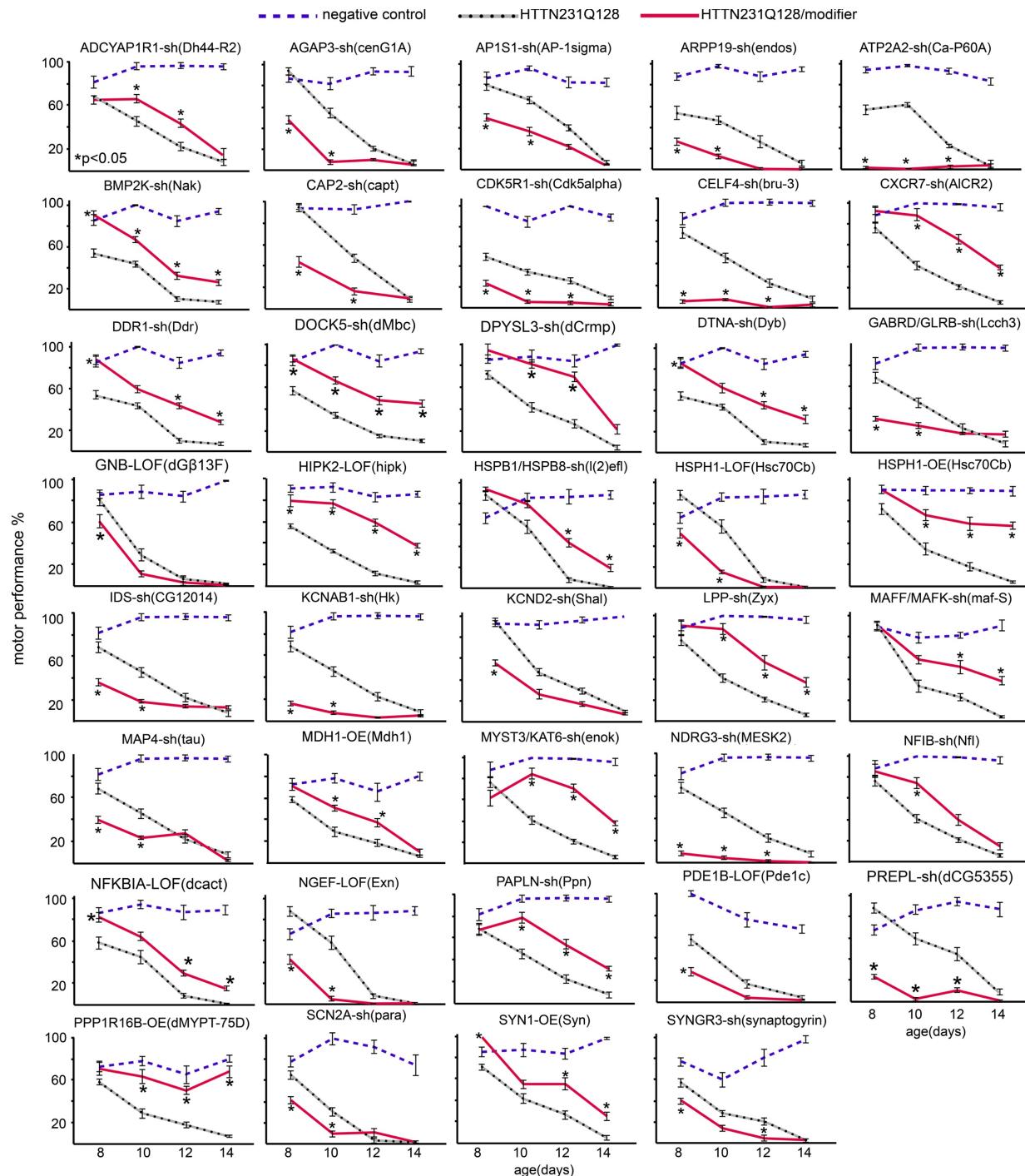


**Figure S1. *Drosophila* motor performance assay data for genes in the compensatory network.**

**Related to Figure 1.**

Charts show motor performance as a function of age in *Drosophila* negative controls (blue dashed lines,  $elav^{C155}>GAL4/w1118$ ), positive controls expressing HTTN231Q128 in the nervous system (black dotted line,  $elav^{C155}>GAL4/w1118; UAS-HTTN231Q128/+$ ) and experimental animals (red line,  $elav^{C155}>GAL4/w1118; UAS-HTTN231Q128/+; modifier/+$ ). sh: shRNA; LOF: loss of function; OE: overexpression. Error bars in motor performance charts: s.e.m. Significant differences identified using Anova followed by Tukey's post hoc test for each time point ( $\alpha=0.05$ ).

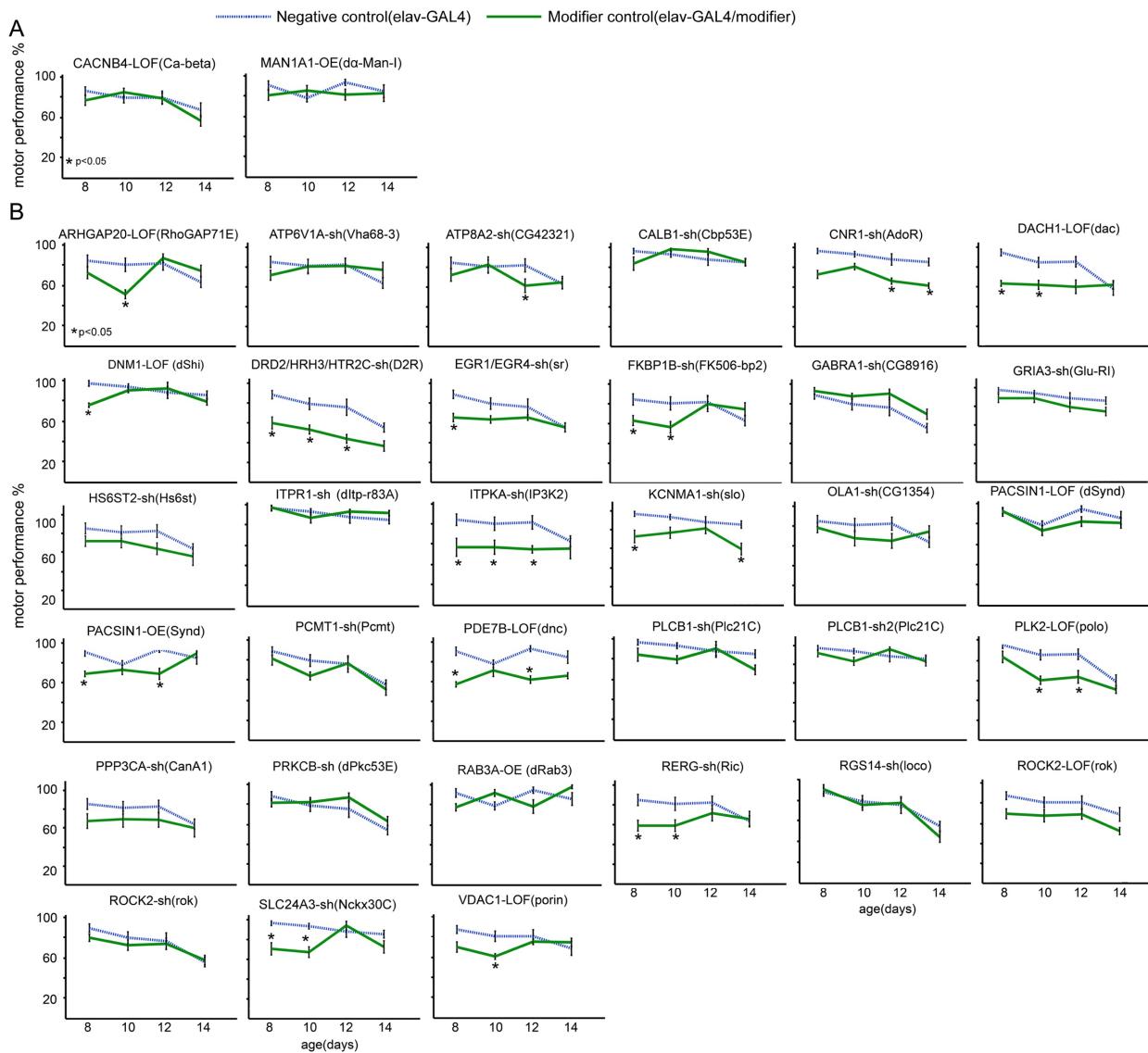
**Figure S2. *Drosophila* motor performance assay data for genes in the pathogenic networks Related to Figure 1.**



**Figure S2. *Drosophila* motor performance assay data for genes in the pathogenic networks Related to Figure 1.**

Charts show motor performance as a function of age in *Drosophila* negative controls (blue dashed lines, *elav<sup>C155</sup>>GAL4/w1118*), positive controls expressing HTTN231Q128 in the nervous system (black dotted line, *elav<sup>C155</sup>>GAL4/w1118; UAS-HTTN231Q128/+*) and experimental animals (red line, *elav<sup>C155</sup>>GAL4/w1118; UAS-HTTN231Q128/+; modifier/+*). sh: shRNA; LOF: loss of function; OE: overexpression. Error bars in motor performance charts: s.e.m. Significant differences identified using Anova followed by Tukey's post hoc test for each time point ( $\alpha=0.05$ ).

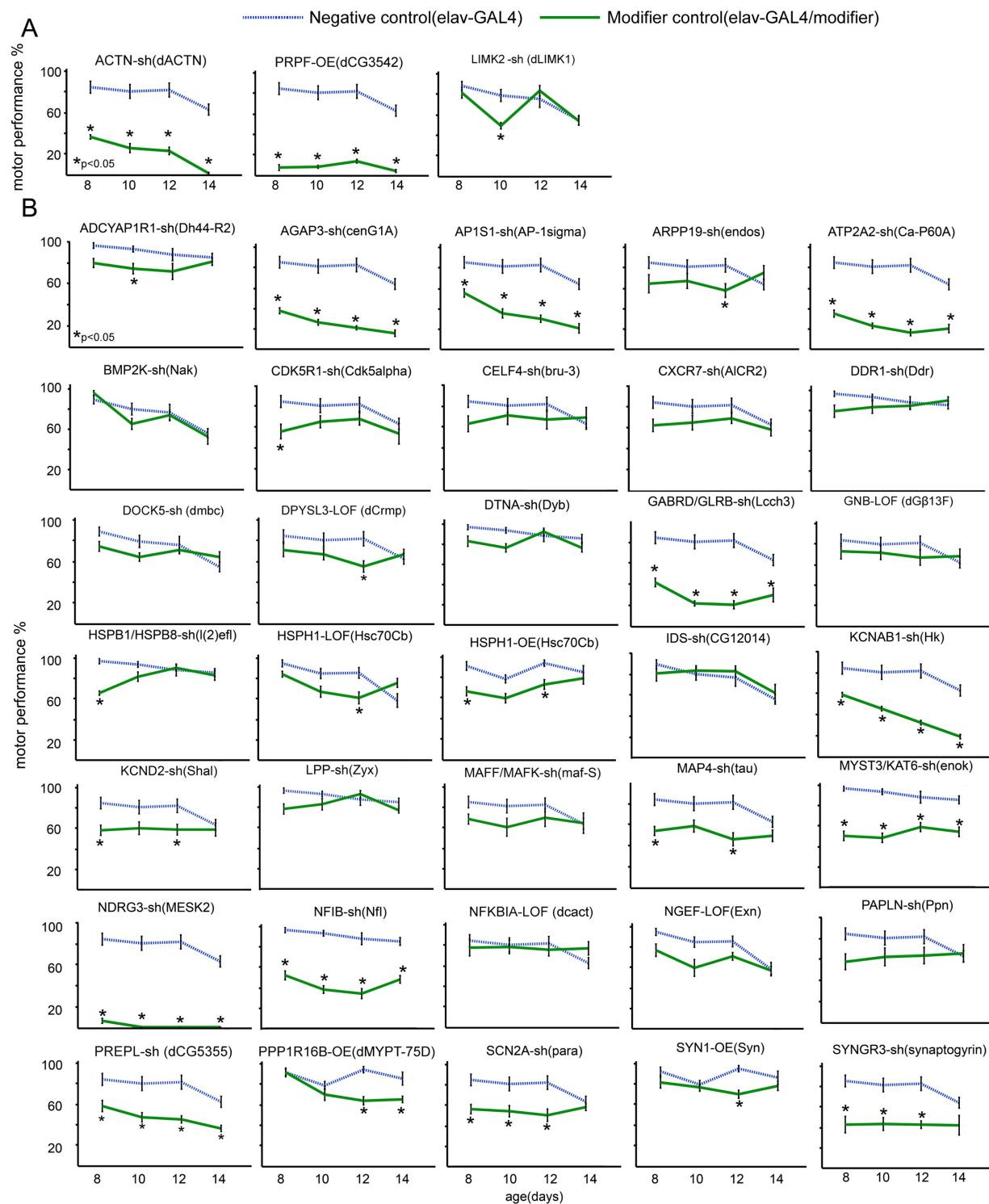
**Figure S3. *Drosophila* motor performance tests for the modifier alleles of the compensatory network in control animals. Related to Figure 1.**



**Figure S3. *Drosophila* motor performance tests for the modifier alleles of the compensatory network in control animals. Related to Figure 1.**

Blue lines indicate the motor performance of negative controls ( $elav^{C155}/w1118$ ). Green line shows motor performance of animals carrying the indicated modifier alleles but not expressing HTTN231Q128 ( $elav^{C155}/w1118$ ; *modifier*/+). Note that some modifiers impair the motor performance of the negative controls ( $elav^{C155}/w1118$ ) but these modifiers nonetheless improve the performance of HD animals ( $elav^{C155}>GAL4/w1118$ ; *UAS-HTTN231Q128*/+; see **Figure S1**). sh: shRNA; LOF: loss of function; OE: overexpression. Green error bars in gene expression scatter plots: average and standard deviation. Error bars in motor performance charts: s.e.m. Significant differences identified using Anova followed by Tukey's post hoc test for each time point ( $\alpha=0.05$ ). Panel S3A shows the control assays for the modifiers shown in Figure 1A. Panel S3B shows the control assays for the modifiers shown in Figure S1.

**Figure S4. *Drosophila* motor performance tests for the modifier alleles of the pathogenic network in control animals. Related to Figure 1.**

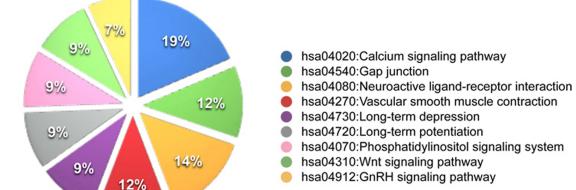


**Figure S4. *Drosophila* motor performance tests for the modifier alleles of the pathogenic network in control animals. Related to Figure 1.**

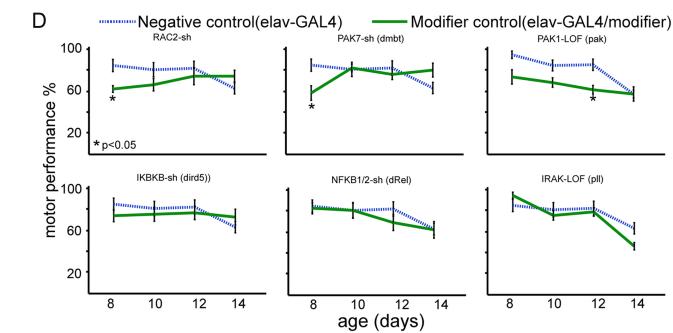
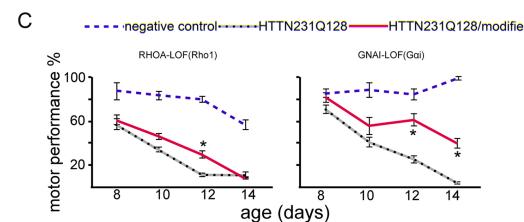
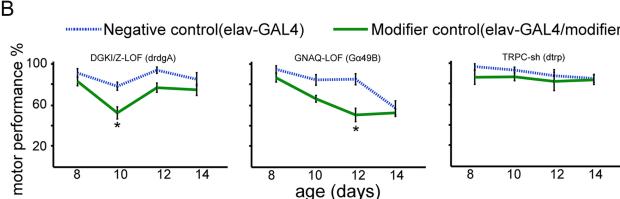
Blue lines indicate the motor performance of negative controls ( $elav^{C155}/w1118$ ). Green line shows motor performance of animals carrying the indicated modifier alleles but not expressing HTTN231Q128 ( $elav^{C155}/w1118$ ; *modifier*/+). Note that some modifiers impair motor performance of both the negative controls ( $elav^{C155}/w1118$ ) and the HD animals ( $elav^{C155}>GAL4/w1118$ ; *UAS-HTTN231Q128*/+; see **Figure S2**); dysregulation of these genes is likely to contribute to HD pathogenesis. sh: shRNA; LOF: loss of function; OE: overexpression. Green error bars in gene expression scatter plots: average and standard deviation. Error bars in motor performance charts: s.e.m. Significant differences identified using Anova followed by Tukey's post hoc test for each time point ( $\alpha=0.05$ ). Panel S4A shows the control assays for the modifiers shown in Figure 1C. Panel S4B shows the control assays for the modifiers shown in Figure S2.

Figure S5. Pathway enrichment, additional climbing data and additional pathway analysis related to Figure 4.

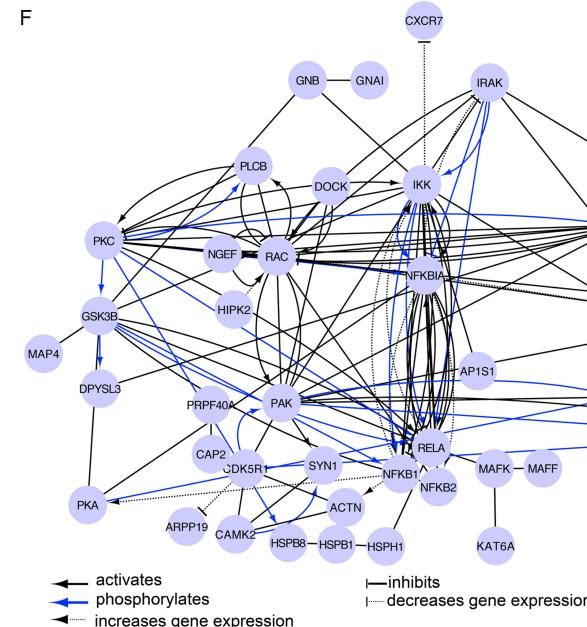
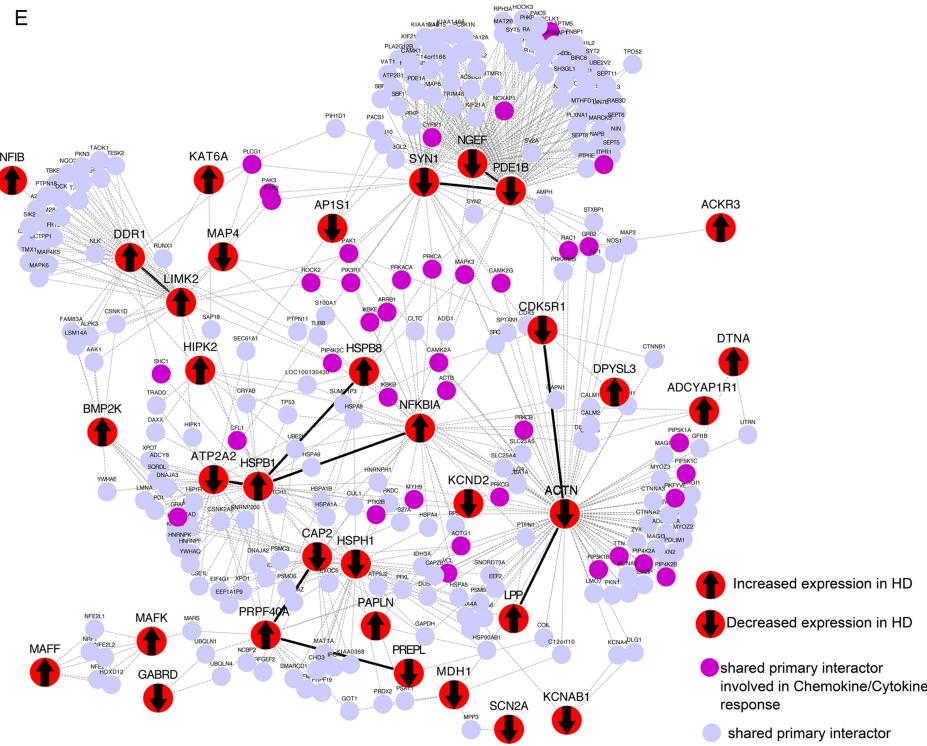
A. Pathway enrichment analysis of genes in the compensatory network.



B. Climbing data for controls and additional modifiers shown in Figure 4.



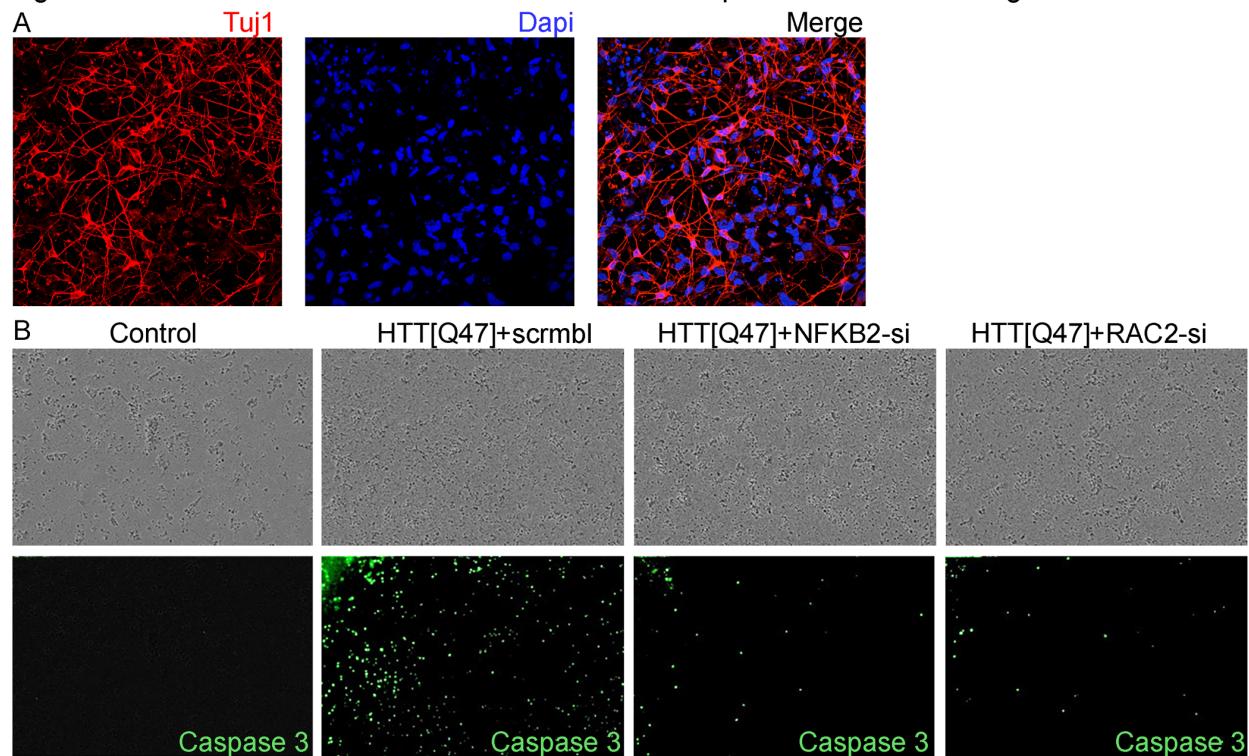
Additional network analysis of the pathogenic subnetwork reveals enrichment in genes involved in inflammation and actin cytoskeleton



**Figure S5. Pathway enrichment, additional climbing data and additional pathway analysis related to Figure 4.**

**(A)** Pie chart illustrating the pathway enrichment within the compensatory category compared to the background of genes expressed in the striatum, using DAVID. The gene category most enriched is KEGG-hsa04020: “Ca<sup>2+</sup> signaling pathway” (EASE score p=9.21E-06, Benjamini 4.70E-04). **(B-D)** Motor performance data for controls and additional modifiers shown in Figure 4. **(B and D)** Green line in charts shows motor performance of animals carrying the indicated modifier allele but not expressing HTTN231Q128 (*elav*<sup>C155</sup>/*w1118*; *modifier*/+) compared to *elav*<sup>C155</sup>/*w1118* negative controls (blue line). **(C)** Additional modifiers of HTTN231Q128-induced neuronal dysfunction identified through pathway expansion in the pathogenic network. sh: shRNA; LOF: loss of function; OE: overexpression. Green error bars in gene expression scatter plots: average and standard deviation. Error bars in motor performance charts: s.e.m. Significant differences identified using Anova followed by Tukey’s post hoc test for each time point ( $\alpha=0.05$ ). **Table S5** lists the specific modifier alleles identified through pathway expansion. **(E-F)** Additional network analysis of the pathogenic network reveals an enrichment in genes involved in inflammation and actin cytoskeleton regulation. **(E)** Network of genes categorized as pathogenic showing *Drosophila* modifiers altered in the human HD transcriptome (red circles with arrows) and their shared primary interactors. Note abundance of genes involved in inflammatory response and actin cytoskeleton regulation, downstream of chemokine/cytokine receptors (dark purple nodes). **(F)** Complete representation of all the interactions of the *Drosophila* modifier genes in the pathogenic network and their primary interacting partners generated using Ingenuity Pathway Analysis. This is the basis for the summary scheme shown in **Figure 4C**.

**Figure S6 . Validation in iPS-derived neurons from HD patients. Related to Figure 5.**



**Figure S6. Validation in iPS-derived neuron-like cells from HD patients. Related to Figure 5.**

**(A)** Immunofluorescence staining of iPS-derived cells showing positive signal for the Tuj1 neuronal marker (red), which also highlights their neuron-like morphology. **(B)** Bright field and fluorescence images of control and HTT[Q47] iPS-derived neuron-like cells transfected with a scramble, RAC2 and NFKB2 targeting siRNAs 12 hours after BDNF deprivation. Caspase-3 activation signal is shown in green.

**Supplementary Tables.**

**Table S1. Genes considered for this study because their expression is altered early during human HD progression (Vonsattel grade 0-2 patients) and in at least one HD mouse model. Related to Figure 1.** For the majority of these genes (67%), the direction of the transcriptional alteration (up or down) is the same between HD patients and at least one of the mouse HD models, i.e., not all models show transcriptional change in the same direction.

**Table S2. Human genes whose expression alterations in HD we categorized as compensatory.**

**Related to Figure 1.** Also shown: the *Drosophila* homolog, blast e-value, direction of transcriptomic change in human, class of the modifier allele and specific allele that modified the HTTN231Q128-induced behavioral deficits.

**Table S3. Human genes whose expression alterations in HD we categorized as pathogenic. Related to Figure 1.** Also shown: the *Drosophila* homolog, blast e-value, direction of transcriptomic change in human, class of the modifier allele and specific allele that modified the HTTN231Q128-induced behavioral deficits.

**Table S4. Pathways enriched in the compensatory and pathogenic networks. Related to Figure 4.**

Analysis performed using primary modifier genes from the screen and shared primary protein interactors using either Inweb or STRING databases. Query performed using DAVID/KEGG databases. P-value represents EASE enrichment score.

**Table S5. Additional modifiers identified through pathway analysis. Related to Figure 4.**

**Table S6. Complete list of siRNAs used for the HTRF screen in human HD fibroblasts. Related to Figures 5 and 6.**

Table S2. Human genes whose gene expression alterations in HD are categorized as compensatory. Related to Figure 1.

Human/ Mouse Gene Symbol	Drosophila Homologous Gene	Protein Blast e-value	Transcriptomic change in human	Allele Class	Modifier Allele	Effect on <i>Drosophila</i> HD model	Effect of allele on control motor performance
ARHGAP20	RhoGAP71E	9E-22	DOWN	LOF	RhoGAP71E[j6b9]	S	N
ATP6V1A	Vha68-3	0	DOWN	shRNA*	Vha68-3[GD11527]-v34926	S	N
ATP6V1A	Vha68-3	0	DOWN	shRNA	Vha68-3[GD11527]-v41646	S	N
ATP8A2	CG42321	0	DOWN	LOF	CG42321[KG03082]	S	N
ATP8A2	CG42321	0	DOWN	LOF*	CG42321[KG05950]	S	N
CACNA1B	cac	0	DOWN	LOF*	cac[HC129]	S	N
CACNB4	Ca-beta	6E-178	DOWN	LOF	Ca-beta[I(2)04008]	S	N
CALB1	Cbp53E	1E-68	DOWN	shRNA	Cbp53E[GD10584]	S	N
CNR1	AdoR	4E-21	DOWN	shRNA	AdoR[GD380]	S	I
DACH1	dac	1E-54	DOWN	LOF	dac[9]	S	I
DNM1	shi	0	DOWN	LOF	shi[1]	S	N
DRD2	D2R	2E-79	DOWN	shRNA	D2R[GD732]	S	I
EGR1	sr	1E-64	DOWN	shRNA	sr[KK107677]	S	N
EGR4	sr	1E-46	DOWN	shRNA	sr[KK107677]	S	N
FKBP1B	FK506-bp2	3E-54	DOWN	shRNA*	FK506-bp2[GD6658]-v45014	S	N
FKBP1B	FK506-bp2	3E-54	DOWN	shRNA	FK506-bp2[GD6658]-v45015	S	N
GABRA1	CG8916	4E-96	DOWN	shRNA	CG8916[GD3383]	S	N
GRIA3	Glu-RI	0	DOWN	shRNA	Glu-RI[KK101533]	S	N
GRIA3	Glu-RI	0	DOWN	shRNA*	Glu-RI[GD3582]	S	N
HRH3	D2R	3E-37	DOWN	shRNA	D2R[GD732]	S	I
HS6ST2	Hs6st	6E-108	DOWN	shRNA	Hs6st[KK101636]	S	N
HTR2C	D2R	1E-54	DOWN	shRNA	D2R[GD732]	S	I
ITPKA	IP3K2	7E-120	DOWN	shRNA	IP3K2[GD8778]	S	I
ITPR1	Itp-r83A	0	DOWN	shRNA	Itp-r83A[GD1676]	S	N
KCNMA1	slo	0	DOWN	shRNA	slo[GD244]	S	I
MAN1A1	alpha-Man-I	0	DOWN	OE	alpha-Man-I[EY00697]	E	N
OLA1	CG1354	4E-113	DOWN	shRNA	CG1354[GD8048]	S	N
PACSIN1	Synd	2E-121	DOWN	LOF	Synd[EY07010]	S	N
PACSIN1	Synd	2E-121	DOWN	OE	Synd[EP877]	E	N
PCMT1	Pcmt	4E-83	DOWN	shRNA	Pcmt[GD8637]	S	N
PDE7B	dnc	2E-52	DOWN	LOF	dnc[M14]	S	N
PLCB1	Plc21C	1E-145	DOWN	shRNA	Plc21C[GD11359]-v26557	S	N
PLCB1	Plc21C	1E-145	DOWN	shRNA	Plc21C[GD11359]-v26558	S	N
PLK2	polo	3E-132	DOWN	LOF	polo[01673]	S	I
PPP3CA	CanA1	0	DOWN	shRNA	CanA1[GD8366]	S	N
PRKCB	Pkc53E	0	DOWN	shRNA	Pkc53E[GD11984]	S	N
RAB3A	Rab3	1E-125	DOWN	OE	P{UASp-YFP.Rab3.Q80L}11	E	N
RERG	Ric	2E-35	DOWN	RNAi	Ric[GD13919]	S	I
RGS14	loco	4E-37	DOWN	shRNA	loco[GD1282]	S	N
ROCK2	rok	0	DOWN	shRNA	rok[GD1522]	S	N
ROCK2	rok	0	DOWN	LOF	rok[1]	S	N
SLC24A3	Nckx30C	2E-114	DOWN	shRNA	Nckx30C[GD1040]	S	N
STXBP1	Rop	0	DOWN	OE*	Rop[G4478]	E	N
STXBP1	Rop	0	DOWN	LOF*	Rop[G27]	S	N
SYNE1	Msp-300	0	DOWN	LOF	Msp-300[DG03312]	S	N
VDAC1	porin	5E-121	DOWN	LOF	porin[k05123]	S	N

LOF: loss of function; OE: overexpression; shRNA: inducible hairpin RNAi; S: suppressor; E: enhancer; N: no effect; I: impaired; \* data not shown

Table S3. Human genes whose gene expression alterations in HD are categorized as pathogenic. Related to Figure 1.

Human/ Mouse Gene Symbol	Drosophila Homologous Gene	Protein Blast e-value	Transcriptomic change in human	Allele Class	Modifier Allele	Effect on <i>Drosophila</i> HD model	Effect of allele on control motor performance
ACTN1	Actn	0	DOWN	shRNA	Actn[GD7761]	E	I
ACTN2	Actn	0	DOWN	shRNA	Actn[GD7761]	E	I
ADCYAP1R1	Dh44-R2	2E-47	UP	shRNA	Dh44-R2[GD15731]	S	N
AGAP3	cenG1A	8E-121	DOWN	shRNA	cenG1A[KK104806]	E	I
AP1S1	AP-1sigma	1E-86	DOWN	shRNA	AP-1sigma[KK108869]	E	I
ARPP19	endos	9E-20	DOWN	shRNA	endos[KK108642]	E	N
ARPP19	endos	9E-20	DOWN	shRNA*	endos[GD10576]	E	N
ATP2A2	Ca-P60A	0	DOWN	shRNA	Ca-P60A[KK107371]	E	I
BMP2K	Nak	6E-136	UP	shRNA	Nak[GD12597]	S	N
CAP2	capt	5E-134	DOWN	shRNA	capt[KK103925]	E	N
CDK5R1	Cdk5alpha	8E-71	DOWN	shRNA	Cdk5alpha[KK100567]	E	N
CELF4	bru-3	1E-110	DOWN	shRNA	bru-3[KK111663]	E	N
CXCR7	AICR2	2E-32	UP	shRNA	AICR2[GD17055]	S	N
DDR1	Ddr	8E-64	UP	shRNA	Ddr[GD7334]	S	N
DOCK5	mbc	0	UP	shRNA	mbc[GD6965]	S	N
DPYSL3	CRMP	0	UP	LOF	CRMP[supl2]	S	N
DPYSL3	CRMP	0	UP	shRNA*	CRMP[GD4716]	S	N
DTNA	Dyb	2E-137	UP	shRNA	Dyb[KK109268]	S	N
GABRD	Lcch3	1E-98	DOWN	shRNA	Lcch3[KK100854]	E	I
GLRB	Lcch3	2E-80	DOWN	shRNA	Lcch3[KK100854]	E	I
HIPK2	hipk	0	UP	LOF	hipk[EY12599]	S	N
HSPB1	I(2)efl	1E-28	UP	shRNA	I(2)efl[GD11341]	S	N
HSPB8	I(2)efl	4E-24	UP	shRNA	I(2)efl[KK108279]	S	N
HSPB8	I(2)efl	4E-24	UP	shRNA	I(2)efl[GD11341]	S	N
HSPH1	Hsc70Cb	0	DOWN	LOF	Hsc70Cb[00082]	E	N
HSPH1	Hsc70Cb	0	DOWN	OE	Hsc70Cb[EY00671]	S	N
IDS	CG12014	6E-140	DOWN	shRNA	CG12014[KK101391]	E	N
KCNAB1	Hk	1E-112	DOWN	shRNA	Hk[KK109058]	E	I
KCND2	Shal	0	DOWN	shRNA	Shal[KK100264]	E	I
LIMK2	LIMK1	2E-148	UP	shRNA	LIMK1[GD9586]-v25343	S	N
LIMK2	LIMK1	2E-148	UP	shRNA	LIMK1[GD9586]-v25344	S	N
LPP	Zyx	6E-92	UP	shRNA	Zyx[GD10696]	S	N
MAFF	maf-S	1E-26	UP	shRNA	maf-S[KK102044]	S	N
MAFK	maf-S	3E-30	UP	shRNA	maf-S[KK102044]	S	N
MAP4	tau	8E-29	DOWN	shRNA	tau[KK109359]	E	I
MDH1	Mdh1	6E-150	DOWN	OE	Mdh1[EY08761]	S	N
MYST3/KAT6	enok	2E-111	UP	shRNA	enok[GD4037]	S	I
NDRG3	MESK2	9E-79	DOWN	shRNA	MESK2[GD9083]	E	I
NFIB	Nfl	2E-92	UP	shRNA	Nfl[GD4283]	S	I
NFKBIA	cact	2E-35	UP	LOF*	cact[KG00376]	S	N
NFKBIA	cact	2E-35	UP	LOF	cact[KG07677]	S	N
NGEF	Exn	7E-76	DOWN	LOF	Exn[EP3042]	E	N
PAPLN	Ppn	4E-89	UP	shRNA	Ppn[KK101228]	S	N
PDE1B	Pde1c	3E-161	DOWN	LOF	Pde1c[c04487]*	E	N
PPP1R16B	MYPT-75D	4E-102	DOWN	OE	UAS-MYPT-75D.HM}B9	S	I
PREPL	CG5355	5E-23	DOWN	shRNA	CG5355[KK100768]	E	I
PRPF40A	CG3542	6E-172	UP	OE	CG3542[EP719]	E	I
SCN2A	para	0	DOWN	shRNA	para[GD3392]	E	I
SYN1	Syn	7E-118	DOWN	OE	Syn[EY01930]	S	N
SYNGR3	synaptogyrin	1E-41	DOWN	shRNA	synaptogyrin[KK108648]	E	I

LOF: loss of function; OE: overexpression; shRNA: inducible hairpin RNAi; S: suppressor; E: enhancer; N: no effect; I: impaired; \* data not shown

**Table S4. Pathways enriched in the compensatory and pathogenic networks. Related to Figure 4.**

Compensatory						
	David Source	Pathway	Gene Count	%	P-Value	Benjamini
Inweb	KEGG_PATHWAY	hsa04720:Long-term potentiation	4	44.44	4.45E-05	8.00E-04
	KEGG_PATHWAY	hsa04070:Phosphatidylinositol signaling system	4	44.44	5.74E-05	6.88E-04
	KEGG_PATHWAY	hsa04020:Calcium signaling pathway	6	66.67	2.74E-07	9.87E-06
Pathogenic						
	David Source	Pathway	Gene Count	%	P-Value	Benjamini
Inweb	KEGG_PATHWAY	Regulation of actin cytoskeleton	26	8.4	7.4E-09	2.9E-07
	PANTHER_PATHWAY	P00031:Inflammation mediated by chemokine and cytokine signaling pathway	25	8.1	1.8E-03	3.9E-02
	KEGG_PATHWAY	MAPK signaling pathway	22	7.1	7.2E-05	5.3E-04
	KEGG_PATHWAY	Tight junction	21	6.8	3.7E-09	2.2E-07
	KEGG_PATHWAY	Calcium signaling pathway	21	6.8	4E-07	6.8E-06
	KEGG_PATHWAY	Focal adhesion	21	6.8	3.4E-06	3.6E-05
	PANTHER_PATHWAY	P00005:Angiogenesis	19	6.1	5.3E-03	6.6E-02
	KEGG_PATHWAY	Pathways in cancer	19	6.1	1.3E-02	4.7E-02
	KEGG_PATHWAY	Adherens junction	18	5.8	1.1E-10	1.3E-08
	KEGG_PATHWAY	Leukocyte transendothelial migration	18	5.8	1E-07	2.4E-06
	David Source	Pathway	Gene Count	%	P-Value	Benjamini
STRING	KEGG_PATHWAY	Regulation of actin cytoskeleton	57	11	2.1E-23	9E-22
	KEGG_PATHWAY	Focal adhesion	56	10.8	3.9E-24	2.5E-22
	KEGG_PATHWAY	Pathways in cancer	56	10.8	1.2E-13	2.2E-12
	PANTHER_PATHWAY	P00031:Inflammation mediated by chemokine and cytokine signaling pathway	52	10	3.3E-07	1.1E-05
	KEGG_PATHWAY	MAPK signaling pathway	45	8.7	8.8E-11	1.3E-09
	PANTHER_PATHWAY	P00034:Integrin signalling pathway	45	8.7	1E-08	5E-07
	KEGG_PATHWAY	Dilated cardiomyopathy	42	8.1	1.9E-27	2.5E-25
	KEGG_PATHWAY	Chemokine signaling pathway	34	6.6	4.3E-09	3.1E-08
	KEGG_PATHWAY	Hypertrophic cardiomyopathy (HCM)	32	6.2	7.3E-18	1.9E-16
	KEGG_PATHWAY	Tight junction	32	6.2	1E-11	1.7E-10

**Table S5. Additional modifiers identified through pathway analysis. Related to Figure 4.**

Human/Mouse Gene Symbol	Drosophila Homologous Gene	Protein Blast e-value	Allele Class	Modifier Allele
GRM3	mGlurA	0	shRNA	GD707
GNAQ	Gα49B	0	LOF	KG07290
TRPC4/5	trp	1.46288E-172	shRNA	GD372
DGKI/Z	rdgA	0	LOF	EY11543
IKBKB	ird5	8.11278E-47	shRNA	GD11248
NFKB1/2	rel	7E-56	shRNA	HM05154
IRAK	pll	1.5784E-39	LOF	pll <sup>2</sup>
PAK1	pak	0	LOF	pak <sup>11</sup>
PAK7	mbt	2E-176	shRNA	GD9608
RAC2	Rac2	4.47228E-99	shRNA	GD17536

**Table S6. Complete list of siRNAs used for the HTRF screen in human HD fibroblasts. related to Figures 5 and 6.**

Gene Symbol	siRNA	Source									
ACTN1	J-011195-05	Dharmacon	DDR1	SI00030513	Qiagen	IKBKE	J-003723-12	Dharmacon	NFKBIA	J-004765-05	Dharmacon
ACTN1	J-011195-06	Dharmacon	DDR1	SI00605437	Qiagen	IKBKE	SI00300538	Qiagen	NFKBIA	J-004765-06	Dharmacon
ACTN1	J-011195-07	Dharmacon	DDR1	SI00605444	Qiagen	IKBKE	SI02622319	Qiagen	NFKBIA	J-004765-07	Dharmacon
ACTN1	J-011195-08	Dharmacon	DDR1	SI03649254	Qiagen	IKBKE	SI02622326	Qiagen	NFKBIA	J-004765-08	Dharmacon
ACTN1	SI02654407	Qiagen	DNM1	J-003940-05	Dharmacon	IKBKE	SI02655324	Qiagen	NFKBIA	SI0126812	Qiagen
ACTN1	SI02654414	Qiagen	DNM1	J-003940-06	Dharmacon	IRAK1	J-004760-12	Dharmacon	NFKBIA	SI0126819	Qiagen
ACTN1	SI02757223	Qiagen	DNM1	J-003940-07	Dharmacon	IRAK1	J-004760-13	Dharmacon	NFKBIA	SI0126826	Qiagen
ACTN1	SI02757230	Qiagen	DNM1	J-003940-08	Dharmacon	IRAK1	J-004760-14	Dharmacon	NFKBIA	SI03114930	Qiagen
ACTN2	J-011196-06	Dharmacon	DNM1	SI00603378	Qiagen	IRAK1	J-004760-15	Dharmacon	NEGF	J-009354-09	Dharmacon
ACTN2	J-011196-07	Dharmacon	DNM1	SI00603385	Qiagen	IRAK1	SI02626491	Qiagen	NEGF	J-009354-10	Dharmacon
ACTN2	J-011196-08	Dharmacon	DNM1	SI00603392	Qiagen	IRAK1	SI03099166	Qiagen	NEGF	J-009354-11	Dharmacon
ACTN2	J-011196-09	Dharmacon	DNM1	SI03071894	Qiagen	IRAK1	SI03649429	Qiagen	NEGF	J-009354-12	Dharmacon
ACTN2	SI00291508	Qiagen	DOCK5	J-018931-05	Dharmacon	IRAK1	SI04897011	Qiagen	NEGF	SI04345453	Qiagen
ACTN2	SI04171699	Qiagen	DOCK5	J-018931-06	Dharmacon	IRAK4	J-003302-11	Dharmacon	NEGF	SI04354553	Qiagen
ACTN2	SI04205544	Qiagen	DOCK5	J-018931-07	Dharmacon	IRAK4	J-003302-12	Dharmacon	NEGF	SI04356555	Qiagen
ACTN2	SI04315304	Qiagen	DOCK5	J-018931-08	Dharmacon	IRAK4	J-003302-13	Dharmacon	NEGF	SI04374839	Qiagen
ADCYAP1R1	J-005413-07	Dharmacon	DOCK5	SI03123645	Qiagen	IRAK4	J-003302-14	Dharmacon	OLA1	J-015680-09	Dharmacon
ADCYAP1R1	J-005413-08	Dharmacon	DOCK5	SI04227559	Qiagen	IRAK4	SI01113617	Qiagen	OLA1	J-015680-10	Dharmacon
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ADCYAP1R1	SI00022162	Qiagen	DPP6	J-010129-05	Dharmacon	IRAK4	SI03025771	Qiagen	OLA1	J-034873-11	Dharmacon
ADCYAP1R1	SI00022169	Qiagen	DPP6	J-010129-06	Dharmacon	ITPKA	J-006742-08	Dharmacon	OLA1	SI04178321	Qiagen
ADCYAP1R1	SI00022176	Qiagen	DPP6	J-010129-07	Dharmacon	ITPKA	J-006742-09	Dharmacon	OLA1	SI04220846	Qiagen
ADCYAP1R1	SI04894694	Qiagen	DPP6	J-010129-08	Dharmacon	ITPKA	J-006742-10	Dharmacon	OLA1	SI04348806	Qiagen
AGAP3	J-019127-09	Dharmacon	DPP6	SI00149835	Qiagen	ITPKA	J-006742-11	Dharmacon	OLA1	SI04374643	Qiagen
AGAP3	J-019127-10	Dharmacon	DPP6	SI02627324	Qiagen	ITPKA	SI00034822	Qiagen	OPRK1	J-005684-06	Dharmacon
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AGAP3	SI00343504	Qiagen	DYPSL3	J-009821-09	Dharmacon	ITPKA	SI04378796	Qiagen	OPRK1	J-005684-09	Dharmacon
AGAP3	SI04205152	Qiagen	DYPSL3	J-009821-10	Dharmacon	ITPR1	J-006207-05	Dharmacon	OPRK1	SI00018347	Qiagen
AGAP3	SI04300352	Qiagen	DYPSL3	J-009821-11	Dharmacon	ITPR1	J-006207-06	Dharmacon	OPRK1	SI03074162	Qiagen
AGAP3	SI04358599	Qiagen	DYPSL3	J-009821-12	Dharmacon	ITPR1	J-006207-07	Dharmacon	OPRK1	SI04897319	Qiagen
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AP1S1	J-011241-10	Dharmacon	DYPSL3	SI04210549	Qiagen	ITPR1	SI00034531	Qiagen	PACSN1	J-007735-05	Dharmacon
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AP1S1	SI00024738	Qiagen	DRD2	J-005478-05	Dharmacon	ITPR1	SI03063607	Qiagen	PACSN1	J-007735-08	Dharmacon
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AQP4	J-011578-06	Dharmacon	DRD2	SI00015813	Qiagen	KCNAB1	J-008831-09	Dharmacon	PACSN1	SI04901225	Qiagen
AQP4	J-011578-07	Dharmacon	DRD2	SI03060876	Qiagen	KCNAB1	SI00157360	Qiagen	PAK1	J-003521-09	Dharmacon
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AQP4	J-011578-09	Dharmacon	DRD2	SI04895681	Qiagen	KCNAB1	SI03029047	Qiagen	PAK1	J-003521-11	Dharmacon
AQP4	SI00026831	Qiagen	DTNA	SI03024182	Qiagen	KCNAB1	SI03050971	Qiagen	PAK1	J-003521-12	Dharmacon
AQP4	SI00026838	Qiagen	DTNA	SI03037629	Qiagen	KCN2D	J-006225-05	Dharmacon	PAK1	SI00039774	Qiagen
AQP4	SI03040324	Qiagen	DTNA	SI03045497	Qiagen	KCN2D	J-006225-06	Dharmacon	PAK1	SI00039781	Qiagen
AQP4	SI03068961	Qiagen	DTNA	SI03048262	Qiagen	KCN2D	J-006225-07	Dharmacon	PAK1	SI06056969	Qiagen
ARHGAP20	J-026514-09	Dharmacon	DTNA	SI03056263	Qiagen	KCN2D	J-006225-08	Dharmacon	PAK1	SI04713520	Qiagen
ARHGAP20	J-026514-10	Dharmacon	DTNA	SI03081841	Qiagen	KCN2D	SI00098889	Qiagen	PAK2	J-003597-10	Dharmacon
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ARHGAP20	J-026514-12	Dharmacon	DTNA	SI04382070	Qiagen	KCN2D	SI03033534	Qiagen	PAK2	J-003597-12	Dharmacon
ARHGAP20	SI04259752	Qiagen	EFDH1	J-010673-17	Dharmacon	KCN2D	SI03046722	Qiagen	PAK2	J-003597-13	Dharmacon
ARHGAP20	SI04265632	Qiagen	EFDH1	J-010673-18	Dharmacon	KCNMA1	J-006267-06	Dharmacon	PAK2	SI0301077	Qiagen
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ARHGAP5	J-009580-05	Dharmacon	EFDH1	SI0376621	Qiagen	KCNMA1	J-006267-09	Dharmacon	PAK2	SI04897431	Qiagen
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ARHGAP5	SI00302519	Qiagen	EGR1	J-006526-07	Dharmacon	KCNQ2	J-006272-05	Dharmacon	PAK3	SI00039837	Qiagen
ARHGAP5	SI03229492	Qiagen	EGR1	J-006526-08	Dharmacon	KCNQ2	J-006272-07	Dharmacon	PAK3	SI00039844	Qiagen
ARHGAP5	SI04245759	Qiagen	EGR1	J-006526-09	Dharmacon	KCNQ2	J-006272-08	Dharmacon	PAK3	SI00039851	Qiagen
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ARPP19	J-015338-05	Dharmacon	EGR1	SI03052511	Qiagen	KCNQ2	SI00157297	Qiagen	PAPLN	J-030307-05	Dharmacon
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ARPP19	SI00090447	Qiagen	EGR4	J-006529-08	Dharmacon	KCNQ5	J-006275-06	Dharmacon	PAPLN	SI04202541	Qiagen
ARPP19	SI00090454	Qiagen	EGR4	SI03376999	Qiagen	KCNQ5	J-006275-07	Dharmacon	PAPLN	SI04270903	Qiagen
ARPP19	SI00090461	Qiagen	EGR4	SI03377006	Qiagen	KCNQ5	J-006275-08	Dharmacon	PAPLN	SI04372592	Qiagen
ATP2A2	J-004082-05	Dharmacon	EGR4	SI04318644	Qiagen	KCNQ5	SI00124383	Qiagen	PCMT1	J-010000-05	Dharmacon
ATP2A2	J-004082-06	Dharmacon	EGR4	SI04323669	Qiagen	KCNQ5	SI00124390	Qiagen	PCMT1	J-010000-06	Dharmacon
ATP2A2	J-004082-07	Dharmacon	EGR4	SI04318615	Qiagen	KCNQ5	SI00124397	Qiagen	PCMT1	J-010000-07	Dharmacon
ATP2A2	J-004082-08	Dharmacon	EGRB2IP	J-031861-06	Dharmacon	KCNQ5	SI03076577	Qiagen	PCMT1	J-010000-08	Dharmacon
ATP2A2	SI02626673	Qiagen	EGRB2IP	J-031861-07	Dharmacon	KCNQ5	SI03666586	Qiagen	PCMT1	SI00076398	Qiagen
ATP2A2	SI02626680	Qiagen	EGRB2IP	J-031861-08	Dharmacon	LAMP2	J-011715-05	Dharmacon	PCMT1	SI00076405	Qiagen
ATP2A2	SI03053337	Qiagen	EGRB2IP	SI02625042	Qiagen	LAMP2	J-011715-06	Dharmacon	PCMT1	SI00076412	Qiagen
ATP2A2	SI03062710	Qiagen	EGRB2IP	SI02778188	Qiagen	LAMP2	J-011715-07	Dharmacon	PCMT1	SI03074743	Qiagen
ATP6V1A	J-017590-09	Dharmacon	EGRB2IP	SI02778195	Qiagen	LAMP2	J-011715-08	Dharmacon	PDE7B	J-007659-06	Dharmacon
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ATP6V1A	J-017590-11	Dharmacon	FAM126B	J-018148-17	Dharmacon	LAMP2	SI03038819	Qiagen	PDE7B	J-007659-08	Dharmacon
ATP6V1A	J-017590-12	Dharmacon	FAM126B	J-018148-18	Dharmacon	LAMP2	SI03049949	Qiagen	PDE7B	J-007659-09	Dharmacon
ATP6V1A	SI00307489	Qiagen	FAM126B	J-018148-19	Dharmacon	LAMP2	SI03065944	Qiagen	PDE7B	SI0123249	Qiagen
ATP6V1A	SI00307496	Qiagen	FAM126B	J-018148-20	Dharmacon	LATS2	J-003865-09	Dharmacon	PDE7B	SI0123256	Qiagen
ATP6V1A	SI00307503	Qiagen	FAM126B	SI00640612	Qiagen	LATS2	J-003865-10	Dharmacon	PDE7B	SI0123263	Qiagen
ATP6V1A	SI00307510	Qiagen	FAM126B	SI04178839	Qiagen	LATS2	J-003865-11	Dharmacon	PDE7B	SI0123270	Qiagen
ATP6V1A	SI04247838	Qiagen	FAM1								

Gene Symbol	siRNA	Source									
B4GALT6	Si03150259	Qiagen	GABRD	J-006171-08	Dharmacon	LIX1	Si04318069	Qiagen	PLK2	J-003325-20	Dharmacon
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B4GALT6	Si04238962	Qiagen	GABRD	Si00016240	Qiagen	LPP	J-020012-05	Dharmacon	PLK2	Si0090363	Qiagen
B4GALT6	Si04330508	Qiagen	GABRD	Si00016247	Qiagen	LPP	J-020012-06	Dharmacon	PLK2	Si02288386	Qiagen
BMP2K	J-005071-05	Dharmacon	GABRD	Si00016254	Qiagen	LPP	J-020012-07	Dharmacon	PLK2	Si04438770	Qiagen
BMP2K	J-005071-06	Dharmacon	GDAP1	J-021225-05	Dharmacon	LPP	J-020012-08	Dharmacon	PLK2	Si04438777	Qiagen
BMP2K	J-005071-07	Dharmacon	GDAP1	J-021225-06	Dharmacon	Si00623119	Qiagen	PPP1R16B	J-004065-05	Dharmacon	
BMP2K	J-005071-08	Dharmacon	GDAP1	J-021225-07	Dharmacon	Si03687439	Qiagen	PPP1R16B	J-004065-06	Dharmacon	
BMP2K	Si02224803	Qiagen	GDAP1	J-021225-08	Dharmacon	Si04173659	Qiagen	PPP1R16B	J-004065-07	Dharmacon	
BMP2K	Si02224810	Qiagen	GDAP1	Si00123445	Qiagen	Si04264022	Qiagen	PPP1R16B	J-004065-08	Dharmacon	
BMP2K	Si04440051	Qiagen	GDAP1	Si00123466	Qiagen	Si04300114	Qiagen	PPP1R16B	Si00111447	Qiagen	
BMP2K	Si04901757	Qiagen	GDAP1	Si02777831	Qiagen	Si04371129	Qiagen	PPP1R16B	Si00111461	Qiagen	
CA11	J-009336-09	Dharmacon	GDAP1	Si02777838	Qiagen	LRIG1	J-013940-05	Dharmacon	PPP1R16B	Si00111468	Qiagen
CA11	J-009336-10	Dharmacon	GLRB	J-006183-05	Dharmacon	LRIG1	J-013940-06	Dharmacon	Si0315469	Qiagen	
CA11	J-009336-11	Dharmacon	GLRB	J-006183-06	Dharmacon	LRIG1	J-013940-07	Dharmacon	PPP3CA	J-008300-06	Dharmacon
CA11	J-009336-12	Dharmacon	GLRB	J-006183-07	Dharmacon	LRIG1	J-013940-08	Dharmacon	PPP3CA	J-008300-07	Dharmacon
CA11	Si00023548	Qiagen	GLRB	J-006183-08	Dharmacon	LRIG1	Si04316767	Qiagen	PPP3CA	J-008300-08	Dharmacon
CA11	Si00023555	Qiagen	GLRB	Si00016436	Qiagen	LRIG1	Si04341022	Qiagen	PPP3CA	Si0018942	Qiagen
CA11	Si03047877	Qiagen	GLRB	Si00016443	Qiagen	LRIG1	Si04363667	Qiagen	PPP3CA	Si02624636	Qiagen
CA11	Si03062206	Qiagen	GLRB	Si00016450	Qiagen	LTBP3	J-014144-05	Dharmacon	PPP3CA	Si04436572	Qiagen
CA12	J-003634-06	Dharmacon	GNB5	J-017235-05	Dharmacon	LTBP3	J-014144-06	Dharmacon	PREPL	J-005904-05	Dharmacon
CA12	J-003634-07	Dharmacon	GNB5	J-017235-06	Dharmacon	LTBP3	J-014144-07	Dharmacon	PREPL	J-005904-06	Dharmacon
CA12	J-003634-08	Dharmacon	GNB5	J-017235-07	Dharmacon	LTBP3	J-014144-08	Dharmacon	PREPL	J-005904-07	Dharmacon
CA12	J-003634-09	Dharmacon	GNB5	J-017235-08	Dharmacon	LTBP3	Si00625121	Qiagen	PREPL	J-005904-08	Dharmacon
CA12	Si00023583	Qiagen	GNB5	Si02637607	Qiagen	LTBP3	Si04204613	Qiagen	PRKCB	J-003758-13	Dharmacon
CA12	Si03041913	Qiagen	GNB5	Si03025876	Qiagen	LTBP3	Si04274158	Qiagen	PRKCB	J-003758-14	Dharmacon
CA12	Si03106306	Qiagen	GNB5	Si03049242	Qiagen	LTBP3	Si04285505	Qiagen	PRKCB	J-003758-15	Dharmacon
CA12	Si03115273	Qiagen	GNB5	Si03073973	Qiagen	MAFF	J-003903-05	Dharmacon	PRKCB	J-003758-16	Dharmacon
CACNA1B	J-006122-07	Dharmacon	GPR155	J-005494-05	Dharmacon	MAFF	J-003903-06	Dharmacon	PRKCB	Si0305948	Qiagen
CACNA1B	J-006122-08	Dharmacon	GPR155	J-005494-06	Dharmacon	MAFF	J-003903-07	Dharmacon	PRKCB	Si03092915	Qiagen
CACNA1B	J-006122-09	Dharmacon	GPR155	J-005494-08	Dharmacon	MAFF	J-003903-08	Dharmacon	PRKCB	Si03043747	Qiagen
CACNA1B	J-006122-10	Dharmacon	GPR155	Si04903682	Qiagen	MAFF	Si02662058	Qiagen	PRKCB	J-003758-17	Dharmacon
CACNA1B	Si00337141	Qiagen	GPR155	Si04903686	Qiagen	MAFF	Si03026611	Qiagen	PRKCB	J-003758-18	Dharmacon
CACNA1B	Si00337148	Qiagen	GPR155	Si04903689	Qiagen	MAFF	Si03067372	Qiagen	PRKCB	Si0042273	Qiagen
CACNA1B	Si00337155	Qiagen	GPR155	Si04903696	Qiagen	MAFF	Si03116456	Qiagen	PRKCB	Si0605948	Qiagen
CACNA1B	Si00337162	Qiagen	GPR155	Si04903703	Qiagen	MAFK	J-008580-05	Dharmacon	PRKCB	Si03036278	Qiagen
CACNB4	J-009062-05	Dharmacon	GRIA3	J-006186-07	Dharmacon	MAFK	J-008580-06	Dharmacon	PRKF40A	Si0313674	Qiagen
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CACNB4	J-009062-08	Dharmacon	GRIA3	J-006186-10	Dharmacon	MAN1A1	J-012174-05	Dharmacon	PSD3	Si04131295	Qiagen
CACNB4	Si00014294	Qiagen	GRIA3	Si00016541	Qiagen	MAN1A1	J-012174-06	Dharmacon	PSD3	Si04195604	Qiagen
CACNB4	Si00014308	Qiagen	GRIA3	Si03043642	Qiagen	MAN1A1	J-012174-07	Dharmacon	PSD3	Si04289740	Qiagen
CACNB4	Si00014315	Qiagen	GRIA3	Si03046967	Qiagen	MAN1A1	J-012174-08	Dharmacon	PTBP1	J-003528-06	Dharmacon
CACNB4	Si03113179	Qiagen	GRIA3	Si03111024	Qiagen	MAN1A1	Si04172826	Qiagen	PTBP1	J-003528-07	Dharmacon
CALB1	J-011989-05	Dharmacon	HIPK2	J-003266-10	Dharmacon	MAN1A1	Si04273689	Qiagen	PTBP1	J-003528-08	Dharmacon
CALB1	J-011989-06	Dharmacon	HIPK2	J-003266-11	Dharmacon	MAN1A1	Si04341316	Qiagen	PTBP1	J-003528-09	Dharmacon
CALB1	J-011989-07	Dharmacon	HIPK2	J-003266-12	Dharmacon	MAP4	J-013561-05	Dharmacon	PTBP1	Si0043631	Qiagen
CALB1	J-011989-08	Dharmacon	HIPK2	Si003266-13	Dharmacon	MAP4	Si04239214	Qiagen	PTBP1	Si0141638	Qiagen
CALB1	Si04132933	Qiagen	HIPK2	Si01343340	Qiagen	MAP4	Si04315731	Qiagen	PTBP1	Si02649206	Qiagen
CALB1	Si04308087	Qiagen	HIPK2	Si00134337	Qiagen	MAP4	Si04337053	Qiagen	PTPRO	J-008500-06	Dharmacon
CALB1	Si04329143	Qiagen	HIPK2	Si04439386	Qiagen	MDH1	J-009264-09	Dharmacon	PTPRO	J-008500-07	Dharmacon
CALB1	Si04362183	Qiagen	HIPK2	Si04439393	Qiagen	MDH1	J-009264-10	Dharmacon	PTPRO	Si03164140	Qiagen
CAP2	J-012211-05	Dharmacon	HS6ST2	J-015558-05	Dharmacon	MDH1	J-009264-11	Dharmacon	PTPRO	Si04136307	Qiagen
CAP2	J-012211-06	Dharmacon	HS6ST2	J-015558-06	Dharmacon	MDH1	J-009264-12	Dharmacon	PTPRO	J-008500-08	Dharmacon
CAP2	J-012211-07	Dharmacon	HS6ST2	J-015558-07	Dharmacon	MDH1	J-009264-13	Dharmacon	PTPRO	J-008500-09	Dharmacon
CAP2	J-012211-08	Dharmacon	HS6ST2	J-015558-08	Dharmacon	MDH1	J-009264-14	Dharmacon	PTPRO	J-008500-10	Dharmacon
CAP2	Si00337939	Qiagen	HS6ST2	Si04145414	Qiagen	MDH1	J-009264-15	Dharmacon	PTPRO	J-008500-11	Dharmacon
CAP2	Si04226663	Qiagen	HS6ST2	Si04269020	Qiagen	MDH1	J-009264-16	Dharmacon	PTPRO	J-008500-12	Dharmacon
CAP2	Si04235490	Qiagen	HS6ST2	Si04296208	Qiagen	MDH1	J-009264-17	Dharmacon	PTPRO	J-008500-13	Dharmacon
CAP2	Si04237331	Qiagen	HS6ST2	Si04335051	Qiagen	MDH1	J-009264-18	Dharmacon	PTPRO	J-008500-14	Dharmacon
CDHR1	J-013677-09	Dharmacon	HSPB1	J-005269-06	Dharmacon	MDH1	J-009265-06	Dharmacon	RAB3A	J-009668-07	Dharmacon
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CDHR1	J-013677-11	Dharmacon	HSPB1	J-005269-08	Dharmacon	MDH1	J-009265-08	Dharmacon	RAB3A	J-009668-09	Dharmacon
CDHR1	J-013677-12	Dharmacon	HSPB1	J-005269-09	Dharmacon	MDH1	J-009265-09	Dharmacon	RAB3A	J-009668-10	Dharmacon
CDHR1	Si00678867	Qiagen	HSPB1	Si02650585	Qiagen	MYLK	J-005351-18	Dharmacon	RAB3A	Si0044793	Qiagen
CDHR1	Si00678874	Qiagen	HSPB1	Si02650606	Qiagen	MYLK	J-005351-19	Dharmacon	RAB3A	Si0044800	Qiagen
CDHR1	Si00678881	Qiagen	HSPB1	Si04231129	Qiagen	MYLK	J-005351-20	Dharmacon	RAB3A	Si0044807	Qiagen
CDHR1	Si00678888	Qiagen	HSPB1	Si04354490	Qiagen	MYLK	J-005351-21	Dharmacon	RAB3A	Si0313411	Qiagen
CDK5R1	J-008988-05	Dharmacon	HSPB8	J-005006-05	Dharmacon	MYLK	Si0083202	Qiagen	RAB40B	J-008353-05	Dharmacon
CDK5R1	J-008988-06	Dharmacon	HSPB8	J-005006-06	Dharmacon	MYLK	Si0083209	Qiagen	RAB40B	J-008353-06	Dharmacon
CDK5R1	J-008988-07	Dharmacon	HSPB8	J-005006-07	Dharmacon	MYLK	Si02224040	Qiagen	RAB40B	J-008353-07	Dharmacon
CDK5R1	J-008988-08	Dharmacon	HSPB8	J-005006-08	Dharmacon	MYLK	Si03067960	Qiagen	RAB40B	J-008353-08	Dharmacon
CDK5R1	Si00057372	Qiagen	HSPB8	Si02224502	Qiagen	MYST3	J-019849-05	Dharmacon	RAB40B	Si04076357	Qiagen
CDK5R1	Si00057393	Qiagen	HSPB8	Si03021921	Qiagen	MYST3	J-019849-06	Dharmacon	RAB40B	Si02622212	Qiagen
CDK5R1	Si02659580	Qiagen	HSPB8	Si03100083	Qiagen	MYST3	J-019849-07	Dharmacon	RAB40B	Si02622772	Qiagen
CDK5R1	Si04379543	Qiagen	HSPB8	Si04900854	Qiagen	MYST3	J-019849-08	Dharmacon	RAB40B	Si03078600	Qiagen
CELF4	J-013824-18	Dharmacon	HSPH1	J-004972-05	Dharmacon	MYST3	Si00901714	Qiagen	RAB40B	Si04385304	Qiagen
CELF4	J-013824-19	Dharmacon	HSPH1	J-004972-06	Dharmacon	MYST3	Si03020836	Qiagen	RAC1	J-003560-14	Dharmacon
CELF4	J-013824-20	Dharmacon	HSPH1	J-004972-07	Dharmacon	MYST3	Si04388293	Qiagen	RAC1	J-003560-15	Dharmacon
CELF4	J-013824-21	Dharmacon	HSPH1	J-004972-08	Dharmacon	MYST3	Si04388300	Qiagen	RAC1	J-003560-16	Dharmacon
CELF4	Si00313621	Qiagen	HSPH1	Si01582821	Qiagen	MYCN	J-012835-18	Dharmacon	RAC1	J-003560-17	Dharmacon
CELF4	Si00313628	Qiagen	HSPH1	Si04159764	Qiagen	MYCN	J-012835-19	Dharmacon	RAC1	J-02655051	Qiagen
CELF4	Si04159463	Qiagen	HSPH1	Si04314905	Qiagen	MYCN	J-012835-20	Dharmacon	RAC1	Si03040884	Qiagen
CELF4	Si04162179	Qiagen	HSPH1	Si04367643	Qiagen	MYCN	J-012835-21	Dharmacon	RAC1	Si03065531	Qiagen
CHI3L1	J-012568-09	Dharmacon	HTR2C	J-005640-05	Dharmacon	MYCN	J-012835-22	Dharmacon	RAC2	J-007741-09	Dharmacon
CHI3L1	J-012568-11	Dharmacon	HTR2C	J-005640-07	Dharmacon	MYCN	J-012835-23	Dharmacon	RAC2	J-007741-10	Dharmacon
CHI3L1	J-012568-12	Dharmacon	HTR2C	J-005640-08	Dharmacon	MYCN	J-012835-24	Dharmacon	RAC2	J-007741-11	Dharmacon
CHI3L1	Si00345842	Qiagen	HTR2C	Si00173747	Qiagen	MYCN	J-012835-25	Dharmacon	RAC2	J-007741-12	Dharmacon
CHI3L1	Si03125227	Qiagen	HTR2C	Si03083164	Qiagen	NECAP1	J-017872-09	Dharmacon	RAC2	J-00836-08	Dharmacon
CHI3L1	Si04254187	Qiagen	HTR2C	Si04986927	Qiagen	NECAP1	J-017872-10	Dharmacon	RAC2	J-00836-09	Dharmacon
CLMN	J-017131-18	Dharmacon	IDS	J-009254-05	Dharmacon	NECAP1	J-017872				

Gene Symbol	siRNA	Source
CORO1C	SI03168578	Qiagen
CORO1C	SI04138631	Qiagen
CORO1C	SI04310481	Qiagen
CRISPLD1	J-016681-17	Dharmacon
CRISPLD1	J-016681-18	Dharmacon
CRISPLD1	J-016681-19	Dharmacon
CRISPLD1	J-016681-20	Dharmacon
CRISPLD1	SI04179049	Qiagen
CRISPLD1	SI04249886	Qiagen
CRISPLD1	SI04295774	Qiagen
CRISPLD1	SI04339370	Qiagen
CXCR7	J-013212-10	Dharmacon
CXCR7	J-013212-11	Dharmacon
CXCR7	J-013212-12	Dharmacon
CXCR7	J-013212-13	Dharmacon
CXCR7	SI00125790	Qiagen
CXCR7	SI02660644	Qiagen
CXCR7	SI04901925	Qiagen
CXCR7	SI04901932	Qiagen
DACH1	J-013222-05	Dharmacon
DACH1	J-013222-06	Dharmacon
DACH1	J-013222-07	Dharmacon
DACH1	J-013222-08	Dharmacon
DACH1	SI00359212	Qiagen
DACH1	SI03649107	Qiagen
DACH1	SI04175073	Qiagen
DACH1	SI04200553	Qiagen
DACH1	SI04238255	Qiagen
DDR1	J-003111-12	Dharmacon
DDR1	J-003111-13	Dharmacon
DDR1	J-003111-14	Dharmacon
DDR1	J-003111-15	Dharmacon

Gene Symbol	siRNA	Source
IKBKAP	SI00287462	Qiagen
IKBKAP	SI03045749	Qiagen
IKBKAP	SI04899181	Qiagen
IKBKAP	SI04899188	Qiagen
IKBKB	J-003503-11	Dharmacon
IKBKB	J-003503-12	Dharmacon
IKBKB	J-003503-13	Dharmacon
IKBKB	J-003503-14	Dharmacon
IKBKB	SI00300545	Qiagen
IKBKB	SI02626442	Qiagen
IKBKB	SI02626456	Qiagen
IKBKE	SI02777376	Qiagen
IKBKE	J-003723-09	Dharmacon
IKBKE	J-003723-10	Dharmacon
IKBKE	J-003723-11	Dharmacon

Gene Symbol	siRNA	Source
NELL2	J-012185-10	Dharmacon
NELL2	J-012185-11	Dharmacon
NELL2	J-012185-12	Dharmacon
NELL2	SI04167205	Qiagen
NELL2	SI04205565	Qiagen
NELL2	SI04292183	Qiagen
NELL2	SI04352901	Qiagen
NFIB	J-008456-05	Dharmacon
NFIB	J-008456-06	Dharmacon
NFIB	J-008456-07	Dharmacon
NFIB	J-008456-08	Dharmacon
NFIB	SI00079352	Qiagen
NFIB	SI00079359	Qiagen
NFIB	SI03032197	Qiagen
NFIB	SI03101141	Qiagen
NFIB	SI04300947	Qiagen
NFKB1	J-003520-06	Dharmacon
NFKB1	J-003520-07	Dharmacon
NFKB1	J-003520-08	Dharmacon
NFKB1	J-003520-09	Dharmacon
NFKB1	SI02654932	Qiagen
NFKB1	SI02654939	Qiagen
NFKB1	SI02662618	Qiagen
NFKB2	J-003918-05	Dharmacon
NFKB2	J-003918-06	Dharmacon
NFKB2	J-003918-07	Dharmacon
NFKB2	J-003918-08	Dharmacon
NFKB2	SI00300965	Qiagen
NFKB2	SI02650949	Qiagen
NFKB2	SI04219852	Qiagen
NFKB2	SI04224290	Qiagen
NFKB2	SI04245318	Qiagen

Gene Symbol	siRNA	Source
RERG	SI03068107	Qiagen
RGS14	J-008826-05	Dharmacon
RGS14	J-008826-06	Dharmacon
RGS14	J-008826-07	Dharmacon
RGS14	J-008826-08	Dharmacon
RGS14	J-022559-18	Dharmacon
RGS14	SI00088879	Qiagen
RGS14	SI00088886	Qiagen
RGS14	SI03054737	Qiagen
RGS14	SI03076367	Qiagen
ROCK1	J-003536-06	Dharmacon
ROCK1	J-003536-07	Dharmacon
ROCK1	J-003536-08	Dharmacon
ROCK1	J-003536-09	Dharmacon
ROCK1	SI02622095	Qiagen
ROCK1	SI02622102	Qiagen
ROCK1	SI03095484	Qiagen
ROCK1	SI04898362	Qiagen
ROCK2	J-004610-06	Dharmacon
ROCK2	J-004610-07	Dharmacon
ROCK2	J-004610-08	Dharmacon
ROCK2	J-004610-09	Dharmacon
ROCK2	SI02223753	Qiagen
ROCK2	SI04899720	Qiagen
ROCK2	SI04899727	Qiagen
TBK1	J-003788-08	Dharmacon
TBK1	J-003788-09	Dharmacon
TBK1	J-003788-10	Dharmacon
TBK1	J-003788-11	Dharmacon
TBK1	SI00100961	Qiagen
TBK1	SI02224411	Qiagen
TBK1	SI04379606	Qiagen
TBK1	SI04901169	Qiagen