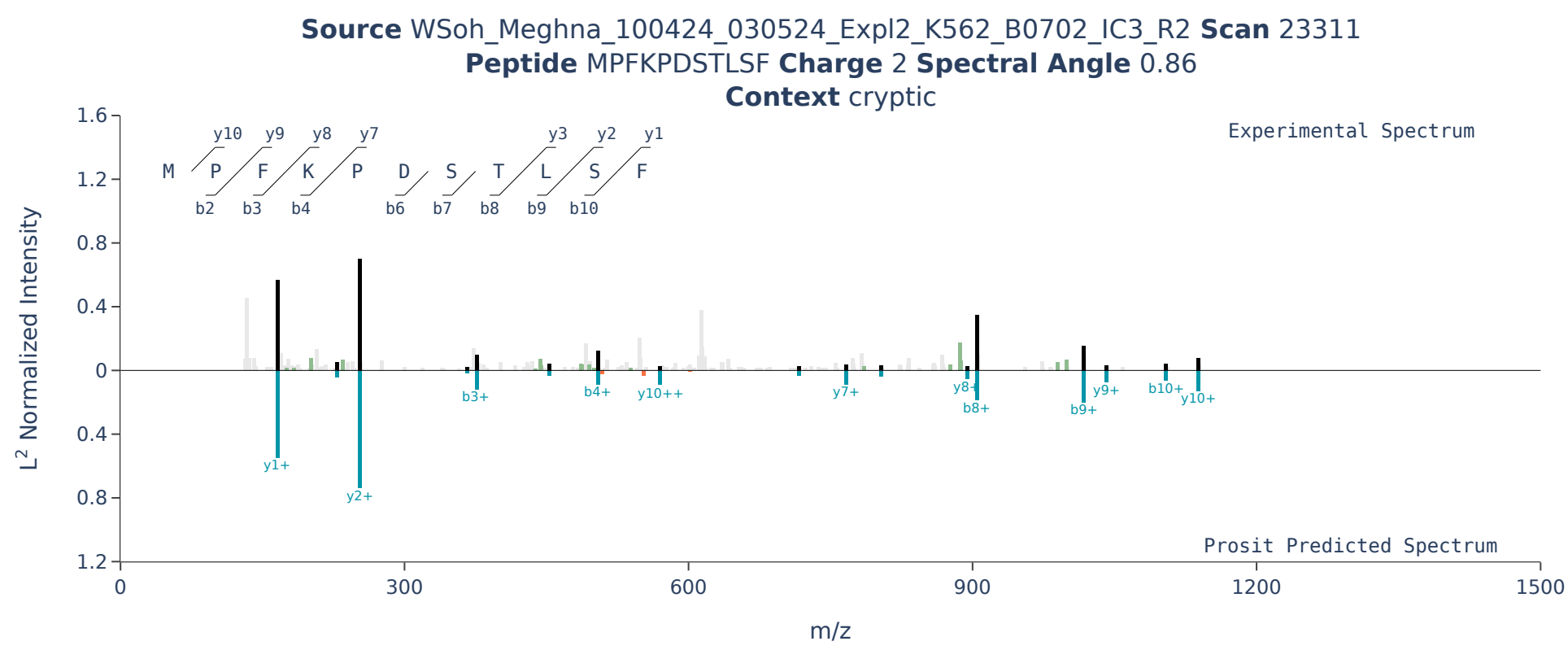
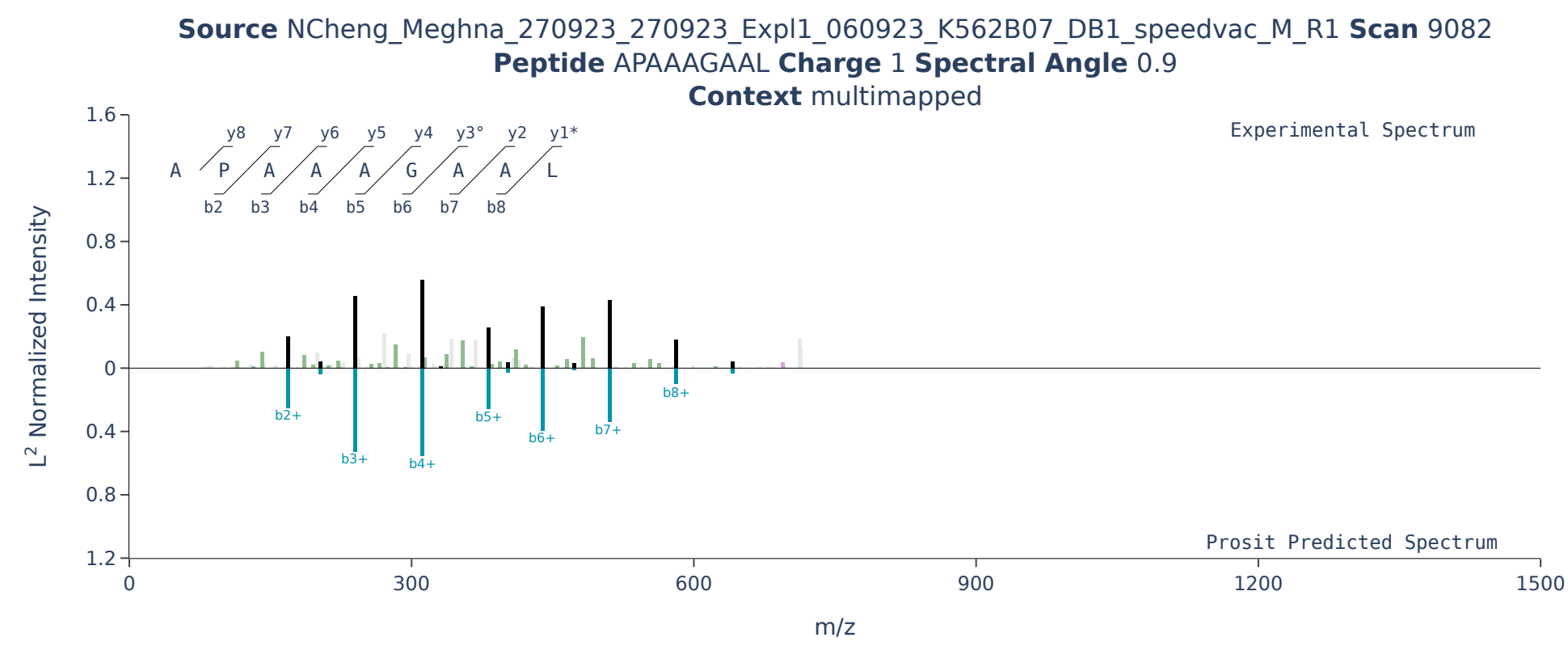
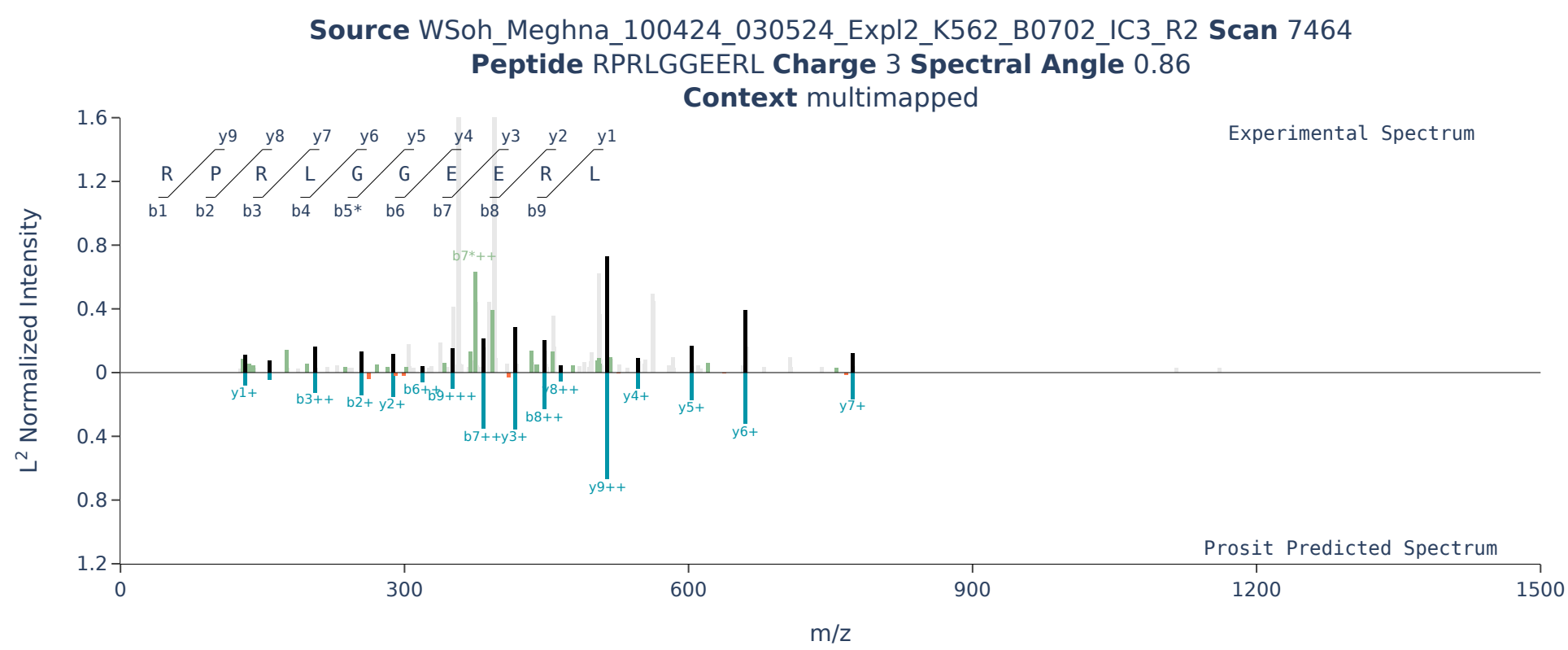
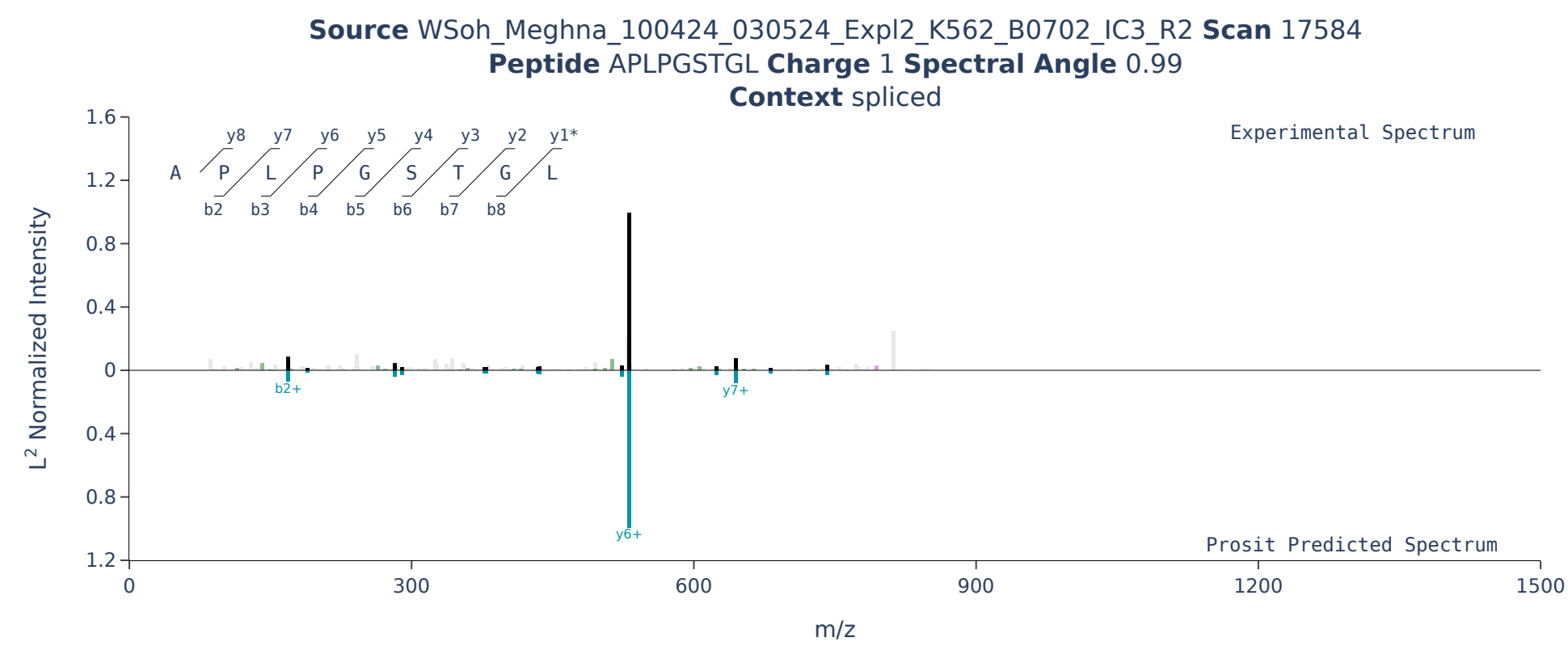
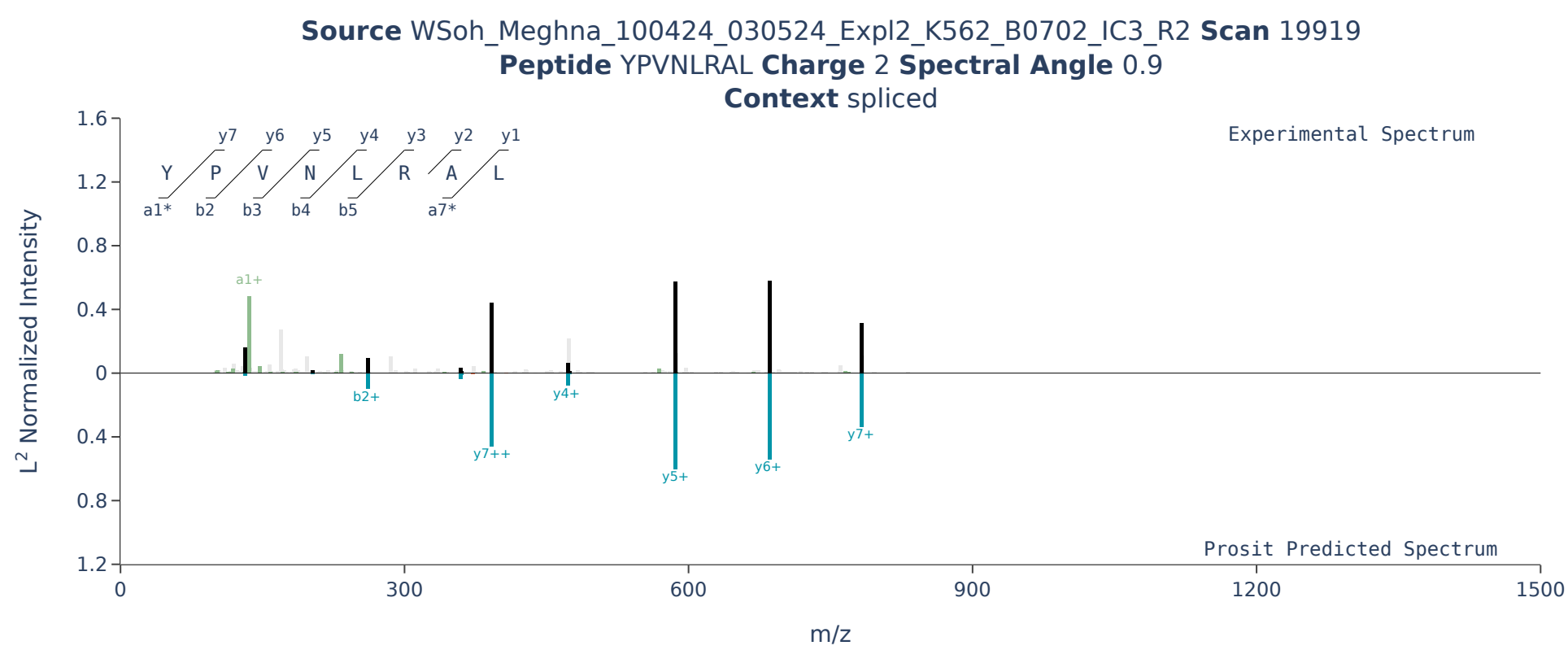
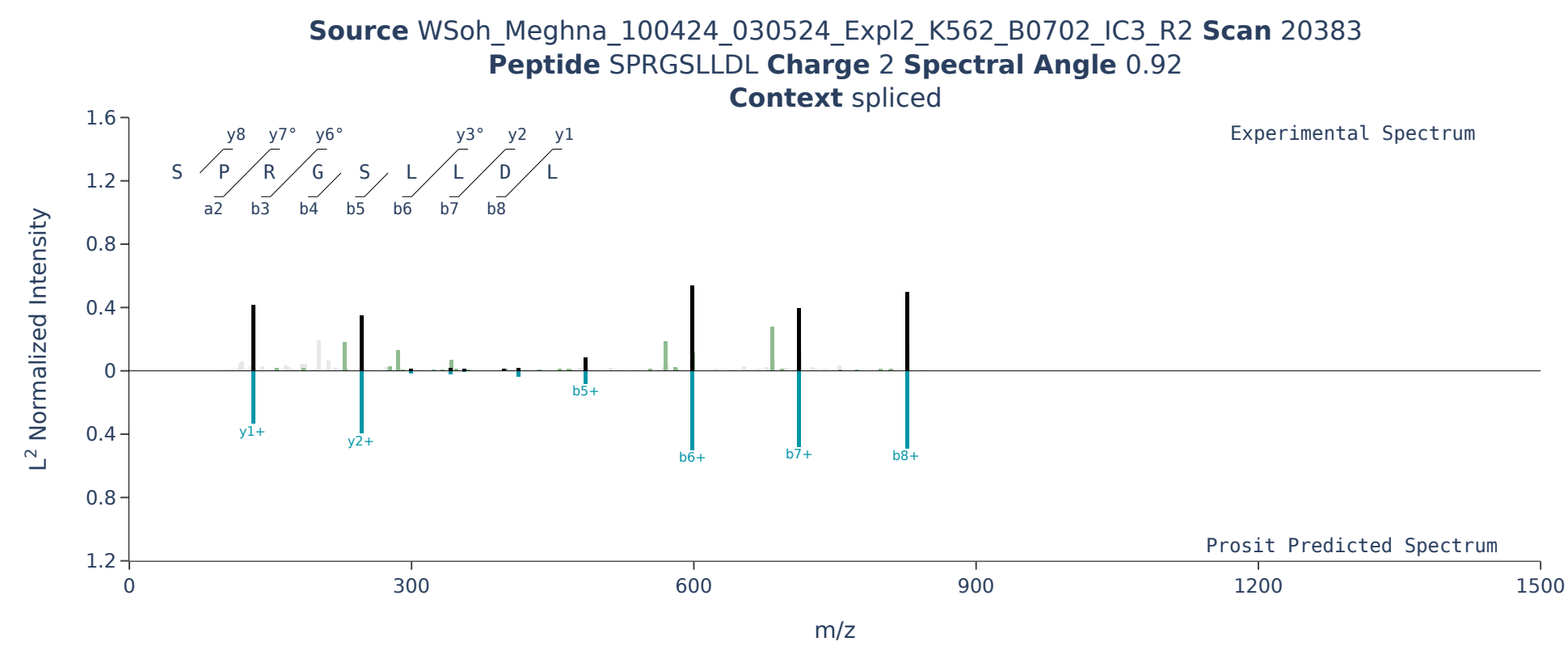
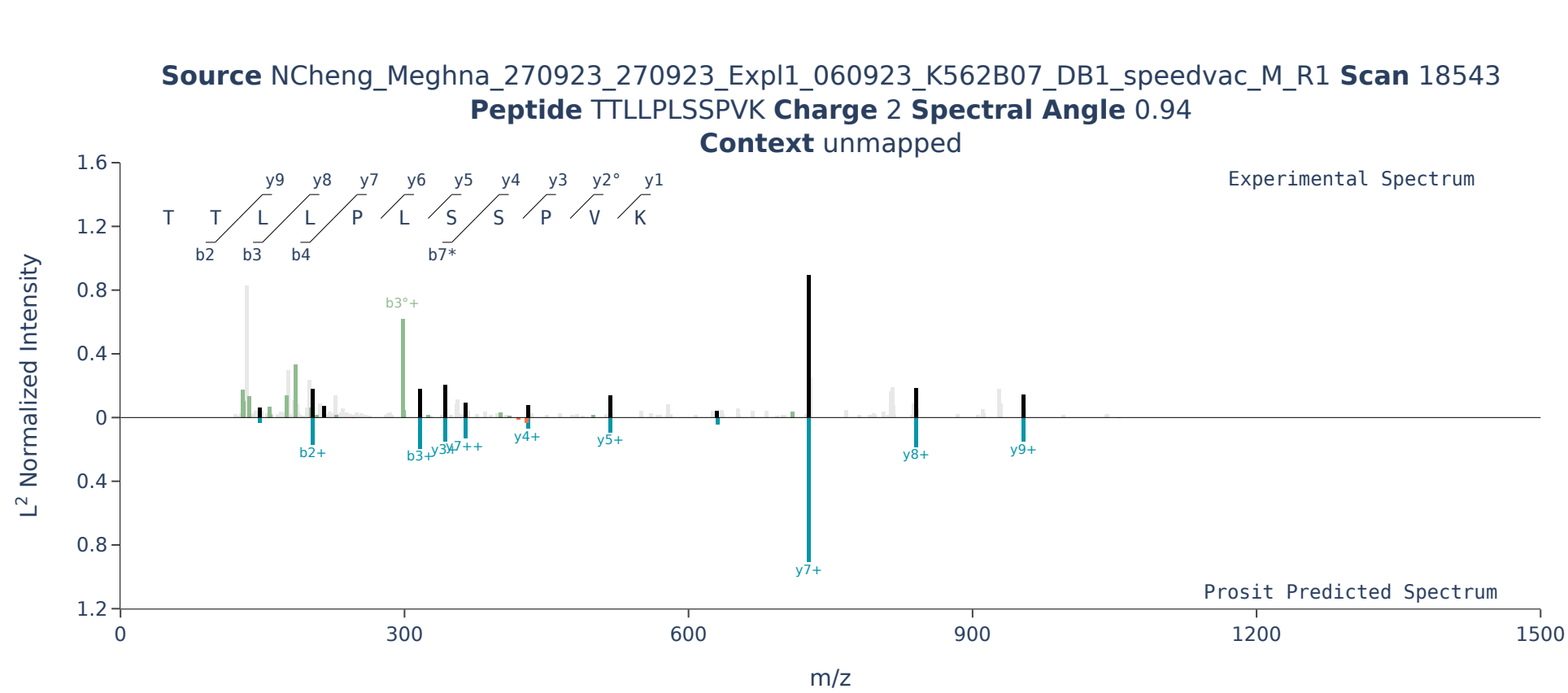
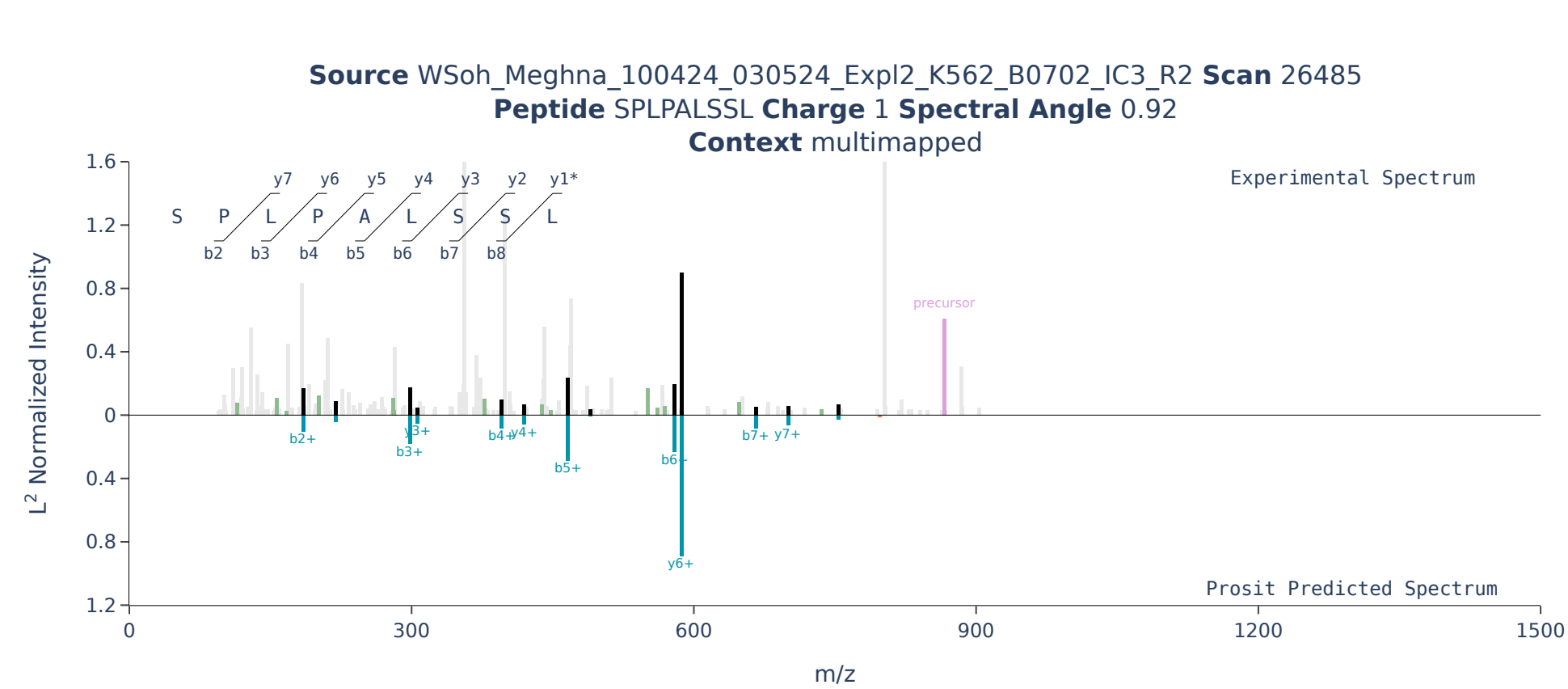


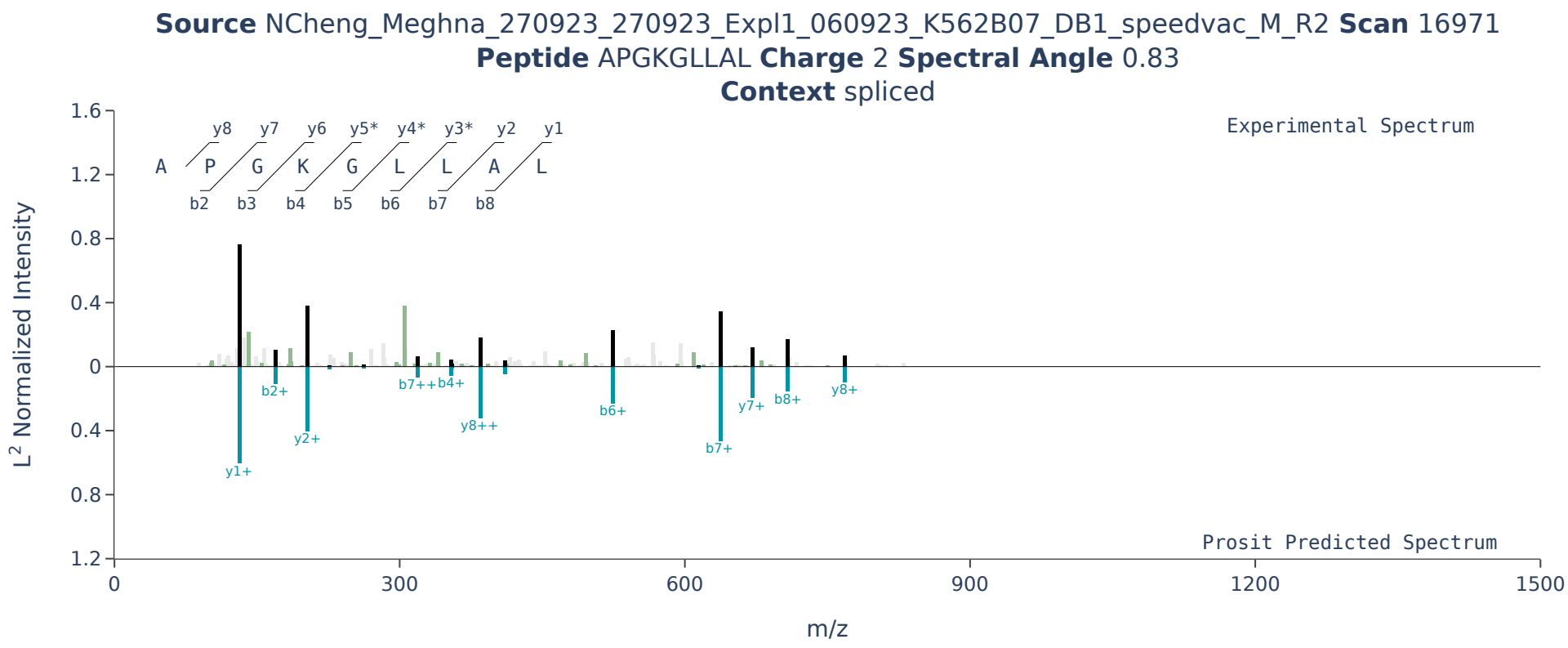
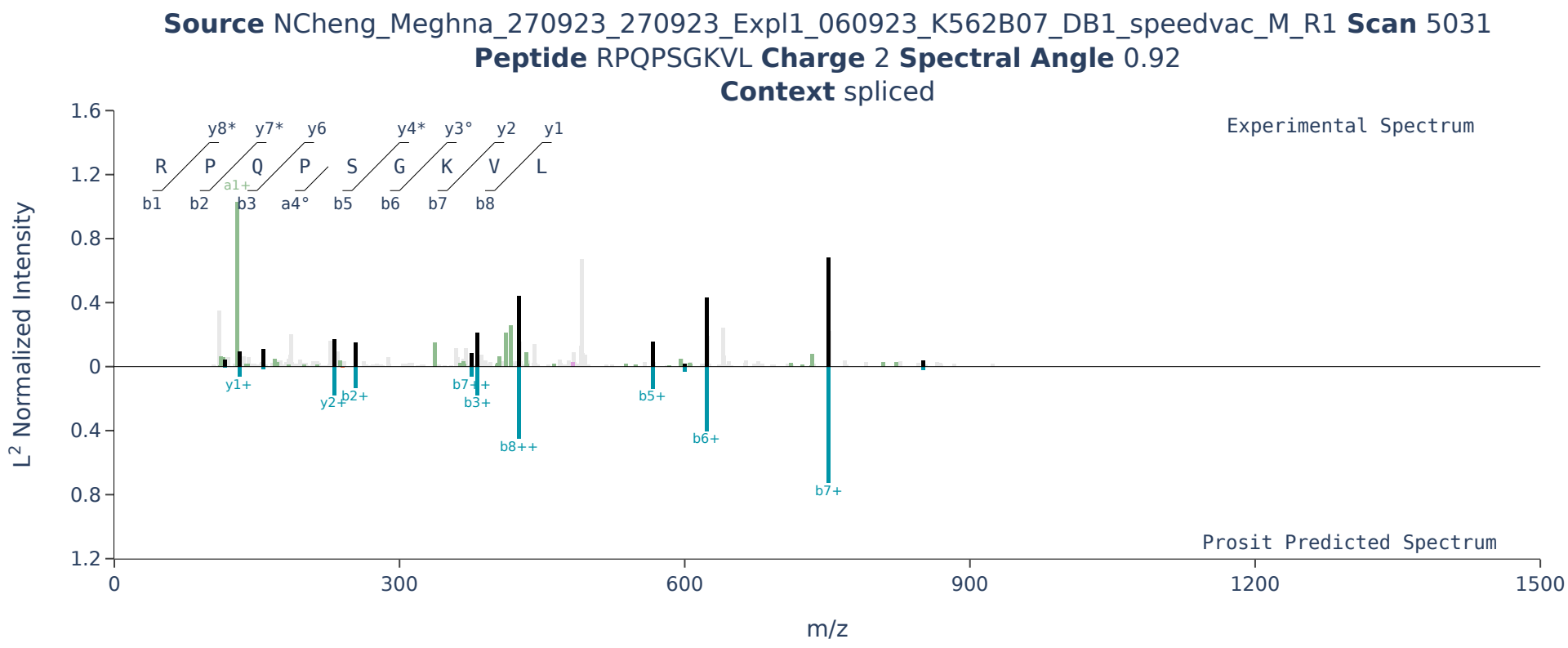
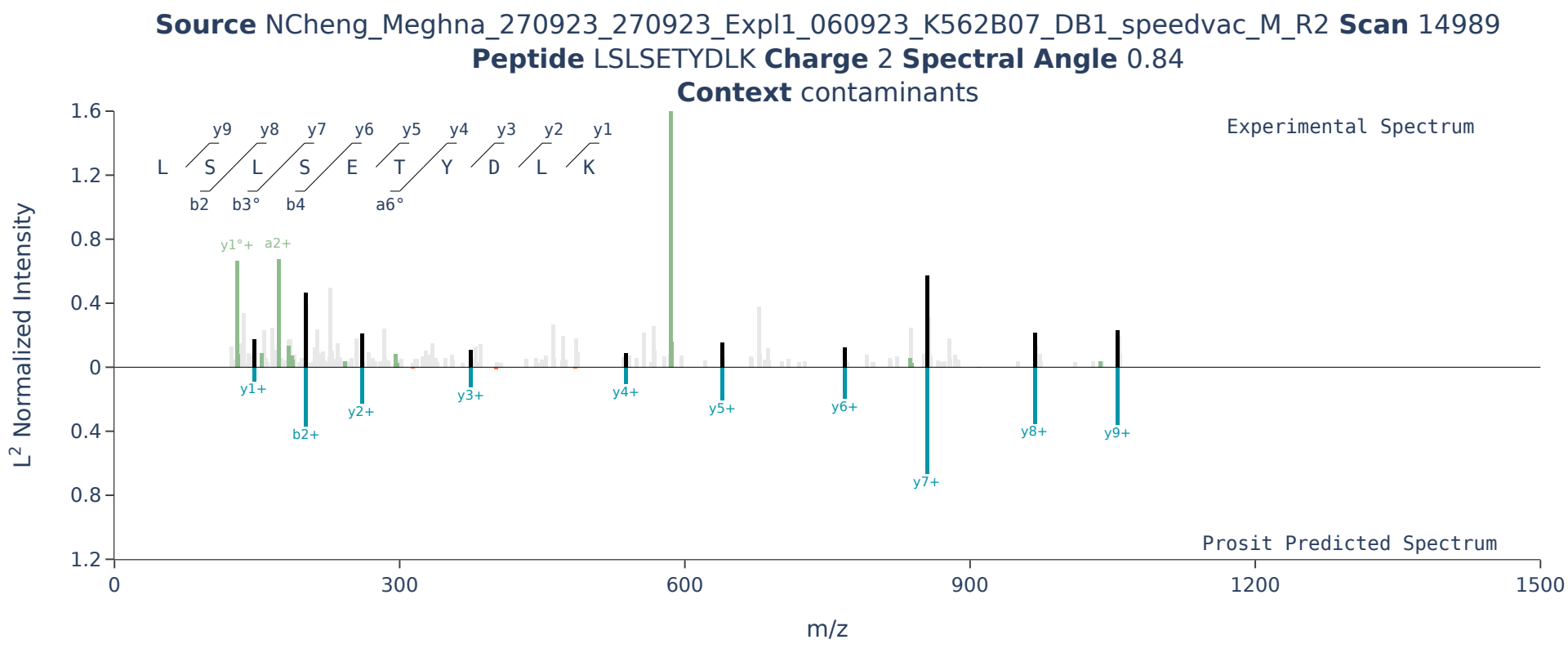
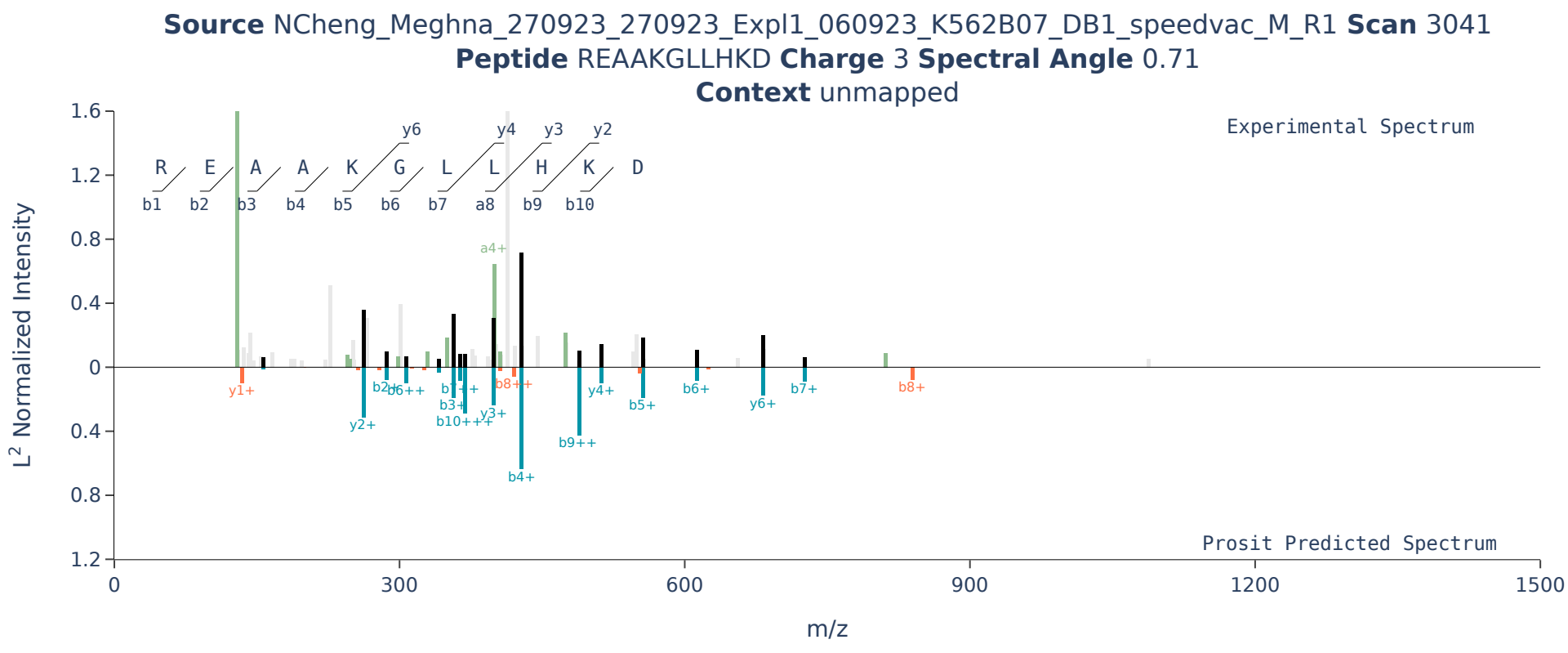
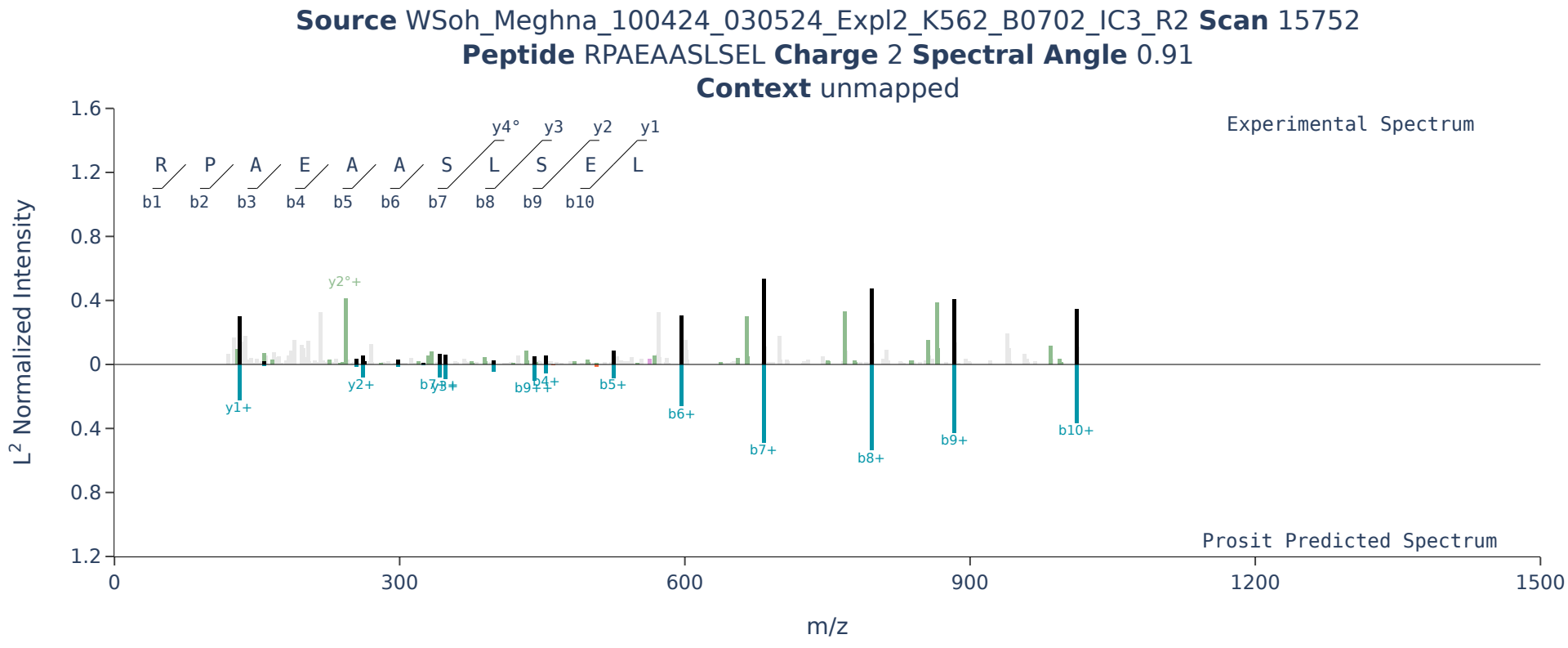
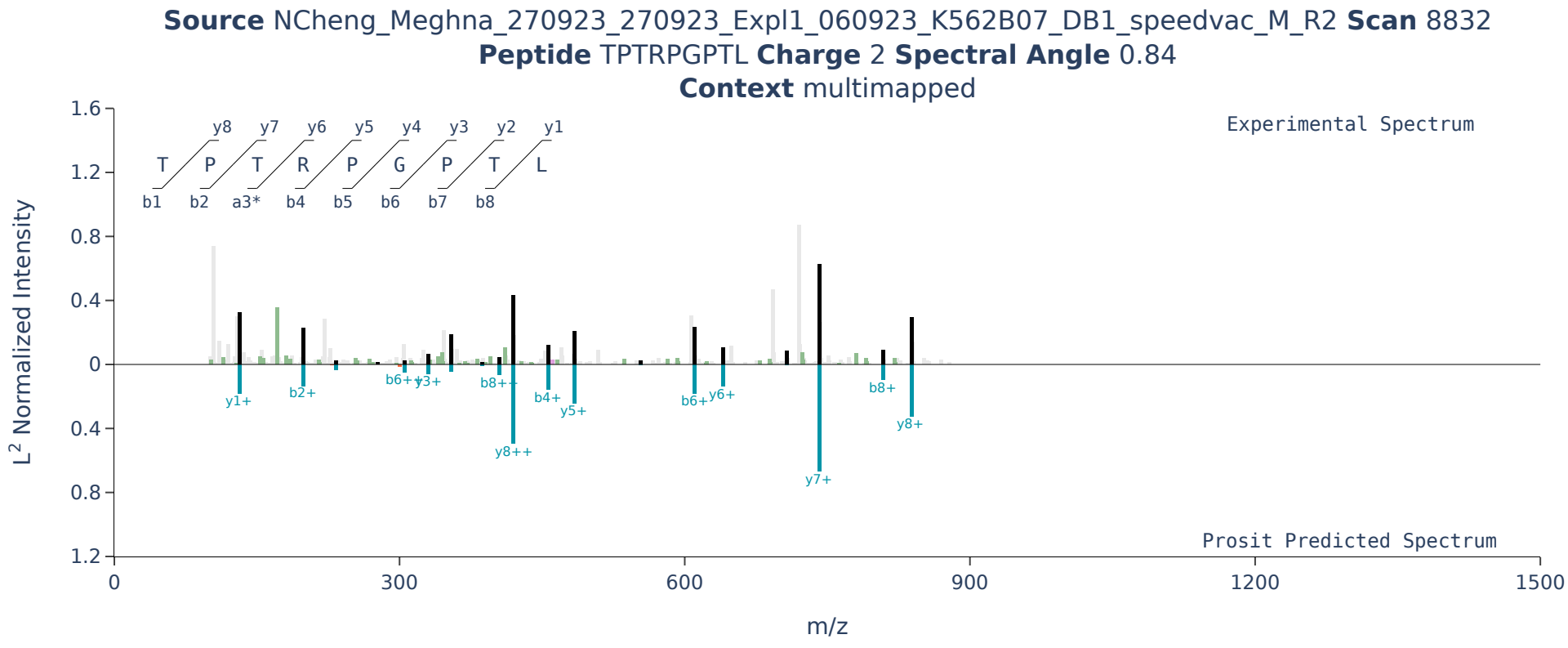
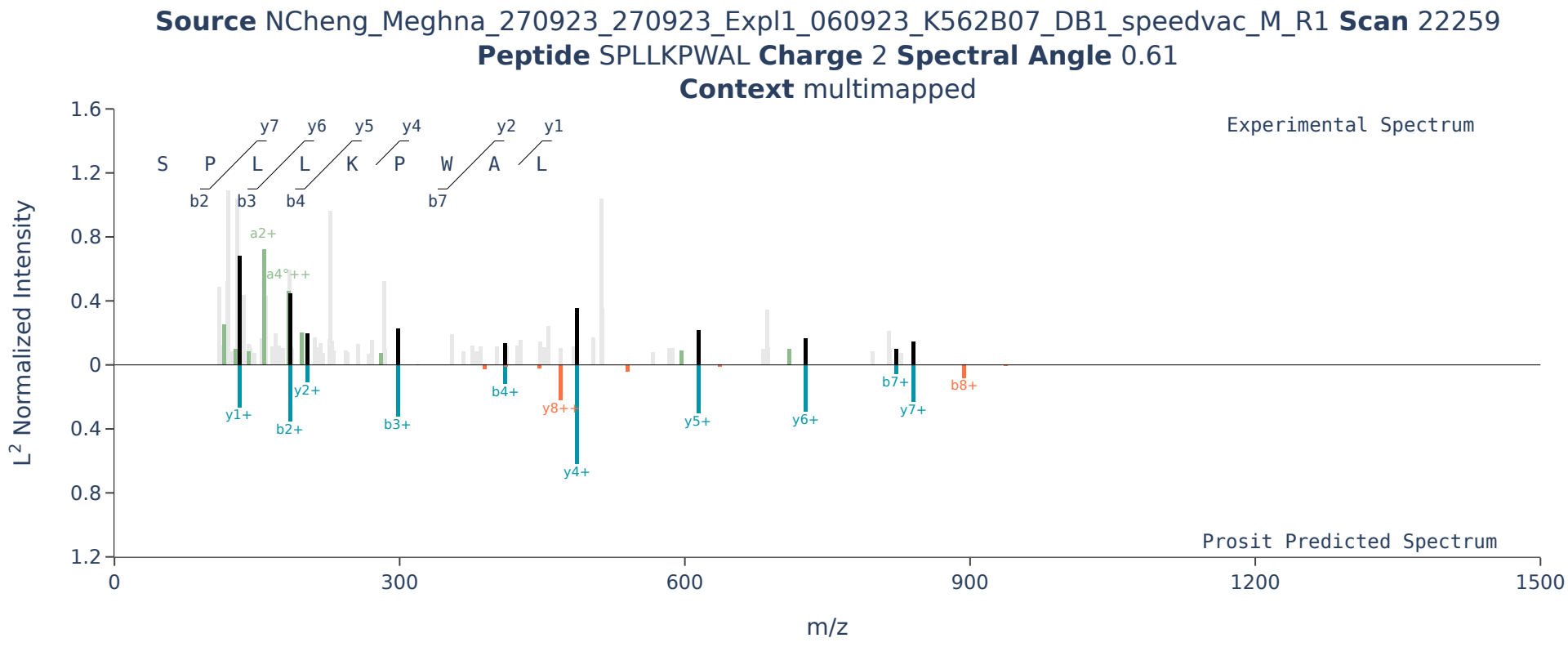
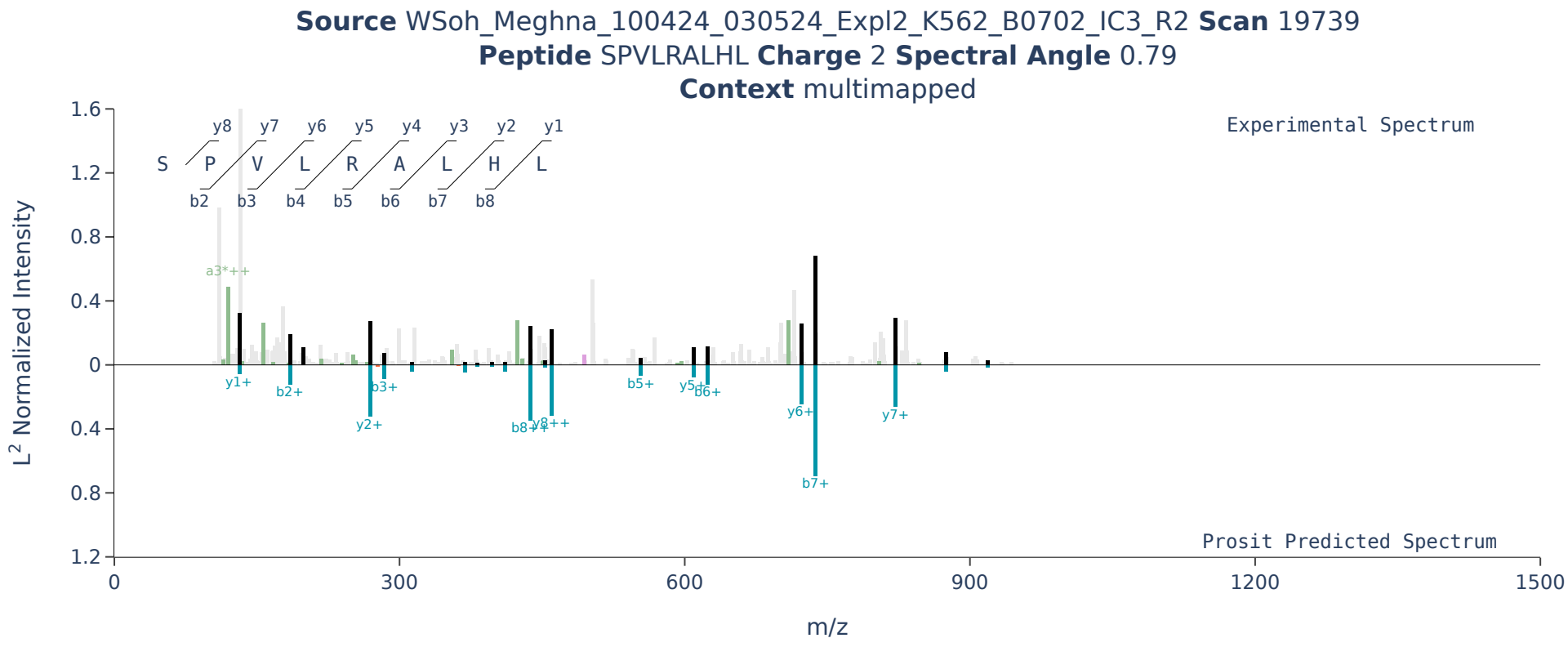
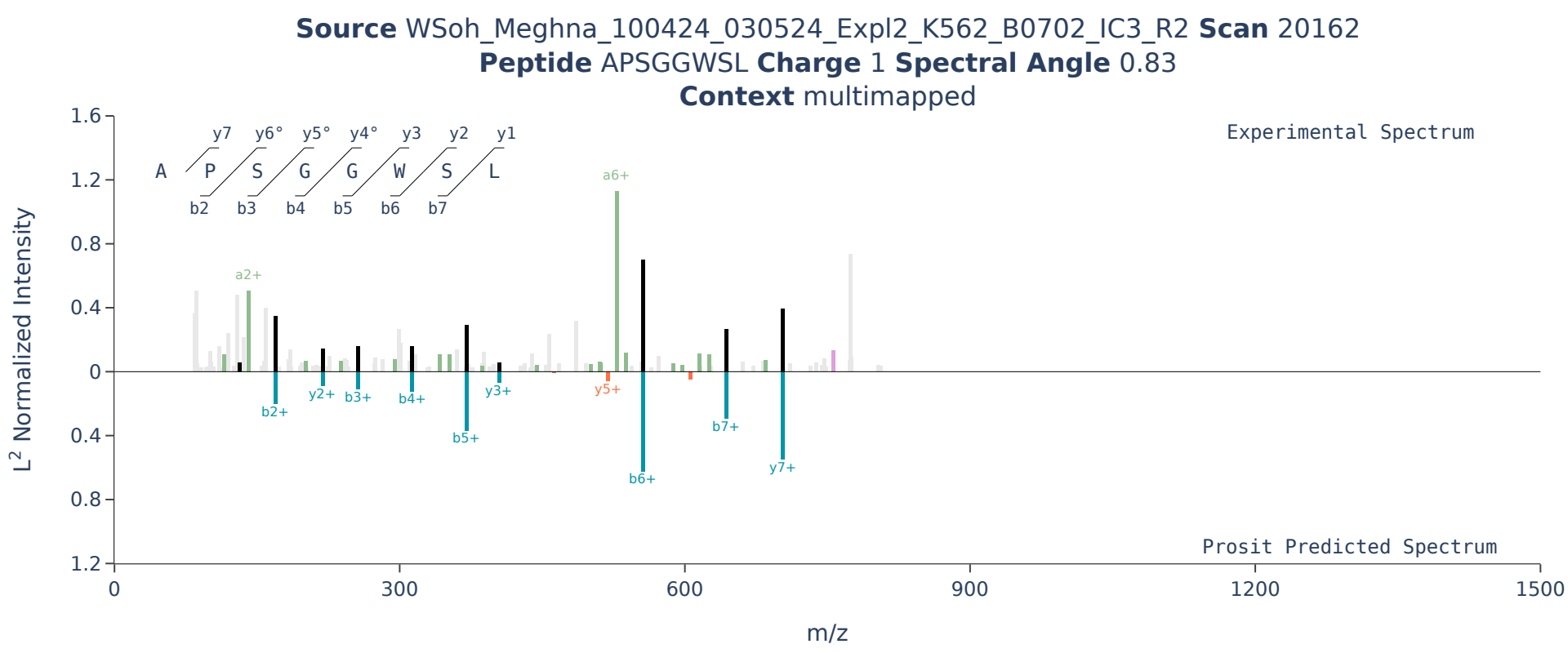
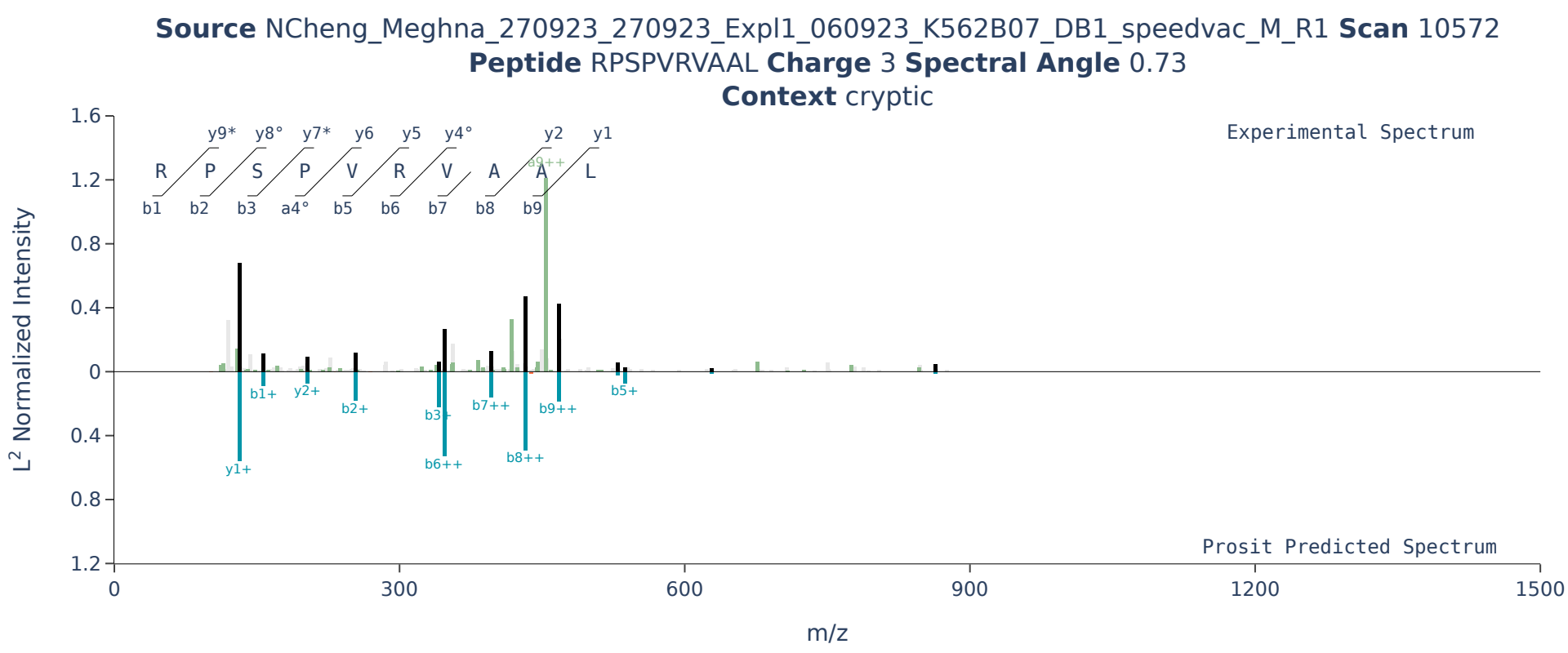
Source NCheng_Meghna_270923_270923_Exp1_060923_K562B07_DB1_speedvac_M_R2 **Scan** 5005
Peptide RPYSHGKFM **Charge** 3 **Spectral Angle** 0.78

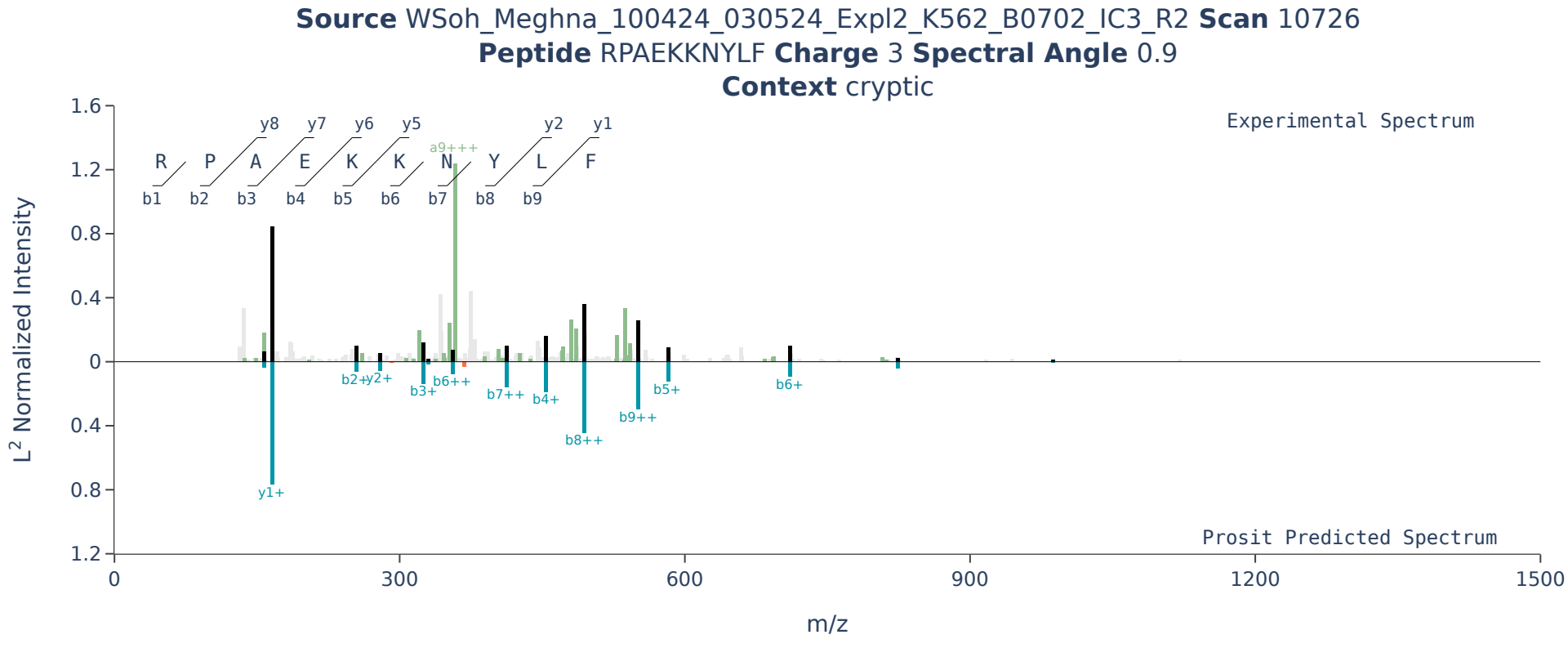
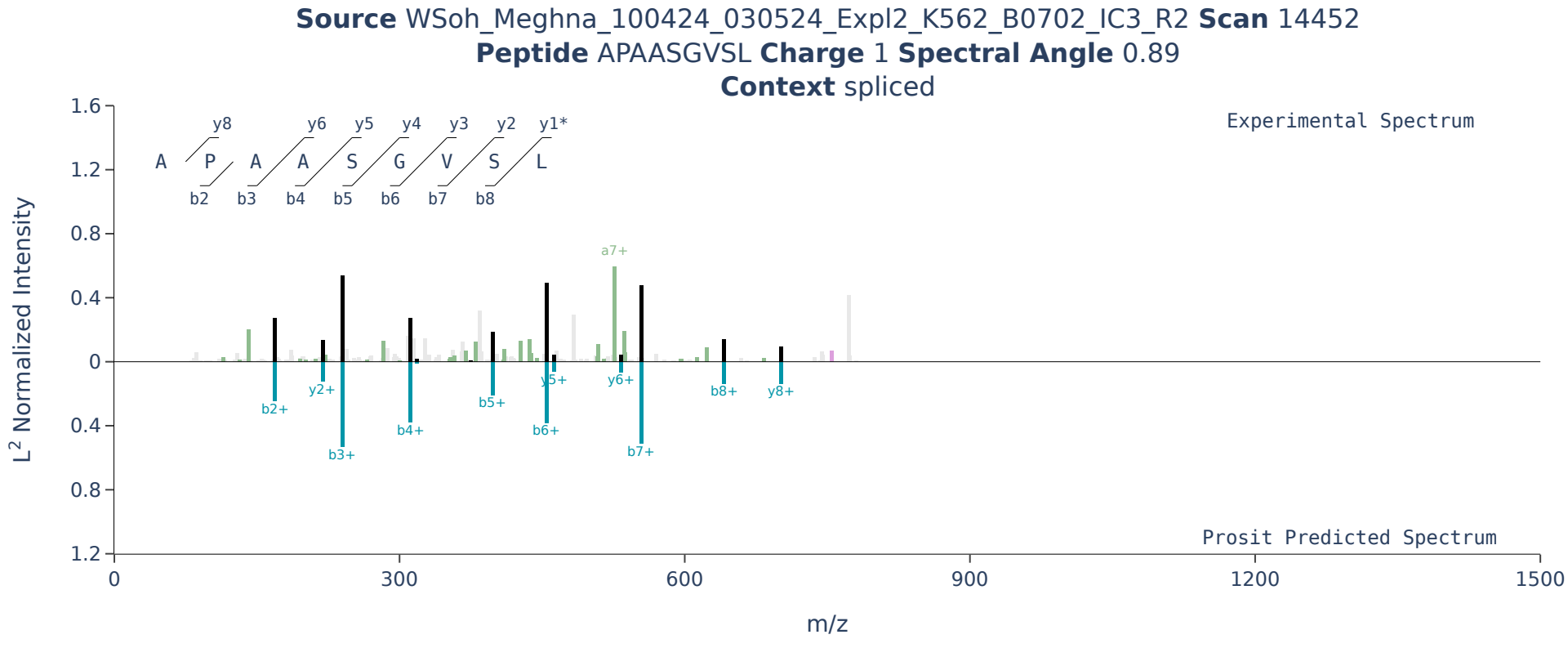
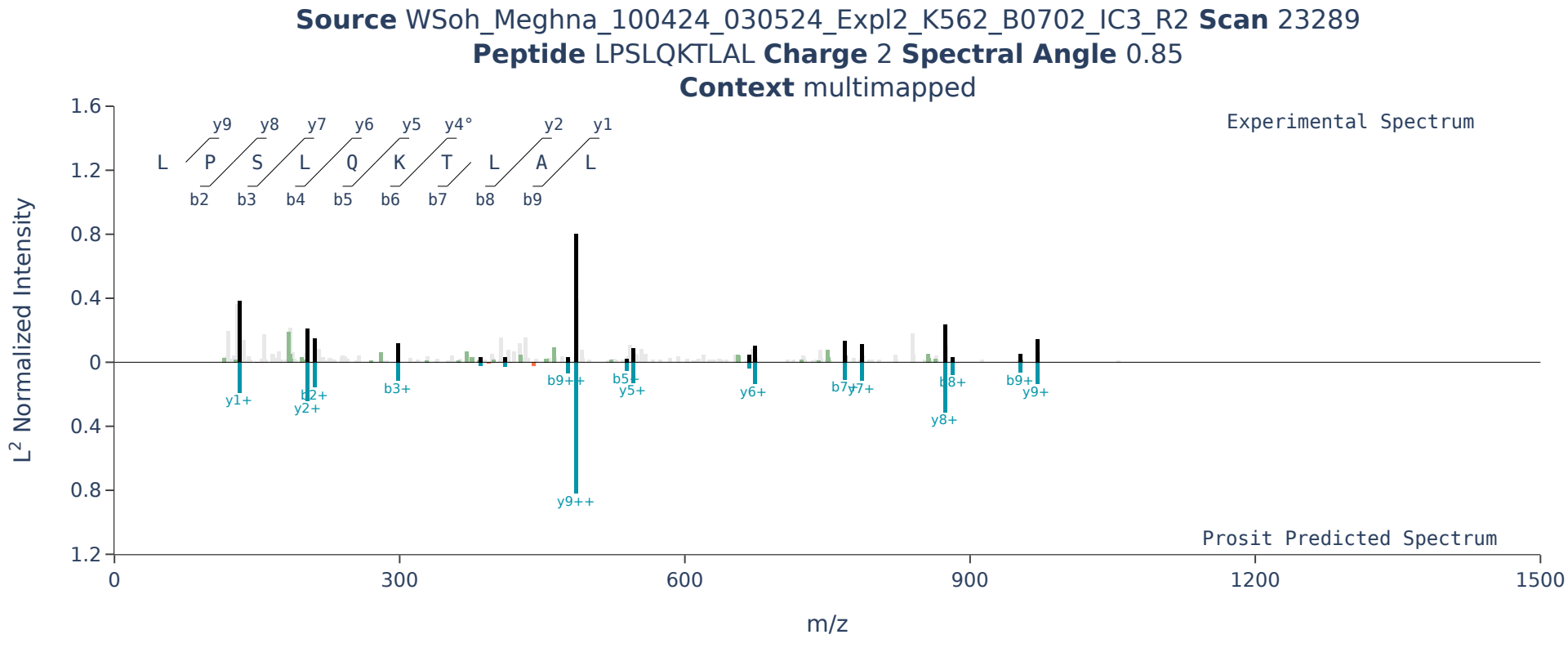
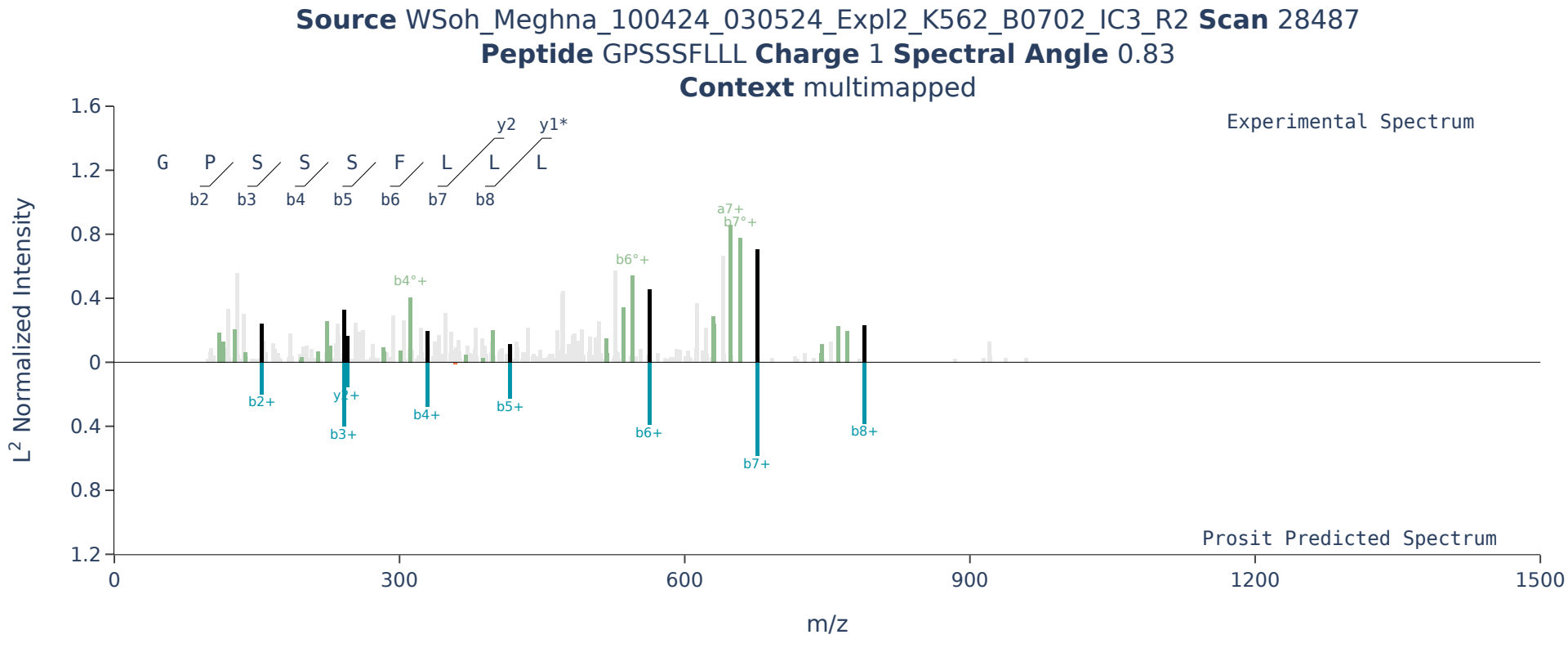
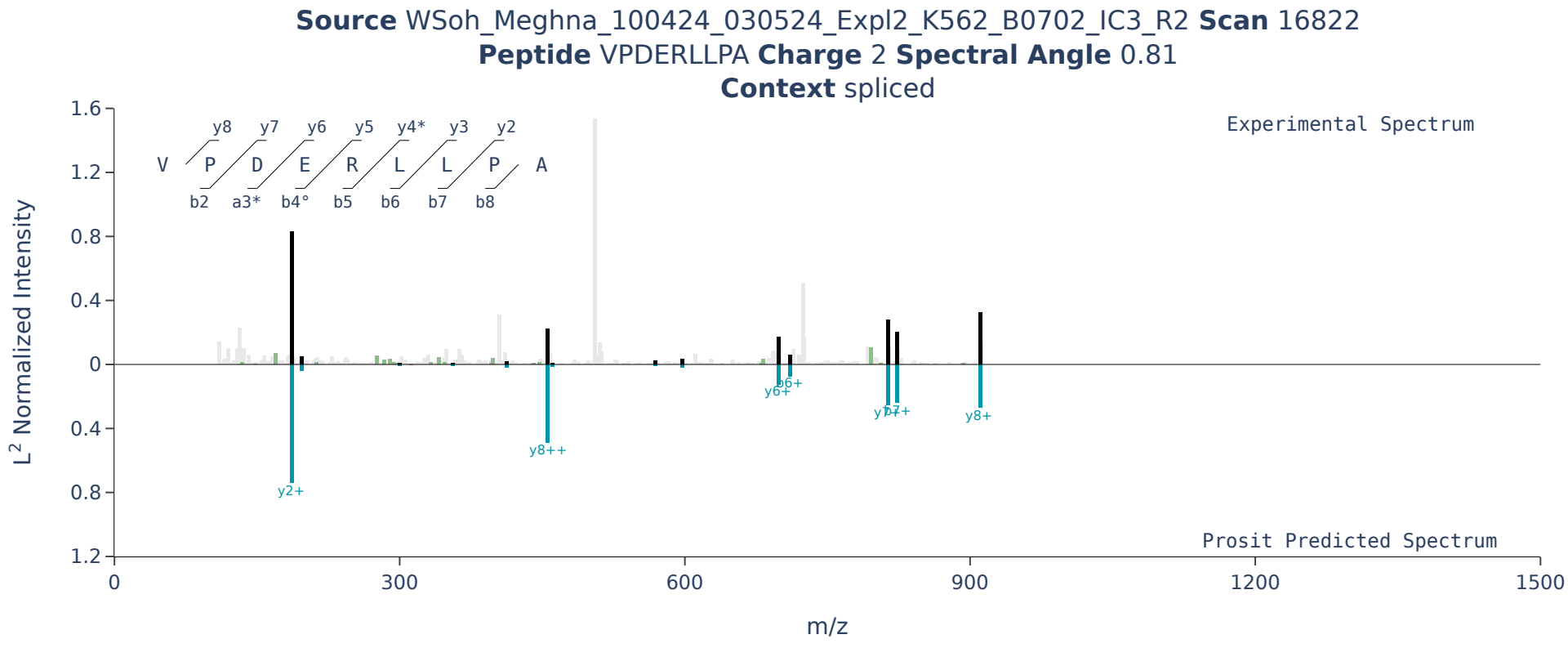
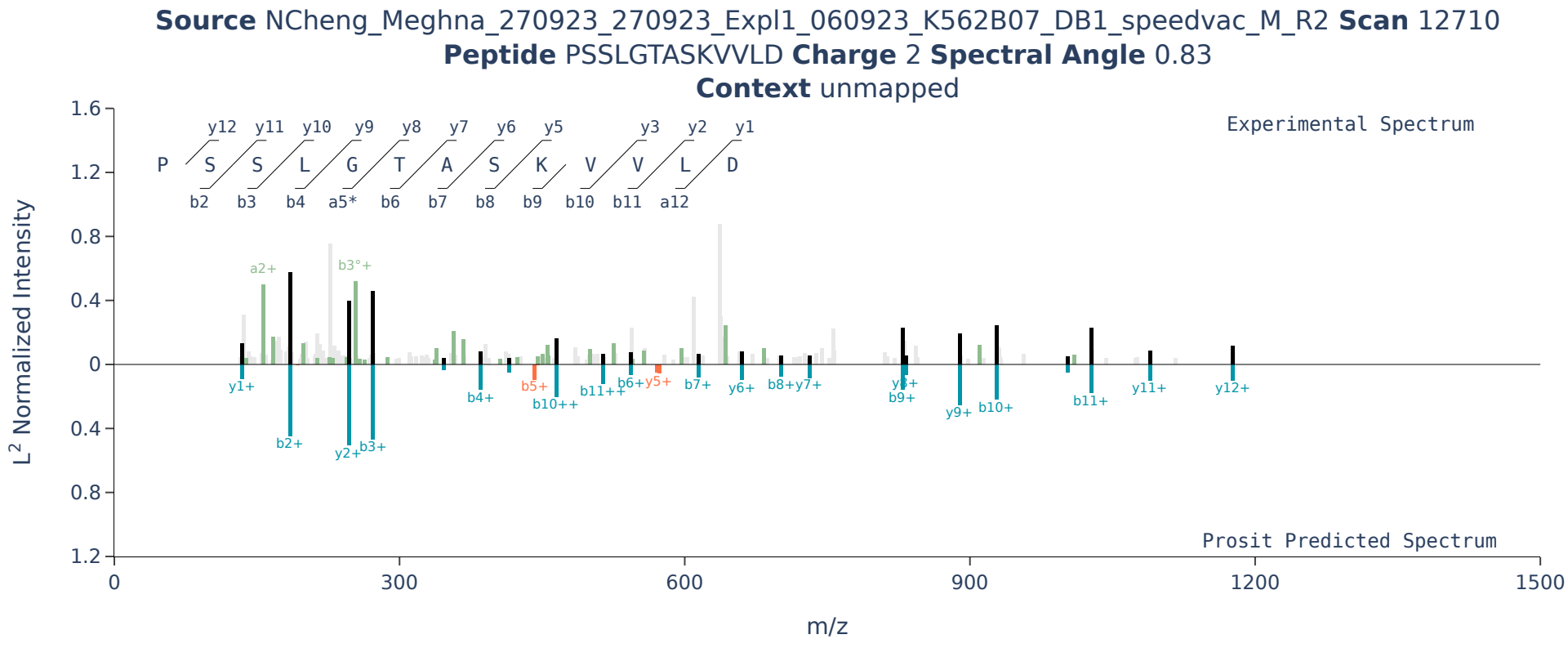
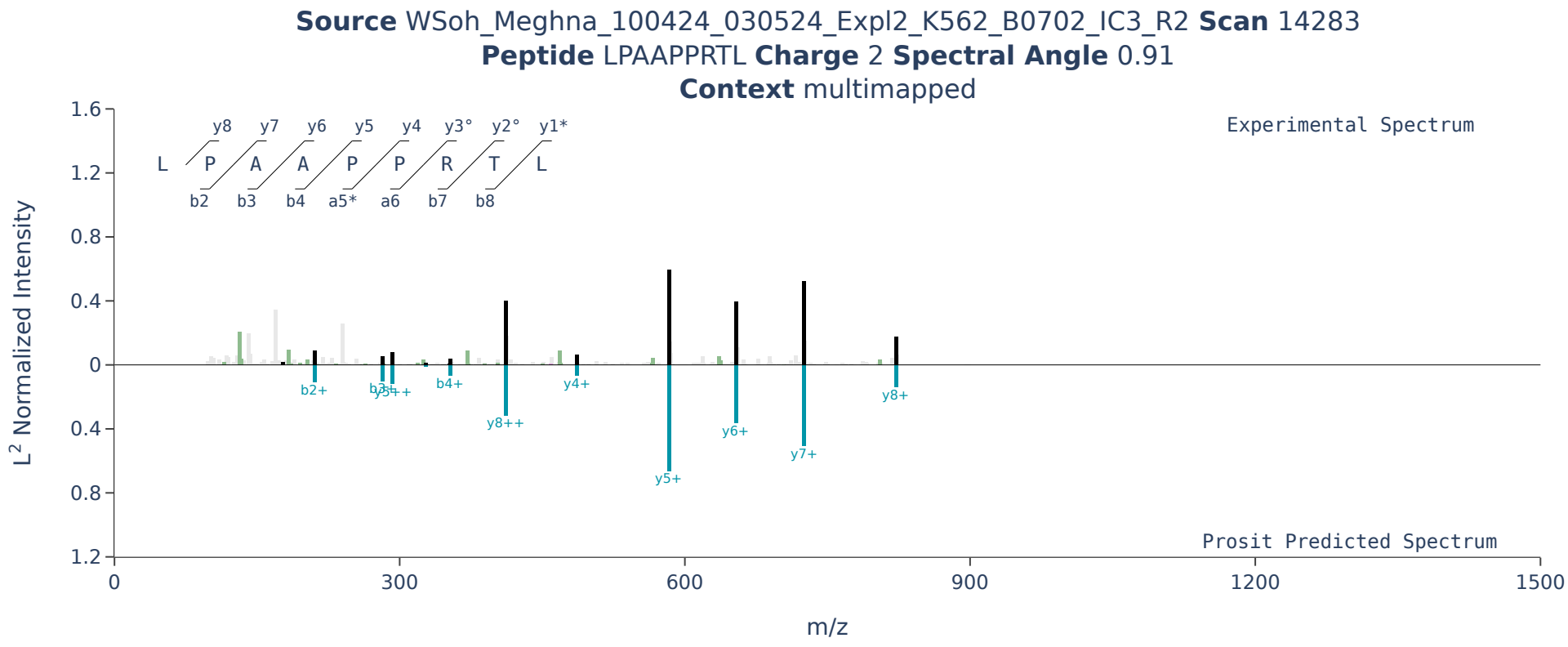
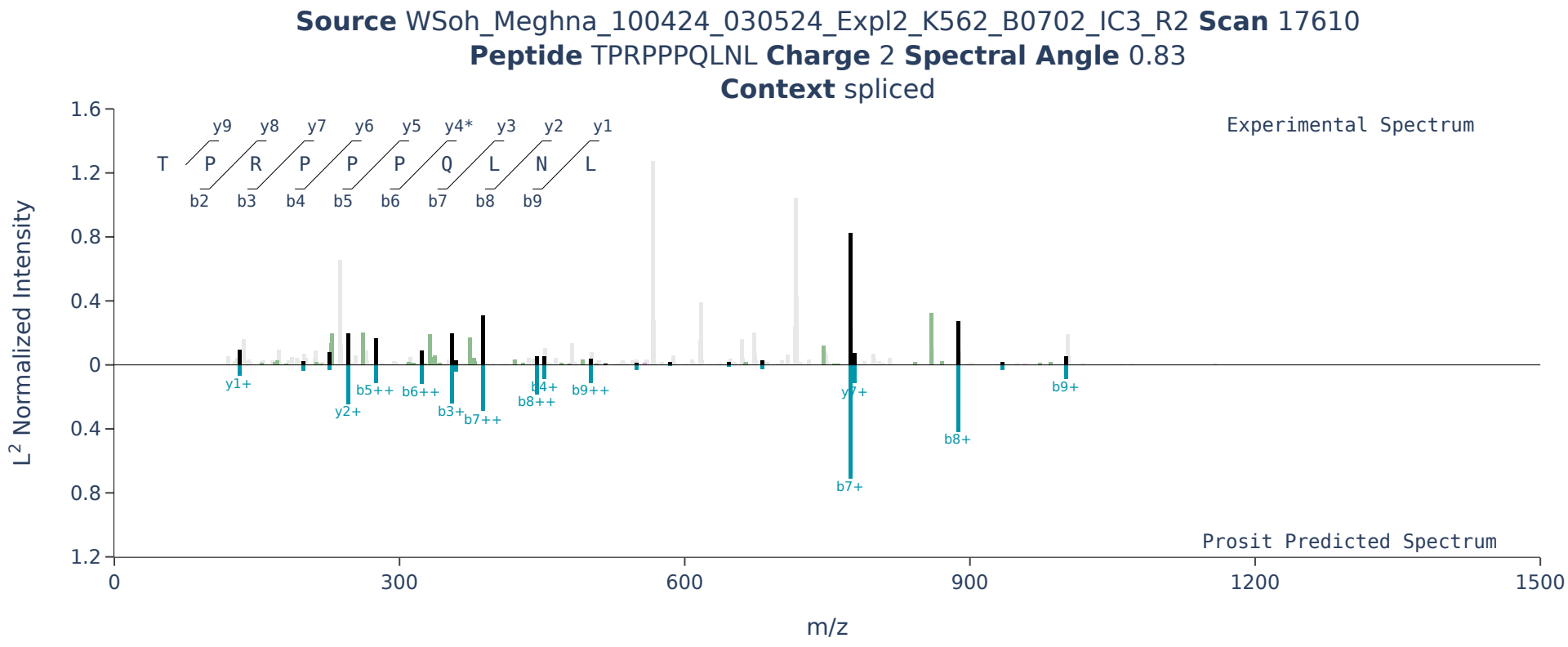
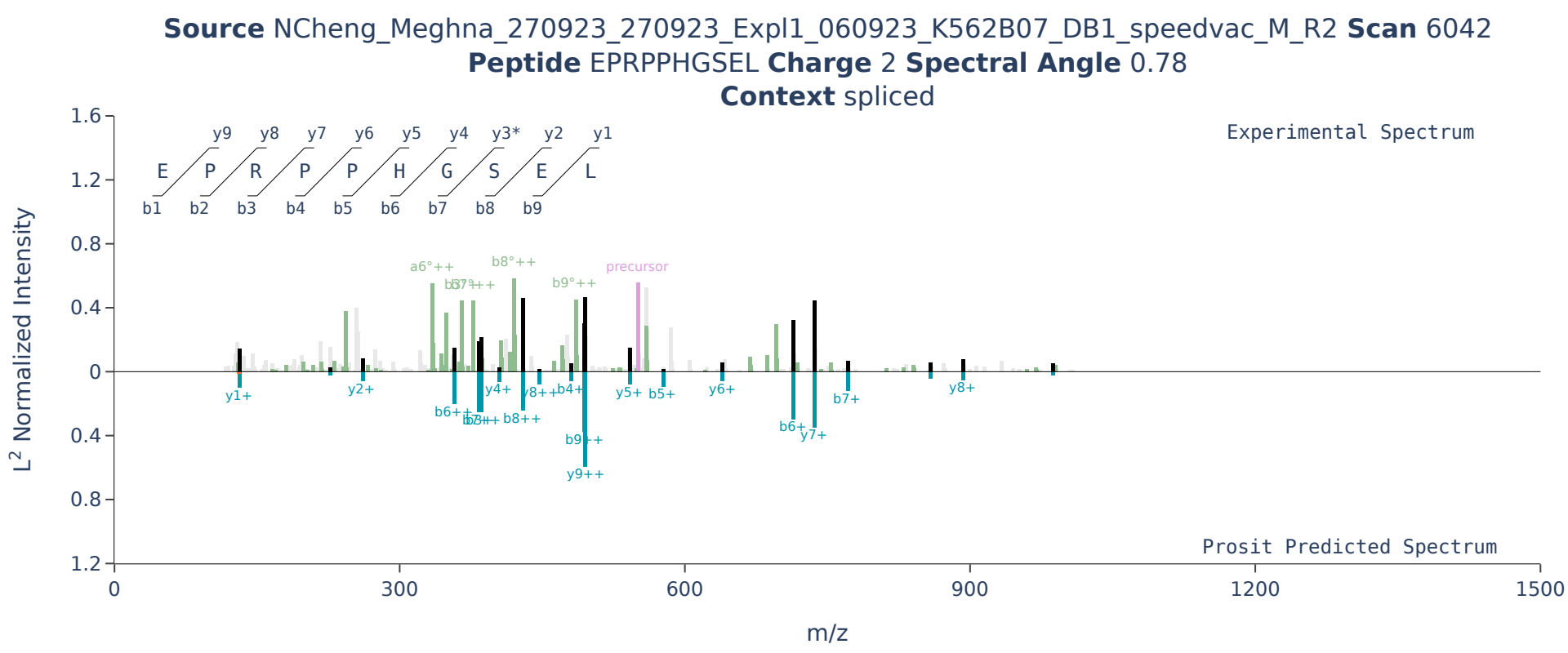
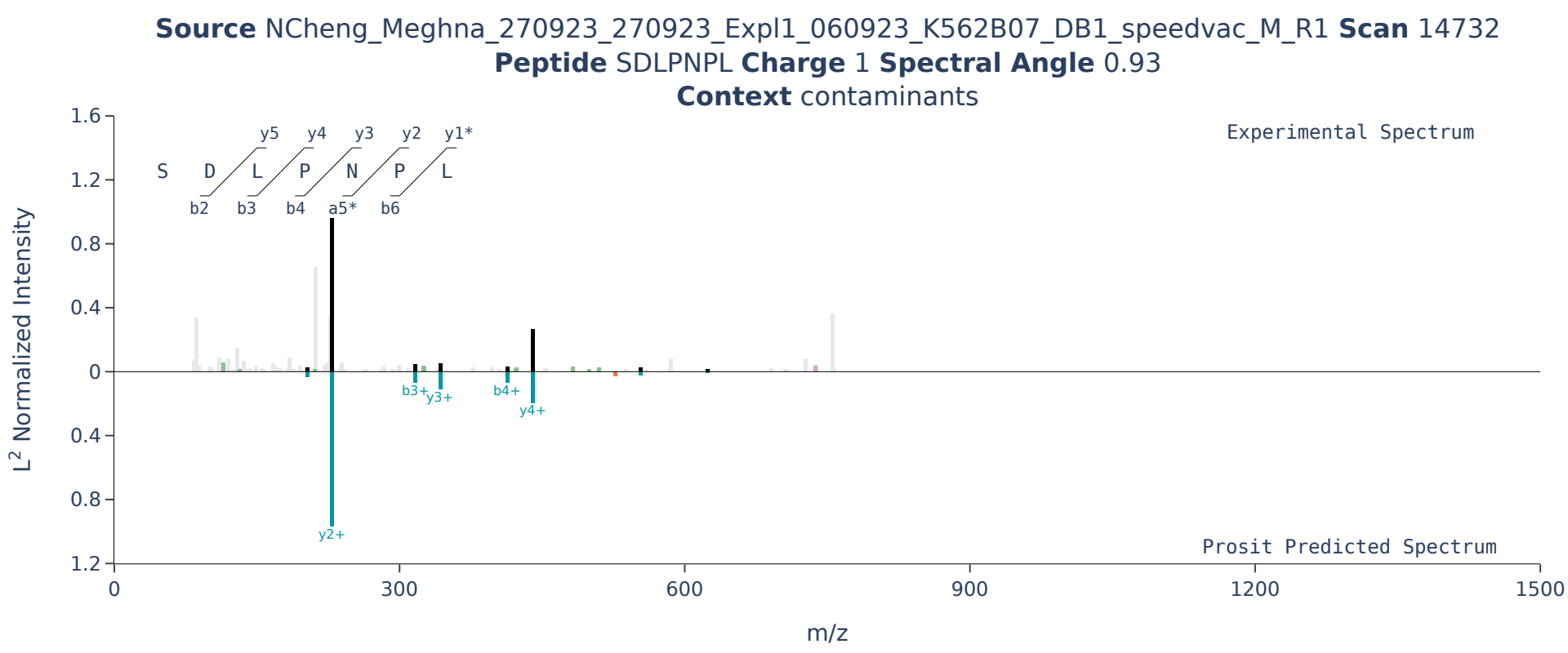


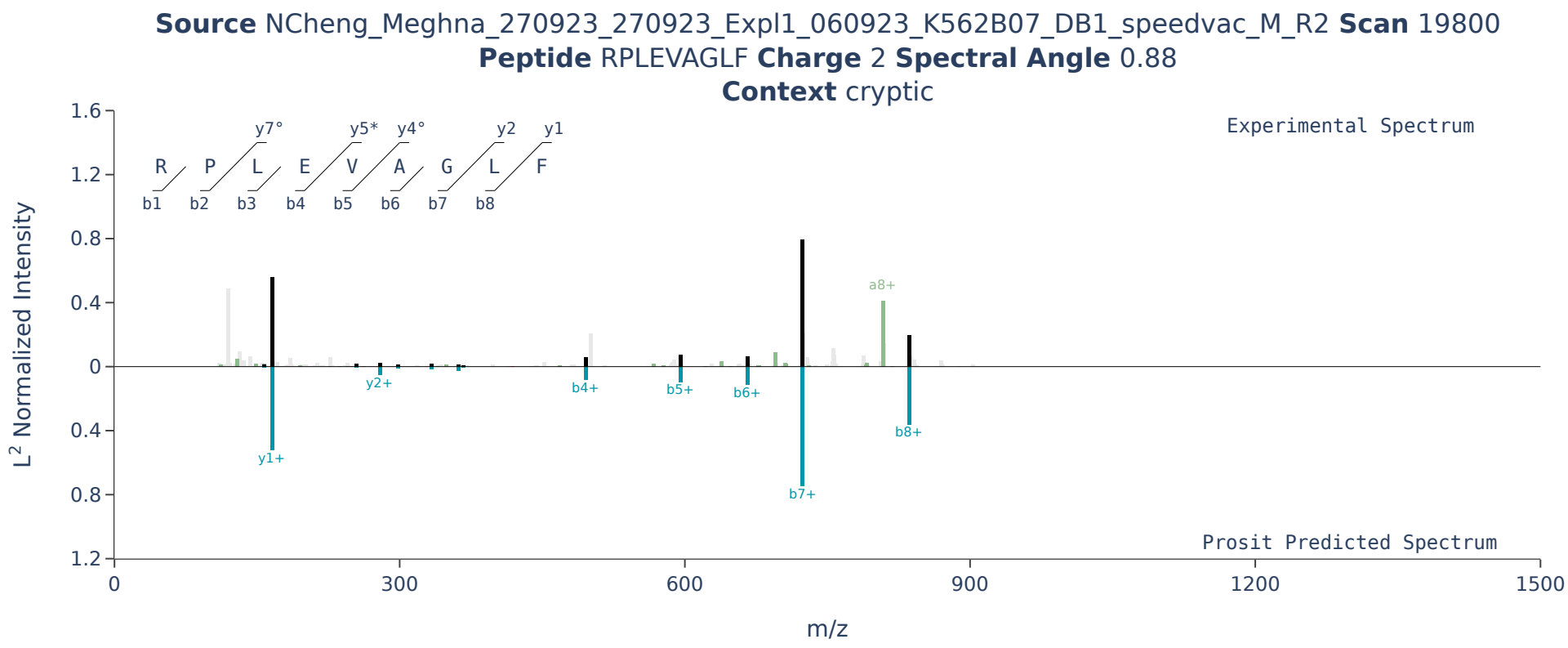
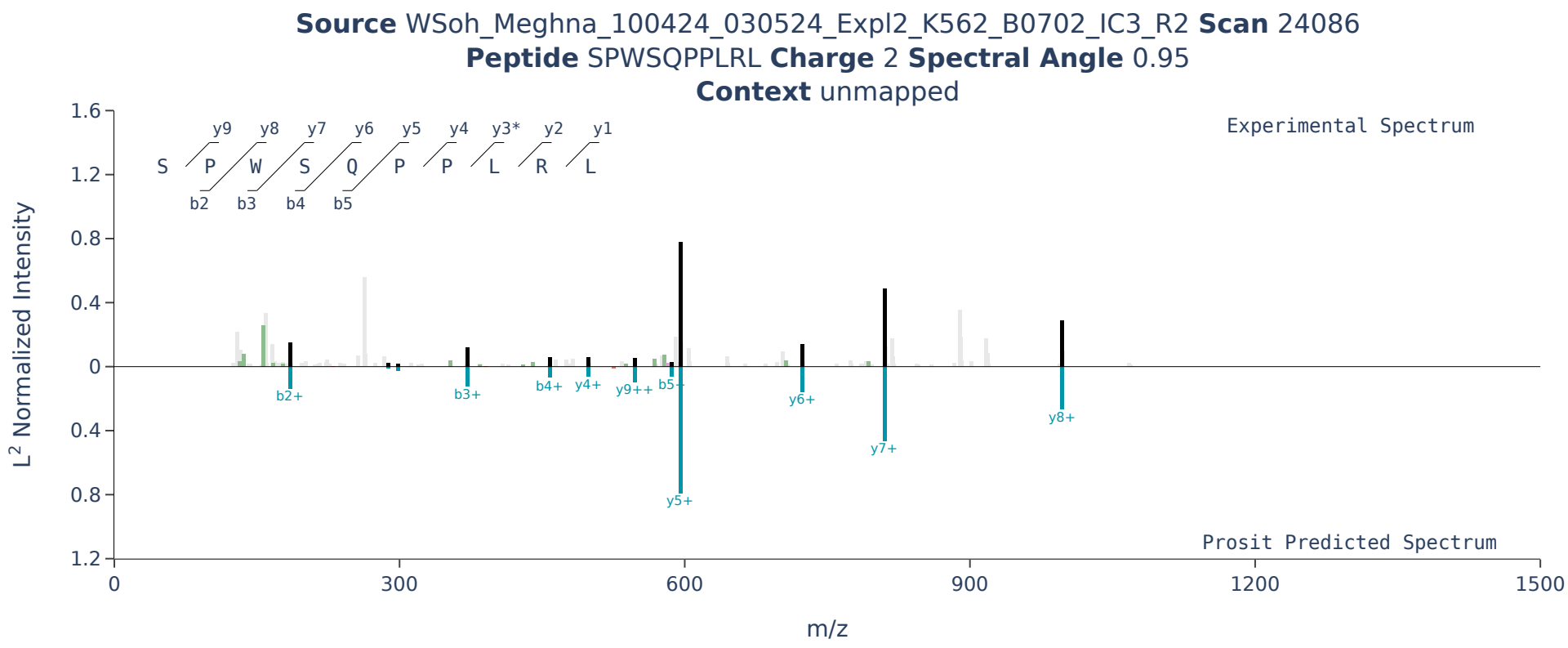
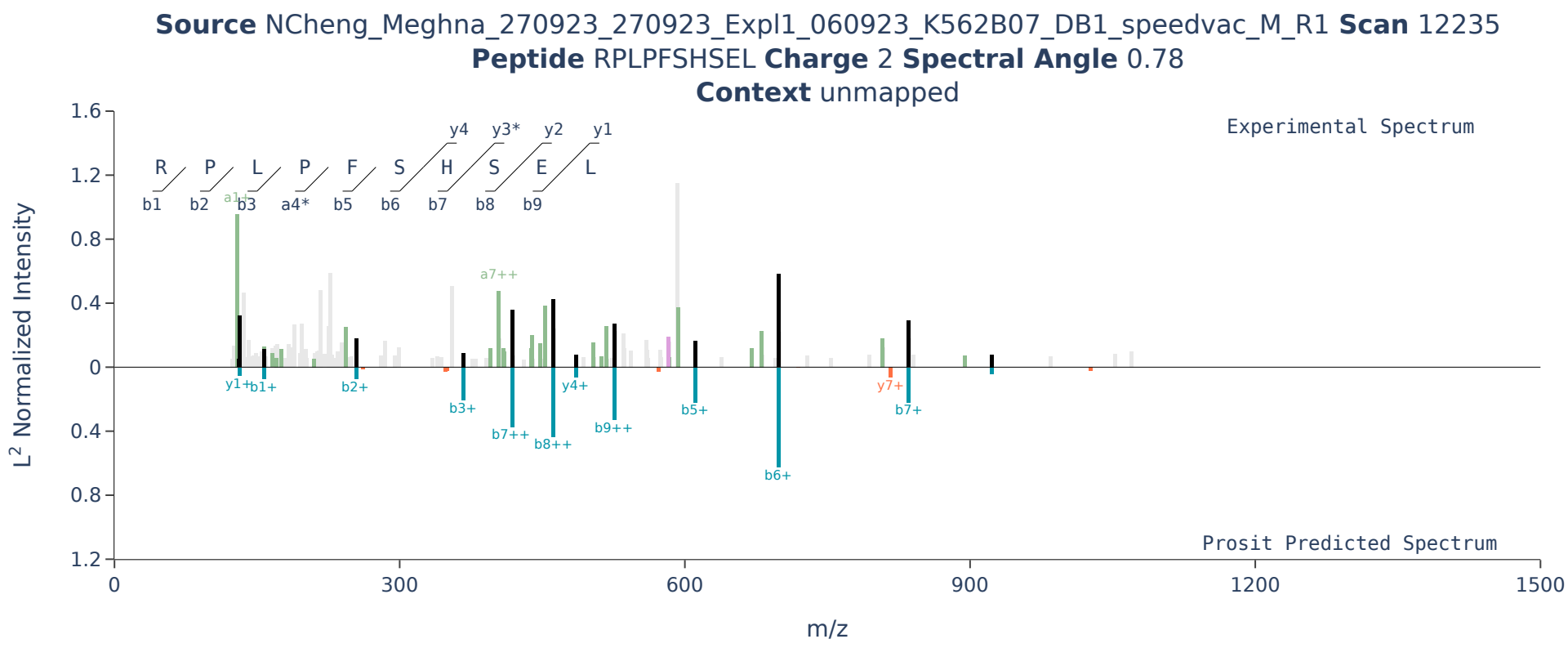
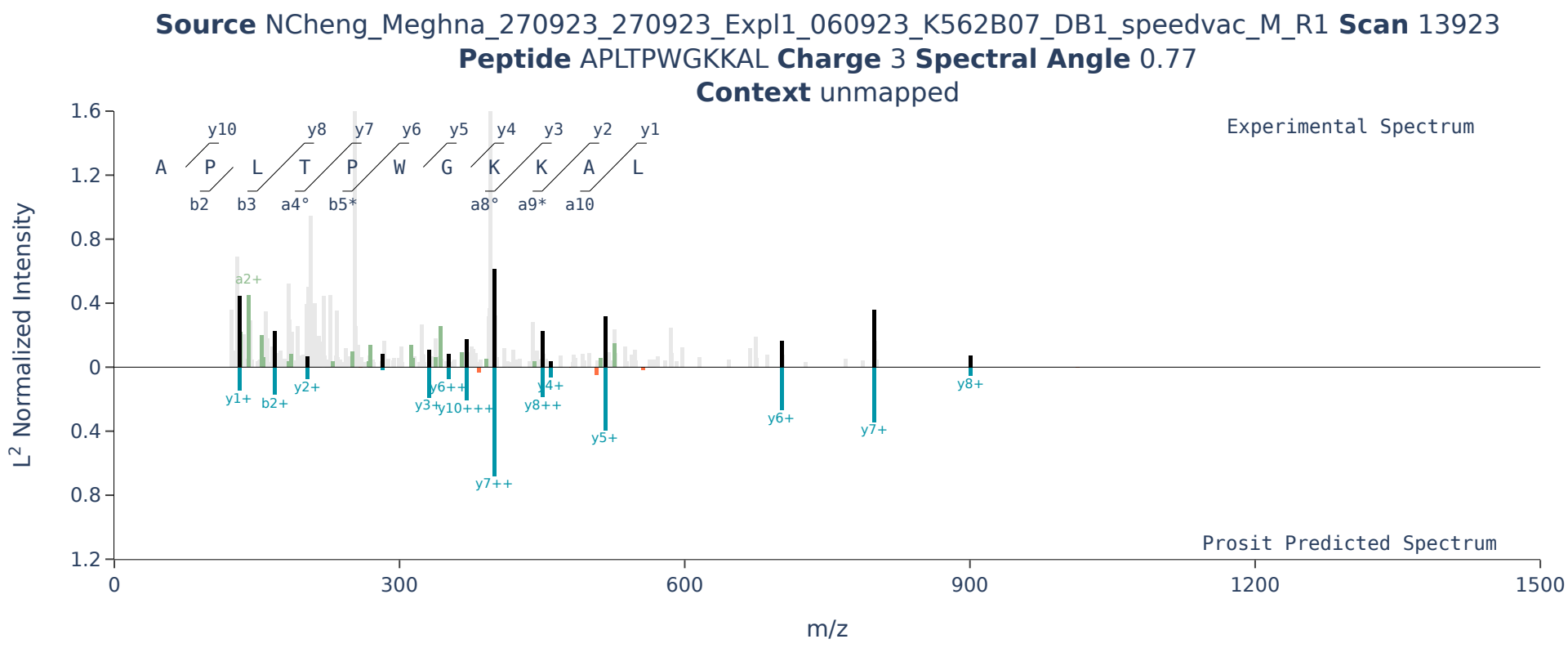
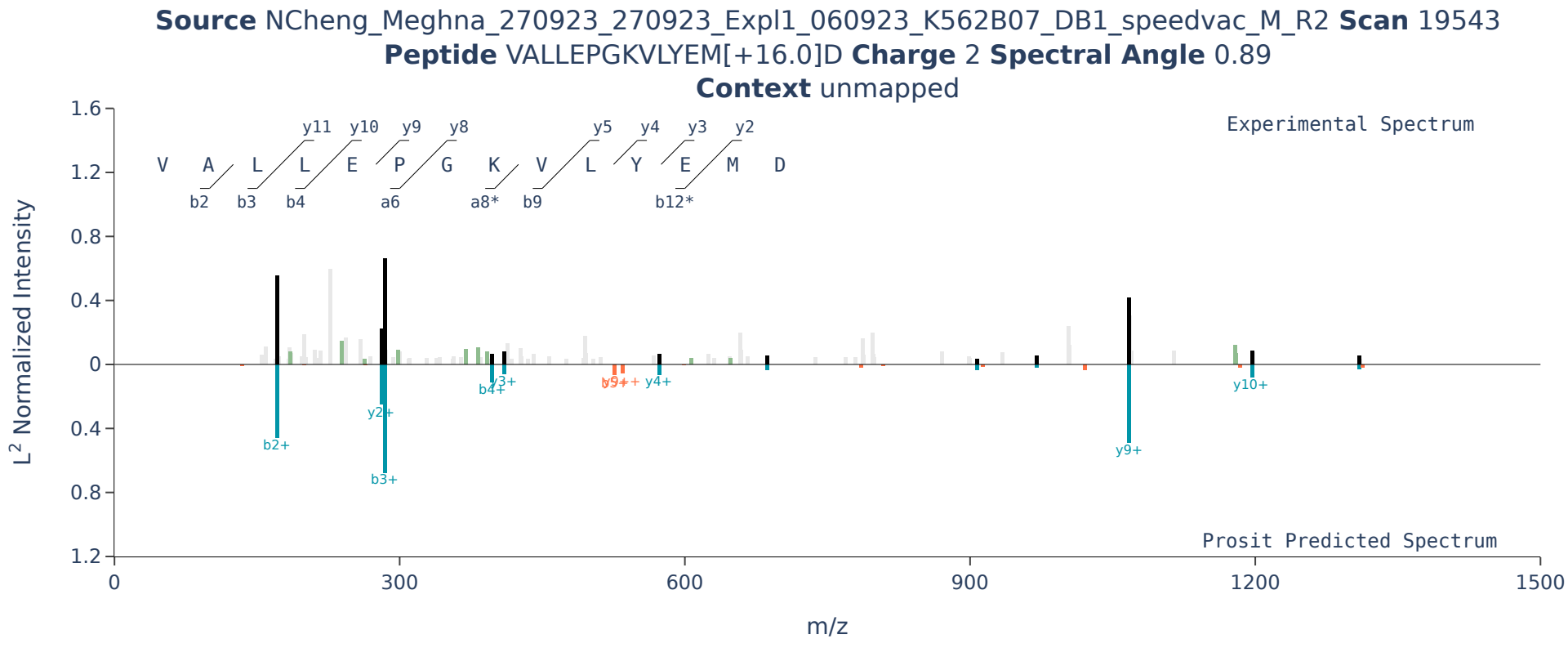
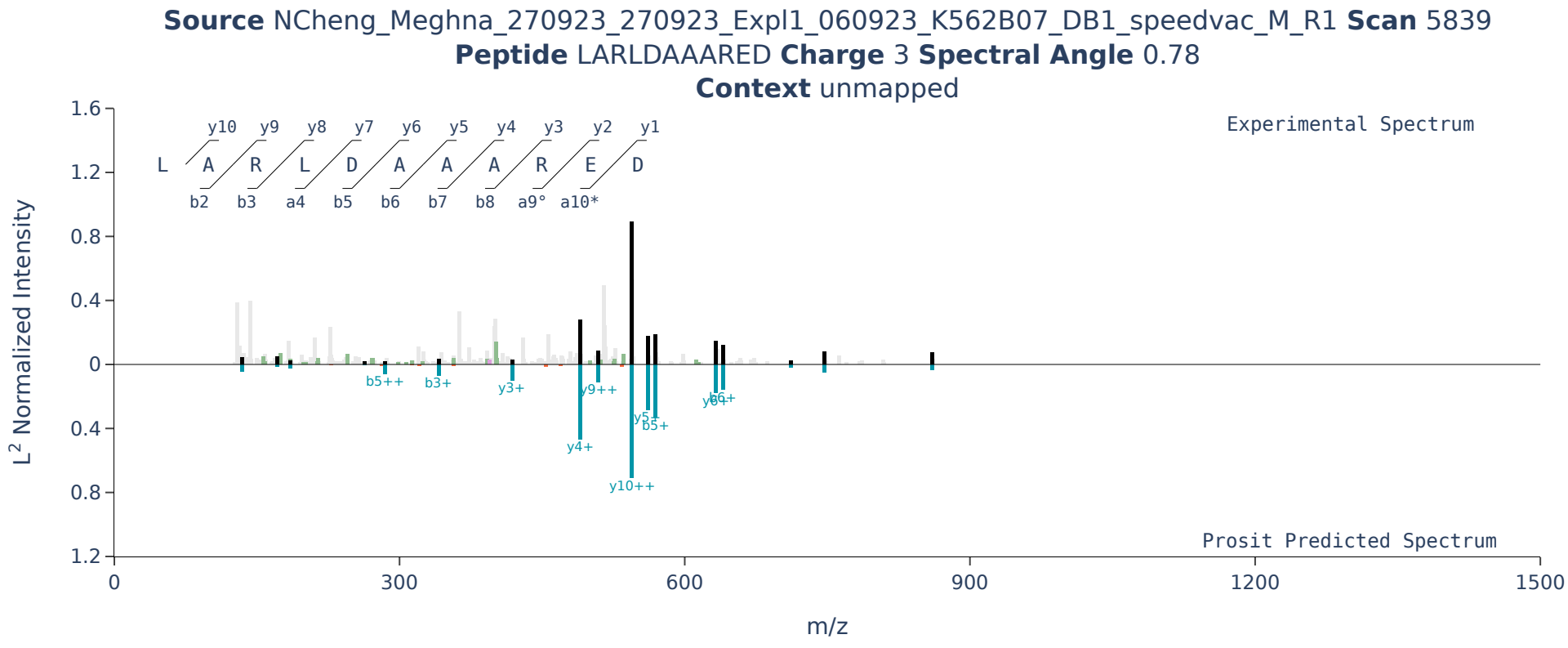
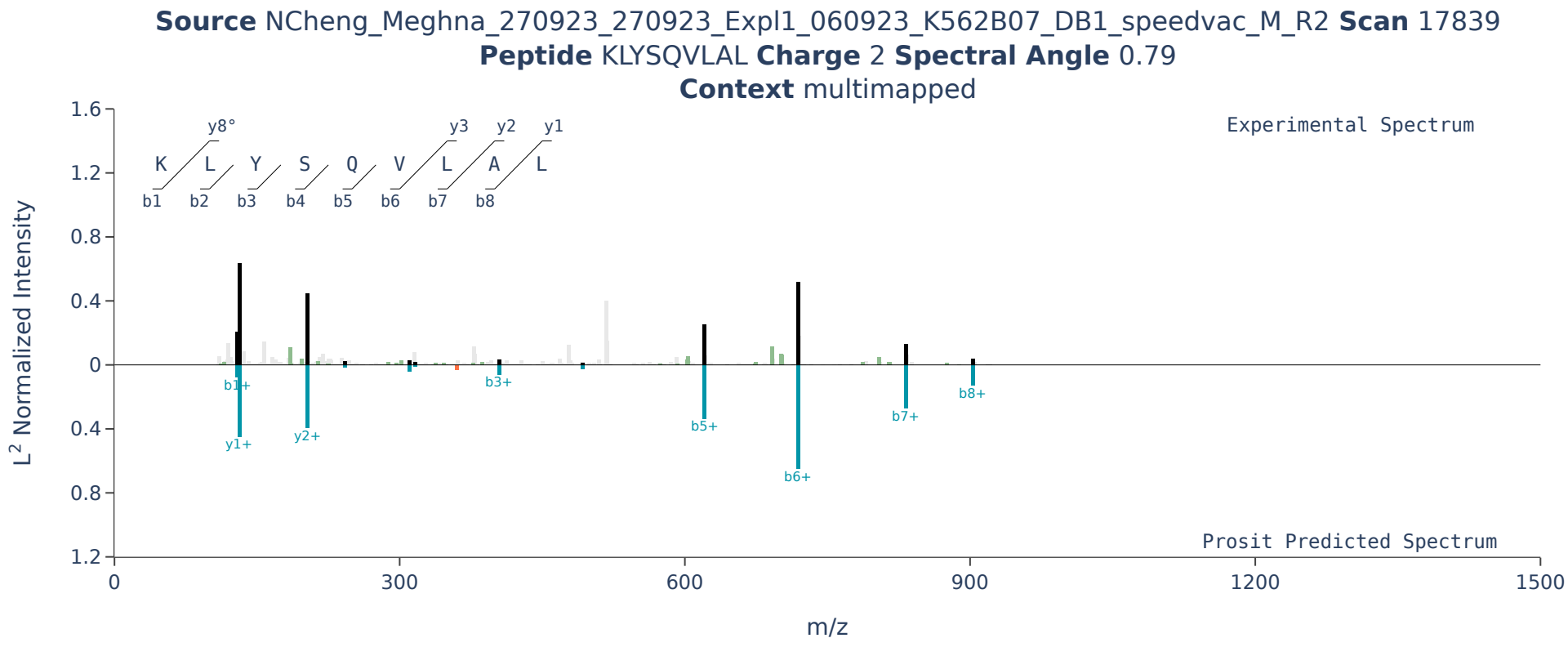
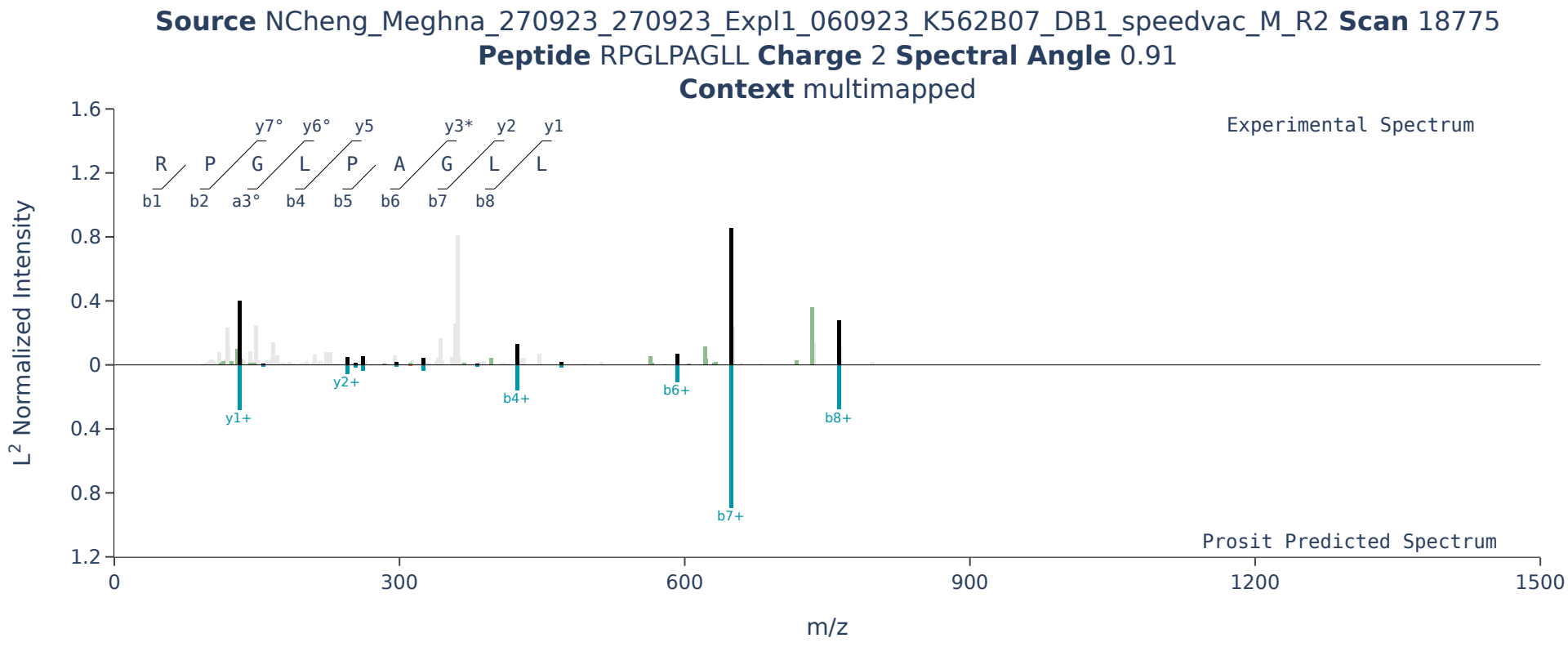
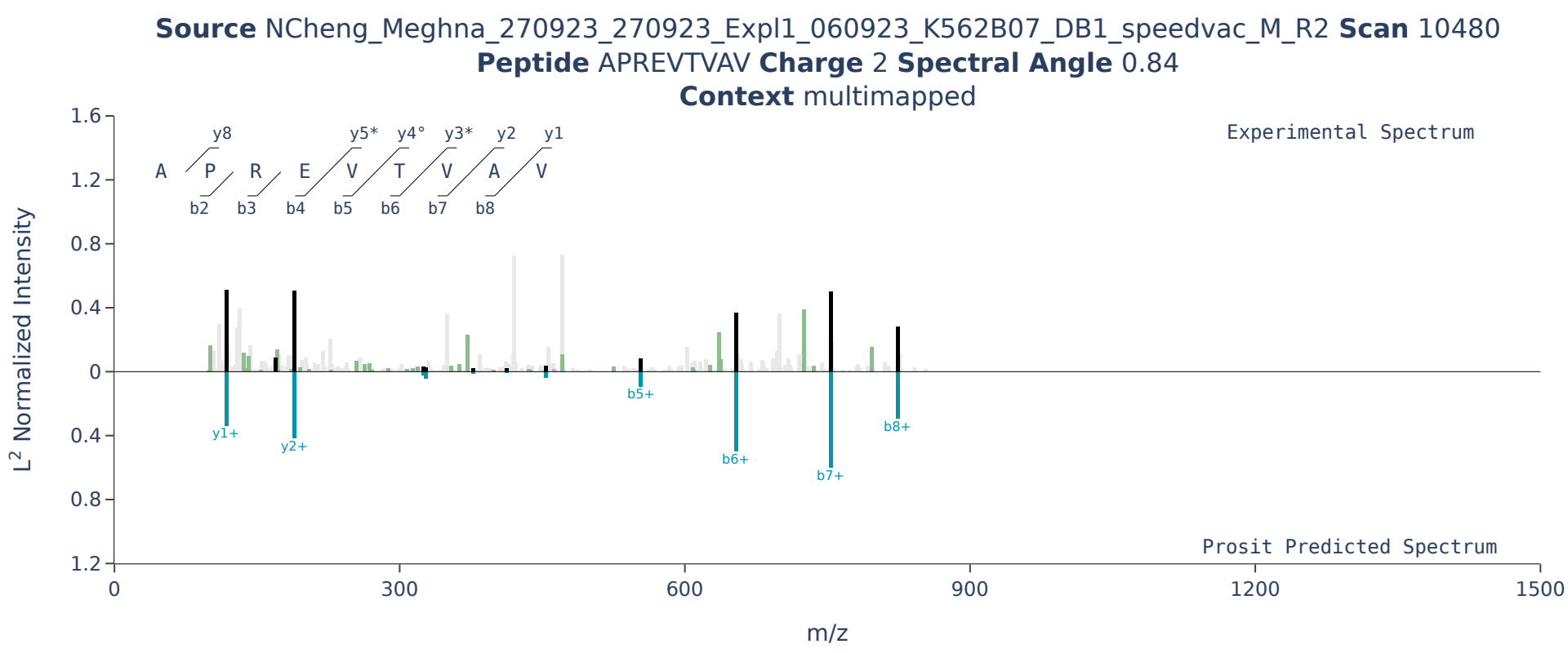
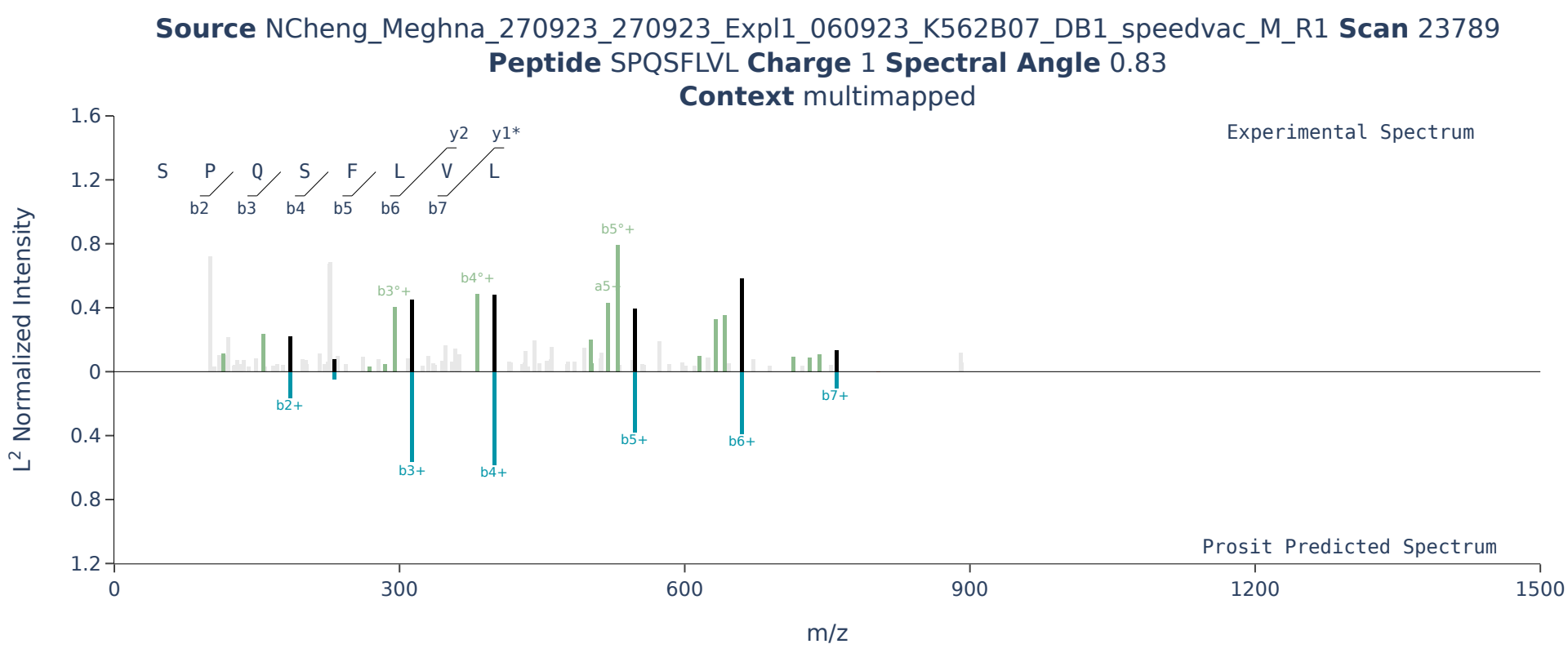
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Peptide APAVGLLL **Charge** 1 **Spectral Angle** 0.95

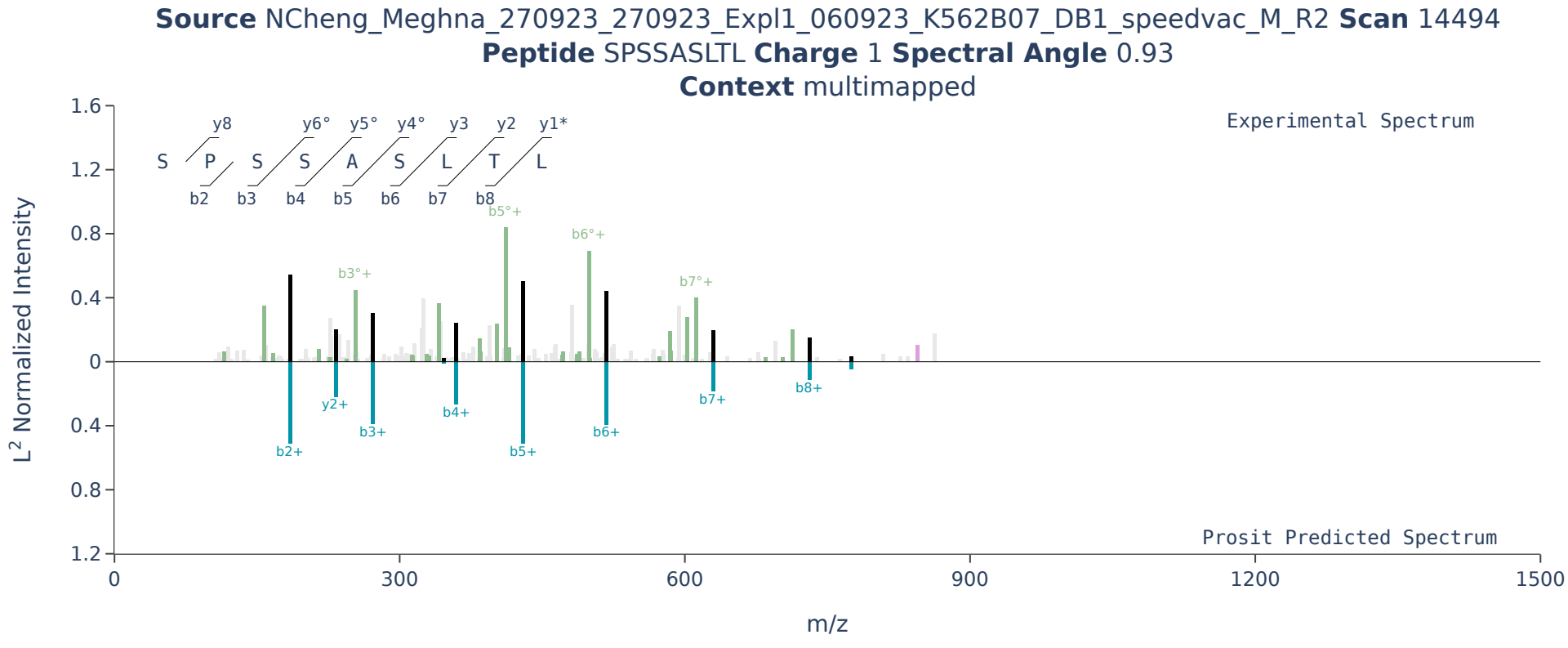
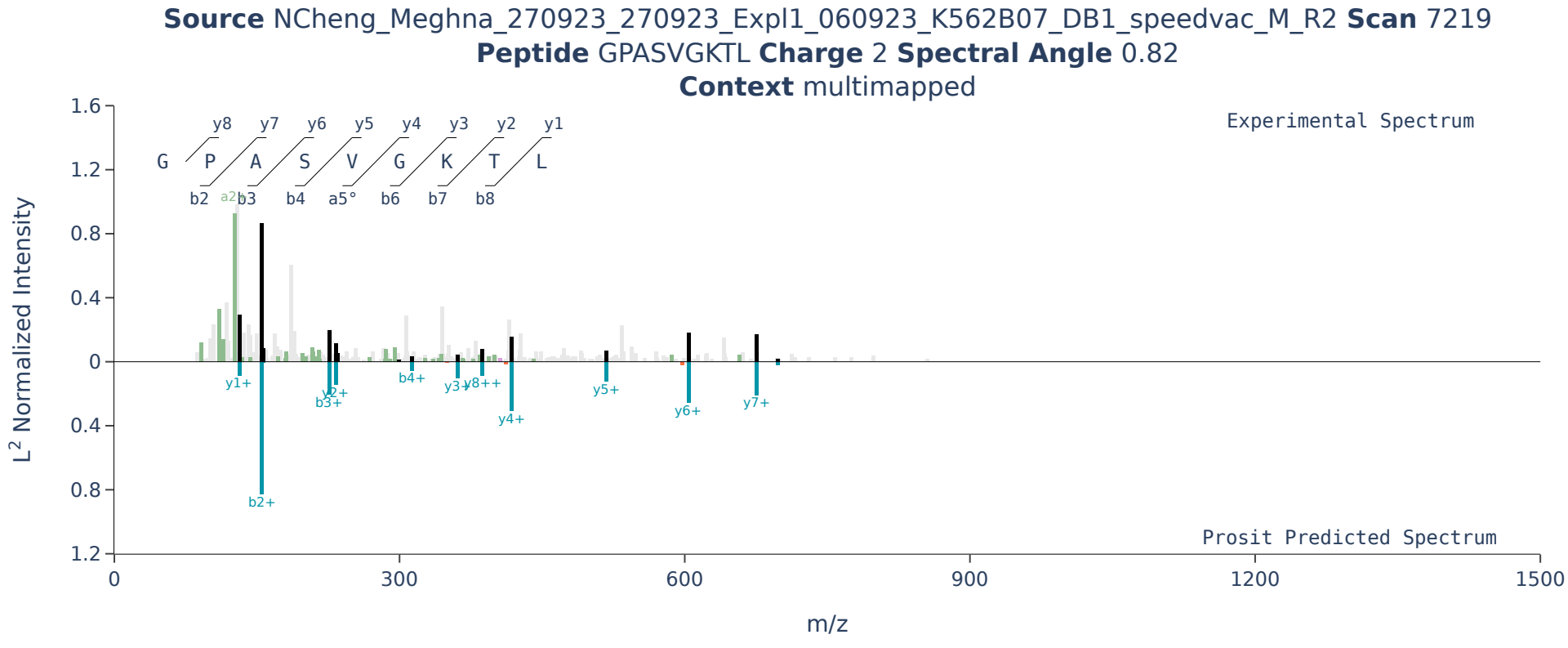
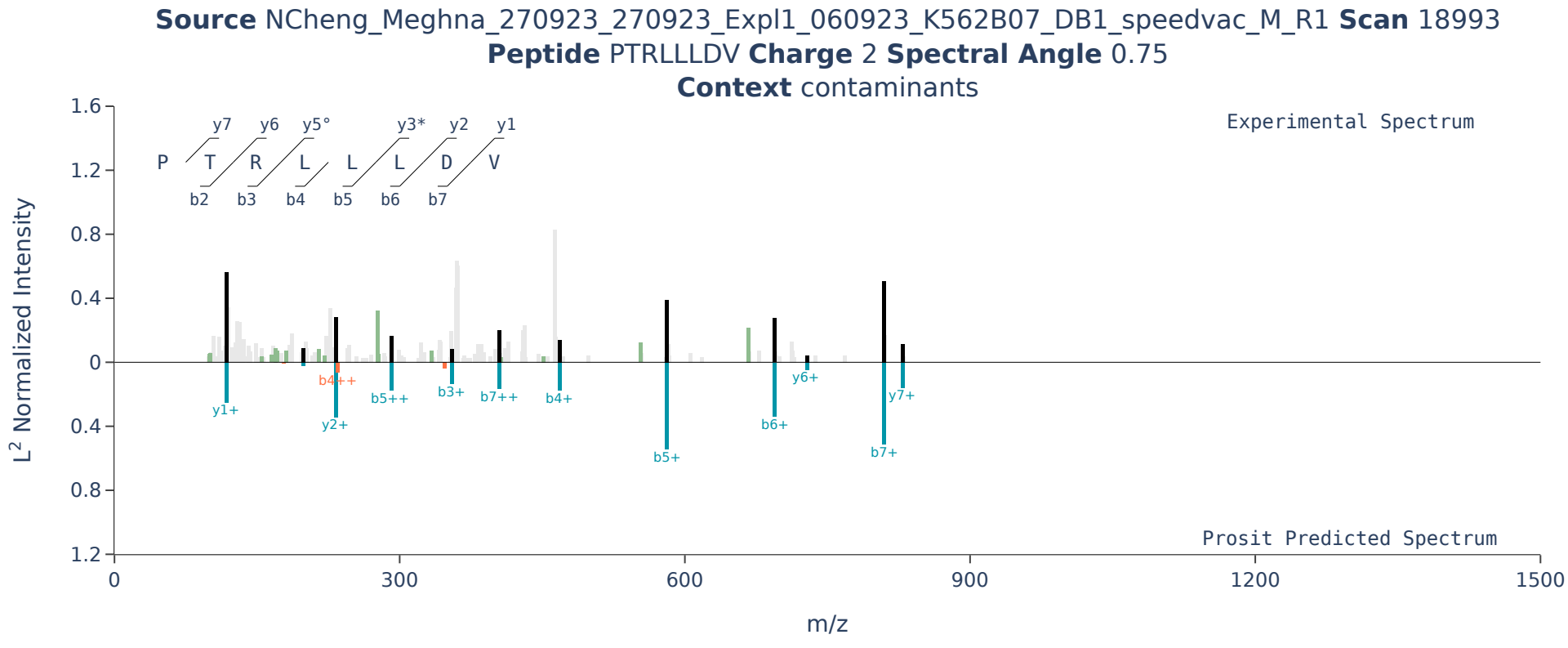
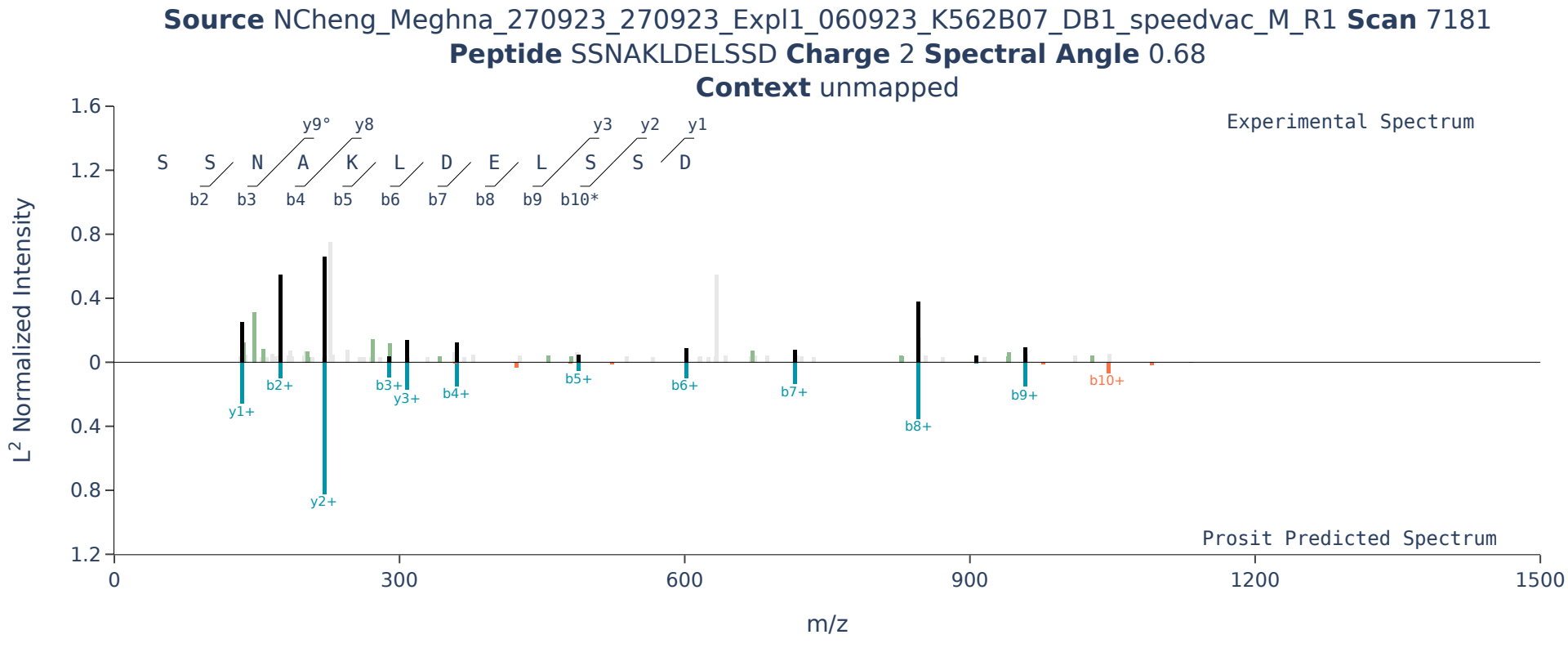
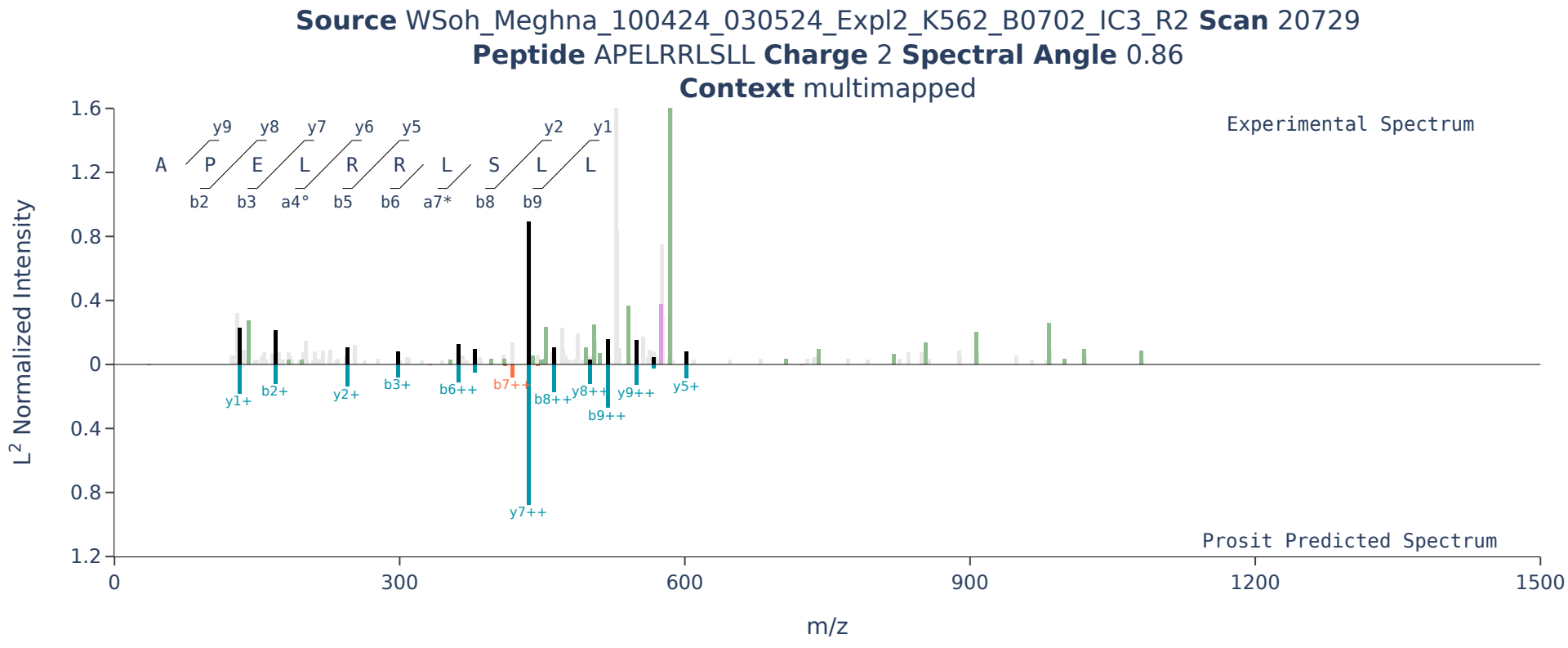
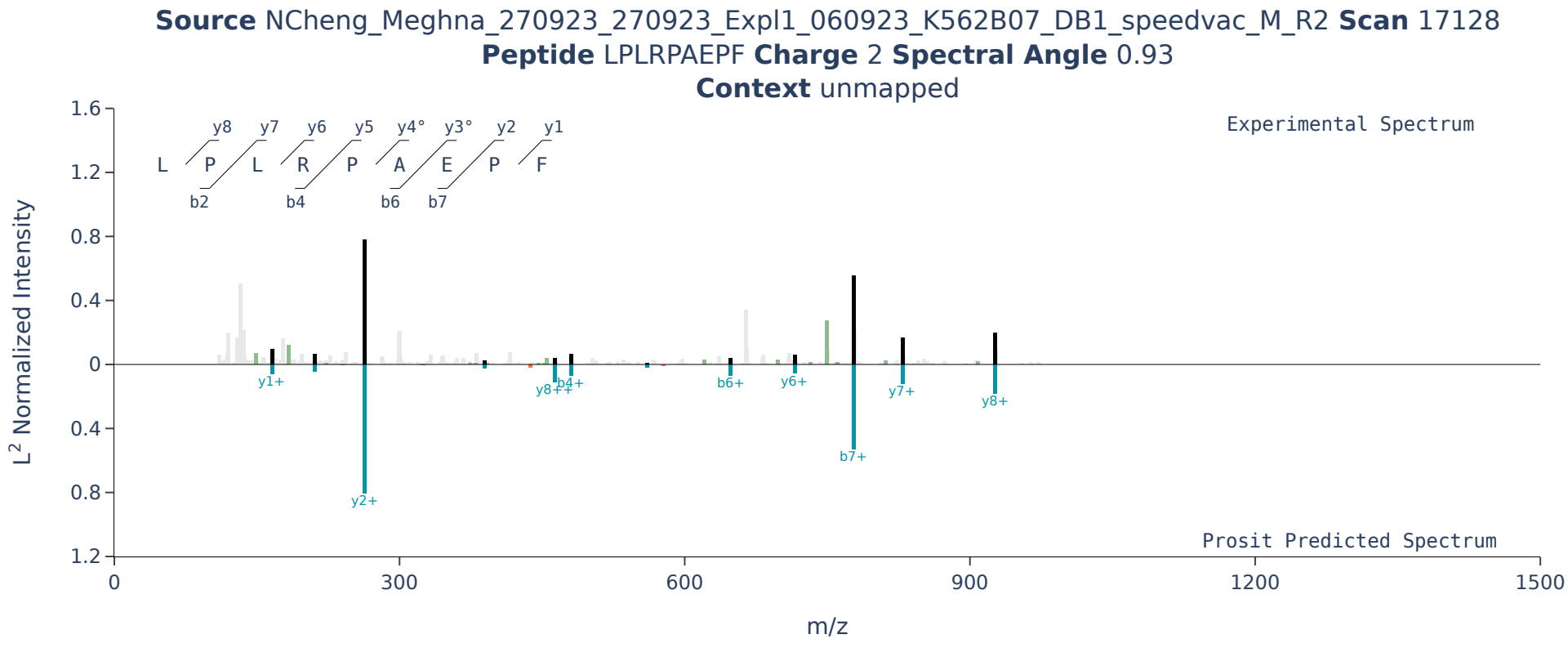
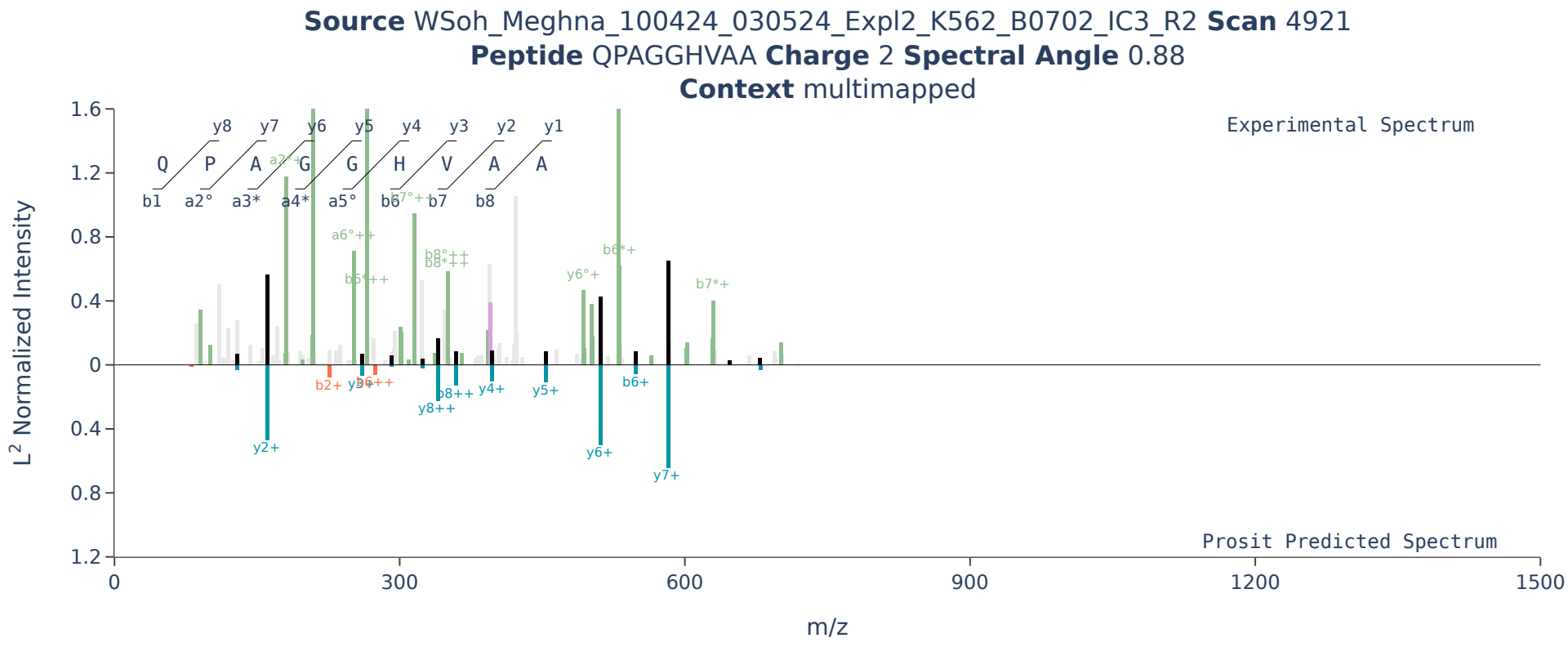
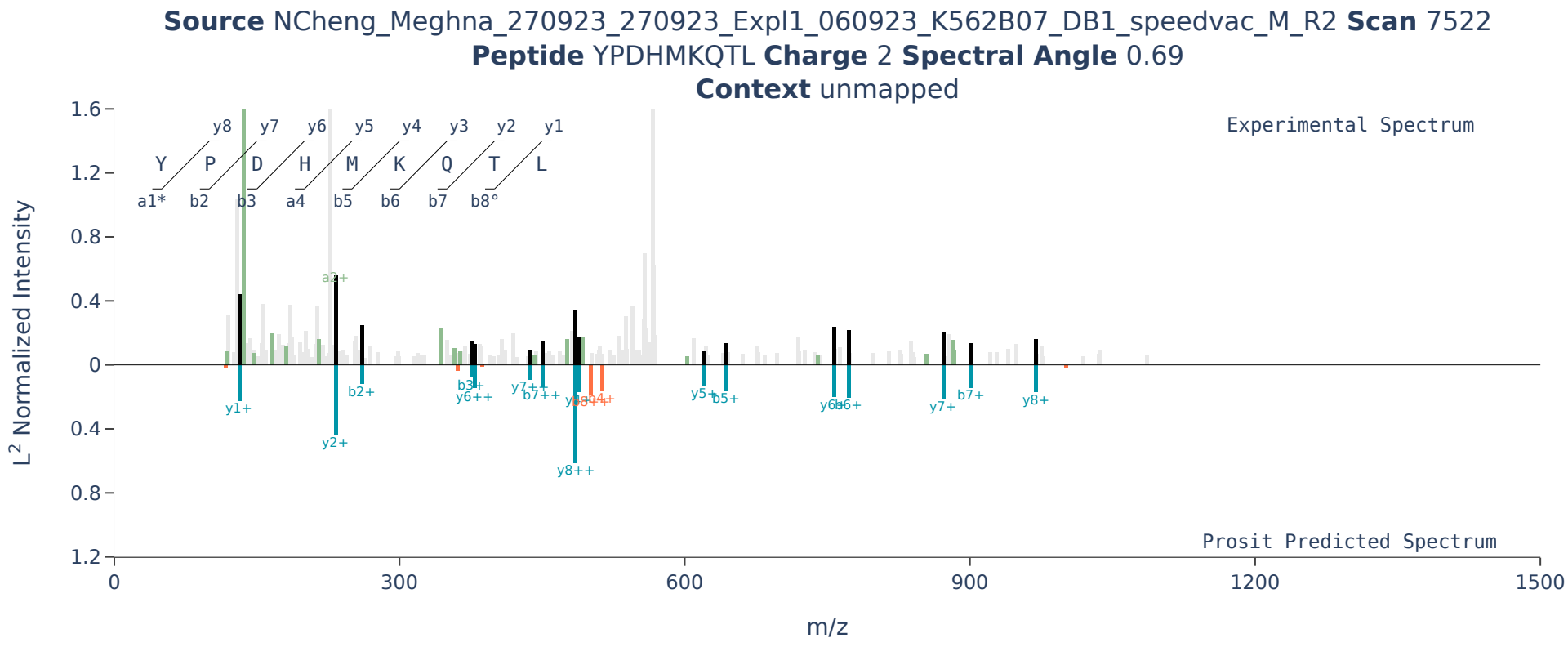
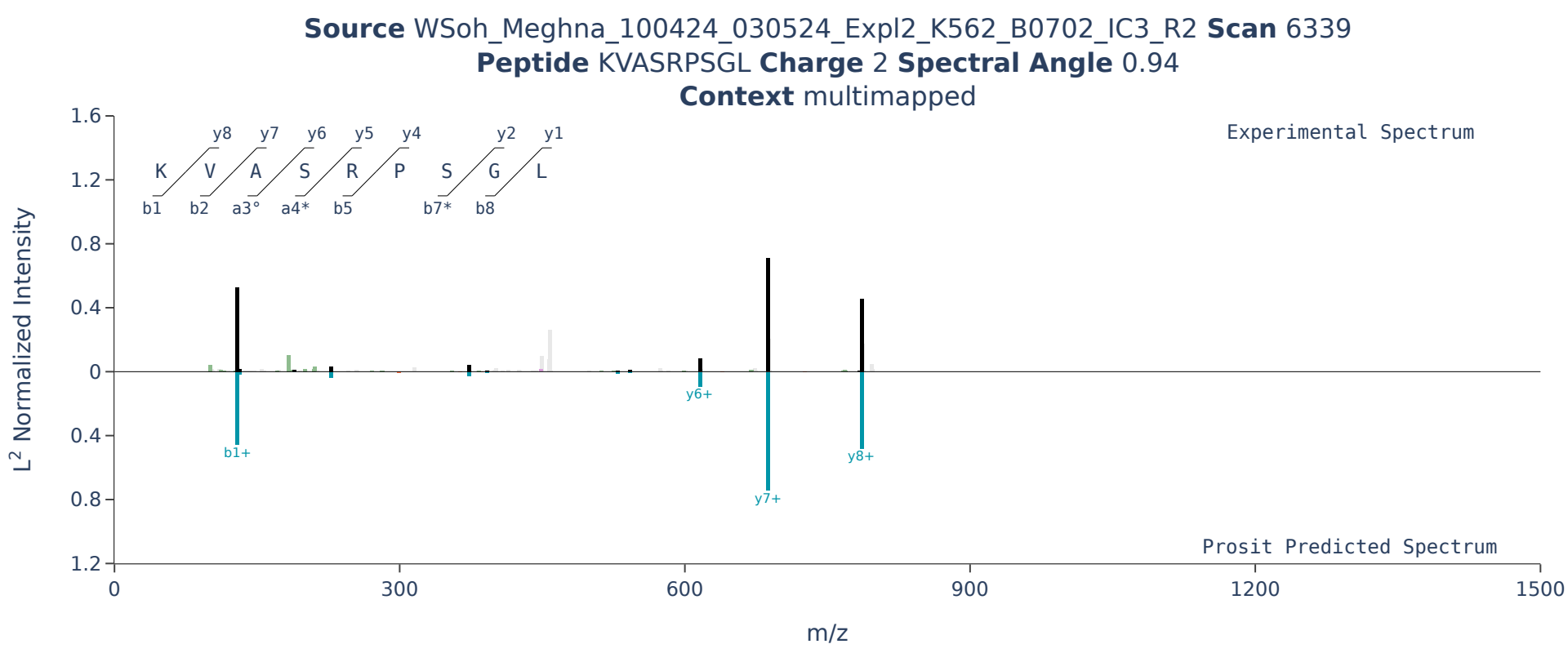
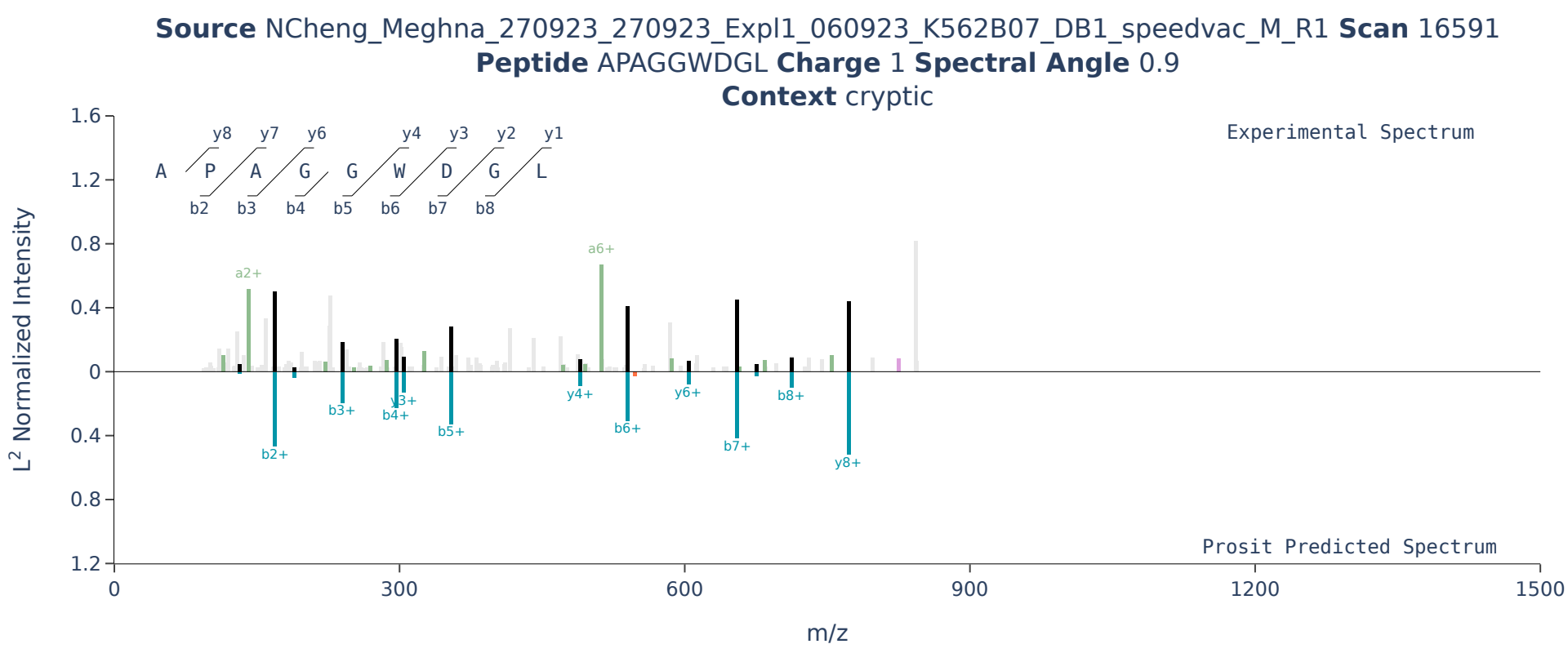


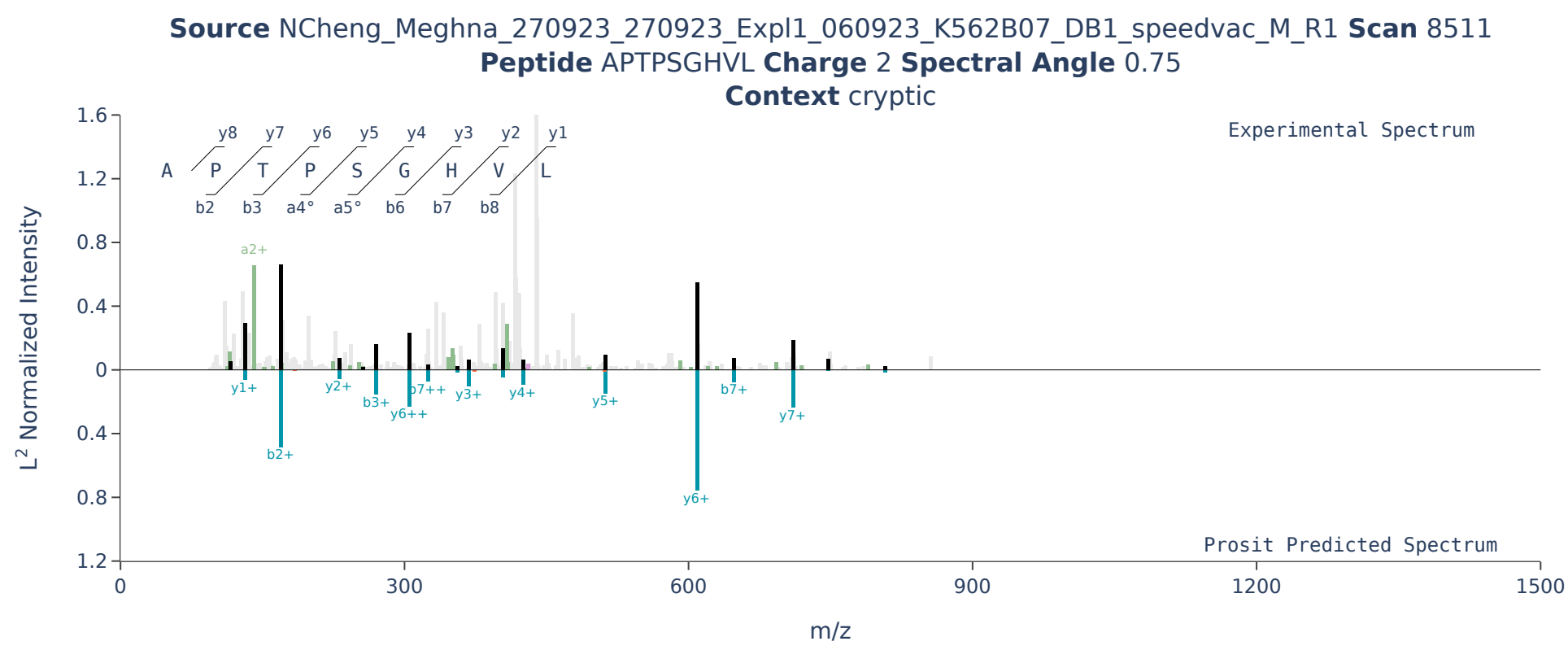
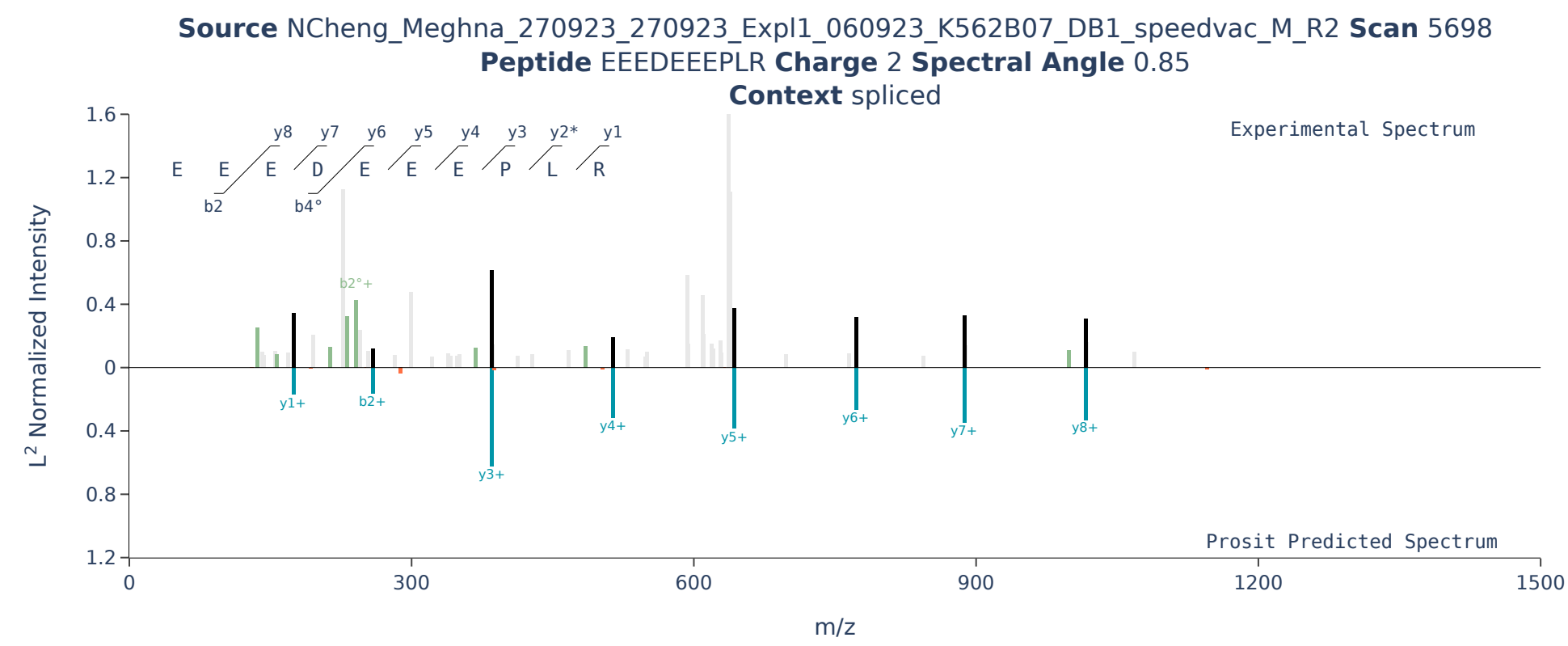
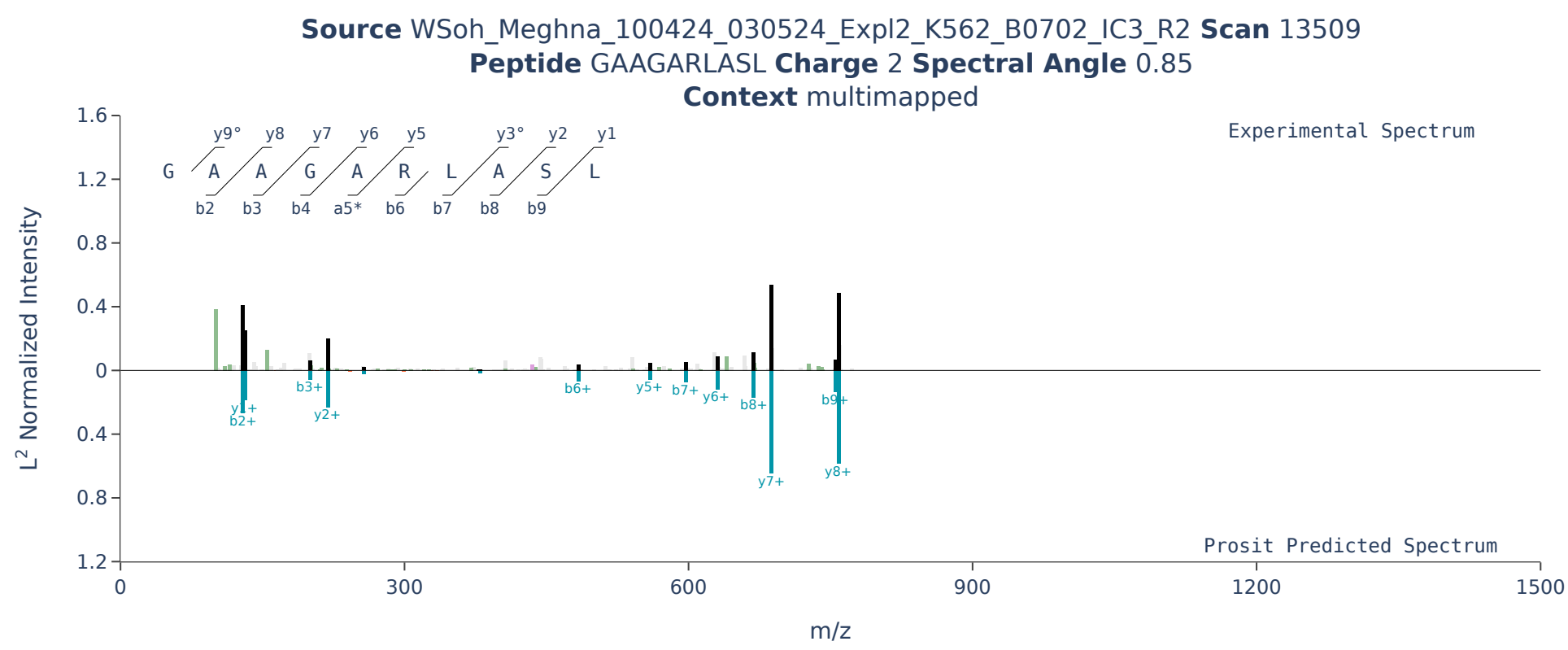
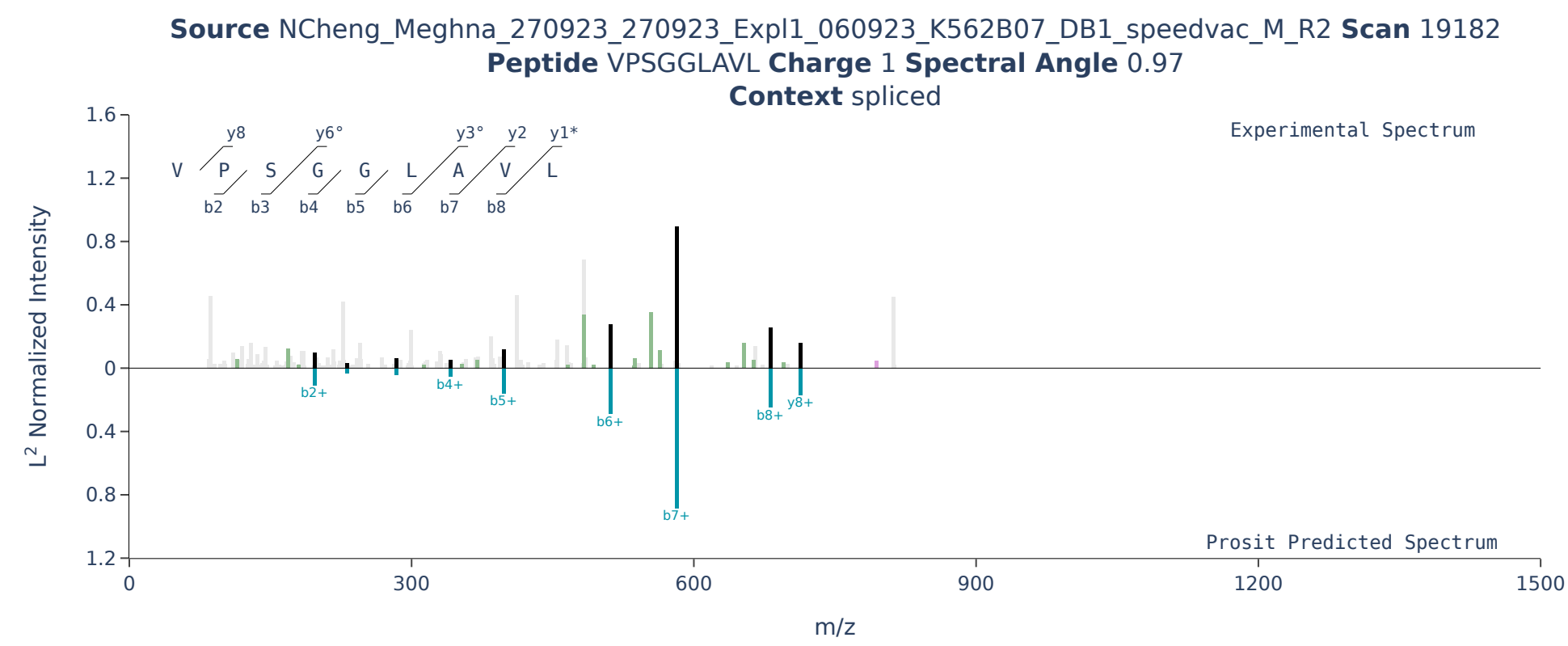
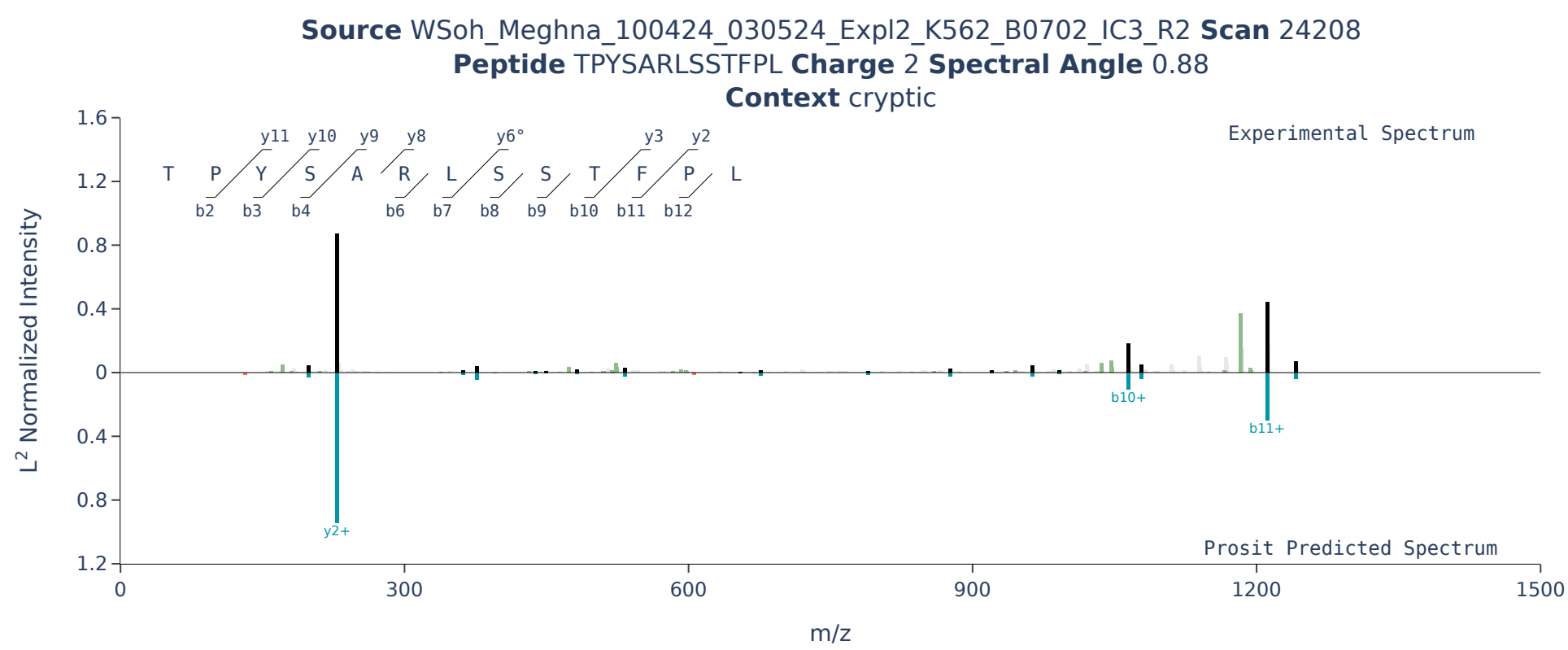
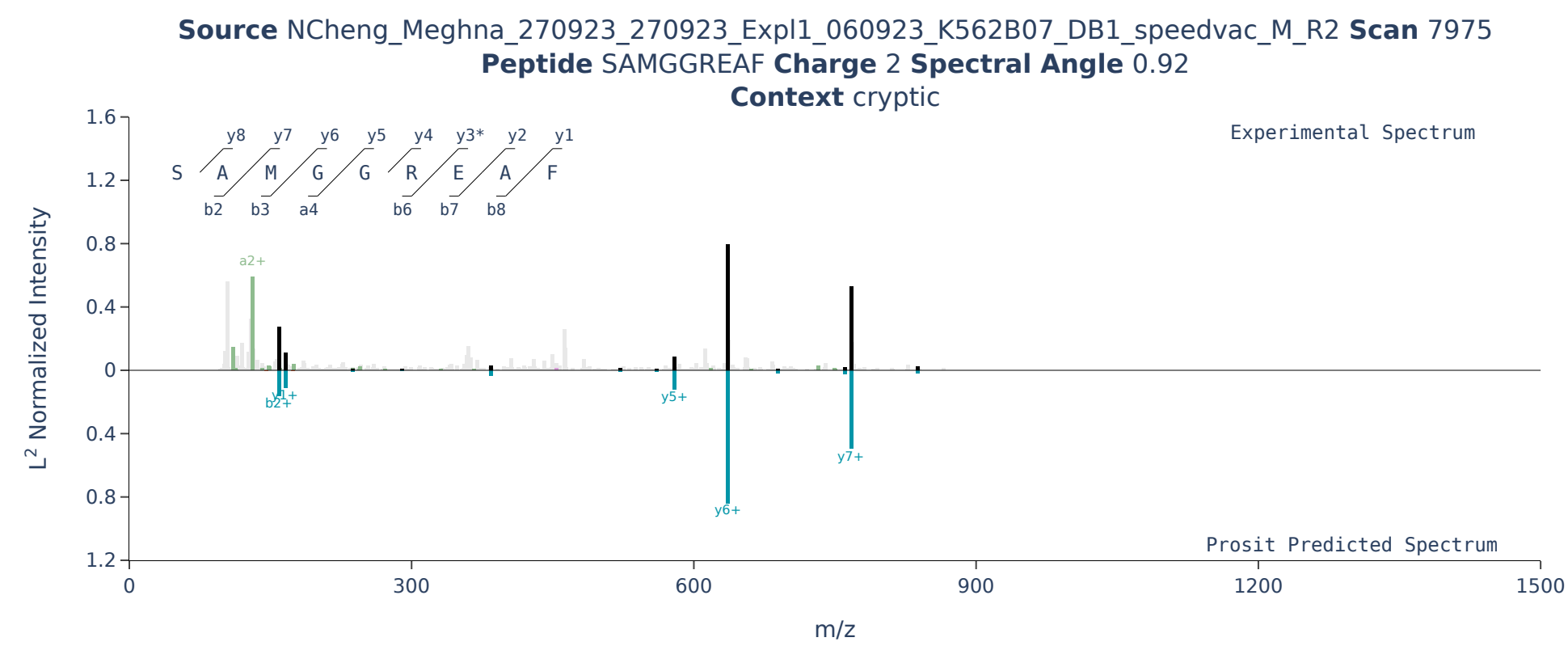
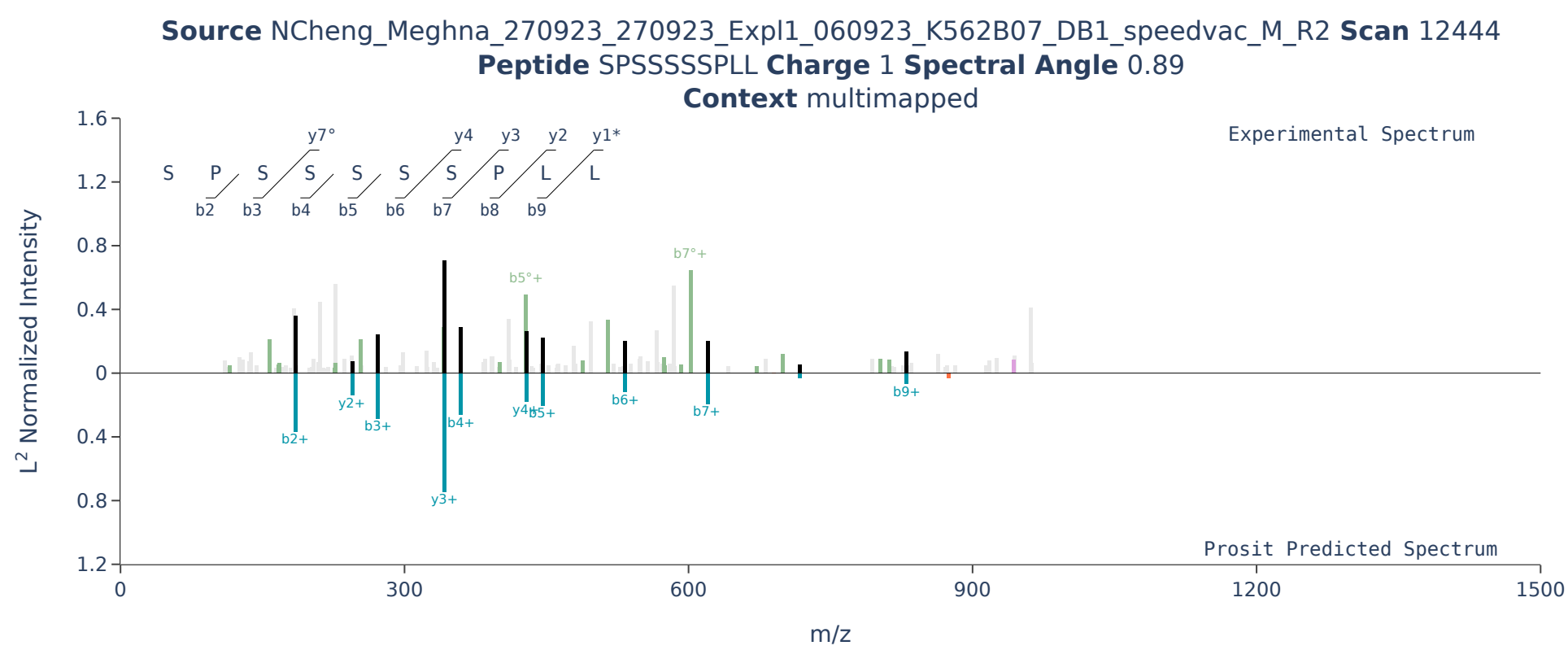
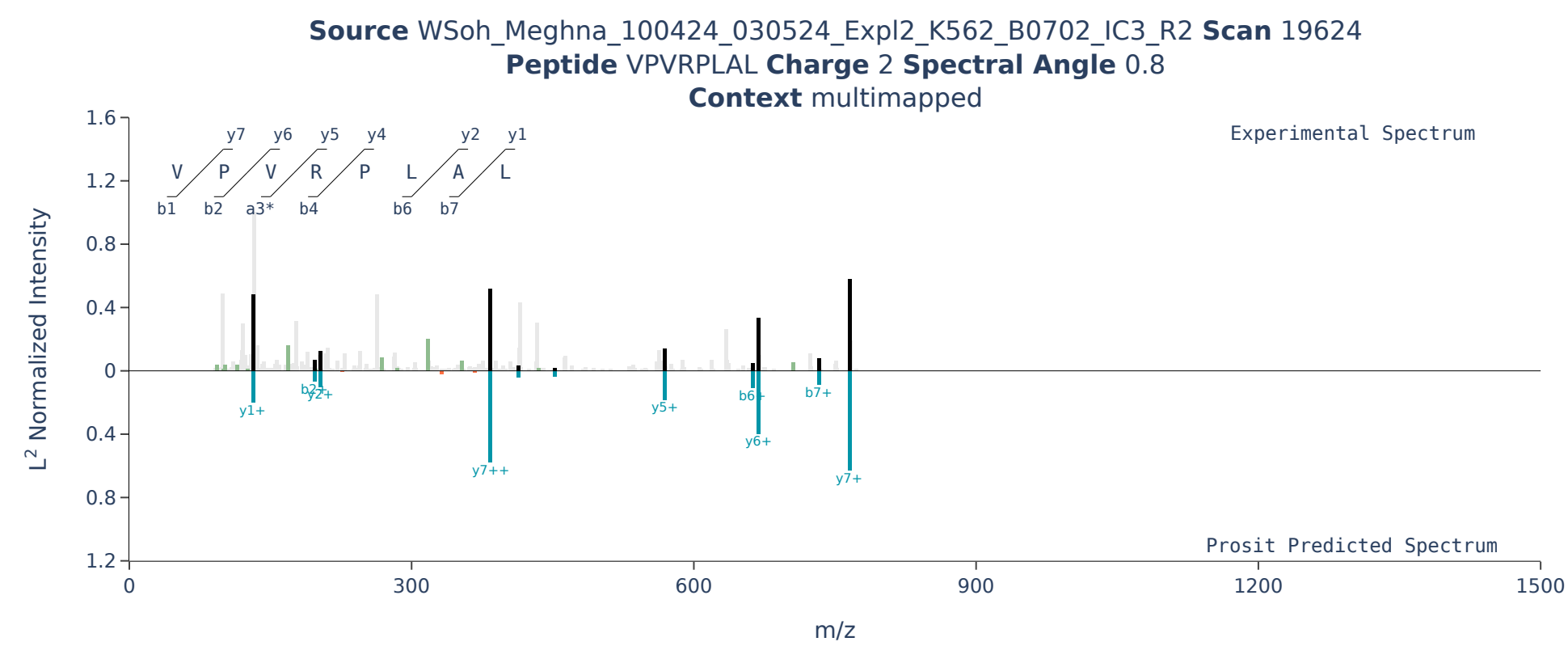
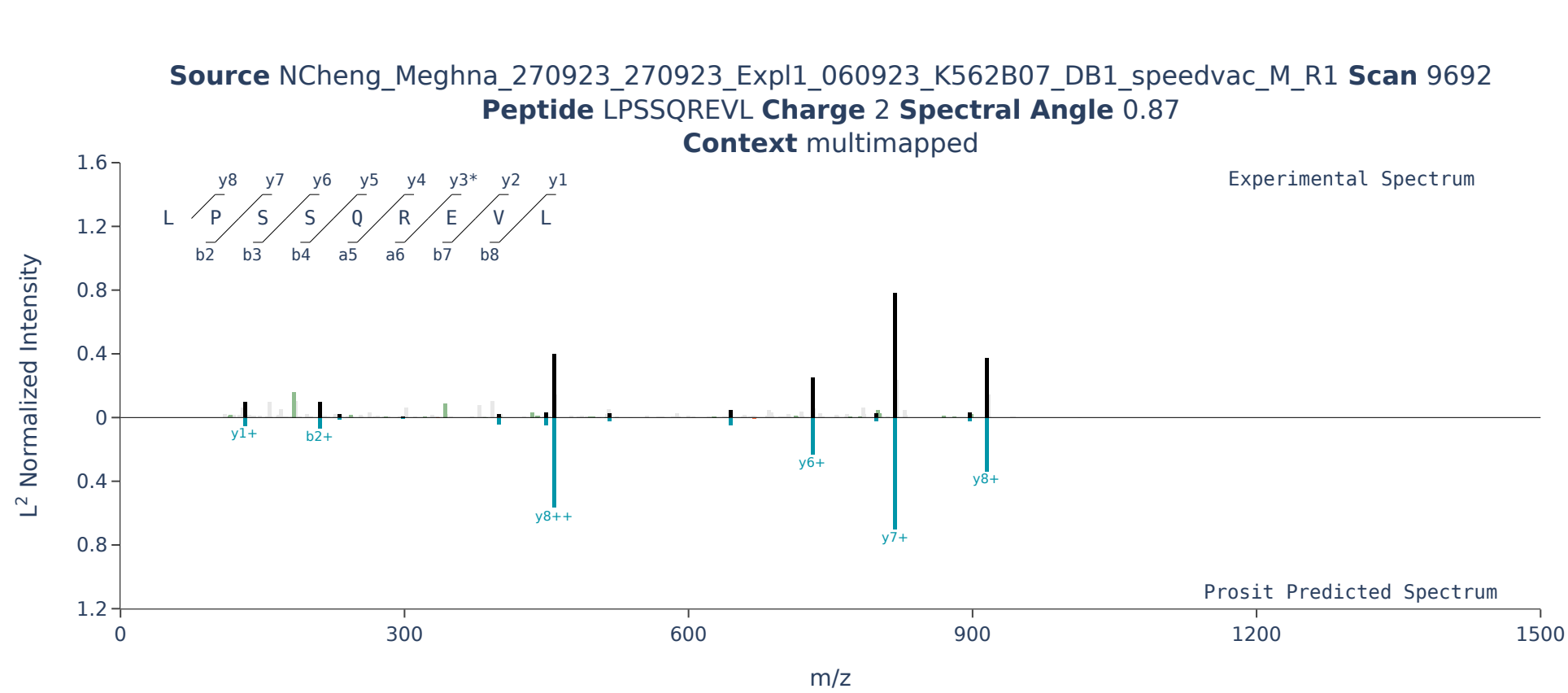
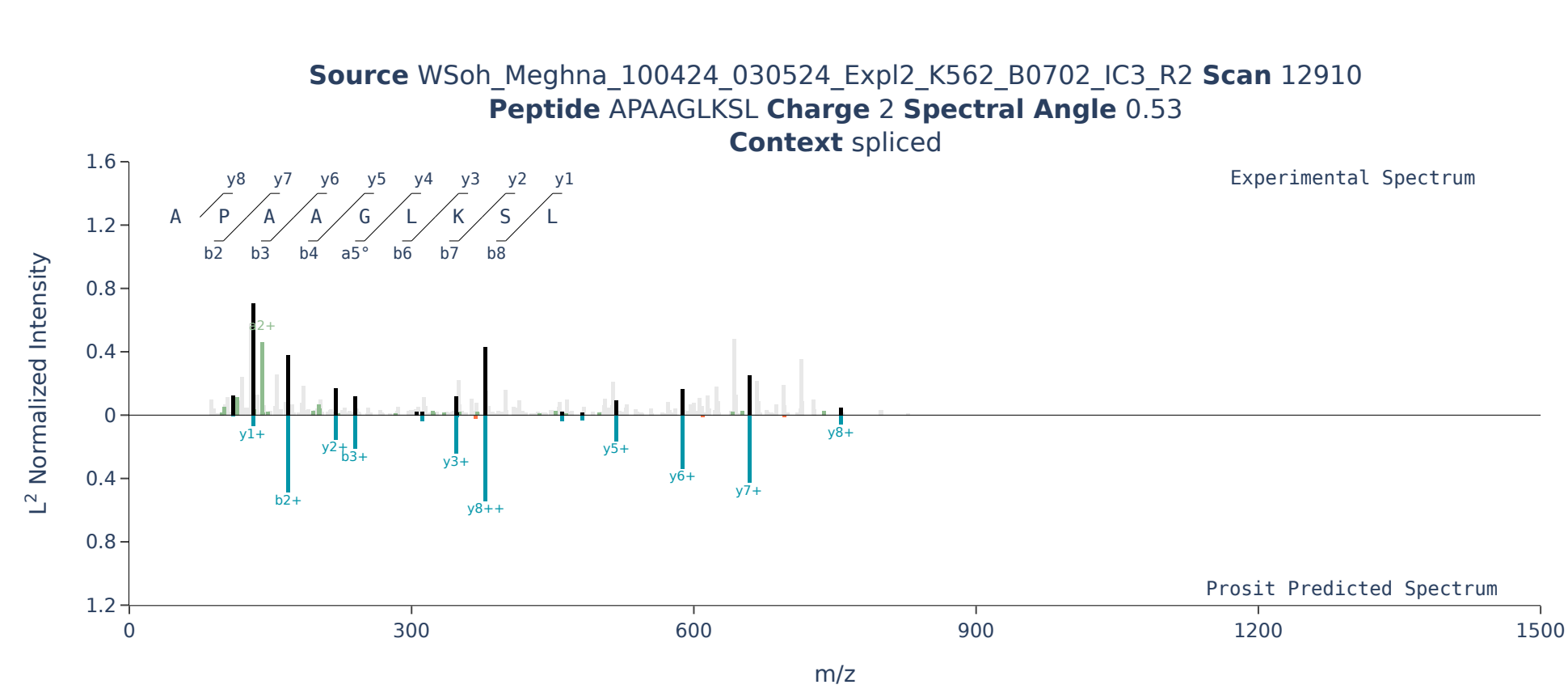


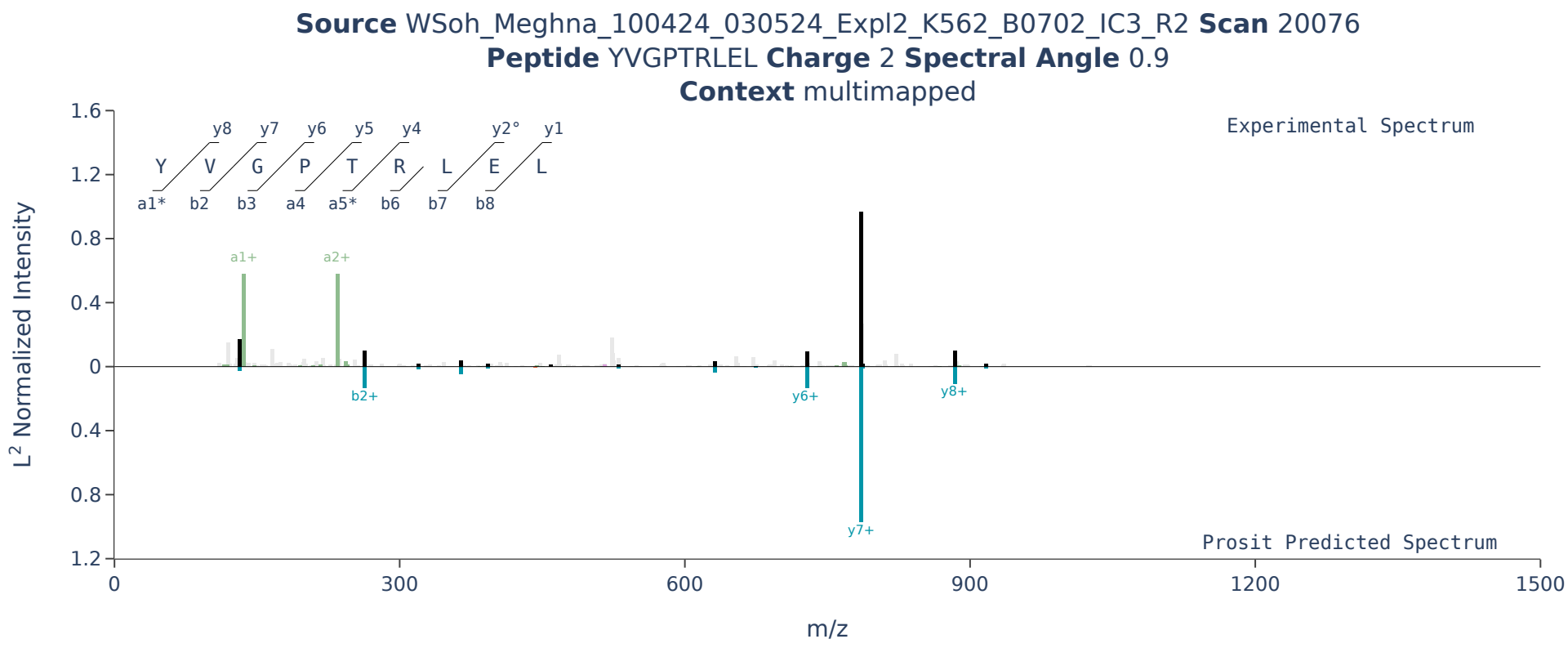
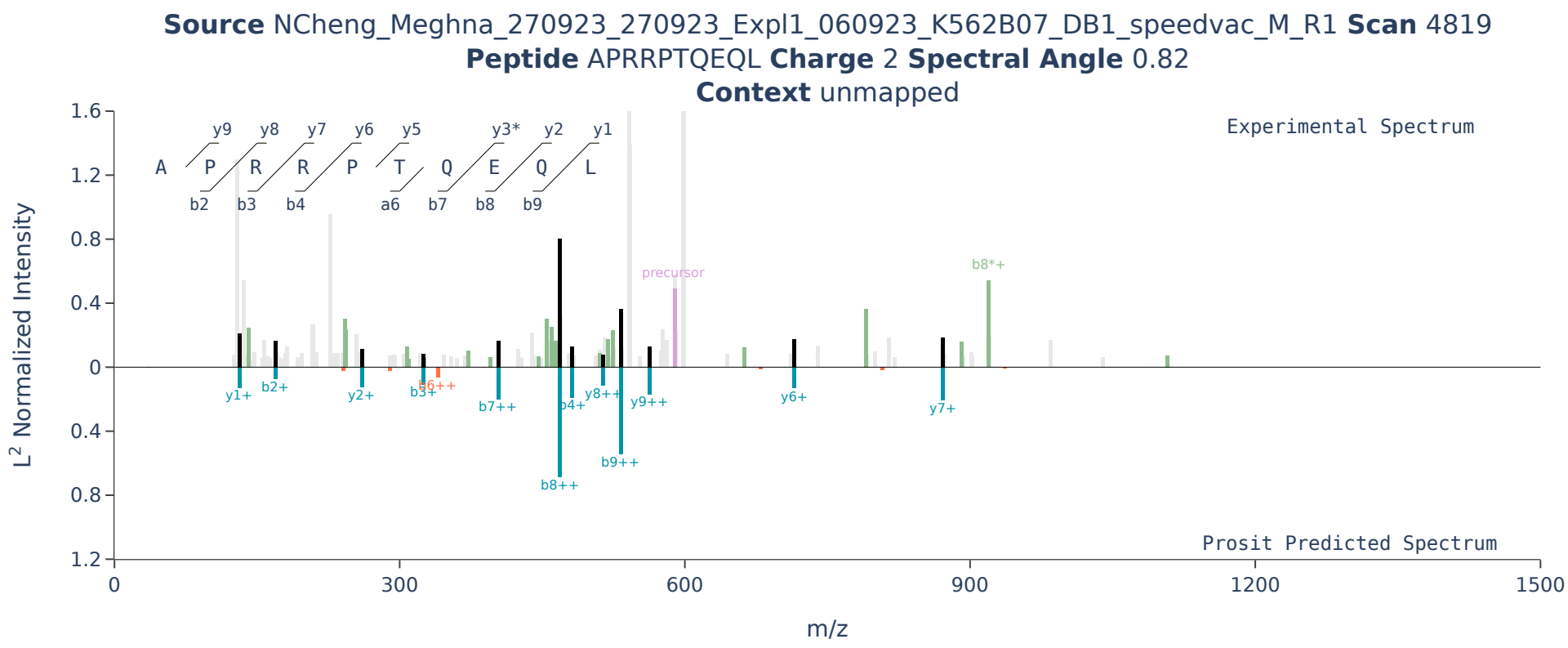
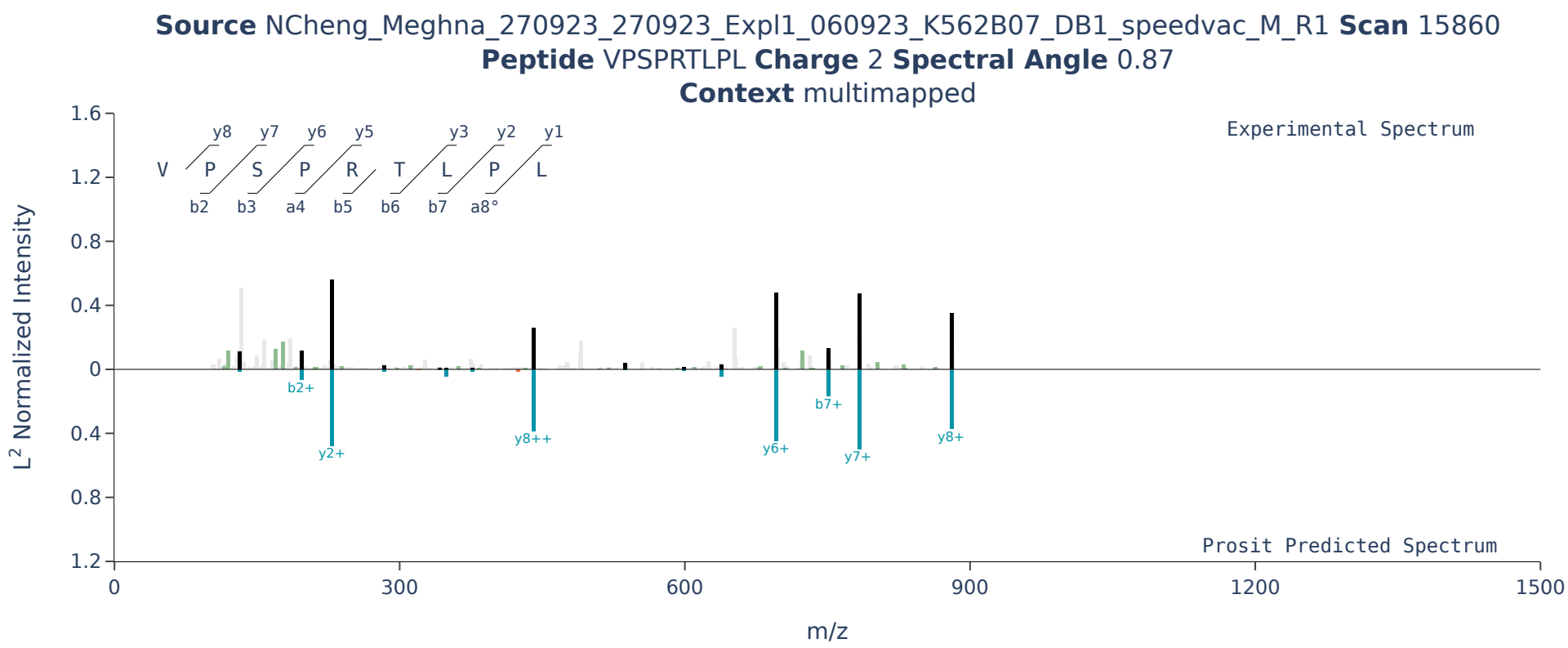
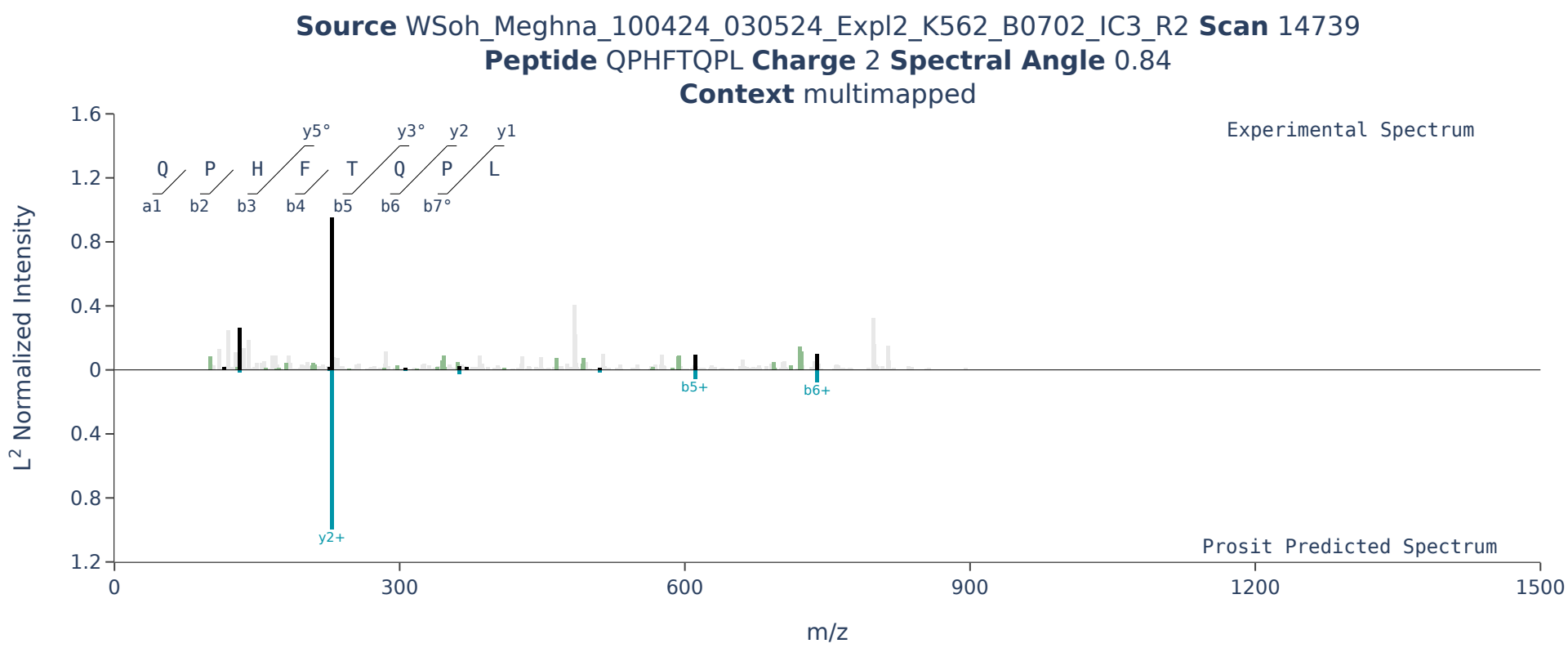
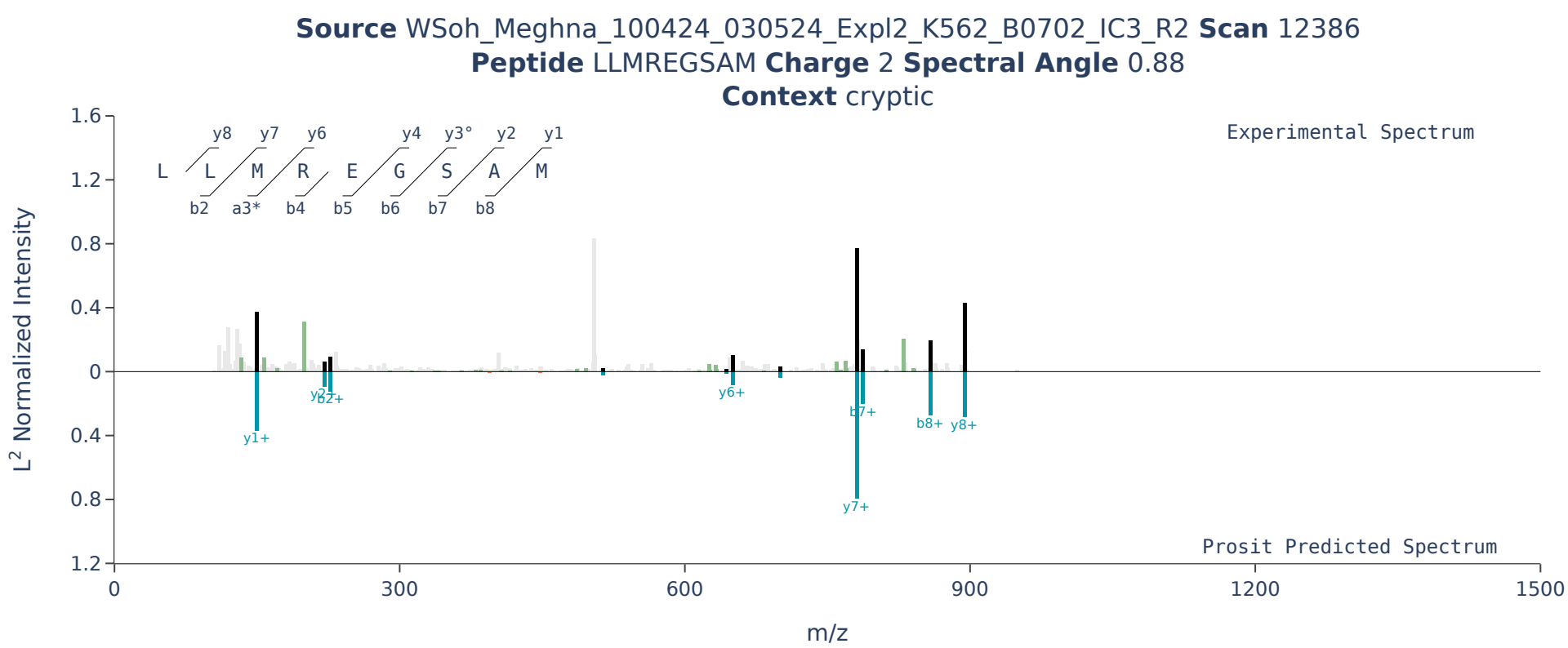


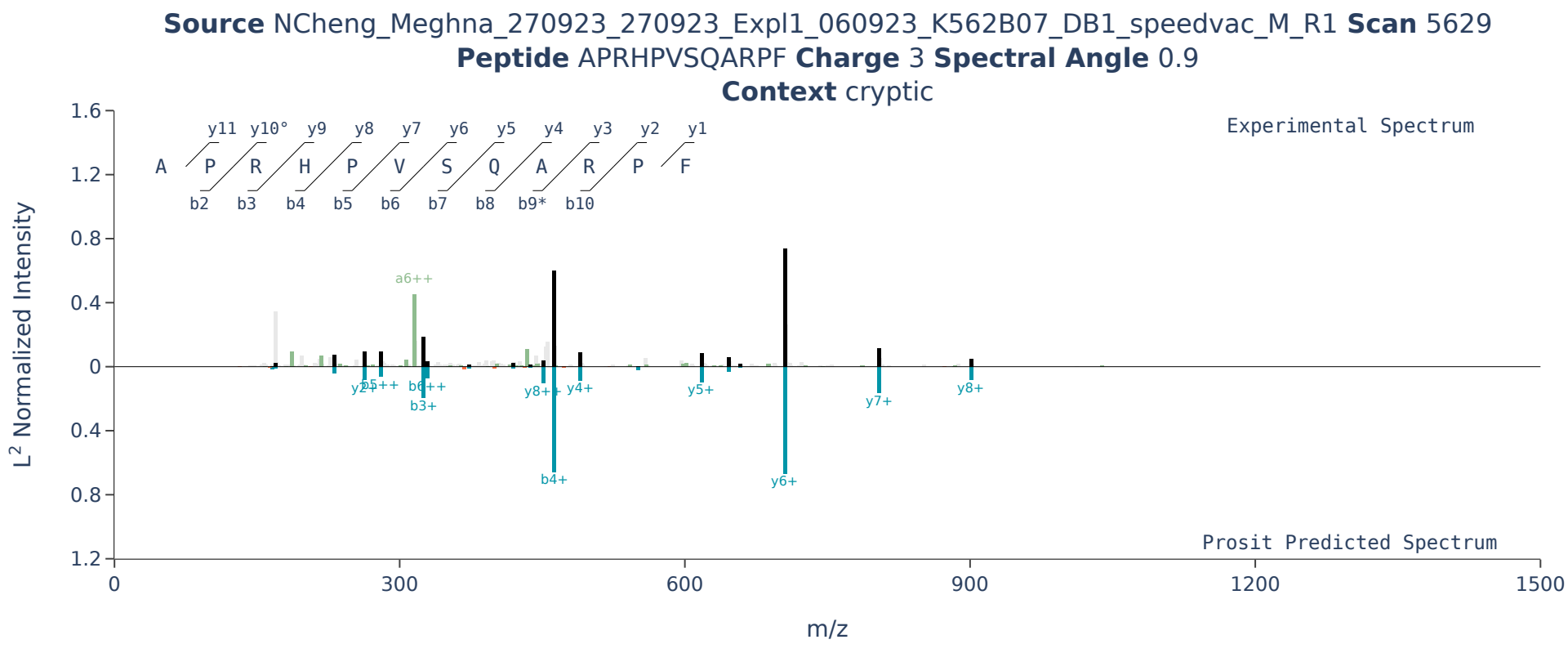
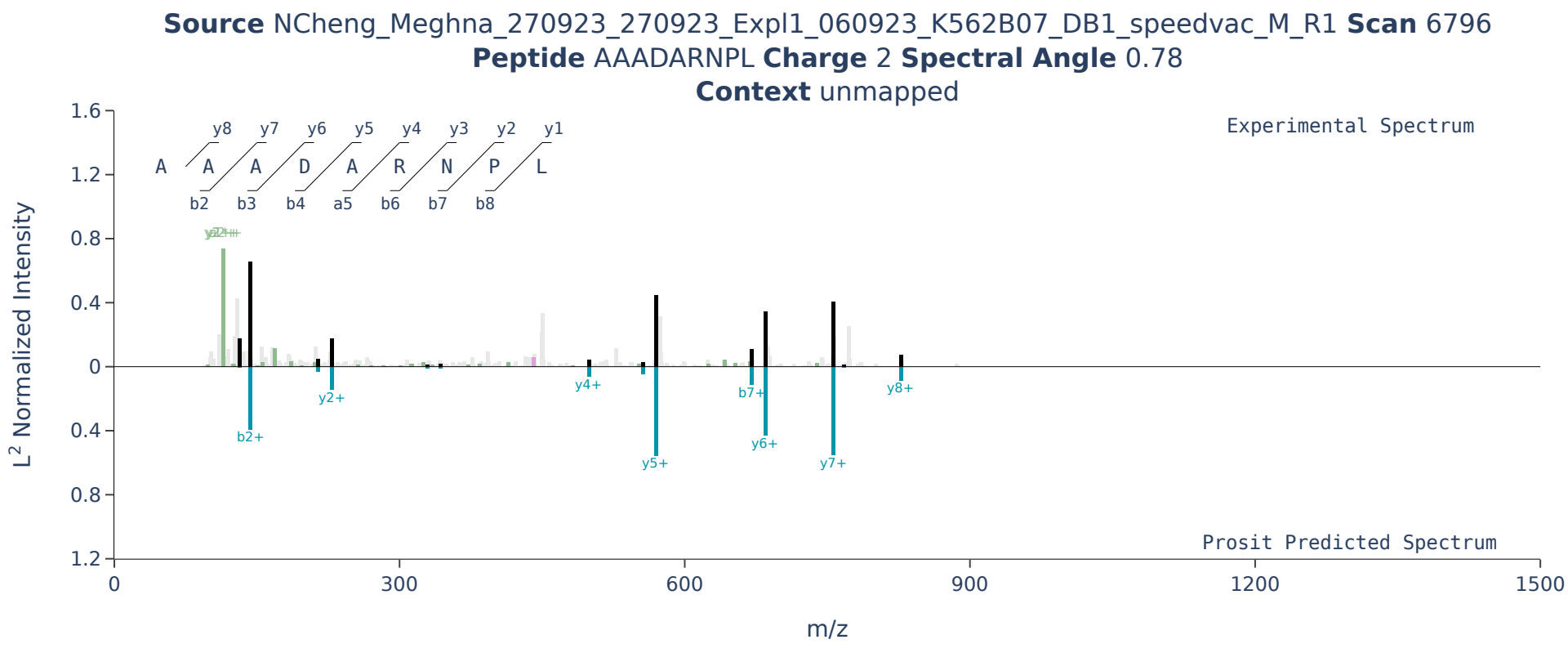
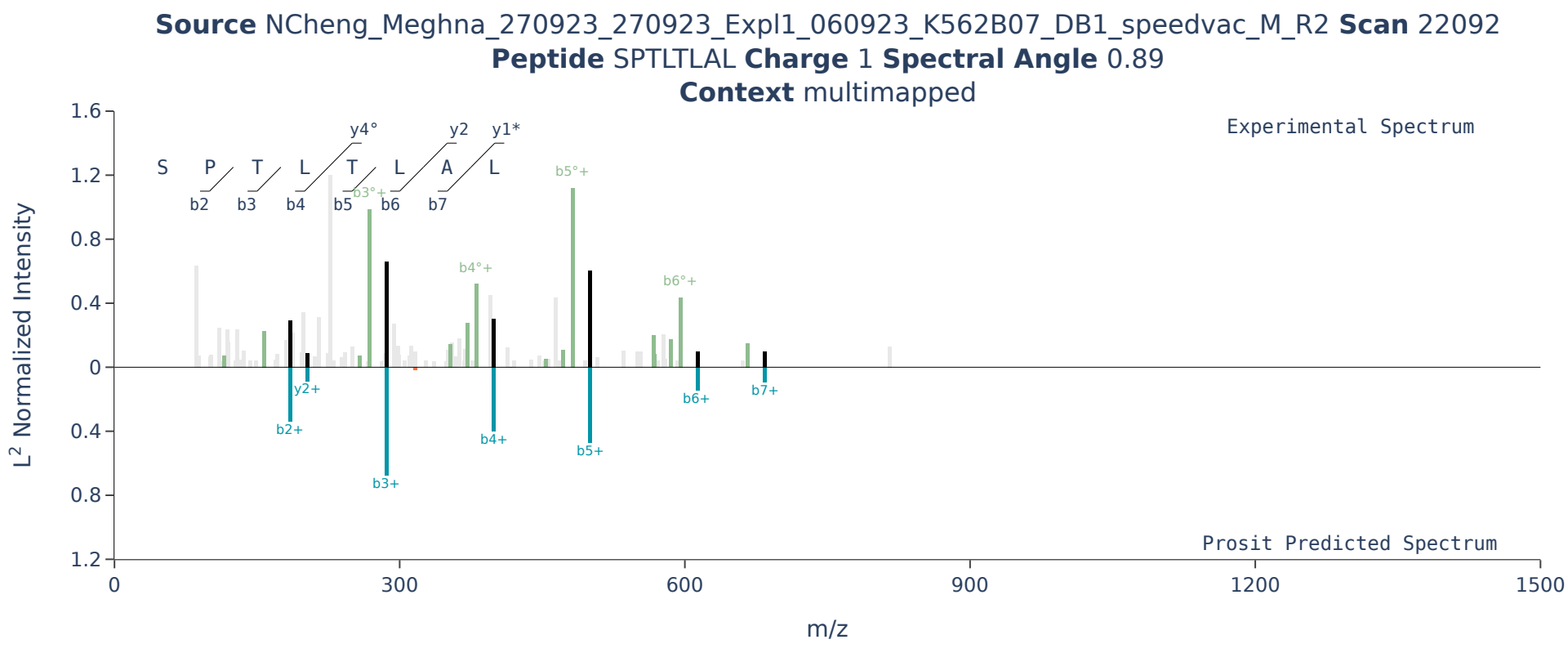
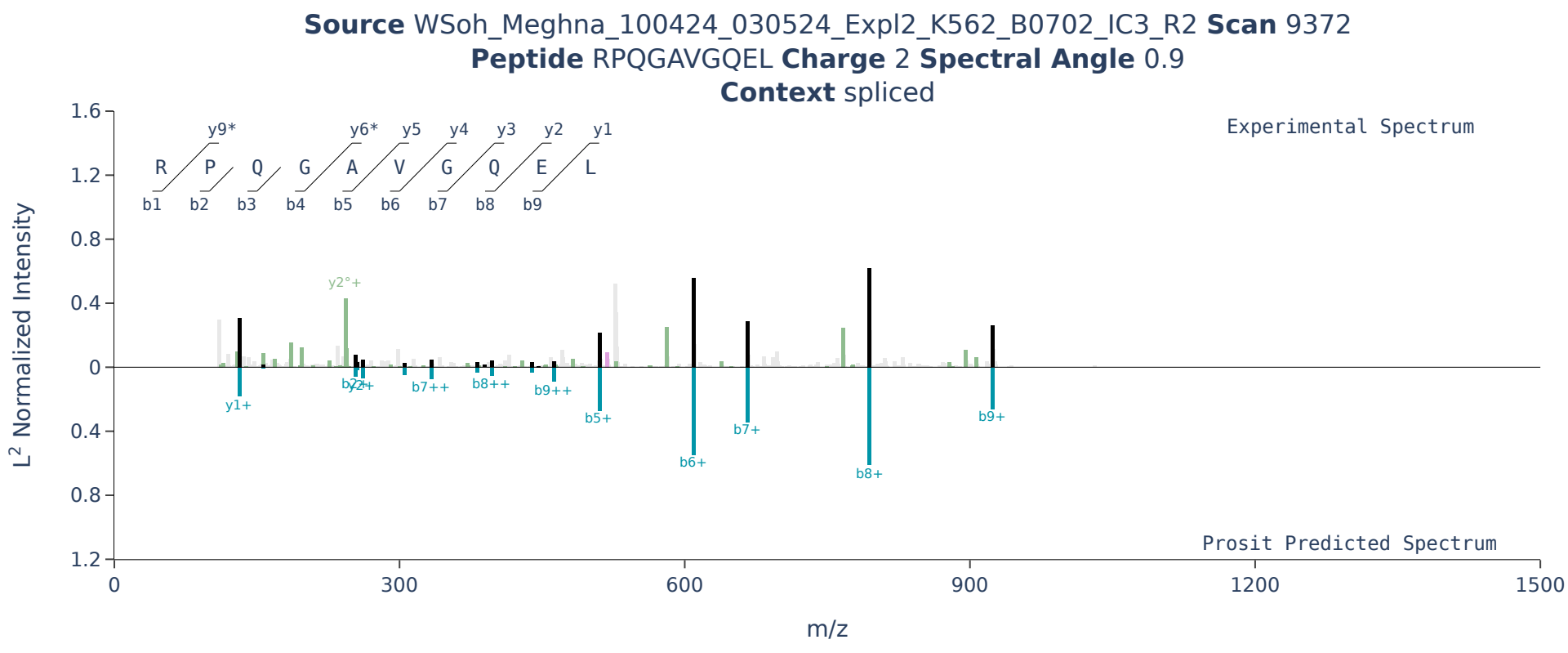
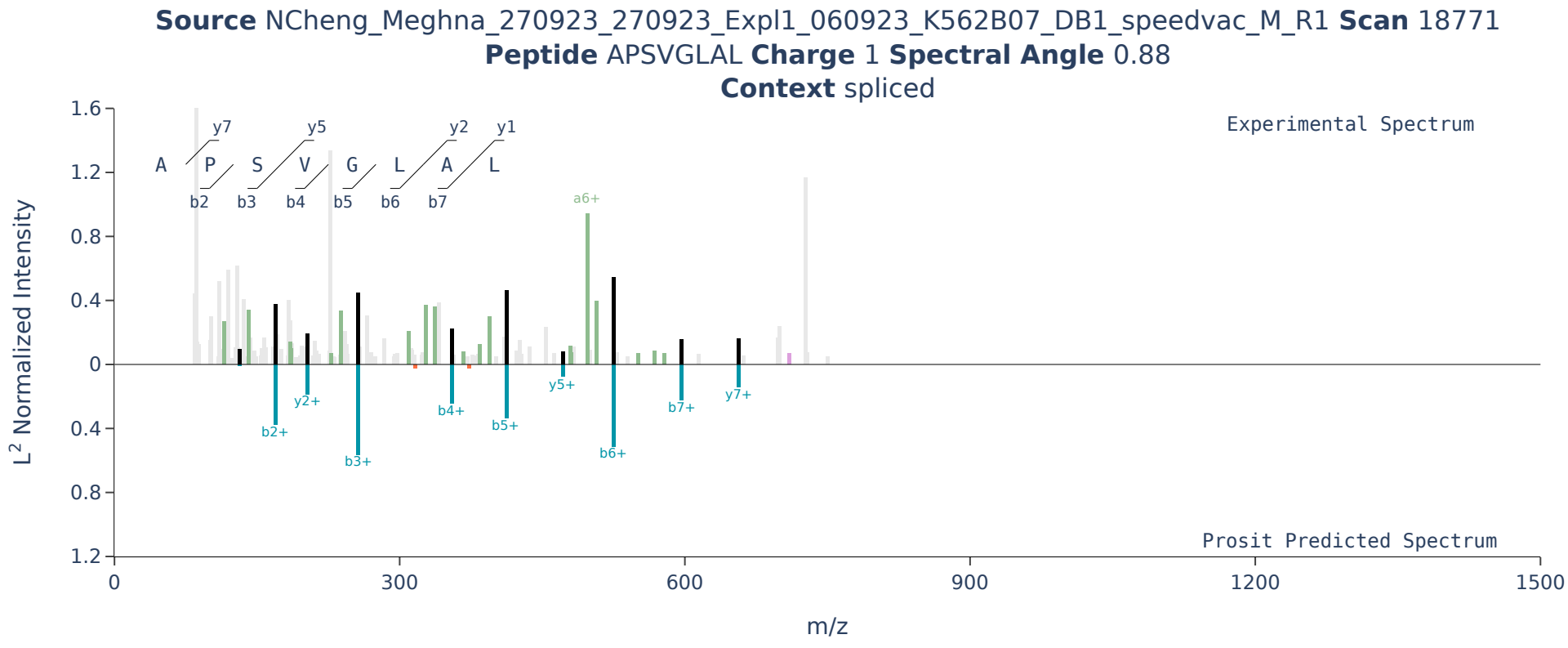
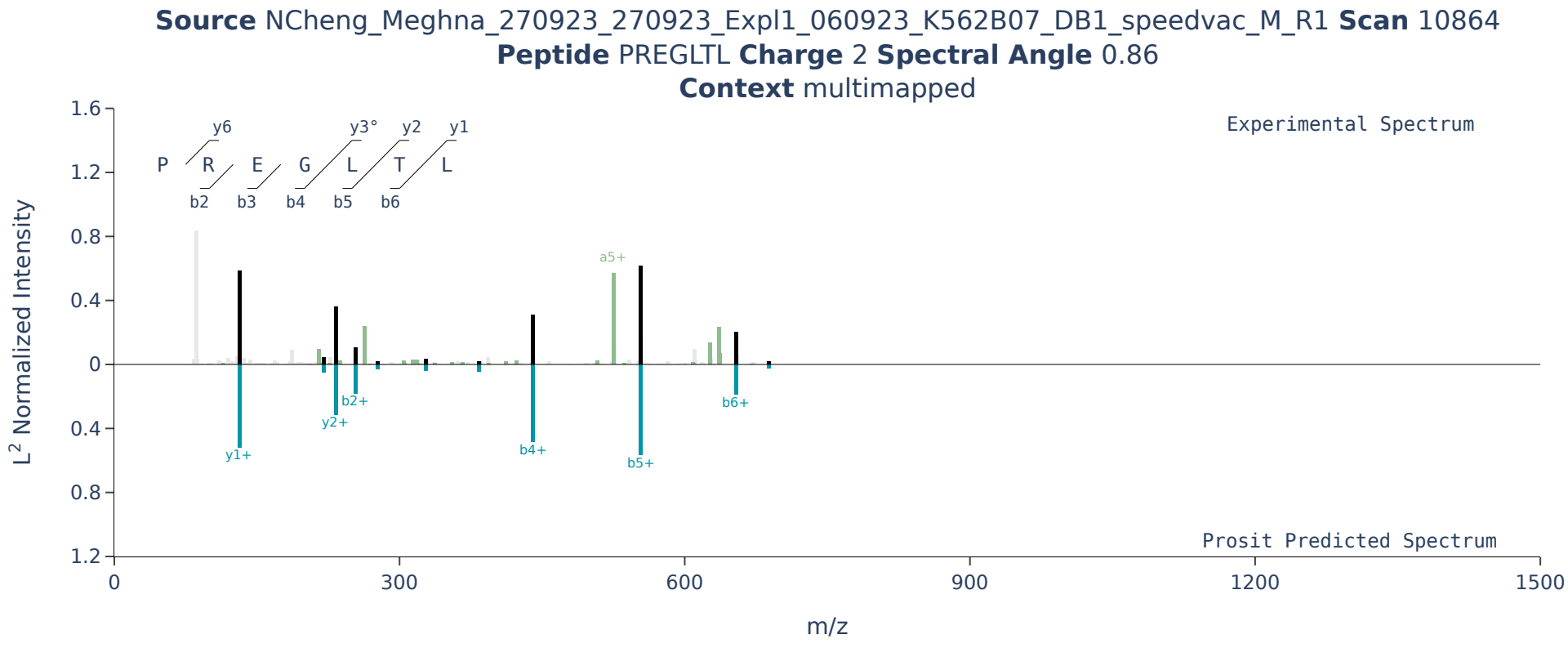
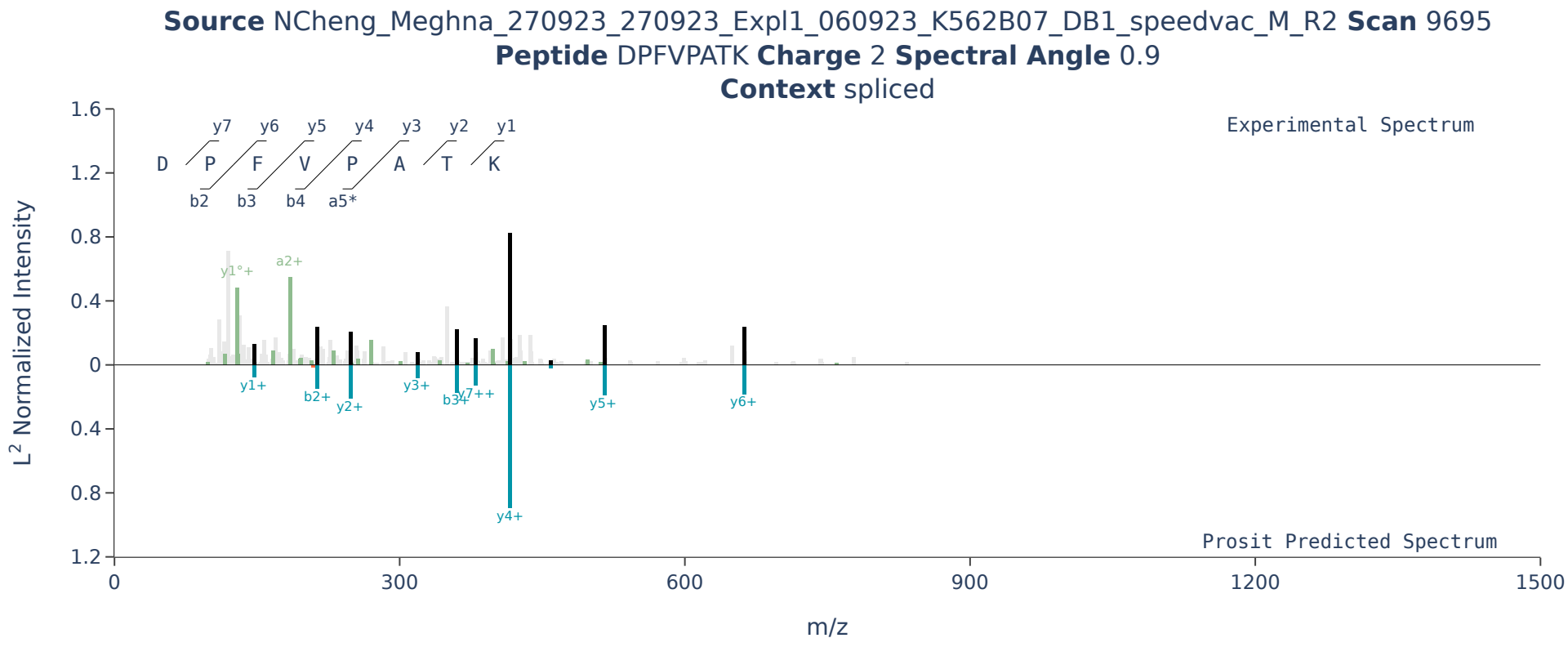
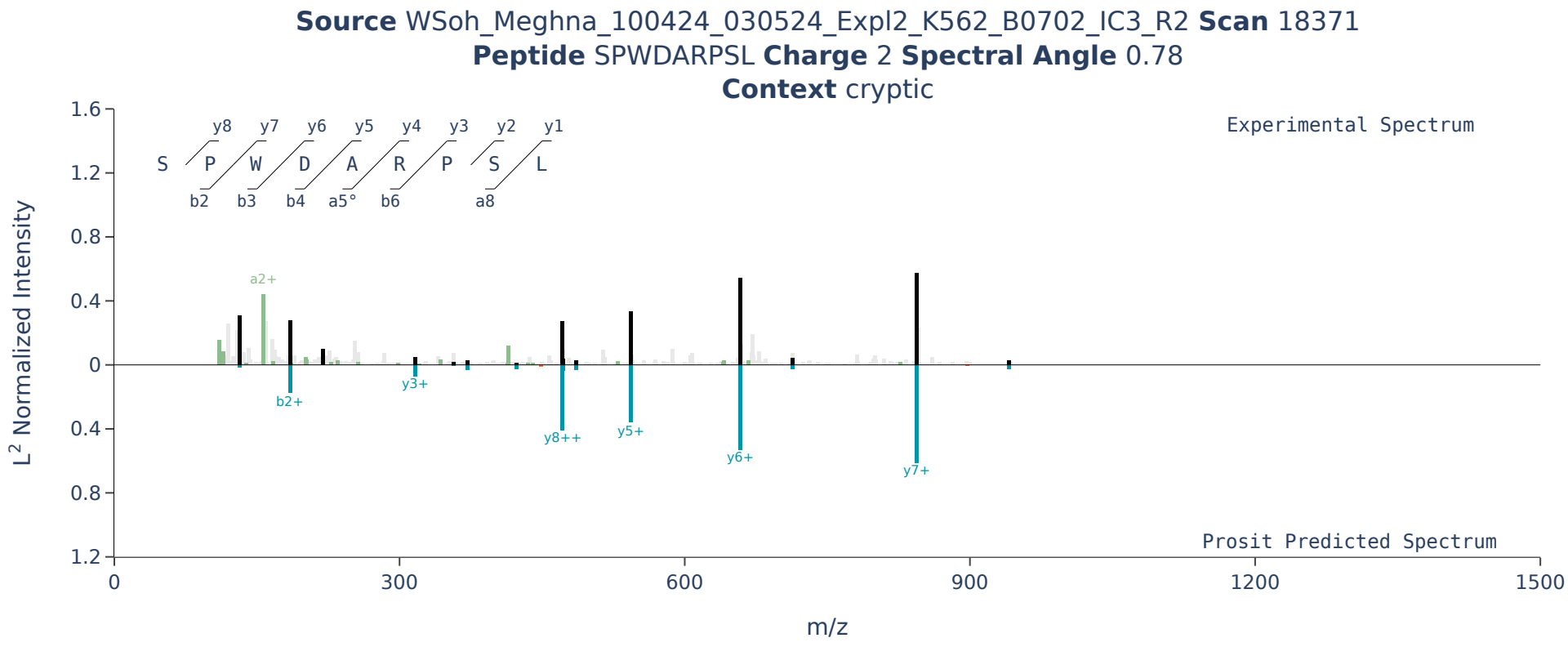
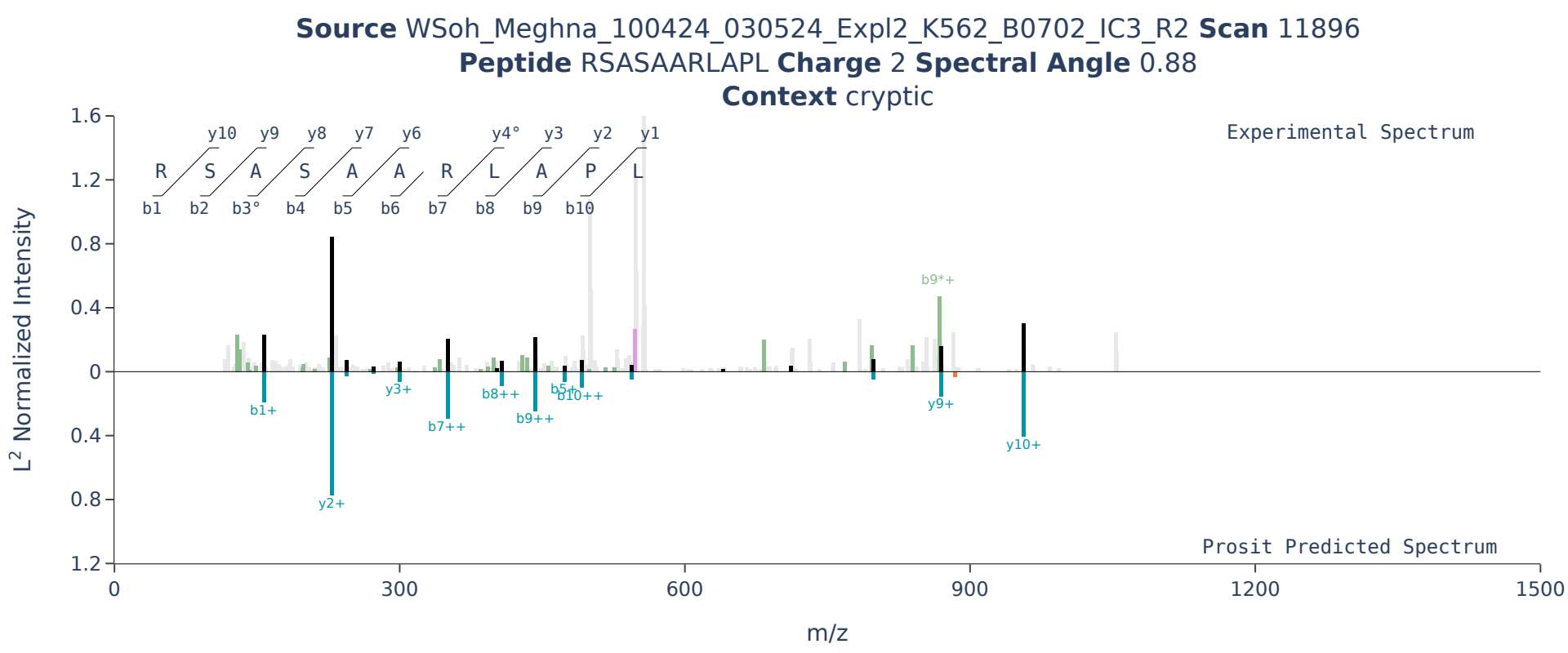
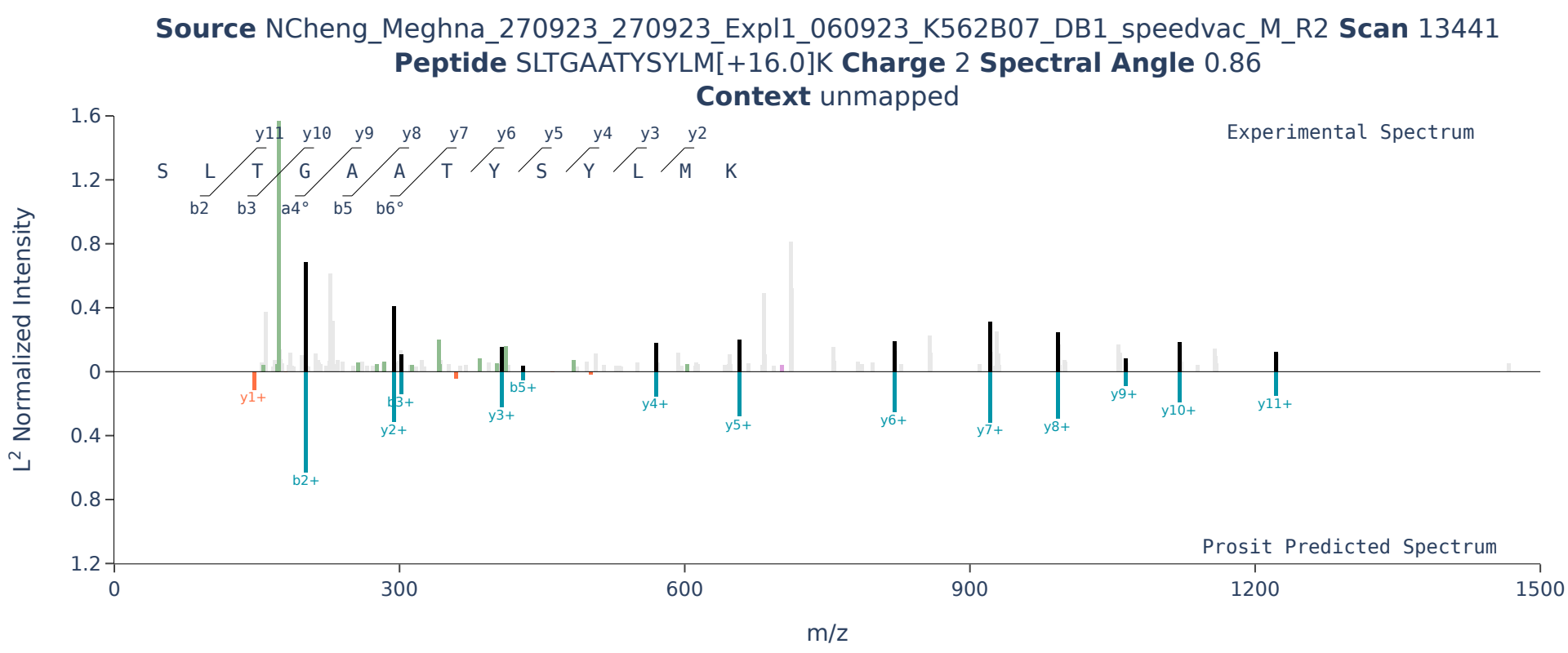


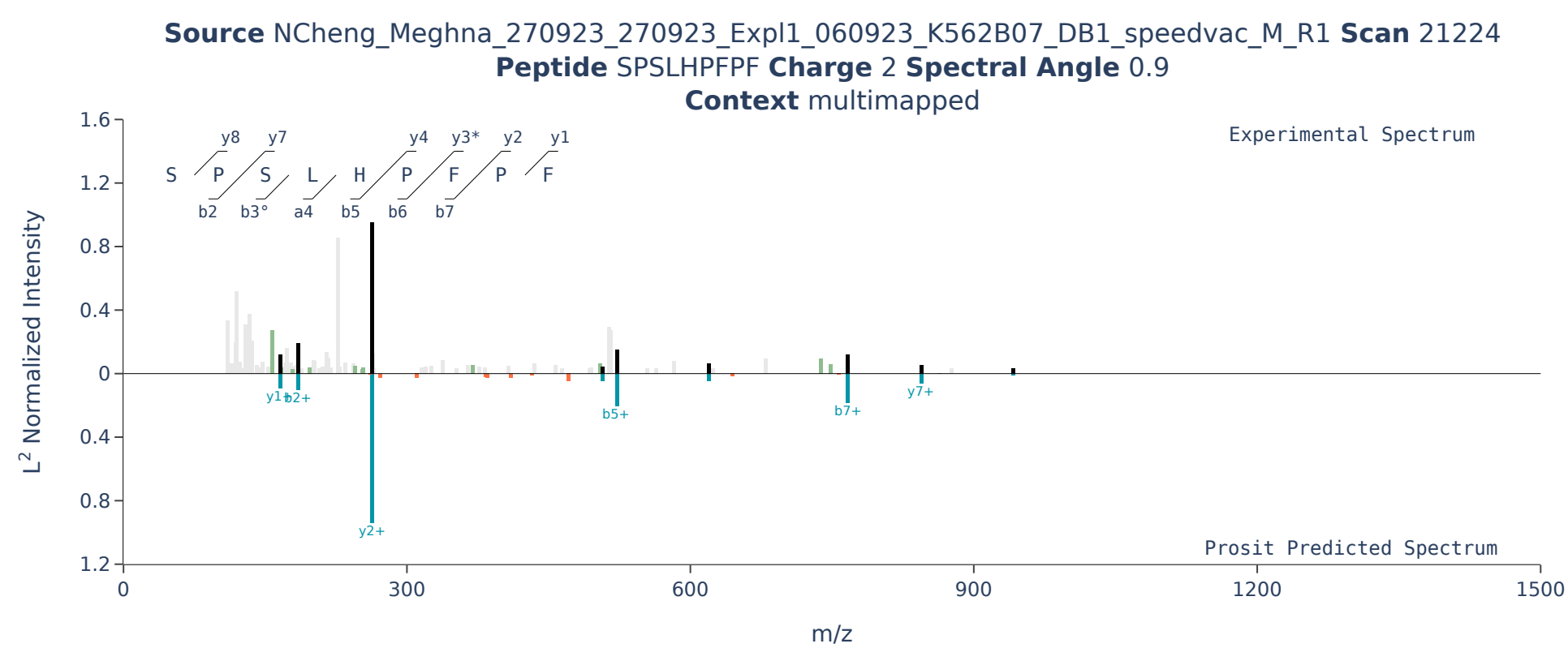
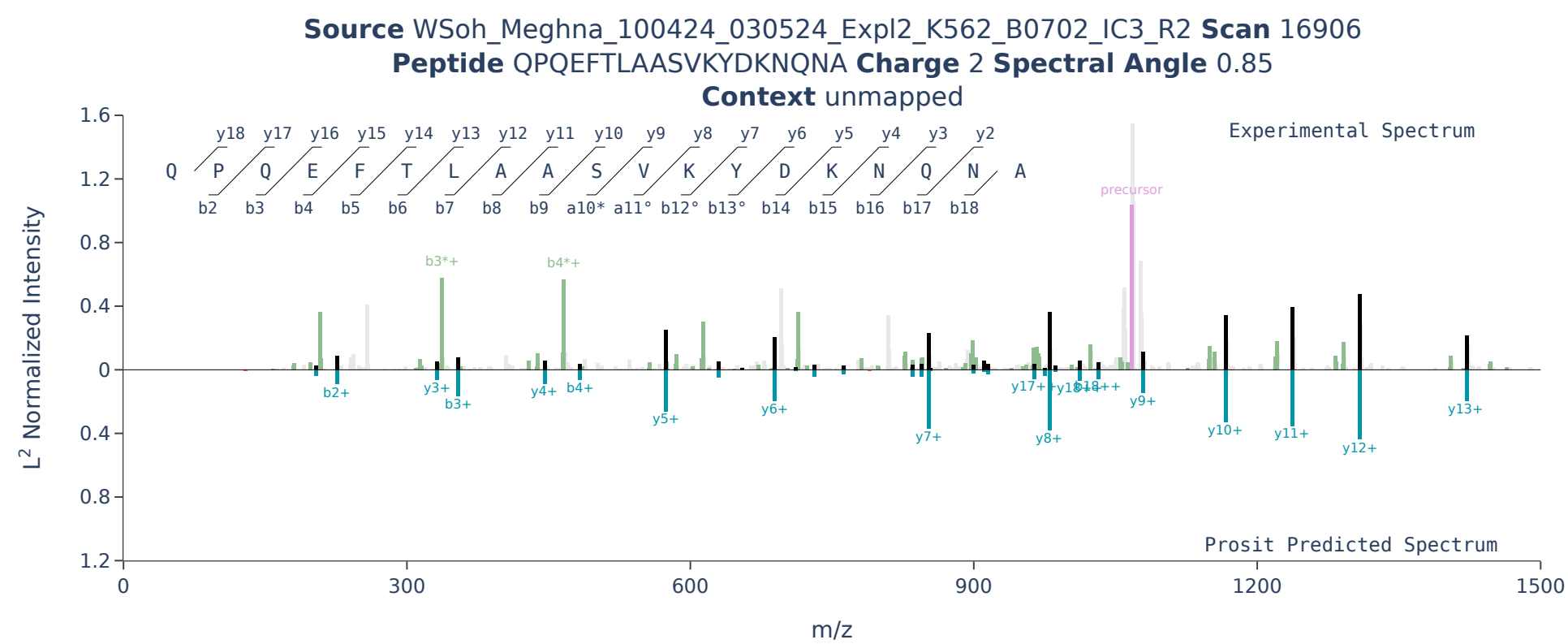
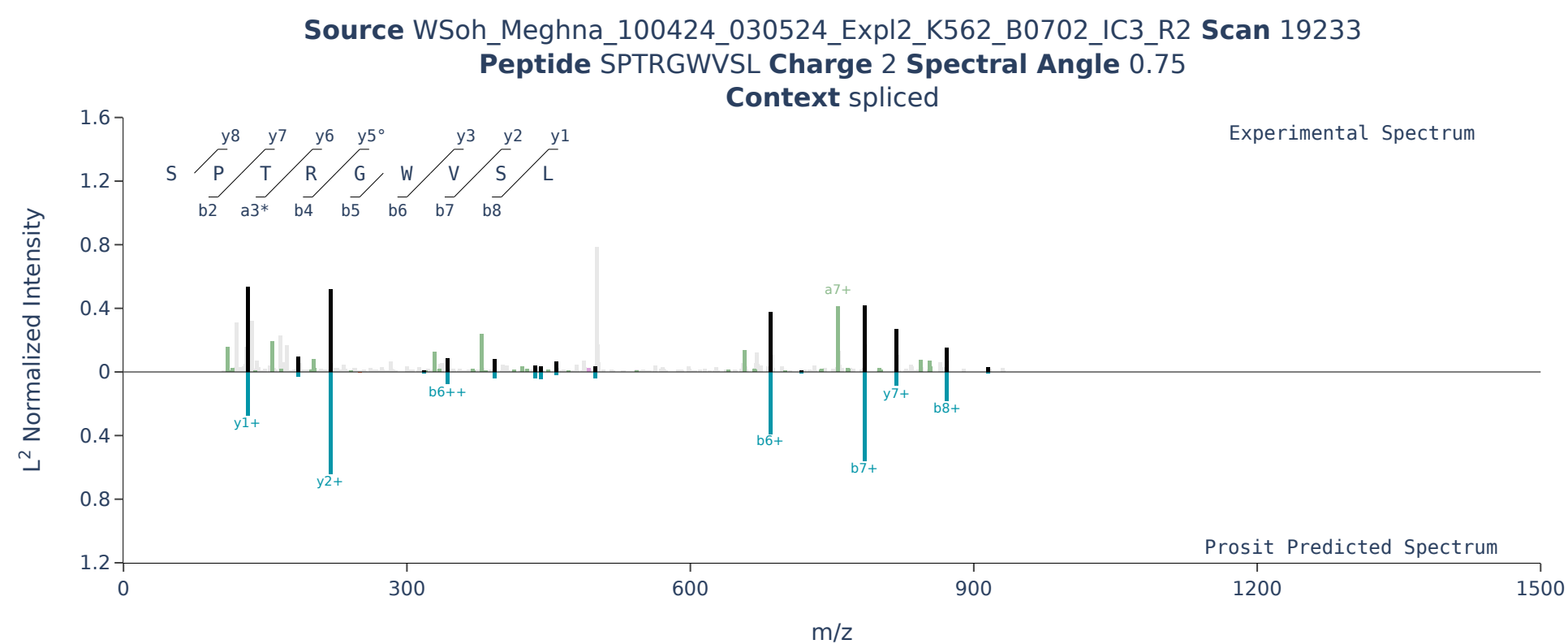
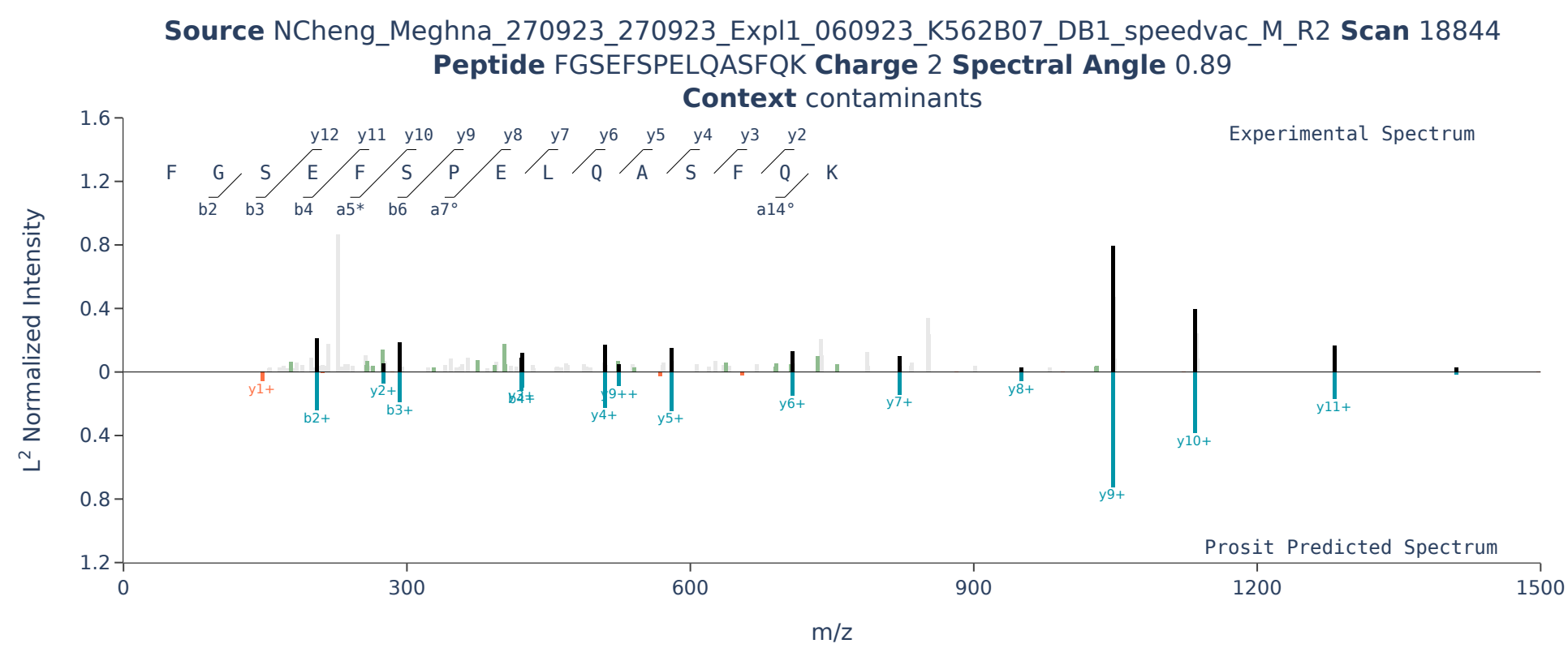
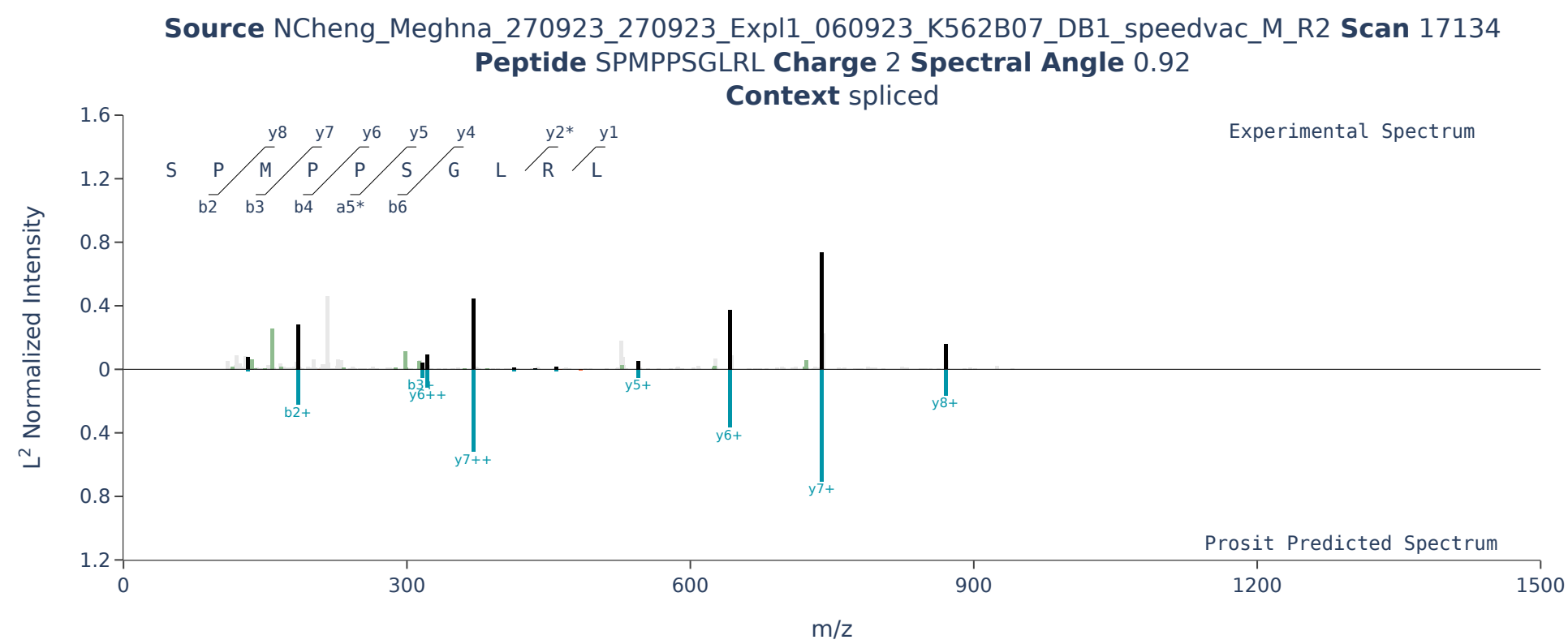


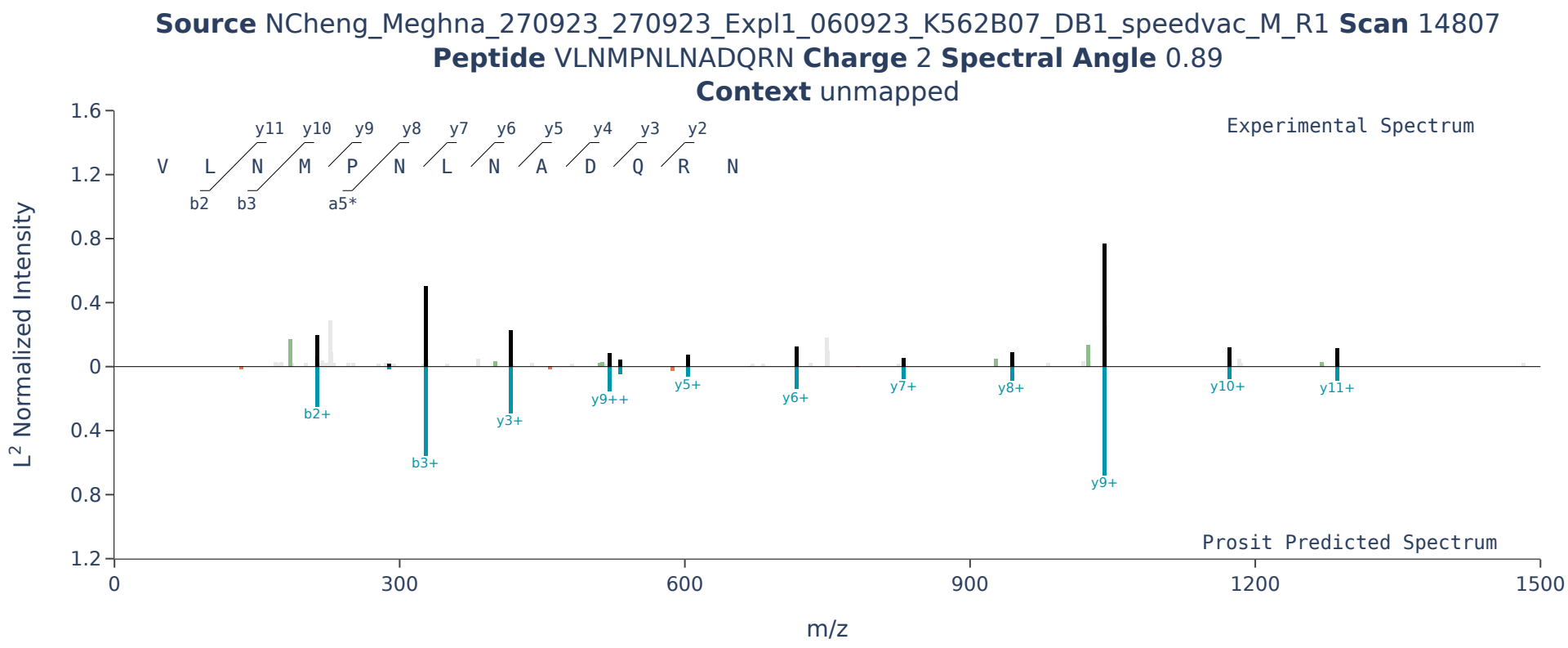
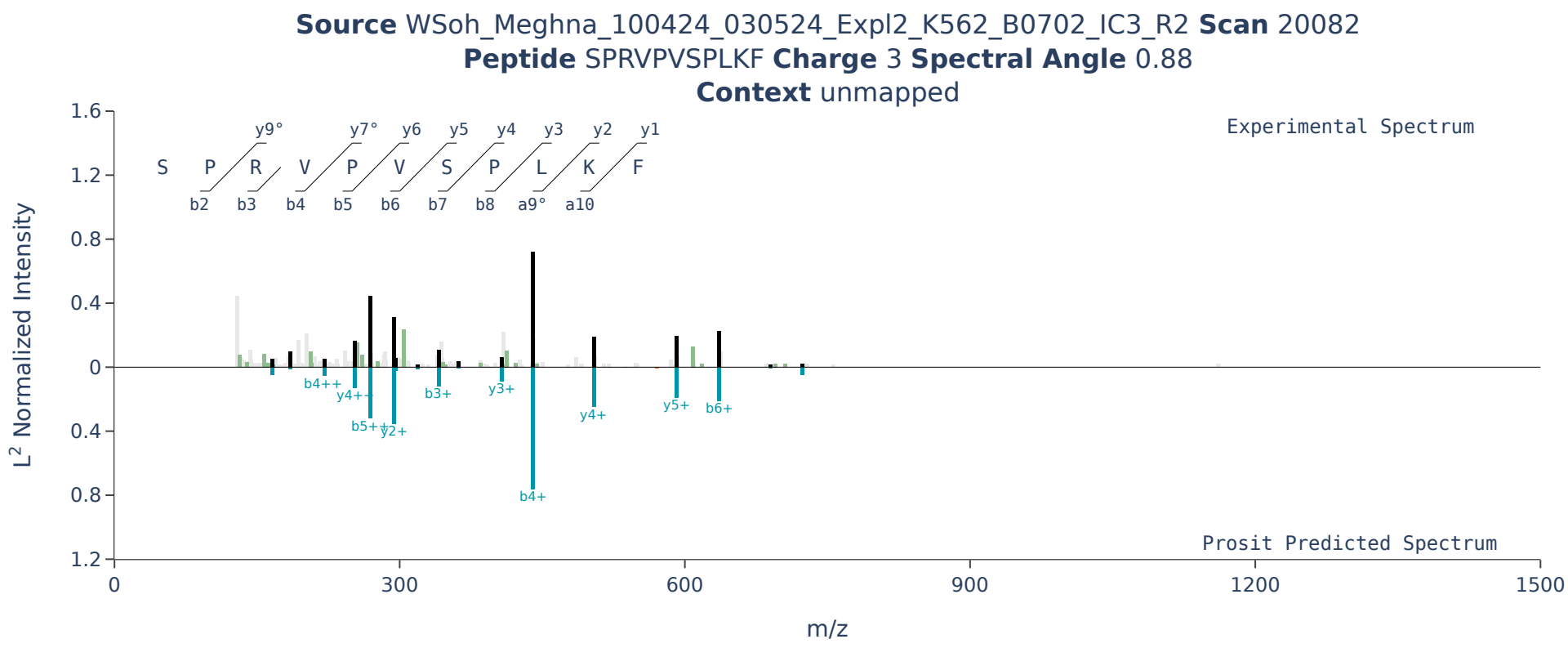
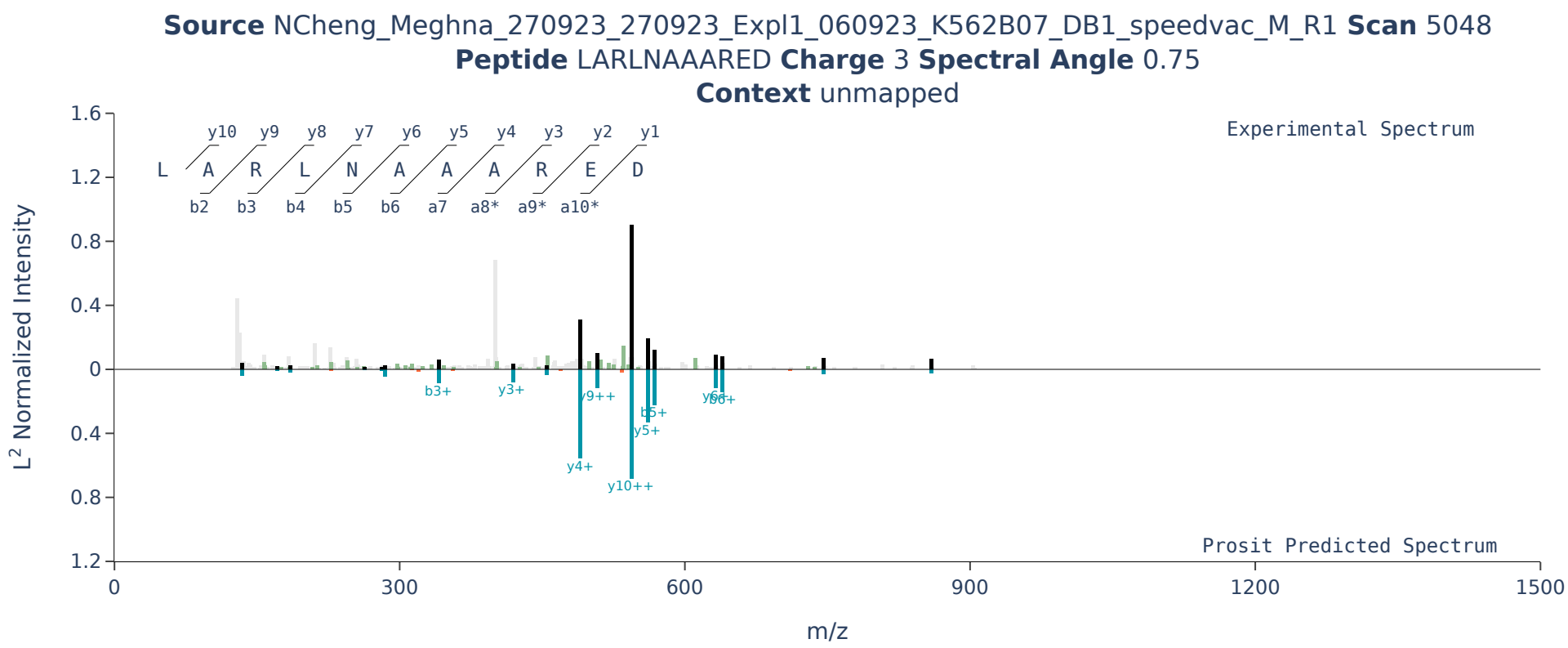
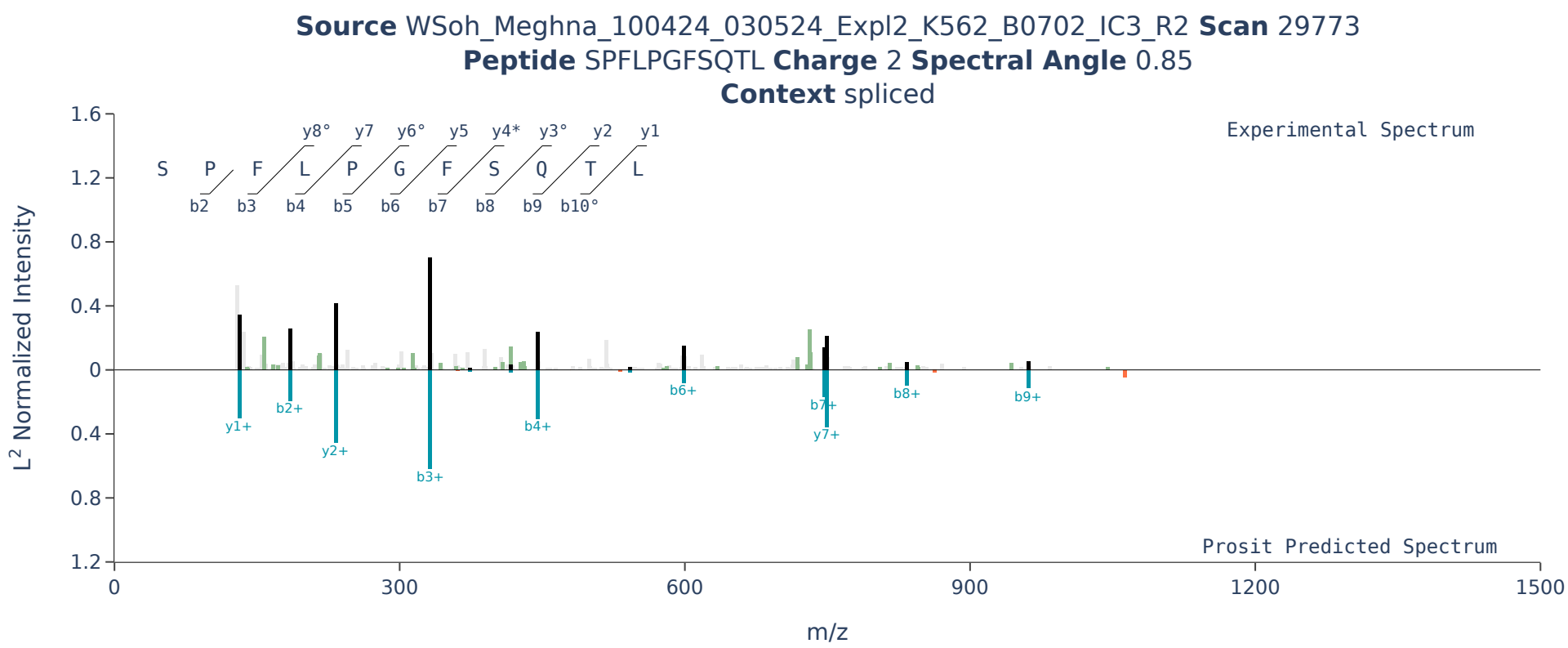
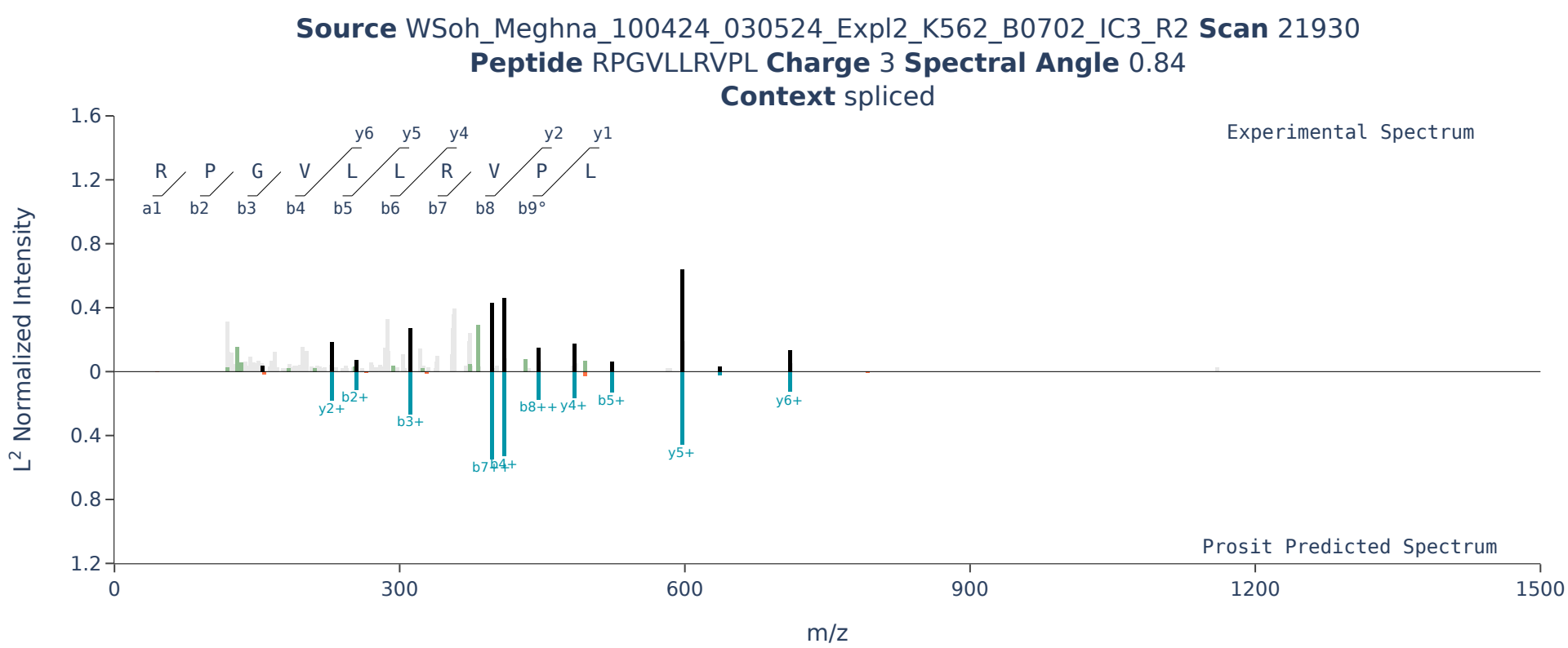


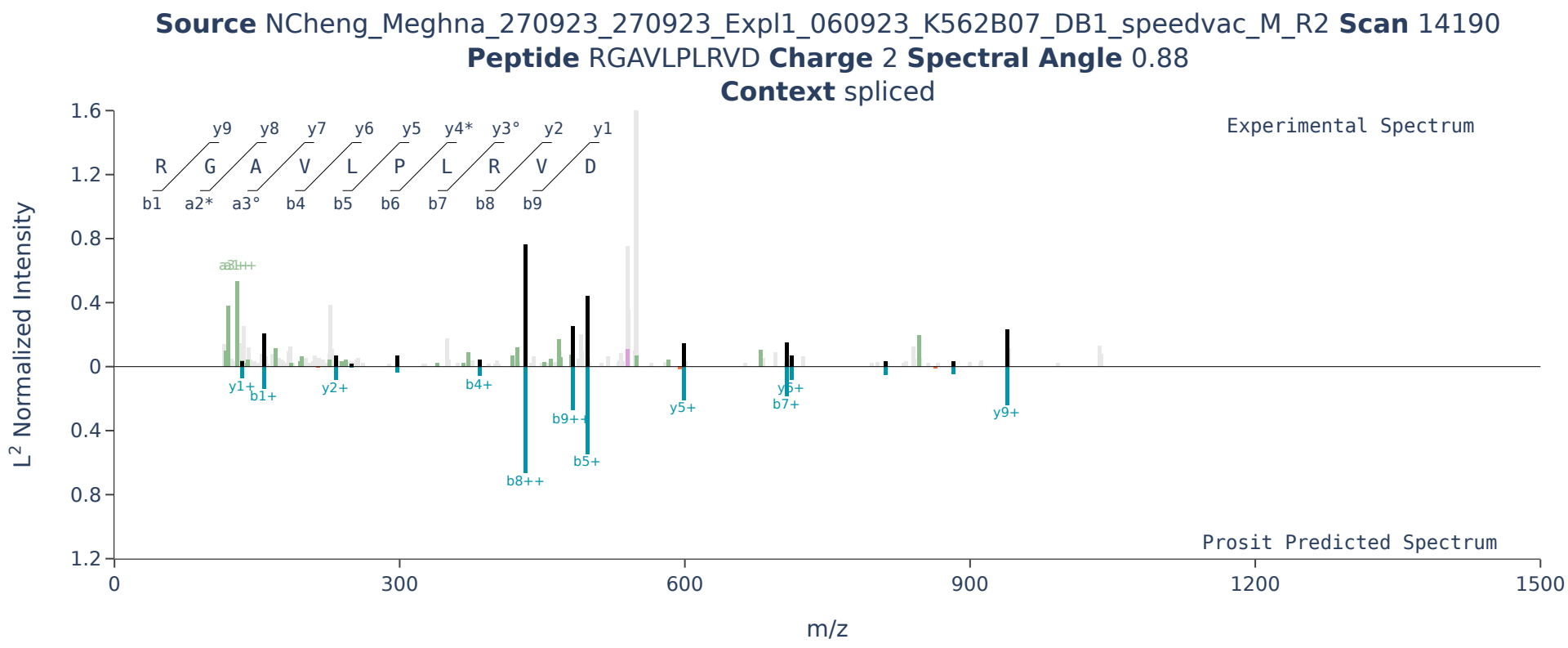
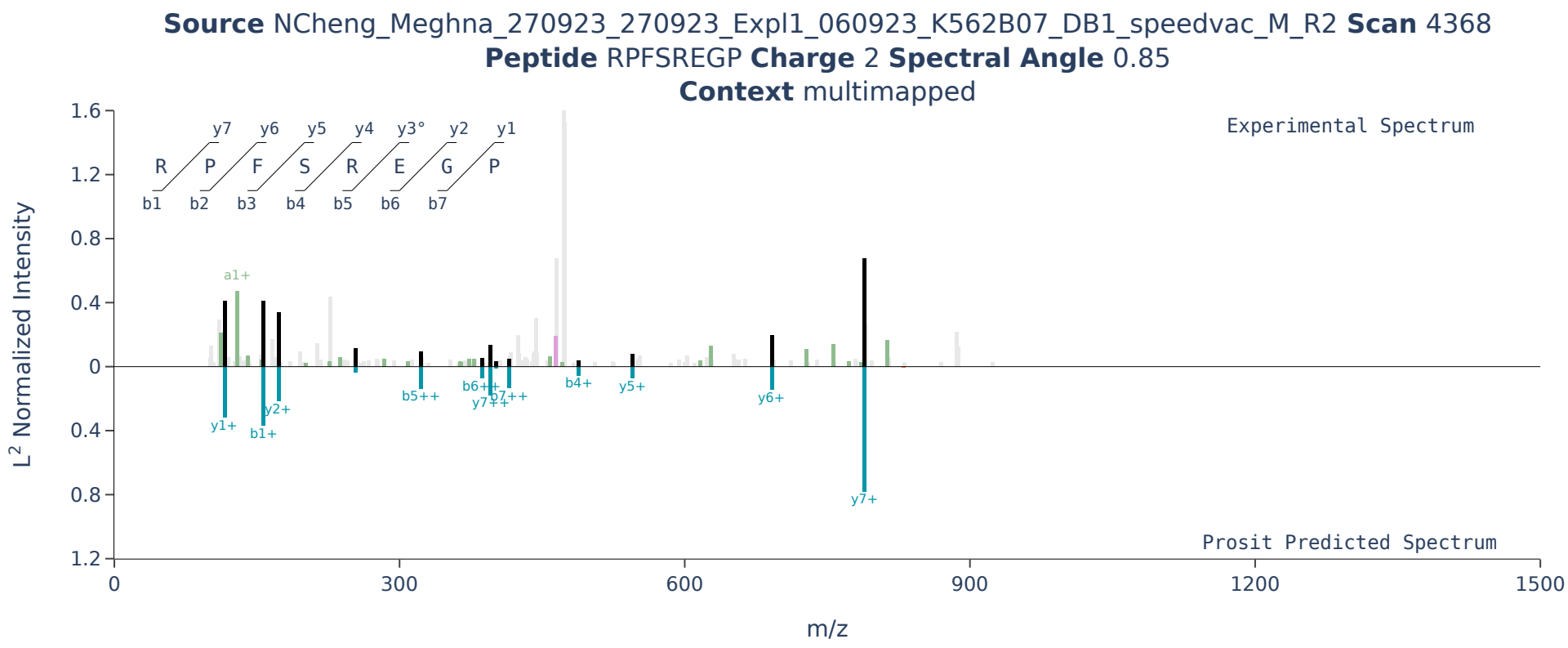
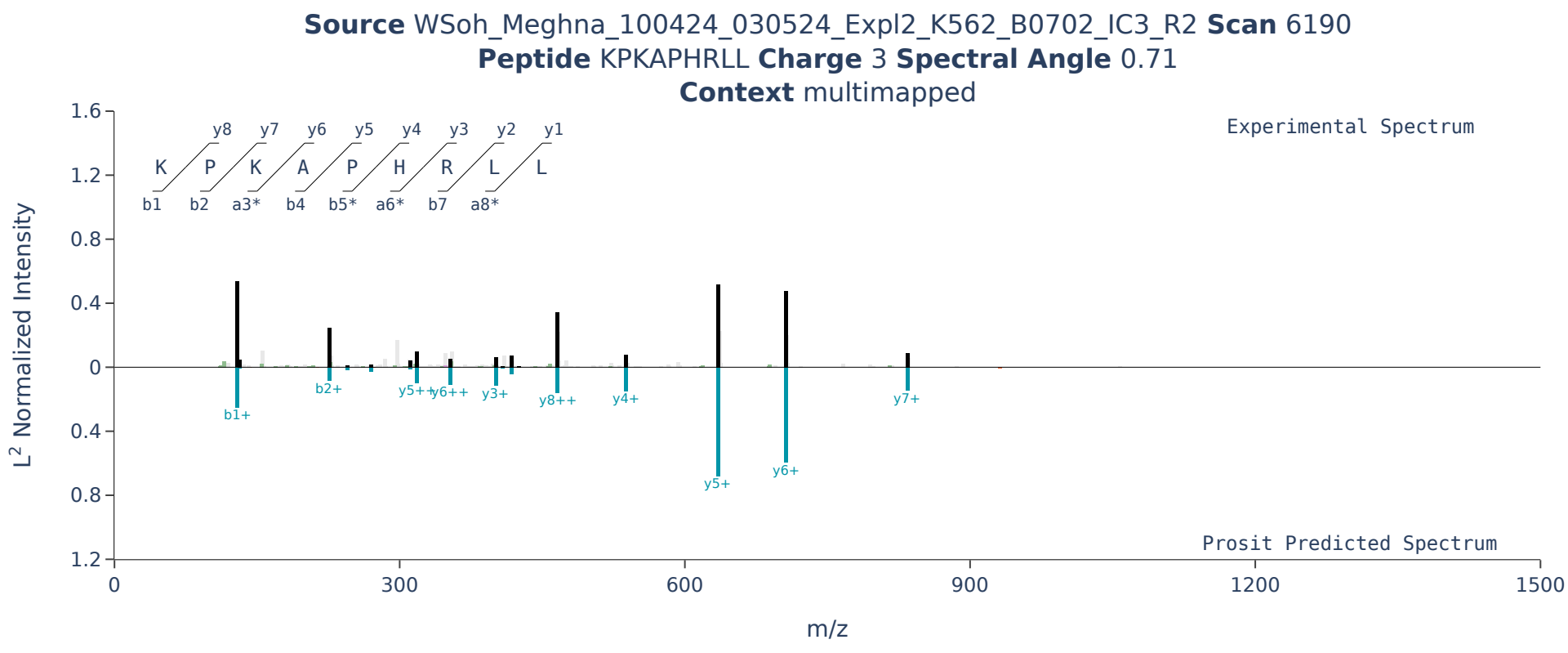
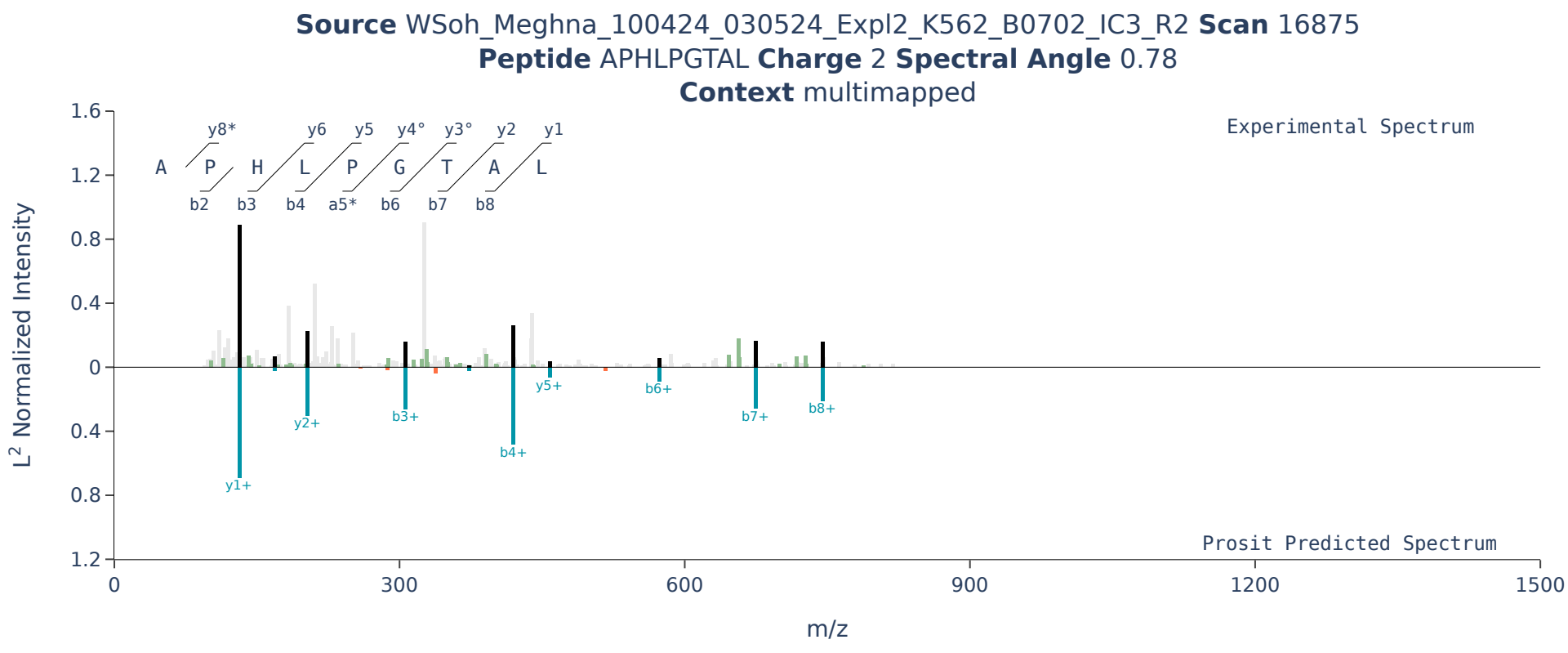
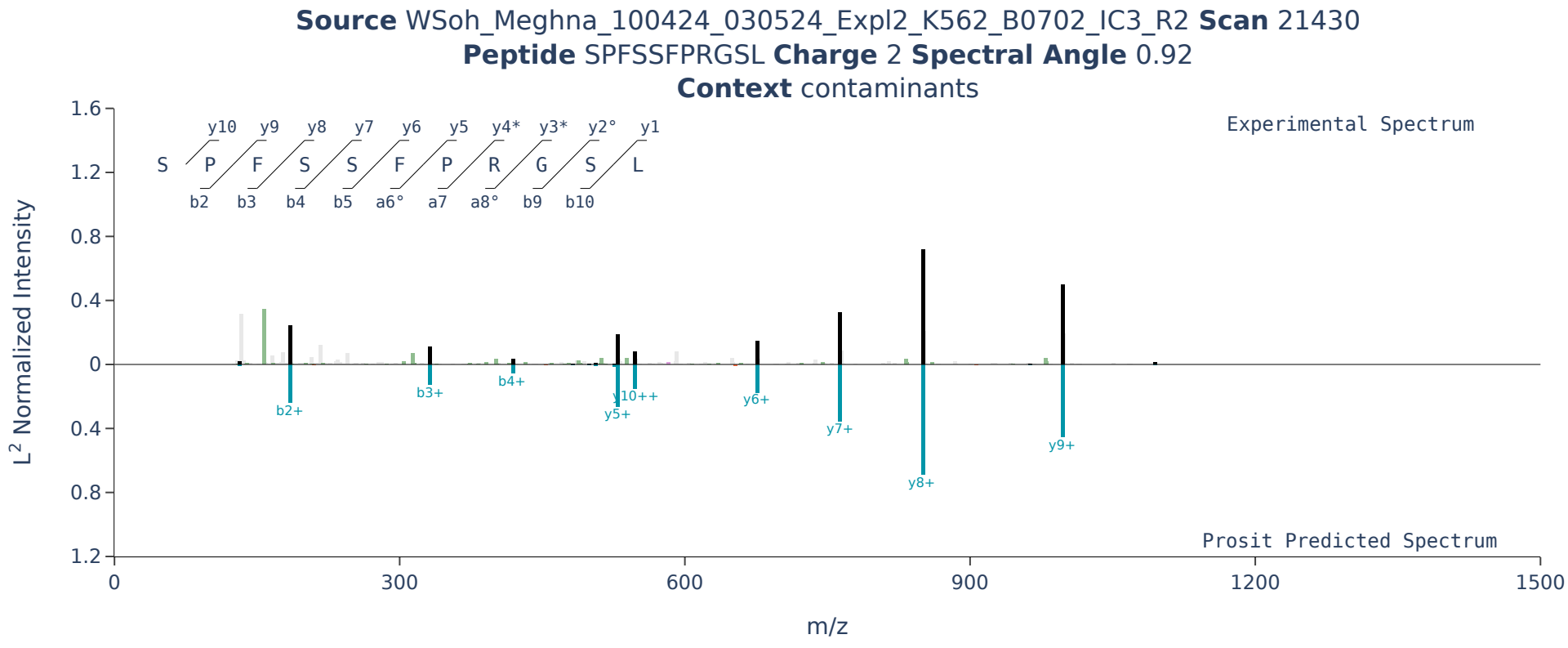
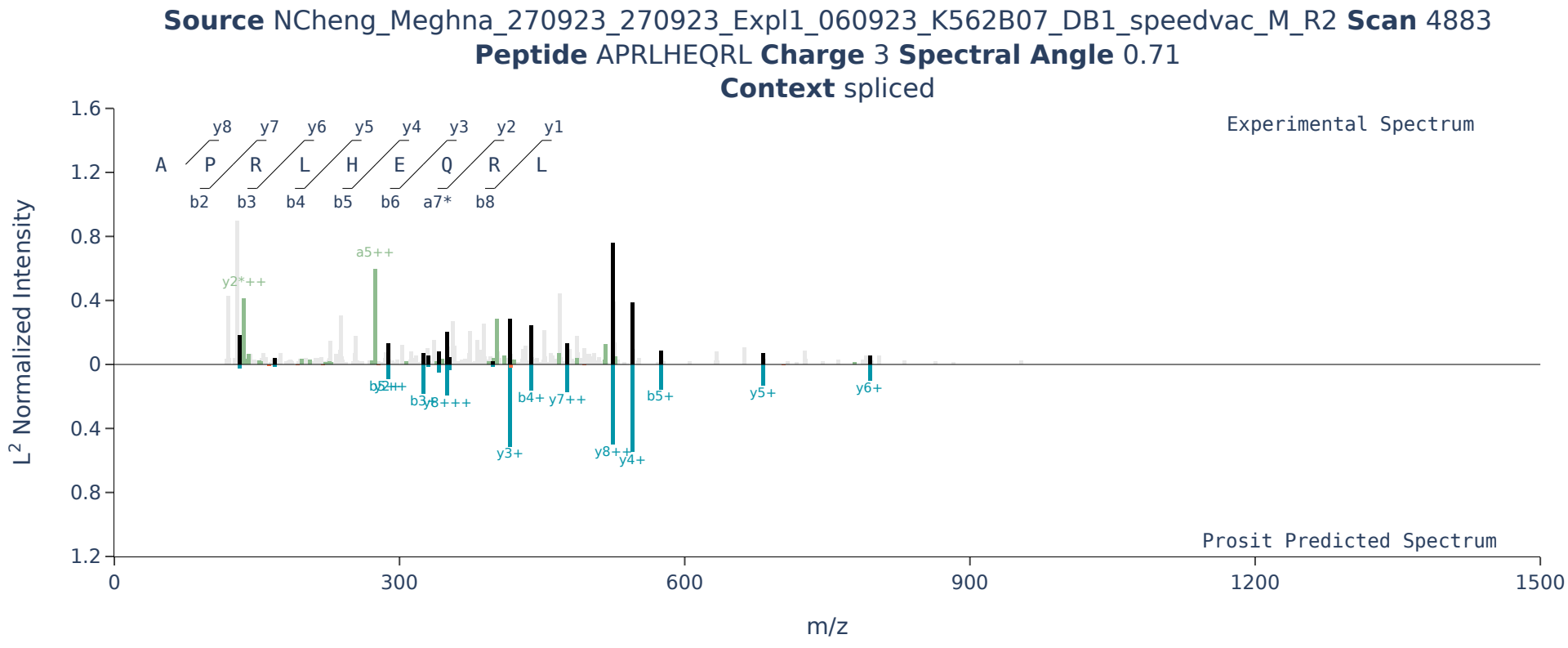
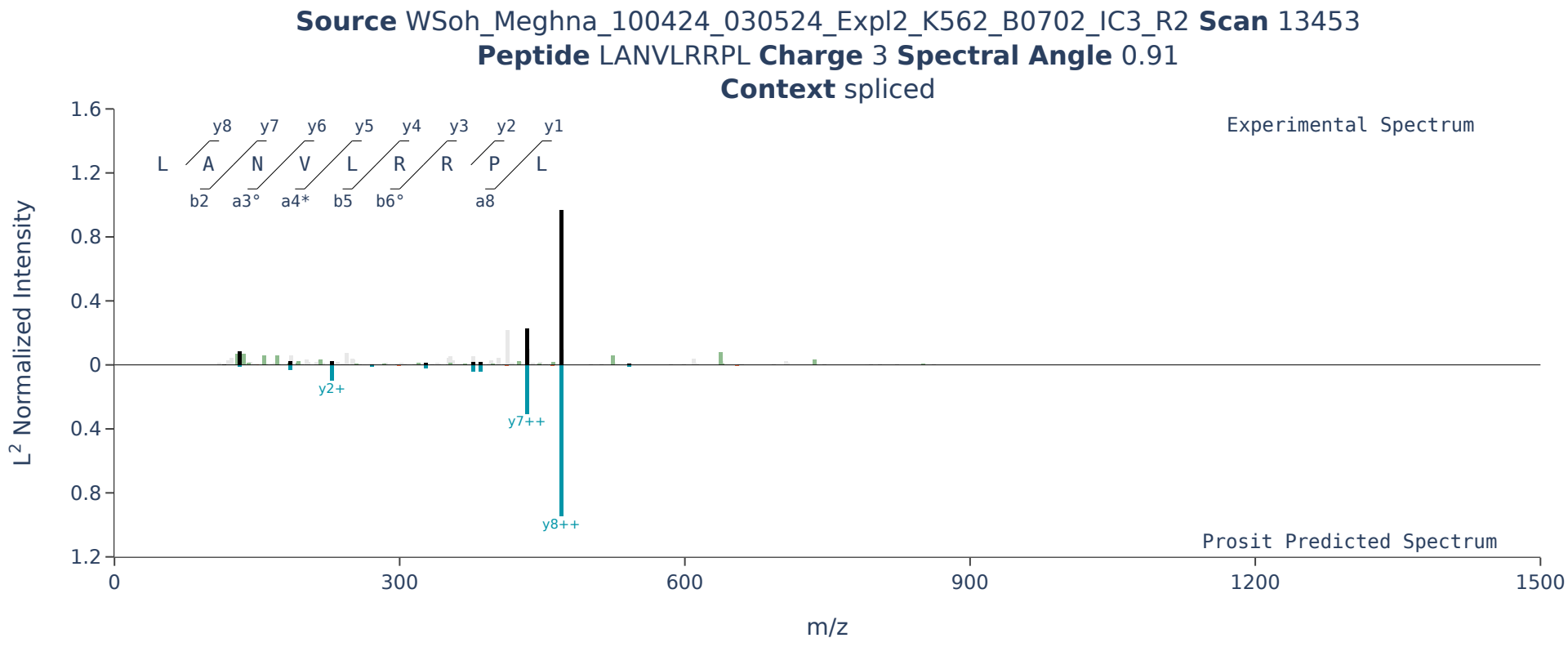
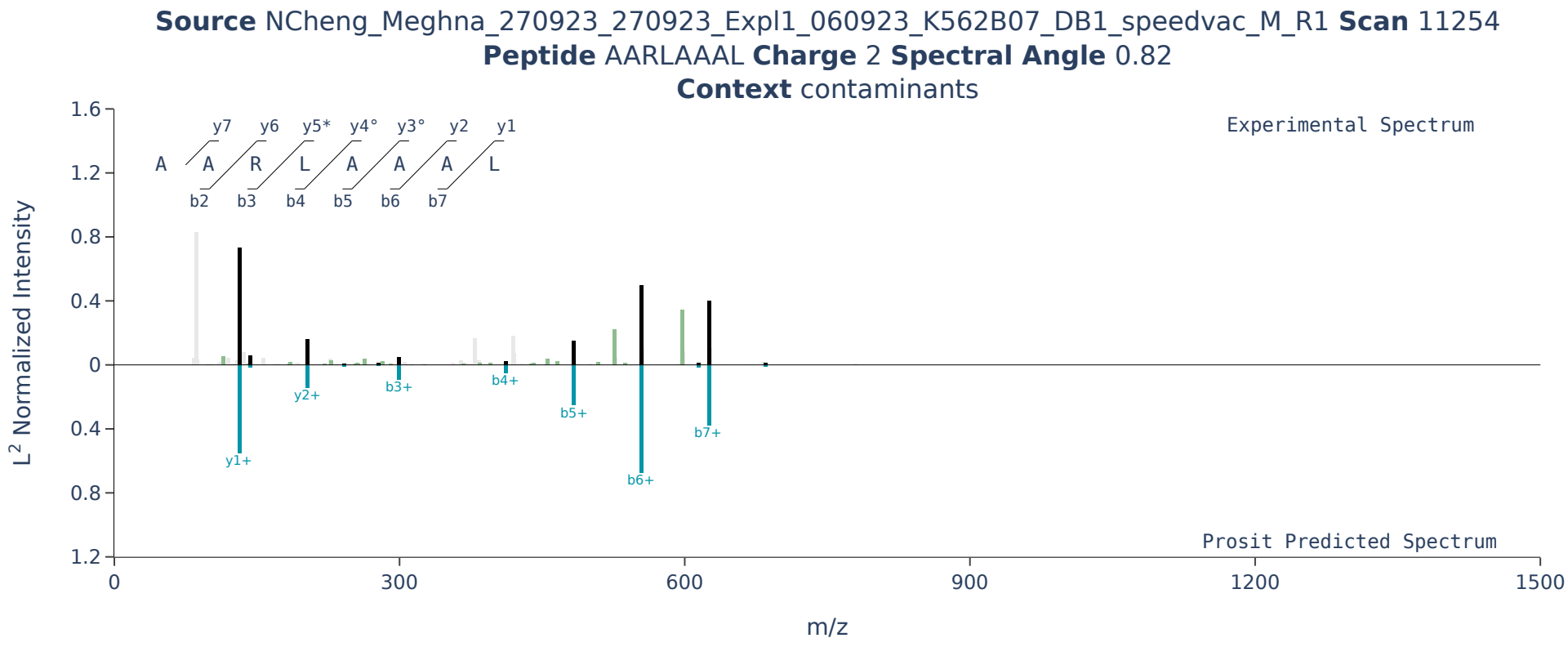
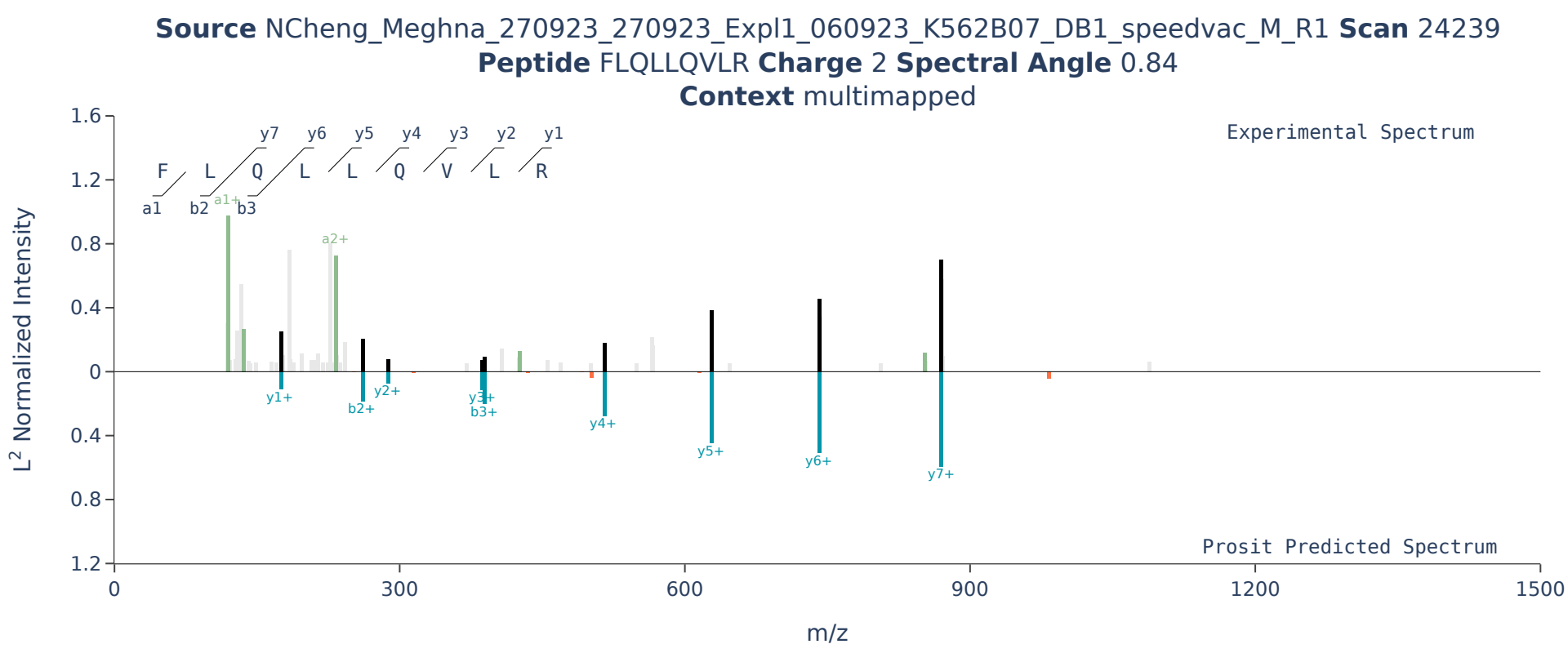
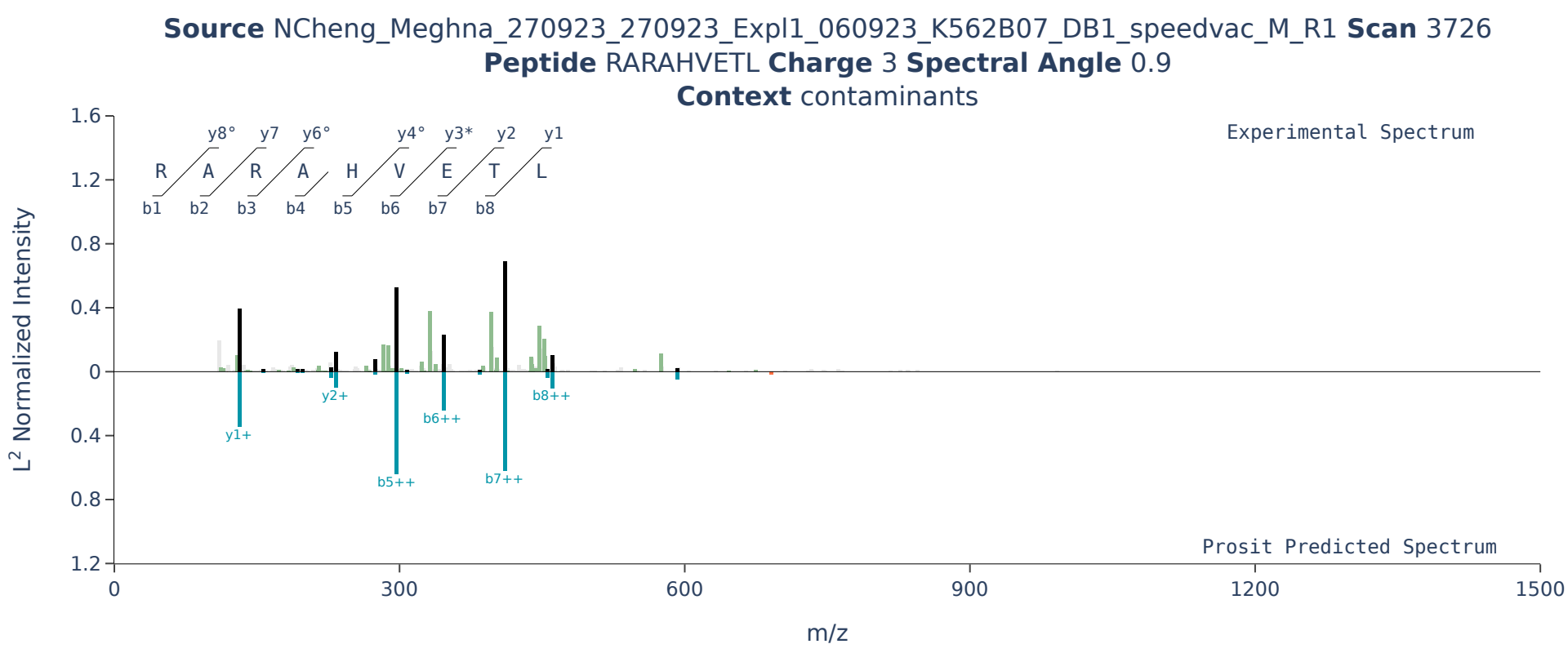


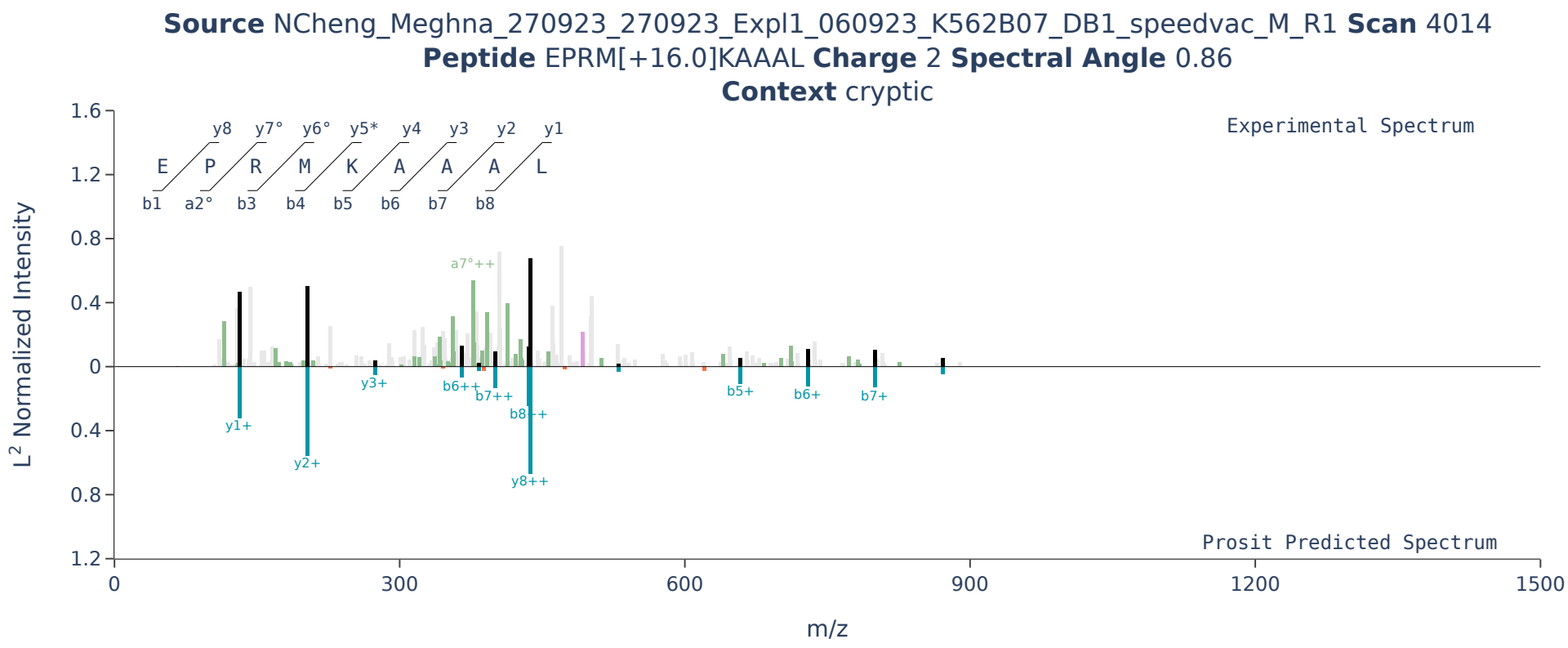
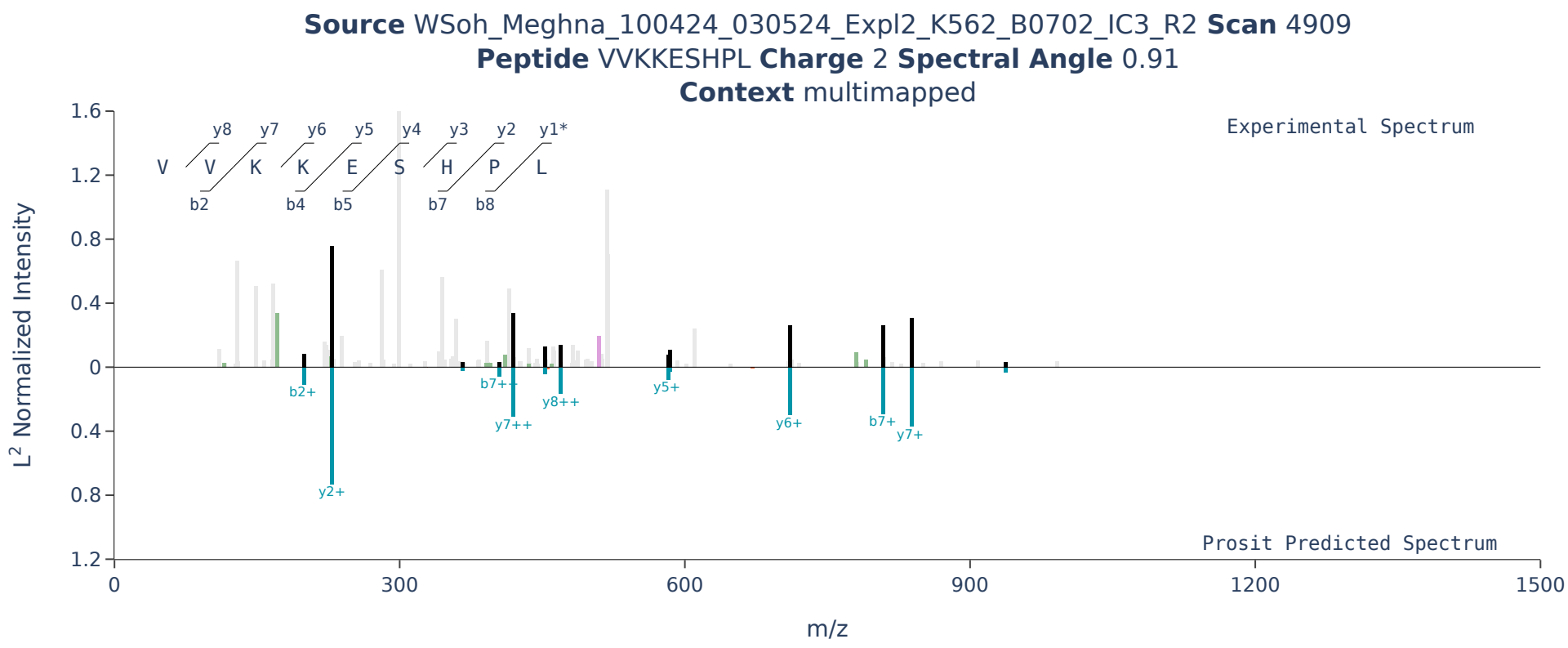
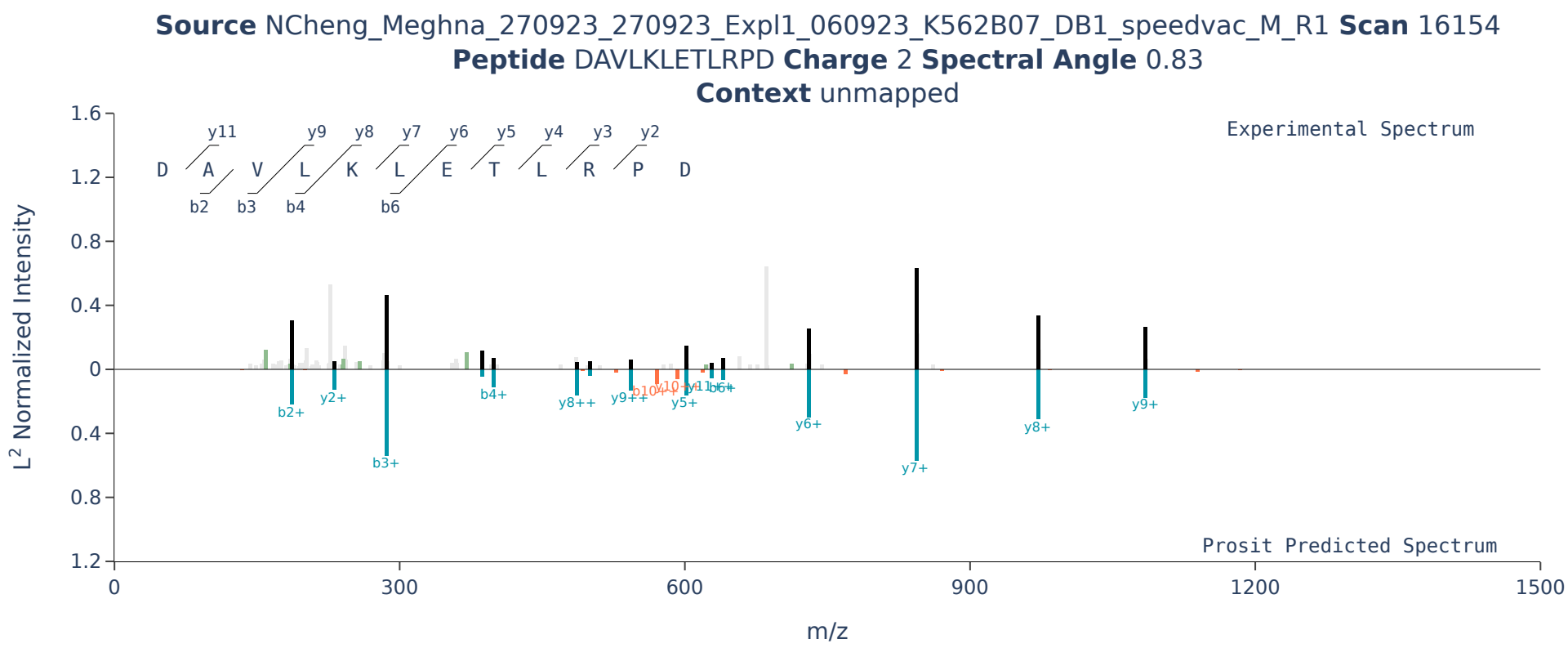
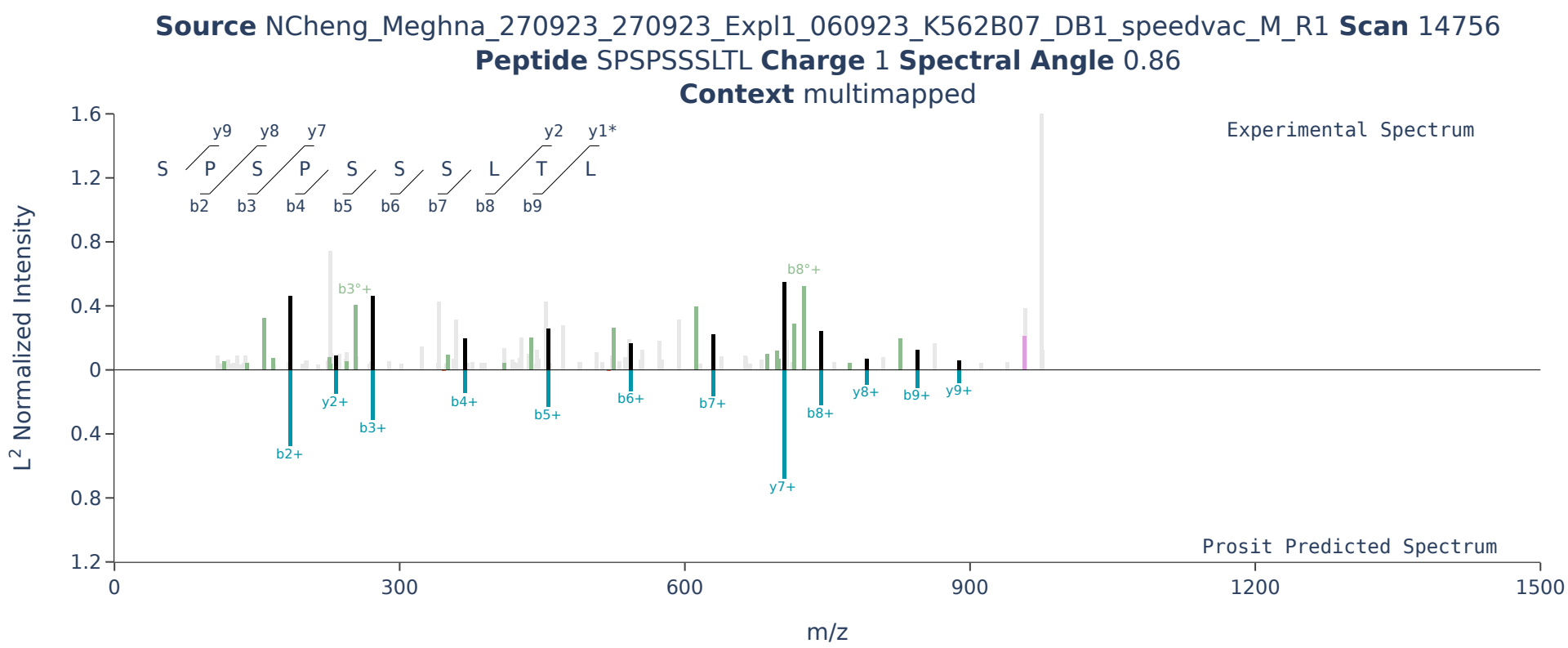
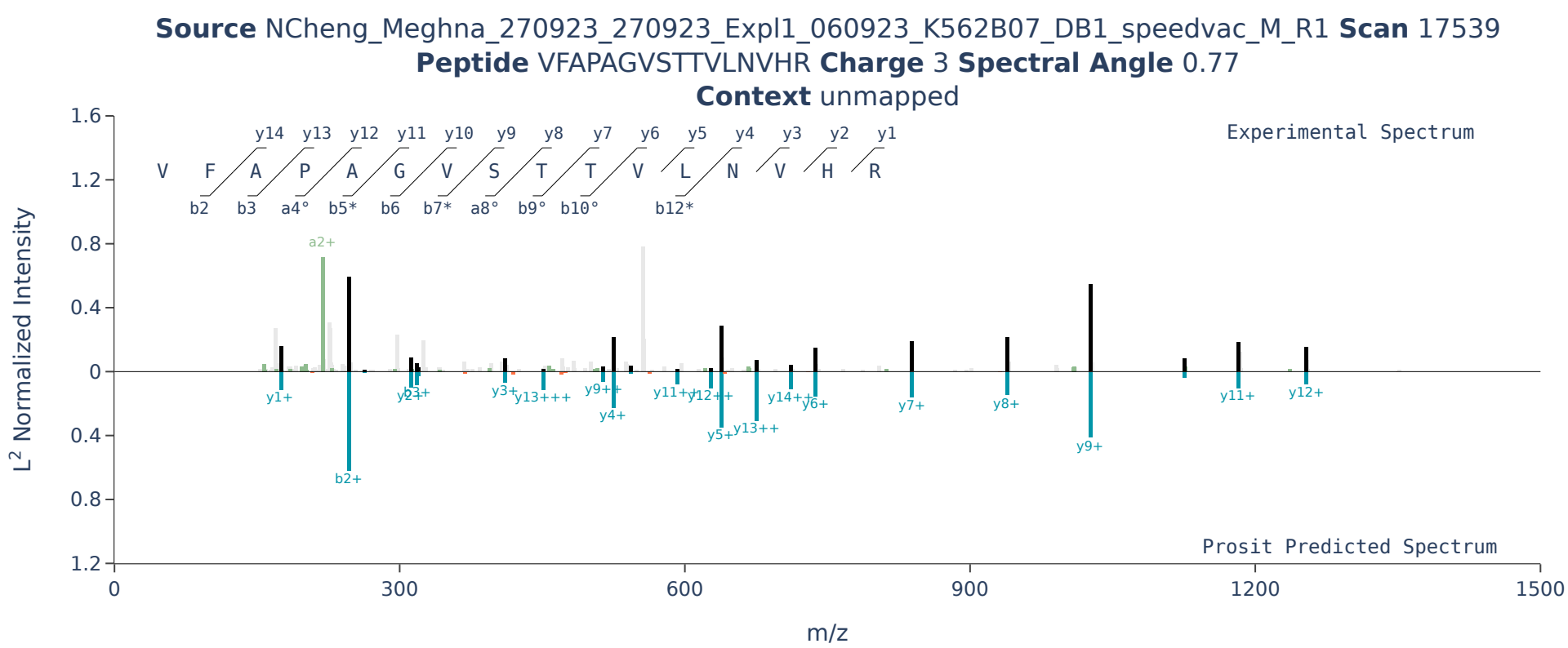


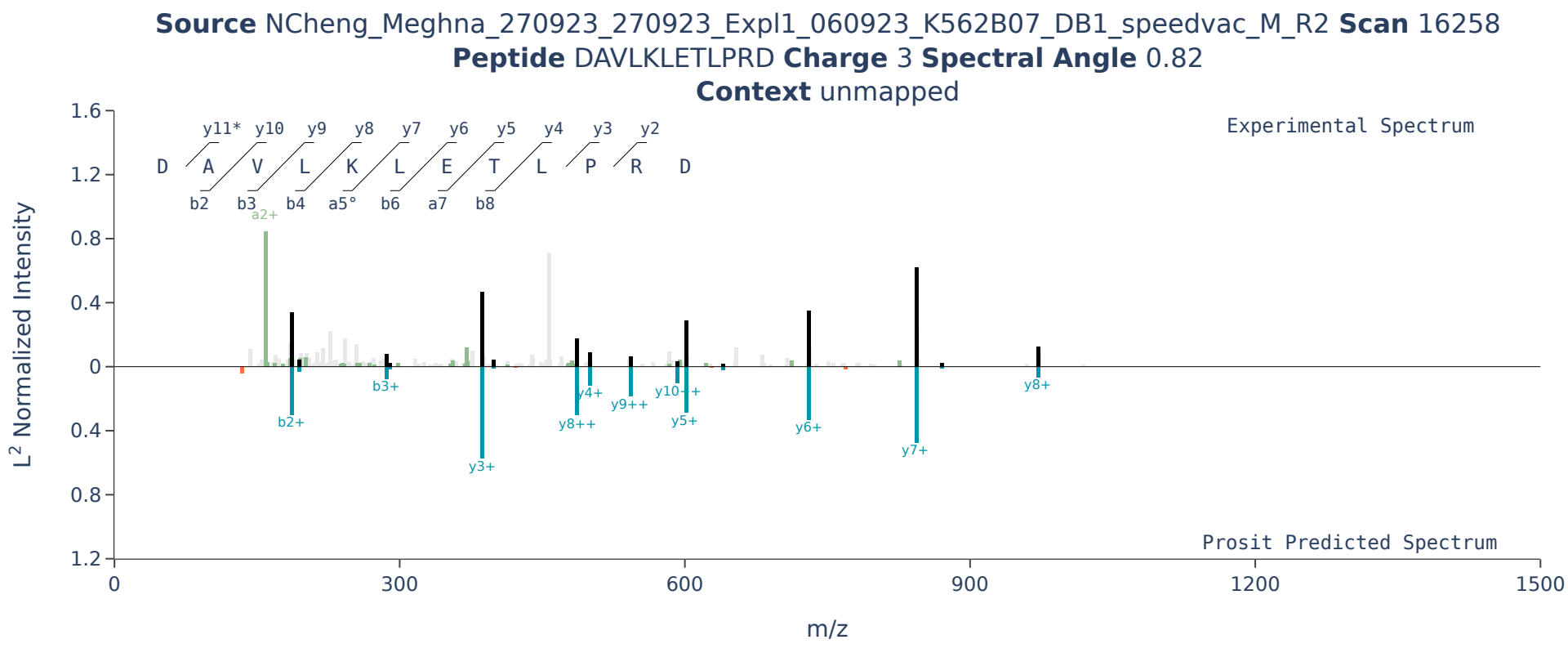
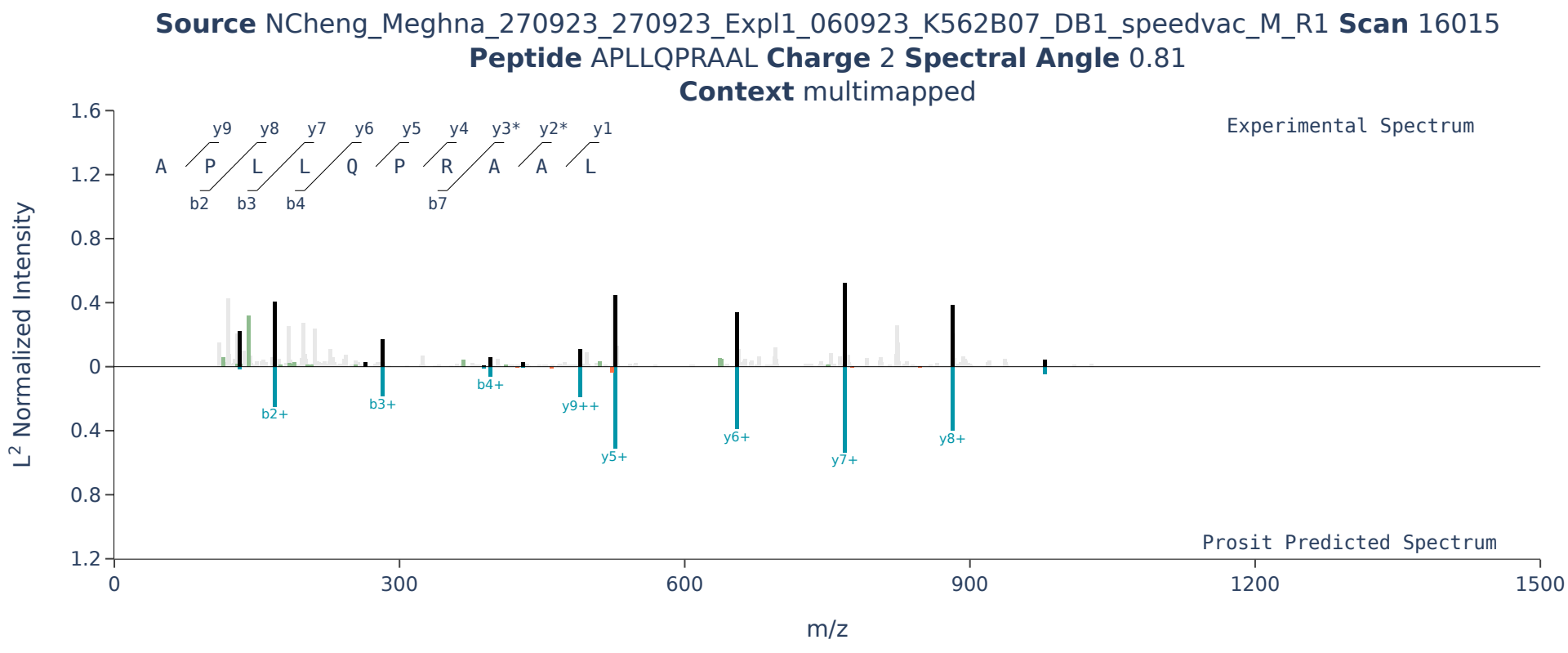
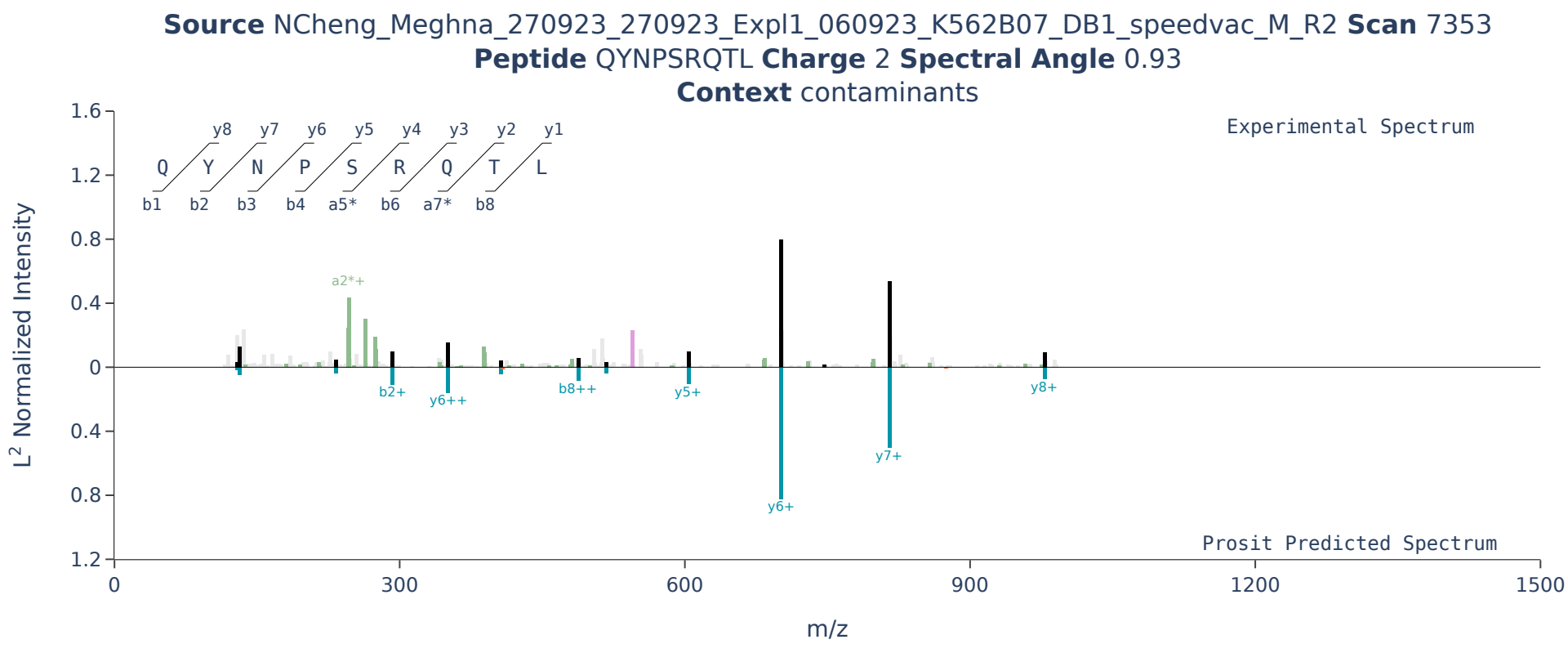
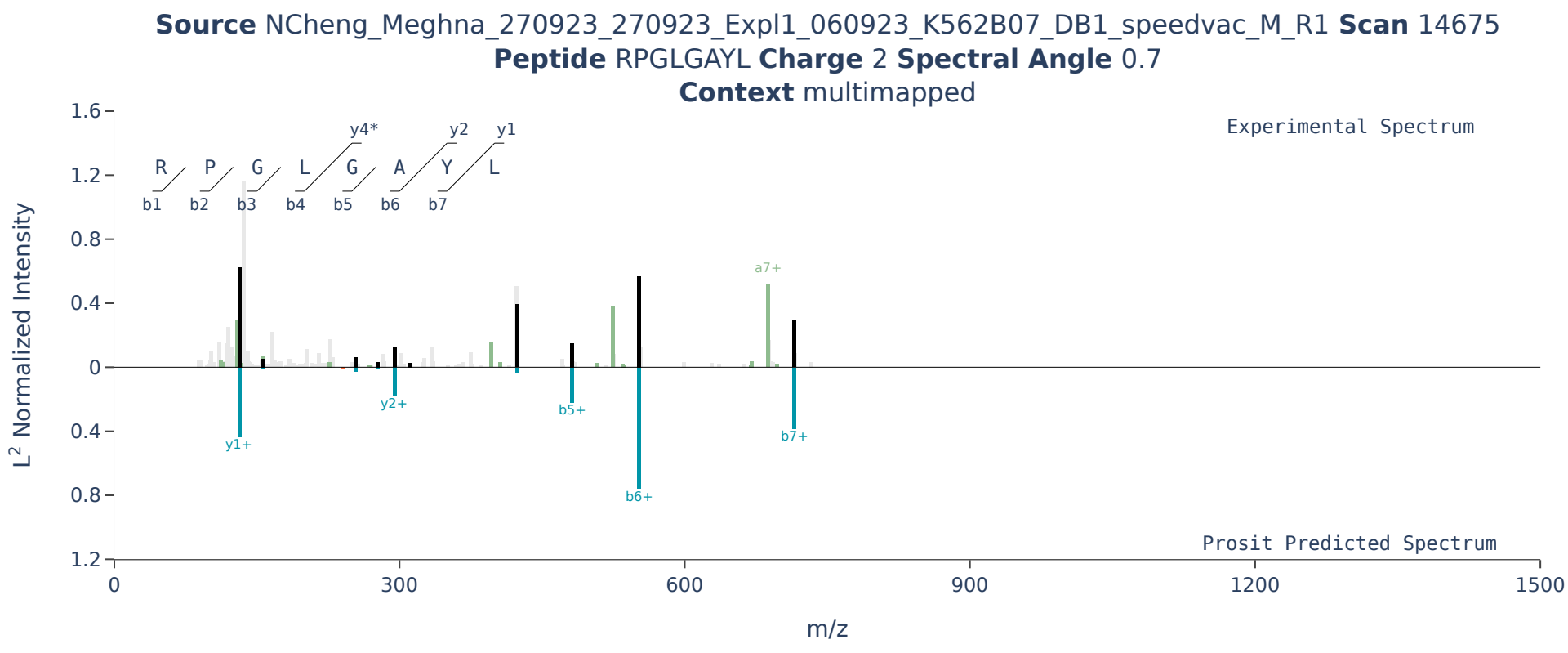
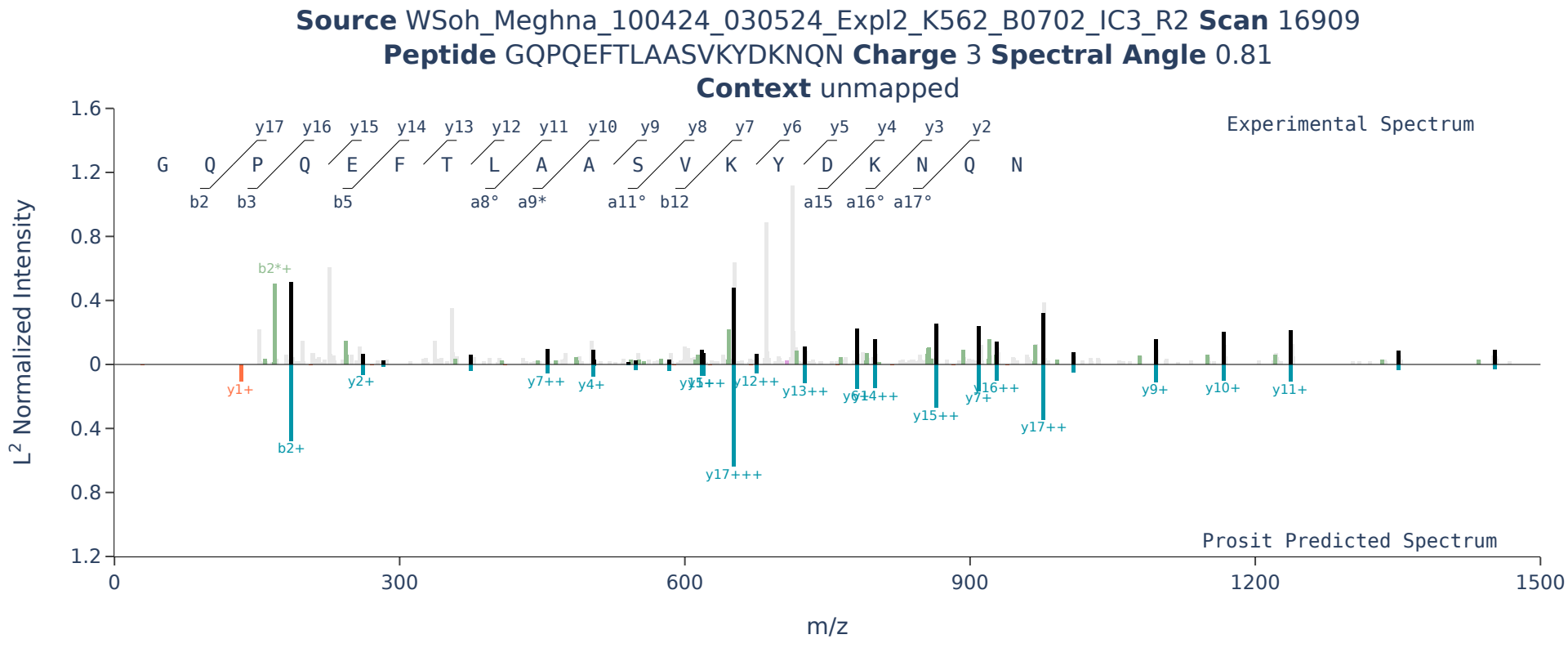
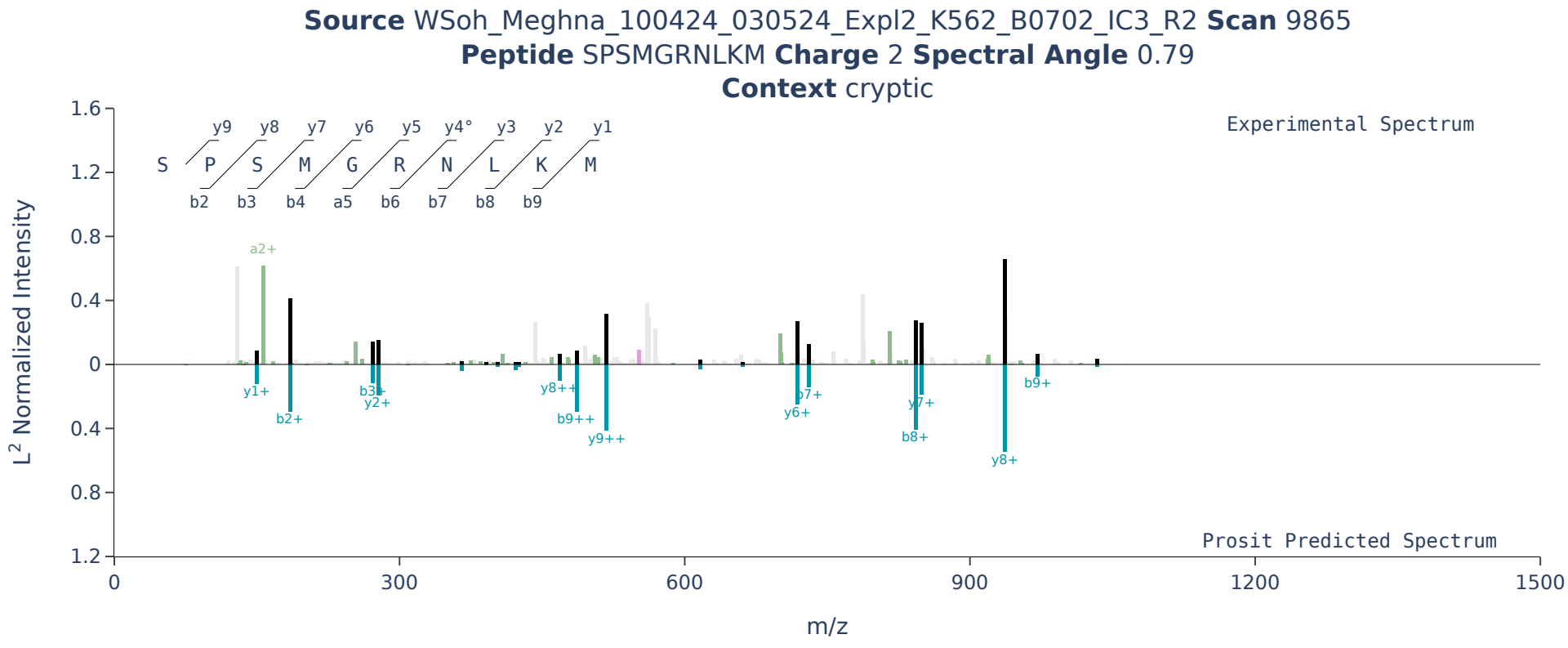
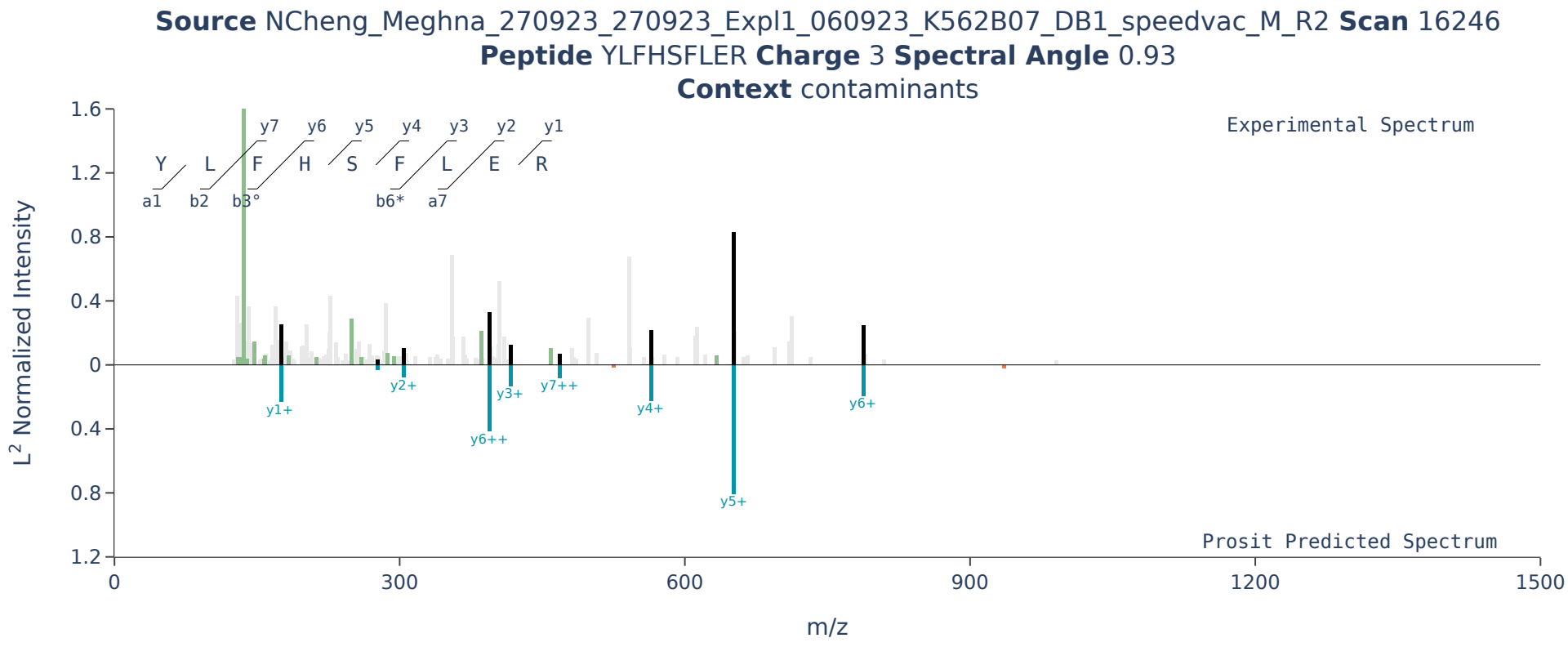
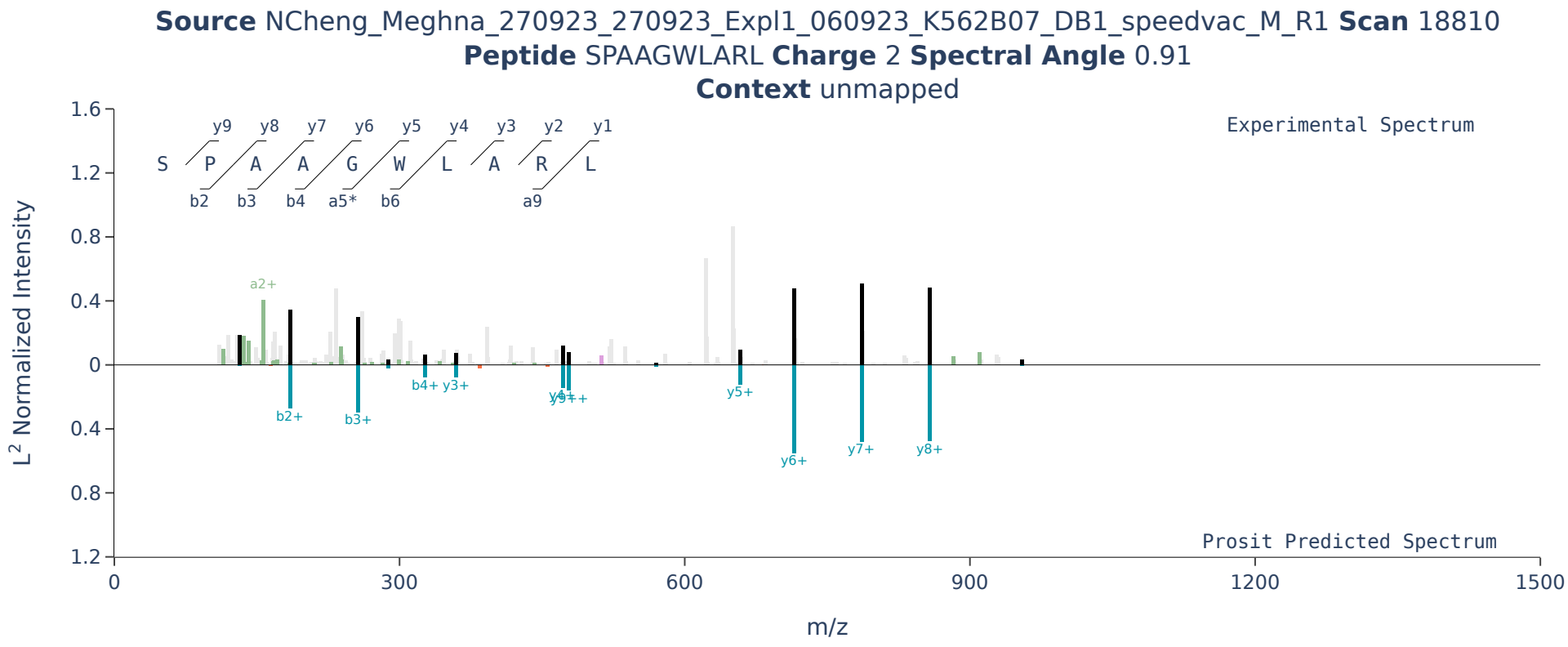
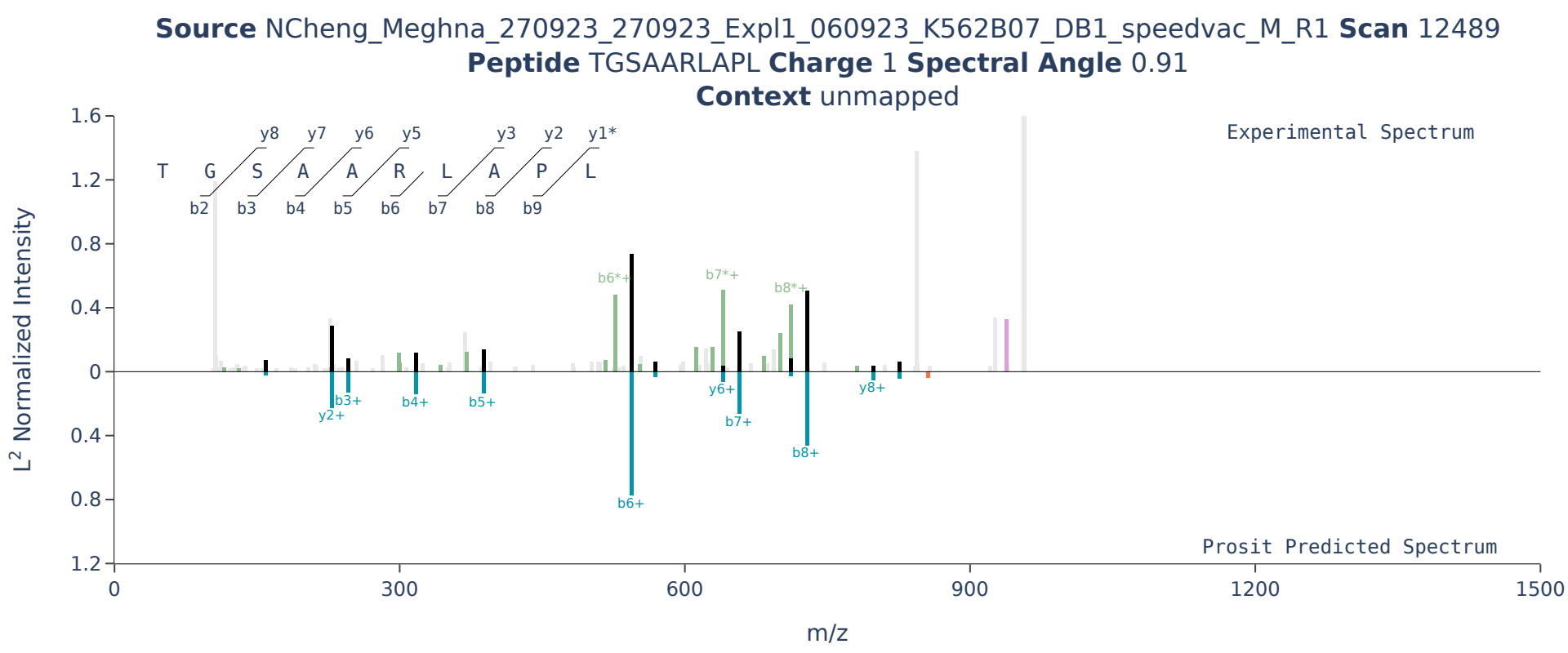
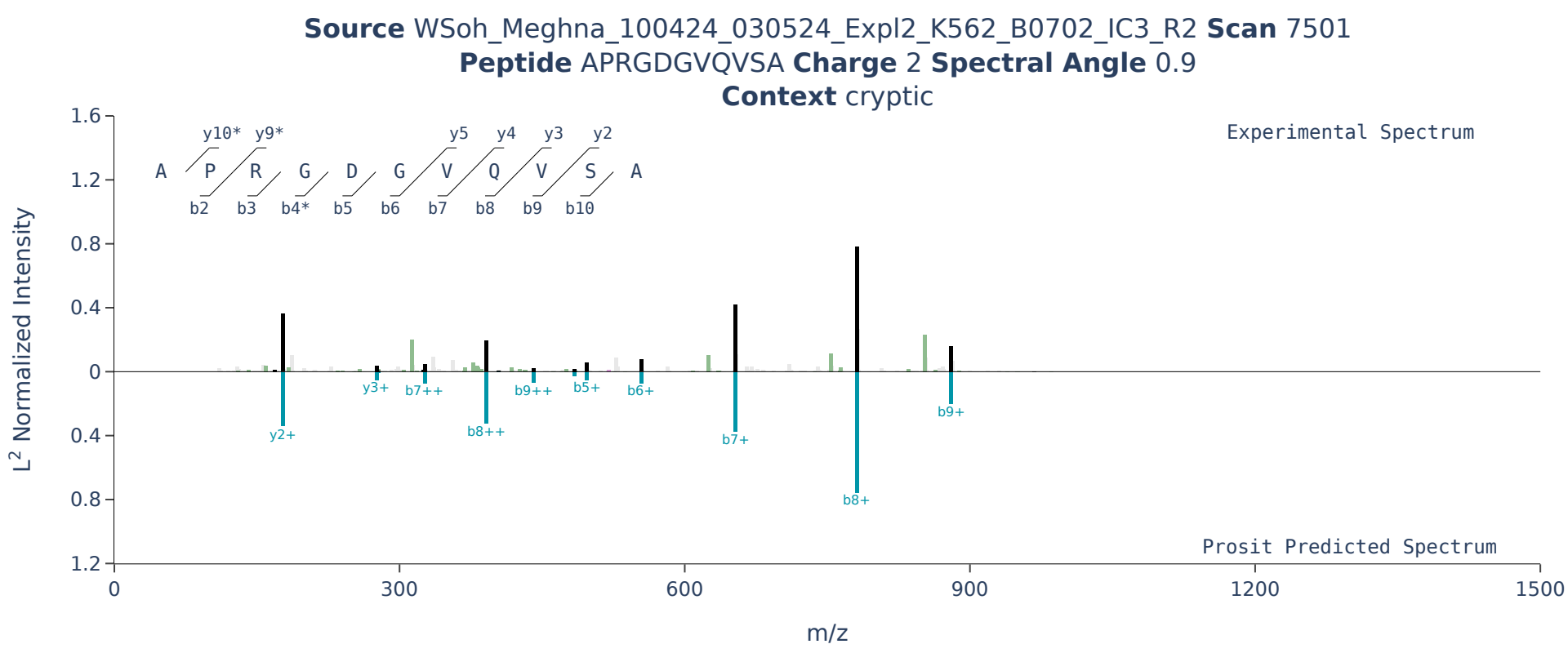


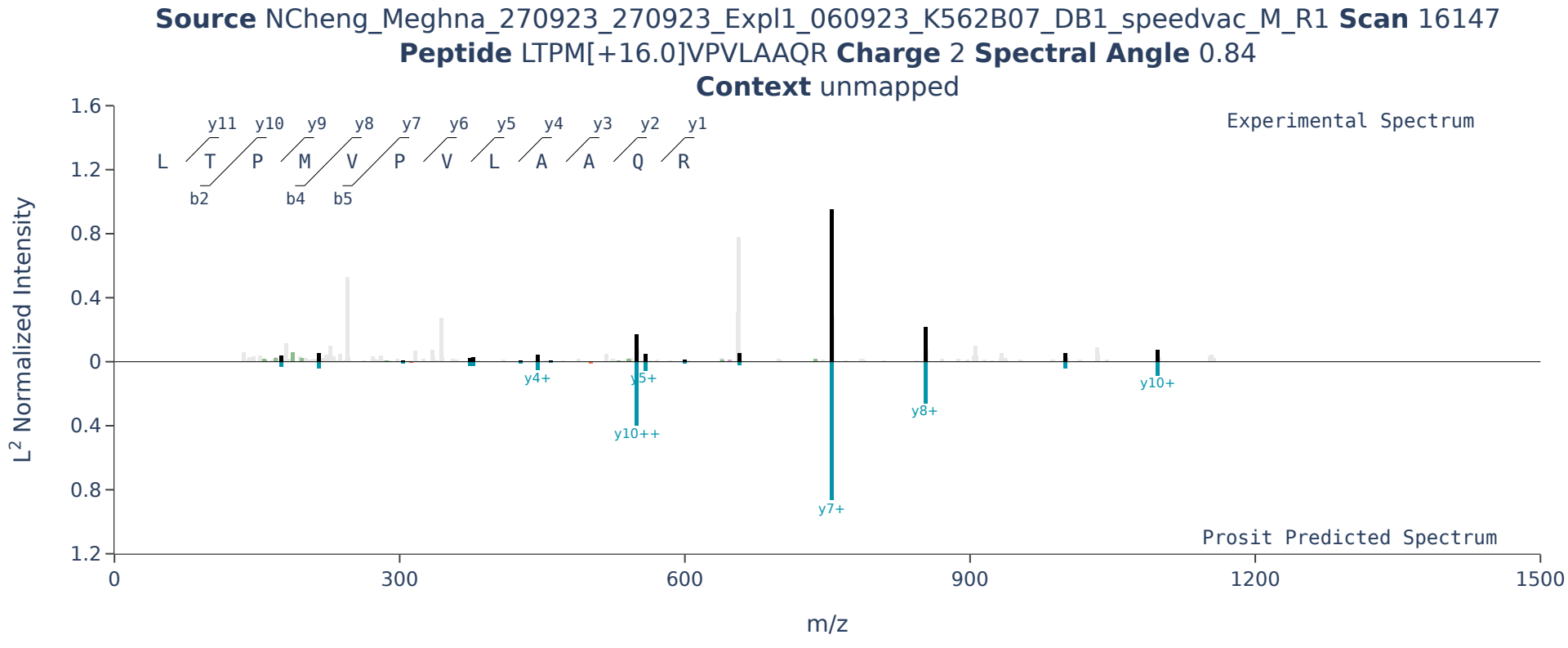
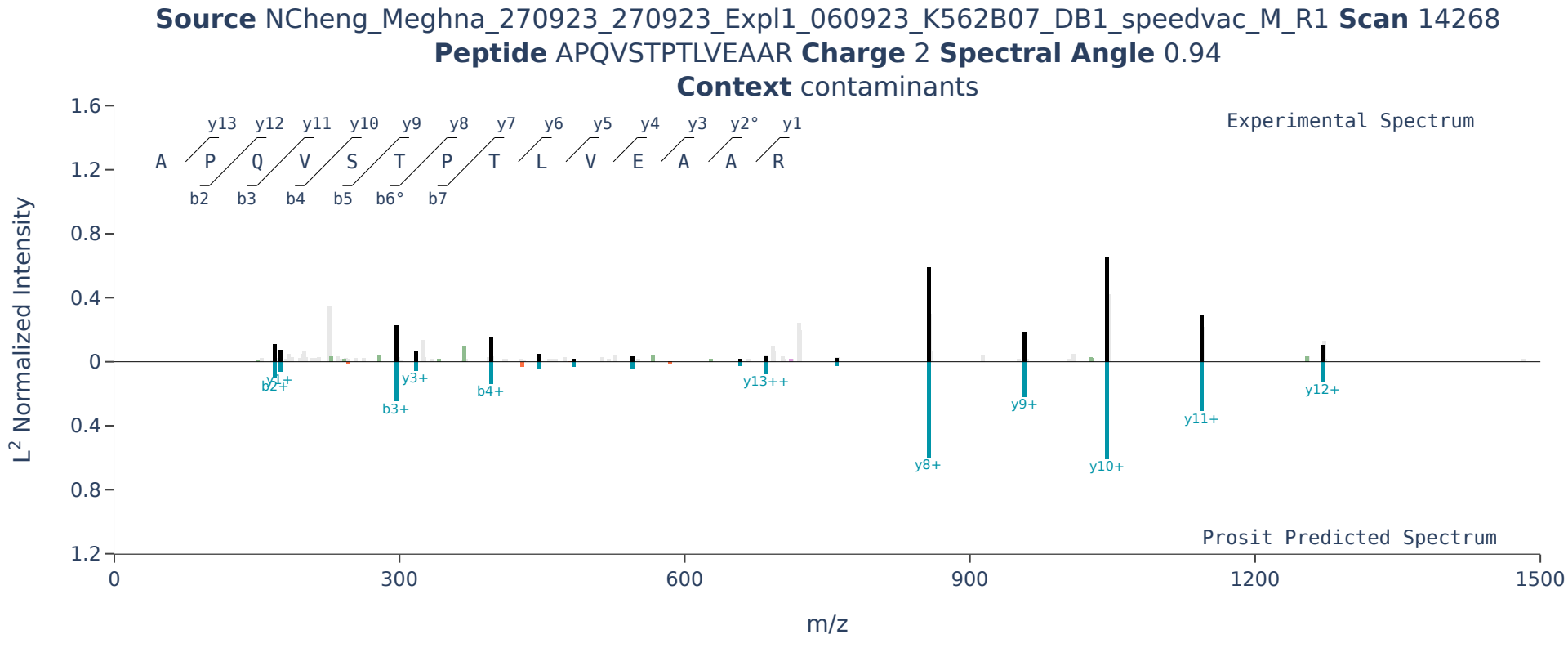
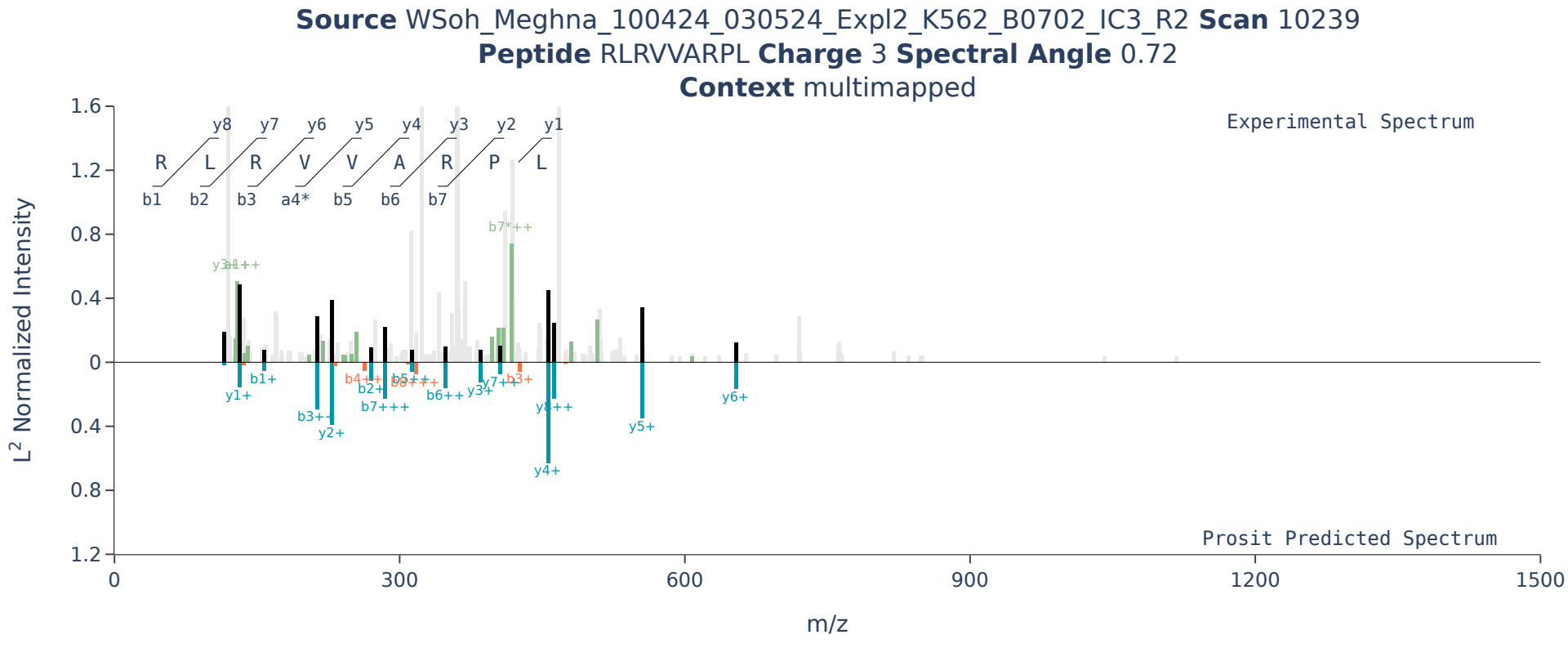
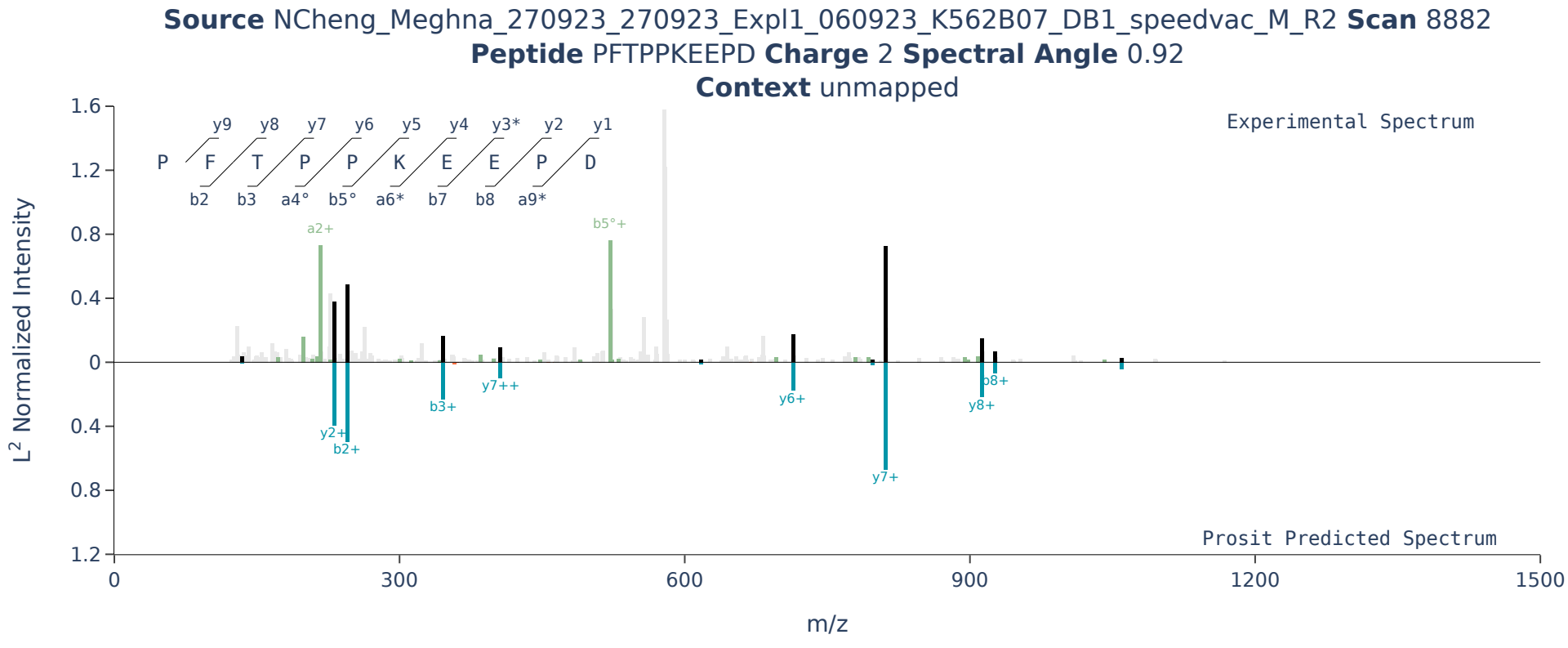
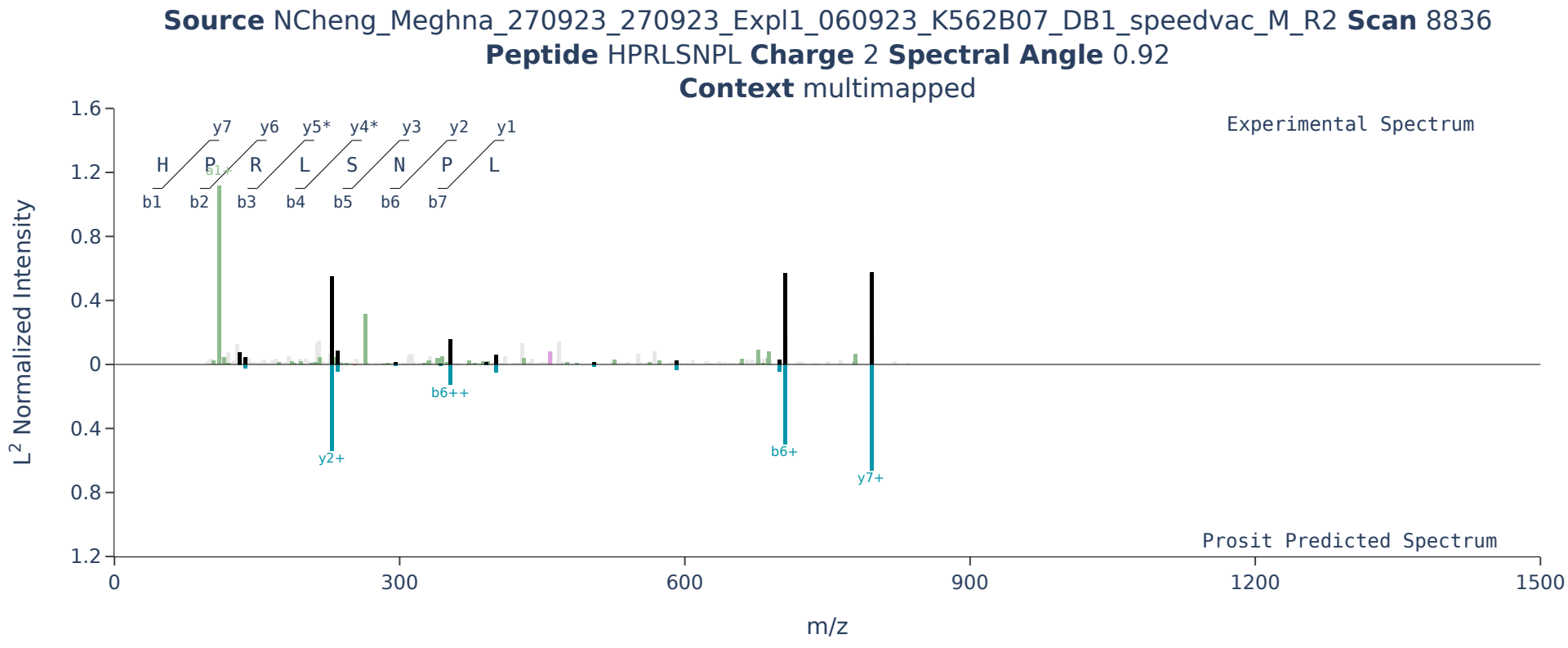
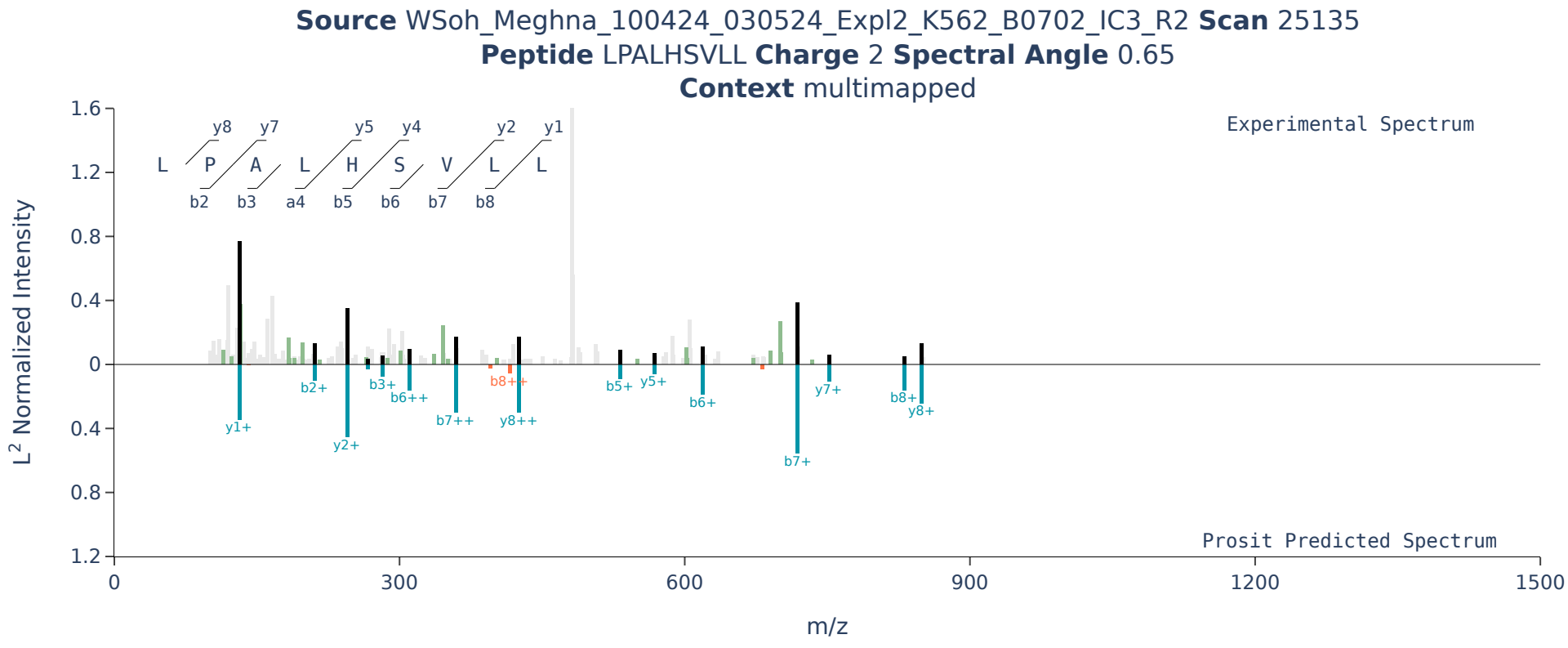
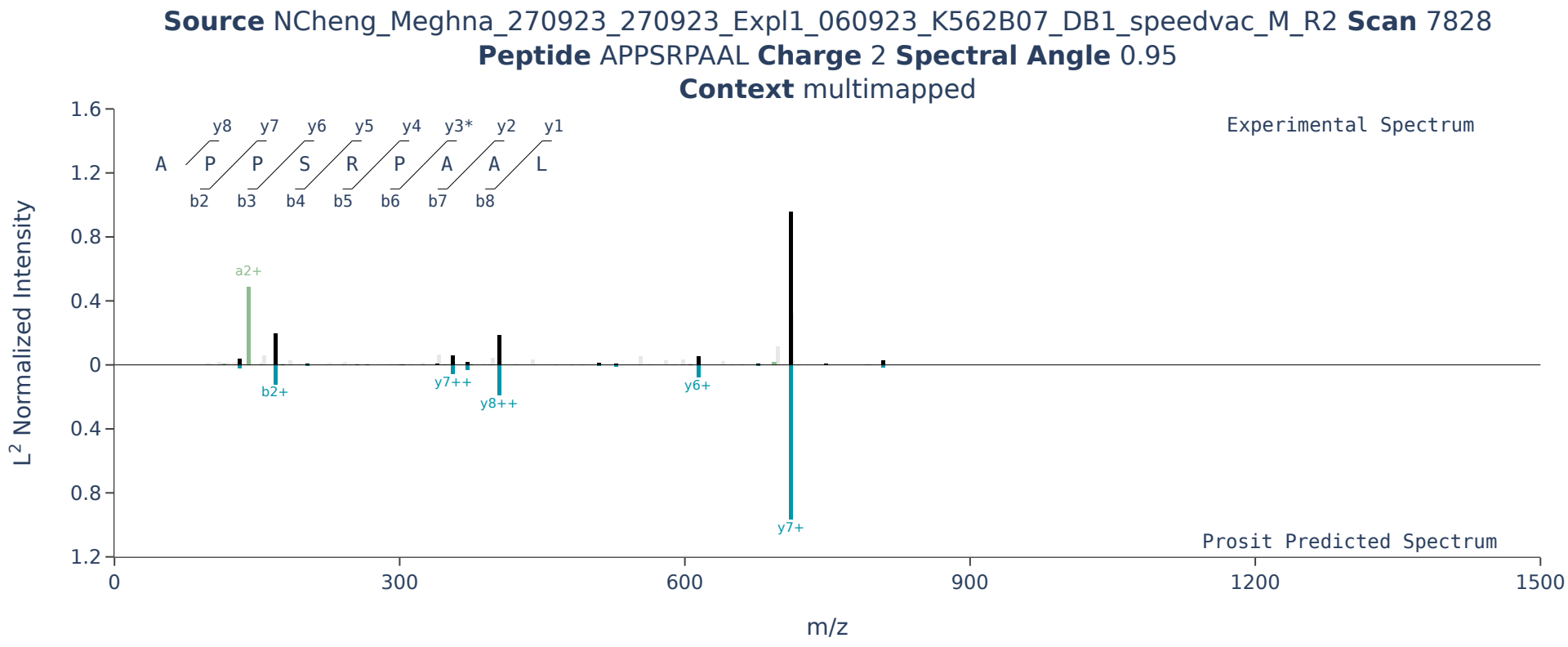
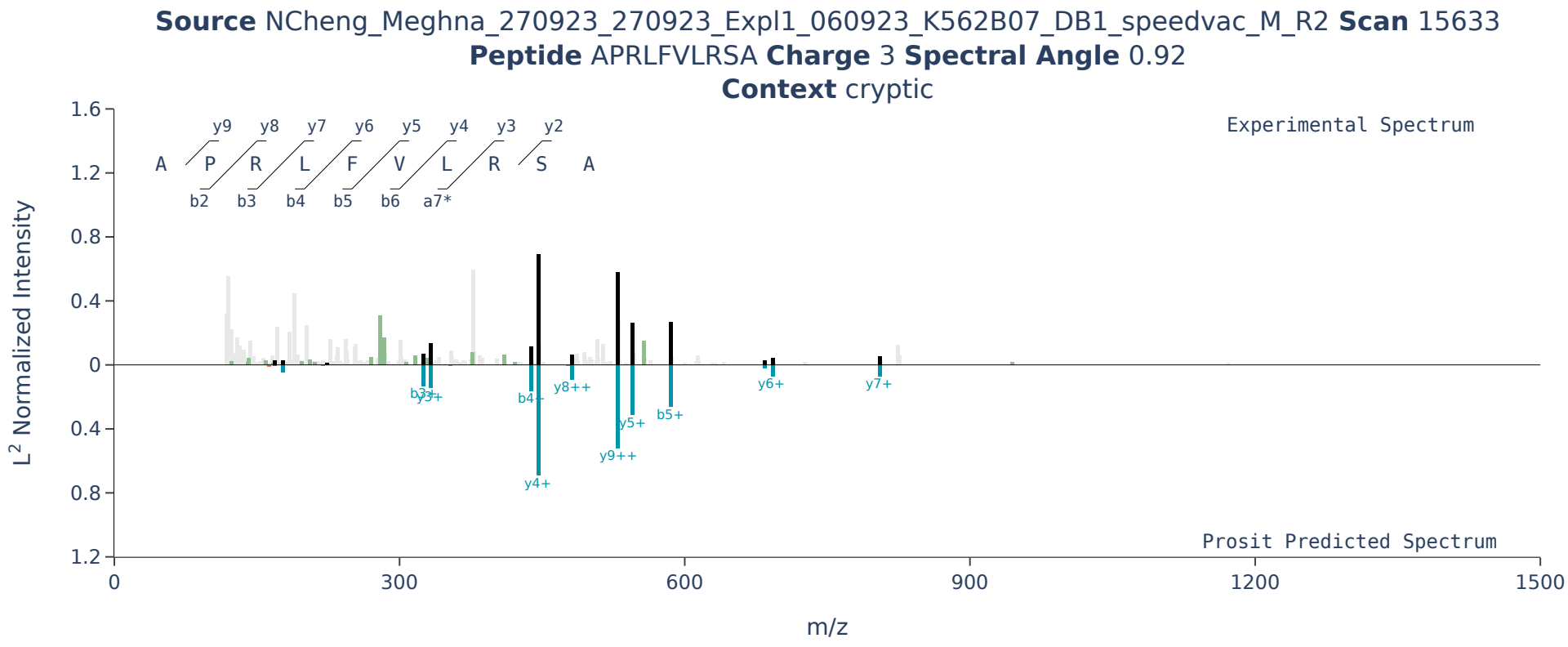
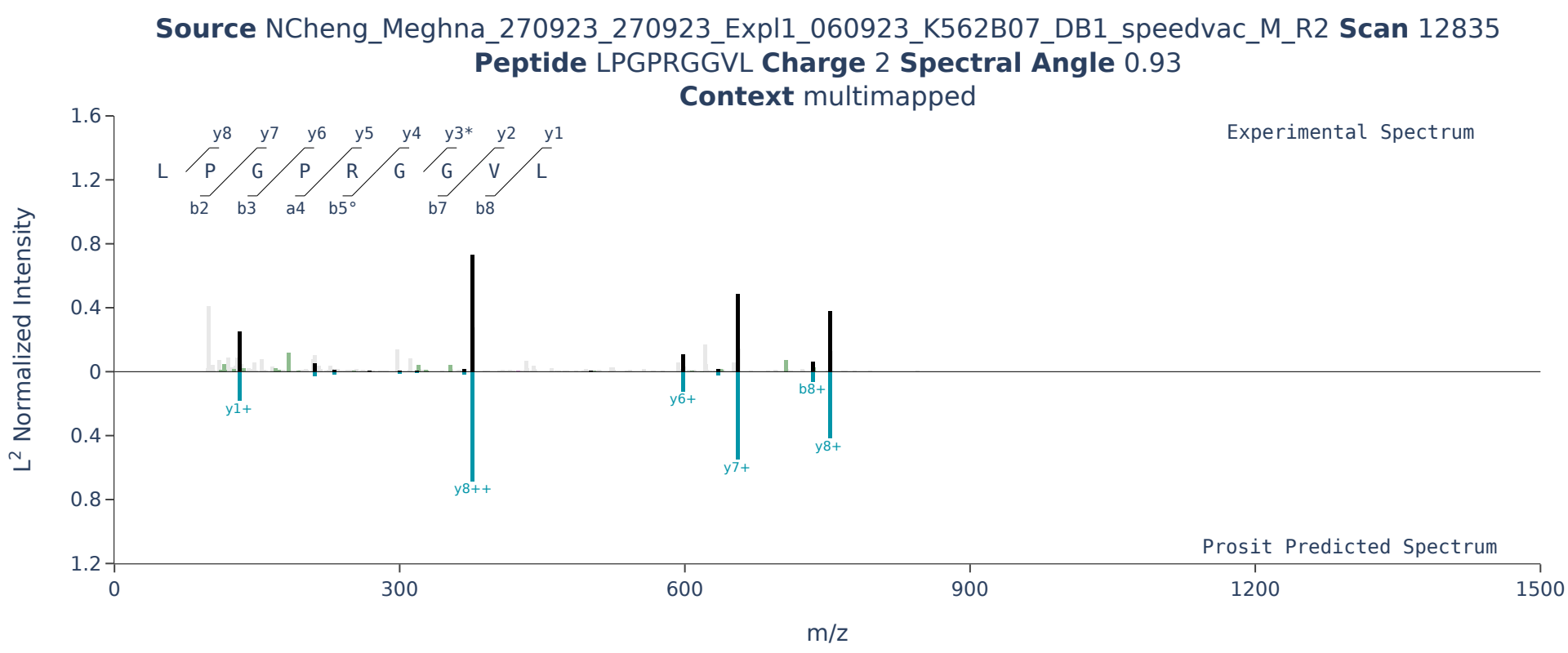
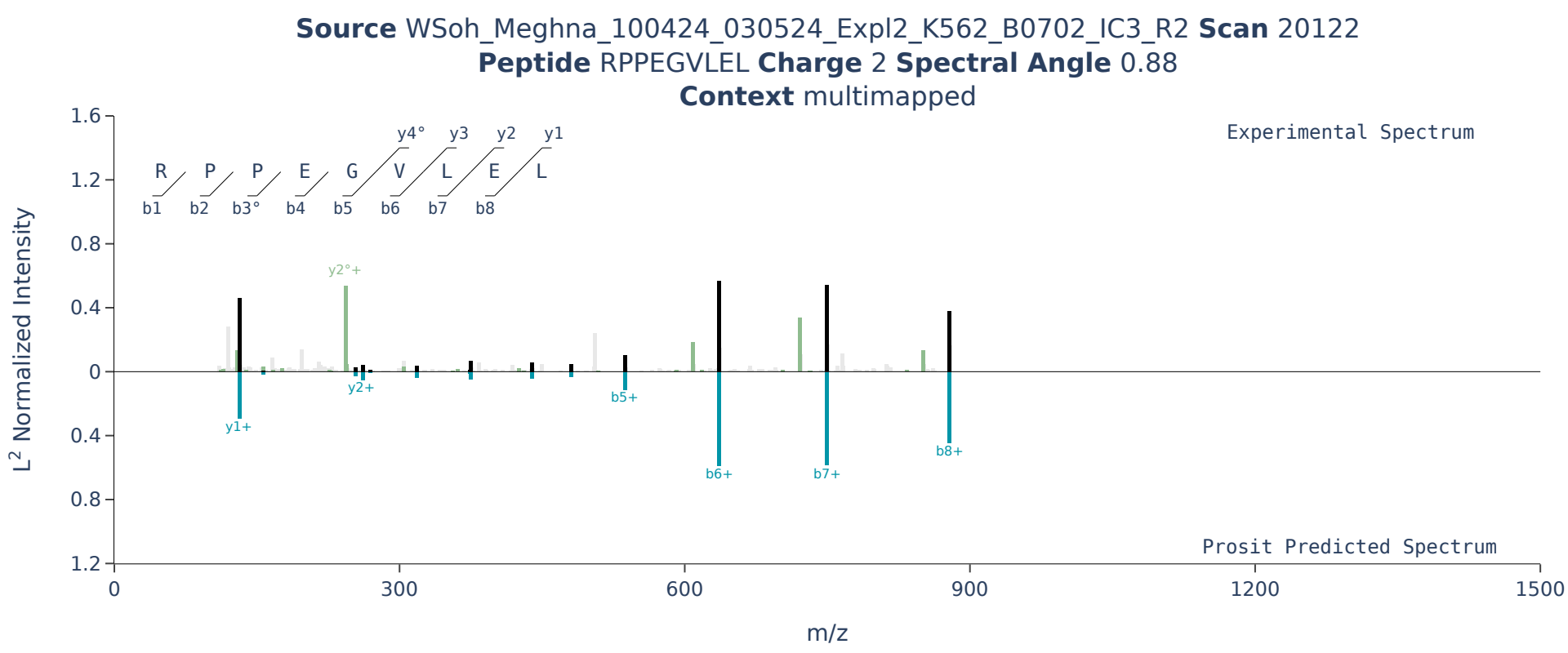


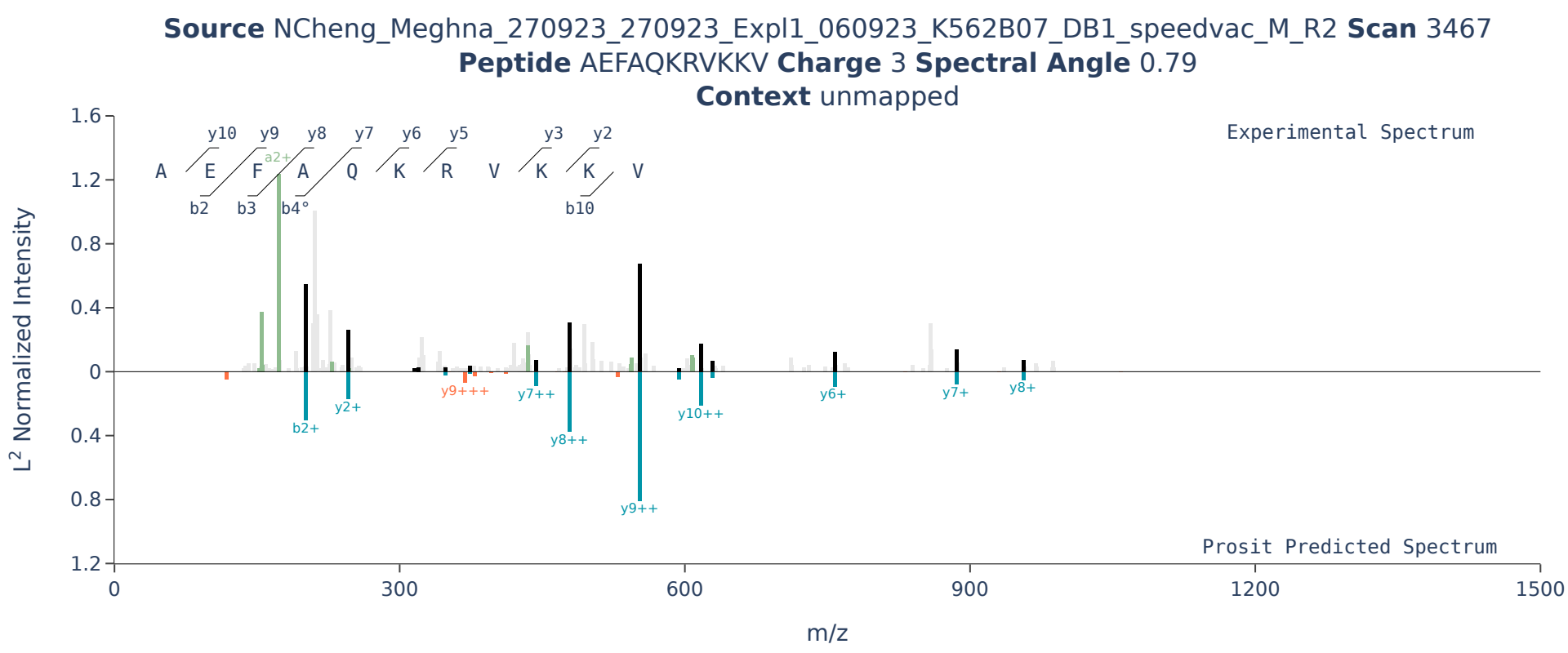
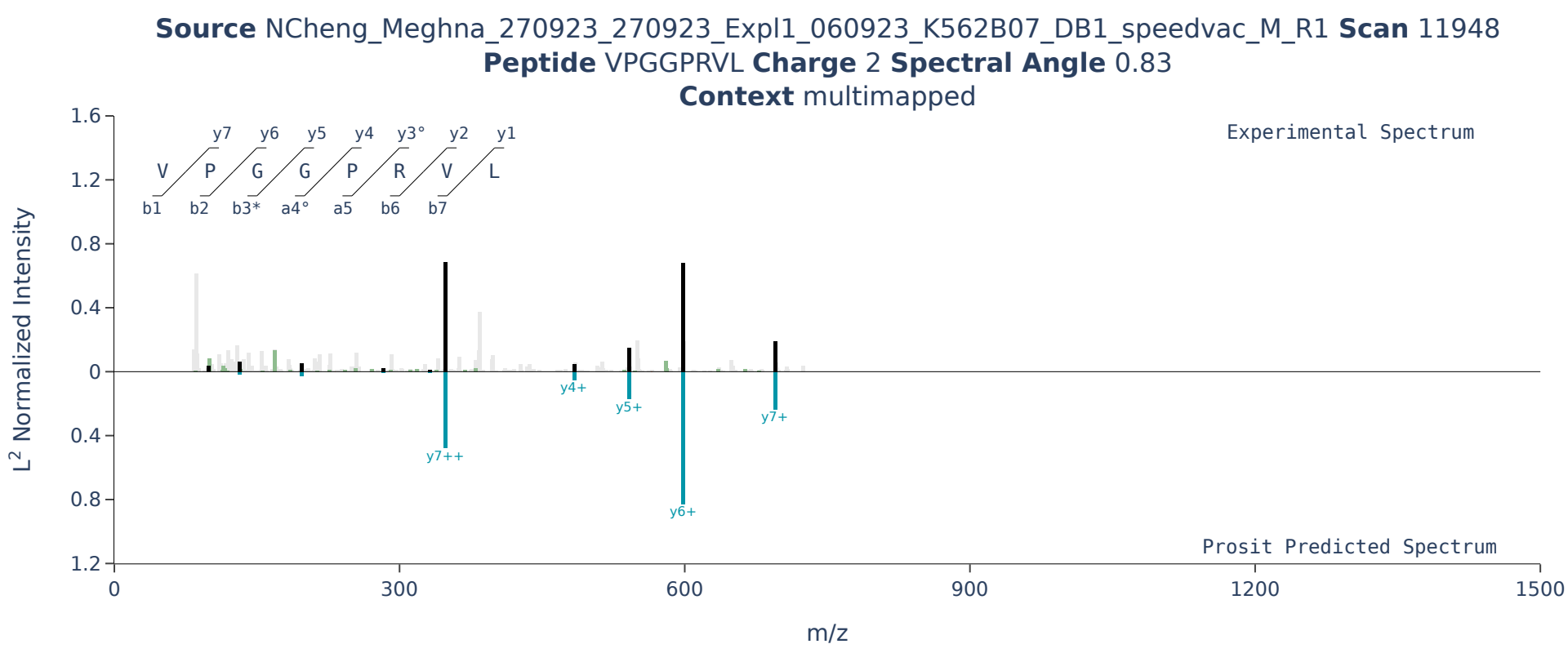


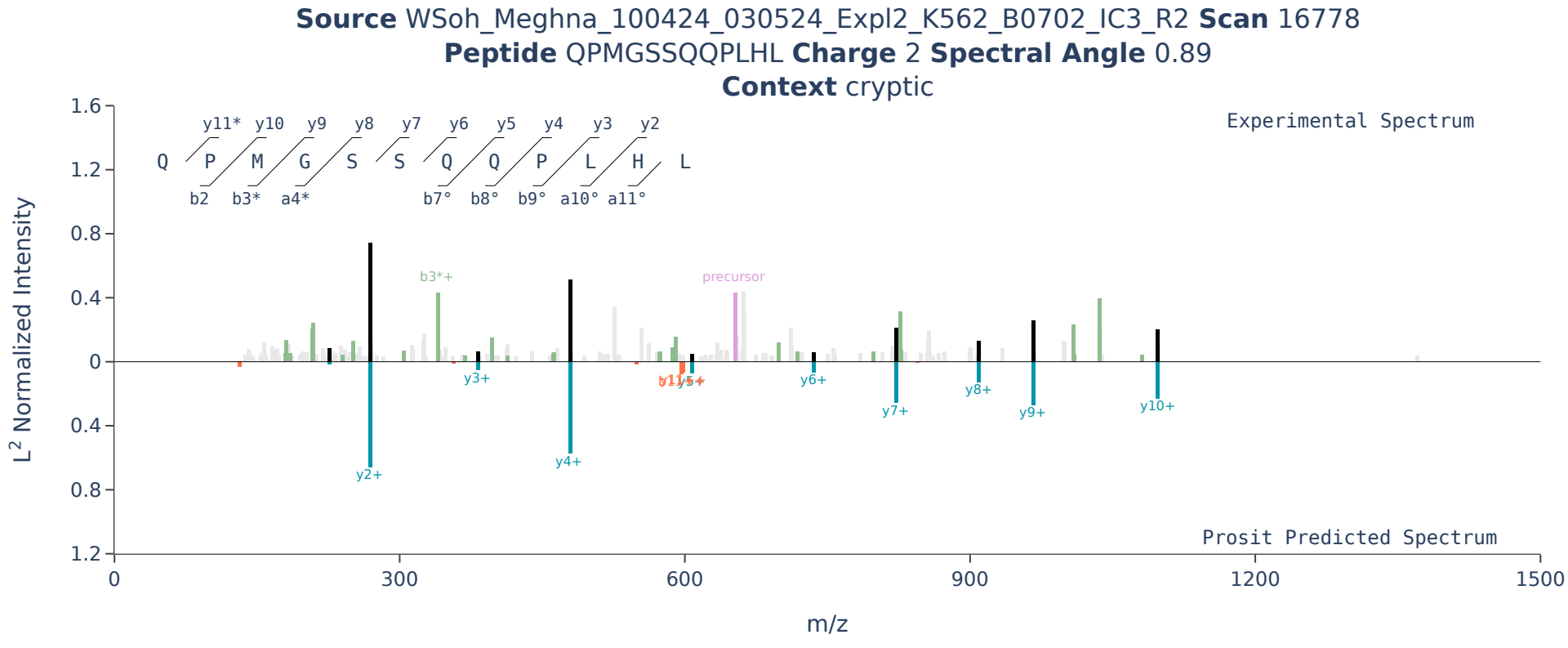
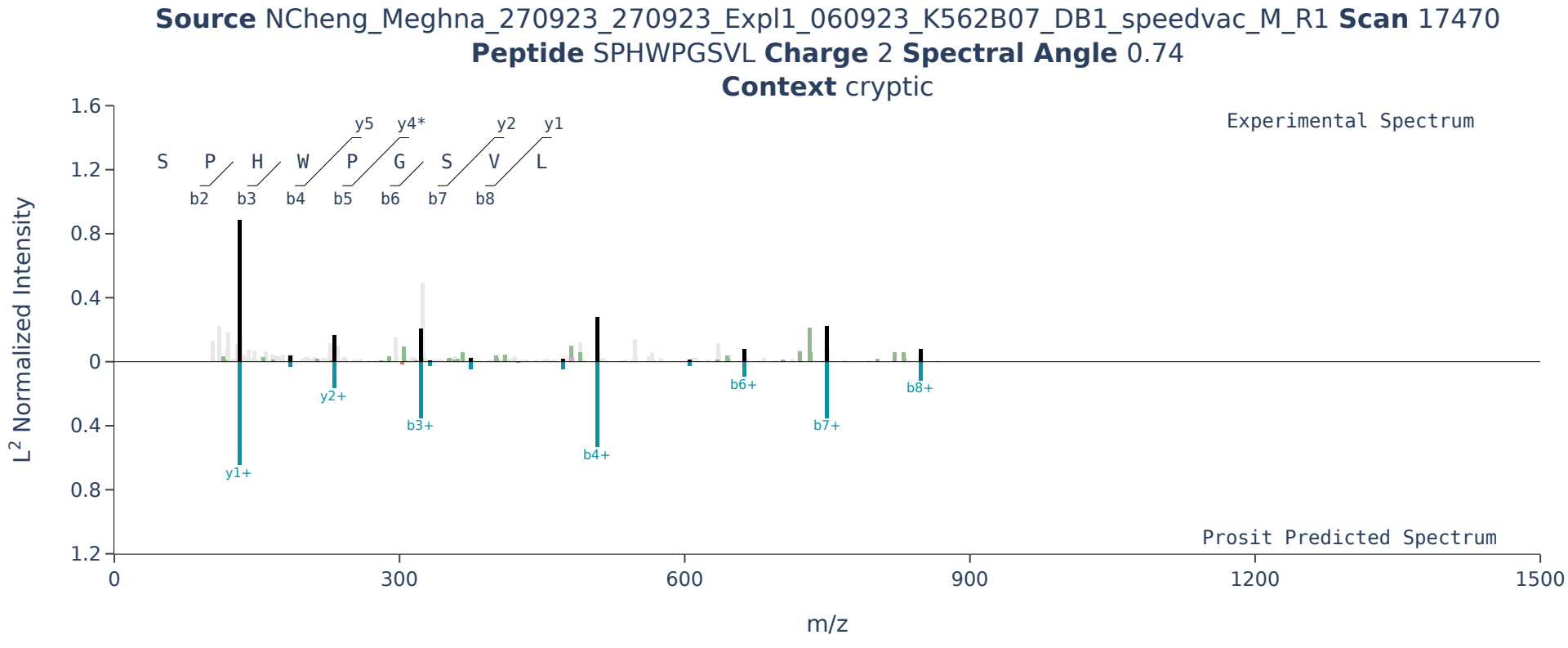
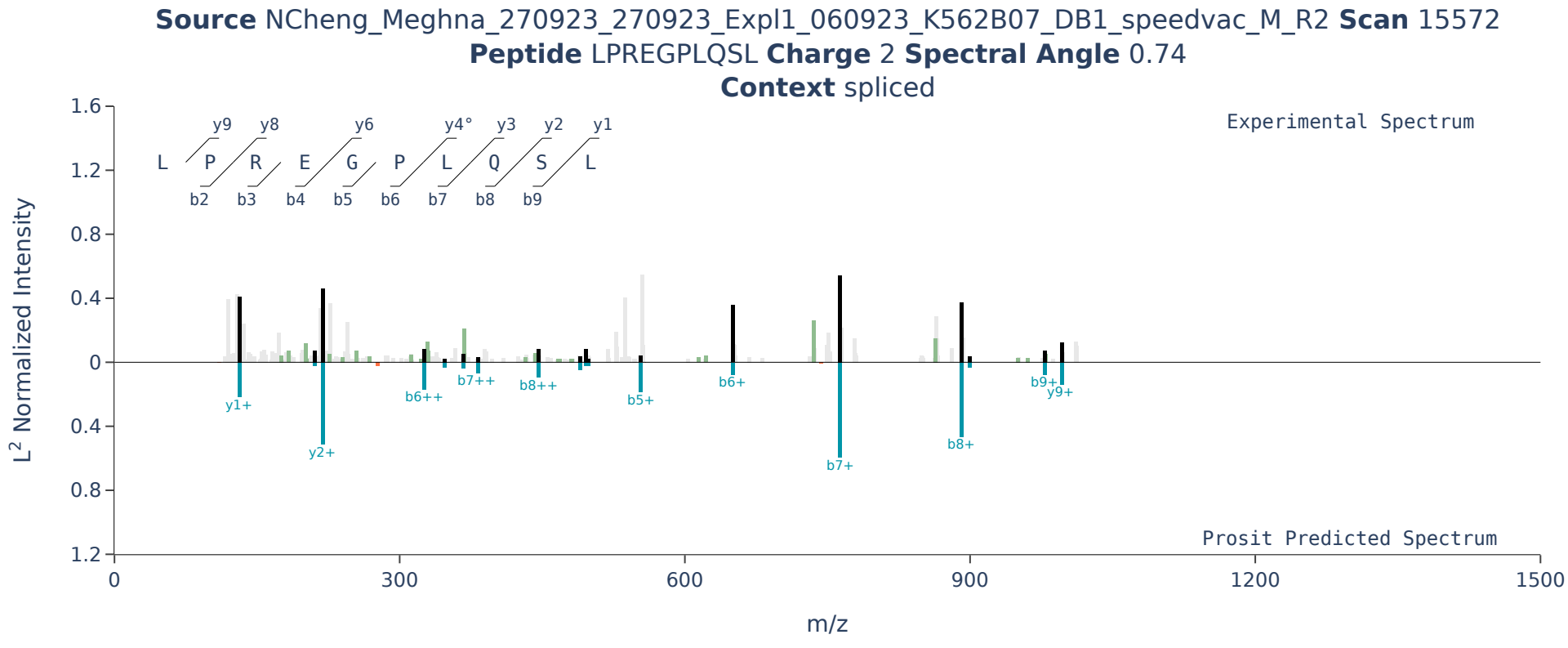
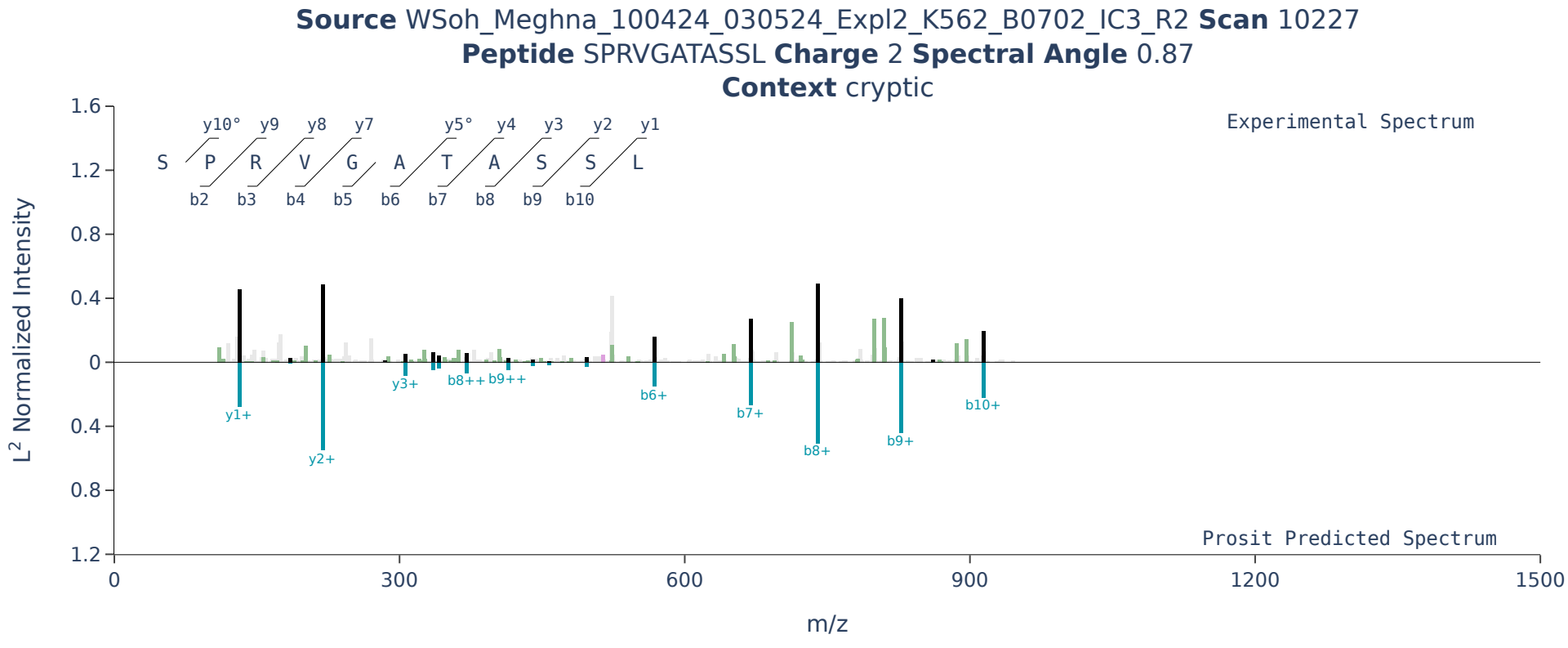
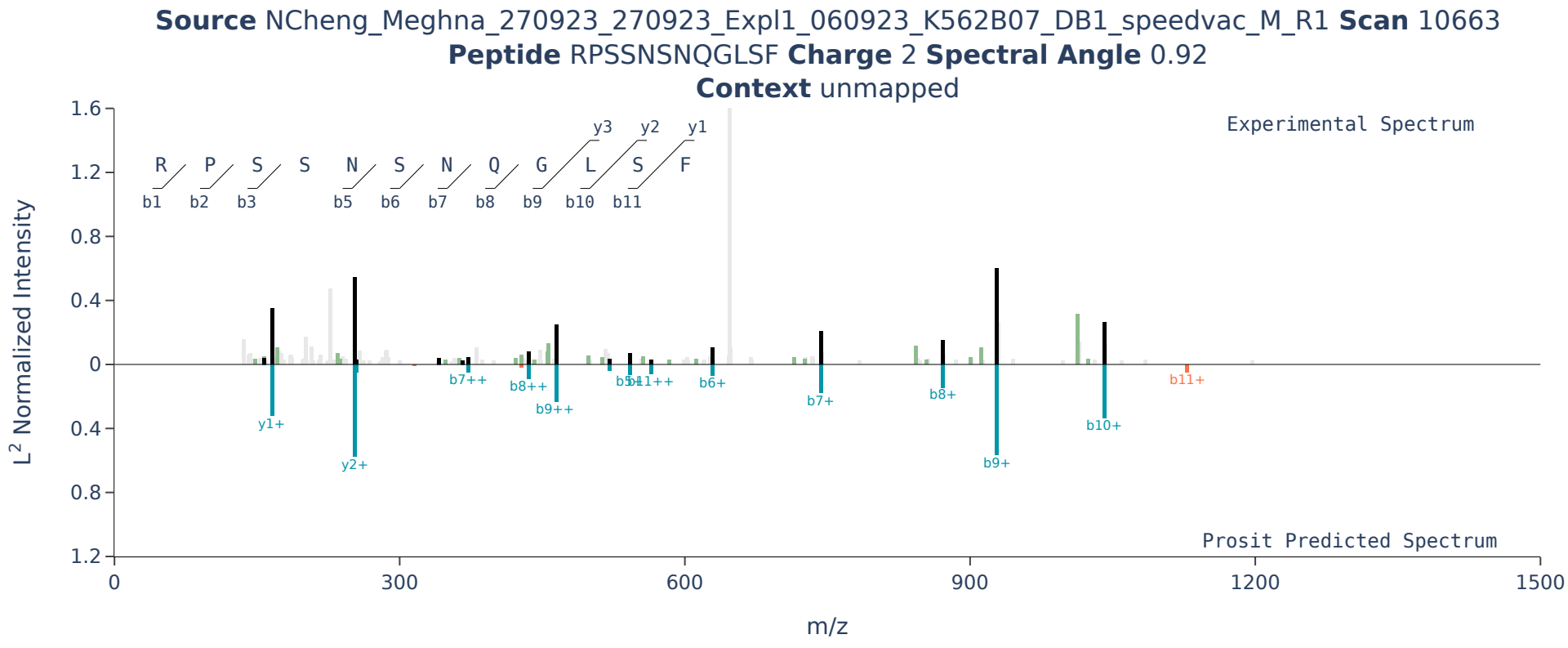
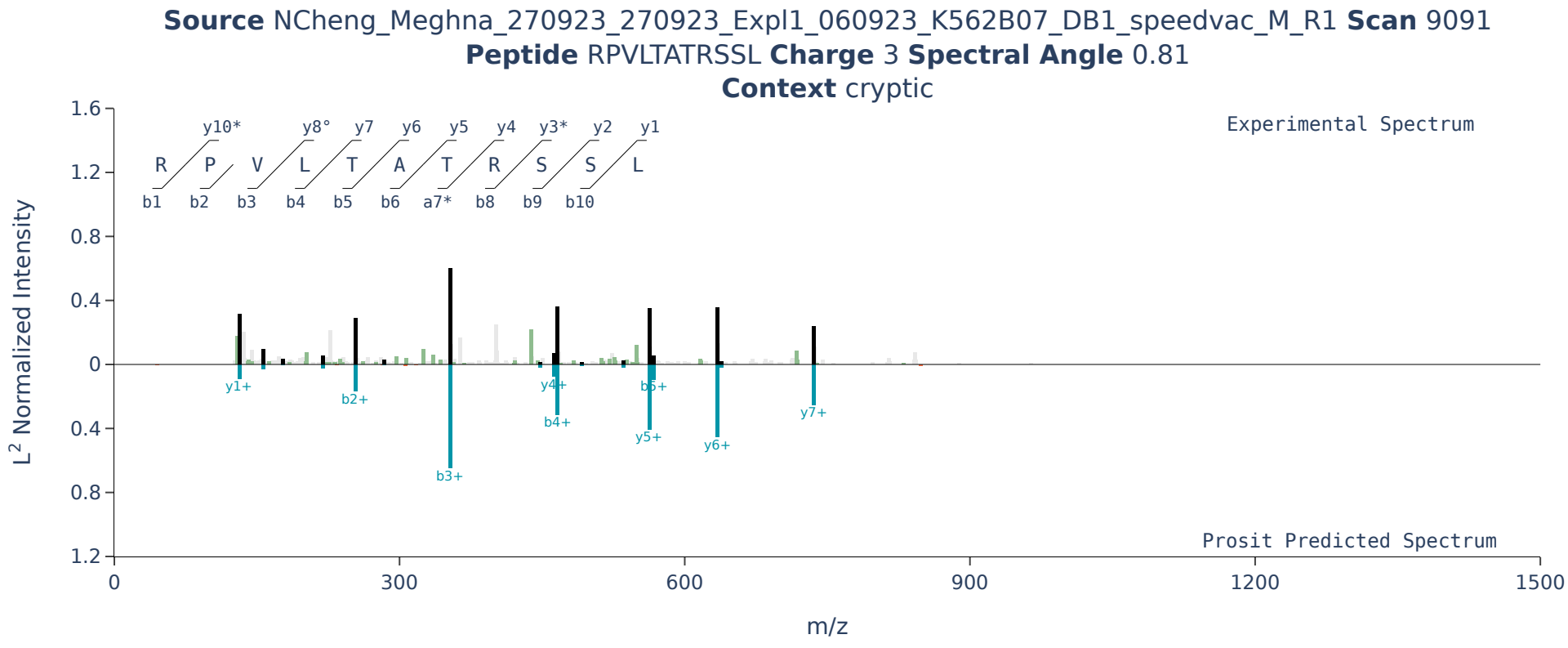
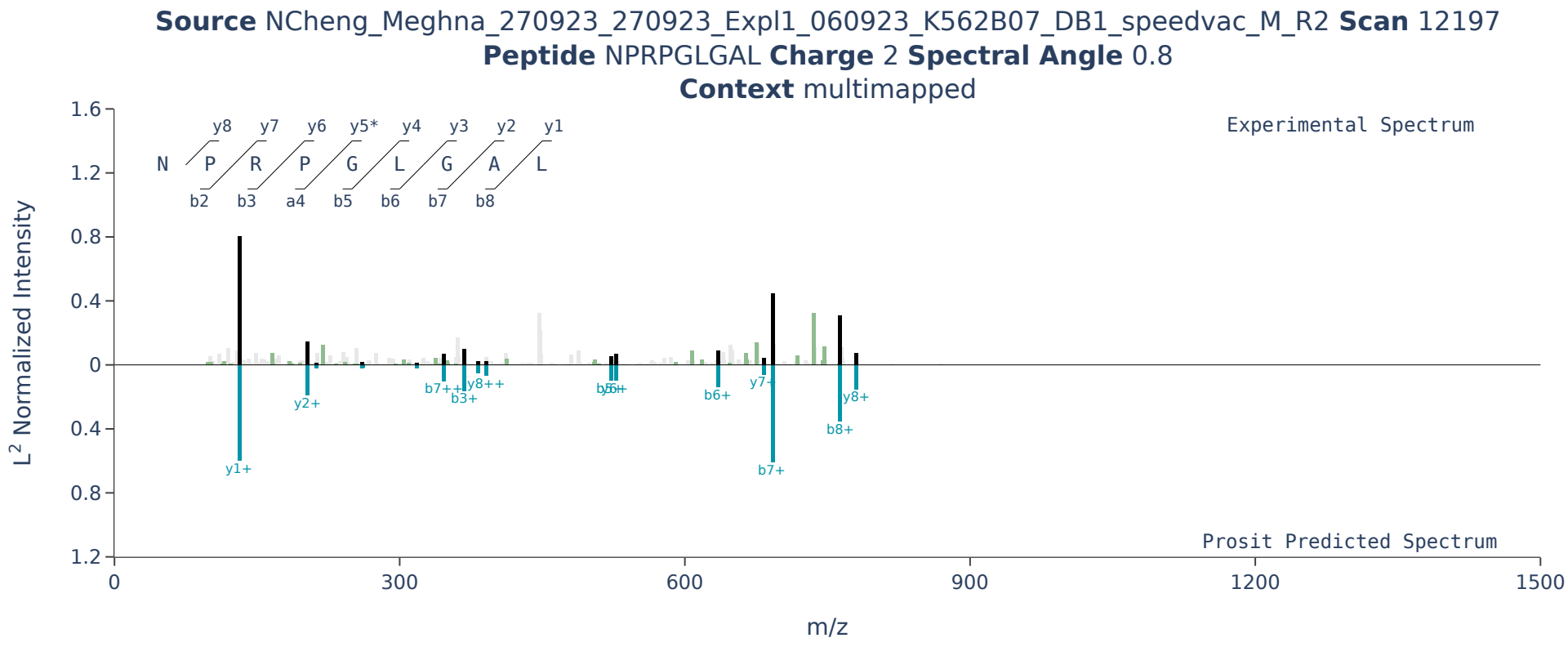
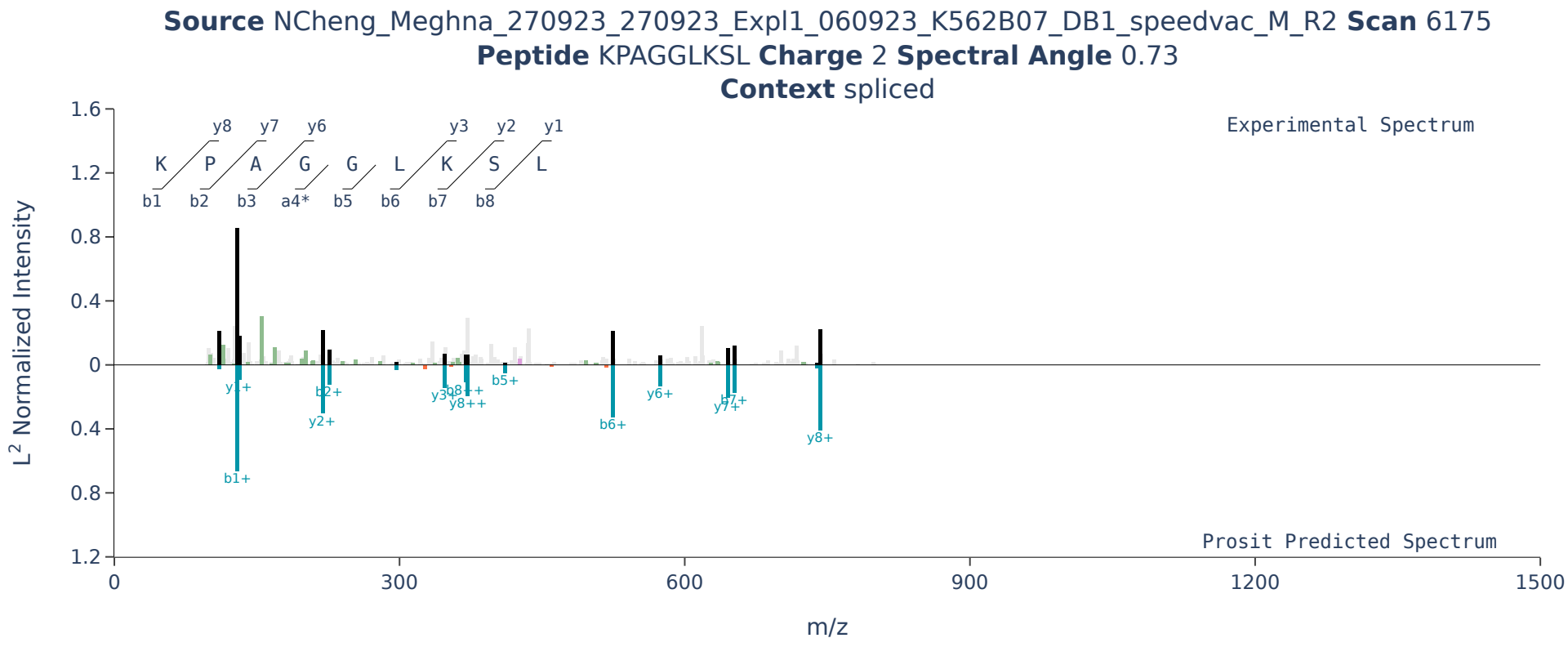
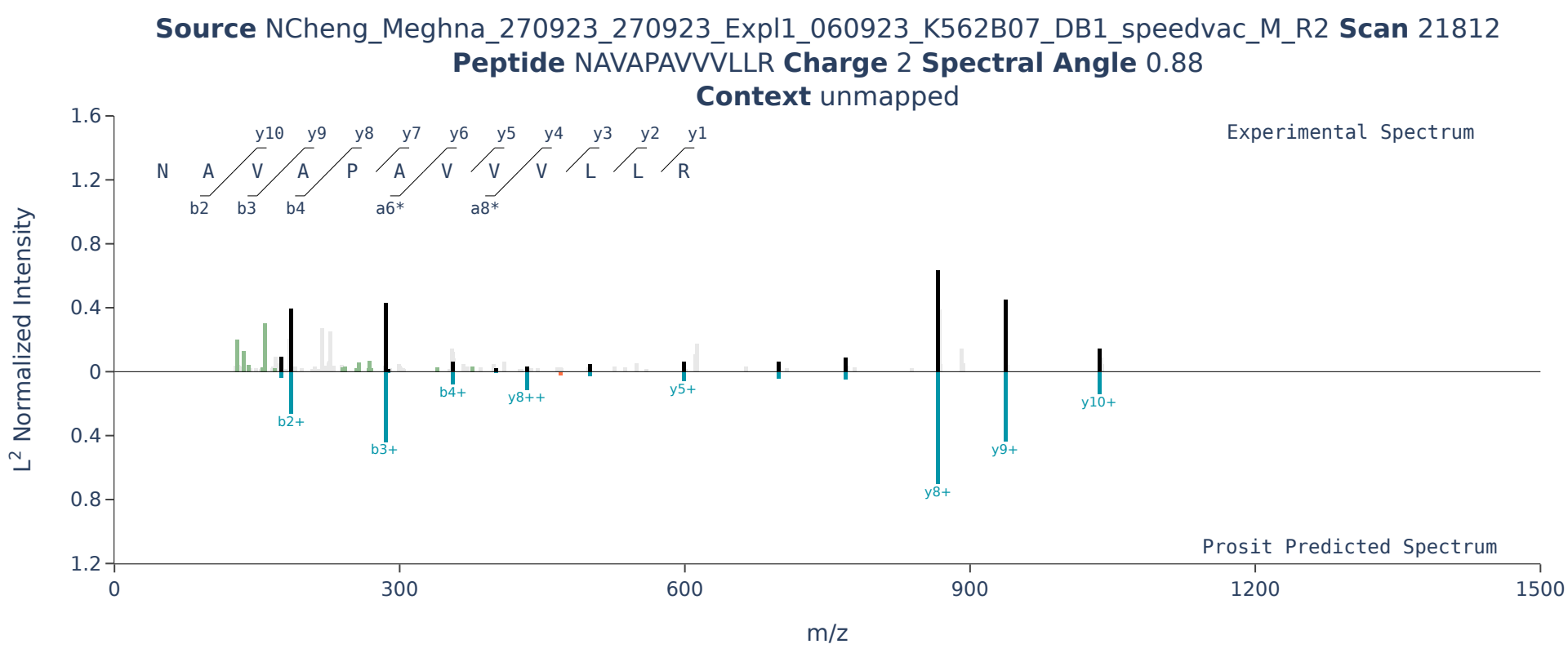
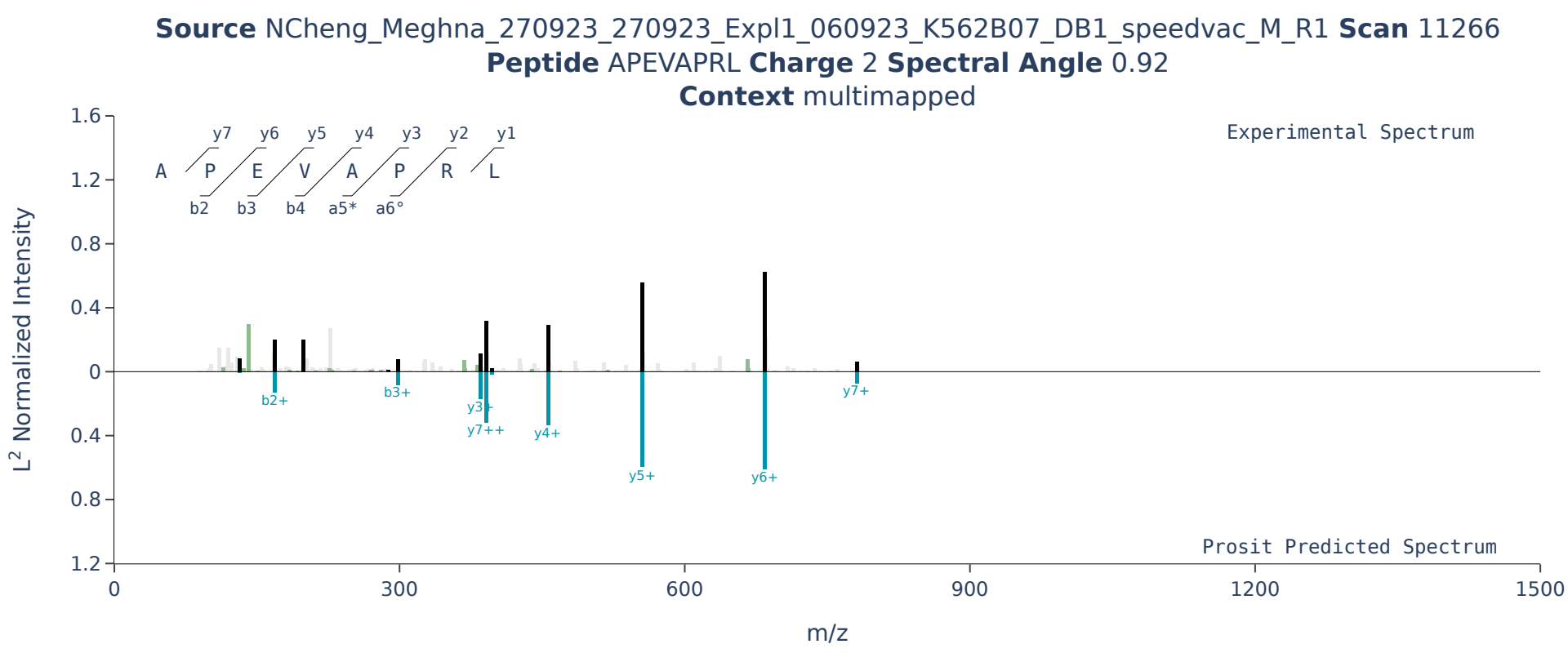


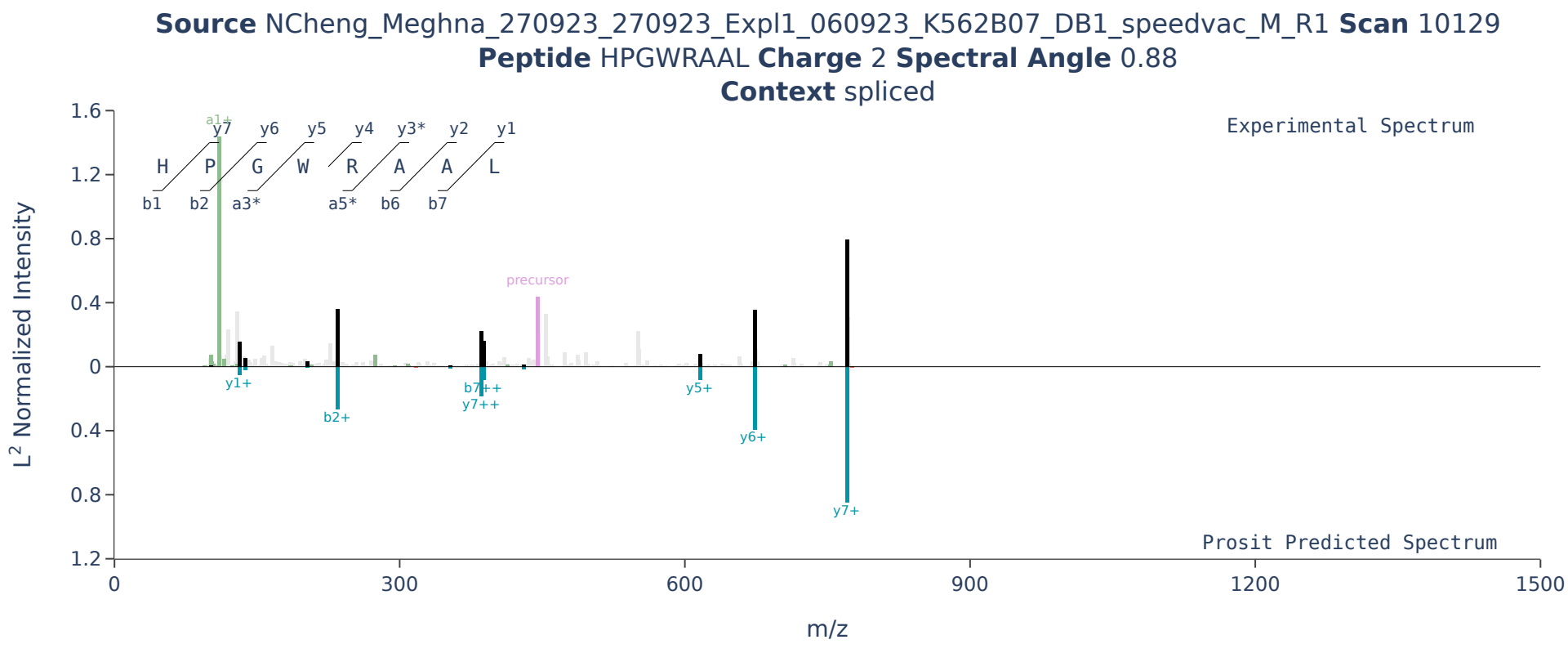
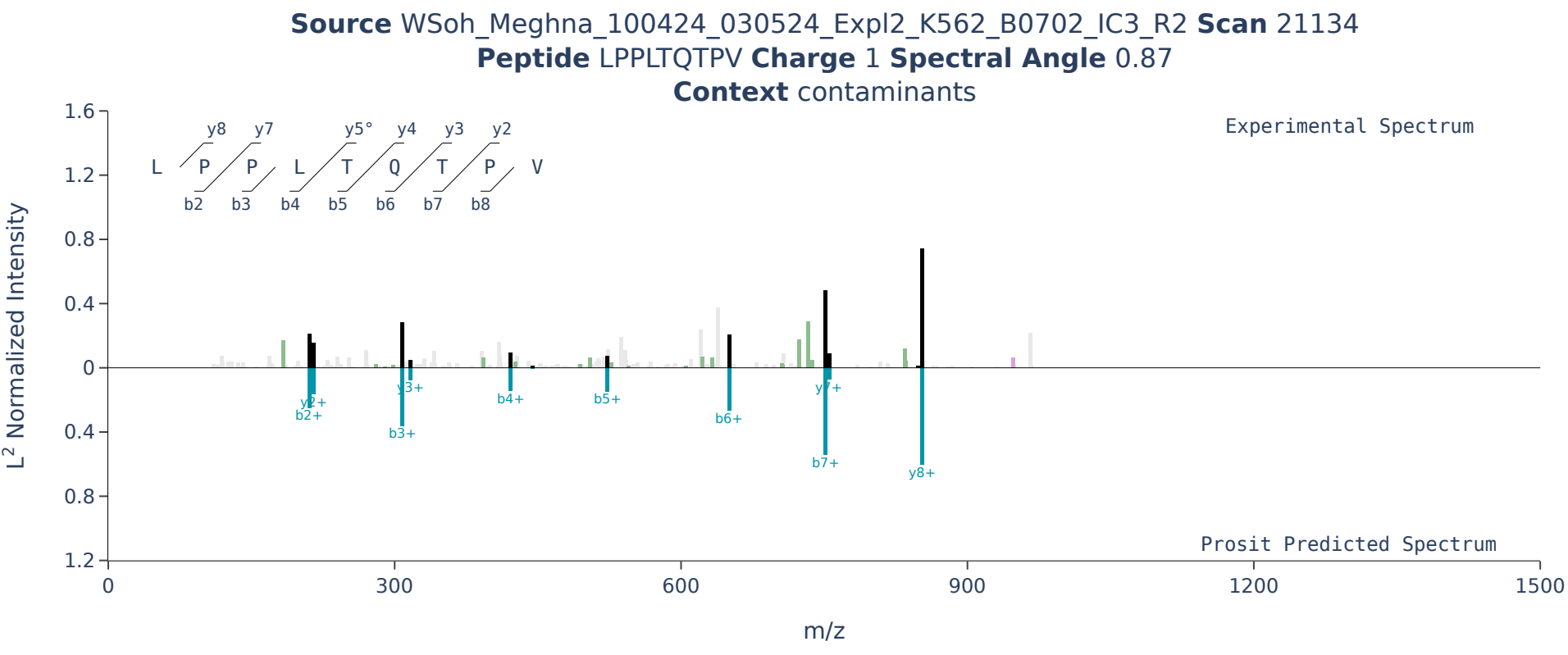
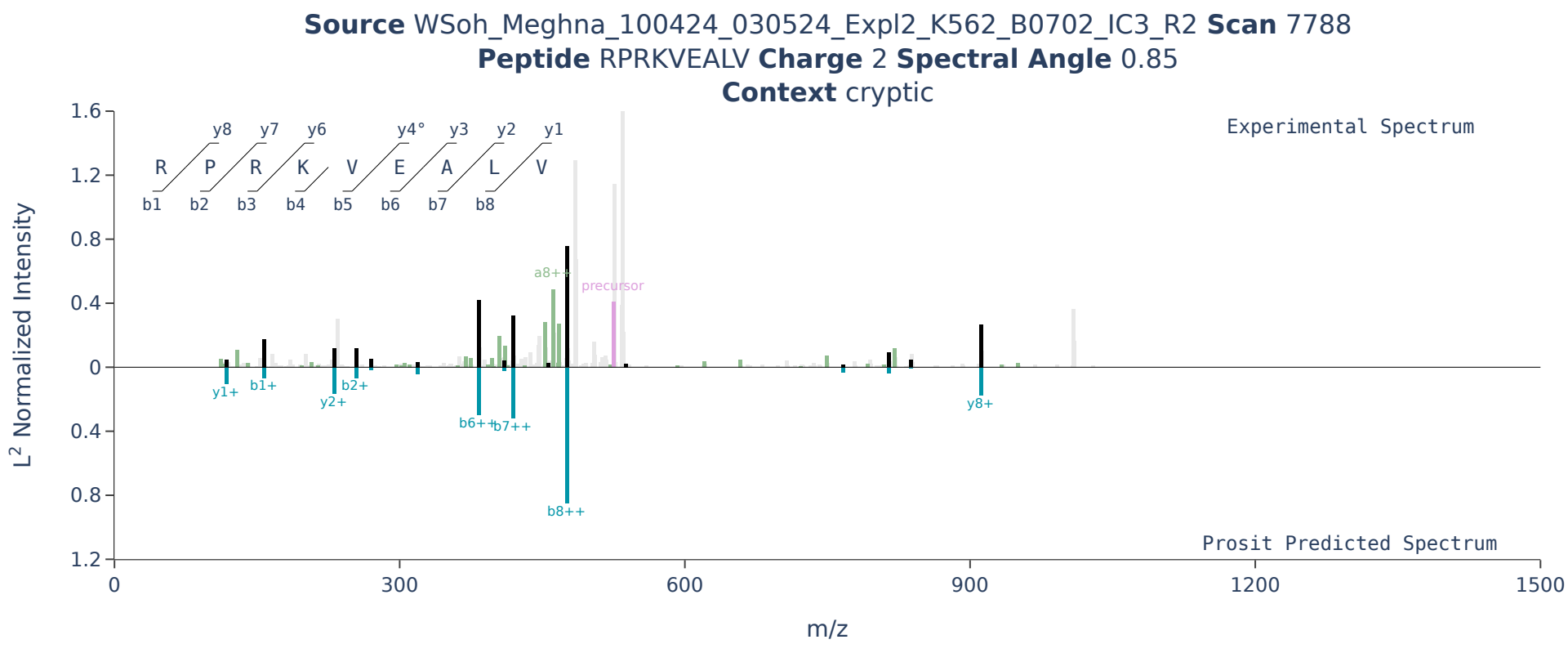
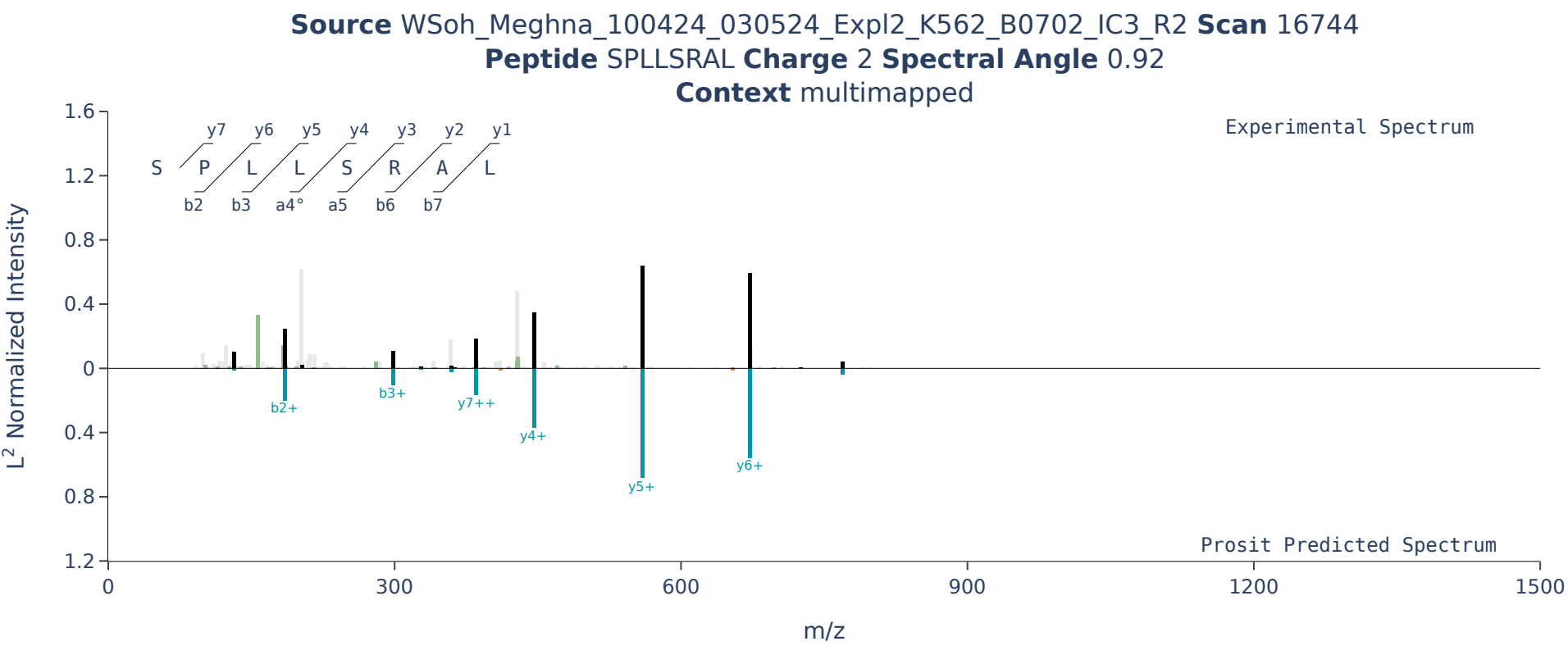
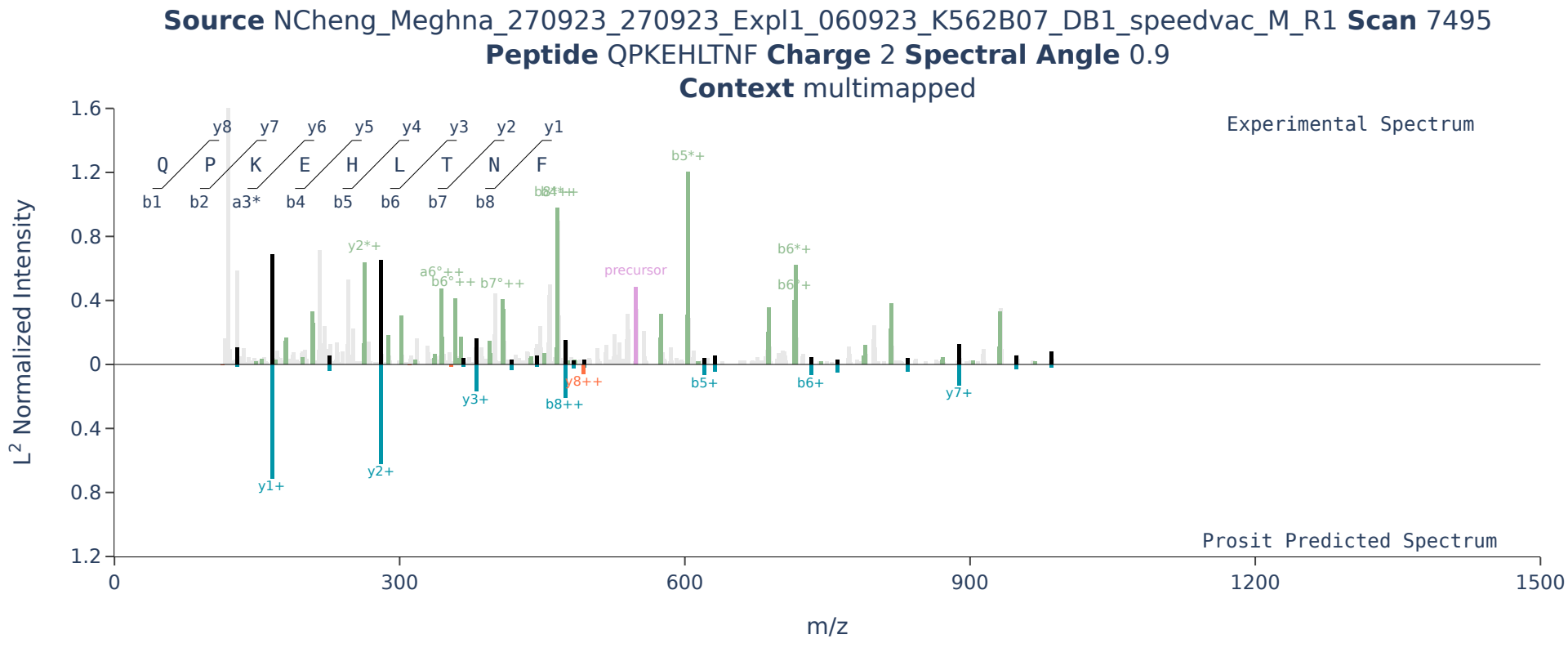
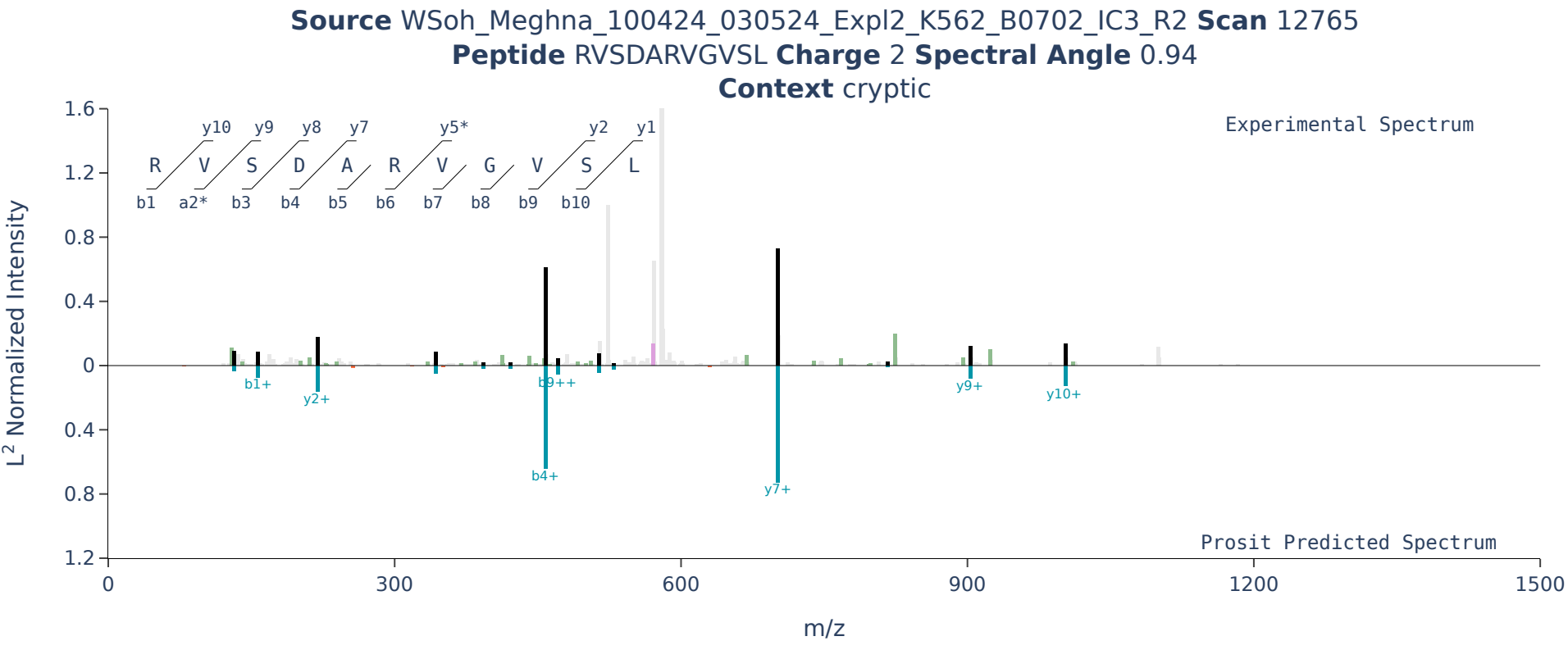
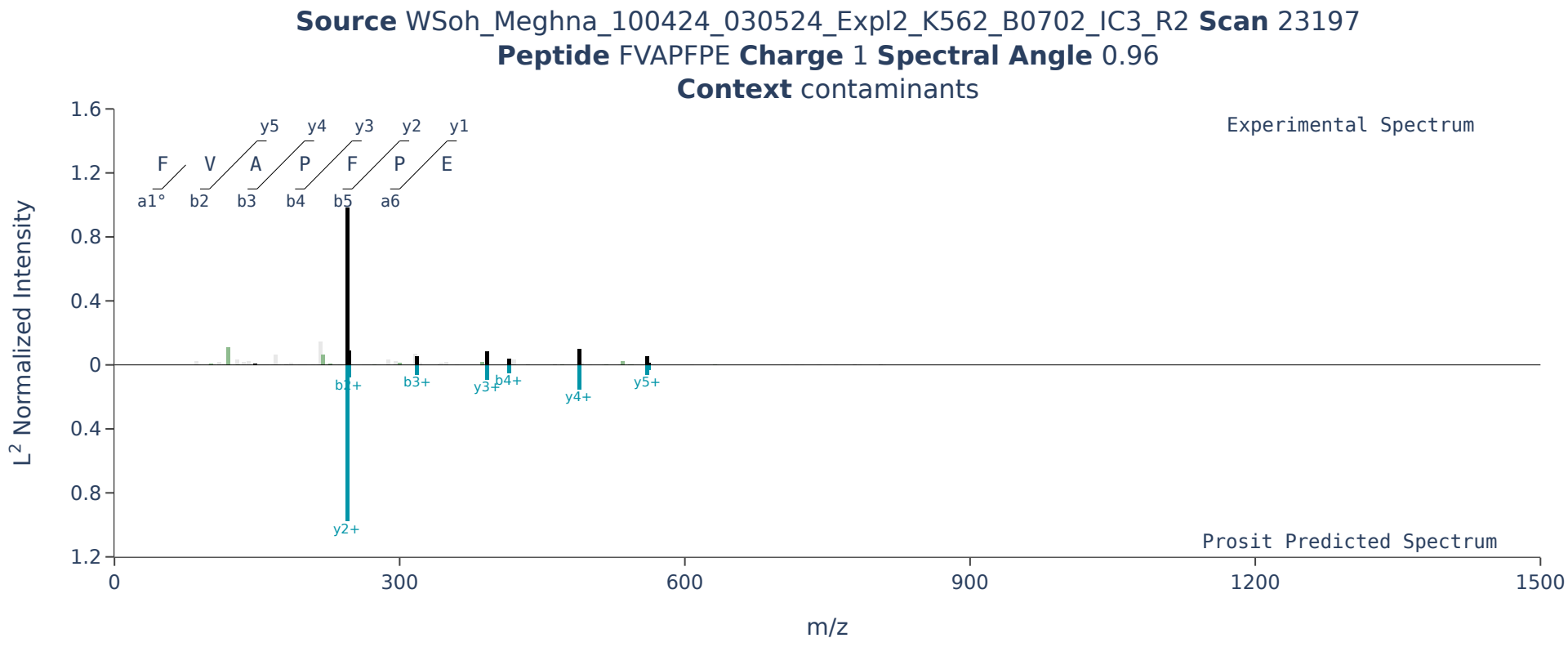
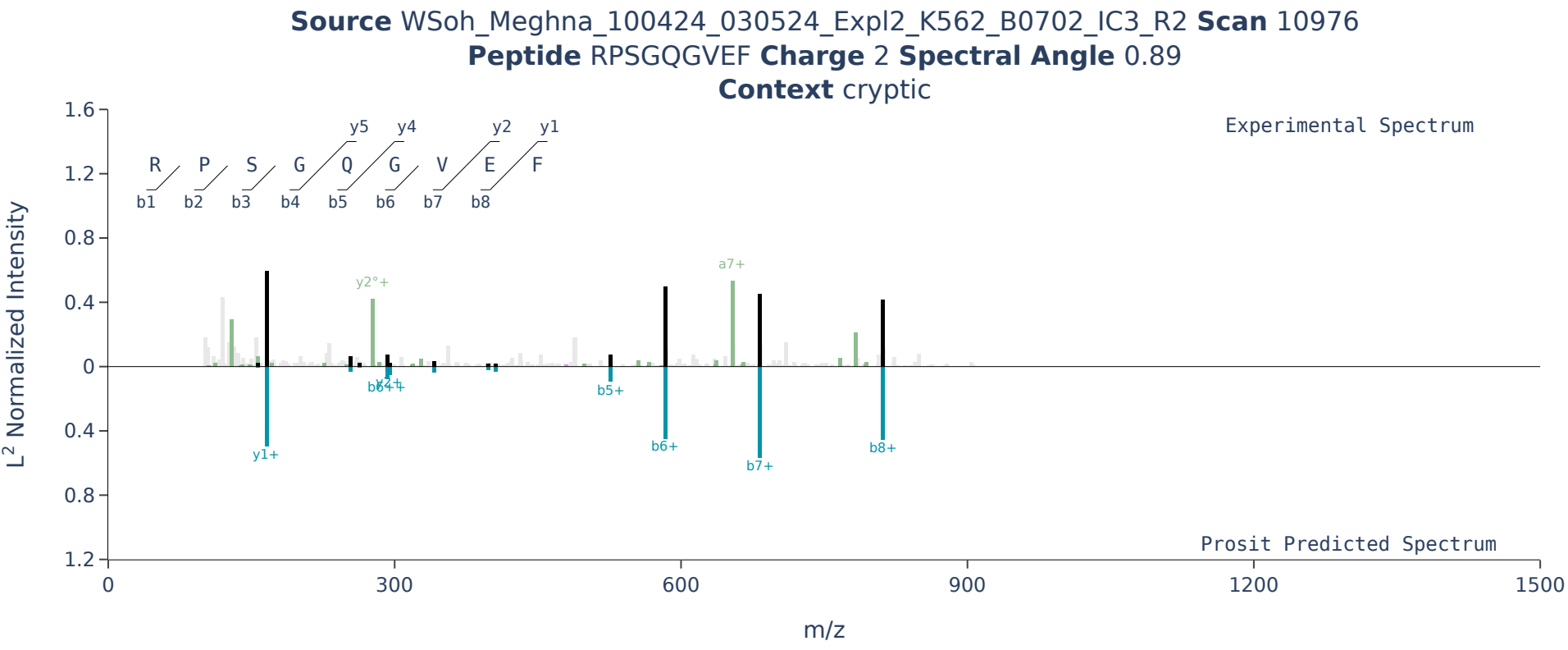
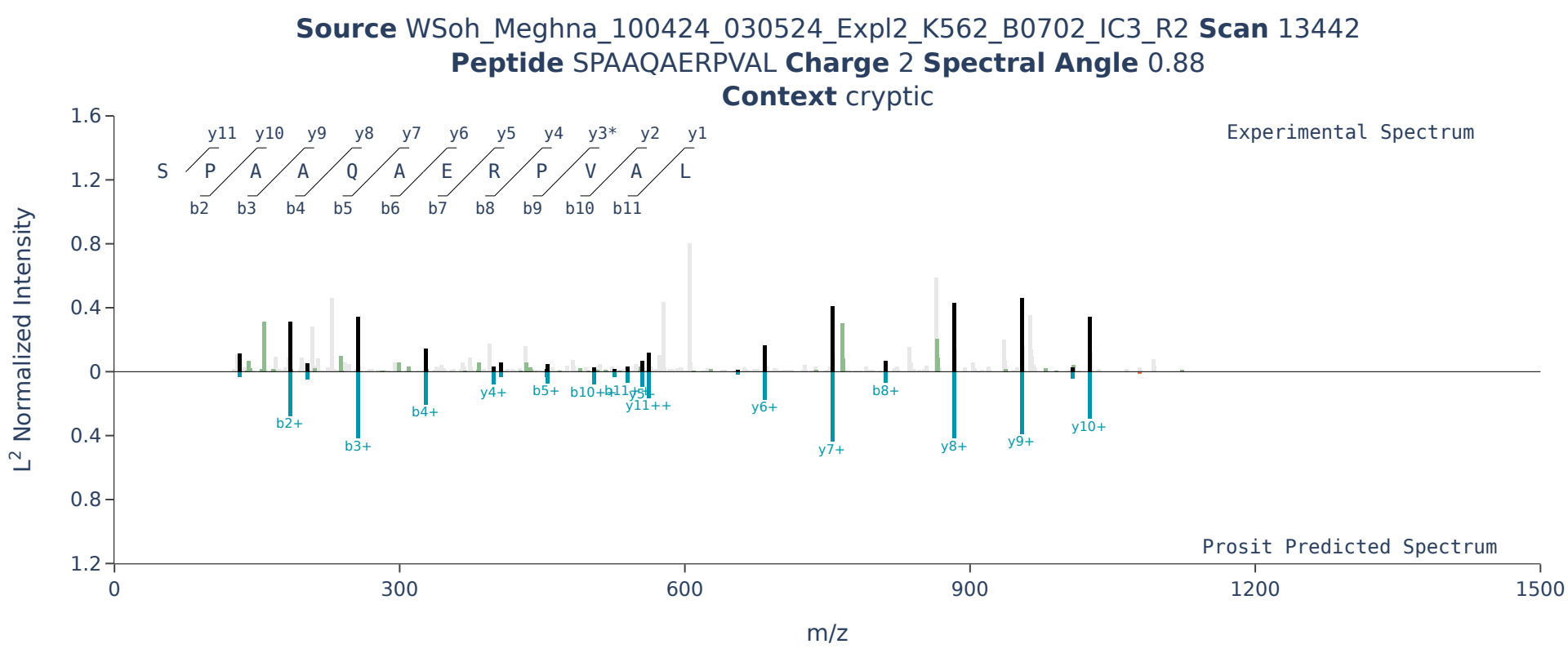
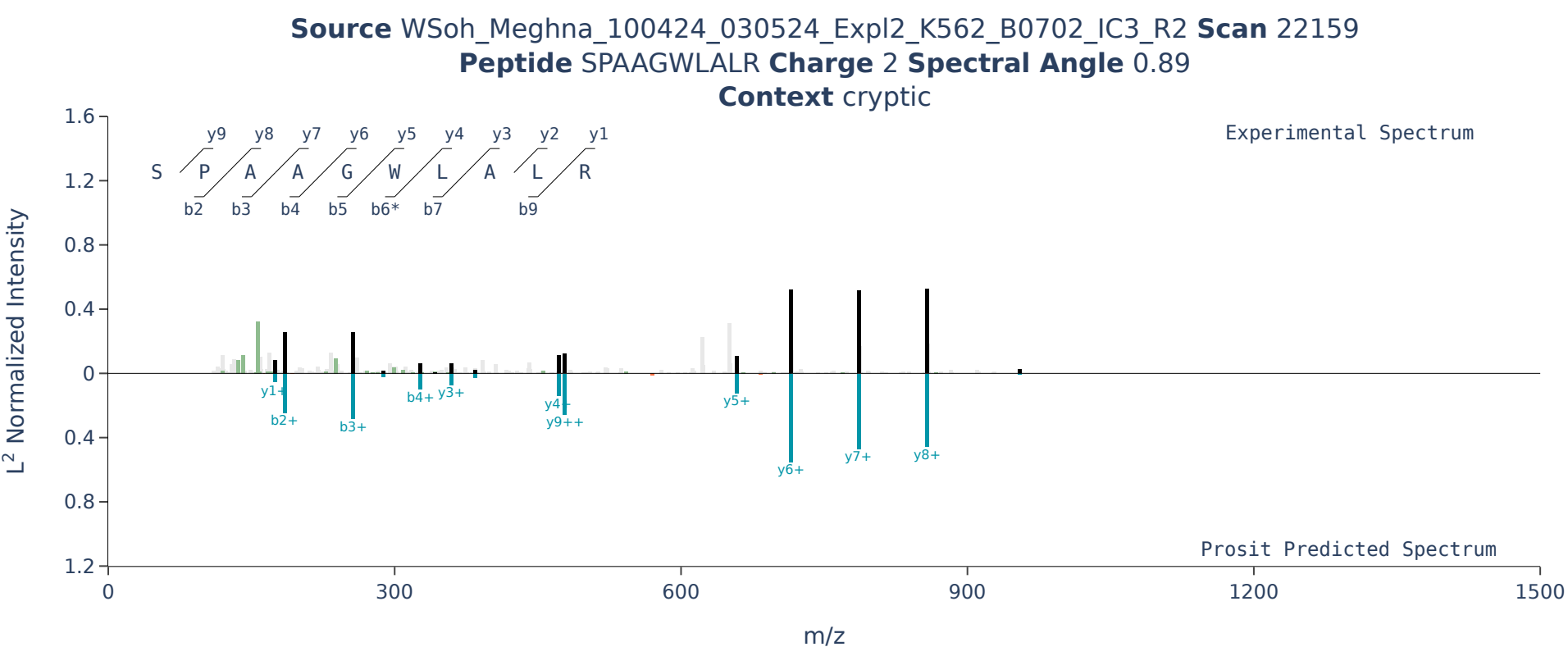


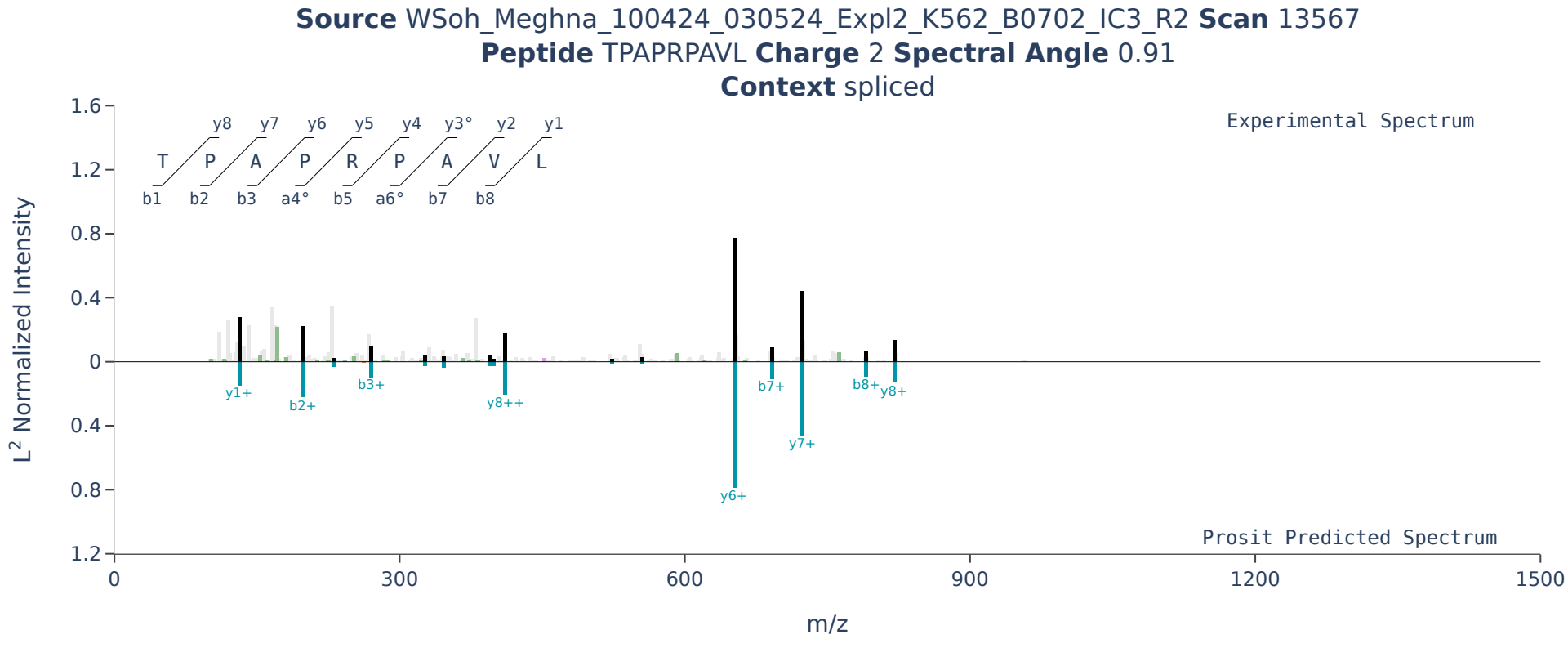
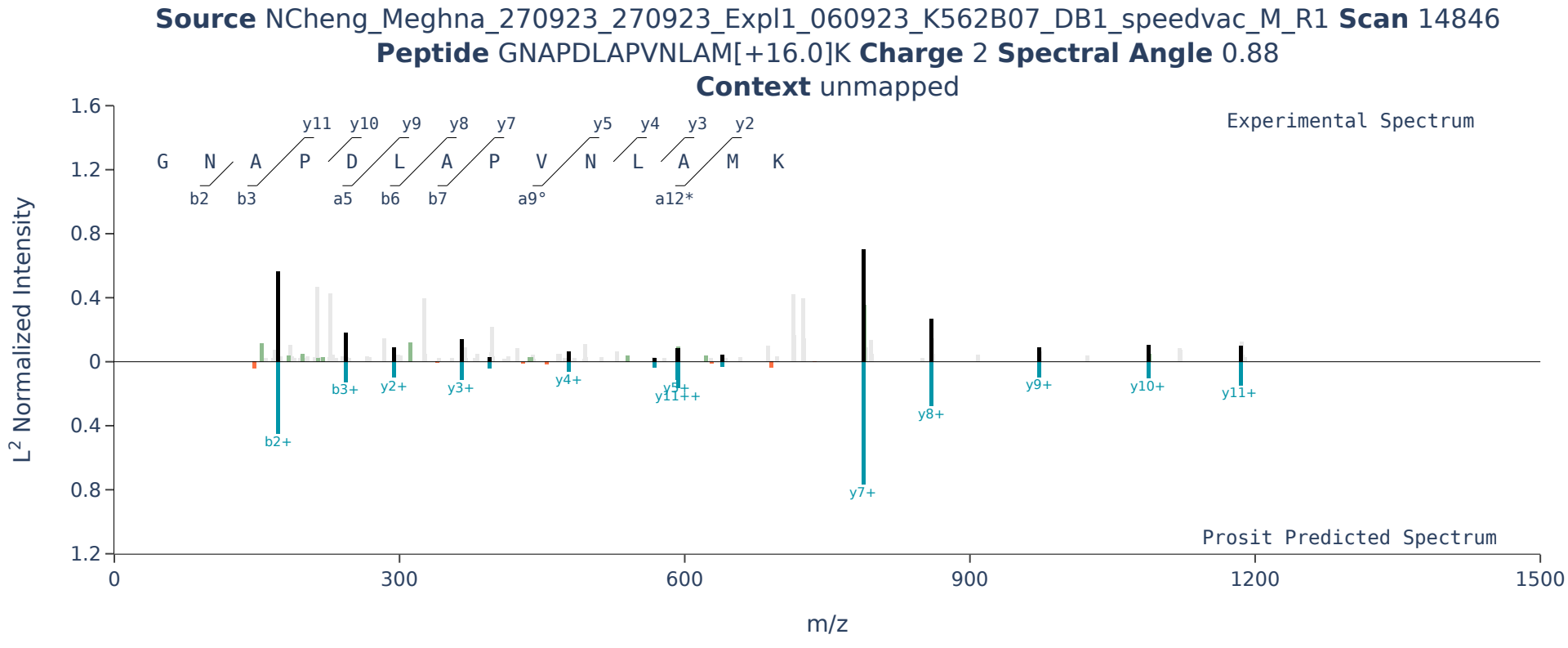
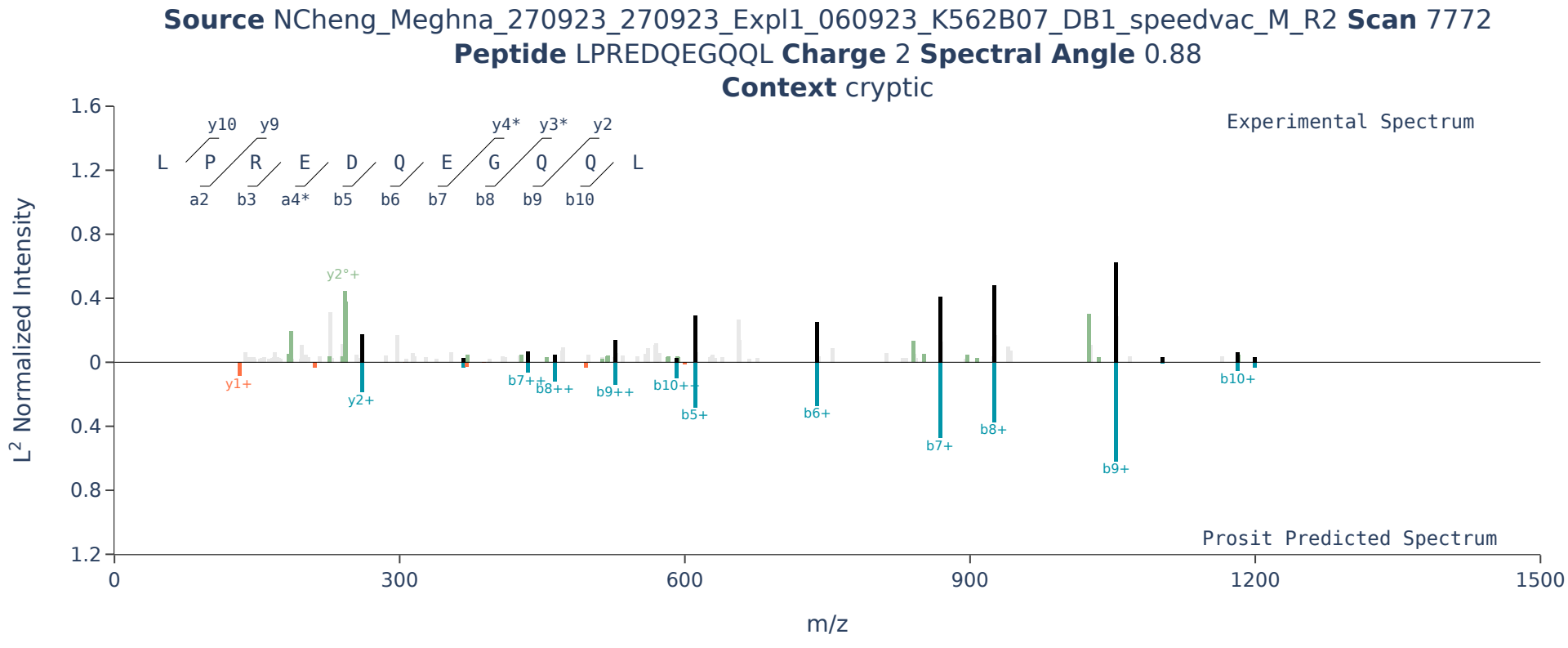
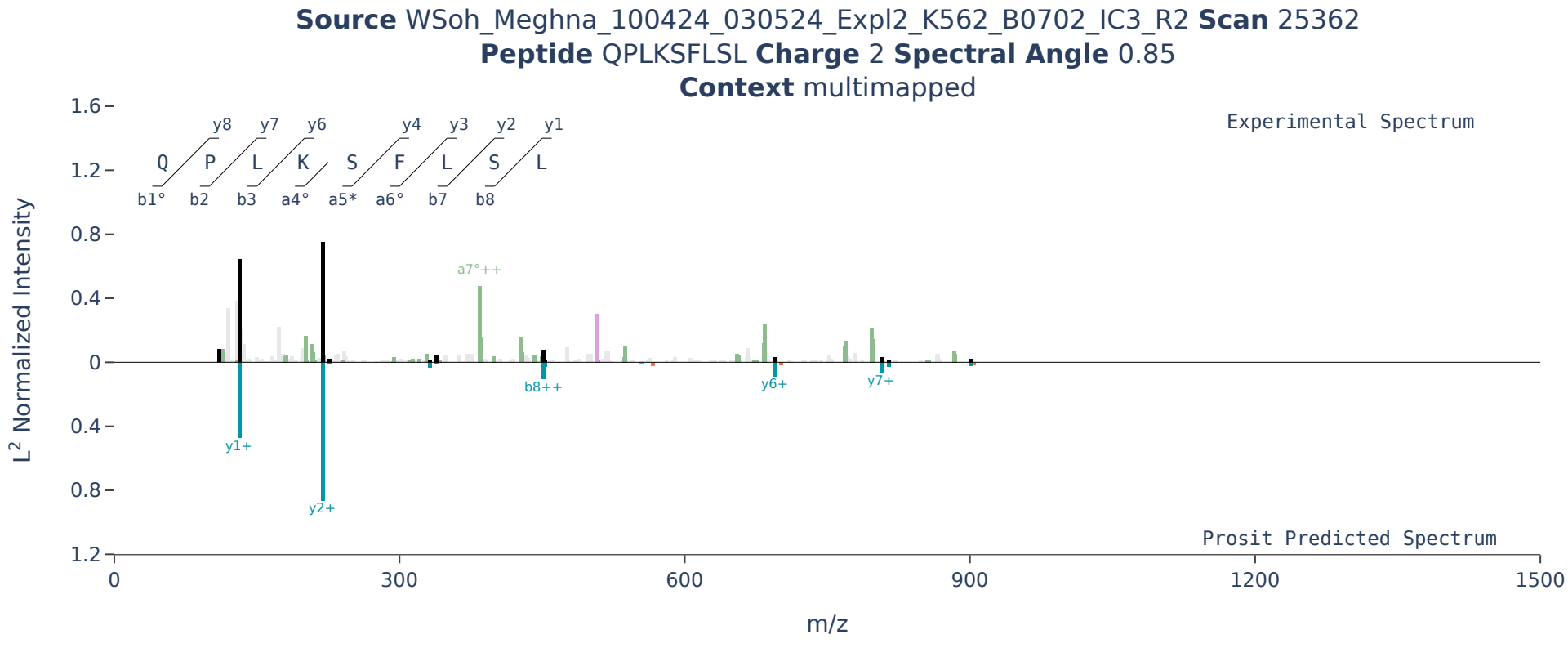
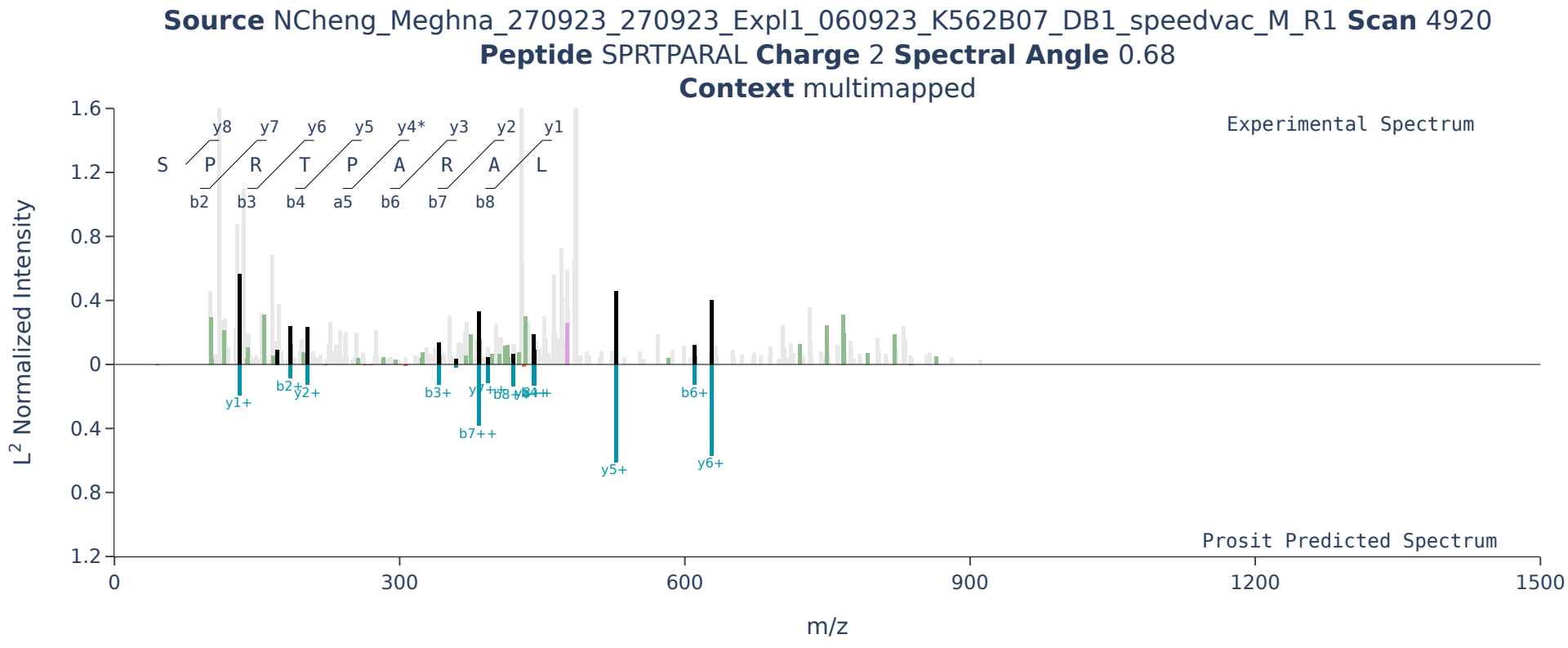
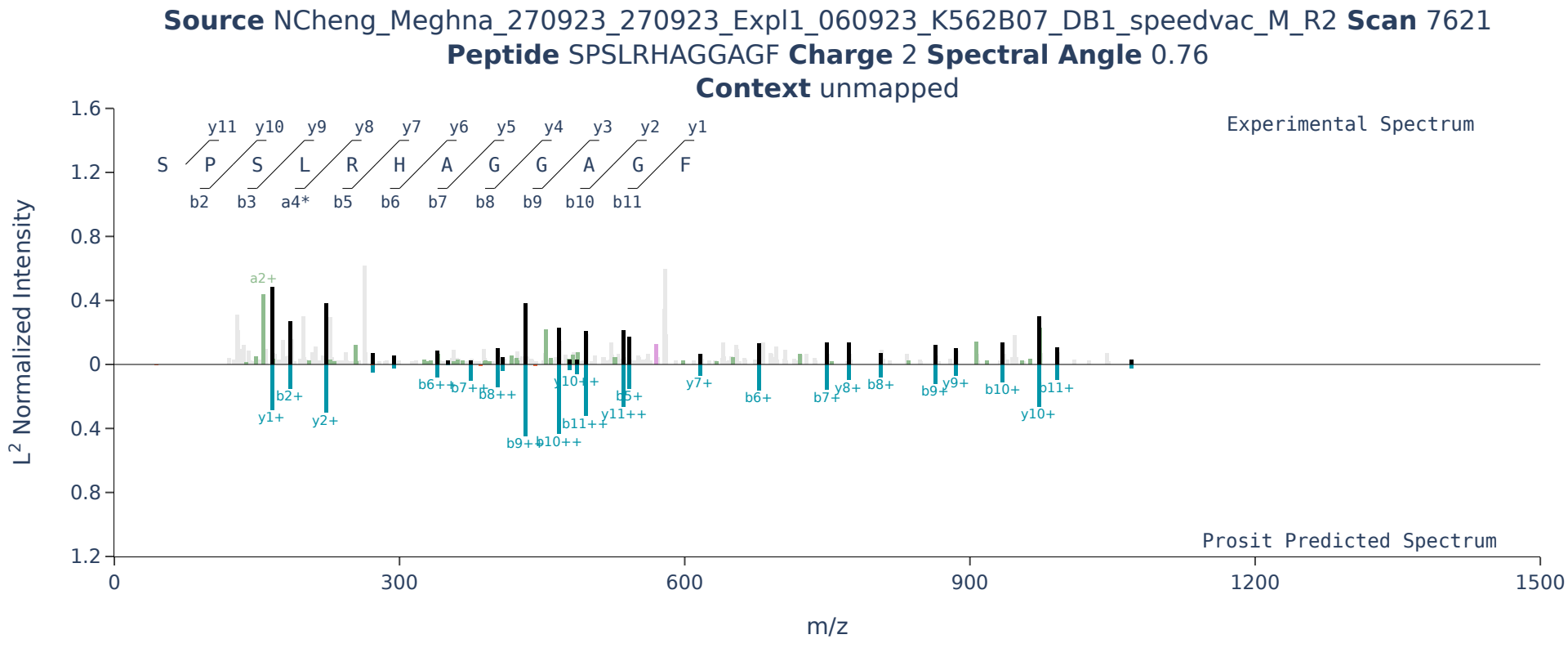
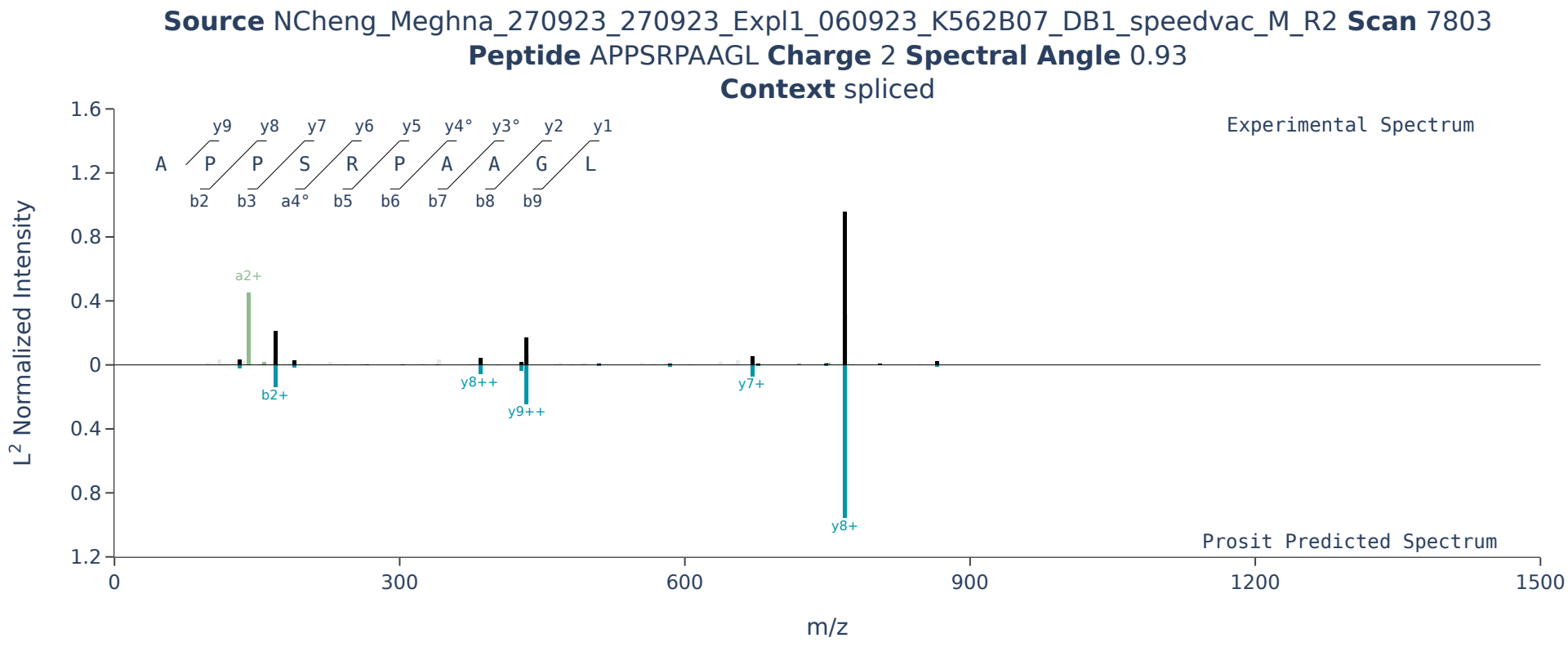
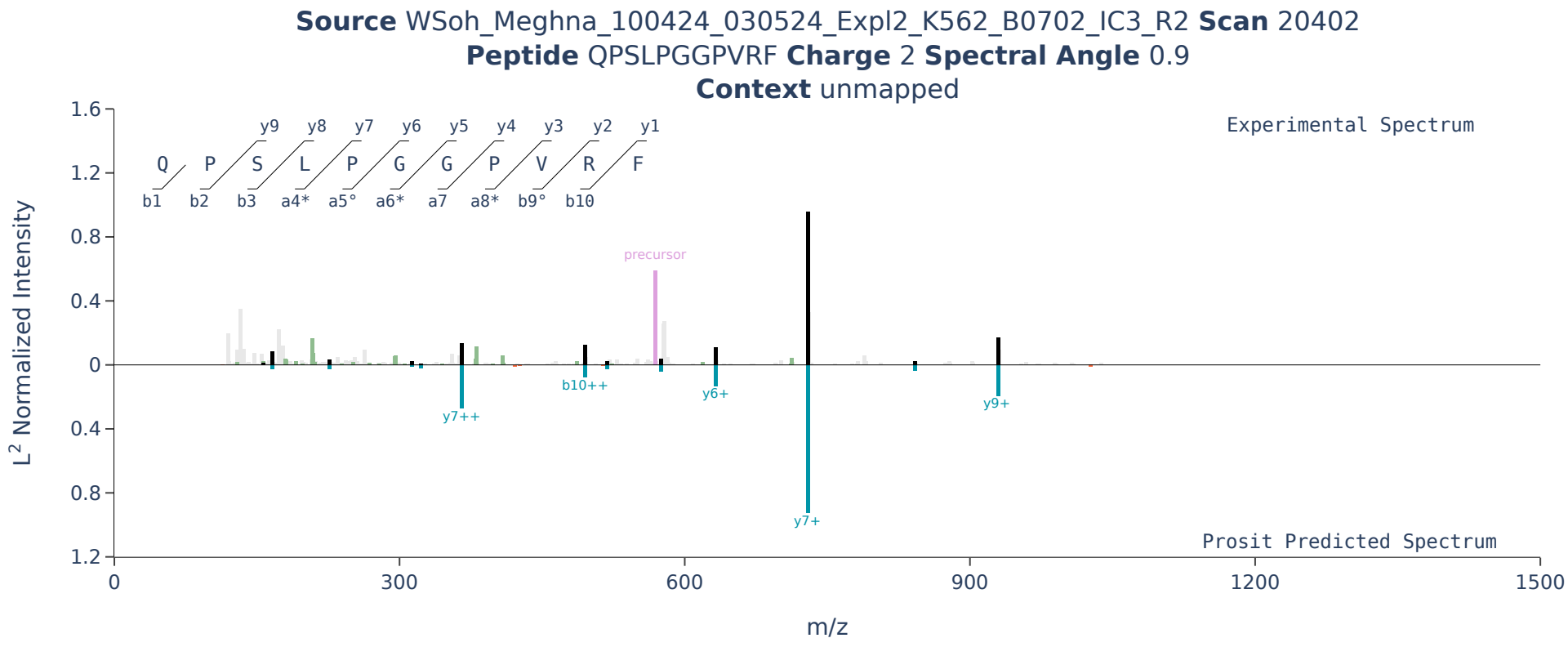
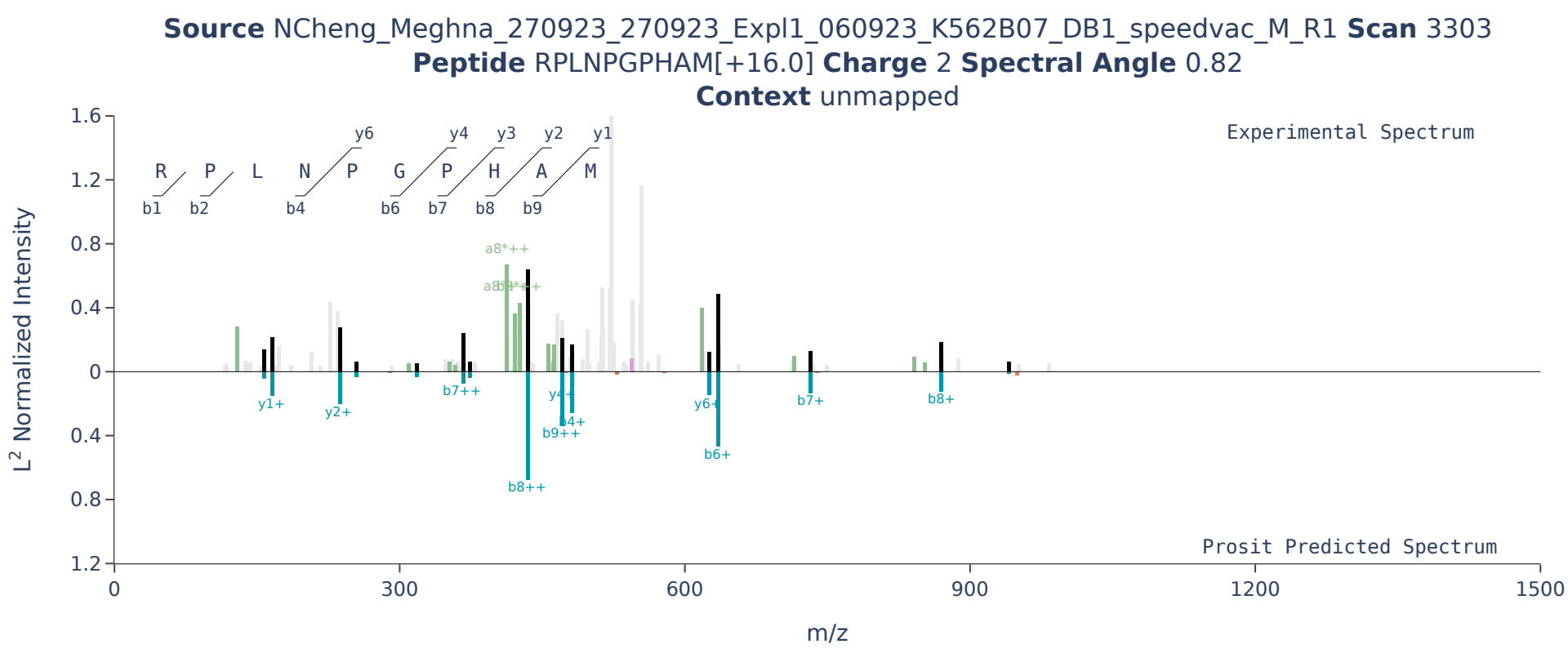
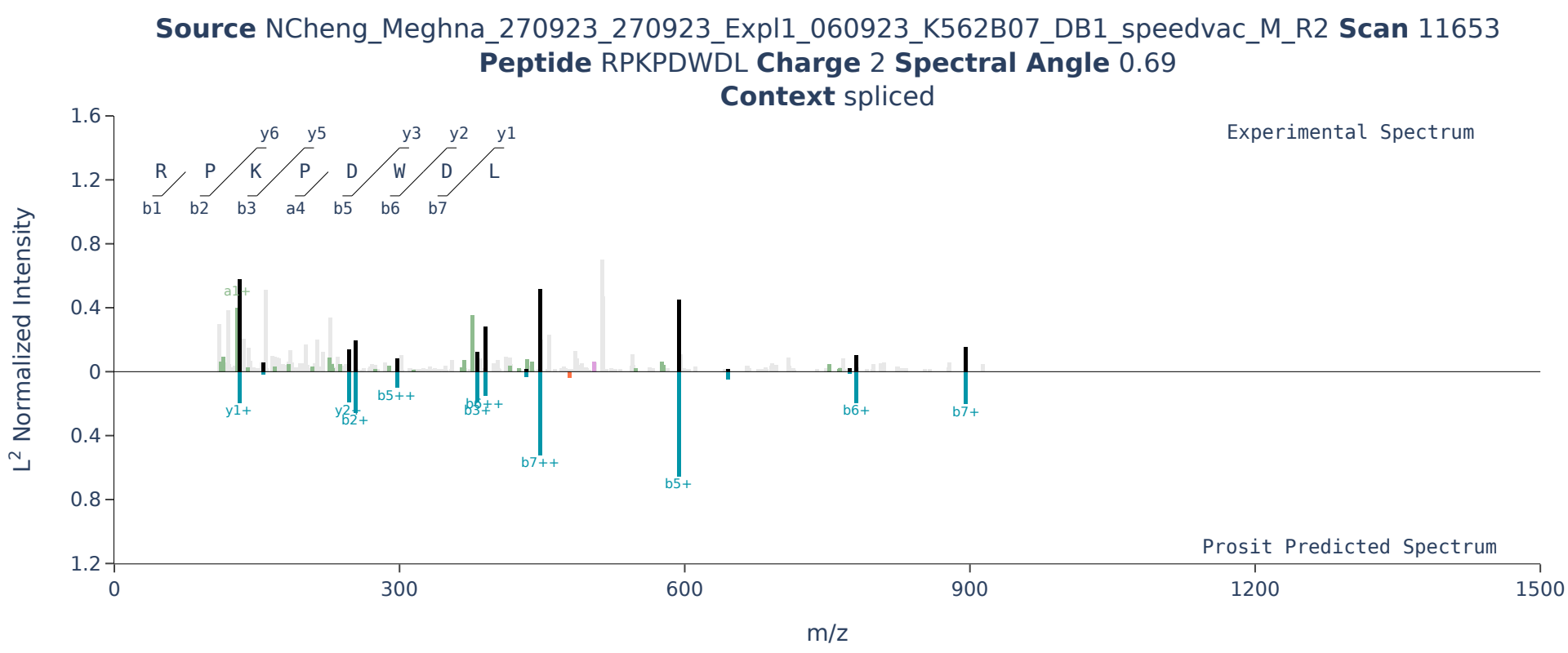


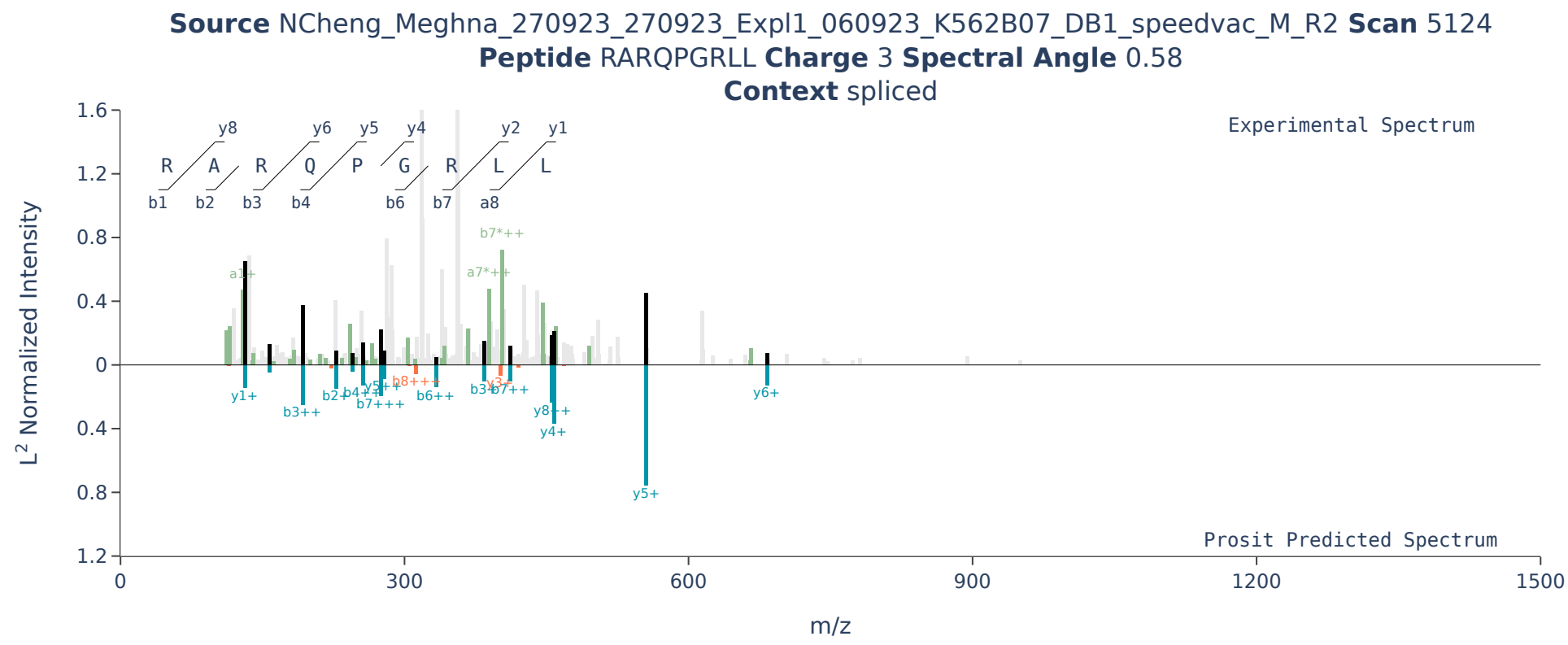
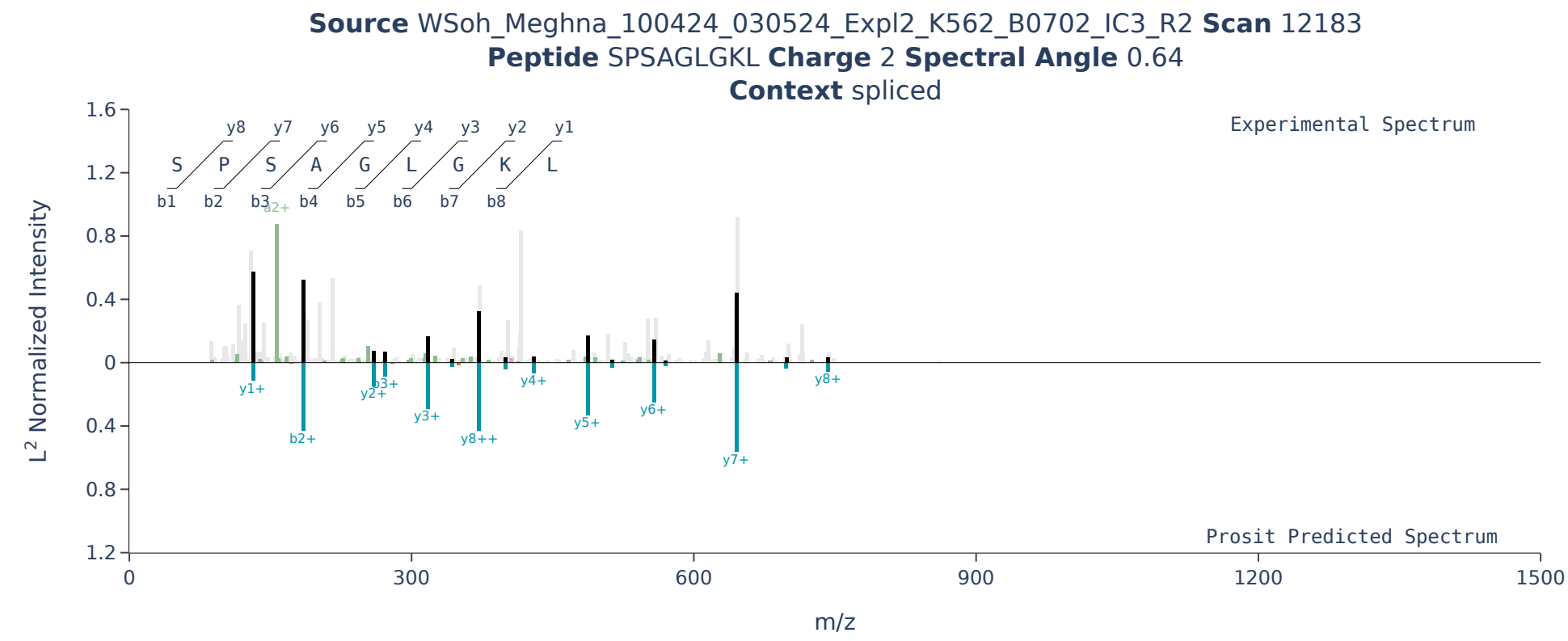
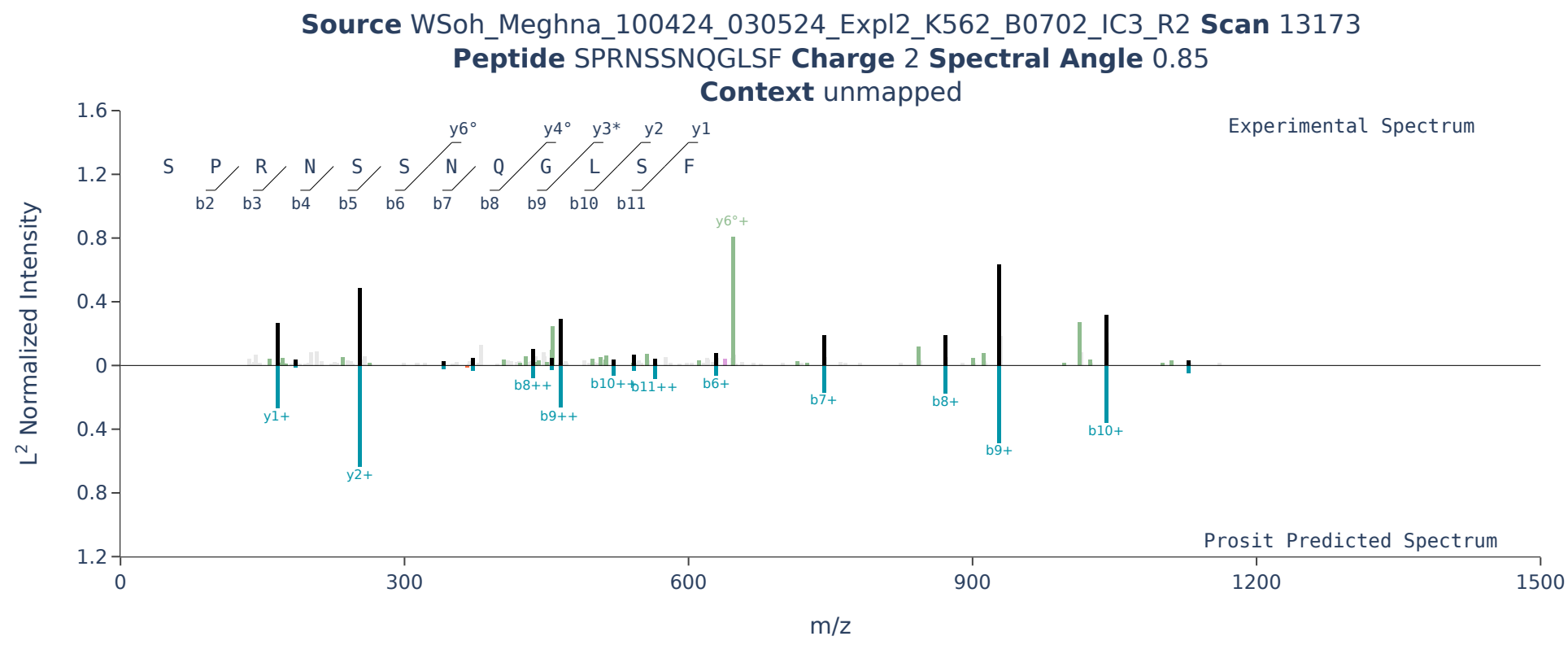
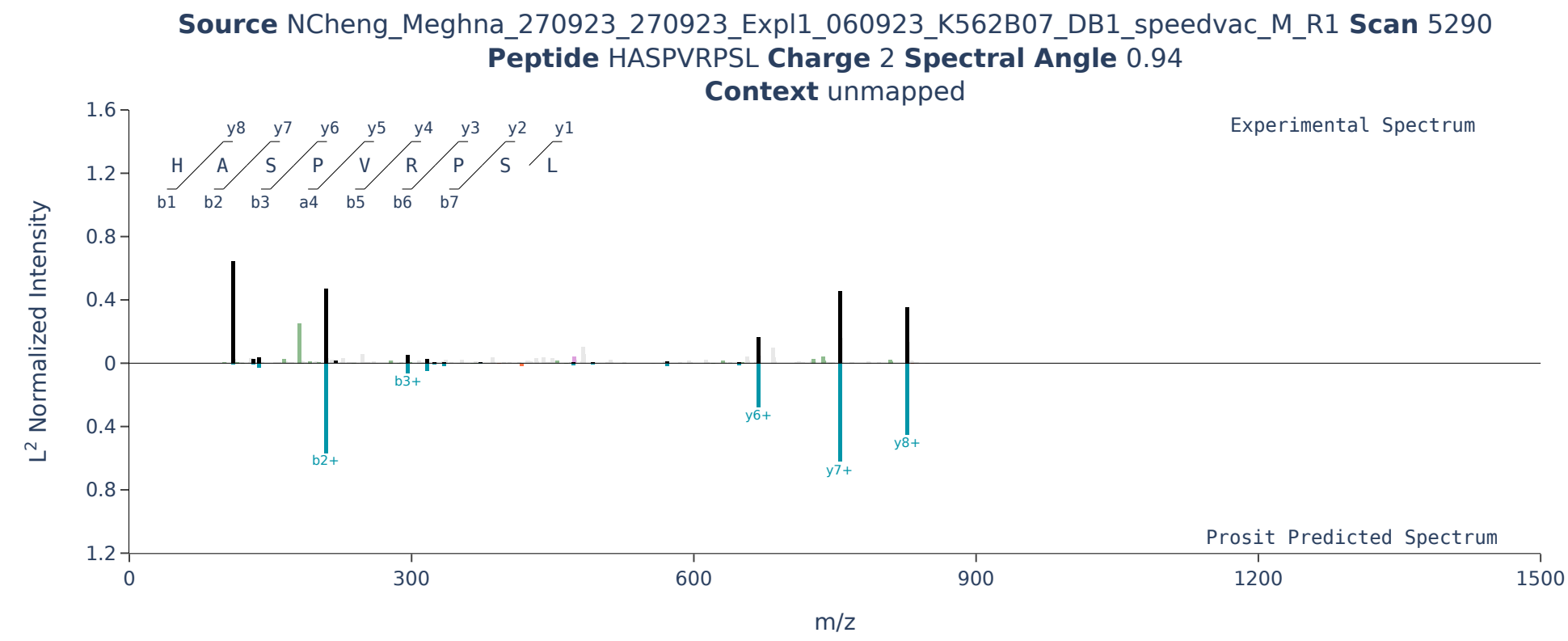
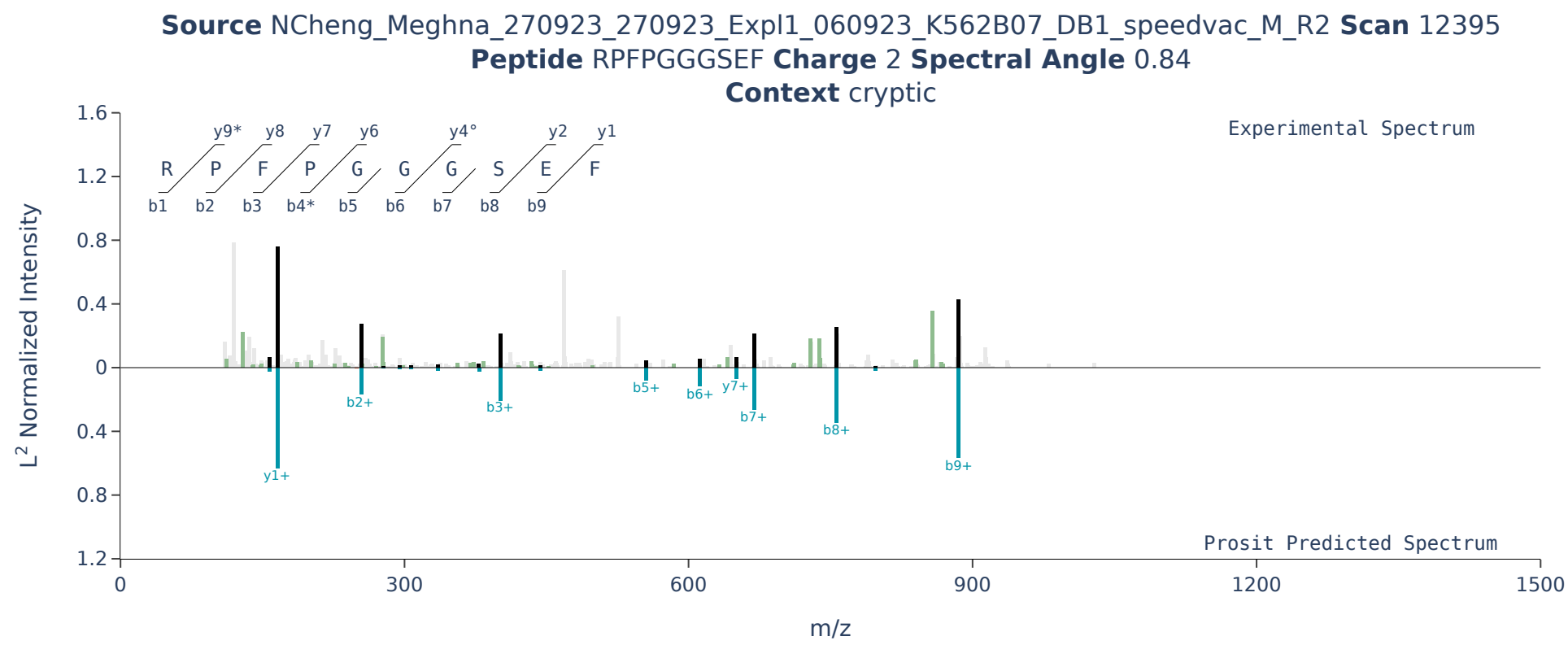
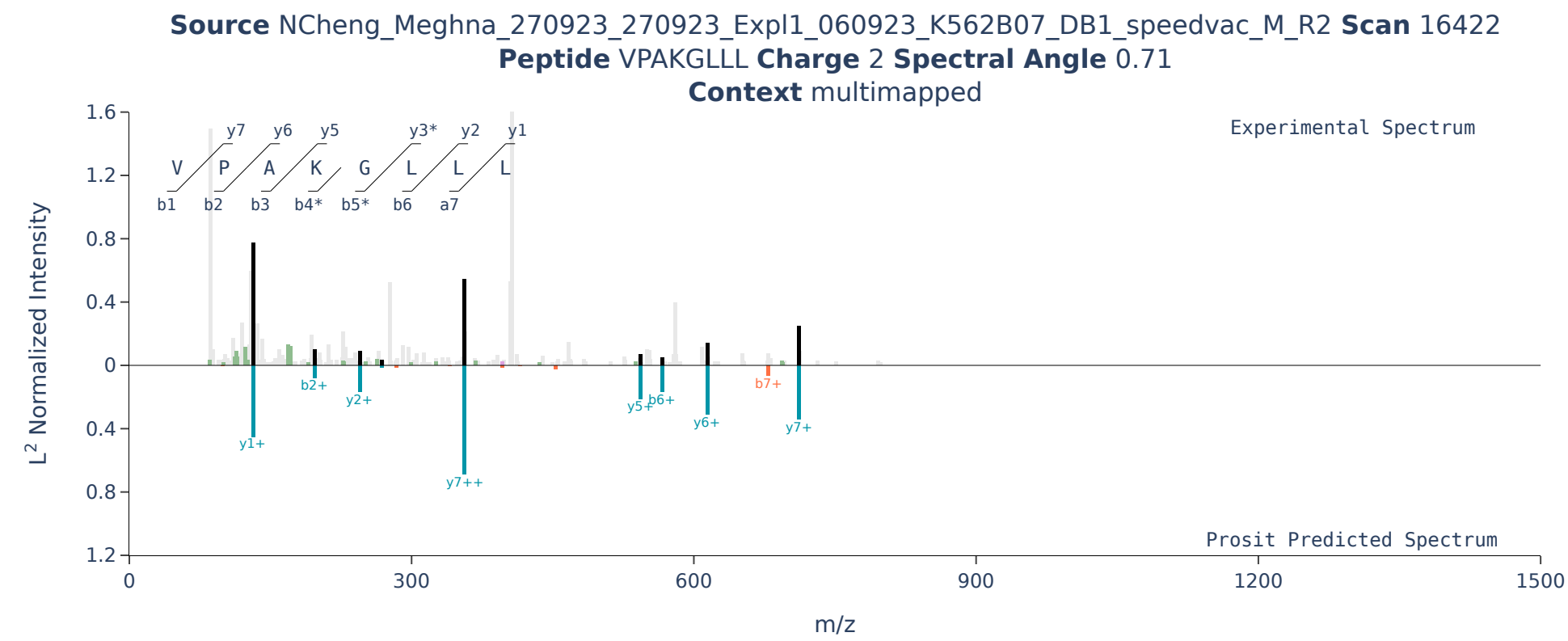
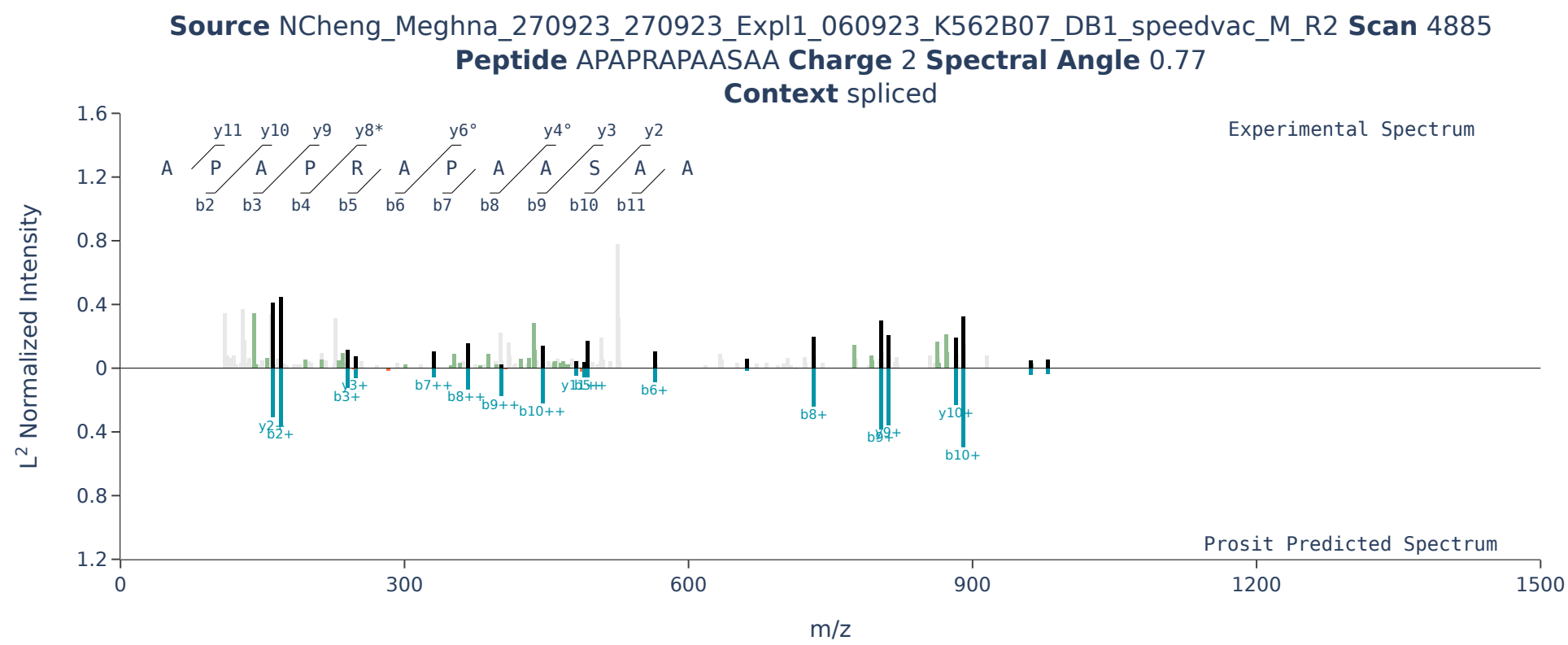
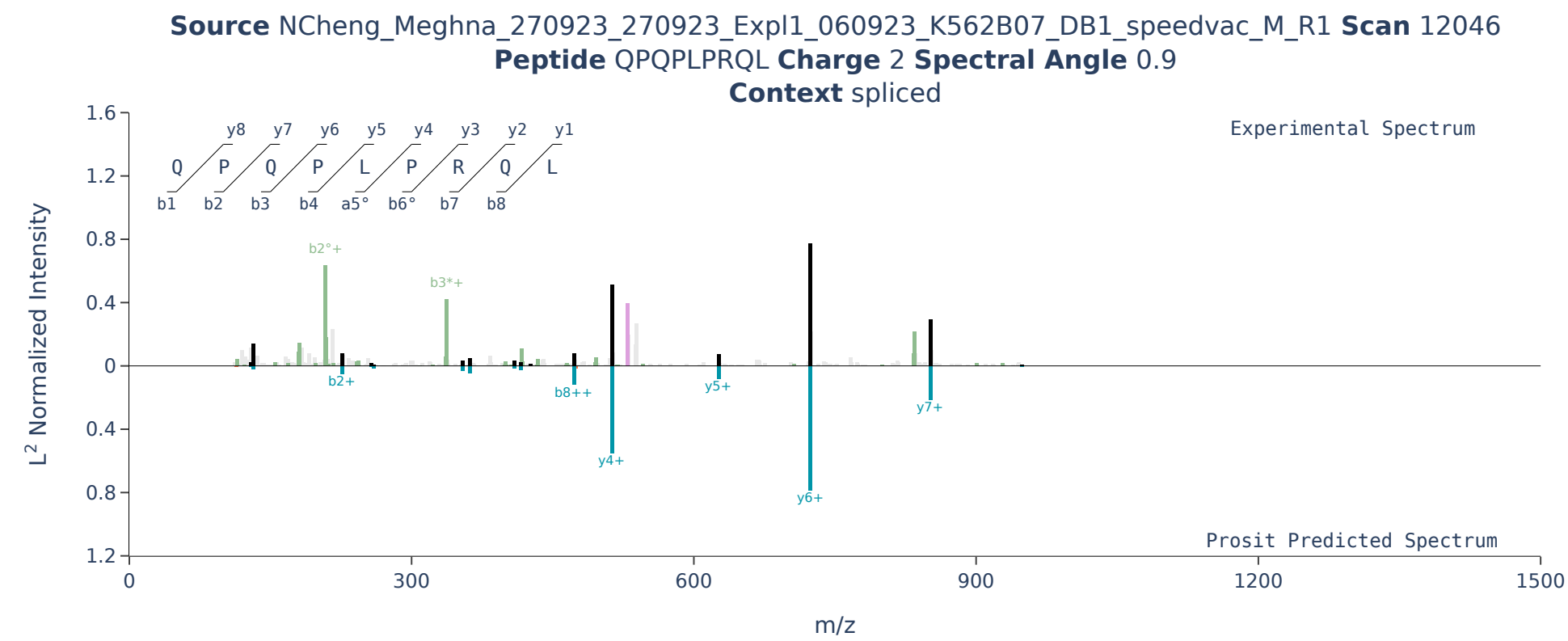
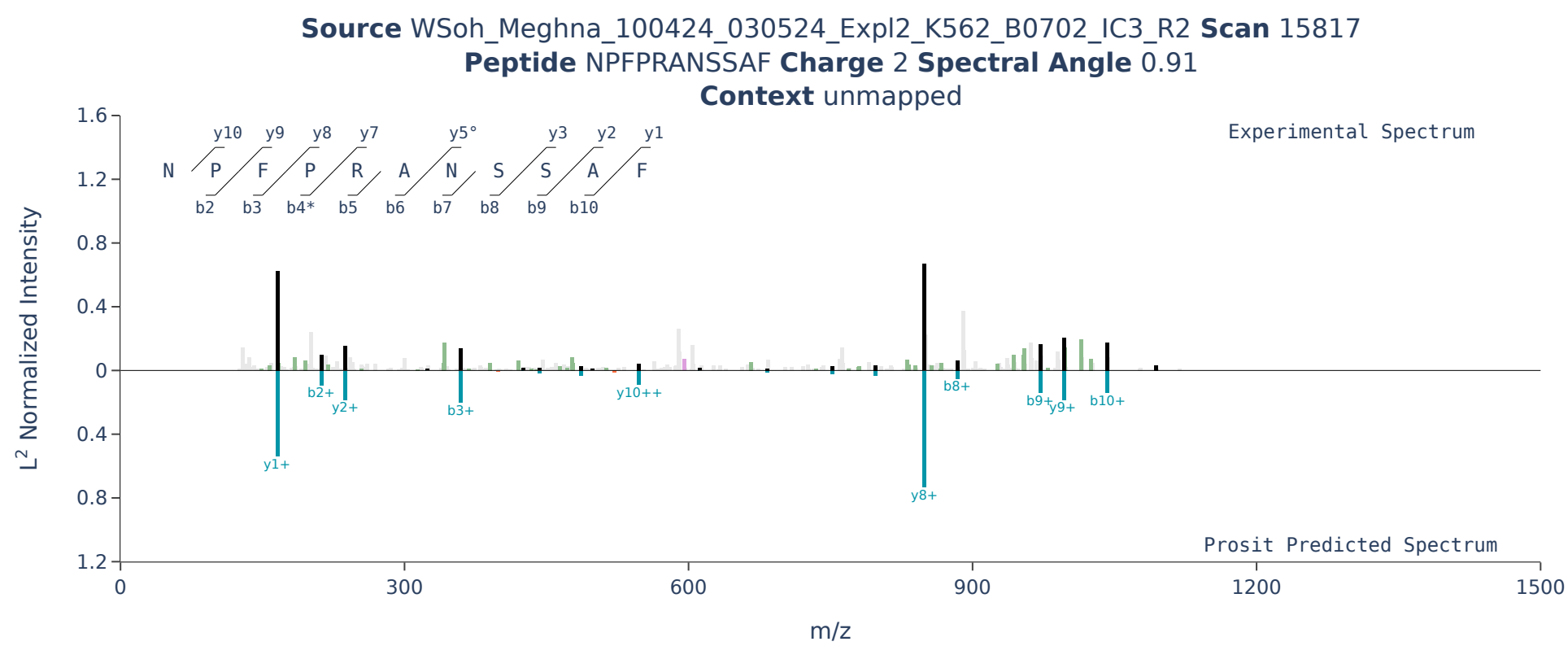
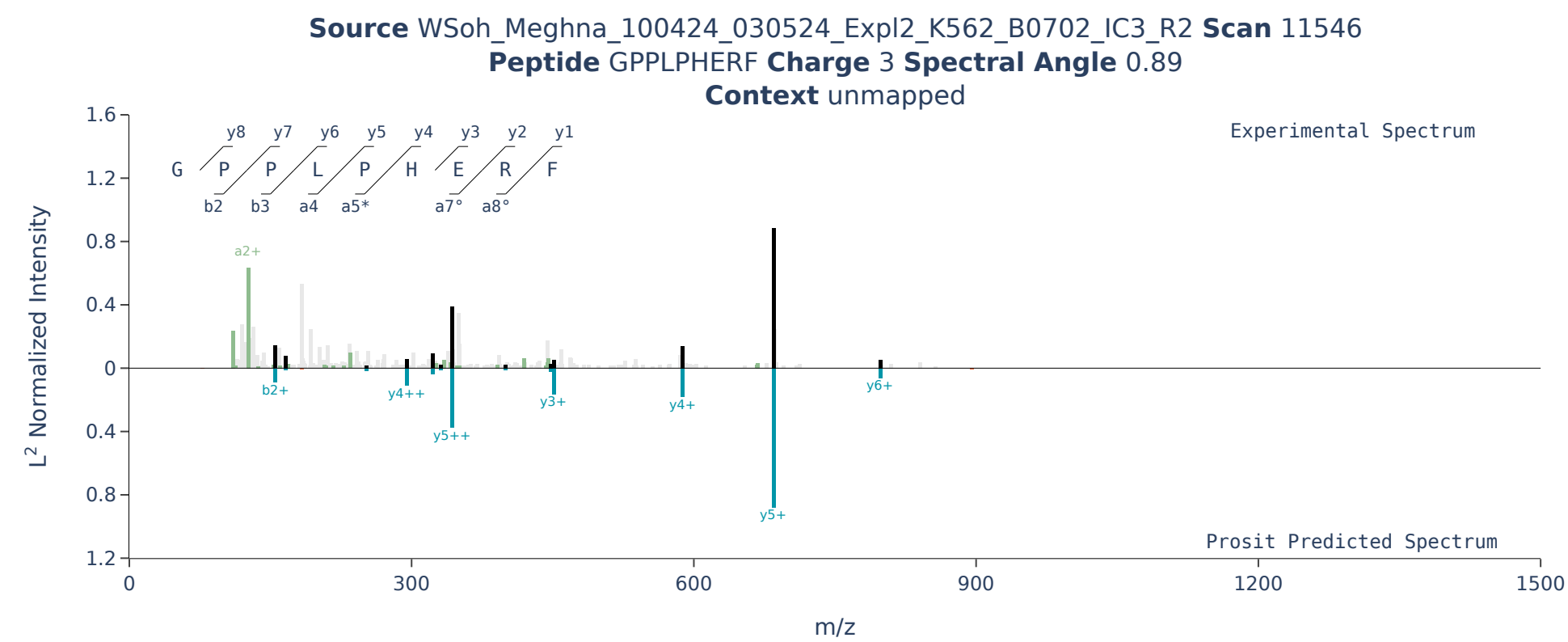


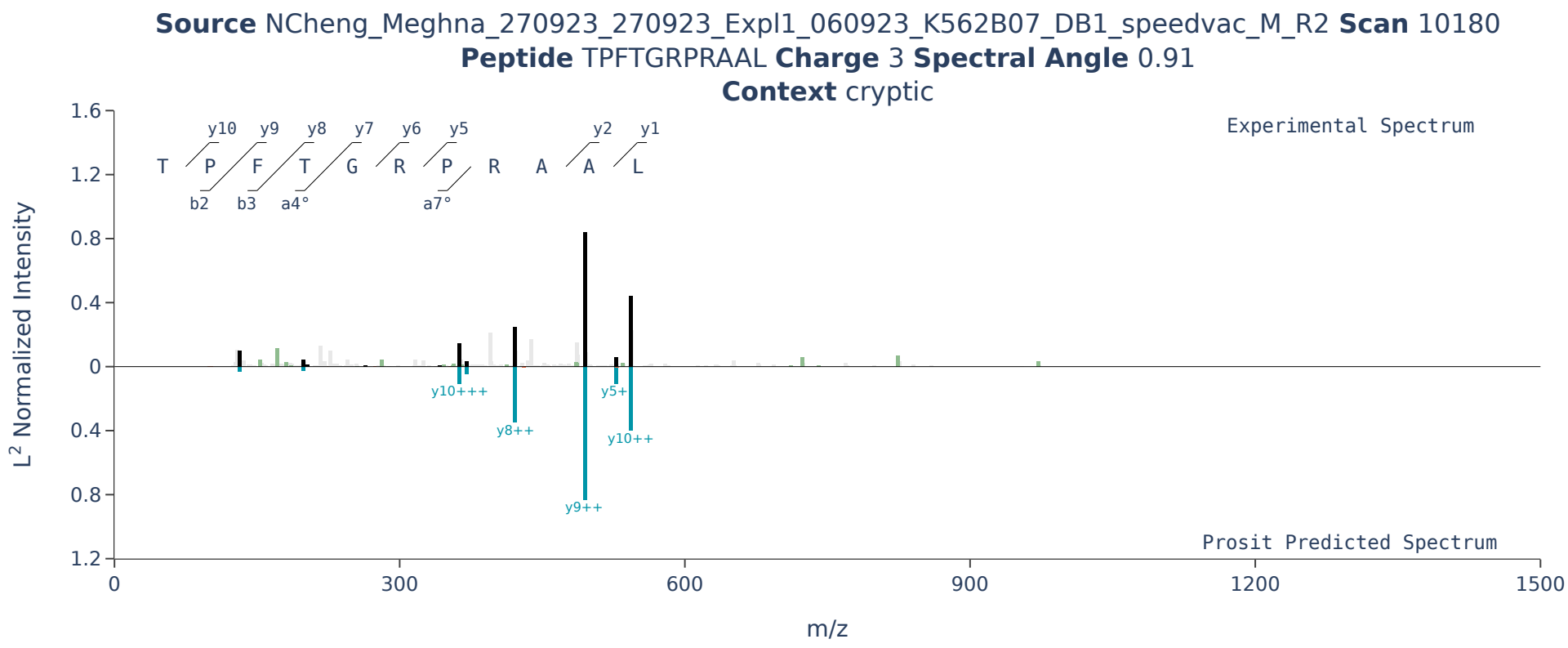
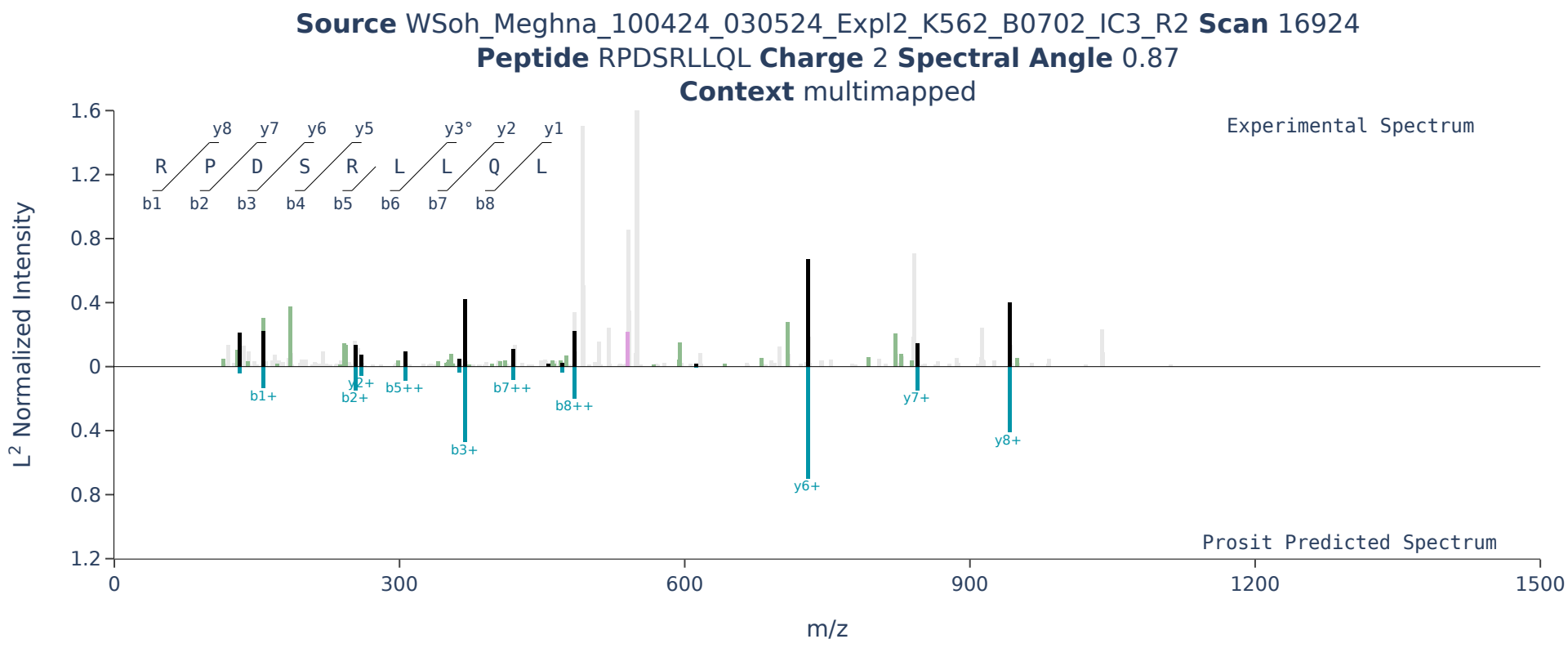
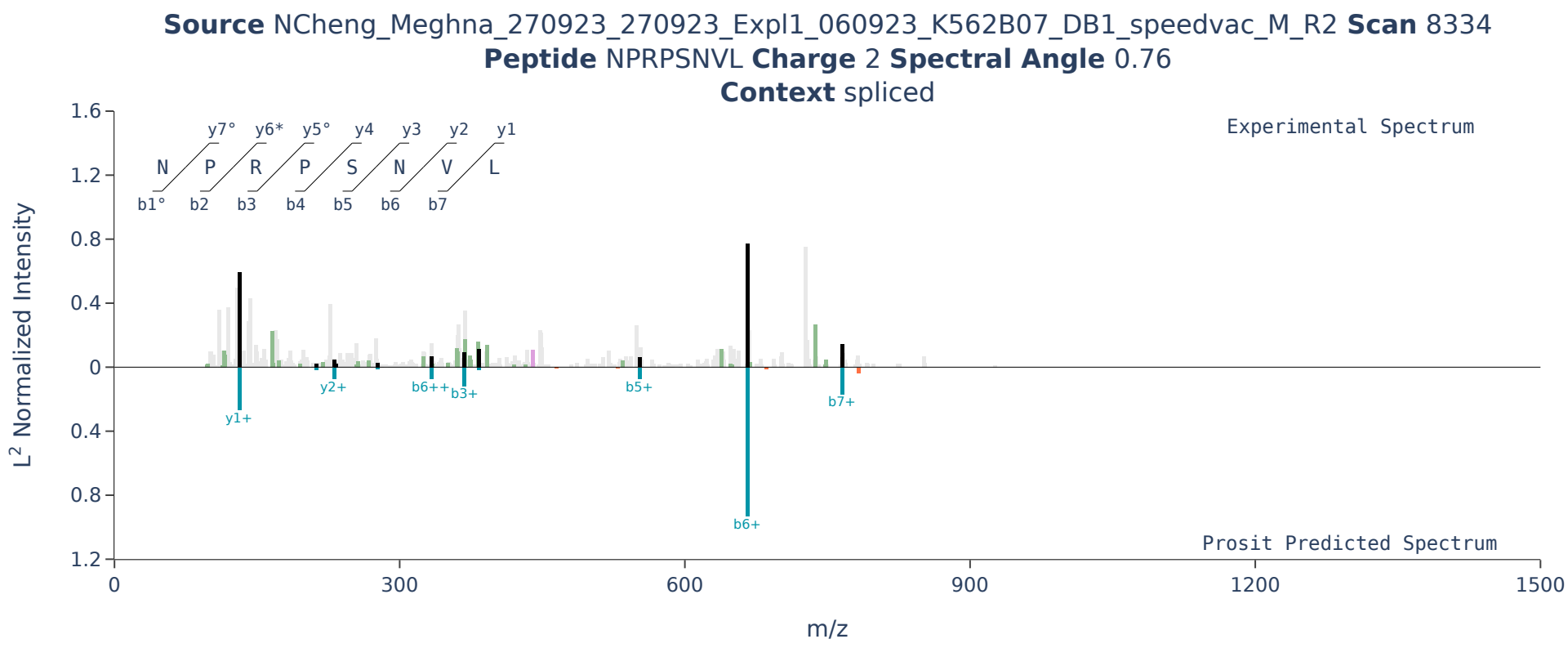
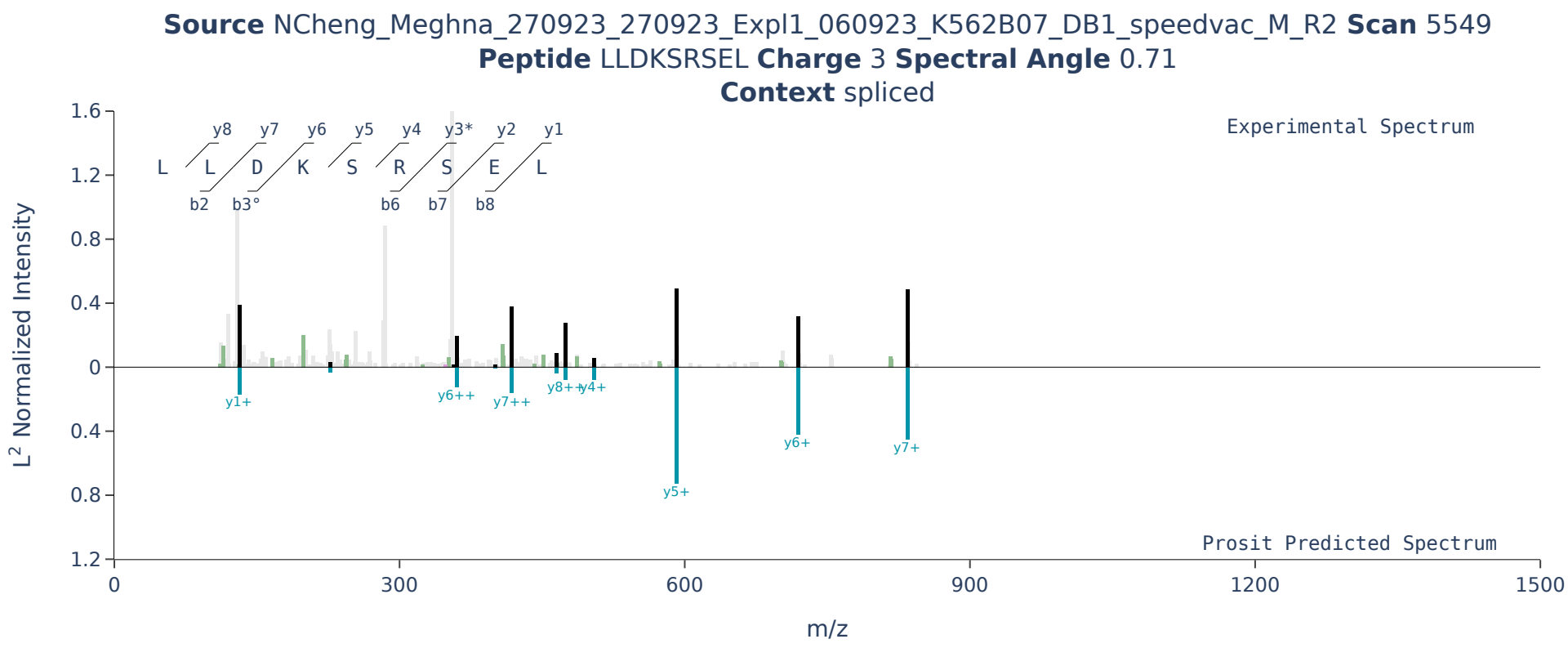
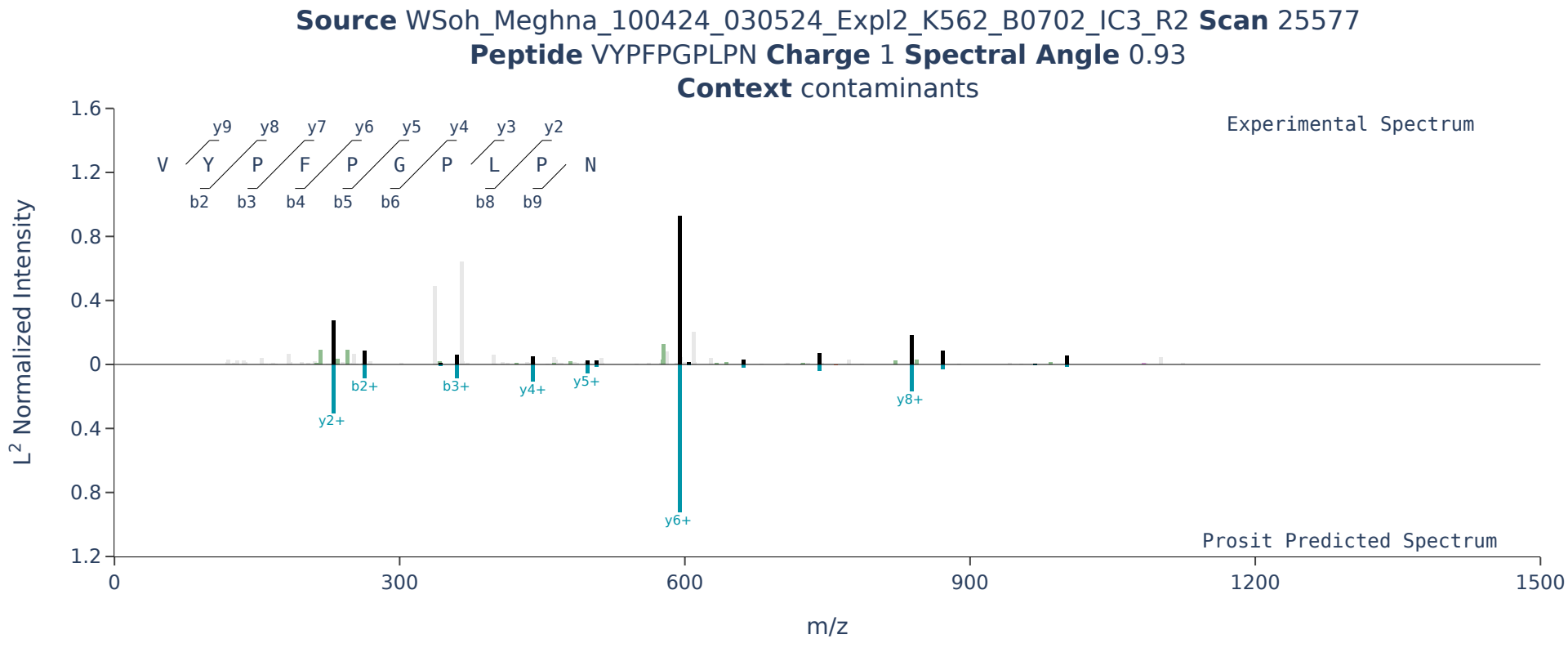
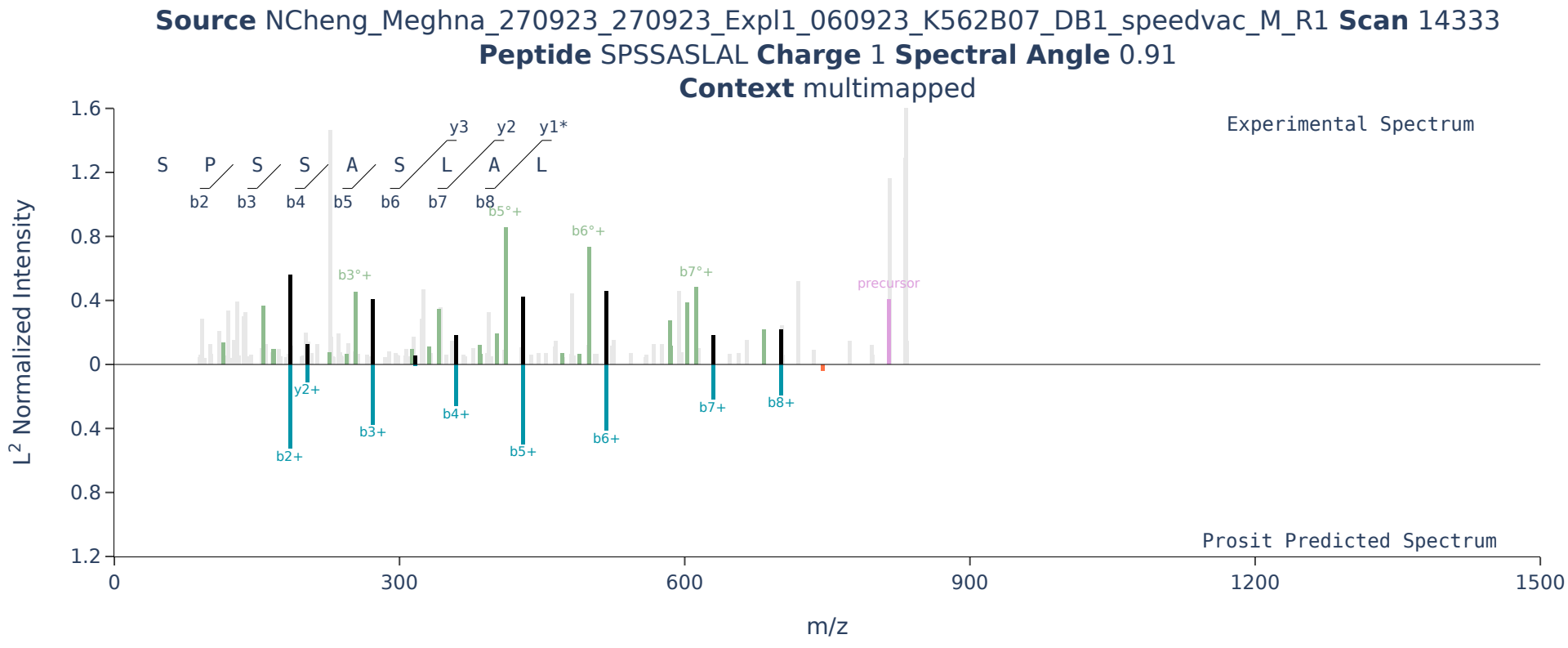
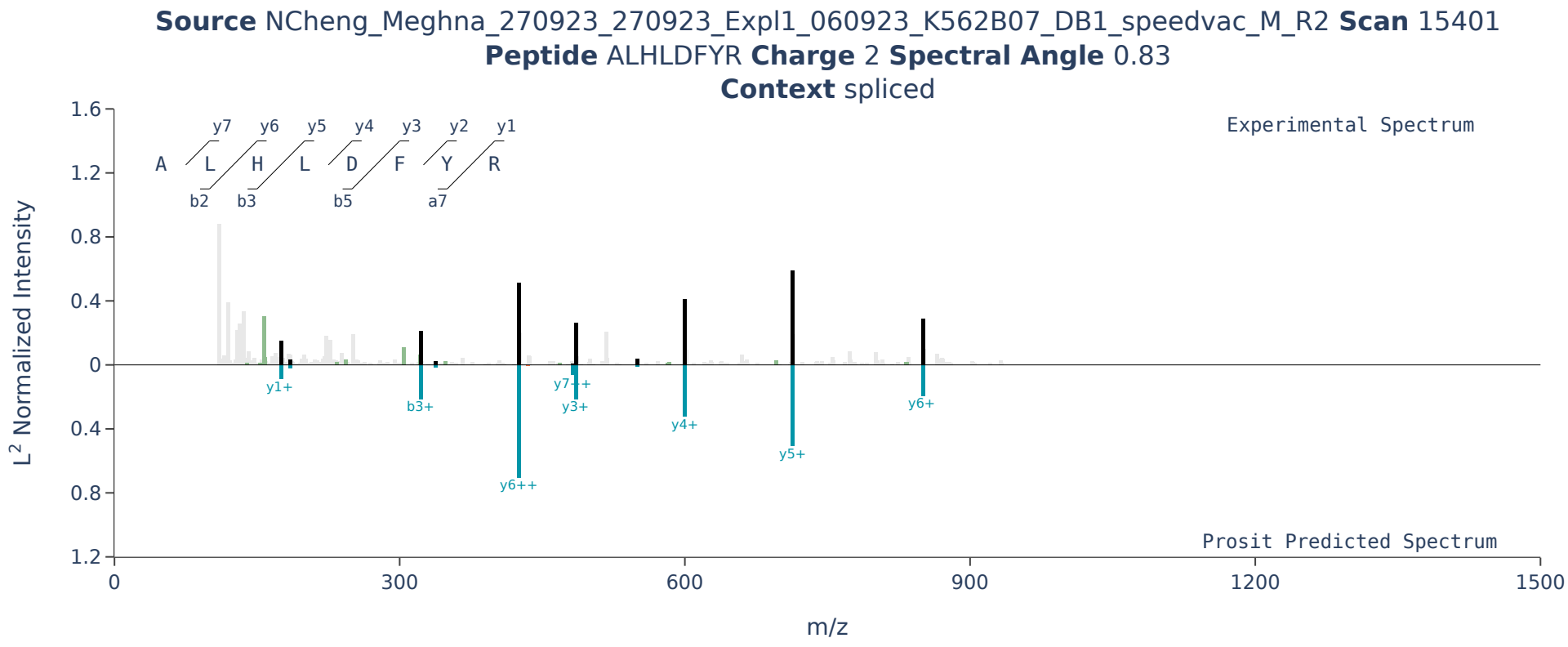
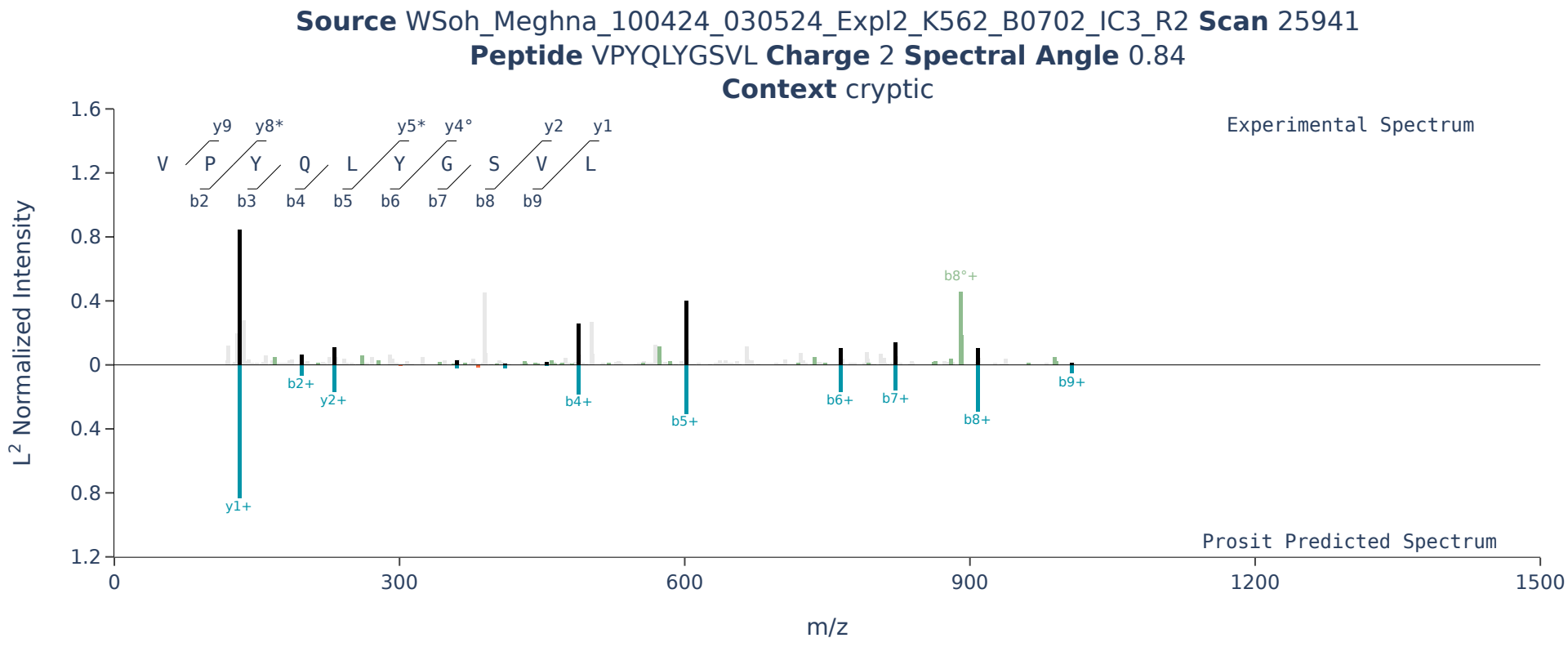
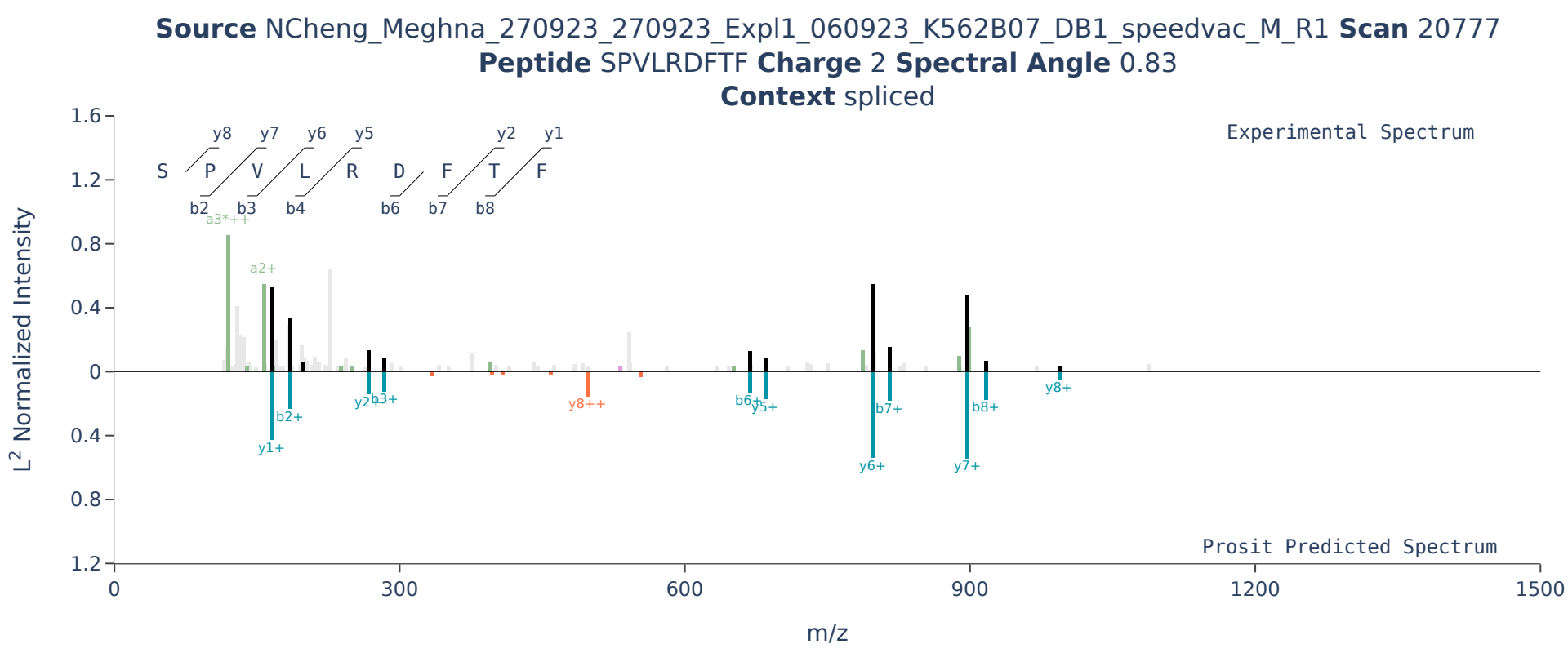
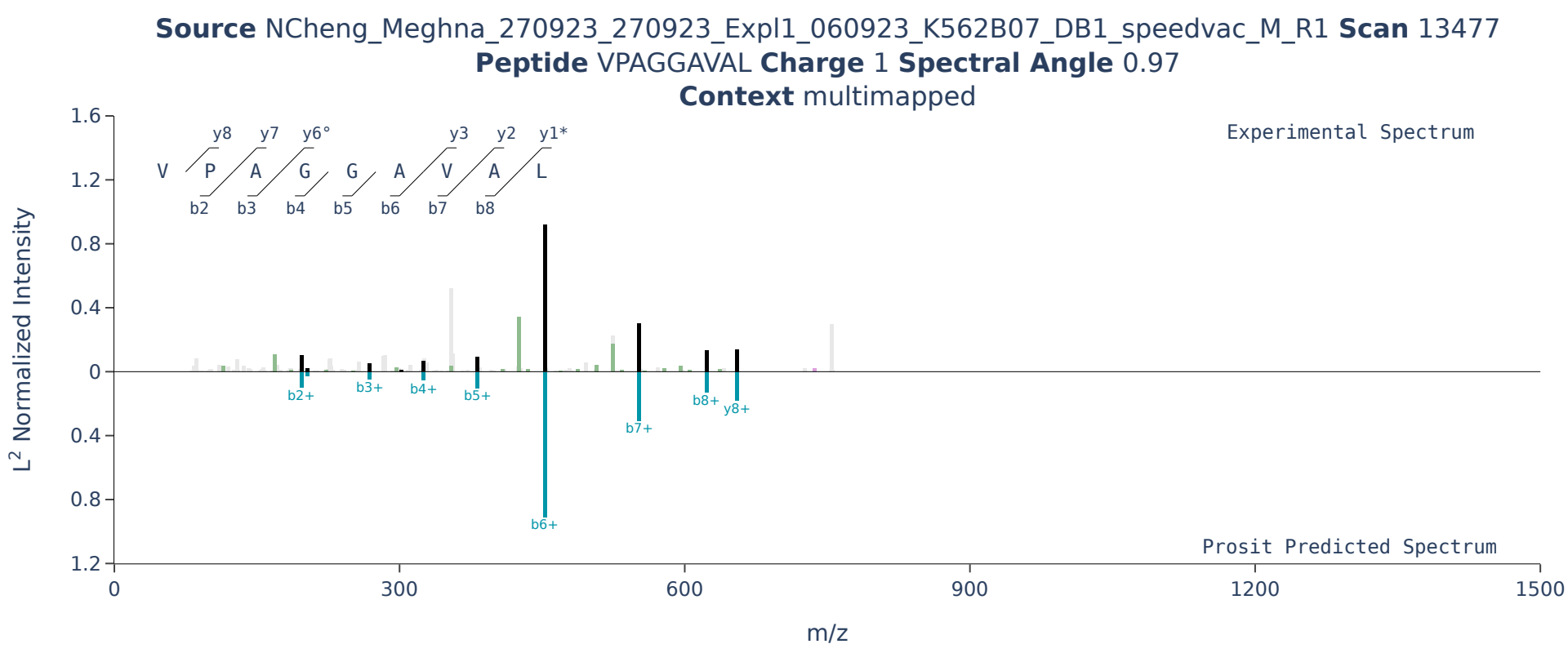


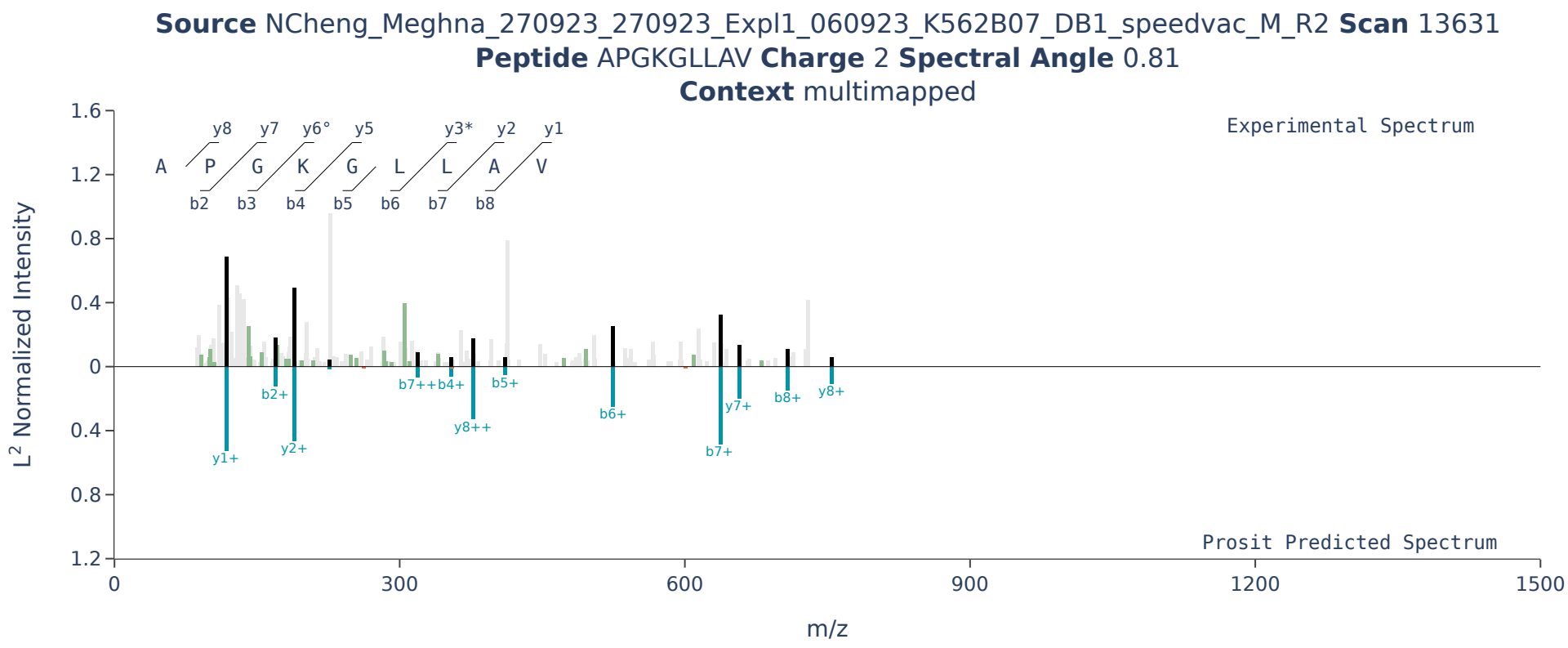
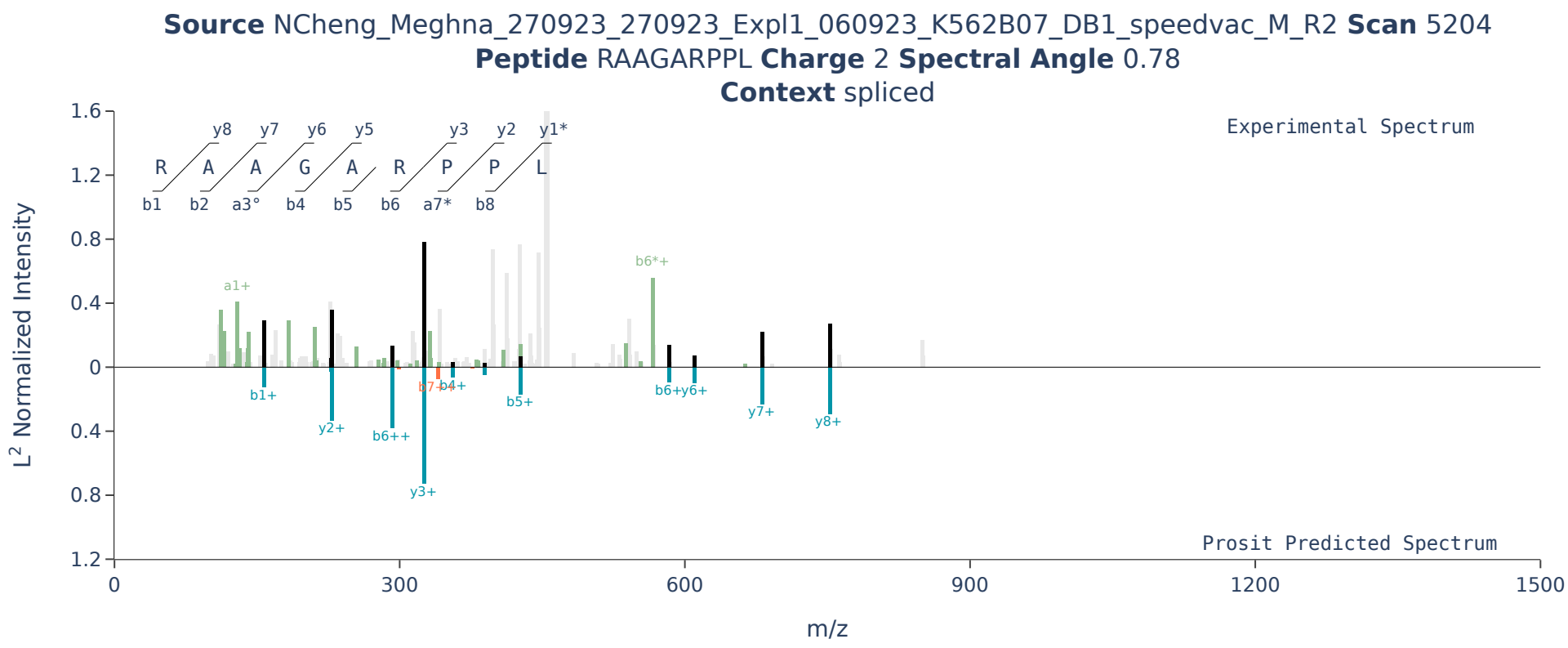
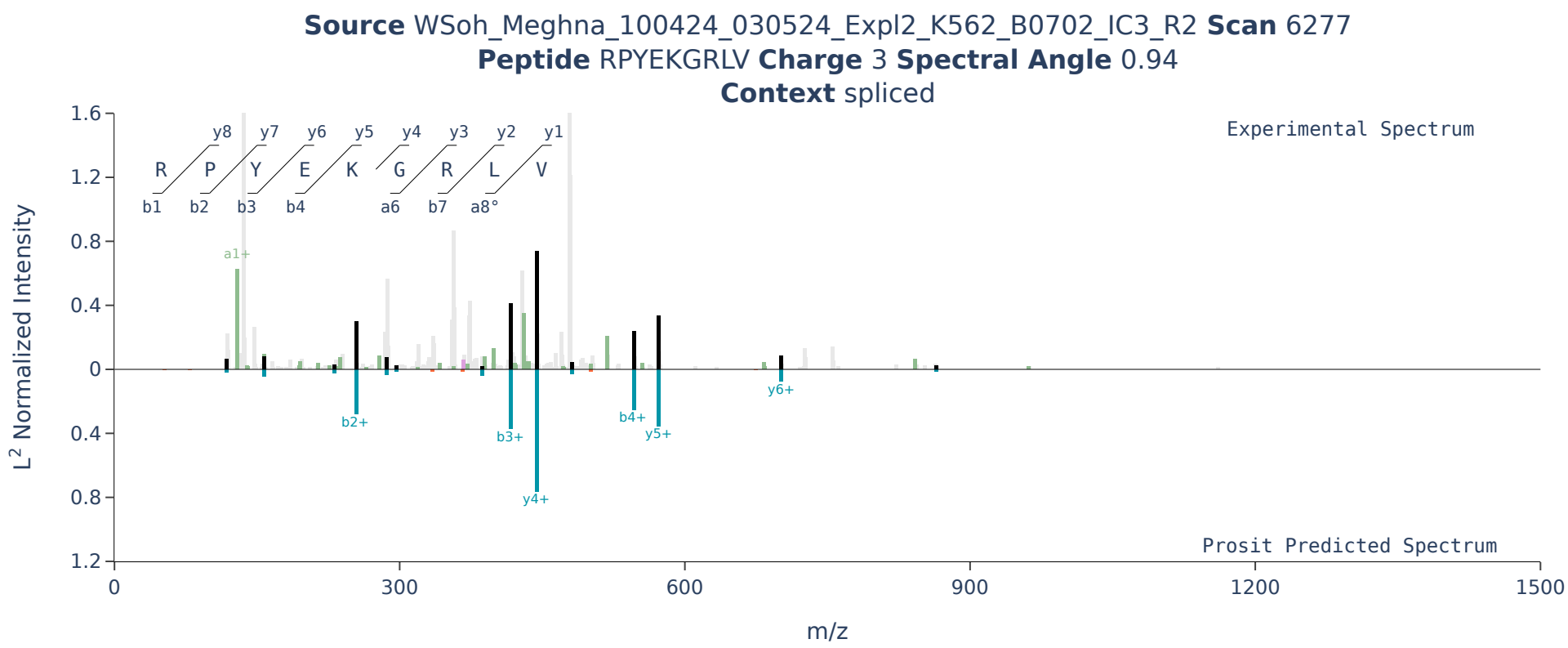
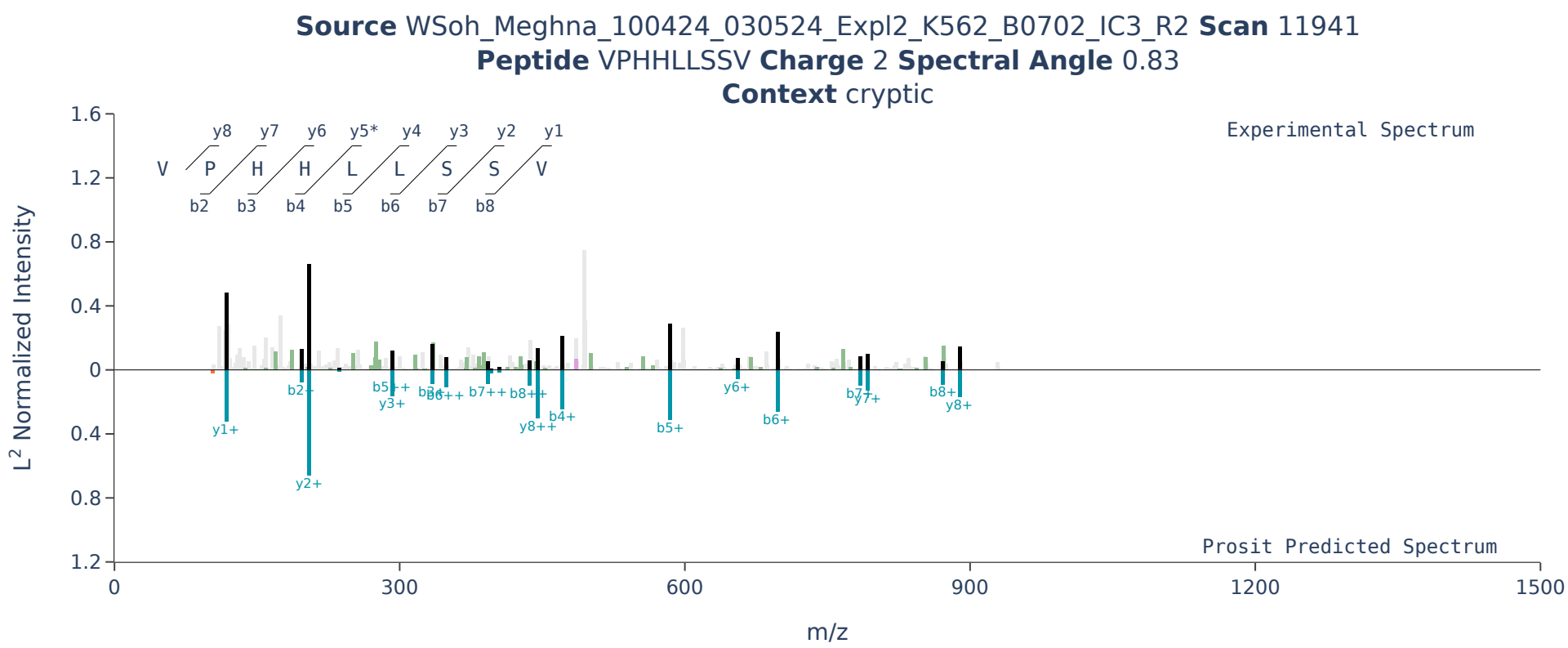
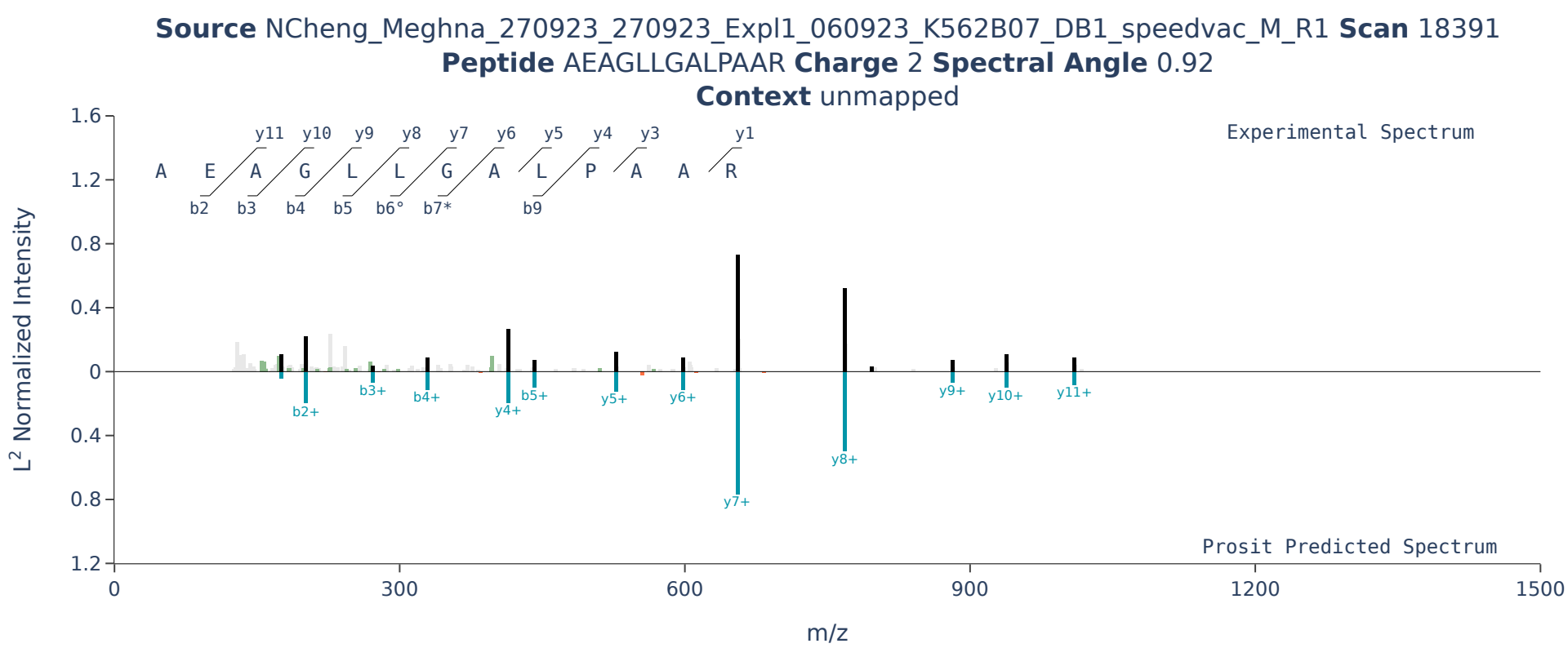


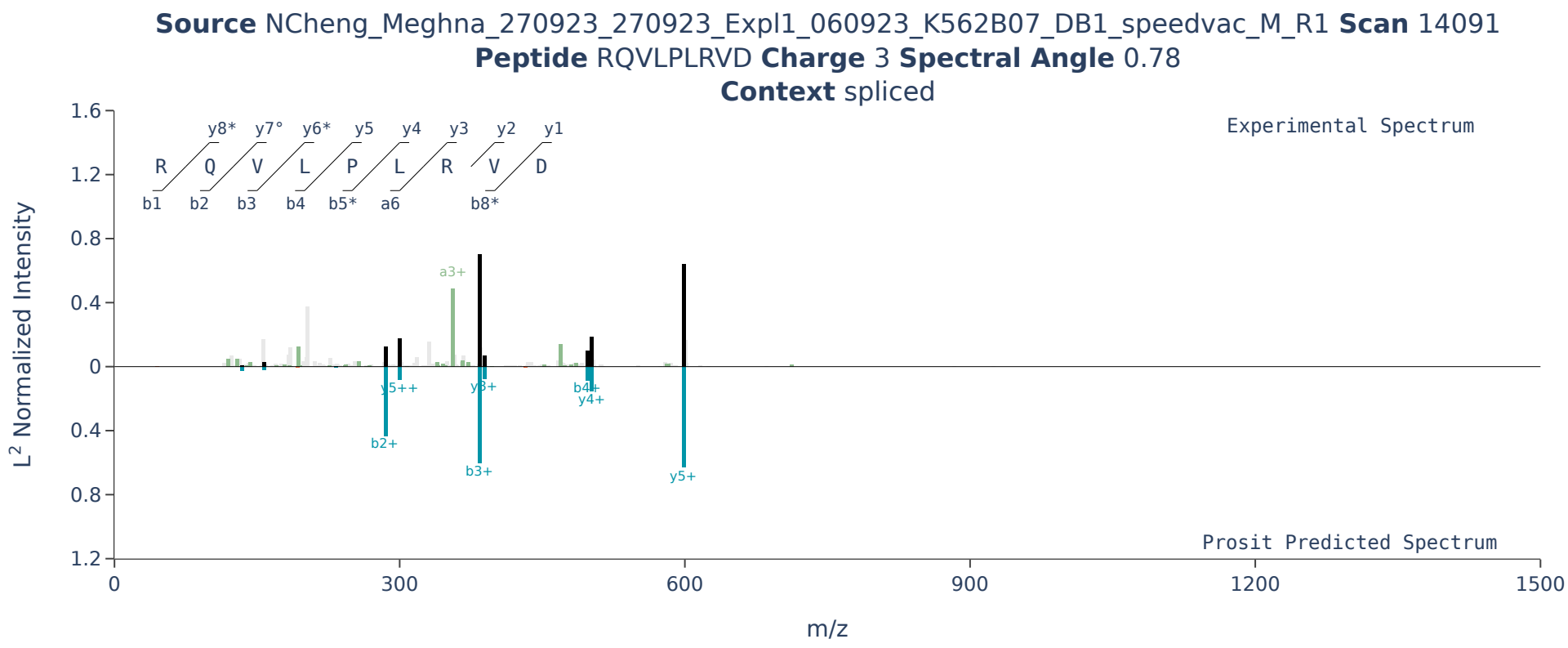
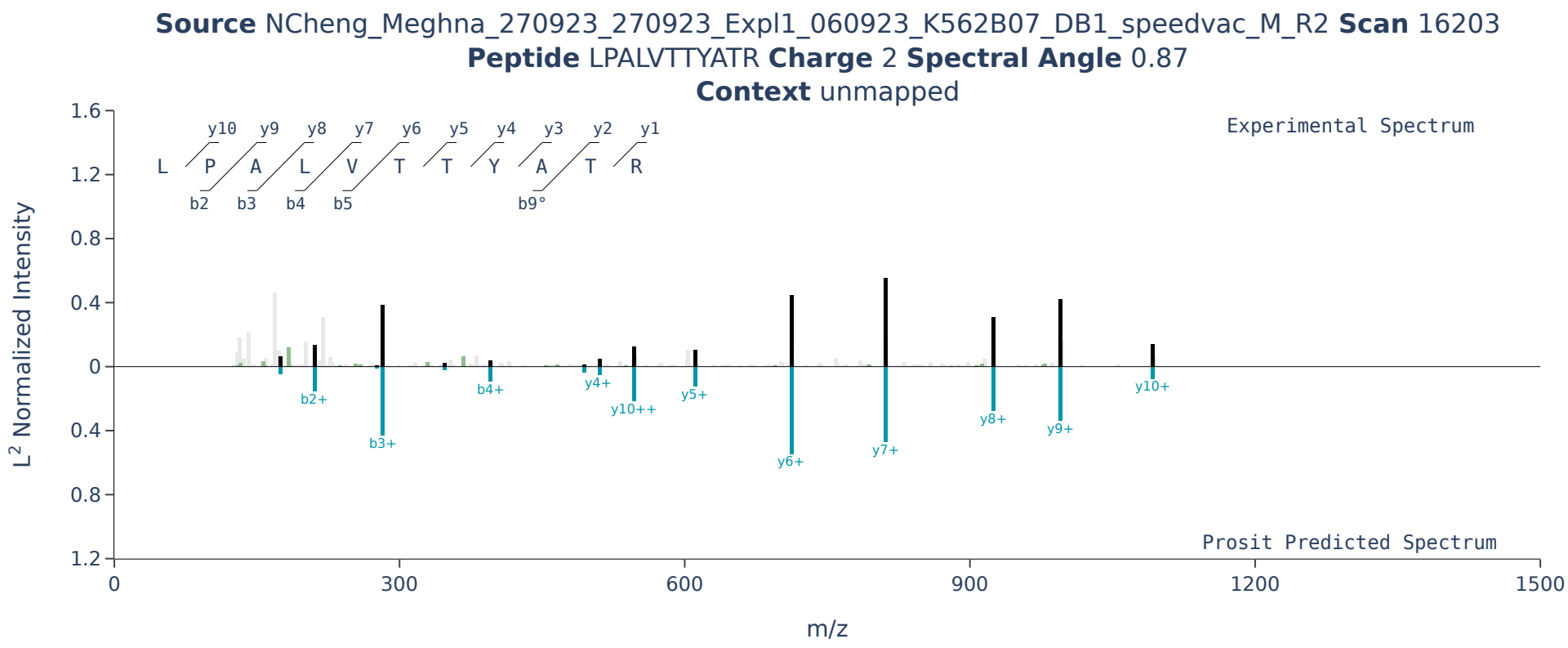
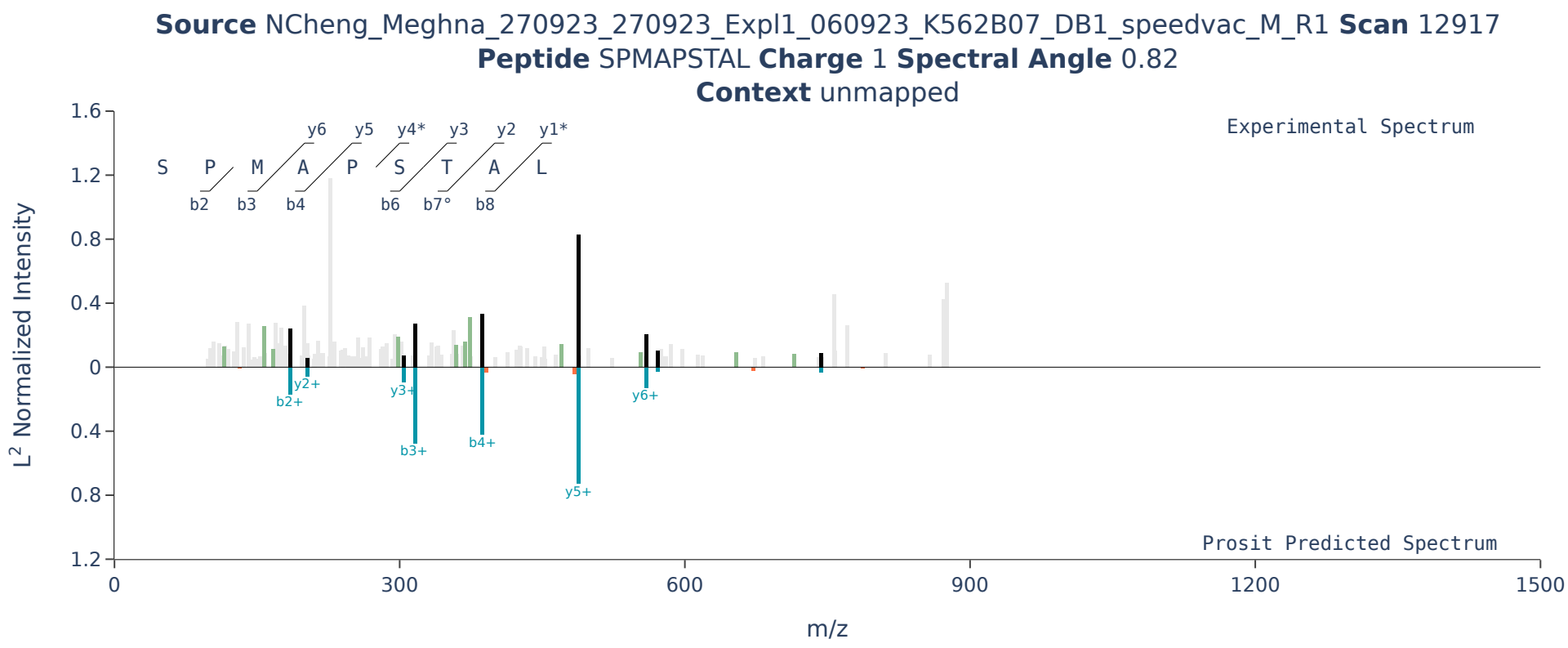
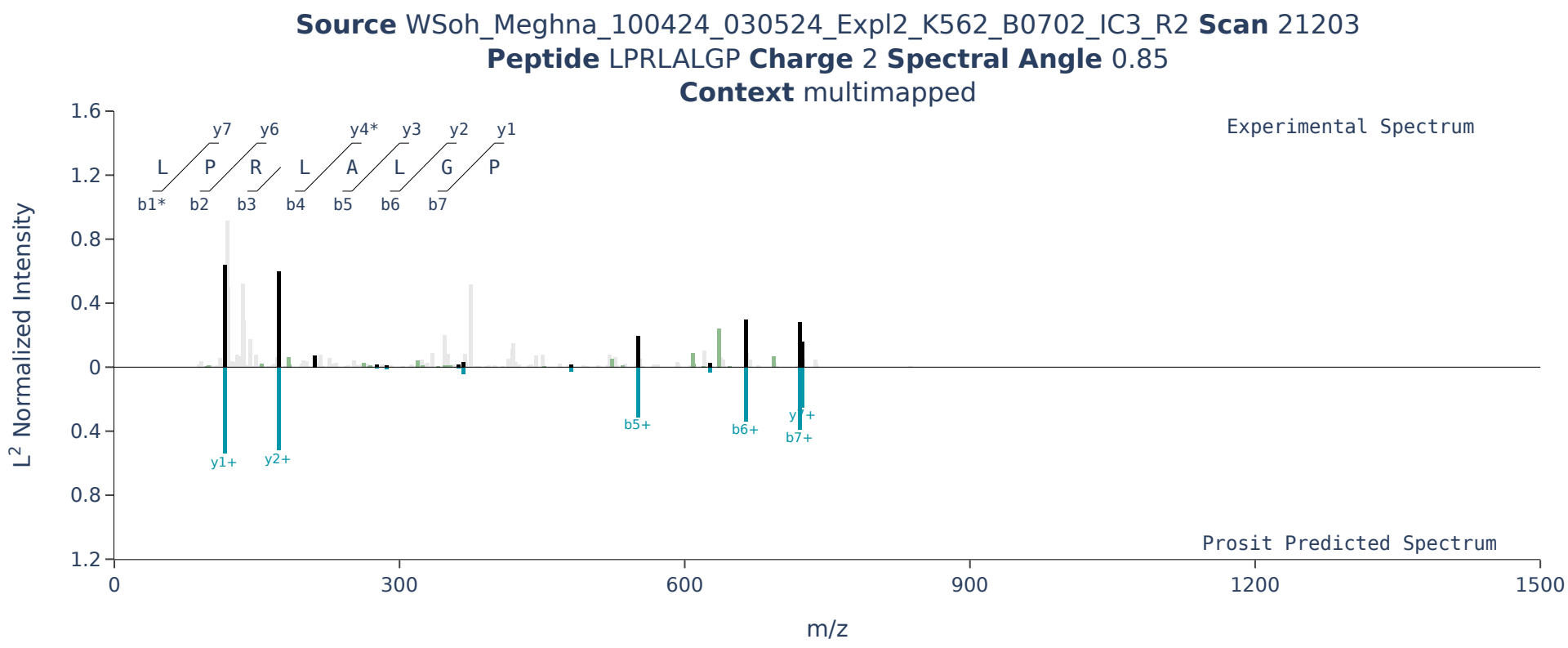
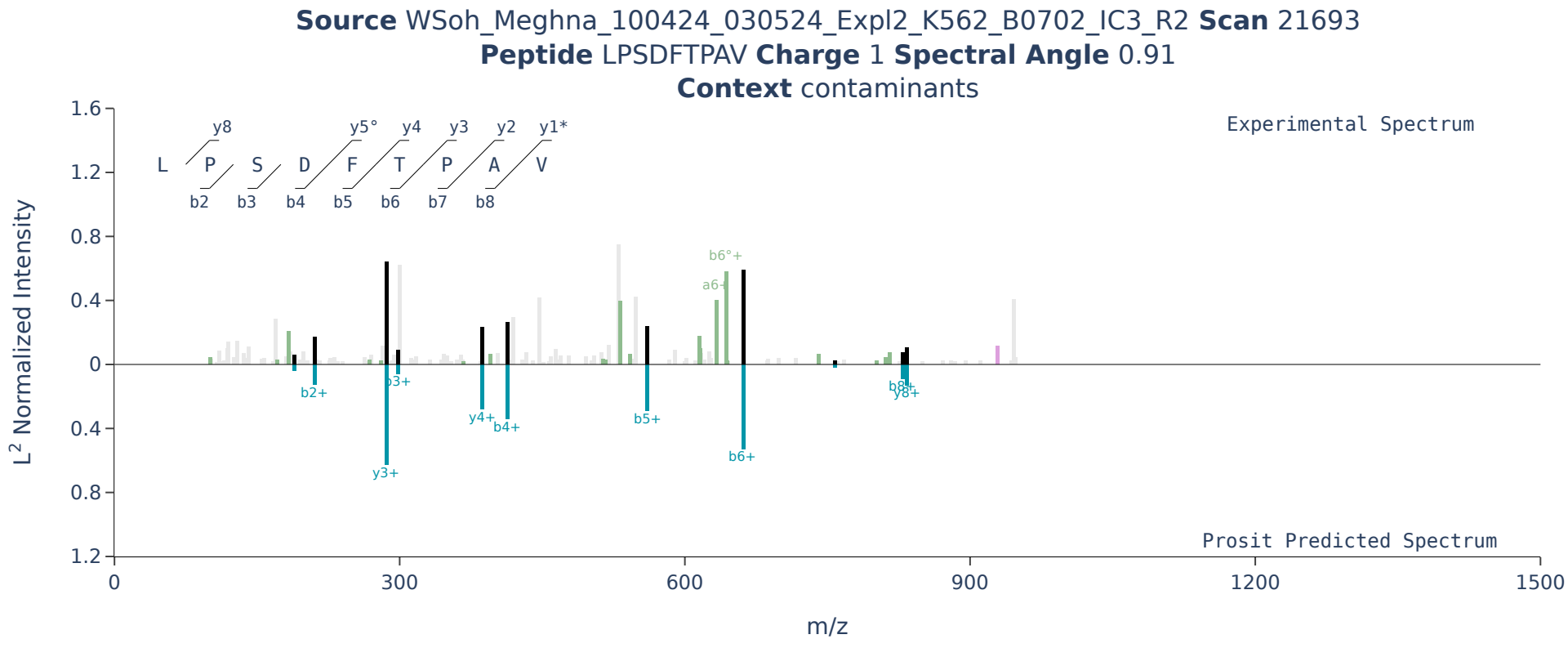
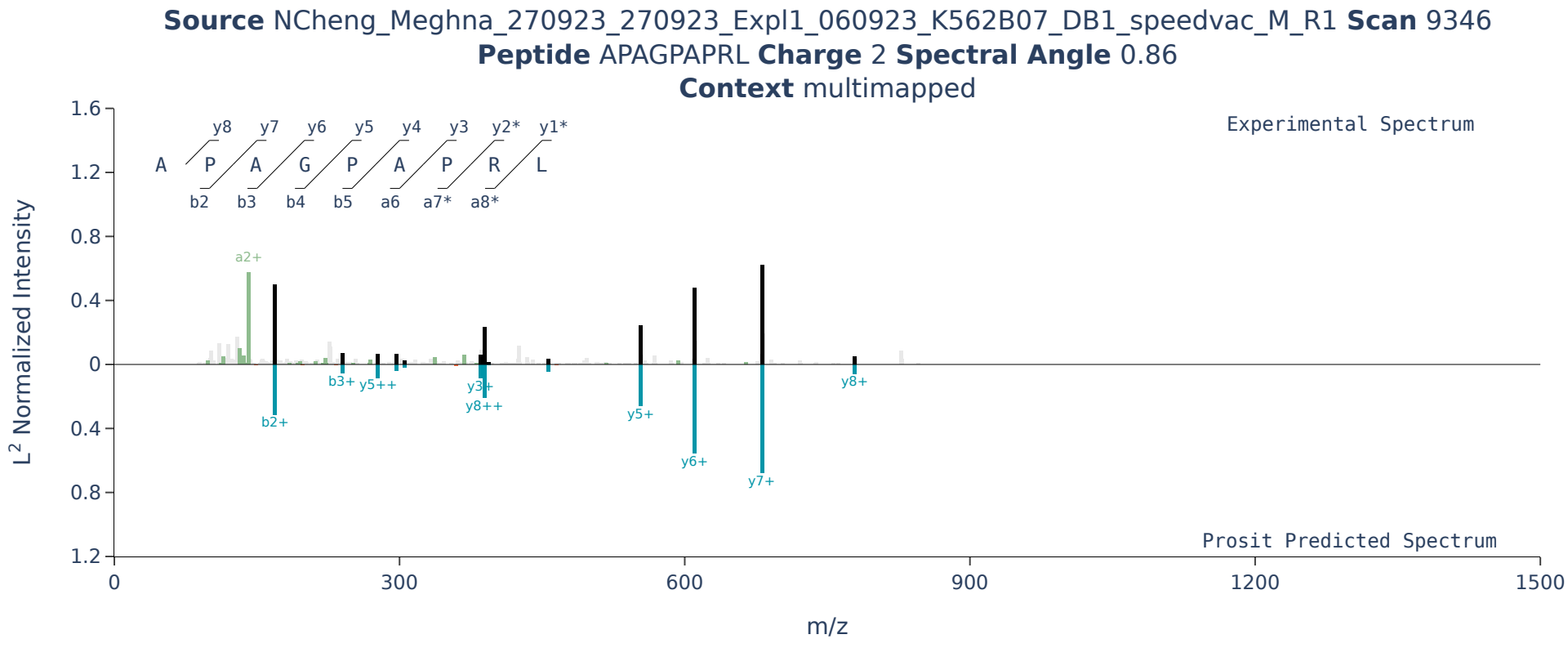
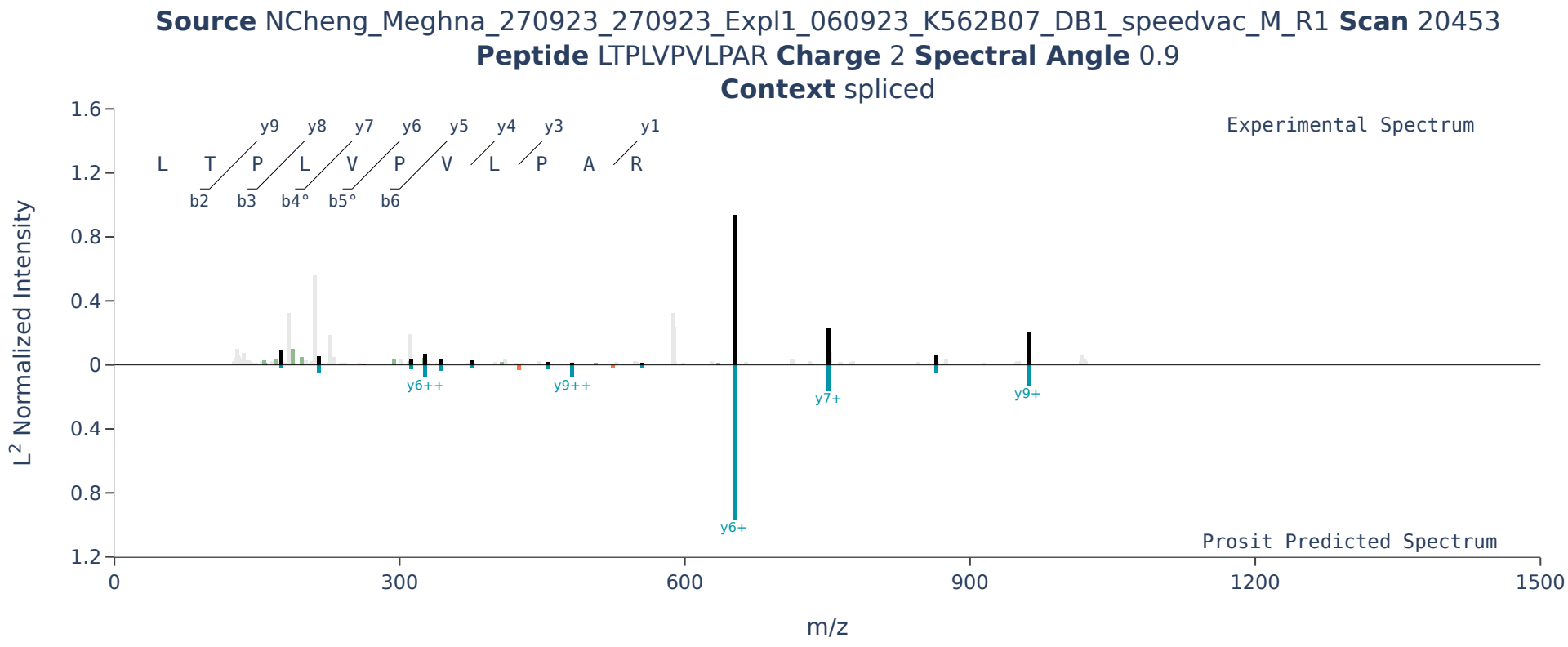
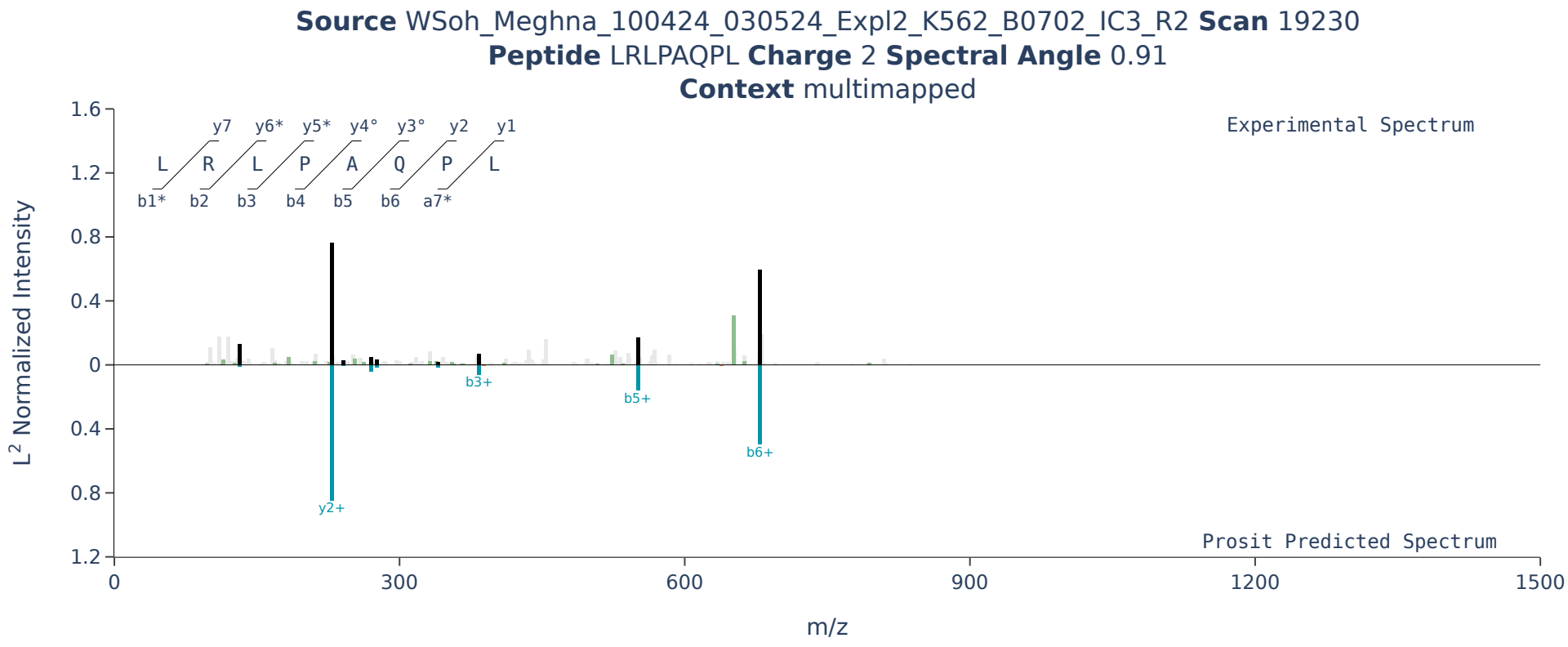
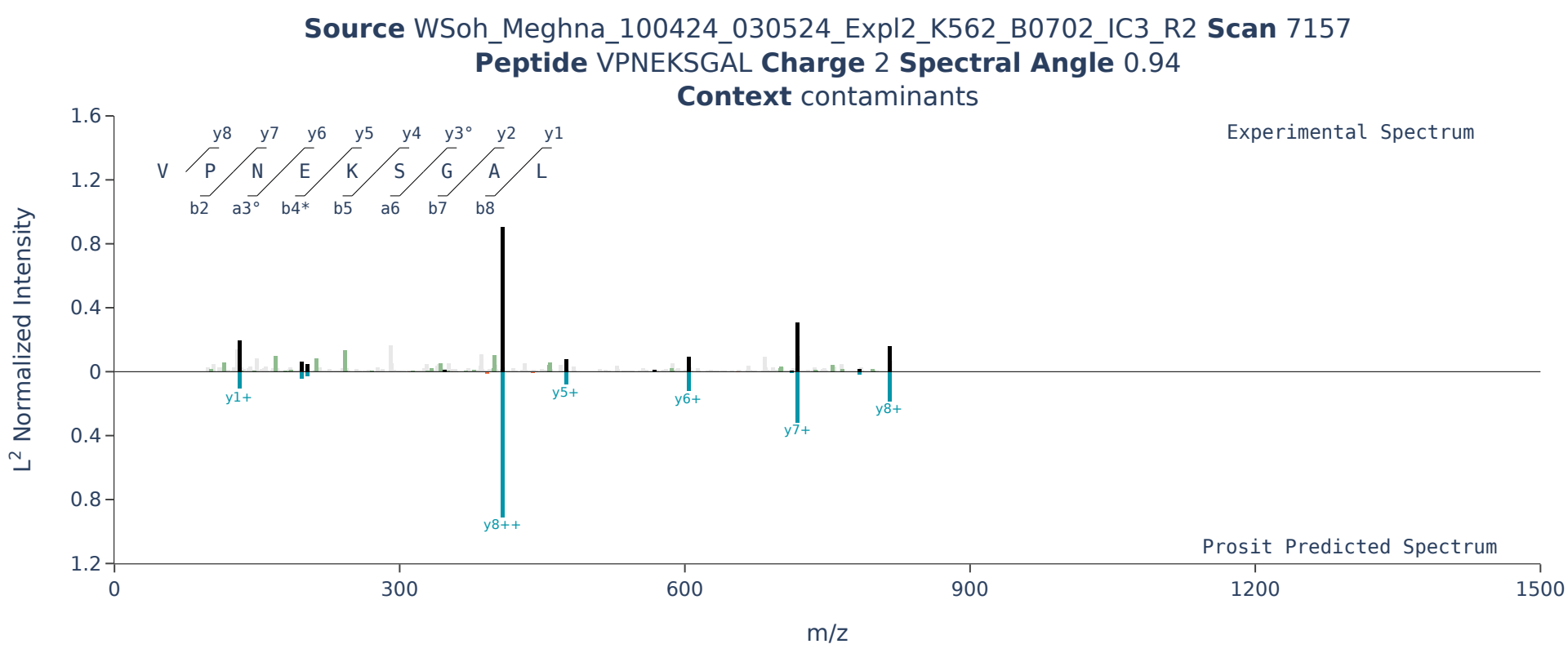
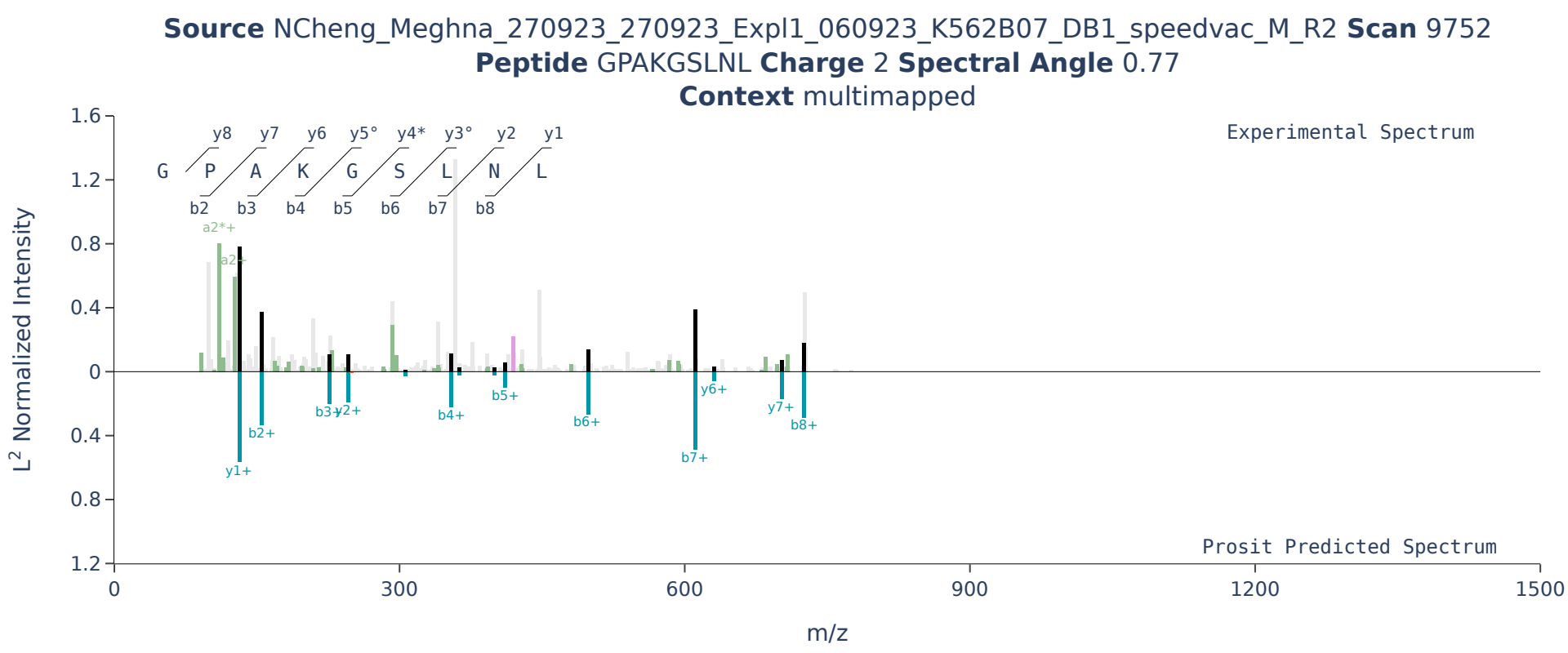


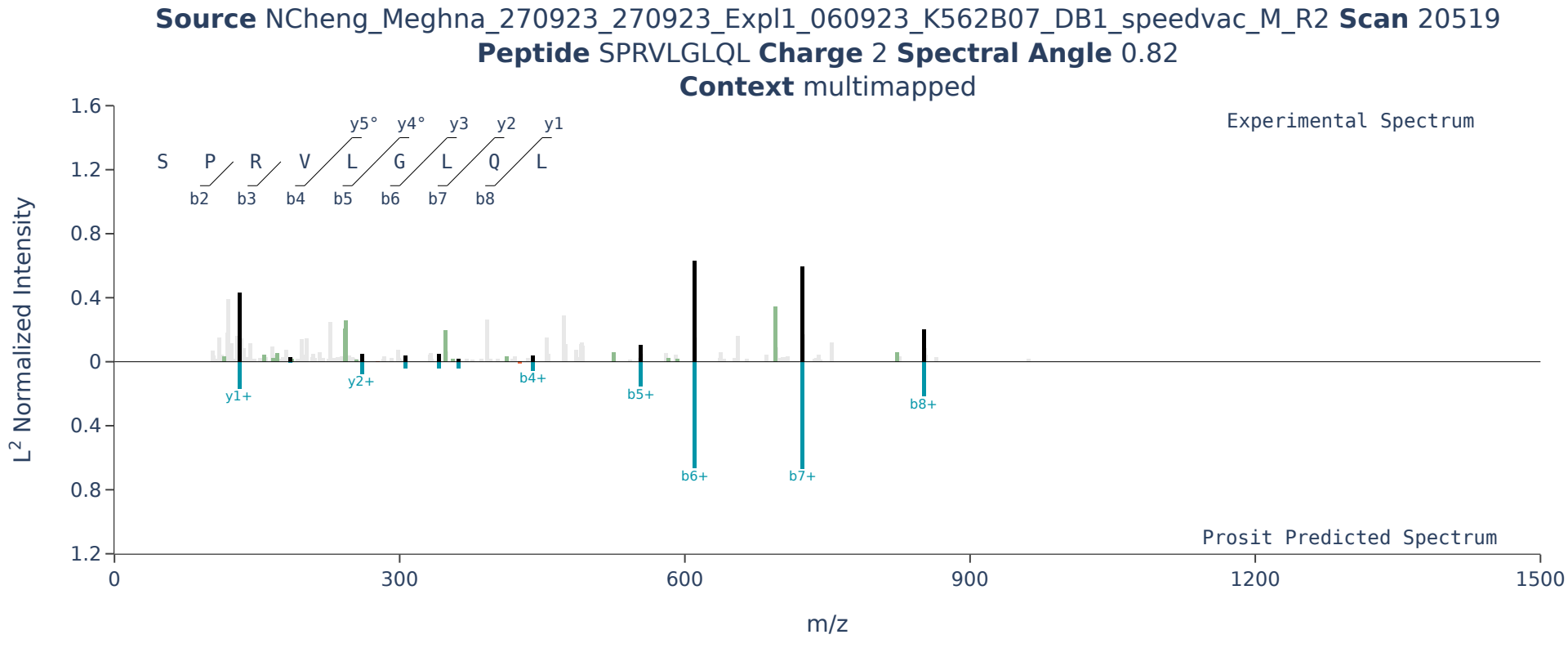
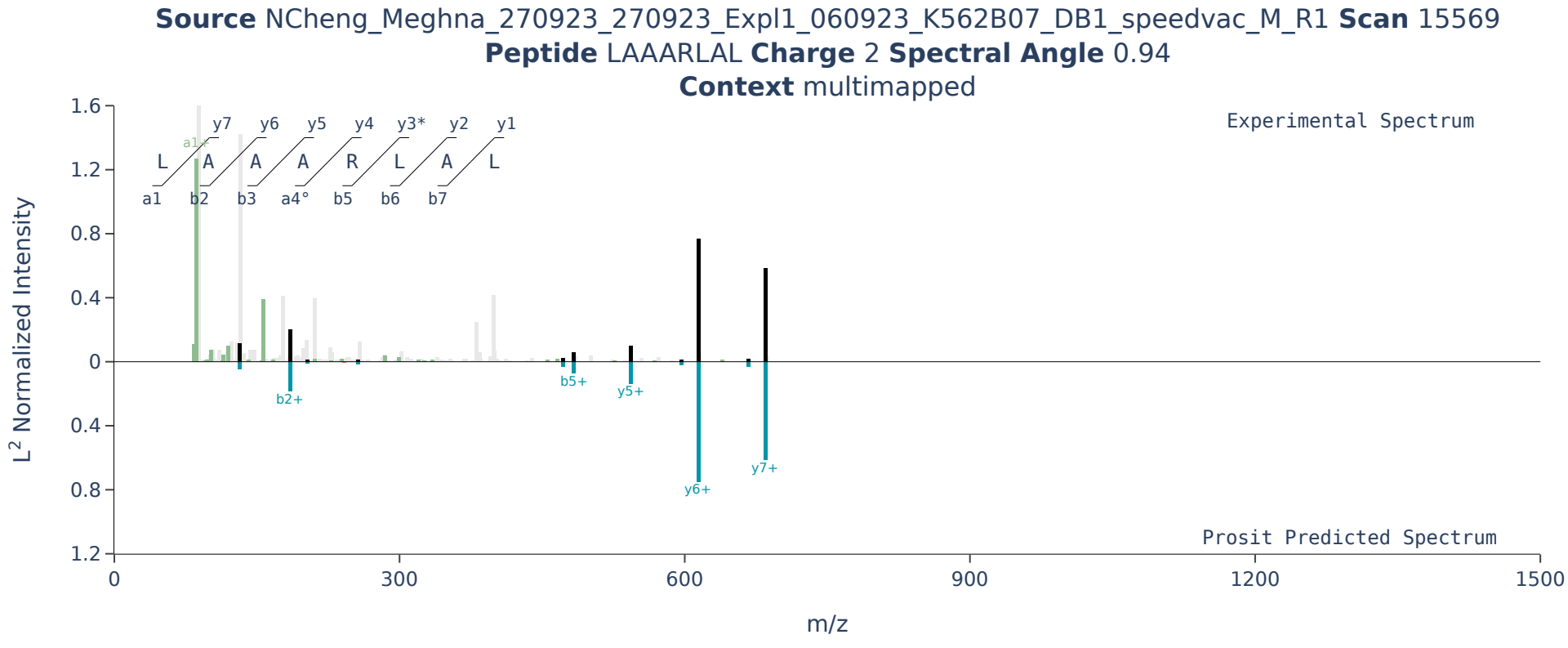
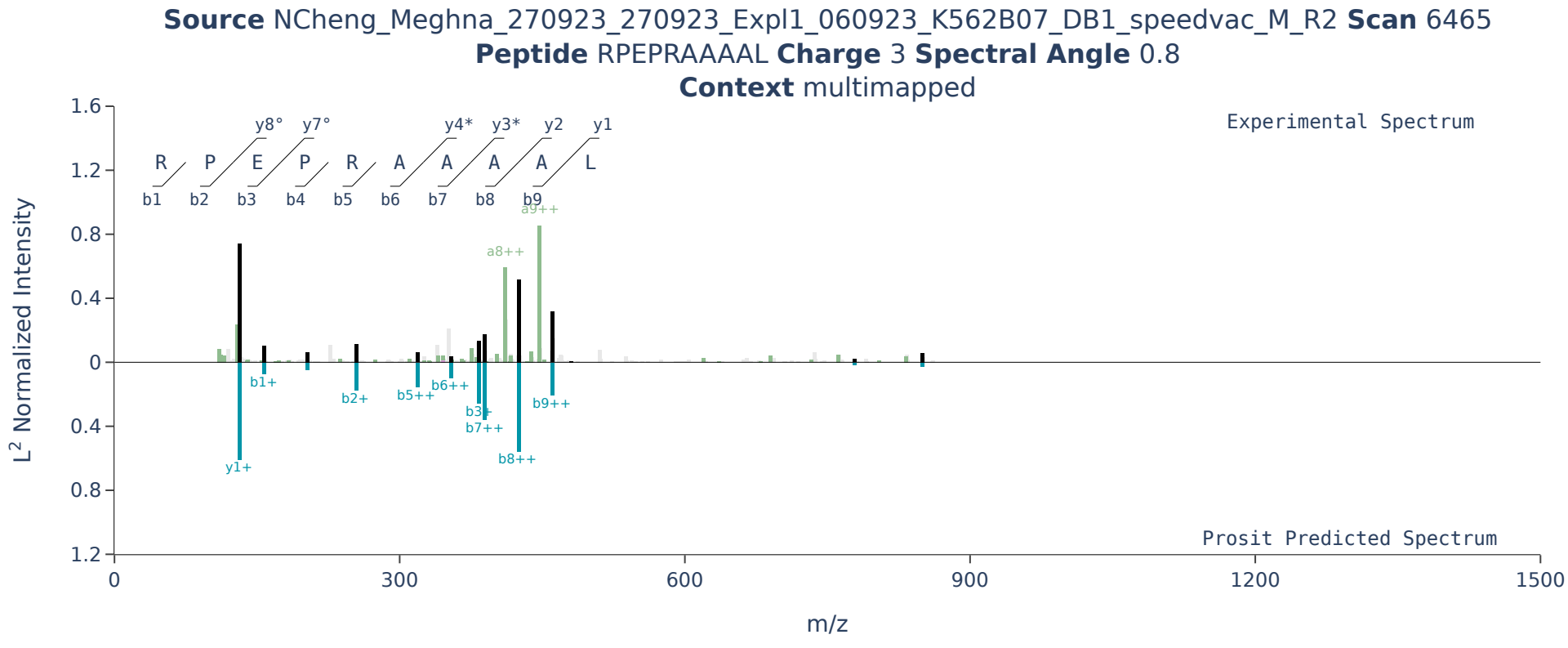
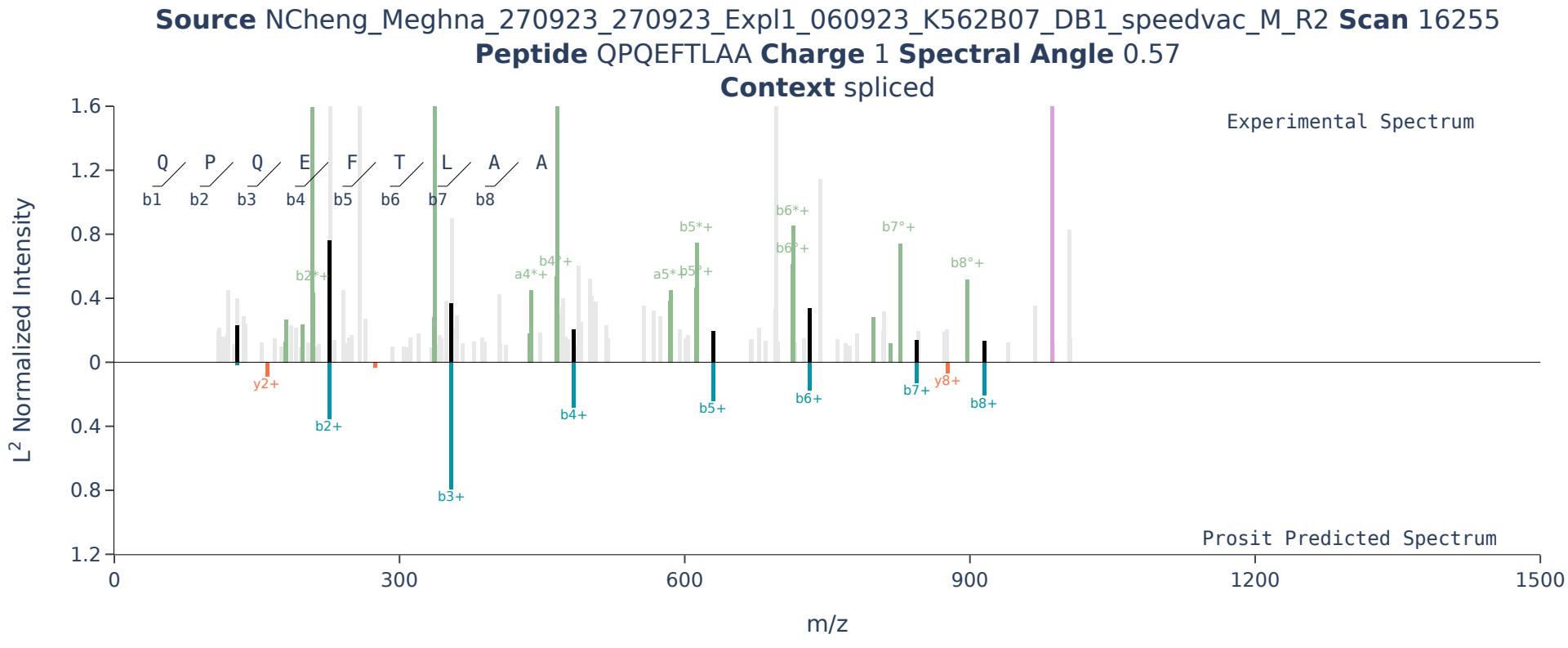
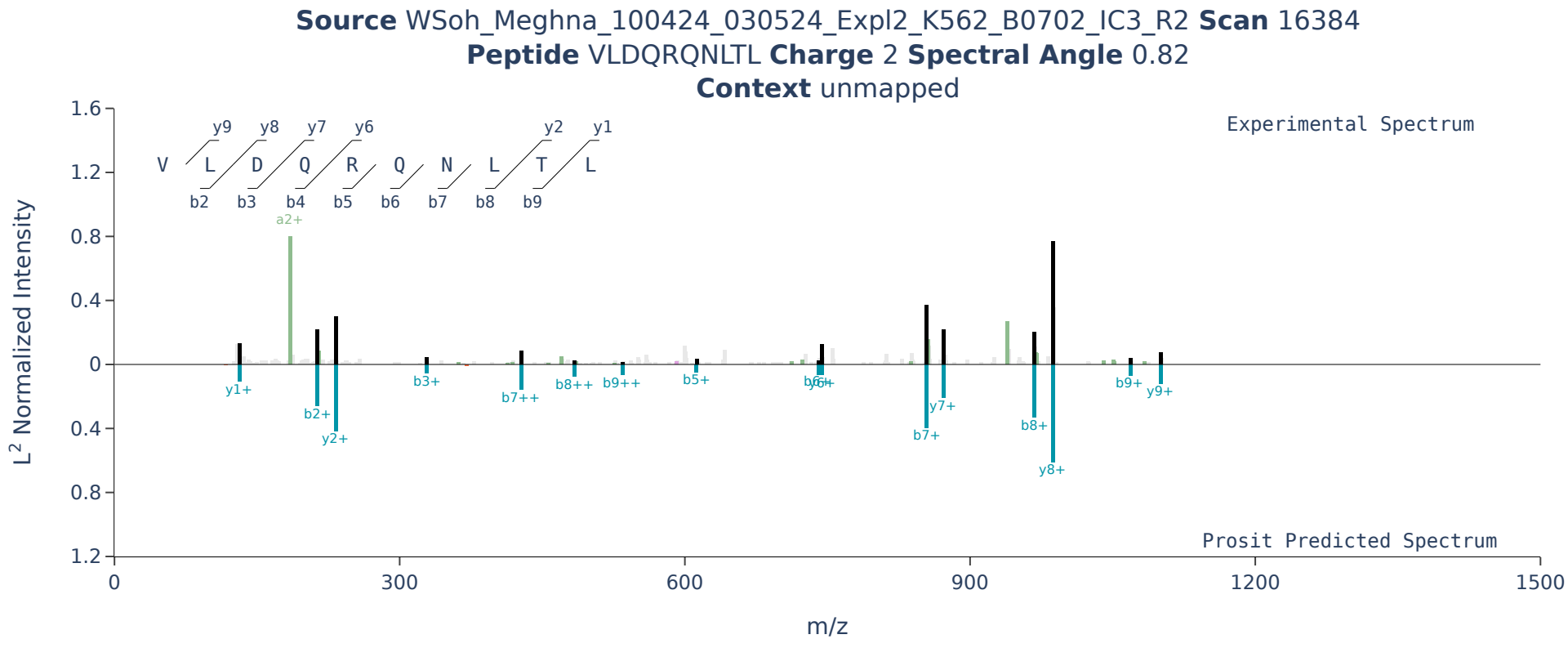
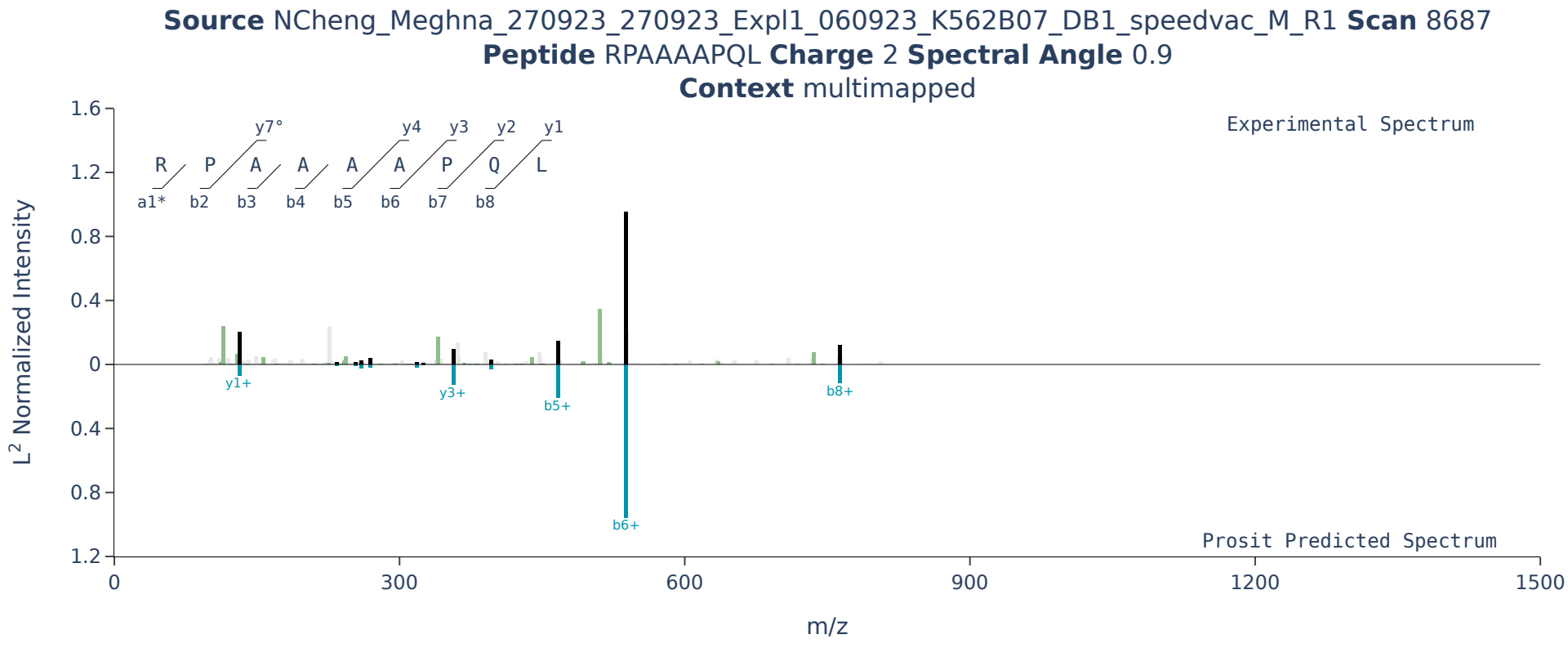
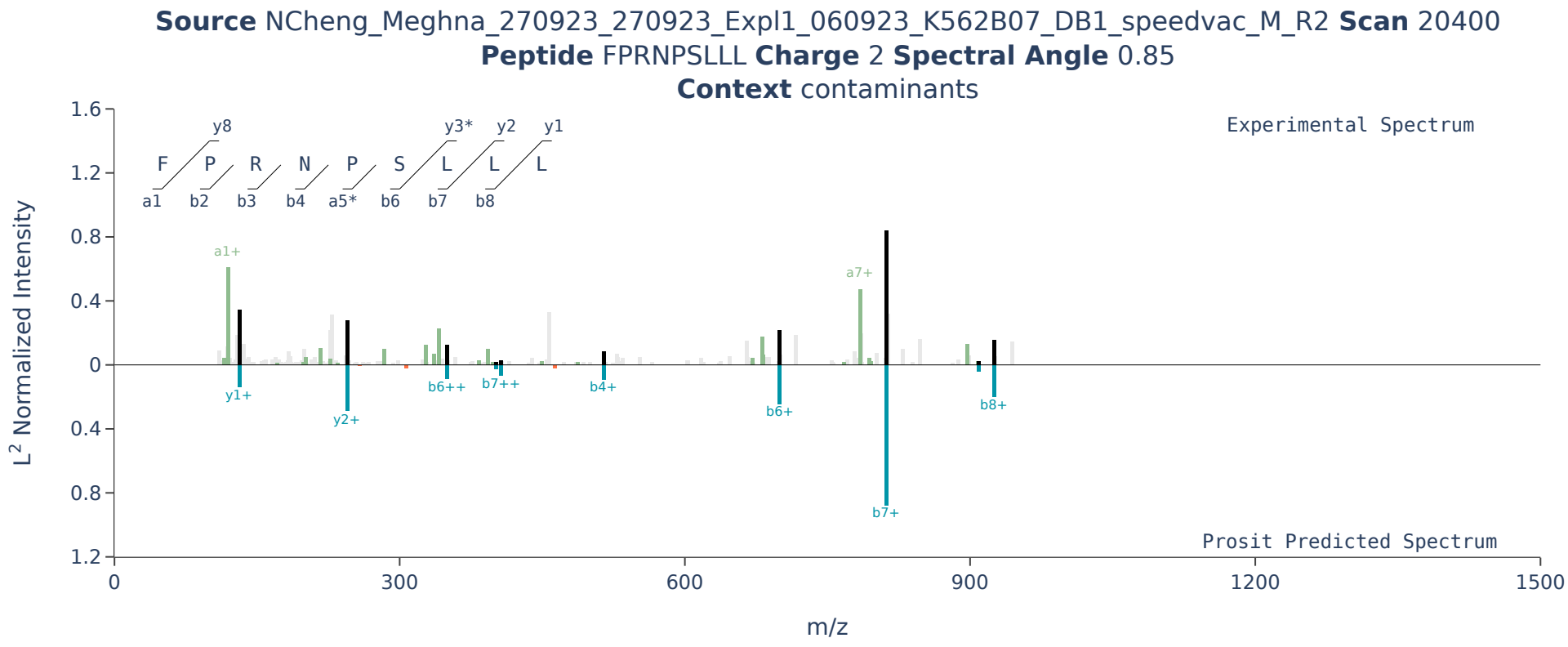
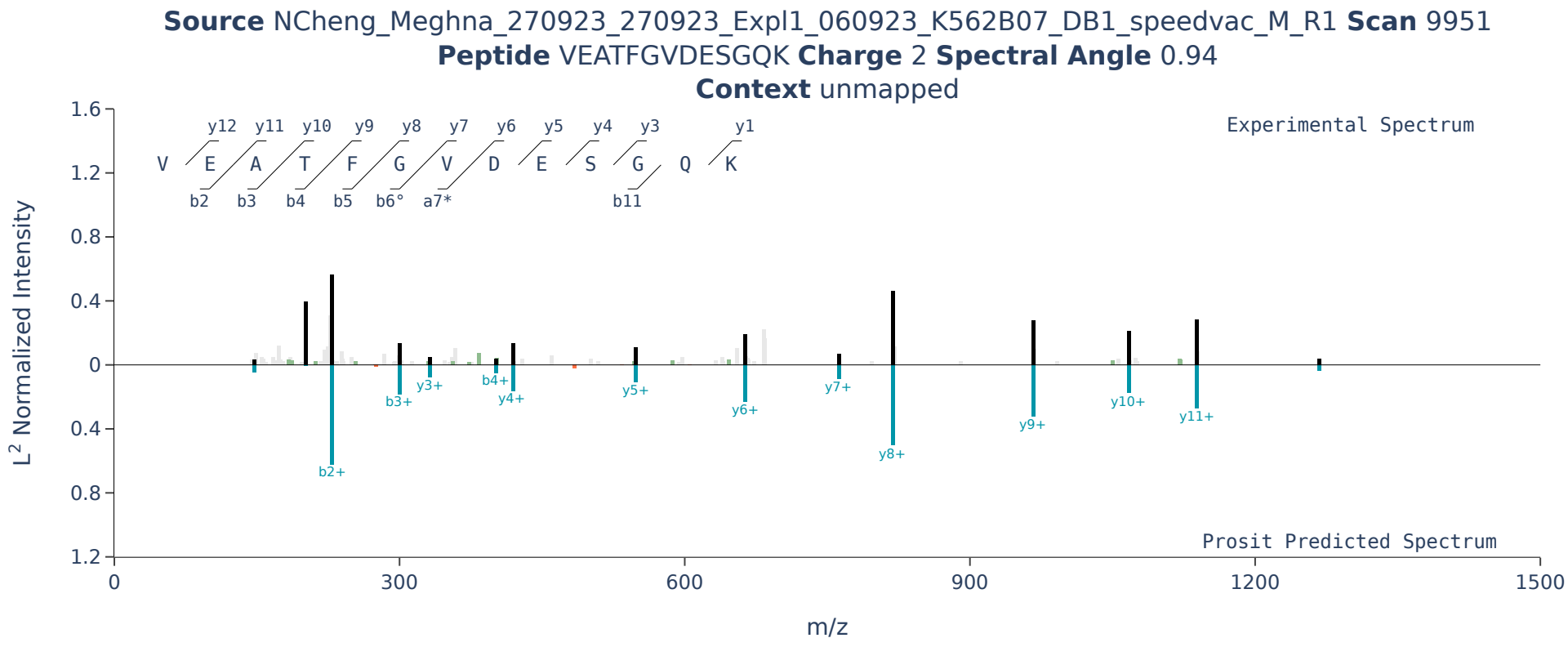
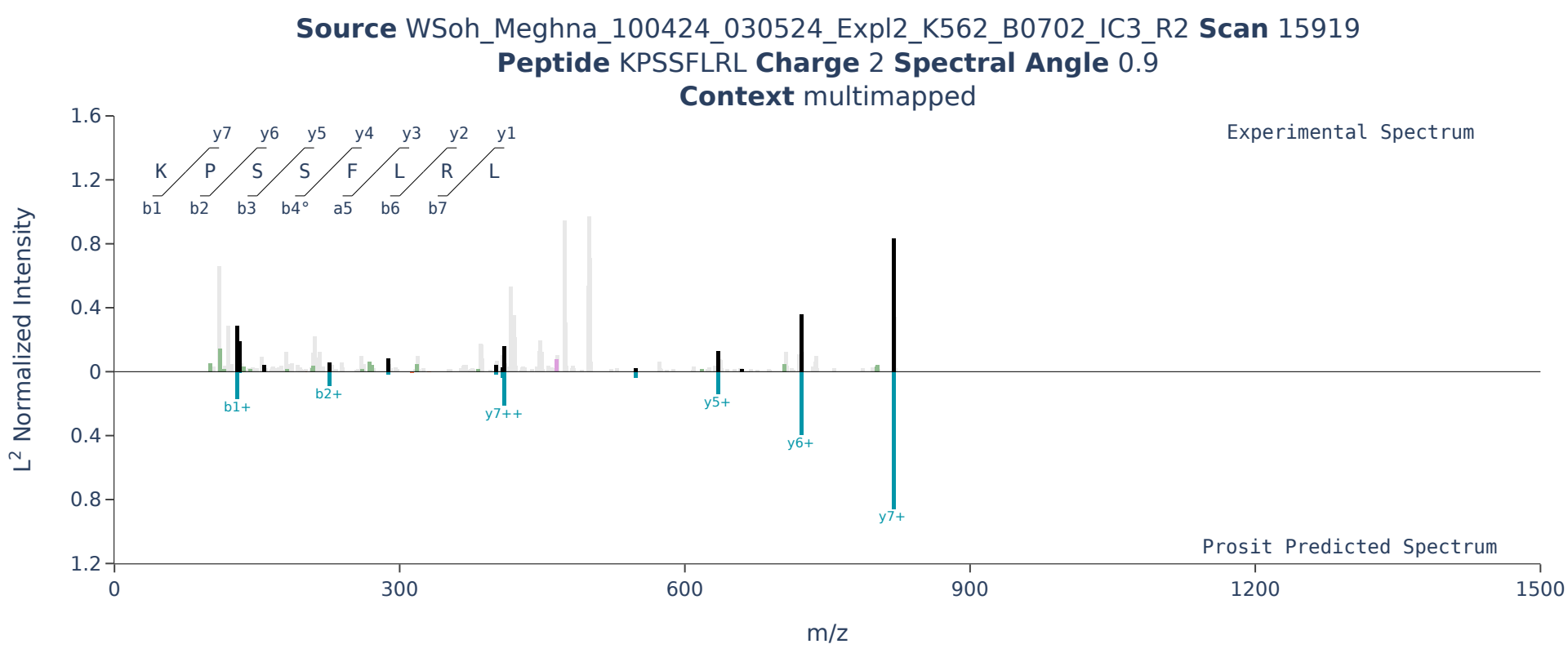
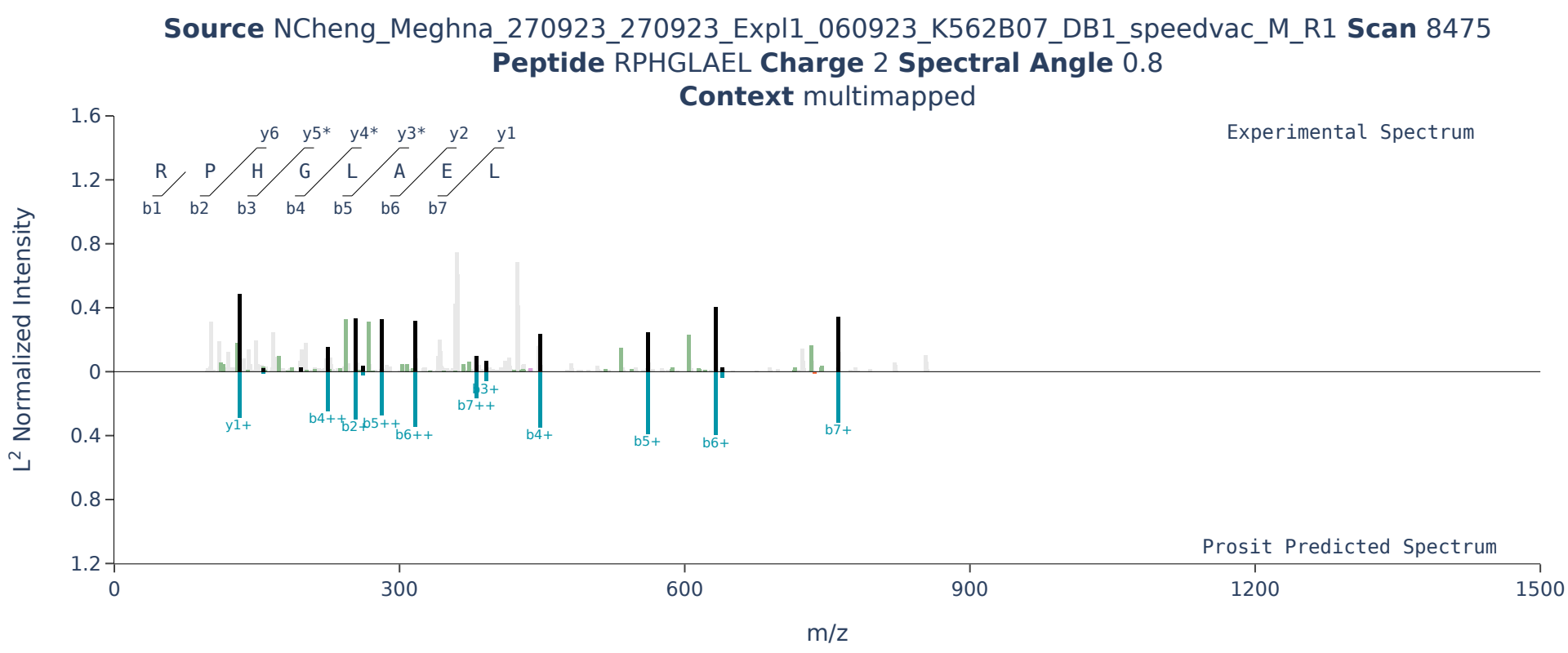


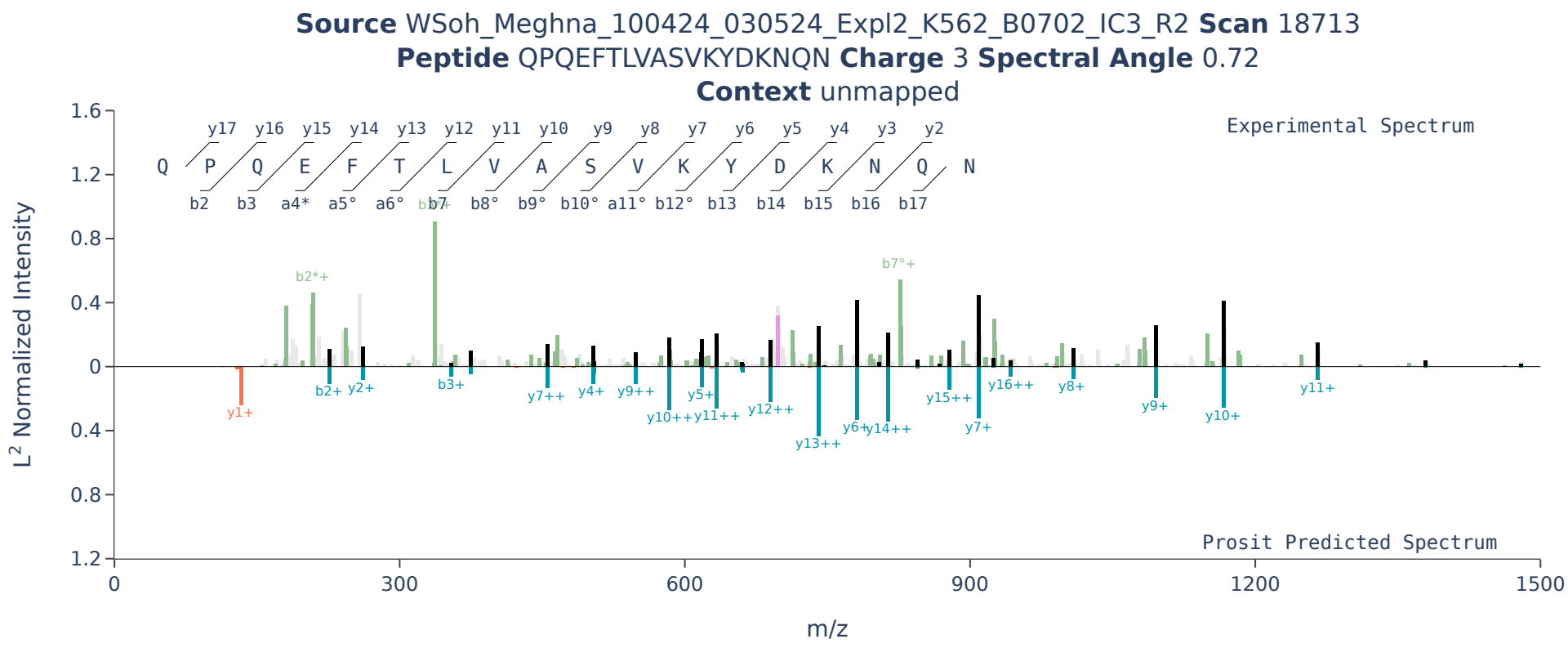
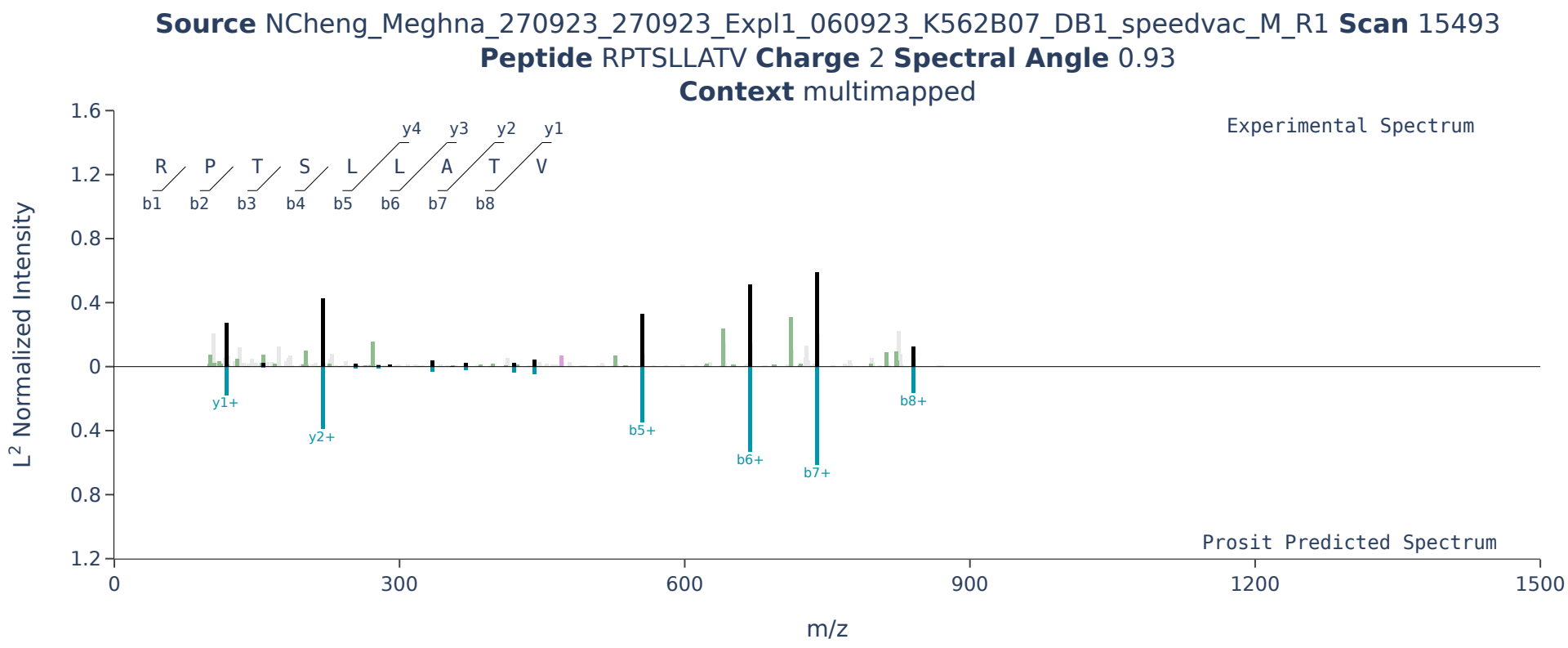
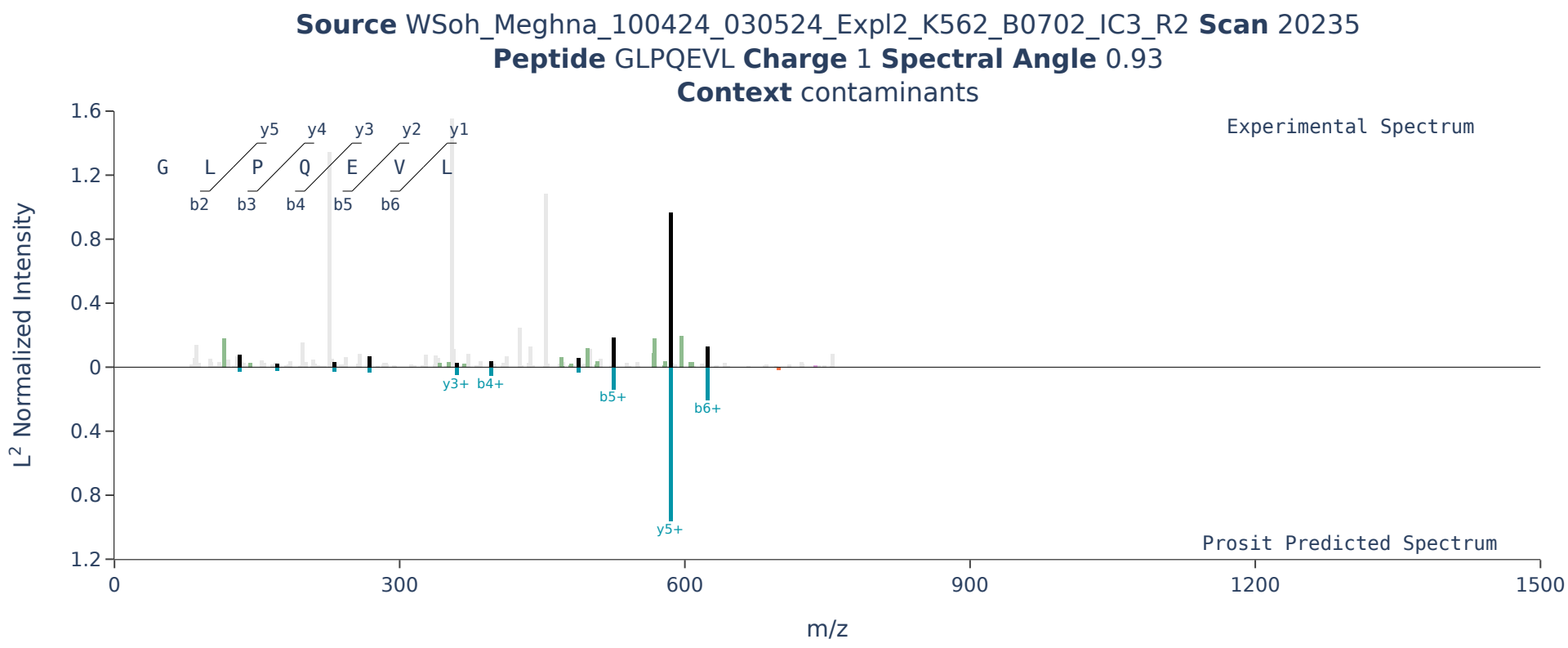
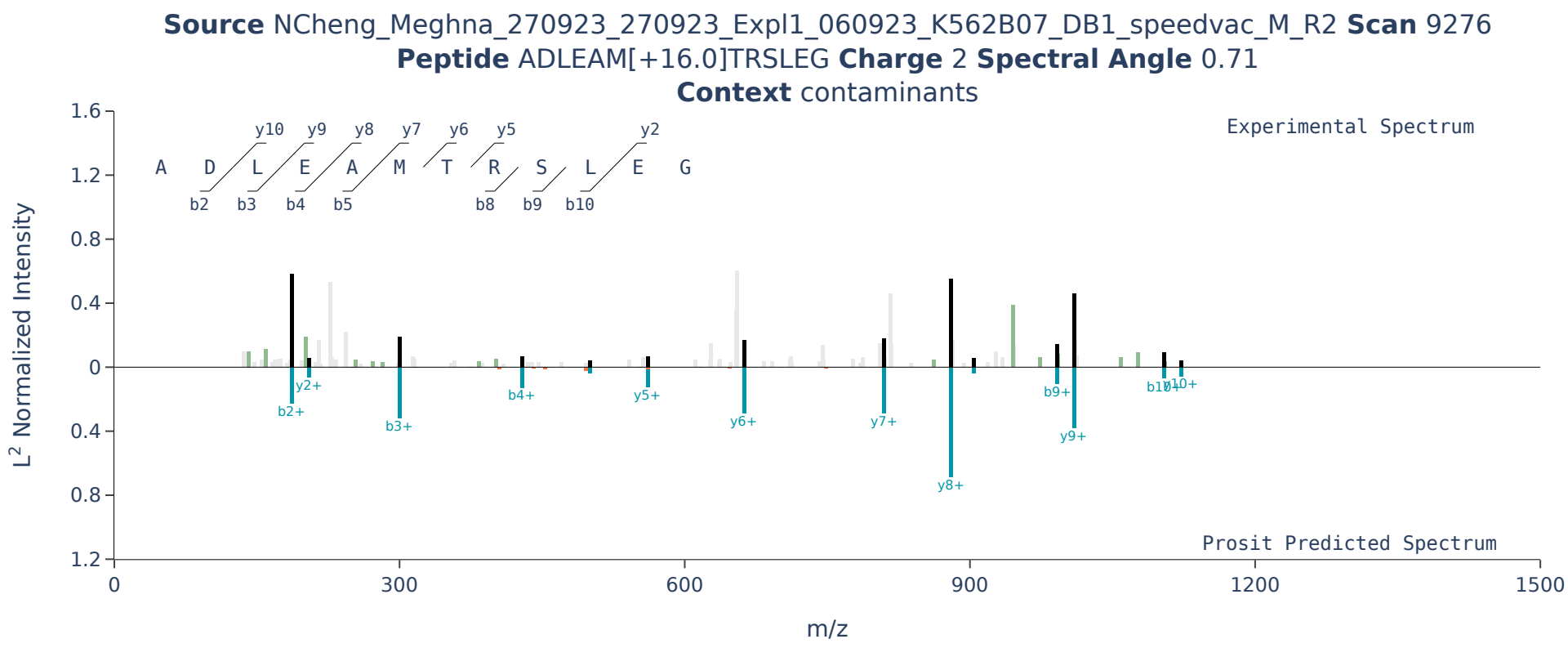
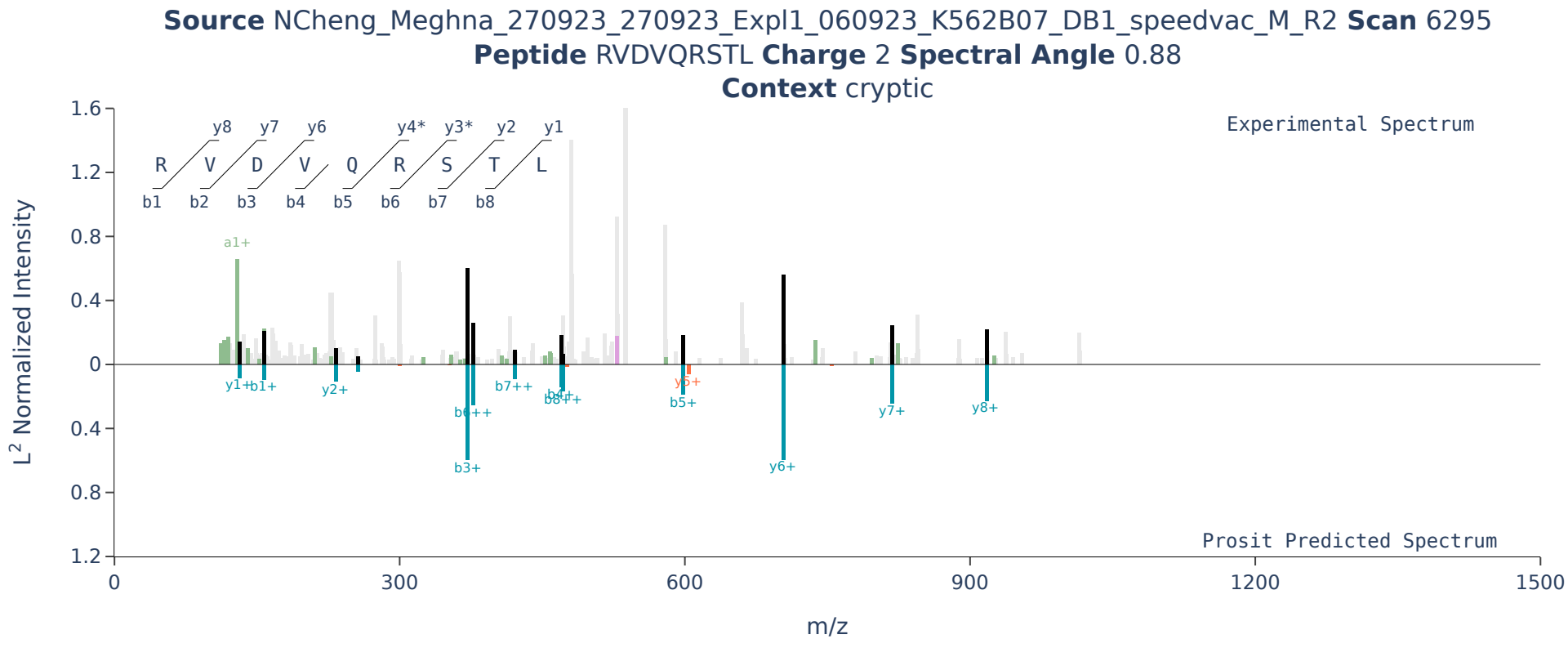
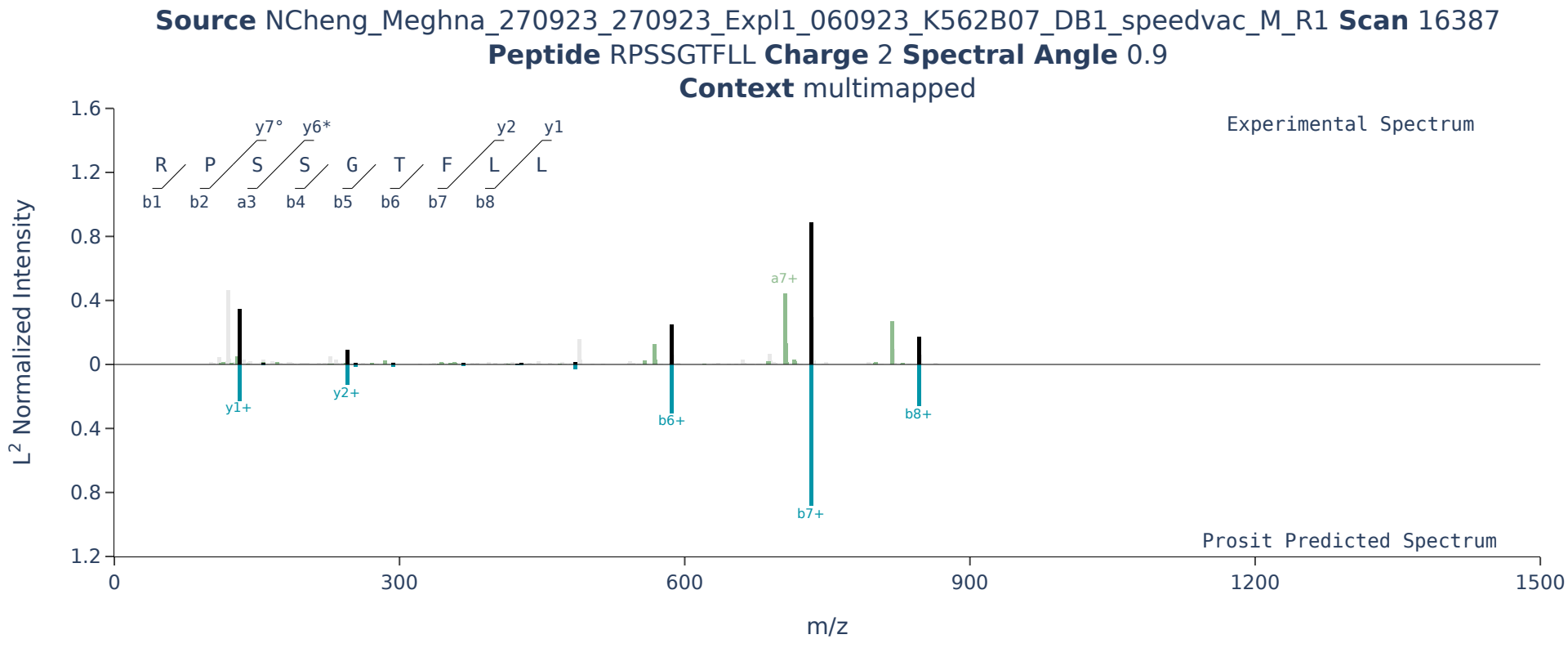
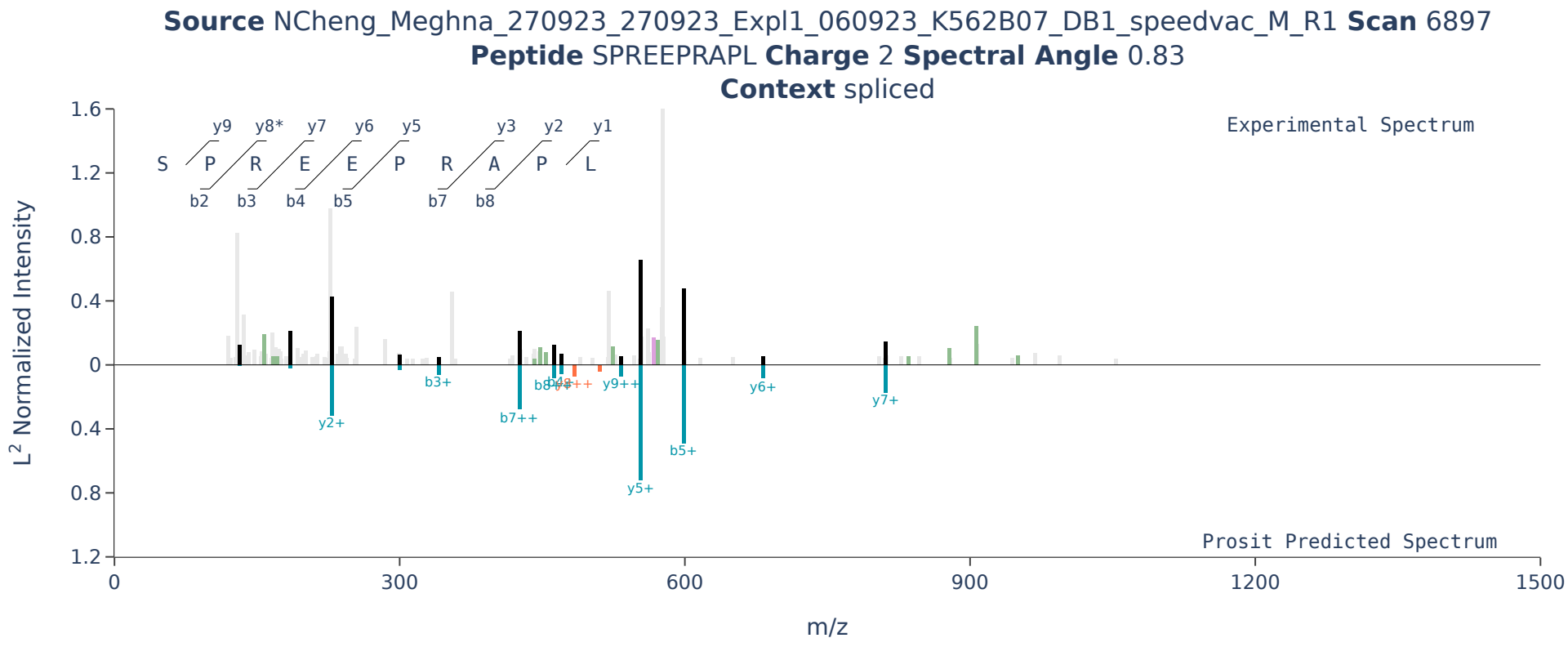
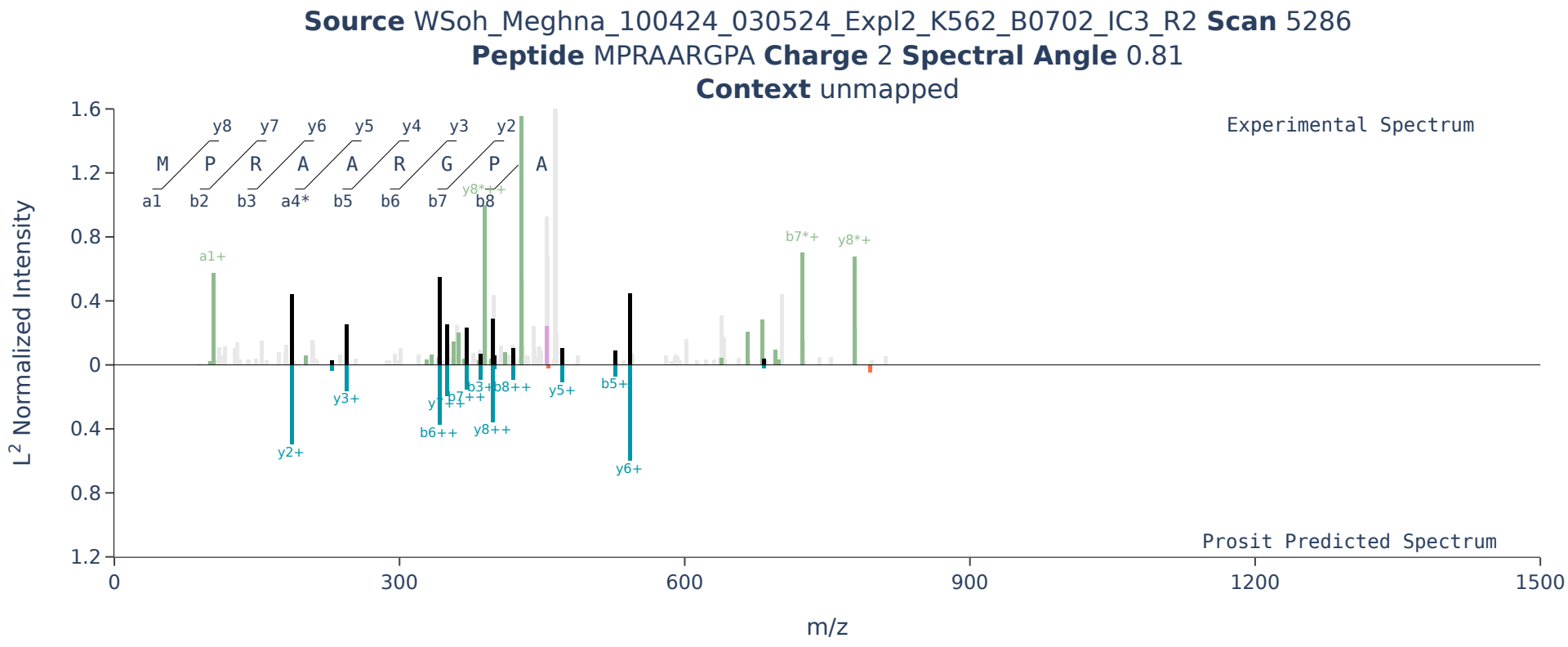
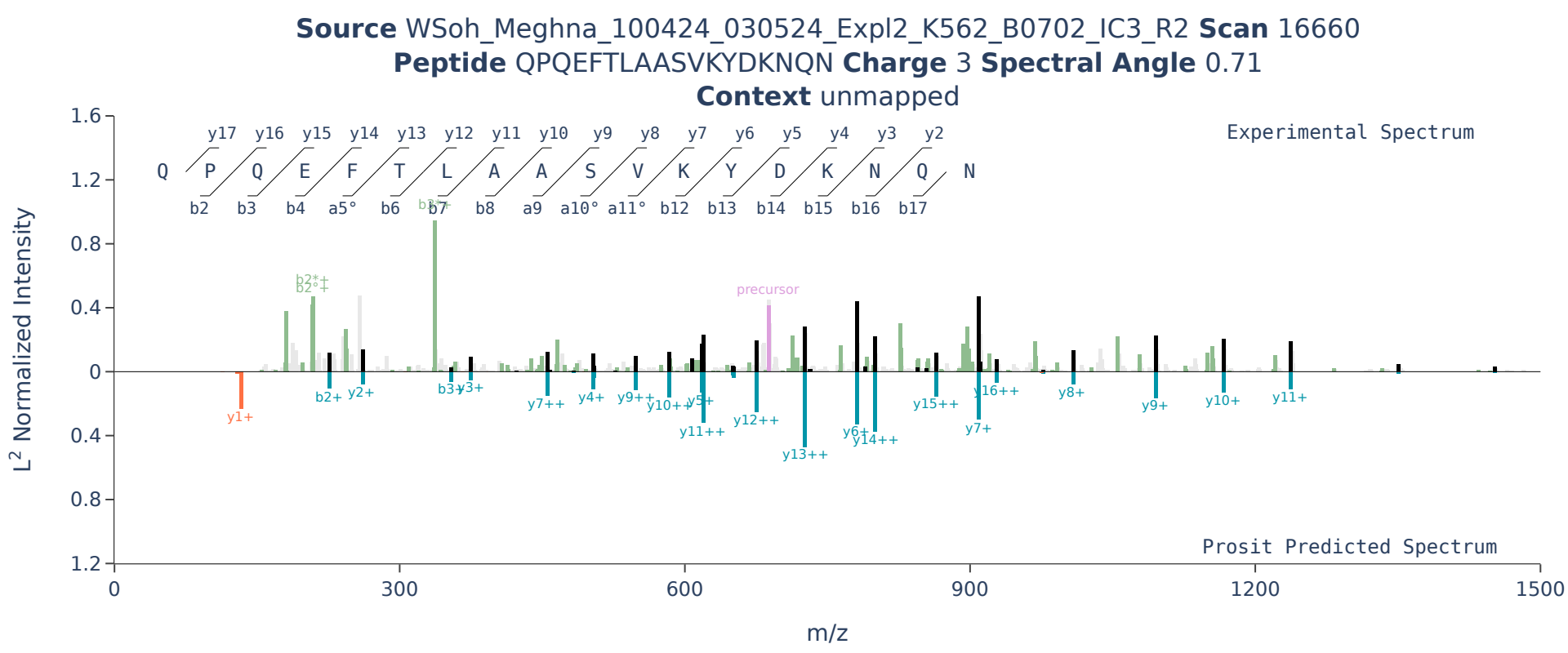
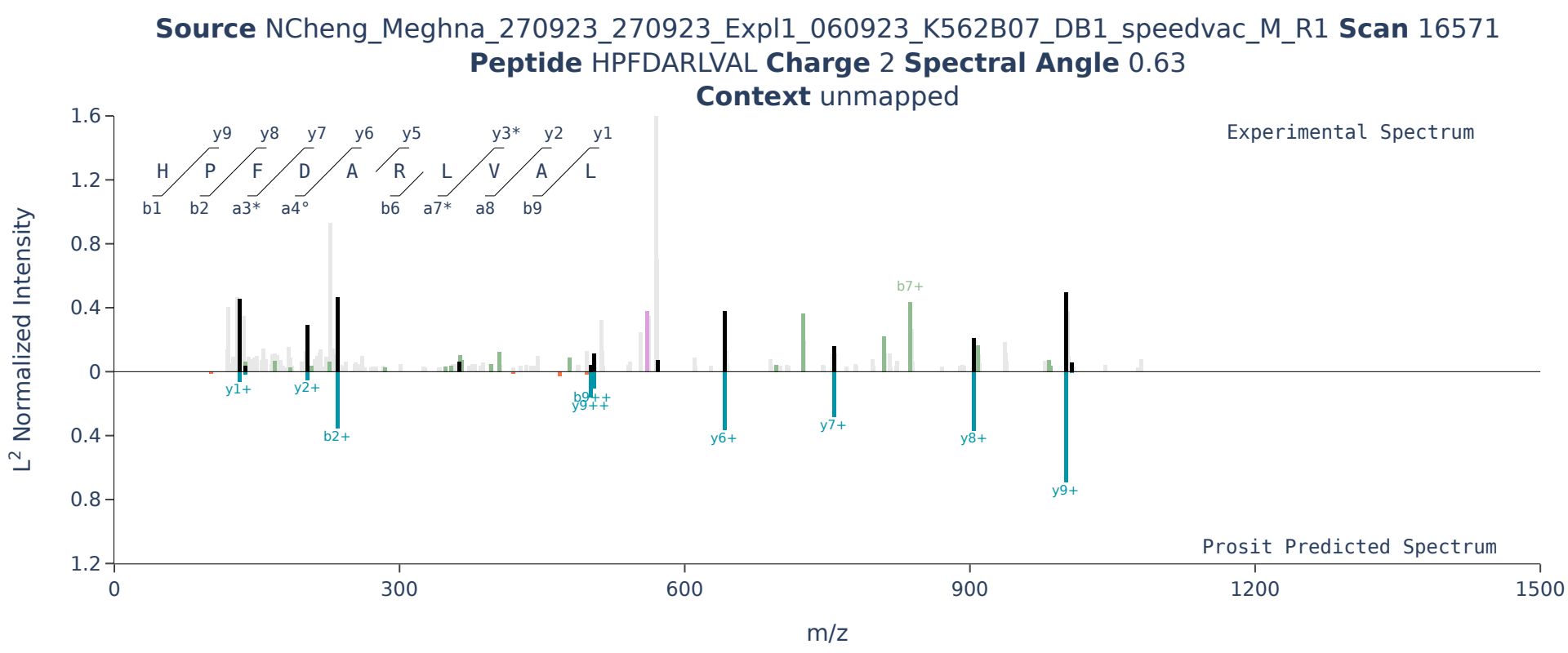


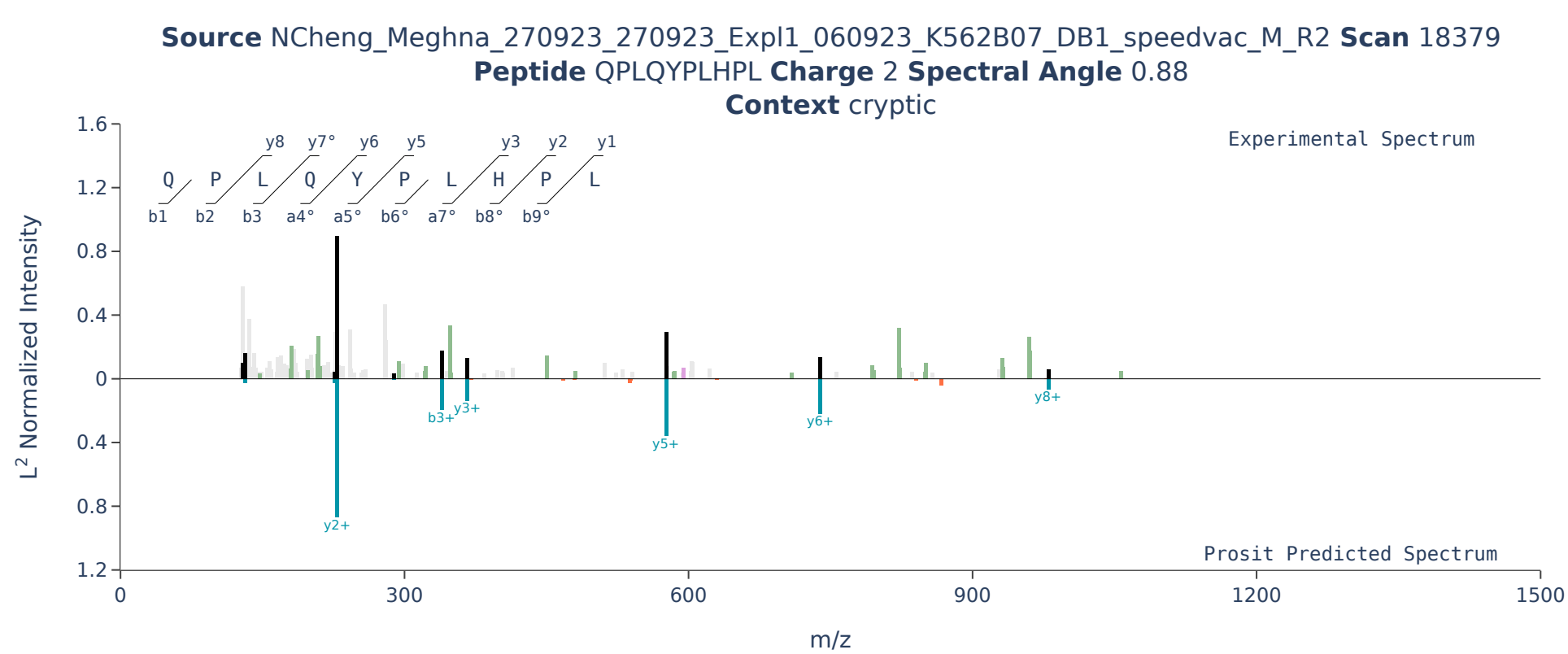
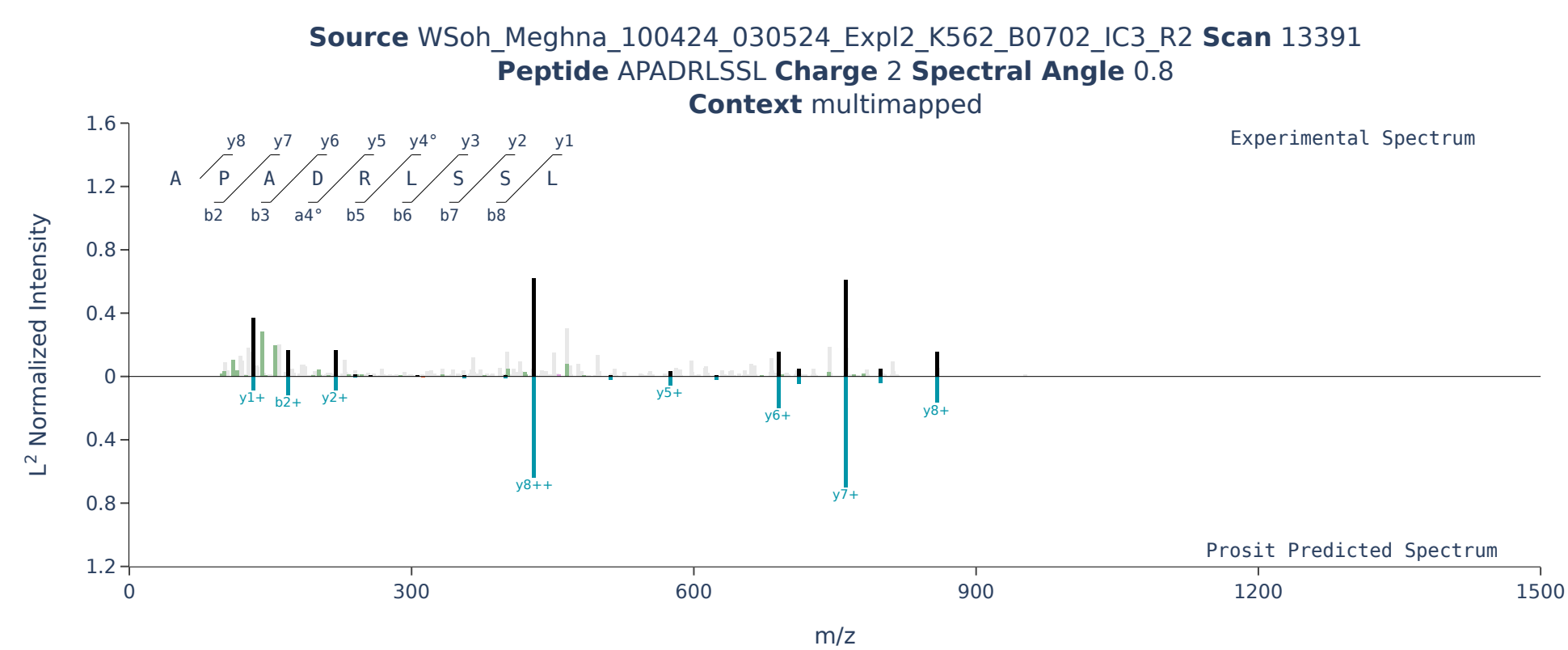
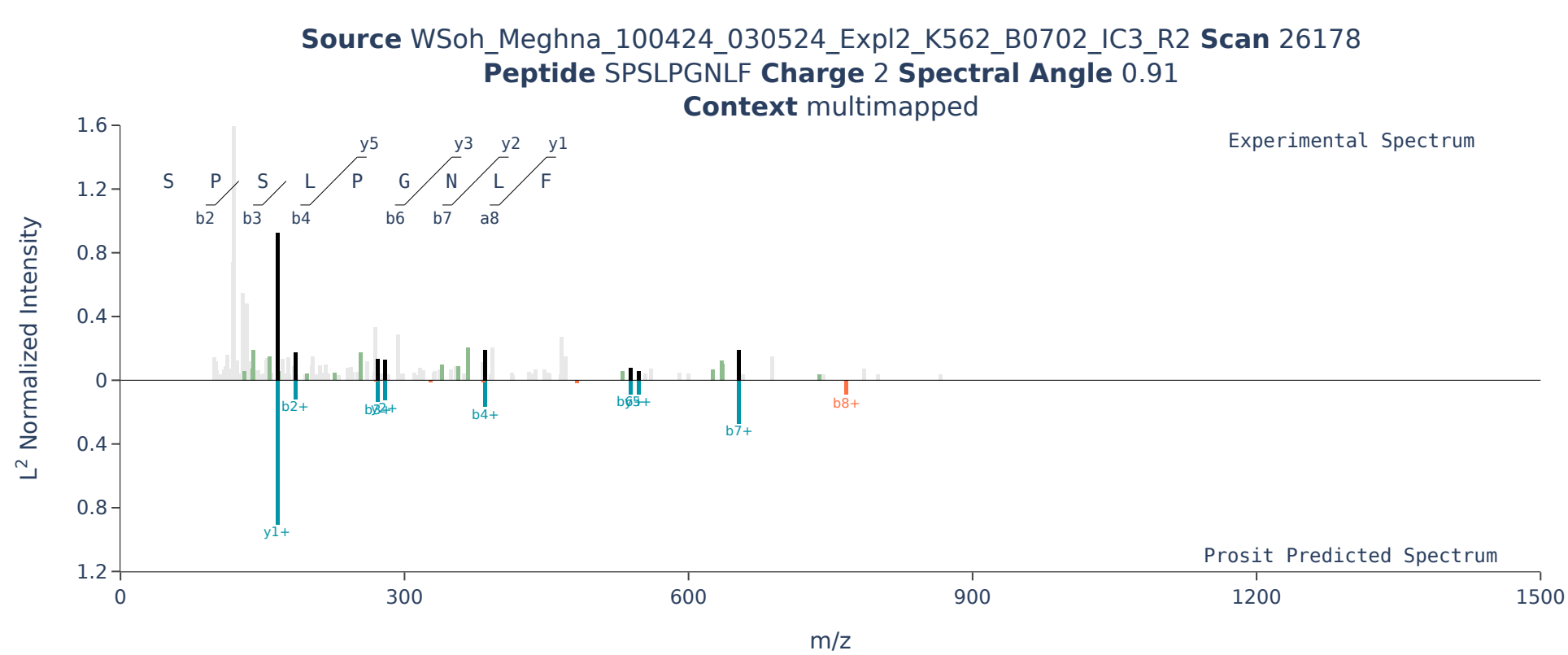
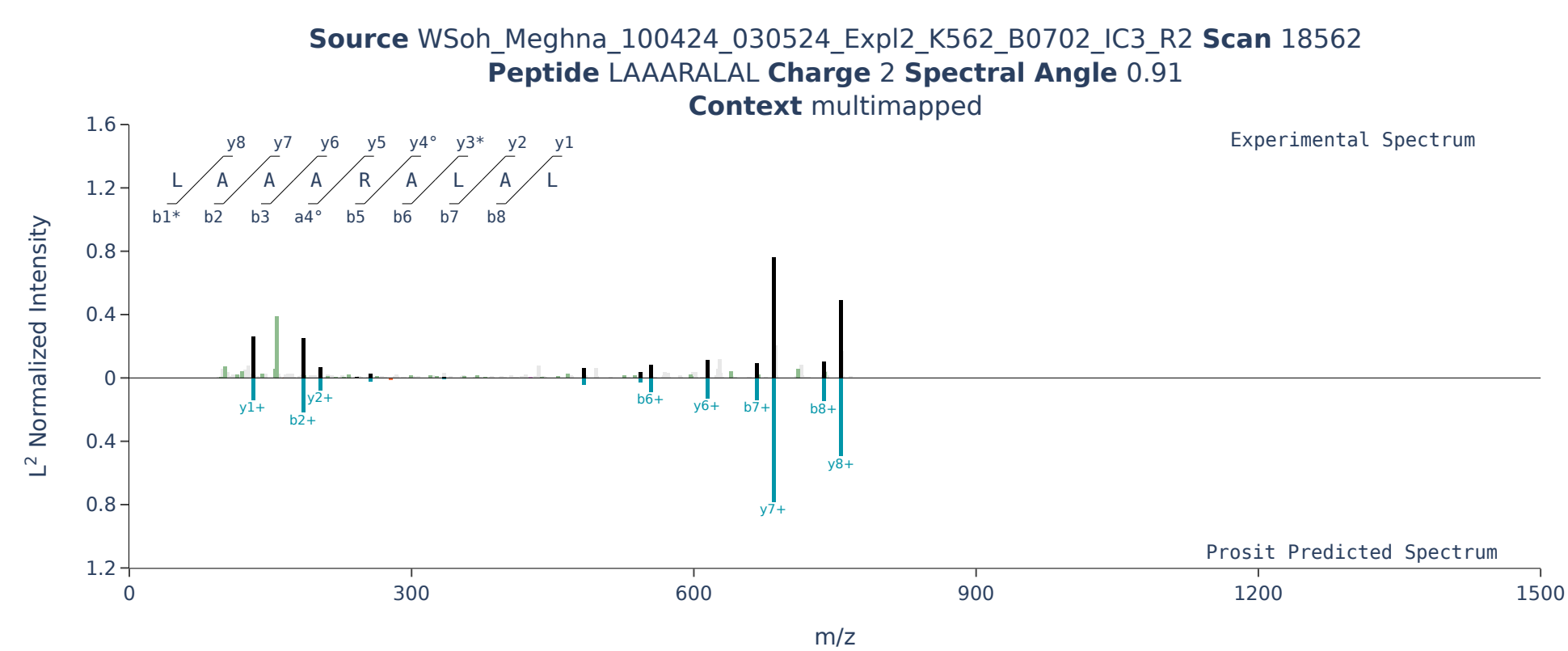
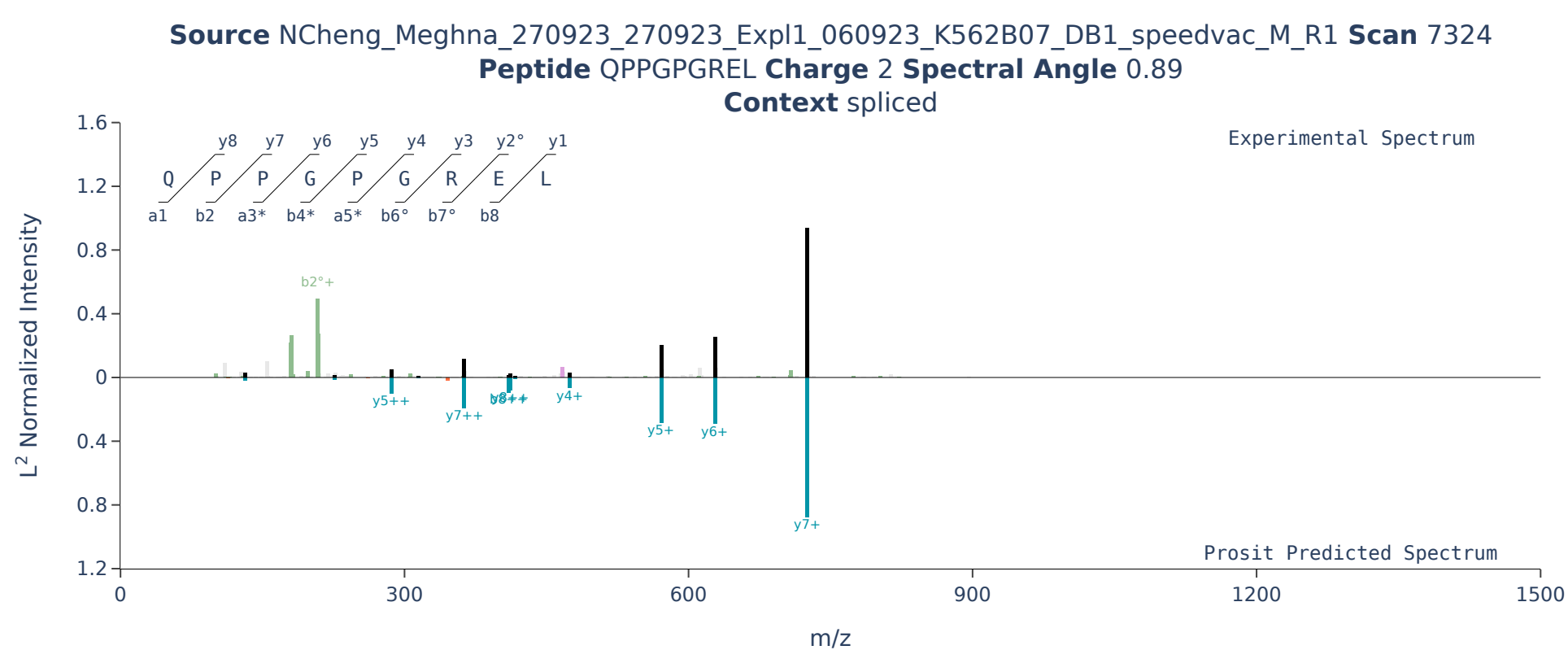
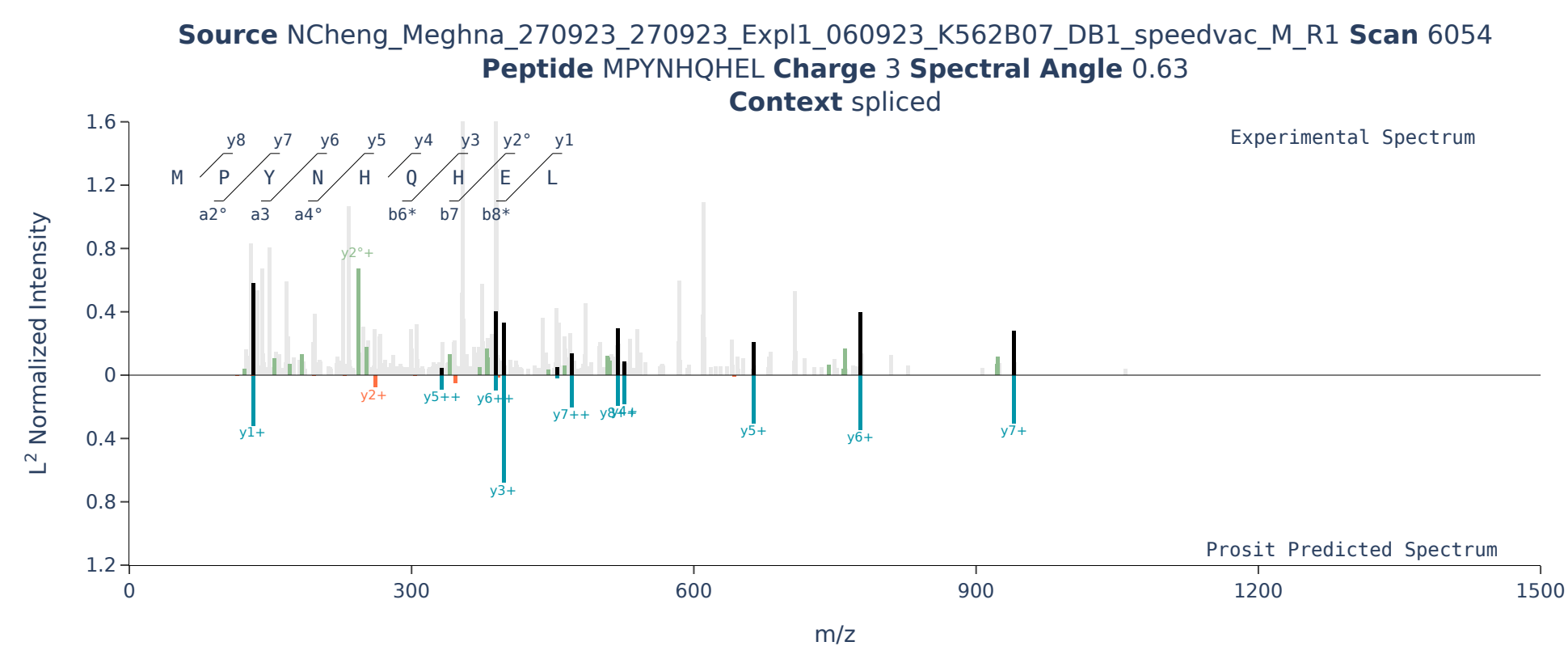
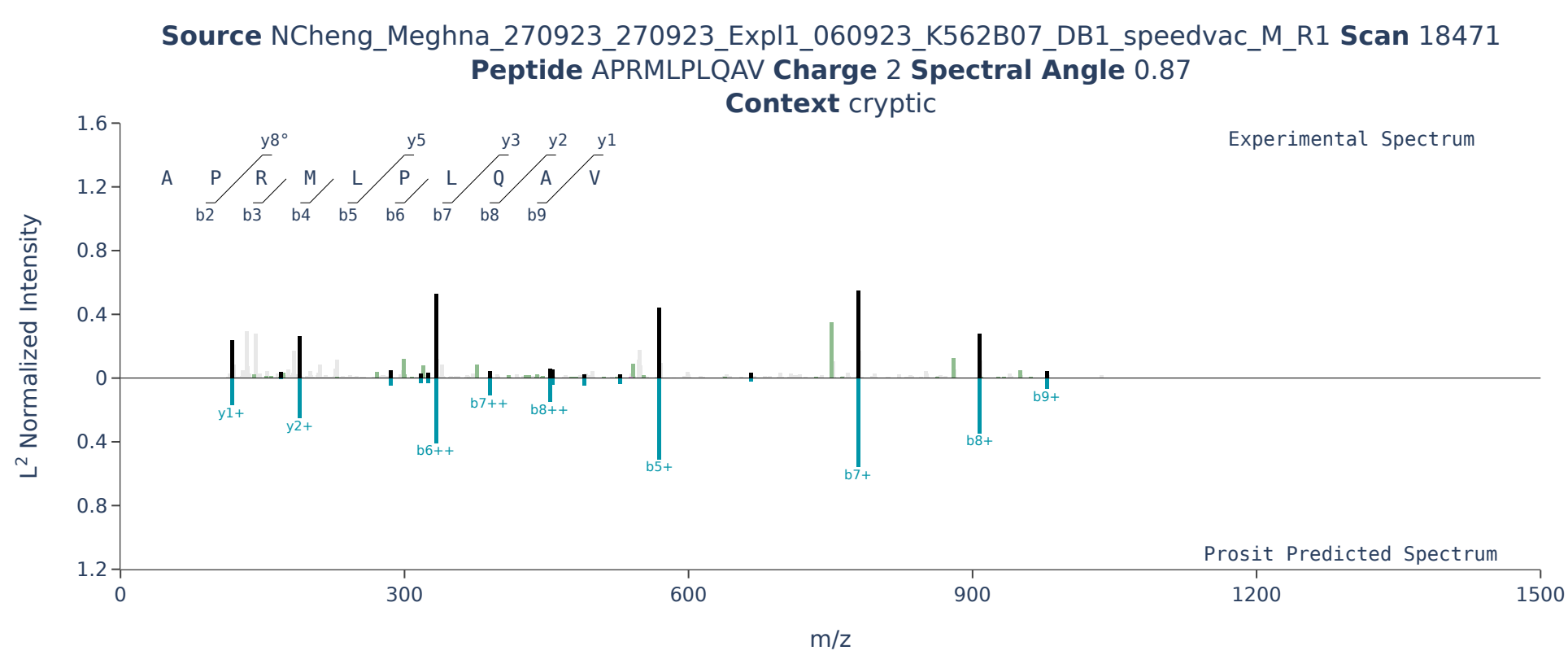
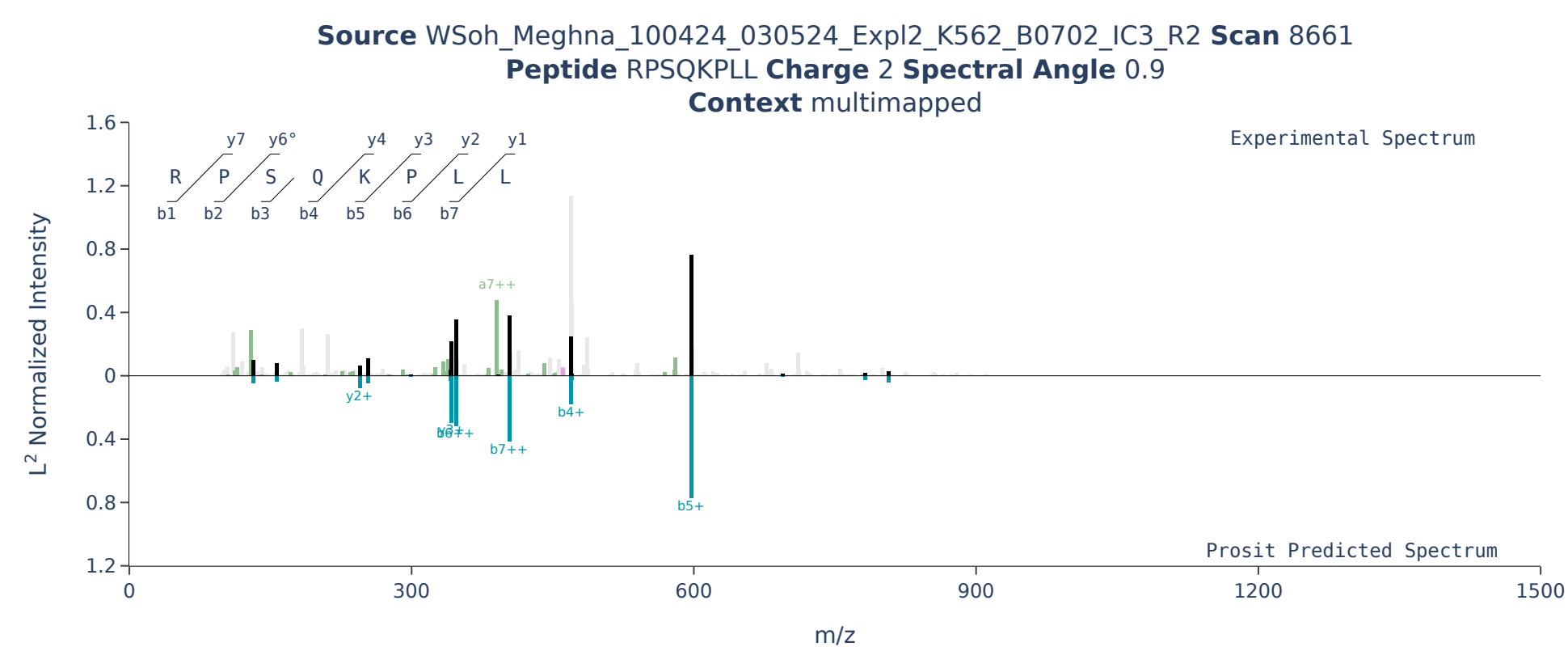
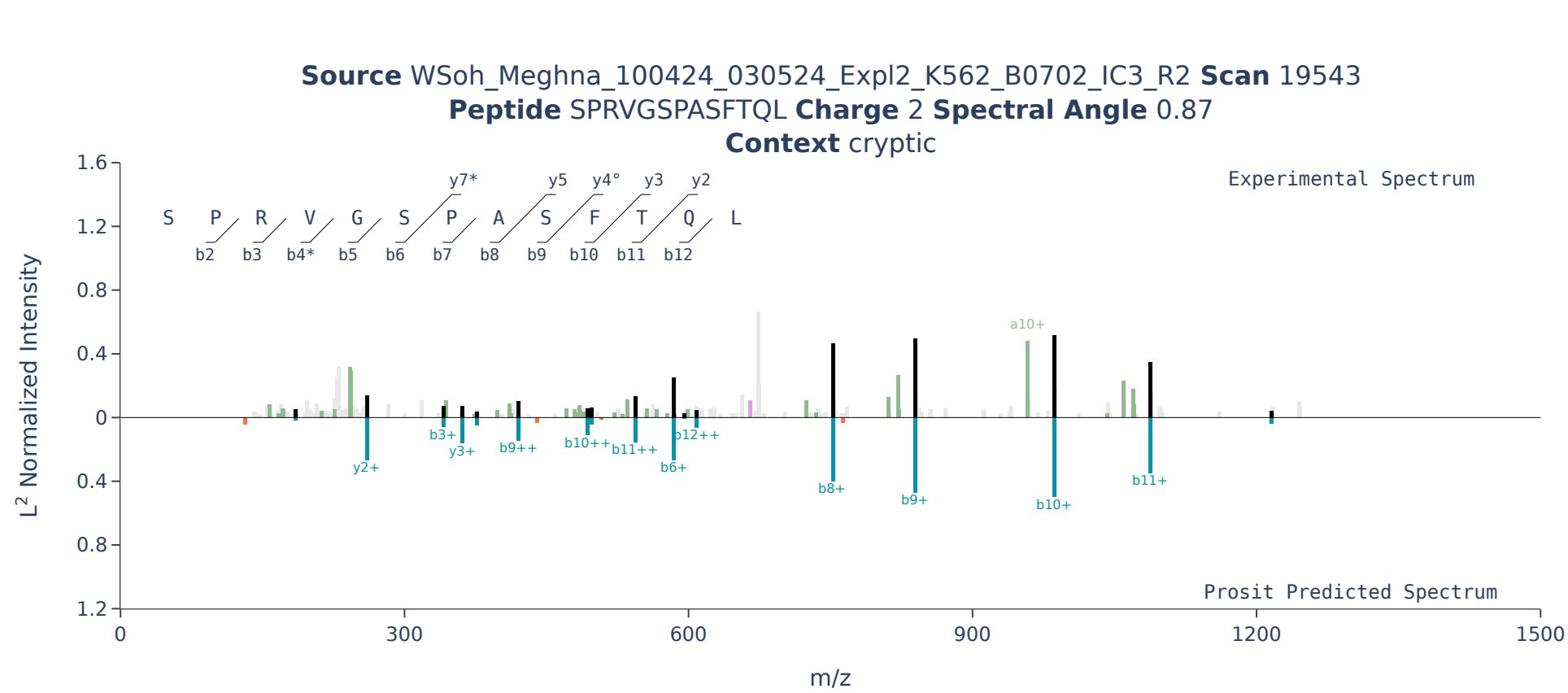
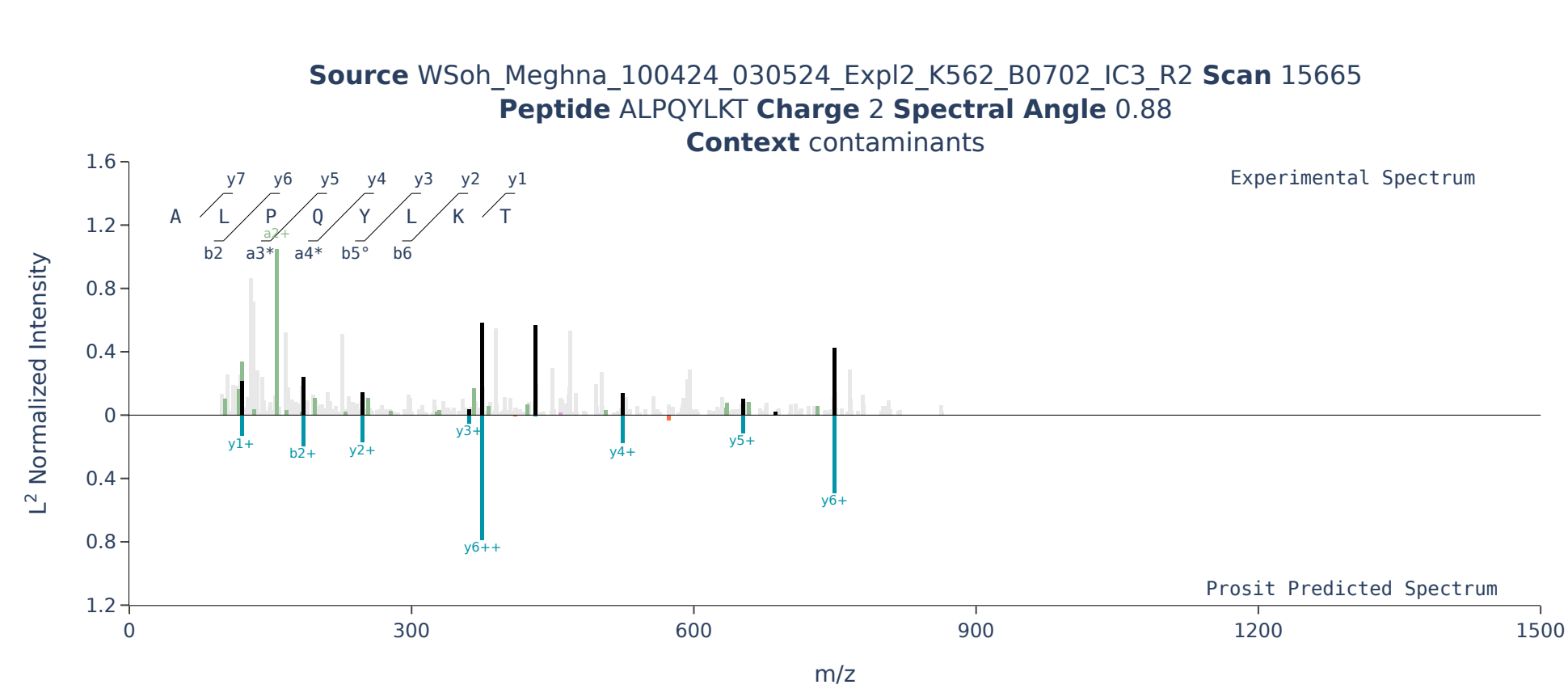


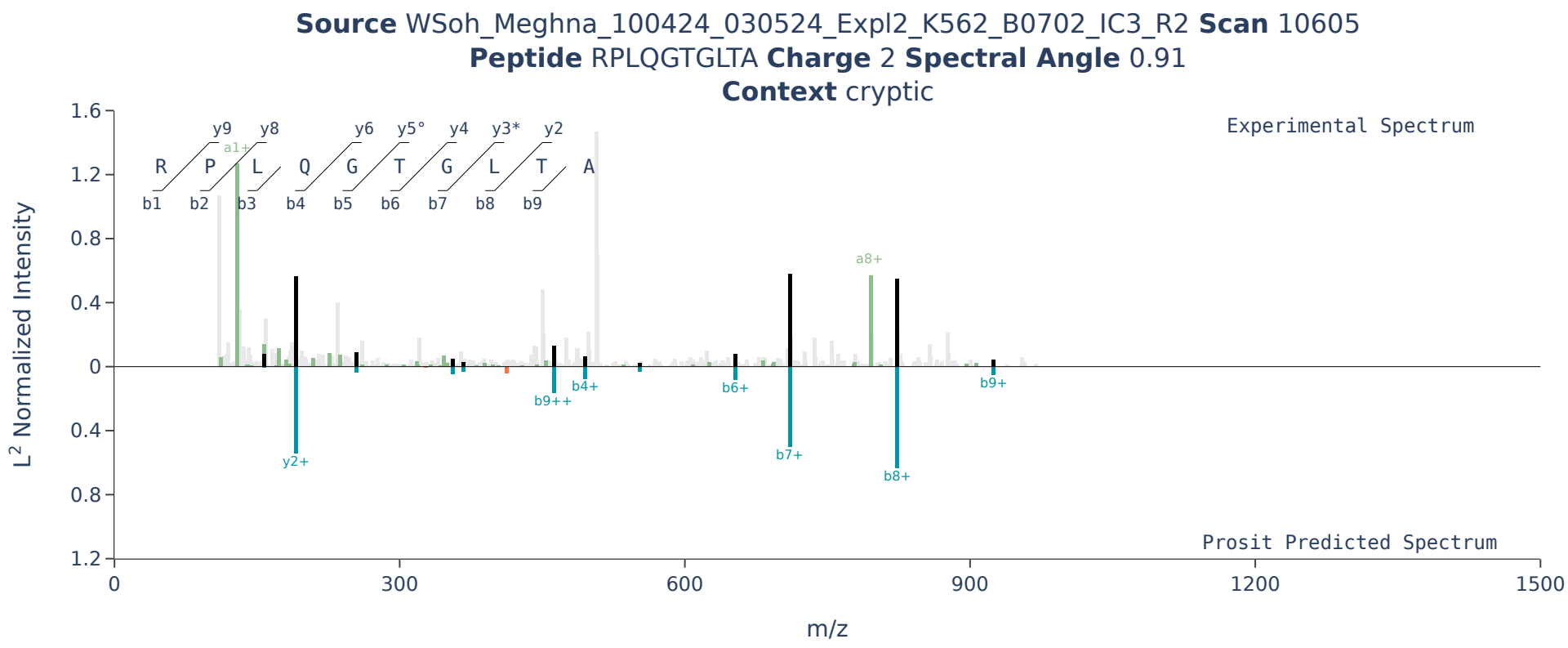
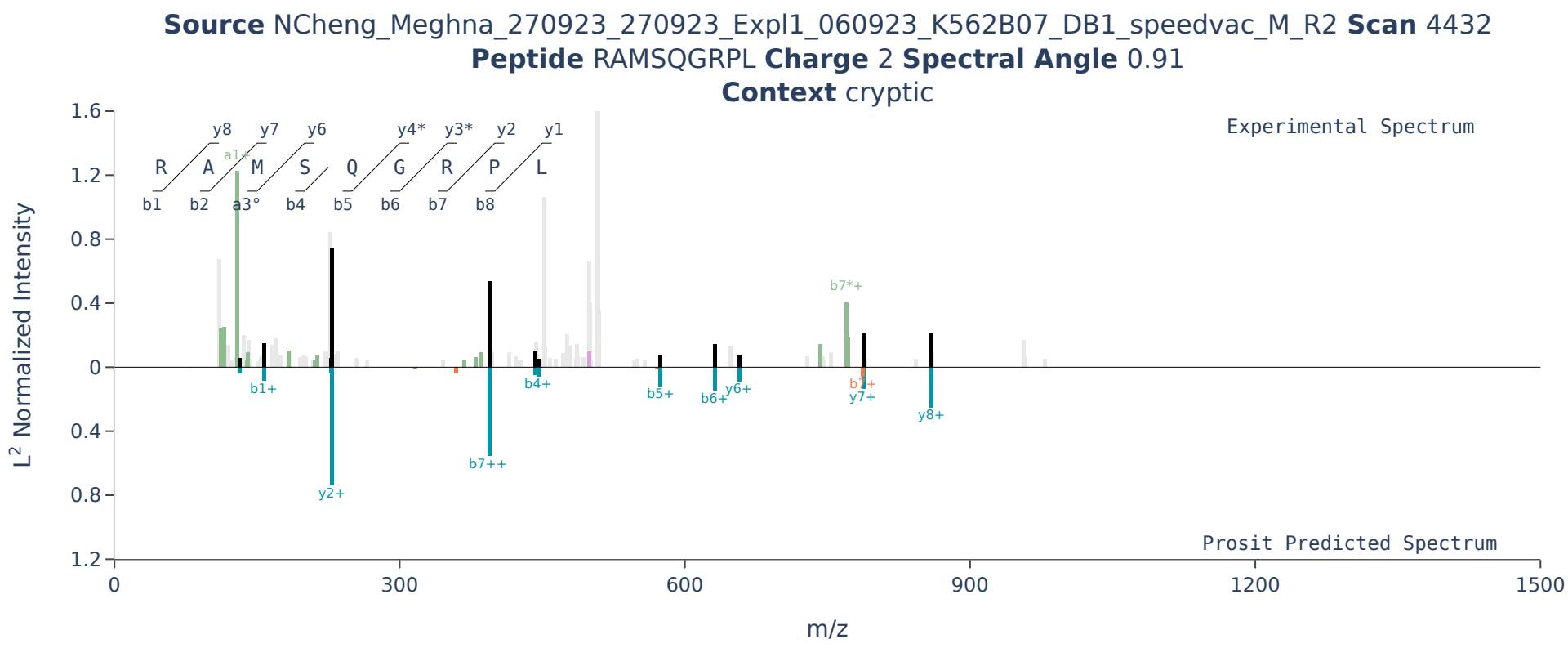
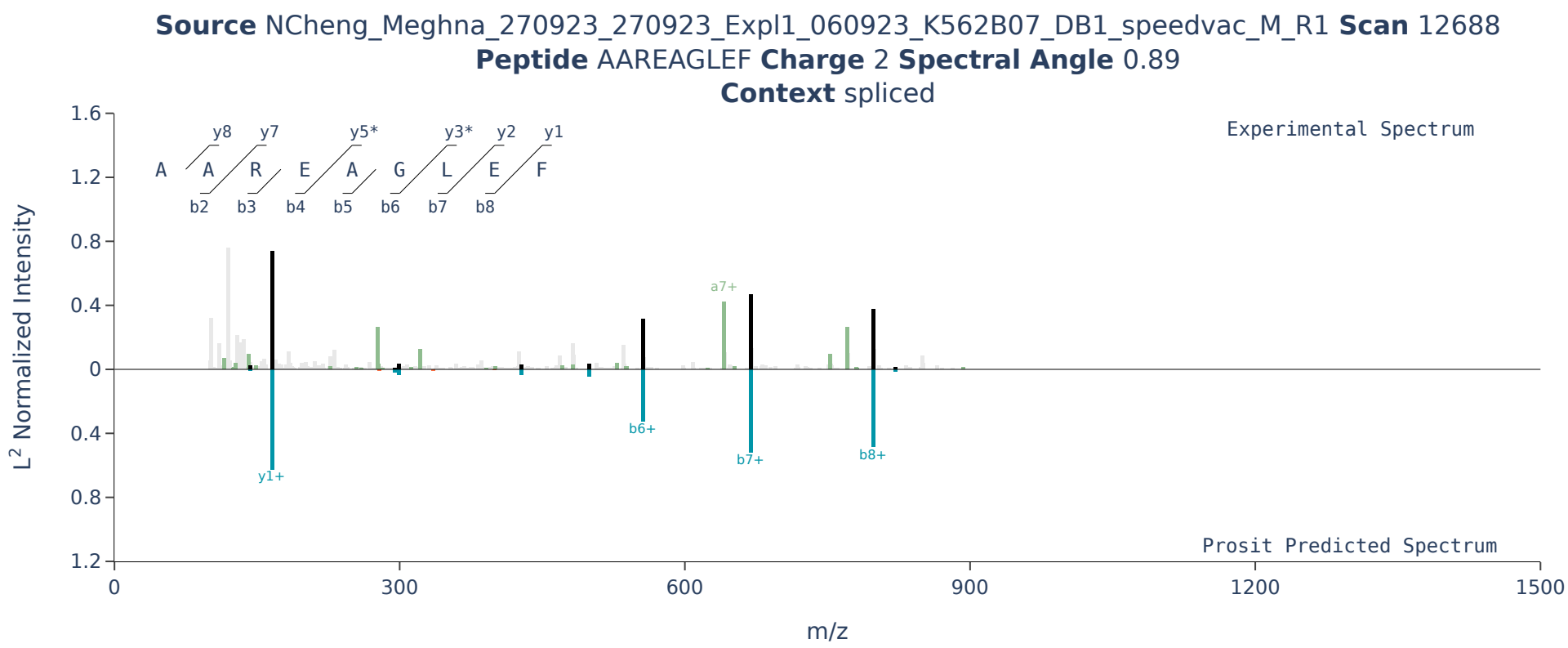
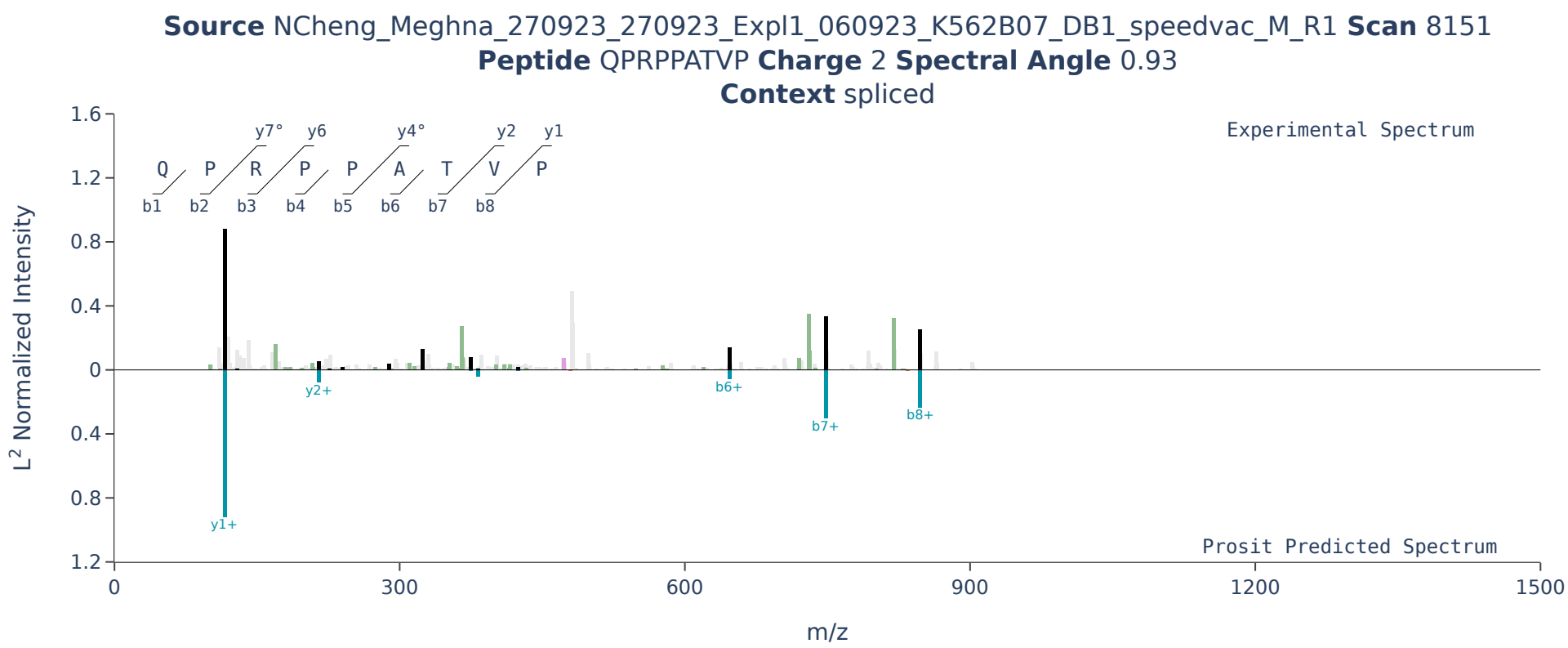
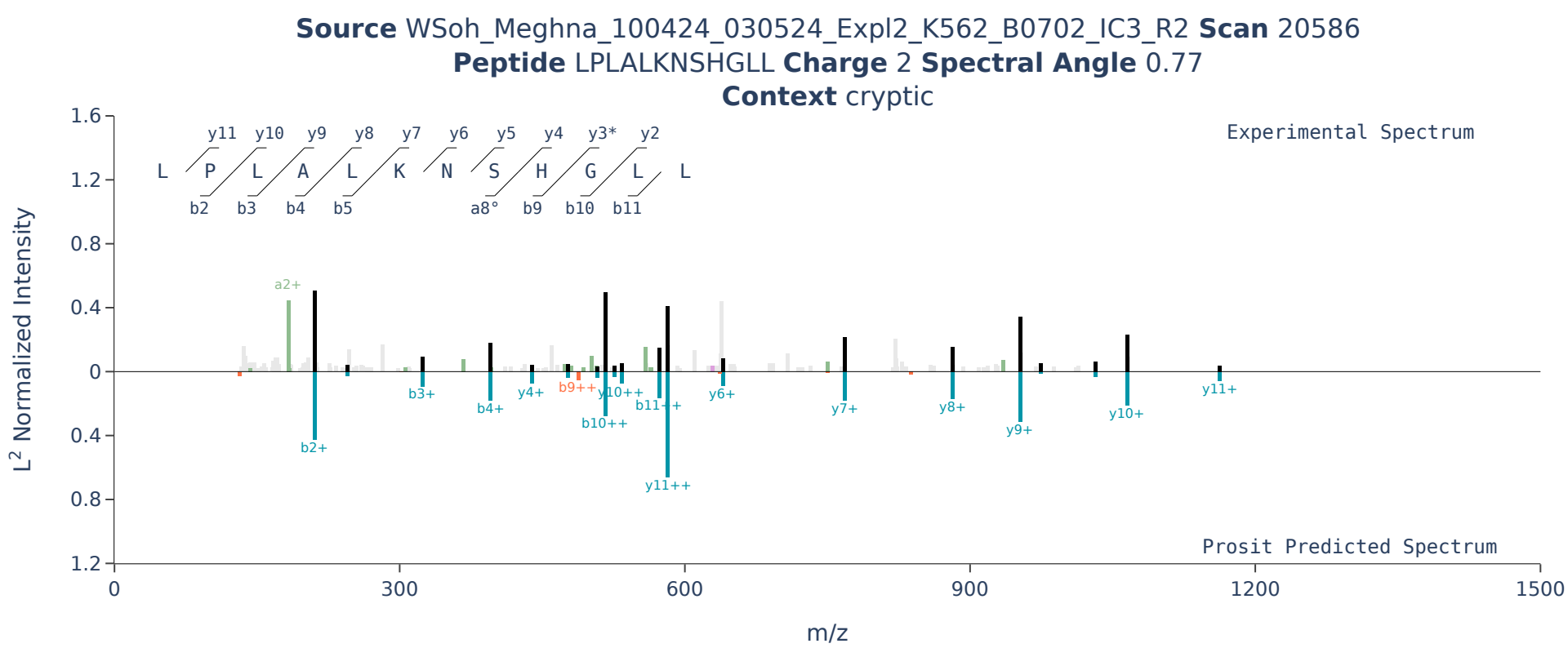


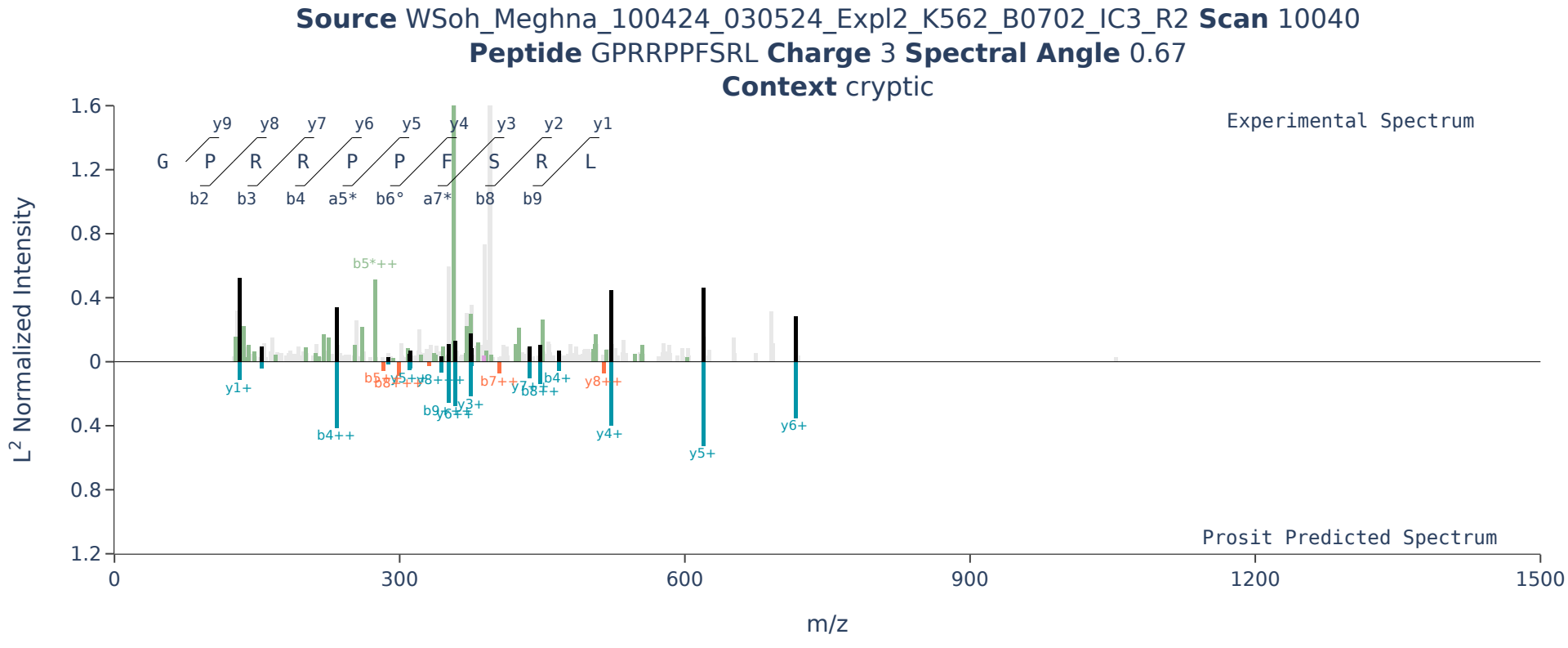
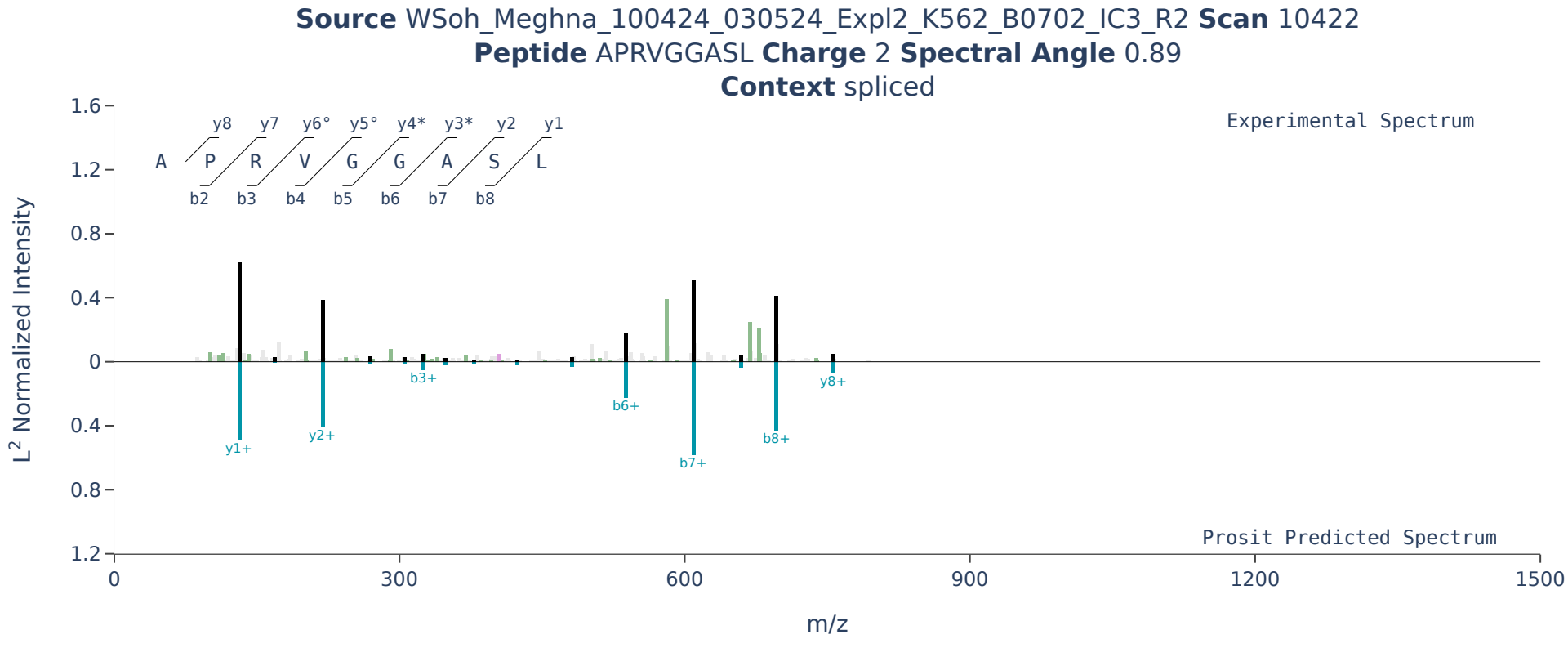
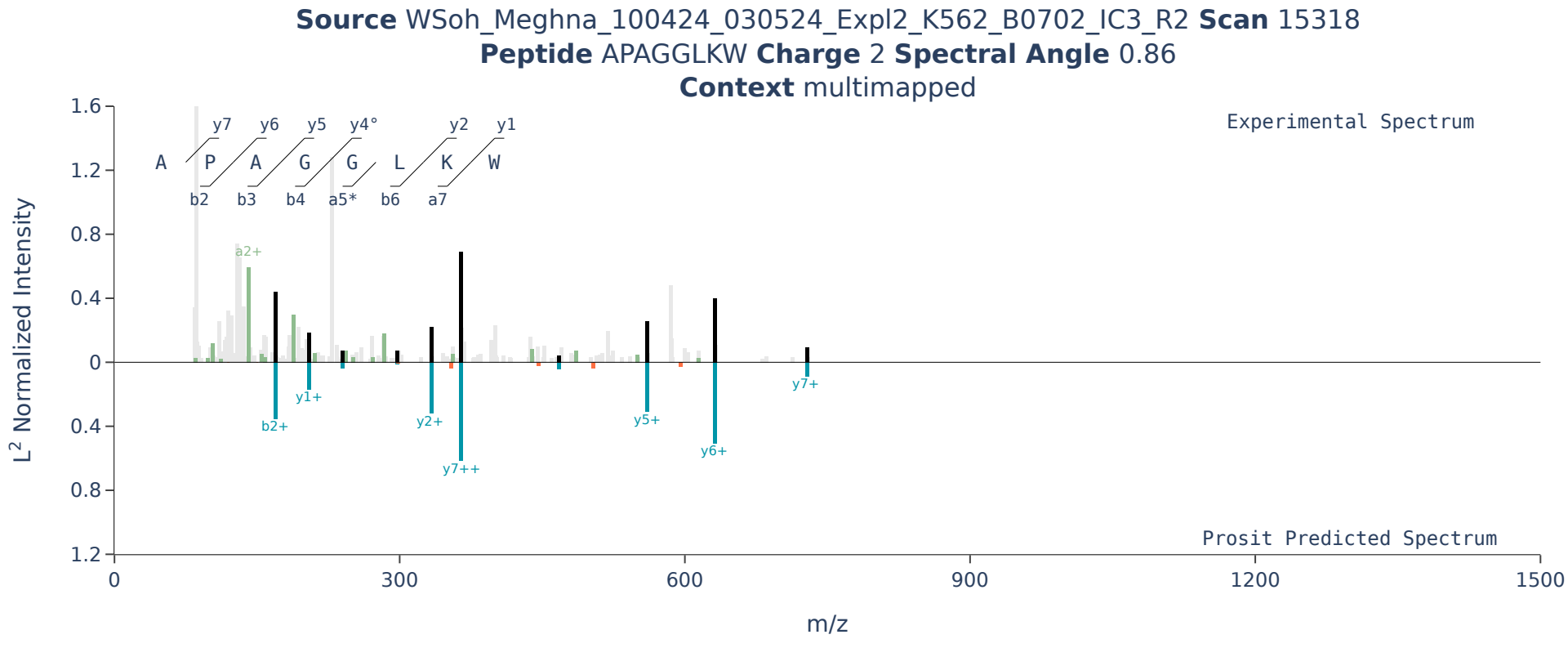
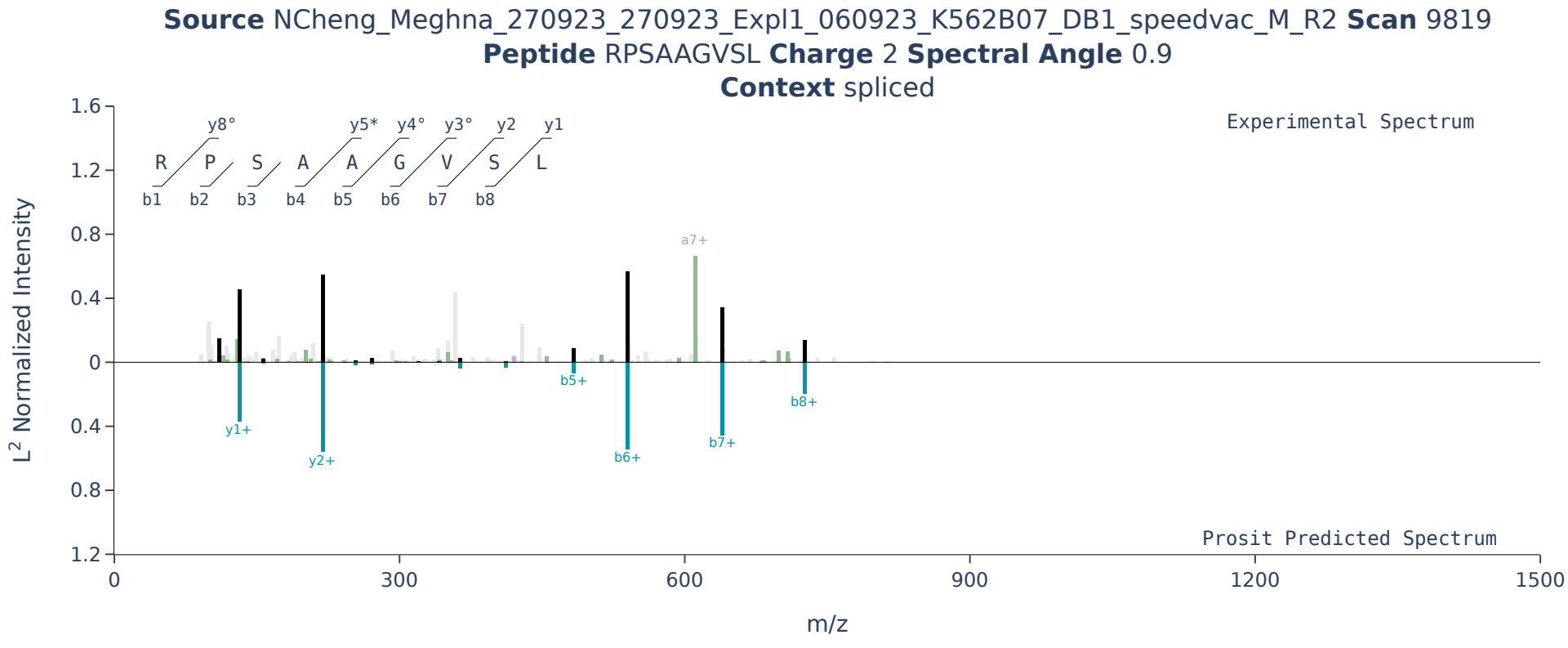
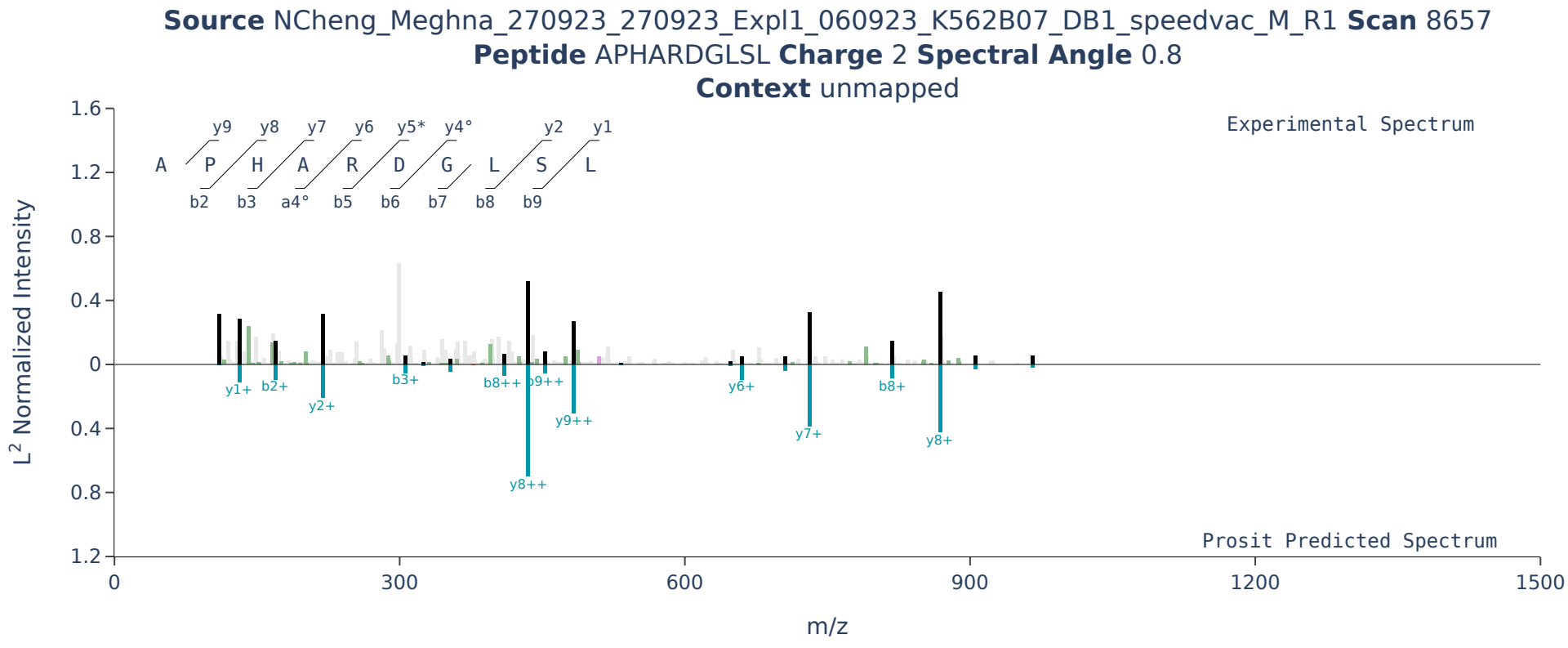
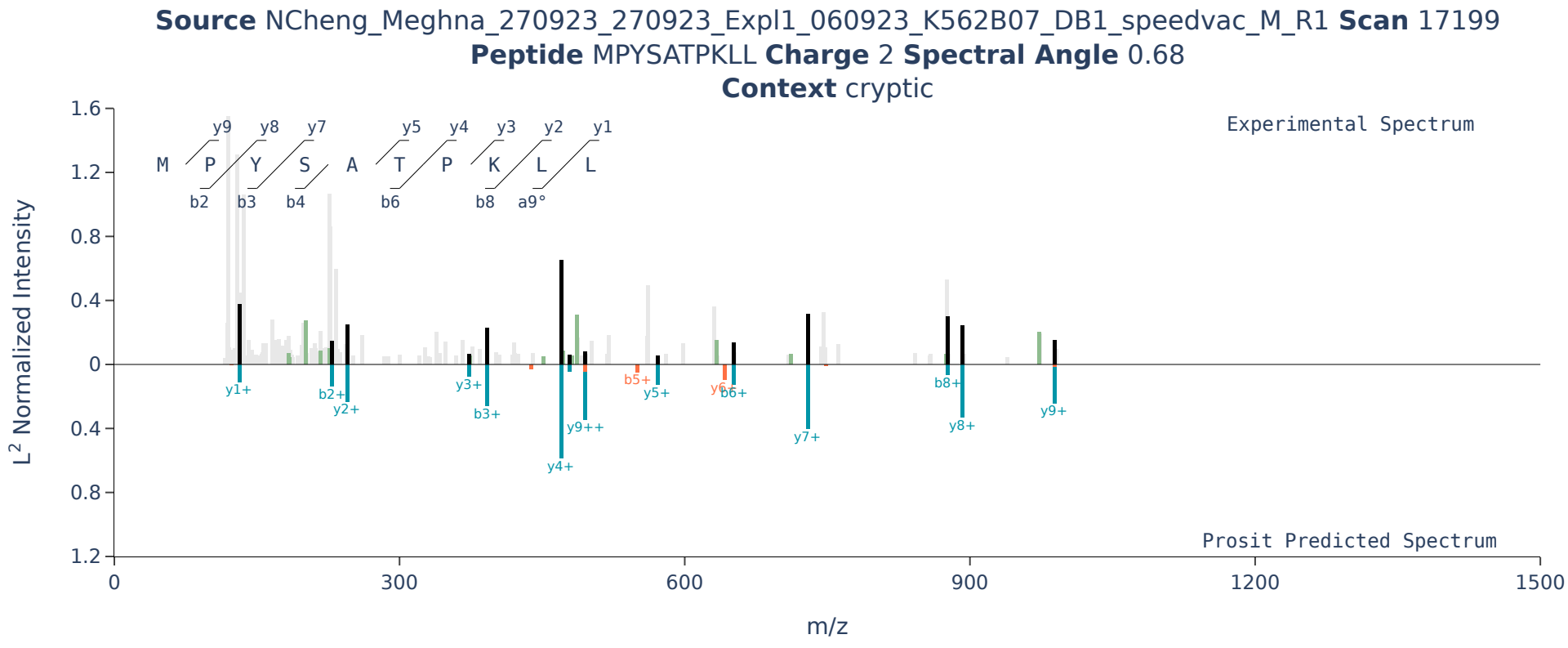
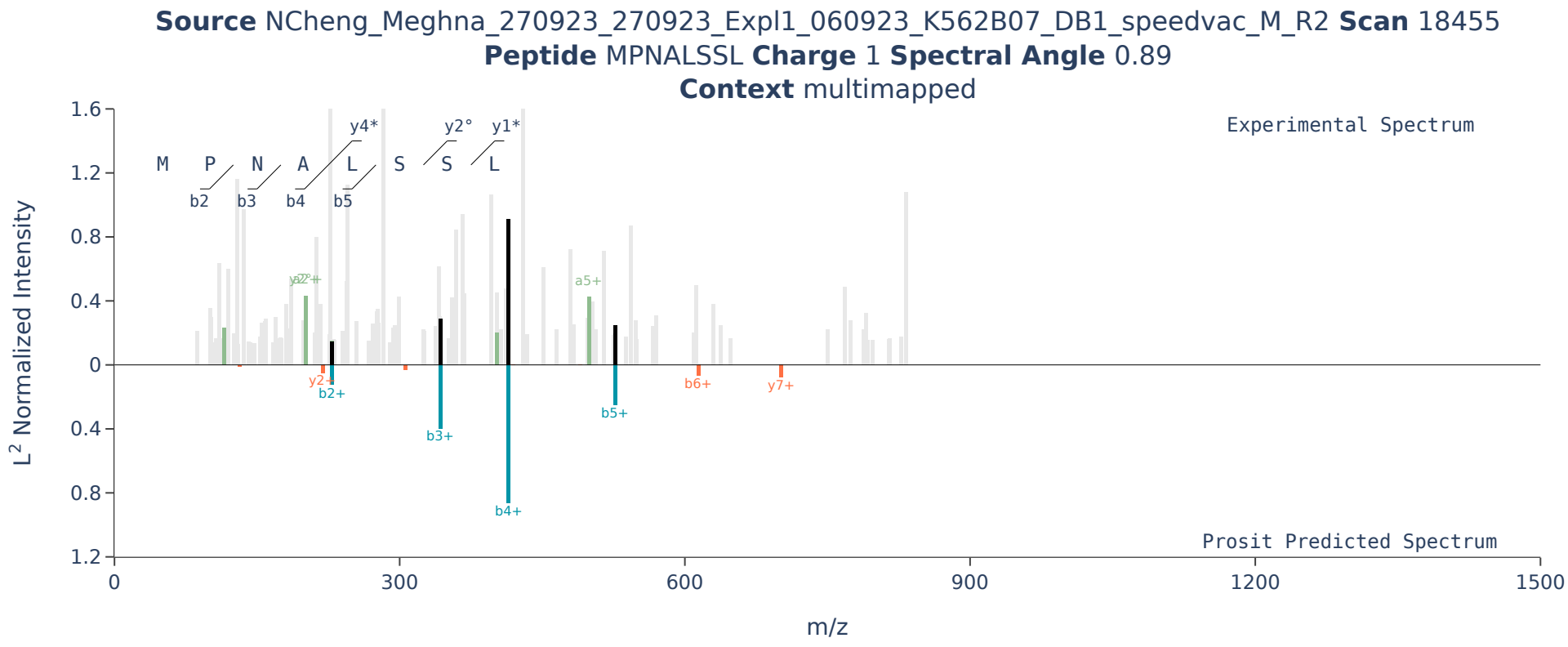
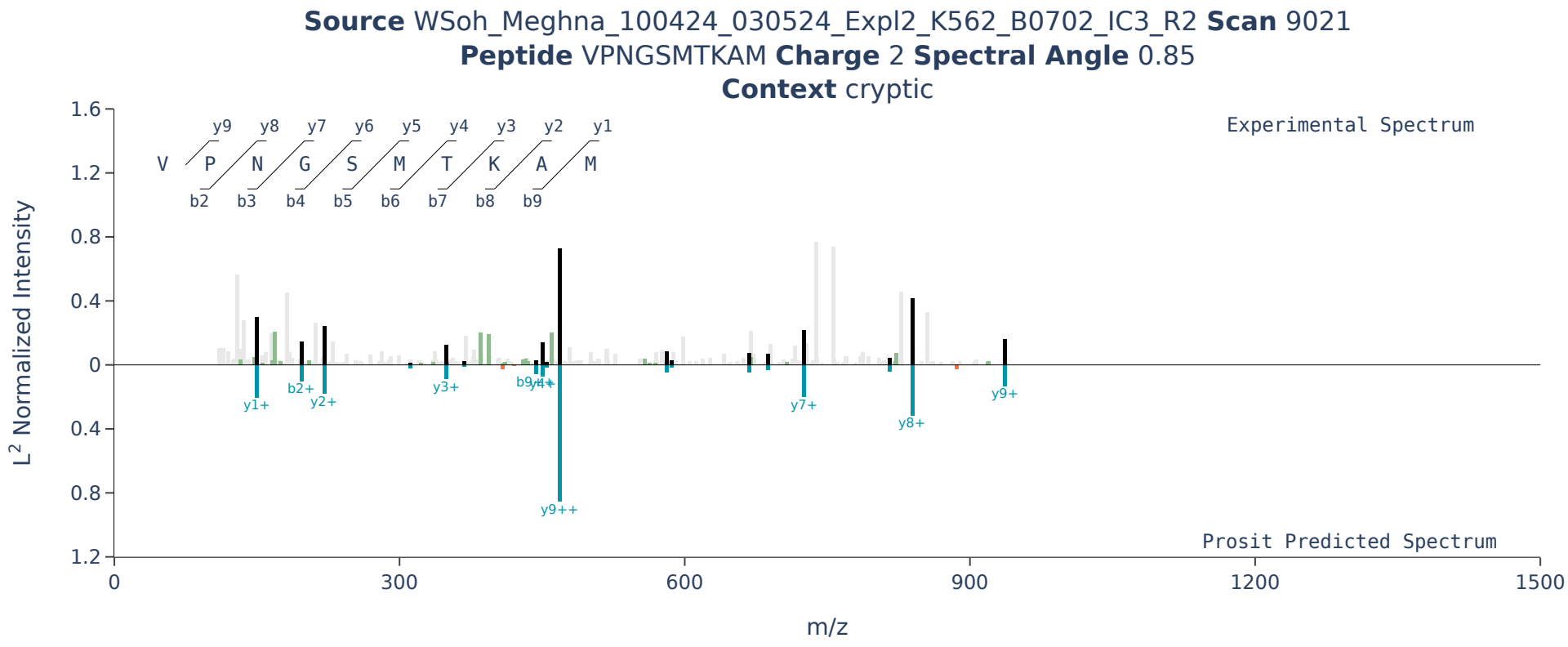
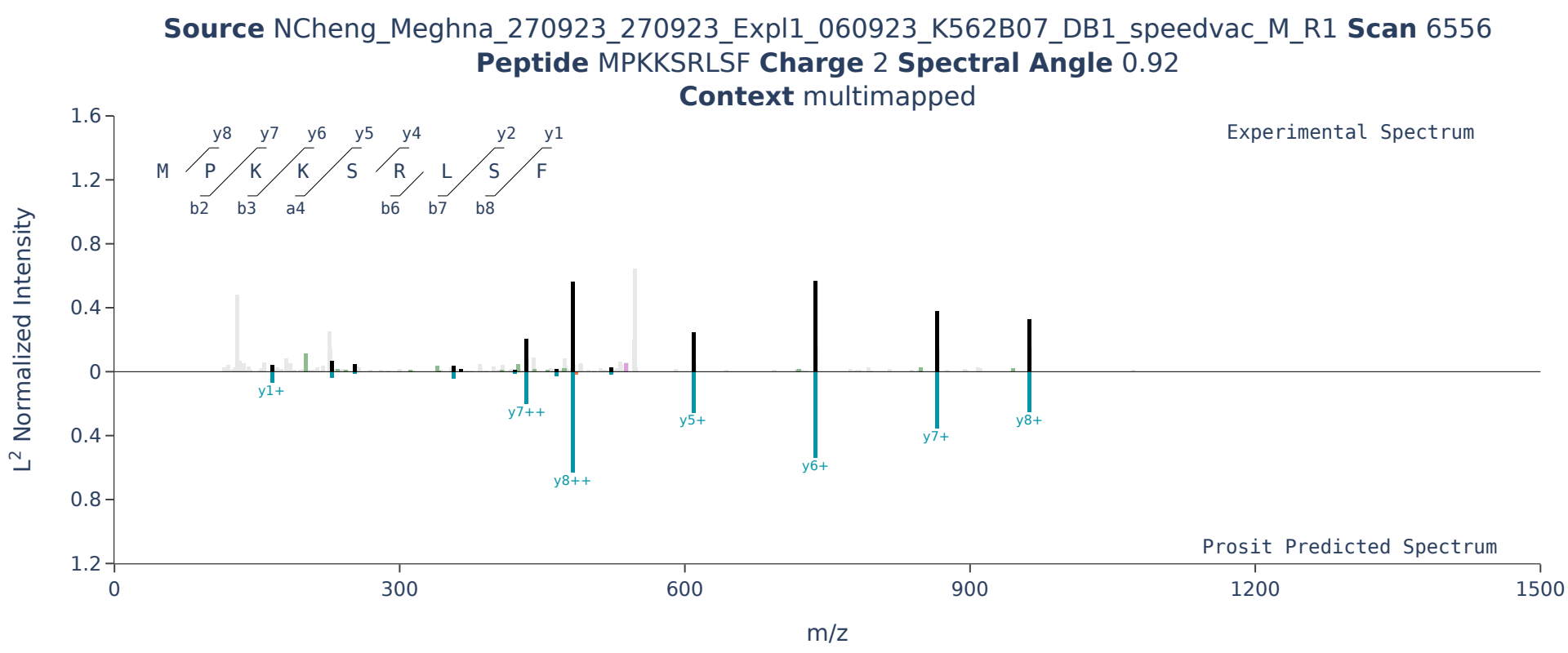
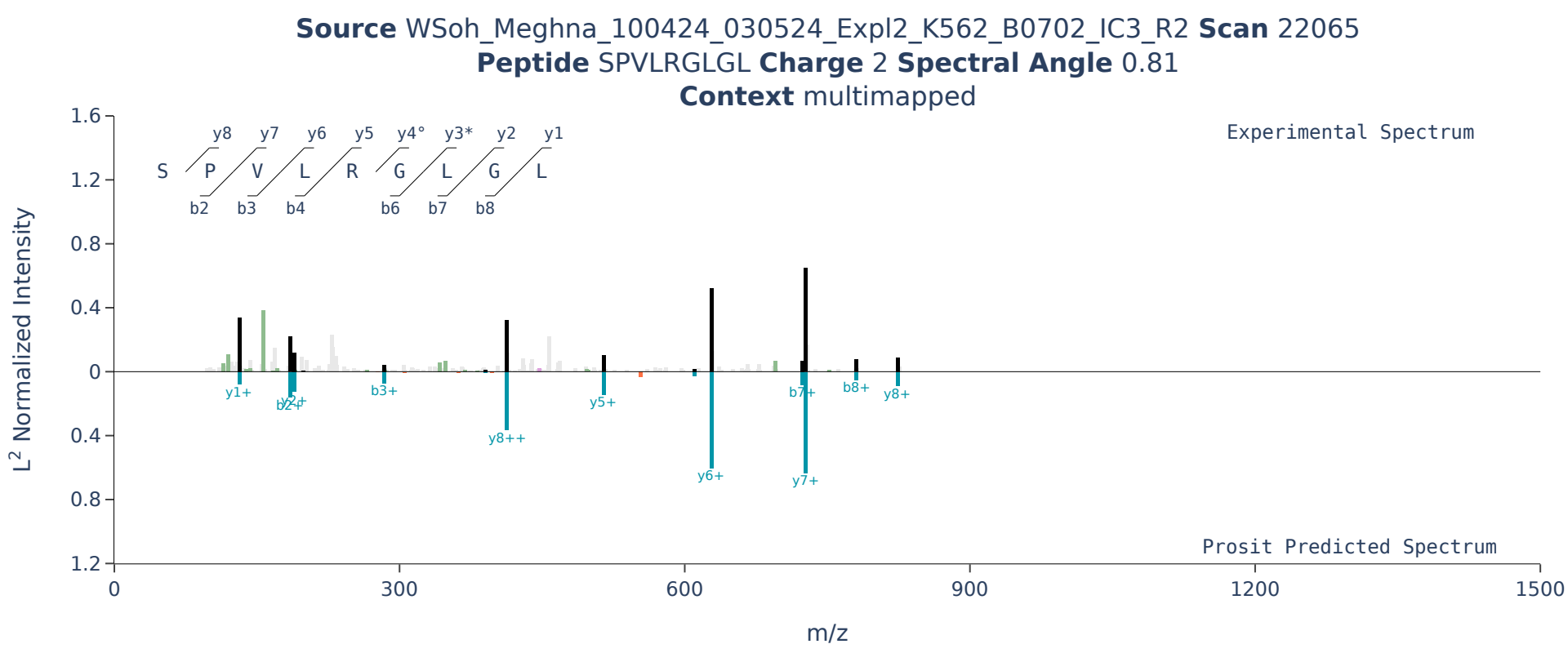




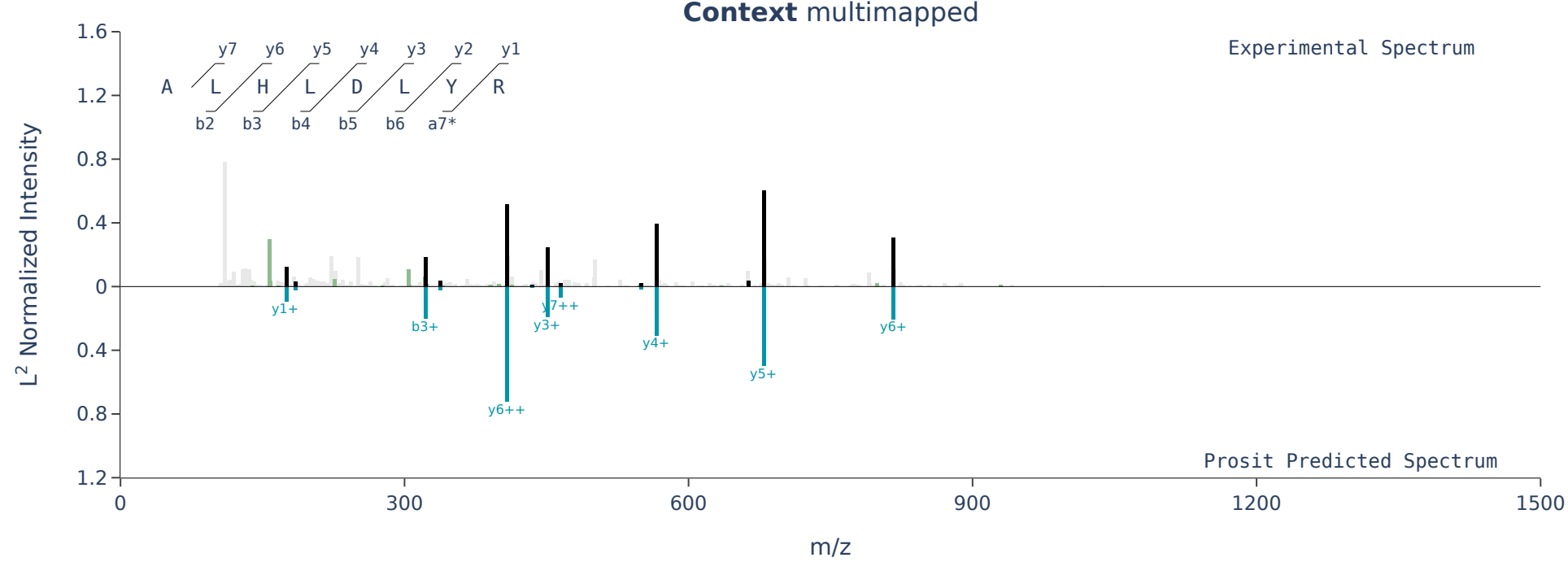




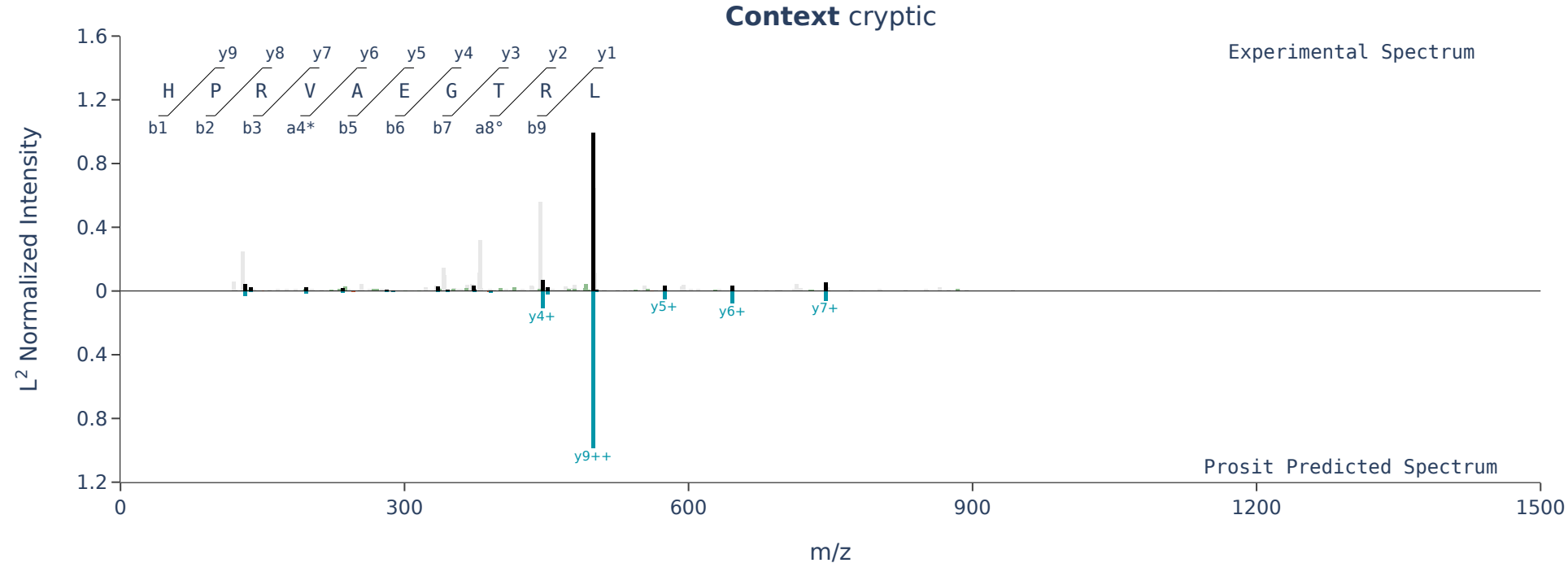




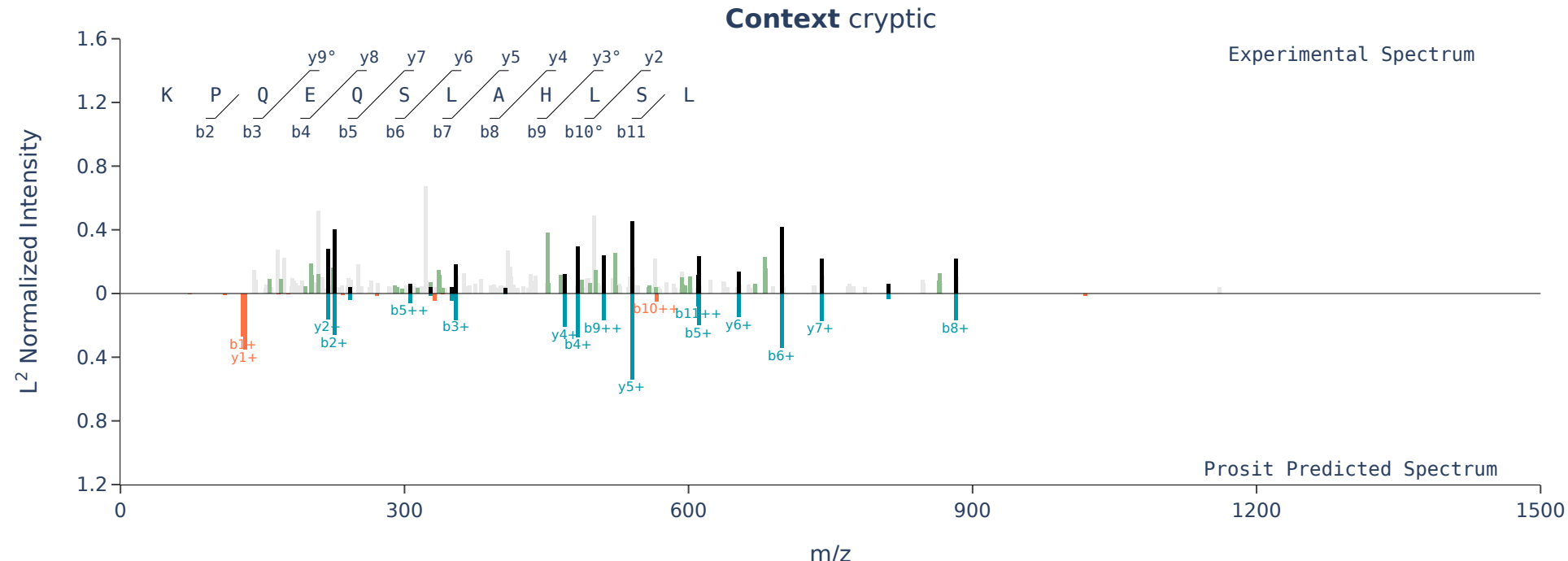
Source NCheng_Meghna_270923_270923_Expl1_060923_K562B07_DB1_speedvac_M_R1 **Scan** 13503
Peptide ALHLDLYR **Charge** 2 **Spectral Angle** 0.82



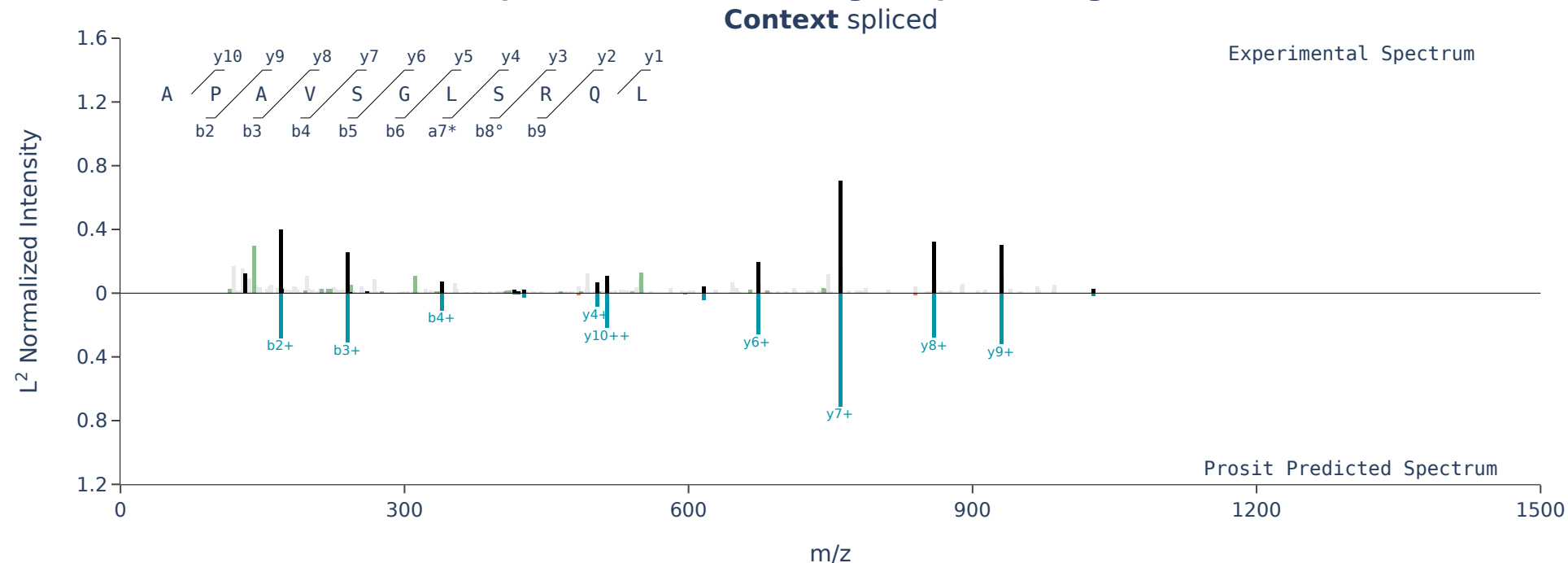
Source WSoh_Meghna_100424_030524_Expl2_K562_B0702_IC3_R2 **Scan** 6364
Peptide HPRVAEGTRL **Charge** 3 **Spectral Angle** 0.96



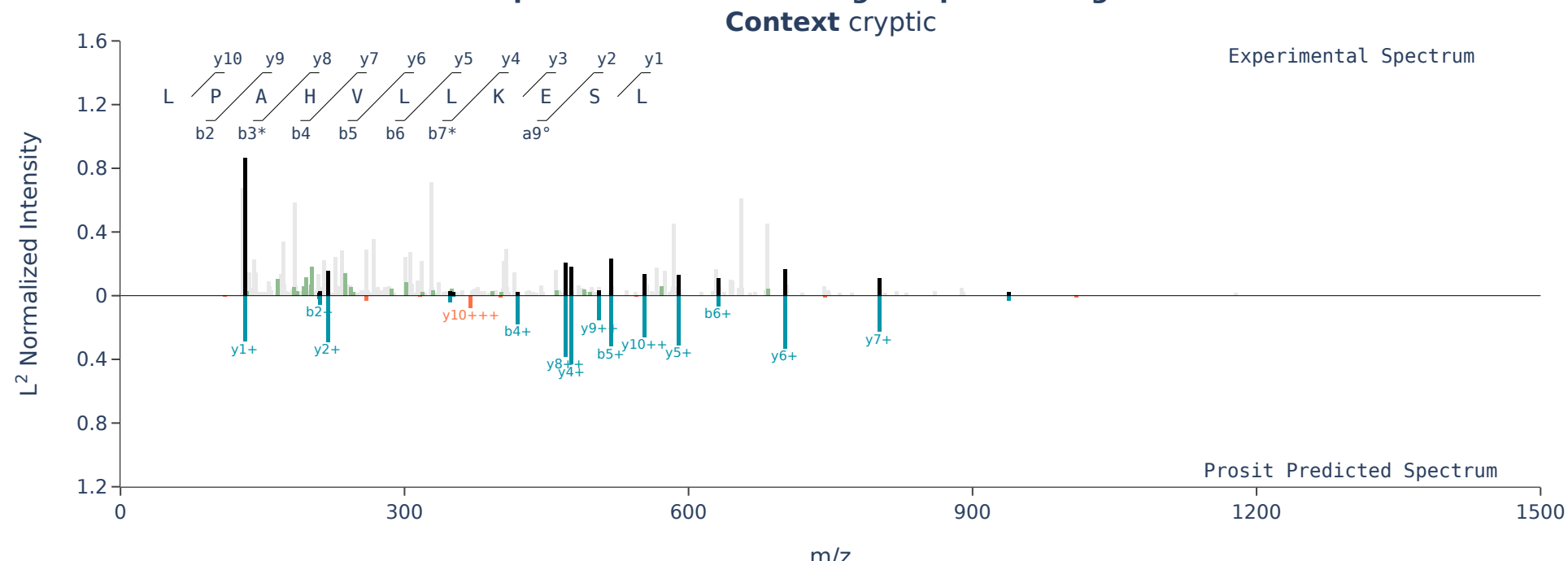
Source WSoh_Meghna_100424_030524_Exp12_K562_B0702_IC3_R2 **Scan** 17129
Peptide KPQEQSLAHL **Charge** 3 **Spectral Angle** 0.66



Source WSoh_Meghna_100424_030524_Exp12_K562_B0702_IC3_R2 **Scan** 15718
Peptide APAVSGLSRQL **Charge** 2 **Spectral Angle** 0.88



Source NCheng_Meghna_270923_270923_Expl1_060923_K562B07_DB1_speedvac_M_R2 **Scan** 14912
Peptide LPAHVLLKESL **Charge** 3 **Spectral Angle** 0.5



Peptide EPRPPHGKEL **Charge** 3 **Spectral Angle** 0.7



Peptide RVAASVQRL Charge 2 Spectral Angle 0.79



Peptide RHPEYAVSVLLR Charge 3 Spectral Angle 0.84



Peptide SQPLASLYVK Charge 2 Spectral Angle 0.86



Peptide RPNKPVLKP Charge 2 Spectral Angle 0.8



Peptide NPRQRLAL **Charge** 3 **Spectral Angle** 0.65



Peptide LFLQGAQGSPFLK Charge 2 Spectral Angle 0.93



Peptide APVRSHGLVL Charge 3 Spectral Angle 0.89



Peptide AGAPGLPSL Charge 1 Spectral Angle 0.93

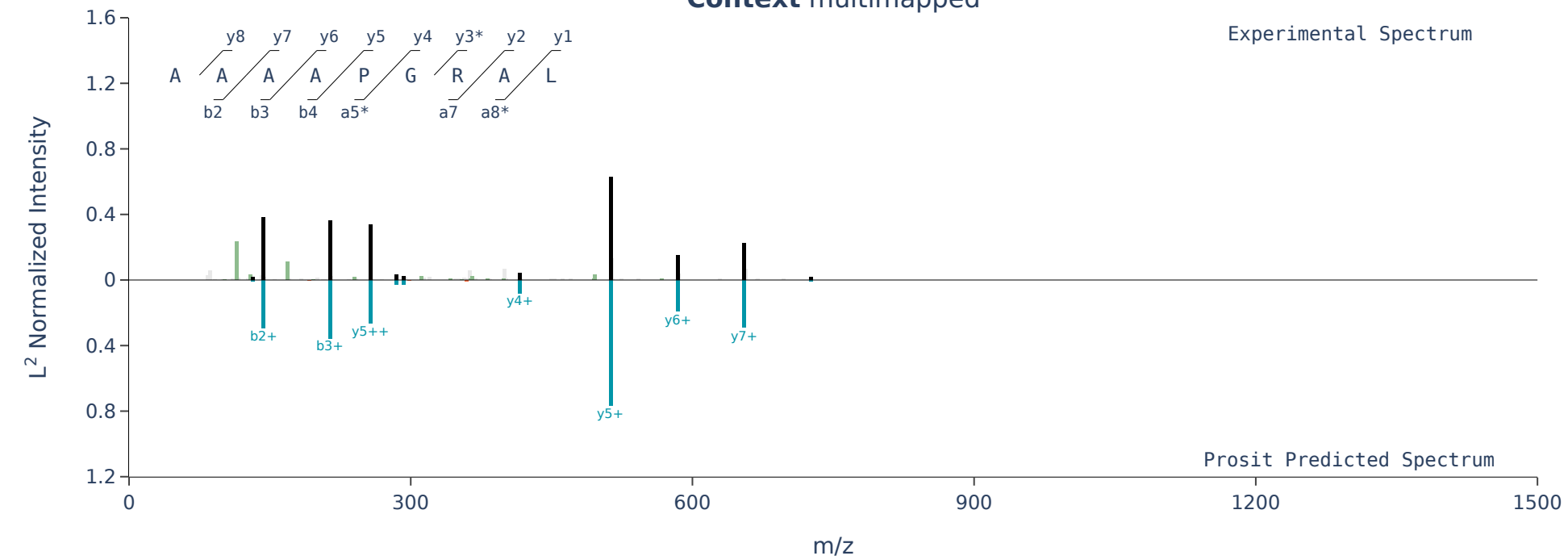


Peptide RQDEVRRAPL Charge 3 Spectral Angle 0.8



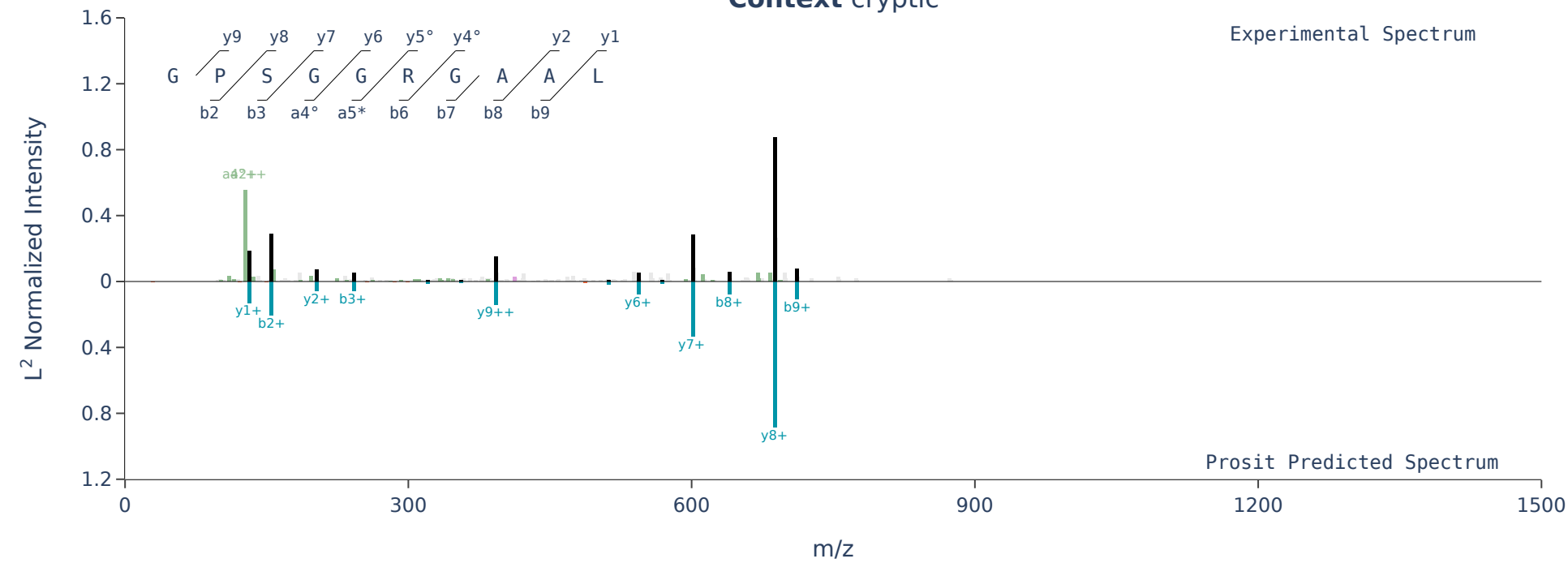
Peptide AAAAPGRAL **Charge** 2 **Spectral Angle** 0.88

Context multimapped



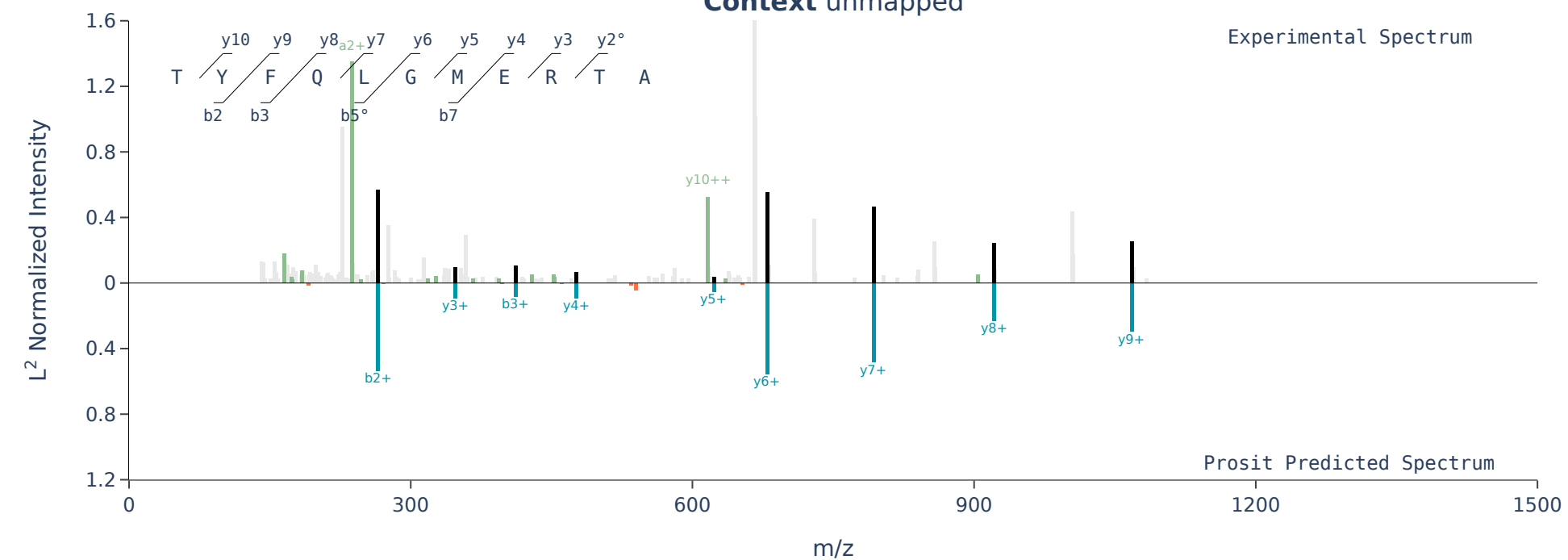
Peptide GPSGGRGAAL Charge 2 Spectral Angle 0.92

Context cryptic



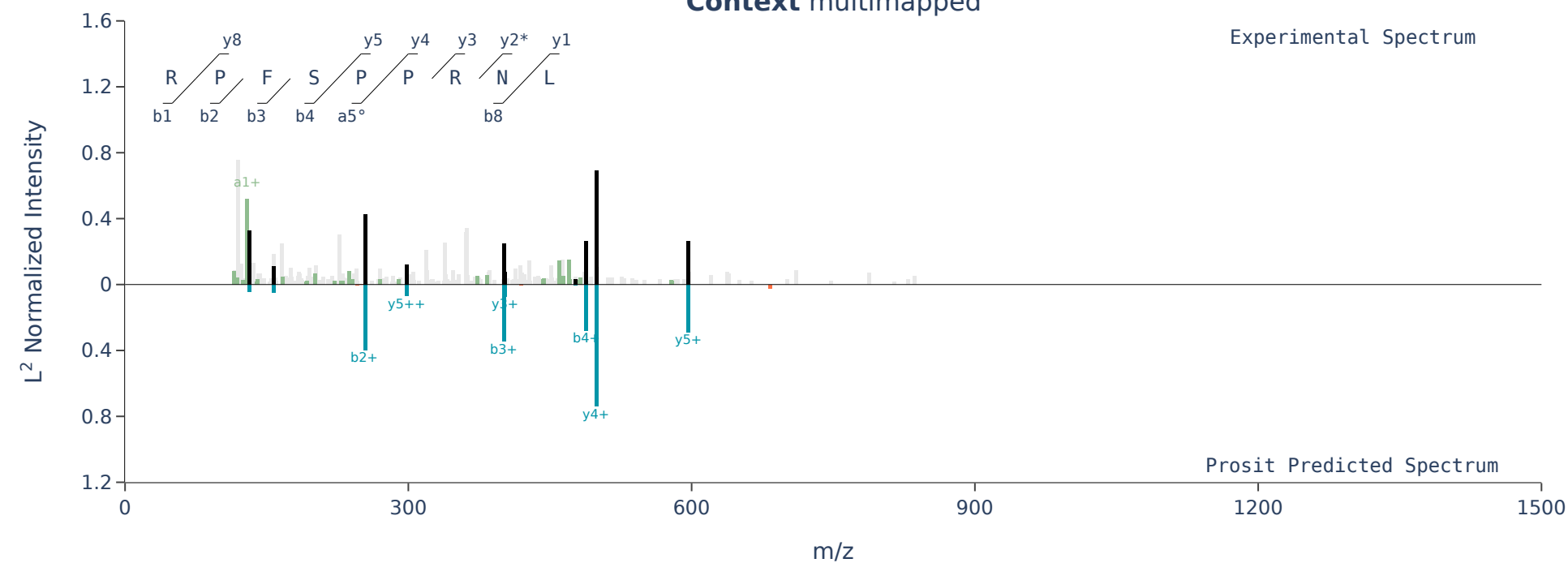
Peptide TYFQLGM[+16.0]ERTA **Charge** 2 **Spectral Angle** 0.94

Context unmapped



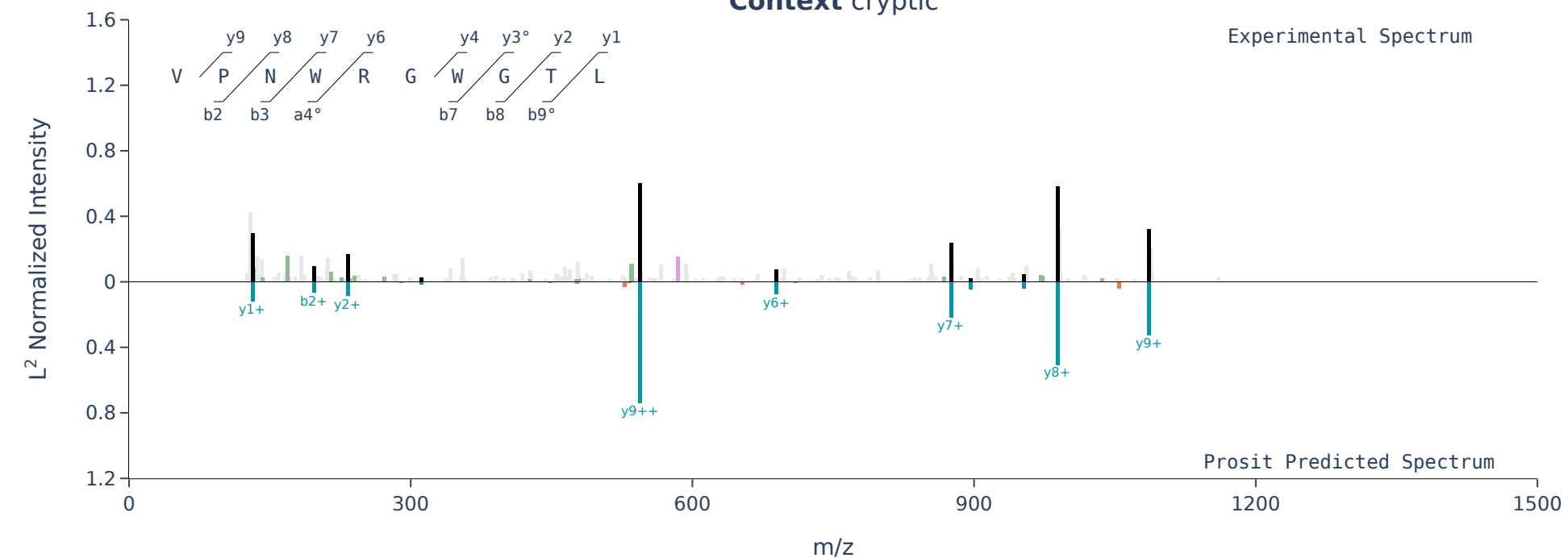
Peptide RPFSPPRNL Charge 3 Spectral Angle 0.8

Context multimapped



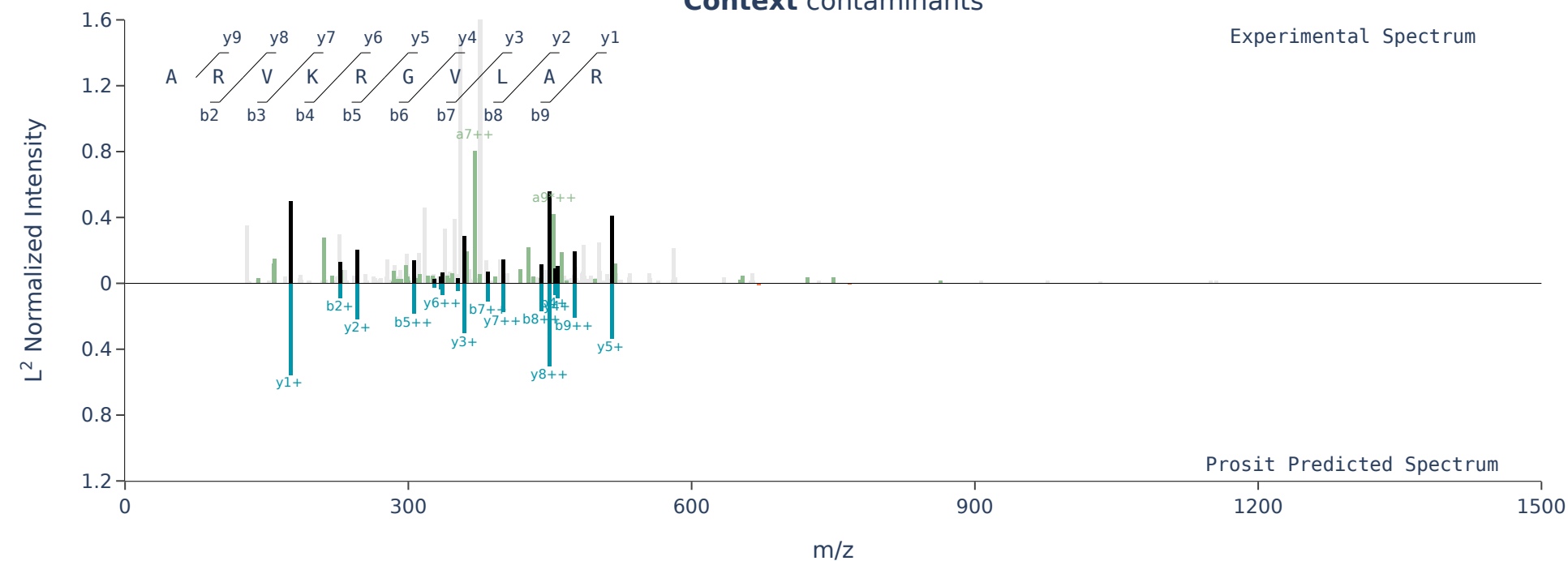
Peptide VPNWRGWGTL **Charge** 2 **Spectral Angle** 0.83

Context cryptic



Peptide ARVKRGVLAR Charge 3 Spectral Angle 0.9

Context contaminants



Peptide RPRPPQAVFL Charge 2 Spectral Angle 0.79

Context spliced

