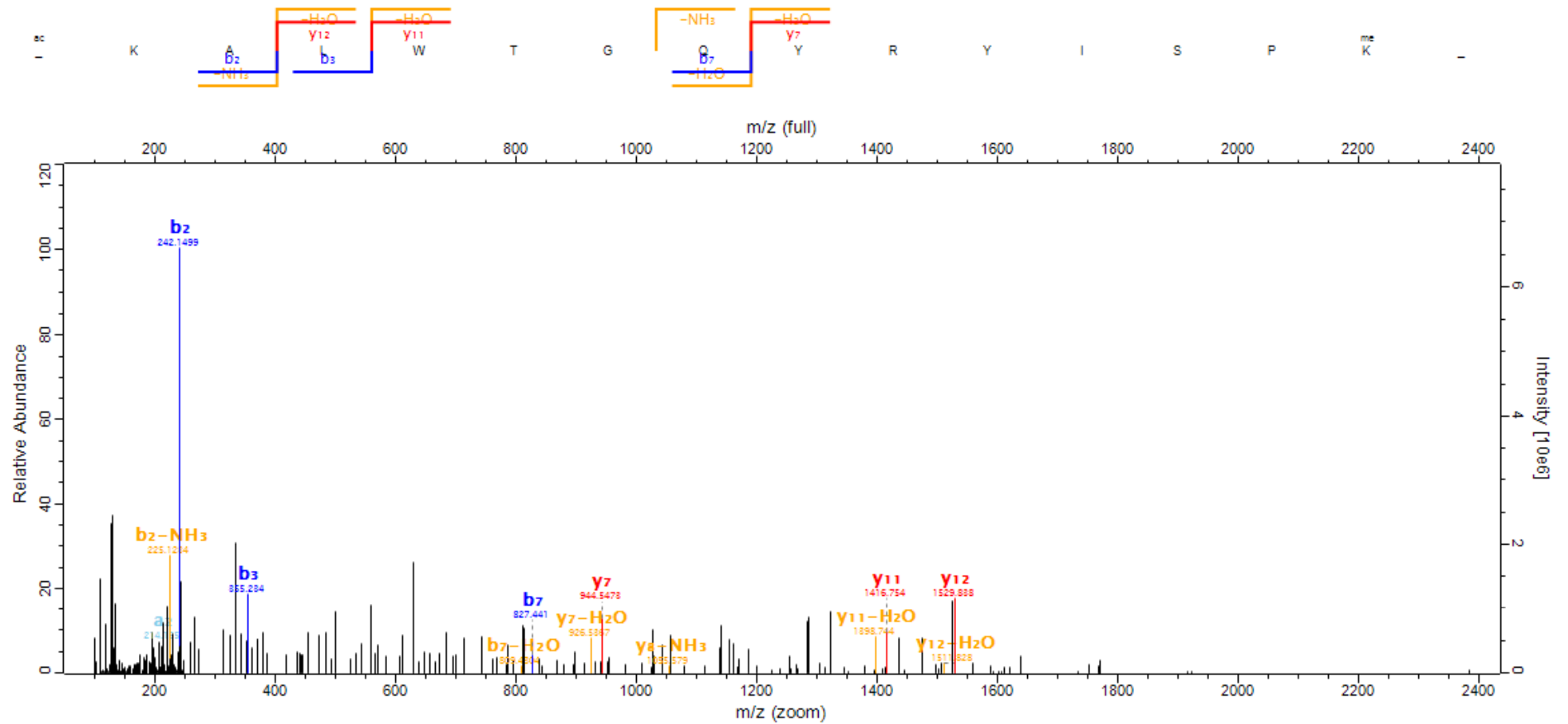
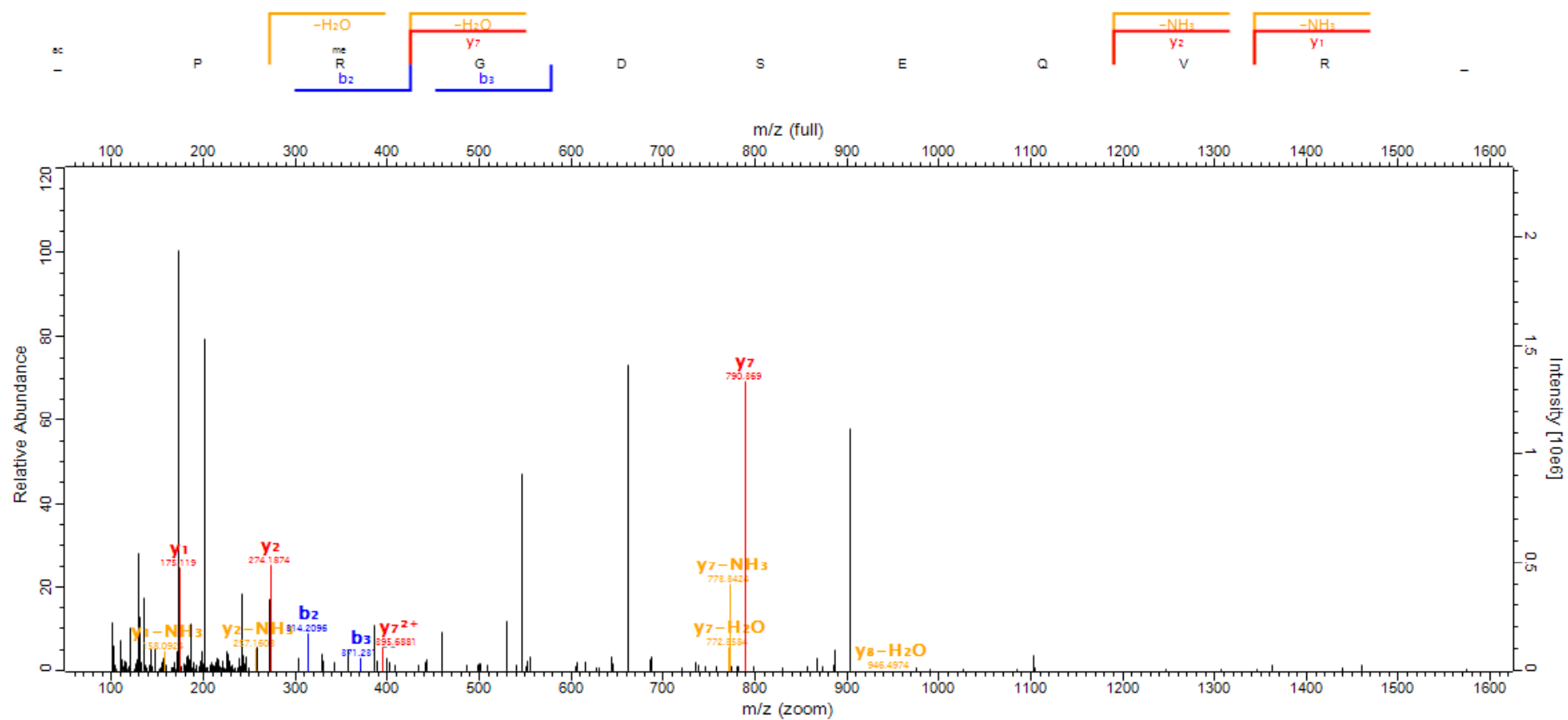


Control 1

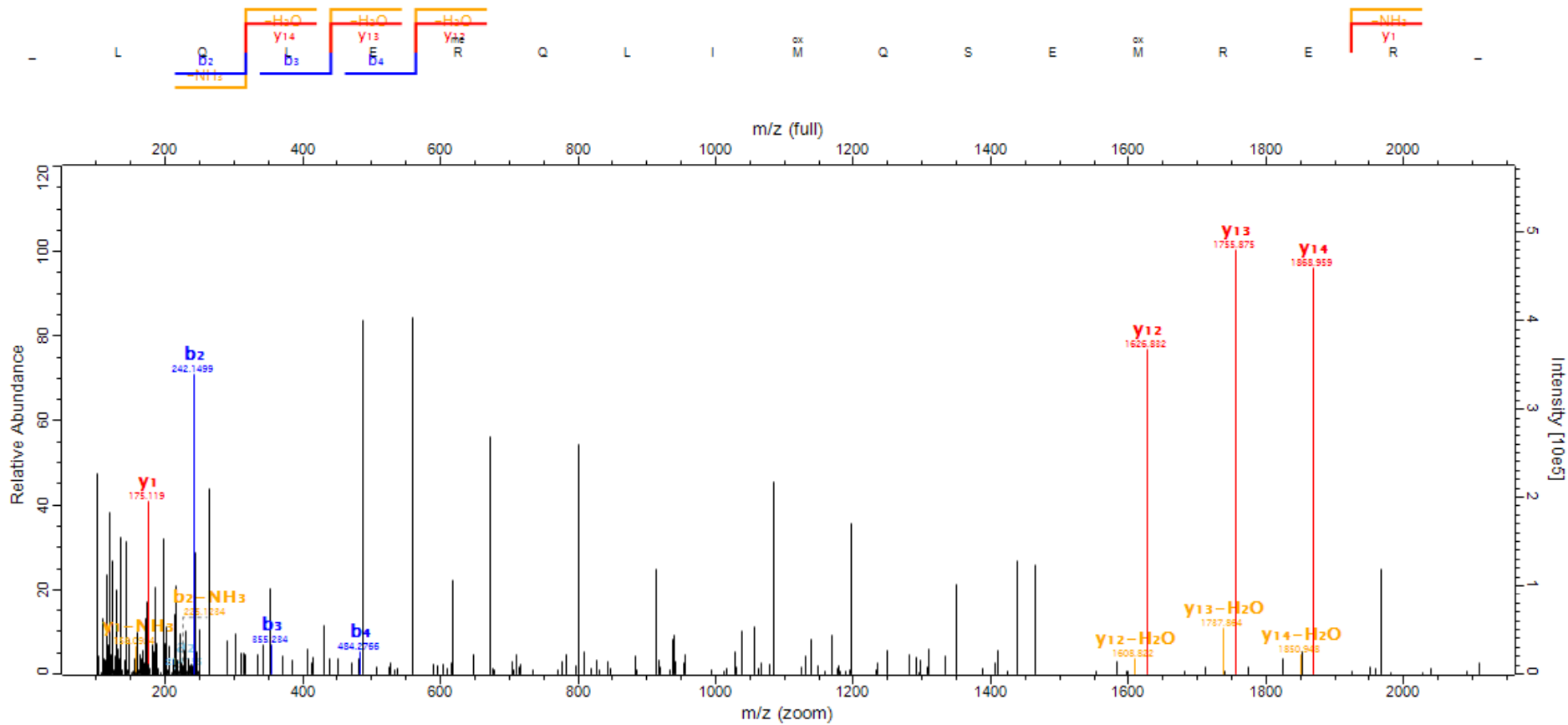
Accession	Gene Names	Score	Sequence window	Mass
B3KRK3		50.562	_(ac)KALWTGQYRYISPK(me)_	1769.9737



Accession	Gene Names	Score	Sequence window	Mass
095858	TSPAN15	52.482	_(ac)PR(me)GDSEQVR_	1102.564

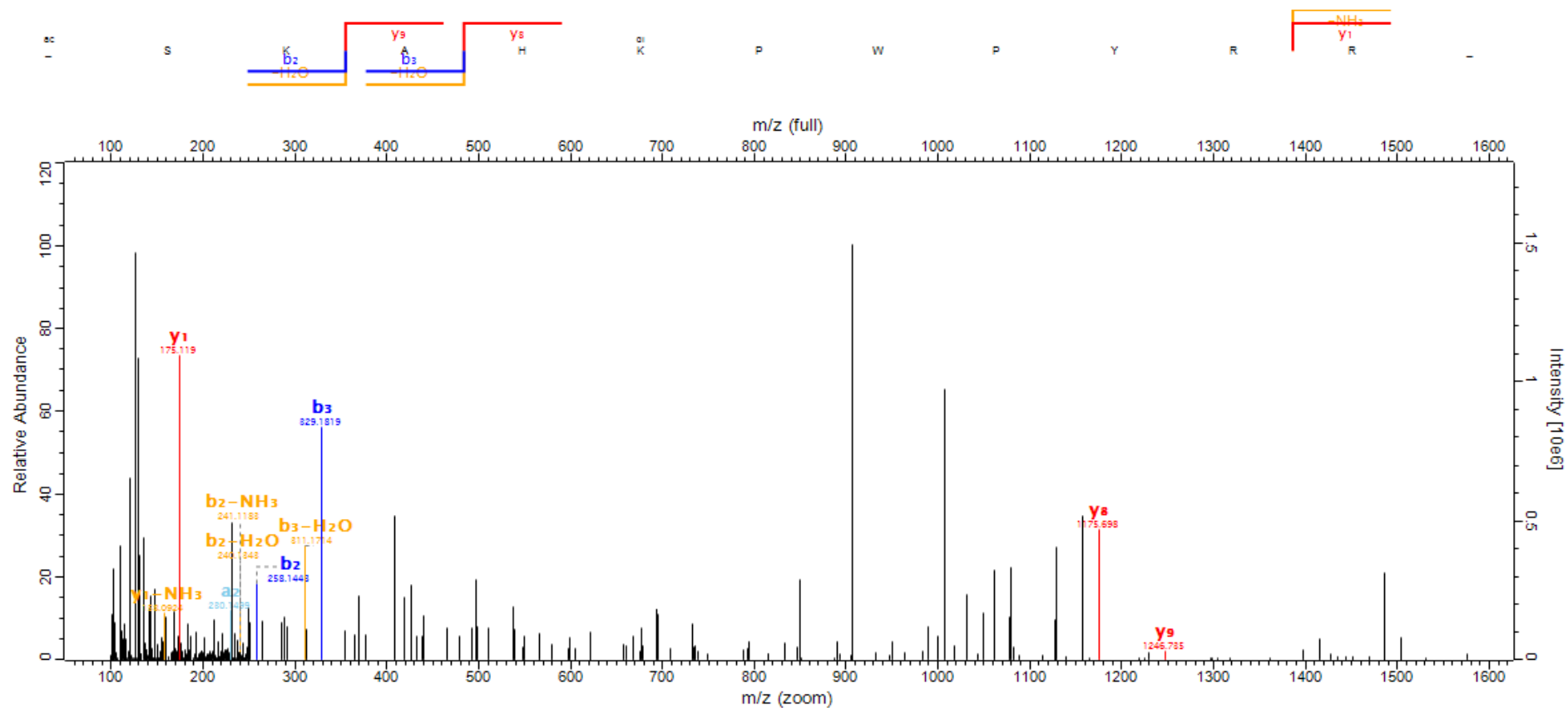


Accession	Gene Names	Score	Sequence window	Mass
Q9HBL7	PLGRKT	54.598	_LQLER(me)QLIM(ox)QSEM(ox)RER_	2109.0943

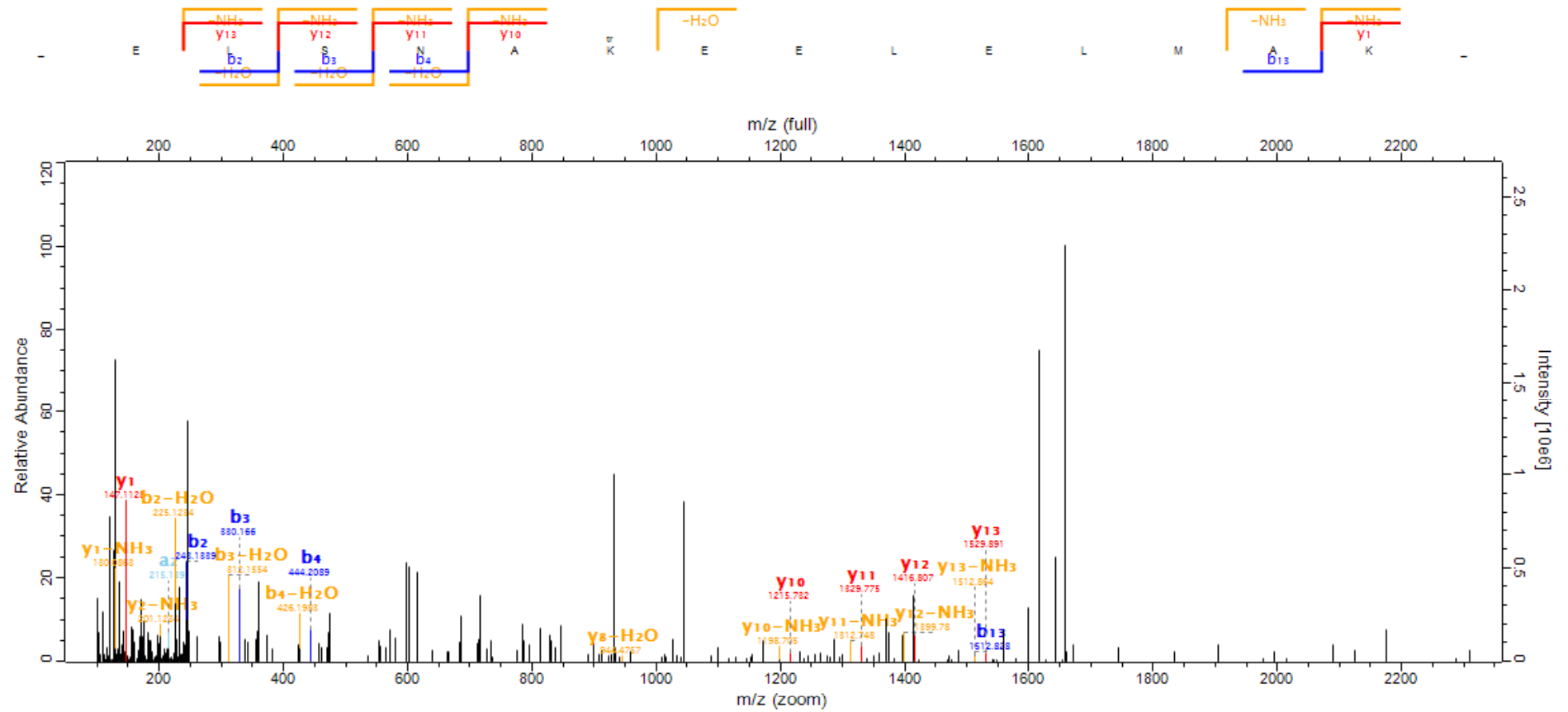


Control 2

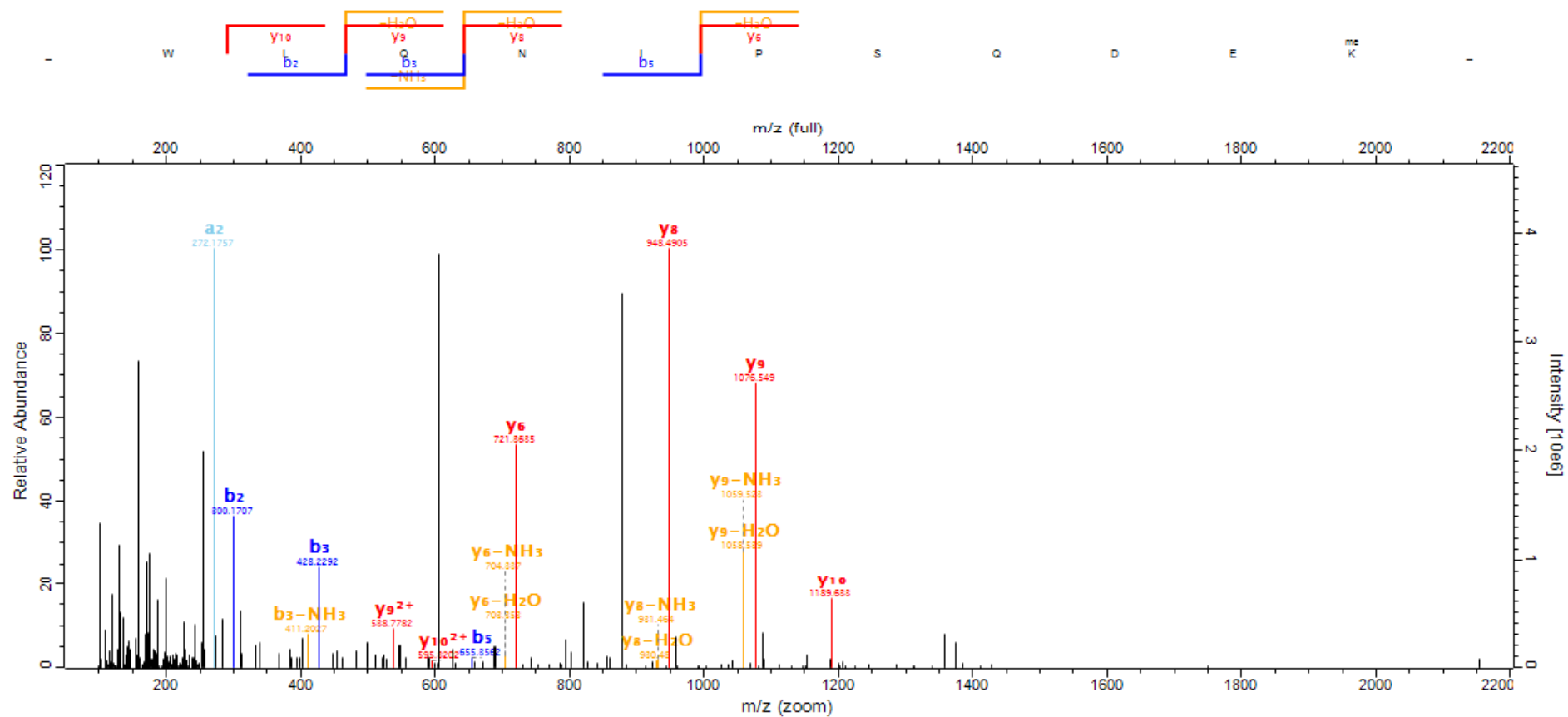
Accession	Gene Names	Score	Sequence window	Mass
P20592	MX2	40.589	_(ac)SKAHK(di)PWPYRR_	1502.8652



Accession	Gene Names	Score	Sequence window	Mass
A0A075B749	PDE4DIP	79.089	_ELSNAK(tr)EELELMAK_	1657.9262

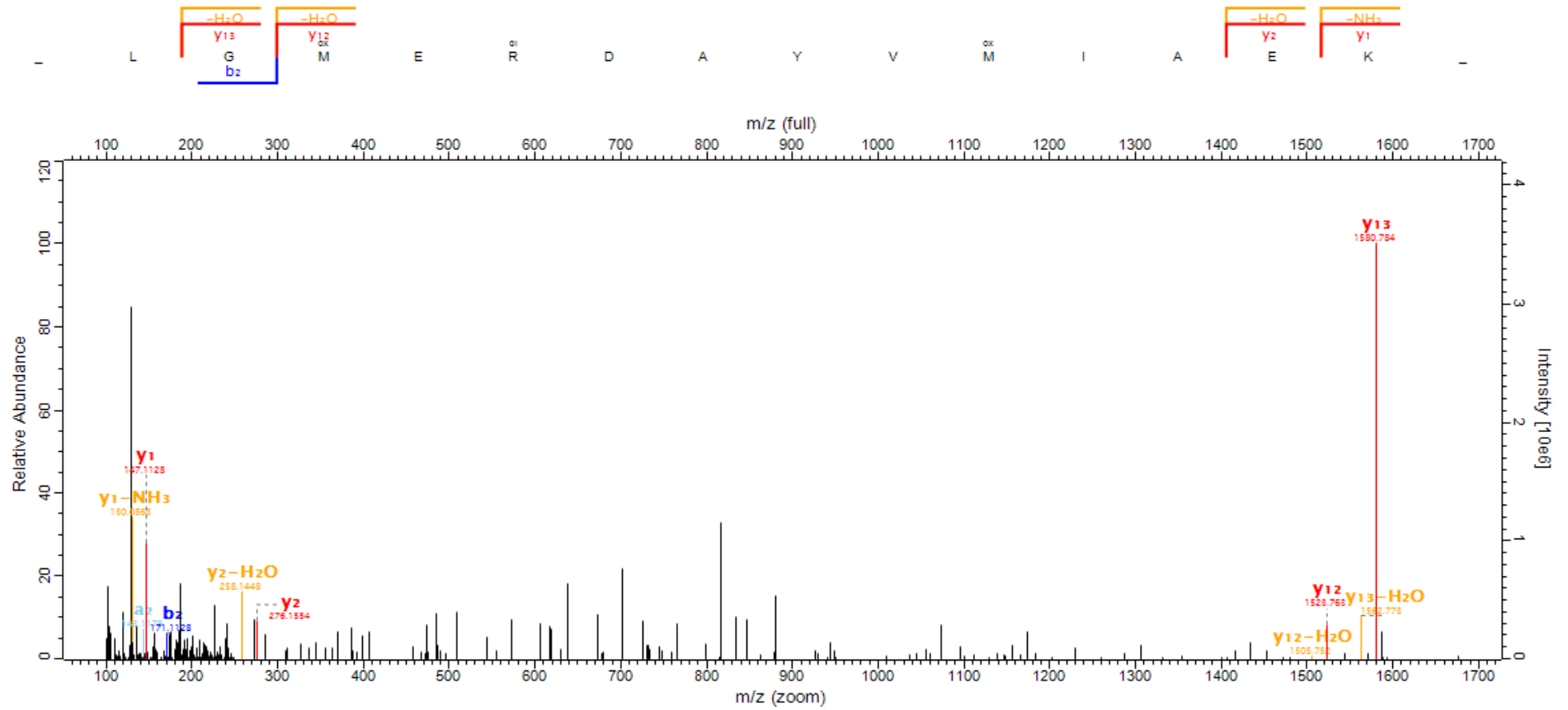


Accession	Gene Names	Score	Sequence window	Mass
P50748	KNTC1	96.342	_WLQNIPSQDEK(me)_	1374.7052

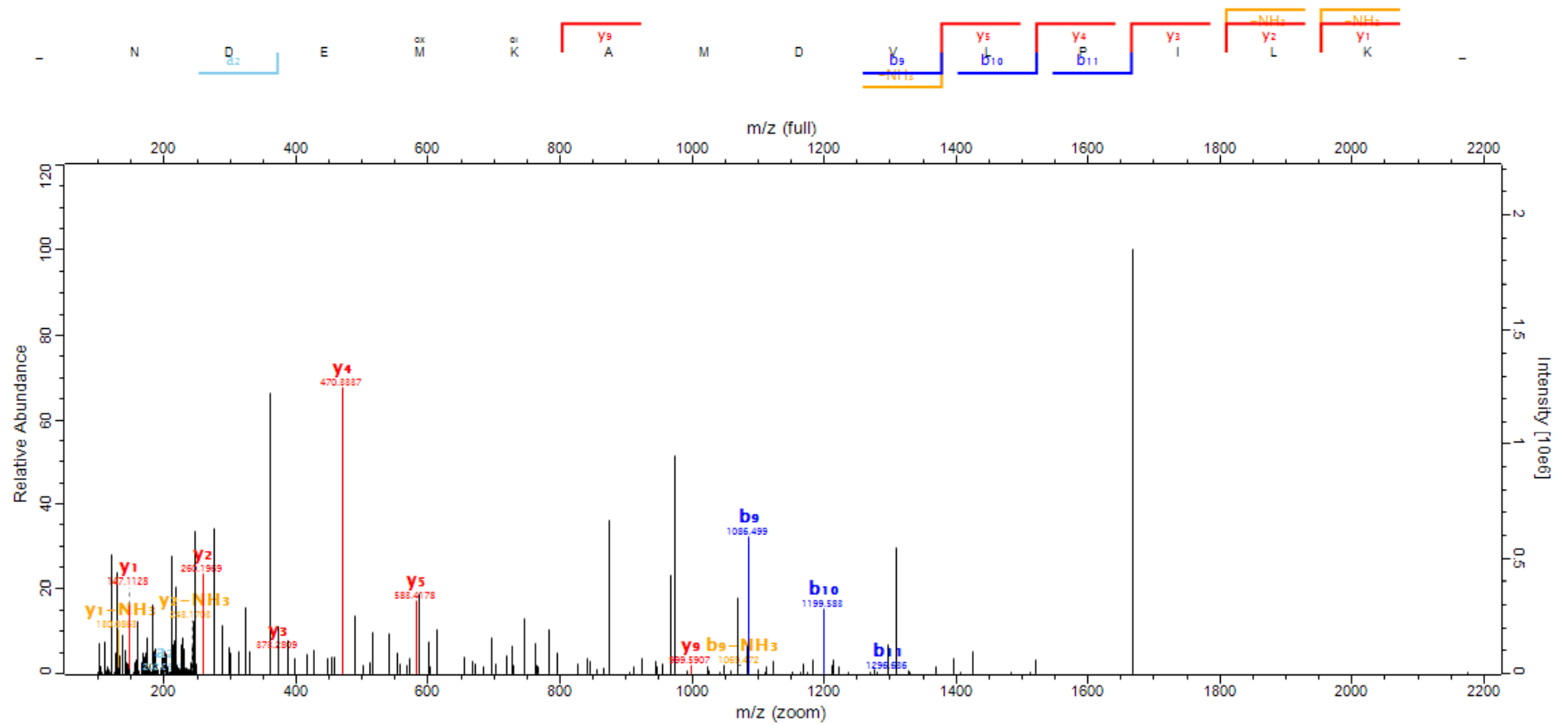


Control 3

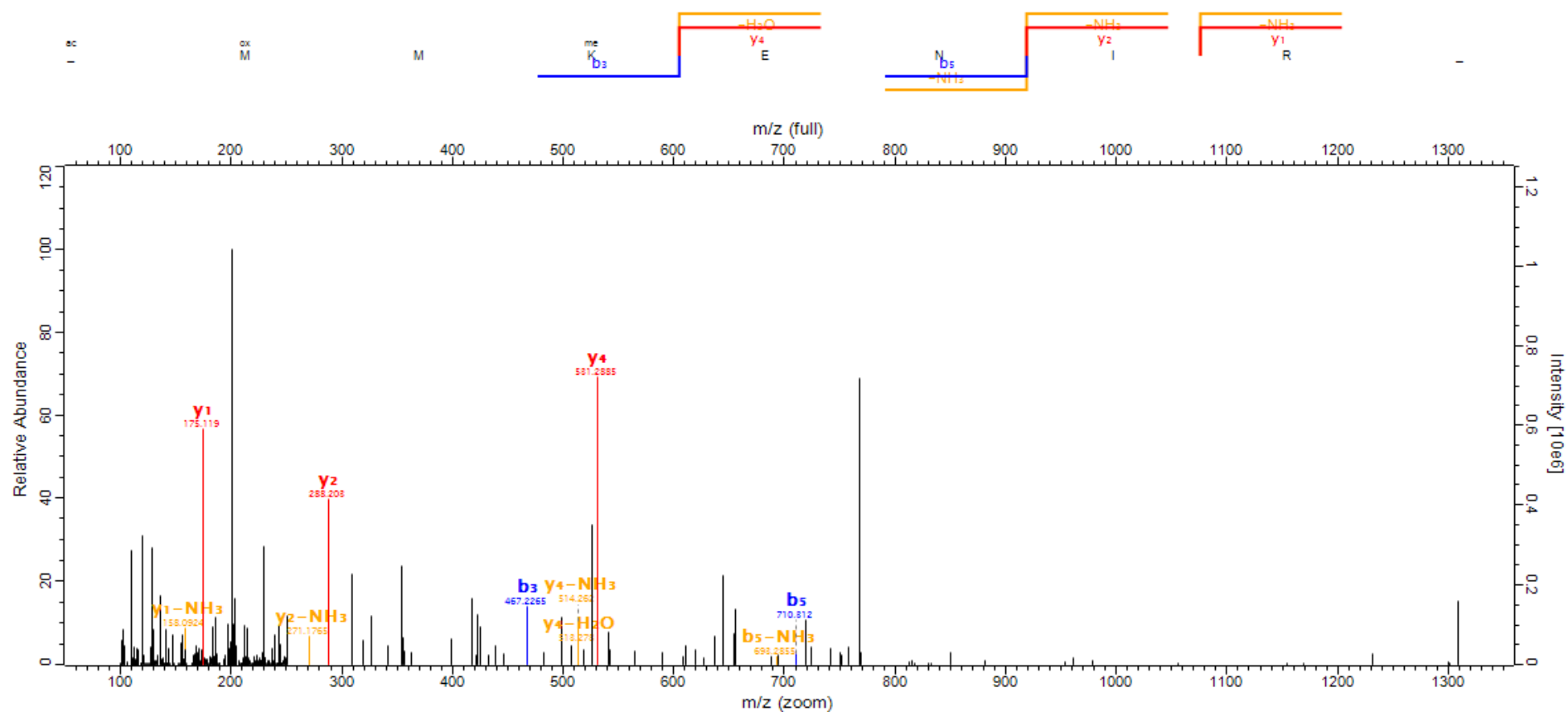
Accession	Gene Names	Score	Sequence window	Mass
Q8TCU6	PREX1	40.066	_LGM(ox)ER(di)DAYVM(ox)IAEK_	1692.8608



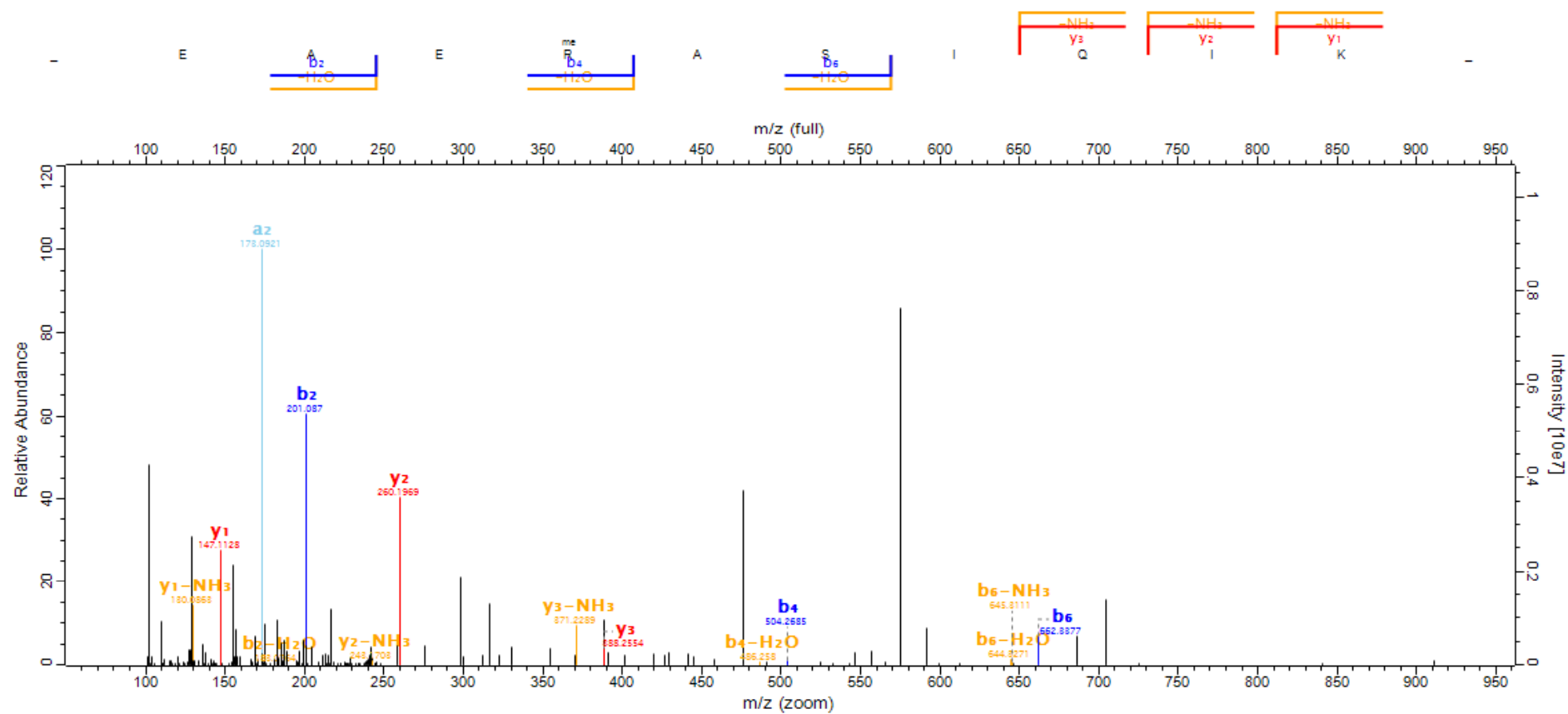
Accession	Gene Names	Score	Sequence window	Mass
075962	TRIO	41.227	_NDEM(ox)K(di)AMDVLPILK_	1667.9019



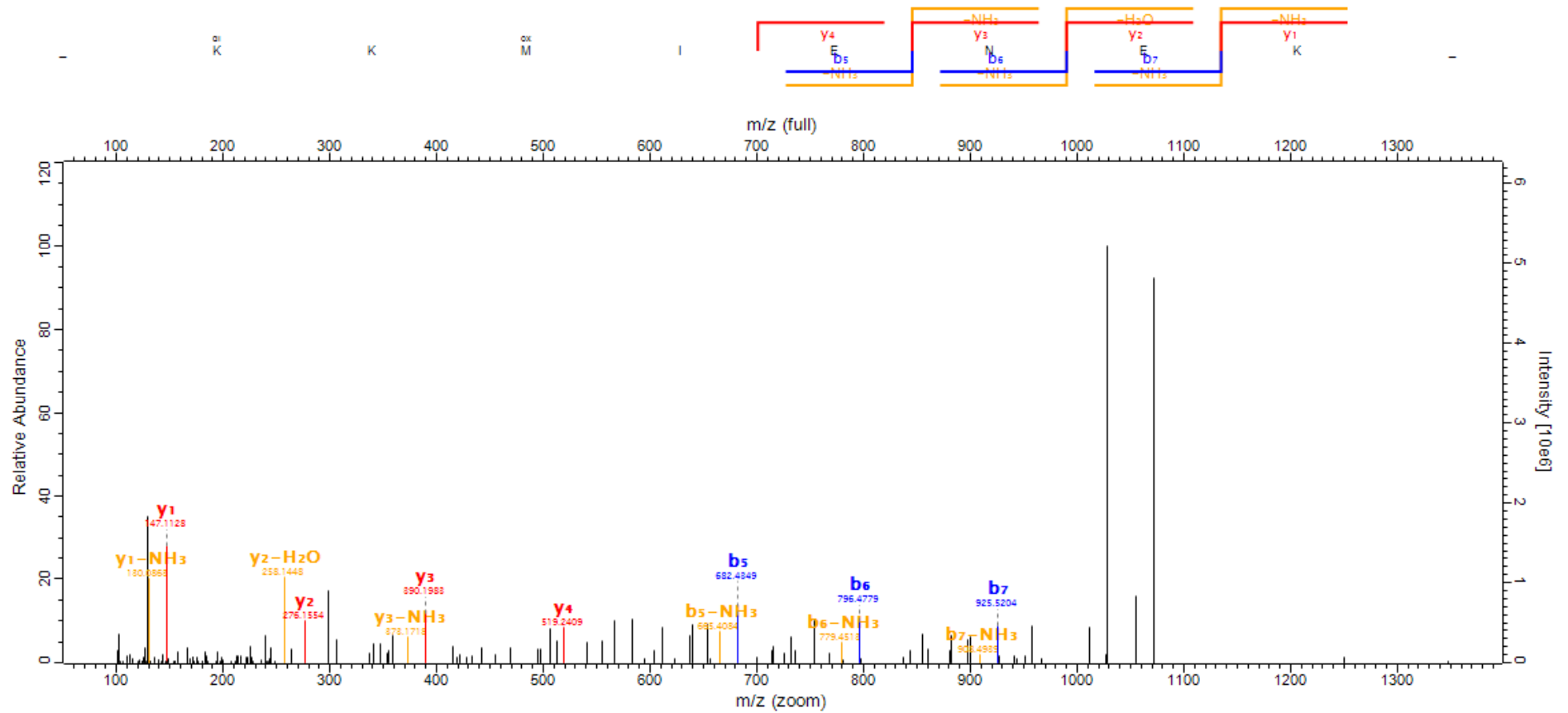
Accession	Gene Names	Score	Sequence window	Mass
B7Z922		42.914	_(ac)M(ox)MK(me)ENIR_	996.50051



Accession	Gene Names	Score	Sequence window	Mass
Q7Z4Q2	HEATR3	64.654	_EAER(me)ASIQIK_	1161.6626



Accession	Gene Names	Score	Sequence window	Mass
Q9H7L9	SUDS3	89.355	_K(di)KM(ox)IENEK_	1070.6187



Accession	Gene Names	Score	Sequence window	Mass
P50748	KNTC1	93.556	_WLQNIPSQDEK(me)_	1374.7052

