

SRF File: C:\Xcalibur\data\100730RS\100730RS-P04.srf

Database... 100211 uniprot\_sprot\_human.fasta

Filter(s)... peptide probability&lt;=5e-003 ; # distinct peptides&gt;=2 ; protein probability&lt;=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
KCRB_HUMAN	Creatine kinase B-type OS=Homo sapiens GN=CKB PE=1			4e-011	2.9	30.2	0.0	0
19.96	R.LEQGQAIDDLM*PAQK.-	1672.8211	2	7e-008	0.94	2.582	0.834	941.8 1 17/28
24.72	K.LAVEALSSLDGDLAGR.Y	1586.8384	2	9e-011	0.94	2.501	0.815	1113.6 1 17/30
27.79	R.LGFSEVELVQM*VVDGVK.L	1864.9725	2	8e-011	0.98	4.416	0.862	2140.0 1 22/32
27.87	R.LGFSEVELVQM*VVDGVK.L	1864.9725	2	4e-011	0.98	3.991	0.845	1830.8 1 21/32

4 of 7 peptide matches reported, 3 removed due to filtering

SIRT2_HUMAN	NAD-dependent deacetylase sirtuin-2 OS=Homo sapien			6e-007	2.9	30.2	0.0	0
20.17	R.SPSTGLYDNLEK.Y	1323.6427	2	6e-007	0.93	2.317	0.776	869.9 1 15/22
20.27	R.SPSTGLYDNLEK.Y	1323.6427	2	1e-006	0.65	1.509	0.861	318.5 1 9/22
24.25	R.LLDELTLLEGVAR.Y	1328.7420	2	7e-004	0.98	3.631	0.836	1910.4 1 18/22
24.55	R.NLFSQTLSLGSQK.E	1422.7587	2	7e-005	0.97	2.903	0.860	1593.6 1 17/24

4 of 7 peptide matches reported, 3 removed due to filtering

SNAG_HUMAN	Gamma-soluble NSF attachment protein OS=Homo sapien			4e-005	1.8	20.1	0.0	0
18.79	K.LIENVDPK.A	1056.5572	2	0.0001	0.88	2.059	0.632	541.6 1 12/16
20.70	R.FDEAALSIQK.E	1121.5837	2	4e-005	0.96	2.648	0.682	1431.9 1 16/18

2 of 3 peptide matches reported, 1 removed due to filtering