

Target gene	Representa	Gene name	3P-seq tags	Conserved	Conserved	Conserved	Conserved	Poorly con
MEX3C	ENST00000	mex-3 RNA	5	1	1	0	0	2
SERINC4	ENST00000	serine inco	9	0	0	0	0	2
GREM1	ENST00000	gremlin 1, l	752	0	0	0	0	2
MS4A12	ENST00000	membrane	10	0	0	0	0	1
PANK2	ENST00000	pantothen	1244	1	1	0	0	0
IRF7	ENST00000	interferon	10	0	0	0	0	1
CLK1	ENST00000	CDC-like kin	3430	1	1	0	0	0
AC119673.	ENST00000	598338.1	5	0	0	0	0	1
CYP7B1	ENST00000	cytochrom	53	1	1	0	0	3
FBXL3	ENST00000	F-box and l	842	0	0	0	0	2
STXBP4	ENST00000	syntaxin bi	62	0	0	0	0	3
ME1	ENST00000	malic enzyr	586	0	0	0	0	1
KRTAP2-2	ENST00000	keratin asso	5	0	0	0	0	1
ZNF345	ENST00000	zinc finger	24	0	0	0	0	2
TRA2B	ENST00000	transforme	1868	1	1	0	0	0
CEP95	ENST00000	centrosom	46	0	0	0	0	1
ZMAT1	ENST00000	zinc finger,	5	1	0	1	0	1
NUDT4	ENST00000	nudix (nucl	41	0	0	0	0	2
SOCS5	ENST00000	suppressor	72	1	1	0	0	1
UCHL1	ENST00000	ubiquitin c	21879	0	0	0	0	1
UPP2	ENST00000	uridine phc	5	1	1	0	0	0
ATF1	ENST00000	activating t	1599	0	0	0	0	1
GABRA6	ENST00000	gamma-am	5	0	0	0	0	1
RAD54B	ENST00000	RAD54 hon	36	0	0	0	0	2
NKAIN3	ENST00000	Na <sup>+</sup> /K <sup>+</sup> tra	5	0	0	0	0	1
CEACAM5	ENST00000	carcinoemb	5	0	0	0	0	1
SLC26A2	ENST00000	solute carri	748	0	0	0	0	4
PCP4L1	ENST00000	Purkinje ce	23	0	0	0	0	3
CASP5	ENST00000	caspase 5,	5	0	0	0	0	1
NEUROD4	ENST00000	neuronal d	5	0	0	0	0	2
PAPOLG	ENST00000	poly(A) pol	49	1	0	0	1	2
RPE	ENST00000	ribulose-5-	333	0	0	0	0	2
HTR1F	ENST00000	5-hydroxyt	5	0	0	0	0	1
SIX1	ENST00000	SIX homeol	1264	1	1	0	0	0
PTPN12	ENST00000	protein tyr	167	0	0	0	0	1
FAM120AC	ENST00000	family with	258	0	0	0	0	1
AP005482.	ENST00000	Uncharacte	44	0	0	0	0	2
RAB22A	ENST00000	RAB22A, m	484	0	0	0	0	3
C9orf38	ENST00000	chromosom	5	0	0	0	0	1
C1orf95	ENST00000	chromosom	28	0	0	0	0	1
NIPAL2	ENST00000	NIPA-like d	7	1	1	0	0	1
PFN2	ENST00000	profilin 2	2184	1	1	0	0	0
KCNH8	ENST00000	potassium	5	0	0	0	0	2
COL25A1	ENST00000	collagen, ty	270	0	0	0	0	2
AQP4	ENST00000	aquaporin	10	0	0	0	0	1
FAM169A	ENST00000	family with	90	0	0	0	0	2

ZIK1	ENST00000 zinc finger	64	0	0	0	0	1
LIG4	ENST00000 ligase IV, D	113	0	0	0	0	1
ZNF24	ENST00000 zinc finger	1796	0	0	0	0	2
WFDC11	ENST00000 WAP four-c	5	0	0	0	0	1
AVP11	ENST00000 arginine va	4404	0	0	0	0	1
PGAM1	ENST00000 phosphogly	927	0	0	0	0	1
GFM2	ENST00000 G elongatic	19	0	0	0	0	1
RP11-382J1	ENST00000 Uncharacte	5	0	0	0	0	1
BTLA	ENST00000 B and T lym	5	0	0	0	0	1
RFESD	ENST00000 Rieske (Fe-	15	1	0	1	0	2
HMG2	ENST00000 high mobili	184	0	0	0	0	1
KDEL2	ENST00000 KDEL (Lys-/	667	0	0	0	0	2
LRRC37A3	ENST00000 leucine rich	5	0	0	0	0	1
MAP3K19	ENST00000 mitogen-ac	5	0	0	0	0	1
KLHL4	ENST00000 kelch-like f	58	0	0	0	0	2
DCTN4	ENST00000 dynactin 4	153	0	0	0	0	1
SMCO1	ENST00000 single-pass	5	0	0	0	0	1
LUC7L	ENST00000 LUC7-like (C	898	0	0	0	0	1
YTHDF3	ENST00000 YTH domain	444	1	1	0	0	1
AFF4	ENST00000 AF4/FMR2	2755	0	0	0	0	3
KIAA1217	ENST00000 KIAA1217	320	0	0	0	0	3
WNT6	ENST00000 wingless-ty	7	0	0	0	0	1
PPP2CB	ENST00000 protein pho	11	0	0	0	0	1
HSDL1	ENST00000 hydroxyste	699	0	0	0	0	1
HES7	ENST00000 hairy and e	416	0	0	0	0	1
CCDC67	ENST00000 coiled-coil	5	1	1	0	0	0
RYBP	ENST00000 RING1 and	342	1	0	1	0	2
CAPZA2	ENST00000 capping pro	2065	0	0	0	0	1
SLA	ENST00000 Src-like-ada	5	0	0	0	0	1
PGAM4	ENST00000 phosphogly	5	0	0	0	0	1
ADAM7	ENST00000 ADAM met	5	0	0	0	0	1
RERG	ENST00000 RAS-like, es	162	0	0	0	0	2
KRT38	ENST00000 keratin 38	5	0	0	0	0	1
CRK	ENST00000 v-crk avian	1841	1	0	1	0	1
ADIPOQ	ENST00000 adiponectin	5	0	0	0	0	1
ZNF772	ENST00000 zinc finger	22	0	0	0	0	1
URM1	ENST00000 ubiquitin re	345	0	0	0	0	2
KRTAP20-4	ENST00000 keratin asso	5	0	0	0	0	1
MTPN	ENST00000 myotrophin	1296	0	0	0	0	2
GLIPR1L2	ENST00000 GLI pathoge	5	0	0	0	0	1
AL354808.1	ENST00000601204.1	5	0	0	0	0	1
MLANA	ENST00000 melan-A	5	0	0	0	0	1
A2ML1	ENST00000 alpha-2-ma	5	0	0	0	0	1
HPDL	ENST00000 4-hydroxyp	418	0	0	0	0	1
APOL6	ENST00000 apolipopro	5	0	0	0	0	2
IKZF3	ENST00000 IKAROS fan	7	0	0	0	0	2
IL26	ENST00000 interleukin	5	0	0	0	0	1

ANKRD44	ENST00000C ankyrin rep	187	0	0	0	0	1
ZNF799	ENST00000C zinc finger	17	0	0	0	0	3
NAMPT	ENST00000C nicotinamir	375	0	0	0	0	1
SULT4A1	ENST00000C sulfotransfe	25	0	0	0	0	2
RBM7	ENST00000C RNA bindin	46	0	0	0	0	1
PANX3	ENST00000C pannexin 3	5	0	0	0	0	1
C6orf106	ENST00000C chromosom	80	1	1	0	0	0

Poorly con:	Poorly con:	Poorly con:	6mer sites	Representa	Cumulative	Total	conte	Aggregate	Previous Ta
0	1	1	1	hsa-miR-15	-0.73	-0.73	N/A		
2	0	0	0	hsa-miR-15	-0.72	-0.72	N/A		
1	0	1	0	hsa-miR-15	-0.71	-0.72	N/A		
1	0	0	0	hsa-miR-15	-0.71	-0.71	N/A		
0	0	0	0	hsa-miR-15	-0.7	-0.7	N/A		
1	0	0	0	hsa-miR-15	-0.67	-0.67	N/A		
0	0	0	0	hsa-miR-15	-0.64	-0.64	N/A		2007
1	0	0	0	hsa-miR-15	-0.64	-0.64	N/A		
0	1	2	0	hsa-miR-15	-0.64	-0.68	N/A		
1	1	0	1	hsa-miR-15	-0.62	-0.7	N/A		
1	0	2	1	hsa-miR-15	-0.62	-0.72	N/A		
1	0	0	0	hsa-miR-15	-0.62	-0.62	N/A		
1	0	0	0	hsa-miR-15	-0.61	-0.61	N/A		
1	0	1	0	hsa-miR-15	-0.61	-0.61	N/A		
0	0	0	0	hsa-miR-15	-0.59	-0.59	N/A		
1	0	0	0	hsa-miR-15	-0.59	-0.59	N/A		
0	1	0	2	hsa-miR-15	-0.57	-0.57	N/A		
1	1	0	1	hsa-miR-15	-0.57	-0.57	N/A		
1	0	0	1	hsa-miR-15	-0.57	-0.57	N/A		
1	0	0	0	hsa-miR-15	-0.56	-0.56	N/A		
0	0	0	0	hsa-miR-15	-0.56	-0.56	N/A		2007
1	0	0	2	hsa-miR-15	-0.55	-0.64	N/A		
1	0	0	0	hsa-miR-15	-0.54	-0.54	N/A		
1	0	1	0	hsa-miR-15	-0.54	-0.59	N/A		
1	0	0	0	hsa-miR-15	-0.53	-0.53	N/A		
1	0	0	0	hsa-miR-15	-0.53	-0.53	N/A		
0	1	3	2	hsa-miR-15	-0.53	-0.53	N/A		
0	0	3	0	hsa-miR-15	-0.53	-0.53	N/A		
1	0	0	0	hsa-miR-15	-0.53	-0.53	N/A		
1	0	1	0	hsa-miR-15	-0.52	-0.52	N/A		
1	0	1	1	hsa-miR-15	-0.51	-0.59	N/A		
1	0	1	0	hsa-miR-15	-0.5	-0.55	N/A		
1	0	0	1	hsa-miR-15	-0.5	-0.5	N/A		
0	0	0	0	hsa-miR-15	-0.5	-0.5	N/A		
1	0	0	0	hsa-miR-15	-0.5	-0.5	N/A		
1	0	0	1	hsa-miR-15	-0.49	-0.49	N/A		
0	2	0	0	hsa-miR-15	-0.49	-0.59	N/A		
1	0	2	1	hsa-miR-15	-0.48	-0.86	N/A		
1	0	0	0	hsa-miR-15	-0.48	-0.48	N/A		
1	0	0	0	hsa-miR-15	-0.48	-0.48	N/A		
0	0	1	1	hsa-miR-15	-0.48	-0.48	N/A		
0	0	0	0	hsa-miR-15	-0.48	-0.5	N/A		2007
1	1	0	0	hsa-miR-15	-0.47	-0.47	N/A		
2	0	0	0	hsa-miR-15	-0.47	-0.57	N/A		
1	0	0	1	hsa-miR-15	-0.47	-0.47	N/A		
0	2	0	2	hsa-miR-15	-0.47	-0.51	N/A		

1	0	0	1 hsa-miR-15	-0.47	-0.47 N/A	
1	0	0	0 hsa-miR-15	-0.47	-0.47 N/A	
1	0	1	0 hsa-miR-15	-0.47	-0.61 N/A	
0	1	0	0 hsa-miR-15	-0.47	-0.47 N/A	
1	0	0	0 hsa-miR-15	-0.47	-0.47 N/A	
1	0	0	0 hsa-miR-15	-0.47	-0.47 N/A	
1	0	0	0 hsa-miR-15	-0.47	-0.66 N/A	
1	0	0	0 hsa-miR-15	-0.47	-0.47 N/A	
1	0	0	0 hsa-miR-15	-0.47	-0.47 N/A	
0	1	1	0 hsa-miR-15	-0.46	-0.82 N/A	
1	0	0	0 hsa-miR-15	-0.46	-0.49 N/A	
1	0	1	0 hsa-miR-15	-0.46	-0.48 N/A	
1	0	0	0 hsa-miR-15	-0.46	-0.46 N/A	
1	0	0	0 hsa-miR-15	-0.46	-0.46 N/A	
0	2	0	1 hsa-miR-15	-0.45	-0.45 N/A	
1	0	0	0 hsa-miR-15	-0.45	-0.45 N/A	
1	0	0	0 hsa-miR-15	-0.45	-0.45 N/A	
1	0	0	0 hsa-miR-15	-0.45	-0.51 N/A	
0	0	1	1 hsa-miR-15	-0.45	-0.45 N/A	2007
0	0	3	0 hsa-miR-15	-0.44	-0.46 N/A	
0	2	1	0 hsa-miR-15	-0.44	-0.47 N/A	
1	0	0	0 hsa-miR-15	-0.44	-0.44 N/A	
1	0	0	0 hsa-miR-15	-0.44	-0.44 N/A	
1	0	0	0 hsa-miR-15	-0.43	-0.43 N/A	
1	0	0	0 hsa-miR-15	-0.43	-0.46 N/A	
0	0	0	0 hsa-miR-15	-0.42	-0.42 N/A	
0	2	0	0 hsa-miR-15	-0.42	-0.58 N/A	2007
1	0	0	0 hsa-miR-15	-0.42	-0.47 N/A	
1	0	0	0 hsa-miR-15	-0.42	-0.42 N/A	
1	0	0	0 hsa-miR-15	-0.42	-0.42 N/A	
1	0	0	0 hsa-miR-15	-0.42	-0.42 N/A	
0	1	1	1 hsa-miR-15	-0.41	-0.45 N/A	
1	0	0	0 hsa-miR-15	-0.4	-0.4 N/A	
0	1	0	0 hsa-miR-15	-0.4	-0.4 N/A	2007
1	0	0	1 hsa-miR-15	-0.4	-0.4 N/A	
1	0	0	1 hsa-miR-15	-0.4	-0.54 N/A	
1	0	1	0 hsa-miR-15	-0.4	-0.6 N/A	
0	1	0	0 hsa-miR-15	-0.4	-0.4 N/A	
0	1	1	0 hsa-miR-15	-0.4	-0.4 N/A	
1	0	0	0 hsa-miR-15	-0.4	-0.4 N/A	
0	1	0	0 hsa-miR-15	-0.39	-0.39 N/A	
1	0	0	0 hsa-miR-15	-0.39	-0.39 N/A	
1	0	0	0 hsa-miR-15	-0.39	-0.39 N/A	
0	1	0	0 hsa-miR-15	-0.39	-0.39 N/A	
0	0	2	1 hsa-miR-15	-0.38	-0.38 N/A	
1	1	0	0 hsa-miR-15	-0.38	-0.46 N/A	
0	1	0	0 hsa-miR-15	-0.38	-0.38 N/A	

1	0	0	0 hsa-miR-15	-0.38	-0.38 N/A	
0	0	3	1 hsa-miR-15	-0.38	-0.54 N/A	
1	0	0	1 hsa-miR-15	-0.38	-0.44 N/A	
0	1	1	0 hsa-miR-15	-0.38	-0.4 N/A	
1	0	0	0 hsa-miR-15	-0.38	-0.38 N/A	
0	1	0	1 hsa-miR-15	-0.37	-0.37 N/A	
0	0	0	0 hsa-miR-15	-0.37	-0.37 N/A	2007

rgetScan publication(s)