

**Table 1. Mass measured and the peptide assignments of identified proteins for “Chemical rescue of  $\Delta$ F508-CFTR mimics Genetic Repair in Cystic Fibrosis Bronchial Epithelial Cells”, Singh, O.V. et al.**

Data is processed peak lists of MALDI-TOF mass spectra that was collected and searched in MS-Fit search engine using SwissProt database (release date: 07.05.2006) as described in Supplemental material 4.4.

Spot No.	Protein Description	Mass Measured	Mass accuracy (ppm)	Peptide assignments
IP-1	Glucose regulated protein 94 kDa/ TRA1	1031.4909	-7.3	<a href="#">YFMAGSSR</a>
		1139.5863	4.7	<a href="#">LGVIEDHSNR</a>
		1275.6226	-16	<a href="#">ELISNASDALDK</a>
		1306.6560	4.2	<a href="#">EFEPLLNWMK</a>
		1485.7143	-27	<a href="#">GVVDSDDLPLNVS</a>
		1513.7701	-6.3	<a href="#">NLLHVTDTGVMTR</a>
		1525.7168	-6.9	<a href="#">EEASDYLELDTIK</a>
		1595.7252	0.87	<a href="#">VFITDDFHDMMPK</a>
		1866.7046	-11	EESDDEAAVEEEEEK
		1878.9495	3.4	<a href="#">YSQFINFPIYVWSSK</a>
		2046.0101	-25	<a href="#">LISLTDENALSGNEELTVK</a>
		2287.0263	1.5	<a href="#">ESDDPMA YIHFTAEGEVTFK</a>
		2542.2211	-28	<a href="#">TVWDWELMNDIKPIWQRPSK</a>
	2711.2090	0.14	<a href="#">HNNDTQHIWESDSNEFSVIADPR</a>	
IP-2	Heat shock protein 84 kDa / HSPCB	730.4407	-7.7	<a href="#">LSELLR</a>
		891.4227	-2.9	<a href="#">FYEAFSK</a>
		1160.5934	8.2	<a href="#">SIYYITGESK</a>
		1348.6604	-3.4	<a href="#">HFSVEGQLEFR</a>
		1349.7144	-15	<a href="#">TLTLVDTGIGMTK</a>

		1416.6346	-2.5	<a href="#">EGLELPEDEEEK</a>
		1494.6578	33	<a href="#">IEDVGSDEEDDSGK</a>
		1527.7498	3.6	<a href="#">SLTNDWEDHLAVK</a>
		1782.9457	-2.6	<a href="#">HLEINPDHPIVETLR</a>
		1808.9081	-28	<a href="#">HSQFIGYPITLYLEK</a>
		1847.7920	-3.0	<a href="#">NPDDITQEEYGEFYK</a>
		2176.9151	-14	<a href="#">YHTSQSGDEMTSLSEYVSR</a>
		2255.9558	-1.6	<a href="#">HNDDEQYAWESSAGGSFTVR</a>
		2988.5912	20	<a href="#">DLVLLFETALLSSGFSLEDPQTHSNR</a>
		3284.6046	1.0	<a href="#">AFMEALQAGADISMIGQFGVGFYSAYLVAEK</a>
		3532.6148	-8.7	<a href="#">LGLGIDEDEVA</a> <a href="#">AEEPNA</a> <a href="#">AVPDEIPPLEGDEDASR</a>
IP-3	<b>Glucose regulated protein 78 kDa (BiP)/ HSPA5</b>	728.4695	3.4	<a href="#">IQQLVK</a>
		806.3743	3.1	<a href="#">MVNDAEK</a>
		903.4707	-6.1	<a href="#">VMEHFIK</a>
		918.4718	-7.2	<a href="#">VLEDSDLK</a>
		918.5027	26	<a href="#">VLEDSDLK</a>
		981.4857	-3.7	<a href="#">ETAEAYLGK</a>
		986.5193	3.5	<a href="#">LTPEEIER</a>
		997.5073	-11	<a href="#">ALSSQHQAR</a>
		1074.5578	3.2	<a href="#">ITITNDQNR</a>
		1191.6768	33	<a href="#">VYEGERPLTK</a>
		1313.6104	-7.3	<a href="#">FEELNMDLFR</a>
		1316.6168	-16	<a href="#">NELESYAYSLK</a>
		1316.6378	0.29	<a href="#">NELESYAYSLK</a>
		1397.7806	-6.1	<a href="#">ELEEIVQPIISK</a>
		1460.7491	-7.2	<a href="#">SDIDEIVLVGGSTR</a>
		1536.7968	-1.0	<a href="#">TFAPEEISAMVLTK</a>

		1566.7298	-32	<a href="#">ITPSYVAFTPEGER</a>
		1604.8672	1.5	<a href="#">TKPYIQVDIGGGOTK</a>
		1677.8098	0.83	<a href="#">NQLTSNPENTVFDAK</a>
		1818.8292	-5.8	<a href="#">LYGSAGPPPTGEEDTAEK</a>
		1836.9308	-1.9	<a href="#">SQIFSTASDNQPTVTIK</a>
		1887.9761	2.3	<a href="#">VTHAVVTVPAYFNDAQR</a>
		1934.0230	4.9	<a href="#">DNHLLGTFDLTGIPPAPR</a>
		1974.9059	-1.3	<a href="#">IEWLESHODADIEDFK</a>
		1999.0808	-2.8	<a href="#">GVPQIEVTFEIDVNGILR</a>
		2164.9910	-0.76	<a href="#">IEIESFYEGEDFSETLTR</a>
IP-4	<b>dnaK-type molecular chaperone/ HSPA1</b>	704.3553	-3.7	<a href="#">DISQNK</a>
		723.4135	13	<a href="#">VQVSYK</a>
		760.4179	-3.4	<a href="#">NSTIPTK</a>
		902.4819	-1.8	<a href="#">STLEPVEK</a>
		1109.5538	-19	<a href="#">LLQDFFNCR</a>
		1183.6479	0.31	<a href="#">FELSGIPPAPR</a>
		1197.6939	-1.4	<a href="#">DAGVIAGLNVLR</a>
		1465.8181	3.7	<a href="#">AQIHDLVLVGGSTR</a>
		1487.7112	6.3	<a href="#">TTPSYVAFTDTER</a>
		1614.8083	-0.38	<a href="#">AFYPEEISSMVLTK</a>
		2304.1560	0.62	<a href="#">SINPDEAVAYGAAVQAAILMGDK</a>
		2530.3235	7.7	<a href="#">GVPQIEVTFDIDANGILNVTATDK</a>
		2786.3661	0.86	<a href="#">QTQIFTTYSNQPGLIQVYEGER</a>
		2981.4686	1.8	<a href="#">TLSSSTQASLEIDSLFEGIDFYTSITR</a>
IP-5	<b>Heat shock 70 kDa protein 2/ HSPA2</b>	848.4207	24	<a href="#">QTVEDEK</a>
		1076.5097	7.8	<a href="#">DEYEHKQK</a>

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	1111.5094	6.7	<a href="#">YKSEDEANR</a>
	1175.6206	-5.6	<a href="#">ENKITITNDK</a>
	1190.5804	-6.4	<a href="#">MVSHLAEEFK</a>
	1253.6120	-3.7	<a href="#">FEELNADLFR</a>
	1270.5520	-8.3	<a href="#">FEDATVQSDMK</a>
	1309.6890	4.9	<a href="#">MKEIAEAYLGGK</a>
	1315.6558	1.8	<a href="#">NALESYTYNIK</a>
	1450.8152	6.5	<a href="#">IQKLLQDFFNGK</a>
	1476.6797	3.0	<a href="#">DDIDRMVQEAER</a>
	1480.7243	-2.1	<a href="#">ARFEELNADLFR</a>
	1487.7092	5.0	<a href="#">TTPSYVAFTDTER</a>
	1646.9161	2.6	<a href="#">VVSEGGKPKVQVEYK</a>
	1657.9291	3.3	<a href="#">QATKDAGTITGLNVL</a>
	1787.9971	3.6	<a href="#">IINEPTAAAIAYGLDKK</a>
	1795.0406	-2.0	<a href="#">GQIQEIVLVGGSTRIPK</a>
	1813.0117	-3.7	<a href="#">LDKGQIQEIVLVGGSTR</a>
	1819.9169	2.4	<a href="#">NQVAMNPTNTIFDAKR</a>
	1887.9504	-5.1	<a href="#">TFFPEEISSMVLTKMK</a>
	1938.0423	-1.3	<a href="#">DNNLLGKFDLTGIPPAPR</a>
	1993.9295	3.2	<a href="#">FEDATVQSDMKHWPFR</a>
	2156.0839	-1.7	<a href="#">TTPSYVAFTDTERLIGDAAK</a>
	2332.1915	-2.4	<a href="#">LIGDAAKNQVAMNPTNTIFDAK</a>
	2697.3134	0.51	<a href="#">VEIANDQGNRTTPSYVAFTDTER</a>
	2873.4820	-2.7	<a href="#">GVPOIEVTFDIDANGILNVTAADKSTGK</a>
	2935.4988	1.2	<a href="#">EIAEAYLGGKVHSAVITVPAFENDSOR</a>
	2983.4438	0.46	<a href="#">TLSSSTQASIEIDSLYEGVDFYTSITR</a>
	3139.5479	1.4	<a href="#">RTLSSSTQASIEIDSLYEGVDFYTSITR</a>
	3211.5232	0.45	<a href="#">QTQFTTYSDNQSSVLVQVYEGERAMTK</a>
	3236.7607	-3.0	<a href="#">SENVQDLLLLDVTPLSLGIETAGGVMTPLIK</a>
	3392.8718	0.098	SENVQDLLLLDVTPLSLGIETAGGVMTPLIKR

IP-6	Heat shock 60 kDa Protein 1/ HSPD1	743.4287	-2.2	<a href="#">GIIDPTK</a>
		785.5121	-2.0	<a href="#">IGIEIIK</a>
		833.3910	0.49	<a href="#">APGFGDNR</a>
		844.4951	-1.8	<a href="#">VGEVIVTK</a>
		901.5353	-0.62	<a href="#">LSDGVAVLK</a>
		912.5876	-0.69	<a href="#">VGLQVVAVK</a>
		960.5209	9.9	<a href="#">VTDALNATR</a>
		1190.6075	-1.3	<a href="#">EIGNIISDAMK</a>
		1215.6549	-3.0	<a href="#">NAGVEGSLIVEK</a>
		1344.7158	-0.40	<a href="#">TVIIEQSWGSPK</a>
		1389.7149	6.8	<a href="#">GYISPYFINTSK</a>
		1428.8130	-0.43	<a href="#">GVMLAVDAVIAELK</a>
		1504.7263	-20	<a href="#">TLNDELEIIEGMK</a>
		1674.6231	1.4	<a href="#">DPGMGAMGGMGGGMGGGMF</a>
		1919.0708	-0.33	<a href="#">ISSIQSIVPALEIANHR</a>
		2038.0236	0.19	<a href="#">IQEIEQLDVTTSEYEK</a>
		2113.1495	4.4	<a href="#">ALMLQGVDLLADAVAVTMGPK</a>
2365.3346	0.15	<a href="#">KPLVIIAEDVDGEALSTLVLNR</a>		
2482.4114	3.8	TALLDAAGVASLLTTAEVVVTEIPK		
IP-7	Glucose regulated protein 75 kDa/ MTHSP75	715.4151	-3.6	<a href="#">LVGMPAK</a>
		764.3143	-14	<a href="#">EMAGDNK</a>
		782.3365	5.7	<a href="#">YAEEDR</a>
		847.4190	11	<a href="#">DNMALQR</a>
		978.4590	2.4	<a href="#">AMQDAEVSK</a>
		993.4513	-1.6	<a href="#">YDDPEVQK</a>
		1123.4662	-18	<a href="#">EGSGSSGTGEQK</a>

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		1231.6621	-2.1	<a href="#">QAASSLQOASLK</a>
		1290.6810	0.29	<a href="#">VQOTVQDLFGR</a>
		1333.6282	-8.0	<a href="#">ETAENYLGHTAK</a>
		1341.6142	-3.4	<a href="#">ASNGDAWVEAHGK</a>
		1361.7403	-1.9	<a href="#">AQFEGIVTDLIR</a>
		1446.7520	-7.4	<a href="#">SDIGEVILVGGMTR</a>
		1450.7162	-1.1	<a href="#">TTPSVVAFTADGER</a>
		1473.7917	0.27	<a href="#">EQQIVIQSSGGLSK</a>
		1553.8296	-6.8	<a href="#">LYSPSQIGAFVLMK</a>
		1568.7713	0.26	<a href="#">QAVTNPNTFYATK</a>
		1592.9552	1.5	<a href="#">LLGQFTLIGIPPAPR</a>
		1645.8595	-13	<a href="#">VINEPTAAALAYGLDK</a>
		1694.8426	-4.5	<a href="#">NAVITVPAYFNSQR</a>
		1808.9124	5.2	<a href="#">SQVFSTAADGOTOVEIK</a>
		1856.9088	1.3	<a href="#">VEAVNMAEGIIHDTETK</a>
		2055.9217	-20	<a href="#">STNGDTFLGGEDFDQALLR</a>
		2251.2280	2.4	<a href="#">VIAVYDLGGGTFDISILEIQK</a>
		2309.2535	17	<a href="#">GVPQIEVTFDIDANGIVHWSAK</a>
IP-8	Heat shock 70 kDa protein 1-HOM/ HSP-HOM	804.4451	-2.0	<a href="#">ITITNDK</a>
		805.4104	-3.2	<a href="#">MVLDAEK</a>
		872.4773	5.0	<a href="#">GTLEPVEK</a>
		933.4807	-9.2	<a href="#">SVVSDEGLK</a>
		1125.5617	-6.7	<a href="#">LLODYFNDR</a>
		1183.6369	-9.0	<a href="#">FDLTGIPPAPR</a>
		1197.6909	-3.9	<a href="#">DAGVIAGLNVLRL</a>
		1234.6166	-25	<a href="#">LVSHFVEEFK</a>
		1263.6048	0.36	<a href="#">FNDPVVQADMK</a>

		1266.7564	31	<a href="#">IHDIVLVGGSTR</a>
		1287.6098	4.2	<a href="#">NALESYAFNMK</a>
		1487.7092	5.0	<a href="#">TTPSYVAFTDTER</a>
		1614.8483	24	<a href="#">AFYPEEISSMVLTK</a>
		1675.7386	4.4	<a href="#">ATAGDTHLGGEDFDNR</a>
		1676.8261	12	<a href="#">NQVAMNPQNTVFDAK</a>
IP-9	<b>Glucose regulated protein 58 kDa/ PDIA3</b>	782.3480	-12	<a href="#">TEEEFK</a>
		782.3556	-2.1	<a href="#">GSNYWR</a>
		787.4197	12	<a href="#">FLDAGHK</a>
		877.4880	-1.8	<a href="#">LNFAVASR</a>
		939.4772	-1.6	<a href="#">TVAYTEQK</a>
		995.5532	-11	<a href="#">QAGPASVPLR</a>
		1040.5724	-1.6	<a href="#">TADGIVSHLK</a>
		1084.5573	-9.8	<a href="#">YGVSGYPTLK</a>
		1172.5304	-9.1	<a href="#">FVMQEEFSR</a>
		1191.6014	0.36	<a href="#">LAPEYEAAATR</a>
		1236.5117	-1.3	<a href="#">DGEEAGAYDGPR</a>
		1341.6737	-7.9	<a href="#">GFPTIYFSPANK</a>
		1359.6569	-1.2	<a href="#">FLQDYFDGNLK</a>
		1368.6631	-1.2	<a href="#">SEPIPESNDGPVK</a>
		1370.6840	-8.5	<a href="#">ELSDFISYLQR</a>
		1619.7819	-1.6	<a href="#">DLLIAYYDVDYEK</a>
		1664.7574	-0.97	<a href="#">MDATANDVPSPYEVR</a>
		1832.9124	5.1	<a href="#">VVVAENFDEIVNNENK</a>
		2348.0727	0.18	<a href="#">DASIVGFFDDSFSEAHSEFLK</a>
IP-	<b>Heat shock cognate protein 8 isoform 1, 71</b>	705.3453		<a href="#">DISENK</a>

10	<b>kDa/ HSPA8</b>		4.8	
		765.4101	-6.0	<a href="#">VQVEYK</a>
		774.4306	-7.2	<a href="#">NTTIPTK</a>
		791.3701	-11	<a href="#">ATVEDEK</a>
		804.4161	-38	<a href="#">ITITNDK</a>
		858.4557	-1.8	<a href="#">GTLDPVEK</a>
		993.5551	30	<a href="#">EIAEAYLGK</a>
		1074.4950	8.7	<a href="#">EEFEHQOK</a>
		1081.5676	-0.57	<a href="#">LLQDFENGK</a>
		1197.6676	3.7	<a href="#">FELTGIPPAPR</a>
		1199.6942	16	<a href="#">DAGTIAGLNVLR</a>
		1228.6230	-4.5	<a href="#">VEIANDQGNR</a>
		1235.6201	-3.7	<a href="#">MVNHFAEFK</a>
		1253.6460	23	<a href="#">FEELNADLFR</a>
		1254.5680	0.30	<a href="#">FDDAVVQSDMK</a>
		1303.5937	-4.3	<a href="#">NSLESYAFNMK</a>
		1481.8170	6.3	<a href="#">SQIHDIVLVGGSTR</a>
		1487.7082	4.3	<a href="#">TTPSYVAFTDTER</a>
		2260.1465	0.17	<a href="#">SINPDEAVAYGAAVQAAILSGDK</a>
		2514.3385	12	<a href="#">GVPQIEVTFDIDANGILNVSAVDK</a>
		2774.3257	-0.59	<a href="#">QTQFTTYSNQPGLIQVYEGER</a>
		2997.4175	-14	<a href="#">TLSSSTQASIEIDSLYEGIDFYTSITR</a>
		3346.4940	-1.7	<a href="#">LYQSAGGMPGGMPGGFPGGGAPPSGGASSGPTIEEVD</a>
IP-11	<b>BAG-3</b>	801.4706	-2.0	<a href="#">VEAILEK</a>
		990.4629	-1.6	<a href="#">TTTWNDPR</a>
		1151.6195	8.2	<a href="#">DPLPPGWEIK</a>
		1302.6640	-1.2	<a href="#">YLMIEEYLTK</a>
		1413.7119	-7.5	<a href="#">ELLALDSVDPEGR</a>

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		1481.7485	-1.1	<a href="#">IQGDDWEPRPLR</a>
		1915.9185	4.9	<a href="#">IDPQTGWPFVVDHNSR</a>
IP-12	<b><math>\alpha</math>-tubulin isoform 1/ TUBA1B</b>	775.4057	-5.9	<a href="#">GHYTIGK</a>
		781.4190	12	<a href="#">LSVDYGK</a>
		887.4341	0.45	<a href="#">FDLMYAK</a>
		906.4277	3.8	<a href="#">EDMAALEK</a>
		1015.5482	-30	<a href="#">DVNAAIATIK</a>
		1023.4460	-3.5	<a href="#">EDAANNYAR</a>
		1085.6501	27	<a href="#">EIIDLVLDR</a>
		1410.7769	1.7	<a href="#">QLFHPEQLITGK</a>
		1487.8771	-1.8	<a href="#">LISQIVSSITASLR</a>
		1701.9157	5.5	<a href="#">AVFVDLEPTVIDEVR</a>
		1718.8870	2.6	<a href="#">NLDIERPTYTNLNR</a>
		1756.9232	-23	<a href="#">IHFPLATYAPVISA EK</a>
		1824.9894	1.9	<a href="#">VGINYQPPTVVPGGDLAK</a>
		2007.8910	-1.3	<a href="#">TIGGGDDSFNTFFSETGAGK</a>
		2330.0382	8.3	<a href="#">AFVHWYVGEGMEEGEFSEAR</a>
		2348.9028	-1.1	<a href="#">DYEEVGVDSEGE GEEEGEEY</a>
		2409.2184	3.9	<a href="#">FDGALNVDLTEFQTNLVPYPR</a>
IP-13	<b><math>\beta</math> 5-tubulin/ TUBB</b>	738.3509	-3.5	<a href="#">GSQQYR</a>
		808.3451	-3.2	<a href="#">EEYPDR</a>
		1039.5925	-1.5	<a href="#">YLTVA AVER</a>
		1143.6393	3.9	<a href="#">LAVNMVPEPR</a>
		1301.6302	-5.8	<a href="#">ISVYYNEATGGK</a>

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		1319.7128	7.2	<a href="#">IMNTFSVVPSPK</a>
		1446.6803	-6.6	<a href="#">EVDEQMLNVQNK</a>
		1615.8259	-6.6	<a href="#">AILVDLEPGTMDSVR</a>
		1620.8345	-0.97	<a href="#">LHFFMPGFAPLTSR</a>
		1659.8752	-12	<a href="#">ALTVPELTQQVFDAK</a>
		1696.8309	-1.5	<a href="#">NSSYFVEWIPNNVK</a>
		1869.9679	-5.6	<a href="#">MAVTFIGNSTAIQELFK</a>
		1958.9827	0.18	<a href="#">GHYTEGAELVDSVLDVVR</a>
		2798.3533	3.4	<a href="#">SGPFGQIFRPDNFVFGQSGAGNNWAK</a>
		3102.4775	22	<a href="#">FWEVISDEHGIDPTGTYHGSDQLDR</a>
IP-14	Vimentin/ VIM	701.3950	0.55	<a href="#">SSVPGVR</a>
		870.4306	-6.4	<a href="#">FANYIDK</a>
		905.4023	-34	<a href="#">GTNESLER</a>
		906.4619	-7.3	<a href="#">FLEQQNK</a>
		912.4027	-3.9	<a href="#">QESTEYR</a>
		914.4558	-2.8	<a href="#">SYVTTSTR</a>
		1023.5155	4.3	<a href="#">QQYESVAAK</a>
		1046.5209	-9.1	<a href="#">LOBEMLQR</a>
		1060.5769	47	<a href="#">KLEGEESR</a>
		1076.5041	-0.50	<a href="#">DNLAEDIMR</a>
		1115.5491	-18	<a href="#">VELQELNDR</a>
		1121.5897	4.8	<a href="#">EYQDLLNVK</a>
		1169.7540	34	<a href="#">ILLAELEOLK</a>
		1254.5680	0.30	<a href="#">LGDLYEEEMR</a>
		1295.6164	-39	<a href="#">MALDIEIATYR</a>
		1309.6099	2.6	<a href="#">NLQEAEWYK</a>
		1323.6275	7.1	<a href="#">EEAENTLQSF</a>

		1405.7503	-5.4	<a href="#">VESLQEEIAFLK</a>
		1428.7147	1.7	<a href="#">SLYASSPGGVYATR</a>
		1494.7188	4.3	<a href="#">MFGGPGTASRPSSSR</a>
		1495.7963	6.3	<a href="#">TYSLGSALRPSTSR</a>
		1570.8981	1.5	<a href="#">ISLPLPNFSSLNLR</a>
		1668.8539	38 1Met-ox	LGDLYEEEMRELR
		1836.7294	-38	<a href="#">DGOVINETSQHDDLE</a>
		2186.9698	1.5	<a href="#">EMEENFAVEAANYQDTIGR</a>
		3923.0244	2.4	LHEEEIQELQAQIQEQHVQIDVDVSKPDLTAALR
IP-15	Cytokeratin 18/ KRT18	718.3716	-29	<a href="#">IMADIR</a>
		807.3925	-9.4	<a href="#">LAADDFR</a>
		884.4168	6.1	<a href="#">NHEEEVK</a>
		965.4667	-2.6	<a href="#">AQYDELAR</a>
		975.4230	-31	<a href="#">STFSTNYR</a>
		992.4875	-2.5	<a href="#">VVSETNDTK</a>
		1012.4748	-2.5	<a href="#">YETELAMR</a>
		1012.4748	-31	<a href="#">KNHEEEVK</a>
		1046.5176	-29	<a href="#">VIDDTNITR</a>
		1065.5585	0.42	<a href="#">LEAEIATYR</a>
		1267.6289	-8.4	<a href="#">QSVENDIHGLR</a>
		1286.7212	0.34	<a href="#">LQLETEIEALK</a>
		1506.7498	1.6	<a href="#">TVQSLEIDLDSMR</a>
		1884.0172	5.0	<a href="#">GLOAQIASSGLTVEVDAPK</a>
		2261.1062	3.4	<a href="#">YETELAMROSVENDIHGLR</a>
		2261.1062	1.9	<a href="#">GGMGSGGLATGIAGGLAGMGGIQNEK</a>
		2670.3959	1.3	<a href="#">YALQMEQLNGILLHLESELAQTR</a>
		2740.2627	0.13	LLEDGEDFNLDALDSSNSMQTIQK

... [9]

IP-16	Heat shock 27kDa protein 1/ HSPB1	721.3302	-9.2	<a href="#">SDETAAK</a>
		831.5287	23	<a href="#">VPFSLLR</a>
		917.4978	3.7	<a href="#">DGVVEITGK</a>
		941.4710	0.45	<a href="#">AQLGGPEAAK</a>
		987.6198	9.6	<a href="#">RVPFSLLR</a>
		1075.5752	0.41	<a href="#">QLSSGVSEIR</a>
		1104.5168	8.5	<a href="#">QDEHGYISR</a>
		1146.6394	2.1	<a href="#">TKDGVVEITGK</a>
		1163.6217	0.33	<a href="#">LFDQAFGLPR</a>
		1468.7290	-7.2	<a href="#">DGVVEITGKHEER</a>
		1655.7580	3.2	<a href="#">HEERQDEHGYISR</a>
		1773.9234	-12	<a href="#">VPFSLLRGPSWDPER</a>
		2013.0601	-2.8	<a href="#">VSLDVNHFAPDELTVKTK</a>
		2105.0321	-1.7	<a href="#">DWYPHSRLFDQAFGLPR</a>

... [10]

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1659.8152	-49	IINEPTAAAIAAYGLDK
Page 3: [2] Deleted	osingh1	12/7/2007 8:10 AM
1091.4212	-46	SEDEANRDR
Page 4: [3] Deleted	osingh1	12/7/2007 8:10 AM
1456.8818	48	GQIQEIVLVGGSTR
Page 4: [4] Deleted	osingh1	12/7/2007 8:11 AM
1659.8252	-43	IINEPTAAAIAAYGLDK
Page 5: [5] Deleted	osingh1	12/7/2007 8:11 AM
1233.5457	-41	VGGTSDVEVNEK
Page 8: [6] Deleted	osingh1	12/7/2007 8:12 AM
1981.9177	-41	TVTNAVVTVPAYFNDSQR
Page 9: [7] Deleted	osingh1	12/7/2007 8:12 AM
1130.5352	-54	FPGQLNADLR
Page 9: [8] Deleted	osingh1	12/7/2007 8:12 AM
1229.5383	-49	ISEQFTAMFR
Page 11: [9] Deleted	osingh1	12/7/2007 8:13 AM
848.4508	46	SLETENR
Page 12: [10] Deleted	osingh1	12/7/2007 8:13 AM
960.4722	41	DWYPHSR