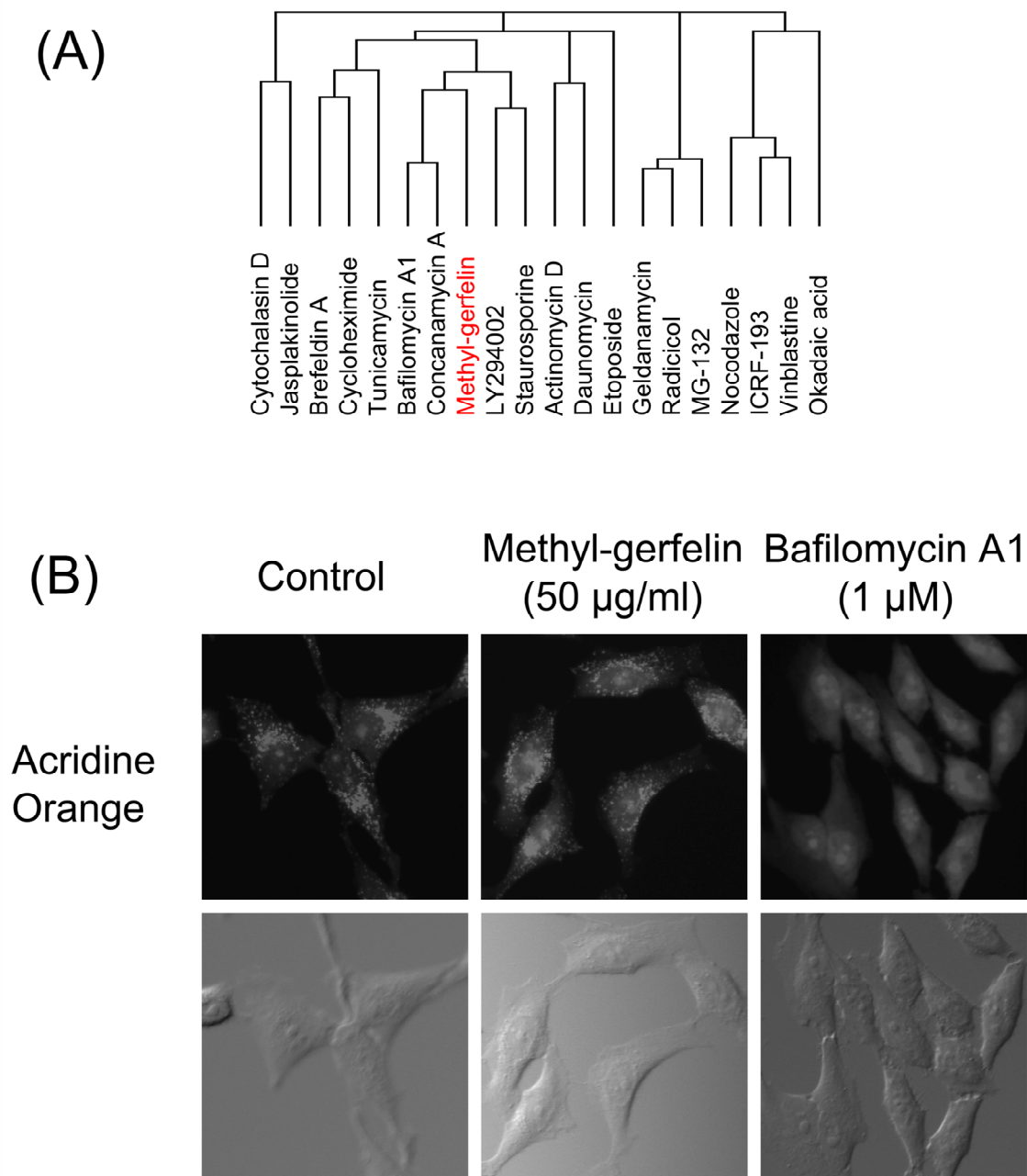


Figure S1



A) Clustering of 19 compounds and methyl-gerfelin by proteomic profiling analysis.

HeLa cells treated with 50 μ g/ml methyl-gerfelin for 18 h were analyzed by 2D-DIGE. The data was combined with the data of 19 compounds and the

data of 312 spots matched to the all gels were used for hierarchical clustering. Slight correlation of the pattern of V-ATPase inhibitor-treated HeLa cells was observed.

B) Effect of methyl-gerfelin on organelles acidification.

Acidic organelles such as lysosome are acidified by V-ATPase. After 2 h incubation with a compound, HeLa cells were stained with 5 μ M acridine orange for 15 min. The medium was then replaced to acridine orange free medium and accumulation of acridine orange in acidic organelles was observed by a fluorescent microscopy (Olympus, Tokyo). No inhibitory effect of methyl-gerfelin on organelles acidification was observed. Upper panel, WIG filter (ex. 520-550, em. > 580; Olympus, Tokyo); Lower panel, Nomarski optics.

Result of Figure 1B (Geldanamycin and Radicicol)

Master No.	Control	Geldanamycin					Radicicol
		0.005 μ M	0.05 μ M	0.5 μ M	5 μ M	10 μ M	Rad 10 μ M
257	1	0.873925283	0.943677984	0.784290057 **	0.724564554 **	0.719296695 **	0.870815425
297	1	1.015998799	1.04121973	1.278737437 *	1.315568736 **	1.146612215	0.815041374
412	1	0.970571797	0.884633562	1.223686369 **	1.108495608	1.110246157	1.064525099
453	1	1.019390084	0.85556685 **	1.033679277	1.040195392	0.929174605	0.874267595 *
461	1	0.966122115	1.155067789 **	1.219707867 **	1.163596218 **	1.112966062 *	1.116092784 *
508	1	0.919433225	1.369871771 *	1.51614718 **	1.77940585 **	1.704832543 **	1.952166706 **
513	1	0.906762743	1.442304378 *	1.736066308 **	2.436340971 **	2.085068324 **	2.711879948 **
564	1	1.040031407	0.92022118	0.881519193	0.778968631 **	0.746753859 **	0.896959139
576	1	1.066846334	1.055228652	1.302789095 **	1.312686372 **	1.271788104 **	1.091886139
603	1	1.077729437	0.994748819	1.175993477 *	1.280038245 **	1.433572733 **	1.05278334
605	1	1.012987594	1.111662115	1.128704465	1.142466024	1.263983678 **	1.301814461 **
607	1	1.208424146	1.15976399	1.457718912 *	1.73547962 **	1.8515377 **	1.191591635
608	1	1.018520638	1.16962163 *	1.257491409 **	1.294745051 **	1.212194759 *	1.200041208 *
623	1	0.942402963	0.962874019	0.809411866 **	0.826291514 **	0.793082388 **	0.98864867
641	1	0.984700653	0.978555345	1.351220809 **	1.397116224 **	1.477046382 **	1.147953279
642	1	0.996941276	0.94069723	1.254746013 **	1.309727745 **	1.397351226 **	1.206326107 *
643	1	0.916786019	0.944057026	0.717793029 **	0.764885897 **	0.722081175 **	0.790182718 **
645	1	1.038520865	1.033011046	1.326846865 *	1.490405327 **	1.570319851 **	1.173937962
647	1	1.016972315	0.968759297	1.294788437 **	1.307735 **	1.409169761 **	1.192079078 *
650	1	1.033084879	1.08313397	1.429951253 **	2.265064083 **	2.230578587 **	1.357977802 *
669	1	1.040122638	1.073155491	1.324718133 **	1.230466432 *	1.200213309 *	1.187050161 *
683	1	1.004908402	0.844337131	1.003150891	1.261177385 **	1.125498225	0.86575043
687	1	1.059656335	1.027289062	1.227094983 **	1.213365096 **	1.141346637	0.970354003
696	1	0.965138476	1.016899249	0.830486428 **	0.872035577 *	0.89434964	0.932262485
698	1	1.112140206	0.959150436	0.863651178 *	0.937003021	0.821209016 **	0.897234192
700	1	1.029333153	0.969118263	0.672198848 **	0.66328058 **	0.652455109 **	0.839135234 *
701	1	1.025813848	0.982852672	0.727914365 **	0.705266025 **	0.671186793 **	0.900569189
708	1	1.039021421	1.025670862	1.207017697 **	1.199029639 **	1.158939417 **	1.040266907
713	1	0.972139971	0.934029994	0.714137643 **	0.658738615 **	0.70481901 **	0.823334198 *
714	1	0.949940127	0.954934259	0.63335145 **	0.594149424 **	0.548598125 **	0.716249327 **
728	1	0.962069683	0.910814885 *	0.696624142 **	0.674481169 **	0.627133669 **	0.704036849
747	1	1.031296622	1.001197612	0.913857908 *	0.8594471 **	0.904272933 *	0.873436964 **
761	1	1.045999391	0.919542917	0.7848582 **	0.7248204 **	0.738762353 **	0.674325912 **
780	1	1.06914249	1.117467669 *	1.242091827 **	1.161666558 **	1.241941527 **	1.17363328 **
784	1	0.860099132	0.998599857	1.296594821 **	1.258003561 *	1.379272707 **	1.412532089 **
788	1	1.036405034	1.004108462	1.19788258 **	1.141394128 *	1.094667964	1.013191628
794	1	1.004015574	1.048066019	1.22059625 **	1.160476669 *	1.107253867	1.056567784
813	1	1.02282288	1.152613444	1.691579283 **	1.813066662 **	1.76759592 **	1.605611071 **
852	1	1.046097971	1.044048436	1.203163698 **	1.220286207 **	1.153387626 *	1.026254106
856	1	0.939484691	1.036712226	1.282955129 **	1.344323579 **	1.410069772 **	1.19635279 **
863	1	1.061642672	1.001408975	1.212362517 **	1.151428908 **	1.11314608 *	0.925243668
865	1	0.998616487	0.981672746	0.927888367	0.949472255	0.946688706	0.845789033 **
866	1	1.015116072	0.998692275	1.151076609 *	1.253747983 **	1.247518295 **	1.176513356 *
867	1	1.029370347	0.979893045	1.154838028 **	1.115379035 *	1.139343803 **	0.97838696
873	1	1.025281085	0.981846158	0.768247788 **	0.705179973 **	0.804390466 *	0.818475497 *
877	1	0.900893403	0.849324461	0.719958622 **	0.908213281	0.79207379 *	0.813432054 *
889	1	1.053961047	1.021503083	0.929607774	0.896723858	0.875108407 *	0.812616178 **
907	1	0.956591372	1.067735383	1.110142114	1.263667398 **	1.284974361 **	1.313103861 **
909	1	1.025847353	0.982046613	0.905092587 **	0.845232541 **	0.811427045 **	0.668618263 **
914	1	0.975992212	0.999578328	1.014420212	1.101035775 *	1.168723608 **	1.034124229
916	1	0.96549165	0.945485415 *	0.93037947 *	0.990184963	1.028295598	0.898678387 **
921	1	1.074000017	1.065222926	0.879872626	1.311004662 *	1.340778238 **	0.810274309
928	1	1.040781602	1.057950658	1.255537435 **	1.207692911 *	1.227327663 *	0.969551894
929	1	1.040672296	0.925688347	0.844821504 *	0.927399239	0.890698831	0.755004607 **
937	1	0.999302718	1.003463524	1.084946752	1.429932099 **	1.40427608 **	0.906227734
939	1	0.862027738 **	0.902492625	0.949655188	0.963472874	0.973763056	0.807487933 **
944	1	1.03650141	1.029547276	1.244953243 **	1.255722132 **	1.301714413 **	1.06365052
951	1	0.997025201	0.953701405	0.866957381 **	0.862232516 **	0.842882223 **	0.749489979 **
971	1	1.028203885	1.071165488	2.259898763 **	5.90325774 **	5.720081278 **	2.60235366 **
983	1	1.006277284	1.040710137	1.289900085 *	2.701901957 **	2.709074895 **	1.284927162 *

998	1	0.953723156	1.034025409	1.385511438 **	1.306259419 **	1.406792396 **	1.281523982 *
1006	1	1.049399904	1.066961139	1.409720333 **	1.349471745 **	1.329754741 **	1.165908351 *
1009	1	1.022769994	1.055961658	1.46613842 **	1.429128035 **	1.409670648 **	1.179306568 *
1020	1	0.957144935	0.971387885	0.82243203 *	0.732689094 **	0.719061299 **	0.85621227 *
1034	1	1.004902184	1.033917657	0.68872867 **	0.75296886 **	0.692618891 **	0.910848043
1036	1	0.957821528	0.99283698	1.049459485	1.369349849 **	1.276495885 **	1.023151038
1040	1	0.972522298	1.122824189	1.427694346 **	1.387438051 **	1.470089407 **	1.50036558 **
1045	1	1.030711267	1.227031823 **	1.631045491 **	1.538737441 **	1.606034607 **	1.646044839 **
1046	1	0.983055841	0.920337115	0.691966035 **	0.719272618 **	0.677710078 **	0.730142856 **
1047	1	0.980796832	1.110376872	1.304364977 **	1.241025777 **	1.269966318 **	1.329673346 **
1069	1	1.067751087	1.028622922	1.021806387	0.997501107	0.928679466	0.829528457 **
1073	1	1.015680288	0.977728251	0.985293624	0.677164054 **	0.58771955 **	0.677745029 **
1076	1	0.989549559	0.883217092 *	0.888470458 *	0.85531782 **	0.824028697 **	0.80337174 **
1081	1	1.111997408	1.121462045	1.971910144 **	2.243249997 **	1.622434068	1.652732568
1082	1	1.020667032	0.934252574	0.89761797	0.833339952 **	0.753068681 **	0.831623165 **
1084	1	1.034921797	1.095401935	0.887590427	0.844045614 **	0.934274846	0.826865736 **
1109	1	1.082698741	2.107400502 **	4.460597197 **	4.347874524 **	4.82247445 **	5.279213633 **
1110	1	1.089007744	2.729972624 **	7.299329972 **	7.397572179 **	6.995505625 **	8.759012438 **
1111	1	0.923165552	1.541776129	4.110073667 **	4.262372595 **	3.983808647 **	3.857172428 **
1113	1	1.055948979	0.955077208	1.13469809	1.595407877 **	1.52867207 **	1.360830896 **
1114	1	1.054317967	2.736477643 **	7.216407984 **	7.329337431 **	8.052530781 **	7.42779164 **
1127	1	1.104323516	3.404717155 **	8.943202929 **	9.202001876 **	9.26454239 **	9.392162145 **
1129	1	0.947860659	0.969333027	1.060220364	1.451911249 **	1.586709448 **	1.050856239
1135	1	1.103946739	1.396211492	1.902582408 **	1.95071494 **	2.022745604 **	2.240773788 **
1160	1	1.01155521	1.003422497	0.871018887 **	0.872066047 **	0.922782421	0.852121303 **
1174	1	0.954669424	1.060625887	1.220261352 **	1.167523907 *	1.111514362	1.119200366
1190	1	1.062255268	1.006009139	1.235589009 **	1.31537193 **	1.248573145 **	1.056327016
1191	1	1.03403368	0.984081522	1.24490893 **	1.27272913 **	1.256761278 **	0.981202737
1192	1	1.104871518	1.041494159	1.342654091 **	1.32192985 **	1.276631706 **	1.018173284
1203	1	0.955772408	0.973252124	1.003582978	1.152480166	1.17426081 *	1.401875883 **
1206	1	1.057253926	1.012398086	1.110691531 *	1.158649786 **	1.098999497 *	1.004435184
1210	1	0.954953654	0.999168915	1.129508474 *	1.130968695 *	1.057217888	1.264701541 **
1211	1	0.91516427 *	0.943266193	0.815526881 **	0.829710414 **	0.770368935 **	0.953766543
1213	1	0.932317512	0.961098938	1.023517263	1.056489412	1.077950529	1.267099048 **
1214	1	0.959716174	1.104572173	1.385642744 **	1.30218585 **	1.395044247 **	1.268834794 **
1219	1	1.035385644	0.966481472	1.197815231 **	1.208707042 **	1.242734485 **	1.097351597
1221	1	0.942949742	1.145619313 *	1.121077527 *	1.208742986 **	1.149032452 *	1.222287298 **
1229	1	0.988367359	0.980509642	1.026511659	1.329458428 **	1.251113204 *	0.992568557
1280	1	0.925044209	0.986722204	1.199630205	1.322869731 *	1.495181979 **	1.164365319
1284	1	1.014344636	1.043109398	1.425901815 **	1.239366646 *	1.272025533 *	1.168644281
1292	1	1.010835109	1.106879424	1.684039443 **	1.57417884 **	1.680909093 **	1.38223157 **
1293	1	1.043991799	1.174827758	1.643348103 **	1.466559366 **	1.536884398 **	1.241481138
1295	1	1.022660991	1.193960771 *	1.749965053 **	1.616481783 **	1.640567463 **	1.344759594 **
1300	1	0.978225976	0.945949447	0.868813254 **	0.867997043 **	0.888909077 *	0.939310488
1306	1	1.043626183	1.109992371	1.270862173 **	1.177648376 *	1.157901183 *	1.227101096 **
1319	1	1.037249028	0.970933611	0.915756959	0.832243576 **	0.836033743 **	0.754193318 **
1323	1	0.936662338	0.945972681	0.753553589 **	0.779243997 **	0.751205509 **	0.85672576 *
1326	1	1.040943986	1.022748254	0.801517116 **	0.911357621	0.816311505 *	0.782354805 **
1342	1	0.988921541	0.984836429	1.175537982 **	1.266072369 **	1.246899839 **	1.015889848
1353	1	1.011685517	0.977406259	1.299926536 **	1.423504907 **	1.547406207 **	1.119303055
1358	1	1.209802012	1.398130603 *	1.393575403 *	1.364631278 *	1.493290409 **	1.010336777
1361	1	1.001547324	1.001470518	1.237423328 **	1.210205332 *	1.208040942 *	1.051897873
1362	1	1.038418794	0.986528464	1.210601681 **	1.487112712 **	1.462999074 **	1.021995893
1365	1	1.029161968	1.017362772	1.193789054 **	1.300397383 **	1.308064209 **	1.021755033
1376	1	0.993856337	0.951200083	0.823229637 **	0.877573926 **	0.889895315 **	0.977923213
1382	1	0.99565273	0.969287302	0.815451117 **	0.899795107 *	0.87918329 *	1.072855908
1386	1	1.081541372	0.941870963	1.254028665 *	1.258731915 *	1.328231732 **	1.033810699
1399	1	1.090123146	1.05910539	1.205915048 **	1.261031752 **	1.287037984 **	1.002525041
1405	1	1.045621017	1.130055127	1.267387779 *	1.246973232 *	1.334032017 **	1.160463961
1406	1	1.058056087	1.023001336	1.484417917 **	1.325807325 **	1.306849884 **	1.277727762 **
1408	1	0.847147343	1.041405531	1.104657783	1.118889262	1.271903206 **	1.115949796
1409	1	0.982209039	0.976087069	1.427899735 **	1.214973591 **	1.086221827	1.679449625 **
1417	1	1.102529787	1.010284563	1.2715438 **	1.27470779 **	1.201194935 *	0.995135077
1420	1	1.031304138	0.992042335	0.632097334 **	0.505897741 **	0.495557376 **	0.641428639 **

1421	1	0.982323707	0.936074828	1.193685012	1.320544219 **	1.260764701 *	1.08017654
1423	1	0.924752152	0.968514142	0.842146866 *	0.872958693 *	0.808428301 **	1.020246312
1429	1	0.962503809	0.959969849	0.612269658 **	0.392938859 **	0.400020849 **	0.551111044 **
1443	1	0.939487028	0.940214517	1.061248645	1.528325991 **	1.514400754 **	1.197622618 *
1445	1	1.024461716	0.983311439	0.964524867	1.354493995 **	1.268682586 **	1.030175675
1446	1	1.001786739	0.963731188	0.736177344 **	0.744091831 **	0.703918895 **	0.897972301 *
1448	1	0.936909621	0.9535136	0.762977157 **	0.740029758 **	0.730025579 **	0.771585452 **
1453	1	0.966854638	1.019414457	0.874330253 *	0.806544479 **	0.83932759 **	0.806035268 **
1460	1	0.992475296	1.041989892	1.325950706 **	1.117435841	1.219652549 **	1.018713052
1462	1	1.014339462	0.959063534	1.320990707 **	1.312593893 **	1.261207939 **	1.166065343 *
1466	1	1.085366749	1.041916851	1.508729393 **	1.479905951 **	1.496410661 **	1.136674775
1469	1	1.032344459	1.006843674	0.866287734	0.82839966	0.753293648 **	0.832450207 *
1472	1	1.116318548	1.093683544	1.496574426 **	1.471166655 **	1.46403214 **	1.225734917 *
1474	1	1.075883932	1.070822366	1.457272502 **	1.415754275 **	1.397130067 **	1.169801199 *
1475	1	1.070534523	1.044016034	1.438045913 **	1.349727515 **	1.25803295 **	1.068430694
1480	1	1.04642127	1.042911502	1.599396113 **	1.516624085 **	1.525022868 **	1.346997228 **
1481	1	1.103706825	1.045998129	1.493147587 **	1.551052891 **	1.467863714 **	1.076627645
1482	1	1.081355447	0.995597917	1.281036659 **	1.264092318 **	1.216069595 *	1.073267902
1484	1	0.965903159	0.960155464	0.711102385 **	0.722801295 **	0.707567347 **	0.751948313 **
1485	1	0.960645703	0.944600827	0.792468603 **	0.834967039 **	0.856863778 **	0.855973827 **
1488	1	1.039809086	0.975011675	1.173900282 **	1.164602409 **	1.12114999 **	0.939026228
1489	1	0.999727271	0.946788802	1.189946934 **	1.226308187 **	1.193443359 **	0.905654502
1491	1	1.006582513	0.932364285	0.739849224 **	0.744230659 **	0.725295655 **	0.898116866 *
1498	1	1.053740877	0.98107789	1.207681493 **	1.150537753 **	1.140304885 **	0.913356821
1499	1	1.031519462	0.999793163	1.281058799 **	1.259556339 **	1.19045735 *	1.102505966
1511	1	0.943564405	0.940937867	0.860605533 **	0.885006554 **	0.83582014 **	0.849389925 **
1518	1	1.058206679	1.015521301	1.131785526 **	1.124001101 *	1.074320187	0.99214311
1548	1	0.95758026	0.937851102	0.884922779 *	0.895268807 *	0.887101266 *	0.837125072 **
1562	1	0.933251967	0.984596814	0.849255237 **	0.896429989 *	0.857205593 **	0.836630419
1563	1	1.003117944	1.009544925	0.862383412 **	0.877575018 **	0.905160224 *	0.846911737
1570	1	0.886311154	0.992193481	0.774534503 **	0.733314227 **	0.814783606 *	0.711089124 **
1579	1	1.035037927	1.023528964	1.191253648 **	1.442258341 **	1.422386348 **	1.069648553
1596	1	1.006940679	0.980929553	1.240916935 **	1.20822614 **	1.252009 **	0.920681597
1619	1	1.077214079	1.101491461	1.354108981 **	1.326743396 **	1.310155901 **	1.042621929
1632	1	1.060098672	1.082132756	1.427906034 **	1.410060151 **	1.381717785 **	1.064252502
1634	1	1.062376373	0.981975704	1.468082192 **	1.249272662 **	1.285084272 **	1.449639785 **
1635	1	1.042331508	1.046406568	1.40612721 **	1.46051817	1.444687347 **	1.205757752 **
1639	1	0.945145064	0.965782981	0.912162538 *	0.815506676 **	0.831607561 **	0.917458684 *
1649	1	0.952368526	0.974736191	0.879760765 *	0.843222152 **	0.84504656 **	1.010410498
1651	1	0.942614337	0.950846412	0.81615129 **	0.832928951 **	0.839077125 **	0.910592345 *
1652	1	0.930831085	0.947500756	0.763113049 **	0.782195045 **	0.75553176 **	0.924039966
1674	1	0.943360533	0.959413703	0.795568423 **	0.826796677 **	0.825032056 **	0.794598645 **
1685	1	0.997300324	0.994918555	0.816477799 **	0.85507672 **	0.834951172 **	0.8956025 *
1690	1	1.008932469	1.175099196 **	1.27411243 **	1.290051543 **	1.270197846 **	1.270961621 **
1696	1	1.082084982	1.255809055 **	1.951958354 **	1.984567267 **	2.081671413 **	1.621377022 **
1700	1	0.9408498	0.939118738	0.793316532 **	0.832910413 **	0.778941201 **	0.945103708
1732	1	1.04037258	0.984403518	1.173346359 **	1.155931295 **	1.1146369 *	0.92248693
1737	1	1.010190009	1.017618451	0.888372629 *	0.841507455 **	0.828898599 **	0.880070359 *
1738	1	1.081334943	0.924325596	1.212688429 *	1.28716749 **	1.214709129 *	0.973213947
1744	1	0.948819758	0.979779858	0.800039463 **	0.781475849 **	0.758539451 **	0.952963328
1746	1	0.969702584	0.951376185	0.765781939 **	0.836801749 **	0.815090318 **	0.800785167 **
1761	1	1.066365837	0.965254267	1.11419425	1.1338282	1.118161182	0.856923714
1780	1	1.121937025	1.080254324	1.242252311 *	1.433775251 **	1.361278326 **	0.970360071
1788	1	0.908344715	0.950383362	0.789534692 **	0.828520777 **	0.790764966 **	0.932984299
1795	1	1.017465472	1.004701756	0.882587867 **	0.841103066 **	0.831001722 **	0.906801885 *
1808	1	0.928037588	0.992600697	0.748682215 **	0.73632815 **	0.706124194 **	0.862059468
1815	1	1.043021023	1.031312758	1.241887868 **	1.398553891 **	1.388978051 **	0.985471406
1816	1	0.921936292	0.978685612	0.798104093 **	0.84621689 **	0.79527629 **	0.91150068
1837	1	0.947179172	0.940508779	0.725970352 **	0.804649137 **	0.764703274 **	0.721989063 **
1840	1	0.918096779	0.935346329	0.730125259 **	0.784043412 **	0.739978709 **	0.861795141
1847	1	0.951899691	0.936819642	0.725273324 **	0.733821637 **	0.685712473 **	0.85792347 **
1849	1	1.010627881	0.987507905	0.763191833 **	0.814143998 **	0.765484176 **	0.894625906
1853	1	0.978307999	0.90952953	0.70297251 **	0.71003492 **	0.63511686 **	0.885428075
1860	1	0.956074258	0.902584956 *	0.759138385 **	0.81173032 **	0.812488104 **	0.695381372 **

1861	1	1.003639928	0.963048004	1.128754219 **	1.046534836	1.011247454	0.943637332 *
1869	1	0.950936889	0.973353554	0.813599262 **	0.829501369 **	0.869277422 *	0.90313366
1874	1	0.861412467 *	0.802574921 *	0.623445921 **	0.6023673 **	0.631928366 **	0.805001439 **
1881	1	1.012863863	0.990545498	0.836942119 **	0.85587483 **	0.827049671 **	0.908806443
1905	1	0.983956878	1.061640243	1.01939595	1.058160118	0.975955197	1.3212637
1914	1	1.079068318	0.998137627	0.980115881	0.951738214	0.911913968	0.747892385 **
1915	1	1.040943775	0.975830314	1.249204806 **	1.347194715 **	1.319610667 **	1.057213141
1916	1	0.957782882	0.974670999	1.353922016 **	1.428272497 **	1.440001562 **	1.096011292
1925	1	0.916282413	0.916823864	1.008854421	1.116654037	1.217028525 **	1.143218123
1930	1	0.928502769	0.975653432	1.15879075 *	1.089020767	1.062239984	1.373032333 **
1932	1	0.916808864	0.930303541	0.829399218 *	0.797239321 **	0.760144006 **	0.97973839
1942	1	0.962500739	0.923414356	0.796891658 **	0.795726492 **	0.767993296 **	0.88226844
1943	1	0.971116694	0.925852322	0.810242199 **	0.90706263	0.848501289 **	0.896784778
1946	1	0.997342089	0.989494655	0.915896082	0.859915101	0.862708616 **	0.993372161
1956	1	1.005350841	0.931952793	0.853250254 **	0.861226938 **	0.82892241 **	0.855708779 **
1969	1	0.999569846	1.192988692	1.69820783 **	1.691954853 **	1.667958529 **	2.098865432 **
1990	1	0.894790776	0.96495905	0.741087529 *	0.666424566 **	0.620475898 **	0.86271083
2014	1	0.951400268	0.908276845 *	0.655604521 **	0.688431619 **	0.661710103 **	1.058064853
2020	1	0.915972582 *	0.911235551 *	0.820732726 **	0.87127537 **	0.910480109 *	0.848636009 **
2029	1	1.026008343	0.971452704	0.739580146 **	0.770265545 **	0.720884804 **	0.948548516
2033	1	0.920011902	0.934060515	0.786971352 **	0.764576227 **	0.753110036 **	0.954278517
2036	1	0.983370683	0.924235521	0.72522246 **	0.746951624 **	0.713506479 **	0.855025002 *
2038	1	0.914825331	1.00104514	0.706631088 **	0.701558313 **	0.749965399 **	0.810274665 **
2044	1	0.948117831	0.952949095	0.705822564 **	0.714505288 **	0.698295097 **	0.830592552 **
2048	1	0.981689858	0.997044865	0.739898227 **	0.715167838 **	0.698292093 **	0.853793021 *
2060	1	0.945045183	0.944176822	0.834748984 **	0.867760268 *	0.861258661 *	0.827112898 **
2073	1	0.978235402	0.953200983	0.927403956	0.918028408	0.897840889	0.78232084 **
2075	1	1.038820829	0.992636787	1.254138157 **	1.209225276 **	1.204547239 **	1.0240087
2085	1	1.049338475	1.012061081	0.803773004 *	0.700381106 **	0.667010592 **	0.848883488
2087	1	0.943132231	0.985225625	0.801883134 *	0.787095022 **	0.807762217 *	0.994379071
2092	1	1.067088764	0.991985038	1.112794528	1.200528352 **	1.205118418 **	0.933227899
2095	1	1.046225123	1.004284945	0.831105463 *	0.826429162 *	0.745234231 **	0.991458033
2107	1	0.945917986	0.915779454	0.721734261 **	0.762477735 **	0.706365188 **	0.866676098
2108	1	1.058445381	0.964972422	1.06008411	1.096622223	1.076061184	0.838477884 **
2123	1	1.105271844	1.076963775	0.7869779 *	0.787194385 *	0.695265485 **	0.951735378
2124	1	0.894963907	0.912529674	0.760242912 **	0.783437346 **	0.80265351 **	0.886981425
2125	1	0.918759872	0.947113123	0.742367986 **	0.782688227 **	0.735427986 **	0.887938717 *
2138	1	0.955179692	0.945659146	0.777109844 *	0.797383974 *	0.765563475 **	0.864494571
2140	1	0.951503863	0.965351076	0.786813207 **	0.750646984 **	0.72911288 **	0.86782299 **
2144	1	0.992717494	0.92868078	0.837785341 *	0.843534387 *	0.77213643 **	0.926744342
2159	1	1.057012906	1.024063945	1.178749912 **	1.101049193	1.139218861 *	0.801995778 **
2167	1	0.990455139	1.019220822	0.768774721 **	0.763566549 **	0.811969868 *	0.888901665
2177	1	0.971949195	0.968460686	0.872586895 **	0.849394432 **	0.85077678 **	0.934764682
2193	1	1.029057353	0.946775496	0.826659652 *	0.839214544 *	0.769677046 **	0.952393339
2199	1	1.026717049	0.989420365	0.808651115 **	0.887280497	0.813227034 **	0.848492989 *
2207	1	0.987853842	0.961041975	0.751696278 **	0.798343726 *	0.805735886 *	0.751451419 *
2210	1	0.920995514	1.032163779	0.707191386 **	0.805966723 **	0.808543319 **	1.015941041
2237	1	0.949928688	0.964996106	0.760299477 **	0.789868947 **	0.761939803 **	0.867118488 *
2249	1	0.989445804	0.998513301	1.057673349	1.022154973	1.082021005	0.76625271 **
2262	1	0.907186706	0.902928282	0.735115511 **	0.751807793 **	0.713030139 **	0.918055654
2267	1	0.916586318	1.013837462	0.797892448 **	0.875812482 *	0.83741585 **	1.006959622
2283	1	0.984013283	0.991204561	0.817415054 **	0.811951918 **	0.812205832 **	0.848684117 **
2289	1	0.967134108	0.992201362	0.799109954 **	0.842702765 *	0.837880774 *	0.921403718
2290	1	1.003314716	1.0232075	0.854505874 **	0.877761343 *	0.867892772 *	0.936255707
2294	1	0.978268953	0.990121106	0.820006088 **	0.842422924 **	0.822570474 **	0.868571607 **
2300	1	0.955207945	0.91762332	0.827944101 **	0.839307659 **	0.791266267 **	0.885301049 *
2307	1	0.879295768 *	0.907909543	0.741509691 **	0.727928157 **	0.745377136 **	0.949532996
2309	1	0.921434817	0.980844441	0.710544907 **	0.705484097 **	0.712438472 **	0.864551471 **
2321	1	1.019390379	0.920121733	0.966619636	1.027202992	1.050594201	0.800567674 **
2330	1	0.974387501	0.987626873	0.766357715 **	0.833896531 **	0.833775681 **	0.889718512
2331	1	0.885513231 *	1.002097299	0.716495792 **	0.722008312 **	0.781897167 **	0.76343078 **
2333	1	0.886770104	0.931270199	0.744706582 **	0.846371467	0.853139216	1.049982498
2334	1	0.996457617	0.913991487	1.267115666 **	1.333716748 **	1.447853277 **	1.024324437
2345	1	0.986408324	0.971001078	1.367711167 **	1.354661888 **	1.371118312 **	1.035188426

2356	1	0.926418349	0.961587599	0.709666196 **	0.720605719 **	0.671035976 **	0.880526115 *
2365	1	1.002280971	1.045749993	1.320797281 **	1.229150642 **	1.215813488 **	0.966036947
2366	1	0.962982256	0.907166495 *	0.98531294	0.850257378 **	0.835748587 **	0.959484609
2368	1	0.95478015	1.14951964 *	1.693194016 **	1.744821939 **	1.98740848 **	1.468099934 **
2372	1	1.037770319	1.412025792 *	2.377484431 **	2.479273466 **	2.637947134 **	2.108662084 **
2378	1	1.054506403	1.461588971 **	2.062888857 **	1.797087347 **	1.693574337 **	1.62090956 **
2382	1	0.978062291	1.42507559 **	1.900307374 **	1.816275101 **	1.75307431 **	1.646572245 **
2385	1	0.979305883	0.994065834	1.204195076 **	1.142243395 **	1.173249624 **	1.111782408 *
2395	1	0.960358168	0.980434317	0.862103439 **	0.885961602 *	0.845914659 **	1.005415379
2396	1	0.988735826	0.994475419	0.834310338 **	0.821872266 **	0.841081086 **	0.863517299 *
2403	1	0.944160558	1.019433996	0.855744766 **	0.861833575 **	0.855552524 **	0.898957388
2406	1	0.910289133	0.913711191 *	0.733286608 **	0.758919904 **	0.749558536 **	0.908609027
2412	1	0.927481072	1.309500961 **	1.663209649 **	1.586228504 **	1.583875034 **	1.41583467 **
2420	1	0.934702028	1.01231586	0.799825426 **	0.787292245 **	0.761689007 **	0.967993492
2450	1	0.978538492	1.003653922	0.803878291 **	0.818636815 **	0.84162512 **	0.917508245
2455	1	0.929763363	0.980875266	0.831143036 **	0.868597113 *	0.85718655 **	0.945613672
2462	1	0.949839337	0.937768081	0.811104307 **	0.837141481 **	0.82377987 **	0.890118742 *
2472	1	0.982323953	0.961304962	0.831989969 **	0.878957555 *	0.8509144 **	0.924383895
2475	1	0.987422887	0.874406308	1.140512849	1.459924342 **	1.31584943 **	1.079175135
2477	1	1.00265394	0.974206744	0.843673849 **	0.908980091 *	0.883524503 **	1.005816522
2482	1	0.94619354	0.977431671	0.768171888 **	0.819554173 **	0.816403803 **	0.871043168 *
2492	1	1.028703759	0.995836765	1.200900586 **	1.153682458 *	1.147406098 *	0.948834736
2493	1	0.916069578	0.961187119	0.789057676 **	0.901452534	0.844178163 *	1.013096546
2494	1	0.970042836	0.99852025	0.781918186 **	0.774787374 **	0.765182578 **	0.846329165 *
2498	1	1.045015026	1.006032433	0.823208071	0.793292387 *	0.742555313 **	0.960791016
2502	1	0.979263028	0.985211797	1.12563672 **	1.135864645 **	1.101448551 *	1.016121069
2541	1	1.146059678	1.109758632	1.37957701 **	1.483632669 **	1.265334543 *	1.079180394
2550	1	0.884677173	0.904744261	0.62049134 **	0.594001086 **	0.607665373 **	0.817951548 **
2563	1	0.959783573	0.972686153	0.823475423 **	0.846506255 *	0.807703186 **	0.90874489
2566	1	0.93071974	1.027742819	0.825209225 **	0.730180516 **	0.766189365 **	0.924181582

The mean ratios of identified spots between control and inhibitor-treated cells are listed. Non-repeated measures ANOVA and Dunnett's test for post hoc analysis were performed. Asterisks indicate significant differences from respective controls (*P < 0.05, ** P < 0.01).

Table S2. Identified Spots by Peptide Mass Fingerprinting (PMF)

Master No.	Accession No. of Swiss-Prot	Gene	Protein Name	Score	Sequence Coverage (%)
647	P08238	<i>HSP90AB1</i>	Heat shock protein HSP 90-beta	105	39
713	P13639	<i>EEF2</i>	Elongation factor 2	175	40
714	P13639	<i>EEF2</i>	Elongation factor 2	170	37
970	P02545	<i>LMNA</i>	Lamin-A/C	325	53
971	P11021	<i>HSPA5</i>	78 kDa glucose-regulated protein precursor (GRP78)	179	31
983	P11021	<i>HSPA5</i>	78 kDa glucose-regulated protein precursor (GRP78)	237	44
987	P02545	<i>LMNA</i>	Lamin-A/C	158	32
998	P38646	<i>HSPA9B</i>	Stress-70 protein, mitochondrial precursor	192	47
1114	P08107	<i>HSPA1B</i>	Heat shock 70 kDa protein 1 (HSP70)	129	26
1127	P08107	<i>HSPA1B</i>	Heat shock 70 kDa protein 1 (HSP70)	195	44
1227	P61978	<i>HNRNPK</i>	Heterogeneous nuclear ribonucleoprotein K	176	41
1295	P10809	<i>HSPD1</i>	60 kDa heat shock protein, mitochondrial precursor	270	69
1429	Q01518	<i>CAP1</i>	Adenylyl cyclase-associated protein 1	104	29
1484	Q16658	<i>FSCN1</i>	Fascin	157	38
1579	Q15084	<i>PDIA6</i>	Protein disulfide-isomerase A6 precursor	66	32
1758	P02571/ P60709	<i>ACTG1/ ACTB</i>	Actin, cytoplasmic 2 / 1	160	68
1767	P02571	<i>ACTG1</i>	Actin, cytoplasmic 2	135	55
2014	P52895	<i>AKRIC2</i>	Aldo-keto reductase family 1 member C2	136	50
2372	P04792	<i>HSPB1</i>	Heat-shock protein beta-1 (HSP27)	92	56
2382	P04792	<i>HSPB1</i>	Heat-shock protein beta-1 (HSP27)	128	40

Selected protein spots were digested with trypsin and identified by PMF as described in Materials and Methods. The Mascot search program was used to search the Swiss-Prot database for peptide masses.

Methods for Table S2

Protein Determination by Peptide Mass Fingerprinting (PMF)

To identify the protein in the spots, we performed PMF with matrix-assisted laser desorption/ionization time-of-flight (MALDI-TOF) mass analysis with a slight modification (Kanoh, et al., 2005). After HeLa cell lysates (500 µg of protein) were subjected to two-dimensional electrophoresis, the gels were stained with SYPRO Ruby (Invitrogen, Carlsbad, CA) according to the manufacturer's protocol. Spot maps of the images were matched to the master gel image using the DIA and BVA modules of DeCyder 6.5.

The protein-containing region was picked using an Ettan Spot Picker (GE Healthcare) and washed successively with water, decolorant (50% acetonitrile and 50 mM NH₄HCO₃), and acetonitrile. The samples were then dried *in vacuo*. An aqueous alkylating solution (100 mM iodoacetamide and 100 mM NH₄HCO₃) was added to the dried gels, and the mixture was incubated at 37°C for 30 min. The gels were then washed with 100 mM NH₄HCO₃ solution, dehydrated with acetonitrile, and dried *in vacuo*. The dried gels were seeded in digestion buffer (10 mM Tris-HCl (pH 8.8) and 25 ng/mL trypsin) at 4°C for 30 min and incubated at 37°C overnight.

Samples were analyzed using a Bruker Reflex II MALDI-TOF mass spectrometer, operated in the positive ion reflector mode. Here, α-Cyano-4-hydroxycinnamic acid (10 mg/mL) dissolved in 50% acetonitrile/0.1% trifluoroacetic acid was used as the matrix. The Mascot search program (Matrix Science Inc., Boston, MA) was used to search the Swiss-Prot database for peptide masses.

Reference

Kanoh, N., Honda, K., Simizu, S., Muroi, M., and Osada, H. (2005). Photo-cross-linked small molecule affinity matrix for facilitating forward and reverse chemical genetics. *Angew. Chem. Int. Ed.* 44, 3559-3562.

Results of protein determination by PMF

Search Parameters on MASCOT Peptide Mass Fingerprint

Type of search : Peptide Mass Fingerprint
Database : SwissProt
Taxonomy : Homo sapiens (human)
Enzyme : Trypsin
Fixed modifications : Carbamidomethyl (C)
Variable modifications : Oxidation (M)
Mass values : Monoisotopic
Protein Mass : Unrestricted
Peptide Mass Tolerance : ± 150 ppm
Peptide Charge State : 1+
Max Missed Cleavages : 1

Master No. 647

Match to: HS90B_HUMAN

Score: 105

Expect: 5e-007

Heat shock protein HSP 90-beta

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
822.3542	821.3469	821.3953	-58.8	1	553	-	559	K.MEESKAK.F	
829.4785	828.4712	828.5221	-61.5	0	331	-	337	R.ALLFIPR.R	
901.4813	900.474	900.5181	-49	0	285	-	291	K.TKPIWTR.N	
951.4318	950.4245	950.457	-34.1	0	169	-	177	R.ADHGEPIGR.G	
1045.5406	1044.5333	1044.5927	-56.9	1	575	-	583	K.VEKVTISNR.L	
1080.5049	1079.4976	1079.5287	-28.8	0	339	-	347	R.APFDLFENK.K	
1141.5443	1140.537	1140.5523	-13.4	0	439	-	448	K.LGIHEDSTNR.R	
1151.5483	1150.541	1150.5506	-8.33	0	276	-	284	K.YIDQEELNK.T	
1194.6359	1193.6286	1193.6404	-9.87	0	73	-	82	K.IDIIPNPQER.T	
1236.6296	1235.6223	1235.6299	-6.11	1	338	-	347	R.RAPFDLFENK.K	
1249.6087	1248.6014	1248.6098	-6.78	0	492	-	502	K.EQVANSAFVER.V	
1264.6377	1263.6304	1263.4934	108	0	613	-	623	R.DNSTMGYMMAK.K	Oxidation
1311.5646	1310.5573	1310.5626	-4.06	0	187	-	196	K.EDQTEYLEER.R	
1348.6625	1347.6552	1347.6572	-1.42	0	320	-	330	K.HFSVEGQLEFRA	
1513.7787	1512.7714	1512.7784	-4.59	0	379	-	392	R.GVVDSEDLPLNISRE	
1527.7434	1526.7362	1526.7365	-0.24	0	307	-	319	K.SLTNDWEDHLAVK.H	
1782.9398	1781.9325	1781.9424	-5.57	0	625	-	639	K.HLEINPDHPIVETLR.Q	
1808.9517	1807.9444	1807.9509	-3.55	0	205	-	219	K.HSQFIGYPITLYLEK.E	
1847.7891	1846.7818	1846.7897	-4.29	0	292	-	306	R.NPDDITQEEYGEFYK.S	
2192.9625	2191.9552	2191.9328	10.2	0	457	-	475	R.YHTSQSGDEMTSLSEYVSR.M	Oxidation
2255.9538	2254.9466	2254.9516	-2.21	0	149	-	168	K.HNDDEQYAWESSAGGSFTVR.A	
2448.1561	2447.1488	2447.1389	4.07	0	507	-	526	R.GFEVVYMTPEIDYCVQQLK.E	

Master No. 713

Match to: EF2_HUMAN

Score: 175

Expect: 5e-014

Elongation factor 2

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
890.5946	889.5874	889.5022	95.8	0	499	-	506	K.FSVSPVVR.V	
912.5057	911.4984	911.4092	97.9	1	253	-	259	K.VEDMMKK.L	2 Oxidation
922.5402	921.5329	921.4556	83.9	0	573	-	580	K.SDPVVSYLE	
969.6474	968.6401	968.5403	103	0	717	-	726	R.GGGQIIP.TAR.R	
1013.618	1012.6107	1012.4938	116	0	240	-	249	K.GEGQLGPAERA	
1091.7091	1090.7019	1090.5771	114	0	2	-	10	M.VNFTVDQIR.A	
1100.6478	1099.6405	1099.5372	93.9	0	300	-	308	K.VFDAIMNFK.K	Oxidation
1107.7217	1106.7144	1106.6336	73	0	416	-	426	R.VFSGLVSTGLK.V	
1138.6682	1137.6609	1137.5666	82.9	1	273	-	283	K.FSKSATSPGK.K	
1224.6582	1223.6509	1223.491	131	0	689	-	698	K.EGALCEENMR.G	Oxidation
1274.8566	1273.8493	1273.703	115	0	440	-	449	K.EDLYLKPIQR.T	
1308.8132	1307.8059	1307.651	118	0	677	-	688	K.DSVVAGFQWATK.E	
1323.7849	1322.7776	1322.6401	104	0	21	-	32	R.NMSVIAHVDHGK.S	Oxidation
1378.8767	1377.8695	1377.7075	118	0	728	-	739	R.CLYASVLTAPRL	
1402.9566	1401.9493	1401.798	108	1	439	-	449	K.EDLYLKPIQR.T	
1444.8787	1443.8714	1443.7609	76.6	1	846	-	858	K.EGIPALDNFLDKL-	
1543.935	1542.9278	1542.7678	104	0	606	-	619	K.ARPFPDGLAEDIDK.G	
1567.9209	1566.9137	1566.7712	90.9	1	560	-	572	K.DLEEDHACIPK.K	
1615.9789	1614.9716	1614.7573	133	0	482	-	495	K.TGTITTFEHAHNM.R.V	
1631.9023	1630.8951	1630.7522	87.6	0	482	-	495	K.TGTITTFEHAHNM.R.V	Oxidation
1800.0608	1799.0535	1798.889	91.5	0	786	-	801	K.AYLPVNESFGFTADLR.S	
1978.0884	1977.0811	1976.9666	57.9	0	768	-	785	R.GHVFEESQVAGTPMFVVK.A	Oxidation
2143.2056	2142.1984	2142.0705	59.7	1	606	-	625	K.ARPFPDGLAEDIDKGEVSAR.Q	
2204.229	2203.2217	2203.1048	53	0	72	-	90	K.STAISLFYELSENDLNFIK.Q	
2220.308	2219.3007	2219.1474	69.1	0	163	-	180	R.ALLELQLEPEELYQTFQRI	
2233.2816	2232.2743	2232.1249	66.9	1	648	-	667	R.KIWCFGPDGTGNILTDITK.G	
2343.1946	2342.1874	2342.0558	56.2	1	367	-	386	K.YRCELLYEGPPDDEAAMGIK.S	Oxidation
2576.4418	2575.4345	2575.3219	43.7	1	288	-	308	R.TFCQLIDPIFKVFDAIMNFK.K	Oxidation
2801.5756	2800.5684	2800.4032	59	0	94	-	120	K.DGAGFLINLIDSPGHVDFSSEVTAALR.V	

Master No. 714

Match to: EF2_HUMAN

Score: 170

Expect: 1.6e-013

Elongation factor 2

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
890.4837	889.4764	889.5022	-29	0	499	-	506	K.FSVSPVVR.V	
911.3738	910.3665	910.4185	-57.1	0	265	-	272	R.YFDPANGK.F	
922.4431	921.4358	921.4556	-21.4	0	573	-	580	K.SDPVVSYLE	
969.5122	968.5049	968.5403	-36.6	0	717	-	726	R.GGGQIIPAR.R	
1013.4999	1012.4926	1012.4938	-1.12	0	240	-	249	K.GEGQLGPAER.A	
1063.494	1062.4867	1062.5709	-79.2	0	668	-	676	K.GVQYLNEIK.D	
1091.5628	1090.5555	1090.5771	-19.8	0	2	-	10	M.VNFTVDQIR.A	
1100.5006	1099.4934	1099.5372	-39.9	0	300	-	308	K.VFDAIMNF.K	Oxidation
1102.4901	1101.4829	1101.5165	-30.5	0	227	-	235	K.QFAEMYVAK.F	Oxidation
1107.5846	1106.5774	1106.6336	-50.8	0	416	-	426	R.VFSGLVSTGLK.V	
1138.4985	1137.4912	1137.5091	-15.7	0	639	-	647	K.YEWDVAEARK	
1274.6997	1273.6924	1273.703	-8.34	0	440	-	449	K.EDLYLKPIQR.T	
1378.7096	1377.7023	1377.7075	-3.73	0	728	-	739	R.CLYASVLTAPRL	
1402.7352	1401.728	1401.798	-49.9	1	439	-	449	K.KEDLYLKPIQR.T	
1444.6814	1443.6741	1443.7609	-60.1	1	846	-	858	K.EGIPALDNFLDKL-	
1475.7222	1474.7149	1474.7822	-45.6	1	151	-	162	R.IKPVLMMNKMDRA	
1494.7169	1493.7097	1493.7952	-57.3	0	288	-	299	R.TFCQLILDPIFK.V	
1631.6991	1630.6918	1630.7522	-37	0	482	-	495	K.TGTITTFEHAHNMR.V	Oxidation
1799.8376	1798.8304	1798.889	-32.6	0	786	-	801	K.AYLPVNESFGFTADLR.S	
2142.9783	2141.971	2142.0705	-46.5	1	606	-	625	K.ARPFPDGLAEDIDKGEVSAR.Q	
2203.9784	2202.9712	2203.1048	-60.7	0	72	-	90	K.STAISLFYELSENDLNFIK.Q	
2220.0725	2219.0653	2219.1474	-37	0	163	-	180	R.ALLELQLEPEELYQTFQRI	
2576.2287	2575.2214	2575.2986	-30	0	121	-	144	R.VTDGALVVVDCVSGVCVQTETVLR.Q	
2801.3551	2800.3478	2800.4032	-19.8	0	94	-	120	K.DGAGFLINLIDSPGHVDFSSEVTAAALR.V	
3005.5046	3004.4973	3004.5402	-14.3	0	740	-	765	R.LMEPIYLVEIQCPQVVGIIYGLNR.K	Oxidation

Master No. 970

Match to: LMNA_HUMAN

Score: 325

Expect: 5e-029

Lamin-A/C

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
807.4807	806.4734	806.4286	55.5	1	109	-	114	K.VREEFK.E	
849.5269	848.5196	848.4756	51.9	0	42	-	48	R.LAVYIDR.V	
859.4774	858.4701	858.4308	45.9	1	190	-	196	R.RVDAENR.L	
919.4682	918.4609	918.4308	32.8	0	428	-	435	R.SSFSQHAR.T	
949.5205	948.5133	948.4876	27.1	1	420	-	427	R.KLESTESR.S	
974.4986	973.4913	973.4716	20.3	0	379	-	386	K.LLEGEER.L	
1023.5209	1022.5137	1022.5032	10.2	0	209	-	216	K.NIYSEELR.E	
1028.5871	1027.5799	1027.5662	13.3	0	241	-	249	R.LADALQELR.A	
1029.7433	1028.736	1028.5866	145	1	172	-	181	K.LEAALGEAKK.Q	
1043.558	1042.5507	1042.5407	9.6	0	124	-	133	K.EGDLIAAQR.L	
1048.5154	1047.5082	1047.5019	6.01	0	182	-	189	K.QLQDEMLR.R	Oxidation
1089.5589	1088.5516	1088.5462	5.02	0	51	-	60	R.SLETENAGLR.L	
1131.5159	1130.5086	1130.5204	-10.4	0	33	-	41	K.EDLQELNDR.L	
1148.5728	1147.5655	1147.5721	-5.69	0	63	-	72	R.ITESEEVVSR.E	
1165.5385	1164.5312	1164.5411	-8.46	0	79	-	89	K.AAYEALGDAR.K	
1171.6189	1170.6117	1170.6357	-20.5	1	123	-	133	K.KEGDLIAAQR.L	
1176.6005	1175.5932	1175.5968	-3.08	1	181	-	189	K.QLQDEMLR.R	Oxidation
1182.5943	1181.587	1181.604	-14.4	0	157	-	166	R.TLEGELHDLR.G	
1187.6207	1186.6134	1186.6306	-14.5	1	320	-	329	K.LRDLEDLAR.E	
1243.6975	1242.6902	1242.7183	-22.6	1	134	-	144	R.LKDLEALLNSK.E	
1291.5899	1290.5826	1290.6204	-29.3	0	472	-	482	R.QNGDDPLLTYR.F	
1293.7084	1292.7012	1292.636	50.4	1	79	-	90	K.AAYEALGDAR.K	
1331.625	1330.6177	1330.6703	-39.5	0	367	-	377	K.LALDMEIHAYR.K	
1347.6241	1346.6169	1346.6652	-35.9	0	367	-	377	K.LALDMEIHAYR.K	Oxidation
1359.645	1358.6377	1358.679	-30.4	0	12	-	25	R.SGAQASSTPLSPTR.I	
1363.576	1362.5687	1362.6099	-30.2	0	516	-	527	K.AQNTWGCNSLR.T	
1430.7239	1429.7167	1429.7776	-42.7	0	299	-	311	R.IDSLSAQLSQLQK.Q	
1502.6455	1501.6382	1501.7161	-51.9	1	250	-	261	R.AQHEDQVEQYKK.E	
1507.6597	1506.6524	1506.7348	-54.7	0	528	-	541	R.TALINSTGEEVAMR.K	Oxidation
1525.6828	1524.6755	1524.7494	-48.4	1	197	-	208	R.LQTMKEELDFQK.N	Oxidation
1566.67	1565.6627	1565.7434	-51.6	0	628	-	644	R.SVGGSGGSGFDNLVTR.S	
1605.7232	1604.7159	1604.8046	-55.3	1	440	-	453	R.VAVEEVDEEGKFVR.L	
1629.7149	1628.7077	1628.8005	-57	1	29	-	41	R.LQEKEDLQELNDR.L	
1752.7387	1751.7314	1751.855	-70.6	0	281	-	296	R.NSNLVGAAHEELQQSR.I	
1794.6848	1793.6775	1793.8002	-68.4	1	456	-	470	R.NKSNEDQSMGNWQIK.R	Oxidation
2364.9714	2363.9642	2364.1517	-79.3	0	598	-	624	K.ASASGSGAQVGGPISSGSSASSVTVTR.S	

Master No. 971

Match to: GRP78_HUMAN

Score: 179

Expect: 2.5e-014

78 kDa glucose-regulated protein

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
919.4897	918.4824	918.4633	20.8	0	262	-	268	R.VMEHFIK.L	Oxidation
986.5055	985.4982	985.508	-9.96	0	533	-	540	R.LTPEEIER.M	
997.5076	996.5003	996.5101	-9.81	0	298	-	306	R.ALSSQHQARI	
1074.5372	1073.53	1073.5465	-15.4	0	524	-	532	K.ITITNDQNR.L	
1191.591	1190.5838	1190.6295	-38.4	0	465	-	474	K.VYEGERPLTK.D	Oxidation
1210.5304	1209.5231	1209.5778	-45.2	1	377	-	386	K.EFFNGKEPSR.G	
1228.5744	1227.5672	1227.6207	-43.7	0	50	-	60	R.VEIANDQGNRI	
1313.5233	1312.516	1312.6122	-73.3	0	327	-	336	K.FEELNMDLFR.S	
1316.5674	1315.5602	1315.6295	-52.7	0	563	-	573	R.NELESYAYSLK.N	Oxidation
1329.5497	1328.5425	1328.6071	-48.6	0	327	-	336	K.FEELNMDLFR.S	
1460.6585	1459.6512	1459.7518	-68.9	0	354	-	367	K.SDIDEIVLVGGSTR.I	
1512.6002	1511.5929	1511.7442	-100	1	325	-	336	R.AKFEELNMDLFR.S	
1528.5947	1527.5874	1527.7391	-99.3	1	325	-	336	R.AKFEELNMDLFR.S	Oxidation
1566.6647	1565.6574	1565.7726	-73.5	0	61	-	74	R.ITPSYVAFTPEGER.L	
1588.7337	1587.7264	1587.8468	-75.8	1	353	-	367	K.KSDIDEIVLVGGSTR.I	
1604.7575	1603.7502	1603.857	-66.6	0	124	-	138	K.TKPYIQVDIGGGQTK.T	
1677.6776	1676.6703	1676.8006	-77.7	0	82	-	96	K.NQLTSNPENTVFDAK.R	Oxidation
1815.8203	1814.8131	1814.989	-96.9	1	198	-	214	R.IINEPTAAAIAYGLDKR.E	
1887.7987	1886.7914	1886.9639	-91.4	0	165	-	181	K.VTHAVVTVPAYFNDAQR.Q	
1933.8342	1932.8269	1933.0058	-92.5	0	475	-	492	K.DNHLLGTFDLTGIPPAPR.G	
1974.7242	1973.7169	1973.9007	-93.1	0	602	-	617	K.IEWLESHQDADIEDFK.A	Oxidation
2164.7698	2163.7625	2163.9848	-103	0	307	-	324	R.IEIESFYEGEDFSETLTRA	
2175.7792	2174.7719	2174.9855	-98.2	1	634	-	654	K.LYGSAGPPPTGEEDTAEKDEL.	

Master No. 983

Match to: GRP78_HUMAN (P11021)

Score: 237

Expect: 3.1e-020

78 kDa glucose-regulated protein precursor

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
1228.6066	1227.5994	1227.6207	-17	0	50	-	60	R.VEIIANDQGNRI	
1566.7173	1565.71	1565.7726	-40	0	61	-	74	R.ITPSYVAFTPEGERL	
1677.736	1676.7288	1676.8006	-43	0	82	-	96	K.NQLTSNPENTVFDAK.R	
1430.6339	1429.6266	1429.6838	-40	0	102	-	113	R.TWNDPSVQQDIK.F	
1604.7912	1603.784	1603.857	-46	0	124	-	138	K.TKPYIQVDIGGQTK.T	
1887.8754	1886.8682	1886.9639	-51	0	165	-	181	K.VTHAVVTVPAYFNDAQR.Q	
1217.6106	1216.6033	1216.6234	-17	0	186	-	197	K.DAGTIAGLNVMRI	
1233.5924	1232.5851	1232.6183	-27	0	186	-	197	K.DAGTIAGLNVMRI	Oxidation
1815.9044	1814.8972	1814.989	-51	1	198	-	214	R.IINEPTAAAIAYGLDKR.E	
919.476	918.4687	918.4633	6	0	262	-	268	R.VMEHFIK.L	Oxidation
997.5293	996.522	996.5101	12	0	298	-	306	R.ALSSQHQARI	
2164.887	2163.8797	2163.9848	-49	0	307	-	324	R.IEIESFYEGEDFSETLTRA	
1512.6971	1511.6899	1511.7442	-36	1	325	-	336	R.AKFEELNMDLFR.S	
1528.6851	1527.6778	1527.7391	-40	1	325	-	336	R.AKFEELNMDLFR.S	Oxidation
1313.5831	1312.5758	1312.6122	-28	0	327	-	336	K.FEELNMDLFR.S	
1329.5624	1328.5552	1328.6071	-39	0	327	-	336	K.FEELNMDLFR.S	Oxidation
1588.7828	1587.7755	1587.8468	-45	1	353	-	367	K.KSDIDEIVLVGGSTR.I	
1460.7055	1459.6982	1459.7518	-37	0	354	-	367	K.SDIDEIVLVGGSTR.I	
1210.561	1209.5537	1209.5778	-20	1	377	-	386	K.EFFNGKEPSR.G	
1836.8961	1835.8888	1835.9265	-21	0	448	-	464	K.SQIFSTASDNQPTVTIK.V	
1191.6131	1190.6058	1190.6295	-20	0	465	-	474	K.VYEGERPLTK.D	
1933.9149	1932.9076	1933.0058	-51	0	475	-	492	K.DNHLLGTFDLTGIPPAPR.G	
1074.5433	1073.536	1073.5465	-10	0	524	-	532	K.ITITNDQNR.L	
986.5213	985.5141	985.508	6	0	533	-	540	R.LTPEEIER.M	
1316.5939	1315.5866	1315.6295	-33	0	563	-	573	R.NELESYAYSLK.N	
1974.8067	1973.7995	1973.9007	-51	0	602	-	617	K.IEWLESHQDADIEDFK.A	
2175.9134	2174.9062	2174.9855	-36	1	634	-	654	K.LYGSAGPPPTGEEDTAEKDEL-	

Master No. 987

Match to: LMNA_HUMAN

Score: 158

Expect: 2.5e-012

Lamin-A/C (70 kDa lamin)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
1023.6618	1022.6545	1022.5032	0	148	209	-	216	K.NIYSEELRE	
1028.7189	1027.7117	1027.5662	0	142	241	-	249	R.LADALQELRA	
1165.7018	1164.6945	1164.5411	0	132	79	-	89	K.AAYEAEELGDARK	
1171.7955	1170.7883	1170.6357	1	130	226	-	235	R.LVEIDNGKQRE	
1182.781	1181.7737	1181.604	0	144	157	-	166	R.TLEGELHDLRG	
1187.7893	1186.782	1186.6306	1	128	320	-	329	K.LRDLEDSLARE	
1243.885	1242.8777	1242.7183	1	128	134	-	144	R.LKDLEALLNSKE	
1291.8011	1290.7939	1290.6204	0	134	472	-	482	R.QNGDDPLLTYRF	
1293.8117	1292.8045	1292.636	1	130	79	-	90	K.AAYEAEELGDARK.T	
1347.8375	1346.8303	1346.6652	0	123	367	-	377	K.LALDMEIHAYRK	Oxidation
1475.9613	1474.954	1474.7602	1	131	367	-	378	K.LALDMEIHAYRK.L	Oxidation
1502.9197	1501.9125	1501.7161	1	131	250	-	261	R.AQHEDQVEQYKKE	
1507.9326	1506.9253	1506.7348	0	126	528	-	541	R.TALINSTGEEVAMRK	Oxidation
1525.9553	1524.948	1524.7494	1	130	197	-	208	R.LQTMKEELDFQKN	Oxidation
1566.9693	1565.962	1565.7434	0	140	628	-	644	R.SVGGSGGGSFGDNLVTRS	
1606.0389	1605.0316	1604.8046	1	141	440	-	453	R.VAVEEVDEEGKFVRL	
1630.0424	1629.0351	1628.8005	1	144	29	-	41	R.LQEKEDLQELNDR.L	
1666.1317	1665.1245	1664.8846	1	144	157	-	171	R.TLEGELHDLRGQVAK.L	
1700.2207	1699.2134	1698.9628	1	147	297	-	311	R.IRIDSLSAQLSQLQK.Q	
1761.1529	1760.1457	1759.8893	1	146	472	-	486	R.QNGDDPLLTYRFPPK.F	
1910.2012	1909.1939	1908.9139	0	147	352	-	366	R.MQQQLDEYQELLDIK.L	Oxidation

Master No. 998

Match to: GRP75_HUMAN

Score: 192

Expect: 9.9e-016

Stress-70 protein, mitochondrial precursor

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
938.4454	937.4381	937.4253	13.6	1	568	-	574	K.YAEEDRR.K	
958.5312	957.5239	957.4879	37.5	0	77	-	85	K.VLENAEGAR.T	
1045.5718	1044.5645	1044.5314	31.7	1	647	-	654	K.LFEMAYKK.M	Oxidation
1149.5528	1148.5455	1148.5462	-0.62	1	127	-	135	R.RYDDPEVQK.D	
1242.6713	1241.664	1241.6728	-7.08	0	207	-	218	K.DAGQISGLNVL.R.V	
1290.6784	1289.6711	1289.6728	-1.33	0	395	-	405	K.VQQTIVQDLFGR.A	
1333.6244	1332.6171	1332.631	-10.4	0	176	-	187	K.ETAENYLGHAK.N	
1341.5991	1340.5918	1340.6109	-14.3	0	147	-	159	R.ASNGDAWVEAHGK.L	
1361.7404	1360.7331	1360.7351	-1.46	0	349	-	360	R.AQFEGIVTDLIR.R	
1446.7644	1445.7571	1445.7548	1.57	0	378	-	391	K.SDIGEVILVGGMTR.M	
1450.719	1449.7118	1449.71	1.22	0	86	-	99	R.TTPSVVAFTADGER.L	
1462.7538	1461.7466	1461.7497	-2.18	0	378	-	391	K.SDIGEVILVGGMTR.M	Oxidation
1568.8018	1567.7946	1567.7631	20.1	0	108	-	121	R.QAVTNPNTFYATK.R	
1592.9679	1591.9606	1591.945	9.8	0	499	-	513	K.LLGQFTLIGIPPAPR.G	
1608.7877	1607.7804	1607.7613	11.9	1	174	-	187	K.MKETAENYLGHAK.N	Oxidation
1645.9021	1644.8948	1644.8723	13.7	0	219	-	234	R.VINEPTAAALAYGLDK.S	
1694.8744	1693.8671	1693.8424	14.6	0	188	-	202	K.NAVITVPAYFNDSQR.Q	
1808.938	1807.9307	1807.8952	19.6	0	469	-	485	K.SQVFSTAADGQTQVEIK.V	
1872.9269	1871.9197	1871.8935	14	0	579	-	595	R.VEAVNMAEGIIHDTETK.M	Oxidation
2055.992	2054.9847	2054.9545	14.7	0	266	-	284	K.STNGDTFLGGEDFDQALLR.H	
2233.1458	2232.1385	2232.0982	18.1	1	626	-	646	K.DSETGENIRQAASSLQQASLK.L	
2251.2585	2250.2512	2250.2148	16.2	0	239	-	259	K.VIAYVDLGGGTFDISILEIQK.G	
2309.25	2308.2427	2308.2063	15.8	0	514	-	535	R.GVPQIEVTFDIDANGIVHSAK.D	
2658.3024	2657.2951	2657.22	28.3	0	317	-	340	K.CELSSSVQTDINLPYLTMDSSGPK.H	Oxidation

Master No. 1114

Match to: HSP71_HUMAN (P08107)

Score: 129

Expect: 2e-009

Heat shock 70 kDa protein 1 (HSP70.1)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
1228.5866	1227.5794	1227.6207	-34	0	26	-	36	K.VEIIANDQGNRT	
1487.6169	1486.6096	1486.694	-57	0	37	-	49	R.TTPSYVAFTDTERL	
1658.7357	1657.7285	1657.8424	-69	0	57	-	71	K.NQVALNPQNTVFDAK.R	
1197.6365	1196.6292	1196.6877	-49	0	160	-	171	K.DAGVIAGLNVLR.I	
1687.8165	1686.8092	1686.894	-50	0	172	-	187	R.IINEPTAAAIAYGLDR.T	
1675.6449	1674.6376	1674.7234	-51	0	221	-	236	K.ATAGDTHLGGEDFDNR.L	
1417.6794	1416.6721	1416.7514	-56	1	237	-	247	R.LVNHFVEEFKR.K	
2981.5576	2980.5504	2980.4553	32	0	273	-	299	R.TLSSSTQASLEIDSLFEGIDFYTSITR.A	
1542.6492	1541.642	1541.7296	-57	1	300	-	311	R.ARFEELCSDLFR.S	
1315.5437	1314.5365	1314.5914	-42	0	302	-	311	R.FEELCSDLFR.S	
1465.7337	1464.7264	1464.8049	-54	0	329	-	342	K.AQIHDLVLVGGSTR.I	
1109.5372	1108.5299	1108.5665	-33	0	349	-	357	K.LLQDFFNGR.D	
801.4184	800.4111	800.414	-4	0	452	-	458	K.DNNLLGR.F	
1137.5119	1136.5046	1136.5462	-37	1	525	-	533	K.YKAEDEVQRE	

Master No. 1127

Match to: HSP71_HUMAN (P08107)

Score: 195

Expect: 5e-016

Heat shock 70 kDa protein 1 (HSP70.1)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
1228.6076	1227.6003	1227.6207	-17	0	26	-	36	K.VEIIANDQGGR.T	
1487.6613	1486.6541	1486.694	-27	0	37	-	49	R.TTPSYVAFTDTER.L	
1658.8256	1657.8184	1657.8424	-14	0	57	-	71	K.NQVALNPQNTVFDK.R	
1680.7906	1679.7833	1679.842	-35	0	89	-	102	K.HWPFQVINDGDKPK.V	
1630.7659	1629.7586	1629.796	-23	0	113	-	126	K.AFYPEEISSMVLTK.M	Oxidation
3001.7838	3000.7765	3000.4869	97	0	129	-	155	K.EIAEAYLGYPVTNAVITVPAYFNDSQR.Q	
1197.6644	1196.6571	1196.6877	-26	0	160	-	171	K.DAGVIAGLNVLR.I	
1687.855	1686.8478	1686.894	-27	0	172	-	187	R.IINEPTAAAIAYGLDR.T	
1675.691	1674.6837	1674.7234	-24	0	221	-	236	K.ATAGDTHLGGEDFDNR.L	
1261.6421	1260.6349	1260.6503	-12	0	237	-	246	R.LVNHFVEEFK.R	
1417.725	1416.7177	1416.7514	-24	1	237	-	247	R.LVNHFVEEFKR.K	
2981.7148	2980.7076	2980.4553	85	0	273	-	299	R.TLSSSTQASLEIDSLFEGIDFYTSITR.A	
1542.6927	1541.6854	1541.7296	-29	1	300	-	311	R.ARFEELCSDLFR.S	
1315.5731	1314.5658	1314.5914	-19	0	302	-	311	R.FEELCSDLFR.S	
1465.7777	1464.7704	1464.8049	-24	0	329	-	342	K.AQIHDLVLVGGSTR.I	
1109.5644	1108.5572	1108.5665	-8	0	349	-	357	K.LLQDFFNGR.D	
2786.4773	2785.47	2785.3559	41	0	424	-	447	K.QTQIFTTYSNQPGLVLIQVYEGE.R.A	
1248.6733	1247.6661	1247.6292	30	1	448	-	458	R.AMTKDNNLLGR.F	Oxidation
1183.6179	1182.6106	1182.6397	-25	0	459	-	469	R.FELSGIPPAPR.G	
1017.5971	1016.5899	1016.5614	28	1	501	-	509	K.IITITNDKGR.L	
1137.551	1136.5437	1136.5462	-2	1	525	-	533	K.YKAEDEVQRE	
1303.5859	1302.5786	1302.5914	-10	0	540	-	550	K.NALESYAFNMK.S	Oxidation

Master No. 1227

Match to: HSPB1_HUMAN

Score: 176

Expect: 3.9e-014

Heterogeneous nuclear ribonucleoprotein K (hnRNP K)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
871.3599	870.3527	870.4712	-136	0	288	-	296	R.GPPPPPPGR.G	
997.3432	996.336	996.4335	-97.9	0	317	-	325	R.GGDL MAYDR.R	
998.3108	997.3035	997.3811	-77.8	0	279	-	286	R.DYDDMSPR.R	
1013.3388	1012.3315	1012.4284	-95.7	0	317	-	325	R.GGDL MAYDR.R	Oxidation
1014.2876	1013.2803	1013.376	-94.4	0	279	-	286	R.DYDDMSPR.R	Oxidation
1053.5631	1052.5558	1052.6342	-74.5	0	192	-	201	R.VVLIGGKPDR.V	
1098.3734	1097.3661	1097.4448	-71.7	0	140	-	148	K.GSDFDCELR.L	
1106.4428	1105.4355	1105.5074	-65	0	38	-	46	R.NTDEMVELR.I	
1122.4469	1121.4396	1121.5023	-55.9	0	38	-	46	R.NTDEMVELR.I	Oxidation
1169.4541	1168.4468	1168.5295	-70.8	1	317	-	326	R.GGDL MAYDRR.G	Oxidation
1194.6312	1193.6239	1193.6921	-57.1	0	306	-	316	R.NLPLPPPPPR.G	
1259.4983	1258.491	1258.5677	-60.9	0	423	-	433	K.IDEPLGSEDRI	
1340.7281	1339.7208	1339.7962	-56.3	0	208	-	219	K.IILD LISEPIK.G	
1549.5733	1548.5661	1548.645	-51	0	180	-	191	K.LFQECPPHSTDV	
1579.6376	1578.6303	1578.6984	-43.1	0	22	-	34	K.RPAEDMEEEQAFK.R	
1595.6218	1594.6145	1594.6933	-49.4	0	22	-	34	K.RPAEDMEEEQAFK.R	Oxidation
1714.883	1713.8758	1713.9764	-58.7	0	87	-	102	R.ILSISADIETIGELK.K	
1735.6882	1734.6809	1734.7995	-68.4	1	22	-	35	K.RPAEDMEEEQAFK.RS	
1751.7102	1750.7029	1750.7944	-52.3	1	22	-	35	K.RPAEDMEEEQAFK.RS	Oxidation
1780.6833	1779.676	1779.7911	-64.7	0	70	-	86	R.TDYNASVSPDSSGPER.I	
1842.9749	1841.9676	1842.0713	-56.3	1	87	-	103	R.ILSISADIETIGELKK.I	
1916.9119	1915.9047	1916.0255	-63.1	0	378	-	396	R.GSYGDLGGPIITQTIPK.D	

Master No. 1295

Match to: CH60.HUMAN

Score: 270

Expect: 1.6e-023

60 kDa heat shock protein, mitochondrial precursor (Hsp60)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
833.4097	832.4025	832.3828	23.7	0	302	-	309	K.APGFGDNR.K	
844.5442	843.5369	843.5066	36	0	345	-	352	K.VGEVIVTK.D	
855.4952	854.4879	854.461	31.5	0	134	-	141	K.GANPVEIR.R	
912.5891	911.5818	911.5804	1.54	0	293	-	301	K.VGLQVVAVK.A	
941.6272	940.62	940.6069	13.9	1	463	-	470	K.IGIEIIKR.T	
960.4997	959.4924	959.5036	-11.7	0	421	-	429	R.VTDALNATRA	
961.4849	960.4776	960.4777	-0.14	1	302	-	310	K.APGFGDNRK.N	
1153.7044	1152.6971	1152.7594	-54	1	291	-	301	R.LKVGLQVVAVK.A	
1190.5187	1189.5114	1189.6012	-75.5	0	181	-	191	K.EIGNIISDAMK.K	
1215.5952	1214.5879	1214.6507	-51.7	0	482	-	493	K.NAGVEGSLIVEK.I	
1233.5036	1232.4964	1232.5885	-74.7	0	406	-	417	K.VGGTSDVEVNEK.K	
1344.632	1343.6247	1343.7085	-62.4	0	61	-	72	R.TVIEQSWGSPK.V	
1389.6243	1388.617	1388.6976	-58	0	222	-	233	R.GYISPYFINTSK.G	
1428.7216	1427.7144	1427.8058	-64	0	143	-	156	R.GVMLAVDAVIAELK.K	
1444.7256	1443.7183	1443.8007	-57	0	143	-	156	R.GVMLAVDAVIAELK.K	Oxidation
1504.6738	1503.6666	1503.749	-54.8	0	206	-	218	K.TLNDELEIIIEGMK.F	
1520.6688	1519.6615	1519.7439	-54.2	0	206	-	218	K.TLNDELEIIIEGMK.F	Oxidation
1556.8233	1555.816	1555.9007	-54.4	1	143	-	157	R.GVMLAVDAVIAELKK.Q	
1572.8268	1571.8195	1571.8956	-48.5	1	143	-	157	R.GVMLAVDAVIAELKK.Q	Oxidation
1584.8281	1583.8209	1583.9069	-54.3	1	142	-	156	R.RGVMLAVDAVIAELK.K	
1601.6695	1600.6622	1600.7443	-51.3	0	237	-	249	K.CEFQDAYVLLSEK.K	
1684.8343	1683.827	1683.8978	-42.1	0	430	-	446	R.AAVEEGIVLGGGCALLR.C	
1771.7882	1770.7809	1770.8458	-36.6	0	447	-	462	R.CIPALDSLTPANEDQK.I	
1919.0052	1917.998	1918.0636	-34.2	0	251	-	268	K.ISSIQSIVPALEIANHR.K	
1938.8933	1937.886	1937.9404	-28.1	1	206	-	221	K.TLNDELEIIIEGMKFDR.G	Oxidation
2037.9479	2036.9407	2037.0153	-36.7	0	371	-	387	R.IQEIIIEQLDVTTSSEYEK.E	
2113.0593	2112.052	2112.1323	-38	0	38	-	58	R.ALMLQGVDLLADAVAVTMGPK.G	
2129.0503	2128.043	2128.1272	-39.6	0	38	-	58	R.ALMLQGVDLLADAVAVTMGPK.G	Oxidation
2145.055	2144.0478	2144.1221	-34.7	0	38	-	58	R.ALMLQGVDLLADAVAVTMGPK.G	2 Oxidation
2365.2383	2364.231	2364.3264	-40.3	0	269	-	290	R.KPLVIIAEDVDGEALSTLVNRL	
2482.3201	2481.3128	2481.3942	-32.8	0	527	-	551	R.TALLDAAGVASLLTTAEVVVTEIPKE.E	
2508.015	2507.0077	2507.1018	-37.5	0	494	-	516	K.IMQSSSEVGYDAMAGDFVNMVEK.G	
2540.0632	2539.056	2539.0917	-14.1	0	494	-	516	K.IMQSSSEVGYDAMAGDFVNMVEK.G	2 Oxidation
2560.1303	2559.123	2559.2413	-46.2	0	97	-	121	K.LVQDVANNTNEEAGDGTATVLR.S	
2868.4753	2867.468	2867.5743	-37.1	1	527	-	554	R.TALLDAAGVASLLTTAEVVVTEIPKEEK.D	
3097.4057	3096.3985	3096.5074	-35.2	0	315	-	344	K.DMAIATGGAVFGEEGLTLNLEDVQPHDLGK.V	
3113.4077	3112.4005	3112.5023	-32.7	0	315	-	344	K.DMAIATGGAVFGEEGLTLNLEDVQPHDLGK.V	Oxidation
3555.5625	3554.5552	3554.8138	-72.7	1	158	-	191	K.QSKPVTTPPEIIAQVATISANGDKEIGNIISDAMK.K	

Master No. 1429

Match to: CAP1_HUMAN

Score: 104

Expect: 6.3e-007

Adenylyl cyclase-associated protein 1 (CAP 1)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
884.4756	883.4683	883.4803	-13.6	0	349	-	355	K.QVAYIYK.C	
1076.5526	1075.5454	1075.5696	-22.5	0	356	-	364	K.CVNTTLQIK.G	
1138.5615	1137.5542	1137.5601	-5.14	0	72	-	81	K.HAEMVHTGLK.L	Oxidation
1151.5121	1150.5048	1150.5077	-2.55	0	413	-	422	K.TDGCHAYLSK.N	
1212.6811	1211.6738	1211.6761	-1.93	1	318	-	328	K.EPAVLELEGKK.W	
1235.6137	1234.6064	1234.5863	16.3	0	423	-	433	K.NSLDCEIVSAK.S	
1276.6061	1275.5988	1275.6248	-20.4	0	199	-	209	K.EFHITGLAWSK.T	
1382.659	1381.6517	1381.6408	7.85	0	18	-	29	R.LEAVSHTSDMHR.G	
1398.641	1397.6337	1397.6358	-1.47	0	18	-	29	R.LEAVSHTSDMHR.G	Oxidation
1426.7985	1425.7912	1425.8079	-11.7	0	101	-	113	K.LSDLLAPISEQIK.E	
1478.5832	1477.5759	1477.5966	-14	0	156	-	167	K.EMNDAAMFYTN.R.V	Oxidation
1494.5894	1493.5821	1493.5915	-6.26	0	156	-	167	K.EMNDAAMFYTN.R.V	2 Oxidation
1757.872	1756.8647	1756.8778	-7.44	0	85	-	100	R.ALLVTASQCQPAENK.L	
2073.024	2072.0167	2072.0273	-5.13	0	331	-	348	R.VENQENVSNLVIETELK.Q	
2415.1723	2414.165	2414.2077	-17.7	1	329	-	348	K.WRVENQENVSNLVIETELK.Q	

Master No. 1484

Match to: FSCN1_HUMAN

Score: 157

Expect: 3.1e-012

Fascin (Singed-like protein)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
944.3333	943.326	943.4036	-82.2	0	111	-	118	R.YFGGTEDRL	
1032.3971	1031.3898	1031.4607	-68.8	0	390	-	398	R.GEHGFIGCR.K	
1076.4447	1075.4374	1075.5047	-62.5	0	186	-	194	R.YSVQTADHR.F	
1113.4629	1112.4556	1112.5363	-72.5	0	101	-	109	R.WSLQSEHR.R	
1142.5186	1141.5114	1141.588	-67.1	0	91	-	100	R.FLIVAHDDGR.W	
1146.4781	1145.4708	1145.5757	-91.6	0	23	-	32	K.YLTAEAFGFK.V	
1190.5603	1189.553	1189.6343	-68.3	0	230	-	241	R.YLAPSGPSGTLK.A	
1200.5631	1199.5558	1199.6662	-92	1	469	-	479	R.YLKGDHAGVLK.A	
1240.7251	1239.7178	1239.7815	-51.4	0	380	-	389	K.LINRPIIVFR.G	
1460.5668	1459.5596	1459.619	-40.7	0	331	-	341	K.NASCYFDIEWR.D	
1611.6218	1610.6145	1610.7359	-75.4	1	69	-	82	R.YLAADKDGNTVCER.E	
1819.8567	1818.8494	1818.9628	-62.3	0	202	-	217	R.LVARPEPATGYTLFR.S	
2109.8166	2108.8093	2108.9691	-75.8	0	409	-	426	R.SSYDVFQLEFNDGAYNIK.D	
2224.9457	2223.9384	2224.1045	-74.7	1	359	-	379	K.KNGQLAASVETAGDSEFLMK.L	Oxidation
2675.1953	2674.188	2674.3384	-56.3	1	248	-	271	K.VGKDELFALEQSCAQVVLQAANER.N	

Master No. 1579

Match to:PDIA6_HUMAN

Score:66

Expect: 0.0039

Protein disulfide-isomerase A6 precursor (EC 5.3.4.1)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
1078.6354	1077.6281	1077.607	19.6	0	314	-	322	R.NSYLEVLLK.L	
1151.5481	1150.5408	1150.4891	45	0	246	-	256	K.GESPVDYDGGR.T	
1191.5684	1190.5611	1190.5428	15.4	0	109	-	118	K.NRPEDYQGGR.T	
1386.7392	1385.732	1385.7514	-14	0	119	-	132	R.TGEAIVDAALSALR.Q	
1483.6895	1482.6822	1482.7103	-18.9	0	374	-	386	K.GSFSEQGINEFLR.E	
1514.7286	1513.7213	1513.7412	-13.1	0	195	-	208	K.NLEPEWAAAASEVK.E	
1527.8161	1526.8088	1526.8416	-21.5	0	217	-	231	K.LAAVDATVNQVLASR.Y	
1615.7968	1614.7895	1614.8366	-29.2	0	393	-	409	R.GSTAPVGGGAFPTIVER.E	
1825.8505	1824.8432	1824.9271	-46	0	86	-	102	K.HHSLGGQYGVQGFPTIK.I	
2637.321	2636.3138	2636.3585	-17	0	265	-	288	R.ALDFSDNAPPELLEIINEDIAK.R	

Master No. 1758

Match to : ACTG_HUMAN, ACTB_HUMAN*

Score: 160

Expect: 1.6e-012

Actin, cytoplasmic 2 (Gamma-actin), Actin, cytoplasmic 1 (Beta-actin)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
976.4337	975.4264	975.441	-15	0	19	-	28	K.AGFAGDDAPRA	
1198.6466	1197.6393	1197.6982	-49	0	29	-	39	R.AVFPSIVGRPR.H	
1171.5191	1170.5118	1170.5638	-44	0	40	-	50	R.HQGVMVGMGQK.D	
1187.5038	1186.4965	1186.5587	-52	0	40	-	50	R.HQGVMVGMGQK.D	Oxidation
1203.5114	1202.5041	1202.5536	-41	0	40	-	50	R.HQGVMVGMGQK.D	2 Oxidation
1354.5819	1353.5746	1353.6161	-31	1	51	-	62	K.DSYVGDEAQS.KR.G	
800.5403	799.533	799.528	6	1	62	-	68	K.RGILTLK.Y	
1515.6945	1514.6872	1514.7419	-36	0	85	-	95	K.IWHHTFYNELR.V	
1954.0574	1953.0501	1953.0571	-4	0	96	-	113	R.VAPEEHPVLLTEAPLNPK.A	
3183.5835	3182.5762	3182.6071	-10	0	148	-	177	R.TTGIVMDSGDGVTHTVPIYEGYALPHAILR.L	
3199.596	3198.5887	3198.602	-4	0	148	-	177	R.TTGIVMDSGDGVTHTVPIYEGYALPHAILR.L	Oxidation
998.4597	997.4525	997.479	-27	0	184	-	191	R.DLTDYLMK.I	
1014.4595	1013.4522	1013.4739	-21	0	184	-	191	R.DLTDYLMK.I	Oxidation
1132.4784	1131.4711	1131.5197	-43	0	197	-	206	R.GYSFTTTAER.E	
2550.1776	2549.1703	2549.1665	1	0	216	-	238	K.LCYVALDFEQEMATAASSSSLEK.S	
2566.1895	2565.1822	2565.1614	8	0	216	-	238	K.LCYVALDFEQEMATAASSSSLEK.S	Oxidation
1790.8755	1789.8683	1789.8846	-9	0	239	-	254	K.SYELPDGQVITIGNER.F	
3231.443	3230.4357	3230.4545	-6	0	257	-	284	R.CPEALFQPSFLGMESCGIHETTFNSIMK.C	
3247.491	3246.4837	3246.4494	11	0	257	-	284	R.CPEALFQPSFLGMESCGIHETTFNSIMK.C	Oxidation
2343.1248	2342.1175	2342.1576	-17	1	291	-	312	R.KDLYANTVLSGGTTMYPGIADR.M	
2359.1376	2358.1303	2358.1526	-9	1	291	-	312	R.KDLYANTVLSGGTTMYPGIADR.M	Oxidation
2215.0391	2214.0319	2214.0627	-14	0	292	-	312	K.DLYANTVLSGGTTMYPGIADR.M	
2231.0574	2230.0502	2230.0576	-3	0	292	-	312	K.DLYANTVLSGGTTMYPGIADR.M	Oxidation
1161.5541	1160.5468	1160.6111	-55	0	316	-	326	K.EITALAPSTMK.I	
923.5749	922.5677	922.56	8	1	329	-	336	K.IIAPPERK.Y	
2730.3048	2729.2975	2729.4251	-47	1	336	-	359	R.KYSVWIGGSILASLSTFQQMWISK.Q	
2746.3269	2745.3196	2745.42	-37	1	336	-	359	R.KYSVWIGGSILASLSTFQQMWISK.Q	Oxidation

* All sequences were overlapped in gamma-actin and beta-actin.

Master No. 1767

Match to: ACTG_HUMAN (P02571)

Score: 135

Expect: 5e-010

Actin, cytoplasmic 2 (Gamma-actin)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
1794.7623	1793.755	1793.8175	-35	0	2	-	18	M.EEEIAALVIDNGSGMCK.A	Oxidation
976.4188	975.4116	975.441	-30	0	19	-	28	K.AGFAGDDAPRA	
1198.6289	1197.6216	1197.6982	-64	0	29	-	39	R.AVFPSIVGRPR.H	
1171.4905	1170.4832	1170.5638	-69	0	40	-	50	R.HQGVMMVGMGQK.D	
1187.4823	1186.475	1186.5587	-71	0	40	-	50	R.HQGVMMVGMGQK.D	Oxidation
1203.4839	1202.4766	1202.5536	-64	0	40	-	50	R.HQGVMMVGMGQK.D	2 Oxidation
800.5371	799.5298	799.528	2	1	62	-	68	K.RGILTLK.Y	
1515.6704	1514.6632	1514.7419	-52	0	85	-	95	K.IWHHTFYNELR.V	
1954.015	1953.0077	1953.0571	-25	0	96	-	113	R.VAPEEHPVLLTEAPLNPK.A	
3183.4799	3182.4726	3182.6071	-42	0	148	-	177	R.TTGIVMDSGDGVTHTVPIYEGYALPHAILR.L	Oxidation
3199.5258	3198.5185	3198.602	-26	0	148	-	177	R.TTGIVMDSGDGVTHTVPIYEGYALPHAILR.L	
1132.4642	1131.4569	1131.5197	-55	0	197	-	206	R.GYSFTTTAERE	
1790.8467	1789.8394	1789.8846	-25	0	239	-	254	K.SYELPDGQVITIGNER.F	
2343.0651	2342.0578	2342.1576	-43	1	291	-	312	R.KDLYANTVLSGGTTMYPGIADR.M	Oxidation
2359.1172	2358.1099	2358.1526	-18	1	291	-	312	R.KDLYANTVLSGGTTMYPGIADR.M	
2215.0061	2213.9988	2214.0627	-29	0	292	-	312	K.DLYANTVLSGGTTMYPGIADR.M	
923.5349	922.5276	922.56	-35	1	329	-	336	K.IIAPPERK.Y	
2730.2553	2729.248	2729.4251	-65	1	336	-	359	R.KYSVWIGGSILASLSTFQQMWISK.Q	Oxidation
2746.2786	2745.2713	2745.42	-54	1	336	-	359	R.KYSVWIGGSILASLSTFQQMWISK.Q	

Master No. 2014

Match to: AK1C2_HUMAN

Score: 136

Expect: 3.9e-010

Aldo-keto reductase family 1 member C2

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
854.5383	853.531	853.5385	-8.78	0	251	-	258	R.TPALIALR.Y	
1130.5709	1129.5636	1129.5629	0.7	0	162	-	171	K.SIGVSNFNHRL	
1158.5827	1157.5754	1157.5717	3.24	1	76	-	84	K.REDIFYTSK.L	
1384.8087	1383.8014	1383.8159	-10.5	0	172	-	183	R.LLEMILNKPGLK.Y	Oxidation
1500.8114	1499.8042	1499.8096	-3.63	0	210	-	223	K.DIVLVAYSALGSHRE	
1715.9222	1714.9149	1714.9366	-12.7	1	208	-	223	K.SKDIVLVAYSALGSHR.E	
1972.874	1971.8667	1971.9248	-29.5	0	279	-	294	R.QNVQVFEFQLTSEEMK.A	Oxidation
2239.0306	2238.0233	2238.0099	6	0	184	-	200	K.YKPVCNQVECHPYFNQR.K	
2376.1183	2375.111	2375.162	-21.5	0	10	-	31	K.LNDGHFMPVLGFGTYAPAEVPK.S	Oxidation
3003.5253	3002.518	3002.4998	6.05	0	40	-	66	K.LAIEAGFHHIDSAHVYNNEEQVGLAIR.S	
3125.7171	3124.7098	3124.6849	7.99	0	105	-	131	K.NLQLDYVDLYLIHFPVSVKPGEEVIPK.D	

Master No. 2372

Match to: HSPB1_HUMAN

Score: 92

Expect: 1.1e-005

Heat-shock protein beta-1 (HSP 27)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
831.4723	830.465	830.5014	-43.9	0	6	-	12	R.VPFSLLR.G	
960.4303	959.4231	959.425	-1.97	0	21	-	27	R.DWYPHSR.L	
961.4552	960.4479	960.4454	2.64	0	13	-	20	R.GPSWDPFR.D	
1075.5771	1074.5698	1074.5669	2.66	0	80	-	89	R.QLSSGVSEIR.H	
1104.5135	1103.5063	1103.4996	6.06	0	128	-	136	R.QDEHGYISR.C	
1163.6423	1162.635	1162.6135	18.5	0	28	-	37	R.LFDQAFGLPR.L	
1783.9321	1782.9248	1782.9152	5.39	0	97	-	112	R.VSLDVNHFAPDELTVK.T	
1906.0153	1905.008	1904.9843	12.4	0	172	-	188	K.LATQSNITIPVTFESRA	
3242.7054	3241.6981	3241.6428	17	1	141	-	171	R.KYTLPPGVDPQTQVSSSLSPGTLTVEAPMPK.I	Oxidation

Master No. 2382
 Match to: HSPB1_HUMAN
 Score: 128
 Expect: 2.5e-009
 Heat-shock protein beta-1 (HSP 27)

Observed	Mr(expt)	Mr(calc)	ppm	Miss	Start	-	End	Sequence	Methionine oxidation
831.6192	830.6119	830.5014	133	0	6	-	12	R.VPFSLLR.G	
961.5476	960.5403	960.4454	99	0	13	-	20	R.GPSWDPFR.D	
960.5343	959.527	959.425	106	0	21	-	27	R.DWYPHSR.L	
1163.6807	1162.6735	1162.6135	52	0	28	-	37	R.LFDQAFGLPR.L	
1075.6409	1074.6336	1074.5669	62	0	80	-	89	R.QLSSGVSEIR.H	
1783.8911	1782.8838	1782.9152	-18	0	97	-	112	R.VSLDVNHFAPDELTVK.T	
1104.5779	1103.5707	1103.4996	64	0	128	-	136	R.QDEHGYISR.C	
1905.9628	1904.9555	1904.9843	-15	0	172	-	188	K.LATQSNEITIPVTFESRA	

Result of Figure 2

Master No.	Control	Actinomycin D	Bafilomycin A	Brefeldin A	Concanamycin A	Cycloheximide	Cytocalsin D	Daunomycin	Geldanamycin	IGRF-193	Jaspilactone	LY294002	MK-137	Nocodazole	Okadaic acid	Radiolol	Staurosporine	Tunicamycin	VP-16	
420	1	0.921222663	1.350417112 **	1.083669313	1.111666028	0.876454926	0.952343876	0.85456517	1.12476183	0.95963574	0.88344749	0.86403163	0.84211959	0.78157145 *	0.975268812	0.914181114	0.833793294 **	0.702362239 **	0.876901886	0.934372605
453	1	0.908680256 *	1.100223431	0.98293065	1.144922562 *	0.885579168	0.98259561 *	0.921179626	1.036392727	0.957814127	0.908463849	0.83431205	0.966448046	0.932736344 **	0.911103853	0.824604427 *	0.871708392	1.009511212	0.944766505	1.024376505
505	1	1.065318089	0.834685898	1.166865822	0.934255057	1.248703216 **	0.882399952	0.987456648	0.889953802	1.086909384	1.095581378	1.136480398	0.881846094	0.845178066	0.824878987 *	1.000308820	0.721722726 **	1.208175561 *	1.0248833	0.964655534
576	1	0.935440224	1.149333274	0.948320561	1.075817382	0.986337106	1.191411651	0.98630825	1.302789265 **	0.989716812	1.155918972 *	1.045132739	0.901325358	0.960138327	0.960138327	1.019188139	0.931743396	1.091199203	1.092570537	1.092570537
594	1	0.777267512 **	1.082989802	0.837767576 *	0.948689854	0.760384084 *	0.807878924 *	0.780775912 **	0.805957265 *	0.989697107	0.884403374	0.79834984 *	0.804467472	0.811227341 *	0.965223737	0.912412299	0.84758429	0.981269788	0.90482124	0.90482124
608	1	0.864681588 *	0.906777718	0.965924505	0.908411481	0.921132782	0.795448897 **	0.928318695	1.257491409 *	1.02634160	0.925485783	0.831050810 *	1.01738022	0.845624391 **	0.994421223	1.200041208 **	0.542263509 **	0.927647203	0.98112329	1.008899194
609	1	0.998253387	1.061244076	1.254293924 **	0.923056834	0.965143192	1.127004427	0.90725682	1.227084529 **	1.028312198	1.02262106	1.002062296	1.277443081 **	0.94505091	0.87307298 **	1.102324282	0.88523795	0.852139629	0.937436009	0.949652571
623	1	0.937892591	0.889997996	0.928462037	0.82741907 **	1.101333814	0.833540045 *	0.893477122	0.809411866 **	1.008143548	0.989692274	0.945654348	0.876071145	0.999709688	0.897014371	0.98864887	1.251719075 **	1.061945508	0.920293219	0.904141071
642	1	1.038100581	1.155441571	1.120902006	1.228105554 *	1.2879889243	0.994686937	1.104441797	1.254746013 **	0.981110277	0.946823859	0.992685704	0.798445494 *	0.927440883	1.206326107 *	0.878542603	1.341769684 *	0.987434074	1.063589610	0.987434074
651	1	0.940675591	0.804159917 **	0.921624584	0.826883823 *	0.93786562	0.832400394 *	0.878217385	0.717793209 *	0.976351648	0.909911643 **	0.896547635	0.726649894 *	0.788067656 **	0.842440264 *	0.790182718 **	0.98450821	1.005925595	0.902903310	0.960778904
647	1	0.953256766	1.102364064	1.072953965	1.243193951 **	1.292075129 **	1.120337514	1.101898915	1.294785413	1.00248371	1.037384933	1.177911266	1.054379413	0.840213731	0.924749265	1.192079078 *	0.673709045 **	1.356345693 **	1.053920132	1.07422633
650	1	0.972667754	1.068697158	0.876558269	0.850119235	0.91764137	0.855081516	0.995956585	1.429412533 **	0.918980909	0.840924138	1.059391717	0.914791857	0.877131056 *	1.069953009	1.359778002 **	1.100137011	1.453912065 **	1.046216285	1.044976728
657	1	0.972501751	0.833026566 **	0.955907073	1.058873395	1.121239605	0.880035858	1.084176567	0.974357122	0.986622871	0.912837904	0.823705036 **	1.011434332	0.796382013 *	0.942155799	0.992557021	0.602189005 **	0.928460595	0.954138282	1.26518277 **
659	1	0.971297202	0.915444219	0.979407346	0.806541177	1.010175579	0.790219146 **	0.662075628 **	0.918055659	0.848322137 **	0.937403499	0.773807455 *	0.080178627 *	0.526318644 **	0.683644802 **	0.856850645	0.869199824 **	0.841205032 *	0.503836758 **	0.904051975 **
660	1	0.849871494 **	0.823373771	0.962835905	0.836656791	0.759311372 **	0.709840655 *	0.777462351 *	0.940210498	0.774362177 *	0.846301983	0.9724922	0.787200213 *	0.506320573 *	0.758628605 *	0.917711777	0.849874039	0.597555052	0.987035157	0.987035157
663	1	0.91464379	0.792256236 **	1.001367775	0.939461566	1.078005306	0.816594236 **	1.058109261	0.933405757 *	0.653622677 *	0.889338284	0.643696262 *	0.842549735 *	0.545245952 **	0.736413815 *	0.85770036 *	0.294837307 *	0.839951022 *	0.489712329 *	0.882058291
690	1	0.959024554	0.895094067	0.950206658	1.018127079	0.93196778	0.819704532 *	0.986832654	0.91377719	0.956334779	0.782595046 **	1.03784024	0.789592797	0.784952797	0.936135573	0.893987578	0.761700238 **	1.033556001	0.94413576	0.988405309
696	1	0.895801629	0.91416118	0.948172686	0.926972744	1.024649234	0.807059003 **	0.96959507	0.830846428 *	0.743381273 **	0.890679226	0.814145336 **	0.915492743	0.509425527	0.786833775 **	0.932224485	0.766646505 **	0.987091573	0.580425168 **	0.989174395
713	1	1.101705001	0.789123378 *	1.06925914	0.846529514	1.207437336 *	0.167964933	1.180154583	1.014713763 **	0.927395957	1.081813471	1.055800694	0.920109928	0.954015203	0.879441519	0.822334198	0.802131003	1.193667171	0.96263272	1.05448241
714	1	1.123830017	0.71314759 **	0.984124414	0.735414546 *	1.090999457	0.797690386 *	1.079846805	0.63335145	0.8987734	0.938116915	0.871232842	0.817353005 *	0.869266694	0.787498519	0.673663667 *	0.716249327 **	1.00237993	1.01277068	1.045594219
728	1	0.946306005	0.838347905 *	0.998568811	0.916274782	0.955348301	0.768787223 **	0.994453553	0.896624122 *	0.949745599	0.957869314	0.843368674	0.780524052 *	0.74803459	0.834085247 **	0.704034895 **	0.848158464 *	0.809137384	0.885059849 *	1.165574608 **
740	1	1.051741515	0.827164656 *	1.08416505	0.977838033	1.02644137	0.91216973	0.950626618	0.914845337	1.201399177 **	1.027856433	0.837484793	0.710898468	0.91243766	0.738267417 **	1.052424144	0.935974178	0.866215011	0.989282342	0.887868923
747	1	0.887130833	0.83779897	0.882322228	0.838906795 **	0.855216041 *	0.93896705	0.855216041 *	0.93896705	1.16873057	0.970204682	0.782598806	0.983373755	0.939715934	0.82022455	0.871349894	1.14445959	0.836843037 *	1.003174753	0.954118918
760	1	1.016897383	1.220273599 **	1.05745444	1.0514515	1.102356586	0.976298794	1.00192373	0.961495915	0.949037316	1.02028089	0.96694079	1.08888348	0.900033448	0.979912353	0.904223513	1.173667681 *	1.04382266	0.942936132	1.003935448
780	1	1.0252325914	1.069878797	1.07214466	0.946233499	0.982819092	1.05218633	1.242091977 **	1.107363754	1.047920689	0.959322356	1.183257148 **	1.05642224	1.15633841	0.956462224	1.173832611	0.806182647 **	0.950309409	0.971979705	1.09187696
788	1	0.92880195	1.155746965 **	0.876980535	1.035480194	0.89533137	0.883212059 *	1.007109779	1.19788258	0.958197088	0.915212057	0.968414698	1.017536227	0.729399005	0.925264271	1.013191826	0.981019121	0.843797087 **	0.955116094	1.071775874
791	1	0.988835321	1.045083586	0.989077657	1.115856565	0.948830869	0.968222605 *	0.9383133	0.92595825	1.03447092	0.958600365	1.229487159	0.937871556	0.985093729	0.953707299	1.059457784	0.954984958	0.986042825	0.986042825	0.986042825
795	1	0.96176708	0.887848564	0.84656241 *	0.8285274321 *	0.894013327	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805	0.952470805
798	1	0.854226683	0.896578459	0.941303499	0.807574299	0.899071501	0.697938321 *	0.880520494	1.07298544	0.967902199	0.824216052 *	0.583085767 *	0.871882002	0.700712123	0.947588826	1.129466462	0.192593303 **	0.952710333 *	0.921738085	1.035484235
852	1	0.933550781	1.032172848	0.789355982 **	0.855500426 **	0.855245462 *	0.728441503	1.203163988 **	0.950622317	0.773773982	0.852545611 *	1.074034502	0.937025002	0.983002977	0.965504064	1.026254106	0.562030274 *	0.744353038	0.921738085	1.02639071 **
861	1	0.810753 *	1.121528951 **	0.917405461	1.0814088	0.887891021	0.90481953	0.932770873	1.21262517	0.999558801	1.116145399 *	0.92607769	0.893026927	0.982460544	0.952438688	1.120033381 *	0.857398734 *	0.960751775	1.063434452	1.063434452
863	1	0.988835321	1.045083586	0.989077657	1.115856565	0.948830869	0.968222605 *	0.9383133	0.92595825	1.03447092	0.958600365	1.229487159	0.937871556	0.985093729	0.953707299	1.059457784	0.954984958	0.986042825	0.986042825	0.986042825
865	1	1.16385933 *	1.012974662 *	0.849195313 **	0.931230574	0.848357203 **	0.775027853 *	0.74747629	0.938838937	0.855209001 *	0.817146716 *	0.910787661	0.877034079	0.845338033	0.466335933 *	0.720855053 *	0.921738085	1.063434452	1.063434452	1.063434452
867	1	0.837744 **	1.03155028	0.886836638 *	0.9953609	0.849187097 **	0.996103539	0.941675914	1.154838028 **	0.974089	1.0420493	0.91679979	1.02265019	0.98827493	1.085433657	0.97838696	0.97622778	0.880999538 **	1.02482591	0.97265362
879	1	1.042383598	0.887291268 **	1.027282727	1.02051177	1.238393052 *	1.117815844 *	0.928703299	0.94225751	0.716614824 **	1.4402714 **	0.819009582	0.805572428 *	0.582652272	0.834985116 **	1.059557657	0.840652135 *	1.02499951	1.05255085	1.04652752
890	1	0.94074668	1.175734535 *	0.988655421	1.056228243	0.886084929	0.894818272	0.956214575	1.135861471 *	1.042283003	0.916174912	1.048380928	1.045983967	0.918960835	0.926868997	0.866702626 *	1.099249847	0.873970248 *	0.907974761	1.103075768
898	1	0.78101661 **	1.205836633 **	1.000459594	1.046098642	0.877489821	0.773882592 *	0.928607774	0.902384372	0.784430985	0.650207386 *	0.369643003	0.90927136	0.812616178 **	0.361408016 *	0.60265772	0.956565096 **	0.96864666	0.96864666	0.96864666
900	1	0.988278725	0.758743269 **	1.007561503	0.791629504 *	1.17382856 *	1.195054121 *	1.028474578	1.11014214	1.065512303	1.328846733 *	0.901313728	1.424380111 **	1.1424131027 *	0.730298894 *	1.031232568	0.73			

1253	1	0.924296772	0.881424643	**	0.866311124	0.768259175	**	0.844363388	**	0.798738956	**	0.841037262	**	0.9985892725	0.844108703	**	0.80871551	**	0.883013523	0.852761648	**	0.946309147	1.009319593	0.593014986	**	0.91774424	0.92734828	0.983691949		
1258	1	1.56860324	0.821587173	**	1.12726605	1.020018418	**	1.045303217	**	0.974145705	**	1.228784558	**	1.194774573	1.106934084	**	0.99441096	1.810341161	**	1.032566668	**	1.017550824	0.903654805	2.520300913	**	1.805931782	**	1.029770723	1.149489206	
1272	1	0.8665636	0.906273953	**	0.863611681	0.832038219	**	0.832038219	**	0.70891931	**	0.857120125	**	0.968398984	0.841188961	**	0.828813800	**	0.829426235	**	0.7318265	1.033270298	0.941731666	**	0.931458466	**	0.833716236	0.93172028		
1280	1	1.161980263	1.08846065	**	0.757513015	1.076034080	**	0.94763607	**	0.911018099	**	0.935306468	**	1.199630205	0.902409554	**	0.944158722	**	0.9861552	**	1.072043452	1.268853331	**	1.164365319	1.471887148	**	1.007820743	0.891102068		
1285	1	0.98845671	2.319198801	**	0.979415913	1.651957938	**	0.844917068	**	0.950211260	**	1.20665595	**	1.093122514	1.04683668	**	0.976378628	**	0.84354322	**	0.981658065	1.01346783	**	1.10599843	1.176925177	**	0.891244252	0.962737825		
1290	1	0.987857358	0.85517054	**	0.903347054	0.822189233	**	0.857226038	**	1.025923248	**	1.069102956	**	1.002548174	1.106619892	**	0.947702248	**	1.03205476	**	1.042173956	1.042208992	0.914449559	**	0.58036237	**	0.940898201	1.187208818		
1295	1	0.987127145	1.394086127	**	0.793717474	1.030470261	**	0.819192981	**	0.930470261	**	0.951904242	**	1.039480251	1.04830251	**	1.039480251	**	1.04830251	**	1.039480251	1.039480251	1.039480251	**	1.039480251	**	1.039480251	1.039480251		
1298	1	1.063904807	1.117213794	**	0.722660015	0.919310664	**	0.874004991	**	0.78463932	**	0.906067245	**	1.843348103	0.898759457	**	0.790824126	**	0.950918576	**	1.129018765	1.100935253	1.241148133	**	1.408900241	**	0.760006076	0.886609097		
1295	1	0.98996054	1.232920137	**	0.783267288	**	0.98738257	**	0.7921833	**	0.926287328	**	1.74995055	**	1.009597866	0.800804052	**	0.931732187	**	1.089515476	**	0.921242416	1.123423657	1.344759594	**	0.416864743	**	0.816265746	0.917621131	
1316	1	1.017130512	0.988087733	**	0.784377092	0.93314635	**	1.00364566	**	1.106383284	**	0.974291812	**	0.943841551	0.975842804	**	1.085215155	**	0.943294389	**	1.06422241	1.103675231	0.954572177	**	0.88606329	**	1.126963225	0.93811006		
1317	1	0.907373771	0.741867488	**	1.021442551	0.930043823	**	0.89611203	**	0.89990683	**	0.911439695	**	0.846450891	1.020070703	**	0.95918993	**	0.854319113	**	0.729179908	**	0.935130698	0.932610098	0.804743288	**	0.017602743	1.001297665		
1319	1	1.05673038	0.91025431	**	0.858122534	**	0.918175388	**	0.77801102	**	0.943617402	**	0.96147595	**	0.951756859	0.796928714	**	1.035414299	**	0.936822501	**	0.732610803	**	0.96597971	1.064482719	**	0.754183318	**	1.002062344	
1323	1	1.015367343	0.691006254	**	1.022483283	**	0.90503417	**	0.948848026	**	0.795076677	**	0.89044673	**	0.755353598	0.929867744	**	0.794032511	**	0.786452439	**	0.975546666	0.99036451	0.85672576	**	0.532722135	**	1.01891958	0.92776642	
1328	1	1.058247979	0.902547404	**	1.01882585	0.904202303	**	0.991721134	**	1.07683981	**	1.112609264	**	0.81265154	1.072542955	**	1.172097448	**	0.980021165	**	0.787891195	**	1.331919328	**	1.199427416	**	0.904899412	0.827204723		
1329	1	0.972845456	0.949339501	**	1.173642176	**	0.910718436	**	0.968712387	**	0.947706504	**	1.076105586	**	1.175189211	**	1.006702691	**	0.900108585	**	1.106584728	**	1.058241161	**	1.153799766	**	1.119647882	**	0.930777816	
1330	1	0.940250441	1.065314268	**	0.88751357	1.015666551	**	0.922561326	**	0.994613024	**	0.902955531	**	0.90870665	1.013719772	**	1.105730737	**	1.035969689	**	0.906625385	**	1.11162524	**	1.280645434	**	0.807001722	**	1.048968021	
1341	1	0.92986707	0.890864238	**	1.080274866	**	0.854195341	**	0.944201075	**	0.93811473	**	0.885612574	**	0.912165541	0.902524551	**	0.901781596	**	1.033250796	**	0.953914274	**	0.923809533	1.01438622	**	0.875722521	**	1.189641922	
1342	1	1.08150101	1.33396977	**	1.023881412	1.039833198	**	0.888156047	**	0.934218404	**	1.053286613	**	0.946272782	0.911755378	**	0.88779625	**	0.905482603	**	0.889448428	**	0.02806254	1.01588948	1.150195745	**	0.010982396	0.951485154		
1343	1	0.05640379	0.763790367	**	0.937777065	0.768996321	**	1.13840217	**	1.078013891	**	1.198533838	**	0.74770077	1.320668192	**	1.085473729	**	0.789351578	**	0.92063837	**	0.914762844	0.914298324	1.19080983	**	1.21259976	**	1.04988793	
1353	1	1.124034354	1.323803761	**	0.957703449	0.988198458	**	1.017517832	**	1.178201156	**	1.089306033	**	1.299626536	**	1.043573427	0.986673695	**	1.27737059	**	1.007749143	**	1.054433384	1.153167179	**	1.11930365	**	1.486509727	**	1.227297055
1361	1	0.927281172	1.269666682	**	0.897580467	0.986452736	**	1.01962679	**	1.10706298	**	0.99522303	**	1.237423328	**	0.916579071	1.047116096	**	1.074249412	**	1.03626615	**	1.066876998	0.944635882	1.051897973	**	1.128259746	**	1.108995198	
1362	1	0.94605005	1.220170884	**	0.906702446	**	0.941508113	**	1.01802236	**	0.884268394	**	1.02744769	**	1.219081681	**	0.967178096	**	0.850961068	**	0.79292375	**	0.937713155	0.911660681	1.120188754	**	1.02199583	**	1.274557817	
1365	1	1.10923997	1.153500441	**	0.93578457	0.959765719	**	0.896292206	**	0.894547366	**	0.9641054	**	1.03789054	**	0.880527344	**	1.07449611	**	0.986988362	**	1.07449611	**	0.839442684	**	0.9112863	**	1.02175503	**	0.96728955
1375	1	1.01489206	0.84648144	**	0.765037659	**	0.874727388	**	0.82419161	**	0.85233705	**	0.94035614	**	0.93055414	**	0.932011957	**	0.956448809	**	0.705987349	**	0.822924004	**	0.835452284	**	0.080103125	**	0.805619144	
1376	1	1.161492871	0.978686512	**	0.785975533	0.858621657	**	0.971900427	**	0.923474951	**	1.052250683	**	0.94710681	1.043439157	**	0.958214395	**	0.937007715	**	0.91633764	**	0.998179915	1.068467929	0.85055785	**	0.763794544	**	0.959884849	
1383	1	0.90463754	1.046453124	**	0.908463754	1.001543453	**	0.908463754	**	0.912381572	**	0.908463754	**	0.912381572	**	0.908463754	**	0.912381572	**	0.908463754	**	0.912381572	**	0.908463754	**	0.912381572	**	0.908463754	**	0.912381572
1382	1	0.951212822	0.73524435	**	1.001185346	0.85858366	**	0.965524696	**	0.904549316	**	1.006331901	**	0.815451117	**	0.874686159	**	0.905635138	**	0.923691864	**	0.9008284692	**	0.881348414	**	0.888535791	**	1.072855908	**	1.25836331
1386	1	1.05588047	1.297134209	**	0.89461287	1.029155887	**	0.91964255	**	1.129712531	**	1.105564235	**	1.197725321	**	1.105564235	**	1.197725321	**	1.105564235	**	1.197725321	**	1.105564235	**	1.197725321	**	1.105564235	**	1.197725321
1387	1	1.09282507	1.199341917	**	1.135457939	1.129074845	**	1.022899783	**	0.997931571	**	1.06825989	**	1.198129345	**	1.074592199	**	1.02637402	**	1.338336235	**	1.24164821	**	1.103061098	1.033536766	1.033007713	**	1.066324078	**	1.091399303
1398	1	1.06078599	0.98717475	**	1.03355058	**	0.933097803	**	0.92362644	**	0.974318293	**	0.94525268	**	0.745184948	**	0.88146948	**	1.017495125	**	0.87188907	**	1.034989903	0.89302663	0.774662283	**	1.06224946	**	0.98609874	
1401	1	0.98401905	0.966714143	**	1.129437152	**	1.10029891	**	0.912982917	**	1.10029891	**	0.912982917	**	1.10029891	**	0.912982917	**	1.10029891	**	0.912982917	**	1.10029891	**	0.912982917	**	1.10029891	**	0.912982917	
1405	1	1.04728436	1.010881966	**	1.027400406	1.069109425	**	1.046663636	**	0.760274263	**	0.93750127	**	1.26738779	**	0.88749188	**	0.93421977	**	0.908782444	**	0.884741519	**	0.889297122	1.160463961	**	1.059584023	**	1.04058882	
1408	1	1.011789341	0.862101617	**	1.00061486	0.998962968	**	1.036393082	**	1.091727308	**	1.087614622	**	1.100567783	**	1.176598241	**	1.092689863	**	0.915072748	**	0.91756699	**	0.973443649	1.019427272	**	1.019342272	**	1.016350718	
1413	1	0.989672607	**	0.93180321	0.784744742	**	0.981409588	**	0.837248239	**	0.98706954	**	0.98639258	**	1.00963558	**	1.03439255	**	1.00862342	**	1.012738306	**	0.995135077	1.054383637	0.977596322	**	1.05288037	**	1.05288037	
1414	1	0.92625664	0.656806313	**	0.745737881	**	0.637255563	**	0.65958758	**	0.655017204	**	0.630293094	**	0.712539267	**	0.90778652	**	0.738759740	**	0.522227409	**	0.83158427	**	0.947482655	**	0.759805335	**	0.80216661	
1421	1	0.995397941	1.153058617	**	0.889780217	0.988674026	**	0.867720809	**	0.925128981	**	1.008057504	**	1.193860512	**	0.900323689	**	0.989108045	**	0.901870004	**	0.995956945	**	0.933073771	0.918788398	1.08017654	**	1.428875915	**	1.070674482
1423	1	1.070935042	0.827800374	**	1.188367879	**	0.884661919	**	0.708758864	**	0.888908155	**	0.941246866	**	1.095866406	**	0.934557036	**	0.887905708	**	1.01123701	**	0.987628911	1.020246312	0.941					

1746	1	0.906456282	**	0.886594967	**	0.913764566		0.906293331		0.860957892		0.870346335		0.901263399		0.765781939	**	0.920723284		1.194627553		0.915959914		0.880494026		1.177015281	**	0.980635888		0.800785167	**	0.944875593		1.033470255		1.072837181			
1754	1	0.75831483		0.84613016		0.182242498	**	0.929041612		0.91822968		0.734203191	**	1.027569441		0.96655229		0.930067355		0.898927524		0.88223961		0.842044626		0.861167152	**	1.068489336		0.969707296		1.023607399		0.849183255		0.823250204	**	0.72078927	
1758	1	0.942190754		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675		0.94443675	
1760	1	0.80301707	**	0.750397863	**	1.003818804		0.83476829	**	0.945667757		0.843348048	**	0.73265721	**	0.801321241	**	0.85076612		0.826419492	**	0.836939202	**	0.981579135		1.065220352		0.789739204	**	0.98769904	**	1.022785169		1.03202811		0.832054001		0.825005484	
1761	1	0.768796457	**	1.229729304	**	1.01873886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886		0.91073886	
1764	1	0.081922342		0.845406146		1.02658263		0.996471268		1.305794773	**	1.435787477	**	0.970144648		1.100022671		1.10388519		1.623637769	**	1.283498839	**	1.100229181		1.168912427		0.727467502	**	1.028949357		0.978208155		1.313912721	**	1.002889358		0.995387513	
1767	1	0.039633064		0.08029015		0.89302756		0.90271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212		1.00271212	
1768	1	1.164811617	**	0.795681991	**	0.99611366		0.90515778		0.98951588		1.204529979	**	0.971159953		1.067510026		1.150379195		1.387118632	**	1.040626137		1.03962692		1.206472951	**	0.813037337	**	1.048467696		0.931882225		1.022247682		0.972710648		0.960504152	
1777	1	0.795314486	**	0.742201999	**	1.03421977		0.841926975		1.096582193		0.947688847		0.729860674	**	0.944429634		0.98834362		1.000446061		0.918318461		0.893324951		0.905050514		0.931237457		0.90620538		0.906779649		0.867621943		0.992513933		0.966517454	
1783	1	1.159423695	**	0.81440707	**	1.017612794		0.957923688		0.845476984	**	0.61377016	**	0.91142806		0.989579702		0.800651834	**	0.91736402		0.792495281	**	0.98186386		0.868485811		0.606207118	**	1.260004376	**	0.917280806		0.97439633		0.97439633		0.97439633	
1788	1	0.044658105		0.795726533	**	0.897357091		1.041381918		0.919934249		0.795128954	**	0.954242084		0.789534692	**	0.950676764		0.975943062	**	0.819006528	**	0.86141819		0.935238895		0.94933829		0.932984299		0.600265927	**	1.064891957		0.964270762		0.889723239	
1795	1	0.089655205	**	0.83160396		1.02432084		0.96432694		0.92854591		0.749660247	**	0.9347281		0.882587667		0.943440066		0.7496648015	**	0.904460505	**	0.91228496		1.01291401		0.906801895	**	0.6653307	**	0.889889508	**	0.933241109		0.907487146		0.907487146	
1799	1	0.044386191		0.97855369		0.85696515		1.07093526	**	0.98813215		1.322799356	**	0.98813215		1.322799356	**	0.98813215		1.322799356	**	0.98813215		1.322799356	**	0.98813215		1.322799356	**	0.98813215		1.322799356	**	0.98813215		1.322799356	**	0.98813215	
1815	1	1.17759308	**	1.13975598	**	0.929035686		1.026013286		0.96606617		0.987865359		1.196695392	**	1.241887668	**	1.085265959		0.934744833		1.116618217	**	1.006224611		1.090406489		0.992131961		0.985471406		1.511710683	**	0.7112057	**	0.944436938		1.067337678	
1837	1	0.871193301	**	0.814587214	**	1.09547414	**	0.840843762	**	0.874510785	**	1.03864366		0.75970352	**	0.927841226		0.948764108		1.008686946		0.969916748		0.588190849	**	0.8111857	**	0.721989063	**	0.637667722	**	0.972132236		0.763698986	**	0.984945228		0.984945228	
1846	1	1.219113001	**	0.986395392	**	0.86962287		1.035746113		1.007799736		0.902480231	**	1.01188475		0.948995506	**	0.937651326		0.971252533		0.994098577	**	0.947681084		0.99696817		1.07303436		0.97976695		0.76379752	**	1.055925957	**	0.921717331		0.982175928	
1947	1	1.192026821	**	0.83364391	**	0.839061914	**	0.856346714		0.845818899	**	0.684827042	**	1.010217353		0.752373234	**	0.945305429		0.879327584	**	0.84105847	**	0.800403064	**	0.887138756	**	1.002697204		0.85792347	**	0.59864435	**	0.970705379		0.981221165		0.921217346	
1850	1	0.105584925	**	1.178493625	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**	0.9960748579	**
1853	1	1.198829117	**	0.928795517	**	0.885975944		1.004392144		0.931391186		0.634708983	**	1.043599144		0.70297251	**	0.930713538		0.814340454	**	1.001346655		0.845869175		0.852200027		0.934578541		0.885428075		0.640738841	**	1.057953321		0.993885066		1.00436763	
1854	1	0.104758107		1.099450729	**	1.2231834		1.099415663		1.091707669		2.080049992	**	1.204835907		0.807109982	**	1.15847172		1.668943542	**	1.20421367		1.221437672		1.000813532	**	0.951566889		1.168156925		1.904231116	**	1.168595545		1.117760519		1.089095106	
1855	1	1.144021778	**	1.195510288	**	0.741639363	**	1.027076944		0.981052748	**	1.01008627	**	1.145354261	**	1.079741468		0.988925151		0.935632698		1.063944907		1.024287754		0.943399308		1.004983473		0.986295514		0.97862128		0.744263083	**	0.885736279	**	1.067728559	
1859	1	0.993368262		1.01798269		1.319160626	**	1.080476874		0.898981507		1.521446164	**	1.055231254		0.966650026		1.043858324		1.440066583		1.082718724		1.202376175	**	1.426731543	**	1.06424523		1.0329795		1.377821827	**	0.997329107		1.118125796		0.860000293	**
1860	1	0.103850408		0.94238947		0.94482756		0.917523319		0.948725756		0.72291062	**	0.938956556		0.791533835	**	0.735726562	**	0.87196929	**	0.809937197	**	0.91839847	**	0.762306481	**	1.046734545	**	0.695381372	**	1.159824541	**	0.941601437		0.502099885		0.502099885	
1877	1	0.881605208	**	0.696346725	**	0.951367313		0.763255294	**	0.865695484	**	0.9257735079		0.97643134		0.813599262	**	1.034049195		1.040737184		0.76223035	**	0.873156235	**	0.399994301		0.981958543		0.90313366		0.529980708	**	0.932788307		1.009647274		0.975372665	
1878	1	1.167369098	**	1.15853031	**	1.04148475	**	1.090211286		1.101451009	**	1.008324069	**	1.071458335		1.027781757	**	1.048446589		1.007930263		0.913174767	**	1.047686074	**	1.004806171		1.00911568		1.01259708		1.039881405	**	1.101735744	**	0.9881858		1.045062913	
1879	1	1.15181873	**	0.796052915	**	1.21670687	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**	0.796052915	**
1889	1	0.89705496		0.903352501	**	0.9348011		0.975995565		0.779164378	**	0.870345494	**	0.869975802	**	0.94789795	**	1.08225876		0.899761925	**	0.919137398		0.997792914		1.068145859		0.92819431		0.964900989		1.041816146		0.935714526		0.9542036		1.089152897	
1898	1	1.009576294		0.806293185	**	1.035426967		1.024710903		1.058326915		0.987810735	**	1.038359955		1.017821388		0.91564696	**	1.0151853	**	0.903676082		0.915679652	**	0.903676082		0.915679652	**	0.903676082		0.915679652	**	0.903676082		0.915679652	**	0.903676082	
1902	1	1.13167899	**	1.084264833	**	0.968113499	**	1.120863426	**	1.075132351	**	0.829252297	**	0.957399543	**	1.103775236	**	0.846331243	**	0.858420305	**	0.938857508	**	0.989251925	**	0.750126031	**	0.96989581	**	0.884188963	**	0.868417285	**	1.061670807	**	0.767744642	**	1.10383819	**
1915	1	0.8845555	**	1.04504269	**	1.135282978	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**	1.17518173	**
1925	1	0.951216579	**	0.83364391	**	0.839061914	**	0.913111221	**	0.738173337	**	0.887158674	**	0.913111221	**	0.738173337	**	0.887158674	**	0.913111221	**	0.738173337	**	0.887158674	**	0.913111221	**	0.738173337	**	0.887158674	**</								

2378	1	1.156224718 **	1.119965652 *	1.186054183 **	0.930283228	0.993131288	0.779249372 **	1.207054613 **	2.062888857 **	1.158412646 **	0.997130724	1.331178953 **	1.076602229	0.876983873 *	0.72903818 **	1.62090956 **	2.314786619 **	1.020753229	0.97769085	0.993651103
2382	1	1.07963357	0.994191583	0.883226339	0.847210936 **	0.853042882 *	0.798800363 **	1.031730573	1.900307374 **	1.035019361	1.065535307	0.986113289	0.92772866	0.867603957	0.774942036 **	1.646572245 **	2.303103085 **	0.859572575 *	0.910419961	0.930879554
2383	1	1.329190118 **	1.110735384	1.014836602	0.920945346	0.94245229	0.995719175	1.266055715 **	1.023533597	1.130784053	1.01442455	1.071629289	1.050503388	0.984558888	1.084350775	1.034433959	0.939812027	0.969016788	1.056934945	**
2395	1	1.151638067 **	0.746676335 **	1.120094973 **	0.817335106 **	0.985809795	0.843688844 **	1.075429369 *	0.862103439 **	1.077956955 *	0.911710548 **	0.991521554	0.912819002 **	0.966639955	1.003970892	1.005415379	0.82216933 **	1.052698435	1.007264479	0.965171462
2396	1	1.16904634 *	0.896449382	1.061818047	0.832647133 *	0.815481564 **	0.878992065	1.069806939	0.834310338 *	1.084932683	0.927190744	0.88051067	0.842050953 *	0.80975111 **	0.998057471	0.863517299	0.491840664 **	0.951471408	0.807142359 **	1.046483719
2401	1	0.986611226	0.683426523 **	0.84082333 *	0.828890855 *	0.95372248	0.839938221 *	0.954625331	0.90576684	0.939391846	0.922069713	0.754711319 **	1.050087264	0.924297049	1.208835171 **	0.950653782	1.163608505 *	0.907577236	0.978945189	1.052651082
2406	1	1.163054923 *	0.923241236	1.039107262	0.945677387	0.985064391	1.048677256	1.016559144	0.733286808 **	1.12337112	1.020537313	0.955396676	0.835405635 *	0.948262382	1.044751249	0.908609027	0.691298633 **	1.124196958	0.978768144	1.029106421
2412	1	1.106467674	1.014082893	0.919075247	0.869421474	0.829637747 *	0.8850322	1.065046865	1.663209649 **	1.005869694	1.036814226	1.037694937	1.031754845	1.020534696	0.811274415 **	1.41583467 **	2.218703293 **	0.906607421	0.921099726	0.935189251
2420	1	1.282914891 **	0.735079329 **	1.134853968	0.77859556 *	1.242140818 **	0.903972837	1.200842468 *	0.799825426 *	1.063757623	1.049306277	1.086280678	0.863535265	0.996188603	0.973477052	0.967993492	1.248265014 **	1.202287889 *	0.919985853	1.002307872
2438	1	0.960817348	0.547788307 **	1.068342147	0.847055823	0.931809533	0.874262322	1.224807238	1.131603822	1.179332691	1.02488419	1.170941367	1.126088017	1.118513846	0.998367558	1.014587933	0.91368798	1.017449829	0.991639355	1.00863258
2450	1	0.967307941	0.69809128 **	0.828766659 **	0.878771827 *	0.879924515 *	0.931517583	0.957159898	0.803878291 **	0.891027385 *	1.011821916	0.834381524 **	0.999739997	0.96226942	1.05479112	0.917508245	0.626959851 **	0.978109657	0.911279892	0.971025563
2455	1	1.034571489	0.724073999 **	0.915180301	0.791390622 **	0.900957328	0.765053805 **	0.955290459	0.831143036 **	1.064441062	0.903170576	0.804630537 **	0.864508137 *	0.958577912	1.022894014	0.945613672	0.600777202 **	0.97790886	1.00663188	0.967057578
2462	1	1.129785584	0.749278873 **	1.028398957	0.801728017 **	1.001951053	0.909024355	1.053223516	0.811104307 **	1.121796852	0.960122011	0.960814484	1.031106382	1.058275173	1.045468731	0.890118742	0.7717347 **	1.131995459	1.054247317	1.057437745

The mean ratios of identified spots between control and inhibitor-treated cells are listed. Non-repeated measures ANOVA and Dunnett's test for post hoc analysis were performed. Asterisks indicate significant differences from respective controls (*P < 0.05, ** P < 0.01).