Jump to: <u>Multiple Alignment Motif Tree Motif Matching</u> <u>Input file: 509 motifs loaded</u>

Settings: Metric=PCC, Alignment=SWU, Gap-open=1000, Gap-extend=1000, -nooverlapalign
Multiple Alignment=IR, Tree=UPGMA, Matching against: JASPAR
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Multiple Alignment

(Consensus sequence representations shown, but multiple alignment was carried out on the matrices)

PHDP:	NTAATNNNATTAN
Prrx1:	NNNAATTA
Pknox1:	NGACANNN
LHX1:	NYNATTAN
LHX2:	NNNNTTNNNN
LHX4:	NTAATTAN
LHX6:	NYAATCAN
CBG04010:	NNNMATTAANN
LHX9:	NNTNATTAN
PK24181.1:	NNNNNNNNNNNN
IRX5:	NNNTGTNNNACAYNN
IRX2:	NWACAYGACANN
pal-1:	TTWATTAC
GSC2:	NNNATTAN
Isl2:	NNNTNNN
WOX13:	NNTGATTGAN
Q0N4F9_NEMVE:	MTAATTAN
hox12:	TTWATKAC
eve:	NSTAATTAGN
ZHD7:	NNNNNTTANN
OdsH:	NYAATTAN
unc-62:	NNNTNTNNN
NEMVEDRAFT_v1g113057:	NNGGATTARN
DLX3:	NYAATTAN
HBX4_ECHGR:	NNGGATTANN
Q7M3U4_ACRFO:	NNNAATTANNN
Hoxb8:	NNNNATWNN
Hoxb9:	NNNNRTWAAN
Hoxb3:	NNNNATTAN
Six6:	NNKATNN

GSX1:	NTNATKAN
Hoxb6:	NNNNNNN
Hoxb7:	NNNNTNNN
Hoxb4:	NNNNATTAN
ENSCING00000016045:	NCYAATTARNN
Hoxc11:	NNNNRTWAAN
ceh-36:	NNGATTAN
Hoxc13:	NTTTAYNNNN
Hoxc12:	NNTWAYNNNN
VSX2:	CTAATTAN
HOXB8:	GYMATTAN
HOXB9:	NTTTTACGAC
SNAPOd2T00006145001:	NNTRATTRNN
HOXB2:	NYAATTAN
HOXB3:	NNYMATTANN
HOXB6:	GYMATTAN
HOXB7:	NYAATTAN
HOXB5:	NYMATTAN
HOXC11:	NNTTTACGAYN
HOXC13:	NTTTTATTGSN
HOXC12:	NTTTTATTACN
Hoxd12_CONSTRUCT:	NNTNAYNNNN
six3:	NNNATMNNNN
hbn:	NTAATTTAATTAN
Hoxa6:	NNNNATTAN
HDG11:	NCATTWAWKN
barx1:	NNNNATTAN
pbx4:	NNCATCAA
Duxbl1:	NTGATTRNN
Hmx2:	NNNCRNTTNN
ceh-24:	NNYNCTYNN
ceh-23:	NNNTAATTAN
ceh-22:	NNNCACTTNA
vis:	TGACAN
Dfd:	NNYCATTANN
BSX:	NNCGTTAN
В-Н2:	NNCGTTTANN
B-H1:	NNCGTTTANN
CG5369:	TAATNNAATTA
ceh-90:	NGTTTTANN
BARX1:	NNATTAAAWANCNATTA
CNK00090:	ATAATTAA
Gbx2:	NNYAATTANN

Gbx1:	NNYAATTANN
Hlx:	NNNAWNN
SNOG_08903:	TTGATTTR
EMX2:	NTAATTAG
EMX1:	NTAATTAN
Gsx2:	NNNNATTAN
GRMZM2G087741:	NNTGACRKCN
Pdx1:	NNNATTAN
cad:	NTTTTATKGCNN
C4B866_NEMVE:	NNTTAATTANNN
oc:	NGGATTAN
Q0N4H2_NEMVE:	ANYAATTA
Q9GP48_9TURB:	NTAATTAN
Pph13:	NYAATTAN
Q24782_9CNID:	TTAATTAA
Esx1:	NAATTANN
Q70HR3_PLEPI:	NNNTACTTAN
Nobox:	NNYNATNNN
Bsx:	NNNNATTAN
ventx1.1.L:	NNNNNNNNNNNN
ceh-5:	NNNATTANN
HNF1B:	GTTAATNATTAAY
POU5F1B:	NNNTTNN
AAZ08039:	NNYAATTANN
PROP1:	NAAYNNAATTA
Optix:	TAATTARATTA
YHP1:	NWWATTRNN
Hoxb13:	NTTWAYNNNN
ATEG_00670:	NNNNACNNN
TVAG_451170:	NNCRCATCAN
lab:	STAATTAN
Nkx2-9:	NNRAGTRNNN
Prop1:	NRATNNN
PHOX2A:	NNAATTAN
Nkx2-3:	NNRAGNRNN
PHOX2B:	NYAATTAN
Nkx2-6:	NNNCACTTNN
CG2808:	TAATSNAATTA
HOXB13:	NTTTTATTGGN
HMBOX1:	NCTAGTTAN
Ptx1:	NGGATTAN
caup:	NWNTAACA
NEMVEDRAFT_v1g101664:	GATYGATT

zfhx4:	NYAATTANN
unc-4:	TAATTNAATTA
Ubx:	NYAATTAN
Rx:	NYAATTAN
SNAPOd2T00005194001:	NNACATGTANN
CBG24578:	NNNTGACAGSTN
Nkx6-3:	NNNATNANN
Nkx6-1:	NYNATTANN
MTR_2g038000:	NNNNNAGNNN
Q9C1N5_COPSC:	NNGATTGNN
SIX6:	NNNTATCRNN
SIX3:	NNNTATCRNN
Barx2:	NNNNATTAN
HNF1A:	NRTTAATNATTAACN
Vsx1:	NCAATTAN
Vsx2:	NYAATTAN
RAX:	YTAATTAN
GRMZM2G135447:	NTGACRKS
Hoxa13:	NTTWAYNNN
DPRX:	NGGATTATCN
ESX1:	NYAATTAN
SHOX2:	NYAATTAN
Hoxa10:	NNNNNTNNN
hoxd13a:	WTTTATKA
Tlx2:	ATTAAWTA
ceh-62:	NNTAATTANN
Hoxa5:	NNNNATTAN
$Nk \times 2-1$:	NNRAGNRN
en:	NYAATTAN
NOBOX:	NNNNATNRNN
Otx1:	NNTAATCGGATTANN
abd-A:	NGTAATTANN
Dll:	NYAATTAN
Lhx6_CONSTRUCT:	NYAATYANN
pha-2:	NNNATNNNNN
MNX1:	NTAATTAN
Cphx1:	WGATNNNNN
CG11085:	NCRATTAN
ftz:	GTAATTAA
uncx:	NYAATTANN
EN1:	NYAATTAN
EN2:	NTAATTAN
E5:	NNNATTAN

ro:	TAATTARN
lms:	NYAATTAN
TOS8:	NNNTGACGTMW
Hoxa7:	NNNNATWNN
En1:	
En2:	NSTAATTANN
Hoxa4:	NNNNATTAN
Hoxa3:	NNNYNATTAN
Hoxa2:	NNTAATTANN
Hoxal:	NNYNATTNN
NANOG:	NCNRTYAN
Hoxa9:	
Nanog:	NNNNATNNN
HAT1:	NNATGATTRN
HAT3:	NTGATTRN
NCU03266:	NYNATTWNNN
Q0N4H9_NEMVE:	NNNNTYRNN
PDX1:	NTMATTAR
NK7.1:	NACCGTTTAAN
ISX:	CTAATTAN
Q70LF3_ARTSF:	NTAATTAN
Hoxall:	NNNTTTACGANN
ZFHX2:	NNTAATTANN
Nkx2-2:	NNNRAGNRN
Q9BJW6_9CNID:	NNTAATTAGN
CAD26261:	TWWATTANN
NEMVEDRAFT_v1g129868:	NNNNTNRN
KNAT6:	NNNGACNNNN
Hmx:	NCAMTTAN
Lim3:	NTAATTAN
Lim1:	NYAATTAN
PRH_PETCR:	ATATATWA
Abd-B:	NWTTTACGACN
Nkx2-4:	NNRAGNRNNN
Nkx3-1:	NNAAGTGNNN
Otp:	NNATTANN
Nkx2-5:	NNNRAGNRNN
Nkx3-2:	NNAAGTRNN
ceh-19:	NNCNATTAN
Meis3:	NTGACAN
Q24780_9CNID:	NNTTAATTANN
ATHB-12:	NNNNNNNN
ATHB-16:	AATKATTA

Q00354_COPCI:	NGATTGGT
AGAP009986:	NNNACANNNN
Mnx1:	NNNNNATTAN
Q86SD7_PODCA:	NNTAATTANN
Antp:	TMATTAAR
HOXA7:	NYMATTAN
HOXA6:	NGYMATTANN
HOXA5:	STAATTAS
HOXA4:	RTMATTAN
HOXA2:	NTAATTAN
HOXA1:	NNTAATTANN
Q8VHG7_MOUSE:	NGGATTAW
HOXA9:	NNNTTTAYGAYN
Obox6:	NNRGATTAN
Obox5:	NNNATNAN
Obox3:	NNATNWN
Obox1:	NNNGGATTAN
lbe:	NNTCGTTA
PITX2:	NGGATTAN
PITX3:	NGGATTAN
PITX1:	NGGATTAN
lbl:	CTMATTAN
CBG07641:	NNTAATTANN
HXD12_HETFR:	TTAATTAC
DMBX1:	NGGATTAN
ENSTNIG00000004116:	CTAATTAA
ALX3:	NYAATTAN
ALX1:	NTAAYNYAATTAN
Emx2:	NNNNATTAN
ALX4:	NYAATTAN
Dr:	NNYAATTA
NEMVEDRAFT_v1g114268:	AYCNATTAN
vnd:	TYTCAAGTR
Smp_158000:	CYAATTAA
CG32105:	YTAATTAA
DRGX:	NYAATTAN
Q68SS9_PLEDJ:	NGATTGNN
bsh:	TAATYTAATTA
Dlx5:	NNNATTW
PBX4:	NNNNTNNNN
exex:	NNNNYAATTNNCNN
PBX1:	NTGATTGAYR
SNAPOd2T00003340001:	NWWATTANNN

exd:	NTTTGACA
CBG10835:	NNNNTCANN
TLX2:	NYAATTAN
TLX3:	NAATTGNNNNNNNNNNNNNCAATTN-
GSX2:	SYMATTAN
TGIF2:	TGACANNTGTCA
TGIF1:	TGACAGCTGTCA
Dmbx1:	NNRGATTAN
Rhox6:	NNNATTAN
CG18599:	TAATTAN
Rhox8:	NTTACACCN
Uncx:	NTAATYNAATTAN
Hoxb5:	NNNNATNAN
Tgif2:	NNTGACANNN
Tgif1:	NNTGACANNN
slou:	NNYAATTANN
Hoxd8:	NNNNNNNN
Hoxd9:	NCCATWAAA
Hoxc10:	NTTTTAYGAC
yox1:	NNNNATNNNN
Hoxd3:	NNYAATTANN
POU6F1:	NNTAATTANN
Q0N4H7_NEMVE:	NNNNNATNNN
HOXD8:	GYMATTAN
HOXD9:	NTTTTATTGC
HOXD4:	NYMATTAN
zen:	NTMATTAN
EVX1:	NTAATTAN
EVX2:	NTAATTAN
repo:	NYTAATTAAN
hbx4:	NNTGACRNNN
hbx2:	NNATATATWN
vsx2:	NYTAATTAANN
hbx9:	NNTGACAN
Six3:	NKATNNN
bap:	NNCACTTANN
Hnf1b:	NNNGTNNN
NKX2-8:	NNRAGNRN
Otx2:	NNGATTAN
SNAPOd2T00006620001:	NYAATTAN
ems:	NTAATTAN
PRRX1:	NYAATTAN
CG15696:	NNNATTRNN

PRRX2:	NNCRTNAN
CRX:	NSGGATTAGN
hbx5-2:	WTGATTRN
Meis2:	NTGACAN
Meis1:	NTGACANNN
hbx5-1:	TTGATTGR
ara:	NNAWAACA
VENTX:	CTAATTAN
btn:	NNTMATTANN
CAGL0H02959g:	NNTGACRNNN
MEIS2:	TGACAGCTGTCA
MEIS1:	NTGACAN
tup:	CMATTAWS
ceh-58:	NNYAATTANN
Awh:	NTAATTAN
ceh-54:	NNYAATTAN
Dobox5_CONSTRUCT:	NNRGATTAN
ceh-57:	NNNNNTWAN
ceh-51:	NNNATNRNN
ceh-53:	NRGATTAN
PHO2:	NNNNNATNNN
Q6WL90_TRICY:	NNNATTAN
LBX2:	CTCRTTAA
LBX1:	CTAATTAN
PKNOX1:	TGACASSTGTCA
PKNOX2:	TGACANNTGTCA
CBG23031:	NNNNATWAAN
HOXC10:	NNTTTATKRCN
NEMVEDRAFT_v1g81436:	NNNNTNNNNN
Hnfla:	NNNGTTAN
ENSTNIG00000017092:	NTGATTAN
H2.0:	TTWATNA
NEMVEDRAFT_v1g80394:	NNNATNNNN
unpg:	NNCAATTANN
NEMVEDRAFT_v1g164989:	YTAATTAG
Noto:	NNNNATNNNN
MEOX1:	NSTAATTANN
MEOX2:	NNTAATTANN
Nkx1-1:	NNNNATTANN
pdm3:	CTAATTAN
UM04928:	NNNATNNNNN
Hmbox1:	NNNGTNNN
Alx3:	NNNATTA

CG11617:	WTTTWACA
Alx1:	NNTAATTANN
HOXA13:	GTTTTATTGGN
HOXA11:	NTTTTACGACN
HOXA10:	NTTTTACGACN
pb:	GTMATTAN
Barx1:	NNNNATTANN
ATHB-4:	NNMTGCATRNN
ATHB-6:	NATNATTNN
ATHB-7:	NNATGATTGN
Barhl1:	NNCRTTTANN
Rhox11:	NNNTGTWNNN
ventx3.2:	TAATTAAT
Crx:	NNGATTAN
ENSCING0000007395:	NWNNNNNN
otp:	NYAATTAN
Prrx2:	NYAATTAN
P79681_ACIBE:	NTAATTAA
si_dkey-43p13.5:	NNNNWWNNNN
Lbx2:	NNNNTTNN
CG9876:	NNYRATTANS
NEMVEDRAFT_v1g8907:	NNNYMATTAN
Phox2a:	NNATTNNN
C15:	YMWTTAAN
UNCX:	NNAATTNN
TGIF2LY:	TGACANNTGTCA
TGIF2LX:	TGACAGNTGTCA
CG34031:	NNNNNNRN
mix11.S:	NNTAATTAANN
NEMVEDRAFT_v1g205636:	NTAATTAN
ARX:	YTAATTAN
dharma:	GGATNNN
Dlx4:	NNNATTAN
ATEG_04171:	NNNYAATTWN
Dlx1:	NYAATTAN
D1x2:	NYAATTAN
Dlx3:	NNNNATTAN
Hmx1:	NNNCRNTTNN
CDX1:	NTTTTATGGCN
ISL1:	SCACTTAN
ISL2:	NCACTTAN
CDX2:	NNTTAYRAYN
SPU_026099:	NTAATTAA

GSC:	NGGATTAN
ARGFX:	NCTAATTARN
NKX2-3:	NNCACTTNNN
CAD25284:	TTAATTAN
NKX2-5:	YWCTTNN
Phyra73700:	NNNNNTCAN
prd-b:	NTRATTRNN
ANIA_00885:	NNTKACANNN
CBG00023:	NNNNNATWAN
Msx1:	NNATTAN
Msx2:	NNYAATTA
Msx3:	NCAATTANNNNSCAATTA
CBG00029:	TAATTAAT
Vax2:	NNNNATTAN
Vax1:	NNNNATTAN
YOX1:	YNATWWN
Hoxd13:	NTTTTATTRS
Hoxd10:	NNNNNNNN
Hoxd11:	NNNNNNNN
NKX6-2:	NYAATTAN
NKX6-3:	NTCATTAWN
Hoxc9:	NTTAATTACN
Hoxc8:	NNNNNNNNN
BARHL2:	ANCGTTTAN
BARHL1:	NCGTTTAN
Arx:	NTAATTNAATTAN
Hoxc5:	NNATTNN
Hoxc4:	NNNNATTAN
Hoxc6:	RTAATTAA
Shox2:	NYAATTAN
vab-7:	MTAATTAA
ENSTNIG00000006040:	NNNYNATTANNNN
lim-4:	NNTAATTAR
Evx1:	NNNNATTAN
HOXC4:	NTAATTAA
Evx2:	NNATTANN
Isx:	NNATTNNN
NEMVEDRAFT_v1g136480:	NNNNNNNNNNNN
Lhx1:	NNNNATTA
Lhx2:	NYMATTAN
Lhx3:	NNATTANNN
Lhx4:	NTAATTAN
Lhx5:	NNNYAATTA

Lhx6:	NTRATTRN
Lhx8:	-NTRATTANNNNTRATTAN
GRMZM2G038252:	NCAATCAN
OTX1:	NGGATTAN
OTX2:	NNTAATCGGATTANN
Pknox2:	TGACAGGTGTCA
DLX4:	NCAATTAN
DLX5:	NCAATTAN
DLX6:	NCAATTAN
DLX1:	NYAATTAN
DLX2:	NYAATTAN
ATML1:	NNNANTNANN
ceh-37:	NGGATTAN
LMX1A:	TTAATTAN
LMX1B:	NNNNTAATTAN
Gsc:	NGGATTA
achi:	TGACAN
RLT1:	NANGAANN
HOXD12:	NTTTACGAYN
HOXD13:	GKTTTATTGSS
HOXD10:	NTTTTATKRC
HOXD11:	NTWANNNTTTACGAYN
CUP9:	NTGACACAWN
Rax:	NNAATTANN
Scr:	NYMATTAN
VAX2:	NNATTANN
Q4RRY4_TETNG:	NTRATTRNN
Q0ZPQ4_NEMVE:	CTTCTTAA
Q00352_COPCI:	TGATTGGTT
Alx4:	NTAATYNAATTAN
GBX2:	NYAATTAN
GBX1:	NSYAATTANN
kal-1:	NNNTNNNN
HESX1:	NCYAATTAN
AAEL012091:	NNGATNNNN
Pitx2:	NGGATTAN
Pitx3:	NGGATTAN
Pitx1:	NNGGATTAN
scro:	NCACTYNNN
HHEX:	WTNAATTA
PK25011.1:	NNWTNNNNN
CG32532:	NTAATNNNATTAN
Q0ZPQ8_NEMVE:	NTGACRNNN

NCU09556:	TATATTAA
MIXL1:	NNYAATTANN
CG4328:	NWTWATTR
CNG03450:	NNWATTAAN
bcd:	NNGGATTANNN
Q0N4M8_NEMVE:	GTAATTAN
NKX3-2:	NCCACTTANN
NKX3-1:	NCCACTTANN
Irx5:	NNNNACANN
CBG20882:	NYNATNNNN
Irx6:	
Irx2:	NNNACANNN
HMX1:	NCAMTTANN
HMX3:	NNCAMTTANNN
HMX2:	NCACTTRNC
NEMVEDRAFT_v1g81309:	NNNNATTAN
Cdx1:	NNYNATWNN
inv:	TAATTARW
Cdx2:	NWWTANWN
hbx10:	NWWATTRN
al:	TAATNNNATTA
RHOXF1:	GGATNATCN
ap:	TAATTARN
ind:	TAATTARNR
Sebox:	NNTTAATTAN
Phox2b:	NNATTNNN
XP_003689450.1:	NNNANMCNNN
Lmx1a:	NNNATNRN
Lmx1b:	NNATNAN
tin:	NCCACTTGAN
ANL2:	NCATTAATTNN
MSX1:	NCAATTAN
MSX2:	NCAATTAN
AGAP006642:	GGATTANN
Meox1:	NNYMATTAN
Meox2:	NTAATTAN
RAX2:	NYAATTAN
SHOX:	NYAATTAN
CBG01820:	NGGATTAN
zen2:	NTMATTAN
NEMVEDRAFT_v1g105595:	NNAATTAN
CG11294:	TAATNNAATTA
Nkx1-2:	NNNATTANN

Barhl2:	NNYNNTTAN
HGTX:	NTMATTAN
dve:	NNTAATTAGN
Q8I9J5_9BILA:	NRTWATTANN
NOTO:	NYAATTAN
KLLA0_D10043g:	NNNNNNNNNNNN

Familial Profile:
 (click for matrix)

Motif Tree

Tree (drawn by Phylip) Click here for Newick-format tree (viewable with ME	Input Motif	Best match in JASPAR
	<u>PHDP</u>	MA0229.1_inv (E val: 1.9109e-06)
	<u>hbn</u>	MA0229.1_inv (E val: 1.1169e-09)
	CG5369	MA0194.1_Lim1 (E val: 1.2872e-09)
	<u>Optix</u>	MA0229.1_inv (E val: 3.1478e-11)

<u>bsh</u>	MA0229.1_inv (E val: 3.9607e-11)
unc-4	MA0178.1_CG32105 (E val: 1.8305e-08)
<u>Arx</u>	MA0194.1_Lim1 (E val: 1.0070e-08)
<u>Unex</u>	MA0172.1_CG11294 (E val: 3.7957e-09)
Alx4	MA0194.1_Lim1 (E val: 6.3149e-08)
CG2808	MA0188.1_Dr (E val: 1.4849e-08)
CG11294	MA0229.1_inv (E val: 6.6941e-10)
ALX1	MA0229.1_inv (E val: 1.5023e-08)

<u>CG32532</u>	MA0232.1_lbl (E val: 5.3536e-07)
<u>al</u>	MA0172.1_CG11294 (E val: 1.2922e-05)
PROP1	MA0188.1_Dr (E val: 2.8776e-09)
Otx1	MA0234.1_oc (E val: 2.4512e-07)
OTX2	MA0234.1_oc (E val: 2.2972e-07)
Prrx1	MA0075.1_Prrx2 (E val: 1.7693e-06)
<u>Dlx5</u>	MA0075.1_Prrx2 (E val: 1.2097e-06)
Alx3	MA0075.1_Prrx2 (E val: 8.5973e-07)

Msx2	MA0075.1_Prrx2 (E val: 1.9540e-06)
<u>Pph13</u>	MA0075.1_Prrx2 (E val: 1.7716e-06)
ALX4	MA0075.1_Prrx2 (E val: 1.8830e-06)
PHOX2A	MA0075.1_Prrx2 (E val: 1.9368e-06)
NEMVEDRAFT v1g105595	MA0075.1_Prrx2 (E val: 2.2012e-06)
<u>Rax</u>	MA0075.1_Prrx2 (E val: 3.8995e-06)
<u>Lhx 1</u>	MA0075.1_Prrx2 (E val: 5.0294e-06)
LHX1	MA0426.1_YHP1 (E val: 3.7149e-08)

<u>Lim1</u>	MA0426.1_YHP1 (E val: 4.6222e-08)
PHOX2B	MA0426.1_YHP1 (E val: 5.2955e-08)
ESX1	MA0426.1_YHP1 (E val: 9.7762e-08)
<u>OdsH</u>	MA0426.1_YHP1 (E val: 2.3854e-08)
<u>otp</u>	MA0426.1_YHP1 (E val: 2.8789e-08)
<u>DII</u>	MA0426.1_YHP1 (E val: 2.2713e-08)
Vsx2	MA0426.1_YHP1 (E val: 2.3819e-08)
DLX5	MA0426.1_YHP1 (E val: 2.1206e-08)

DLX4	MA0426.1_YHP1 (E val: 2.6660e-08)
DLX6	MA0426.1_YHP1 (E val: 2.4257e-08)
DLX1	MA0426.1_YHP1 (E val: 3.6604e-08)
GBX2	MA0426.1_YHP1 (E val: 4.0831e-08)
<u>lms</u>	MA0426.1_YHP1 (E val: 3.2934e-08)
<u>Rx</u>	MA0426.1_YHP1 (E val: 2.5179e-08)
<u>EN1</u>	MA0426.1_YHP1 (E val: 3.7473e-08)
<u>Vsx1</u>	MA0426.1_YHP1 (E val: 1.3173e-08)

MSX1	MA0426.1_YHP1 (E val: 1.1714e-08)
MSX2	MA0426.1_YHP1 (E val: 1.1888e-08)
<u>en</u>	MA0426.1_YHP1 (E val: 6.3159e-08)
<u>Dlx1</u>	MA0426.1_YHP1 (E val: 7.3537e-08)
<u>Dlx2</u>	MA0426.1_YHP1 (E val: 8.3123e-08)
<u>Dr</u>	MA0426.1_YHP1 (E val: 2.9088e-08)
SNAPOd2T00006620001	MA0426.1_YHP1 (E val: 5.3836e-08)
DLX3	MA0232.1_lbl (E val: 1.1929e-07)

SHOX2	MA0232.1_lbl (E val: 1.2365e-07)
Shox2	MA0232.1_lbl (E val: 9.8309e-08)
TLX2	MA0232.1_lbl (E val: 8.2750e-08)
DLX2	MA0232.1_lbl (E val: 1.1128e-07)
RAX2	MA0232.1_lbl (E val: 1.1674e-07)
PRRX1	MA0232.1_lbl (E val: 6.1316e-08)
<u>ceh-54</u>	MA0426.1_YHP1 (E val: 3.0546e-07)
Gbx2	MA0426.1_YHP1 (E val: 2.9462e-07)

Gbx1	MA0426.1_YHP1 (E val: 3.9716e-07)
<u>CG11085</u>	MA0426.1_YHP1 (E val: 2.0173e-08)
<u>Hmx</u>	MA0426.1_YHP1 (E val: 4.3245e-08)
<u>AAZ08039</u>	MA0426.1_YHP1 (E val: 2.2824e-07)
GBX1	MA0232.1_lbl (E val: 4.9552e-07)
<u>CG9876</u>	MA0426.1_YHP1 (E val: 4.0548e-07)
unpg	MA0426.1_YHP1 (E val: 2.0776e-07)
<u>ceh-58</u>	MA0232.1_lbl (E val: 4.0935e-07)

Barhl2	MA0426.1_YHP1 (E val: 5.7920e-07)
<u>ceh-19</u>	MA0426.1_YHP1 (E val: 4.4632e-07)
YHP1	MA0232.1_lbl (E val: 5.2801e-06)
ATEG 04171	MA0232.1_lbl (E val: 4.6903e-06)
NCU03266	MA0196.1_NK7.1 (E val: 1.9654e-08)
<u>tup</u>	MA0248.1_tup (E val: 2.1446e-11)
LHX4	MA0232.1_lbl (E val: 1.2360e-07)
EMX1	MA0232.1_lbl (E val: 1.3018e-08)

<u>lab</u>	MA0232.1_lbl (E val: 1.0849e-08)
2210200	MA0232.1_lbl
CG18599	(E val: 1.9400e-09)
Meox2	MA0232.1_lbl (E val: 7.1346e-09)
<u>ems</u>	MA0232.1_lbl (E val: 9.6821e-09)
EVX1	MA0232.1_lbl (E val: 1.2369e-08)
EVX2	MA0232.1_lbl (E val: 1.2336e-08)
Lim3	MA0232.1_lbl (E val: 2.1239e-08)
<u>Lhx4</u>	MA0232.1_lbl (E val: 2.0476e-08)

Q9GP48 9TURB	MA0232.1_lbl (E val: 3.8246e-08)
Q70LF3 ARTSF	MA0232.1_lbl (E val: 2.7533e-08)
NEMVEDRAFT v1g205636	MA0232.1_lbl (E val: 3.7288e-08)
Esx1	MA0232.1_lbl (E val: 3.5689e-08)
MNX1	MA0232.1_lbl (E val: 2.6559e-08)
<u>DRGX</u>	MA0232.1_lbl (E val: 3.1653e-08)
<u>EN2</u>	MA0232.1_lbl (E val: 3.0403e-08)
Prrx2	MA0232.1_lbl (E val: 2.7448e-08)

ALX3	MA0232.1_lbl (E val: 2.1881e-08)
SHOX	MA0232.1_lbl (E val: 2.2701e-08)
<u>Awh</u>	MA0232.1_lbl (E val: 1.6501e-08)
I hu 2	MA0232.1_lbl
Lhx3	(E val: 6.3907e-08)
uncx	MA0232.1_lbl (E val: 9.0879e-08)
<u>Lhx5</u>	MA0232.1_lbl (E val: 7.9458e-08)
zfhx4	MA0232.1_lbl (E val: 5.4071e-08)
OARRYA TETNO	MA0232.1_lbl
Q4RRY4 TETNG	(E val: 2.1430e-07)

Hoxa1	MA0232.1_lbl (E val: 3.8787e-08)
HOXA1	MA0232.1_lbl (E val: 1.4927e-07)
<u>prd-b</u>	MA0232.1_lbl (E val: 7.3149e-07)
<u>Lhx6</u>	MA0232.1_lbl (E val: 1.2291e-06)
<u>ceh-23</u>	MA0232.1_lbl (E val: 1.3081e-07)
<u>Alx1</u>	MA0232.1_lbl (E val: 1.4374e-07)
MIXL1	MA0232.1_lbl (E val: 2.4876e-07)
<u>En2</u>	MA0232.1_lbl (E val: 6.0426e-08)

slou	MA0232.1_lbl (E val: 1.4101e-07)
GSX1	MA0232.1_lbl (E val: 6.6879e-08)
HOXB2	MA0232.1_lbl (E val: 1.5599e-08)
<u>NOTO</u>	MA0232.1_lbl (E val: 1.6063e-08)
HOXB7	MA0232.1_lbl (E val: 1.8210e-08)
NKX6-2	MA0232.1_lbl (E val: 1.8473e-08)
<u>Ubx</u>	MA0232.1_lbl (E val: 1.5127e-08)
<u>PDX1</u>	MA0232.1_lbl (E val: 1.1612e-08)

HOXA2	MA0232.1_lbl (E val: 8.6640e-09)
Lhx2	MA0232.1_lbl (E val: 9.1589e-09)
<u>HGTX</u>	MA0232.1_lbl (E val: 6.2738e-09)
HOXB5	MA0232.1_lbl (E val: 5.6550e-08)
HOXA7	MA0232.1_lbl (E val: 1.0318e-07)
HOXD4	MA0232.1_lbl (E val: 6.8867e-08)
<u>Scr</u>	MA0232.1_lbl (E val: 4.3130e-08)
HOXB8	MA0232.1_lbl (E val: 1.3096e-07)

HOXD8	MA0232.1_lbl (E val: 2.2035e-07)
HOXB6	MA0232.1_lbl (E val: 3.1744e-08)
HOXA5	MA0223.1_exex (E val: 3.9288e-10)
Q0N4M8 NEMVE	MA0223.1_exex (E val: 2.3856e-09)
HOXA6	MA0223.1_exex (E val: 8.5000e-08)
abd-A	MA0223.1_exex (E val: 1.9332e-08)
HOXA4	MA0232.1_lbl (E val: 8.1005e-08)
<u>pb</u>	MA0232.1_lbl (E val: 6.8619e-08)

HOXB3	MA0232.1_lbl (E val: 5.6179e-08)
NEMVEDRAFT v1g8907	MA0232.1_lbl (E val: 1.4668e-07)
Hoxd3	MA0232.1_lbl (E val: 1.0207e-07)
Hoxa2	MA0232.1_lbl (E val: 3.9527e-08)
MEOX2	MA0232.1_lbl (E val: 7.0922e-08)
<u>eve</u>	MA0241.1_ro (E val: 7.8255e-10)
GSX2	MA0232.1_lbl (E val: 8.5976e-09)
MEOX1	MA0223.1_exex (E val: 5.9674e-09)

VSX2	MA0241.1_ro (E val: 2.7339e-11)
<u>ARX</u>	MA0241.1_ro (E val: 3.0360e-11)
ISX	MA0241.1_ro (E val: 8.0527e-11)
<u>VENTX</u>	MA0241.1_ro (E val: 9.6463e-11)
EMX2	MA0241.1_ro (E val: 2.5200e-11)
<u>ro</u>	MA0241.1_ro (E val: 1.2763e-11)
LBX1	MA0241.1_ro (E val: 4.2517e-10)
pdm3	MA0241.1_ro (E val: 1.8173e-10)

<u>lbl</u>	MA0241.1_ro (E val: 1.9839e-09)
<u>dve</u>	MA0241.1_ro (E val: 9.7944e-10)
Q9BJW6 9CNID	MA0241.1_ro (E val: 8.0265e-10)
HESX1	MA0188.1_Dr (E val: 1.7446e-09)
<u>exex</u>	MA0198.1_OdsH (E val: 1.1913e-08)
<u>pal-1</u>	MA0448.1_H2.0 (E val: 2.2106e-10)
<u>hox12</u>	MA0174.1_CG42234 (E val: 1.3600e-10)
HXD12 HETFR	MA0206.1_abd-A (E val: 9.0750e-11)

<u>ftz</u>	MA0226.1_hbn (E val: 2.0192e-11)
Hoxc6	MA0194.1_Lim1 (E val: 1.9802e-11)
Hoxc9	MA0240.1_repo (E val: 7.4846e-10)
<u>H2.0</u>	MA0448.1_H2.0 (E val: 2.5989e-11)
Antp	MA0166.1_Antp (E val: 1.2763e-11)
<u>C15</u>	MA0170.1_C15 (E val: 1.2763e-11)
<u>Q24780 9CNID</u>	MA0194.1_Lim1 (E val: 7.3681e-10)
Q0N4F9 NEMVE	MA0180.1_Vsx2 (E val: 1.1622e-12)

P79681 ACIBE	MA0180.1_Vsx2 (E val: 6.4537e-13)
<u>SPU 026099</u>	MA0180.1_Vsx2 (E val: 8.9306e-13)
<u>vab-7</u>	MA0180.1_Vsx2 (E val: 1.7780e-12)
ENSTNIG00000004116	MA0180.1_Vsx2 (E val: 5.9242e-13)
Smp 158000	MA0180.1_Vsx2 (E val: 4.7229e-13)
NEMVEDRAFT v1g164989	MA0180.1_Vsx2 (E val: 7.4326e-12)
<u>ARGFX</u>	MA0180.1_Vsx2 (E val: 1.2971e-10)
vsx2	MA0180.1_Vsx2 (E val: 3.4137e-10)

Q24782 9CNID	MA0194.1_Lim1 (E val: 1.4038e-11)
<u>CG32105</u>	MA0236.1_otp (E val: 3.4164e-11)
<u>repo</u>	MA0240.1_repo (E val: 2.8598e-10)
<u>CNK00090</u>	MA0240.1_repo (E val: 2.3502e-11)
HOXC4	MA0240.1_repo (E val: 2.0401e-11)
<u>CAD25284</u>	MA0194.1_Lim1 (E val: 1.5897e-11)
LMX1A	MA0172.1_CG11294 (E val: 1.7100e-11)
<u>Tlx2</u>	MA0194.1_Lim1 (E val: 2.7621e-10)

ventx3.2	MA0194.1_Lim1 (E val: 1.7841e-11)
<u>CBG00029</u>	MA0194.1_Lim1 (E val: 1.6173e-11)
<u>HHEX</u>	MA0172.1_CG11294 (E val: 2.1773e-10)
ENSCING00000016045	MA0180.1_Vsx2 (E val: 5.3273e-10)
RAX	MA0202.1_Rx (E val: 5.1507e-11)
<u>inv</u>	MA0229.1_inv (E val: 6.5503e-15)
<u>ap</u>	MA0209.1_ap (E val: 1.2763e-11)
<u>lim-4</u>	MA0181.1_Vsx1 (E val: 9.2564e-11)

<u>ind</u>	MA0228.1_ind (E val: 7.6496e-11)
<u>Sebox</u>	MA0194.1_Lim1 (E val: 2.5830e-10)
mix11.S	MA0194.1_Lim1 (E val: 6.0393e-10)
C4B866 NEMVE	MA0194.1_Lim1 (E val: 1.2649e-09)
CAD26261	MA0178.1_CG32105 (E val: 2.6988e-08)
LHX9	MA0232.1_lbl (E val: 2.4650e-07)
POU6F1	MA0232.1_lbl (E val: 1.9095e-07)
Q86SD7_PODCA	MA0232.1_lbl (E val: 2.5359e-07)

CBG07641	MA0232.1_lbl (E val: 1.8462e-07)
<u>ceh-62</u>	MA0232.1_lbl (E val: 2.4811e-07)
ZFHX2	MA0232.1_lbl (E val: 2.6731e-07)
<u>btn</u>	MA0232.1_lbl (E val: 7.5444e-08)
LMX1B	MA0232.1_lbl (E val: 2.8185e-07)
SNAPOd2T00006145001	MA0232.1_lbl (E val: 2.6419e-06)
HMX3	MA0426.1_YHP1 (E val: 2.8398e-07)
SNAPOd2T00003340001	MA0232.1_lbl (E val: 6.4014e-06)

CG4328	MA0182.1_CG4328 (E val: 1.4526e-07)
nbx10	MA0182.1_CG4328 (E val: 1.1851e-05)
<u> </u>	MA0448.1_H2.0 (E val: 3.6600e-05)
Q7M3U4 ACRFO	MA0075.1_Prrx2 (E val: 1.9488e-05)
ENSTNIG00000006040	MA0426.1_YHP1 (E val: 8.7845e-07)
<u>CBG04010</u>	MA0206.1_abd-A (E val: 7.7487e-06)
<u>CNG03450</u>	MA0444.1_CG34031 (E val: 7.7102e-07)
HDG11	MA0151.1_ARID3A (E val: 3.6185e-06)

ANL2	MA0206.1_abd-A (E val: 4.3694e-06)
ATML1	MA0186.1_Dfd (E val: 8.4622e-05)
<u>Dfd</u>	MA0232.1_lbl (E val: 1.7692e-06)
<u>zen</u>	MA0232.1_lbl (E val: 7.9820e-08)
zen2	MA0232.1_lbl (E val: 2.1091e-08)
NKX6-3	MA0215.1_btn (E val: 3.0070e-09)
HNF1B	MA0153.1_HNF1B (E val: 5.5378e-13)
<u>HNF1A</u>	MA0153.1_HNF1B (E val: 6.3727e-14)

<u>ATHB-16</u>	MA0008.1_HAT5 (E val: 1.8398e-09)
<u>NANOG</u>	MA0214.1_bsh (E val: 1.0306e-04)
NEMVEDRAFT v1g113057	MA0212.1_bcd (E val: 2.6743e-08)
HBX4 ECHGR	MA0190.1_Gsc (E val: 3.2658e-08)
<u>oc</u>	MA0212.1_bcd (E val: 3.8618e-09)
<u>GSC</u>	MA0234.1_oc (E val: 3.9560e-09)
<u>Gsc</u>	MA0234.1_oc (E val: 9.9606e-10)
Q8VHG7 MOUSE	MA0201.1_Ptx1 (E val: 2.4983e-10)

PITX1	MA0234.1_oc (E val: 3.8797e-09)
CBG01820	MA0212.1_bcd (E val: 3.9289e-09)
PITX3	MA0234.1_oc (E val: 3.7633e-09)
OTX1	MA0234.1_oc (E val: 3.7603e-09)
<u>Ptx1</u>	MA0234.1_oc (E val: 5.0663e-09)
<u>Pitx2</u>	MA0234.1_oc (E val: 4.9980e-09)
Pitx3	MA0234.1_oc (E val: 6.8909e-09)
<u>PITX2</u>	MA0234.1_oc (E val: 7.9895e-09)

DMBX1	MA0234.1_oc (E val: 8.8704e-09)
<u>ceh-53</u>	MA0190.1_Gsc (E val: 4.6874e-09)
ceh-37	MA0190.1_Gsc (E val: 4.0372e-09)
AGAP006642	MA0190.1_Gsc (E val: 4.0356e-09)
Pitx1	MA0234.1_oc (E val: 1.8154e-08)
Dobox5 CONSTRUCT	MA0190.1_Gsc (E val: 2.6002e-08)
<u>Obox6</u>	MA0190.1_Gsc (E val: 2.0177e-08)
Obox1	MA0190.1_Gsc (E val: 3.5326e-08)

<u>DPRX</u>	MA0190.1_Gsc (E val: 2.8082e-08)
RHOXF1	MA0234.1_oc (E val: 4.4835e-08)
CRX	MA0212.1_bcd (E val: 2.6743e-08)
<u>bcd</u>	MA0234.1_oc (E val: 5.4501e-08)
<u>ceh-36</u>	MA0234.1_oc (E val: 2.6099e-05)
Otx2	MA0234.1_oc (E val: 2.8635e-05)
<u>Crx</u>	MA0234.1_oc (E val: 2.9955e-05)
Dmbx1	MA0234.1_oc (E val: 4.4797e-05)

<u>dharma</u>	MA0234.1_oc (E val: 4.8065e-04)
LHX6	MA0235.1_onecut (E val: 2.8624e-04)
<u>Duxbl1</u>	MA0235.1_onecut (E val: 4.4831e-04)
<u>WOX13</u>	MA0070.1_PBX1 (E val: 5.5638e-06)
PBX1	MA0070.1_PBX1 (E val: 1.9530e-07)
Q00354 COPCI	MA0316.1_HAP5 (E val: 3.4094e-09)
Q00352 COPCI	MA0060.1_NFYA (E val: 4.3583e-09)
Q68SS9 PLEDJ	MA0315.1_HAP4 (E val: 9.4207e-07)

HAT1	MA0235.1_onecut (E val: 1.4678e-04)
hbx5-2	MA0235.1_onecut (E val: 9.7419e-08)
hbx5-1	MA0070.1_PBX1 (E val: 4.1702e-10)
ATHB-7	MA0070.1_PBX1 (E val: 1.0559e-04)
HAT3	MA0235.1_onecut (E val: 2.3651e-05)
GRMZM2G038252	MA0426.1_YHP1 (E val: 9.8215e-06)
ENSTNIG00000017092	MA0235.1_onecut (E val: 1.9058e-05)
SNOG 08903	MA0235.1_onecut (E val: 1.5205e-10)

NEMVEDRAFT v1g1016	664 MA0070.1_PBX1 (E val: 5.5008e-06)
Lhx6 CONSTRUCT	MA0075.1_Prrx2 (E val: 2.2088e-04)
Cphx1	MA0235.1_onecut (E val: 1.1410e-03)
<u>ATHB-6</u>	MA0110.1_ATHB-5 (E val: 6.6140e-05)
Q9C1N5 COPSC	MA0037.1_GATA3 (E val: 5.0444e-03)
AAEL012091	MA0212.1_bcd (E val: 1.3723e-02)
BARX1	MA0151.1_ARID3A (E val: 1.2246e-06)
Msx3	MA0192.1_Hmx (E val: 1.0690e-08)

Q0ZPQ4 NEMVE	MA0127.1_PEND (E val: 1.6660e-08)
TLX3	MA0188.1_Dr (E val: 1.1926e-03)
<u>Lhx8</u>	MA0198.1_OdsH (E val: 9.5457e-08)
ventx1.1.L	MA0075.1_Prrx2 (E val: 1.2557e-02)
<u>Hlx</u>	MA0075.1_Prrx2 (E val: 9.7967e-03)
Lmx1b	MA0075.1_Prrx2 (E val: 2.9556e-03)
Lmx1a	MA0075.1_Prrx2 (E val: 5.3755e-03)
<u>kal-1</u>	MA0075.1_Prrx2 (E val: 6.1057e-03)

<u>PHO2</u>	MA0075.1_Prrx2 (E val: 1.2104e-02)
<u>ceh-51</u>	MA0075.1_Prrx2 (E val: 4.0765e-02)
<u>UM04928</u>	MA0075.1_Prrx2 (E val: 1.3088e-02)
LHX2	MA0075.1_Prrx2 (E val: 8.0794e-03)
Q0N4H9 NEMVE	MA0075.1_Prrx2 (E val: 2.6414e-02)
NEMVEDRAFT v1g129868	MA0075.1_Prrx2 (E val: 5.9220e-03)
NEMVEDRAFT v1g136480	MA0094.1_Ubx (E val: 7.7218e-03)
GSC2	MA0094.1_Ubx (E val: 2.0696e-03)

<u>Isl2</u>	MA0094.1_Ubx (E val: 3.3589e-03)
Obox3	MA0094.1_Ubx (E val: 1.7438e-03)
Msx1	MA0094.1_Ubx (E val: 1.2509e-03)
Hoxc5	MA0094.1_Ubx (E val: 1.3094e-03)
POU5F1B	MA0075.1_Prrx2 (E val: 3.0457e-03)
<u>Dlx4</u>	MA0075.1_Prrx2 (E val: 2.1990e-03)
<u>E5</u>	MA0075.1_Prrx2 (E val: 1.9975e-03)
<u>Isx</u>	MA0094.1_Ubx (E val: 2.0861e-03)

Q6WL90 TRICY	MA0075.1_Prrx2 (E val: 1.9260e-03)
VAX2	MA0075.1_Prrx2 (E val: 2.0276e-03)
<u>Pdx 1</u>	MA0094.1_Ubx (E val: 2.5080e-03)
Rhox6	MA0094.1_Ubx (E val: 2.0443e-03)
Evx2	MA0075.1_Prrx2 (E val: 2.3487e-03)
<u>Otp</u>	MA0075.1_Prrx2 (E val: 2.0878e-03)
<u>Phox2a</u>	MA0094.1_Ubx (E val: 2.1722e-03)
Phox2b	MA0075.1_Prrx2 (E val: 2.3015e-03)

Obox5	MA0094.1_Ubx (E val: 2.4391e-03)
Barx2	MA0094.1_Ubx (E val: 3.1310e-03)
<u>Vax1</u>	MA0075.1_Prrx2 (E val: 3.3455e-03)
Dlx3	MA0075.1_Prrx2 (E val: 3.2354e-03)
barx1	MA0075.1_Prrx2 (E val: 3.1848e-03)
Vax2	MA0075.1_Prrx2 (E val: 3.3359e-03)
Hoxb3	MA0075.1_Prrx2 (E val: 3.1781e-03)
Hoxa5	MA0094.1_Ubx (E val: 3.3308e-03)

Hoxb4	MA0094.1_Ubx (E val: 3.3626e-03)
Hoxc4	MA0094.1_Ubx (E val: 3.5159e-03)
Evx1	MA0075.1_Prrx2 (E val: 3.1475e-03)
Hoxa4	MA0094.1_Ubx (E val: 3.2531e-03)
<u>Gsx2</u>	MA0075.1_Prrx2 (E val: 3.0869e-03)
Emx2	MA0075.1_Prrx2 (E val: 3.2531e-03)
<u>Nkx1-2</u>	MA0075.1_Prrx2 (E val: 3.3875e-03)
<u>Nkx6-1</u>	MA0075.1_Prrx2 (E val: 3.4714e-03)

NEMVEDRAFT v1g81309	MA0075.1_Prrx2 (E val: 3.3645e-03)
<u>Bsx</u>	MA0075.1_Prrx2 (E val: 3.4670e-03)
<u>Lbx2</u>	MA0094.1_Ubx (E val: 5.3979e-03)
<u>ceh-5</u>	MA0075.1_Prrx2 (E val: 3.4502e-03)
<u>Hoxb5</u>	MA0094.1_Ubx (E val: 3.7479e-03)
<u>Nkx6-3</u>	MA0075.1_Prrx2 (E val: 3.8618e-03)
<u>Nobox</u>	MA0094.1_Ubx (E val: 4.8980e-03)
Meox1	MA0075.1_Prrx2 (E val: 4.5297e-03)

Nanog	MA0075.1_Prrx2 (E val: 4.5812e-03)
CBG20882	MA0094.1_Ubx (E val: 4.1346e-03)
<u>Hoxb8</u>	MA0094.1_Ubx (E val: 6.4583e-03)
<u>Hoxb6</u>	MA0094.1_Ubx (E val: 2.5913e-03)
<u>Hoxb7</u>	MA0094.1_Ubx (E val: 3.4932e-03)
<u>Hoxd8</u>	MA0094.1_Ubx (E val: 4.7931e-03)
<u>Hoxa7</u>	MA0094.1_Ubx (E val: 5.6775e-03)
YOX1	MA0094.1_Ubx (E val: 1.1609e-02)

<u>Hoxa6</u>	MA0094.1_Ubx (E val: 3.7532e-03)
<u>Hoxc8</u>	MA0094.1_Ubx (E val: 8.9132e-03)
Barx1	MA0094.1_Ubx (E val: 4.4172e-03)
<u>Hoxa3</u>	MA0075.1_Prrx2 (E val: 4.2577e-03)
Mnx1	MA0075.1_Prrx2 (E val: 4.4611e-03)
ZHD7	MA0075.1_Prrx2 (E val: 5.0331e-03)
<u>Nkx1-1</u>	MA0075.1_Prrx2 (E val: 4.6337e-03)
<u>CBG00023</u>	MA0094.1_Ubx (E val: 5.4028e-03)

<u>Hmx2</u>	MA0426.1_YHP1 (E val: 3.6198e-04)
<u>Hmx1</u>	MA0426.1_YHP1 (E val: 3.2593e-04)
<u>Prop1</u>	MA0075.1_Prrx2 (E val: 1.9940e-03)
<u>En1</u>	MA0075.1_Prrx2 (E val: 2.1945e-03)
<u>UNCX</u>	MA0075.1_Prrx2 (E val: 1.8805e-03)
<u>Noto</u>	MA0075.1_Prrx2 (E val: 6.3461e-03)
<u>yox1</u>	MA0075.1_Prrx2 (E val: 7.7965e-03)
<u>Q0N4H7 NEMVE</u>	MA0094.1_Ubx (E val: 2.9184e-02)

NOBOX	MA0426.1_YHP1 (E val: 3.2827e-02)
<u>CG15696</u>	MA0426.1_YHP1 (E val: 2.0724e-02)
<u>CG34031</u>	MA0426.1_YHP1 (E val: 2.0324e-02)
NEMVEDRAFT v1g80394	MA0426.1_YHP1 (E val: 1.3574e-02)
pha-2	MA0426.1_YHP1 (E val: 4.0154e-02)
<u>Hoxb9</u>	MA0165.1_Abd-B (E val: 1.0286e-03)
CBG23031	MA0174.1_CG42234 (E val: 8.4823e-04)
Hoxc11	MA0165.1_Abd-B (E val: 2.3036e-03)

Hoxc13	MA0165.1_Abd-B (E val: 1.2421e-03)
Hoxb13	MA0165.1_Abd-B (E val: 1.5614e-03)
Hoxa13	MA0165.1_Abd-B (E val: 9.6505e-04)
HOXC13	MA0216.1_cad (E val: 1.0465e-08)
HOXA13	MA0216.1_cad (E val: 2.1130e-08)
HOXD13	MA0216.1_cad (E val: 3.3250e-08)
HOXB13	MA0216.1_cad (E val: 1.6334e-08)
Hoxd13	MA0216.1_cad (E val: 3.4315e-10)

Hoxd9	MA0165.1_Abd-B (E val: 4.1215e-07)
HOXC12	MA0216.1_cad (E val: 4.7854e-09)
hoxd13a	MA0216.1_cad (E val: 8.3883e-11)
<u>cad</u>	MA0216.1_cad (E val: 2.2162e-08)
HOXD9	MA0216.1_cad (E val: 6.3975e-10)
HOXD10	MA0216.1_cad (E val: 5.7397e-10)
CDX1	MA0216.1_cad (E val: 1.0263e-07)
HOXC10	MA0216.1_cad (E val: 2.9939e-09)

<u>ceh-90</u>	MA0393.1_STE12 (E val: 1.3427e-03)
HOXB9	MA0165.1_Abd-B (E val: 2.2251e-06)
Hoxc10	MA0165.1_Abd-B (E val: 2.4823e-08)
Abd-B	MA0165.1_Abd-B (E val: 1.4554e-06)
HOXA10	MA0165.1_Abd-B (E val: 7.2371e-06)
HOXA11	MA0165.1_Abd-B (E val: 2.0449e-05)
Hoxa11	MA0165.1_Abd-B (E val: 6.8791e-06)
HOXD11	MA0153.1_HNF1B (E val: 2.3770e-05)

HOXC11	MA0165.1_Abd-B (E val: 6.4582e-06)
HOXD12	MA0165.1_Abd-B (E val: 1.5126e-05)
HOXA9	MA0165.1_Abd-B (E val: 4.9453e-07)
CDX2	MA0165.1_Abd-B (E val: 1.8242e-04)
Hoxc12	MA0151.1_ARID3A (E val: 3.9486e-02)
Hoxd12 CONSTRUCT	MA0151.1_ARID3A (E val: 4.2453e-02)
<u>B-H2</u>	MA0168.1_B-H1 (E val: 3.6424e-04)
<u>B-H1</u>	MA0168.1_B-H1 (E val: 3.8028e-04)

BARHL1	MA0168.1_B-H1 (E val: 5.0670e-05)
Barhl1	MA0168.1_B-H1 (E val: 2.9661e-04)
Q0N4H2 NEMVE	MA0188.1_Dr (E val: 5.7239e-10)
NEMVEDRAFT v1g114268	MA0188.1_Dr (E val: 6.9261e-05)
NK7.1	MA0054.1_myb.Ph3 (E val: 5.3691e-06)
BARHL2	MA0054.1_myb.Ph3 (E val: 6.5933e-07)
<u>Hoxa10</u>	MA0307.1_GLN3 (E val: 8.7847e-03)
<u>Hoxa9</u>	MA0307.1_GLN3 (E val: 6.1601e-03)

Hoxd10	MA0307.1_GLN3 (E val: 4.1907e-03)
Hoxd11	MA0307.1_GLN3 (E val: 1.4126e-02)
Cdx2	MA0307.1_GLN3 (E val: 8.2090e-03)
Cdx1	MA0307.1_GLN3 (E val: 7.8786e-03)
<u>BSX</u>	MA0217.1_caup (E val: 4.8362e-03)
PRRX2	MA0186.1_Dfd (E val: 2.6340e-03)
HMBOX1	MA0231.1_lbe (E val: 4.3822e-07)
<u>lbe</u>	MA0186.1_Dfd (E val: 4.2043e-04)

LBX2	MA0186.1_Dfd (E val: 5.8346e-08)
<u>Hnf1b</u>	MA0217.1_caup (E val: 4.2044e-03)
<u>Hnfla</u>	MA0217.1_caup (E val: 2.6865e-03)
<u>ceh-57</u>	MA0217.1_caup (E val: 6.2734e-03)
<u>Hmbox1</u>	MA0231.1_lbe (E val: 9.7179e-02)
MTR 2g038000	MA0132.1_Pdx1 (E val: 2.8654e-02)
<u>ceh-24</u>	MA0211.1_bap (E val: 5.8321e-02)
ENSCING00000007395	MA0211.1_bap (E val: 3.0174e-02)

<u>ceh-22</u>	MA0211.1_bap (E val: 4.4240e-08)
<u>vnd</u>	MA0253.1_vnd (E val: 1.1102e-16)
<u>tin</u>	MA0264.1_ceh-22 (E val: 3.1586e-09)
<u>bap</u>	MA0211.1_bap (E val: 2.1113e-06)
<u>ISL1</u>	MA0211.1_bap (E val: 7.7296e-07)
ISL2	MA0211.1_bap (E val: 6.2608e-07)
HMX1	MA0063.1_Nkx2-5 (E val: 2.2710e-06)
HMX2	MA0211.1_bap (E val: 1.1516e-08)

NKX3-2	MA0211.1_bap (E val: 2.0976e-06)
NKX3-1	MA0211.1_bap (E val: 2.0256e-06)
<u>Nkx2-6</u>	MA0211.1_bap (E val: 7.4956e-04)
NKX2-3	MA0211.1_bap (E val: 1.0660e-03)
<u>Nkx2-9</u>	MA0211.1_bap (E val: 1.9645e-03)
<u>Nkx2-3</u>	MA0211.1_bap (E val: 9.1394e-04)
Nkx2-1	MA0211.1_bap (E val: 9.7504e-04)
<u>NKX2-5</u>	MA0211.1_bap (E val: 7.6109e-05)

<u>NKX2-8</u>	MA0211.1_bap (E val: 2.4048e-03)
<u>Nkx2-2</u>	MA0211.1_bap (E val: 2.2528e-03)
Nkx3-2	MA0211.1_bap (E val: 6.6546e-04)
<u>scro</u>	MA0211.1_bap (E val: 1.4217e-03)
<u>Nkx2-5</u>	MA0211.1_bap (E val: 1.5526e-03)
Nkx2-4	MA0211.1_bap (E val: 2.2018e-03)
<u>Nkx3-1</u>	MA0211.1_bap (E val: 8.3180e-04)
Q70HR3 PLEPI	MA0124.1_NKX3-1 (E val: 1.7532e-06)

<u>Six6</u>	MA0246.1_so (E val: 1.1002e-02)
six3	MA0246.1_so (E val: 5.1282e-02)
XP 003689450.1	MA0036.1_GATA2 (E val: 5.8039e-02)
SIX6	MA0199.1_Optix (E val: 1.5619e-05)
SIX3	MA0246.1_so (E val: 4.5416e-07)
Six3	MA0199.1_Optix (E val: 3.4735e-03)
PRH PETCR	MA0345.1_NHP6A (E val: 4.6352e-09)
<u>NCU09556</u>	MA0379.1_SIG1 (E val: 2.9552e-06)

hbx2	MA0379.1_SIG1 (E val: 6.3617e-06)
KLLA0 D10043g	MA0379.1_SIG1 (E val: 8.4022e-03)
NEMVEDRAFT v1g81436	MA0216.1_cad (E val: 9.7454e-02)
Pknox1	MA0227.1_hth (E val: 7.0473e-03)
<u>unc-62</u>	MA0207.1_achi (E val: 1.5344e-02)
<u>ATEG 00670</u>	MA0227.1_hth (E val: 3.4042e-02)
IRX2	MA0207.1_achi (E val: 6.9992e-04)
<u>vis</u>	MA0252.1_vis (E val: 8.8622e-07)

<u>achi</u>	MA0252.1_vis (E val: 8.7480e-07)
Meis2	MA0252.1_vis (E val: 6.4391e-06)
MEIS1	MA0252.1_vis (E val: 6.6419e-06)
Meis3	MA0227.1_hth (E val: 6.6077e-06)
<u>hbx9</u>	MA0288.1_CUP9 (E val: 9.3246e-06)
CUP9	MA0288.1_CUP9 (E val: 3.7091e-11)
<u>exd</u>	MA0222.1_exd (E val: 2.5740e-10)
<u>Meis1</u>	MA0252.1_vis (E val: 4.1357e-05)

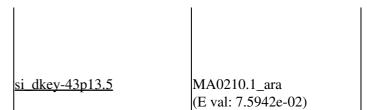
Tgif2	MA0227.1_hth (E val: 6.9961e-05)
<u>Tgif1</u>	MA0227.1_hth (E val: 6.8698e-05)
<u>hbx4</u>	MA0227.1_hth (E val: 1.0419e-04)
<u>CAGL0H02959g</u>	MA0227.1_hth (E val: 1.1990e-04)
ANIA 00885	MA0207.1_achi (E val: 1.9445e-04)
<u>GRMZM2G087741</u>	MA0227.1_hth (E val: 1.3141e-06)
<u>GRMZM2G135447</u>	MA0227.1_hth (E val: 5.9773e-09)
<u>CBG24578</u>	MA0207.1_achi (E val: 1.4139e-07)

TGIF2	MA0227.1_hth (E val: 3.1704e-07)
TGIF1	MA0207.1_achi (E val: 1.8587e-07)
MEIS2	MA0207.1_achi (E val: 3.0941e-07)
PKNOX1	MA0227.1_hth (E val: 5.8002e-07)
PKNOX2	MA0227.1_hth (E val: 5.7190e-07)
TGIF2LY	MA0227.1_hth (E val: 3.0965e-07)
TGIF2LX	MA0227.1_hth (E val: 2.5463e-07)
Pknox2	MA0207.1_achi (E val: 1.0777e-07)

IRX5	MA0328.1_MATALPHA2 (E val: 3.9553e-05)
<u>caup</u>	MA0217.1_caup (E val: 1.6159e-06)
<u>CG11617</u>	MA0173.1_CG11617 (E val: 1.4526e-07)
Rhox8	MA0392.1_STB5 (E val: 2.3740e-05)
<u>ara</u>	MA0210.1_ara (E val: 1.6159e-06)
<u>AGAP009986</u>	MA0210.1_ara (E val: 2.8411e-02)
<u>Irx5</u>	MA0210.1_ara (E val: 1.3053e-02)
<u>Irx6</u>	MA0210.1_ara (E val: 5.7942e-03)

<u>Irx2</u>	MA0210.1_ara (E val: 1.6628e-02)
Rhox11	MA0210.1_ara (E val: 2.0801e-02)
SNAPOd2T00005194001	MA0328.1_MATALPHA2 (E val: 1.7656e-04)
ATHB-4	MA0254.1_vvl (E val: 8.1236e-04)
<u>PK24181.1</u>	MA0227.1_hth (E val: 4.3791e-02)
pbx4	MA0235.1_onecut (E val: 3.6885e-04)
TVAG 451170	MA0246.1_so (E val: 1.2975e-03)
TOS8	MA0018.2_CREB1 (E val: 8.6372e-07)

PBX4	MA0204.1_Six4 (E val: 3.4604e-02)
<u>CBG10835</u>	MA0199.1_Optix (E val: 1.5682e-02)
<u>Phyra73700</u>	MA0199.1_Optix (E val: 1.5829e-02)
KNAT6	MA0207.1_achi (E val: 1.6623e-02)
Q0ZPQ8 NEMVE	MA0207.1_achi (E val: 1.1053e-02)
RLT1	MA0127.1_PEND (E val: 3.5933e-02)
<u>PK25011.1</u>	MA0089.1_NFE2L1_MafG (E val: 1.5545e-01)
<u>ATHB-12</u>	MA0089.1_NFE2L1_MafG (E val: 2.0267e-01)



Motif Similarity Matches

PHDP

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{lll} MA0229.1_inv & 1.9109e\text{-}06 & & \text{NTAATNNNATTAN} \\ & ---\text{WYTAATTA-} \end{array}$

hbn

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0229.1_inv & 1.1169e-09 & & NTAATTTAATTAN \\ & & ----WYTAATTA- \end{array}$

CG5369

forward

reverse compliment

Name E value Alignment Motif

Optix

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0229.1_inv} & 3.1478\text{e-}11 & & \text{TAATTARATTA} \\ & & \text{TAATTARW}-- \end{array}$

bsh

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0229.1_inv} & 3.9607\text{e-}11 & & \text{TAATTARATTA} \\ & & \text{TAATTARW}-- \end{array}$

unc-4

forward reverse compliment

Name E value Alignment Motif

MA0178.1_CG32105 1.8305e-08 TAATTNAATTA TAATTAA----

Arx

forward reverse compliment

Name E value Alignment Motif

MA0194.1_Lim1 1.0070e-08 NTAATTNAATTAN -TAATTAA----

Uncx

forward reverse compliment

Name E value Alignment Motif

MA0172.1_CG11294 3.7957e-09 NTAATTNRATTAN
-TAATTAA----

Alx4

forward reverse compliment

Name E value Alignment Motif

MA0194.1_Lim1 6.3149e-08 NTAATYNAATTAN -TAATTAA----

CG2808

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0188.1_Dr} & \text{1.4849e-08} & & \text{TAATTNSATTA} \\ & & \text{TAATTGG----} \end{array}$

CG11294

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0229.1_inv} & \text{6.6941e-10} & & \text{TAATNNAATTA} \\ & & --\text{WYTAATTA} \end{array}$

ALX1

forward reverse compliment

MA0229.1_inv 1.5023e-08

NTAAYNYAATTAN
----WYTAATTA-

CG32532

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 5.3536e-07 NTAATNNNATTAN ----TMATTA-

al

forward reverse compliment

Name E value Alignment Motif

MA0172.1_CG11294 1.2922e-05 TAATNNNATTA TAATTAA----

PROP1

forward reverse compliment

MA0188.1_Dr 2.8776e-09

NAAYNNAATTA ----CCAATTA

Otx1

forward reverse compliment

Name E value Alignment Motif

MA0234.1_oc 2.4512e-07 NNTAATCCGATTANN --TAATCC-----

OTX2

forward reverse compliment

Name E value Alignment Motif

MA0234.1_oc 2.2972e-07 NNTAATCGGATTANN ------GGATTA--

Prrx1

forward reverse compliment

MA0075.1_Prrx2 1.7693e-06

TAATTNNN TAATT---

Dlx5

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 1.2097e-06 WAATNNN TAATT--

Alx3

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 8.5973e-07 TAATNNN
TAATT--

Msx2

forward reverse compliment

MA0075.1_Prrx2 1.9540e-06

TAATTRNN TAATT---

Pph13

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 1.7716e-06 NTAATTRN
-TAATT--

ALX4

forward reverse compliment

Name E value Alignment Motif

PHOX2A

forward reverse compliment

MA0075.1_Prrx2 1.9368e-06

NTAATTNN -TAATT--

NEMVEDRAFT_v1g105595

forward reverse compliment

Name E value Alignment Motif

NTAATTNN MA0075.1_Prrx2 2.2012e-06 -TAATT--

Rax

forward reverse compliment

E value Name Alignment Motif

NNTAATTNN MA0075.1_Prrx2 3.8995e-06

--TAATT--

Lhx1

forward reverse compliment

E value Motif Name Alignment

MA0075.1_Prrx2 5.0294e-06 TAATNNNNN

TAATT----

LHX1

forward reverse compliment

Name E value Alignment Motif

Lim1

forward reverse compliment

Name E value Alignment Motif

PHOX2B

forward reverse compliment

MA0426.1_YHP1 5.2955e-08

NYAATTAN -CAATTA-

ESX1

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 9.7762e-08 NYAATTAN -CAATTA-

OdsH

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 2.3854e-08 NYAATTAN -CAATTA-

otp

forward reverse compliment

MA0426.1_YHP1 2.8789e-08

NYAATTAN -CAATTA-

Dll

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 2.2713e-08 NYAATTAN -CAATTA-

Vsx2

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 2.3819e-08 NYAATTAN -CAATTA-

DLX5

forward reverse compliment

MA0426.1_YHP1 2.1206e-08

NTAATTGN -TAATTG-

DLX4

forward reverse compliment

Name E value Alignment Motif

NCAATTAN MA0426.1_YHP1 2.6660e-08 -CAATTA-

DLX6

forward reverse compliment

E value Motif Name Alignment

NCAATTAN MA0426.1_YHP1 2.4257e-08 -CAATTA-

DLX1

forward reverse compliment

Name E value Motif Alignment

MA0426.1_YHP1 3.6604e-08 NYAATTAN

-CAATTA-

GBX2

forward reverse compliment

Name E value Alignment Motif

lms

forward reverse compliment

Name E value Alignment Motif

Rx

forward reverse compliment

MA0426.1_YHP1 2.5179e-08

NYAATTAN -CAATTA-

EN1

forward reverse compliment

Name E value Alignment Motif

Vsx1

forward reverse compliment

Name E value Alignment Motif

MSX1

forward reverse compliment

MA0426.1_YHP1 1.1714e-08

NCAATTAN -CAATTA-

MSX2

forward

reverse compliment

Name E value

Alignment Motif

MA0426.1_YHP1 1.1888e-08

NCAATTAN -CAATTA-

en

forward

reverse compliment

Name

E value

Alignment

Motif

MA0426.1_YHP1 6.3159e-08

NTAATTRN -TAATTG-

Dlx1

forward

reverse compliment

Name

E value

Alignment

Motif

MA0426.1_YHP1 7.3537e-08

NYAATTAN -CAATTA-

Dlx2

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 8.3123e-08 NYAATTAN -CAATTA-

Dr

forward reverse compliment

Name E value Alignment Motif

SNAPOd2T00006620001

forward reverse compliment

MA0426.1_YHP1 5.3836e-08

NTAATTRN -TAATTG-

DLX3

forward reverse compliment

E value Name Alignment Motif

NYAATTAN MA0232.1_lbl 1.1929e-07 -TMATTA-

SHOX2

forward reverse compliment

E value Motif Name Alignment

NYAATTAN MA0232.1_lbl 1.2365e-07

-TMATTA-

Shox2

forward reverse compliment

Name E value Motif Alignment

MA0232.1_lbl 9.8309e-08 NYAATTAN

-TMATTA-

TLX2

	forward	reverse compliment	
Name	E value	Alignment	Motif
MA0232.1_lbl	8.2750e-08	NYAATTAN -TMATTA-	

DLX2

	forward	reverse compliment	
Name	E value	Alignment	Motif
MA0232.1_lbl	1.1128e-07	NYAATTAN -TMATTA-	

RAX2

	jorwara	reverse compument	
Name	E value	Alignment	Motif
MA0232.1_lbl	1.1674e-07	NTAATTRN	
		-TAATKA-	

PRRX1

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 6.1316e-08 NTAATTRN -TAATKA-

ceh-54

forward reverse compliment

Name E value Alignment Motif

Gbx2

forward reverse compliment

MA0426.1_YHP1 2.9462e-07

NNYAATTANN --CAATTA--

Gbx1

forward reverse compliment

Name E value Alignment Motif

CG11085

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 2.0173e-08 NCRATTAN -CAATTA-

Hmx

forward reverse compliment

MA0426.1_YHP1 4.3245e-08

NTAAKTGN -TAATTG-

AAZ08039

forward

reverse compliment

Name E value

Alignment

MA0426.1_YHP1 2.2824e-07

NNTAATTRNN --TAATTG--

GBX1

forward

reverse compliment

Name E value

Alignment

Motif

Motif

MA0232.1_lbl 4.9552e-07

NSYAATTANN --TMATTA--

CG9876

forward

reverse compliment

Name E value

Alignment

Motif

MA0426.1_YHP1 4.0548e-07

NNYRATTANS
--CAATTA--

unpg

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 2.0776e-07 NNCAATTANN --CAATTA--

ceh-58

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 4.0935e-07 NNYAATTANN --TMATTA--

Barhl2

forward reverse compliment

MA0426.1_YHP1 5.7920e-07

NNYNNTTAN
--CAATTA-

ceh-19

forward reverse compliment

Name E value Alignment Motif

YHP1

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 5.2801e-06 NNYAATWWN --TMATTA-

ATEG_04171

forward reverse compliment

MA0232.1_lbl 4.6903e-06

NNNYAATTWN
---TMATTA-

NCU03266

forward

reverse compliment

Name E value

Alignment

MA0196.1_NK7.1 1.9654e-08

NYNATTWNNN -CAATTAA--

tup

forward

reverse compliment

Name

E value

Alignment

Motif

Motif

MA0248.1_tup 2.1446e-11

SWTAATKG -WTAATKG

LHX4

forward

reverse compliment

Name

E value

Alignment

Motif

MA0232.1_lbl 1.2360e-07

NTAATTAN -TMATTA-

EMX1

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 1.3018e-08 NTAATTAN -TMATTA-

lab

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 1.0849e-08 STAATTAN -TMATTA-

CG18599

forward reverse compliment

MA0232.1_lbl 1.9400e-09

TAATTAN TMATTA-

Meox2

forward

reverse compliment

Name E value

Alignment

MA0232.1_lbl 7.1346e-09

NTAATTAN -TMATTA-

ems

forward

reverse compliment

Name E value

Alignment

Motif

Motif

MA0232.1_lbl 9.6821e-09

NTAATTAN -TMATTA-

EVX1

forward

E value

reverse compliment

Name

Alignment

Motif

NTAATTAN MA0232.1_lbl 1.2369e-08 -TMATTA-

EVX2

E value Name Alignment Motif

NTAATTAN MA0232.1_lbl 1.2336e-08 -TAATKA-

forward

Lim3

reverse compliment

reverse compliment

E value Motif Name Alignment

NTAATTAN MA0232.1_lbl 2.1239e-08 -TAATKA-

forward

Lhx4

forward reverse compliment

Name E value Motif Alignment

MA0232.1_lbl 2.0476e-08 NTAATTAN

-TMATTA-

Q9GP48_9TURB

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 3.8246e-08 NTAATTAN -TMATTA-

Q70LF3_ARTSF

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0232.1_lbl} & 2.7533\text{e-}08 & & \text{NTAATTAN} \\ & & -\text{TMATTA-} \end{array}$

NEMVEDRAFT_v1g205636

forward reverse compliment

MA0232.1_lbl 3.7288e-08

NTAATTAN -TMATTA-

Esx1

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 3.5689e-08 NNTAATTN --TMATTA

MNX1

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 2.6559e-08 NTAATTAN
-TMATTA-

DRGX

forward reverse compliment

MA0232.1_lbl 3.1653e-08

NYAATTAN -TMATTA-

EN2

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 3.0403e-08 NTAATTAN -TMATTA-

Prrx2

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 2.7448e-08 NTAATTRN -TAATKA-

ALX3

forward reverse compliment

MA0232.1_lbl 2.1881e-08 NTAATTRN -TAATKA-

SHOX

forward reverse compliment

E value Name Alignment Motif

NYAATTAN MA0232.1_lbl 2.2701e-08 -TMATTA-

Awh

forward reverse compliment

E value Motif Name Alignment

NTAATTAN 1.6501e-08 MA0232.1_lbl

-TMATTA-

Lhx3

forward reverse compliment

Name E value Motif Alignment

MA0232.1_lbl 6.3907e-08 NNATTANNN

TAATKA---

uncx

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 9.0879e-08 NNTAATTRN --TAATKA-

Lhx5

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 7.9458e-08 NNNYAATTA ---TMATTA

zfhx4

forward reverse compliment

MA0232.1_lbl 5.4071e-08

NYAATTANN -TMATTA--

Q4RRY4_TETNG

forward

reverse compliment

Name E value

Alignment

MA0232.1_lbl 2.1430e-07

NNYAATYAN --TMATTA-

Hoxa1

forward

reverse compliment

Name

E value

Alignment

Motif

Motif

MA0232.1_lbl 3.8787e-08

NNYNATTNN --TMATTA-

HOXA1

forward

reverse compliment

Name

E value

Alignment

Motif

09/21/20

MA0232.1_lbl 1.4927e-07 NNTAATTANN --TAATKA--

prd-b

forward reverse compliment

Name E value Alignment Motif

NTRATTRNN MA0232.1_lbl 7.3149e-07 -TAATKA--

Lhx6

forward reverse compliment

Name E value Alignment Motif

NYAATYAN MA0232.1_lbl 1.2291e-06

-TMATTA-

ceh-23

forward reverse compliment

Alignment Name E value Motif

09/21/20

MA0232.1_lbl 1.3081e-07

NTAATTANNN -TAATKA---

Alx1

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 1.4374e-07 NNTAATTANN --TMATTA--

MIXL1

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0232.1_lbl} & 2.4876\text{e-07} & & \text{NNYAATTANN} \\ & & -\text{TMATTA--} \end{array}$

En2

forward reverse compliment

MA0232.1_lbl 6.0426e-08 NSTAATTANN --TMATTA--

slou

forward reverse compliment

Name E value Alignment Motif

NNYAATTANN MA0232.1_lbl 1.4101e-07 --TMATTA--

GSX1

forward reverse compliment

Name E value Alignment Motif

NTMATNAN MA0232.1_lbl 6.6879e-08

-TAATKA-

HOXB2

forward reverse compliment

Alignment Name E value Motif MA0232.1_lbl 1.5599e-08

NYAATTAN -TMATTA-

NOTO

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0232.1_lbl} & 1.6063\text{e-}08 & & \text{NYAATTAN} \\ & & -\text{TMATTA-} \end{array}$

HOXB7

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 1.8210e-08 NYAATTAN -TMATTA-

NKX6-2

forward reverse compliment

NTAATTRN MA0232.1_lbl 1.8473e-08 -TAATKA-

Ubx

forward reverse compliment

E value Name Alignment Motif

NYAATTAN MA0232.1_lbl 1.5127e-08 -TMATTA-

PDX1

forward reverse compliment

E value Motif Name Alignment

YTAATKAN MA0232.1_lbl 1.1612e-08

-TAATKA-

HOXA2

forward reverse compliment

Name E value Motif Alignment

MA0232.1_lbl 8.6640e-09 NTAATTAN

-TMATTA-

Lhx2

	forward	reverse compliment	
Name	E value	Alignment	Motif
MA0232.1_lbl	9.1589e-09	NYMATTAN -TMATTA-	

HGTX

	forward	reverse compliment	
Name	E value	Alignment	Motif
MA0232.1_lbl	6.2738e-09	NTMATTAN -TMATTA-	

HOXB5

	forward	reverse compliment	
Name	E value	Alignment	Motif

MA0232.1_lbl	5.6550e-08	NYMATTAN
	3.03306-08	-TMATTA-

HOXA7

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 1.0318e-07 NTAATKRN -TAATKA-

HOXD4

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 6.8867e-08 NYMATTAN -TMATTA-

Scr

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 4.3130e-08 NYMATTAN

-TMATTA-

HOXB8

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 1.3096e-07 NTAATKRC -TAATKA-

HOXD8

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 2.2035e-07 NTAATKRC -TAATKA-

HOXB6

forward reverse compliment

MA0232.1_lbl 3.1744e-08

GYMATTAN -TMATTA-

HOXA5

forward reverse compliment

Name E value Alignment Motif

Q0N4M8_NEMVE

forward reverse compliment

Name E value Alignment Motif

MA0223.1_exex 2.3856e-09 GTAATTAN NTAATTA-

HOXA6

forward reverse compliment

Name E value Alignment Motif

MA0223.1_exex 8.5000e-08 NNTAATKRCN --TAATTAN-

abd-A

reverse compliment forward

E value Motif Name Alignment

NGTAATTANN MA0223.1_exex 1.9332e-08 -NTAATTA--

HOXA4

forward reverse compliment

Alignment Name E value Motif

NTAATKAY MA0232.1_lbl 8.1005e-08

-TAATKA-

pb

forward reverse compliment

MA0232.1_lbl 6.8619e-08

GTMATTAN
-TMATTA-

HOXB3

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 5.6179e-08 NNYMATTANN --TMATTA--

NEMVEDRAFT_v1g8907

forward reverse compliment

Name E value Alignment Motif

Hoxd3

forward reverse compliment

09/21/20

MA0232.1_lbl 1.0207e-07

NNYAATTANN --TMATTA--

Hoxa2

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 3.9527e-08 NNTAATTANN --TMATTA--

MEOX2

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0232.1_lbl} & 7.0922\text{e-}08 & & \text{NNTAATTANN} \\ & & -\text{TMATTA--} \end{array}$

eve

forward reverse compliment

MA0241.1_ro 7.8255e-10 NSTAATTAGN --TAATTAR-

GSX2

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0232.1_lbl & 8.5976e\text{-}09 & & \text{NTAATKRS} \\ & & -\text{TAATKA-} \end{array}$

MEOX1

forward reverse compliment

Name E value Alignment Motif

MA0223.1_exex 5.9674e-09 NNTAATTASN --TAATTAN-

VSX2

forward reverse compliment

Name E value Alignment Motif

MA0241.1_ro 2.7339e-11 NTAATTAG -TAATTAR

ARX

forward reverse compliment

Name E value Alignment Motif

MA0241.1_ro 3.0360e-11 YTAATTAN YTAATTA-

ISX

forward reverse compliment

Name E value Alignment Motif

MA0241.1_ro 8.0527e-11 CTAATTAN YTAATTA-

VENTX

forward reverse compliment

MA0241.1_ro 9.6463e-11

CTAATTAN YTAATTA-

EMX2

forward reverse compliment

Name E value Alignment Motif

MA0241.1_ro 2.5200e-11 NTAATTAG -TAATTAR

ro

forward reverse compliment

Name E value Alignment Motif

MA0241.1_ro 1.2763e-11 TAATTARN TAATTAR-

LBX1

forward reverse compliment

MA0241.1_ro 4.2517e-10

NTAATTAG -TAATTAR

pdm3

forward reverse compliment

Name E value Alignment Motif

lbl

forward reverse compliment

Name E value Alignment Motif

MA0241.1_ro 1.9839e-09 NTAATKAG -TAATTAR

dve

forward reverse compliment

MA0241.1_ro 9.7944e-10

NCTAATTANN -YTAATTA--

Q9BJW6_9CNID

forward reverse compliment

Name E value Alignment Motif

MA0241.1_ro 8.0265e-10 NNTAATTAGN --TAATTAR-

HESX1

forward reverse compliment

Name E value Alignment Motif

MA0188.1_Dr 1.7446e-09 NTAATTRGN -TAATTGG-

exex

forward reverse compliment

Name E value Alignment Motif

MA0198.1_OdsH 1.1913e-08 NNNNYAATTNNCNN ----YAATTAR---

pal-1

forward reverse compliment

Name E value Alignment Motif

MA0448.1_H2.0 2.2106e-10 TTWATTAC NTWATNA-

hox12

forward reverse compliment

Name E value Alignment Motif

MA0174.1_CG42234 1.3600e-10 GTMATWAA -TMATWAA

HXD12_HETFR

forward reverse compliment

Name E value Alignment Motif

MA0206.1_abd-A 9.0750e-11 GTAATTAA -TAATTAA

ftz

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0226.1_hbn & 2.0192e\text{-}11 & & \text{GTAATTAA} \\ & & -\text{YAATTAA} \end{array}$

Hoxc6

forward reverse compliment

Name E value Alignment Motif

Hoxc9

forward reverse compliment

7.4846e-10 MA0240.1_repo

NGTAATTAAN --TAATTAA-

H2.0

forward reverse compliment

Name E value Alignment Motif

TNATWAA MA0448.1_H2.0 2.5989e-11 TNATWAN

Antp

forward reverse compliment

E value Motif Name Alignment

TMATTAAR MA0166.1_Antp 1.2763e-11

TMATTAA-

C15

forward reverse compliment

E value Name Motif Alignment

MA0170.1_C15 1.2763e-11 NTTAAWKR

-TTAAWKR

Q24780_9CNID

forward reverse compliment

Name E value Alignment Motif

MA0194.1_Lim1 7.3681e-10 NNTAATTAANN --TAATTAA--

Q0N4F9_NEMVE

forward reverse compliment

Name E value Alignment Motif

P79681_ACIBE

forward reverse compliment

MA0180.1_Vsx2 6.4537e-13

NTAATTAA-CTAATTAAM

SPU_026099

forward reverse compliment

Name E value Alignment Motif

MA0180.1_Vsx2 8.9306e-13 NTAATTAA-CTAATTAAM

vab-7

forward reverse compliment

Name E value Alignment Motif

ENSTNIG00000004116

forward reverse compliment

MA0180.1_Vsx2 5.9242e-13

CTAATTAA— CTAATTAAM

Smp_158000

forward reverse compliment

Name E value Alignment Motif

MA0180.1_Vsx2 4.7229e-13 CYAATTAA-CTAATTAAM

NEMVEDRAFT_v1g164989

forward reverse compliment

Name E value Alignment Motif

MA0180.1_Vsx2 7.4326e-12 CTAATTAR-CTAATTAAM

ARGFX

forward reverse compliment

Name E value Alignment Motif

MA0180.1_Vsx2 1.2971e-10 NCTAATTARN -CTAATTAAM

vsx2

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0180.1_Vsx2} & 3.4137\text{e-}10 & & \text{NYTAATTAANN} \\ & -\text{CTAATTAAM-} \end{array}$

Q24782_9CNID

forward reverse compliment

Name E value Alignment Motif

MA0194.1_Lim1 1.4038e-11 TTAATTAA -TAATTAA

CG32105

forward reverse compliment

MA0236.1_otp 3.4164e-11

TTAATTAR -TAATTAR

repo

forward

reverse compliment

Name E value

Alignment

MA0240.1_repo 2.8598e-10

NYTAATTAAN --TAATTAA-

CNK00090

forward

reverse compliment

Name

E value

Alignment

Motif

Motif

MA0240.1_repo 2.3502e-11

ATAATTAA -TAATTAA

HOXC4

forward

reverse compliment

Name

E value

Alignment

Motif

MA0240.1_repo 2.0401e-11

NTAATTAA -TAATTAA

CAD25284

forward

reverse compliment

Name E value

Alignment Motif

MA0194.1_Lim1 1.5897e-11

TTAATTAN TTAATTA-

LMX1A

forward

reverse compliment

Name

E value

Alignment

Motif

MA0172.1_CG11294 1.7100e-11

TTAATTAN TTAATTA-

Tlx2

forward

reverse compliment

Name

E value

Alignment

Motif

MA0194.1_Lim1 2.7621e-10

TAWTTAAT TAATTAA-

ventx3.2

forward reverse compliment

Name E value Alignment Motif

CBG00029

forward reverse compliment

Name E value Alignment Motif

HHEX

forward reverse compliment

MA0172.1_CG11294 2.1773e-10

TAATTNAW TAATTAA-

ENSCING00000016045

forward reverse compliment

Name E value Alignment Motif

MA0180.1_Vsx2 5.3273e-10 NCYAATTARNN -CTAATTAAM-

RAX

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0202.1_Rx & 5.1507e\text{-}11 & & \text{NTAATTAR} \\ & & -\text{TAATTAR} \end{array}$

inv

forward reverse compliment

MA0229.1_inv 6.5503e-15

TAATTARW TAATTARW

ap

forward reverse compliment

Name E value Alignment Motif

MA0209.1_ap 1.2763e-11 TAATTARN TAATTAR-

lim-4

forward reverse compliment

Name E value Alignment Motif

ind

forward reverse compliment

MA0228.1_ind 7.6496e-11

TAATTARNR TAATTAR--

Sebox

forward reverse compliment

Name E value Alignment Motif

MA0194.1_Lim1 2.5830e-10 NTAATTAANN -TAATTAA--

mixl1.S

forward reverse compliment

Name E value Alignment Motif

MA0194.1_Lim1 6.0393e-10 NNTAATTAANN --TAATTAA--

C4B866_NEMVE

forward reverse compliment

MA0194.1_Lim1 1.2649e-09

NNNTAATTAANN
---TAATTAA--

CAD26261

forward reverse compliment

Name E value Alignment Motif

MA0178.1_CG32105 2.6988e-08 TWWATTANN TTAATTA--

LHX9

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 2.4650e-07 NNTNATTAN --TMATTA-

POU6F1

forward reverse compliment

MA0232.1_lbl 1.9095e-07

NNTAATTANN --TAATKA--

Q86SD7_PODCA

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 2.5359e-07 NNTAATTANN --TMATTA--

CBG07641

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0232.1_lbl} & 1.8462\text{e-07} & & \text{NNTAATTANN} \\ & -\text{TMATTA}-- & & & \end{array}$

ceh-62

forward reverse compliment

MA0232.1_lbl 2.4811e-07

NNTAATTANN --TMATTA--

ZFHX2

forward

reverse compliment

Name E value

Alignment

MA0232.1_lbl 2.6731e-07

NNTAATTANN --TAATKA--

btn

forward

reverse compliment

Name E value

Alignment

Motif

Motif

MA0232.1_lbl 7.5444e-08

NNTMATTANN
--TMATTA--

LMX1B

forward

reverse compliment

Name E value

Alignment

Motif

MA0232.1_lbl 2.8185e-07

NNNNTAATTAN
----TMATTA-

SNAPOd2T00006145001

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 2.6419e-06 NNYAATYANN --TMATTA--

HMX3

forward reverse compliment

Name E value Alignment Motif

SNAPOd2T00003340001

forward reverse compliment

MA0232.1_lbl 6.4014e-06 NWWATTANNN -TAATKA---

CG4328

forward reverse compliment

Name E value Alignment Motif

YAATWAWN MA0182.1_CG4328 1.4526e-07 YAATWAW-

hbx10

forward reverse compliment

Name E value Alignment Motif

NWWATTRN MA0182.1_CG4328 1.1851e-05

WTWATTR-

Q8I9J5_9BILA

forward reverse compliment

E value Name Alignment Motif

Motif

MA0448.1_H2.0 3.6600e-05

NRTWATTANN
-NTWATNA--

Q7M3U4_ACRFO

forward

reverse compliment

Name E value

Alignment

MA0075.1_Prrx2 1.9488e-05

NNNTAATTNNN ---TAATT---

ENSTNIG00000006040

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 8.7845e-07 NNNYNATTANNNN ---CAATTA----

CBG04010

forward reverse compliment

MA0206.1_abd-A 7.7487e-06

NNTTAATKNNN --TTAATTA--

CNG03450

forward reverse compliment

Name E value Alignment Motif

MA0444.1_CG34031 7.7102e-07 NNWATTAAN -CAATTAA-

HDG11

forward reverse compliment

Name E value Alignment Motif

MA0151.1_ARID3A 3.6185e-06 NCATTWAWKN --ATTAAA--

ANL₂

forward reverse compliment

MA0206.1_abd-A 4.3694e-06

NNAATTAATGN
-TAATTAA---

ATML1

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0186.1_Dfd & 8.4622e\text{-}05 & & \text{NNNANTNANN} \\ & & -\text{TCATTAA}-- & & \end{array}$

Dfd

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0232.1_lbl} & 1.7692\text{e-}06 & & \text{NNYCATTANN} \\ & -\text{TMATTA}-- & & \end{array}$

zen

forward reverse compliment

MA0232.1_lbl 7.9820e-08

NTAATKAN -TAATKA-

zen2

forward reverse compliment

Name E value Alignment Motif

MA0232.1_lbl 2.1091e-08 NTMATTAN -TMATTA-

NKX6-3

forward reverse compliment

Name E value Alignment Motif

MA0215.1_btn 3.0070e-09 NTCATTAWN
-TCATTAN-

HNF1B

forward reverse compliment

MA0153.1_HNF1B 5.5378e-13

RTTAATNATTAAC GTTAAWYATTAA-

HNF1A

forward reverse compliment

Name E value Alignment Motif

MA0153.1_HNF1B 6.3727e-14 NRTTAATNATTAACN --TTAATRWTTAAC-

ATHB-16

forward reverse compliment

Name E value Alignment Motif

NANOG

forward reverse compliment

MA0214.1_bsh 1.0306e-04

NCNRTYAN -YMATTAA

NEMVEDRAFT_v1g113057

forward reverse compliment

Name E value Alignment Motif

MA0212.1_bcd 2.6743e-08 NNGGATTARN --GGATTA--

HBX4_ECHGR

forward reverse compliment

Name E value Alignment Motif

MA0190.1_Gsc 3.2658e-08 NNGGATTANN
--GGATTA--

--GGATTA--

 \mathbf{oc}

forward reverse compliment

MA0212.1_bcd 3.8618e-09

NGGATTAN -GGATTA-

GSC

forward reverse compliment

Name E value Alignment Motif

MA0234.1_oc 3.9560e-09 NTAATCCN -TAATCC-

Gsc

forward reverse compliment

Name E value Alignment Motif

MA0234.1_oc 9.9606e-10 NGGATTA -GGATTA

Q8VHG7_MOUSE

forward reverse compliment

MA0201.1_Ptx1 2.4983e-10

NGGATTAW -GGATTAA

PITX1

forward reverse compliment

Name E value Alignment Motif

MA0234.1_oc 3.8797e-09 NGGATTAN -GGATTA-

CBG01820

forward reverse compliment

Name E value Alignment Motif

MA0212.1_bcd 3.9289e-09 NGGATTAN -GGATTA-

PITX3

forward reverse compliment

MA0234.1_oc 3.7633e-09

NGGATTAN -GGATTA-

OTX1

forward reverse compliment

Name E value Alignment Motif

MA0234.1_oc 3.7603e-09 NTAATCCN -TAATCC-

Ptx1

forward reverse compliment

Name E value Alignment Motif

MA0234.1_oc 5.0663e-09 NGGATTAN
-GGATTA-

Pitx2

forward reverse compliment

MA0234.1_oc 4.9980e-09

NGGATTAN -GGATTA-

Pitx3

forward

reverse compliment

Name E value Alignment

Motif

MA0234.1_oc 6.8909e-09

NGGATTAN -GGATTA-

PITX2

forward

reverse compliment

Name E value

Alignment

Motif

MA0234.1_oc 7.9895e-09

NTAATCCN -TAATCC-

DMBX1

forward

reverse compliment

Name E value

Alignment

Motif

MA0234.1_oc 8.8704e-09

NGGATTAN -GGATTA-

ceh-53

forward reverse compliment

Name E value Alignment Motif

MA0190.1_Gsc 4.6874e-09 NRGATTAN -GGATTA-

ceh-37

forward reverse compliment

Name E value Alignment Motif

MA0190.1_Gsc 4.0372e-09 NGGATTAN
-GGATTA-

AGAP006642

forward reverse compliment

MA0190.1_Gsc 4.0356e-09

GGATTANN GGATTA--

Pitx1

forward reverse compliment

Name E value Alignment Motif

MA0234.1_oc 1.8154e-08 NNGGATTAN --GGATTA-

Dobox5_CONSTRUCT

forward reverse compliment

Name E value Alignment Motif

MA0190.1_Gsc 2.6002e-08 NTAATCYNN -TAATCC--

Obox6

forward reverse compliment

MA0190.1_Gsc 2.0177e-08

NNRGATTAN
--GGATTA-

Obox1

forward reverse compliment

Name E value Alignment Motif

MA0190.1_Gsc 3.5326e-08 NNNGGATTAN ---GGATTA-

DPRX

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0190.1_Gsc & 2.8082e\text{-}08 & \\ & -\text{GGATTA---} \end{array}$

-GGATTA--

RHOXF1

forward reverse compliment

4.4835e-08 MA0234.1_oc

GGATNATCN GGATTA---

CRX

forward reverse compliment

Name E value Alignment Motif

NSGGATTAGN MA0212.1_bcd 2.6743e-08 --GGATTA--

bcd

forwardreverse compliment

Name E value Alignment Motif

NNGGATTANNN MA0234.1_oc 5.4501e-08

--GGATTA---

ceh-36

forward reverse compliment

MA0234.1_oc 2.6099e-05

NNGATTAN -GGATTA-

Otx2

forward

reverse compliment

Name E value

Alignment Motif

MA0234.1_oc 2.8635e-05

NNGATTAN -GGATTA-

Crx

forward

reverse compliment

Name

E value

Alignment

Motif

MA0234.1_oc 2.9955e-05

NNGATTAN -GGATTA-

Dmbx1

forward

reverse compliment

Name E value

Alignment

Motif

MA0234.1_oc 4.4797e-05

NNRGATTAN
--GGATTA-

dharma

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0234.1_oc & 4.8065e\text{-}04 & & \text{NNNATCC} \\ & -\text{TAATCC} \end{array}$

LHX6

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0235.1_onecut} & 2.8624\text{e-}04 & & \text{NYAATCAN} \\ & & -\text{NAATCAA} \end{array}$

Duxbl1

forward reverse compliment

NNYAATCAN --NAATCAA

WOX13

forward reverse compliment

Name E value Alignment Motif

MA0070.1_PBX1 5.5638e-06 NNTGATTGAN-NTTGATTGATNN

PBX1

forward reverse compliment

Name E value Alignment Motif

MA0070.1_PBX1 1.9530e-07 -NTGATTGAYR-NTTGATTGATNN

Q00354_COPCI

forward reverse compliment

MA0316.1_HAP5 3.4094e-09

ACCAATCN----ACCAATNAGMNNSSN

Q00352_COPCI

forward

reverse compliment

Name E value

Alignment

Motif

MA0060.1_NFYA 4.3583e-09

---TGATTGGTT---NNNCYSATTGGYYNNN

Q68SS9_PLEDJ

forward

reverse compliment

Name

E value

Alignment

Motif

MA0315.1_HAP4 9.4207e-07

NNCAATCN----NCCAATCARNNNNN

HAT1

forward

reverse compliment

Name E value

Alignment

Motif

NYAATCATNN -NAATCAA--

hbx5-2

forward reverse compliment

Name E value Alignment Motif

MA0235.1_onecut 9.7419e-08 NYAATCAW -NAATCAA

hbx5-1

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \textbf{MA0070.1_PBX1} & \textbf{4.1702e-10} & & - \textbf{TTGATTGR---} \\ & & \textbf{NTTGATTGATNN} \end{array}$

ATHB-7

forward reverse compliment

MA0070.1_PBX1 1.0559e-04

NNATGATTGN---NTTGATTGATNN

HAT3

forward reverse compliment

Name E value Alignment Motif

MA0235.1_onecut 2.3651e-05 NYAATCAN
-NAATCAA

GRMZM2G038252

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 9.8215e-06 NCAATCAN -CAATTA-

ENSTNIG00000017092

forward reverse compliment

MA0235.1_onecut 1.9058e-05

NTAATCAN -NAATCAA

SNOG_08903

forward reverse compliment

Name E value Alignment Motif

NEMVEDRAFT_v1g101664

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0070.1_PBX1 & 5.5008e\text{-}06 & & & --\text{-}GATYGATT-\\ & & & NTTGATTGATNN \end{array}$

Lhx6_CONSTRUCT

forward reverse compliment

MA0075.1_Prrx2 2.2088e-04

NNTRATTRN
--TAATT--

Cphx1

forward reverse compliment

Name E value Alignment Motif

MA0235.1_onecut 1.1410e-03 NNNNNNATCW----NAATCAA

ATHB-6

forward reverse compliment

Name E value Alignment Motif

MA0110.1_ATHB-5 6.6140e-05 NNAATNATN---AATMATTGN

Q9C1N5_COPSC

forward reverse compliment

MA0037.1_GATA3 5.0444e-03

NNCAATCNN --YTATCN-

AAEL012091

forward reverse compliment

Name E value Alignment Motif

MA0212.1_bcd 1.3723e-02 NNGATNNNN
-GGATTA--

BARX1

forward reverse compliment

Name E value Alignment Motif

Msx3

forward reverse compliment

MA0192.1_Hmx 1.0690e-08

NCAATTANNNNSCAATTA -CAATTAA----

Q0ZPQ4_NEMVE

forward

reverse compliment

Name E value

Alignment

MA0127.1_PEND 1.6660e-08

-TTAAGAAG-MATAAGAANT

TLX3

forward

reverse compliment

Name E value

Alignment

Motif

Motif

MA0188.1_Dr 1.1926e-03

NAATTGNNNNNNNNNNNNNCAATTN TAATTGG-----

Lhx8

forward

reverse compliment

Name

E value

Alignment

Motif

MA0198.1_OdsH 9.5457e-08

NTRATTANNNTRATTAN
-----YTAATTR-

ventx1.1.L

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 1.2557e-02 NNNNNNNNN --TAATT---

Hlx

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 9.7967e-03 NNWTNNN TAATT--

Lmx1b

forward reverse compliment

MA0075.1_Prrx2 2.9556e-03

NTNATNN -TAATT-

Lmx1a

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 5.3755e-03 NYNATNNN -TAATT--

kal-1

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 6.1057e-03 NNNNANNN --TAATT-

PHO₂

forward reverse compliment

Name E value Alignment Motif

ceh-51

forward reverse compliment

Name E value Alignment Motif

UM04928

forward reverse compliment

Name E value Alignment Motif

LHX2

forward reverse compliment

MA0075.1_Prrx2 8.0794e-03

NNNNAANNNN ---TAATT--

Q0N4H9_NEMVE

forward

reverse compliment

Name E value

Alignment

MA0075.1_Prrx2 2.6414e-02

NNYRANNNN --TAATT--

NEMVEDRAFT_v1g129868

forward

reverse compliment

Name

E value

Alignment

Motif

Motif

MA0075.1_Prrx2 5.9220e-03

NYNANNNNN -TAATT---

NEMVEDRAFT_v1g136480

forward

reverse compliment

Name

E value

Alignment

Motif

MA0094.1_Ubx 7.7218e-03

NNNNNNNNN ----ATTA--

GSC₂

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 2.0696e-03 NTAATNNN -TAAT---

Isl2

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 3.3589e-03 NNNTNNN --ATTA-

Obox3

forward reverse compliment

MA0094.1_Ubx 1.7438e-03

NNATNWN
--ATTA-

Msx1

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 1.2509e-03 NNATTAN --ATTA-

Hoxc5

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 1.3094e-03 NNATTNN --ATTA-

POU5F1B

forward reverse compliment

MA0075.1_Prrx2 3.0457e-03

NNAANNN -TAATT-

Dlx4

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 2.1990e-03 NTAATNNN -TAATT--

E5

forward reverse compliment

Name E value Alignment Motif

Isx

forward reverse compliment

MA0094.1_Ubx 2.0861e-03

NNATTNNN --ATTA--

Q6WL90_TRICY

forward reverse compliment

Name E value Alignment Motif

VAX2

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 2.0276e-03 NNTAATNN --TAATT-

Pdx1

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 2.5080e-03 NNNATTAN ---ATTA-

Rhox6

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 2.0443e-03 NTAATNNN -TAAT---

Evx2

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 2.3487e-03 NNATTANN -AATTA--

Otp

forward reverse compliment

MA0075.1_Prrx2 2.0878e-03

NNTAATNN --TAATT-

Phox2a

forward reverse compliment

E value Name Alignment Motif

NNATTNNN MA0094.1_Ubx 2.1722e-03 --ATTA--

Phox2b

reverse compliment forward

Motif E value Name Alignment

NNNAATNN MA0075.1_Prrx2 2.3015e-03 --TAATT-

Obox5

forward reverse compliment

Name E value Motif Alignment

MA0094.1_Ubx 2.4391e-03 NNNATNAN

---ATTA-

Barx2

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 3.1310e-03 NNNNATTAN ----ATTA-

Vax1

forward reverse compliment

Name E value Alignment Motif

Dlx3

forward reverse compliment

MA0075.1_Prrx2 3.2354e-03

NTAATNNNN -TAATT---

barx1

forward reverse compliment

Name E value Alignment Motif

Vax2

forward reverse compliment

Name E value Alignment Motif

Hoxb3

forward reverse compliment

MA0075.1_Prrx2 3.1781e-03

NTAATNNNN -TAATT---

Hoxa5

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 3.3308e-03 NTAATNNNN -TAAT----

Hoxb4

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 3.3626e-03 NTAATNNNN -TAAT---

Hoxc4

forward reverse compliment

MA0094.1_Ubx 3.5159e-03

NNNNATTAN
----ATTA-

Evx1

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 3.1475e-03 NTAATNNNN -TAATT---

Hoxa4

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 3.2531e-03 NTAATNNNN -TAAT----

Gsx2

forward reverse compliment

MA0075.1_Prrx2 3.0869e-03

NTAATNNNN -TAATT---

Emx2

forward reverse compliment

Name E value Alignment Motif

Nkx1-2

forward reverse compliment

Name E value Alignment Motif

Nkx6-1

forward reverse compliment

MA0075.1_Prrx2 3.4714e-03

NNTAATNRN --TAATT--

NEMVEDRAFT_v1g81309

forward reverse compliment

Name E value Alignment Motif

Bsx

forward reverse compliment

Name E value Alignment Motif

Lbx2

forward reverse compliment

MA0094.1_Ubx 5.3979e-03

NNAANNNN -TAAT---

ceh-5

forward reverse compliment

Name E value Alignment Motif

Hoxb5

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 3.7479e-03 NTNATNNNN -TAAT----

Nkx6-3

forward reverse compliment

MA0075.1_Prrx2 3.8618e-03

NNNATNANN --AATTA--

Nobox

forward reverse compliment

Name E value Alignment Motif

Meox1

forward reverse compliment

Name E value Alignment Motif

Nanog

forward reverse compliment

MA0075.1_Prrx2 4.5812e-03

NNNATNNNN -TAATT---

CBG20882

forward

reverse compliment

Name E value

Alignment

MA0094.1_Ubx 4.1346e-03

NYNATNNNN ---ATTA--

Hoxb8

forward

reverse compliment

Name

E value

Alignment

Motif

Motif

MA0094.1_Ubx 6.4583e-03

NNNNATWNN ----ATTA-

Hoxb6

forward

reverse compliment

Name

E value

Alignment

Motif

MA0094.1_Ubx 2.5913e-03

NNNNNNN TAAT---

Hoxb7

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 3.4932e-03 NNNNTNNN ---ATTA-

Hoxd8

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 4.7931e-03 NNNNNNNN ---ATTA-

Hoxa7

forward reverse compliment

MA0094.1_Ubx 5.6775e-03

NNWATNNNN -TAAT----

YOX1

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 1.1609e-02 YNATWWN --ATTA-

Hoxa6

forward reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 3.7532e-03 NNNNATTAN ----ATTA-

Hoxc8

forward reverse compliment

MA0094.1_Ubx 8.9132e-03

NNNNNNNN ----ATTA-

Barx1

forward reverse compliment

Name E value Alignment Motif

Hoxa3

forward reverse compliment

Name E value Alignment Motif

Mnx1

forward reverse compliment

MA0075.1_Prrx2 4.4611e-03

NTAATNNNNN -TAATT----

ZHD7

forward reverse compliment

Name E value Alignment Motif

Nkx1-1

forward reverse compliment

Name E value Alignment Motif

CBG00023

forward reverse compliment

MA0094.1_Ubx 5.4028e-03

NNNNNATWAN
----ATTA-

Hmx2

forward reverse compliment

Name E value Alignment Motif

Hmx1

forward reverse compliment

Name E value Alignment Motif

Prop1

forward reverse compliment

MA0075.1_Prrx2 1.9940e-03

NNNATYN -TAATT-

En1

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 2.1945e-03 NNNANNN -TAATT-

UNCX

forward reverse compliment

Name E value Alignment Motif

Noto

forward reverse compliment

MA0075.1_Prrx2 6.3461e-03

NNNNATNNNN ---AATTA--

yox1

forward reverse compliment

Name E value Alignment Motif

MA0075.1_Prrx2 7.7965e-03 NNNNATNNNN --TAATT---

Q0N4H7_NEMVE

forward

reverse compliment

Name E value Alignment Motif

MA0094.1_Ubx 2.9184e-02 NNNNNATNNN -----ATTA-

NOBOX

forward reverse compliment

MA0426.1_YHP1 3.2827e-02

NNYNATNNNN --CAATTA--

CG15696

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 2.0724e-02 NNYAATNNN --CAATTA-

CG34031

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 2.0324e-02 NYNNNNNN -CAATTA-

NEMVEDRAFT_v1g80394

forward reverse compliment

Name E value Alignment Motif

MA0426.1_YHP1 1.3574e-02 NNNNATNNN --CAATTA-

pha-2

forward reverse compliment

Name E value Alignment Motif

Hoxb9

forward reverse compliment

Name E value Alignment Motif

MA0165.1_Abd-B 1.0286e-03 NNNNRTWAAN --TMATAAA-

CBG23031

forward reverse compliment

MA0174.1_CG42234 8.4823e-04

NNNNATWAAN --TMATWAA-

Hoxc11

forward reverse compliment

Name E value Alignment Motif

NNNNRTWAAN MA0165.1_Abd-B 2.3036e-03 --TMATAAA-

Hoxc13

forward reverse compliment

E value Name Alignment Motif

NNNNRTAAAN MA0165.1_Abd-B 1.2421e-03

--TMATAAA-

Hoxb13

forward reverse compliment

Name E value Motif Alignment

MA0165.1_Abd-B 1.5614e-03 NNNNRTWAAN

--TMATAAA-

Hoxa13

forward reverse compliment

Name E value Alignment Motif

MA0165.1_Abd-B 9.6505e-04 NNNRTWAAN
-TMATAAA-

HOXC13

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0216.1_cad} & \text{1.0465e-08} & & \text{NSCAATAAAAN} \\ & & -\text{YAATAAA--} \end{array}$

HOXA13

forward reverse compliment

MA0216.1_cad 2.1130e-08

GTTTTATTGGN
--TTTATTR--

HOXD13

forward reverse compliment

Name E value Alignment Motif

HOXB13

forward reverse compliment

Name E value Alignment Motif

Hoxd13

forward reverse compliment

Name E value Alignment Motif

MA0216.1_cad 3.4315e-10 SYAATAAAAN -YAATAAA--

Hoxd9

forward reverse compliment

Name E value Alignment Motif

MA0165.1_Abd-B 4.1215e-07 NCCATWAAA -TMATAAA-

HOXC12

forward reverse compliment

Name E value Alignment Motif

hoxd13a

forward reverse compliment

MA0216.1_cad 8.3883e-11

TMATAAAW YAATAAA-

cad

forward reverse compliment

Name E value Alignment Motif

MA0216.1_cad 2.2162e-08 NNGCMATAAAAN ---YAATAAA--

HOXD9

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0216.1_cad} & 6.3975\text{e-}10 & & \text{GCAATAAAAN} \\ & -\text{YAATAAA--} & & \end{array}$

HOXD10

forward reverse compliment

MA0216.1_cad 5.7397e-10

GYMATAAAAN -YAATAAA--

CDX1

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0216.1_cad} & \text{1.0263e-07} & & \text{NTTTTATGGCN} \\ & & -\text{TTTATR--} \end{array}$

HOXC10

forward reverse compliment

Name E value Alignment Motif

MA0216.1_cad 2.9939e-09 NNTTTATKRCN --TTTATTR--

ceh-90

forward reverse compliment

MA0393.1_STE12 1.3427e-03

NGTTTTANN YGTTTCA--

HOXB9

forward reverse compliment

Name E value Alignment Motif

MA0165.1_Abd-B 2.2251e-06 GTCGTAAAAN
-TMATAAA--

Hoxc10

forward reverse compliment

Name E value Alignment Motif

MA0165.1_Abd-B 2.4823e-08 NTTTTAYGAC --TTTATKA-

Abd-B

forward reverse compliment

MA0165.1_Abd-B 1.4554e-06

NGTCGTAAAWN --TMATAAA--

HOXA10

forward reverse compliment

Name E value Alignment Motif

NTTTTACGACN MA0165.1_Abd-B 7.2371e-06 --TTTATKA--

HOXA11

forward reverse compliment

E value Motif Name Alignment

NGTCGTAAAAN MA0165.1_Abd-B 2.0449e-05

--TMATAAA--

Hoxa11

forward reverse compliment

MA0165.1_Abd-B 6.8791e-06

NNTCGTAAANNN --TMATAAA---

HOXD11

forward reverse compliment

Name E value Alignment Motif

MA0153.1_HNF1B 2.3770e-05

NRTCGTAAANNNTWAN
----GTTAAWYATTAA

HOXC11

forward reverse compliment

Name E value Alignment Motif

MA0165.1_Abd-B 6.4582e-06 NNTTTACGAYN --TTTATKA--

HOXD12

forward reverse compliment

Name E value Alignment Motif

MA0165.1_Abd-B 1.5126e-05 NRTCGTAAAN --TMATAAA-

HOXA9

forward reverse compliment

Name E value Alignment Motif

NNNTTTAYGAYN MA0165.1_Abd-B 4.9453e-07 ---TTTATKA--

CDX2

forward reverse compliment

E value Motif Name Alignment

NRTYRTAANN MA0165.1_Abd-B 1.8242e-04 --TMATAAA-

Hoxc12

forward reverse compliment

Name E value Motif Alignment

MA0151.1_ARID3A 3.9486e-02 NNTWAYNNNN -TTTAAT---

Hoxd12_CONSTRUCT

forward reverse compliment

Name E value Alignment Motif

MA0151.1_ARID3A 4.2453e-02 NNTNAYNNNN
-TTTAAT---

B-H2

forward reverse compliment

Name E value Alignment Motif

MA0168.1_B-H1 3.6424e-04 NNCGTTTANN --CAATTAA-

B-H1

forward reverse compliment

MA0168.1_B-H1 3.8028e-04

NNCGTTTANN --CAATTAA-

BARHL1

forward reverse compliment

Name E value Alignment Motif

MA0168.1_B-H1 5.0670e-05 NCGTTTAN -CAATTAA

Barhl1

forward reverse compliment

Name E value Alignment Motif

MA0168.1_B-H1 2.9661e-04 NNCRTTTANN --CAATTAA-

Q0N4H2_NEMVE

forward reverse compliment

MA0188.1_Dr 5.7239e-10

TAATTRNT TAATTGG-

NEMVEDRAFT_v1g114268

forward reverse compliment

Name E value Alignment Motif

MA0188.1_Dr 6.9261e-05 NTAATNGRT -TAATTGG-

NK7.1

forward reverse compliment

Name E value Alignment Motif

MA0054.1_myb.Ph3 5.3691e-06
-NACCGTTTAAN
TAACNGTTW---

BARHL2

forward reverse compliment

Name E value Alignment Motif

Hoxa10

forward reverse compliment

Name E value Alignment Motif

MA0307.1_GLN3 8.7847e-03 NNNNNTNNN ---GATAA-

Hoxa9

forward reverse compliment

Name E value Alignment Motif

MA0307.1_GLN3 6.1601e-03 NNNNNNNN -TTATC--

Hoxd10

forward reverse compliment

MA0307.1_GLN3 4.1907e-03

NNNNNNN -TTATC--

Hoxd11

forward reverse compliment

Name E value Alignment Motif

NNNNNNN MA0307.1_GLN3 1.4126e-02 -TTATC--

Cdx2

forward reverse compliment

E value Motif Name Alignment

NWWATNRNN MA0307.1_GLN3 8.2090e-03

-TTATC---

Cdx1

forward reverse compliment

Name E value Motif Alignment

MA0307.1_GLN3 7.8786e-03 NNWATNRNN

-TTATC---

BSX

forward reverse compliment

Name E value Alignment Motif

PRRX2

forward reverse compliment

Name E value Alignment Motif

MA0186.1_Dfd 2.6340e-03 NNCRTNAN -TCATTAA

HMBOX1

forward reverse compliment

MA0231.1_lbe 4.3822e-07

NTAACTAGN -TAAYNA--

lbe

forward reverse compliment

Name E value Alignment Motif

LBX2

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0186.1_Dfd & 5.8346e\text{-}08 & & \texttt{CTCRTTAA} \\ & & -\texttt{TCATTAA} \end{array}$

Hnf1b

forward reverse compliment

MA0217.1_caup 4.2044e-03

NNNGTNNN --TGTTA-

Hnf1a

forward reverse compliment

Name E value Alignment Motif

ceh-57

forward reverse compliment

Name E value Alignment Motif

Hmbox1

forward reverse compliment

MA0231.1_lbe 9.7179e-02

NNNACNNN -TAAYNA-

MTR_2g038000

forward

reverse compliment

Name E value

Alignment

Motif

MA0132.1_Pdx1 2.8654e-02

NNNNNAGNNN -AATTAG---

ceh-24

forward

reverse compliment

Name

E value

Alignment

Motif

MA0211.1_bap 5.8321e-02

NNYNCTYNN --CACTTAA

ENSCING00000007395

forward

reverse compliment

Name

E value

Alignment

Motif

MA0211.1_bap 3.0174e-02

NWNNNNNN CACTTAA-

ceh-22

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0211.1_bap} & \text{4.4240e-08} & & \text{NNNCACTTNA} \\ & --\text{CACTTAA} & & \end{array}$

vnd

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0253.1_vnd} & 1.1102\text{e-}16 & & \text{YACTTGARA} \\ & & \text{YACTTGARA} \end{array}$

tin

forward reverse compliment

Stamp F	Results
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MA0264.1_ceh-22 3.1586e-09

NCCACTTGAN-NCCACTTGANN

reverse compliment

reverse compliment

bap

Name	E value	Alignment	Motif

NNCACTTANN MA0211.1_bap 2.1113e-06 --CACTTAA-

forward

ISL1

E value Motif Name Alignment

SCACTTAN 7.7296e-07 MA0211.1_bap -CACTTAA

forward

ISL₂

forward reverse compliment Name E value Motif Alignment MA0211.1_bap 6.2608e-07 NCACTTAN -CACTTAA

HMX1

forward reverse compliment

Name E value Alignment Motif

MA0063.1_Nkx2-5 2.2710e-06 NCAMTTANN -CAMTTAW-

HMX2

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 1.1516e-08 NCACTTRNC -CACTTAA-

NKX3-2

forward reverse compliment

MA0211.1_bap 2.0976e-06

NCCACTTANN
--CACTTAA-

NKX3-1

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 2.0256e-06 NNTAAGTGGN -TTAAGTG--

Nkx2-6

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 7.4956e-04 NNNCACTTNN ---CACTTAA

NKX2-3

forward reverse compliment

MA0211.1_bap 1.0660e-03

NNCACTTNNN --CACTTAA-

Nkx2-9

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 1.9645e-03 NNRAGTRNNN TTAAGTG---

Nkx2-3

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 9.1394e-04 NNRAGNRNN TTAAGTG--

Nkx2-1

forward reverse compliment

MA0211.1_bap 9.7504e-04

NYNCTYNN -CACTTAA

NKX2-5

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0211.1_bap} & 7.6109\text{e-}05 & & \frac{\text{NNAAGWR}}{\text{TTAAGTG}} \end{array}$

NKX2-8

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 2.4048e-03 NYNCTYNN -CACTTAA

Nkx2-2

forward reverse compliment

MA0211.1_bap 2.2528e-03

NYNCTYNNN -CACTTAA-

Nkx3-2

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 6.6546e-04 NNYACTTNN --CACTTAA

scro

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 1.4217e-03 NCACTYNNN -CACTTAA-

Nkx2-5

forward reverse compliment

MA0211.1_bap 1.5526e-03

NNYNCTYNNN --CACTTAA-

Nkx2-4

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 2.2018e-03 NNRAGNRNNN TTAAGTG---

Nkx3-1

forward reverse compliment

Name E value Alignment Motif

MA0211.1_bap 8.3180e-04 NNNCACTTNN ---CACTTAA

Q70HR3_PLEPI

forward reverse compliment

MA0124.1_NKX3-1 1.7532e-06

NTAAGTANNN -TAAGTAT--

Six6

forward reverse compliment

Name E value Alignment Motif

MA0246.1_so 1.1002e-02 NNKATNN -GTATCA

six3

forward reverse compliment

Name E value Alignment Motif

MA0246.1_so 5.1282e-02 NNNATMNNNN
-TGATAC---

XP_003689450.1

forward reverse compliment

MA0036.1_GATA2 5.8039e-02

NNNGKNTNNN ---NGATN--

SIX6

forward reverse compliment

Name E value Alignment Motif

NNNTATCRNN MA0199.1_Optix 1.5619e-05 ---TATCA--

SIX3

forward reverse compliment

E value Motif Name Alignment

NNNTATCRNN MA0246.1_so 4.5416e-07 --GTATCA--

Six3

forward reverse compliment

Name E value Motif Alignment

MA0199.1_Optix 3.4735e-03 NKATNNN -TATCA-

PRH_PETCR

forward reverse compliment

Name E value Alignment Motif

NCU09556

forward reverse compliment

Name E value Alignment Motif

MA0379.1_SIG1 2.9552e-06 TTAATATA ---ATATA

hbx2

forward reverse compliment

MA0379.1_SIG1 6.3617e-06

NNATATATWN
---TATAT--

KLLA0_D10043g

forward

reverse compliment

Name E value

Alignment

Motif

MA0379.1_SIG1 8.4022e-03

NNNNNNNNN --TATAT---

NEMVEDRAFT_v1g81436

forward

reverse compliment

Name

E value

Alignment

Motif

MA0216.1_cad 9.7454e-02

NNNNNANNNN ---YAATAAA

Pknox1

forward

reverse compliment

Name

E value

Alignment

Motif

MA0227.1_hth 7.0473e-03

NNNTGTCN
--CTGTCA

unc-62

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0207.1_achi} & 1.5344\text{e-02} & & \text{NNNTNTNNN} \\ & & -\text{CTGTCA-} \end{array}$

ATEG_00670

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0227.1_hth} & 3.4042\text{e-}02 & & \text{NNNGTNNNN} \\ & & -\text{CTGTCA--} \end{array}$

IRX2

forward reverse compliment

MA0207.1_achi 6.9992e-04

NNTGTCRTGTWN
-CTGTCA----

vis

forward reverse compliment

Name E value Alignment Motif

MA0252.1_vis 8.8622e-07 NTGTCA NTGTCA

achi

forward reverse compliment

Name E value Alignment Motif

MA0252.1_vis 8.7480e-07 NTGTCA NTGTCA

Meis2

forward reverse compliment

6.4391e-06 MA0252.1_vis

NTGTCAN NTGTCA-

MEIS1

forward reverse compliment

E value Name Alignment Motif

NTGACAN MA0252.1_vis 6.6419e-06 -TGACAN

Meis3

forward reverse compliment

E value Motif Name Alignment

NTGTCAN 6.6077e-06 MA0227.1_hth CTGTCA-

hbx9

forward reverse compliment

Name E value Motif Alignment

MA0288.1_CUP9 9.3246e-06 ---NTGTCANN

WWTGTGTCA--

CUP9

forward reverse compliment

Name E value Alignment Motif

MA0288.1_CUP9 3.7091e-11 NWTGTGTCAN WWTGTGTCA-

exd

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0222.1_exd} & \text{2.5740e-10} & & \text{TGTCAAAN} \\ & & \text{TGTCAAAN} \end{array}$

Meis1

forward reverse compliment

NNNTGTCAN 4.1357e-05 MA0252.1_vis --NTGTCA-

Tgif2

forward reverse compliment

Name E value Alignment Motif

NNNTGTCANN MA0227.1_hth 6.9961e-05 --CTGTCA--

Tgif1

forward reverse compliment

Motif E value Name Alignment

NNNTGTCANN MA0227.1_hth 6.8698e-05

--CTGTCA--

hbx4

forward reverse compliment

Name E value Motif Alignment

MA0227.1_hth 1.0419e-04 NNNYGTCANN

--CTGTCA--

CAGL0H02959g

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0227.1_hth & 1.1990e\text{-}04 & & NNNYGTCANN \\ & & --\text{CTGTCA}\text{-}- \end{array}$

ANIA_00885

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} \text{MA0207.1_achi} & 1.9445\text{e-04} & & \text{NNNTGTMANN} \\ & & -\text{CTGTCA--} \end{array}$

GRMZM2G087741

forward reverse compliment

MA0227.1_hth 1.3141e-06

NGMYGTCANN --CTGTCA--

GRMZM2G135447

forward

reverse compliment

Name E value

Alignment

MA0227.1_hth 5.9773e-09

SMYGTCAN -CTGTCA-

CBG24578

forward

reverse compliment

Name

E value

Alignment

Motif

Motif

MA0207.1_achi 1.4139e-07

NNNTGACAGSTN
---TGACAG---

TGIF2

forward

reverse compliment

Name

E value

Alignment

Motif

MA0227.1_hth 3.1704e-07

TGACANNTGTCA
----CTGTCA

TGIF1

forward reverse compliment

Name E value Alignment Motif

MEIS2

forward reverse compliment

Name E value Alignment Motif

MA0207.1_achi 3.0941e-07 TGACAGCTGTCA TGACAG-----

PKNOX1

forward reverse compliment

MA0227.1_hth 5.8002e-07

TGACASSTGTCA
----CTGTCA

PKNOX2

forward reverse compliment

Name E value Alignment Motif

MA0227.1_hth 5.7190e-07 TGACANNTGTCA ----CTGTCA

TGIF2LY

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{lll} MA0227.1_hth & 3.0965e\text{-}07 & & \texttt{TGACANNTGTCA} \\ & -----\texttt{CTGTCA} & & \end{array}$

TGIF2LX

forward reverse compliment

MA0227.1_hth 2.5463e-07

TGACANCTGTCA
----CTGTCA

Pknox2

forward reverse compliment

Name E value Alignment Motif

 $\begin{array}{ccc} MA0207.1_achi & 1.0777e\text{-}07 & & \texttt{TGACAGGTGTCA} \\ & & \texttt{TGACAG-----} \end{array}$

IRX5

forward reverse compliment

Name E value Alignment Motif

MA0328.1_MATALPHA2 3.9553e-05

NNNTGTNNNACAYNN
----AATTACATG-

caup

forward reverse compliment

MA0217.1_caup 1.6159e-06

TGTTANWN TGTTA---

CG11617

forward reverse compliment

Name E value Alignment Motif

MA0173.1_CG11617 1.4526e-07 -TGTWAAAW ATGTWAA--

Rhox8

forward reverse compliment

Name E value Alignment Motif

MA0392.1_STB5 2.3740e-05 NTTACACCN -TANNNCCG

ara

forward reverse compliment

MA0210.1_ara 1.6159e-06

TGTTWTNN
TGTTW---

AGAP009986

forward

reverse compliment

Name E value

Alignment Motif

MA0210.1_ara 2.8411e-02

NNNNTGTNNN ----TGTTW-

Irx5

forward

reverse compliment

Name

E value

Alignment

Motif

MA0210.1_ara 1.3053e-02

NNTGTNNNN --TGTTW--

Irx6

forward

 $reverse\ compliment$

Name E value

Alignment

Motif

MA0210.1_ara 5.7942e-03

NTNTNNN -TGTTW-

Irx2

forward reverse compliment

Name E value Alignment Motif

Rhox11

forward reverse compliment

Name E value Alignment Motif

MA0210.1_ara 2.0801e-02 NNNTGTWNNN ---TGTTW--

SNAPOd2T00005194001

forward reverse compliment

MA0328.1_MATALPHA2 1.7656e-04

-NNTACATGTNN AATTACATG---

ATHB-4

forward reverse compliment

Name E value Alignment Motif

MA0254.1_vvl 8.1236e-04 NNYATGCAKNN --TATKCA---

PK24181.1

forward reverse compliment

Name E value Alignment Motif

pbx4

forward reverse compliment

NNCATCAA -NAATCAA

TVAG_451170

forward reverse compliment

Name E value Alignment Motif

NNCRCATCAN MA0246.1_so 1.2975e-03 ---GTATCA-

TOS8

forward reverse compliment

E value Name Alignment Motif

WKACGTCANNN MA0018.2_CREB1 8.6372e-07

TGACGTCA---

PBX4

forward reverse compliment

E value Motif Name Alignment

MA0204.1_Six4 3.4604e-02 NNNNNTNNNN

--GTNTCA--

CBG10835

forward reverse compliment

Name E value Alignment Motif

MA0199.1_Optix 1.5682e-02 NNNNTCANN --TATCA--

Phyra73700

forward reverse compliment

Name E value Alignment Motif

KNAT6

forward reverse compliment

MA0207.1_achi 1.6623e-02

NNNNGTCNNN --CTGTCA--

Q0ZPQ8_NEMVE

forward

reverse compliment

Name E value

Alignment

MA0207.1_achi 1.1053e-02

NTGACRNNN -TGACAG--

RLT1

forward

reverse compliment

Name

E value

Alignment

Motif

Motif

MA0127.1_PEND 3.5933e-02

--NANGAANN MATAAGAANT

PK25011.1

forward

reverse compliment

Name

E value

Alignment

Motif

MA0089.1_NFE2L1_MafG 1.5545e-01

NNNNNNAWNN
---GTCATN-

ATHB-12

forward reverse compliment

Name E value Alignment Motif

MA0089.1_NFE2L1_MafG 2.0267e-01

NNNNNNN
-NATGAC-

si_dkey-43p13.5

forward reverse compliment

Name E value Alignment Motif

MA0210.1_ara 7.5942e-02 NNNNWWNNNN --TGTTW---

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