## Berg, et al Supplementary Information

Supplemental Figure Legends

#### Figure S1. Anti-EJC monoclonal antibodies immunoprecipitate spliced mRNA

Denaturing PAGE analysis for splicing of <sup>32</sup>P-labeled Ad2ΔIVS pre-mRNA (Input) followed by immunoprecipitations with Control (SP2/0), Y14 (4C4), eIF4AIII (3F1) and hnRNP C1/C2 (4F4) mouse monoclonal antibodies. Identities of splicing intermediates are depicted to the right of the gel.

#### Figure S2. Optimization of the EJIPT assay parameters

(A) Splicing reactions with ATP (0.5 mM), without ATP, or with 5 mM AMP-PNP were immunoprecipitated with an anti-EJC antibody. SP2/0 is a control anti-mouse antibody substituted for the anti-EJC antibody. Ad2 mRNA produced without splicing was immunoprecipitated with an anti-EJC antibody. Parameters for the EJIPT assay were examined using increasing amounts of splicing extract (B) and increasing amounts of biotin-labeled Ad2ΔIVS pre-mRNA (C) in a 10 μl reaction volume.

# Figure S3. NSC663284 and BN82685 are more selective splicing inhibitors compared to NSC95397

*In vitro* snRNP assembly was performed in the presence of increasing amounts of compounds. Chemiluminescence signals are the averages of quadruplicate measurements. The red dashed line indicates the average for DMSO controls  $(6.0 \pm 1.1 \times 10^4 \text{ RLU})$ .

### Supplemental Table S1. Association of individual proteins with drug-treated spliceosomes.

Peptide counts for mass spectrometry of DMSO and BN82685 streptavidin pull downs are shown for three independent experiments, along with a technical repeat of experiment 3. The ratio of BN82685 to DMSO is shown in dark grey columns and the average and standard deviation of the four data points is listed on the right in light grey columns. Data points not used in the calculation are in grey type with italics.

Top 200 Identified Proteins	A ccession Number	M olecular Weight	Exp 1 DM SO	Exp 1 BN82685	ratio	Exp 2 DM SO	Exp 2	ratio	Exp 3A DM SO	Exp 3A BN82685	ratio			ratio	Average	Sed
soform 1 of A cetyl-CoA carboxylase 1	IPI00011569	266 kDa	51	73	1.43	11	2	0.18	376	401	1.07	910	984	1.08	0.94	
Splicing factor 3B subunit 1	IP100011309	146 kDa	188	175	0.93	305	231	0.76	137	114	0.83	307	204	0.66	0.80	
Dihydrolipoyllysine-residue acetyltransferase component	IP100021338	69 kDa	109	110	1.01	251	260	1.04	120	135	1.13	312	337	1.08	1.06	
Pre-mRNA-processing-splicing factor 8	IP100027938	274 kDa	170	143	0.84	232	106	0.46	174	188	1.08	200	213	1.07	0.86	
Isoform 1 of Splicing factor 3B subunit 3	IP100300371	136 kDa	104	105	1.01	199	169	0.40	118	96	0.81	246	196	0.80	0.87	
116 kDa U5 small nuclear ribonucleoprotein component	IP100300371	109 kDa	132	99	0.75	203	122	0.60	94	98	1.04	196	218	1.11	0.88	
	IP100003319	53 kDa	80	61	0.75	136	168	1.24	54	69	1.28	128	147	1.15	1.11	
Lipoamide acyltransferase component		89 kDa	106	77	0.76	163	108	0.66	64	60	0.94	130	107	0.82	0.79	
Splicing factor 3A subunit 1	IPI00017451															
Splicing factor 3B subunit 2	IPI00221106	100 kDa	82	70	0.85	95	98	1.03	78	60	0.77	162	108	0.67	0.83	
soform 1 of U5 small nuclear ribonucleoprotein 200 kDa helicase	IPI00420014	245 kDa	78	92	1.18	77	39	0.51	94	83	0.88	137	190	1.39	0.99	
J2 small nuclear ribonucleoprotein A	IP100297477	28 kDa	58	48	0.83	92	82	0.89	48	40	0.83	149	129	0.87	0.85	
soform 2 of U5 small nuclear ribonucleoprotein 200 kDa helicase	IPI00168235	71 kDa	80	56	0.70	73	38	0.52	55	60	1.09	74	91	1.23	0.89	
Putative uncharacterized protein A L B	IP100022434	72 kDa	40	34	0.85	62	76	1.23	43	46	1.07	109	107	0.98	1.03	
soform 2 of Splicing factor 3B subunit 3	IPI00179138	30 kDa	46	31	0.67	69	65	0.94	28	30	1.07	85	85	1.00	0.92	
soform 1 of Polyadeny late-binding protein 1	IP100008524	71 kDa	48	45	0.94	25	40	1.60	46	62	1.35	124	102	0.82	1.18	
Cell division cycle 5-like protein	IPI00465294	92 kDa	51	58	1.14	61	34	0.56	40	50	1.25	70	90	1.29	1.06	
ntron-binding protein aquarius	IP100297572	171 kDa	53	51	0.96	42	23	0.55	54	52	0.96	40	69	1.73	1.05	
soform Long of Splicing factor, proline- and glutamine-rich	IPI00010740	76 kDa	5	8	1.60	0	1	n/a	74	64	0.86	237	80	0.34	0.93	
Ceratin, type II cytoskeletal 1	IP100220327	66 kDa	52	95	1.83	39	57	1.46	37	42	1.14	49	75	1.53	1.49	
Splicing factor 3A subunit 3	IP100029764	59 kDa	47	45	0.96	80	50	0.63	39	22	0.56	73	47	0.64	0.70	
soform 1 of Heterogeneous nuclear ribonucleoprotein M	IP100023704	78 kDa	53	49	0.92	30	34	1.13	77	53	0.50	72		1.01	0.94	
	IP100171903	55 kDa	35	43	1.23	80	53	0.66	32	42	1.31	71	96	1.35	1.14	
Pre-mRNA-processing factor 19																
Py ruvate carboxy lase, mitochondrial	IP100299402	130 kDa	18	27	1.50	5	3	0.60	78	75	0.96	109	89	0.82	0.97	
Pre-mRNA-splicing factor SYF1	IPI00163084	100 kDa	62	63	1.02	50	23	0.46	39	63	1.62	45	59	1.31	1.10	
y ruvate dehy drogenase protein X component, mitochondrial	IP100298423	54 kDa	51	31	0.61	43	72	1.67	34	58	1.71	61	52	0.85	1.21	
soform 1 of Crooked neck-like protein 1	IPI00177437	100 kDa	55	39	0.71	48	24	0.50	30	49	1.63	41	63	1.54	1.09	
Ceratin, type I cytoskeletal 10	IP100009865	59 kDa	66	74	1.12	23	40	1.74	44	39	0.89	22	26	1.18	1.23	
nsulin-like growth factor 2 mRNA-binding protein 1	IP100008557	63 kDa	33	43	1.30	22	28	1.27	44	32	0.73	81	60	0.74	1.01	
Propiony I-CoA carboxy lase beta chain, mitochondrial	IP100007247	58 kDa	25	13	0.52	17	10	0.59	48	47	0.98	95	89	0.94	0.76	
Non-POU domain-containing octamer-binding protein	IP100304596	54 kDa	0	1	n/a	0	0	0.00	66	50	0.76	147	48	0.33	0.36	
propiony I-CoA carboxy lase alpha chain	IP100552419	75 kDa	22	13	0.59	11	9	0.82	47	58	1.23	71	80	1.13	0.94	
Methylcrotonoyl-CoA carboxylase subunit alpha, mitochondrial	IP100024580	80 kDa	4	2	0.50	0	0	0.00	69	78	1.13	68	82	1.21	0.71	
SM -B' of Small nuclear ribonucleoprotein-associated proteins B and B'	IP100027285	25 kDa	28	5	0.18	19	23	1.21	31	23	0.74		51	0.63	0.69	
	IP100027203	96 kDa	21	12	0.10	8	2	0.25	40	36	0.90	80	32	0.40	0.53	
Heterogeneous nuclear ribonucleoprotein U-like protein 1				49												
Keratin, type II cytoskeletal 2 epidermal	IP100021304	66 kDa	36	-	1.36	24	38	1.58	21	16	0.76	31	33	1.06	1.19	
ukary otic initiation factor 4A-III	IP100009328	47 kDa	22	4	0.18	20	1	0.05	21	42	2,00	17	99	5.82	2.01	
NW domain-containing protein 1	IPI00013830	61 kDa	25	10	0.40	22	18	0.82	25	30	1.20	30		1.10	0.88	
soform 1 of Terminal uridy ly Itransferase 4	IPI00289861	185 kDa	8	6	0.75	28	21	0.75	5	9	1.80	31	76	2.45	1.44	
:DNA FLJ56825, highly similar to WD repeat protein 57	IP100006723	45 kDa	17	12	0.71	11	5	0.45	17	30	1.76	56	56	1.00	0.98	
soform 1 of Pleiotropic regulator 1	IP100002624	57 kDa	40	16	0.40	14	8	0.57	29	34	1.17	22	32	1.45	0.90	
Pre-mRNA-processing factor 17	IP100026307	66 kDa	28	22	0.79	10	7	0.70	28	33	1.18	22	30	1.36	1.01	
ubulin alpha-4A chain	IP100007750	50 kDa	7	6	0.86	4	9	2.25	19	27	1.42	44	58	1.32	1.46	
ctin, cytoplasmic 1	IPI00021439	42 kDa	17	4	0.24	19	5	0.26	7	20	2.86	21	32	1.52	1.22	
M atrin-3	IP100017297	95 kDa	4	15	3.75	0	0	0.00	42	35	0.83	52	28	0.54	1.28	
Tubulin beta-2C chain	IP100007752	50 kDa	5	4	0.80	1	0	0.00	33	53	1.61	25		1.92	1.08	
soform 1 of RNA-binding protein 14	IPI00013174	69 kDa	11	4	0.36	3	0	0.00	27	27	1.00	51	25	0.49	0.46	
Heterogeneous nuclear ribonucleoprotein C-like 1	IP100013174	32 kDa	17	13	0.76	10	10	1.00	26	23	0.88	28	29	1.04	0.92	
The state of the s																
soform C1 of Heterogeneous nuclear ribonucleoproteins C1/C2	IP100216592	32 kDa	13	5	0.38	10	11	1.10	20	22	1.10	40		0.48	0.76	
by ruvate dehy drogenase E1 alpha 1 isoform 2 precursor	IPI00306301	48 kDa	13	17	1.31	26	32	1.23	8	9	1.13	29	22	0.76	1.11	
Dihydrolipoy I dehydrogenase, mitochondrial	IP100015911	54 kDa	27	17	0.63	14	20	1.43	12	14	1.17	21	28	1.33	1.14	
soform 1 of Pre-mRNA-splicing factor ISY1 homolog	IP100063673	35 kDa	17	5	0.29	11	10	0.91	15	29	1.93	25	29	1.16	1.07	
Ceratin, type I cytoskeletal 9	IP100019359	62 kDa	18	46	2.56	14	20	1.43	8	12	1.50	2	18	9.00	3.62	
tibonuclease inhibitor	IP100550069	50 kDa	24	15	0.63	14	16	1.14	12	12	1.00	29	11	0.38	0.79	
PHD finger-like domain-containing protein 5A	IPI00005511	12 kDa	16	12	0.75	34	25	0.74	9	10	1.11	11	9	0.82	0.85	
y ruvate dehy drogenase E1 component subunit beta, mitochondrial	IP100003925	39 kDa	23	13	0.57	12	16	1.33	9	16	1.78	8	7	0.88	1.14	
TUBA1C protein	IP100166768	37 kDa	6	11	1.83	13	12	0.92	6	13	2.17	33	35	1.06	1.50	
soform 1 of Pre-mRNA-splicing factor RBM 22	IPI00019046	47 kDa	11	6	0.55	6	4	0.67	12	22	1.83	6		3.33	1.59	
VD40 repeat-containing protein SM U1	IP100305833	58 kDa	25	8	0.32	5	3	0.60	11	21	1.91	19		1.00	0.96	
eta-actin-like protein 2	IP100303833	42 kDa	14	4	0.32	20	9	0.60	6	15	2.50	20		1.10		
rutative uncharacterized protein FUBP3	IP100063245	35 kDa	14	9	0.64	5	13	2.60	20	18	0.90	18		0.39	1.13	
imentin	IPI00418471	54 kDa	18	13	0.72	3	6	2.00	15	21	1.40			2.38	1.62	
Actin, aortic smooth muscle	IP100008603	42 kDa	5	3	0.60	19	8	0.42	6	15	2.50	19		1.58	1.28	
DNA FLJ55034	IPI00384122	40 kDa	15	18	1.20	24	25	1.04	7	1	0.14	10		0.60	0.75	
12 small nuclear ribonucleoprotein B"	IP100029267	25 kDa	12	9	0.75	1	7	7,00	14	11	0.79	27	28	1.04	0.86	
soform A of Peptidyl-prolyl cis-trans isomerase E	IPI00009316	33 kDa	11	2	0.18	6	4	0.67	19	16	0.84	29	23	0.79	0.62	
H-type splicing regulatory protein	IP100479786	73 kDa	14	23	1.64	1	11	11.00	23	8	0.35	11	1	0.09	0.69	
soform 1 of Paraspeckle component 1	IP100103525	59 kDa	- 1	0		0	0		31	19	0.61	42	13	0.31	0.46	
F3A2 protein (Fragment)	IPI00017341	51 kDa	12	11	0.92	5	13	2.60	12	13	1.08	15		1.27	1.47	
DNA FLJ60076, highly similar to ELAV-like protein 1	IP100301936	39 kDa	9		1.00	3		1.67	22		0.45	28		0.79		

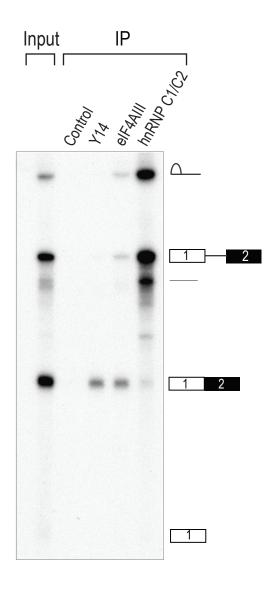
**Table S1.** Peptide counts for top proteins (1-69) detected by mass spectrometry of DMSO and BN82685 streptavidin pull downs.

Top 200 Identified Proteins	A ccession Number	M olecular Weight	Exp 1 DM SO	Exp 1 BN82685	ratio	Exp 2 DM SO B	Exp 2 N82685	ratio	Exp 3A DM SO	Exp 3A BN82685	ratio	Exp 3B DM SO	Exp 3B BN82685	ratio	A verage S	td D
Pre-mRNA branch site protein p14	IP100032827	15 kDa	14	21	1.50	26	17	0.65	6	7	1.17	3	2	0.67	1.00	0.4
Pre-mRNA-splicing factor SPF27	IPI00025178	26 kDa	7	9	1.29	9	7	0.78	10	12	1.20	22	16	0.73	1.00	0.3
Coiled-coil domain-containing protein 12	IP100453463	19 kDa	11	14	1.27	10	7	0.70	14	11	0.79	13	10	0.77	0.88	0.3
Tubulin, beta	IP100433463	48 kDa	3	4	1.33	1	1	1.00	6	12		33	42	1.27	1.40	0.4
U1 small nuclear ribonucleoprotein A	IP100012382	31 kDa	9	6	0.67	12	11	0.92	8	12	1.50	18		0.94	1.01	0.
Pre-mRNA-processing factor 6	IP100305068	107 kDa	19	9	0.47	21	2	0.10	6	9	1.33	11	4	0.36	0.31	0.
Isoform 2 of RNA-binding motif, single stranded-interacting protein 1	IPI00303068	44 kDa	5	3	0.60	10	9	0.10	12	12		25	20	0.30	0.83	0.1
Isoform 1 of RNA-binding protein 4	IP100003704	40 kDa	12	4	0.33	1	0	0.00	7	12		30	23	0.77	0.94	0.
Isoform 1 of Insulin-like growth factor 2 mRNA-binding protein 3	IP100658000	64 kDa	6	7	1.17	0	0		16	18	1.13	22	18	0.82	1.04	0.
Isoform 2 of Double-stranded RNA-specific adenosine deaminase	IP100035057	133 kDa	5	8	1.60	0	0	0.00	24	25	3000	8		1.88	1.51	0.
Peptidyl-prolyl cis-trans isomerase-like 1	IP100023037	18 kDa	19	10	0.53	12	3	0.25	14	4	0.29	7		0.29	0.34	0.
Heterogeneous nuclear ribonucleoprotein H	IPI00007019			14		3	4						8	0.29		
		49 kDa	10		1.40			1.33	13	12		18			1.03	0.
Trypsin-1	IP100011694	27 kDa	10	12	1.20	10	13	1.30	3	3	1.00	3	24	8.00	1.17	0.
Isoform 1 of Polyadenylate-binding protein 4	IP100012726	71 kDa	13	6	0.46	4	3	0.75	16	8	0.50	14		0.57	0.57	0.
60S acidic ribosomal protein P0	IP100008530	34 kDa	3	3	1.00		0	0.00	18	15		28	12	0.43	0.75	0.:
Small nuclear ribonucleoprotein Sm D3	IP100017964	14 kDa	7	5	0.71	14	16	1.14	4	3	0.75	8	9	1.13	0.93	0
Small nuclear ribonucleoprotein Sm D 1	IP100302850	13 kDa	16	16	1,00	7	8	1.14	5	4	0.80	13	11	0.85	0.95	0.
Protein CWC15 homolog	IP100009009	27 kDa	9	4	0.44	2	3	1.50	11	12		12		0.67	0.93	0.4
cDNA FLJ77422	IPI00011268	33 kDa	4	2	0.50	0	1	0.00	10	7	0.70	13	10	0.77	0.66	0.1
Isoform 1 of Protein lin-28 homolog B	IPI00398131	27 kDa	2	1	0.50	1	0	0.00	16	9	0.56	27	12	0.44	0.50	0.0
Small nuclear ribonucleoprotein Sm D2	IP100017963	14 kDa	11	9	0.82	12	8	0.67	7	7	1.00	8	9	1.13	0.90	0.3
Isoform 1 of Heterogeneous nuclear ribonucleoprotein K	IPI00216049	51 kDa	5		2.20	0	3	n/a	15	8	0.53	21	6	0.29	1.01	1.0
Heterogeneous nuclear ribonucleoprotein F	IP100003881	46 kDa	6	13	2.17	0	3	n/a	12	9	0.75	11	10	0.91	1.28	0.7
K eratin, type II cytoskeletal 75	IP100005859	60 kDa	8	8	1.00	5	9	1.80	8	5	0.63	10		1.20	1.16	0.4
DEAD (Asp-Glu-Ala-Asp) box polypeptide 39 transcript variant	IP100062206	35 kDa	1	4	4.00	0	2	n/a	7	3	0.43	25	16	0.64	1.69	2.0
Splicing factor 3B subunit 4	IPI00017339	44 kDa	7	12	1.71	8	10	1.25	4	4	1.00	4		2.25	1.55	0.5
ubiquitin and ribosomal protein S27a precursor	IPI00179330	18 kDa	5	8	1.60	2	9	4.50	5	12	2.40	9	7	0.78	2.32	1.0
Small nuclear ribonucleoprotein E	IP100029266	11 kDa	13	13	1.00	9	7	0.78	6	7	1.17	2	1	0.50	0.86	0.2
605 ribosomal protein L7	IPI00030179	29 kDa	2	1	0.50	0	2	rva	17	13	0.76	8	8	1.00	0.75	0.2
Tubulin beta-1 chain	IPI00006510	50 kDa	2	.0	0.00	0	.0		12	12	1.00	11	17	1.55	1.27	0.3
Propionyl Coenzyme A carboxylase, alpha polypeptide	IPI00552081	17 kDa	0	3	n/a	8	6	0.75	1	4	4.00	17	18	1.06	1.94	1.7
Isoform 1 of Poly pyrimidine tract-binding protein 1	IPI00179964	57 kDa	2	4	2.00	0	0		7	7	1.00	12	1	0.08	0.77	0.9
poly (rC)-binding protein 2 isoform b	IP100012066	38 kDa	0	1	TV/s	0	2	rvis	14	8	0.57	17	.11	0.65	0.61	0.0
Full-length cDNA 5-PRIME end of clone CS0DJ009YL13	IPI00384016	30 kDa	8	9	1.13	4	16	4.00	5	5	1.00	6	1	0.17	1.57	1.6
pre-mRNA-splicing factor ATP-dependent RNA helicase DHX 15	IP100396435	91 kDa	3	1	0.33	0	0	0.00	16	5	0.31	9	6	0.67	0.44	0.2
Splicingfactor 3B subunit 5	IPI00010404	10 kDa	9	12	1.33	17	6	0.35	5	3	0.60	0	0	0.00	0.76	0.5
Protein BUD31 homolog	IPI00013180	17 kDa	13	6	0.46	9	7	0.78	5	2	0.40	1	1	1.00	0.66	0.2
Isoform 1 of Putative helicase M OV-10	IP100444452	114 kDa	3	8	2.67	0	.1	rva	18	13	0.72	2	1	0.50	1.30	1.
Protein Red	IPI00011875	66 kDa	7	3	0.43	5	2	0.40	5	8	1.60	6	8	1.33	0.94	0.6
Complement C4-A	IP100032258	193 kDa	5	2	0.40	2	2	1.00	6	8	1.33	9	14	1.56	1.07	0.5
Isoform 1 of GTP-binding protein 8	IPI00107246	32 kDa	8	3	0.38	5	4	0.80	10	7	0.70	4	4	1.00	0.72	0.2
Isoform 1 of Heterogeneous nuclear ribonucleoprotein D-like	IPI00011274	46 kDa	4	3	0.75	1	2	2.00	6	6	1.00	15	8	0.53	1.07	0.6
Isoform 1 of U4/U6 small nuclear ribonucleoprotein Prp4	IPI00150269	58 kDa	16	2	0.13	10	5	0.50	6	7		1	0		0.31	0.2
Guanine nucleotide-binding protein subunit beta-2-like 1	IP100848226	35 kDa	0	0	0.00	0	-0	0.00	14	12	0.86	14	6	0.43	0.64	0.3
Heterogeneous nuclear ribonucleoprotein L.	IP100027834	64 kDa	0.	1	rva	3	1	0.33	6	5	0.83	3	1	0.33	0.50	0.2
RNA binding motif protein, X -linked-like 1	IPI00061178	42 kDa	3	3	1.00	0	2	n/a	6	2	0.33	19	5	0.26	0.53	0.4
Isoform 1 of DNA-binding protein A	IPI00031801	40 kDa	4	6	1.50	3	2	0.67	5	8	1.60	9	5	0.56	1.08	0.9
Isoform 2 of Spliceosome RNA helicase BAT1	IP100641829	51 kDa	4	6	1.50	1	1	1.00	4	5	1.25	12	10	0.83	1.15	0.2
K eratin, type I cytoskeletal 24	IP100004550	55 kDa	2	6	3.00	8	10	1.25	1	2	2.00	5	8	1.60	1.96	0.7
Isoform 2 of RISC-loading complex subunit TARBP2	IP100472583	37 kDa	3	3	1.00	0	2	n/a	7	0	0.00	15	- 1	0.67	0.83	0.3
Pre-mRNA-splicing factor CWC22 homolog	IPI00177381	105 kDa	6	4	0.67	0	0		8	5	0.63	6		1.17	0.82	0.3
40S ribosomal protein S3	IPI00011253	27 kDa	1	0	0.00	0	0		10	6	0.60	16	7	0.44	0.52	0.1
405 ribosomal protein S4, X isoform	IP100217030	30 kDa	1	1	1.00	0	0	0.00	13	6		13	- 1	0.23	0.56	0.3
A lpha-ketoglutarate dehy drogenase complex dihy drolipoy I succiny Itransferase		49 kDa	6	8	1.33	3	8		1	3		2		1.00	2.00	0.9
Isoform 1 of Insulin-like growth factor 2 mRNA-binding protein 2	IP100033034	66 kDa	5	10	2.00	1	0	0.00	19	14		15		0.73	1.16	0.
Isoform 1 of ELAV-like protein 2	IP100030250	40 kDa	4	2	0.50	0	0		6	5	0.83	12		0.73	0.61	0.
Poly (rC)-binding protein 1	IP100030250	37 kDa		2	0.50	1	2		0	3	0.83	13	14	1.08	1.54	0.
1 - 1 C(1)				1					2	7		13	14			
Protein mago nashi homolog 2	IP100059292	17 kDa	10		0.10	11	0	0.00			3.50		-	0.64	0.05	0.0
ADP/ATP translocase 2	IP100007188	33 kDa	2	1	0.50	0	0	0.00	6	6	1.00	11	7	0.64	0.71	0
Isoform Long of A cetyl-CoA carboxylase 2	IP100013003	277 kDa	0	0	0.00	1	0		12	10		3		1.67	1.25	0.
Iso 1 of M ethylcrotonoyl-CoA carboxylase beta chain, mitochondrial	IP100784044	61 kDa	12	0	0.00	0	0	0.00	7	12		1		1.00	0.90	0.
Probable ATP-dependent RNA helicase DDX 5	IP100017617	69 kDa	0	3	n/a	1	2		8	3	0.38	13		0.23	0.87	0.
Heterogeneous nuclear ribonucleoprotein G	IP100304692	42 kDa	0	1	r/a	0	.0	0.00	7	1	0.14	19	5	0.26	0.20	0.
Pre-mRNA-splicing factor SYF2	IP100022963	29 kDa	6	3	0.50	0	5	rva	4	6	1.50	3	3	1.00	1.00	0.
ATP-dependent RNA helicase A	IP100844578	141 kDa	3	2	0.67	0	.0	0.00	10	5	0.50	9	1	0.11	0.43	0.
RNA-binding motif protein, X-linked-like-2	IP100004450	43 kDa	0	3	n/a	0	0	0.00	5	4	0.80	16	6	0.38	0.59	

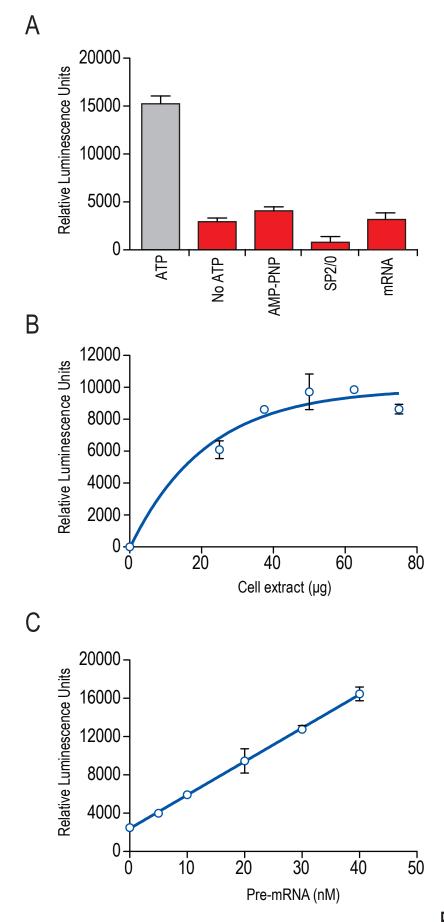
**Table S1.** Peptide counts for top proteins (70-137) detected by mass spectrometry of DMSO and BN82685 streptavidin pull downs.

Top 200 Identified Proteins	A ccession Number	M olecular Weight	Exp 1 Exp 1 DM SO BN 82685	s patie	Exp 2	Exp 2 BN82685	ratio	Exp 3A DM SO	Exp 3A BN82685	ratio		Exp 3B BN82685	ratio	Augram	td D-
Top 200 Identified Proteins  Isoform 1 of Far upstream element-binding protein 3	IP100377261	62 kDa	1 1 1			B N 82685	2.00	DM SO	BN82685	0.57	DM 50		0.33		ota De
Survival of motor neuron-related-splicing factor 30	IP100377201	27 kDa	4		0	3	11/8	7	8	1.14	5		0.20		0.5
Nuclear cap-binding protein subunit 1	IPI00019380	92 kDa	3		0	0		11	10		2		0.50		0.3
K eratin, type II cytoskeletal 5	IP100009867	62 kDa	8 7		1		2.00	2	2	1.00	2		1.00		0.5
Isoform 2 of Insulin-like growth factor 2 mRNA-binding protein 3	IPI00165467	22 kDa	3 0		0	0	0.00	6	3	0.50	10		0.70		0.1
Isoform F of Protein SON	IP100000192	264 kDa	4 3		0	0		9	6		2		1.00		0.1
Small nuclear ribonucleoprotein F	IP100220528	10 kDa	3 3	3 1.00	9	7	0.78	2	1	0.50	1	2	2.00	1.07	0.6
Nucleoly sin TIAR	IP100005615	42 kDa	8 (	0.00	0	0		2	2	1.00	7	4	0.57	0.52	0.5
Isoform 1 of DNA-dependent protein kinase catalytic subunit	IP100296337	469 kDa	3 1	0.33	2	2	1.00	2	3	1.50	4	2	0.50	0.83	0.5
M icrofibrillar-associated protein 1	IP100022790	52 kDa	4 0	0.00	0	0	0.00	8	8	1.00	4	1	0.25	0.63	0.5
Peptidyl-prolyl cis-trans isomerase H	IP100007346	19 kDa	9 5	0.56	5	1	0.20	2	0	0.00	0	0	0.00	0.38	0.2
Small nuclear ribonucleoprotein G	IPI00016572	8 kDa	7 7	7 1.00	4	1	0.25	0	1	n/a	0	. 1	rva	0.63	0.5
Isoform 1 of 60S ribosomal protein L12	IP100024933	18 kDa	1 3	2.00	0	0	0.00	10	4	0.40	8	0	0.00	0.80	1.0
Interferon-inducible dsRNA-dependent protein kinase activator A	IP100021167	34 kDa	4 (		0		0.00	7	6	0.86	5	2	0.40	0.63	0.3
Isoform 1 of U4/U6 small nuclear ribonucleoprotein Prp3	IP100005861	78 kDa	14	0.07	3		0.00	. 1	5	5.00	. 0	0	0.00	0.04	0.0
Isoform 2 of NLR family member X 1	IPI00377214	102 kDa	0 1	n/a	5		0.00	.0	0	0.00	3		0.67	0.33	0.4
Isoform 4 of Heterogeneous nuclear ribonucleoprotein A/B	IP100106509	31 kDa	0 0		0	0		3	1	0.33	10		0.80	0.57	0.3
A -kinase anchor protein 8	IPI00014474	76 kDa	0 1	0,00	0	0	0.00	5	6		2		3.00	2.10	1.2
Isoform Short of Heterogeneous nuclear ribonucleoprotein U	IP100479217	89 kDa	0 2	0/0	0		0.00	6	. 7	1.17	-1	0	0.00	1.17	0.0
Ribosomal protein L14 variant	IP100555744	24 kDa	1 0		. 0	0		9	8		2		1.00		0.0
Tubulin beta-2A chain	IPI00013475	50 kDa	0 2	n/a	0	0		9	10		4		0.00		0.7
Endoribonuclease Dicer	IP100219036	219 kDa	0 1	17/4	0	0		19	0		2		0.00	0.00	0.0
60S ribosomal protein L24	IP100306332	18 kDa	5 1	100000	0	0		4	5		7		0.00		0.6
Nuclease-sensitive element-binding protein 1 Isoform 1 of Serine/arginine repetitive matrix protein 2	IPI00031812 IPI00782992	36 kDa 300 kDa	7 7		0	0		5	5	0.50	0		3.00	2.00 0.75	0.3
Isoform 1 of Bcl-2-associated transcription factor 1	IP100762992	106 kDa	0 0		0	0		7	12		0			1.71	0.0
Zinc finger matrin-type protein 2	IP100069734	24 kDa	0 0		- 1			6	8		1		1.00		0.2
Interleukin enhancer-binding factor 2	IP100005734	43 kDa	1 1		0			9	0		7		0.14		0.5
Superkiller viralicidic activity 2-like 2	IP100647217	118 kDa	0 0		0	0		6	6		3		1.00		0.0
NHP2-like protein 1	IPI00026167	14 kDa	7		7		0.00	0	0	0.00	0		0.00	0.21	0.3
60S ribosomal protein L13a	IPI00304612	24 kDa	0 0		0	0		10	6		0	0		0.60	0.0
Py ruvate dehy drogenase E1 component subunit alpha, testis-specific	IP100024087	43 kDa	2 0		1	6	6.00	0	2	n/a	1	5	5.00	5.50	0.7
Isoform ASF-1 of Splicing factor, arginine/serine-rich 1	IPI00215884	28 kDa	0 0		0	0		6	6	1.00	3		0.33	0.67	0.4
Iso 2 of Serine/threonine-protein phosphatase PGAM 5, mitochondrial	IP100063242	28 kDa	0 0		0	0		10	4	0.40	5		0.00	0.20	0.2
Isoform 1 of Clathrin heavy chain 1	IP100024067	192 kDa	0 0		0	0	0.00	4	4	1.00	0	0		1.00	0.0
DEAH (Asp-Glu-Ala-His) box polypeptide 16	IPI00292510	63 kDa	4 2	0.50	0	0	0.00	7	- 1	0.14	2	1	0.50	0.38	0.2
Propionyl Coenzyme A carboxylase, alpha polypeptide	IPI00553241	13 kDa	0 0	0.00	.1	3	3.00	- 1	2	2.00	5	5	1.00	2.00	1.0
40S ribosomal protein S11	IP100025091	18 kDa	1 1	1 1.00	. 0	0		3	2	0.67	6	0	0.00	0.56	0.5
Isoform 2 of Heterogeneous nuclear ribonucleoprotein A/B	IP100334587	36 kDa	0 0	0.00	0	0		5	3	0.60	5	5	1.00	0.40	0.4
60S ribosomal protein L21	IP100247583	19 kDa	1 0	0.00	0	0		2	2	1.00	6	2	0.33	0.67	0.4
Isoform 1 of RNA-binding protein 8A	IP100001757	20 kDa	4 (	0.00	1	0		2	3	1.50	0	2	n/a	0.75	1.0
Poly merase delta interacting protein 46	IP100429180	20 kDa	0 0	0.00	0	0		4	3	0.75	2	5	2.50	1.62	1.2
40S ribosomal protein S8	IP100216587	24 kDa	0 0	0.00	0	0		5	6	1.20	4	0	0.00	0.60	0.8
Elongation factor 1-alpha 2	IPI00014424	50 kDa	4 5		1	1	1.00	2	2	1.00	0	0		1.08	0.1
40S ribosomal protein S15a	IP100221091	15 kDa	1 1		- 1	0		8	3	0.38	- 1	0	0.00	0.69	0.4
Isoform 1 of Peptidyl-prolyl cis-trans isomerase CWC27 homolog	IPI00025174	54 kDa		0.75	2			2	5		0			1.62	1.2
Isoform 4 of Dynein heavy chain 6, axonemal	IP100884959	290 kDa	0 0		5		1.00	.1	2	2.00	0			1.50	0.7
cDNA FLJ54030, highly similar to Polymerase delta-interacting protein 3	IP100440688	48 kDa	0 0		0		0.00	5	11	2.20	0		0.00	2.20	0.0
Isoform 3 of Probable ATP-dependent RNA helicase DDX 17	IP100651653	73 kDa	1 1		0			6	4	0.67	2		0.50		0.2
Isoform 1 of Peptidyl-prolyl cis-trans isomerase-like 2	IP100003824	59 kDa	2 0		0	0		3	5	1.67	1		1.00	1.33	0.4
Protein DGCR14	IP100165171	53 kDa	0 0		0	0	0.00	3	9		0	0	0.00	3.00	0.0
Putative uncharacterized protein ENSP00000382160	IP100180956	49 kDa	1 3			1	1.00	2	1	0.50	0	4	0.50	1.50	1.3
T argeting protein for X klp2 WD repeat domain 57 (U5 snRNP specific), isoform CRA_b	IP100008477	86 kDa	3 (			0		4 2	3		1	0	0.50		0.1
WD repeat domain 57 (US SHKNP specific), isoform CKA_B HLA-B associated transcript 1	IPI00385642 IPI00641634	14 kDa 26 kDa	0 0		0	0	0.00	2	5		5		1.00	1.25	0.0
TDP43	IP100041634	45 kDa	1 1					2	4		2		1.00		0.5
U6 snRNA-associated Sm-like protein LSm3	IP100023813	12 kDa		0.00			0.40	1	1	1.00	0		0.00	0.46	0.5
U4/U6.U5 tri-snRNP-associated protein 1	IP100219229	90 kDa	8 1		1		1.00	0	- 1	1.00	0			0.46	0.6
Isoform 1 of Heterogeneous nuclear ribonucleoprotein A3	IP100021417	40 kDa	0 2		0		0.00	3	1	0.33	6		0.00		0.2
ATP-dependent RNA helicase DHX8	IP100419373	139 kDa	0 1	n/a	1			0	- 5	n/a	1				0.0
Isoform 3 of PCI domain-containing protein 2	IP100072541	43 kDa			0	0		1	3		2	0	0.00	1.66	1.8
32 kDa protein	IP100399077	32 kDa	0 0		0			3	3		5		0.20		0.5
HSPA 5 protein	IP100003362	72 kDa	5 3		1		1.00	0	0	0.00	0		0.00	0.80	0.2
33 kDa protein	IPI00413108	33 kDa	0 0		0		0.00	5	2		5		0.00		0.2
Protein CASC3	IP100289491	76 kDa			0			0	8		0		n/a	0.00	0.0
Isoform 1 of Heterogeneous nuclear ribonucleoprotein D0	IP100028888	38 kDa	0 0		0			4	5		9				0.4
Splicing factor, arginine/serine-rich 9	IPI00012340	26 kDa			0			6	1	0.17	2		0.00	0.17	0.0
Isoform 1 of Splicing factor U2AF 65 kDa subunit	IPI00031556	54 kDa		0.00	0			7			0		0.00	0.14	0.0

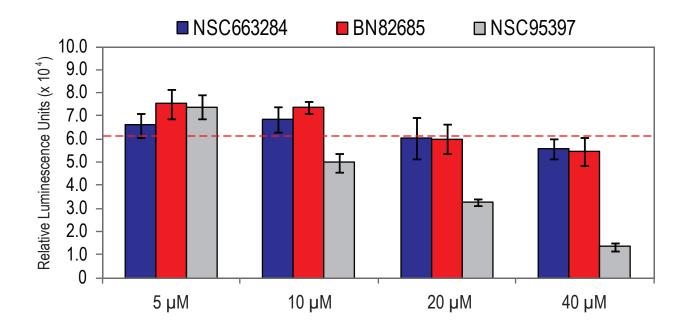
**Table S1.** Peptide counts for top proteins (138-203) detected by mass spectrometry of DMSO and BN82685 streptavidin pull downs.



Berg et al. Figure S1



Berg et al. Figure S2



Berg et al. Figure S3