

HOXD-AS1	HOXD-AS1	HOXD-AS1	chr2:1770379 shneg	shCHD1	OK	0,867928	0,23625	-1,87726	-3,68947	5,00E-05	0,0004193	yes
LINC00478	LINC00478	LINC00478	chr21:174428 shneg	shCHD1	OK	4,65674	1,25593	-1,89057	-6,97149	5,00E-05	0,0004193	yes
C19orf77	C19orf77	C19orf77	chr19:347440 shneg	shCHD1	OK	4,50397	1,21045	-1,89565	-3,83722	5,00E-05	0,0004193	yes
ABCA1	ABCA1	ABCA1	chr9:1075432 shneg	shCHD1	OK	1,60729	0,429173	-1,905	-7,29114	5,00E-05	0,0004193	yes
LOC10013041	LOC10013041	LOC10013041	chr1:852952-1 shneg	shCHD1	OK	3,86921	1,03226	-1,90623	-2,10601	0,0034	0,0162243	yes
PADI3	PADI3	PADI3	chr1:1757559 shneg	shCHD1	OK	52,5499	13,9552	-1,91288	-10,3732	5,00E-05	0,0004193	yes
LILRB1	LILRB1	LILRB1	chr19:551286 shneg	shCHD1	OK	6,06691	1,60351	-1,91973	-7,25182	5,00E-05	0,0004193	yes
PIP5K1L	PIP5K1L	PIP5K1L	chr9:1306838 shneg	shCHD1	OK	8,44134	2,22836	-1,92149	-5,7099	5,00E-05	0,0004193	yes
COL9A3	COL9A3	COL9A3	chr20:614484 shneg	shCHD1	OK	30,3258	7,89154	-1,94216	-6,80619	5,00E-05	0,0004193	yes
MCHR1	MCHR1	MCHR1	chr22:410751 shneg	shCHD1	OK	1,04053	0,264599	-1,97543	-3,26856	5,00E-05	0,0004193	yes
FAT4	FAT4	FAT4	chr4:1262375 shneg	shCHD1	OK	0,869511	0,217808	-1,99715	-7,2862	5,00E-05	0,0004193	yes
TMEM88B	TMEM88B	TMEM88B	chr1:1361507 shneg	shCHD1	OK	17,7548	4,43279	-2,00192	-4,87937	5,00E-05	0,0004193	yes
CXorf57	CXorf57	CXorf57	chrX:1058551 shneg	shCHD1	OK	0,804038	0,199909	-2,00792	-3,54667	5,00E-05	0,0004193	yes
GPR56	GPR56	GPR56	chr16:576539 shneg	shCHD1	OK	4,01344	0,997338	-2,00869	-7,05613	5,00E-05	0,0004193	yes
PLCB4	PLCB4	PLCB4	chr20:904970 shneg	shCHD1	OK	7,87406	1,95479	-2,0101	-9,64815	5,00E-05	0,0004193	yes
SMARCA2	SMARCA2	SMARCA2	chr9:2015341 shneg	shCHD1	OK	2,33899	0,578857	-2,01461	-7,09627	5,00E-05	0,0004193	yes
GLI1	GLI1	GLI1	chr12:578539 shneg	shCHD1	OK	27,1843	6,67142	-2,02671	-10,2725	5,00E-05	0,0004193	yes
NIPSNAP3B	NIPSNAP3B	NIPSNAP3B	chr9:1075264 shneg	shCHD1	OK	0,903187	0,221319	-2,0289	-2,40524	0,0033	0,0158557	yes
PRPH	PRPH	PRPH	chr12:496889 shneg	shCHD1	OK	2,0501	0,472387	-2,11765	-4,13866	5,00E-05	0,0004193	yes
HOXD-AS2	HOXD-AS2	HOXD-AS2	chr2:1769995 shneg	shCHD1	OK	4,8129	1,10816	-2,11874	-3,17766	5,00E-05	0,0004193	yes
KRT20	KRT20	KRT20	chr17:390321 shneg	shCHD1	OK	6,51106	1,45311	-2,16375	-6,70073	5,00E-05	0,0004193	yes
SSBP2	SSBP2	SSBP2	chr5:8025650 shneg	shCHD1	OK	1,09734	0,242936	-2,17536	-2,1259	0,00045	0,00293585	yes
DMRT2	DMRT2	DMRT2	chr9:1050345 shneg	shCHD1	OK	1,01939	0,222106	-2,19839	-2,93162	5,00E-05	0,0004193	yes
KLF8	KLF8	KLF8	chrX:5625882 shneg	shCHD1	OK	1,19421	0,258459	-2,20805	-5,38582	5,00E-05	0,0004193	yes
CACNA1A	CACNA1A	CACNA1A	chr19:133172 shneg	shCHD1	OK	19,333	4,11771	-2,23116	-11,9388	5,00E-05	0,0004193	yes
JSRP1	JSRP1	JSRP1	chr19:225234 shneg	shCHD1	OK	1,67328	0,355693	-2,23398	-2,59277	0,00185	0,00975716	yes
C9orf125	C9orf125	C9orf125	chr9:1042376 shneg	shCHD1	OK	6,8899	1,46257	-2,23598	-7,57731	5,00E-05	0,0004193	yes
ADAMTS15	ADAMTS15	ADAMTS15	chr11:130318 shneg	shCHD1	OK	14,3224	2,89769	-2,3053	-11,8322	5,00E-05	0,0004193	yes
AK5	AK5	AK5	chr1:7774766 shneg	shCHD1	OK	8,51097	1,71913	-2,30764	-9,47418	5,00E-05	0,0004193	yes
STARD13	STARD13	STARD13	chr13:336772 shneg	shCHD1	OK	0,821538	0,163958	-2,325	-3,23416	5,00E-05	0,0004193	yes
HOXD8	HOXD8	HOXD8	chr2:1769944 shneg	shCHD1	OK	2,86972	0,563235	-2,3491	-4,56564	5,00E-05	0,0004193	yes
BAHCC1	BAHCC1	BAHCC1	chr17:793735 shneg	shCHD1	OK	2,15779	0,401016	-2,42782	-9,06946	5,00E-05	0,0004193	yes
NEUROG2	NEUROG2	NEUROG2	chr4:1134346 shneg	shCHD1	OK	19,0302	3,4787	-2,45167	-10,9818	5,00E-05	0,0004193	yes
SCG5	SCG5	SCG5	chr15:329338 shneg	shCHD1	OK	7,152	1,29694	-2,46324	-6,26403	5,00E-05	0,0004193	yes
NES	NES	NES	chr1:1566385 shneg	shCHD1	OK	17,5164	3,15265	-2,47407	-12,6063	5,00E-05	0,0004193	yes
ZNF462	ZNF462	ZNF462	chr9:1096253 shneg	shCHD1	OK	3,13746	0,532536	-2,55865	-9,9039	5,00E-05	0,0004193	yes
FAM46C	FAM46C	FAM46C	chr1:1181486 shneg	shCHD1	OK	0,987062	0,167471	-2,55923	-5,62262	5,00E-05	0,0004193	yes
SCEL	SCEL	SCEL	chr13:781098 shneg	shCHD1	OK	8,77028	1,48701	-2,56021	-9,88864	5,00E-05	0,0004193	yes
H19	H19	H19	chr11:201640 shneg	shCHD1	OK	39,17	6,49447	-2,59247	-12,9506	5,00E-05	0,0004193	yes
SFRP5	SFRP5	SFRP5	chr10:995265 shneg	shCHD1	OK	0,749249	0,123031	-2,60643	-2,54274	0,01265	0,0482745	yes
IFI27	IFI27	IFI27	chr14:945770 shneg	shCHD1	OK	120,055	19,46	-2,62511	-12,3472	5,00E-05	0,0004193	yes
RNF128	RNF128	RNF128	chrX:1059370 shneg	shCHD1	OK	20,9839	3,2436	-2,69362	-12,6553	5,00E-05	0,0004193	yes
CXXC11	CXXC11	CXXC11	chr2:2428118 shneg	shCHD1	OK	0,75147	0,105819	-2,82812	-2,32771	0,0088	0,0358932	yes
SAMD11	SAMD11	SAMD11	chr1:861120-1 shneg	shCHD1	OK	4,80576	0,64728	-2,8923	-2,3725	0,0034	0,0162243	yes
VIT	VIT	VIT	chr2:3692383 shneg	shCHD1	OK	14,768	1,83891	-3,00555	-11,1988	5,00E-05	0,0004193	yes
PLCH2	PLCH2	PLCH2	chr1:2407753 shneg	shCHD1	OK	30,583	3,68911	-3,05139	-15,4903	5,00E-05	0,0004193	yes
LILRA2	LILRA2	LILRA2	chr19:550852 shneg	shCHD1	OK	27,7526	3,30164	-3,07136	-12,5892	5,00E-05	0,0004193	yes
NOTUM	NOTUM	NOTUM	chr17:799103 shneg	shCHD1	OK	169,935	19,9807	-3,08831	-16,927	5,00E-05	0,0004193	yes
KSR2	KSR2	KSR2	chr12:117890 shneg	shCHD1	OK	1,99548	0,232059	-3,10417	-11,8184	5,00E-05	0,0004193	yes
ACY3	ACY3	ACY3	chr11:674100 shneg	shCHD1	OK	1,37624	0,152754	-3,17144	-2,53222	0,0058	0,0254211	yes
FLRT3	FLRT3	FLRT3	chr20:139761 shneg	shCHD1	OK	1,11806	0,12048	-3,21413	-4,26459	5,00E-05	0,0004193	yes
GLI3	GLI3	GLI3	chr7:4200054 shneg	shCHD1	OK	1,26182	0,127218	-3,31013	-7,68974	5,00E-05	0,0004193	yes
NRP2	NRP2	NRP2	chr2:2065472 shneg	shCHD1	OK	2,13926	0,192695	-3,47272	-5,1153	5,00E-05	0,0004193	yes
LOC375295	LOC375295	LOC375295	chr2:1774943 shneg	shCHD1	OK	6,92168	0,419817	-4,04329	-6,27169	5,00E-05	0,0004193	yes