

**Annotated peptide spectral matches of single peptide hits**

Translation ID

Peptide

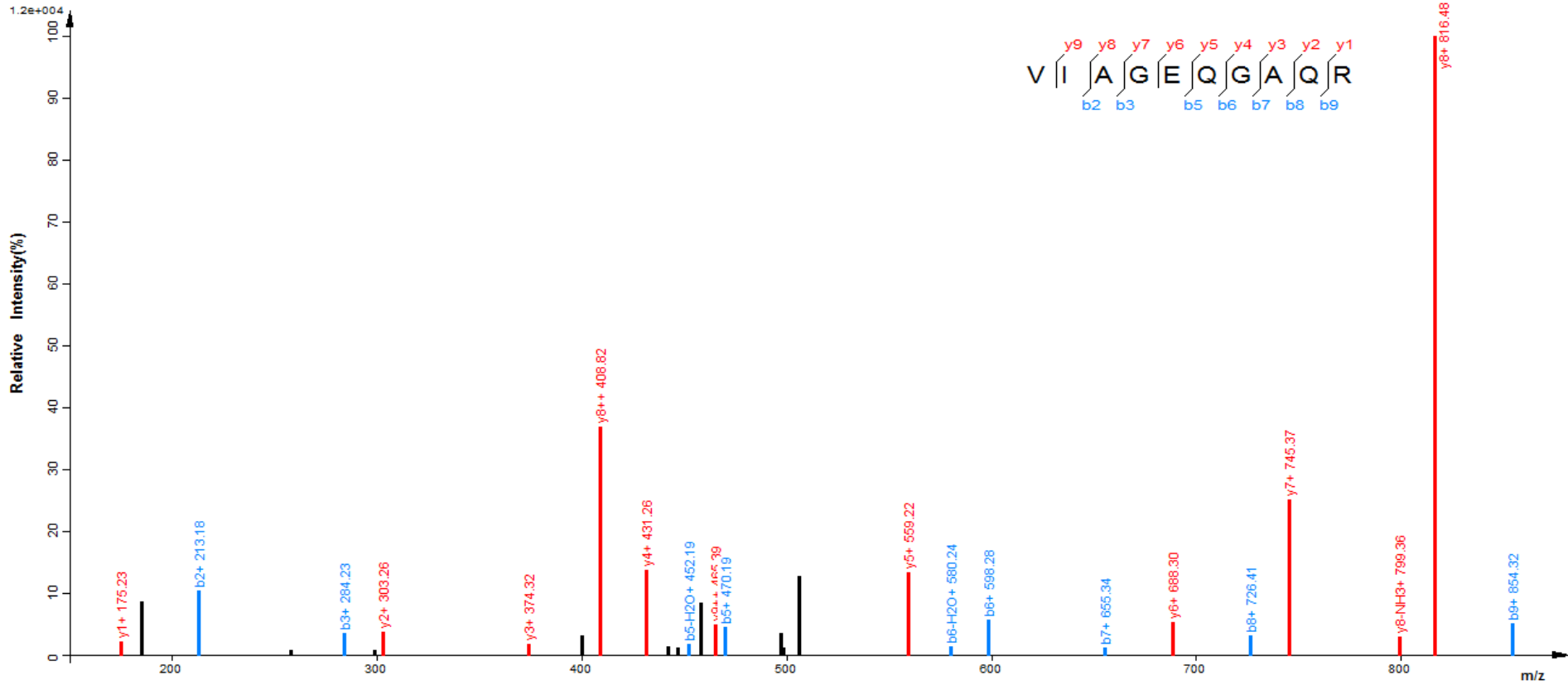
Total PSM

chrGI:27375111:ref:NC004463.1:\_4603653\_4603312\_-1

VIAGEQGAQR

23

E:\PROJECTS\Bradyrhizobium\_japonicum\INPUT\10115.mgf-63507 SEQ:VIAGEQGAQR Charge:2 MH+:1028.5483



Translation ID

Peptide

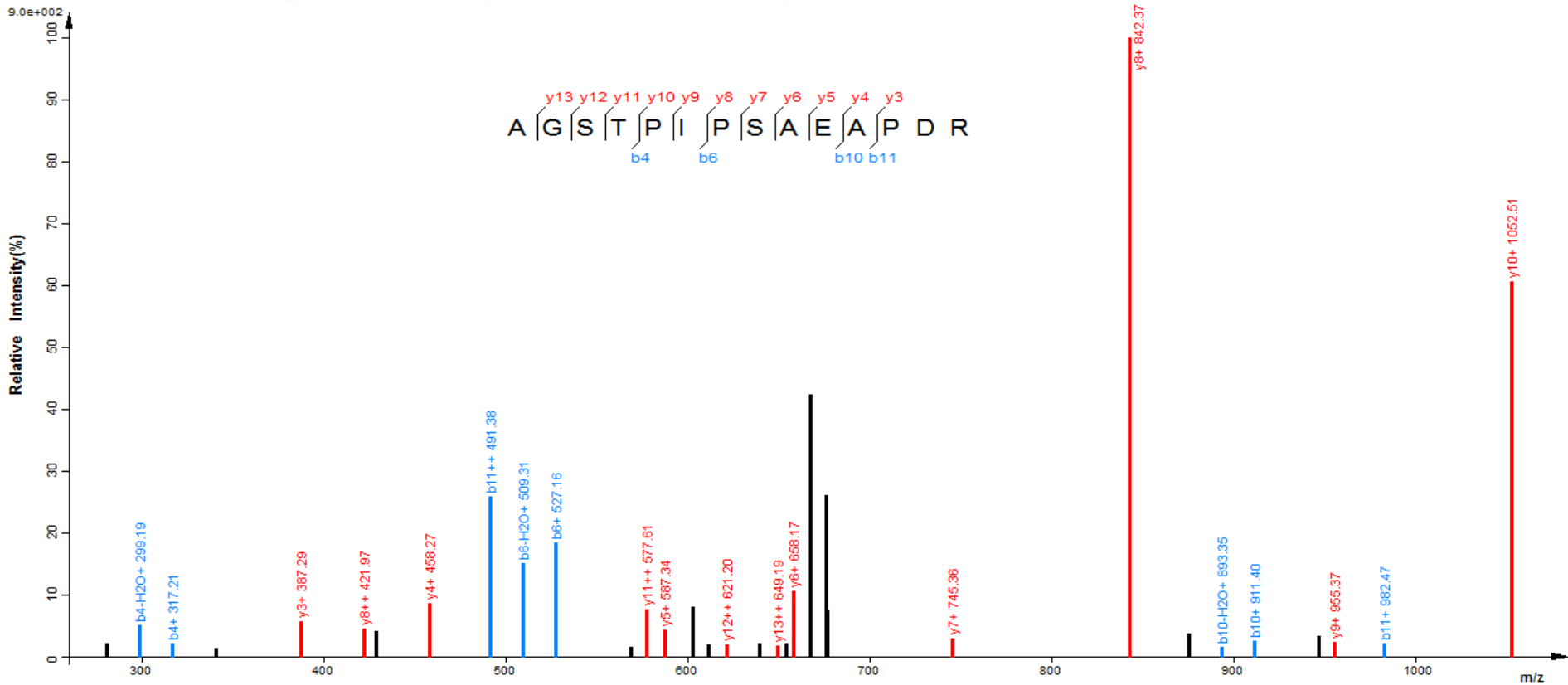
Total PSM

chrGI:27375111:ref:NC004463.1:\_6676608\_6676399\_-1

AGSTIPSAEAPDR

27

E:\PROJECTS\Bradyrhizobium\_japonicum\INPUT\10116.mgf-54525 SEQ:AGSTIPSAEAPDR Charge:2 MH+:1368.6771



Translation ID

Peptide

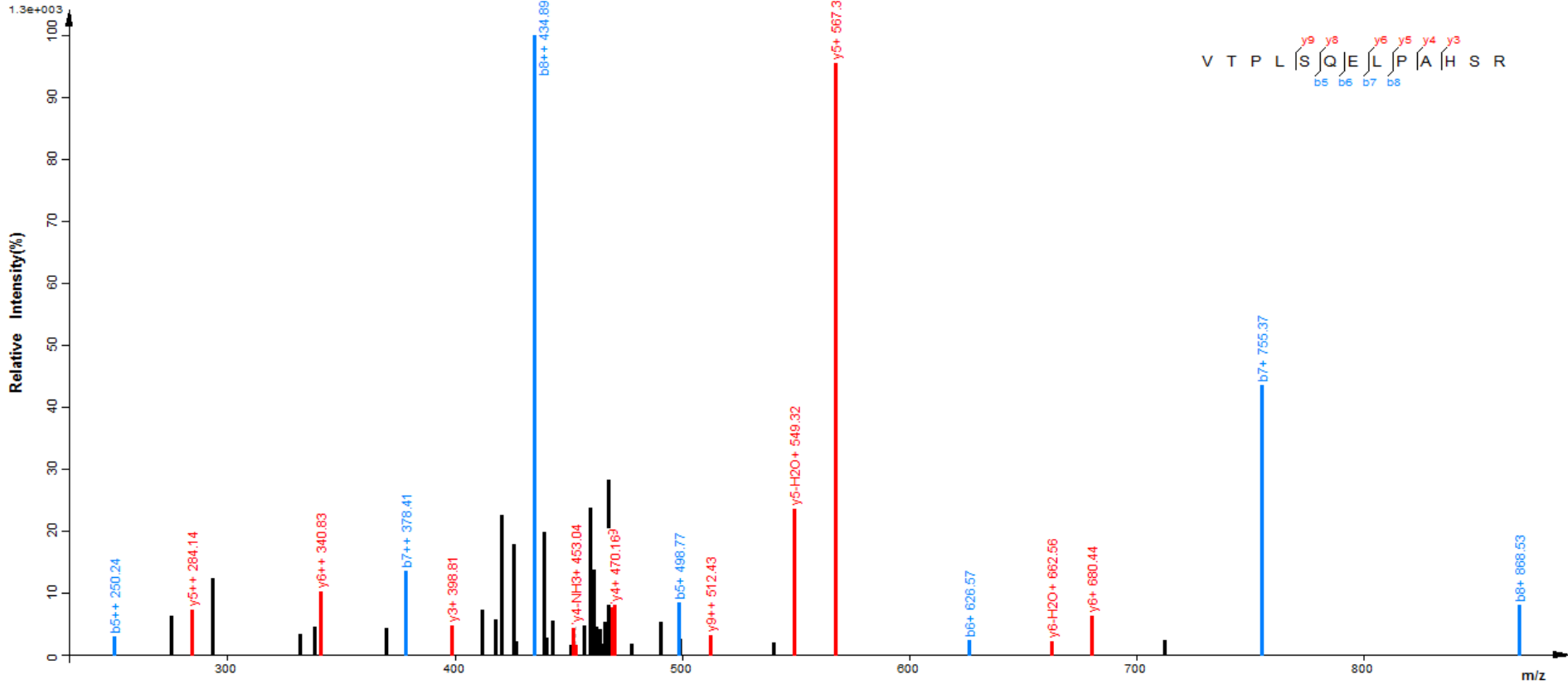
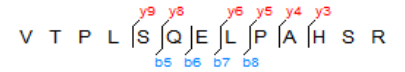
Total PSM

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VTPLSQELPAHSR

18

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Translation ID

Peptide

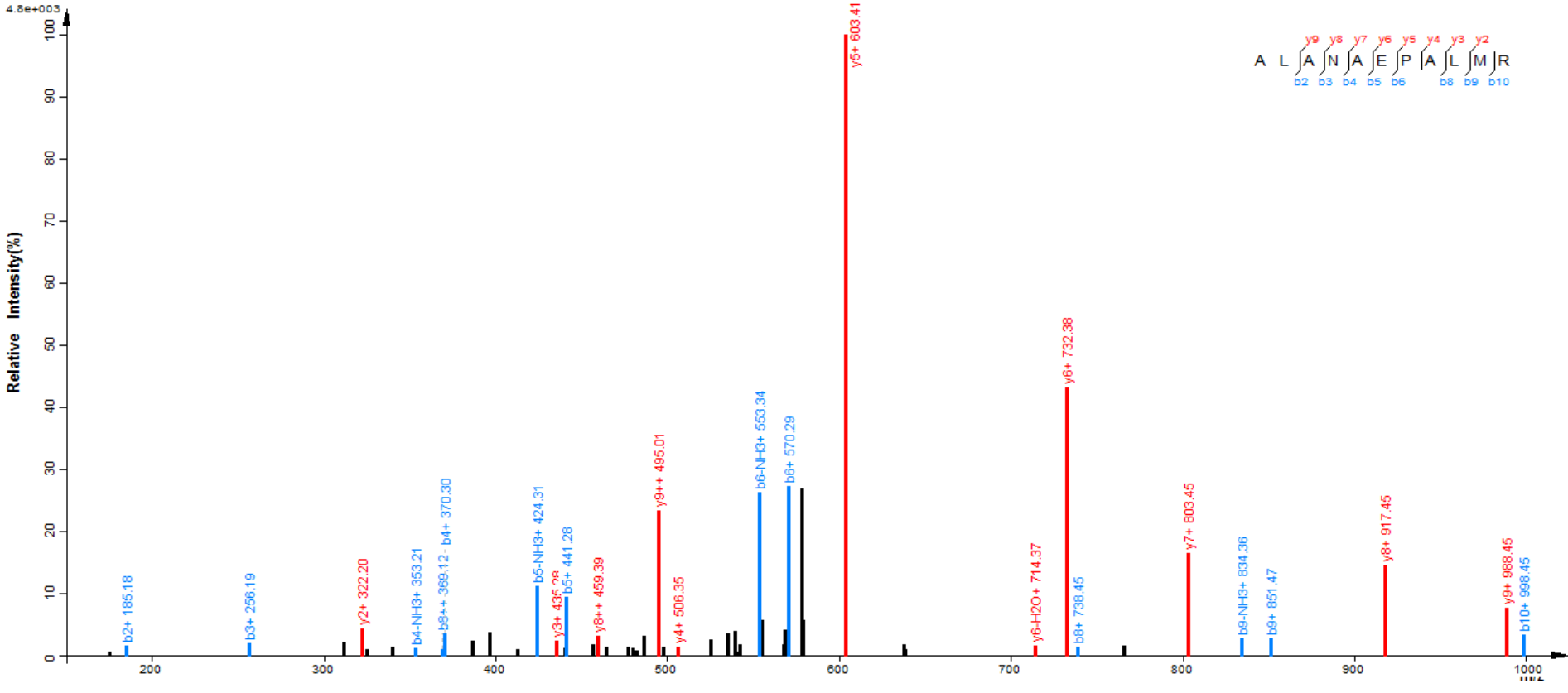
Total PSM

chrqi:27375111:ref:NC004463.1:\_896768\_896406\_-2

ALANAEPALMR

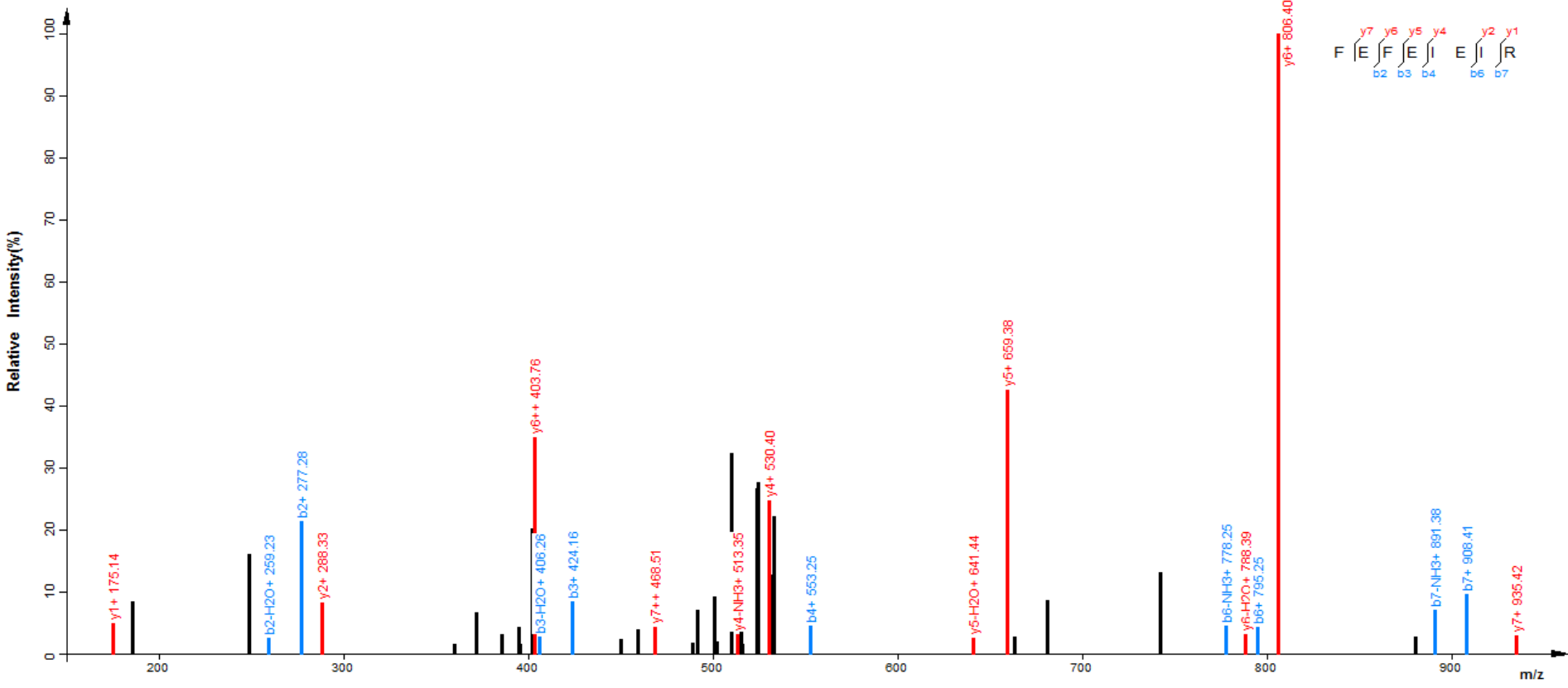
26

E:\PROJECTS\Bradyrhizobium\_japonicum\INPUT\10116.mgf-47380 SEQ:ALANAEPALMR 10\_Oxidation\_M\_15.995 Charge:2 MH+:1172.6089



Translation ID	Peptide	Total PSM
chrGI:27375111:ref:NC004463.1:_302817_302362_-1	FEFEIEIR	13

E:\PROJECTS\Bradyrhizobium\_japonicum\INPUT\10116.mgf-58614 SEQ:FEFEIEIR Charge:2 MH+:1082.4907



Translation ID

Peptide

Total PSM

chrqi:27375111:ref:NC004463.1:\_7834140\_7833187\_-1

WLSEPTGVVAGAK

14

E:\PROJECTS\Bradyrhizobium\_japonicum\INPUT\10114.mgf-1367 SEQ:WLSEPTGVVAGAK Charge:2 MH+:1443.7494

