

Supplemental D – List of all peptides, transition ions, and collision energies chosen for MRM analysis of the second set of patient and control pooled sweat samples.

Precursor	Transition	CE	Peptide	Protein
704.87	668.36	27.28	YSLTYIYTGLSK	alpha-2-glycoprotein1_zinc
704.87	781.44	27.28	YSLTYIYTGLSK	alpha-2-glycoprotein1_zinc
704.87	944.51	27.28	YSLTYIYTGLSK	alpha-2-glycoprotein1_zinc
704.87	1045.56	27.28	YSLTYIYTGLSK	alpha-2-glycoprotein1_zinc
704.87	1158.64	27.28	YSLTYIYTGLSK	alpha-2-glycoprotein1_zinc
726.34	557.34	28.01	AYLEEECPATLR	alpha-2-glycoprotein1_zinc
726.34	717.37	28.01	AYLEEECPATLR	alpha-2-glycoprotein1_zinc
726.34	846.41	28.01	AYLEEECPATLR	alpha-2-glycoprotein1_zinc
726.34	975.46	28.01	AYLEEECPATLR	alpha-2-glycoprotein1_zinc
726.34	1104.5	28.01	AYLEEECPATLR	alpha-2-glycoprotein1_zinc
501.3	333.19	20.36	VLTEIIASR	annexin5
501.3	446.27	20.36	VLTEIIASR	annexin5
501.3	559.36	20.36	VLTEIIASR	annexin5
501.3	688.4	20.36	VLTEIIASR	annexin5
501.3	789.45	20.36	VLTEIIASR	annexin5
670.81	621.29	26.12	GTVTDFPGFDER	annexin5
670.81	720.33	26.12	GTVTDFPGFDER	annexin5
670.81	867.4	26.12	GTVTDFPGFDER	annexin5
670.81	982.43	26.12	GTVTDFPGFDER	annexin5
670.81	1083.47	26.12	GTVTDFPGFDER	annexin5
852.95	476.28	32.31	GLGTDEESILTLLTSR	annexin5
852.95	589.37	32.31	GLGTDEESILTLLTSR	annexin5
852.95	690.41	32.31	GLGTDEESILTLLTSR	annexin5
852.95	803.5	32.31	GLGTDEESILTLLTSR	annexin5
852.95	1003.61	32.31	GLGTDEESILTLLTSR	annexin5
557.3	472.2	22.26	GGVEEGPTVLR	arginase_typel
557.3	585.37	22.26	GGVEEGPTVLR	arginase_typel
557.3	642.39	22.26	GGVEEGPTVLR	arginase_typel
557.3	771.44	22.26	GGVEEGPTVLR	arginase_typel
557.3	900.48	22.26	GGVEEGPTVLR	arginase_typel
889.48	563.33	33.56	TVNTAVAITLACFGLAR	arginase_typel
889.48	723.36	33.56	TVNTAVAITLACFGLAR	arginase_typel
889.48	794.4	33.56	TVNTAVAITLACFGLAR	arginase_typel
889.48	907.48	33.56	TVNTAVAITLACFGLAR	arginase_typel
889.48	1008.53	33.56	TVNTAVAITLACFGLAR	arginase_typel
636.36	574.33	24.95	VGGVQSLGGTGALR	aspartate_aminotransferase1
636.36	631.35	24.95	VGGVQSLGGTGALR	aspartate_aminotransferase1
636.36	744.44	24.95	VGGVQSLGGTGALR	aspartate_aminotransferase1

636.36	831.47	24.95	VGGVQSLGGTGALR	aspartate_aminotransferase1
636.36	959.53	24.95	VGGVQSLGGTGALR	aspartate_aminotransferase1
815.92	622.29	31.06	EPESILQVLSQMEK	aspartate_aminotransferase1
815.92	735.37	31.06	EPESILQVLSQMEK	aspartate_aminotransferase1
815.92	797.4	31.06	EPESILQVLSQMEK	aspartate_aminotransferase1
815.92	962.5	31.06	EPESILQVLSQMEK	aspartate_aminotransferase1
815.92	1075.58	31.06	EPESILQVLSQMEK	aspartate_aminotransferase1
720.32	472.2	27.8	DGEAVWFGCDVGK	bleomycin_hydrolase [homo_sapiens]
720.32	635.28	27.8	DGEAVWFGCDVGK	bleomycin_hydrolase [homo_sapiens]
720.32	658.28	27.8	DGEAVWFGCDVGK	bleomycin_hydrolase [homo_sapiens]
720.32	782.35	27.8	DGEAVWFGCDVGK	bleomycin_hydrolase [homo_sapiens]
720.32	968.43	27.8	DGEAVWFGCDVGK	bleomycin_hydrolase [homo_sapiens]
801.4	289.16	30.56	LYTVEYLSNMVGGR	bleomycin_hydrolase [homo_sapiens]
801.4	633.31	30.56	LYTVEYLSNMVGGR	bleomycin_hydrolase [homo_sapiens]
801.4	720.35	30.56	LYTVEYLSNMVGGR	bleomycin_hydrolase [homo_sapiens]
801.4	996.49	30.56	LYTVEYLSNMVGGR	bleomycin_hydrolase [homo_sapiens]
801.4	1313.64	30.56	LYTVEYLSNMVGGR	bleomycin_hydrolase [homo_sapiens]
514.24	377.18	20.8	VNYEEFAR	calmodulin-like_skin_protein
514.24	393.22	20.8	VNYEEFAR	calmodulin-like_skin_protein
514.24	522.27	20.8	VNYEEFAR	calmodulin-like_skin_protein
514.24	651.31	20.8	VNYEEFAR	calmodulin-like_skin_protein
514.24	814.37	20.8	VNYEEFAR	calmodulin-like_skin_protein
609.83	322.19	24.05	AGLEDLQVAFR	calmodulin-like_skin_protein
609.83	393.22	24.05	AGLEDLQVAFR	calmodulin-like_skin_protein
609.83	492.29	24.05	AGLEDLQVAFR	calmodulin-like_skin_protein
609.83	848.46	24.05	AGLEDLQVAFR	calmodulin-like_skin_protein
609.83	977.5	24.05	AGLEDLQVAFR	calmodulin-like_skin_protein
943.41	679.34	35.39	AADTDGDGQVNYEEFVR	calmodulin-like3
943.41	842.4	35.39	AADTDGDGQVNYEEFVR	calmodulin-like3
943.41	1055.52	35.39	AADTDGDGQVNYEEFVR	calmodulin-like3
943.41	1240.6	35.39	AADTDGDGQVNYEEFVR	calmodulin-like3
943.41	1412.64	35.39	AADTDGDGQVNYEEFVR	calmodulin-like3
482.74	262.15	19.73	FQQAIDSR	caspase14_precursor
482.74	377.18	19.73	FQQAIDSR	caspase14_precursor
482.74	561.3	19.73	FQQAIDSR	caspase14_precursor
482.74	689.36	19.73	FQQAIDSR	caspase14_precursor
482.74	703.34	19.73	FQQAIDSR	caspase14_precursor
652.84	375.2	25.51	LENLFEALNNK	caspase14_precursor
652.84	559.32	25.51	LENLFEALNNK	caspase14_precursor
652.84	688.36	25.51	LENLFEALNNK	caspase14_precursor
652.84	835.43	25.51	LENLFEALNNK	caspase14_precursor
652.84	1062.56	25.51	LENLFEALNNK	caspase14_precursor
782.36	647.32	29.91	DPTAEQFQEELEK	caspase14_precursor

782.36	922.45	29.91	DPTAEQFQEELEK	caspase14_precursor
782.36	1050.51	29.91	DPTAEQFQEELEK	caspase14_precursor
782.36	1179.55	29.91	DPTAEQFQEELEK	caspase14_precursor
782.36	1250.59	29.91	DPTAEQFQEELEK	caspase14_precursor
508.28	227.17	20.6	IILQPCGSK	corneodesmosin_precursor
508.28	451.2	20.6	IILQPCGSK	corneodesmosin_precursor
508.28	548.25	20.6	IILQPCGSK	corneodesmosin_precursor
508.28	676.31	20.6	IILQPCGSK	corneodesmosin_precursor
508.28	789.39	20.6	IILQPCGSK	corneodesmosin_precursor
827.92	727.4	31.46	GSPGVPSFAAGPPISEGK	corneodesmosin_precursor
827.92	784.42	31.46	GSPGVPSFAAGPPISEGK	corneodesmosin_precursor
827.92	855.46	31.46	GSPGVPSFAAGPPISEGK	corneodesmosin_precursor
827.92	926.49	31.46	GSPGVPSFAAGPPISEGK	corneodesmosin_precursor
827.92	1257.65	31.46	GSPGVPSFAAGPPISEGK	corneodesmosin_precursor
670.36	844.48	26.11	PATPEIQEIVDK	cystatin_A
670.36	973.52	26.11	PATPEIQEIVDK	cystatin_A
670.36	1070.57	26.11	PATPEIQEIVDK	cystatin_A
670.36	1171.62	26.11	PATPEIQEIVDK	cystatin_A
670.36	1242.66	26.11	PATPEIQEIVDK	cystatin_A
678.86	801.41	26.4	TQVVAGTNYYIK	cystatin_A
678.86	858.44	26.4	TQVVAGTNYYIK	cystatin_A
678.86	929.47	26.4	TQVVAGTNYYIK	cystatin_A
678.86	1028.54	26.4	TQVVAGTNYYIK	cystatin_A
678.86	1127.61	26.4	TQVVAGTNYYIK	cystatin_A
564.77	513.31	22.52	ENAGEDPGLAR	dermcidin_preproprotein
564.77	628.34	22.52	ENAGEDPGLAR	dermcidin_preproprotein
564.77	757.38	22.52	ENAGEDPGLAR	dermcidin_preproprotein
564.77	814.4	22.52	ENAGEDPGLAR	dermcidin_preproprotein
564.77	885.44	22.52	ENAGEDPGLAR	dermcidin_preproprotein
818.42	760.36	31.14	YQGTILSIDDNLQR	desmoglein1_preproprotein
818.42	873.44	31.14	YQGTILSIDDNLQR	desmoglein1_preproprotein
818.42	960.47	31.14	YQGTILSIDDNLQR	desmoglein1_preproprotein
818.42	1073.56	31.14	YQGTILSIDDNLQR	desmoglein1_preproprotein
818.42	1344.71	31.14	YQGTILSIDDNLQR	desmoglein1_preproprotein
437.75	286.14	18.2	DGLILTSR	DJ-1_protein
437.75	363.2	18.2	DGLILTSR	DJ-1_protein
437.75	399.22	18.2	DGLILTSR	DJ-1_protein
437.75	476.28	18.2	DGLILTSR	DJ-1_protein
437.75	589.37	18.2	DGLILTSR	DJ-1_protein
830.42	861.47	31.55	DVVICPDASLEDAKK	DJ-1_protein
830.42	976.49	31.55	DVVICPDASLEDAKK	DJ-1_protein
830.42	1073.55	31.55	DVVICPDASLEDAKK	DJ-1_protein
830.42	1233.58	31.55	DVVICPDASLEDAKK	DJ-1_protein

830.42	1346.66	31.55	DVVICPDASLEDAKK	DJ-1_protein
373.23	276.15	16	IATAIEK	Enolase(Yeast)
373.23	389.24	16	IATAIEK	Enolase(Yeast)
373.23	460.28	16	IATAIEK	Enolase(Yeast)
373.23	561.32	16	IATAIEK	Enolase(Yeast)
373.23	632.36	16	IATAIEK	Enolase(Yeast)
404.22	288.2	17.06	TFAEALR	Enolase(Yeast)
404.22	359.24	17.06	TFAEALR	Enolase(Yeast)
404.22	488.28	17.06	TFAEALR	Enolase(Yeast)
404.22	559.32	17.06	TFAEALR	Enolase(Yeast)
404.22	706.39	17.06	TFAEALR	Enolase(Yeast)
407.76	373.28	17.18	AADALLLK	Enolase(Yeast)
407.76	486.36	17.18	AADALLLK	Enolase(Yeast)
407.76	557.4	17.18	AADALLLK	Enolase(Yeast)
407.76	672.43	17.18	AADALLLK	Enolase(Yeast)
407.76	743.47	17.18	AADALLLK	Enolase(Yeast)
625.82	516.23	24.59	TGEEVGFVVDK	filaminB_beta_(actin binding protein 278)
625.82	531.31	24.59	TGEEVGFVVDK	filaminB_beta_(actin binding protein 278)
625.82	735.4	24.59	TGEEVGFVVDK	filaminB_beta_(actin binding protein 278)
625.82	834.47	24.59	TGEEVGFVVDK	filaminB_beta_(actin binding protein 278)
625.82	963.51	24.59	TGEEVGFVVDK	filaminB_beta_(actin binding protein 278)
765.9	549.28	29.35	VPTANVSVDLTCR	glyceraldehyde-3-phosphate_dehydrogenase
765.9	664.31	29.35	VPTANVSVDLTCR	glyceraldehyde-3-phosphate_dehydrogenase
765.9	862.44	29.35	VPTANVSVDLTCR	glyceraldehyde-3-phosphate_dehydrogenase
765.9	949.48	29.35	VPTANVSVDLTCR	glyceraldehyde-3-phosphate_dehydrogenase
765.9	1048.55	29.35	VPTANVSVDLTCR	glyceraldehyde-3-phosphate_dehydrogenase
882.4	596.28	33.32	LISWYDNEFGYSNR	glyceraldehyde-3-phosphate_dehydrogenase
882.4	743.35	33.32	LISWYDNEFGYSNR	glyceraldehyde-3-phosphate_dehydrogenase
882.4	986.43	33.32	LISWYDNEFGYSNR	glyceraldehyde-3-phosphate_dehydrogenase
882.4	1101.46	33.32	LISWYDNEFGYSNR	glyceraldehyde-3-phosphate_dehydrogenase
882.4	1264.52	33.32	LISWYDNEFGYSNR	glyceraldehyde-3-phosphate_dehydrogenase
592.84	343.21	23.47	LLCGATLIAPR	kallikrein11_isoform1_preproprotein
592.84	456.29	23.47	LLCGATLIAPR	kallikrein11_isoform1_preproprotein
592.84	670.42	23.47	LLCGATLIAPR	kallikrein11_isoform1_preproprotein
592.84	798.48	23.47	LLCGATLIAPR	kallikrein11_isoform1_preproprotein
592.84	958.51	23.47	LLCGATLIAPR	kallikrein11_isoform1_preproprotein
658.36	459.27	25.7	SLDLDSIINAVR	keratin_1B
658.36	572.35	25.7	SLDLDSIINAVR	keratin_1B
658.36	685.44	25.7	SLDLDSIINAVR	keratin_1B
658.36	772.47	25.7	SLDLDSIINAVR	keratin_1B
658.36	887.49	25.7	SLDLDSIINAVR	keratin_1B
668.36	749.42	26.04	VDTLTGEVNFLK	keratin_1B
668.36	806.44	26.04	VDTLTGEVNFLK	keratin_1B

668.36	907.49	26.04	VDTLTGEVNFLK	keratin_1B
668.36	1020.57	26.04	VDTLTGEVNFLK	keratin_1B
668.36	1121.62	26.04	VDTLTGEVNFLK	keratin_1B
665.37	446.26	25.94	NLDLDSIIAEVK	keratin_2
665.37	559.34	25.94	NLDLDSIIAEVK	keratin_2
665.37	874.49	25.94	NLDLDSIIAEVK	keratin_2
665.37	987.57	25.94	NLDLDSIIAEVK	keratin_2
665.37	1102.6	25.94	NLDLDSIIAEVK	keratin_2
704.36	549.31	27.26	ADTLTDEINFLR	keratin_6A
704.36	662.4	27.26	ADTLTDEINFLR	keratin_6A
704.36	791.44	27.26	ADTLTDEINFLR	keratin_6A
704.36	906.47	27.26	ADTLTDEINFLR	keratin_6A
704.36	1007.52	27.26	ADTLTDEINFLR	keratin_6A
618.27	520.25	24.34	FSSSSGYGGGSSR	keratin_9
618.27	683.31	24.34	FSSSSGYGGGSSR	keratin_9
618.27	740.33	24.34	FSSSSGYGGGSSR	keratin_9
618.27	827.36	24.34	FSSSSGYGGGSSR	keratin_9
618.27	1001.43	24.34	FSSSSGYGGGSSR	keratin_9
497.25	359.24	20.22	YENEVALR	keratin10
497.25	458.31	20.22	YENEVALR	keratin10
497.25	587.35	20.22	YENEVALR	keratin10
497.25	701.39	20.22	YENEVALR	keratin10
497.25	830.44	20.22	YENEVALR	keratin10
691.33	575.3	26.82	ALEESNYELEGK	keratin10
691.33	852.41	26.82	ALEESNYELEGK	keratin10
691.33	939.44	26.82	ALEESNYELEGK	keratin10
691.33	1068.48	26.82	ALEESNYELEGK	keratin10
691.33	1197.53	26.82	ALEESNYELEGK	keratin10
617.36	417.25	24.3	IFGVTTLDIVR	mitochondrial_malate_dehydrogenase_precursor
617.36	502.3	24.3	IFGVTTLDIVR	mitochondrial_malate_dehydrogenase_precursor
617.36	716.43	24.3	IFGVTTLDIVR	mitochondrial_malate_dehydrogenase_precursor
617.36	817.48	24.3	IFGVTTLDIVR	mitochondrial_malate_dehydrogenase_precursor
617.36	973.57	24.3	IFGVTTLDIVR	mitochondrial_malate_dehydrogenase_precursor
897.05	783.53	33.81	VAVLGASGGIGQPLSLLK	mitochondrial_malate_dehydrogenase_precursor
897.05	968.61	33.81	VAVLGASGGIGQPLSLLK	mitochondrial_malate_dehydrogenase_precursor
897.05	1138.72	33.81	VAVLGASGGIGQPLSLLK	mitochondrial_malate_dehydrogenase_precursor
897.05	1195.74	33.81	VAVLGASGGIGQPLSLLK	mitochondrial_malate_dehydrogenase_precursor
897.05	1282.77	33.81	VAVLGASGGIGQPLSLLK	mitochondrial_malate_dehydrogenase_precursor
482.74	317.19	19.73	FEEELAAR	nucleobindin1
482.74	406.16	19.73	FEEELAAR	nucleobindin1
482.74	430.28	19.73	FEEELAAR	nucleobindin1
482.74	559.32	19.73	FEEELAAR	nucleobindin1
482.74	688.36	19.73	FEEELAAR	nucleobindin1

753.91	404.22	28.95	LVTLEEF LASTQR	nucleobindin1
753.91	491.26	28.95	LVTLEEF LASTQR	nucleobindin1
753.91	562.29	28.95	LVTLEEF LASTQR	nucleobindin1
753.91	945.53	28.95	LVTLEEF LASTQR	nucleobindin1
753.91	1016.57	28.95	LVTLEEF LASTQR	nucleobindin1
460.76	262.14	18.98	GLFIIDDK	peroxiredoxin1
460.76	377.17	18.98	GLFIIDDK	peroxiredoxin1
460.76	490.25	18.98	GLFIIDDK	peroxiredoxin1
460.76	603.33	18.98	GLFIIDDK	peroxiredoxin1
460.76	750.4	18.98	GLFIIDDK	peroxiredoxin1
598.82	510.26	23.67	LVQAFQFTDK	peroxiredoxin1
598.82	638.31	23.67	LVQAFQFTDK	peroxiredoxin1
598.82	785.38	23.67	LVQAFQFTDK	peroxiredoxin1
598.82	856.42	23.67	LVQAFQFTDK	peroxiredoxin1
598.82	984.48	23.67	LVQAFQFTDK	peroxiredoxin1
811.42	545.31	30.9	VTNGAFTGEISPGMIK	PREDICTED_similar_to_Triosephosphate_isomerase(TI
811.42	632.34	30.9	VTNGAFTGEISPGMIK	PREDICTED_similar_to_Triosephosphate_isomerase(TI
811.42	931.49	30.9	VTNGAFTGEISPGMIK	PREDICTED_similar_to_Triosephosphate_isomerase(TI
811.42	1032.54	30.9	VTNGAFTGEISPGMIK	PREDICTED_similar_to_Triosephosphate_isomerase(TI
811.42	1179.61	30.9	VTNGAFTGEISPGMIK	PREDICTED_similar_to_Triosephosphate_isomerase(TI
642.39	502.3	25.16	TVQIAAVVDVIR	prolactin-induced_protein
642.39	601.37	25.16	TVQIAAVVDVIR	prolactin-induced_protein
642.39	700.43	25.16	TVQIAAVVDVIR	prolactin-induced_protein
642.39	771.47	25.16	TVQIAAVVDVIR	prolactin-induced_protein
642.39	842.51	25.16	TVQIAAVVDVIR	prolactin-induced_protein
780.92	642.32	29.87	LYTLVLTPDAPSR	prostatic_binding_protein
780.92	757.35	29.87	LYTLVLTPDAPSR	prostatic_binding_protein
780.92	858.39	29.87	LYTLVLTPDAPSR	prostatic_binding_protein
780.92	971.48	29.87	LYTLVLTPDAPSR	prostatic_binding_protein
780.92	1070.55	29.87	LYTLVLTPDAPSR	prostatic_binding_protein
1035.47	922.43	38.52	APVAGTCYQAEWDDYVPK	prostatic_binding_protein
1035.47	1122.51	38.52	APVAGTCYQAEWDDYVPK	prostatic_binding_protein
1035.47	1148.5	38.52	APVAGTCYQAEWDDYVPK	prostatic_binding_protein
1035.47	1250.57	38.52	APVAGTCYQAEWDDYVPK	prostatic_binding_protein
1035.47	1413.63	38.52	APVAGTCYQAEWDDYVPK	prostatic_binding_protein
818.95	630.35	31.16	GVNLPGAAVDLPAVSEK	pyruvate_kinase3_isoform1
818.95	858.46	31.16	GVNLPGAAVDLPAVSEK	pyruvate_kinase3_isoform1
818.95	957.52	31.16	GVNLPGAAVDLPAVSEK	pyruvate_kinase3_isoform1
818.95	1028.56	31.16	GVNLPGAAVDLPAVSEK	pyruvate_kinase3_isoform1
818.95	1253.67	31.16	GVNLPGAAVDLPAVSEK	pyruvate_kinase3_isoform1
1088.06	808.42	40.31	LAPITSDPTEATAVGAVEASFK	pyruvate_kinase3_isoform1
1088.06	907.49	40.31	LAPITSDPTEATAVGAVEASFK	pyruvate_kinase3_isoform1
1088.06	978.52	40.31	LAPITSDPTEATAVGAVEASFK	pyruvate_kinase3_isoform1

1088.06	1150.61	40.31	LAPITSDPTEATAVGAVEASFK	pyruvate_kinase3_isoform1
1088.06	1477.75	40.31	LAPITSDPTEATAVGAVEASFK	pyruvate_kinase3_isoform1
628.81	423.22	24.69	GTNYLADVFEK	S100 calcium-binding_protein_A7
628.81	620.3	24.69	GTNYLADVFEK	S100 calcium-binding_protein_A7
628.81	708.36	24.69	GTNYLADVFEK	S100 calcium-binding_protein_A7
628.81	821.44	24.69	GTNYLADVFEK	S100 calcium-binding_protein_A7
628.81	984.5	24.69	GTNYLADVFEK	S100 calcium-binding_protein_A7
374.21	234.14	16.04	ELTIGSK	S100A6
374.21	291.17	16.04	ELTIGSK	S100A6
374.21	404.25	16.04	ELTIGSK	S100A6
374.21	505.3	16.04	ELTIGSK	S100A6
374.21	618.38	16.04	ELTIGSK	S100A6
446.21	290.15	18.49	LMEDLDR	S100A6
446.21	403.23	18.49	LMEDLDR	S100A6
446.21	518.26	18.49	LMEDLDR	S100A6
446.21	647.3	18.49	LMEDLDR	S100A6
446.21	778.34	18.49	LMEDLDR	S100A6
458.25	359.24	18.89	LQDAEIAR	S100A6
458.25	488.28	18.89	LQDAEIAR	S100A6
458.25	559.32	18.89	LQDAEIAR	S100A6
458.25	674.35	18.89	LQDAEIAR	S100A6
458.25	802.4	18.89	LQDAEIAR	S100A6
727.87	708.35	28.06	EEIVYLPCIYR	selenium_binding_protein1
727.87	821.43	28.06	EEIVYLPCIYR	selenium_binding_protein1
727.87	984.5	28.06	EEIVYLPCIYR	selenium_binding_protein1
727.87	1083.57	28.06	EEIVYLPCIYR	selenium_binding_protein1
727.87	1196.65	28.06	EEIVYLPCIYR	selenium_binding_protein1
454.73	389.21	18.77	VGEFSGANK	thioredoxin
454.73	476.25	18.77	VGEFSGANK	thioredoxin
454.73	623.31	18.77	VGEFSGANK	thioredoxin
454.73	752.36	18.77	VGEFSGANK	thioredoxin
454.73	809.38	18.77	VGEFSGANK	thioredoxin
668.82	576.26	26.05	TAFQEALDAAGDK	thioredoxin
668.82	689.35	26.05	TAFQEALDAAGDK	thioredoxin
668.82	760.38	26.05	TAFQEALDAAGDK	thioredoxin
668.82	889.43	26.05	TAFQEALDAAGDK	thioredoxin
668.82	1017.48	26.05	TAFQEALDAAGDK	thioredoxin
869.92	461.22	32.89	LVVVDFSATWCGPCK	thioredoxin
869.92	621.25	32.89	LVVVDFSATWCGPCK	thioredoxin
869.92	807.33	32.89	LVVVDFSATWCGPCK	thioredoxin
869.92	908.37	32.89	LVVVDFSATWCGPCK	thioredoxin
869.92	1066.44	32.89	LVVVDFSATWCGPCK	thioredoxin
793.87	585.27	30.31	DCGATWVVLGHSEK	triosephosphate_isomerase1

793.87	698.36	30.31	DCGATWVVLGHSER	triosephosphate_isomerase1
793.87	896.49	30.31	DCGATWVVLGHSER	triosephosphate_isomerase1
793.87	1082.57	30.31	DCGATWVVLGHSER	triosephosphate_isomerase1
793.87	1183.62	30.31	DCGATWVVLGHSER	triosephosphate_isomerase1
1096.54	1042.49	40.6	VPADTEVVCAPPTAYIDFAR	triosephosphate_isomerase1
1096.54	1150.59	40.6	VPADTEVVCAPPTAYIDFAR	triosephosphate_isomerase1
1096.54	1221.63	40.6	VPADTEVVCAPPTAYIDFAR	triosephosphate_isomerase1
1096.54	1381.66	40.6	VPADTEVVCAPPTAYIDFAR	triosephosphate_isomerase1
1096.54	1480.73	40.6	VPADTEVVCAPPTAYIDFAR	triosephosphate_isomerase1