

## Supplementary Materials

### Nucleolar and nuclear stress-induced membrane-less organelles: A proteome analysis through the prism of liquid-liquid phase separation

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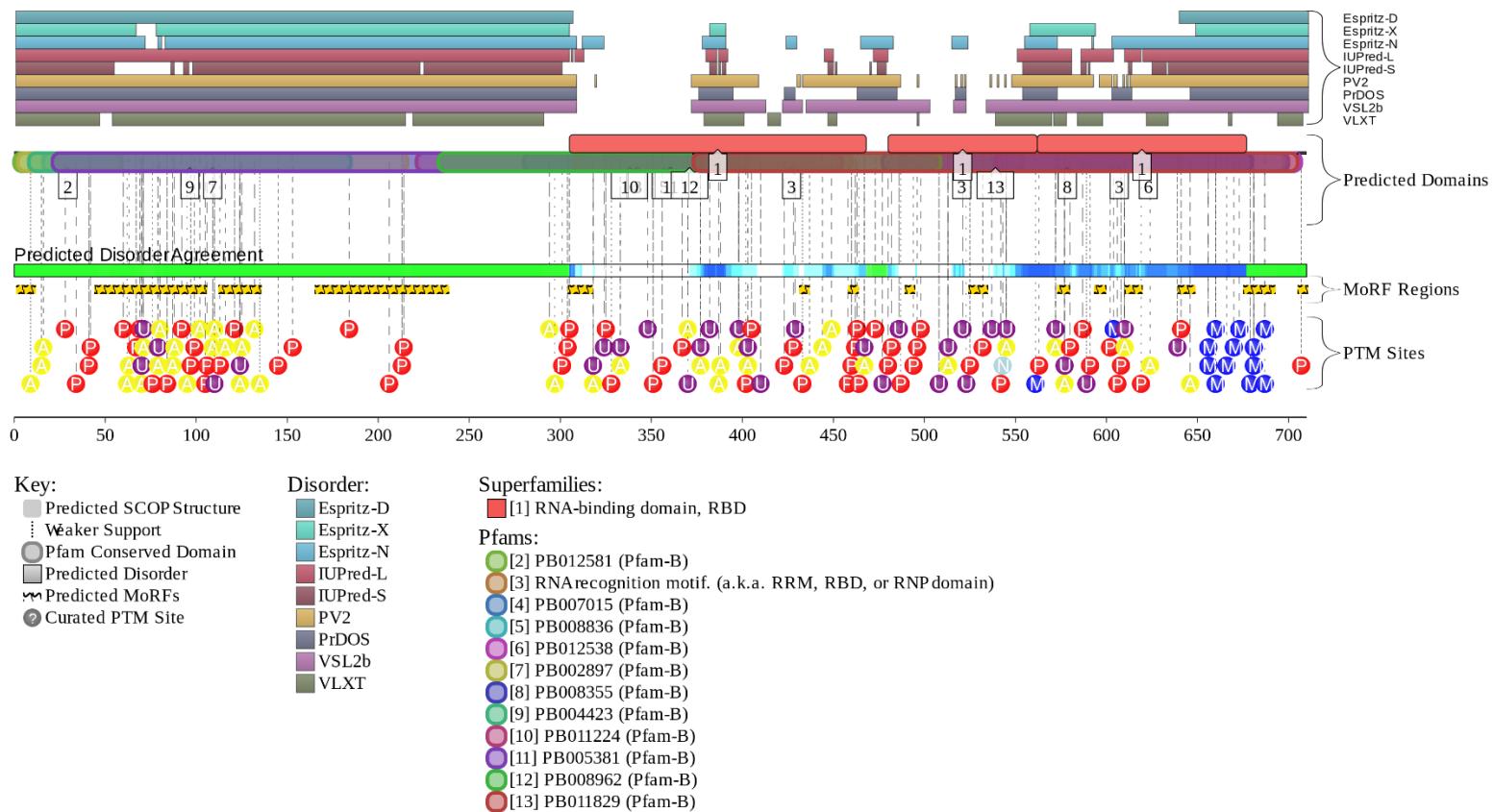
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Supplementary Figures S1-S10 represent amino acid sequences and functional disorder profiles of 10 human proteins shared by the acidosis-induced A-bodies, heat-induced A-bodies, and nSBs. In each figure, the upper plot represents amino acid sequence in FASTA format, whereas the bottom plot shows corresponding functional profile generated by the D<sup>2</sup>P<sup>2</sup> platform. In these profiles, the IDPR localization predicted by IUPred, PONDR<sup>®</sup> VLXT, PONDR<sup>®</sup> VSL2, PrDOS, PV2, and ESpritz are shown by 9 differently colored bars on the top of the plot, whereas the blue-green-white bar in the middle of the plots shows the agreement between the outputs of these disorder predictors, with disordered regions by consensus being shown by blue and green. The two lines with colored and numbered bars above the disorder consensus bar show the positions of functional SCOP domains [40, 41] predicted using the SUPERFAMILY predictor [42]. Positions of the predicted disorder-based binding sites (molecular recognition feature (MoRFs) identified by the ANCHOR algorithm are shown by yellow zigzagged bars [43]. Locations of the sites of different posttranslational modifications (PTMs) identified by the PhosphoSitePlus platform [44] are shown at the bottom of the plots by the differently colored circles.

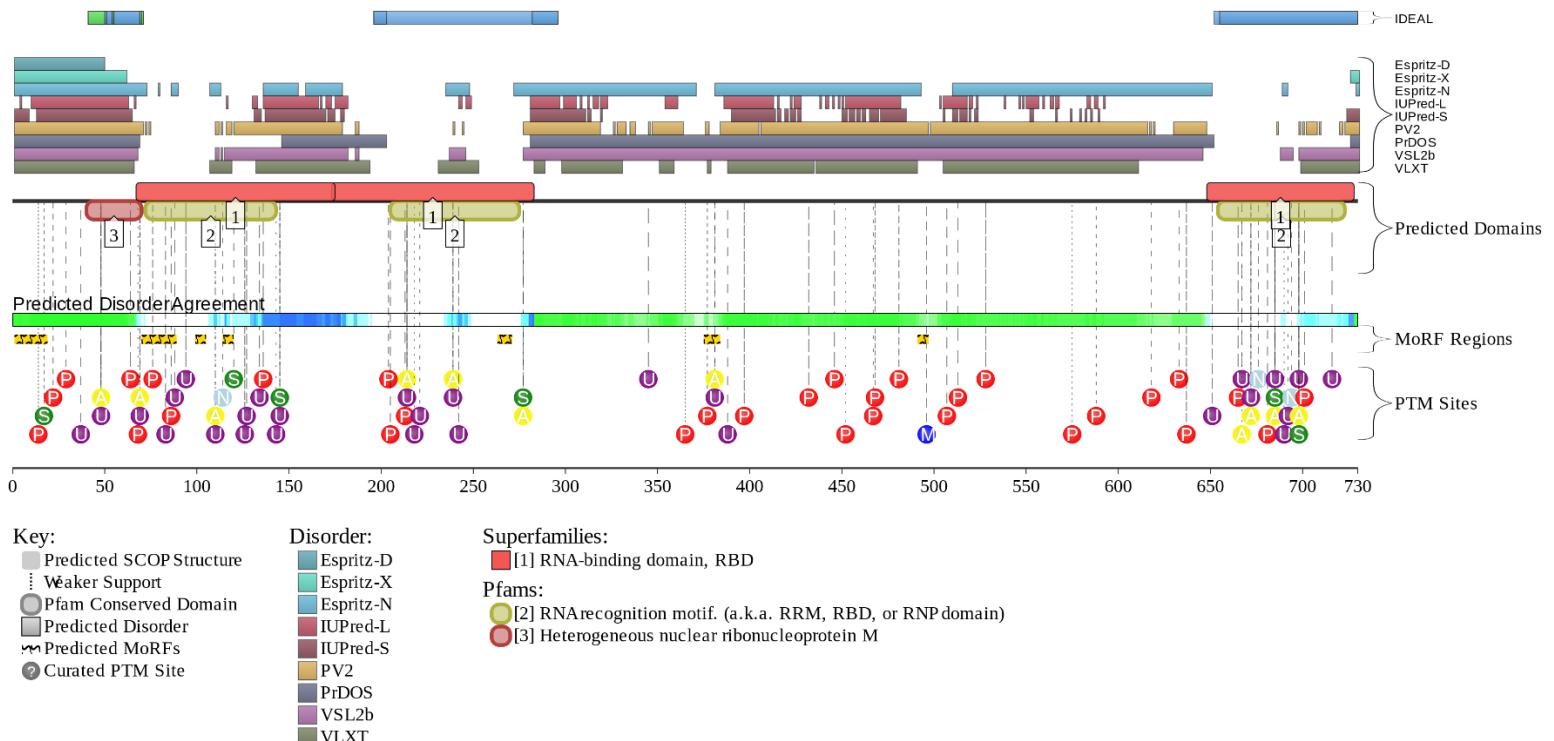
**Figure S1.** Amino acid sequence and functional disorder profile of Nucleolin (NCL; also known as Protein C23; UniProt ID: P19338).

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>sp|P19338|NUCL_HUMAN Nucleolin OS=Homo sapiens OX=9606 GN=NCL PE=1 SV=3
MVKLAQAGKNQGDPKKMAPPPKEVEEDSEDEEMSEDEEDSSGEEVVIHQKKGKAAATSAKVVSPTKVAATP
AKKAATPGKKAATPAKTVTPAKVTPGKKGATPGKALVATPGKGAIIPAKGAKNGKNAKKEDSDEEEDDDSE
EDEEDDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDE
VAEDEDEDEEEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDEDE
LFVGNLNFKNSAPELKTGISDVFAKNDLAVDVRIGMTRKFGYVDFESAEDLEKALELTGLKVFVGNIEKLEKPKGD
SKKERDARTLLAKNLPYKVTQDELKEVFEDAAEIIRLVSKDGKSKGIAYIEFKTEADAETKFEEKQGTEIDGRSISLY
YTGEKGQNQDYRGGKNTWSGESKTLVLSNLSSATEETLQEVFEKATFIKVQPQNCNGSKGYAFIEFASFEDAKEY
LNSCNKREIEGRAIRLELQGPRGSPNARSQPSKTLFVKGLSEDTEETLKESFDGSVRARIVTDRETGSSKGFGFVD
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RGGGGDHKPQGKKTKE
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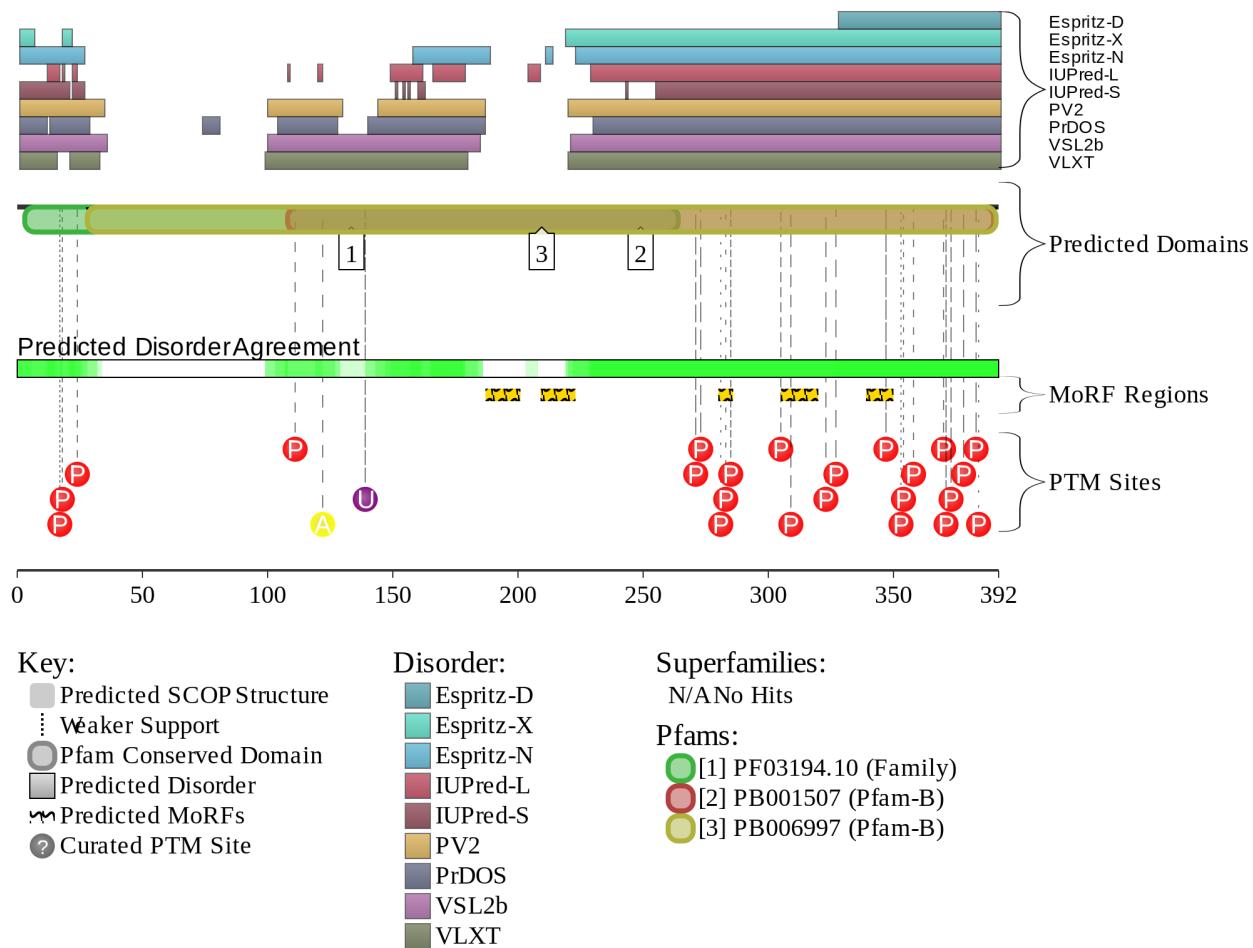
**Figure S2.** Amino acid sequence and functional disorder profile of heterogeneous nuclear ribonucleoprotein M (HNRNPM; UniProt ID: P52272).

```
>sp|P52272|HNRPM_HUMAN Heterogeneous nuclear ribonucleoprotein M OS=Homo sapiens OX=9606 GN=HNRNPM PE=1 SV=3
MAAGVEAAAEVAATEIKMEEESGAPGVPSGNAGAPGPKGEGERPAQNEKRKEKNIKRGGRNFEPYANPTKRYRAFITN
IPFDVKWQSLKDLVKEVGTVYVELLMDAEGKSRGCAVVEFKMEESMKAAEVLNKHSLSGRPLKVKEDPGEHAR
RAMQKVMATTGGGMGPGGGPGMITIPPSILNNNPNIPEIIHALQAGRLGSTVFVANLDYKVGWKKLKEVFSMAGVVV
RADILEDKDGSRGIGTVTFEQSIEAVQAISMFNGQLLFDRPMHVKMDERALPKGDFFPERPQQLPHGLGGIMGL
GPGGQPIDANHLNKIGMGNIGPAGMGMEGIFGINKMGGMEGFPGGGMENMGRFGSGNMGRINEILSNALKREI
IAKQGGGGGGGSVPGIERMGPVIDLGGAGMERMGAGLGHGMDRVGSEIERMGLVMDRMSVERMSGIERMGPLGL
DHMASSIERMGTMERIGSGVERMGAGMGLERMAAPIDRVQQTIERMGSVERMGPAPIERMGLSMERMVPGMGA
GLERMGPVMDRMTGLERMGANNLERMGLERMGANSLERMGLERMGPAAMGPALGAGIERMGLAMGGGG
ASFDRAIEMERGNFGGSFAGSFAGSFAGGHAAPGVARKACQIFVRNLPDFTWKMLDKFNECGHVLYADIKMENGKSKG
CGVVKFESPEVAERACRMNMGKLSGREIDVRIDRNA
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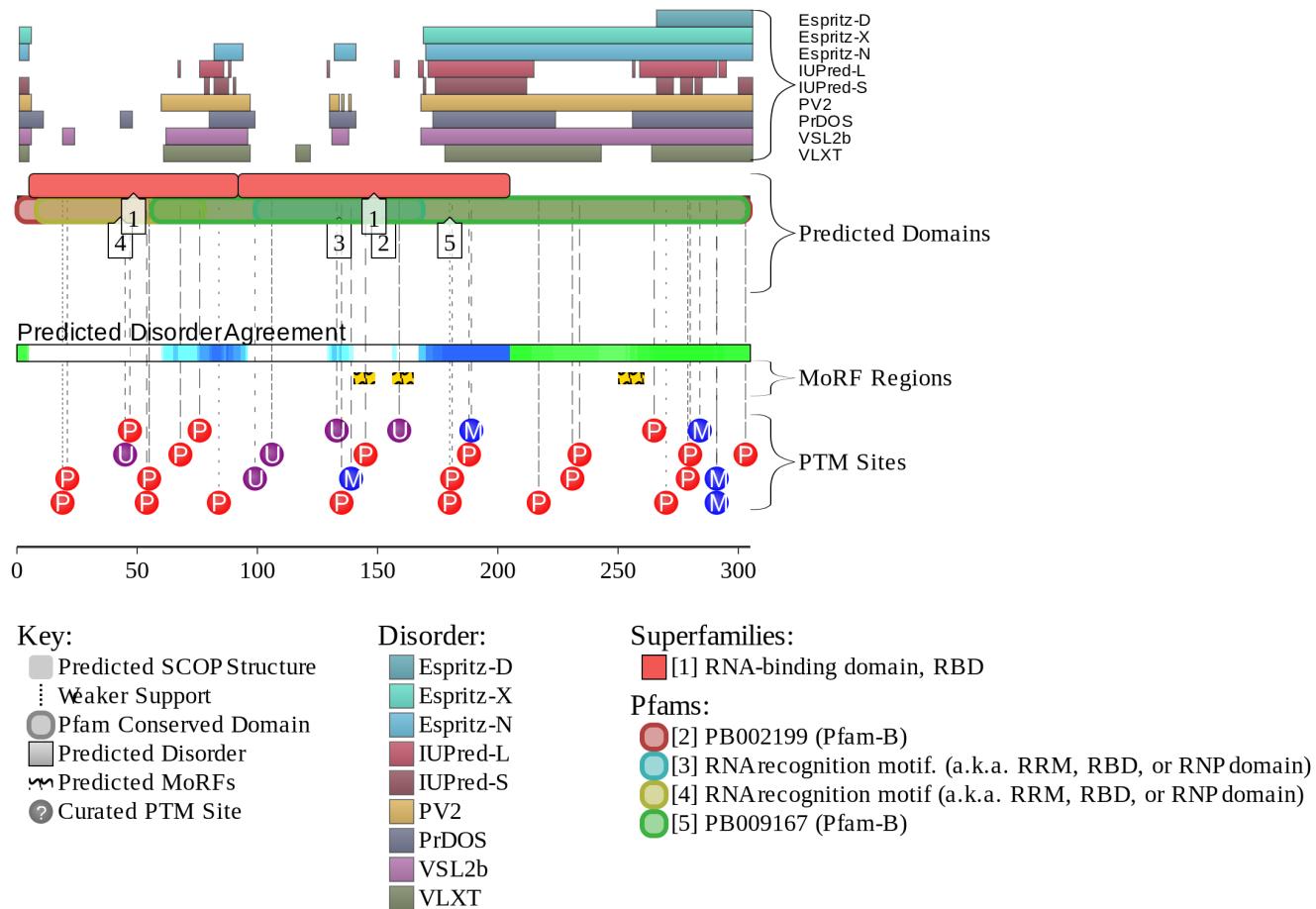
**Figure S3.** Amino acid sequence and functional disorder profile of putative RNA-binding protein Luc7-like 2 (LUC7L2, also known as CGI-74; UniProt ID: Q9Y383).

```
>sp|Q9Y383|LC7L2_HUMAN Putative RNA-binding protein Luc7-like 2 OS=Homo sapiens OX=9606 GN=LUC7L2 PE=1 SV=2
MSAQAAQMRAMLDQLMGTSRDQGDTTRQRIKFSDDRVCKSHLLNCCPHDVLSGTRMDLGECLKVHDLDALRADYEIASKE
QDFFFFELDAMDHLQSFIAACDRRTEVAKKRLAETQEEISAEVAAKAERVHELNNEIGKLLAKVEQLGAEGNVEESQK
VMDEVEKARAKKREAEEVYRNNSMPASSFQQQKLRLVCEVCASAYLGLHDNDRRRLADHFGGKLHLGFIEIREKLEELKRV
VAEKQEKRNRQERLKRREEREREKLRSSRSHSKNPKRSRSREHRRHRSRSMSRERKRRTRSRSREKRHRHRSRSS
SRSRSRSRHQRSRHSSRDRSRERSKRRSSKERFRDQDLASCDRDRSSRDRSPRDRDRDKKRSYESANGRSEDRRSSE
EREAGEI
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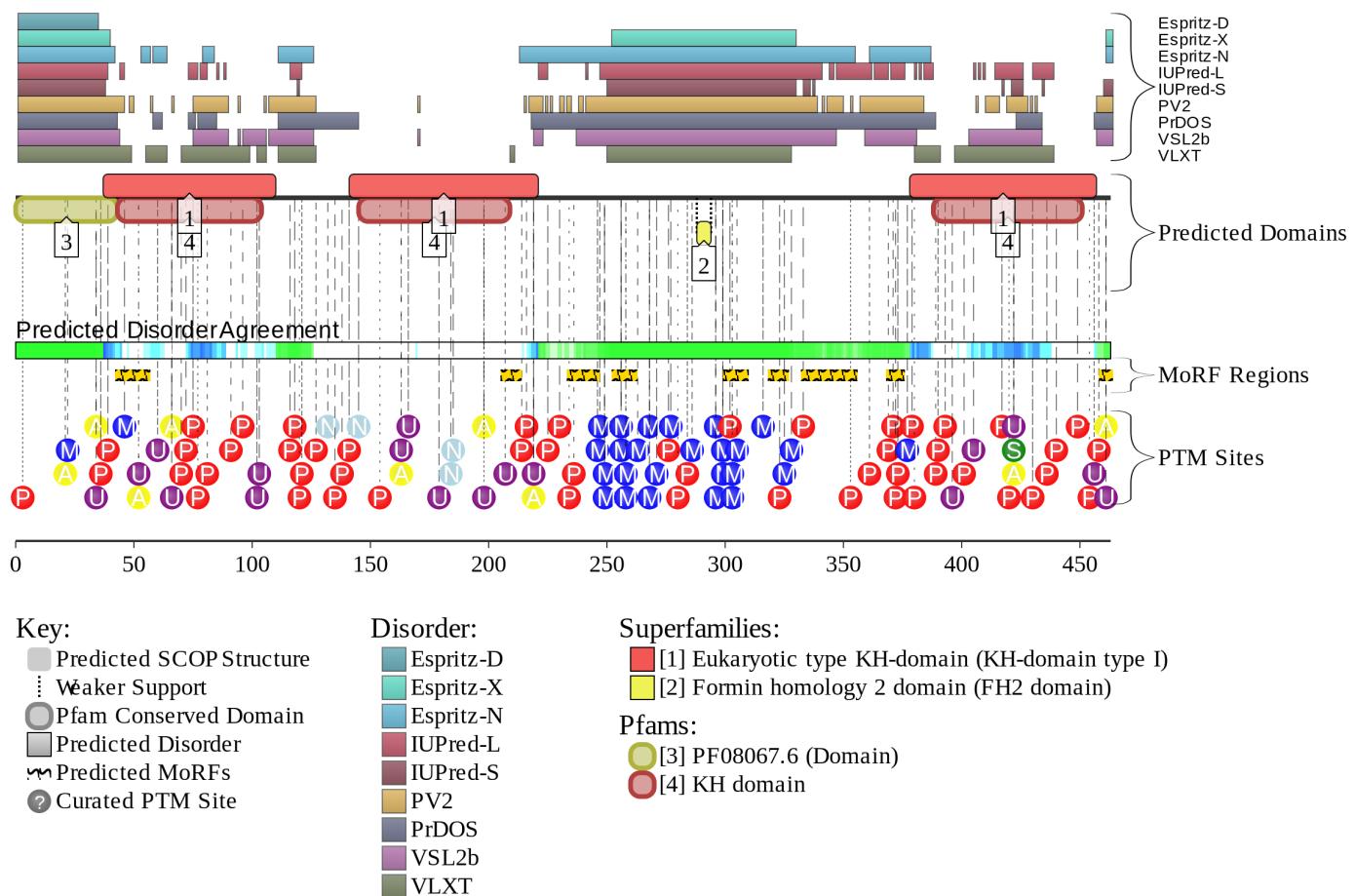
**Figure S4.** Amino acid sequence and functional disorder profile of heterogeneous nuclear ribonucleoprotein A0 (HNRNPA0; UniProt ID: Q13151).

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>sp|Q13151|ROA0_HUMAN Heterogeneous nuclear ribonucleoprotein A0 OS=Homo sapiens OX=9606 GN=HNRNPA0 PE=1 SV=1
MENSQLCKLFIGGLNVQTSESGRLRGHFEAFGTLTDCVVVNPQTKRSRCFGFVTYSNVEEADAAMAASPHAVDGNTV
ELKRAVSREDSARPGAHAKVKKLFGVGLKGDVAEGDLIEHFSQFGTVEKAETIADKQSGKKRGRGFVYFQNHDAAADK
AAVVKFHPIQGHRVEVKKAVPKEDIYSGGGGGSRSSRGGRGGGRGGGRDQNGLSKGGGGGYNSYGGYGGGGGGY
NAYGGGGGGSSYGGSDYGNFGFFGFSYSQHQSSYGPMSGGGGGGGSSWGRSNSGPYRGYYGGGGYGGSSF
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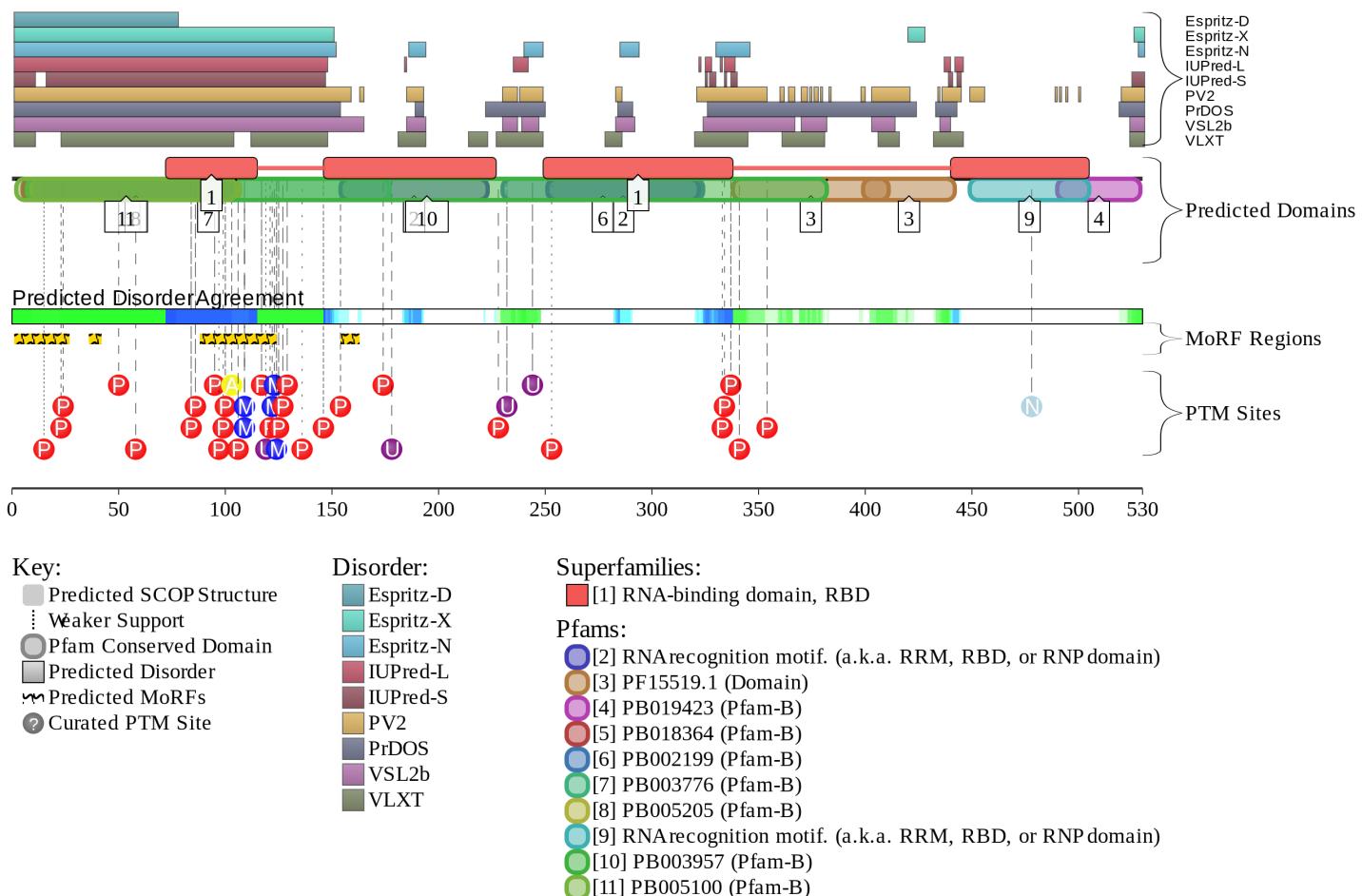
**Figure S5.** Amino acid sequence and functional disorder profile of heterogeneous nuclear ribonucleoprotein K (HNRNPK; UniProt ID: P61978).

```
>sp|P61978|HNRPK_HUMAN Heterogeneous nuclear ribonucleoprotein K OS=Homo sapiens OX=9606 GN=HNRNPK PE=1 SV=1
METEQQPEETFPNTETNGEFGKRPAMDMEEQAFKRSRNTDEMVELRILLQSKNAGAVIGKGGKNIKALRTDYNASVS
VPDSSGPERILSISADIEITIGEILKKIIPITLEEGLQLPSPTATSQLPLESDAVECLNYQHYKGSDFDCELRLLIHQS
LAGGIIGVKGAKIKELRENTQTTIKLFQECCPHSTDVVILGGKPDRVVECIKIILDLISESPIKGRAQPYDPNFYD
ETYDYGGFTMMFDDRRGRPVGFPMRGRGGFDRMPPGRGPMPSSRDRYDDMSPRRGPPPPPGRGGRRGGSRARNLP
LPPPPPPRGDLMAYDRRGRPGDRYDGMOVFSADETWDSAIDTWSPEWQMAYEHQGGSGYDYSYAGGRGSYGDLLGG
PIITTQVTIPKDLAGSIIKGQQRIKQIRHESGASIKIDEPLEGEDRIITITGTQDQIQNAQYLLQNSVKQYSGKF
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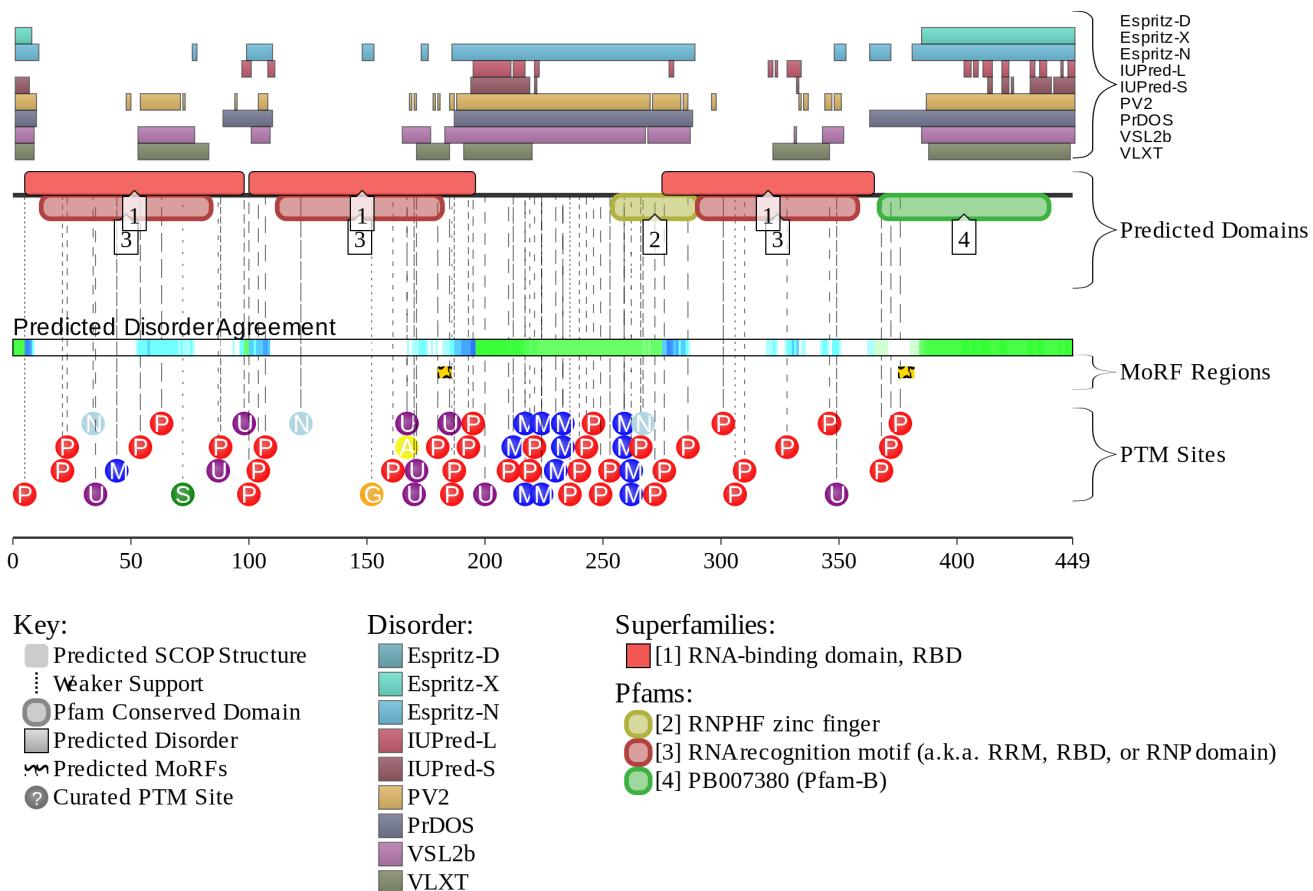
**Figure S6.** Amino acid sequence and functional disorder profile of RNA-binding protein 39 (RBM39; also known as CAPER alpha; Hepatocellular carcinoma protein 1; RNA-binding motif protein 39; RNA-binding region-containing protein 2; and Splicing factor HCC1UniProt ID: Q14498).

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>sp|Q14498|RBM39_HUMAN RNA-binding protein 39 OS=Homo sapiens OX=9606
GN=RBM39 PE=1 SV=2
MADDIDIEAMLEAPYKDENKLSSANGHEERSKKRSRSRSHRKRSKSKERKRSRDRERKKSKSRERKRSRSK
ERRRSRSRSRDRFRGRYRSPYSGPKFNSAIRGKIGLPHSIKLSRRRSRKSPFRKDSPVREPIDNLTEERDART
VFCMQLAARIRPRDLEEFFSTVGKVRDVRMISDRNSRSKGIAVYEFVDVSSVPLAIGLTGQRVLGVPIIVQASQE
KNRAAAMANNLQKGSAGPMRLYVGSLHFNITEDMLRGIFEPEFGRIESIQLMMDSETGRSKGYGFITFSDSECAKKAL
EQLNGFELAGRPMKVGHVTERTDASSASSFLDSDEERTGIDLGGTGRQLMARLAEGTGLQIPPAQQALQMSGSL
AFGAVAESFSVIFDLQTRLSQQTEASALAAAASVQPLATQCFCQLSNMFNPQTEEEVGWDTEIKDDVIEECNKHGGVIH
IYVDKNSAQGNVYVKCPSIAAAIAAVNALHGRWFAGKMITAAYVPLPTYHNLFPDSMTATQLLVPSRR
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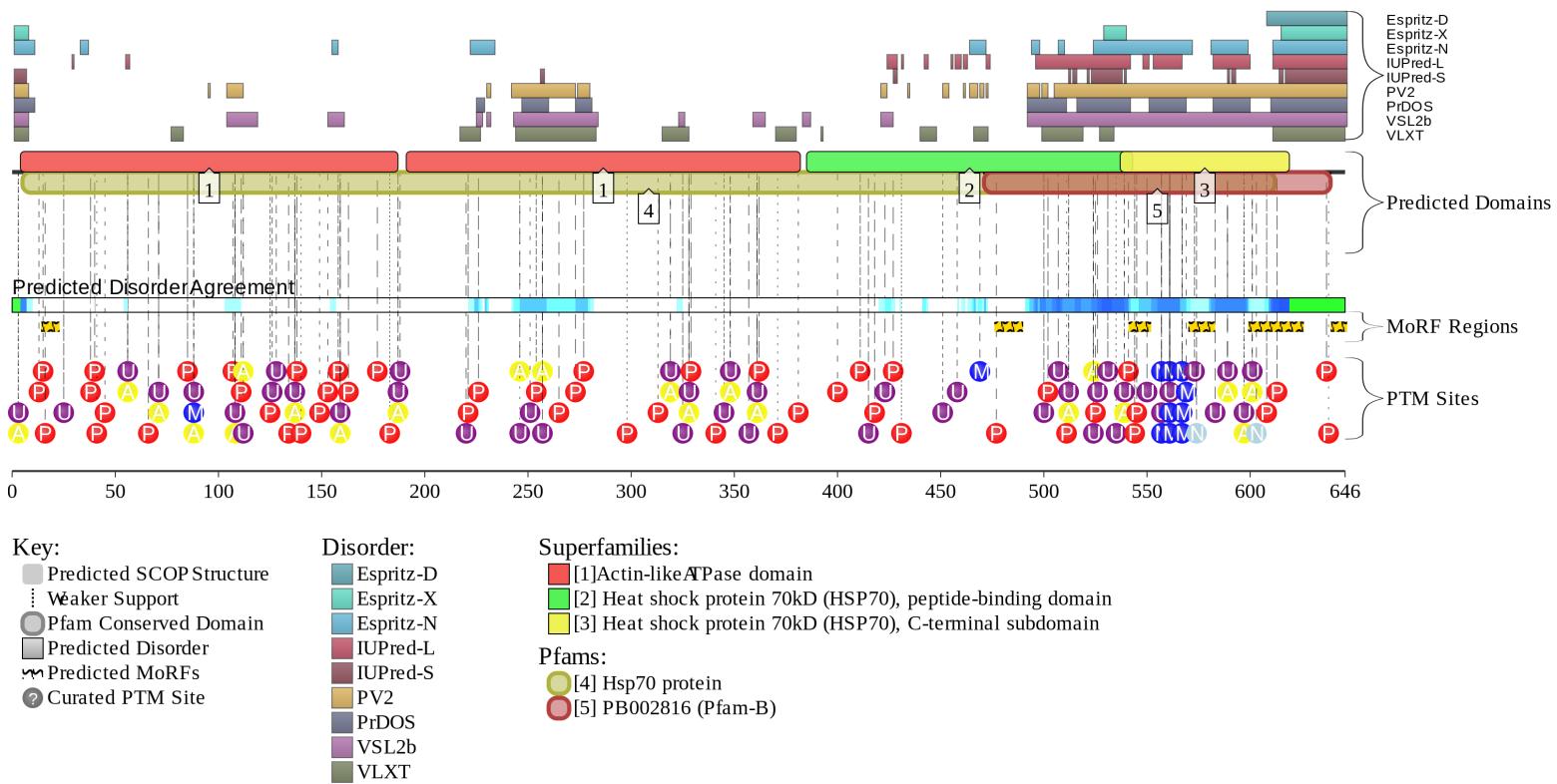
**Figure S7.** Amino acid sequence and functional disorder profile of heterogeneous nuclear ribonucleoprotein H (HNRNPH1; UniProt ID: P31943).

>sp|P31943|HNRH1\_HUMAN Heterogeneous nuclear ribonucleoprotein H OS=Homo sapiens OX=9606 GN=HNRNPH1 PE=1 SV=4  
MMLGTEGGEGFVVKVRGLPWCSADEVQRFFSDCKIQNGAQGIRFIYTREGRPSGEAFVELESEDEVKLALKKDRET  
MGHRYVEVFKSNNVEMDWLKHTGPNSPDTANDGFVRLRGLPFGCSKEIIVQFFSGLIEIVPNGITLPVDFQGRSTGE  
AFVQFASQEIAEKALKKHKERIGHRYIEIFKSSRAEVRTHYDPPRKLMMQRPGPYDRPGAGRGYNSIGRGAGFERM  
RRGAYGGYGGYDDYNGYNDGYGFGSDRFGRDLNYCFSGMSDHRYGDGGSTFQSTGHCVHMRLPYRATENDIYNF  
FSPLNPVRVHIEIGPDRVTGEADVEFATHEDAVAAMSKDKANMQHRYVELFLNSTAGSGGAYEHRYVELFLNSTA  
GASGGAYGSQMMGGMGLSNQSSYGGPASQQLSGGGGGYGGQSSMSGYDQVLQENSSDFQSNIA



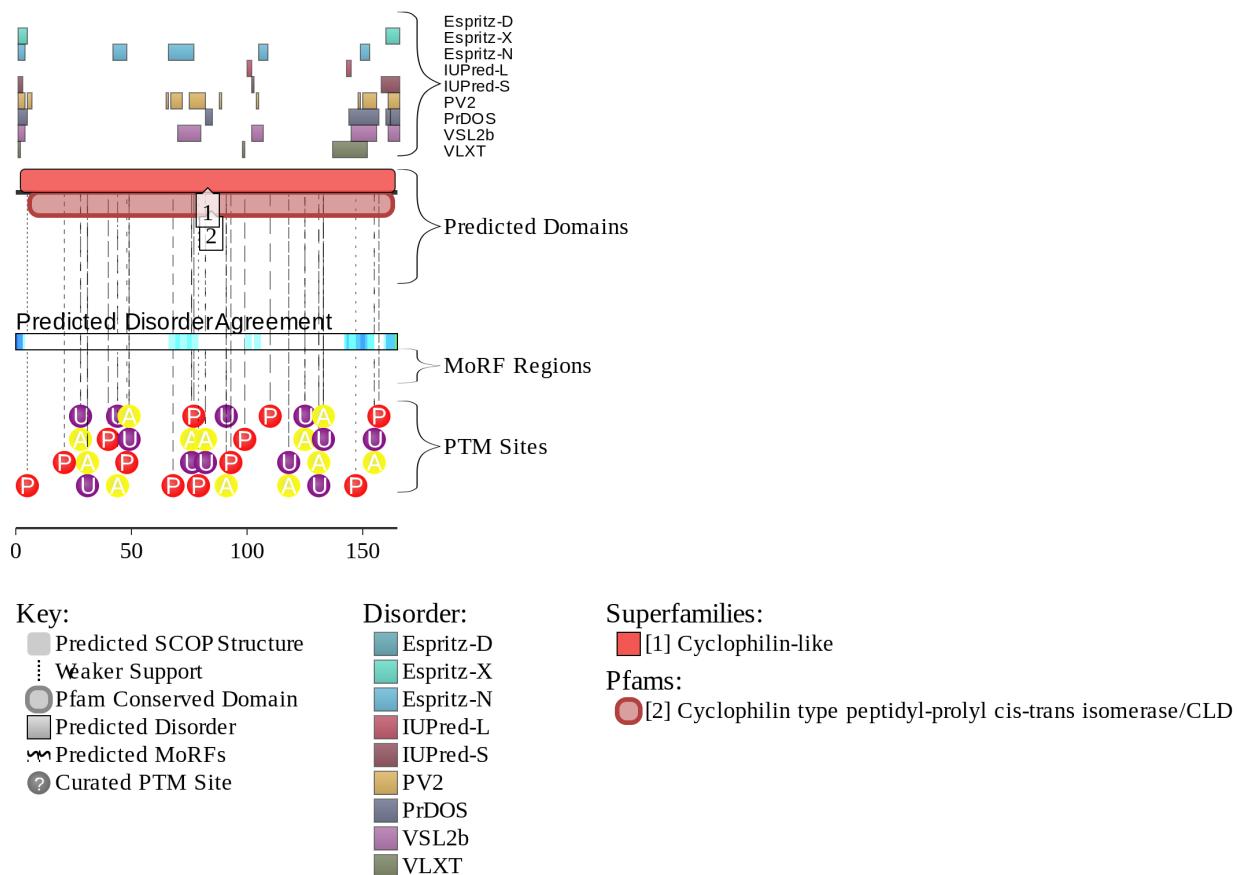
**Figure S8.** Amino acid sequence and functional disorder profile of heat shock cognate 71 kDa protein (also known as heat shock 70 kDa protein 8 (HSPA8) and Lipopolysaccharide-associated protein 1 (LAP-1; LPS-associated protein 1); UniProt ID: P11142).

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>sp|P11142|HSP7C_HUMAN Heat shock cognate 71 kDa protein OS=Homo sapiens
OX=9606 GN=HSPA8 PE=1 SV=1
MSKGPAVGIDLGTTYSCGVFQHGKVEIIANDQGNRTTPSYVAFTDTERLIGDAAKNQVAMNPTNTVDAKRLIGRR
FDDAVVQSDMKHWPFMVVNDAGRPKVQVEYKGETKSFYPEEVSSMVLTKMKEIAEAYLGKTVNAVVTVPAYFNDSQ
RQATKDAGTIAGLNVLRIINEPTAAIAYGLDKKVGAEARNVLIFDLGGTFDVSILTIEDGIFEVKSTAGDTHLGGE
DFDNRMVNHFIAEFKRKHKKDISENKRAVRLRTACERAKRTLSSSTQASIEIDSPLYEGIDFYTSITRARFEELNAD
LFRGTLDPVEKALRDAKLDKSQIHIDIVLVGGSTRIPKIQKLLQDFNGKELNKSINPDEAVAYGAAVQAAILSGDKS
ENVQDLLLLDVTPSLGIETAGGVMVTLIKRNNTIPTKQTQFTTYSNDQPGVLIQVYGERAMTKDNNLLGFELT
GIPPAPRGVPQIEVTFDIDANGILNVSAVDKSTGKENKITITNDKGRLSKEDIERMVQEAEKYKAEDEKQRDKVSSK
NSLESYAFNMKATVEDEKLQGKINDEDKQKILDKCNEIINWLDKNQTAEEKEFEHQQKELEKVCNPPIITKLYQSAGG
MPGGMPGGFPGGAPPSSGGASSGPTIEEV
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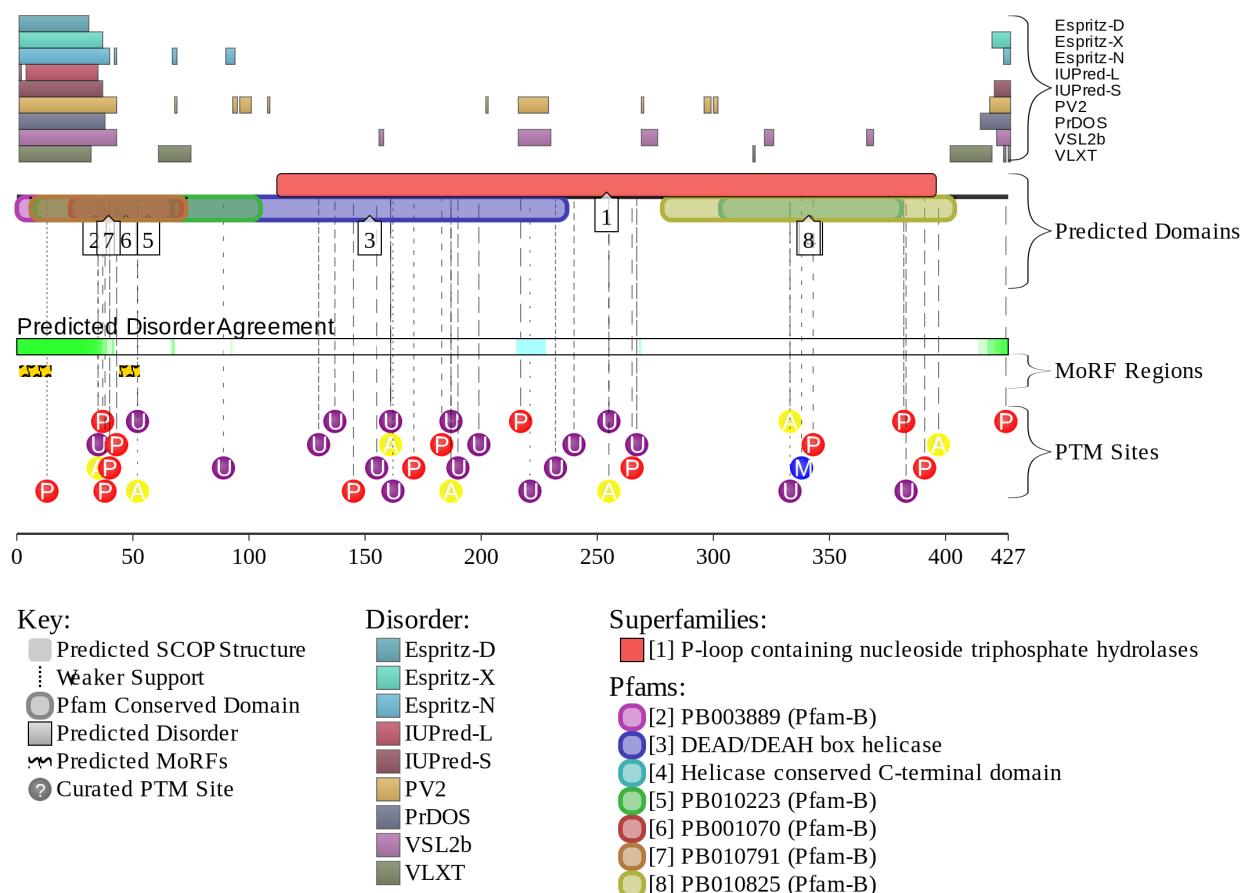
**Figure S9.** Amino acid sequence and functional disorder profile of peptidyl-prolyl cis-trans isomerase A (PPIA, UniProt ID: P62937).

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>sp|P62937|PPIA_HUMAN Peptidyl-prolyl cis-trans isomerase A OS=Homo sapiens
OX=9606 GN=PPIA PE=1 SV=2
MVNPPTVFFDIAVDGEPLGRVSFELFADKVPKTAENFRALSTGEKFGFYKGSCFHRIIPGFMCQGGDFTRHNGGKS
IYGEKFEDENFILKHTGPGILSMANAGPNTNGSQFFICTAKTEWLGDGHVVFGKVKEGMNIVEAMERFGSRNGKTSK
KITIADCQLE
```



**Figure S10.** Amino acid sequence and functional disorder profile of ATP-dependent RNA helicase DDX39A (also known as DEAD box protein 39 (DDX39A) and Nuclear RNA helicase URH49 (UniProt ID: O00148).

```
>sp|O00148|DX39A_HUMAN ATP-dependent RNA helicase DDX39A OS=Homo sapiens
OX=9606 GN=DDX39A PE=1 SV=2
MAEQDVENDLLDYDEEEEPQAPQESTPAPPKKDIKGSYVSIHSSGFRDFLLKPELLRAIVDCGFEHPSEVQHECIPQ
AILGMDVLQCQAKSGMGKTAVFVLATLQQIEPVNGQVTVLVMCHTRELAFQISKEYERFSKYMPSVKVSVFFGGLSIK
KDEEVLKKNCPHVVGTPGRILALVRNRSFSLKNVKHFLDECDKMLEQLDMRRDVQEIRLTPHEKQCMMFSATLS
KDIRPVCRKFMQDPMEVFVDDETKLT LHGLQQYYVKLKDSEKNRKLFDLLDVL EFQNQVII FVKS VQRCM ALAQLLVE
QNFPAIAIH RGMAQEER LSR YQQFKDFQ RRI L VATNL FGR GMDI ER VNIV FNYD MPED SDT YLHR VARAG RFGTKGL
AITFVSDENDAKILNDVQDRF EVNVAEL PEEIDISTY IEQSR
```



**Table S1.** Characteristics of the proteome of the heat shock-induced A-bodies. DPR are droplet-promoting regions, nHS are number of aggregation hot spots, Na4vSS are Normalized a4v Sequence Sum for 100 residues used for aggregation prediction.

Gene Name	UniProtID	FastA	Sequence	PER(VSL2B)	Disorder Classifications	FuzDrop Score	FuzDrop LLPS	PSPredict or Score	PSPredict or LLPS	Fuz&PSP LLPS	Fuz&PSP No LLPS	Number of DPRs	Role in MLO	nHS	Na4vSS	Aggregation Predicted	Aggregation DB scanning	Hydrophobicity	Charge
VHL	P40337	>sp P40337 PGTGRRIH	55.87	IDP	0.6579	Yes	0.0102	No				2	Driver	6	-23	No	Yes	-0.78	-0.073
GSTM3	P21266	>sp P21266 MSCESSM	14.22	Partial IDP	0.1175	No	0.0047	No		+		0	No	9	-2.8	No	No	-0.41	-0.016
FEN1	P39748	>sp P39748 MGIQGLA	44.47	IDP	0.2197	No	0.2899	No		+		2	Client	11	-14.1	No	No	-0.53	0.026
MPP10	O00566	>sp O00566 MAPQVW	76.95	IDP	0.9394	Yes	0.2793	No				7	Driver	9	-36.6	No	No	-1.11	-0.070
PCBP2	Q15366	>sp Q15366 MDTGVI	44.93	IDP	0.9028	Yes	0.2466	No				3	Driver	10	-3.4	No	No	-0.14	0.003
AN32B	Q92688	>sp Q92688 MMDKRR	61.75	IDP	0.4995	No	0.5683	Yes				1	Client	3	-47.7	No	No	-1.16	-0.263
MYL6	P60660	>sp P60660 MCDFTED	19.21	Partial IDP	0.1325	No	0.0072	No		+		0	No	7	-13.2	No	No	-0.39	-0.079
LC7L2	Q9Y383	>sp Q9Y383 M3AAQAN	74.49	IDP	0.9512	Yes	0.0666	No				3	Driver	2	-50.2	No	No	-1.49	0.088
RRP12	Q5JTH9	>sp Q5JTH9 MGRSGKL	36.24	IDP	0.5879	No	0.5229	Yes				7	Client	44	-4.9	No	No	-0.23	0.032
COMT	P21964	>sp P21964 MPEAPPL	7.01	Ordered	0.1043	No	0.0046	No		+		0	No	9	9.1	Yes	No	0.16	-0.022
POP1	Q99575	>sp Q99575 MSNAKER	50.88	IDP	0.9448	Yes	0.6756	Yes		+		8	Driver	37	-16	No	No	-0.62	0.056
UT14A	Q9BVJ6	>sp Q9BVJ6 MTANRLA	81.06	IDP	0.9196	Yes	0.3773	No				6	Driver	5	-29.4	No	No	-0.94	0.011
HSPB1	P04792	>sp P04792 MTERRVPI	57.56	IDP	0.6398	Yes	0.0151	No				2	Driver	3	-13.1	No	No	-0.57	-0.002
CDC73	Q6P1J9	>sp Q6P1J9 MADVLSV	53.67	IDP	0.4019	No	0.4929	No		+		4	Client	14	-12.5	No	No	-0.58	0.046
SUMO2	P61956	>sp P61956 MADEKPK	52.63	IDP	0.1814	No	0.4944	No		+		1	Client	2	-30.2	No	No	-0.89	-0.021
ML12A	P19105	>sp P19105 MSSKRTK	41.52	IDP	0.6107	Yes	0.0338	No				1	Driver	5	-25	No	No	-0.83	-0.064
XPO2	P55060	>sp P55060 MELSDAN	12.05	Partial IDP	0.1253	No	0.1683	No		+		0	No	39	6.8	Yes	No	-0.03	-0.013
NOM1	Q5C9Z4	>sp Q5C9Z4 MAASRSA	55.47	IDP	0.9475	Yes	0.8288	Yes		+		5	Driver	19	-21.1	No	No	-0.64	0.016
BYST	Q13895	>sp Q13895 MPKFKAA	48.97	IDP	0.2822	No	0.1084	No		+		2	Client	15	-8.2	No	No	-0.37	0.015
SSF1	Q9NQ55	>sp Q9NQ55 MGQSGRS	54.55	IDP	0.7386	Yes	0.0358	No				4	Driver	12	-24.2	No	No	-0.76	0.079
NUMA1	Q14980	>sp Q14980 MTLHATR	94.42	IDP	1	Yes	0.6606	Yes		+		26	Driver	20	-34.9492	No	No	-0.92	-0.013
HS71B	P0DMV9	>sp P0DMV9 MAKAAAIC	29.64	Partial IDP	0.3928	No	0.1251	No		+		2	Client	18	-12.3	No	No	-0.38	-0.012
PPIA	P62937	>sp P62937 MVNPNTVF	20.61	Partial IDP	0.2089	No	0.0057	No		+		0	No	7	-6.7	No	No	-0.32	0.018
NOL10	Q9BSC4	>sp Q9BSC4 MQVSSLN	44.62	IDP	0.2473	No	0.0748	No		+		2	Client	19	-15.6	No	No	-0.75	0.022
1433Z	P63104	>sp P63104 MDKNELV	53.88	IDP	0.2169	No	0.0048	No		+		1	Client	6	-18.2	No	No	-0.62	-0.055
AATF	Q9NY61	>sp Q9NY61 MAGQPQL	65.71	IDP	0.7654	Yes	0.2619	No				4	Driver	8	-31.5	No	No	-0.91	-0.058
RRP1	P56182	>sp P56182 MVSRVQL	50.76	IDP	0.7152	Yes	0.0183	No				2	Driver	10	-26.9	No	No	-0.84	0.029
PHF6	Q8IW50	>sp Q8IW50 MSSVEQI	54.52	IDP	0.8738	Yes	0.9423	Yes		+		4	Driver	6	-30.5	No	No	-1.01	0.067
NOC4L	Q9BV14	>sp Q9BV14 MEREPGA	21.71	Partial IDP	0.3262	No	0.0045	No		+		2	Client	21	3.7	Yes	No	-0.08	0.018
ESF1	Q9H501	>sp Q9H501 MSSKQEIN	81.67	IDP	0.9412	Yes	0.43	No				10	Driver	11	-38.5	No	No	-1.23	-0.044
KRR1	Q13601	>sp Q13601 MASPSSLER	55.64	IDP	0.6197	Yes	0.2739	No				4	Driver	10	-18.5	No	No	-0.78	0.084
HS90B	P08238	>sp P08238 MPEEVHH	45.03	IDP	0.3869	No	0.2768	No		+		3	Client	18	-17.3	No	No	-0.68	-0.046
HNRPF	P52597	>sp P52597 MMLGPEC	40.48	IDP	0.5243	No	0.0968	No		+		3	Client	10	-10.7	No	No	-0.47	-0.020
NOP14	P78316	>sp P78316 MAKAKKV	70.36	IDP	0.859	Yes	0.8195	Yes		+		7	Driver	16	-23.3	No	No	-0.76	0.013
DX39A	O00148	>sp O00148 MAEQDVE	18.27	Partial IDP	0.1354	No	0.0079	No		+		1	Client	16	-4	No	No	-0.30	-0.016
HNRH1	P31943	>sp P31943 MMLGTEC	51.22	IDP	0.9486	Yes	0.9727	Yes		+		3	Driver	9	-16	No	No	-0.60	-0.004
U2AF1	Q01081	>sp Q01081 MAEYLASI	45.83	IDP	0.3606	No	0.542	Yes				1	Client	6	-31.9	No	No	-1.06	0.048
K0020	Q15397	>sp Q15397 MEVKGKK	34.41	IDP	0.2025	No	0.3415	No		+		2	Client	23	-12	No	No	-0.56	0.076
BCCIP	Q9P287	>sp Q9P287 MARSRSKR	35.99	IDP	0.3828	No	0.6835	Yes				1	Client	8	-20.5	No	No	-0.69	-0.083
SERA	O43175	>sp O43175 MAFANLR	8.26	Ordered	0.1623	No	0.0149	No		+		1	Client	21	1.2	Yes	No	0.09	0.002
IF5A2	Q9GV4	>sp Q9GV4 MADEIDFT	35.95	IDP	0.2524	No	0.0061	No		+		1	Client	6	-9.4	No	No	-0.35	-0.020
ROAO	Q13151	>sp Q13151 MENSQLC	61.97	IDP	0.9818	Yes	0.9959	Yes		+		1	Driver	5	-20.5	No	No	-0.59	0.043
DDX49	Q9Y6V7	>sp Q9Y6V7 MAGFAEL	21.95	Partial IDP	0.1582	No	0.0188	No		+		1	Client	19	-2.2	No	No	-0.18	0.041
TIF1B	Q13263	>sp Q13263 MAASAAA	63.11	IDP	0.9946	Yes	0.9829	Yes		+		6	Driver	18	-17.4	No	No	-0.39	-0.013
HNRH2	P55795	>sp P55795 MMLTEG	52.34	IDP	0.9768	Yes	0.9661	Yes		+		2							

MGN2	Q96A72	>sp Q96A MAVASDF	16.22	Partial IDP	0.1468	No	0.0254	No	+ +	0	No	5	-1.5	No	No	-0.38	-0.003
NONO	Q15233	>sp Q152 MQSNKTF	76.86	IDP	0.9869	Yes	0.609	Yes	+ +	4	Driver	8	-30.6	No	No	-1.00	0.025
RS26	P62854	>sp P6285 MTKKRRN	50.43	IDP	0.8969	Yes	0.0222	No	+ +	2	Driver	3	-16.5	No	No	-0.66	0.204
NPM	P06748	>sp P067 MEDSMDN	61.22	IDP	0.9553	Yes	0.9884	Yes	+ +	2	Driver	8	-35.6	No	Yes	-0.97	-0.073
RRP8	O43159	>sp O431 MFFEEPEW	62.06	IDP	0.9406	Yes	0.9612	Yes	+ +	4	Driver	7	-24.3	No	No	-0.76	0.058
RBM8A	Q9Y5S9	>sp Q9Y5 MADVLDL	61.49	IDP	0.9709	Yes	0.0354	No	+ +	3	Driver	3	-38	No	No	-1.15	-0.020
MCM3	P25205	>sp P2520 MAGTVL	46.66	IDP	0.6128	Yes	0.2455	No	+ +	6	Driver	23	-16.8	No	No	-0.56	-0.014
RBM28	Q9NW13	>sp Q9NW MAGLTLF	62.19	IDP	0.9274	Yes	0.4076	No	+ +	6	Driver	12	-30.8	No	No	-0.96	0.041
MATR3	P43243	>sp P4324 MSKSFQQ	72.02	IDP	0.9982	Yes	0.9685	Yes	+ +	7	Driver	18	-29.2	No	No	-0.93	-0.006
RL4	P36578	>sp P3657 MACARPL	50.35	IDP	0.3958	No	0.0503	No	+ +	4	Client	11	-14.3	No	No	-0.61	0.177
RS9	P46781	>sp P4678 MPVARSW	28.87	Partial IDP	0.2294	No	0.0171	No	+ +	1	Client	7	-13.4	No	No	-0.64	0.124
MCM4	P33991	>sp P3399 MSSPASTP	35.46	IDP	0.324	No	0.6159	Yes	+ +	3	Client	29	-10.7	No	No	-0.41	0.003
DDX24	Q9GZR7	>sp Q9GZ MKLKRTK	49.36	IDP	0.8808	Yes	0.8971	Yes	+ +	9	Driver	20	-19.1	No	No	-0.66	0.034
COPB	P53618	>sp P5361 MTAAENV	23.82	Partial IDP	0.282	No	0.156	No	+ +	2	Client	34	1	Yes	No	-0.09	-0.008
HNRPC	P07910	>sp P0791 MASNVTN	75.82	IDP	0.9961	Yes	0.7935	Yes	+ +	3	Driver	5	-35.4	No	No	-0.98	-0.036
SIN3A	Q96ST3	>sp Q96S MKRLLDD	48.55	IDP	0.851	Yes	0.9604	Yes	+ +	8	Driver	29	-16.2	No	No	-0.69	0.012
RS24	P62847	>sp P6284 MNDTNTF	47.37	IDP	0.202	No	0.2207	No	+ +	1	Client	3	-21	No	No	-0.99	0.203
DDX50	Q9BQ39	>sp Q9BQ MPGKLLW	38.81	IDP	0.5348	No	0.9234	Yes	+ +	2	Client	22	-20.2	No	No	-0.71	0.037
ETFA	P13804	>sp P13804 MFRAAAP	19.52	Partial IDP	0.2221	No	0.0106	No	+ +	1	Client	14	2.9	Yes	No	0.15	0.020
RL10A	P62906	>sp P62906 MSSKVSRC	32.26	IDP	0.1548	No	0.0108	No	+ +	1	Client	6	-5.5	No	No	-0.41	0.111
MRT4	Q9UKD2	>sp Q9UK MPKSKRD	32.22	IDP	0.2086	No	0.0172	No	+ +	1	Client	6	-15.8	No	No	-0.71	0.015
RL15	P61313	>sp P61313 MAGAYKI	51.47	IDP	0.6722	Yes	0.0107	No	+ +	2	Driver	5	-16.8	No	No	-0.93	0.218
SF3B3	Q15393	>sp Q15393 MFLYNLTL	16.68	Partial IDP	0.1376	No	0.0812	No	+ +	5	Client	47	-0.6	No	No	-0.17	-0.025
TBA1B	P68363	>sp P68363 MRECISHV	16.63	Partial IDP	0.1976	No	0.0132	No	+ +	1	Client	19	-3.1	No	No	-0.23	-0.039
NOL11	Q9H8H0	>sp Q9H8H MAALEEFP	21.97	Partial IDP	0.1517	No	0.1976	No	+ +	1	Client	29	4.1	Yes	No	-0.11	-0.008
RBM19	Q9Y4C8	>sp Q9Y4C8 MSRLIVKNA	58.96	IDP	0.9431	Yes	0.898	Yes	+ +	5	Driver	26	-22.7	No	No	-0.75	-0.001
RL17	P18621	>sp P18621 MVRYSLDR	45.11	IDP	0.1848	No	0.0159	No	+ +	0	No	3	-21.2	No	No	-0.90	0.141
DDX54	Q8TDD1	>sp Q8TDD1 MAADKGP	57.32	IDP	0.9391	Yes	0.6687	Yes	+ +	8	Driver	22	-21.1	No	No	-0.68	0.065
RL12	P30050	>sp P30050 MPPKFDP	23.64	Partial IDP	0.1717	No	0.0066	No	+ +	0	No	5	-12.6	No	No	-0.35	0.055
RPF2	Q9H7B2	>sp Q9H7B2 MDTLDVR	47.39	IDP	0.2631	No	0.0298	No	+ +	1	Client	7	-11	No	No	-0.68	0.119
WDR3	Q9UNX4	>sp Q9UNX4 MGLTKQY	28.53	Partial IDP	0.2985	No	0.0466	No	+ +	4	Client	36	-3.2	No	No	-0.26	0.002
OZF	Q15072	>sp Q15072 MSHLSQQ	69.52	IDP	0.269	No	0.1973	No	+ +	2	Client	4	-22.1	No	No	-0.88	0.106
DDX56	Q9NY93	>sp Q9NY93 MDESEALC	30.71	IDP	0.2624	No	0.0198	No	+ +	2	Client	12	-4.8	No	No	-0.27	0.043
HSP76	P17066	>sp P17066 MQAPREL	32.66	IDP	0.2901	No	0.2186	No	+ +	2	Client	17	-15	No	No	-0.46	-0.006
RPAB3	P52434	>sp P52434 MAGILFED	25.33	Partial IDP	0.1667	No	0.0027	No	+ +	1	Client	6	-5.8	No	No	-0.34	-0.073
IF4A3	P38919	>sp P38919 MATTATM	18.73	Partial IDP	0.1758	No	0.017	No	+ +	1	Client	15	-2.9	No	No	-0.26	0.001
MYH14	Q7Z406	>sp Q7Z406 MAAVTMS	69.27	IDP	0.9776	Yes	0.5611	Yes	+ +	18	Driver	28	-29.2	No	No	-0.81	-0.016
CMS1	Q9BQ75	>sp Q9BQ75 MADDLGD	42.65	IDP	0.2969	No	0.0918	No	+ +	1	Client	9	-10.7	No	No	-0.50	0.043
WDR75	Q8IWA0	>sp Q8IWA0 MVEENIR	22.77	Partial IDP	0.2123	No	0.6001	Yes	+ +	1	Client	28	0.3	Yes	No	-0.25	-0.011
HNRPK	P61978	>sp P61978 Meteqpe	56.8	IDP	0.9031	Yes	0.2392	No	+ +	4	Driver	9	-19.7	No	No	-0.70	-0.013
RL9	P32969	>sp P32969 MKTILSN	7.81	Ordered	0.092	No	0.0241	No	+ +	0	No	8	-4.3	No	No	-0.35	0.076
MK67I	Q9BYG3	>sp Q9BYG3 MATFSGP	69.62	IDP	0.587	No	0.8185	Yes	+ +	4	Client	5	-21.1	No	No	-0.95	0.082
SMD3	P62318	>sp P62318 MSIGVPIK	41.27	IDP	0.2632	No	0.0185	No	+ +	1	Client	6	-11.4	No	No	-0.43	0.103
RL18A	Q02543	>sp Q02543 MKA5GTL	38.64	IDP	0.1807	No	0.0071	No	+ +	1	Client	7	-7.4	No	No	-0.67	0.182
RS4X	P62701	>sp P62701 MARGPKK	13.31	Partial IDP	0.1382	No	0.0053	No	+ +	0	No	10	-3.3	No	No	-0.33	0.106
ELAV1	Q15717	>sp Q15717 MSNGYED	25.46	Partial IDP	0.1844	No	0.0132	No	+ +	0	No	14	-6	No	No	-0.36	0.034
RL6	Q02878	>sp Q02878 MAGEKVE	61.46	IDP	0.5433	No	0.1071	No	+ +	2	Client	10	-12.9	No	No	-0.73	0.194
RL7A	P62424	>sp P62424 MPKGKKA	41.73	IDP	0.3147	No	0.566	Yes	+ +	3	Client	10	-12.1	No	No	-0.55	0.162
WDR12	Q9GZL7	>sp Q9GZL7 MAQLQTR	20.09	Partial IDP	0.2042	No	0.0197	No	+ +	1	Client	11	-6	No			

**Table S2.** Characteristics of the proteome of the acidosis-induced A-bodies. DPR are droplet-promoting regions, nHS are number of aggregation hot spots, Na4vSS are Normalized a4v Sequence Sum for 100 residues used for aggregation prediction.

Gene Name	UniProtID	Fasta	Sequence	PER(VSL2B)	Disorder Classifications	FuzDrop Score	FuzDrop LLPS	PSPredictor Score	PSPredictor LLPS	Fuz&PSP LLPS	Fuz&PSP No LLPS	Number of DPRs	Role in MLO	nHS	Na4vSS	Aggregation Predicted	Aggregation DB scanning	Hydrophobicity	Charge
VHL	P40337	>sp P40337 MPRRAENP	55.87	IDP	0.6579	Yes	0.0102	No				2	Driver	6	-23.0	No	Yes	-0.78	-0.073
AKAP13	Q12802	>sp Q12802 MKNLNPQQ	75.97	IDP	1.0000	Yes	0.7561	Yes	+			30	Driver	46	-22.7	No	No	-0.64	-0.030
HERC2	O95714	>sp O95714 MPSESFCL	28.57	Partial IDP	0.9976	Yes	0.3504	No				28	Driver	153	-5.0	No	No	-0.19	-0.005
NOSIP	Q9Y314	>sp Q9Y314 MTRHGNK	54.49	IDP	0.8464	Yes	0.0313	No				2	Driver	6	-15.1	No	No	-0.52	0.040
HSPB1	P04792	>sp P04792 MTERRVPF	57.56	IDP	0.6398	Yes	0.0151	No				2	Driver	3	-13.1	No	No	-0.57	-0.002
PPIA	P62937	>sp P62937 MVNPTVFP	20.61	Partial IDP	0.2089	No	0.0057	No		+	0	0	No	7	-6.7	No	No	-0.32	0.018
HSPA1A	P08107	>sp P08107 MAKAAAEG	29.64	Partial IDP	0.3928	No	0.1251	No			2	Client	18	-12.3	No	No	-0.38	-0.012	
CIRBP	Q14011	>sp Q14011 MASDEGKU	77.91	IDP	0.9977	Yes	0.9806	Yes	+			1	Driver	4	-32.7	No	No	-1.06	0.032
LYZ	P61626	>sp P61626 MKA1IVLG	8.11	Ordered	0.1266	No	0.0241	No			0	No	5	-2.6	No	Yes	-0.19	0.064	
PABPN1	Q86U42	>sp Q86U42 MAAAAAAA	76.14	IDP	0.9558	Yes	0.9338	Yes	+			2	Driver	4	-30.7	No	No	-0.82	-0.036
HNRNPA1	P09651	>sp P09651 MSKSESPK	70.16	IDP	0.9998	Yes	0.9977	Yes	+			1	Driver	7	-27.8	No	No	-0.89	0.030
ANP32B	Q92688	>sp Q92688 MDMKRRII	61.75	IDP	0.4995	No	0.5683	Yes				1	Client	3	-47.7	No	No	-1.16	-0.263
PTBP1	P26599	>sp P26599 MDGIVPDI	43.27	IDP	0.5386	No	0.9415	Yes				4	Client	25	-2.7	No	No	-0.12	0.030
PTGES3	Q15185	>sp Q15185 MQPASAKW	56.88	IDP	0.7660	Yes	0.1522	No				1	Driver	4	-35.0	No	No	-1.05	-0.116
EEF1B2	P24534	>sp P24534 MGFGDLKS	40.89	IDP	0.4031	No	0.1355	No			2	Client	7	-17.1	No	No	-0.49	-0.076	
YBX1	P67809	>sp P67809 MSSEAETQ	82.41	IDP	0.9970	Yes	0.984	Yes	+			2	Driver	1	-47.4	No	No	-1.48	0.049
CCT4	P50991	>sp P50991 MPENVAPR	22.82	Partial IDP	0.2193	No	0.1655	No				1	Client	23	-0.1	No	No	0.03	0.009
SYMPK	Q92797	>sp Q92797 MAGSGGD	35.56	IDP	0.9426	Yes	0.5602	Yes	+			7	Driver	49	-3.9	No	No	-0.20	-0.005
ATRX	P46100	>sp P46100 MTAEPMSI	72.03	IDP	0.9989	Yes	0.9671	Yes	+			17	Driver	37	-32.6	No	No	-1.06	0.000
TAF15	Q92804	>sp Q92804 MSDSGSY	92.57	IDP	1.0000	Yes	0.9952	Yes	+			4	Driver	3	-50.1	No	No	-1.52	0.009
BLVRB	P30043	>sp P30043 MAVKKIAIR	16.99	Partial IDP	0.1735	No	0.0317	No			1	Client	9	-1.2	No	No	-0.06	0.022	
HNRNPH1	P31943	>sp P31943 MMLGTEG	51.22	IDP	0.9486	Yes	0.9727	Yes	+			3	Driver	9	-16.0	No	No	-0.60	-0.004
HNRNPC	P07910	>sp P07910 MASVNTN	75.82	IDP	0.9961	Yes	0.7935	Yes	+			3	Driver	5	-35.4	No	No	-0.98	-0.036
WDR46	O15213	>sp O15213 METAPKPG	47.05	IDP	0.7993	Yes	0.7594	Yes	+			4	Driver	19	-16.9	No	No	-0.60	0.062
NCL	P19338	>sp P19338 MVKLAKAG	86.2	IDP	0.9872	Yes	0.941	Yes	+			7	Driver	14	-45.4	No	No	-1.13	-0.084
RSBN1	Q5VWQ0	>sp Q5VWQ0 MFISGRRT	51.87	IDP	0.8958	Yes	0.9117	Yes	+			8	Driver	15	-18.7	No	No	-0.66	0.032
YWHAZ	P63104	>sp P63104 MDKNELVQ	53.88	IDP	0.2169	No	0.0048	No			1	Client	6	-18.2	No	No	-0.62	-0.055	
FOXA1	P55317	>sp P55317 MLGTVKML	83.9	IDP	0.9998	Yes	0.9673	Yes	+			4	Driver	12	-14.2	No	No	-0.48	0.028
ACTBL2	Q562R1	>sp Q562R1 MTDNELSA	14.1	Partial IDP	0.1639	No	0.0223	No			0	No	16	0.8	Yes	No	-0.19	-0.016	
XRCC6	P12956	>sp P12956 MSGWESY	43.19	IDP	0.3735	No	0.4906	No			3	Client	19	-9.8	No	No	-0.54	0.001	
HNRNPD	Q14103	>sp Q14103 MSEEQFGC	83.1	IDP	0.9960	Yes	0.9949	Yes	+			2	Driver	7	-27.1	No	No	-0.92	0.010
HNRNPA3	P51991	>sp P51991 MEVKPPPG	70.11	IDP	0.9994	Yes	0.9966	Yes	+			2	Driver	6	-30.9	No	No	-0.99	0.030
RPL29	P47914	>sp P47914 MAKSKNHT	100	IDP	0.8048	Yes	0.6051	Yes	+			2	Driver	1	-34.1	No	No	-1.09	0.267
ZC3H18	Q86VM9	>sp Q86VM9 MDVAESPE	92.86	IDP	0.9998	Yes	0.9929	Yes	+			4	Driver	5	-54.1	No	No	-1.59	0.010
ISY1	Q9ULR0	>sp Q9ULR0 MARNAEK	72.98	IDP	0.5749	No	0.2225	No			1	Client	2	-31.0	No	No	-1.00	-0.037	
HNRNPK	P61978	>sp P61978 MTEEQPEE	56.8	IDP	0.9031	Yes	0.2392	No			4	Driver	9	-19.7	No	No	-0.70	-0.013	
SEC22B	O75396	>sp O75396 MVLLTMI	32.09	IDP	0.2068	No	0.0114	No			1	Client	6	1.8	Yes	No	-0.18	0.021	
HNRNPA2	P22626	>sp P22626 MEKLTLET	69.97	IDP	0.9808	Yes	0.9977	Yes	+			2	Driver	4	-26.5	No	No	-0.93	0.024
TTLL12	Q14166	>sp Q14166 MEAERGPB	15.22	Partial IDP	0.2019	No	0.2262	No			3	Client	20	-5.3	No	No	-0.39	-0.025	
PCBP1	Q15365	>sp Q15365 MDAGVTE	36.52	IDP	0.3965	No	0.5417	Yes				2	Client	8	-4.9	No	No	-0.11	0.007
HNRNPA0	Q13151	>sp Q13151 MENSQLC	61.97	IDP	0.9818	Yes	0.9959	Yes	+			1	Driver	5	-20.5	No	No	-0.59	0.043
RBMB3	P98179	>sp P98179 MSSEEGKL	65.61	IDP	0.9952	Yes	0.9542	Yes	+			1	Driver	3	-32.4	No	No	-1.09	0.025
HNRNPCL1	O60812	>sp O60812 MASVNTN	67.58	IDP	0.9744	Yes	0.75	Yes	+			3	Driver	4	-34.8	No	No	-0.96	-0.039
RBMB39	Q14498	>sp Q14498 MADDIDIE	51.89	IDP	0.4127	No	0.5286	Yes				1	Client	17	-18.5	No	No	-0.65	0.069
TAF6	P49848	>sp P49848 MAEKKLKG	49.78	IDP	0.9420	Yes	0.9754												

HBA1	P69905	>sp P69905 MVLSPADK	20.42	Partial IDP	0.1494	No	0.0034	No		+ +	0	No	5	1.1	Yes	No	0.05	0.049
SNRPD3	P62318	>sp P62318 MSIGVPVK	41.27	IDP	0.2632	No	0.0185	No		+ +	1	Client	6	-11.4	No	No	-0.43	0.103
ZGPAT	Q8N5A5	>sp Q8N5A5 MDDEESLES	70.62	IDP	0.9725	Yes	0.746	Yes		+ +	6	Driver	12	-24.0	No	No	-0.59	-0.022
GATA2B	Q8WXI9	>sp Q8WXI9 MDRMITED	83.14	IDP	0.9217	Yes	0.8894	Yes		+ +	6	Driver	11	-19.7	No	No	-0.61	0.046
UBE2E1	P51965	>sp P51965 MSDDDSRA	45.08	IDP	0.7531	Yes	0.1172	No		+ +	1	Driver	6	-12.5	No	No	-0.58	0.026
NOP9	Q86U38	>sp Q86U38 MGQGPRSA	32.86	IDP	0.4991	No	0.6737	Yes		+ +	3	Client	24	-4.7	No	No	-0.12	0.013
CDK1	P06493	>sp P06493 MEDYTKIE	12.12	Partial IDP	0.1467	No	0.0056	No		+ +	0	No	11	2.2	Yes	No	-0.28	0.020
CHTOP	Q9Y3Y2	>sp Q9Y3Y2 MAAQSAPI	100	IDP	0.9270	Yes	0.9882	Yes		+ +	3	Driver	2	-27.8	No	No	-0.71	0.159
SMNDC1	O75940	>sp O75940 MSEDLAKC	76.89	IDP	0.8486	Yes	0.1218	No		+ +	3	Driver	3	-28.6	No	No	-0.92	0.004
SNRPGP15	A8MWD9	>sp A8MWD9 MSKAHPPE	34.21	IDP	0.1100	No	0.0127	No		+ +	0	No	4	-5.5	No	No	-0.22	0.039
SNRPG	P62308	>sp P62308 MSKAHPPE	32.89	IDP	0.1086	No	0.0068	No		+ +	0	No	3	-1.1	No	No	-0.07	0.039
RBM8A	Q9V5S9	>sp Q9V5S9 MADVLDLH	61.49	IDP	0.9709	Yes	0.0354	No		+ +	3	Driver	3	-38.0	No	No	-1.15	-0.020
GSTM3	P21266	>sp P21266 MSCESSMV	14.22	Partial IDP	0.1175	No	0.0047	No		+ +	0	No	9	-2.8	No	No	-0.41	-0.016
RPS17	P08708	>sp P08708 MGRVRKT	31.85	IDP	0.1601	No	0.009	No		+ +	1	Client	5	-11.4	No	No	-0.58	0.074
RPS17L	POCW22	>sp POCW22 MGRVRKT	31.85	IDP	0.1601	No	0.009	No		+ +	1	Client	5	-11.4	No	No	-0.58	0.074
HNRNPUL	Q9BUJ2	>sp Q9BUJ2 MDVRRLK	63.79	IDP	0.9979	Yes	0.9796	Yes		+ +	4	Driver	14	-29.5	No	No	-1.08	0.003
RPS25	P62851	>sp P62851 MPPKDDK	46.4	IDP	0.8546	Yes	0.0689	No		+ +	2	Driver	5	-26.9	No	No	-0.86	0.156
C1QBP	Q07021	>sp Q07021 MLPLLCRV	62.41	IDP	0.3114	No	0.332	No		+ +	1	Client	12	-14.6	No	No	-0.46	-0.055
EWSR1	Q01844	>sp Q01844 MASTDYST	93.75	IDP	0.9999	Yes	0.9991	Yes		+ +	5	Driver	2	-34.4	No	No	-1.12	0.027
SNRPD2	P62316	>sp P62316 MSLLNPK	54.24	IDP	0.5876	No	0.177	No		+ +	2	Client	4	-15.7	No	No	-0.71	0.097
IGLC1	POCG04	>sp POCG04 GQPKANPT	71.7	IDP	0.2484	No	0.0161	No		+ +	0	No	4	-12.1	No	No	-0.52	0.019
IGLC2	POCG05	>sp POCG05 GQPKAAAPS	74.53	IDP	0.3838	No	0.0419	No		+ +	1	Client	4	-9.8	No	No	-0.44	0.009
IGLC3	POCG06	>sp POCG06 GQPKAAAPS	76.42	IDP	0.5929	No	0.0774	No		+ +	1	Client	4	-11.0	No	No	-0.46	0.009
IGLC6	POCF74	>sp POCF74 GQPKAAAPS	73.58	IDP	0.2728	No	0.0324	No		+ +	0	No	4	-11.1	No	No	-0.47	0.009
IGLC7	A0M8Q6	>sp A0M8Q6 GQPKAAAPS	74.53	IDP	0.3412	No	0.0185	No		+ +	0	No	5	-11.6	No	No	-0.46	0.028
IGLL5	B9A064	>sp B9A064 MRPKTGQ	69.16	IDP	0.7742	Yes	0.1969	No		+ +	2	Driver	7	-7.7	No	No	-0.40	0.035
POLA1	P09884	>sp P09884 MAPVHGD	37.69	IDP	0.6683	Yes	0.1415	No		+ +	7	Driver	42	-8.7	No	No	-0.46	-0.011
RBBP4	Q09028	>sp Q09028 MADKEAA	27.29	Partial IDP	0.3177	No	0.1343	No		+ +	3	Client	16	-15.6	No	No	-0.56	-0.064
IGKC	P01834	>sp P01834 RTVAAPSV	61.68	IDP	0.2708	No	0.0898	No		+ +	1	Client	4	-12.4	No	No	-0.54	0.000
ZFR	Q96KR1	>sp Q96KR1 MIPICPVVS	74.49	IDP	0.9935	Yes	0.9789	Yes		+ +	11	Driver	24	-16.2	No	No	-0.56	0.037
HSPA8	P11142	>sp P11142 MSKGPAVQ	38.7	IDP	0.3211	No	0.5824	Yes		+ +	2	Client	17	-13.8	No	No	-0.46	-0.015
SRSF1	Q07955	>sp Q07955 MSGGGVIR	63.71	IDP	0.9496	Yes	0.9711	Yes		+ +	3	Driver	5	-31.8	No	No	-1.15	0.087
RPLP1	P05386	>sp P05386 MASVSELAK	50	IDP	0.9423	Yes	0.1345	No		+ +	1	Driver	5	-8.0	No	No	0.11	-0.092
DUSP3	P51452	>sp P51452 MSGSFELNS	17.84	Partial IDP	0.1135	No	0.0024	No		+ +	1	Client	6	-7.8	No	No	-0.31	0.014
EEF1D	P29692	>sp P29692 MATNFLAH	62.99	IDP	0.9193	Yes	0.4908	No		+ +	3	Driver	6	-22.9	No	No	-0.59	-0.044
RPL31	P62899	>sp P62899 MAPAKKKG	30.4	IDP	0.2933	No	0.0207	No		+ +	1	Client	2	-16.7	No	No	-0.81	0.144
HMX3	A6NHNT5	>sp A6NHNT5 MPEPGPD	69.75	IDP	0.9069	Yes	0.6825	Yes		+ +	2	Driver	8	-18.8	No	No	-0.49	0.017
RPL11	P62913	>sp P62913 MAQDQGE	16.29	Partial IDP	0.1553	No	0.0102	No		+ +	1	Client	5	-11.4	No	No	-0.53	0.070
SF3A1	Q15459	>sp Q15459 MPAGPVQ	77.68	IDP	0.9627	Yes	0.9809	Yes		+ +	6	Driver	14	-18.8	No	No	-0.74	-0.026
SNRNP40	Q96DI7	>sp Q96DI7 MIEQQKRK	24.65	Partial IDP	0.2387	No	0.1868	No		+ +	2	Client	9	-8.1	No	No	-0.35	0.022
ESRP2	Q9H6T0	>sp Q9H6T0 MTPPPPPP	32.6	IDP	0.3368	No	0.849	Yes		+ +	2	Client	25	-0.9	No	No	-0.09	0.002
RPS3	P23396	>sp P23396 MAVQISKK	25.1	Partial IDP	0.2066	No	0.0055	No		+ +	1	Client	8	-0.1	No	No	-0.15	0.056
TRIM33	Q9UPN9	>sp Q9UPN9 MAENKGQ	65.04	IDP	0.9995	Yes	0.8915	Yes		+ +	9	Driver	21	-18.3	No	No	-0.54	0.003
FIP1L1	Q6UN15	>sp Q6UN15 MSAGEVER	77.1	IDP	0.9978	Yes	0.9323	Yes		+ +	4	Driver	6	-39.4	No	No	-1.25	-0.021
SRPRB	Q9Y5M8	>sp Q9Y5M8 MASADSRP	29.52	Partial IDP	0.3636	No	0.0331	No		+ +	2	Client	9	-0.1	No	No	-0.02	0.024
CTDP1	Q9Y5B0	>sp Q9Y5B0 MEVPAAG	67.12	IDP	0.9981	Yes	0.972	Yes		+ +	5	Driver	16	-28.0	No	No	-0.74	-0.030
THOC6	Q86W42	>sp Q86W42 MERAAPLA	17.3	Partial IDP	0.1452	No	0.0652	No		+ +	0	No	12	-1.6	No	No	-0.16	0.015
DEK	P35659	>sp P35659 MSASAPAA	80.53	IDP	0.													

**Table S3.** Characteristics of the proteome of the nSBs. DPR are droplet-promoting regions, nHS are number of aggregation hot spots, Na4vSS are Normalized a4v Sequence Sum for 100 residues used for aggregation prediction.

Gene Name	UniProtID	Fasta	Sequence	PER(VSL2B)	Disorder Classification	FuzDrop Score	FuzDrop LLPS	PSPredictor Score	PSPredictor LLPS	Fuz&PSP LLPS	Fuz&PSP No LLPS	Number of DPRs	Role in MLO	nHS	Na4vSS	Aggregation Predicted	Aggregation DB	Hydrophobicity	Charge
HSF1	Q00613	>sp Q00613 MDLPVPGP	72.97	IDP	0.9884	Yes	0.9457	Yes	+			5	Driver	15	-0.112	No	No	-0.39	-0.029
HSF2	Q03933	>sp Q03933 MKQSSNVP	67.35	IDP	0.8074	Yes	0.8322	Yes	+			5	Driver	11	-0.16	No	No	-0.58	-0.054
SLTM	Q9NW9H	>sp Q9NW9H MAAATGAV	92.65	IDP	0.9998	Yes	0.9665	Yes	+			6	Driver	6	-50	No	No	-1.45	0.015
NCOA5	Q9HCD5	>sp Q9HCD5 MNTAPSRRPS	81.52	IDP	0.9973	Yes	0.7903	Yes	+			5	Driver	8	-35.4	No	No	-1.10	0.032
DHX9	Q08211	>sp Q08211 MGDVKNFL	32.28	IDP	0.7863	Yes	0.1976	No				5	Driver	39	-5.4	No	No	-0.34	0.004
ZNF638	Q14966	>sp Q14966 MSRPRFNPR	81.24	IDP	0.9954	Yes	0.9458	Yes	+			20	Driver	34	-24.3	No	No	-0.81	-0.003
HNRNPM	P52272	>sp P52272 MAAGVEAA	76.30	IDP	0.9997	Yes	0.8675	Yes	+			8	Driver	18	-14	No	No	-0.34	0.015
SRSF1	Q07955	>sp Q07955 MSGGGVIRQ	63.71	IDP	0.9496	Yes	0.9711	Yes	+			3	Driver	5	-31.8	No	No	-1.15	0.087
TRA2B	P62995	>sp P62995 MSDSGEQQN	79.17	IDP	0.9948	Yes	0.9957	Yes	+			3	Driver	4	-42.9	No	No	-1.62	0.158
SAFB2	Q14151	>sp Q14151 MAETLPGSG	93.18	IDP	0.9996	Yes	0.7464	Yes	+			7	Driver	6	-45.5	No	No	-1.29	-0.009
RBMX	P38159	>sp P38159 MVEADRPNG	86.96	IDP	0.9984	Yes	0.9984	Yes	+			1	Driver	3	-40.5	No	No	-1.37	0.063
RBMXL1	Q96E39	>sp Q96E39 MVEADRPNG	86.92	IDP	0.9984	Yes	0.998	Yes	+			2	Driver	3	-38.2	No	No	-1.29	0.056
SRSF9	Q13242	>sp Q13242 MSGWADERP	52.04	IDP	0.8195	Yes	0.0665	No				3	Driver	4	-25.3	No	No	-1.07	0.023
DDX39A	O00148	>sp O00148 MAEQDVEN	18.27	Partial IDP	0.1354	No	0.0079	No		+		1	Client	16	-4	No	No	-0.30	-0.016
DDX39B	Q13838	>sp Q13838 MAENDVDN	20.79	Partial IDP	0.115	No	0.0115	No		+		2	Client	14	-4.4	No	No	-0.27	-0.018
SRSF7	Q16629	>sp Q16629 MSRYGRYGC	81.09	IDP	0.9857	Yes	0.9955	Yes	+			2	Driver	5	-36.7	No	No	-1.40	0.200
THRAP3	Q9Y2W1	>sp Q9Y2W1 MSKTNKSKS	97.91	IDP	0.9996	Yes	0.998	Yes	+			2	Driver	4	-48.8	No	No	-1.63	0.097
POLDIP3	Q9BY77	>sp Q9BY77 MADISLDEL	76.48	IDP	0.3363	No	0.8247	Yes				2	Client	10	-15.1	No	No	-0.47	0.071
RBM15	Q96T37	>sp Q96T37 MRTAGRDP	68.78	IDP	0.9996	Yes	0.891	Yes	+			6	Driver	14	-27.8	No	No	-0.87	0.058
BCLAF1	Q9NYF8	>sp Q9NYF8 MGRSNSRSR	96.96	IDP	0.9995	Yes	0.9979	Yes	+			11	Driver	4	-47.3	No	No	-1.63	0.088
SRSF2	Q01130	>sp Q01130 MSYGRPPP	83.71	IDP	0.9964	Yes	0.9881	Yes	+			1	Driver	2	-45.8	No	No	-1.62	0.206
ALYREF	Q86V81	>sp Q86V81 MADKMDM	66.54	IDP	0.996	Yes	0.986	Yes	+			2	Driver	3	-35	No	No	-0.82	0.080
HNRNPA2	P22626	>sp P22626 MEKLTETVP	69.97	IDP	0.9808	Yes	0.9977	Yes	+			2	Driver	4	-26.5	No	No	-0.93	0.024
SRSF3	P84103	>sp P84103 MRHDSCPLD	74.39	IDP	0.9945	Yes	0.8645	Yes	+			1	Driver	4	-43.2	No	No	-1.52	0.165
RBMXL2	O75526	>sp O75526 MVEADRPNG	86.73	IDP	0.9981	Yes	0.9385	Yes	+			1	Driver	3	-38.5	No	No	-1.25	0.080
RBMXL3	Q8N7X1	>sp Q8N7X1 MMEA DRPE	93.35	IDP	0.9997	Yes	0.996	Yes	+			3	Driver	7	-42.1	No	No	-1.37	0.047
LUC7L3	O95232	>sp O95232 MISAAQLLD	77.31	IDP	0.9957	Yes	0.6483	Yes	+			2	Driver	4	-55.1	No	No	-1.70	0.074
YTHDC1	Q96MU7	>sp Q96MU7 MAADSREEK	81.71	IDP	0.9987	Yes	0.9083	Yes	+			3	Driver	8	-47.8	No	No	-1.52	-0.008
SRSF5	Q13243	>sp Q13243 MSGCRVFIG	69.49	IDP	0.9488	Yes	0.648	Yes	+			2	Driver	3	-38.3	No	No	-1.31	0.153
RBM12B	Q8IXT5	>sp Q8IXT5 MAVVIRLLG	64.84	IDP	0.9936	Yes	0.7464	Yes	+			6	Driver	23	-28.8	No	No	-1.08	0.006
DDX5	P17844	>sp P17844 MSGYSSDR	41.04	IDP	0.5261	No	0.4192	No		+		3	Client	18	-16	No	No	-0.64	0.029
C7orf55-LU	Q96HJ9	>sp Q96HJ9 MAALGSPSH	35.40	IDP	0.6512	Yes	0.0023	No				1	Driver	3	-10.8	No	No	-0.47	0.115
SRSF4	Q08170	>sp Q08170 MPPRVYIGRL	77.33	IDP	0.9972	Yes	0.9944	Yes	+			2	Driver	5	-55.3	No	No	-1.84	0.197
SRSF6	Q13247	>sp Q13247 MPPRVYIGRL	70.35	IDP	0.9978	Yes	0.998	Yes	+			2	Driver	4	-43.4	No	No	-1.55	0.170
PPHLN1	Q8NEY8	>sp Q8NEY8 MWSSEG RYE	85.59	IDP	0.9978	Yes	0.91	Yes	+			4	Driver	4	-36.1	No	No	-1.30	0.032
CPSF7	Q8N684	>sp Q8N684 MSEGVDLID	74.95	IDP	0.9987	Yes	0.9906	Yes	+			5	Driver	12	-22.6	No	No	-0.78	0.020
SPEN	Q96T58	>sp Q96T58 MVRETRHLV	84.12	IDP	1	Yes	0.8979	Yes	+			24	Driver	55	-27.3501	No	No	-0.86	0.011
ZC3H13	Q5T200	>sp Q5T200 MSKIRRKV	94.18	IDP	0.9997	Yes	0.8977	Yes	+			5	Driver	10	-61.8	No	No	-1.86	0.041
SRSF10	Q75494	>sp Q75494 MSRYLRPPN	73.28	IDP	0.9947	Yes	0.979	Yes	+			1	Driver	4	-45.3	No	No	-1.76	0.179
PIPIA	P62937	>sp P62937 MVNP TVFFF	20.61	Partial IDP	0.2089	No	0.0057	No		+		0	No	7	-6.7	No	No	-0.32	0.018
U2AF2	P26368	>sp P26368 MSDFDEFER	42.74	IDP	0.3891	No	0.0708	No		+		2	Client	14	-15	No	No	-0.59	0.031
DDX17	Q92841	>sp Q92841 MPTGFVAPI	46.36	IDP	0.8739	Yes	0.856	Yes	+			3	Driver	21	-16.8	No	No	-0.61	0.016
HNRNPA1	P09651	>sp P09651 MSKSESPKE	70.16	IDP	0.9998	Yes	0.9977	Yes	+			1	Driver	7	-27.8	No	No	-0.89	0.030
HNRNPA1	Q32P51	>sp Q32P51 MSKSASPKE	63.13	IDP	0.9711	Yes	0.9958	Yes	+			1	Driver	7	-25.6	No	No	-0.84	0.031
HNR																			

DDX21	Q9NR30	>sp Q9NR30 MPGKLRSDD	45.34	IDP	0.4512	No	0.9921	Yes		4	Client	22	-21.8	No	No	-0.76	0.042
DDX3X	O00571	>sp O00571 MSHVALEN	40.94	IDP	0.8355	Yes	0.998	Yes	+	2	Driver	18	-17.3	No	No	-0.63	0.008
DDX3Y	O15523	>sp O15523 MSHVVVKN	41.21	IDP	0.8518	Yes	0.9453	Yes	+	3	Driver	18	-17.2	No	No	-0.64	0.011
FAM208A	Q9UK61	>sp Q9UK61 MATAVETEA	49.94	IDP	0.9916	Yes	0.7529	Yes	+	16	Driver	44	-15.1	No	No	-0.64	-0.015
HSPA8	P11142	>sp P11142 MSKGPAVG	38.70	IDP	0.3211	No	0.5824	Yes		2	Client	17	-13.8	No	No	-0.46	-0.015
LMNA	P02545	>sp P02545 METPSQRRA	87.65	IDP	0.9517	Yes	0.9161	Yes	+	6	Driver	8	-31.5	No	No	-0.86	0.005
ZC3H11A	O75152	>sp O75152 MPNQGEDC	87.53	IDP	0.9982	Yes	0.8823	Yes	+	10	Driver	16	-23.5	No	No	-0.75	0.020
PRPF38B	Q5VTL8	>sp Q5VTL8 MANNSPALT	69.60	IDP	0.9991	Yes	0.9106	Yes	+	3	Driver	6	-51.9	No	No	-1.70	0.127
RBM23	Q86U06	>sp Q86U06 MASDDFDIV	65.83	IDP	0.681	Yes	0.0835	No		1	Driver	14	-23.7	No	No	-0.72	0.051
RBMX2	Q9Y388	>sp Q9Y388 MNPLTKVKL	67.70	IDP	0.9958	Yes	0.9853	Yes	+	1	Driver	5	-45.1	No	No	-1.53	0.096
RPS27A	P62979	>sp P62979 MQIFVKTLT	57.05	IDP	0.1763	No	0.0069	No		1	Client	7	-19.2	No	No	-0.84	0.115
UBA52	P62987	>sp P62987 MQIFVKTLT	39.84	IDP	0.1393	No	0.0758	No		1	Client	4	-17.5	No	No	-0.70	0.121
UBB	POCG47	>sp POCG47 MQIFVKTLT	30.57	IDP	0.1455	No	0.0047	No		0	No	4	-12.2	No	No	-0.48	0.007
UBC	POCG48	>sp POCG48 MQIFVKTLT	28.76	Partial IDP	0.1563	No	0.0106	No		0	No	10	-12.3	No	No	-0.48	0.007
SRP14	P37108	>sp P37108 MVILLESEQF	74.26	IDP	0.5059	No	0.1114	No		1	Client	5	-16.8	No	No	-0.44	0.110
SUB1	P53999	>sp P53999 MPKSKELVS	86.61	IDP	0.9209	Yes	0.8756	Yes	+	1	Driver	2	-32.1	No	No	-1.09	0.055
DDX1	Q92499	>sp Q92499 MAAFSEMG	16.08	Partial IDP	0.2648	No	0.802	Yes		2	Client	23	-8.8	No	No	-0.40	0.013
FUBP1	Q96AE4	>sp Q96AE4 MADYSTVPP	84.47	IDP	0.9962	Yes	0.9922	Yes	+	11	Driver	10	-24.9	No	No	-0.74	0.009
HIST1H1C	P16403	>sp P16403 MSETAPAAP	97.65	IDP	0.9966	Yes	0.9886	Yes	+	2	Driver	3	-25.6	No	No	-0.68	0.258
HNRNPA0	Q13151	>sp Q13151 MENSQLCKI	61.97	IDP	0.9818	Yes	0.9959	Yes	+	1	Driver	5	-20.5	No	No	-0.59	0.043
HSPA1A	P31948	>sp P31948 MEQVNELK	55.80	IDP	0.4991	No	0.0334	No		5	Client	5	-26.6	No	No	-0.94	0.003
HSPA1B	V9GZ37	>tr V9GZ37 MAKAAAIGI	37.18	IDP	0.4139	No	0.4516	No		2	Client	13	-14.6	No	No	-0.41	-0.017
HSPA1L	P34931	>sp P34931 MATAKGIAI	28.39	Partial IDP	0.2245	No	0.1552	No		1	Client	21	-10.1	No	No	-0.34	-0.006
HSPA2	P54652	>sp P54652 MSARGPAIG	38.03	IDP	0.3147	No	0.4757	No		1	Client	18	-15.1	No	No	-0.49	-0.010
HSPA5	P11021	>sp P11021 MKLSLVAAN	41.28	IDP	0.4337	No	0.3872	No		3	Client	18	-14.8	No	No	-0.49	-0.029
KIAA1429	Q69YN4	>sp Q69YN4 MAVDSAME	41.78	IDP	0.9241	Yes	0.7061	Yes	+	9	Driver	57	-10.4	No	No	-0.40	-0.041
PPIAL4C	AOA0B4	>tr AOA0B4 MTTPVHFSS	10.20	Partial IDP	0.1476	No	0.0681	No		0	No	4	30.8	Yes	No	0.89	0.010
PPIG	Q13427	>sp Q13427 MGIKVQRPP	83.95	IDP	0.9974	Yes	0.9977	Yes	+	6	Driver	7	-57.3	No	No	-1.88	0.135
PRPF4B	Q13523	>sp Q13523 MAAAETQS	66.63	IDP	0.9961	Yes	0.9923	Yes	+	3	Driver	15	-37.8	No	No	-1.29	0.116
PTBP1	P26599	>sp P26599 MDGIVPDIA	43.27	IDP	0.5386	No	0.9415	Yes		4	Client	25	-2.7	No	No	-0.12	0.030
PTBP3	O95758	>sp O95758 MDGVVTDL	39.31	IDP	0.8663	Yes	0.7874	Yes	+	5	Driver	22	0.1	Yes	No	-0.10	0.030
PURB	Q96QR8	>sp Q96QR8 MADGDSGS	62.18	IDP	0.9803	Yes	0.5841	Yes	+	5	Driver	6	-24.6	No	No	-0.67	-0.014
RBM14	Q96PK6	>sp Q96PK6 MKitVGNV	81.32	IDP	0.9999	Yes	0.9903	Yes	+	6	Driver	7	-15.4	No	No	-0.45	0.043
RBM15B	Q8NDT2	>sp Q8NDT2 MKRQSERD	61.24	IDP	0.9935	Yes	0.7781	Yes	+	5	Driver	16	-27.8	No	No	-0.85	0.058
RNF20	Q5VTR2	>sp Q5VTR2 MSGIGNKRA	77.85	IDP	0.7779	Yes	0.5943	Yes	+	7	Driver	10	-34.3	No	No	-1.04	-0.011
SNRNP27	Q8WVK2	>sp Q8WVK2 MGRSRSRS	83.87	IDP	0.9856	Yes	0.8507	Yes	+	1	Driver	3	-59.2	No	No	-2.00	0.194
TOP1	P11387	>sp P11387 MSGDHLHN	60.52	IDP	0.3929	No	0.2234	No		3	Client	10	-34.5	No	No	-1.29	0.060