

SRF File: C:\Xcalibur\data\100730RS\100730RS-P05.srf

Database... 100211 uniprot_sprot_human.fasta

Filter(s)... peptide probability<=5e-003 ; # distinct peptides>=2 ; protein probability<=1e-003

Mods: (M* +15.99492) C=160.03064

Reference	Sequence	MH+	z	P	Sf	Score	Coverage	Accession		
Time (s)				P	Sf	XC	DeltaCn	Sp	RSp	Ions
SIRT2_HUMAN	NAD-dependent deacetylase sirtuin-2 OS=Homo sapien			4e-012	4.5	60.2	0.0	0		
19.37	K.IFSEVTPK.C	920.5088	2	7e-005	0.77	1.383	0.397	452.6	1	12/14
20.08	R.SPSTGLYDNLEK.Y	1323.6427	2	5e-009	0.96	2.708	0.795	1163.1	1	17/22
20.25	R.SPSTGLYDNLEK.Y	1323.6427	2	4e-012	0.94	2.332	0.696	1140.0	1	17/22
24.12	R.LLDELTLEGVAR.Y	1328.7420	2	3e-009	0.98	4.337	0.727	1672.4	1	18/22
24.27	R.LLDELTLEGVAR.Y	1328.7420	2	4e-009	0.98	3.801	0.768	1984.0	1	19/22
24.41	R.NLFSQTLSLGSQK.E	1422.7587	2	1e-007	0.98	4.491	0.775	1605.2	1	18/24
24.59	R.NLFSQTLSLGSQK.E	1422.7587	2	2e-009	0.98	3.871	0.824	2072.1	1	20/24
24.83	K.AGQSDPFLGM*IM*GLGGGM*DFDSK.K	2379.0301	2	3e-007	0.05	1.499	-	229.5	1	12/44
27.48	K.YHLPYPEAIFFEISYFK.K	2017.0106	3	0.0004	0.76	2.089	0.833	445.5	1	16/60

9 of 12 peptide matches reported, 3 removed due to filtering