

Number Description

	Scan	Sequence	Charge	MH+theo	MH+obs	DeltaMH+	N-gap	C-gap	Sequence score	Filter score
2	sp Q01517 ALFC2_PEA	Fructose-bisphosphate aldolase 2, chlor								
	172	YLGDWSEEAQK	2	1325.6011	1325.7328	-0.131713	0.0	0.0	150.432	0.965
	201	SAAYYEQQR	2	1115.5122	1115.1581	0.354125	0.0	0.0	75.139	0.791
	340	SLAKLGK	2	917.32324	918.1846	ND	200.856	0.0	123.382	0.509
	639	LGLENTEANR	2	1432.631	1434.1942	ND	316.066	0.0	88.24	0.947
	675	LAMDSENA	2	1465.9894	1466.7211	ND	169.643	363.956	97.537	0.754
	770	TLNLLHR	3	1484.5223	1485.9032	ND	618.001	0.0	88.683	0.66
	829	QEALLFR	2	1018.2915	1018.7406	ND	141.797	0.0	108.978	0.908
	869	EYTLNLLHR	2	1328.4332	1329.2635	ND	169.806	0.0	145.998	0.887
	899	VSLPNDYFGLK	2	1452.3898	1452.8031	ND	199.732	0.0	99.358	0.223
	1170	LVDVLVLEELL	3	1992.569	1993.3284	ND	0.0	756.832	105.177	0.73
	1251	GSNNESWCQGL	2	2401.9492	2401.568	ND	550.15	618.328	81.011	0.717
3	tr A9RX76 A9RX76_PHYPA	Fructose-bisphosphate aldolase OS=Ph								
	524	RLDSLGLNTEANR	2	1587.8088	1587.6835	0.125366	0.0	0.0	134.653	0.921
	639	LGLENTEANR	2	1432.631	1434.1942	ND	316.066	0.0	88.24	0.947
	701	VAEYTLNTYQKR	3	1485.77	1486.0815	-0.311523	0.0	0.0	75.876	0.552
	1177	LVEELL	2	1777.3734	1778.7916	ND	553.887	413.997	91.616	0.29