

Accession	Description	Score	Coverage	# Proteins	# Unique Peptides	# Peptides	# PSMs	# AAs	MW [kDa]	calc. pI					
46402201	Jim Morris Clemson [human]	1516.92	76.43	1	30	30	459	471	51.4	8.82					
	A3	Sequence	# PSMs	# Proteins	# Protein Groups	Protein Group Accessions	Modifications	ΔCn	XCorr	Charge	MH+ [Da]	ΔM [ppm]	RT [min]	# Missed Cleavages	
High		AAQLSASFcaAPLVK	18	1		1.46402201	C9(Cys->Dha); C10(Dioxidation)	0.0000	4.00	2	1506.74639	-10.48	94.46	0	
High		ATAIDGSVFEK	7	1		1.46402201		0.0000	3.54	2	1250.66509	1.86	86.79	0	
High		ATAIDGSVFEKIPSPR	1	1		1.46402201		0.0000	2.53	3	1851.00199	0.44	100.99	1	
High		DGSGGAIFISAMVNDK	26	1		1.46402201		0.0000	4.79	2	1751.86406	0.39	109.29	0	
High		DVAMAALTNQFTmSVESMR	28	1		1.46402201	M13(Oxidation)	0.0000	3.12	3	2117.97013	1.69	111.95	0	
High		DVAMAALTNQFTmSVESMR	21	1		1.46402201		0.0000	5.72	2	2101.97588	2.02	113.32	0	
High		EGAVVDSSTSAFK	11	1		1.46402201		0.0000	3.54	2	1297.62847	1.05	67.24	0	
High		FAGmISADR	25	1		1.46402201	M4(Oxidation)	0.0000	3.07	2	983.46269	1.16	61.49	0	
High		FAGmISADR	6	1		1.46402201		0.0000	2.92	2	967.46697	0.34	72.40	0	
High		FAGmISADRMpGLQFTR	1	1		1.46402201	M4(Oxidation); M10(Oxidation)	0.0000	2.67	3	1929.93436	1.70	80.52	1	
High		FAGmISADRMpGLQFTR	1	1		1.46402201	M4(Oxidation)	0.0000	2.63	3	1913.93698	0.43	90.14	1	
High		FDLIDDASLNK	4	1		1.46402201		0.0000	4.19	2	1365.65544	1.55	90.59	0	
High		FDLIDDASLNKGGQALEK	7	1		1.46402201		0.0000	5.41	2	2120.05351	1.32	89.46	1	
High		FVLPTTKFDLIDDASLNKGGQALEK	6	1		1.46402201		0.0000	4.29	4	2906.50722	-2.58	98.95	2	
High		GSALTPINmESGNFDSK	28	1		1.46402201	M9(Oxidation)	0.0000	3.78	2	1783.82146	2.61	75.06	0	
High		GSALTPINmESGNFDSK	18	1		1.46402201		0.0000	3.88	2	1767.82451	1.49	85.85	0	
High		GVQGNVDVIALQAFAFR	22	1		1.46402201		0.0000	5.37	2	1728.94011	0.50	127.50	0	
High		ILGPEcDVR	6	1		1.46402201	C6(Trioxidation)	0.0000	2.74	2	1049.49370	0.43	72.67	0	
High		LNNILEHISIQGNDGETVR	6	1		1.46402201		0.0000	5.92	3	2122.09201	1.51	90.03	0	
High		mISGMYLGEIAR	8	1		1.46402201	M1(Oxidation); M5(Oxidation)	0.0000	3.59	2	1372.66216	1.56	82.40	0	
High		mISGMYLGEIAR	10	1		1.46402201	M1(Oxidation)	0.0000	3.83	2	1356.66557	0.35	90.53	0	
High		MISGMYLGEIAR	9	1		1.46402201		0.0000	4.35	2	1340.67107	0.66	94.70	0	
High		mPGLQFTR	5	1		1.46402201	M1(Oxidation)	0.0000	2.49	2	965.48802	0.68	73.05	0	
High		mPGLQFTR	7	1		1.46402201		0.0000	2.76	2	949.49254	0.09	82.79	0	
High		QImTYLLYEMVEGLEGR	4	1		1.46402201	M3(Oxidation)	0.0000	4.95	2	2061.00493	0.83	125.87	0	
High		QImTYLLYEMVEGLEGR	15	1		1.46402201		0.0000	5.52	2	2045.00957	0.62	137.92	0	
High		RDVAMAALTNQFTmSVESMR	2	1		1.46402201		0.0000	3.64	3	2258.07730	2.02	106.10	1	
High		RLNILEHISIQGNDGETVR	18	1		1.46402201		0.0000	5.18	3	2278.19583	2.59	87.99	1	
High		RVIVHSSINcLPAALQ TALG NR	2	1		1.46402201	C11(Dioxidation)	0.0000	4.23	3	2478.36435	1.33	109.62	1	
High		RVIVHSSINcLPAALQ TALG NR	5	1		1.46402201	C11(Delta:S(-1)Se(1))	0.0000	5.12	3	2494.35678	16.48	111.33	1	
High		RVIVHSSINcLPAALQ TALG NR	5	1		1.46402201	C11(Trioxidation)	0.0000	3.91	3	2494.35807	0.84	111.98	1	
High		RVLQDNINR	7	1		1.46402201		0.0000	3.50	2	1127.63005	1.79	55.58	1	
High		TVPLGFTSFVPEQTK	22	1		1.46402201		0.0000	4.08	2	1797.94353	0.72	109.70	0	
High		VACKEGAVDSSTSAFK	15	1		1.46402201	C3(Dioxidation)	0.0000	3.80	3	1730.83042	2.19	72.05	1	
High		VACKEGAVDSSTSAFK	15	1		1.46402201	C3(Trioxidation)	0.0000	5.54	2	1746.82573	2.40	73.06	1	
High		VACKEGAVDSSTSAFK	7	1		1.46402201	C3(Delta:S(-1)Se(1))	0.0000	5.19	2	1746.82292	23.86	72.43	1	
High		VACKEGAVDSSTSAFKIPK	2	1		1.46402201	C3(Dioxidation)	0.0000	3.07	3	2069.06046	0.98	80.37	2	
High		VACKEGAVDSSTSAFKIPK	2	1		1.46402201	C3(Delta:S(-1)Se(1))	0.0000	4.49	3	2085.05307	19.19	81.13	2	
High		VACKEGAVDSSTSAFKIPK	2	1		1.46402201	C3(Trioxidation)	0.0000	4.49	3	2085.05307	-0.14	81.13	2	
High		VcGVdVQSIEDLR	3	1		1.46402201	C2(Dioxidation)	0.0000	2.32	2	1464.70036	0.28	91.71	0	
High		VcGVdVQSIEDLR	3	1		1.46402201	C2(Trioxidation)	0.0000	3.41	2	1480.69670	1.23	92.21	0	
High		VIVHSSINcLPAALQ TALG NR	6	1		1.46402201	C10(Dioxidation)	0.0000	4.06	3	2322.26053	0.26	114.16	0	
High		VIVHSSINcLPAALQ TALG NR	3	1		1.46402201	C10(Trioxidation)	0.0000	5.28	3	2338.25772	1.23	116.34	0	
High		VIVHSSINcLPAALQ TALG NR	3	1		1.46402201	C10(Delta:S(-1)Se(1))	0.0205	5.25	3	2338.25699	18.15	116.38	0	
High		VNVVALcNDVTGLLISHYFKDPEVQVG	4	1		1.46402201	C7(Dioxidation); C3(Cys->Dha)	0.0000	3.29	5	5608.81269	-2.57	120.71	2	
High		YALEGNATLDFDIASNVK	20	1		1.46402201		0.0000	6.09	2	2088.03081	1.11	122.56	0	
High		YALEGNATLDFDIASNVK	13	1		1.46402201		0.0000	4.28	3	2216.12766	1.89	117.66	1	