

**Supplementary Table 6: Peptides in the AMT tag database: sequence, molecular weight, modification, corresponding protein groups and proteotypicity.**

Protein Group	AMT_ID	Mono-isotopic Mr	Sequence	Modification	Protein Group Count	Proteotypicity
1433B_HUMAN	21	902.507	VISSIEQK		4	Shared
1433B_HUMAN	33	906.517	NLLSVAYK		7	Shared
1433B_HUMAN	599	1078.395	YDDMAAAMK	Dioxidation (M); Dioxidation (M)	2	Shared
1433B_HUMAN	848	1139.539	EMQPThPIR	Dioxidation (M)	3	Shared
1433B_HUMAN	991	1181.556	YLSEVASGDNK		1	Proteotypic
1433B_HUMAN	1136	1220.643	DSTLIMQLLR	Dioxidation (M)	7	Shared
1433B_HUMAN	1249	1251.602	TMDKSELVQK	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
1433B_HUMAN	1296	1267.634	KEMQPThPIR	Dioxidation (M)	3	Shared
1433B_HUMAN	1928	1597.733	AVTEQGHLSNEER		1	Proteotypic
1433E_HUMAN	33	906.517	NLLSVAYK		7	Shared
1433E_HUMAN	345	1008.480	QMVETELK	Dioxidation (M)	1	Proteotypic
1433E_HUMAN	1136	1220.643	DSTLIMQLLR	Dioxidation (M)	7	Shared
1433E_HUMAN	1772	1478.692	VAGMDVELTVEER	Dioxidation (M)	1	Proteotypic
1433E_HUMAN	1880	1555.682	MDDREDLVYQAK	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
1433F_HUMAN	21	902.507	VISSIEQK		4	Shared
1433F_HUMAN	33	906.517	NLLSVAYK		7	Shared
1433F_HUMAN	913	1155.600	GDREQLLQR	Acetyl (Protein N-term)	1	Proteotypic
1433F_HUMAN	1136	1220.643	DSTLIMQLLR	Dioxidation (M)	7	Shared
1433F_HUMAN	1294	1267.598	EQMQPThPIR	Dioxidation (M)	1	Proteotypic
1433F_HUMAN	1915	1585.758	AVTELNEPLSNEDR		1	Proteotypic
1433G_HUMAN	21	902.507	VISSIEQK		4	Shared
1433G_HUMAN	33	906.517	NLLSVAYK		7	Shared
1433G_HUMAN	599	1078.395	YDDMAAAMK	Dioxidation (M); Dioxidation (M)	2	Shared
1433G_HUMAN	914	1155.625	VDREQLVQK	Acetyl (Protein N-term)	1	Proteotypic
1433G_HUMAN	1136	1220.643	DSTLIMQLLR	Dioxidation (M)	7	Shared
1433G_HUMAN	1974	1642.780	NVTELNEPLSNEER		1	Proteotypic
1433S_HUMAN	22	902.507	VLSSIEQK		1	Proteotypic
1433S_HUMAN	33	906.517	NLLSVAYK		7	Shared
1433S_HUMAN	1136	1220.643	DSTLIMQLLR	Dioxidation (M)	7	Shared
1433T_HUMAN	21	902.507	VISSIEQK		4	Shared
1433T_HUMAN	33	906.517	NLLSVAYK		7	Shared
1433T_HUMAN	274	988.519	EKVESELR		1	Proteotypic
1433T_HUMAN	848	1139.539	EMQPThPIR	Dioxidation (M)	3	Shared
1433T_HUMAN	1136	1220.643	DSTLIMQLLR	Dioxidation (M)	7	Shared
1433T_HUMAN	1296	1267.634	KEMQPThPIR	Dioxidation (M)	3	Shared
1433T_HUMAN	1844	1531.711	AVTEQGAELSNEER		1	Proteotypic
1433Z_HUMAN	33	906.517	NLLSVAYK		7	Shared
1433Z_HUMAN	848	1139.539	EMQPThPIR	Dioxidation (M)	3	Shared
1433Z_HUMAN	984	1177.565	MDKNELVQK	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic

1433Z_HUMAN	1136	1220.643	DSTLIMQLLR	Dioxidation (M)	7	Shared
1433Z_HUMAN	1296	1267.634	KEMQPTHPIR	Dioxidation (M)	3	Shared
1433Z_HUMAN	1331	1278.646	YLAEVAAGDDKK		1	Proteotypic
1433Z_HUMAN	1402	1303.677	FLIPNASQAESK		1	Proteotypic
1433Z_HUMAN	1630	1408.676	DICNDVLSLLEK	Trioxidation (C)	1	Proteotypic
1433Z_HUMAN	1871	1547.706	SVTEQGAELSNEER		1	Proteotypic
1433Z_HUMAN	2272	2039.980	GIVDQSQQAYQEAFEISK		1	Proteotypic
1B07_HUMAN	2393	964.425	FDSDAASPR		1	Proteotypic
2B1B_HUMAN	2411	1070.467	FSDVGEFR		1	Proteotypic
3HIDH_HUMAN	2493	1755.792	MGAVFMDAPVSGGVGAAR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A0A5C9_HUMAN	280	989.503	AGVETTPSK		1	Proteotypic
A0A5C9_HUMAN	2249	1985.011	AAPSVTLFPPSSEELQANK		1	Proteotypic
A0A5E5_HUMAN	2120	1787.851	SGTASVCLLNNFYPR	Trioxidation (C)	1	Proteotypic
A0A5E5_HUMAN	2183	1865.883	VYACEVTHQGLSSPVTK	Trioxidation (C)	1	Proteotypic
A0A5E5_HUMAN	2233	1945.020	TVAAPSVFIFPPSDEQLK		1	Proteotypic
A0A5E5_HUMAN	2300	2134.961	VDNALQSGNSQESVTEQDSK		1	Proteotypic
A0AVI3_HUMAN	163	952.596	LLLPGELAK		2	Shared
A0AVI3_HUMAN	952	1167.588	QVHPDTGISSK		1	Proteotypic
A0AVI3_HUMAN	2088	1742.812	AMGIMNSFVNDIFER		1	Proteotypic
A0AVI3_HUMAN	2123	1790.797	AMGIMNSFVNDIFER	Dioxidation (M); Oxidation (M)	1	Proteotypic
A0AVI3_HUMAN	2134	1806.792	AMGIMNSFVNDIFER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A0JNT2_HUMAN	29	905.461	FLEQQNK		14	Shared
A0JNT2_HUMAN	214	972.488	LLEGEEQR		1	Proteotypic
A0JNT2_HUMAN	1233	1244.636	TKEEINELNR		2	Shared
A0JNT2_HUMAN	1453	1319.708	ATAENEFVALKK		1	Proteotypic
A0S2P1_HUMAN	715	1105.686	VVLPPIEAPIR		1	Proteotypic
A0S2P1_HUMAN	2191	1882.923	ILYMTDEVNDPSLTIK	Dioxidation (M)	2	Shared
A1A4E9_HUMAN	380	1016.575	VDELTLISK		1	Proteotypic
A1A4E9_HUMAN	463	1041.582	VILEIDNAR		2	Shared
A1A4E9_HUMAN	771	1119.534	ITMQNLNDR	Oxidation (M)	2	Shared
A1A4E9_HUMAN	779	1121.572	LEQEIATYR		6	Shared
A1A4E9_HUMAN	836	1135.529	ITMQNLNDR	Dioxidation (M)	2	Shared
A1A4E9_HUMAN	865	1143.625	ILTATIENNR		1	Proteotypic
A1A4E9_HUMAN	1066	1200.610	QSVEADINGLR		2	Shared
A1A4E9_HUMAN	1393	1300.651	ALEEANADLEVK		4	Shared
A1A4E9_HUMAN	1536	1356.711	QSVEADINGLRR		2	Shared
A1A4E9_HUMAN	1694	1440.677	EVSTNTAMIQTSK	Dioxidation (M)	1	Proteotypic
A1A4E9_HUMAN	2174	1848.950	TLQGLEIIEIQSLSMK	Dioxidation (M)	2	Shared
A1A5C4_HUMAN	382	1017.509	LQSSEAEVR		1	Proteotypic
A1A5C4_HUMAN	488	1046.499	AENSQLTER		1	Proteotypic
A1A5C4_HUMAN	571	1072.540	VGAEEELQK		1	Proteotypic
A1A5C4_HUMAN	687	1099.599	LQQENSILR		1	Proteotypic
A1A5C4_HUMAN	863	1143.588	EQEITAVQAR		1	Proteotypic
A1A5C4_HUMAN	1263	1256.672	SIEALLEAGQAR		1	Proteotypic
A1A5C4_HUMAN	1539	1357.709	LLATEQEDAATAVAK		1	Proteotypic
A1A5C4_HUMAN	1601	1389.674	DALNQATSQVESK		1	Proteotypic

A1A5C4_HUMAN	1865	1542.825	QLLLESQSQLDAAK		1	Proteotypic
A1AT_HUMAN	374	1014.607	SVLQGQGITK		1	Proteotypic
A1AT_HUMAN	2163	1832.916	VFSNGADLSGVTEEAPLK		1	Proteotypic
A1L3W5_HUMAN	2446	1233.635	VAGQDGSVVQFK		1	Proteotypic
A1L400_HUMAN	2497	1840.884	STGEAFVQFASQEIAEK		1	Proteotypic
A2A274_HUMAN	1738	1462.742	SQFTITPGSEQIR		1	Proteotypic
A2A274_HUMAN	1932	1600.785	NAVVTQEFQVVPDTR		1	Proteotypic
A2A2R6_HUMAN	2406	1056.618	LLLLGAGESGK		1	Proteotypic
A2A418_HUMAN	52	912.576	RTPITVVK		1	Proteotypic
A2A418_HUMAN	582	1073.510	EGGQTAPASTR		1	Proteotypic
A2A418_HUMAN	1451	1318.688	AGALNSNDAFVLK		1	Proteotypic
A2A418_HUMAN	2158	1828.957	QTQVSVLPEGGETPLFK		1	Proteotypic
A2AB90_HUMAN	2417	1089.585	VSLMPATLAR	Dioxidation (M)	1	Proteotypic
A2ACR1_HUMAN	320	1000.530	VSAGEAVVNR		1	Proteotypic
A2ACR1_HUMAN	1473	1327.644	FTTDAIALMSR	Dioxidation (M)	1	Proteotypic
A2BDF6_HUMAN	180	958.581	STELLIRK		1	Proteotypic
A2BDF6_HUMAN	1488	1334.683	EIAQDFKTDLR		1	Proteotypic
A2BEY0_HUMAN	2431	1162.635	LSAAVTEAFVR		1	Proteotypic
A2BFA2_HUMAN	452	1038.586	YLTVAAVFR		2	Shared
A2BFA2_HUMAN	592	1076.525	IREEYPDR		4	Shared
A2BFA2_HUMAN	769	1119.363	NMMAACDPR	Dioxidation (M); Dioxidation (M); Trioxidation (C)	5	Shared
A2BFA2_HUMAN	808	1129.588	FPGQLNADLR		5	Shared
A2BFA2_HUMAN	928	1158.622	LAVNMVPFPR	Oxidation (M)	5	Shared
A2BFA2_HUMAN	971	1174.617	LAVNMVPFPR	Dioxidation (M)	5	Shared
A2BFA2_HUMAN	1264	1257.683	FPGQLNADLRK		5	Shared
A2BFA2_HUMAN	1275	1260.581	ISEQFTAMFR	Dioxidation (M)	4	Shared
A2BFA2_HUMAN	1398	1302.712	KLAVNMVPFPR	Dioxidation (M)	5	Shared
A2BFA2_HUMAN	1472	1327.641	INVYYNEATGGK		1	Proteotypic
A2BFA2_HUMAN	1522	1350.685	IMNTFSVVPSPK	Dioxidation (M)	3	Shared
A2BFA2_HUMAN	1769	1477.672	EVDEQMLNVQNK	Dioxidation (M)	3	Shared
A2BFA2_HUMAN	1965	1632.803	AVLVDLEPGTMDSVR	Dioxidation (M)	1	Proteotypic
A2BFA2_HUMAN	2068	1722.850	ALTVPELTQQMFDAK	Dioxidation (M)	3	Shared
A2BFA2_HUMAN	2267	2018.859	MSMKEVDEQMLNVQNK	Dioxidation (M); Dioxidation (M); Dioxidation (M)	3	Shared
A2RRC9_HUMAN	414	1027.603	LEALGLVSAR		1	Proteotypic
A2RRC9_HUMAN	892	1149.585	LIFQMPQNK	Dioxidation (M)	3	Shared
A2RTY6_HUMAN	580	1073.473	MATTTMIQSK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A2RTY6_HUMAN	1063	1200.508	SSALDMENFR	Dioxidation (M)	1	Proteotypic
A2RTY6_HUMAN	1910	1581.847	IQPSGGTNINEALLR		1	Proteotypic
A2RUU6_HUMAN	136	943.524	AGLQFPVGR		3	Shared
A2RUU6_HUMAN	1312	1271.672	NDEELNKLLGK		1	Proteotypic
A2RUU6_HUMAN	2221	1930.162	VTIAQGGVLPNIQAVLLPK		2	Shared
A2VCL0_HUMAN	276	988.571	VFLENVIR		1	Proteotypic
A2VCL0_HUMAN	828	1133.535	DAVTYTEHAK		1	Proteotypic
A2VCL0_HUMAN	989	1179.614	ISGLIYEETR		1	Proteotypic
A2VCL0_HUMAN	1468	1324.746	DNIQGITKPAIR		1	Proteotypic
A2VCL0_HUMAN	1504	1341.685	TVTAMDVVYALK	Dioxidation (M)	1	Proteotypic

A2VCL5_HUMAN	835	1135.529	ARAEAEQMAK	Dioxidation (M)	1	Proteotypic
A2VCL5_HUMAN	1118	1214.687	VTGEVLDILTR		1	Proteotypic
A3KLL5_HUMAN	2475	1488.866	VAPPGLTQIPQIQK		1	Proteotypic
A3RJH1_HUMAN	2422	1110.676	ELLIIGGVAAR		1	Proteotypic
A4D0V4_HUMAN	2437	1196.688	LLLNNNDNLLR		1	Proteotypic
A4D0Z3_HUMAN	646	1088.623	DAVLLVFANK		2	Shared
A4D0Z3_HUMAN	767	1118.601	ILMVGLDAAGK	Dioxidation (M)	3	Shared
A4D0Z3_HUMAN	868	1145.556	VQESADELQK		1	Proteotypic
A4D0Z3_HUMAN	2302	2135.093	MLQEDELDRDAVLLVFANK	Dioxidation (M)	1	Proteotypic
A4D0Z3_HUMAN	2336	2322.236	LGEIVTTIPTIGFNVETVEYK		3	Shared
A4D177_HUMAN	2495	1775.832	IIGATDSSGELMFLMK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A4D1N4_HUMAN	138	944.493	LSENVIDR		1	Proteotypic
A4D1N4_HUMAN	216	972.499	AAANEQLTR		1	Proteotypic
A4D1N4_HUMAN	1061	1199.640	VAEELALEQAK		1	Proteotypic
A4D1P1_HUMAN	449	1036.544	ATFVELSTK	Acetyl (Protein N-term)	1	Proteotypic
A4D1P1_HUMAN	711	1105.613	REDLFIVSK		1	Proteotypic
A4D1P1_HUMAN	1523	1350.718	IVENIQVDFK		1	Proteotypic
A4D1P1_HUMAN	2000	1656.766	LSDEEMATILSFNR	Dioxidation (M)	1	Proteotypic
A4D1Y2_HUMAN	2412	1073.547	ILEQQNSSR		1	Proteotypic
A4D204_HUMAN	2405	1056.618	ILLLGAGESGK		1	Proteotypic
A4D208_HUMAN	649	1089.585	VTLMQLPTR	Dioxidation (M)	1	Proteotypic
A4D208_HUMAN	1546	1359.601	AQAVSEDAGGNEGR		1	Proteotypic
A4D2J6_HUMAN	340	1006.475	AMEAVAAQ GK	Dioxidation (M)	1	Proteotypic
A4D2J6_HUMAN	833	1134.570	KAMEAVAAQ GK	Dioxidation (M)	1	Proteotypic
A4D2J6_HUMAN	894	1149.662	VLIAAHGNSLR		1	Proteotypic
A4GX73_HUMAN	887	1148.667	VVAGVANALAHK		1	Proteotypic
A4GX73_HUMAN	1317	1273.718	LLVVYPWTQR		1	Proteotypic
A4GX73_HUMAN	1431	1313.658	VNVDEVGGEALGR		2	Shared
A4GX73_HUMAN	1577	1377.693	EFTPPVQAAYQK		1	Proteotypic
A4GX73_HUMAN	1751	1468.650	GTFATLSELHCDK	Trioxidation (C)	2	Shared
A4GX73_HUMAN	2020	1668.884	VLGAFSDGLAHLNLIK		2	Shared
A4GX73_HUMAN	2103	1766.950	LLGNVLVCVLAHHFGK	Trioxidation (C)	1	Proteotypic
A4GX73_HUMAN	2286	2073.935	FFESFGDLSTPDVAVMGNPK	Oxidation (M)	2	Shared
A4GX73_HUMAN	2288	2089.930	FFESFGDLSTPDVAVMGNPK	Dioxidation (M)	2	Shared
A4QPB0_HUMAN	405	1024.603	ALQSPALGLR		1	Proteotypic
A4QPB0_HUMAN	568	1071.581	LTELGTVDPK		1	Proteotypic
A4QPB0_HUMAN	788	1124.466	LTAEEMDER	Dioxidation (M)	1	Proteotypic
A4QPB0_HUMAN	892	1149.585	LIFQMPQNK	Dioxidation (M)	3	Shared
A4QPB0_HUMAN	1363	1290.697	MREEVITLIR	Dioxidation (M)	1	Proteotypic
A4QPB0_HUMAN	2190	1882.005	ILAIGLINEALDEGDAQK		1	Proteotypic
A4QPB4_HUMAN	6	899.482	VNAELQAR		1	Proteotypic
A4QPB4_HUMAN	709	1105.536	SATEQSGTGIR		1	Proteotypic
A4QPC1_HUMAN	29	905.461	FLEQQNK		14	Shared
A4QPC1_HUMAN	141	944.504	GRLDSELR		4	Shared
A4QPC1_HUMAN	354	1011.499	SGFSSVSVSR		1	Proteotypic
A4QPC1_HUMAN	376	1015.530	QLDSIVGER		3	Shared

A4QPC1_HUMAN	510	1057.439	DVDAAYMNK	Dioxidation (M)	5	Shared
A4QPC1_HUMAN	548	1065.509	YEDEINKR		8	Shared
A4QPC1_HUMAN	614	1081.592	FASFIDKVR		11	Shared
A4QPC1_HUMAN	717	1106.536	AQYEEIAQR		4	Shared
A4QPC1_HUMAN	785	1123.518	KLLEGECCR	Trioxidation (C)	6	Shared
A4QPC1_HUMAN	900	1151.546	NKYEDEINK		6	Shared
A4QPC1_HUMAN	954	1168.543	EYQELMNVK	Oxidation (M)	4	Shared
A4QPC1_HUMAN	1000	1184.538	EYQELMNVK	Dioxidation (M)	4	Shared
A4QPC1_HUMAN	1415	1307.647	NKYEDEINKR		5	Shared
A4QPC1_HUMAN	1476	1328.719	NLDLDSIAEVK		6	Shared
A4QPC1_HUMAN	1628	1406.704	ADTLTDEINFLR		2	Shared
A4QPC1_HUMAN	1711	1446.768	AIGGGLSSVGGGSSTIK		2	Shared
A4ZU86_HUMAN	104	930.466	GPSSVEDIK		1	Proteotypic
A4ZU86_HUMAN	809	1129.598	GPSSVEDIKAK		1	Proteotypic
A4ZU86_HUMAN	2323	2258.198	MSVQPTVSLGGFEITPPVLR	Dioxidation (M)	1	Proteotypic
A5YM50_HUMAN	468	1042.602	VVLIGDSGVGK		1	Proteotypic
A5YM50_HUMAN	607	1079.561	STIGVEFATR		1	Proteotypic
A5YM50_HUMAN	1358	1288.739	GAVGALLVYDIAK		1	Proteotypic
A5YM50_HUMAN	1888	1562.689	GTRDDEYDYLK	Acetyl (Protein N-term)	1	Proteotypic
A6H8Y6_HUMAN	666	1092.556	ADQFLVGTSR		1	Proteotypic
A6H8Y6_HUMAN	1673	1428.746	VQQQEDEITVLK		1	Proteotypic
A6NCZ6_HUMAN	140	944.493	SGDTPITVR		1	Proteotypic
A6NCZ6_HUMAN	1570	1370.752	NLLLSGAQLEASR		1	Proteotypic
A6ND55_HUMAN	2470	1427.683	DFSPSGIFGAFQR		1	Proteotypic
A6ND64_HUMAN	2472	1458.747	FGQGSPIVLDDVR		1	Proteotypic
A6NDN7_HUMAN	395	1021.511	VIMITGDNK	Dioxidation (M)	1	Proteotypic
A6NDN7_HUMAN	943	1164.566	LDEFGEQLSK		1	Proteotypic
A6NDN7_HUMAN	1403	1304.600	EFDELNPSAQR		1	Proteotypic
A6NE14_HUMAN	947	1165.682	AVAQALEVIPR		1	Proteotypic
A6NE14_HUMAN	1005	1184.713	TAVETAVLLLR		1	Proteotypic
A6NE65_HUMAN	1491	1336.848	AIIIFVPVPLK		1	Proteotypic
A6NE65_HUMAN	1590	1383.635	DVNFEPFQQL		1	Proteotypic
A6NEC4_HUMAN	2410	1069.624	ALAAGGVGSIVR		1	Proteotypic
A6NFM4_HUMAN	2445	1229.662	VTSEGAGLQLQK		1	Proteotypic
A6NG06_HUMAN	109	932.533	VNLAELFK		1	Proteotypic
A6NG06_HUMAN	645	1088.587	LLADPTGAFGK		1	Proteotypic
A6NG06_HUMAN	1257	1253.633	FSMVVQDGIVK	Dioxidation (M)	1	Proteotypic
A6NG06_HUMAN	2155	1826.905	VGDAIPAVEVFEGEPGNK		1	Proteotypic
A6NG06_HUMAN	2206	1905.968	ETDLLLLDDSLVSIFGNR		1	Proteotypic
A6NG06_HUMAN	2345	2373.173	ALNVEPDGTGLTCSLAPNIISQL	Trioxidation (C)	1	Proteotypic
A6NG51_HUMAN	71	915.528	LLEATELK		1	Proteotypic
A6NG51_HUMAN	144	945.513	LTVLSEER		1	Proteotypic
A6NG51_HUMAN	281	989.529	LFGAAEVQR		1	Proteotypic
A6NG51_HUMAN	368	1014.546	DLTGVQNLNLR		1	Proteotypic
A6NG51_HUMAN	566	1071.567	KLDPAQASR		1	Proteotypic
A6NG51_HUMAN	572	1072.551	SLQQLAEER		1	Proteotypic

A6NG51_HUMAN	624	1084.624	DLASVQALLR		1	Proteotypic
A6NG51_HUMAN	692	1100.608	DLAALEDKVK		1	Proteotypic
A6NG51_HUMAN	765	1118.572	VNEVNQFAAK		1	Proteotypic
A6NG51_HUMAN	816	1130.532	QEQIDNQTR		1	Proteotypic
A6NG51_HUMAN	907	1153.525	SSEEIESAFR		1	Proteotypic
A6NG51_HUMAN	963	1172.604	DVTGAEALLER		1	Proteotypic
A6NG51_HUMAN	1082	1203.671	DLSSVQTLTK		1	Proteotypic
A6NG51_HUMAN	1395	1301.646	VLETAEDIQER		1	Proteotypic
A6NG51_HUMAN	1423	1309.728	KVEDLFLTFK		1	Proteotypic
A6NG51_HUMAN	1465	1323.689	SQLLGSHEVQR		1	Proteotypic
A6NG51_HUMAN	1487	1334.650	GVIDMGNSLIER	Dioxidation (M)	1	Proteotypic
A6NG51_HUMAN	1603	1392.700	LGESQTLQQFSR		1	Proteotypic
A6NG51_HUMAN	2299	2126.085	ALINADELASDVAGAEALLDR		1	Proteotypic
A6NGV1_HUMAN	2440	1207.597	VEVTEFEDIK		1	Proteotypic
A6NHU4_HUMAN	294	992.504	SIGVSNFNR		1	Proteotypic
A6NHU4_HUMAN	922	1157.572	REDIFYTSK		1	Proteotypic
A6NHU4_HUMAN	1645	1414.785	QLEMILNKPLK	Dioxidation (M)	1	Proteotypic
A6NIS2_HUMAN	29	905.461	FLEQQNK		14	Shared
A6NIS2_HUMAN	548	1065.509	YEDEINKR		8	Shared
A6NIS2_HUMAN	614	1081.592	FASFIDKVR		11	Shared
A6NIS2_HUMAN	1654	1417.745	VDALIDEIDFLR		1	Proteotypic
A6NJG9_HUMAN	166	954.451	LMVMEIR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A6NJG9_HUMAN	1571	1371.634	VDVFREDLCTK	Trioxidation (C)	1	Proteotypic
A6NJG9_HUMAN	1815	1516.850	APLDIPVPDPVKEK		1	Proteotypic
A6NJG9_HUMAN	1819	1518.731	IEDGNNFGVAVQEK		1	Proteotypic
A6NKD8_HUMAN	2477	1508.678	NDLSPPTVMSEGAR	Dioxidation (M)	1	Proteotypic
A6NKT1_HUMAN	303	995.456	FELDTSER		1	Proteotypic
A6NKT1_HUMAN	598	1077.582	YIANTVELR		1	Proteotypic
A6NKT1_HUMAN	1572	1371.725	SIVEEIEDLVAR		1	Proteotypic
A6NL03_HUMAN	2397	988.482	LAEAEETAR		1	Proteotypic
A6NLE7_HUMAN	2400	1015.591	LTLSALVDGK		1	Proteotypic
A6NLF8_HUMAN	149	948.466	FDENAVVR		2	Shared
A6NLF8_HUMAN	1697	1440.830	SIILSGTVLPSAQR		1	Proteotypic
A6NLK4_HUMAN	100	928.534	ALLTLADGR		1	Proteotypic
A6NLK4_HUMAN	384	1017.524	NTPAFLAER		1	Proteotypic
A6NLK4_HUMAN	575	1072.576	SEIDLLDIR		1	Proteotypic
A6NLK4_HUMAN	625	1084.635	ALLTLADGRR		1	Proteotypic
A6NLK4_HUMAN	720	1106.601	MLISILTER	Dioxidation (M)	1	Proteotypic
A6NLK4_HUMAN	1142	1221.599	GIGTDEFTLNR		1	Proteotypic
A6NLK4_HUMAN	2022	1672.863	GAGTNEDALIEILTTR		1	Proteotypic
A6NMY0_HUMAN	867	1144.609	EVANSTANLVK		1	Proteotypic
A6NMY0_HUMAN	1224	1242.657	VLVQNAAGSQEK		1	Proteotypic
A6NNB0_HUMAN	328	1002.535	LSKEEIER		2	Shared
A6NNB0_HUMAN	379	1016.561	ITITNDKGR		4	Shared
A6NNB0_HUMAN	840	1136.546	YKAEDDEVQR		2	Shared
A6NNB0_HUMAN	1049	1196.688	DAGVIAGLNVLR		2	Shared

A6NNB0_HUMAN	1162	1227.621	VEIANDQGNR		4	Shared
A6NNB0_HUMAN	1448	1318.586	NALESYAFNMK	Dioxidation (M)	2	Shared
A6NNB0_HUMAN	1784	1486.694	TTPSYVAFTDTER		4	Shared
A6NNB0_HUMAN	1977	1645.791	AFYPEEISSMVLTK	Dioxidation (M)	2	Shared
A6NNB0_HUMAN	2006	1658.888	IINEPTAAAIAYGLDK		3	Shared
A6NNB0_HUMAN	2023	1674.723	ATAGDTHLGGEDFDNR		3	Shared
A6NNE8_HUMAN	390	1020.491	VELSTGMPR	Dioxidation (M)	1	Proteotypic
A6NNE8_HUMAN	1950	1620.757	NPPGFVFEFEDPR		2	Shared
A6NNT3_HUMAN	812	1130.455	VEFMDDTSR	Dioxidation (M)	1	Proteotypic
A6NNT3_HUMAN	1547	1359.688	EGDVLTLLESER		1	Proteotypic
A7E236_HUMAN	1355	1288.601	SGSDEVQVGGQR		1	Proteotypic
A7E236_HUMAN	1753	1469.773	IPIEDGSGEVVLSR		1	Proteotypic
A7VJG6_HUMAN	149	948.466	FDENAVVR		2	Shared
A7VJG6_HUMAN	1698	1440.830	SILLSGTVLPSAQR		1	Proteotypic
A7Y9J9_HUMAN	443	1035.583	AQAQPGVPLR		1	Proteotypic
A7Y9J9_HUMAN	2244	1965.010	EEGLILFDQIPVSSGFSK		1	Proteotypic
A8C1Z0_HUMAN	492	1046.551	LAGLFNEQR		1	Proteotypic
A8C1Z0_HUMAN	1320	1275.581	FFGDSAASMAIK	Dioxidation (M)	1	Proteotypic
A8C1Z0_HUMAN	1374	1294.786	LPATEKPVLLSK		1	Proteotypic
A8JZY7_HUMAN	28	905.461	NFDEILR		1	Proteotypic
A8JZY7_HUMAN	342	1006.549	VVVFVGPDK		1	Proteotypic
A8JZY7_HUMAN	622	1084.592	LPFPIIDDR		1	Proteotypic
A8JZY7_HUMAN	834	1134.644	VVVFVGPDKK		1	Proteotypic
A8JZY7_HUMAN	1034	1190.666	LSILYPATTGR		1	Proteotypic
A8JZY7_HUMAN	1594	1385.613	DFTPVCTTELGR	Trioxidation (C)	1	Proteotypic
A8JZY7_HUMAN	2179	1860.866	DGDSVMVLPTIPEEEAK	Dioxidation (M)	1	Proteotypic
A8JZY7_HUMAN	2212	1915.981	ELAILLGMLDPAEKDEK	Dioxidation (M)	1	Proteotypic
A8JZY7_HUMAN	2289	2097.085	PGLLLLGDVAPNFEANTTVGR		1	Proteotypic
A8JZY9_HUMAN	370	1014.571	DVNAAIATIK		2	Shared
A8JZY9_HUMAN	398	1022.442	EDAANNYAR		3	Shared
A8JZY9_HUMAN	623	1084.613	EIIDLVLDR		2	Shared
A8JZY9_HUMAN	1635	1409.767	QLFHPEQLITGK		3	Shared
A8JZY9_HUMAN	1786	1486.872	LISQIVSSITASLR		3	Shared
A8JZY9_HUMAN	2054	1700.899	AVFVDLEPTVIDEVR		2	Shared
A8JZY9_HUMAN	2066	1717.875	NLDIERPTYTNLNR		4	Shared
A8JZY9_HUMAN	2154	1823.978	VGINYQPPTVVPGGDLAK		3	Shared
A8JZY9_HUMAN	2260	2006.886	TIGGGDDSFNTFFSETGAGK		2	Shared
A8JZY9_HUMAN	2347	2408.201	FDGALNVDLTFEQTNLVPYPR		3	Shared
A8K008_HUMAN	901	1151.586	NQVSLTCLVK	Trioxidation (C)	1	Proteotypic
A8K008_HUMAN	1007	1185.639	GPSVFPLAPSSK		1	Proteotypic
A8K008_HUMAN	1427	1311.634	STSGGTAALGCLVK	Trioxidation (C)	1	Proteotypic
A8K008_HUMAN	2187	1872.915	TTPPVLDSDGSFFLYSK		1	Proteotypic
A8K048_HUMAN	930	1159.572	TGAEGAVLDEAK		1	Proteotypic
A8K048_HUMAN	1436	1314.678	LITDLQDQNQK		1	Proteotypic
A8K068_HUMAN	592	1076.525	IREEYPDR		4	Shared
A8K068_HUMAN	769	1119.363	NMMAACDPR	Dioxidation (M); Dioxidation (M); Trioxidation (C)	5	Shared

A8K068_HUMAN	808	1129.588	FPGQLNADLR		5	Shared
A8K068_HUMAN	928	1158.622	LAVNMVFPFR	Oxidation (M)	5	Shared
A8K068_HUMAN	971	1174.617	LAVNMVFPFR	Dioxidation (M)	5	Shared
A8K068_HUMAN	1264	1257.683	FPGQLNADLRK		5	Shared
A8K068_HUMAN	1275	1260.581	ISEQFTAMFR	Dioxidation (M)	4	Shared
A8K068_HUMAN	1398	1302.712	KLAVNMVFPFR	Dioxidation (M)	5	Shared
A8K068_HUMAN	1641	1414.647	IMNTFSVMPSPK	Dioxidation (M); Dioxidation (M)	2	Shared
A8K068_HUMAN	1769	1477.672	EVDEQMLNVQNK	Dioxidation (M)	3	Shared
A8K068_HUMAN	1980	1646.819	AILVDLEPGTMDSVR	Dioxidation (M)	3	Shared
A8K068_HUMAN	2267	2018.859	MSMKEVDEQMLNVQNK	Dioxidation (M); Dioxidation (M); Dioxidation (M)	3	Shared
A8K089_HUMAN	66	914.555	KLILDSAR		1	Proteotypic
A8K089_HUMAN	1563	1367.777	LDNLVAILDINR		1	Proteotypic
A8K089_HUMAN	2268	2019.064	ILATPPQEDAPSVDIANIR		1	Proteotypic
A8K094_HUMAN	2390	940.545	AVVGVVAGGGR		1	Proteotypic
A8K0B8_HUMAN	370	1014.571	DVNAAIATIK		2	Shared
A8K0B8_HUMAN	398	1022.442	EDAANNYAR		3	Shared
A8K0B8_HUMAN	623	1084.613	EIIDLVLDR		2	Shared
A8K0B8_HUMAN	1635	1409.767	QLFHPEQLITGK		3	Shared
A8K0B8_HUMAN	1727	1456.861	LIGQIVSSITASLR		1	Proteotypic
A8K0B8_HUMAN	2054	1700.899	AVFVDLEPTVIDEVR		2	Shared
A8K0B8_HUMAN	2066	1717.875	NLDIERPTYTNLNR		4	Shared
A8K0B8_HUMAN	2154	1823.978	VGINYQPPTVVPGGDLAK		3	Shared
A8K0B8_HUMAN	2260	2006.886	TIGGGDDSFNTFFSETGAGK		2	Shared
A8K0B8_HUMAN	2347	2408.201	FDGALNVDLTEFQTNLVPYPR		3	Shared
A8K0K3_HUMAN	898	1150.598	IVNSAQTGSFK		1	Proteotypic
A8K0K3_HUMAN	1650	1416.798	VELVPPTPAEIPR		1	Proteotypic
A8K0N2_HUMAN	181	959.492	VVGAVGSDEK		1	Proteotypic
A8K0N2_HUMAN	1322	1275.692	TVESLEETLKK		1	Proteotypic
A8K0N2_HUMAN	1929	1598.728	LKEGDTMMGQQVAK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K0P1_HUMAN	960	1171.595	MFLVNSFLK	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
A8K0P1_HUMAN	1624	1404.575	SMVAVMDSDTTGK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K133_HUMAN	975	1175.655	LVFLGEQSVGK		1	Proteotypic
A8K133_HUMAN	1323	1276.670	GSDVIIMLVGNK	Dioxidation (M)	1	Proteotypic
A8K133_HUMAN	1359	1289.636	STGGDFGNPLRK	Acetyl (Protein N-term)	1	Proteotypic
A8K133_HUMAN	1768	1476.688	VAAALPGMESTQDR	Dioxidation (M)	1	Proteotypic
A8K158_HUMAN	2386	899.446	AAAAAAGEAR	Acetyl (Protein N-term)	1	Proteotypic
A8K1D2_HUMAN	1653	1417.720	GFSVVADTPELQR		1	Proteotypic
A8K1D2_HUMAN	1973	1639.751	QSFTMVADTPENLR	Dioxidation (M)	1	Proteotypic
A8K1G0_HUMAN	148	947.547	VLLTMIAR	Dioxidation (M)	1	Proteotypic
A8K1G0_HUMAN	278	989.503	GEALSALDSK		1	Proteotypic
A8K1G0_HUMAN	1708	1445.755	IMVANIEEVLQR	Dioxidation (M)	1	Proteotypic
A8K1G0_HUMAN	1916	1585.806	NLGSINTELQDVQR		1	Proteotypic
A8K1G0_HUMAN	2258	2004.906	VADGLPLAASMQEDEQSGR	Dioxidation (M)	1	Proteotypic
A8K1G2_HUMAN	2484	1639.784	IPNQFQSDPPAPSDK		1	Proteotypic
A8K1V1_HUMAN	942	1164.552	GFTLNDAANSR		1	Proteotypic
A8K1V1_HUMAN	1177	1232.546	MEEANIQPNR	Dioxidation (M)	1	Proteotypic



A8K1V1_HUMAN	1311	1271.647	TVLDQQQTPSR		1	Proteotypic
A8K202_HUMAN	2483	1638.756	DVSELTGFPEMLGGR	Dioxidation (M)	1	Proteotypic
A8K220_HUMAN	504	1054.534	VSFELFADK		1	Proteotypic
A8K220_HUMAN	729	1109.491	ITIADCGQLE	Trioxidation (C)	1	Proteotypic
A8K220_HUMAN	908	1153.566	FEDENFILK		1	Proteotypic
A8K220_HUMAN	1201	1237.586	KITIADCGQLE	Trioxidation (C)	1	Proteotypic
A8K220_HUMAN	1503	1341.554	EGMNIVEAMER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K220_HUMAN	1580	1378.750	VSFELFADKVPK		1	Proteotypic
A8K220_HUMAN	1897	1568.717	VKEGMNIVEAMER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K220_HUMAN	1949	1620.691	IIPGFMCQGGDFTR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
A8K220_HUMAN	2159	1830.904	SIYGEKFEDENFILK		1	Proteotypic
A8K220_HUMAN	2231	1944.995	VNPTVFFDIAVDGEPLGR		1	Proteotypic
A8K220_HUMAN	2250	1987.005	VNPTVFFDIAVDGEPLGR	Acetyl (Protein N-term)	1	Proteotypic
A8K236_HUMAN	536	1062.538	LLIEMEQR	Dioxidation (M)	1	Proteotypic
A8K236_HUMAN	1399	1302.718	VLTPELYAELR		1	Proteotypic
A8K236_HUMAN	1917	1585.831	LAVEALSSLDGDLAGR		1	Proteotypic
A8K236_HUMAN	2039	1687.809	LEQQQAIDDLMPAQK	Dioxidation (M)	1	Proteotypic
A8K236_HUMAN	2188	1879.960	LGFSLEVELVQMVVDGVK	Dioxidation (M)	1	Proteotypic
A8K2Q1_HUMAN	2409	1068.618	LSPEELLLR		1	Proteotypic
A8K2Z5_HUMAN	2461	1298.694	AAAAAATAAAAAASIR	Acetyl (Protein N-term)	1	Proteotypic
A8K307_HUMAN	2451	1255.677	VPAINVNDSVTK		1	Proteotypic
A8K318_HUMAN	312	998.612	LIELQAGKK		1	Proteotypic
A8K318_HUMAN	1206	1238.581	SLKDMEESIR	Dioxidation (M)	1	Proteotypic
A8K318_HUMAN	1246	1250.581	SLEDQVEMLR	Dioxidation (M)	1	Proteotypic
A8K318_HUMAN	1435	1314.678	TVKEEAEKPER		1	Proteotypic
A8K318_HUMAN	1461	1322.614	ESLQQMAEVTR	Dioxidation (M)	1	Proteotypic
A8K3B4_HUMAN	2438	1202.641	NPFLAAVTTNR		1	Proteotypic
A8K3C3_HUMAN	1537	1356.725	VIDPATATSVDLR		1	Proteotypic
A8K3C3_HUMAN	1872	1549.883	ALIAGGGAPEIELALR		1	Proteotypic
A8K3F6_HUMAN	346	1008.510	AGVLAGHDNR		1	Proteotypic
A8K3F6_HUMAN	378	1016.550	LLVSASQDGK		1	Proteotypic
A8K3F6_HUMAN	2107	1769.891	SELEQLRQEAQLR	Acetyl (Protein N-term)	1	Proteotypic
A8K3I7_HUMAN	2419	1102.577	AAAAVVEFQR	Acetyl (Protein N-term)	1	Proteotypic
A8K3J2_HUMAN	394	1020.576	VFSIGPVFR		1	Proteotypic
A8K3J2_HUMAN	1069	1200.635	ESIVDVEGVVR		1	Proteotypic
A8K3J2_HUMAN	1089	1206.624	FQTEIQTVVK		1	Proteotypic
A8K3P0_HUMAN	2466	1402.807	VLALELFEQITK		1	Proteotypic
A8K3Z8_HUMAN	372	1014.571	LVLVGDGGTGK		1	Proteotypic
A8K3Z8_HUMAN	1230	1243.620	AAQGEPQVQFK	Acetyl (Protein N-term)	1	Proteotypic
A8K401_HUMAN	512	1057.515	QVAQQEAER		1	Proteotypic
A8K401_HUMAN	529	1060.540	AAIIAEGDSK		1	Proteotypic
A8K401_HUMAN	530	1061.499	QVSDDLTER		1	Proteotypic
A8K401_HUMAN	882	1148.583	FDAGELITQR		1	Proteotypic
A8K401_HUMAN	1002	1184.651	DLQNVNITLR		1	Proteotypic
A8K401_HUMAN	1110	1212.733	VLPSITTEILK		1	Proteotypic
A8K401_HUMAN	1611	1395.835	ILFRPVASQLPR		1	Proteotypic

A8K401_HUMAN	1702	1443.652	IFTSIGEDYDER		1	Proteotypic
A8K401_HUMAN	2256	1997.079	AAELIANSLATAGDGLIELR		1	Proteotypic
A8K414_HUMAN	290	991.534	ANTFVAELK		1	Proteotypic
A8K414_HUMAN	576	1072.576	IQEAGTEVVK		1	Proteotypic
A8K414_HUMAN	862	1143.512	EGVVECSFVK	Trioxidation (C)	1	Proteotypic
A8K414_HUMAN	876	1146.651	VNVPVIGGHAGK		1	Proteotypic
A8K414_HUMAN	879	1147.579	MISDAIPELK	Dioxidation (M)	1	Proteotypic
A8K414_HUMAN	1182	1232.713	IFGVTTLDIVR		1	Proteotypic
A8K414_HUMAN	1543	1358.654	FVFSLVDAMNGK	Dioxidation (M)	1	Proteotypic
A8K414_HUMAN	1781	1485.688	AGAGSATLSMAYAGAR	Dioxidation (M)	1	Proteotypic
A8K414_HUMAN	1882	1559.794	VDFPQDQLTALTGR		1	Proteotypic
A8K414_HUMAN	2124	1792.082	VAVLGASGGIGQPLSLLK		1	Proteotypic
A8K430_HUMAN	1210	1240.641	LDQGGAPLAGTNK		1	Proteotypic
A8K430_HUMAN	1651	1417.705	ETTIQGLDGLSER		1	Proteotypic
A8K430_HUMAN	1822	1521.709	GILAADESVMGMGR	Dioxidation (M)	1	Proteotypic
A8K488_HUMAN	420	1028.587	VLDELTLAR		5	Shared
A8K488_HUMAN	708	1105.519	VTMQNLNDR	Oxidation (M)	3	Shared
A8K488_HUMAN	777	1121.514	VTMQNLNDR	Dioxidation (M)	3	Shared
A8K488_HUMAN	779	1121.572	LEQEIATYR		6	Shared
A8K488_HUMAN	1003	1184.688	RVLDELTLAR		5	Shared
A8K488_HUMAN	1130	1219.593	ASLENSLEETK		2	Shared
A8K488_HUMAN	1270	1259.563	EVFTSSSSSSSR		1	Proteotypic
A8K488_HUMAN	1393	1300.651	ALEEANADLEVK		4	Shared
A8K488_HUMAN	1414	1307.628	TDLEMQIEGLK	Dioxidation (M)	2	Shared
A8K488_HUMAN	2281	2063.137	IIAATIENAQPILQIDNAR		1	Proteotypic
A8K4H3_HUMAN	29	905.461	FLEQQNK		14	Shared
A8K4H3_HUMAN	40	910.414	SYTSGPGSR		1	Proteotypic
A8K4H3_HUMAN	108	931.461	LLEGEESR		4	Shared
A8K4H3_HUMAN	316	999.560	LQAEIEGLK		1	Proteotypic
A8K4H3_HUMAN	330	1002.535	TQEKEQIK		1	Proteotypic
A8K4H3_HUMAN	348	1010.434	TEISEMNR	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	477	1044.545	QLETLGQEK		1	Proteotypic
A8K4H3_HUMAN	525	1059.556	KLLEGEESR		3	Shared
A8K4H3_HUMAN	548	1065.509	YEDEINKR		8	Shared
A8K4H3_HUMAN	601	1078.504	AQYEDIANR		2	Shared
A8K4H3_HUMAN	602	1078.529	QLYEEEIR		1	Proteotypic
A8K4H3_HUMAN	614	1081.592	FASFIDKVR		11	Shared
A8K4H3_HUMAN	684	1099.449	DVDEAYMNK	Oxidation (M)	1	Proteotypic
A8K4H3_HUMAN	750	1115.444	DVDEAYMNK	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	802	1128.614	GELAIKDANAK		2	Shared
A8K4H3_HUMAN	803	1128.614	LSELEAALQR		1	Proteotypic
A8K4H3_HUMAN	841	1136.571	YEELQSLAGK		1	Proteotypic
A8K4H3_HUMAN	900	1151.546	NKYEDEINK		6	Shared
A8K4H3_HUMAN	954	1168.543	EYQELMNVK	Oxidation (M)	4	Shared
A8K4H3_HUMAN	964	1172.629	LVSESSDVLPK		1	Proteotypic
A8K4H3_HUMAN	999	1184.538	AEAESMYQIK	Oxidation (M)	1	Proteotypic

A8K4H3_HUMAN	1000	1184.538	EYQELMNVK	Dioxidation (M)	4	Shared
A8K4H3_HUMAN	1064	1200.533	AEAESMYQIK	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1070	1200.646	RQLETLGQEK		1	Proteotypic
A8K4H3_HUMAN	1208	1239.576	TKTEISEMNR	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1228	1243.539	KDVDEAYMNK	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1324	1276.703	LALDIEIATYR		4	Shared
A8K4H3_HUMAN	1415	1307.647	NKYEDEINKR		5	Shared
A8K4H3_HUMAN	1490	1335.659	SLDMDSIIAEVK	Oxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1502	1340.741	LQAEIEGLKGQR		1	Proteotypic
A8K4H3_HUMAN	1509	1343.668	ASLEAAIADAEQR		1	Proteotypic
A8K4H3_HUMAN	1524	1351.654	SLDMDSIIAEVK	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1562	1367.664	TEMENEFVLIK	Oxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1591	1383.659	TEMENEFVLIK	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1625	1404.798	LALDIEIATYRK		4	Shared
A8K4H3_HUMAN	1656	1418.741	LEGLTDEINFLR		1	Proteotypic
A8K4H3_HUMAN	1666	1427.671	SRAEAEESMYQIK	Oxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1703	1443.666	SRAEAEESMYQIK	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1758	1472.772	DGKLVSESSDVLPK		1	Proteotypic
A8K4H3_HUMAN	1808	1511.754	TEMENEFVLIKK	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1826	1522.676	LESGMQNMSIHTK	Dioxidation (M); Oxidation (M)	1	Proteotypic
A8K4H3_HUMAN	1853	1538.670	LESGMQNMSIHTK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2135	1807.866	LEAELGNMQGLVEDFK	Oxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2153	1823.861	LEAELGNMQGLVEDFK	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2156	1828.815	DVDEAYMKNVELESR	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2200	1894.783	SNMDNMFESYINNLR	Dioxidation (M); Oxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2210	1910.777	SNMDNMFESYINNLR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2239	1955.032	ASLEAAIADAEQRGELAIK		1	Proteotypic
A8K4H3_HUMAN	2283	2065.040	LKLEAELGNMQGLVEDFK	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2284	2066.879	SNMDNMFESYINNLR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2295	2124.000	ELQSQISDTSVVLSMDNSR	Oxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2305	2139.995	ELQSQISDTSVVLSMDNSR	Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2350	2452.121	LLEGEESRLESGMQNMSIHTK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K4H3_HUMAN	2351	2454.271	ASLEAAIADAEQRGELAIKDANAK		1	Proteotypic
A8K4H3_HUMAN	2384	3200.514	QLYEEEIRELQSQISDTSVVLSMDNSR	Dioxidation (M)	1	Proteotypic
A8K4I2_HUMAN	219	972.597	SGVSLAALKK		1	Proteotypic
A8K4I2_HUMAN	718	1106.561	ALAAAGYDVEK		1	Proteotypic
A8K4I2_HUMAN	1053	1197.660	ASGPPVSELITK		1	Proteotypic
A8K4I2_HUMAN	1469	1325.755	KASGPPVSELITK		1	Proteotypic
A8K4I2_HUMAN	1820	1519.752	SETAPAAPAAAPPAEK	Acetyl (Protein N-term)	1	Proteotypic
A8K4K6_HUMAN	239	977.602	RSPPIPLAK		1	Proteotypic
A8K4K6_HUMAN	604	1078.602	DKDPPIPVAK		1	Proteotypic
A8K4K6_HUMAN	824	1132.561	VDATAETDLAK		1	Proteotypic
A8K4K6_HUMAN	995	1183.543	TFDSIVMDPK	Dioxidation (M)	1	Proteotypic
A8K4K6_HUMAN	1026	1189.630	IDATSASVLSAR		1	Proteotypic
A8K4K6_HUMAN	1357	1288.662	VDATAETDLAKR		1	Proteotypic
A8K4K6_HUMAN	1458	1322.541	MDATANDVPSDR	Dioxidation (M)	1	Proteotypic

A8K4K6_HUMAN	1761	1473.750	VSQQLVVMQPEK	Dioxidation (M)	1	Proteotypic
A8K4K6_HUMAN	2065	1717.714	FAMEPEEFDSDTLR	Dioxidation (M)	1	Proteotypic
A8K4Q9_HUMAN	528	1059.644	VLALFQLTR		1	Proteotypic
A8K4Q9_HUMAN	543	1064.528	GAAMVTAVASR	Dioxidation (M)	1	Proteotypic
A8K4Q9_HUMAN	667	1093.540	FLSQIESDR		2	Shared
A8K4Q9_HUMAN	1462	1322.698	FLALDLGGTNFR		1	Proteotypic
A8K4T6_HUMAN	2395	967.581	AAQALALLR	Acetyl (Protein N-term)	1	Proteotypic
A8K4T9_HUMAN	271	987.535	SNIDALLSR		1	Proteotypic
A8K4T9_HUMAN	418	1028.550	LEQIAAEQK		1	Proteotypic
A8K4T9_HUMAN	1353	1287.755	VFIGNLNTALVK		1	Proteotypic
A8K4V1_HUMAN	203	966.525	VLQHAASNK		1	Proteotypic
A8K4V1_HUMAN	733	1110.450	LSAEEMDER	Dioxidation (M)	1	Proteotypic
A8K4V1_HUMAN	853	1140.614	SPAIGLNLDK		1	Proteotypic
A8K4V1_HUMAN	892	1149.585	LIFQMPQNK	Dioxidation (M)	3	Shared
A8K4V1_HUMAN	974	1175.630	FQATSSGPILR		1	Proteotypic
A8K4V1_HUMAN	1010	1185.672	GIAEQTVVTLR		1	Proteotypic
A8K4W0_HUMAN	143	945.477	EVQTNDLK		1	Proteotypic
A8K4W0_HUMAN	154	950.464	APAMFNIR	Dioxidation (M)	1	Proteotypic
A8K4W0_HUMAN	338	1006.377	MMEIMTR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K4W0_HUMAN	2236	1950.969	VFEVSLADLQNDEVAFR		1	Proteotypic
A8K4W1_HUMAN	1087	1206.555	MIAAVDTDSPR	Dioxidation (M)	1	Proteotypic
A8K4W1_HUMAN	1873	1550.688	IGDELDNMEIQR	Dioxidation (M)	1	Proteotypic
A8K4W2_HUMAN	13	900.492	LAQLEEAK		1	Proteotypic
A8K4W2_HUMAN	1665	1425.692	NNIAMALEVTYR	Dioxidation (M)	1	Proteotypic
A8K4W6_HUMAN	118	935.463	ALMDEVVK	Dioxidation (M)	1	Proteotypic
A8K4W6_HUMAN	244	980.464	VDFNVPMK	Dioxidation (M)	1	Proteotypic
A8K4W6_HUMAN	475	1043.623	LTLDKLDVK		1	Proteotypic
A8K4W6_HUMAN	677	1096.613	VLPVDALSNI		1	Proteotypic
A8K4W6_HUMAN	689	1100.532	NNQITNNQR		1	Proteotypic
A8K4W6_HUMAN	818	1130.608	AEPAKIEAFR		1	Proteotypic
A8K4W6_HUMAN	1180	1232.643	IQLINNMLDK	Dioxidation (M)	1	Proteotypic
A8K4W6_HUMAN	1238	1246.681	YSLEPVAVELK		1	Proteotypic
A8K4W6_HUMAN	2080	1737.850	DVLFKDCVGPVEVK	Trioxidation (C)	1	Proteotypic
A8K4W6_HUMAN	2097	1758.893	ACANPAAGSVILLENLR	Trioxidation (C)	1	Proteotypic
A8K4W6_HUMAN	2105	1767.988	ALESPERPFLAILGGAK		1	Proteotypic
A8K4W6_HUMAN	2257	1997.925	VLNNMEIGTSLFDEEGAK	Dioxidation (M)	1	Proteotypic
A8K4W6_HUMAN	2270	2022.031	ITLPVDFVTADKFDENAK		1	Proteotypic
A8K4X0_HUMAN	2	898.512	VLDSGAPIK		1	Proteotypic
A8K4X0_HUMAN	229	974.555	IGLFGGAGVGK		1	Proteotypic
A8K4X0_HUMAN	451	1037.587	IPVGPETLGR		1	Proteotypic
A8K4X0_HUMAN	457	1039.664	LVPLKETIK		1	Proteotypic
A8K4X0_HUMAN	643	1087.628	VVDLLAPYAK		1	Proteotypic
A8K4X0_HUMAN	1327	1277.629	TIAMDGTEGLVR	Oxidation (M)	1	Proteotypic
A8K4X0_HUMAN	1372	1293.623	TIAMDGTEGLVR	Dioxidation (M)	1	Proteotypic
A8K4X0_HUMAN	1648	1416.692	IMNVIGEPIDER	Dioxidation (M)	1	Proteotypic
A8K4X0_HUMAN	1686	1434.747	FTQAGSEVSALLGR		1	Proteotypic

A8K4X0_HUMAN	1690	1438.782	VALTGLTVAEYFR		1	Proteotypic
A8K4X0_HUMAN	1759	1472.827	TVLIMELINNVAK	Oxidation (M)	1	Proteotypic
A8K4X0_HUMAN	1789	1488.822	TVLIMELINNVAK	Dioxidation (M)	1	Proteotypic
A8K4X0_HUMAN	1964	1632.793	VALVYQGMNEPPGAR	Dioxidation (M)	1	Proteotypic
A8K4X0_HUMAN	1986	1649.910	LVLEVAQHLGESTVR		1	Proteotypic
A8K4X0_HUMAN	2140	1811.945	IMNVIGEPIDERGPIK	Dioxidation (M)	1	Proteotypic
A8K4X0_HUMAN	2171	1846.852	IMDPNIVGSEHYDVAR	Dioxidation (M)	1	Proteotypic
A8K4X0_HUMAN	2214	1918.089	VLDSGAPIKIPVGPETLGR		1	Proteotypic
A8K4X0_HUMAN	2217	1920.958	DQEGQDVLFFIDNIFR		1	Proteotypic
A8K4X0_HUMAN	2251	1987.026	AIAELGIYPAVDPLDSTSR		1	Proteotypic
A8K4X0_HUMAN	2338	2329.057	IPSAVGYQPTLATDMGTMQER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K4X0_HUMAN	2367	2706.363	SLQDIIALGMDELSEEDKLTVSR	Dioxidation (M)	1	Proteotypic
A8K4X5_HUMAN	2396	986.555	IPLNDLFR		1	Proteotypic
A8K4Z4_HUMAN	1430	1312.703	TSFFQALGITTK		1	Proteotypic
A8K4Z4_HUMAN	1669	1427.824	GTIEILSDVQLIK		1	Proteotypic
A8K4Z4_HUMAN	2371	2751.460	AFLADPSAFVAAAPVAAATTAAPAAAAAPAK		1	Proteotypic
A8K4Z8_HUMAN	1243	1249.622	IEVIEIMTDR	Dioxidation (M)	1	Proteotypic
A8K4Z8_HUMAN	1384	1298.647	SESPKEPEQLR		1	Proteotypic
A8K4Z8_HUMAN	1688	1436.737	EDSQRPGAHLTVK		1	Proteotypic
A8K4Z8_HUMAN	2117	1783.899	LFIGGLSFETTDESLR		1	Proteotypic
A8K500_HUMAN	297	994.403	TDTGPEMGR	Dioxidation (M)	1	Proteotypic
A8K500_HUMAN	393	1020.552	IMKDILEK	Dioxidation (M)	1	Proteotypic
A8K500_HUMAN	724	1107.535	APFDLFENR		1	Proteotypic
A8K500_HUMAN	897	1150.551	YIDQEELNK		2	Shared
A8K500_HUMAN	951	1167.563	LGIHEDSQNR		1	Proteotypic
A8K500_HUMAN	1191	1234.594	DQVANSAFVER		1	Proteotypic
A8K500_HUMAN	1220	1241.698	ADLNNLGTIAK		2	Shared
A8K500_HUMAN	1283	1263.636	RAPFDLFENR		1	Proteotypic
A8K500_HUMAN	1362	1290.630	ELISNSSDALDK		1	Proteotypic
A8K500_HUMAN	1424	1310.563	EDQTEYLEER		2	Shared
A8K500_HUMAN	1584	1380.717	TLTIVDTGIGMTK	Dioxidation (M)	1	Proteotypic
A8K500_HUMAN	1809	1512.778	GVVDEDLPLNISR		2	Shared
A8K504_HUMAN	2434	1169.677	IALTDNALIAR		1	Proteotypic
A8K505_HUMAN	2449	1239.682	AEVQVLVLDGR	Acetyl (Protein N-term)	1	Proteotypic
A8K512_HUMAN	1752	1468.720	DADVQNFVFSISK		1	Proteotypic
A8K512_HUMAN	1785	1486.770	VLVEPDAGAGVAVMK	Dioxidation (M)	1	Proteotypic
A8K519_HUMAN	527	1059.592	RLEVL DSTK		1	Proteotypic
A8K519_HUMAN	2306	2141.067	NAVTFVSSMSASADVLALAK	Dioxidation (M)	1	Proteotypic
A8K525_HUMAN	7	899.486	AGEVFIHK		1	Proteotypic
A8K525_HUMAN	375	1015.468	QQQDQVDR		1	Proteotypic
A8K525_HUMAN	683	1098.615	AVVIVDDRGR		2	Shared
A8K525_HUMAN	694	1101.549	VELDNMPLR	Oxidation (M)	1	Proteotypic
A8K525_HUMAN	758	1117.544	VELDNMPLR	Dioxidation (M)	1	Proteotypic
A8K525_HUMAN	1164	1227.636	AAPGAEFAPNKR		1	Proteotypic
A8K525_HUMAN	1172	1230.672	GIVEFSGKPAAR		1	Proteotypic
A8K525_HUMAN	1205	1238.571	FACHSASLTVR	Trioxidation (C)	1	Proteotypic

A8K525_HUMAN	1616	1399.596	EKLEMEEAAR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K525_HUMAN	2015	1665.621	MGQMAMGGAMGINNR	Dioxidation (M); Dioxidation (M); Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K525_HUMAN	2364	2667.318	NLPQYYSNELLEAFSVFGQVER		1	Proteotypic
A8K533_HUMAN	1816	1517.620	MMLMSTATAFYR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K533_HUMAN	1900	1571.729	KVFANPEDCVAFGK	Trioxidation (C)	1	Proteotypic
A8K534_HUMAN	538	1062.538	VLEGMEVVR	Dioxidation (M)	1	Proteotypic
A8K534_HUMAN	1446	1317.566	DFMIQGGDFTR	Dioxidation (M)	1	Proteotypic
A8K534_HUMAN	1554	1363.698	TVDNFVALATGEK		1	Proteotypic
A8K534_HUMAN	1723	1456.720	DTNGSQFFITTVK		1	Proteotypic
A8K534_HUMAN	2002	1657.813	VIKDFMIQGGDFTR	Dioxidation (M)	1	Proteotypic
A8K588_HUMAN	426	1029.582	LIVENLSSR		1	Proteotypic
A8K588_HUMAN	540	1063.530	TNEGVIEFR		1	Proteotypic
A8K590_HUMAN	640	1087.558	VLGMDPLPSK	Dioxidation (M)	1	Proteotypic
A8K590_HUMAN	1426	1311.602	FVMEVEVDGQK	Dioxidation (M)	1	Proteotypic
A8K590_HUMAN	1762	1474.709	VLQDMGLPTGAAGR	Dioxidation (M)	1	Proteotypic
A8K590_HUMAN	2209	1908.979	VLAGETLSVNDPPDVLDR		1	Proteotypic
A8K5H2_HUMAN	430	1030.541	SSAEVIAQAR		1	Proteotypic
A8K5H2_HUMAN	620	1083.629	GSPLVVISQGK		1	Proteotypic
A8K5H2_HUMAN	1463	1322.756	QIGENLIVPGGVK		1	Proteotypic
A8K5H2_HUMAN	2095	1756.794	MDENQFVAVTSTNAAK	Dioxidation (M)	1	Proteotypic
A8K5I0_HUMAN	328	1002.535	LSKEEIER		2	Shared
A8K5I0_HUMAN	379	1016.561	ITITNDKGR		4	Shared
A8K5I0_HUMAN	727	1108.567	LLQDFFNDR		1	Proteotypic
A8K5I0_HUMAN	840	1136.546	YKADEVQR		2	Shared
A8K5I0_HUMAN	1049	1196.688	DAGVIAGLNVLR		2	Shared
A8K5I0_HUMAN	1162	1227.621	VEIANDQGNR		4	Shared
A8K5I0_HUMAN	1254	1253.560	FGDPVVQSDMK	Dioxidation (M)	1	Proteotypic
A8K5I0_HUMAN	1276	1260.650	LVNHFVEEFK		1	Proteotypic
A8K5I0_HUMAN	1448	1318.586	NALESYAFNMK	Dioxidation (M)	2	Shared
A8K5I0_HUMAN	1784	1486.694	TTPSYVAFTDTER		4	Shared
A8K5I0_HUMAN	1846	1532.693	ARFEELCSDLFR	Trioxidation (C)	2	Shared
A8K5I0_HUMAN	1977	1645.791	AFYPEEISSMVLTK	Dioxidation (M)	2	Shared
A8K5I0_HUMAN	2003	1657.842	NQVALNPQNTVFDKAK		1	Proteotypic
A8K5I0_HUMAN	2023	1674.723	ATAGDTHLGGEDFDNR		3	Shared
A8K5I0_HUMAN	2038	1686.894	IINEPTAAAIAAYGLDR		2	Shared
A8K5J6_HUMAN	2474	1472.835	LGANSLLDLVVFGR		1	Proteotypic
A8K5L2_HUMAN	302	995.434	ADLSGMSGAR	Dioxidation (M)	1	Proteotypic
A8K5L2_HUMAN	1090	1206.624	LGVQDLFNSSK		1	Proteotypic
A8K5L2_HUMAN	1097	1209.530	MEQLSSANTR	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
A8K5S9_HUMAN	1409	1305.693	VAQVAEITYGQK		1	Proteotypic
A8K5S9_HUMAN	1484	1333.688	ISEIEDAAFLAR		1	Proteotypic
A8K5S9_HUMAN	1515	1347.688	SVQTTLQTDEVK		1	Proteotypic
A8K674_HUMAN	553	1066.613	ESTLHLVLR		1	Proteotypic
A8K674_HUMAN	1830	1522.774	IQDKEGIPPDQQR		1	Proteotypic
A8K674_HUMAN	2119	1786.920	TITLEVEPSDTIENVK		1	Proteotypic
A8K685_HUMAN	90	924.507	FLVGFTNK		1	Proteotypic

A8K685_HUMAN	2035	1684.794	GTEDFIVESLDASFR		1	Proteotypic
A8K690_HUMAN	822	1131.632	LMDVGLIAIR	Dioxidation (M)	1	Proteotypic
A8K690_HUMAN	1455	1320.623	MEQVNELKEK	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
A8K6G8_HUMAN	343	1007.459	MLAEDELRL	Dioxidation (M)	1	Proteotypic
A8K6G8_HUMAN	646	1088.623	DAVLLVFANK		2	Shared
A8K6G8_HUMAN	767	1118.601	ILMVGLDAAGK	Dioxidation (M)	3	Shared
A8K6G8_HUMAN	2011	1661.757	QDLPNAMNAAEITDK	Dioxidation (M)	1	Proteotypic
A8K6G8_HUMAN	2287	2078.072	MLAEDELRLDAVLLVFANK	Dioxidation (M)	1	Proteotypic
A8K6G8_HUMAN	2336	2322.236	LGEIVTTIPTIGFNVETVEYK		3	Shared
A8K6H1_HUMAN	2467	1403.741	ILASTQFEPTAAR		1	Proteotypic
A8K6L1_HUMAN	242	978.586	AVLLGPPGAGK		1	Proteotypic
A8K6L1_HUMAN	610	1080.512	AMVASGSELGK	Dioxidation (M)	1	Proteotypic
A8K6L1_HUMAN	1850	1534.780	LVSDEMVELIEK	Dioxidation (M)	1	Proteotypic
A8K6M8_HUMAN	1231	1243.775	SIATLAITLLK		1	Proteotypic
A8K6M8_HUMAN	1878	1553.855	SLEELPVDIILASVG		1	Proteotypic
A8K6N2_HUMAN	248	982.512	IGFGSFVEK		1	Proteotypic
A8K6N2_HUMAN	371	1014.571	SAVTTVVNPK		1	Proteotypic
A8K6N2_HUMAN	652	1090.566	GEVFNELVGK		1	Proteotypic
A8K6N2_HUMAN	1477	1329.554	SLGTDLMNEMR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K6N2_HUMAN	2266	2018.152	LKPEDITQIQPQQLVLR		1	Proteotypic
A8K6V3_HUMAN	176	957.524	QQEIVVSR		1	Proteotypic
A8K6V3_HUMAN	1335	1282.729	IVPGQFLAVDPK		1	Proteotypic
A8K6V3_HUMAN	1400	1302.730	FLAVGLVDNTR		1	Proteotypic
A8K6V3_HUMAN	1787	1487.747	TVLDPVTGDLSDTR		1	Proteotypic
A8K741_HUMAN	746	1114.562	QLEAELGAER		1	Proteotypic
A8K741_HUMAN	2199	1891.053	VPADLGAEAGLQQLLGALR		1	Proteotypic
A8K745_HUMAN	69	915.503	SPVAVQSTK		1	Proteotypic
A8K745_HUMAN	1379	1297.630	YQETFNVIER		1	Proteotypic
A8K745_HUMAN	1803	1499.660	MMADEALGSGLVSR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K745_HUMAN	1860	1541.805	EVDVGLAADVGTLQR		1	Proteotypic
A8K745_HUMAN	1959	1627.755	KMMADEALGSGLVSR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K745_HUMAN	2074	1730.932	VIGNQSLVNELAFTAR		1	Proteotypic
A8K745_HUMAN	2341	2349.214	VFPDKEVMLDAALALAAEISSK	Dioxidation (M)	1	Proteotypic
A8K766_HUMAN	119	936.433	TALAMGADR	Dioxidation (M)	1	Proteotypic
A8K766_HUMAN	696	1101.567	EIDGGLETLR		1	Proteotypic
A8K766_HUMAN	1494	1338.714	LSVISVEDPPQR		1	Proteotypic
A8K797_HUMAN	173	956.565	GDSVIVVLR		1	Proteotypic
A8K797_HUMAN	1139	1221.577	NNTQVLINCR	Trioxidation (C)	1	Proteotypic
A8K797_HUMAN	1145	1222.539	SEMTPEELQK	Dioxidation (M)	1	Proteotypic
A8K7A4_HUMAN	2429	1155.698	VLVTGATGLLGR		1	Proteotypic
A8K7C2_HUMAN	81	922.560	IIAPPERK		2	Shared
A8K7C2_HUMAN	142	944.544	AVFPSIVGR		2	Shared
A8K7C2_HUMAN	230	975.441	AGFAGDDAPR		2	Shared
A8K7C2_HUMAN	362	1013.474	DLTDYLMK	Oxidation (M)	2	Shared
A8K7C2_HUMAN	424	1029.469	DLTDYLMK	Dioxidation (M)	2	Shared
A8K7C2_HUMAN	445	1035.644	IKIAPPER		2	Shared

A8K7C2_HUMAN	820	1131.520	GYSFTTTAER		2	Shared
A8K7C2_HUMAN	939	1163.739	IKIIPPERK		2	Shared
A8K7C2_HUMAN	976	1176.606	EITALAPSTMK	Oxidation (M)	2	Shared
A8K7C2_HUMAN	1037	1192.601	EITALAPSTMK	Dioxidation (M)	2	Shared
A8K7C2_HUMAN	1051	1197.515	DSYVGDEAQS		2	Shared
A8K7C2_HUMAN	1054	1197.698	AVFPSIVGRPR		2	Shared
A8K7C2_HUMAN	1190	1234.543	HQGVVMVGMGQK	Dioxidation (M); Dioxidation (M)	2	Shared
A8K7C2_HUMAN	1529	1353.616	DSYVGDEAQS		2	Shared
A8K7C2_HUMAN	1813	1515.695	QEYDESGPSIVHR		2	Shared
A8K7C2_HUMAN	2122	1789.885	SYELPDGQVITIGNER		2	Shared
A8K7C2_HUMAN	2201	1899.808	EEEEIAALVIDNGSGMCK	Acetyl (Protein N-term); Dioxidation (M); Trioxidation (C)	1	Proteotypic
A8K7C2_HUMAN	2237	1953.057	VAPEEHPVLLTEAPLNPK		2	Shared
A8K7C2_HUMAN	2321	2246.053	DLYANTVLSGGTTMYPGIADR	Dioxidation (M)	2	Shared
A8K7C2_HUMAN	2360	2572.120	LCYVALDFEQEMATAASSSSLEK	Dioxidation (M); Trioxidation (C)	2	Shared
A8K7C2_HUMAN	2374	2829.257	EKLCYVALDFEQEMATAASSSSLEK	Dioxidation (M); Trioxidation (C)	2	Shared
A8K7F6_HUMAN	172	956.529	ELAQQIQK		1	Proteotypic
A8K7F6_HUMAN	743	1113.676	VLITDILLAR		1	Proteotypic
A8K7F6_HUMAN	1015	1186.656	KEELTLEGIR		1	Proteotypic
A8K7F6_HUMAN	1945	1618.685	MFVLDEADEMLSR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K7J7_HUMAN	410	1026.619	GAALITAVGVR		1	Proteotypic
A8K7J7_HUMAN	667	1093.540	FLSQIESDR		2	Shared
A8K7N0_HUMAN	1165	1227.657	GTAAAAAAAAAAAAAK		1	Proteotypic
A8K7N0_HUMAN	1532	1353.762	LVAIVDVIDQNR		1	Proteotypic
A8K7S1_HUMAN	562	1070.634	LLLIGDSGVGK		3	Shared
A8K7S1_HUMAN	1754	1472.704	MGPGAASGGERP	Dioxidation (M)	1	Proteotypic
A8K7S1_HUMAN	2069	1723.867	EFADSLGIPFLETS		2	Shared
A8K7S1_HUMAN	2223	1931.860	NATNVEQAFMTMAAEIK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K7X6_HUMAN	201	966.419	MREESGAR	Dioxidation (M)	1	Proteotypic
A8K7X6_HUMAN	921	1157.568	IANPVEGSTDR		1	Proteotypic
A8K7X6_HUMAN	1328	1278.551	INISEGNCPER	Trioxidation (C)	2	Shared
A8K7X6_HUMAN	2293	2120.964	ESTGAQVQVAGDMLPNSTER	Dioxidation (M)	2	Shared
A8K844_HUMAN	385	1018.457	EGASEEIR		1	Proteotypic
A8K844_HUMAN	554	1067.514	YLGGTDDTVK		1	Proteotypic
A8K844_HUMAN	564	1071.520	TPEELQAER		1	Proteotypic
A8K844_HUMAN	1274	1259.651	FLSLDLQGDPR		1	Proteotypic
A8K844_HUMAN	1406	1304.671	TEEELLETTLK		1	Proteotypic
A8K844_HUMAN	1516	1347.715	AGQLLSELFTR		1	Proteotypic
A8K844_HUMAN	1614	1397.628	TTTSAVMVHCLR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
A8K844_HUMAN	1760	1472.845	AISESGVALTSVLVK		1	Proteotypic
A8K844_HUMAN	1885	1560.825	QKTEEELLETTLK		1	Proteotypic
A8K844_HUMAN	1923	1590.800	EGYLQIGANTQAAQK		1	Proteotypic
A8K844_HUMAN	2087	1742.725	DAGAPTYMYEFQYR	Dioxidation (M)	1	Proteotypic
A8K844_HUMAN	2099	1760.898	GDVKPLAEQIATAGCK	Trioxidation (C)	1	Proteotypic
A8K844_HUMAN	2178	1858.012	ESQPLLGTVIDGMLLK	Dioxidation (M)	1	Proteotypic
A8K844_HUMAN	2197	1888.993	KGDVKPLAEQIATAGCK	Trioxidation (C)	1	Proteotypic
A8K844_HUMAN	2292	2119.003	LSEDCLYLNIYTPADLTK	Trioxidation (C)	1	Proteotypic



A8K844_HUMAN	2308	2148.089	TVIGDHGDELFSVFGAPFLK		1	Proteotypic
A8K8U1_HUMAN	617	1082.670	ALTLIAGSPLK		1	Proteotypic
A8K8U1_HUMAN	1012	1186.609	MLTFLMLVR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
A8K8U1_HUMAN	1261	1255.775	LGTLSDILIK		1	Proteotypic
A8K8U1_HUMAN	1951	1620.825	SVILEAFSSPSEEVK		1	Proteotypic
A8K987_HUMAN	12	900.482	FLQPGSPR		1	Proteotypic
A8K987_HUMAN	291	991.570	AILNYIASK		1	Proteotypic
A8K987_HUMAN	313	998.612	ISNLPTVKK		1	Proteotypic
A8K987_HUMAN	482	1045.540	SAEDLDKLR		1	Proteotypic
A8K989_HUMAN	1185	1233.635	VLFDKDVNER		1	Proteotypic
A8K989_HUMAN	1548	1359.740	ELSAVTFPDIIR		1	Proteotypic
A8K989_HUMAN	1724	1456.723	VMAAENIPENPLK	Dioxidation (M)	1	Proteotypic
A8K989_HUMAN	1733	1461.713	GLNVDQLNMLGEK	Dioxidation (M)	1	Proteotypic
A8K9A4_HUMAN	132	942.565	VPPPPPIAR		1	Proteotypic
A8K9A4_HUMAN	300	994.497	SDVEAIFSK		1	Proteotypic
A8K9A4_HUMAN	783	1122.592	KSDVEAIFSK		1	Proteotypic
A8K9A4_HUMAN	1167	1227.719	LKGDDLQAIKK		1	Proteotypic
A8K9A4_HUMAN	1444	1315.786	VFIGNLNTLVVK		1	Proteotypic
A8K9A4_HUMAN	1644	1414.767	QKVDSLLENLEK		1	Proteotypic
A8K9A4_HUMAN	2062	1713.897	MIAGQVLDINLAAEPK	Dioxidation (M)	1	Proteotypic
A8K9J2_HUMAN	55	913.506	GFGFVLFK		1	Proteotypic
A8K9J2_HUMAN	1664	1425.629	ESESVDKVMQK	Dioxidation (M)	1	Proteotypic
A8K9J2_HUMAN	1788	1487.751	IFVGGLSPDTPPEEK		1	Proteotypic
A8K9U9_HUMAN	2423	1116.654	DALLLIFANK		1	Proteotypic
A8K9W7_HUMAN	2425	1122.556	FAEALGSTEAK		1	Proteotypic
A8K9X4_HUMAN	2456	1289.625	FDGILTEGEGPR		1	Proteotypic
A8K9Z9_HUMAN	2424	1121.550	VATMLATGGNR	Dioxidation (M)	1	Proteotypic
A8KAH9_HUMAN	59	914.457	EQGQNLAR		1	Proteotypic
A8KAH9_HUMAN	260	984.597	LVVLGSGGVGK		1	Proteotypic
A8KAH9_HUMAN	1799	1493.788	INVNEIFYDLVR		1	Proteotypic
A8KAI9_HUMAN	753	1115.582	LLDSLPSDTR		1	Proteotypic
A8KAI9_HUMAN	1004	1184.692	FQPTLLTLPR		1	Proteotypic
A8MTA6_HUMAN	1510	1343.723	LVILANNCPALR	Trioxidation (C)	1	Proteotypic
A8MTA6_HUMAN	2091	1747.866	VCTLAIIDPGDSDIIR	Trioxidation (C)	1	Proteotypic
A8MTE3_HUMAN	19	902.482	ADAETLRK		1	Proteotypic
A8MTE3_HUMAN	164	953.533	FITIFGTR		1	Proteotypic
A8MTE3_HUMAN	207	969.440	GDTSGDYKK		1	Proteotypic
A8MTE3_HUMAN	322	1000.592	VLTEIIASR		1	Proteotypic
A8MTE3_HUMAN	710	1105.577	SEIDLFNIR		1	Proteotypic
A8MTE3_HUMAN	911	1154.568	GAGTDDHTLIR		1	Proteotypic
A8MTE3_HUMAN	929	1158.679	LIVALMKPSR	Dioxidation (M)	1	Proteotypic
A8MTE3_HUMAN	1020	1187.670	MLVLLQANR	Dioxidation (M)	1	Proteotypic
A8MTE3_HUMAN	1188	1233.672	SEIDLFNIRK		1	Proteotypic
A8MTE3_HUMAN	1408	1305.627	NFATSLYSMIK	Dioxidation (M)	1	Proteotypic
A8MTE3_HUMAN	1495	1339.604	GTVTDFPGFDER		1	Proteotypic
A8MTE3_HUMAN	1709	1445.761	DLLDDLKSELTGK		1	Proteotypic

A8MTE3_HUMAN	1940	1612.904	ETSGNLEQLLAVVK		1	Proteotypic
A8MTE3_HUMAN	2055	1703.894	GLGTDEESILTLTSTR		1	Proteotypic
A8MTE3_HUMAN	2175	1849.967	DLLDDLKSELTKFEK		1	Proteotypic
A8MU63_HUMAN	650	1090.508	NPAGSVVMER	Dioxidation (M)	1	Proteotypic
A8MU63_HUMAN	2328	2281.232	VVPLADIITPNQFEALLSGR		1	Proteotypic
A8MUB1_HUMAN	258	984.560	DVNAAIAAIK		1	Proteotypic
A8MUB1_HUMAN	398	1022.442	EDAANNYAR		3	Shared
A8MUB1_HUMAN	558	1068.582	EIIDPVLDR		1	Proteotypic
A8MUB1_HUMAN	1635	1409.767	QLFHPEQLITGK		3	Shared
A8MUB1_HUMAN	1786	1486.872	LISQIVSSITASLR		3	Shared
A8MUB1_HUMAN	2063	1714.914	AVFVDLEPTVIDEIR		1	Proteotypic
A8MUB1_HUMAN	2066	1717.875	NLDIERPTYTNLNR		4	Shared
A8MUB1_HUMAN	2154	1823.978	VGINYQPPTVVPGGDLAK		3	Shared
A8MUB1_HUMAN	2347	2408.201	FDGALNVDLTEFQTNLVPYPR		3	Shared
A8MUJ2_HUMAN	444	1035.633	LLAEPVPGIK		1	Proteotypic
A8MUJ2_HUMAN	1075	1202.589	TNEAQAIETAR		1	Proteotypic
A8MUS3_HUMAN	1212	1240.678	VNTLIRPDGEK		1	Proteotypic
A8MUS3_HUMAN	1622	1403.693	LAPDYDALDVANK		1	Proteotypic
A8MVA6_HUMAN	253	983.515	GLEGERPAR		1	Proteotypic
A8MVA6_HUMAN	728	1108.653	IAIYELLFK		1	Proteotypic
A8MVA6_HUMAN	1693	1440.663	AEAGAGSATEFQFR		1	Proteotypic
A8MVT9_HUMAN	365	1013.551	ETPALTINR		1	Proteotypic
A8MVT9_HUMAN	1642	1414.662	DFTATDLSEFAAK		1	Proteotypic
A8MW15_HUMAN	618	1083.531	DGPALSGPQSR		1	Proteotypic
A8MW15_HUMAN	1112	1213.667	ALQEALVLSDR		1	Proteotypic
A8MW15_HUMAN	1115	1214.626	VAAALDDGSALGR		1	Proteotypic
A8MW15_HUMAN	1705	1444.771	VAAALTAMDKPLGR	Dioxidation (M)	1	Proteotypic
A8MW15_HUMAN	1792	1490.727	FGGAAVFPNQEQR		1	Proteotypic
A8MW15_HUMAN	1840	1528.909	ALPLALVLHELGAQR		1	Proteotypic
A8MXW0_HUMAN	2491	1750.830	AEEYEFLTPVEEAPK		1	Proteotypic
A8MYV9_HUMAN	2392	962.547	VTNILMLK	Dioxidation (M)	1	Proteotypic
A8MZ70_HUMAN	232	975.539	QLFDQVVK		2	Shared
A8MZ70_HUMAN	478	1044.556	ALTSELANAR		1	Proteotypic
A8MZ70_HUMAN	636	1086.680	FVIKPIDKK		2	Shared
A8MZ70_HUMAN	671	1094.491	EKEELMER	Dioxidation (M)	1	Proteotypic
A8MZ70_HUMAN	992	1181.587	APDFVFYAPR		2	Shared
A8MZ70_HUMAN	1138	1221.555	AQMVEDLEK	Dioxidation (M)	1	Proteotypic
A8MZ70_HUMAN	1805	1503.772	RKPDITIEVQQMK	Dioxidation (M)	2	Shared
A8MZ76_HUMAN	27	904.513	KQVEVFR		1	Proteotypic
A8MZ76_HUMAN	784	1122.592	VEAFQTTISK		1	Proteotypic
A8MZ76_HUMAN	1525	1351.735	TKVEAFQTTISK		1	Proteotypic
A8MZ76_HUMAN	1849	1533.731	IEDGNDFGVAIQEK		1	Proteotypic
AACT_HUMAN	182	959.504	ADLSGITGAR		1	Proteotypic
AACT_HUMAN	668	1093.624	NLAVSQVVHK		1	Proteotypic
AATC_HUMAN	682	1098.592	LALGDDSPALK		1	Proteotypic
AATC_HUMAN	1309	1270.699	VGGVQSLGGTGALR		1	Proteotypic

AATM_HUMAN	1293	1266.730	IAAAILNTPDLR		1	Proteotypic
AATM_HUMAN	1715	1448.799	FVTVQTISGTGALR		1	Proteotypic
ACADS_HUMAN	1055	1198.532	IAMQTLDMGR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
ACADS_HUMAN	1365	1291.644	LADMALALESAR	Dioxidation (M)	1	Proteotypic
ACADV_HUMAN	97	927.503	ELVEPVSR		1	Proteotypic
ACADV_HUMAN	111	933.528	ITAFVVER		1	Proteotypic
ACADV_HUMAN	977	1176.614	IFEGTNDILR		1	Proteotypic
ACADV_HUMAN	1271	1259.564	FGMAAALAGTMR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
ACADV_HUMAN	1407	1304.698	ALEQFATVVEAK		1	Proteotypic
ACADV_HUMAN	1442	1315.725	GIVNEQFLLQR		1	Proteotypic
ACADV_HUMAN	1817	1517.773	VPSENVLGEVSGGFK		1	Proteotypic
ACTB_HUMAN	81	922.560	IIAPPERK		2	Shared
ACTB_HUMAN	142	944.544	AVFPSIVGR		2	Shared
ACTB_HUMAN	230	975.441	AGFAGDDAPR		2	Shared
ACTB_HUMAN	362	1013.474	DLTDYLMK	Oxidation (M)	2	Shared
ACTB_HUMAN	424	1029.469	DLTDYLMK	Dioxidation (M)	2	Shared
ACTB_HUMAN	445	1035.644	IKIAPPER		2	Shared
ACTB_HUMAN	820	1131.520	GYSFTTTAER		2	Shared
ACTB_HUMAN	939	1163.739	IKIAPPERK		2	Shared
ACTB_HUMAN	976	1176.606	EITALAPSTMK	Oxidation (M)	2	Shared
ACTB_HUMAN	1037	1192.601	EITALAPSTMK	Dioxidation (M)	2	Shared
ACTB_HUMAN	1051	1197.515	DSYVGDEAQS		2	Shared
ACTB_HUMAN	1054	1197.698	AVFPSIVGRPR		2	Shared
ACTB_HUMAN	1190	1234.543	HQGVVMVGMGQK	Dioxidation (M); Dioxidation (M)	2	Shared
ACTB_HUMAN	1529	1353.616	DSYVGDEAQS		2	Shared
ACTB_HUMAN	1813	1515.695	QEYDESGPSIVHR		2	Shared
ACTB_HUMAN	2122	1789.885	SYELPDGQVITIGNER		2	Shared
ACTB_HUMAN	2167	1843.745	DDDIAALVVDNGSGMCK	Acetyl (Protein N-term); Dioxidation (M); Trioxidation (C)	1	Proteotypic
ACTB_HUMAN	2237	1953.057	VAPEEHPVLLTEAPLNPK		2	Shared
ACTB_HUMAN	2321	2246.053	DLYANTVLSGGTTMYPGIADR	Dioxidation (M)	2	Shared
ACTB_HUMAN	2360	2572.120	LCYVALDFEQEMATAASSSSLEK	Dioxidation (M); Trioxidation (C)	2	Shared
ACTB_HUMAN	2374	2829.257	EKLCYVALDFEQEMATAASSSSLEK	Dioxidation (M); Trioxidation (C)	2	Shared
ACTN1_HUMAN	786	1123.529	AETAANRICK	Trioxidation (C)	2	Shared
ACTN1_HUMAN	1565	1369.593	GISQEQMNEFR	Dioxidation (M)	1	Proteotypic
ACTN1_HUMAN	1623	1403.769	LMLLLEVISGER	Dioxidation (M)	3	Shared
ACTN1_HUMAN	1674	1428.757	TINEVENQILTR		2	Shared
ACTN1_HUMAN	1851	1536.767	FAIQDISVEETSAK		3	Shared
ACTN1_HUMAN	1861	1541.830	LVSIGAEIIVDGNVK		2	Shared
ACTN1_HUMAN	2098	1760.741	ETADTDADQVMASFK	Dioxidation (M)	1	Proteotypic
ACTN2_HUMAN	1623	1403.769	LMLLLEVISGER	Dioxidation (M)	3	Shared
ACTN2_HUMAN	1740	1464.703	ISNRPAFMPSEGK	Dioxidation (M)	1	Proteotypic
ACTN2_HUMAN	1851	1536.767	FAIQDISVEETSAK		3	Shared
ACTN2_HUMAN	1861	1541.830	LVSIGAEIIVDGNVK		2	Shared
ACTN4_HUMAN	41	910.487	IAESNHIK		1	Proteotypic
ACTN4_HUMAN	151	949.425	QGEAEFNR		1	Proteotypic
ACTN4_HUMAN	786	1123.529	AETAANRICK	Trioxidation (C)	2	Shared

ACTN4_HUMAN	927	1158.563	STLPDADRER		1	Proteotypic
ACTN4_HUMAN	1589	1383.609	GISQEQMQEFR	Dioxidation (M)	1	Proteotypic
ACTN4_HUMAN	1623	1403.769	LMLLLEVISGER	Dioxidation (M)	3	Shared
ACTN4_HUMAN	1674	1428.757	TINEVENQILTR		2	Shared
ACTN4_HUMAN	1741	1464.703	LSNRPAFMPSEGK	Dioxidation (M)	1	Proteotypic
ACTN4_HUMAN	1812	1513.799	LVSIGAEIIVDGNK		1	Proteotypic
ACTN4_HUMAN	1851	1536.767	FAIQDISVEETSAK		3	Shared
ACTN4_HUMAN	2025	1674.894	VLGDKNFITAEELR		1	Proteotypic
ACTN4_HUMAN	2085	1740.805	ETTDTDADQVIASF		1	Proteotypic
ACTN4_HUMAN	2172	1847.857	MLDAEDIVNTARPEK	Dioxidation (M)	1	Proteotypic
ACTN4_HUMAN	2204	1902.932	KDDPVTNLNNAFEVAEK		1	Proteotypic
ACTN4_HUMAN	2215	1919.922	AGTQIENIDEDFRDGLK		1	Proteotypic
ADH1G_HUMAN	131	942.502	IDAASPLEK		1	Proteotypic
ADH1G_HUMAN	165	953.591	KPIQEVVK		1	Proteotypic
ADH1G_HUMAN	591	1075.566	INEGFDLLR		1	Proteotypic
ADH1G_HUMAN	797	1127.655	IIAVDINKDK		1	Proteotypic
ADT2_HUMAN	115	935.315	MMMQSGR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	2	Shared
ADT2_HUMAN	155	950.471	TDAAVSFAK	Acetyl (Protein N-term)	1	Proteotypic
ADT2_HUMAN	231	975.539	QIFLGGVDK		2	Shared
ADT2_HUMAN	655	1091.416	RMMMQSGR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	2	Shared
ADT2_HUMAN	823	1131.640	QIFLGGVDKR		1	Proteotypic
ADT2_HUMAN	839	1135.671	LLLQVQHASK		2	Shared
ADT2_HUMAN	1127	1218.661	DFLAGGVAAAISK		1	Proteotypic
ADT2_HUMAN	1707	1445.734	YFPTQALNFAFK		2	Shared
ADT3_HUMAN	78	920.472	IFRDEGGK		1	Proteotypic
ADT3_HUMAN	115	935.315	MMMQSGR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	2	Shared
ADT3_HUMAN	231	975.539	QIFLGGVDK		2	Shared
ADT3_HUMAN	440	1035.524	TEQAISFAK	Acetyl (Protein N-term)	1	Proteotypic
ADT3_HUMAN	655	1091.416	RMMMQSGR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	2	Shared
ADT3_HUMAN	839	1135.671	LLLQVQHASK		2	Shared
ADT3_HUMAN	1181	1232.676	DFLAGGIAAAISK		1	Proteotypic
ADT3_HUMAN	1707	1445.734	YFPTQALNFAFK		2	Shared
AGR2_HUMAN	1084	1204.645	VFAENKEIQK		1	Proteotypic
AGR2_HUMAN	1371	1292.680	IMFVDPSLTVR	Oxidation (M)	1	Proteotypic
AGR2_HUMAN	1418	1308.675	IMFVDPSLTVR	Dioxidation (M)	1	Proteotypic
AGR2_HUMAN	2290	2100.045	LYAYEPADTALLLDNMKK	Dioxidation (M)	1	Proteotypic
AHNK_HUMAN	9	900.455	ADVDISGPK		1	Proteotypic
AHNK_HUMAN	10	900.455	ADIDVSGPK		1	Proteotypic
AHNK_HUMAN	92	925.388	MPEMNIK	Dioxidation (M); Dioxidation (M)	2	Shared
AHNK_HUMAN	740	1113.603	LKGDVDVSGPK		1	Proteotypic
AHNK_HUMAN	915	1155.650	VKGDVDVSLPK		2	Shared
AHNK_HUMAN	1292	1266.646	AEGPEVDVNLPK		1	Proteotypic
AIBP_HUMAN	2399	994.520	KSATQFTGR		1	Proteotypic
AL1A1_HUMAN	267	986.565	ILDIESGK		1	Proteotypic
AL1A1_HUMAN	672	1094.561	VAFTGSTVEGK		1	Proteotypic
AL1A1_HUMAN	2034	1683.893	SSSGTDLPLVLLTDLK	Acetyl (Protein N-term)	1	Proteotypic

AL1B1_HUMAN	2468	1416.765	LAPALATGNTVVMK	Dioxidation (M)	1	Proteotypic
AL3A2_HUMAN	2452	1265.589	IAFGGETDEATR		1	Proteotypic
ALBU_HUMAN	85	923.475	LCTVATLR	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	95	926.486	LYEYIAR		1	Proteotypic
ALBU_HUMAN	121	937.490	LKCASLQK	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	123	939.441	DDNPNLPR		1	Proteotypic
ALBU_HUMAN	152	950.435	DLGEENFK		1	Proteotypic
ALBU_HUMAN	184	959.555	FQNALLVR		1	Proteotypic
ALBU_HUMAN	250	983.481	TYETTLEK		1	Proteotypic
ALBU_HUMAN	318	999.596	QTALVELVK		1	Proteotypic
ALBU_HUMAN	361	1012.592	LVAASQAALGL		1	Proteotypic
ALBU_HUMAN	546	1065.455	NECFLQHK	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	583	1073.535	LDELRDEGK		1	Proteotypic
ALBU_HUMAN	770	1119.417	CCTESLVNR	Trioxidation (C); Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	799	1127.691	KQTALVELVK		1	Proteotypic
ALBU_HUMAN	854	1140.687	KLVAASQAALGL		1	Proteotypic
ALBU_HUMAN	884	1148.608	LVNEVTEFAK		1	Proteotypic
ALBU_HUMAN	1017	1187.513	DVCKNYAEAK	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1154	1225.598	FKDLGEENFK		1	Proteotypic
ALBU_HUMAN	1527	1352.486	AAFTECCQAADK	Trioxidation (C); Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1538	1357.622	AVMDDFAAFVEK	Oxidation (M)	1	Proteotypic
ALBU_HUMAN	1574	1373.617	AVMDDFAAFVEK	Dioxidation (M)	1	Proteotypic
ALBU_HUMAN	1587	1381.597	VGSKCCKHPEAK	Trioxidation (C); Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1680	1433.598	YICENQDSISSK	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1712	1447.442	ETYGEMADCCAK	Dioxidation (M); Trioxidation (C); Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1713	1447.628	VTKCCTESLVNR	Trioxidation (C); Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1748	1466.836	RHPDYSVLLLR		1	Proteotypic
ALBU_HUMAN	1774	1479.498	TCVADESAENCDK	Trioxidation (C); Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1790	1489.588	ADDKETCFAEEGK	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1807	1510.836	VPQVSTPTLVEVSR		1	Proteotypic
ALBU_HUMAN	1837	1527.716	LKECCEKPLLEK	Trioxidation (C); Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1944	1617.683	ADDKETCFAEEGKK	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1971	1638.775	DVFLGMFLYEYAR	Oxidation (M)	1	Proteotypic
ALBU_HUMAN	1972	1638.930	KVPQVSTPTLVEVSR		1	Proteotypic
ALBU_HUMAN	1982	1647.709	QNCELFEQLGEYK	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	1996	1654.770	DVFLGMFLYEYAR	Dioxidation (M)	1	Proteotypic
ALBU_HUMAN	2202	1900.888	RPCFSALEVDETYVPK	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	2218	1921.993	SLHTLFGDKLCTVATLR	Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	2275	2044.088	VFDEFKPLVEEPQNLK		1	Proteotypic
ALBU_HUMAN	2365	2672.210	LVRPEVDVMCTAFHDNEETFLK	Dioxidation (M); Trioxidation (C)	1	Proteotypic
ALBU_HUMAN	2373	2800.305	LVRPEVDVMCTAFHDNEETFLKK	Dioxidation (M); Trioxidation (C)	1	Proteotypic
ALDH2_HUMAN	843	1136.583	VAFTGSTEIGR		1	Proteotypic
ALDH2_HUMAN	1008	1185.672	VIQVAAGSSNLK		1	Proteotypic
ALDH2_HUMAN	1621	1400.770	LGPALATGNVVMK	Dioxidation (M)	1	Proteotypic
ALDH2_HUMAN	1685	1434.743	EEIFGPVMQILK	Dioxidation (M)	1	Proteotypic
ALDOA_HUMAN	124	939.477	ELSDIAHR		1	Proteotypic

ALDOA_HUMAN	473	1043.561	QLLLTADDR		1	Proteotypic
ALDOA_HUMAN	665	1092.556	AAQEYVVKR		1	Proteotypic
ALDOA_HUMAN	781	1122.534	ALANSLACQ GK	Trioxidation (C)	1	Proteotypic
ALDOA_HUMAN	1481	1331.693	GILAADESTGSI AK		1	Proteotypic
ALDOA_HUMAN	1505	1341.704	ADDGRPFQVIK		1	Proteotypic
ALDOA_HUMAN	1978	1645.802	LQSIGTENTEENRR		1	Proteotypic
ALDOA_HUMAN	2058	1706.830	FSHEEIAMATVTALR	Dioxidation (M)	1	Proteotypic
ALDOA_HUMAN	2131	1801.903	RLQSIGTENTEENRR		1	Proteotypic
ALDOA_HUMAN	2304	2138.079	IGEHTPSALAIMENANVLAR	Dioxidation (M)	1	Proteotypic
AMPL_HUMAN	334	1003.607	TLIEFLLR		1	Proteotypic
AMPL_HUMAN	1041	1193.629	GITFDSSGGISIK		1	Proteotypic
AMPL_HUMAN	1176	1231.667	GVLFFASGQNLAR		1	Proteotypic
AMPL_HUMAN	1447	1317.616	TIQVDNTDAEGR		1	Proteotypic
AMPL_HUMAN	2032	1681.729	QLMETPANEMTPTR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
AN32A_HUMAN	139	944.493	ELVLDNSR		1	Proteotypic
AN32A_HUMAN	221	973.519	KLLESDNR		1	Proteotypic
ANT3_HUMAN	2469	1420.738	EVPLNTIIFMGR	Dioxidation (M)	1	Proteotypic
ANX11_HUMAN	2408	1060.540	SETDLLDIR		1	Proteotypic
ANX13_HUMAN	211	971.529	AEVDLQGIK		1	Proteotypic
ANX13_HUMAN	792	1125.687	ILVSLQANR		1	Proteotypic
ANX13_HUMAN	933	1160.567	GAGTDEETLIR		1	Proteotypic
ANX13_HUMAN	982	1177.536	ASSPQGFDVDR		1	Proteotypic
ANX13_HUMAN	1795	1491.695	ASSPQGFDVDRDAK		1	Proteotypic
ANX13_HUMAN	1979	1645.879	TALALLDRPSEYAAR		1	Proteotypic
ANX13_HUMAN	2013	1662.825	GMGTTNEAAIIILSGR	Dioxidation (M)	1	Proteotypic
ANX13_HUMAN	2040	1688.749	NEGDDVDKDLAQDAK		1	Proteotypic
ANX13_HUMAN	2133	1804.821	DIEEAIEETSGDLQK		1	Proteotypic
ANXA2_HUMAN	441	1035.535	DLYDAGVVKR		1	Proteotypic
ANXA2_HUMAN	627	1085.478	AYTNFDAER		1	Proteotypic
ANXA2_HUMAN	633	1086.592	DALNIETAIK		1	Proteotypic
ANXA2_HUMAN	734	1110.546	QDIAFAYQR		1	Proteotypic
ANXA2_HUMAN	1140	1221.588	TPAQYDASELK		1	Proteotypic
ANXA2_HUMAN	1150	1224.562	DIISDTSGDFR		1	Proteotypic
ANXA2_HUMAN	1229	1243.616	TNQELQEINR		1	Proteotypic
ANXA2_HUMAN	1657	1420.687	SLYYIQQDTK		1	Proteotypic
ANXA2_HUMAN	1794	1491.655	SYSPYDMLESIR	Dioxidation (M)	1	Proteotypic
ANXA2_HUMAN	1862	1541.841	GVDEVTIVNILTNR		1	Proteotypic
ANXA2_HUMAN	1987	1649.972	SALSGHLETVILGLLK		1	Proteotypic
ANXA2_HUMAN	2104	1767.820	GLGTDEDSLIEIICSR	Trioxidation (C)	1	Proteotypic
ANXA2_HUMAN	2138	1810.858	TDLEKDIISDTSGDFR		1	Proteotypic
ANXA2_HUMAN	2169	1843.895	LSLEGDHSTPPSAYGSVK		1	Proteotypic
ANXA2_HUMAN	2207	1907.875	AEDGSVIDYELIDQDAR		1	Proteotypic
ANXA2_HUMAN	2227	1938.953	TDLEKDIISDTSGDFRK		1	Proteotypic
ANXA2_HUMAN	2282	2063.976	RAEDGSVIDYELIDQDAR		1	Proteotypic
ANXA4_HUMAN	177	957.561	VLVLSAGGR		1	Proteotypic
ANXA4_HUMAN	261	985.490	FLTVLCSR	Trioxidation (C)	1	Proteotypic

ANXA4_HUMAN	308	997.446	GDTSGDYRK		1	Proteotypic
ANXA4_HUMAN	651	1090.533	AEIDMLDIR	Oxidation (M)	1	Proteotypic
ANXA4_HUMAN	716	1106.528	AEIDMLDIR	Dioxidation (M)	1	Proteotypic
ANXA4_HUMAN	827	1133.481	SDTSFMFQR	Oxidation (M)	1	Proteotypic
ANXA4_HUMAN	888	1149.476	SDTSFMFQR	Dioxidation (M)	1	Proteotypic
ANXA4_HUMAN	967	1173.599	GLGTDDNTLIR		1	Proteotypic
ANXA4_HUMAN	1352	1287.703	ISQKDIEQSIK		1	Proteotypic
ANXA4_HUMAN	1559	1365.605	QDAQDLYEAGEK		1	Proteotypic
ANXA4_HUMAN	1568	1370.658	ISQTYQQQYGR		1	Proteotypic
ANXA4_HUMAN	1578	1378.636	DEGNLDDALVR		1	Proteotypic
ANXA4_HUMAN	1798	1493.700	QDAQDLYEAGEKK		1	Proteotypic
ANXA4_HUMAN	1927	1596.720	AASGFNAMEDAQLTR	Oxidation (M)	1	Proteotypic
ANXA4_HUMAN	1938	1612.715	AASGFNAMEDAQLTR	Dioxidation (M)	1	Proteotypic
ANXA4_HUMAN	1989	1651.772	GAGTDEGCLIEILASR	Trioxidation (C)	1	Proteotypic
ANXA4_HUMAN	2016	1665.846	SETSGSFEDALLAIVK		1	Proteotypic
ANXA4_HUMAN	2044	1691.873	GLGTDEDAIISVLAYR		1	Proteotypic
ANXA4_HUMAN	2086	1740.810	AASGFNAMEDAQLTRK	Dioxidation (M)	1	Proteotypic
ANXA4_HUMAN	2247	1977.874	SLEDDIRSDTSFMFQR	Dioxidation (M)	1	Proteotypic
ANXA4_HUMAN	2265	2013.935	AMKGAGTDEGCLIEILASR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
ANXA4_HUMAN	2303	2135.971	SETSGSFEDALLAIVKCMR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
ANXA4_HUMAN	2334	2318.187	VLVSLSAGRDEGNLDDALVR		1	Proteotypic
ANXA7_HUMAN	2418	1090.525	GAGTDDSTLVR		1	Proteotypic
AOFA_HUMAN	778	1121.547	FVGGSGQVSER		2	Shared
AOFA_HUMAN	934	1162.590	IMDLLGDQVK	Dioxidation (M)	1	Proteotypic
AOFA_HUMAN	1244	1249.642	KFVGGSGQVSER		2	Shared
AOFB_HUMAN	191	963.470	IMDLLGDR	Dioxidation (M)	1	Proteotypic
AOFB_HUMAN	778	1121.547	FVGGSGQVSER		2	Shared
AOFB_HUMAN	970	1174.594	IISTTNGGQER		1	Proteotypic
AOFB_HUMAN	1244	1249.642	KFVGGSGQVSER		2	Shared
AOFB_HUMAN	1966	1633.915	LLHDSGLNVVVLEAR		1	Proteotypic
AP1M2_HUMAN	87	923.515	YMKIIEK		1	Proteotypic
AP1M2_HUMAN	988	1179.596	VFLSGMPELR	Dioxidation (M)	1	Proteotypic
APT_HUMAN	780	1121.572	IDYIAGLDSR		1	Proteotypic
APT_HUMAN	1475	1328.657	ADSELQLVEQR	Acetyl (Protein N-term)	1	Proteotypic
APT_HUMAN	1742	1464.740	SFPDFPTPGVVFR		1	Proteotypic
APT_HUMAN	1831	1522.807	LAPVPFFSLLQYE		1	Proteotypic
ARF4_HUMAN	701	1102.639	DAVLLLFANK		1	Proteotypic
ARF4_HUMAN	767	1118.601	ILMVGLDAAGK	Dioxidation (M)	3	Shared
ARF4_HUMAN	961	1171.608	IQEVADELQK		1	Proteotypic
ARF4_HUMAN	2336	2322.236	LGEIVTTIPTIGFNVETVEYK		3	Shared
ARF6_HUMAN	756	1116.654	DAIILIFANK		1	Proteotypic
ARF6_HUMAN	826	1132.616	ILMLGLDAAGK	Dioxidation (M)	1	Proteotypic
ARK73_HUMAN	896	1150.529	MDAPTSAAVTR	Dioxidation (M)	1	Proteotypic
ARK73_HUMAN	1412	1306.630	RMDAPTSAAVTR	Dioxidation (M)	1	Proteotypic
ARK73_HUMAN	1827	1522.723	ARPATVLGAMEMGR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
ARK73_HUMAN	1896	1567.840	FYAFNPLAGLLTGK		1	Proteotypic

ARL1_HUMAN	222	973.596	AILVVFANK		1	Proteotypic
ARL1_HUMAN	559	1068.654	ILILGLDGAGK		1	Proteotypic
ARL8A_HUMAN	2414	1078.508	MNLSAIQDR	Dioxidation (M)	1	Proteotypic
ARLY_HUMAN	2404	1055.561	NPDSLELIR		1	Proteotypic
ARP3_HUMAN	570	1072.526	FMEQVIFK	Dioxidation (M)	1	Proteotypic
ARP3_HUMAN	1429	1312.645	NIVLSGGSTMFR	Dioxidation (M)	1	Proteotypic
ARP3_HUMAN	1631	1408.771	DITYFIQQLLR		1	Proteotypic
ARPC2_HUMAN	422	1028.612	IIEETLALK		1	Proteotypic
ARPC2_HUMAN	825	1132.591	MILLEVNNR	Dioxidation (M)	1	Proteotypic
ARPC4_HUMAN	347	1008.539	AENFFILR		1	Proteotypic
ARPC4_HUMAN	1009	1185.672	VLIEGSINSVR		1	Proteotypic
ARPC4_HUMAN	1298	1267.750	ELLQPVTISR		1	Proteotypic
ASXL2_HUMAN	2485	1651.784	GMGALIATNTTENSTR	Oxidation (M)	1	Proteotypic
AT1A1_HUMAN	241	978.513	GIVVYTGDR		1	Proteotypic
AT1A1_HUMAN	557	1068.568	GGDRIPADLR		1	Proteotypic
AT1A1_HUMAN	560	1069.486	NSVFQQGMK	Dioxidation (M)	1	Proteotypic
AT1A1_HUMAN	597	1077.549	LIIVEGCQR	Trioxidation (C)	1	Proteotypic
AT1A1_HUMAN	830	1133.550	AAVPDAVGKCR	Trioxidation (C)	1	Proteotypic
AT1A1_HUMAN	1170	1230.646	AVAGDASESALLK		1	Proteotypic
AT1A1_HUMAN	1197	1235.699	LNIPVSQVNPR		1	Proteotypic
AT1A1_HUMAN	1277	1260.671	IVEIPFNSTNK		1	Proteotypic
AT1A1_HUMAN	1390	1299.697	NMVPQQALVIR	Dioxidation (M)	1	Proteotypic
AT1A1_HUMAN	1639	1412.733	VIMVTGDHPITAK	Dioxidation (M)	1	Proteotypic
AT1A1_HUMAN	1818	1518.695	SPDFTNENPLETR		1	Proteotypic
AT1A1_HUMAN	1825	1521.734	ADIGVAMGIAGSDVSK	Dioxidation (M)	1	Proteotypic
AT1A1_HUMAN	1912	1583.867	AVFQANQENLPILK		1	Proteotypic
AT1A1_HUMAN	1946	1618.743	VDNSSLTGESEPQTR		1	Proteotypic
AT1A1_HUMAN	2026	1676.746	NIAFFSTNCVEGTAR	Trioxidation (C)	1	Proteotypic
AT1A1_HUMAN	2139	1810.943	QGAIVAVTGDGVNDSPALK		1	Proteotypic
AT1A1_HUMAN	2157	1828.917	GVGIISEGNETVEDIAR		1	Proteotypic
AT1A1_HUMAN	2180	1861.934	MSINAEVVVVDLVEVK	Dioxidation (M)	1	Proteotypic
AT1A1_HUMAN	2355	2495.185	QAADMILLDDNFASIVTGVEEGR	Dioxidation (M)	1	Proteotypic
ATPA_HUMAN	39	910.367	FNDGSDEK		1	Proteotypic
ATPA_HUMAN	317	999.571	VLSIGDGIAR		1	Proteotypic
ATPA_HUMAN	409	1025.587	AVDSLVPIGR		1	Proteotypic
ATPA_HUMAN	499	1051.497	RLTDADAMK	Dioxidation (M)	1	Proteotypic
ATPA_HUMAN	883	1148.608	GYLDKLEPSK		1	Proteotypic
ATPA_HUMAN	956	1170.624	VVDALGNAIDGK		1	Proteotypic
ATPA_HUMAN	1349	1286.687	HALIYDDLK		1	Proteotypic
ATPA_HUMAN	1422	1309.655	EIVTNFLAGFEA		1	Proteotypic
ATPA_HUMAN	1443	1315.735	TSIAIDIINQK		1	Proteotypic
ATPA_HUMAN	1722	1454.656	TGTAEMSSILEER	Dioxidation (M)	1	Proteotypic
ATPA_HUMAN	1874	1550.834	LKEIVTNFLAGFEA		1	Proteotypic
ATPA_HUMAN	1905	1574.779	ILGADTSVDLEETGR		1	Proteotypic
ATPA_HUMAN	1958	1623.883	TGAIVDVPVGEELLGR		1	Proteotypic
ATPA_HUMAN	2051	1698.777	NVQAEEMVEFSSGLK	Dioxidation (M)	1	Proteotypic



ATPA_HUMAN	2301	2135.020	GMSLNLEPDNVGVVFGNDK	Dioxidation (M)	1	Proteotypic
ATPA_HUMAN	2340	2337.160	EVAAFAGFGSDLDAATQQLLSR		1	Proteotypic
ATPD_HUMAN	263	985.544	IEANEALVK		1	Proteotypic
ATPD_HUMAN	1676	1430.700	AQAELVGTAEATR		1	Proteotypic
ATPG_HUMAN	433	1031.390	MTAMDNASK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
ATPG_HUMAN	493	1046.561	SEVATLTAAGK		1	Proteotypic
ATPG_HUMAN	659	1091.553	EVMLVGIGDK	Dioxidation (M)	1	Proteotypic
ATPG_HUMAN	670	1094.484	ESTTSEQSAR		1	Proteotypic
ATPG_HUMAN	1432	1313.708	ELIEIISGAAALD		1	Proteotypic
ATPO_HUMAN	407	1024.628	VAQILKEPK		1	Proteotypic
ATPO_HUMAN	712	1105.613	SFLSQGQVLK		1	Proteotypic
ATPO_HUMAN	932	1159.660	VAASVLNPPYVK		1	Proteotypic
ATPO_HUMAN	1360	1289.665	TDPSILGGMIVR	Dioxidation (M)	1	Proteotypic
ATPO_HUMAN	1911	1581.899	LVRPPVQVYGIEGR		1	Proteotypic
ATPO_HUMAN	2186	1872.010	FSPLTTNLINLLAENGR		1	Proteotypic
BAP31_HUMAN	135	943.497	LDVGNAEVK		1	Proteotypic
BAP31_HUMAN	285	990.462	QAESASEAAK		1	Proteotypic
BAP31_HUMAN	333	1003.555	LKDELASTK		1	Proteotypic
BAP31_HUMAN	535	1062.513	AENQVLAMR	Dioxidation (M)	1	Proteotypic
BAP31_HUMAN	1886	1561.729	LQAAVDGPMDDKKEE	Dioxidation (M)	1	Proteotypic
CAH2_HUMAN	112	934.451	GGPLDGTYS		1	Proteotypic
CAH2_HUMAN	255	983.576	VGSAKPGLQK		1	Proteotypic
CAH2_HUMAN	268	986.565	VVDVLSIK		1	Proteotypic
CAH2_HUMAN	953	1168.515	SADFTNFDPR		1	Proteotypic
CAH2_HUMAN	1425	1310.647	QSPVDIDHTAK		1	Proteotypic
CAH2_HUMAN	2019	1667.961	AVQQPDGLAVLGIFLK		1	Proteotypic
CALM_HUMAN	1889	1562.746	ADQLTEEQIAEFK	Acetyl (Protein N-term)	1	Proteotypic
CALM_HUMAN	2168	1843.884	EAFSLFDKDGDTITTK		1	Proteotypic
CALR_HUMAN	224	974.478	GLQTSQDAR		2	Shared
CALR_HUMAN	288	991.461	QIDNPDYK		2	Shared
CALR_HUMAN	806	1129.493	FYGDEEKDK		1	Proteotypic
CALR_HUMAN	874	1146.577	LKEEEEDKK		1	Proteotypic
CALR_HUMAN	1936	1606.767	FYALSASFEPFSNK		1	Proteotypic
CALX_HUMAN	323	1001.430	EIEDPEDR		1	Proteotypic
CALX_HUMAN	637	1087.515	AEDEILNR		1	Proteotypic
CALX_HUMAN	880	1148.520	KDDTDDEIAK		1	Proteotypic
CALX_HUMAN	2106	1769.826	APVPTGEVYFADSFDR		1	Proteotypic
CALX_HUMAN	2182	1862.920	KIPNPDFFEDLEPFR		1	Proteotypic
CAN1_HUMAN	336	1005.524	SEQFINLR		1	Proteotypic
CAN1_HUMAN	416	1028.427	SMVNLMDR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
CAN1_HUMAN	454	1039.509	DFFLANASR		1	Proteotypic
CAN1_HUMAN	1521	1350.634	QLAGEDMEISVK	Dioxidation (M)	1	Proteotypic
CAPG_HUMAN	1375	1295.603	VSDATGQMNLTG	Dioxidation (M)	1	Proteotypic
CAPG_HUMAN	1599	1388.741	QAALQVAEGFISR		1	Proteotypic
CAPZB_HUMAN	344	1008.443	LVEDMENK	Dioxidation (M)	1	Proteotypic
CAPZB_HUMAN	725	1107.640	RLPPQQIEK		1	Proteotypic

CATD_HUMAN	464	1041.643	VSTLPAILTK		1	Proteotypic
CATD_HUMAN	480	1044.597	QPGITFIAAK		1	Proteotypic
CATD_HUMAN	1305	1270.602	FDGILGMAYPR	Dioxidation (M)	1	Proteotypic
CATD_HUMAN	2151	1818.794	DPDAQPGGELMLGGTDSK	Dioxidation (M)	1	Proteotypic
CATD_HUMAN	2252	1990.019	ISVNNVLPVFDNLMQQK	Dioxidation (M)	1	Proteotypic
CAYP1_HUMAN	99	928.509	GASGIQGLAR		1	Proteotypic
CAYP1_HUMAN	283	989.555	EAVIAAAFAK		1	Proteotypic
CAYP1_HUMAN	1119	1215.463	MDAVDATMEK	Acetyl (Protein N-term); Dioxidation (M); Dioxidation (M)	1	Proteotypic
CAYP1_HUMAN	1175	1231.604	SGDGVVTVDLLR		1	Proteotypic
CAYP1_HUMAN	1716	1449.710	NGSGTLDLEEFLLR		1	Proteotypic
CAYP1_HUMAN	1948	1619.782	LGLVLDQAEAEGVCR	Trioxidation (C)	1	Proteotypic
CBR1_HUMAN	979	1176.650	LFSGDVLTAR		1	Proteotypic
CBR1_HUMAN	1144	1221.639	VVNVSSIMSVR	Dioxidation (M)	1	Proteotypic
CBR1_HUMAN	1991	1651.864	GQAAVQQLQAEGLSPR		1	Proteotypic
CD166_HUMAN	1033	1190.655	ALFLETEQLK		1	Proteotypic
CD166_HUMAN	1879	1553.903	ESLTLIVEGKPKIK		1	Proteotypic
CH10_HUMAN	360	1012.555	GGEIQPVSVK		1	Proteotypic
CH10_HUMAN	1437	1314.751	VLQATVVAVGSGSK		1	Proteotypic
CH60_HUMAN	16	900.528	LSDGVAVLK		1	Proteotypic
CH60_HUMAN	48	911.580	VGLQVVAVK		1	Proteotypic
CH60_HUMAN	127	940.607	IGIEIIRK		1	Proteotypic
CH60_HUMAN	183	959.504	VTDALNATR		1	Proteotypic
CH60_HUMAN	186	960.478	APGFGDNRK		1	Proteotypic
CH60_HUMAN	351	1010.562	GANPVEIRR		1	Proteotypic
CH60_HUMAN	906	1152.759	LKVGLQVVAVK		1	Proteotypic
CH60_HUMAN	1116	1214.651	NAGVEGSLIVEK		1	Proteotypic
CH60_HUMAN	1178	1232.588	VGGTSDVEVNEK		1	Proteotypic
CH60_HUMAN	1550	1360.683	VGGTSDVEVNEKK		1	Proteotypic
CH60_HUMAN	1920	1587.891	GVMLAVDAVIAELKK	Dioxidation (M)	1	Proteotypic
CH60_HUMAN	1963	1631.811	VGGTSDVEVNEKKDR		1	Proteotypic
CH60_HUMAN	2012	1661.891	VGEVIVTKDDAMLLK	Dioxidation (M)	1	Proteotypic
CH60_HUMAN	2024	1674.861	AAVEEGIVLGGGCALLR	Trioxidation (C)	1	Proteotypic
CH60_HUMAN	2100	1761.809	CIPALDSLTPANEDQK	Trioxidation (C)	1	Proteotypic
CH60_HUMAN	2213	1918.064	ISSIQSIVPALEIANHR		1	Proteotypic
CH60_HUMAN	2238	1953.935	TLNDELEIIEGMKFDR	Dioxidation (M)	1	Proteotypic
CH60_HUMAN	2314	2176.112	ALMLQGVDLLADAVAVTMGPK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
CH60_HUMAN	2330	2294.153	IQEIIQLDVTTSEYEKEK		1	Proteotypic
CH60_HUMAN	2342	2364.326	KPLVIIAEDVDGEALSTLVLR		1	Proteotypic
CH60_HUMAN	2359	2559.241	LVQDVANNNTNEEAGDGTATVLR		1	Proteotypic
CISD1_HUMAN	2394	964.483	SLTSSSSVR	Acetyl (Protein N-term)	1	Proteotypic
CISY_HUMAN	307	996.586	DILADLIPK		1	Proteotypic
CISY_HUMAN	487	1046.426	SMSTEGLMK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
CISY_HUMAN	793	1126.650	ALGFPLERPK		1	Proteotypic
CISY_HUMAN	1096	1208.713	IVPNVLEQGGK		1	Proteotypic
CKAP4_HUMAN	265	985.581	LALQALTEK		1	Proteotypic
CKAP4_HUMAN	870	1145.593	ASVSQVEADLK		1	Proteotypic

CKAP4_HUMAN	1186	1233.645	QTESLESLSK		1	Proteotypic
CKAP4_HUMAN	1757	1472.772	SVGELPSTVESLQK		1	Proteotypic
CKAP4_HUMAN	2125	1795.947	VQSLQATFGTFESILR		1	Proteotypic
CLH1_HUMAN	1297	1267.648	TLQIFNIEMK	Dioxidation (M)	1	Proteotypic
CLH1_HUMAN	1401	1303.652	NNLAGAEELFAR		1	Proteotypic
CLH1_HUMAN	1482	1333.626	IYIDSNNNPER		1	Proteotypic
CLH1_HUMAN	1801	1497.713	VIQCFAETGQVQK	Trioxidation (C)	1	Proteotypic
CLH1_HUMAN	1921	1588.788	RPISADSAIMNPASK	Dioxidation (M)	1	Proteotypic
CLH1_HUMAN	2248	1977.997	AFMTADLPNELIELLEK	Dioxidation (M)	1	Proteotypic
CLIC1_HUMAN	448	1036.534	GFTIPEAFR		1	Proteotypic
CLIC1_HUMAN	555	1068.466	IGNCPFSQR	Trioxidation (C)	1	Proteotypic
CLIC1_HUMAN	1333	1280.661	GVTFNVTTVDTK		1	Proteotypic
CLIC1_HUMAN	1471	1327.637	NSNPALNDNLEK		1	Proteotypic
CLIC1_HUMAN	1729	1457.740	AEEQPQVELFVK	Acetyl (Protein N-term)	1	Proteotypic
CLIC1_HUMAN	2170	1843.968	LAALNPESNTAGLDIFAK		1	Proteotypic
CNDP2_HUMAN	30	905.522	AALTTLFK	Acetyl (Protein N-term)	1	Proteotypic
CNDP2_HUMAN	736	1111.599	TGQEIPVNVR		1	Proteotypic
CNDP2_HUMAN	794	1127.484	MMEVAAADVK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
CNDP2_HUMAN	1179	1232.640	TVFGVEPDLTR		1	Proteotypic
CNDP2_HUMAN	1433	1313.719	QLGGSVELVDIGK		1	Proteotypic
CNDP2_HUMAN	2037	1685.972	LPDGSEIPLPPILLGR		1	Proteotypic
CNDP2_HUMAN	2075	1733.884	EGGSIPVTLTFQEATGK		1	Proteotypic
CNDP2_HUMAN	2229	1942.125	QKLPDGSEIPLPPILLGR		1	Proteotypic
CO1A1_HUMAN	137	944.431	QGSPGASGER		1	Proteotypic
CO1A1_HUMAN	1376	1296.606	GESGSPGAGPTGAR		1	Proteotypic
CO1A1_HUMAN	1869	1545.790	GETGPAGPAGPVGPGAR		1	Proteotypic
CO1A1_HUMAN	2041	1689.771	DGEAGAQQPPGPAGPAGER		1	Proteotypic
CO1A1_HUMAN	2242	1960.972	SGDRGETGPAGPAGPVGPGAR		1	Proteotypic
CO1A2_HUMAN	1192	1234.605	GEAGAAGPAGPAGPR		1	Proteotypic
CO1A2_HUMAN	1887	1561.785	GETGSPGVPAGAVGPR		1	Proteotypic
COF1_HUMAN	506	1056.422	MLPDKDCR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
COF1_HUMAN	821	1131.614	KSSTPEEVKK		1	Proteotypic
COF1_HUMAN	864	1143.614	ASGVAVSDGVIK	Acetyl (Protein N-term)	1	Proteotypic
COF1_HUMAN	1387	1299.638	AVLFCLSEDKK	Trioxidation (C)	1	Proteotypic
COF1_HUMAN	1497	1339.771	LGGSAVISLEGKPL		1	Proteotypic
COF1_HUMAN	2382	3061.586	NIILEEGKEILVGDVGQTVDDPYATFVK		1	Proteotypic
CPSF5_HUMAN	193	963.506	LMTEILGR	Dioxidation (M)	1	Proteotypic
CPSF5_HUMAN	757	1116.654	LFLVQLQEK		1	Proteotypic
CTNA1_HUMAN	1240	1247.654	NLMNAVVTQTVK	Dioxidation (M)	1	Proteotypic
CTNA1_HUMAN	1350	1287.649	AIMAQLPQEQK	Dioxidation (M)	1	Proteotypic
CTNA1_HUMAN	1576	1377.678	IAEQVASFQEEK		1	Proteotypic
CTNA1_HUMAN	1845	1531.788	QIIVDPLSFSEER		1	Proteotypic
DAD1_HUMAN	485	1045.577	SASVVSISR	Acetyl (Protein N-term)	1	Proteotypic
DAD1_HUMAN	763	1118.536	ADFQGISPER		1	Proteotypic
DBLOH_HUMAN	2435	1170.624	LAEAQIEELR		1	Proteotypic
DCMC_HUMAN	431	1030.567	GFGPGLTARR		1	Proteotypic

DCMC_HUMAN	859	1142.604	LASGQAAGALER		1	Proteotypic
DDAH1_HUMAN	673	1094.562	AGLGHPAAFGR	Acetyl (Protein N-term)	1	Proteotypic
DDAH1_HUMAN	931	1159.583	SAKGEEVDVAR		1	Proteotypic
DDX17_HUMAN	286	990.513	QLAEDFLR		1	Proteotypic
DDX17_HUMAN	392	1020.549	LIDFLESGK		1	Proteotypic
DDX17_HUMAN	1045	1194.555	DMVGIAQTGSGK	Dioxidation (M)	1	Proteotypic
DDX17_HUMAN	1156	1225.703	APILIATDVASR		2	Shared
DDX17_HUMAN	1467	1324.699	VLEEANQAINPK		1	Proteotypic
DDX17_HUMAN	1617	1399.611	MLDMGFEPQIR	Dioxidation (M); Dioxidation (M)	2	Shared
DDX5_HUMAN	804	1128.650	QVSDLISVLR		1	Proteotypic
DDX5_HUMAN	1156	1225.703	APILIATDVASR		2	Shared
DDX5_HUMAN	1617	1399.611	MLDMGFEPQIR	Dioxidation (M); Dioxidation (M)	2	Shared
DECR_HUMAN	737	1111.639	FNVIQPGPIK		1	Proteotypic
DECR_HUMAN	980	1176.650	VAFITGGGTGLGK		1	Proteotypic
DECR_HUMAN	1135	1220.586	AMLPPNSFQGK	Dioxidation (M)	1	Proteotypic
DECR_HUMAN	1483	1333.647	ATAEQISSQTGNK		1	Proteotypic
DESP_HUMAN	266	986.540	DLKDEIVR		1	Proteotypic
DESP_HUMAN	325	1001.514	VQEQLTR		1	Proteotypic
DESP_HUMAN	472	1043.561	VLLQEEGTR		1	Proteotypic
DESP_HUMAN	479	1044.556	GIVDSITGQR		1	Proteotypic
DESP_HUMAN	511	1057.450	FGDSNTVMR	Dioxidation (M)	1	Proteotypic
DESP_HUMAN	520	1058.547	SQLQISNNR		1	Proteotypic
DESP_HUMAN	574	1072.551	GAGSIAGASAPK		1	Proteotypic
DESP_HUMAN	801	1128.603	IEVLEEELR		1	Proteotypic
DESP_HUMAN	871	1145.629	LQSLTENLTK		1	Proteotypic
DESP_HUMAN	1128	1219.568	ETQSQLETER		1	Proteotypic
DESP_HUMAN	1255	1253.593	AITGFDDPFSGK		1	Proteotypic
DESP_HUMAN	1308	1270.699	QLQNIQATSR		1	Proteotypic
DESP_HUMAN	1310	1271.647	RVEEDIQQQK		1	Proteotypic
DEST_HUMAN	1215	1241.633	AVIFCLSADKK	Trioxidation (C)	1	Proteotypic
DEST_HUMAN	1459	1322.577	ASGVQVADEVCR	Acetyl (Protein N-term); Trioxidation (C)	1	Proteotypic
DHB11_HUMAN	167	954.477	HGLEETAAK		1	Proteotypic
DHB11_HUMAN	1068	1200.617	LMHGILTEQK	Dioxidation (M)	1	Proteotypic
DHB11_HUMAN	1904	1571.877	TLTDELAALQITGVK		1	Proteotypic
DHB11_HUMAN	2185	1869.991	MIFIPSSIAFLTTLR	Dioxidation (M)	1	Proteotypic
DHB11_HUMAN	2226	1937.969	NPSTSLGPTLEPEEVVNR		1	Proteotypic
DHB2_HUMAN	1281	1262.563	AAVTMFSSVMR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
DHB2_HUMAN	1894	1567.697	LVNVSSMGGGAPMER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
DHB4_HUMAN	2433	1168.693	VVLVTGAGAGLGR		1	Proteotypic
DHE3_HUMAN	188	962.519	YNLGLDLR		1	Proteotypic
DHE3_HUMAN	1209	1240.582	YSTDVSVDEVK		1	Proteotypic
DHX15_HUMAN	192	963.481	VAAMSVAQR	Dioxidation (M)	1	Proteotypic
DHX15_HUMAN	1410	1305.721	TLATDILMGVLK	Dioxidation (M)	1	Proteotypic
DHX9_HUMAN	105	930.513	ISAVSVAER		1	Proteotypic
DHX9_HUMAN	331	1002.586	LGGIGQFLAK		1	Proteotypic
DHX9_HUMAN	397	1022.434	MGGEEAEIR	Dioxidation (M)	1	Proteotypic

DHX9_HUMAN	588	1074.546	LAQFEPSQR		1	Proteotypic
DHX9_HUMAN	634	1086.615	RISAVSVAER		1	Proteotypic
DHX9_HUMAN	1367	1291.669	AAMEALVVEVTK	Dioxidation (M)	1	Proteotypic
DLDH_HUMAN	44	911.530	GRIPVNTR		1	Proteotypic
DLDH_HUMAN	156	951.433	GIEMSEVR	Dioxidation (M)	1	Proteotypic
DLDH_HUMAN	704	1103.546	ADGGTQVIDTK		1	Proteotypic
DLDH_HUMAN	1893	1566.825	NLGLLEELGIELDPR		1	Proteotypic
DSRAD_HUMAN	134	943.472	QEAADAALR		1	Proteotypic
DSRAD_HUMAN	893	1149.595	AMTILLEEAK	Dioxidation (M)	1	Proteotypic
DUS3_HUMAN	2457	1289.662	AADFIDQALAQK		1	Proteotypic
DYHC1_HUMAN	491	1046.551	AATSPALFNR		1	Proteotypic
DYHC1_HUMAN	742	1113.639	SLLQALNEVK		1	Proteotypic
DYHC1_HUMAN	774	1120.518	ANEVEQMIR	Dioxidation (M)	1	Proteotypic
DYHC1_HUMAN	1200	1236.642	LFLTMEINPK	Dioxidation (M)	1	Proteotypic
ECHA_HUMAN	169	955.468	FGELVMTK	Dioxidation (M)	1	Proteotypic
ECHA_HUMAN	226	974.503	DATLTALDR		1	Proteotypic
ECHA_HUMAN	309	997.544	ILQEGVDPK		1	Proteotypic
ECHA_HUMAN	761	1117.598	DTSASAVAVGLK		1	Proteotypic
ECHA_HUMAN	1520	1349.711	MQLLEIITTEK	Dioxidation (M)	1	Proteotypic
ECHA_HUMAN	1636	1410.626	SEVSSDEDIQFR		1	Proteotypic
ECHA_HUMAN	1875	1552.729	DLNSDMDSILASLK	Dioxidation (M)	1	Proteotypic
ECHA_HUMAN	1937	1608.782	FGGGNPELLTQMVSQK	Dioxidation (M)	1	Proteotypic
ECHA_HUMAN	1999	1656.752	MVGVPAALDMMLTGR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	1	Proteotypic
ECHA_HUMAN	2060	1710.839	ADMVIEAVFEDLSLK	Dioxidation (M)	1	Proteotypic
ECHA_HUMAN	2259	2006.116	TVLGTPEVLLGALPGAGGTQR		1	Proteotypic
ECHB_HUMAN	26	904.513	LAAFAVSR		1	Proteotypic
ECHB_HUMAN	170	955.545	NVVVVDGVR		1	Proteotypic
ECHB_HUMAN	337	1005.553	KLMLDLNK	Dioxidation (M)	1	Proteotypic
ECHB_HUMAN	542	1064.514	EAALGAGFSQK		1	Proteotypic
ECHB_HUMAN	838	1135.551	LEQDEYALR		1	Proteotypic
ECHB_HUMAN	1108	1212.639	TPFLLSGTSYK		1	Proteotypic
ECHB_HUMAN	1647	1415.766	DQLLLGPTYATPK		1	Proteotypic
ECHM_HUMAN	1595	1385.617	SLAMEMVLTGDR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
ECHM_HUMAN	1746	1465.836	NNTVGLIQLNRPK		1	Proteotypic
ECHM_HUMAN	2296	2124.133	AQFAQPEILIGTIPGAGGTQR		1	Proteotypic
EF1A1_HUMAN	57	913.560	QTVAVGVK		1	Proteotypic
EF1A1_HUMAN	228	974.544	LPLQDVYK		1	Proteotypic
EF1A1_HUMAN	406	1024.603	IGGIGTVPVGR		1	Proteotypic
EF1A1_HUMAN	773	1119.592	STTTGHLYK		1	Proteotypic
EF1A1_HUMAN	2358	2546.367	VETGVLKPGMVVTFAPVNVVTEVK	Dioxidation (M)	1	Proteotypic
EF1A1_HUMAN	2376	2931.384	NMITGTSQADCAVLIVAAGVGEFEAGISK	Dioxidation (M); Trioxidation (C)	1	Proteotypic
EF1D_HUMAN	565	1071.535	ATNFLAHEK	Acetyl (Protein N-term)	1	Proteotypic
EF1D_HUMAN	630	1085.583	QENGASVILR		1	Proteotypic
EF1D_HUMAN	1385	1298.719	GVVQELQQAIK		1	Proteotypic
EF1D_HUMAN	1540	1357.720	IASLEVENQSLR		1	Proteotypic
EF1G_HUMAN	762	1117.675	ILGLLDAYLK		1	Proteotypic

EF1G_HUMAN	1211	1240.645	STFVLDEFKR		1	Proteotypic
EF1G_HUMAN	1513	1346.731	ALIAAQYSGAQVR		1	Proteotypic
EF1G_HUMAN	1903	1571.815	KLDPGSEETQTLVR		1	Proteotypic
EF2_HUMAN	205	968.540	GGGQIPTAR		1	Proteotypic
EF2_HUMAN	357	1012.494	GEGQLGPAER		1	Proteotypic
EF2_HUMAN	563	1071.494	KVEDMMKK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
EF2_HUMAN	654	1090.577	VNFTVDQIR		1	Proteotypic
EF2_HUMAN	721	1106.634	VFSGLVSTGLK		1	Proteotypic
EF2_HUMAN	752	1115.532	VFDAIMNFK	Dioxidation (M)	1	Proteotypic
EF2_HUMAN	1168	1230.449	EGALCEENMR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
EF2_HUMAN	1704	1443.761	EGIPALDNFLDKL		1	Proteotypic
EF2_HUMAN	1780	1484.758	TFCQLILDPIFK	Trioxidation (C)	1	Proteotypic
EF2_HUMAN	1913	1584.719	ETVSEESNVLCLSK	Trioxidation (C)	1	Proteotypic
EFTU_HUMAN	435	1031.561	GTVVTGTLER		1	Proteotypic
EFTU_HUMAN	1001	1184.615	AEAGDNLGALVR		1	Proteotypic
EFTU_HUMAN	1863	1541.845	LLDAVDTYIPVPAR		1	Proteotypic
ELAV1_HUMAN	327	1002.488	FAANPNQNK		1	Proteotypic
ELAV1_HUMAN	1019	1187.651	VLVDQTTGLSR		1	Proteotypic
ELAV1_HUMAN	1528	1352.682	SLFSSIGEVESAK		1	Proteotypic
ENOA_HUMAN	3	898.549	TIAPALVSK		1	Proteotypic
ENOA_HUMAN	23	903.455	IEEELGSK		1	Proteotypic
ENOA_HUMAN	642	1087.587	KLNVTEQEK		1	Proteotypic
ENOA_HUMAN	860	1142.608	IGAEVYHNLK		1	Proteotypic
ENOA_HUMAN	1508	1343.558	LMIEMDGTENK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
ENOA_HUMAN	1627	1405.709	GNPTVEVDLFTSK		1	Proteotypic
ENOA_HUMAN	1663	1424.719	YISPDQLADLYK		1	Proteotypic
ENOA_HUMAN	1881	1555.770	VVIGMDVAASEFFR	Oxidation (M)	1	Proteotypic
ENOA_HUMAN	1902	1571.765	VVIGMDVAASEFFR	Dioxidation (M)	1	Proteotypic
ENOA_HUMAN	1957	1623.777	VNQIGSVTESLQACK	Trioxidation (C)	1	Proteotypic
ENOA_HUMAN	2053	1699.764	IDKLMIEMDGTENK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
ENOA_HUMAN	2132	1803.937	AAVPSGASTGIYEALER		2	Shared
ENOA_HUMAN	2241	1959.917	DATNVGDEGGFAPNILENK		1	Proteotypic
ENOA_HUMAN	2245	1970.959	LAMQEFMILPVGAANFR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
ENOA_HUMAN	2271	2032.048	FTASAGIQVVGDDTLVTNPK		1	Proteotypic
ENOA_HUMAN	2370	2742.371	DATNVGDEGGFAPNILENKEGLELLK		1	Proteotypic
ENOB_HUMAN	1956	1623.777	VNQIGSVTESIQACK	Trioxidation (C)	1	Proteotypic
ENOB_HUMAN	2132	1803.937	AAVPSGASTGIYEALER		2	Shared
ENPL_HUMAN	76	918.529	NKEIFLR		1	Proteotypic
ENPL_HUMAN	185	960.451	TDDEVVQR		1	Proteotypic
ENPL_HUMAN	190	962.580	LIINSLYK		1	Proteotypic
ENPL_HUMAN	247	981.477	SGTSEFLNK		1	Proteotypic
ENPL_HUMAN	532	1062.480	IYFMAGSSR	Dioxidation (M)	1	Proteotypic
ENPL_HUMAN	612	1080.535	FAFQAEVNR		1	Proteotypic
ENPL_HUMAN	621	1084.576	LSLNIDPDAK		1	Proteotypic
ENPL_HUMAN	846	1138.573	LGVIEDHSNR		1	Proteotypic
ENPL_HUMAN	889	1149.530	EAESSPFVER		1	Proteotypic

ENPL_HUMAN	948	1166.546	EGVKFDESEK		1	Proteotypic
ENPL_HUMAN	1016	1186.671	SILFVPTSAPR		1	Proteotypic
ENPL_HUMAN	1318	1274.635	ELISNASDALDK		2	Shared
ENPL_HUMAN	1326	1277.625	KEAESSPFVER		1	Proteotypic
ENPL_HUMAN	1354	1288.594	DISTNYYASQK		1	Proteotypic
ENPL_HUMAN	1779	1484.747	GVVDSDDLPLNVS		1	Proteotypic
ENPL_HUMAN	1833	1524.719	EEASYLELDTIK		1	Proteotypic
ENPL_HUMAN	1867	1543.821	ELISNASDALDKIR		2	Shared
ENPL_HUMAN	1868	1544.762	NLLHVTDGTGVMTR	Dioxidation (M)	1	Proteotypic
ENPL_HUMAN	2143	1814.797	RVFITDDFHDMMPK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
ENPL_HUMAN	2276	2045.053	LISLTDENALSGNEELTVK		1	Proteotypic
ENPL_HUMAN	2369	2727.331	TDDEVVQREEEAIQLDGLNASQIR		1	Proteotypic
ERP29_HUMAN	832	1133.644	SLNILTAFQK		1	Proteotypic
ERP29_HUMAN	2082	1739.767	ILDQGEDFPASEMTR	Dioxidation (M)	1	Proteotypic
ESYT1_HUMAN	847	1138.671	LLAETVAPAVR		1	Proteotypic
ESYT1_HUMAN	1421	1308.813	ALTLGALTLPLAR		1	Proteotypic
ETFA_HUMAN	573	1072.551	SDRPELTGAK		1	Proteotypic
ETFA_HUMAN	1586	1380.823	LEVAPISDIIAK		1	Proteotypic
ETFA_HUMAN	1962	1629.712	GTSFDAAATSGGSASSEK		1	Proteotypic
ETFA_HUMAN	2078	1735.013	GLLPEELTPLILATQK		1	Proteotypic
EZRI_HUMAN	56	913.523	IALLEEAR		1	Proteotypic
EZRI_HUMAN	232	975.539	QLFDQVVK		2	Shared
EZRI_HUMAN	603	1078.533	EKEELMLR	Dioxidation (M)	1	Proteotypic
EZRI_HUMAN	636	1086.680	FVIKPIDKK		2	Shared
EZRI_HUMAN	992	1181.587	APDFVIFYAPR		2	Shared
EZRI_HUMAN	1706	1444.789	QLLTLSELSQAR		1	Proteotypic
EZRI_HUMAN	1805	1503.772	RKPDTIEVQQMK	Dioxidation (M)	2	Shared
F16P1_HUMAN	2056	1704.796	ADQAPFDTDVNTLTR	Acetyl (Protein N-term)	1	Proteotypic
F16P1_HUMAN	2061	1711.940	APVILGSPDDVLEFLK		1	Proteotypic
FAHD1_HUMAN	2489	1711.961	IITLEEGDIILGTGPK		1	Proteotypic
FIS1_HUMAN	1148	1222.754	GIVLLELLPK		1	Proteotypic
FIS1_HUMAN	1692	1439.737	GLLQTEPQNNQAK		1	Proteotypic
FKB1A_HUMAN	2462	1313.658	GVQVETISPGDGR		1	Proteotypic
FLNA_HUMAN	160	951.466	AEFTVETR		1	Proteotypic
FLNA_HUMAN	272	987.535	KGEITGEVR		1	Proteotypic
FLNA_HUMAN	611	1080.527	LSPFMADIR	Dioxidation (M)	1	Proteotypic
FLNA_HUMAN	681	1098.567	GTVEPQLEAR		1	Proteotypic
FLNA_HUMAN	949	1166.593	VPVHDVTDASK		1	Proteotypic
FLNA_HUMAN	1149	1224.562	EATTEFSVDAR		1	Proteotypic
FLNA_HUMAN	1157	1225.765	LIALLEVLSQK		2	Shared
FLNA_HUMAN	1555	1363.698	VDVGKDQEFTVK		1	Proteotypic
FLNA_HUMAN	1643	1414.742	IANLQTDLSDGLR		1	Proteotypic
FLNA_HUMAN	1672	1428.711	AFGPGLQGGGAGSPAR		1	Proteotypic
FLNA_HUMAN	1681	1433.715	ANLPQSFQVDTSK		1	Proteotypic
FLNA_HUMAN	1898	1569.800	GAGTGGLGLAVEGPSEAK		1	Proteotypic
FLNA_HUMAN	1990	1651.853	VTAQGPGLPSGNIANK		1	Proteotypic

FLNB_HUMAN	91	924.539	AGLAPLEVR		1	Proteotypic
FLNB_HUMAN	249	982.556	IAGPGLGSGVR		1	Proteotypic
FLNB_HUMAN	1099	1209.624	VLQSFTVDSSK		1	Proteotypic
FLNB_HUMAN	1157	1225.765	LIALLEVLSQK		2	Shared
FLNB_HUMAN	1967	1634.761	SPFEVQVGPEAGMQK	Dioxidation (M)	1	Proteotypic
FLOT2_HUMAN	782	1122.556	SAFSEEVNIK		1	Proteotypic
FLOT2_HUMAN	1057	1198.656	QVLLAQAEAEK		1	Proteotypic
FLOT2_HUMAN	1802	1498.843	MALVLEALPQIAAK	Dioxidation (M)	1	Proteotypic
FPPS_HUMAN	2407	1059.531	KQDADSLQR		1	Proteotypic
FUBP2_HUMAN	605	1078.614	IGGGIDVPVPR		1	Proteotypic
FUBP2_HUMAN	857	1142.495	MMLDDIVSR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
FUBP2_HUMAN	996	1183.692	IINDLLQSLR		1	Proteotypic
FUBP2_HUMAN	1232	1244.552	QLEDGDQPESK		1	Proteotypic
FUBP2_HUMAN	1396	1301.683	SVSLTGAPESVQK		1	Proteotypic
FUMH_HUMAN	2447	1235.607	AIEMLGGELGSK	Dioxidation (M)	1	Proteotypic
G3P_HUMAN	36	908.483	AGAHLQGGAK		1	Proteotypic
G3P_HUMAN	545	1064.584	AGAHLQGGAKR		1	Proteotypic
G3P_HUMAN	1287	1264.579	VVDLMAHMASK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
G3P_HUMAN	1607	1393.622	VVDLMAHMASKE	Dioxidation (M); Dioxidation (M)	1	Proteotypic
G3P_HUMAN	1637	1410.783	GALQNIIPASTGAAK		1	Proteotypic
G3P_HUMAN	1821	1520.750	VPTANVSVVDLTCR	Trioxidation (C)	1	Proteotypic
G3P_HUMAN	1939	1612.894	LIVINGNPITIFQER		1	Proteotypic
G3P_HUMAN	2145	1814.839	IISNASCTTNCLAPLAK	Trioxidation (C); Trioxidation (C)	1	Proteotypic
G3P_HUMAN	2273	2040.100	LIVINGNPITIFQERDPSK		1	Proteotypic
G3P_HUMAN	2317	2187.157	VPTANVSVVDLTCRLEKPAK	Trioxidation (C)	1	Proteotypic
G3P_HUMAN	2327	2276.082	VIISAPSADAPMFVMGVNHEK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
G3P_HUMAN	2363	2626.342	VIHDNFGIVEGLMTTVHAIATQK	Dioxidation (M)	1	Proteotypic
G6PI_HUMAN	187	962.438	DVMPEVNK	Dioxidation (M)	1	Proteotypic
G6PI_HUMAN	723	1107.523	NLVTEDVMR	Dioxidation (M)	1	Proteotypic
G6PI_HUMAN	1125	1217.640	HFVALSTNTTK		1	Proteotypic
G6PI_HUMAN	1997	1654.805	TFTTQETITNAETAK		1	Proteotypic
G6PI_HUMAN	2077	1734.934	ILLANFLAQTEALMR	Dioxidation (M)	1	Proteotypic
G6PI_HUMAN	2160	1831.009	TLAQLNPESSLFIASK		1	Proteotypic
GANAB_HUMAN	94	925.560	PAAVVLQTK		1	Proteotypic
GANAB_HUMAN	213	971.576	SLLLSVNR		1	Proteotypic
GANAB_HUMAN	339	1006.472	VTEGGEPYR		1	Proteotypic
GANAB_HUMAN	550	1065.582	NPEPELLVR		1	Proteotypic
GANAB_HUMAN	795	1127.572	GLLEFEHQR		1	Proteotypic
GANAB_HUMAN	968	1173.614	FRIDELEPR		1	Proteotypic
GANAB_HUMAN	1526	1351.844	VLLVLELQGLQK		1	Proteotypic
GANAB_HUMAN	2014	1663.040	VVIIGAGKPAAVVLQTK		1	Proteotypic
GANAB_HUMAN	2309	2153.983	VSQGSKDPAEGDGAQPEETPR		1	Proteotypic
GBLP_HUMAN	158	951.433	TEQMTLR	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
GBLP_HUMAN	1417	1308.631	DVLSVAFSSDNR		1	Proteotypic
GBLP_HUMAN	2121	1787.988	IIVDELKQEVISTSSK		1	Proteotypic
GDIB_HUMAN	917	1156.580	MLLYTEVTR	Dioxidation (M)	1	Proteotypic



GDIB_HUMAN	1124	1216.707	FVSISDLLVPK		1	Proteotypic
GDIB_HUMAN	1557	1364.694	DLGTESQIFISR		1	Proteotypic
GDIB_HUMAN	2262	2010.982	VPSTEEALASSLMGLFEK	Dioxidation (M)	1	Proteotypic
GFAP_HUMAN	25	904.425	GTNESLER		2	Shared
GFAP_HUMAN	29	905.461	FLEQQNK		14	Shared
GFAP_HUMAN	1324	1276.703	LALDIEIATYR		4	Shared
GFAP_HUMAN	1625	1404.798	LALDIEIATYRK		4	Shared
GLRX3_HUMAN	2499	2557.266	AAGAAEAAVAEEVGSAGQFEELLR	Acetyl (Protein N-term)	1	Proteotypic
GRP75_HUMAN	175	957.488	VLENAEGAR		1	Proteotypic
GRP75_HUMAN	877	1147.547	KDSETGENIR		1	Proteotypic
GRP75_HUMAN	1171	1230.657	QAASSLQQASLK		1	Proteotypic
GRP75_HUMAN	1218	1241.673	DAGQISGLNVLR		1	Proteotypic
GRP75_HUMAN	1361	1289.673	VQQTVQDLFGR		1	Proteotypic
GRP75_HUMAN	1552	1360.735	AQFEGIVTDLIR		1	Proteotypic
GRP75_HUMAN	1717	1449.710	TTPSVVAFTADGER		1	Proteotypic
GRP75_HUMAN	1770	1477.745	SDIGEVILVGGMTR	Dioxidation (M)	1	Proteotypic
GRP75_HUMAN	2280	2054.955	STNGDTFLGGEDFDQALLR		1	Proteotypic
GRP78_HUMAN	113	934.458	VMEHFIK	Dioxidation (M)	1	Proteotypic
GRP78_HUMAN	262	985.508	LTPEEIER		1	Proteotypic
GRP78_HUMAN	306	996.510	ALSSQHQR		1	Proteotypic
GRP78_HUMAN	484	1045.566	VLEDSLK		1	Proteotypic
GRP78_HUMAN	585	1073.547	ITITNDQNR		1	Proteotypic
GRP78_HUMAN	764	1118.557	VTAEDKGTGNK		1	Proteotypic
GRP78_HUMAN	1032	1190.630	VYEGERPLTK		1	Proteotypic
GRP78_HUMAN	1098	1209.578	EFFNGKEPSR		1	Proteotypic
GRP78_HUMAN	1162	1227.621	VEIANDQGNR		4	Shared
GRP78_HUMAN	1242	1248.613	DAGTIAGLNVMR	Dioxidation (M)	1	Proteotypic
GRP78_HUMAN	1438	1315.630	NELESYAYSLK		1	Proteotypic
GRP78_HUMAN	1613	1396.781	ELEEIVQPIISK		1	Proteotypic
GRP78_HUMAN	1731	1459.752	SDIDEIVLVGGSTR		1	Proteotypic
GRP78_HUMAN	1866	1543.734	AKFEELNMDLFR	Dioxidation (M)	1	Proteotypic
GRP78_HUMAN	1891	1565.773	ITPSYVAFTPEGER		1	Proteotypic
GRP78_HUMAN	1895	1567.780	TFAPEEISAMVLTK	Dioxidation (M)	1	Proteotypic
GRP78_HUMAN	1919	1587.847	KSDIDEIVLVGGSTR		1	Proteotypic
GRP78_HUMAN	1994	1652.971	KKELEEIVQPIISK		1	Proteotypic
GRP78_HUMAN	2006	1658.888	IINEPTAAAIAAYGLDK		3	Shared
GRP78_HUMAN	2028	1676.801	NQLTSNPENTVFDKAK		1	Proteotypic
GRP78_HUMAN	2147	1814.989	IINEPTAAAIAAYGLDKR		1	Proteotypic
GRP78_HUMAN	2162	1832.902	NQLTSNPENTVFDKAKR		1	Proteotypic
GRP78_HUMAN	2165	1835.927	SQIFSTASDNQPTVTIK		1	Proteotypic
GRP78_HUMAN	2225	1933.006	DNHLLGTFDLTGIPPAPR		1	Proteotypic
GRP78_HUMAN	2312	2163.985	IEIESFYEGEDFSETLTR		1	Proteotypic
GSHR_HUMAN	813	1130.458	MVCANKEEK	Dioxidation (M); Trioxidation (C)	1	Proteotypic
GSHR_HUMAN	902	1151.703	ALLTPVAIAAGR		1	Proteotypic
GSTK1_HUMAN	1739	1463.693	AGMSAEQAQGLLEK	Dioxidation (M)	1	Proteotypic
GSTK1_HUMAN	2072	1727.858	NEDITEPQSILAAAEK		1	Proteotypic

GSTO1_HUMAN	2413	1073.623	VPSLVGSFIR		1	Proteotypic
GSTP1_HUMAN	2193	1882.942	FQDGDLTLYQSNTILR		1	Proteotypic
GSTP1_HUMAN	2298	2125.153	ALPGQLKPFETLLSQNQGGK		1	Proteotypic
GSTP1_HUMAN	2307	2147.964	DQQEAAALVDMVNDGVEDLR	Dioxidation (M)	1	Proteotypic
GSTP1_HUMAN	2348	2427.053	DQQEAAALVDMVNDGVEDLRCK	Dioxidation (M); Trioxidation (C)	1	Proteotypic
H10_HUMAN	524	1059.520	VGENADSQIK		1	Proteotypic
H10_HUMAN	1506	1342.673	TENSTSAPAAKPK	Acetyl (Protein N-term)	1	Proteotypic
H2A1A_HUMAN	136	943.524	AGLQFPVGR		3	Shared
H2A1A_HUMAN	2325	2270.373	LLGGVTIAQGGVLPNIQAVLLPK		1	Proteotypic
H2A1B_HUMAN	136	943.524	AGLQFPVGR		3	Shared
H2A1B_HUMAN	1389	1299.678	NDEELNKLGR		1	Proteotypic
H2A1B_HUMAN	2221	1930.162	VTIAQGGVLPNIQAVLLPK		2	Shared
H2AW_HUMAN	65	914.534	AGVIFPVGR		2	Shared
H2AW_HUMAN	791	1125.651	GVTIASGGVLP		1	Proteotypic
H2AY_HUMAN	65	914.534	AGVIFPVGR		2	Shared
H2AY_HUMAN	259	984.597	QTAAQLILK		1	Proteotypic
H2AY_HUMAN	653	1090.577	SIAFPSIGSGR		1	Proteotypic
H2AY_HUMAN	1081	1203.650	EFVEAVLELR		1	Proteotypic
HBA_HUMAN	631	1086.542	MFLSFPTTK	Oxidation (M)	1	Proteotypic
HBA_HUMAN	635	1086.619	LRVDPVNFK		1	Proteotypic
HBA_HUMAN	698	1102.537	MFLSFPTTK	Dioxidation (M)	1	Proteotypic
HBA_HUMAN	959	1170.661	VLSPADKTNVK		1	Proteotypic
HBA_HUMAN	1250	1251.707	FLASVSTVLTSK		1	Proteotypic
HBA_HUMAN	1839	1528.727	VGAHAGEYGAEALER		1	Proteotypic
HCD2_HUMAN	1122	1215.665	GGIVGMLPIAR	Dioxidation (M)	1	Proteotypic
HCD2_HUMAN	2142	1812.011	GLVAVITGGASGLGLATAER		1	Proteotypic
HCD2_HUMAN	2149	1816.801	LVAGEMGQNEPDQGGQR	Dioxidation (M)	1	Proteotypic
HINT1_HUMAN	157	951.433	CAADLGLNK	Trioxidation (C)	1	Proteotypic
HINT1_HUMAN	1646	1415.752	AQVARPGGDTIFGK		1	Proteotypic
HNMT_HUMAN	24	903.466	TSNLENVK		1	Proteotypic
HNMT_HUMAN	282	989.539	IGDTKSEIK		1	Proteotypic
HNMT_HUMAN	1908	1578.866	VLFNNTLSFIVIEA		1	Proteotypic
HNMT_HUMAN	2009	1660.831	AELGKDLQEPEFSAK		1	Proteotypic
HNMT_HUMAN	2115	1783.009	ILSIGGGAGEIDLQILSK		1	Proteotypic
HNRPF_HUMAN	2498	1866.936	ITGEAFVQFASQELAEK		1	Proteotypic
HNRPG_HUMAN	50	912.466	VEADRPVK	Acetyl (Protein N-term)	1	Proteotypic
HNRPG_HUMAN	234	975.567	IVEVLLMK	Dioxidation (M)	1	Proteotypic
HNRPG_HUMAN	1683	1434.710	VEQATKPSFESGR		1	Proteotypic
HNRPG_HUMAN	1684	1434.735	LFIGGLNTEETNEK		1	Proteotypic
HNRPG_HUMAN	1782	1485.714	GFAFVTFESPADAK		1	Proteotypic
HNRPK_HUMAN	110	933.477	ENTQTTIK		1	Proteotypic
HNRPK_HUMAN	415	1028.423	GGDLMAYDR	Dioxidation (M)	1	Proteotypic
HNRPK_HUMAN	501	1052.634	VVLIGGKPR		1	Proteotypic
HNRPK_HUMAN	845	1137.497	NTDEMVELR	Dioxidation (M)	1	Proteotypic
HNRPK_HUMAN	1043	1193.692	NLPLPPPPPR		1	Proteotypic
HNRPK_HUMAN	1267	1258.568	IDEPLEGSER		1	Proteotypic

HNRPK_HUMAN	1498	1339.796	IILDLISESPIK		1	Proteotypic
HNRPK_HUMAN	2102	1766.789	RPAEDMEEEEQAFKR	Dioxidation (M)	1	Proteotypic
HNRPK_HUMAN	2111	1779.791	TDYNASVSPDSSGPER		1	Proteotypic
HNRPK_HUMAN	2362	2588.381	IITITGTQDQIQNAQYLLQNSVK		1	Proteotypic
HNRPM_HUMAN	35	908.402	MGANSLER	Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	117	935.413	MGANNLER	Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	388	1019.380	FGSGMNMGR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	533	1062.498	FESPEVAER		1	Proteotypic
HNRPM_HUMAN	699	1102.551	ADILEDKDGK		1	Proteotypic
HNRPM_HUMAN	741	1113.639	INEILSNALK		1	Proteotypic
HNRPM_HUMAN	940	1164.465	MGAGLGHGMDR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	1022	1188.490	MGAGMGFGLER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	1111	1213.569	GGERPAQNEK		1	Proteotypic
HNRPM_HUMAN	1151	1224.617	EVFSMAGVVVR	Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	1247	1251.559	MVPAGMGAGLER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	1248	1251.584	VKEDPDGEHAR		1	Proteotypic
HNRPM_HUMAN	1285	1263.686	AFITNIPFDVK		1	Proteotypic
HNRPM_HUMAN	1337	1283.622	QGGGGGGGSPGIER		1	Proteotypic
HNRPM_HUMAN	1710	1446.587	MGLAMGGGGGASFDR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	1791	1490.686	MGPAMGALGAGIER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	1859	1541.794	AAGVEAAAEVAATEIK	Acetyl (Protein N-term)	1	Proteotypic
HNRPM_HUMAN	2027	1676.750	MGPLGLDHMASSIER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	2116	1783.855	VGEVTVYVLLMDAEGK	Dioxidation (M)	1	Proteotypic
HNRPM_HUMAN	2166	1841.632	MGGMEGPFGGGMENMGR	Dioxidation (M); Dioxidation (M); Dioxidation (M); Dioxidation (M)	1	Proteotypic
HNRPQ_HUMAN	1756	1472.772	NLANTVTEEILEK		1	Proteotypic
HNRPQ_HUMAN	1925	1592.797	DLFEDELVPLFEK		1	Proteotypic
HNRPR_HUMAN	2473	1459.777	NLATTVTEEILEK		1	Proteotypic
HNRPU_HUMAN	304	995.504	DIDIHEVR		1	Proteotypic
HNRPU_HUMAN	494	1047.535	NGQDLGVAFK		1	Proteotypic
HNRPU_HUMAN	586	1073.597	VSELKEELK		1	Proteotypic
HNRPU_HUMAN	1073	1201.692	VSELKEELKK		1	Proteotypic
HNRPU_HUMAN	1332	1279.569	KMCLFAGFQR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
HNRPU_HUMAN	1981	1646.838	NFIELDQTNVSAQAQR		1	Proteotypic
HNRPU_HUMAN	2383	3157.422	LQAALDDEEAGGRPAMEPNGSLDLGGDSAGR	Dioxidation (M)	1	Proteotypic
HS90B_HUMAN	153	950.457	ADHGEPIGR		1	Proteotypic
HS90B_HUMAN	897	1150.551	YIDQEELNK		2	Shared
HS90B_HUMAN	1042	1193.640	IDIIPNPQER		1	Proteotypic
HS90B_HUMAN	1196	1235.630	RAPFDLFENK		1	Proteotypic
HS90B_HUMAN	1220	1241.698	ADLNNLGTIAK		2	Shared
HS90B_HUMAN	1241	1248.610	EQVANSAFVER		1	Proteotypic
HS90B_HUMAN	1318	1274.635	ELISNASDALDK		2	Shared
HS90B_HUMAN	1424	1310.563	EDQTEYLEER		2	Shared
HS90B_HUMAN	1585	1380.717	TLTLVDTGIGMTK	Dioxidation (M)	1	Proteotypic
HS90B_HUMAN	1809	1512.778	GVVDSEDLPLNISR		2	Shared
HS90B_HUMAN	1867	1543.821	ELISNASDALDKIR		2	Shared
HSP74_HUMAN	881	1148.546	FQSEERPDK		1	Proteotypic

HSP74_HUMAN	1456	1320.704	VLATAFDTTLGGR		1	Proteotypic
HSP76_HUMAN	379	1016.561	ITITNDKGR		4	Shared
HSP76_HUMAN	613	1080.560	LLQDFFNKG		2	Shared
HSP76_HUMAN	1163	1227.621	VEILANDQGNR		1	Proteotypic
HSP76_HUMAN	1784	1486.694	TTPSYVAFTDTER		4	Shared
HSP76_HUMAN	1846	1532.693	ARFEELCSDLFR	Trioxidation (C)	2	Shared
HSP76_HUMAN	2023	1674.723	ATAGDTHLGGEDFDNR		3	Shared
HSP76_HUMAN	2038	1686.894	IINEPTAAAIAYGLDR		2	Shared
HSP7C_HUMAN	275	988.519	LSKEDIER		1	Proteotypic
HSP7C_HUMAN	379	1016.561	ITITNDKGR		4	Shared
HSP7C_HUMAN	613	1080.560	LLQDFFNKG		2	Shared
HSP7C_HUMAN	990	1179.614	VQVEYKGETK		1	Proteotypic
HSP7C_HUMAN	1059	1198.667	DAGTIAGLNVLR		1	Proteotypic
HSP7C_HUMAN	1162	1227.621	VEIANDQGNR		4	Shared
HSP7C_HUMAN	1252	1252.609	FEELNADLFR		1	Proteotypic
HSP7C_HUMAN	1290	1266.607	MVNHFIAEFK	Dioxidation (M)	1	Proteotypic
HSP7C_HUMAN	1338	1283.643	MKEIAEAYLGK	Dioxidation (M)	1	Proteotypic
HSP7C_HUMAN	1485	1334.581	NSLESYAFNMK	Dioxidation (M)	1	Proteotypic
HSP7C_HUMAN	1699	1441.651	RFDDAVVQSDMK	Dioxidation (M)	1	Proteotypic
HSP7C_HUMAN	1775	1479.747	ARFEELNADLFR		1	Proteotypic
HSP7C_HUMAN	1777	1480.800	SQIHDIIVLVGGSTR		1	Proteotypic
HSP7C_HUMAN	1784	1486.694	TTPSYVAFTDTER		4	Shared
HSP7C_HUMAN	1983	1647.770	SFYPEEVSSMVLTK	Dioxidation (M)	1	Proteotypic
HSP7C_HUMAN	2006	1658.888	IINEPTAAAIAYGLDK		3	Shared
HSP7C_HUMAN	2031	1680.778	NQVAMNPTNTVFDK	Dioxidation (M)	1	Proteotypic
HSP7C_HUMAN	2043	1690.718	STAGDTHLGGEDFDNR		1	Proteotypic
HSPB1_HUMAN	74	916.487	DGVVEITGK		1	Proteotypic
HSPB1_HUMAN	126	940.498	AQLGGPEAAK		1	Proteotypic
HSPB1_HUMAN	590	1074.567	QLSSGVSEIR		1	Proteotypic
HSPB1_HUMAN	703	1103.500	QDEHGYISR		1	Proteotypic
HSPB1_HUMAN	872	1145.629	TKDGVVEITGK		1	Proteotypic
HSPB1_HUMAN	935	1162.613	LFDQAFGLPR		1	Proteotypic
HSPB1_HUMAN	2114	1782.915	VSLDVNHFAPDELTVK		1	Proteotypic
HSPB1_HUMAN	2205	1904.984	LATQSNEITIPVTFESR		1	Proteotypic
IDHC_HUMAN	233	975.550	NILGGTVFR		2	Shared
IDHC_HUMAN	632	1086.571	ATDFVVPGPVK		1	Proteotypic
IDHC_HUMAN	1377	1297.589	LIDDMVAQAMK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
IDHC_HUMAN	1689	1437.699	VEITYTPSDGTQK		1	Proteotypic
IDHC_HUMAN	1856	1540.639	SDYLNTFEFMDK	Dioxidation (M)	1	Proteotypic
IDHC_HUMAN	2096	1758.740	ISGGSVVEMQGDDEMTR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
IDHP_HUMAN	233	975.550	NILGGTVFR		2	Shared
IDHP_HUMAN	1166	1227.657	GKLDGNQDLIR		1	Proteotypic
IDHP_HUMAN	1321	1275.674	LIDDMVAQVLK	Dioxidation (M)	1	Proteotypic
IDHP_HUMAN	2083	1739.771	VAKPVVEMDGDDEMTR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
IF2A_HUMAN	2463	1347.652	VVTDTDETELAR		1	Proteotypic
IF4A3_HUMAN	1104	1211.687	ETQALILAPTR		1	Proteotypic

IF4A3_HUMAN	1381	1298.577	ATTATMATSGSAR	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
IGHA1_HUMAN	106	930.539	TPLTATLSK		2	Shared
IGHA1_HUMAN	125	939.477	SAVQGPPER		1	Proteotypic
IGHA1_HUMAN	2164	1834.943	QEPSQGTTFVAVTSILR		2	Shared
ILF2_HUMAN	459	1040.634	VLQSALAAIR		1	Proteotypic
ILF2_HUMAN	1198	1235.760	ILITTVPPNLR		1	Proteotypic
ILF2_HUMAN	1632	1408.829	ILPTLEAVAALGNK		1	Proteotypic
IMB1_HUMAN	168	954.586	VAALQNLVK		1	Proteotypic
IMB1_HUMAN	936	1162.615	MELITILEK	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
IMB1_HUMAN	1152	1224.657	VLANPGNSQVAR		1	Proteotypic
IMB1_HUMAN	2004	1657.904	AAVENLPTFLVELSR		1	Proteotypic
IMMT_HUMAN	199	965.449	VAMIDETR	Dioxidation (M)	1	Proteotypic
IMMT_HUMAN	522	1058.572	KVQAAQSEAK		1	Proteotypic
IMMT_HUMAN	748	1114.587	AVDEAADALLK		1	Proteotypic
IMMT_HUMAN	904	1152.589	ERPPEEVAAR		1	Proteotypic
K1C10_HUMAN	292	992.493	YENEVALR		1	Proteotypic
K1C10_HUMAN	432	1030.591	VLDELTLTK		1	Proteotypic
K1C10_HUMAN	708	1105.519	VTMQNLNDR	Oxidation (M)	3	Shared
K1C10_HUMAN	777	1121.514	VTMQNLNDR	Dioxidation (M)	3	Shared
K1C10_HUMAN	944	1164.577	LENEIQTYR		1	Proteotypic
K1C10_HUMAN	998	1184.516	SSSSGSGVGESSSK		1	Proteotypic
K1C10_HUMAN	1066	1200.610	QSVEADINGLR		2	Shared
K1C10_HUMAN	1187	1233.672	LKYENEVALR		1	Proteotypic
K1C10_HUMAN	1279	1261.590	SLLEGEGSSGGGGR		1	Proteotypic
K1C10_HUMAN	1479	1331.578	NHEEEMKDLR	Dioxidation (M)	1	Proteotypic
K1C10_HUMAN	1536	1356.711	QSVEADINGLRR		2	Shared
K1C10_HUMAN	1556	1364.632	SQYEQLAEQNR		1	Proteotypic
K1C10_HUMAN	1583	1380.641	ALEESNYELEGK		1	Proteotypic
K1C10_HUMAN	1600	1389.674	QSLEASLAETEGR		1	Proteotypic
K1C10_HUMAN	1682	1433.763	IRLENEIQTYR		1	Proteotypic
K1C10_HUMAN	1797	1492.727	SQYEQLAEQNRK		1	Proteotypic
K1C10_HUMAN	1855	1539.633	SGGGGGGGCGGGGGVSSLR	Trioxidation (C)	1	Proteotypic
K1C10_HUMAN	2057	1706.765	GSLGGGFSSGGFSGGSFSR		1	Proteotypic
K1C10_HUMAN	2126	1796.004	NVQALEIELQSQLALK		1	Proteotypic
K1C10_HUMAN	2255	1995.964	ELTTEIDNIEQISSYK		1	Proteotypic
K1C10_HUMAN	2285	2072.921	AETECQNTEYQQLLDIK	Trioxidation (C)	1	Proteotypic
K1C10_HUMAN	2319	2211.091	SKELTTEIDNIEQISSYK		1	Proteotypic
K1C10_HUMAN	2322	2255.124	ADLEMQIESLTEELAYLKK	Dioxidation (M)	1	Proteotypic
K1C10_HUMAN	2343	2366.255	NQILNLTDDNANILLQIDNAR		1	Proteotypic
K1C10_HUMAN	2378	2935.365	NVSTGDVNVEMNAAPGVDLTQLLNMR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
K1C10_HUMAN	2381	3051.620	TIDDLKNQILNLTDDNANILLQIDNAR		1	Proteotypic
K1C14_HUMAN	420	1028.587	VLDELTLAR		5	Shared
K1C14_HUMAN	708	1105.519	VTMQNLNDR	Oxidation (M)	3	Shared
K1C14_HUMAN	777	1121.514	VTMQNLNDR	Dioxidation (M)	3	Shared
K1C14_HUMAN	779	1121.572	LEQEIATYR		6	Shared
K1C14_HUMAN	1003	1184.688	RVLDELTLAR		5	Shared

K1C14_HUMAN	1129	1219.587	MSVEADINGLR	Oxidation (M)	2	Shared
K1C14_HUMAN	1130	1219.593	ASLENSLEETK		2	Shared
K1C14_HUMAN	1195	1235.582	MSVEADINGLR	Dioxidation (M)	2	Shared
K1C14_HUMAN	1393	1300.651	ALEEANADLEVK		4	Shared
K1C14_HUMAN	1549	1360.683	EVATNSELVQSGK		2	Shared
K1C14_HUMAN	1602	1391.683	MSVEADINGLRR	Dioxidation (M)	2	Shared
K1C14_HUMAN	2291	2116.970	GQVGGDVNVEMDAAPGVLSR	Dioxidation (M)	1	Proteotypic
K1C15_HUMAN	420	1028.587	VLDELTLAR		5	Shared
K1C15_HUMAN	463	1041.582	VILEIDNAR		2	Shared
K1C15_HUMAN	771	1119.534	ITMQNLNDR	Oxidation (M)	2	Shared
K1C15_HUMAN	779	1121.572	LEQEIATYR		6	Shared
K1C15_HUMAN	836	1135.529	ITMQNLNDR	Dioxidation (M)	2	Shared
K1C15_HUMAN	1003	1184.688	RVLDELTLAR		5	Shared
K1C15_HUMAN	1393	1300.651	ALEEANADLEVK		4	Shared
K1C17_HUMAN	293	992.493	FETEQALR		2	Shared
K1C17_HUMAN	420	1028.587	VLDELTLAR		5	Shared
K1C17_HUMAN	779	1121.572	LEQEIATYR		6	Shared
K1C17_HUMAN	1003	1184.688	RVLDELTLAR		5	Shared
K1C17_HUMAN	1143	1221.635	TKFETEQALR		2	Shared
K1C17_HUMAN	1549	1360.683	EVATNSELVQSGK		2	Shared
K1C18_HUMAN	88	923.519	IREHLEK		1	Proteotypic
K1C18_HUMAN	198	964.461	AQYDELAR		1	Proteotypic
K1C18_HUMAN	289	991.482	VVSETNDTK		1	Proteotypic
K1C18_HUMAN	458	1040.598	IVLQIDNAR		2	Shared
K1C18_HUMAN	470	1043.459	YETELAMR	Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	483	1045.540	VIDDTNITR		1	Proteotypic
K1C18_HUMAN	544	1064.550	LEAEIATYR		1	Proteotypic
K1C18_HUMAN	726	1108.482	ETMQSLNDR	Oxidation (M)	1	Proteotypic
K1C18_HUMAN	789	1124.477	ETMQSLNDR	Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	969	1173.635	KVIDDTNITR		1	Proteotypic
K1C18_HUMAN	1258	1254.628	VKYETELAMR	Oxidation (M)	1	Proteotypic
K1C18_HUMAN	1291	1266.632	QSVENDIHGLR		1	Proteotypic
K1C18_HUMAN	1306	1270.623	VKYETELAMR	Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	1345	1285.713	LQLETEIEALK		1	Proteotypic
K1C18_HUMAN	1366	1291.652	SLGSVQAPSYGAR		1	Proteotypic
K1C18_HUMAN	1368	1291.714	VKLEAEIATYR		1	Proteotypic
K1C18_HUMAN	1449	1318.663	AQIFANTVDNAR		1	Proteotypic
K1C18_HUMAN	1609	1394.727	QSVENDIHGLRK		1	Proteotypic
K1C18_HUMAN	1655	1418.740	QAQEYEALLNIK		1	Proteotypic
K1C18_HUMAN	1755	1472.758	ASLENSLREVEAR		1	Proteotypic
K1C18_HUMAN	1824	1521.734	TVQSLEIDLDSMR	Oxidation (M)	1	Proteotypic
K1C18_HUMAN	1852	1537.729	TVQSLEIDLDSMR	Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	1909	1579.759	PVSSAASVYAGAGGSGSR		1	Proteotypic
K1C18_HUMAN	2045	1693.830	RTVQSLEIDLDSMR	Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	2194	1883.000	GLQAQIASSGLTVEVDAPK		1	Proteotypic
K1C18_HUMAN	2318	2208.160	LQLETEIEALKEELFMK	Dioxidation (M)	1	Proteotypic

K1C18_HUMAN	2333	2308.079	GGMGSGGLATGIAGGLAGMGGIQNEK	Dioxidation (M); Oxidation (M)	1	Proteotypic
K1C18_HUMAN	2337	2324.074	GGMGSGGLATGIAGGLAGMGGIQNEK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	2339	2336.255	LQLETEIEALKEELLFMKK	Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	2366	2701.374	YALQMEQLNGILLHLESELAQTR	Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	2372	2771.244	LLEDGEDFNLDALDSSNSMQTIQK	Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	2375	2927.345	RILLEDGEDFNLDALDSSNSMQTIQK	Dioxidation (M)	1	Proteotypic
K1C18_HUMAN	2385	3430.540	GGMGSGGLATGIAGGLAGMGGIQNEKETMQSLNDR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	1	Proteotypic
K1C19_HUMAN	196	964.436	QNQEYQR		1	Proteotypic
K1C19_HUMAN	227	974.514	SEVTDLRR		1	Proteotypic
K1C19_HUMAN	293	992.493	FETEQALR		2	Shared
K1C19_HUMAN	299	994.494	APSIHGGSGGR		1	Proteotypic
K1C19_HUMAN	420	1028.587	VDELTLAR		5	Shared
K1C19_HUMAN	458	1040.598	IVLQIDNAR		2	Shared
K1C19_HUMAN	577	1072.588	ILGATIENSR		1	Proteotypic
K1C19_HUMAN	772	1119.534	LTMQNLNDR	Oxidation (M)	1	Proteotypic
K1C19_HUMAN	779	1121.572	LEQEIATYR		6	Shared
K1C19_HUMAN	837	1135.529	LTMQNLNDR	Dioxidation (M)	1	Proteotypic
K1C19_HUMAN	1003	1184.688	RVLDELTLAR		5	Shared
K1C19_HUMAN	1129	1219.587	MSVEADINGLR	Oxidation (M)	2	Shared
K1C19_HUMAN	1143	1221.635	TKFETEQALR		2	Shared
K1C19_HUMAN	1159	1226.589	NHEEEISTLR		1	Proteotypic
K1C19_HUMAN	1195	1235.582	MSVEADINGLR	Dioxidation (M)	2	Shared
K1C19_HUMAN	1221	1242.646	ALEAANGELEVK		1	Proteotypic
K1C19_HUMAN	1414	1307.628	TDLEMQIEGLK	Dioxidation (M)	2	Shared
K1C19_HUMAN	1441	1315.710	DKILGATIENSR		1	Proteotypic
K1C19_HUMAN	1533	1354.684	KNHEEEISTLR		1	Proteotypic
K1C19_HUMAN	1558	1364.705	SRLEQEIATYR		1	Proteotypic
K1C19_HUMAN	1566	1369.593	SQYEVMAEQNR	Oxidation (M)	1	Proteotypic
K1C19_HUMAN	1593	1385.588	SQYEVMAEQNR	Dioxidation (M)	1	Proteotypic
K1C19_HUMAN	1598	1388.678	AALEDTLAETEAR		1	Proteotypic
K1C19_HUMAN	1602	1391.683	MSVEADINGLRR	Dioxidation (M)	2	Shared
K1C19_HUMAN	1804	1500.699	EVAGHTEQLQMSR	Oxidation (M)	1	Proteotypic
K1C19_HUMAN	1811	1513.683	SQYEVMAEQNRK	Dioxidation (M)	1	Proteotypic
K1C19_HUMAN	1814	1516.694	EVAGHTEQLQMSR	Dioxidation (M)	1	Proteotypic
K1C19_HUMAN	1876	1553.743	QSSATSSFGLGGGSVR		1	Proteotypic
K1C19_HUMAN	2174	1848.950	TLQGLEIELQSQLSMK	Dioxidation (M)	2	Shared
K1C19_HUMAN	2264	2013.001	GQVGGQVSVEVDSAPGTDLAK		1	Proteotypic
K1C19_HUMAN	2310	2154.077	TDLEMQIEGLKEELAYLK	Dioxidation (M)	1	Proteotypic
K1C19_HUMAN	2324	2259.055	TEELNREVAGHTEQLQMSR	Dioxidation (M)	1	Proteotypic
K1C9_HUMAN	49	912.401	MTLDDFR	Oxidation (M)	1	Proteotypic
K1C9_HUMAN	98	928.396	MTLDDFR	Dioxidation (M)	1	Proteotypic
K1C9_HUMAN	526	1059.556	TLLDIDNTR		1	Proteotypic
K1C9_HUMAN	608	1080.487	STMQELNSR	Oxidation (M)	1	Proteotypic
K1C9_HUMAN	675	1096.482	STMQELNSR	Dioxidation (M)	1	Proteotypic
K1C9_HUMAN	678	1097.481	FEMEQLNR	Dioxidation (M)	1	Proteotypic
K1C9_HUMAN	918	1156.584	QGVDAINGLR		1	Proteotypic

K1C9_HUMAN	1141	1221.591	QVLDNLTMEK	Dioxidation (M)	1	Proteotypic
K1C9_HUMAN	1173	1231.591	SGGGGGGGLGSGGSIR		1	Proteotypic
K1C9_HUMAN	1189	1234.521	FSSSSGYGGSSR		1	Proteotypic
K1C9_HUMAN	1493	1338.660	IKFEMEQLNR	Dioxidation (M)	1	Proteotypic
K1C9_HUMAN	1914	1585.758	VQALEEANNDLENK		1	Proteotypic
K1C9_HUMAN	2356	2509.124	EIETYHNLLEGGQEDFESSGAGK		1	Proteotypic
K1H1_HUMAN	311	998.576	LVVQIDNAK		1	Proteotypic
K1H1_HUMAN	997	1184.513	ETMQFLNDR	Dioxidation (M)	1	Proteotypic
K22E_HUMAN	217	972.524	IEISELNR		2	Shared
K22E_HUMAN	296	993.561	LQGEIAHVK		1	Proteotypic
K22E_HUMAN	447	1036.519	YLDGLTAER		1	Proteotypic
K22E_HUMAN	548	1065.509	YEDEINKR		8	Shared
K22E_HUMAN	614	1081.592	FASFIDKVR		11	Shared
K22E_HUMAN	717	1106.536	AQYEEIAQR		4	Shared
K22E_HUMAN	785	1123.518	KLLEGEECR	Trioxidation (C)	6	Shared
K22E_HUMAN	1039	1192.609	YEELQVTVGR		1	Proteotypic
K22E_HUMAN	1093	1207.608	TAAENDFVTLK		1	Proteotypic
K22E_HUMAN	1256	1253.600	GFSSGSAAVVSIGGSR		1	Proteotypic
K22E_HUMAN	1474	1328.632	NVQDAIADAEQR		1	Proteotypic
K22E_HUMAN	1476	1328.719	NLDLDSIIAEVK		6	Shared
K22E_HUMAN	1569	1370.704	LNDLEEFALQQAK		1	Proteotypic
K22E_HUMAN	1732	1459.792	VDLLNQEIEFLK		1	Proteotypic
K22E_HUMAN	1764	1474.778	FLEQQNQVLQTK		2	Shared
K2C1_HUMAN	217	972.524	IEISELNR		2	Shared
K2C1_HUMAN	429	1030.428	DVDGAYMTK	Dioxidation (M)	1	Proteotypic
K2C1_HUMAN	436	1032.509	TLLEGEESR		1	Proteotypic
K2C1_HUMAN	548	1065.509	YEDEINKR		8	Shared
K2C1_HUMAN	656	1091.496	GSGGGSSGGSIGGR		1	Proteotypic
K2C1_HUMAN	790	1124.535	AEAESLYQSK		1	Proteotypic
K2C1_HUMAN	900	1151.546	NKYEDEINK		6	Shared
K2C1_HUMAN	916	1156.507	DYQELMNTK	Oxidation (M)	1	Proteotypic
K2C1_HUMAN	962	1172.502	DYQELMNTK	Dioxidation (M)	1	Proteotypic
K2C1_HUMAN	986	1178.593	YEELQITAGR		1	Proteotypic
K2C1_HUMAN	1072	1200.683	LRSEIDNVKK		1	Proteotypic
K2C1_HUMAN	1288	1264.630	TNAENEFVTIK		1	Proteotypic
K2C1_HUMAN	1325	1276.703	LALDLEIATYR		1	Proteotypic
K2C1_HUMAN	1397	1301.708	SLDLDSIIAEVK		1	Proteotypic
K2C1_HUMAN	1415	1307.647	NKYEDEINKR		5	Shared
K2C1_HUMAN	1496	1339.662	SKAEAESLYQSK		1	Proteotypic
K2C1_HUMAN	1514	1347.507	NMQDMVEDYR	Dioxidation (M); Oxidation (M)	1	Proteotypic
K2C1_HUMAN	1535	1356.688	LNDLEDALQQAK		1	Proteotypic
K2C1_HUMAN	1553	1363.502	NMQDMVEDYR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
K2C1_HUMAN	1588	1382.683	SLNNQFASFIDK		1	Proteotypic
K2C1_HUMAN	1605	1392.725	TNAENEFVTIKK		1	Proteotypic
K2C1_HUMAN	1764	1474.778	FLEQQNQVLQTK		2	Shared
K2C1_HUMAN	1930	1598.826	NKLNDLEDALQQAK		1	Proteotypic



K2C1_HUMAN	1970	1637.853	SLNNQFASFIDKVR		1	Proteotypic
K2C1_HUMAN	2064	1715.844	QISNLQQSISDAEQR		1	Proteotypic
K2C1_HUMAN	2094	1755.691	FSSCGGGGSGFAGGGFGSR	Trioxidation (C)	1	Proteotypic
K2C1_HUMAN	2228	1940.980	LNDLEDALQQAKEDLAR		1	Proteotypic
K2C1_HUMAN	2277	2048.867	LDSELKNMQDMVEDYR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
K2C1_HUMAN	2316	2183.118	NKLNDLEDALQQAKEDLAR		1	Proteotypic
K2C4_HUMAN	717	1106.536	AQYEEIAQR		4	Shared
K2C4_HUMAN	855	1141.533	EYQELMSVK	Oxidation (M)	2	Shared
K2C4_HUMAN	920	1157.527	EYQELMSVK	Dioxidation (M)	2	Shared
K2C4_HUMAN	1324	1276.703	LALDIEIATYR		4	Shared
K2C4_HUMAN	1625	1404.798	LALDIEIATYRK		4	Shared
K2C5_HUMAN	29	905.461	FLEQQNK		14	Shared
K2C5_HUMAN	141	944.504	GRLDSELR		4	Shared
K2C5_HUMAN	376	1015.530	QLDSIVGER		3	Shared
K2C5_HUMAN	510	1057.439	DVDAAYMNK	Dioxidation (M)	5	Shared
K2C5_HUMAN	548	1065.509	YEDEINKR		8	Shared
K2C5_HUMAN	614	1081.592	FASFIDKVR		11	Shared
K2C5_HUMAN	663	1092.520	AQYEEIANR		1	Proteotypic
K2C5_HUMAN	735	1110.567	ISISTSGGSFR		1	Proteotypic
K2C5_HUMAN	785	1123.518	KLLEGEECR	Trioxidation (C)	6	Shared
K2C5_HUMAN	900	1151.546	NKYEDEINK		6	Shared
K2C5_HUMAN	1040	1193.568	YEELQQTAGR		1	Proteotypic
K2C5_HUMAN	1303	1269.555	NMQDLVEDFK	Dioxidation (M)	1	Proteotypic
K2C5_HUMAN	1415	1307.647	NKYEDEINKR		5	Shared
K2C5_HUMAN	1476	1328.719	NLDLDSIIAEVK		6	Shared
K2C6C_HUMAN	29	905.461	FLEQQNK		14	Shared
K2C6C_HUMAN	141	944.504	GRLDSELR		4	Shared
K2C6C_HUMAN	376	1015.530	QLDSIVGER		3	Shared
K2C6C_HUMAN	408	1025.514	SGFSSISVSR		1	Proteotypic
K2C6C_HUMAN	510	1057.439	DVDAAYMNK	Dioxidation (M)	5	Shared
K2C6C_HUMAN	548	1065.509	YEDEINKR		8	Shared
K2C6C_HUMAN	614	1081.592	FASFIDKVR		11	Shared
K2C6C_HUMAN	717	1106.536	AQYEEIAQR		4	Shared
K2C6C_HUMAN	785	1123.518	KLLEGEECR	Trioxidation (C)	6	Shared
K2C6C_HUMAN	900	1151.546	NKYEDEINK		6	Shared
K2C6C_HUMAN	954	1168.543	EYQELMNVK	Oxidation (M)	4	Shared
K2C6C_HUMAN	1000	1184.538	EYQELMNVK	Dioxidation (M)	4	Shared
K2C6C_HUMAN	1415	1307.647	NKYEDEINKR		5	Shared
K2C6C_HUMAN	1476	1328.719	NLDLDSIIAEVK		6	Shared
K2C6C_HUMAN	1628	1406.704	ADTLTDEINFLR		2	Shared
K2C6C_HUMAN	1711	1446.768	AIGGGLSSVGGGSSTIK		2	Shared
K2C7_HUMAN	29	905.461	FLEQQNK		14	Shared
K2C7_HUMAN	108	931.461	LLEGESR		4	Shared
K2C7_HUMAN	277	989.467	QEESQIK		1	Proteotypic
K2C7_HUMAN	363	1013.514	LDADPSLQR		1	Proteotypic
K2C7_HUMAN	366	1014.433	DVDAAYMSK	Oxidation (M)	1	Proteotypic

K2C7_HUMAN	401	1023.429	NEISEMNR	Dioxidation (M)	1	Proteotypic
K2C7_HUMAN	428	1030.428	DVDAAYMSK	Dioxidation (M)	1	Proteotypic
K2C7_HUMAN	467	1042.566	LQAEIDNIK		1	Proteotypic
K2C7_HUMAN	525	1059.556	KLLEGEESR		3	Shared
K2C7_HUMAN	614	1081.592	FASFIDKVR		11	Shared
K2C7_HUMAN	661	1091.561	FETLQAQAGK		1	Proteotypic
K2C7_HUMAN	705	1103.572	SAYGGPVGAGIR		1	Proteotypic
K2C7_HUMAN	855	1141.533	EYQELMSVK	Oxidation (M)	2	Shared
K2C7_HUMAN	920	1157.527	EYQELMSVK	Dioxidation (M)	2	Shared
K2C7_HUMAN	987	1179.552	NKYEDEINR		1	Proteotypic
K2C7_HUMAN	1006	1185.599	QEELEAALQR		1	Proteotypic
K2C7_HUMAN	1065	1200.599	LEAAIAEAEER		1	Proteotypic
K2C7_HUMAN	1133	1219.645	TAAENEFVVLK		2	Shared
K2C7_HUMAN	1216	1241.636	GQLEALQVDGGR		1	Proteotypic
K2C7_HUMAN	1234	1244.636	VRQEESEQIK		1	Proteotypic
K2C7_HUMAN	1313	1271.697	SLDLDGIIAEVK		1	Proteotypic
K2C7_HUMAN	1324	1276.703	LALDIEIATYR		4	Shared
K2C7_HUMAN	1518	1347.740	TAAENEFVVLKK		2	Shared
K2C7_HUMAN	1592	1384.731	AKQEELEAALQR		1	Proteotypic
K2C7_HUMAN	1608	1394.621	NTRNEISEMNR	Dioxidation (M)	1	Proteotypic
K2C7_HUMAN	1619	1399.731	AKLEAAIAEAEER		1	Proteotypic
K2C7_HUMAN	1625	1404.798	LALDIEIATYRK		4	Shared
K2C7_HUMAN	1626	1405.699	SIHFSSPVFTSR	Acetyl (Protein N-term)	1	Proteotypic
K2C7_HUMAN	1652	1417.720	VDALNDEINFLR		1	Proteotypic
K2C7_HUMAN	1696	1440.768	LQAEIDNIKNQR		1	Proteotypic
K2C7_HUMAN	1700	1441.793	LPDIFEAQIAGLR		1	Proteotypic
K2C7_HUMAN	1721	1452.830	EVTINQSL LAPLR		1	Proteotypic
K2C7_HUMAN	1771	1477.753	PGGLGSSSLYGLGASR		1	Proteotypic
K2C7_HUMAN	2073	1730.906	PGGLGSSSLYGLGASRPR		1	Proteotypic
K2C7_HUMAN	2141	1811.963	LEAAIAEAEER GELALK		1	Proteotypic
K2C7_HUMAN	2263	2011.095	AKLEAAIAEAEER GELALK		1	Proteotypic
K2C7_HUMAN	2349	2448.334	EVTINQSL LAPLR LRDADPSLQR		1	Proteotypic
K2C7_HUMAN	2380	3041.435	TLNETELTELQSQISDTSVVL SMDNSR	Dioxidation (M)	1	Proteotypic
K2C72_HUMAN	510	1057.439	DVDAAYMNK	Dioxidation (M)	5	Shared
K2C72_HUMAN	614	1081.592	FASFIDKVR		11	Shared
K2C72_HUMAN	1133	1219.645	TAAENEFVVLK		2	Shared
K2C72_HUMAN	1518	1347.740	TAAENEFVVLKK		2	Shared
K2C75_HUMAN	29	905.461	FLEQQNK		14	Shared
K2C75_HUMAN	510	1057.439	DVDAAYMNK	Dioxidation (M)	5	Shared
K2C75_HUMAN	548	1065.509	YEDEINKR		8	Shared
K2C75_HUMAN	601	1078.504	AQYEDIANR		2	Shared
K2C75_HUMAN	614	1081.592	FASFIDKVR		11	Shared
K2C75_HUMAN	785	1123.518	KLLEGEECR	Trioxidation (C)	6	Shared
K2C75_HUMAN	1454	1319.708	TAAENEFVALKK		1	Proteotypic
K2C75_HUMAN	1476	1328.719	NLDLDSIIAEVK		6	Shared
K2C79_HUMAN	29	905.461	FLEQQNK		14	Shared

K2C79_HUMAN	141	944.504	GRLDSELR		4	Shared
K2C79_HUMAN	614	1081.592	FASFIDKVR		11	Shared
K2C79_HUMAN	842	1136.571	YEELQVTAGK		1	Proteotypic
K2C79_HUMAN	900	1151.546	NKYEDEINK		6	Shared
K2C79_HUMAN	1476	1328.719	NLDLDSIIAEVK		6	Shared
KCY_HUMAN	194	963.518	FLIDGFPR		1	Proteotypic
KCY_HUMAN	1620	1400.555	EMDQTMAANAQK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
KPYM_HUMAN	5	899.465	MQHLIAR	Dioxidation (M)	1	Proteotypic
KPYM_HUMAN	43	911.446	ASDVHEVR		1	Proteotypic
KPYM_HUMAN	162	952.473	IENHEGVR		1	Proteotypic
KPYM_HUMAN	279	989.503	GSGTAEVELK		1	Proteotypic
KPYM_HUMAN	350	1010.486	VNFAMNVGK	Dioxidation (M)	1	Proteotypic
KPYM_HUMAN	387	1018.508	GDYPLEAVR		1	Proteotypic
KPYM_HUMAN	455	1039.541	KASDVHEVR		1	Proteotypic
KPYM_HUMAN	759	1117.598	GSGTAEVELKK		1	Proteotypic
KPYM_HUMAN	852	1140.603	GDLGIEIPA EK		1	Proteotypic
KPYM_HUMAN	1048	1196.640	LDIDSPPIAR		1	Proteotypic
KPYM_HUMAN	1095	1208.632	SVETLKEMIK	Dioxidation (M)	1	Proteotypic
KPYM_HUMAN	1519	1349.661	NTGIICTIGPASR	Trioxidation (C)	1	Proteotypic
KPYM_HUMAN	1735	1461.808	IYVDDGLISLQVK		1	Proteotypic
KPYM_HUMAN	1969	1635.883	GVNLPGAAVDLPVSEK		1	Proteotypic
KPYM_HUMAN	2110	1778.869	GADFLVTEVENGGSLGSK		1	Proteotypic
KPYM_HUMAN	2176	1852.899	RFDEILEASDGIMVAR	Dioxidation (M)	1	Proteotypic
KPYM_HUMAN	2198	1890.882	FGVEQDVMVFASFIR	Dioxidation (M)	1	Proteotypic
KPYM_HUMAN	2222	1930.905	AGKPVICATQMLESMIK	Dioxidation (M); Dioxidation (M); Trioxidation (C)	1	Proteotypic
KPYM_HUMAN	2313	2174.111	LAPITSDPTEATAVGAVEASFK		1	Proteotypic
KPYM_HUMAN	2352	2464.285	TATESFASDPILYRPVAVALDTK		1	Proteotypic
KRT84_HUMAN	29	905.461	FLEQQNK		14	Shared
KRT84_HUMAN	108	931.461	LLEGEESR		4	Shared
KRT84_HUMAN	197	964.461	AQYEEVAR		1	Proteotypic
KRT84_HUMAN	614	1081.592	FASFIDKVR		11	Shared
KRT84_HUMAN	641	1087.562	RLLEGEESR		1	Proteotypic
KRT85_HUMAN	29	905.461	FLEQQNK		14	Shared
KRT85_HUMAN	270	987.524	LTAEIENAK		1	Proteotypic
KRT85_HUMAN	1134	1219.645	ATAENEFVVLK		1	Proteotypic
KRT85_HUMAN	1233	1244.636	TKEEINELNR		2	Shared
KRT85_HUMAN	1517	1347.740	ATAENEFVVLKK		1	Proteotypic
KU70_HUMAN	335	1004.554	ILELDQFK		1	Proteotypic
KU70_HUMAN	706	1104.548	IMATPEQVGK	Dioxidation (M)	1	Proteotypic
KU70_HUMAN	1091	1206.650	DSLIFLVDASK		1	Proteotypic
KU86_HUMAN	579	1072.624	LTIGSNLSIR		1	Proteotypic
KU86_HUMAN	745	1114.548	FFMGNQVLK	Dioxidation (M)	1	Proteotypic
KU86_HUMAN	905	1152.633	KVITMFVQR	Dioxidation (M)	1	Proteotypic
KU86_HUMAN	1606	1393.621	EEASGSSVTAE EAK		1	Proteotypic
KU86_HUMAN	1823	1521.716	EEASGSSVTAE EAKK		1	Proteotypic
LA_HUMAN	2480	1548.840	LTTDFNIVEALSK		1	Proteotypic

LAMP1_HUMAN	2402	1032.545	TVESITDIR		1	Proteotypic
LDHA_HUMAN	51	912.576	LVITAGAR		1	Proteotypic
LDHA_HUMAN	101	928.575	FIIPNVVK		1	Proteotypic
LDHA_HUMAN	349	1010.467	DYNVTANSK		1	Proteotypic
LDHA_HUMAN	768	1118.634	DQLIYNLLK		1	Proteotypic
LDHA_HUMAN	831	1133.556	VTLTSEEEAR		1	Proteotypic
LDHA_HUMAN	1056	1198.649	RVHPVSTMIK	Dioxidation (M)	1	Proteotypic
LDHA_HUMAN	1203	1238.556	VIGSGCNLDSAR	Trioxidation (C)	2	Shared
LDHA_HUMAN	1284	1263.671	QVVESAYEVIK		1	Proteotypic
LDHA_HUMAN	2001	1656.846	DLADELALVDVIEDK		1	Proteotypic
LDHA_HUMAN	2274	2044.025	ITVVGVGAVGMACAISILMK	Dioxidation (M); Dioxidation (M); Trioxidation (C)	1	Proteotypic
LDHB_HUMAN	53	912.576	IVVVTAGVR		1	Proteotypic
LDHB_HUMAN	174	956.606	FIIPQIVK		1	Proteotypic
LDHB_HUMAN	179	958.545	GLTSVINQK		1	Proteotypic
LDHB_HUMAN	925	1157.629	LKDDEVAQLK		1	Proteotypic
LDHB_HUMAN	1203	1238.556	VIGSGCNLDSAR	Trioxidation (C)	2	Shared
LDHB_HUMAN	1346	1285.724	LKDDEVAQLKK		1	Proteotypic
LDHB_HUMAN	1383	1298.643	MVVESAYEVIK	Dioxidation (M)	1	Proteotypic
LDHB_HUMAN	1806	1509.767	IVADKDYSVTANSK		1	Proteotypic
LDHB_HUMAN	1961	1628.851	SLADELALVDVLEDK		1	Proteotypic
LDHB_HUMAN	2046	1693.889	LIAPVAEEEEATVPNNK		1	Proteotypic
LDHB_HUMAN	2254	1995.038	ITVVGVGQVGMACAISILGK	Dioxidation (M); Trioxidation (C)	1	Proteotypic
LEG4_HUMAN	531	1061.587	VVFNTLQGGK		1	Proteotypic
LEG4_HUMAN	1687	1434.758	VGSSGDIALHINPR		1	Proteotypic
LEG4_HUMAN	2230	1944.985	ITHNPFPGPGQFFDLSIR		1	Proteotypic
LG3BP_HUMAN	516	1058.510	LADGGATNQGR		1	Proteotypic
LG3BP_HUMAN	819	1130.666	RIDITLSSVK		1	Proteotypic
LG3BP_HUMAN	1534	1354.771	SDLAVPSELALLK		1	Proteotypic
LG3BP_HUMAN	1924	1591.784	ELSEALGQIFDSQR		1	Proteotypic
LKHA4_HUMAN	2427	1133.611	ELVALMSAIR	Dioxidation (M)	1	Proteotypic
LMAN1_HUMAN	209	971.504	DIDNLVQR		1	Proteotypic
LMAN1_HUMAN	235	976.473	QVFEGQNR		1	Proteotypic
LMNA_HUMAN	14	900.492	LEAALGEAK		1	Proteotypic
LMNA_HUMAN	75	917.445	DLEDSLAR		1	Proteotypic
LMNA_HUMAN	79	921.386	METPSQR	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
LMNA_HUMAN	150	948.488	KLESTESR		1	Proteotypic
LMNA_HUMAN	210	971.504	LSPSPTSQR		1	Proteotypic
LMNA_HUMAN	220	973.472	LLEGEER		3	Shared
LMNA_HUMAN	412	1027.566	LADALQELR		1	Proteotypic
LMNA_HUMAN	419	1028.587	LEAALGEAKK		1	Proteotypic
LMNA_HUMAN	465	1042.541	EGDLIAAQAR		1	Proteotypic
LMNA_HUMAN	539	1063.497	QLQDEMLR	Dioxidation (M)	1	Proteotypic
LMNA_HUMAN	644	1088.546	SLETENAGLR		1	Proteotypic
LMNA_HUMAN	695	1101.567	KLLEGEER		3	Shared
LMNA_HUMAN	744	1114.438	EREMAEMR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
LMNA_HUMAN	766	1118.582	EAALSTALSEK		1	Proteotypic

LMNA_HUMAN	814	1130.520	EDLQELNDR		1	Proteotypic
LMNA_HUMAN	878	1147.572	ITESEEVSR		1	Proteotypic
LMNA_HUMAN	941	1164.541	AAYEAE LGDAR		1	Proteotypic
LMNA_HUMAN	957	1170.636	KEGD LIAA QAR		1	Proteotypic
LMNA_HUMAN	993	1181.604	TLEGEL HDLR		1	Proteotypic
LMNA_HUMAN	1013	1186.631	LRDLED SLAR		1	Proteotypic
LMNA_HUMAN	1035	1191.592	KQLQDEMLR	Dioxidation (M)	1	Proteotypic
LMNA_HUMAN	1074	1202.567	VAVEEVDEEGK		1	Proteotypic
LMNA_HUMAN	1225	1242.718	LKDLEALLNSK		1	Proteotypic
LMNA_HUMAN	1319	1274.683	EAALSTALSEKR		1	Proteotypic
LMNA_HUMAN	1369	1292.636	AAYEAE LGDARK		1	Proteotypic
LMNA_HUMAN	1544	1358.679	SGAQASSTPLSPTR		1	Proteotypic
LMNA_HUMAN	1675	1429.778	IDSLSAQLSQLQK		1	Proteotypic
LMNA_HUMAN	1828	1522.730	TALINSTGEEVAMR	Dioxidation (M)	1	Proteotypic
LMNA_HUMAN	1858	1540.744	LQTMKEELDFQK	Dioxidation (M)	1	Proteotypic
LMNA_HUMAN	1890	1565.743	SVGGSGGGSGFDNLVTR		1	Proteotypic
LMNA_HUMAN	1934	1604.805	VAVEEVDEEGKFVR		1	Proteotypic
LMNA_HUMAN	1960	1628.801	LQEKEDLQELNDR		1	Proteotypic
LMNA_HUMAN	1988	1650.825	TALINSTGEEVAMRK	Dioxidation (M)	1	Proteotypic
LMNA_HUMAN	2052	1698.963	IRIDSLSAQLSQLQK		1	Proteotypic
LMNA_HUMAN	2093	1751.855	NSNLVGA AHEELQQSR		1	Proteotypic
LMNA_HUMAN	2219	1924.909	MQQQ LDEYQELLDIK	Dioxidation (M)	1	Proteotypic
LMNB1_HUMAN	89	924.386	EELMESR	Dioxidation (M)	1	Proteotypic
LMNB1_HUMAN	220	973.472	LLEGEEER		3	Shared
LMNB1_HUMAN	471	1043.561	LQEKEELR		2	Shared
LMNB1_HUMAN	695	1101.567	KLLEGEER		3	Shared
LMNB1_HUMAN	754	1116.516	NSQGEEVAQR		1	Proteotypic
LMNB1_HUMAN	909	1153.609	AGGPTTPLSPTR		1	Proteotypic
LMNB1_HUMAN	958	1170.650	IQELEDLLAK		1	Proteotypic
LMNB1_HUMAN	965	1172.640	DAALATALGDKK		1	Proteotypic
LMNB1_HUMAN	1836	1527.683	LSSEMNTSTVNSAR	Dioxidation (M)	1	Proteotypic
LMNB2_HUMAN	220	973.472	LLEGEEER		3	Shared
LMNB2_HUMAN	471	1043.561	LQEKEELR		2	Shared
LMNB2_HUMAN	517	1058.536	EGELTVAQGR		1	Proteotypic
LMNB2_HUMAN	639	1087.551	GLES DVAELR		1	Proteotypic
LMNB2_HUMAN	657	1091.528	MAQALEELR	Dioxidation (M)	1	Proteotypic
LMNB2_HUMAN	695	1101.567	KLLEGEER		3	Shared
LMNB2_HUMAN	787	1123.599	AGGPATPLSPTR		1	Proteotypic
LMNB2_HUMAN	1031	1190.614	ISEKEEV TTR		1	Proteotypic
LMNB2_HUMAN	1295	1267.600	ATSSSSGSLSATGR		1	Proteotypic
LYPA1_HUMAN	62	914.507	ALIDQEVK		1	Proteotypic
LYPA1_HUMAN	1307	1270.642	ASFPQGP IGGANR		1	Proteotypic
M2OM_HUMAN	145	945.540	NVFNALIR		1	Proteotypic
M2OM_HUMAN	1695	1440.732	AATASAGAGGIDGKPR	Acetyl (Protein N-term)	1	Proteotypic
MCA3_HUMAN	2430	1156.634	AAA AELSLLEK	Acetyl (Protein N-term)	1	Proteotypic
MDHC_HUMAN	70	915.503	LGVTANDVK		1	Proteotypic

MDHC_HUMAN	404	1024.522	KLSSAMSAAK	Dioxidation (M)	1	Proteotypic
MDHC_HUMAN	938	1163.594	GEFVTTVQQR		1	Proteotypic
MDHC_HUMAN	1604	1392.704	FVEGLPINDFSR		1	Proteotypic
MDHC_HUMAN	1649	1416.728	DLDVAILVGSMPR	Dioxidation (M)	1	Proteotypic
MDHC_HUMAN	1718	1449.735	DVIATDKEDVAFK		1	Proteotypic
MDHC_HUMAN	2092	1747.877	VIVVGNPANTNCLTASK	Trioxidation (C)	1	Proteotypic
MDHC_HUMAN	2148	1815.841	ELTEEKESAFELSSA		1	Proteotypic
MET7A_HUMAN	2481	1622.794	ELFSNLQEFAGPSGK		1	Proteotypic
MIF_HUMAN	437	1034.543	LLCGLLAER	Trioxidation (C)	1	Proteotypic
MIF_HUMAN	1450	1318.670	PMFIVNTNVPR	Dioxidation (M)	1	Proteotypic
MLRM_HUMAN	552	1066.496	ELLTTMGDR	Dioxidation (M)	1	Proteotypic
MLRM_HUMAN	1161	1227.610	LNGTDPEDVIR		1	Proteotypic
MLRM_HUMAN	1272	1259.593	GNFNIEFTR		1	Proteotypic
MLRM_HUMAN	1299	1268.546	EAFNMIDQNR	Dioxidation (M)	1	Proteotypic
MLRM_HUMAN	2294	2121.968	ATSNVVFAMFDQSQIQEFK	Dioxidation (M)	1	Proteotypic
MPCP_HUMAN	381	1016.598	GSSASLVLKR		1	Proteotypic
MPCP_HUMAN	439	1035.507	FGFYEVFK		1	Proteotypic
MPCP_HUMAN	994	1181.644	GIFNGFSVTLK		1	Proteotypic
MPCP_HUMAN	1551	1360.710	IQTQPGYANTLR		1	Proteotypic
MPCP_HUMAN	1947	1618.839	LPRPPPEMPESLK	Dioxidation (M)	1	Proteotypic
MTPN_HUMAN	2490	1746.842	GPDGLTAFEATDNQAIK		1	Proteotypic
MVP_HUMAN	377	1015.555	AQLELEVSK		1	Proteotypic
MVP_HUMAN	389	1019.529	ATEEFIIR	Acetyl (Protein N-term)	1	Proteotypic
MVP_HUMAN	434	1031.513	ILDQSEAEK		1	Proteotypic
MVP_HUMAN	739	1113.603	AQQLAEVEVK		1	Proteotypic
MVP_HUMAN	810	1129.598	IEGEGSVLQAK		1	Proteotypic
MVP_HUMAN	1356	1288.633	DLAVAGPEMQVK	Dioxidation (M)	1	Proteotypic
MVP_HUMAN	1943	1616.852	DAQGLVLFVDVTGQVR		1	Proteotypic
MYH14_HUMAN	215	972.488	VAQLEEER		1	Proteotypic
MYH14_HUMAN	581	1073.499	ALEEEQEAR		1	Proteotypic
MYH14_HUMAN	628	1085.547	VAEQAANDLR		1	Proteotypic
MYH14_HUMAN	658	1091.550	FDQLLAEEK		2	Shared
MYH14_HUMAN	1132	1219.645	KFDQLLAEEK		2	Shared
MYH14_HUMAN	1262	1256.672	KEEELQAALAR		2	Shared
MYH14_HUMAN	1405	1304.661	EQADFALEALAK		1	Proteotypic
MYH14_HUMAN	2070	1725.941	QLLQANPILEAFGNAK		2	Shared
MYH9_HUMAN	20	902.507	ASITALEAK		1	Proteotypic
MYH9_HUMAN	73	916.461	DLEGLSQR		1	Proteotypic
MYH9_HUMAN	86	923.487	VVFQEFR		1	Proteotypic
MYH9_HUMAN	264	985.556	GALALEEKR		1	Proteotypic
MYH9_HUMAN	310	998.555	GDLPFVPR		1	Proteotypic
MYH9_HUMAN	453	1038.655	VKPLLQVSR		1	Proteotypic
MYH9_HUMAN	514	1057.552	EKQLAENR		1	Proteotypic
MYH9_HUMAN	547	1065.501	LQEMEGTVK	Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	606	1079.480	ALEEAMEQK	Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	658	1091.550	FDQLLAEEK		2	Shared

MYH9_HUMAN	912	1154.656	RGDLPFVVPR		1	Proteotypic
MYH9_HUMAN	1038	1192.609	ALELDSNLYR		1	Proteotypic
MYH9_HUMAN	1117	1214.662	ASREEILAQAK		1	Proteotypic
MYH9_HUMAN	1132	1219.645	KFDQLLAEK		2	Shared
MYH9_HUMAN	1147	1222.631	AGVLAHLEER		1	Proteotypic
MYH9_HUMAN	1262	1256.672	KEEELQAALAR		2	Shared
MYH9_HUMAN	1334	1281.503	EMEALEDER	Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	1340	1283.708	VEAQLQELQVK		1	Proteotypic
MYH9_HUMAN	1370	1292.665	VKLQEMEGTVK	Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	1404	1304.661	EQADFAIEALAK		1	Proteotypic
MYH9_HUMAN	1478	1330.600	QLEEAEEEAQR		1	Proteotypic
MYH9_HUMAN	1633	1409.598	EMEALEDERK	Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	1638	1411.803	KVEAQLQELQVK		1	Proteotypic
MYH9_HUMAN	1783	1485.892	VISGVLQGNIVFK		1	Proteotypic
MYH9_HUMAN	1841	1529.757	IAQLEEQLDNETK		1	Proteotypic
MYH9_HUMAN	1926	1596.657	ELEDATETADAMNR	Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	1953	1622.684	NTDQASMPDNTAAQK	Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	2029	1677.788	ALEEAMEQKAELER	Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	2030	1678.791	IMGIPEEEQMGLLR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	2070	1725.941	QLLQANPILEAFGNAK		2	Shared
MYH9_HUMAN	2090	1747.713	TEMEDLMSSKDDVVK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	2118	1785.765	QEEEMMAKEEELVK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	2144	1814.828	LTEMETLQSQLMAEK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	2146	1814.901	IAQLEEQLDNETKER		1	Proteotypic
MYH9_HUMAN	2184	1868.959	ANLQIDQINTDLNLER		1	Proteotypic
MYH9_HUMAN	2232	1945.000	LQVELDNVTGLLSQSDDSK		1	Proteotypic
MYH9_HUMAN	2278	2049.045	IIGLDQVAGMSETALPGAFK	Dioxidation (M)	1	Proteotypic
MYH9_HUMAN	2332	2302.129	IRELESQISELQEDLESER		1	Proteotypic
MYH9_HUMAN	2353	2471.166	IAQLEEELEEEQGNTELINDR		1	Proteotypic
MYH9_HUMAN	2354	2492.167	DFSALESQIQDTQELLQEENR		1	Proteotypic
MYL6_HUMAN	403	1024.498	EAFQLFDR		1	Proteotypic
MYL6_HUMAN	1531	1353.725	ALGQNPTNAEVLK		1	Proteotypic
NDUAC_HUMAN	2436	1188.654	MELVQVLKR	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
NFH_HUMAN	775	1120.576	EYQDLLNVK		2	Shared
NFH_HUMAN	785	1123.518	KLLEGECCR	Trioxidation (C)	6	Shared
NH2L1_HUMAN	58	914.435	TEADVNP	Acetyl (Protein N-term)	1	Proteotypic
NH2L1_HUMAN	1725	1456.763	QQIQSIQSIER		1	Proteotypic
NICA_HUMAN	2415	1084.624	ALADVATVLGR		1	Proteotypic
NLTP_HUMAN	680	1098.505	ITGNMGLAMK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
NLTP_HUMAN	1344	1285.576	MNPQSAFFQGK	Dioxidation (M)	1	Proteotypic
NNTM_HUMAN	82	922.560	SAPLLLPGR		1	Proteotypic
NNTM_HUMAN	129	941.529	VAGAQQGAK		1	Proteotypic
NNTM_HUMAN	402	1024.486	KTCDALQAK	Trioxidation (C)	1	Proteotypic
NNTM_HUMAN	569	1071.618	EVLASDLVVK		1	Proteotypic
NNTM_HUMAN	849	1139.598	FFTGQITAAGK		1	Proteotypic
NNTM_HUMAN	910	1153.636	AVVLAANHFGR		1	Proteotypic

NNTM_HUMAN	1280	1261.634	TTVLAMDQVPR	Dioxidation (M)	1	Proteotypic
NNTM_HUMAN	1304	1269.682	SLGAEPLEVDLK		1	Proteotypic
NNTM_HUMAN	1373	1294.761	VALSPAGVQNLVK		1	Proteotypic
NNTM_HUMAN	1832	1523.904	ILIVGGGVAGLASAGAAK		1	Proteotypic
NNTM_HUMAN	1906	1577.769	QGFNVVVEGAGEASK		1	Proteotypic
NQO1_HUMAN	2420	1107.640	ALIVLAHSER		1	Proteotypic
NUCL_HUMAN	120	936.492	TGISDVFAK		1	Proteotypic
NUCL_HUMAN	315	999.535	NDLAVVDVR		1	Proteotypic
NUCL_HUMAN	983	1177.561	EVFEDAAEIR		1	Proteotypic
NUCL_HUMAN	1036	1191.610	IVTDRETGSSK		1	Proteotypic
NUCL_HUMAN	1612	1396.636	TEADAECTFEK		1	Proteotypic
NUCL_HUMAN	1884	1560.673	GFGFVDFNSEEDAK		1	Proteotypic
NUCL_HUMAN	2253	1990.985	VTQDELKEVFEDAAEIR		1	Proteotypic
O60744_HUMAN	34	907.440	VGEFSGANK		1	Proteotypic
O60744_HUMAN	1489	1335.631	TAFQEALDAAGDK		1	Proteotypic
ODO2_HUMAN	1025	1188.650	VEGGTPLFTLR		1	Proteotypic
ODO2_HUMAN	1691	1439.635	NVEAMNFADIER	Dioxidation (M)	1	Proteotypic
OST48_HUMAN	486	1045.592	SSLNPILFR		1	Proteotypic
OST48_HUMAN	1302	1268.745	TLVLLDNLNVR		1	Proteotypic
PA1B2_HUMAN	2398	992.675	IIVLGLLPR		1	Proteotypic
PABP1_HUMAN	476	1044.503	GFGFVSFER		1	Proteotypic
PABP1_HUMAN	919	1156.660	FSPAGPILSIR		1	Proteotypic
PABP1_HUMAN	1378	1297.622	ALDTMNFVVIK	Dioxidation (M)	1	Proteotypic
PABP2_HUMAN	2448	1239.632	AAAAAAAAAAGAAGGR	Acetyl (Protein N-term)	1	Proteotypic
PAIRB_HUMAN	358	1012.494	EAGGGVGGPGAK		1	Proteotypic
PAIRB_HUMAN	1259	1254.632	RPDQQLQGEGK		1	Proteotypic
PAIRB_HUMAN	1730	1459.701	SAAQAAAQTSNAAGK		1	Proteotypic
PARK7_HUMAN	923	1157.604	EILKEQENR		1	Proteotypic
PARK7_HUMAN	1984	1647.825	VTVAGLAGKDPVQCSR	Trioxidation (C)	1	Proteotypic
PARK7_HUMAN	1985	1649.782	DVVICPDASLEDAKK	Trioxidation (C)	1	Proteotypic
PARK7_HUMAN	2081	1738.775	GAEEMETVIPVDVMR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
PARK7_HUMAN	2361	2583.318	EGPYDVVVLPGGNLGAQNLSESAVK		1	Proteotypic
PARP1_HUMAN	2482	1623.799	VVSEDFLQDVSASTK		1	Proteotypic
PCBP1_HUMAN	364	1013.525	QGANINEIR		1	Proteotypic
PCBP1_HUMAN	629	1085.547	IANPVEGSSGR		1	Proteotypic
PCBP1_HUMAN	1328	1278.551	INISEGNCPER	Trioxidation (C)	2	Shared
PCBP1_HUMAN	1597	1387.807	IITLTGPTNAIFK		1	Proteotypic
PCBP1_HUMAN	1678	1432.760	LVVPATQCGSLIGK	Trioxidation (C)	1	Proteotypic
PCBP1_HUMAN	2293	2120.964	ESTGAQVQVAGDMLPNSTER	Dioxidation (M)	2	Shared
PCYOX_HUMAN	2465	1383.653	FLNEMIAPVMR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
PDIA1_HUMAN	38	909.434	FFPASADR		1	Proteotypic
PDIA1_HUMAN	161	952.407	ITEFCHR	Trioxidation (C)	1	Proteotypic
PDIA1_HUMAN	200	965.559	ILEFFGLK		1	Proteotypic
PDIA1_HUMAN	240	978.444	KEECPAVR	Trioxidation (C)	1	Proteotypic
PDIA1_HUMAN	287	990.539	ENLLDFIK		1	Proteotypic
PDIA1_HUMAN	326	1001.551	LKAEGSEIR		1	Proteotypic



PDIA1_HUMAN	446	1036.483	NNFEDEVTK		1	Proteotypic
PDIA1_HUMAN	549	1065.509	TVIDYNGER		1	Proteotypic
PDIA1_HUMAN	1105	1212.530	NFEDVAFDEK		1	Proteotypic
PDIA1_HUMAN	1202	1237.611	LITLEEEMTK	Dioxidation (M)	1	Proteotypic
PDIA1_HUMAN	1416	1308.587	MDSTANEVEAVK	Oxidation (M)	1	Proteotypic
PDIA1_HUMAN	1466	1324.582	MDSTANEVEAVK	Dioxidation (M)	1	Proteotypic
PDIA1_HUMAN	1499	1340.625	NFEDVAFDEKK		1	Proteotypic
PDIA1_HUMAN	1661	1423.771	YQLDKDGVVLFK		1	Proteotypic
PDIA1_HUMAN	1719	1450.694	YKPESEELTAER		1	Proteotypic
PDIA1_HUMAN	2112	1779.828	VDATEESDLAQQYGVV		1	Proteotypic
PDIA1_HUMAN	2377	2934.486	TGPAATLPLDGAAAESLVESSEVAVIGFFK		1	Proteotypic
PDIA3_HUMAN	301	994.556	QAGPASVPLR		1	Proteotypic
PDIA3_HUMAN	1029	1190.593	LAPEYEEAATR		1	Proteotypic
PDIA3_HUMAN	1050	1196.713	LSKDPNIVIAK		1	Proteotypic
PDIA3_HUMAN	1079	1203.523	FVMQEEFSR	Dioxidation (M)	1	Proteotypic
PDIA3_HUMAN	1193	1235.505	DGEEAGAYDGPR		1	Proteotypic
PDIA3_HUMAN	1542	1358.651	FLQDYFDGNLK		1	Proteotypic
PDIA3_HUMAN	1561	1367.657	SEPIPESNDGPPK		1	Proteotypic
PDIA3_HUMAN	1567	1369.688	ELSDFISYLQR		1	Proteotypic
PDIA3_HUMAN	1907	1578.825	EATNPPVIQEEKPK		1	Proteotypic
PDIA3_HUMAN	2048	1695.741	MDATANDVPSPYEVR	Dioxidation (M)	1	Proteotypic
PDIA3_HUMAN	2161	1831.895	VVVAENFDEIVNNENK		1	Proteotypic
PDIA6_HUMAN	373	1014.607	AATALKDVVK		1	Proteotypic
PDIA6_HUMAN	895	1150.489	GESPVDYDGGR		1	Proteotypic
PDIA6_HUMAN	1596	1385.751	TGEAIVDAALSALR		1	Proteotypic
PDIA6_HUMAN	1778	1482.710	GSFSEQGINEFLR		1	Proteotypic
PDIA6_HUMAN	1835	1526.842	LAAVDATVNQVLASR		1	Proteotypic
PDIA6_HUMAN	1942	1614.837	GSTAPVGGAFPTIVER		1	Proteotypic
PEBP1_HUMAN	1883	1559.820	LYTLVLTDPDAPSR		1	Proteotypic
PEBP1_HUMAN	2235	1948.938	GNDISSGTVLSDYVGSPPK		1	Proteotypic
PGM1_HUMAN	648	1089.578	LSGTGSAGATIR		1	Proteotypic
PGM1_HUMAN	1067	1200.610	QEATLVVGGDGR		1	Proteotypic
PHB2_HUMAN	4	898.560	AQVSLLR		1	Proteotypic
PHB2_HUMAN	245	980.481	EYTAHAVEAK		1	Proteotypic
PHB2_HUMAN	295	993.477	LGLDYEER		1	Proteotypic
PHB2_HUMAN	507	1056.531	QVAQQAQR		1	Proteotypic
PHB2_HUMAN	978	1176.625	FNASQLITQR		1	Proteotypic
PHB2_HUMAN	1100	1209.733	VLPSIVNEVLK		1	Proteotypic
PHB2_HUMAN	1113	1214.614	IVQAEGEAEAAK		1	Proteotypic
PHB2_HUMAN	1123	1216.584	ESVFTVEGGHR		1	Proteotypic
PHB2_HUMAN	1131	1219.623	DLQMVNISLR	Dioxidation (M)	1	Proteotypic
PHB2_HUMAN	1269	1258.740	LLLGAGAVAYGVR		1	Proteotypic
PHB2_HUMAN	2177	1852.980	IGGVQQDTILAEGLHFR		1	Proteotypic
PHB2_HUMAN	2216	1919.952	VLSRPNAQELPSMYQR	Dioxidation (M)	1	Proteotypic
PIGR_HUMAN	2479	1538.644	ASVDSGSSEEQGGSSR		1	Proteotypic
PLEC1_HUMAN	15	900.503	LAAIGEATR		1	Proteotypic

PLEC1_HUMAN	17	901.487	LTVDEAVR		1	Proteotypic
PLEC1_HUMAN	63	914.519	GLLSAEVAR		1	Proteotypic
PLEC1_HUMAN	68	915.503	LTVEEAVR		1	Proteotypic
PLEC1_HUMAN	225	974.503	DLSELGSVR		1	Proteotypic
PLEC1_HUMAN	252	983.504	VPDVQDGVR		1	Proteotypic
PLEC1_HUMAN	298	994.464	VQMEELSK	Dioxidation (M)	1	Proteotypic
PLEC1_HUMAN	319	1000.519	AQAEVEGLGK		1	Proteotypic
PLEC1_HUMAN	324	1001.478	QLAEEDAAR		1	Proteotypic
PLEC1_HUMAN	367	1014.546	LSVAAQEAAAR		1	Proteotypic
PLEC1_HUMAN	417	1028.550	AELELELGR		1	Proteotypic
PLEC1_HUMAN	461	1041.582	ALQALEELR		1	Proteotypic
PLEC1_HUMAN	462	1041.582	LAAEQELIR		1	Proteotypic
PLEC1_HUMAN	481	1045.504	SLAAEEEAAR		1	Proteotypic
PLEC1_HUMAN	519	1058.536	GTQGAEVLR		1	Proteotypic
PLEC1_HUMAN	521	1058.572	LKAEATEAAR		1	Proteotypic
PLEC1_HUMAN	523	1059.508	VLADPSDDTK		1	Proteotypic
PLEC1_HUMAN	541	1064.492	MQAVQEATR	Dioxidation (M)	1	Proteotypic
PLEC1_HUMAN	567	1071.581	LAEVEAALEK		1	Proteotypic
PLEC1_HUMAN	647	1089.530	QSSEAEIQAK		1	Proteotypic
PLEC1_HUMAN	685	1099.562	GGAEGELQALR		1	Proteotypic
PLEC1_HUMAN	686	1099.587	QLLEEEELAR		1	Proteotypic
PLEC1_HUMAN	690	1100.557	LQAEAAERR		1	Proteotypic
PLEC1_HUMAN	691	1100.557	QLAEGTAQQR		1	Proteotypic
PLEC1_HUMAN	730	1109.608	AQLEPVASPAK		1	Proteotypic
PLEC1_HUMAN	800	1128.589	NLVDNITGQR		1	Proteotypic
PLEC1_HUMAN	858	1142.593	LQAEVVAQQK		1	Proteotypic
PLEC1_HUMAN	937	1163.582	LTAEDLFEAR		1	Proteotypic
PLEC1_HUMAN	1027	1189.671	LLFNDVQTLK		1	Proteotypic
PLEC1_HUMAN	1030	1190.604	FRELAEEAAR		1	Proteotypic
PLEC1_HUMAN	1044	1194.519	QLEMSAEAER	Dioxidation (M)	1	Proteotypic
PLEC1_HUMAN	1083	1204.619	EGLTSIEEVTK		1	Proteotypic
PLEC1_HUMAN	1114	1214.625	RPELEDSTLR		1	Proteotypic
PLEC1_HUMAN	1160	1226.662	QLQLAQEAAQK		1	Proteotypic
PLEC1_HUMAN	1169	1230.620	SKEQAELEAAR		1	Proteotypic
PLEC1_HUMAN	1194	1235.520	QQQQMEQER	Dioxidation (M)	1	Proteotypic
PLEC1_HUMAN	1204	1238.571	AQLMADFQAGR	Dioxidation (M)	1	Proteotypic
PLEC1_HUMAN	1217	1241.673	QVQVALETAQR		1	Proteotypic
PLEC1_HUMAN	1339	1283.708	SLVPAAELLESR		1	Proteotypic
PLEC1_HUMAN	1348	1286.683	AQVEQELTTLR		1	Proteotypic
PLEC1_HUMAN	1386	1298.756	VTLVQTLQIQR		1	Proteotypic
PLEC1_HUMAN	1388	1299.642	QLAEEDLAQQR		1	Proteotypic
PLEC1_HUMAN	1457	1321.651	SQVEEELFSVR		1	Proteotypic
PLEC1_HUMAN	1668	1427.787	IIITVVEEQEQK		1	Proteotypic
PLEC1_HUMAN	1796	1492.719	SQVMDEATALQLR	Dioxidation (M)	1	Proteotypic
PLEC1_HUMAN	1842	1530.877	VLALPEPSPAAPTLR		1	Proteotypic
PLEC1_HUMAN	1892	1565.878	APVPASELLASGVLSR		1	Proteotypic

PNPH_HUMAN	2426	1124.623	VFGFSLITNK		1	Proteotypic
PP1B_HUMAN	2439	1206.591	IVQMTEAEVR	Dioxidation (M)	1	Proteotypic
PRDX1_HUMAN	77	919.501	GLFIIDDK		2	Shared
PRDX1_HUMAN	719	1106.597	TIAQDYGVLK		1	Proteotypic
PRDX1_HUMAN	1046	1195.554	ATAVMPDGGQFK	Dioxidation (M)	1	Proteotypic
PRDX1_HUMAN	1047	1195.624	LVQAFQFTDK		1	Proteotypic
PRDX1_HUMAN	1103	1210.667	QITVNDLPVGR		2	Shared
PRDX1_HUMAN	1545	1358.792	GLFIIDDKGILR		2	Shared
PRDX1_HUMAN	1995	1653.840	QGGLGPMNIPLVSDPK	Dioxidation (M)	1	Proteotypic
PRDX2_HUMAN	83	923.435	TDEGIAYR		1	Proteotypic
PRDX2_HUMAN	238	977.518	ATAVVDGAFK		1	Proteotypic
PRDX2_HUMAN	1103	1210.667	QITVNDLPVGR		2	Shared
PRDX2_HUMAN	2076	1733.968	EGGLGPLNIPLLDVTR		1	Proteotypic
PRDX2_HUMAN	2181	1862.063	KEGGLGPLNIPLLDVTR		1	Proteotypic
PRDX3_HUMAN	159	951.455	DLSLDDFK		1	Proteotypic
PRDX3_HUMAN	391	1020.524	GTAVVNGEFK		1	Proteotypic
PRDX3_HUMAN	1086	1205.652	HLSVNDLPVGR		1	Proteotypic
PRDX3_HUMAN	1342	1284.744	GLFIIDPNGVIK		1	Proteotypic
PRDX3_HUMAN	1734	1461.783	DYGVLLLEGSLALR		1	Proteotypic
PRDX4_HUMAN	77	919.501	GLFIIDDK		2	Shared
PRDX4_HUMAN	1153	1224.683	QITLNDLPVGR		1	Proteotypic
PRDX4_HUMAN	1545	1358.792	GLFIIDDKGILR		2	Shared
PRKDC_HUMAN	61	914.482	LPSNTLDR		1	Proteotypic
PRKDC_HUMAN	355	1011.587	FVPLLPGNR		1	Proteotypic
PRKDC_HUMAN	500	1051.555	GIFTSEIGTK		1	Proteotypic
PRKDC_HUMAN	972	1174.619	VTELALTASDR		1	Proteotypic
PRKDC_HUMAN	1391	1299.715	QITQSALLAEAR		1	Proteotypic
PRKDC_HUMAN	1530	1353.660	MSTSPEAFLAR	Dioxidation (M)	1	Proteotypic
PRKDC_HUMAN	1747	1466.700	SLGPPQGEEDSVPR		1	Proteotypic
PRKDC_HUMAN	1765	1474.781	VVQMLGSLGGQINK	Dioxidation (M)	1	Proteotypic
PRKDC_HUMAN	1843	1531.675	NLLTVTSSDEMMK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
PROF1_HUMAN	31	905.526	TLVLLMGK	Dioxidation (M)	1	Proteotypic
PROF1_HUMAN	899	1150.646	EGVHGGLINKK		1	Proteotypic
PROF1_HUMAN	1021	1188.454	CYEMASHLR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
PROF1_HUMAN	1245	1249.695	TDKTLVLLMGK	Dioxidation (M)	1	Proteotypic
PROF1_HUMAN	1579	1378.709	STGGAPTFNVTVK		1	Proteotypic
PROF1_HUMAN	1975	1642.929	TFVNITPAEVGVLVGK		1	Proteotypic
PROF1_HUMAN	1998	1656.730	DSLLODGEFSMDLR	Dioxidation (M)	1	Proteotypic
PROF1_HUMAN	2211	1914.057	TFVNITPAEVGVLVGKDR		1	Proteotypic
PSA_HUMAN	64	914.519	QILSADLR		1	Proteotypic
PSA_HUMAN	805	1128.675	AGIISTVEVLK		1	Proteotypic
PSA_HUMAN	1282	1262.629	VALSNMNVDR	Dioxidation (M)	1	Proteotypic
PSA_HUMAN	1512	1345.735	LGLQNDLFSLAR		1	Proteotypic
PSA_HUMAN	1582	1379.839	VLGATLLPDLIQK		1	Proteotypic
PSA3_HUMAN	2444	1216.641	AVENSSTAIGIR		1	Proteotypic
PSA4_HUMAN	18	901.523	VEIATLTR		1	Proteotypic

PSA4_HUMAN	1155	1225.623	LLDEVFFSEK		1	Proteotypic
PSA5_HUMAN	534	1062.509	GVNTFSPEGR		1	Proteotypic
PSA5_HUMAN	1660	1422.776	LFQVEYAIEAIK		1	Proteotypic
PSA5_HUMAN	1737	1462.722	ITSPLMEPSSIEK	Dioxidation (M)	1	Proteotypic
PSA6_HUMAN	1094	1208.607	GKDCAVIVTQK	Trioxidation (C)	1	Proteotypic
PSA6_HUMAN	1341	1284.715	AINQGGLTSAVAVR		1	Proteotypic
PSA7_HUMAN	236	976.501	NIELAVMR	Dioxidation (M)	1	Proteotypic
PSA7_HUMAN	688	1099.624	ALLEVVQSGGK		1	Proteotypic
PSB1_HUMAN	341	1006.508	DVFISAAER		1	Proteotypic
PSB1_HUMAN	1351	1287.678	EGIREETVSLR		1	Proteotypic
PSB2_HUMAN	662	1091.564	VAASNIVQMK	Dioxidation (M)	1	Proteotypic
PSB2_HUMAN	1411	1305.745	FILNLPPTFSVR		1	Proteotypic
PSB3_HUMAN	1976	1644.782	FGIQAQMVTTDFQK	Dioxidation (M)	1	Proteotypic
PSB3_HUMAN	2195	1888.042	LYIGLAGLATDVQTVAQR		1	Proteotypic
PSB4_HUMAN	442	1035.560	FQIATVTEK		1	Proteotypic
PSB4_HUMAN	796	1127.593	QPVLSQTEAR		1	Proteotypic
PSB6_HUMAN	732	1109.644	QVLLGDQIPK		1	Proteotypic
PSB6_HUMAN	749	1114.598	LAAIAESGVER		1	Proteotypic
PSMD2_HUMAN	2432	1166.532	FGGSGSQVDSAR		1	Proteotypic
PTBP1_HUMAN	189	962.547	VTNLLMLK	Dioxidation (M)	1	Proteotypic
PTBP1_HUMAN	223	974.442	TDSSPNQAR		1	Proteotypic
PTBP1_HUMAN	713	1105.613	VLFSNNGGVVK		1	Proteotypic
PTBP1_HUMAN	2326	2274.270	IAIPGLAGAGNSVLLVSNLNPER		1	Proteotypic
PTMA_HUMAN	815	1130.520	EVVEEAENGR		1	Proteotypic
PTMA_HUMAN	1744	1465.678	SDAAVDTSSSEITTK	Acetyl (Protein N-term)	1	Proteotypic
PTMS_HUMAN	589	1074.556	SVEAAAELSAK		1	Proteotypic
PTMS_HUMAN	1541	1358.595	AAEEEEDEADPKR		1	Proteotypic
Q05DE8_HUMAN	2403	1049.510	AASLGSSQSSR		1	Proteotypic
Q06AH7_HUMAN	195	963.525	APNHAVVTR		1	Proteotypic
Q06AH7_HUMAN	237	977.482	DGAGDVAFVK		1	Proteotypic
Q06AH7_HUMAN	946	1165.584	HQTVPQNTGGK		1	Proteotypic
Q06AH7_HUMAN	1158	1226.535	DSGFQMNQLR	Dioxidation (M)	1	Proteotypic
Q0QD40_HUMAN	2443	1213.703	VVVVTGANTGIGK		1	Proteotypic
Q0V924_HUMAN	146	946.560	KGDIFLVR		1	Proteotypic
Q0V924_HUMAN	495	1048.544	DVDLEFLAK		1	Proteotypic
Q0V924_HUMAN	615	1082.507	MDELQLFR	Dioxidation (M)	1	Proteotypic
Q0V924_HUMAN	751	1115.496	GGNIGDGGGAADR		1	Proteotypic
Q0V924_HUMAN	1214	1241.625	EVDIGIPDATGR		1	Proteotypic
Q0V924_HUMAN	1736	1461.811	IVSLLLTLMDGLK	Dioxidation (M)	1	Proteotypic
Q0V924_HUMAN	2137	1809.988	NAPAIIFIDELDAIAPK		1	Proteotypic
Q13382_HUMAN	2389	928.509	KSXPSAVAR		1	Proteotypic
Q13727_HUMAN	11	900.455	ADLDVSGPK		1	Proteotypic
Q13727_HUMAN	92	925.388	MPEMNIK	Dioxidation (M); Dioxidation (M)	2	Shared
Q13727_HUMAN	915	1155.650	VKGDVDVSLPK		2	Shared
Q14C98_HUMAN	856	1141.609	AQADEVLQLR		1	Proteotypic
Q14C98_HUMAN	924	1157.615	ENLQLETRR		1	Proteotypic

Q1WM23_HUMAN	256	983.613	GLVGEIHKR		1	Proteotypic
Q1WM23_HUMAN	460	1041.455	GDFCIQVGR	Trioxidation (C)	1	Proteotypic
Q1WM23_HUMAN	556	1068.556	NIIHGSDSVK		1	Proteotypic
Q1WM23_HUMAN	886	1148.634	DRPFFAGLVK		1	Proteotypic
Q1WM23_HUMAN	973	1174.650	DRPFFPGLVK		1	Proteotypic
Q1WM23_HUMAN	1106	1212.570	FMQASEDLLK	Dioxidation (M)	1	Proteotypic
Q1WM23_HUMAN	1511	1343.756	TFIAIKPDGVQR		1	Proteotypic
Q1WM23_HUMAN	2150	1816.899	VMLGETNPADSKPGTIR	Dioxidation (M)	1	Proteotypic
Q1ZYL5_HUMAN	269	987.499	AELSEGQVR		1	Proteotypic
Q1ZYL5_HUMAN	383	1017.509	AGSSSLEAVR	Acetyl (Protein N-term)	1	Proteotypic
Q1ZYL5_HUMAN	1222	1242.646	IQLVEEELDR		3	Shared
Q1ZYL5_HUMAN	1767	1475.674	LEEAEKAADESER		3	Shared
Q1ZYL5_HUMAN	2071	1726.885	IQLVEEELDRAQER		3	Shared
Q2NKY5_HUMAN	592	1076.525	IREEYPDR		4	Shared
Q2NKY5_HUMAN	769	1119.363	NMMAACDPR	Dioxidation (M); Dioxidation (M); Trioxidation (C)	5	Shared
Q2NKY5_HUMAN	808	1129.588	FPGQLNADLR		5	Shared
Q2NKY5_HUMAN	928	1158.622	LAVNMVFPFR	Oxidation (M)	5	Shared
Q2NKY5_HUMAN	971	1174.617	LAVNMVFPFR	Dioxidation (M)	5	Shared
Q2NKY5_HUMAN	1264	1257.683	FPGQLNADLRK		5	Shared
Q2NKY5_HUMAN	1398	1302.712	KLAVNMVFPFR	Dioxidation (M)	5	Shared
Q2NKY5_HUMAN	1641	1414.647	IMNTFSVMPSPK	Dioxidation (M); Dioxidation (M)	2	Shared
Q2NKY5_HUMAN	2068	1722.850	ALTVPELTQQMFDAK	Dioxidation (M)	3	Shared
Q3KP43_HUMAN	163	952.596	LLLPGELAK		2	Shared
Q3KP43_HUMAN	2079	1735.755	MGIMNSFVNDIFER	Dioxidation (M); Dioxidation (M)	1	Proteotypic
Q3KQZ2_HUMAN	425	1029.545	DVLVGADSVR		1	Proteotypic
Q3KQZ2_HUMAN	926	1157.656	FLTQPQVVAR		1	Proteotypic
Q3ZCW5_HUMAN	1120	1215.610	LEGTNVQEAQK		1	Proteotypic
Q3ZCW5_HUMAN	1207	1239.552	INFDDNAEFR		1	Proteotypic
Q3ZCW5_HUMAN	1413	1306.659	MAENLGFVGPLK	Dioxidation (M)	1	Proteotypic
Q4TWB7_HUMAN	1316	1273.718	LLVVYPWTXR		1	Proteotypic
Q4TWB7_HUMAN	1431	1313.658	VNVDEVGGEALGR		2	Shared
Q4TWB7_HUMAN	1751	1468.650	GTFATLSELHCDK	Trioxidation (C)	2	Shared
Q4TWB7_HUMAN	2020	1668.884	VLGAFSDGLAHLNLIK		2	Shared
Q4TWB7_HUMAN	2286	2073.935	FFESFGDLSTPDVAVMGNPK	Oxidation (M)	2	Shared
Q4TWB7_HUMAN	2288	2089.930	FFESFGDLSTPDVAVMGNPK	Dioxidation (M)	2	Shared
Q4VXC1_HUMAN	869	1145.593	DLAEITLDR		1	Proteotypic
Q4VXC1_HUMAN	1679	1433.584	VDCDQHSDIAQR	Trioxidation (C)	1	Proteotypic
Q4ZJ1_HUMAN	413	1027.603	GELLEAIKR		1	Proteotypic
Q4ZJ1_HUMAN	1662	1423.803	GDVTAQIALQPALK		1	Proteotypic
Q53EP4_HUMAN	243	979.570	LPVALDPGAK		1	Proteotypic
Q53EP4_HUMAN	518	1058.536	QPDSGISSIR		1	Proteotypic
Q53EP4_HUMAN	1394	1300.735	ALTSEIALLQSR		1	Proteotypic
Q53EP4_HUMAN	1677	1431.611	GEDEEENLEVR		1	Proteotypic
Q53EP4_HUMAN	1993	1652.885	VTAEVVLAHLGGGSTR		1	Proteotypic
Q53EP4_HUMAN	2007	1658.888	ATSFLLALEPELEAR		1	Proteotypic
Q53EW8_HUMAN	1720	1451.714	TYEQVLENLESK		1	Proteotypic

Q53EW8_HUMAN	1793	1490.776	KVDLSQPLIATCR	Trioxidation (C)	1	Proteotypic
Q53EW8_HUMAN	2261	2008.985	FLGTEPEPDAVGLDSGHIR		1	Proteotypic
Q53F27_HUMAN	2492	1752.915	FMPAGLIAGASLLMVAK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
Q53F91_HUMAN	84	923.438	MVADTISR	Dioxidation (M)	1	Proteotypic
Q53F91_HUMAN	731	1109.633	DPETPIIVVK		1	Proteotypic
Q53F91_HUMAN	1023	1188.573	TNNLETGPSTR		1	Proteotypic
Q53F91_HUMAN	1286	1264.568	EVQGNESAEFR		1	Proteotypic
Q53F91_HUMAN	1420	1308.729	AAVPDTVVEPALK		1	Proteotypic
Q53FN0_HUMAN	2496	1814.693	DAEDAMDAMDGAVALDGR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
Q53FR4_HUMAN	257	983.613	ILVGTNLVR		1	Proteotypic
Q53FR4_HUMAN	352	1010.576	AELAEPLR		1	Proteotypic
Q53FR4_HUMAN	1223	1242.657	LSQLEGVNVER		1	Proteotypic
Q53FS6_HUMAN	2478	1512.778	GVVDESDIPLNLSR		1	Proteotypic
Q53G21_HUMAN	2471	1457.747	NAIANASTLAEVER		1	Proteotypic
Q53G25_HUMAN	254	983.540	QAVDVSPLR		1	Proteotypic
Q53G25_HUMAN	1314	1272.490	LTNSMMMHR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	1	Proteotypic
Q53G25_HUMAN	1952	1621.787	TIAECLADELINAAK	Trioxidation (C)	1	Proteotypic
Q53G92_HUMAN	769	1119.363	NMMAACDPR	Dioxidation (M); Dioxidation (M); Trioxidation (C)	5	Shared
Q53G92_HUMAN	808	1129.588	FPGQLNADLR		5	Shared
Q53G92_HUMAN	928	1158.622	LAVNMVFPFR	Oxidation (M)	5	Shared
Q53G92_HUMAN	971	1174.617	LAVNMVFPFR	Dioxidation (M)	5	Shared
Q53G92_HUMAN	1264	1257.683	FPGQLNADLRK		5	Shared
Q53G92_HUMAN	1275	1260.581	ISEQFTAMFR	Dioxidation (M)	4	Shared
Q53G92_HUMAN	1398	1302.712	KLAVNMVFPFR	Dioxidation (M)	5	Shared
Q53G92_HUMAN	1522	1350.685	IMNTFSVVPSPK	Dioxidation (M)	3	Shared
Q53G92_HUMAN	1980	1646.819	AILVDLEPGTMDSVR	Dioxidation (M)	3	Shared
Q53G92_HUMAN	2068	1722.850	ALTVPELTQQMFDK	Dioxidation (M)	3	Shared
Q53G92_HUMAN	2196	1888.924	MSSTFIGNSTAIQELFK	Oxidation (M)	1	Proteotypic
Q53GF9_HUMAN	674	1095.665	LKPLEVELR		1	Proteotypic
Q53GF9_HUMAN	1440	1315.706	IPDQLVILDMK	Dioxidation (M)	1	Proteotypic
Q53GR3_HUMAN	811	1129.609	VVQSLEQTAR		1	Proteotypic
Q53GR3_HUMAN	1265	1257.704	KVVQSLEQTAR		1	Proteotypic
Q53GR3_HUMAN	2050	1697.884	EAVTEILGIEPDREK		1	Proteotypic
Q53H42_HUMAN	2494	1764.926	EGEAVVLPVEPGLTAR		1	Proteotypic
Q53HE2_HUMAN	1183	1233.595	SNVSDAVAQSTR		1	Proteotypic
Q53HE2_HUMAN	1445	1316.628	IYGGSVTGATCK	Trioxidation (C)	1	Proteotypic
Q53HE2_HUMAN	1640	1413.783	QSLGELIGTLNAAK		1	Proteotypic
Q53HE2_HUMAN	1728	1457.715	HVFGESDELIGQK		1	Proteotypic
Q53HE2_HUMAN	1745	1465.716	TATPQQAQEVHEK		1	Proteotypic
Q53HE2_HUMAN	1864	1541.878	KQSLGELIGTLNAAK		1	Proteotypic
Q53HE2_HUMAN	1992	1652.808	VTNGAFTGEISPGMIK	Dioxidation (M)	1	Proteotypic
Q53HE2_HUMAN	2129	1797.929	VAHALAEGLGVIACIGEK	Trioxidation (C)	1	Proteotypic
Q53HE2_HUMAN	2315	2182.025	VPADTEVVCAPPTAYIDFAR	Trioxidation (C)	1	Proteotypic
Q53HE2_HUMAN	2379	3028.576	ELASQPDVDGFLVGGASLKPFEVDIINAK		1	Proteotypic
Q53HU0_HUMAN	798	1127.666	LATNAAVTVLR		1	Proteotypic
Q53HU0_HUMAN	891	1149.582	FAEAFEIPR		1	Proteotypic

Q53HU0_HUMAN	1573	1371.736	AIADTGANVVVTGGK		1	Proteotypic
Q53HV1_HUMAN	284	989.591	LSNIFVIGK		1	Proteotypic
Q53HV1_HUMAN	498	1051.468	DANGNSFATR		1	Proteotypic
Q53HV1_HUMAN	1848	1532.875	LRECLPLIIFLR	Trioxidation (C)	1	Proteotypic
Q53HW7_HUMAN	676	1096.580	VILAEFDYK		1	Proteotypic
Q53HW7_HUMAN	1507	1342.720	TAAAVAAQSGILDR		1	Proteotypic
Q53HW7_HUMAN	1714	1447.731	YADALQEIIQER		1	Proteotypic
Q53QE9_HUMAN	421	1028.591	EFPTVPLVK		1	Proteotypic
Q53QE9_HUMAN	1260	1255.677	GLPDNISSVLNK		1	Proteotypic
Q53QE9_HUMAN	1480	1331.683	SFENSLGINVPR		1	Proteotypic
Q53QU2_HUMAN	2454	1268.636	IEGDPQGVQQAK		1	Proteotypic
Q53T32_HUMAN	178	958.458	VFFDLMR	Dioxidation (M)	1	Proteotypic
Q53T32_HUMAN	438	1034.543	VIMVGSQGVGK	Dioxidation (M)	1	Proteotypic
Q53T32_HUMAN	616	1082.583	RQVPVEEAR		1	Proteotypic
Q56UN9_HUMAN	2416	1084.649	ILIIGESGVGK		1	Proteotypic
Q59FA2_HUMAN	72	915.529	LRVEFPR		1	Proteotypic
Q59FA2_HUMAN	596	1077.546	DGTGVVEFVR		1	Proteotypic
Q59FF0_HUMAN	537	1062.538	EGLVMVEVR	Dioxidation (M)	1	Proteotypic
Q59FF0_HUMAN	885	1148.619	LGTLSPAFSTR		1	Proteotypic
Q59FI9_HUMAN	1629	1408.650	CTGGEVGTASALAPK	Trioxidation (C)	1	Proteotypic
Q59FI9_HUMAN	2017	1665.966	QAQIEVVPSASALIK		1	Proteotypic
Q59G24_HUMAN	679	1098.465	DDNMFQIGK	Dioxidation (M)	1	Proteotypic
Q59G24_HUMAN	1273	1259.599	EQISDIDDAVR		1	Proteotypic
Q5JP53_HUMAN	452	1038.586	YLTVAAVFR		2	Shared
Q5JP53_HUMAN	592	1076.525	IREEYPDR		4	Shared
Q5JP53_HUMAN	769	1119.363	NMMAACDPR	Dioxidation (M); Dioxidation (M); Trioxidation (C)	5	Shared
Q5JP53_HUMAN	808	1129.588	FPGQLNADLR		5	Shared
Q5JP53_HUMAN	928	1158.622	LAVNMVFPFR	Oxidation (M)	5	Shared
Q5JP53_HUMAN	971	1174.617	LAVNMVFPFR	Dioxidation (M)	5	Shared
Q5JP53_HUMAN	1264	1257.683	FPGQLNADLRK		5	Shared
Q5JP53_HUMAN	1275	1260.581	ISEQFTAMFR	Dioxidation (M)	4	Shared
Q5JP53_HUMAN	1392	1300.630	ISVYYNEATGGK		1	Proteotypic
Q5JP53_HUMAN	1398	1302.712	KLAVNMVFPFR	Dioxidation (M)	5	Shared
Q5JP53_HUMAN	1522	1350.685	IMNTFSVVPSPK	Dioxidation (M)	3	Shared
Q5JP53_HUMAN	1769	1477.672	EVDEQMLNVQNK	Dioxidation (M)	3	Shared
Q5JP53_HUMAN	1980	1646.819	AILVDLEPGTMDSVR	Dioxidation (M)	3	Shared
Q5JP53_HUMAN	2008	1658.888	ALTVPELTQQVFDK		1	Proteotypic
Q5JP53_HUMAN	2203	1900.960	MAVTFIGNSTAIQELFK	Dioxidation (M)	1	Proteotypic
Q5JP53_HUMAN	2267	2018.859	MSMKEVDEQMLNVQNK	Dioxidation (M); Dioxidation (M); Dioxidation (M)	3	Shared
Q5JR94_HUMAN	2476	1505.773	ISSLLEEQFQQGK		1	Proteotypic
Q5JRR8_HUMAN	697	1101.614	LKSDTAAAVR		1	Proteotypic
Q5JRR8_HUMAN	1226	1242.718	KPLLESGTLGTK		1	Proteotypic
Q5JRR8_HUMAN	1268	1258.725	LQTSSVLVSGLR		1	Proteotypic
Q5JRR8_HUMAN	1452	1319.639	LDQPMTEIVSR	Dioxidation (M)	1	Proteotypic
Q5JRR8_HUMAN	1954	1622.899	LAGTQPLEVLEAVQR		1	Proteotypic
Q5JRR8_HUMAN	2136	1807.943	ALPAVQQNNLDEDLIR		1	Proteotypic

Q5JRR8_HUMAN	2192	1882.938	NEEDAAELVALAQAVNAR		1	Proteotypic
Q5JRR8_HUMAN	2208	1907.988	SLVASLAEPDFVVTDFAK		1	Proteotypic
Q5JRR8_HUMAN	2224	1932.036	AAVATFLQSVQVPEFTPK		1	Proteotypic
Q5JVH5_HUMAN	2391	941.529	NIVEAAAVR		1	Proteotypic
Q5SQT9_HUMAN	873	1145.644	LVFLGLDNAGK		1	Proteotypic
Q5SQT9_HUMAN	2357	2528.341	VELNALMTDETISNPILILGNK	Dioxidation (M)	1	Proteotypic
Q5SUJ3_HUMAN	147	946.585	IAFAITAIK		1	Proteotypic
Q5SUJ3_HUMAN	321	1000.530	VLNTNIDGR		1	Proteotypic
Q5SUJ3_HUMAN	700	1102.580	VITIMQNPR	Dioxidation (M)	1	Proteotypic
Q5SUJ3_HUMAN	1236	1246.568	AGELTEDEVER		1	Proteotypic
Q5SZ71_HUMAN	600	1078.460	AELDDTPMR	Dioxidation (M)	1	Proteotypic
Q5SZ71_HUMAN	683	1098.615	AVVIVDDRGR		2	Shared
Q5SZ71_HUMAN	861	1142.620	FATHAAALSVR		1	Proteotypic
Q5SZ71_HUMAN	1235	1244.688	GIVEFASKPAAR		1	Proteotypic
Q5SZ71_HUMAN	1428	1312.586	RREEEMMIR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
Q5SZ71_HUMAN	1500	1340.659	FGQGGAGPVGGQGPR		1	Proteotypic
Q5SZ71_HUMAN	1968	1635.698	RMEELHNQEMQK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
Q5SZ71_HUMAN	2243	1963.005	LFVGNLPADITEDEFKR		1	Proteotypic
Q5TOP7_HUMAN	399	1022.470	MNDTVTIR	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
Q5TOP7_HUMAN	1615	1397.734	TTPDVIFVFGFR		1	Proteotypic
Q5T663_HUMAN	456	1039.545	FSIIGFSNR		1	Proteotypic
Q5T663_HUMAN	2036	1684.834	NVVFVLDSSASVMGK	Dioxidation (M)	1	Proteotypic
Q5U0D9_HUMAN	466	1042.566	AGLNSLEAVK	Acetyl (Protein N-term)	1	Proteotypic
Q5U0D9_HUMAN	1058	1198.667	AGLNSLEAVKR	Acetyl (Protein N-term)	1	Proteotypic
Q5U0D9_HUMAN	1222	1242.646	IQLVEEELDR		3	Shared
Q5U0D9_HUMAN	1767	1475.674	LEEAEKAADESER		3	Shared
Q5U0D9_HUMAN	2071	1726.885	IQLVEEELDRAQER		3	Shared
Q5U0I6_HUMAN	562	1070.634	LLLIGDSGVGK		3	Shared
Q5U0I6_HUMAN	594	1077.476	MGPATAGGAEK	Dioxidation (M)	1	Proteotypic
Q5U0I6_HUMAN	2069	1723.867	EFADSLGIPFLETSK		2	Shared
Q5U0I6_HUMAN	2234	1947.855	NATNVEQSFMTMAAEIK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
Q5VU58_HUMAN	474	1043.586	AGITTIEAVK	Acetyl (Protein N-term)	1	Proteotypic
Q5VU58_HUMAN	1062	1199.687	AGITTIEAVKR	Acetyl (Protein N-term)	1	Proteotypic
Q5VU58_HUMAN	1222	1242.646	IQLVEEELDR		3	Shared
Q5VU58_HUMAN	1266	1258.492	CREMDEQIR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
Q5VU58_HUMAN	1439	1315.637	EQAEAEVASLNR		1	Proteotypic
Q5VU58_HUMAN	1767	1475.674	LEEAEKAADESER		3	Shared
Q5VU58_HUMAN	2071	1726.885	IQLVEEELDRAQER		3	Shared
Q5VVD0_HUMAN	1766	1475.631	AQDQGEKENPMR	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
Q5VVD0_HUMAN	1870	1545.840	VLEQLTGQTPVFSK		1	Proteotypic
Q5VYA2_HUMAN	102	929.445	EVPTTEAR		1	Proteotypic
Q5VYA2_HUMAN	508	1056.618	VVLIGESGVGK		1	Proteotypic
Q5VYJ4_HUMAN	2464	1360.754	VMVQPINLIFR	Dioxidation (M)	1	Proteotypic
Q6B838_HUMAN	314	999.524	AEELVLER	Acetyl (Protein N-term)	1	Proteotypic
Q6B838_HUMAN	1364	1290.718	LVVNFESDKLK		1	Proteotypic
Q6B838_HUMAN	1670	1428.622	LAEKEETGMAMR	Dioxidation (M); Dioxidation (M)	1	Proteotypic



Q6FG96_HUMAN	2450	1255.677	NIEDVIAQGIGK		1	Proteotypic
Q6FGH5_HUMAN	2428	1153.492	MGESDDSIILR	Dioxidation (M)	1	Proteotypic
Q6FGI1_HUMAN	246	980.515	DVGRPQPGR		1	Proteotypic
Q6FGI1_HUMAN	411	1027.501	IQASTMAFK	Dioxidation (M)	1	Proteotypic
Q6FGI1_HUMAN	1184	1233.614	NVIGLQMG TNR	Dioxidation (M)	1	Proteotypic
Q6FGI1_HUMAN	1237	1246.670	TLMNLGGLAVAR	Dioxidation (M)	1	Proteotypic
Q6FGI1_HUMAN	1329	1278.584	NFSDNQLQEGK		1	Proteotypic
Q6FGI1_HUMAN	2059	1709.804	QMEQISQFLQAAER	Dioxidation (M)	1	Proteotypic
Q6FH55_HUMAN	626	1084.649	LVLGESA V GK		1	Proteotypic
Q6FH55_HUMAN	1102	1210.569	NEPQNATGAPGR		1	Proteotypic
Q6FH55_HUMAN	1382	1298.621	GVDLQENNPASR		1	Proteotypic
Q6FH55_HUMAN	1610	1394.788	QASPNIVIALAGNK		1	Proteotypic
Q6FHT8_HUMAN	332	1003.468	AINDNTNSR		1	Proteotypic
Q6FHT8_HUMAN	1743	1464.753	IVMFTIDIGEAPK	Dioxidation (M)	1	Proteotypic
Q6IAM7_HUMAN	2388	917.533	VGEIVVFR		1	Proteotypic
Q6IB18_HUMAN	1101	1210.521	EDIYSGGGGGSR		1	Proteotypic
Q6IB18_HUMAN	2042	1689.905	LFIGGLNVQTSEGLR		1	Proteotypic
Q6IBD1_HUMAN	1092	1206.650	ITVTSEVPFSK		1	Proteotypic
Q6IBD1_HUMAN	1219	1241.673	AGNLGGGVVTIER		1	Proteotypic
Q6IPX4_HUMAN	669	1093.711	LLEPVLLLGK		1	Proteotypic
Q6IPX4_HUMAN	1014	1186.646	GPLQSVQVFR		1	Proteotypic
Q6N092_HUMAN	93	925.462	SAVQGPPDR		1	Proteotypic
Q6N092_HUMAN	106	930.539	TPLTATLSK		2	Shared
Q6N092_HUMAN	2164	1834.943	QEPSQGTTTFAVTSILR		2	Shared
Q6NSD1_HUMAN	45	911.544	LLVVTDP R		1	Proteotypic
Q6NSD1_HUMAN	46	911.544	LLVVTDP R		1	Proteotypic
Q6NSD1_HUMAN	1078	1202.641	FAAATGATPIAGR		1	Proteotypic
Q6NSD1_HUMAN	2049	1697.853	FTPGTFTNQIAAFR		1	Proteotypic
Q6NSD1_HUMAN	2084	1739.942	AIVAIENPADVSVSSR		1	Proteotypic
Q6NSD1_HUMAN	2173	1847.919	SGALDVLQMKEEDVLK	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
Q6NZ52_HUMAN	2421	1110.640	TGAAPIIDVVR		1	Proteotypic
Q6P3T4_HUMAN	218	972.560	EVVIVSATR		1	Proteotypic
Q6P3T4_HUMAN	1060	1198.692	LGSIAIQGAIEK		1	Proteotypic
Q76M58_HUMAN	551	1065.593	TALIH DGLAR		1	Proteotypic
Q76M58_HUMAN	2329	2288.121	AEEGIAAGGVMDVNTALQEVLK	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
Q85KX8_HUMAN	714	1105.686	VVXPIEAPIR		1	Proteotypic
Q85KX8_HUMAN	2191	1882.923	ILYMTDEVNDPSLTIK	Dioxidation (M)	2	Shared
Q8IWY7_HUMAN	29	905.461	FLEQQNK		14	Shared
Q8IWY7_HUMAN	329	1002.535	TQEEKQIK		1	Proteotypic
Q8IWY7_HUMAN	802	1128.614	GELAIKDANAK		2	Shared
Q8IWY7_HUMAN	954	1168.543	EYQELMNVK	Oxidation (M)	4	Shared
Q8IWY7_HUMAN	1000	1184.538	EYQELMNVK	Dioxidation (M)	4	Shared
Q8IZK8_HUMAN	2441	1208.688	VNPALAE LNLR		1	Proteotypic
Q8TB65_HUMAN	133	942.575	ILEVVKDK		1	Proteotypic
Q8TB65_HUMAN	396	1021.519	LNDFASTVR		1	Proteotypic
Q8TB65_HUMAN	985	1177.620	RLNDFASTVR		1	Proteotypic

Q8WVX7_HUMAN	96	926.530	KLTPQGQR		1	Proteotypic
Q8WVX7_HUMAN	128	941.529	IAGQVAAANK		1	Proteotypic
Q8WVX7_HUMAN	208	969.586	VLQALEGLK		1	Proteotypic
Q8WVX7_HUMAN	829	1133.547	DVNQQEFVR		1	Proteotypic
Q8WVX7_HUMAN	945	1165.503	MVEKDQDGGR	Dioxidation (M)	1	Proteotypic
Q96Q14_HUMAN	2486	1676.898	VLLESEQFLTELTR		1	Proteotypic
Q9BTQ7_HUMAN	8	899.508	GSAITGPVAK		1	Proteotypic
Q9BTQ7_HUMAN	1829	1522.737	LPAGVGDMVMATVK	Dioxidation (M); Dioxidation (M)	1	Proteotypic
Q9H853_HUMAN	1634	1409.767	QIFHPEQLITGK		1	Proteotypic
Q9H853_HUMAN	1786	1486.872	LISQIVSSITASLR		3	Shared
Q9H853_HUMAN	2066	1717.875	NLDIERPTYTNLNR		4	Shared
Q9UDG2_HUMAN	224	974.478	GLQTSQDAR		2	Shared
Q9UDG2_HUMAN	288	991.461	QIDNPDYK		2	Shared
Q9UDG2_HUMAN	1935	1606.767	FYALSAXFEPFSNK		1	Proteotypic
QCR1_HUMAN	515	1057.577	IAEVDASVVR		1	Proteotypic
QCR1_HUMAN	1315	1273.455	EMQENDASMR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
QCR2_HUMAN	2453	1266.603	QVAEQFLNMR	Dioxidation (M)	1	Proteotypic
QOR_HUMAN	1	898.487	GTIEINPR		1	Proteotypic
QOR_HUMAN	122	938.544	SDIAVPIPK		1	Proteotypic
QOR_HUMAN	502	1053.557	DLSLLSHGGR		1	Proteotypic
QOR_HUMAN	1077	1202.614	ILGTAGTEEGQK		1	Proteotypic
QOR_HUMAN	1137	1220.644	VFEFGGPEVLK		1	Proteotypic
QOR_HUMAN	1749	1467.782	ESSIIGVTLFSSTK		1	Proteotypic
QOR_HUMAN	2108	1773.955	GIDIIIEMLANVNLSK	Dioxidation (M)	1	Proteotypic
QOR_HUMAN	2368	2715.412	KPLLPYTPGSDVAGVIEAVGDNASAFK		1	Proteotypic
RAB10_HUMAN	562	1070.634	LLLIGDSGVGK		3	Shared
RAB10_HUMAN	1278	1260.708	AFLTLAEDILR		1	Proteotypic
RAB14_HUMAN	505	1054.541	LTSEPQQR		1	Proteotypic
RAB14_HUMAN	1564	1368.671	GAAGALMVYDITR	Dioxidation (M)	1	Proteotypic
RAB14_HUMAN	1955	1622.936	NLTNPNTVILIGNK		1	Proteotypic
RAB14_HUMAN	2101	1765.889	QFAEENGLLFLEASAK		1	Proteotypic
RAB21_HUMAN	738	1112.532	AAAGGGGGGAAAAGR	Acetyl (Protein N-term)	1	Proteotypic
RAB21_HUMAN	1146	1222.550	MIETAQVDER	Dioxidation (M)	1	Proteotypic
RAB35_HUMAN	561	1070.634	LLIIGDSGVGK		1	Proteotypic
RAB35_HUMAN	2005	1658.797	FAGQMGIQLFETSAK	Dioxidation (M)	1	Proteotypic
RAB7A_HUMAN	509	1056.618	VIIIGDSGVGK		1	Proteotypic
RAB7A_HUMAN	1763	1474.746	DPENFPFVVLGNK		1	Proteotypic
RAB7A_HUMAN	1922	1588.821	EAINVEQAFQTIAR		1	Proteotypic
RALYL_HUMAN	2455	1287.755	VFIGNLNTAIVK		1	Proteotypic
REEP5_HUMAN	1028	1190.524	HESQMDSVVK	Dioxidation (M)	1	Proteotypic
REEP5_HUMAN	1071	1200.671	KATVNLLGEEK		1	Proteotypic
RL24_HUMAN	2458	1292.665	AITGASLADIMAK	Dioxidation (M)	1	Proteotypic
RL31_HUMAN	116	935.402	EMGTPDVR	Dioxidation (M)	1	Proteotypic
RL31_HUMAN	273	987.535	SAINEVVTR		1	Proteotypic
RL31_HUMAN	866	1144.536	NLQTVNVNVDEN		1	Proteotypic
ROA2_HUMAN	42	911.373	GGSDGYGSGR		1	Proteotypic

ROA2_HUMAN	356	1012.436	GGNFGFGDSR		1	Proteotypic
ROA2_HUMAN	1018	1187.640	IDTIEIITDR		1	Proteotypic
ROA2_HUMAN	1199	1236.540	QEMQEVQSSR	Oxidation (M)	1	Proteotypic
ROA2_HUMAN	1251	1252.535	QEMQEVQSSR	Dioxidation (M)	1	Proteotypic
ROA2_HUMAN	1492	1337.694	EESGKPGAHVTVK		1	Proteotypic
ROA2_HUMAN	1575	1376.622	GGGGNFGPGPSNFR		1	Proteotypic
ROA2_HUMAN	1810	1513.675	LTDCVVMRDPASK	Dioxidation (M); Trioxidation (C)	1	Proteotypic
ROA2_HUMAN	2021	1669.776	LTDCVVMRDPASKR	Dioxidation (M); Trioxidation (C)	1	Proteotypic
ROA2_HUMAN	2047	1694.758	GFGFVTFDDHDPVDK		1	Proteotypic
ROA2_HUMAN	2128	1797.915	LFIGGLSFETTEESLR		1	Proteotypic
ROA3_HUMAN	950	1167.504	EDTEEYNLR		1	Proteotypic
ROA3_HUMAN	1289	1265.581	IETIEVMEDR	Dioxidation (M)	1	Proteotypic
ROA3_HUMAN	1581	1379.741	EDSVKPGAHLTVK		1	Proteotypic
ROA3_HUMAN	2152	1820.884	MEVKPPPGRPQPDSCR	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
RS11_HUMAN	578	1072.613	VLLGETGKEK		1	Proteotypic
RS11_HUMAN	850	1140.523	CPFTGNVSIR	Trioxidation (C)	1	Proteotypic
RS11_HUMAN	2246	1975.988	DVQIGDIVTVGECRPLSK	Trioxidation (C)	1	Proteotypic
RS13_HUMAN	204	967.484	DSHGVAQVR		1	Proteotypic
RS13_HUMAN	1253	1252.750	GLTPSQIGVILR		1	Proteotypic
RS14_HUMAN	503	1053.557	TPGPGAQSALR		1	Proteotypic
RS14_HUMAN	1671	1428.710	IEDVTPIPSDSTR		1	Proteotypic
RS20_HUMAN	2459	1293.660	SLEKVCADLIR	Trioxidation (C)	1	Proteotypic
RS25_HUMAN	212	971.565	AALQELLSK		1	Proteotypic
RS25_HUMAN	619	1083.629	LITPAVVSER		1	Proteotypic
RS3_HUMAN	423	1028.623	TEIILATR		1	Proteotypic
RS3_HUMAN	660	1091.561	AELNEFLTR		1	Proteotypic
RS3_HUMAN	1659	1422.663	ELAEDGYSGVEVR		1	Proteotypic
RS3_HUMAN	1901	1571.751	FGFPEGSVELYAEK		1	Proteotypic
RS3_HUMAN	1933	1604.787	GGKPEPPAMPQPPTA	Dioxidation (M)	1	Proteotypic
S10A6_HUMAN	60	914.482	LQDAEIAR		1	Proteotypic
S10A6_HUMAN	80	922.407	LMEDLDR	Dioxidation (M)	1	Proteotypic
S10AB_HUMAN	386	1018.493	ISSPTETER		1	Proteotypic
S10AB_HUMAN	851	1140.589	NQKDPGVLDLDR		1	Proteotypic
S10AB_HUMAN	1380	1297.659	CIESLIAVFQK	Trioxidation (C)	1	Proteotypic
S10AB_HUMAN	2189	1880.887	TEFLSFMNTELAFTK	Dioxidation (M)	1	Proteotypic
S10AB_HUMAN	2331	2298.141	ISSPTETERCIESLIAVFQK	Trioxidation (C)	1	Proteotypic
SCMC1_HUMAN	2442	1211.699	QLLAGGIAGAVSR		1	Proteotypic
SEPT9_HUMAN	489	1046.524	ADTLLEER		1	Proteotypic
SEPT9_HUMAN	497	1050.680	VVNIVPVIK		1	Proteotypic
SEPT9_HUMAN	1052	1197.610	QVENAGAIGPSR		1	Proteotypic
SF3B1_HUMAN	2387	914.519	VGAAEISR		1	Proteotypic
SFRS3_HUMAN	1950	1620.757	NPPGFVFEFEDPR		2	Shared
SFRS3_HUMAN	2335	2319.092	NPPGFVFEFEDPRDAADAVR		1	Proteotypic
SMD1_HUMAN	1301	1268.698	LSHETVTIELK		1	Proteotypic
SMD1_HUMAN	1877	1553.853	NREPVQLETLSIR		1	Proteotypic
SMD3_HUMAN	776	1120.620	FLILPDKL	Dioxidation (M)	1	Proteotypic

SMD3_HUMAN	1126	1217.677	VAQLEQVYIR		1	Proteotypic
SNRPA_HUMAN	37	908.512	GQAFVIFK		1	Proteotypic
SNRPA_HUMAN	490	1046.536	EVSSATNALR		1	Proteotypic
SPTB2_HUMAN	32	906.445	IDDIFER		1	Proteotypic
SPTB2_HUMAN	47	911.555	VQAVVAVAR		1	Proteotypic
SPTB2_HUMAN	103	929.457	QLQEDAAR		1	Proteotypic
SPTB2_HUMAN	206	968.577	VAVVNQIAR		1	Proteotypic
SPTB2_HUMAN	369	1014.571	LLEVLSGER		1	Proteotypic
SPTB2_HUMAN	496	1049.551	LVDTGDKFR		1	Proteotypic
SPTB2_HUMAN	593	1076.554	LEMNLGLQK	Dioxidation (M)	1	Proteotypic
SPTB2_HUMAN	609	1080.512	ETASELLMR	Dioxidation (M)	1	Proteotypic
SPTB2_HUMAN	817	1130.582	VLDNAIETEK		1	Proteotypic
SPTB2_HUMAN	1011	1185.697	LTTLELLEVR		1	Proteotypic
SPTB2_HUMAN	1024	1188.573	LVSDGNINSDR		1	Proteotypic
SPTB2_HUMAN	1076	1202.589	DQNTVETLQR		1	Proteotypic
SPTB2_HUMAN	1174	1231.593	EIGQSVDEVEK		1	Proteotypic
SPTB2_HUMAN	1213	1240.693	LILEVHQFSR		1	Proteotypic
SPTB2_HUMAN	1560	1366.757	TALPAQSAATLPAR		1	Proteotypic
SPTB2_HUMAN	1658	1421.686	FMELLEPLNER	Dioxidation (M)	1	Proteotypic
SPTB2_HUMAN	1750	1468.629	QQMLENQMEVR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
SPTB2_HUMAN	2109	1776.885	SQNIVTDSSLSAEAIR		1	Proteotypic
SPTB2_HUMAN	2269	2021.011	LVSQDNFGFDLPAVEAATK		1	Proteotypic
SPTB2_HUMAN	2311	2161.068	VIESTQDLGNDLAGVMALQR	Dioxidation (M)	1	Proteotypic
SSRG_HUMAN	2488	1706.811	QQSEEDLLLQDFSR		1	Proteotypic
STOM_HUMAN	67	914.555	LLAQTTLR		1	Proteotypic
STOM_HUMAN	450	1037.445	AMAAEAEASR	Dioxidation (M)	1	Proteotypic
STOM_HUMAN	1330	1278.587	VIAAEGEMNASR	Dioxidation (M)	1	Proteotypic
STOM_HUMAN	2089	1746.882	EASMVITESPAALQLR	Dioxidation (M)	1	Proteotypic
STOM_HUMAN	2130	1801.896	GPGLFFILPCTDSFIK	Trioxidation (C)	1	Proteotypic
STOM_HUMAN	2220	1928.992	VQNATLAVANITNADSATR		1	Proteotypic
SYEP_HUMAN	130	941.529	VAVQGDVVR		1	Proteotypic
SYEP_HUMAN	1080	1203.598	LTVAENEAETK		1	Proteotypic
TALDO_HUMAN	107	931.461	AAQASDLEK		1	Proteotypic
TALDO_HUMAN	359	1012.501	TIVMGASFR	Dioxidation (M)	1	Proteotypic
TALDO_HUMAN	584	1073.535	LGGSQEDQIK		1	Proteotypic
TALDO_HUMAN	1109	1212.671	LLGELLQDNAK		1	Proteotypic
TALDO_HUMAN	1336	1283.618	LSFDKAMVAR	Dioxidation (M)	1	Proteotypic
TCPA_HUMAN	875	1146.603	EQLAIAEFAR		1	Proteotypic
TCPA_HUMAN	1085	1204.682	FATEAAITILR		1	Proteotypic
THEM2_HUMAN	171	955.570	ITLVSAAPGK		1	Proteotypic
THEM2_HUMAN	305	996.455	TSMTQSLR	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
THEM2_HUMAN	1419	1308.693	TLAFTSVDLTNK		1	Proteotypic
THOC4_HUMAN	513	1057.549	GRGRAGSQGGR		1	Proteotypic
THOC4_HUMAN	1931	1599.701	ADKMDMSLDDIIK	Acetyl (Protein N-term); Dioxidation (M); Dioxidation (M)	1	Proteotypic
TIF1B_HUMAN	469	1043.423	MNEAFGDTK	Dioxidation (M)	1	Proteotypic
TIF1B_HUMAN	903	1152.508	SERGEVSGLMR	Dioxidation (M)	1	Proteotypic

TIF1B_HUMAN	2279	2050.027	IVAERPSTNSTGPAPMAPPR	Dioxidation (M)	1	Proteotypic
TM109_HUMAN	2460	1296.704	EAPVDVLTQIGR		1	Proteotypic
TMED9_HUMAN	693	1100.619	DKLSELQLR		1	Proteotypic
TMED9_HUMAN	807	1129.559	FSLFAGGMLR	Dioxidation (M)	1	Proteotypic
TMED9_HUMAN	1501	1340.730	QLVEQVEQIQK		1	Proteotypic
TPPP3_HUMAN	2487	1685.757	AASTDMAGLEESFRK	Acetyl (Protein N-term); Dioxidation (M)	1	Proteotypic
UAP56_HUMAN	114	934.512	ELAFQISK		1	Proteotypic
UAP56_HUMAN	702	1102.650	ILVATNLFGR		1	Proteotypic
UAP56_HUMAN	844	1136.659	VAVFFGGLSIK		1	Proteotypic
UAP56_HUMAN	1227	1242.734	DFLLKPELLR		1	Proteotypic
UAP56_HUMAN	1773	1478.725	GLAITFVSDENDAK		1	Proteotypic
UBP5_HUMAN	251	983.504	EVQDGIAPR		1	Proteotypic
UBP5_HUMAN	981	1177.536	SSENPNEVFR		1	Proteotypic
UBP5_HUMAN	1701	1442.725	SAADSISESVPVGP		1	Proteotypic
UBP5_HUMAN	2113	1781.977	AELSEEALLSVLPTIR	Acetyl (Protein N-term)	1	Proteotypic
UGDH_HUMAN	587	1073.598	LAANAFLAQR		1	Proteotypic
UGDH_HUMAN	755	1116.578	VTVVVDVNESR		1	Proteotypic
VDAC1_HUMAN	427	1029.607	LTLALDGGK		1	Proteotypic
VDAC1_HUMAN	1107	1212.614	VTQSNFAVGYK		1	Proteotypic
VDAC1_HUMAN	1618	1399.662	LTFDSSFSPNTGK		1	Proteotypic
VDAC1_HUMAN	1838	1527.757	LTFDSSFSPNTGKK		1	Proteotypic
VDAC1_HUMAN	2240	1958.870	SENGLEFTSSGSANTETTK		1	Proteotypic
VDAC3_HUMAN	2401	1029.607	LTLALDGGK		1	Proteotypic
VIME_HUMAN	25	904.425	GTNESLER		2	Shared
VIME_HUMAN	29	905.461	FLEQQNK		14	Shared
VIME_HUMAN	54	913.451	SYVTTSTR		1	Proteotypic
VIME_HUMAN	108	931.461	LLEGEESR		4	Shared
VIME_HUMAN	202	966.513	TYSLGSALR		1	Proteotypic
VIME_HUMAN	353	1011.444	MFGGPGTASR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	400	1022.503	QQYESVAAK		1	Proteotypic
VIME_HUMAN	525	1059.556	KLLEGEESR		3	Shared
VIME_HUMAN	595	1077.512	LQEEMLQR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	638	1087.526	QDVNASLAR		1	Proteotypic
VIME_HUMAN	664	1092.520	FADLSEANR		1	Proteotypic
VIME_HUMAN	722	1107.487	DNLAEDIMR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	747	1114.562	VELQELNDR		1	Proteotypic
VIME_HUMAN	775	1120.576	EYQDLLNVK		2	Shared
VIME_HUMAN	890	1149.534	LQDEIQNMK	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	955	1168.707	ILLAELEQLK		1	Proteotypic
VIME_HUMAN	1121	1215.621	RQVDQLTNDK		1	Proteotypic
VIME_HUMAN	1343	1285.550	LGDLYEEEMR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	1347	1286.658	QVDQLTNDKAR		1	Proteotypic
VIME_HUMAN	1460	1322.610	EEAENTLQSF		1	Proteotypic
VIME_HUMAN	1470	1326.649	MALDIEIATYR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	1486	1334.650	EKLQEEMLQR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	1667	1427.704	SLYASSPGGVYATR		1	Proteotypic

VIME_HUMAN	1776	1480.708	QVQSLTCEVDALK	Trioxidation (C)	1	Proteotypic
VIME_HUMAN	1800	1494.779	TYSLGSALRPSTSR		1	Proteotypic
VIME_HUMAN	1834	1525.694	MFGGPGTASRPSSSR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	1847	1532.845	KVESLQEEIAFLK		1	Proteotypic
VIME_HUMAN	1854	1538.903	ILLAELEQLKGQ GK		1	Proteotypic
VIME_HUMAN	1899	1569.888	ISLPLPNFSSLNLR		1	Proteotypic
VIME_HUMAN	1918	1586.790	TNEKVELQELNDR		1	Proteotypic
VIME_HUMAN	2010	1660.940	KVESLQEEIAFLKK		1	Proteotypic
VIME_HUMAN	2018	1667.837	ETNLDSLPLVDTHSK		1	Proteotypic
VIME_HUMAN	2033	1683.777	LGDLYEEEMRELR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	2067	1719.810	VEVERDNLAEDIMR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	2127	1797.787	LQDEIQNMKEEMAR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
VIME_HUMAN	2297	2125.058	LLQDSVDFSLADAINTEFK		1	Proteotypic
VIME_HUMAN	2320	2217.948	EMEENFAVEAANYQDTIGR	Dioxidation (M)	1	Proteotypic
VIME_HUMAN	2344	2367.122	QVQSLTCEVDALKGTNESLER	Trioxidation (C)	1	Proteotypic
VIME_HUMAN	2346	2382.112	LQEEMLQREEAENTLQSFR	Dioxidation (M)	1	Proteotypic
VINC_HUMAN	707	1104.603	SLGEISALTSK		1	Proteotypic
VINC_HUMAN	760	1117.598	STVEGIQASVK		1	Proteotypic
VINC_HUMAN	966	1172.640	ALASQLQDSLK		1	Proteotypic
VINC_HUMAN	1088	1206.591	MSAEINEIIR	Dioxidation (M)	1	Proteotypic
VINC_HUMAN	1239	1247.520	MALLMAEMSR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	1	Proteotypic
VINC_HUMAN	1300	1268.672	AVAGNISDPGLQK		1	Proteotypic
VINC_HUMAN	1434	1313.730	QVATALQNLQTK		1	Proteotypic
VINC_HUMAN	1464	1323.623	MTGLVDEAIDTK	Dioxidation (M)	1	Proteotypic
VINC_HUMAN	1726	1456.789	AQQVSQGLDVLTA K		1	Proteotypic
VINC_HUMAN	1857	1540.686	MLGQMTDQVADLR	Dioxidation (M); Dioxidation (M)	1	Proteotypic
VINC_HUMAN	1941	1613.685	AGEVINQPMMAAR	Dioxidation (M); Dioxidation (M); Dioxidation (M)	1	Proteotypic