2	gene score co	oublesex_DM.HG1.18:like:DMRT3 or: 0.04; expression cor: 0.04; max zscore: 8.43 ne score cor: 0.10; expression cor: -0.03; max zscore: 6.93
1- 0		Forkhead: M04857_2.00 Nvec_vc1.1_XM_001627792.3 Foxn4/Foxn1 r: -0.03; expression cor: 0.25; max zscore: 8.37 F; gene score cor: 0.31; expression cor: 0.10; max zscore: 7.21
2 1 0		Forkhead: M03058_2.00 Nvec_vc1.1_XM_001628319.3 Foxj3
2 1 0		: -0.05; expression cor: 0.24; max zscore: 10.16; gene score cor: -0.07; expression cor: 0.49; max zscore: 8.52 [RC] TGTTTATGTTAC Forkhead: M00164_2.00 Nvec_vc1.1_XM_032384126.2
2 1 0	gene score cor ARCH964_Foxc2/Foxc1_Foxg1/G	Nvec_vc1.1_xM_032384126.2 orkhead.HG1.17:like:FOXI1/FOXI2 r: 0.20; expression cor: 0.30; max zscore: 15.05; gene score cor: 0.45; expression cor: 0.17; max zscore: 4.90 [RC]
2 1-	ARCH1095_Foxa3/Foxa1/Foxa2	Forkhead: M03042_2.00 Nvec_vc1.1_XM_048724319.1 Foxd2/Foxd4/Foxd3/Foxd1 r: 0.12; expression cor: 0.27; max zscore: 13.26 g; gene score cor: 0.10; expression cor: 0.46; max zscore: 7.16
0 []]	Forkl gene score cor	Forkhead: M07960_2.00 Nvec_vc1.1_XM_048728051.1 head.HG1.1:FOXA1/FOXA2/FOXA3 r: 0.08; expression cor: 0.34; max zscore: 23.27 ene score cor: 0.42; expression cor: -0.12; max zscore: 9.18
1- 0		Grainyhead_LSF: M03769_2.00 Nvec_vc1.1_XM_001629572.3 Ubp1/Tfcp2/Tfcp2l1 r: 0.06; expression cor: -0.09; max zscore: 7.91
2 1 0		AACÇÇÜTT_AACÇÇÜTT_ HLH: M01726_2.00 Nvec_vc1.1_XM_032375349.2 HLH.HG1.50:HES3
2 1 0		: -0.17; expression cor: 0.35; max zscore: 25.28 .; gene score cor: 0.47; expression cor: 0.14; max zscore: 42.35 [RC] HLH: M08766_2.00 Nvec_vc1.1_XM_032384433.2
2 1 0		Twist2/Twist1 r: 0.64; expression cor: 0.15; max zscore: 40.46 8; gene score cor: 0.34; expression cor: 0.48; max zscore: 5.36 [RC]
2 1		HLH: M05990_2.00 Nvec_vc1.1_XM_032385496.2 Ptf1a r: 0.31; expression cor: 0.21; max zscore: 27.83 .; gene score cor: 0.37; expression cor: 0.21; max zscore: 42.35 [RC]
0 []]		HLH: M04211_2.00 Nvec_v1g210540 Ascl1/Ascl2 r: 0.54; expression cor: 0.40; max zscore: 39.36 .; gene score cor: 0.58; expression cor: 0.36; max zscore: 42.35 [RC]
1 0 2	gene score cor	HLH: M00588_2.00 Nvec_vc1.1_XM_001628300.3 G1.84:NEUROG1/NEUROG2/NEUROG3 r: 0.11; expression cor: 0.15; max zscore: 17.99 score cor: -0.12; expression cor: 0.51; max zscore: 3.81 [RC]
1		CAGE TGT HLH: M05989_2.00 Nvec_vc1.1_XM_001631403.3 Mlxipl/Mlxip : -0.03; expression cor: 0.31; max zscore: 23.69
2 1 0	ARCH1068_bHLH122; gene	score cor: -0.04; expression cor: 0.50; max zscore: 6.93 [RC] CACCTCA HLH: M03721_2.00 Nvec_vc1.1_XM_001634712.3 Ebf1/Ebf4/Ebf2/Ebf3
2 1 0	gene score cor	r: 0.36; expression cor: 0.22; max zscore: 20.21 [RC]
2 1 0		Nvec_vc1.1_XM_001635315.3 HLH.HG1.63:like:HES5/HES7 : -0.18; expression cor: 0.06; max zscore: 16.91 score cor: 0.35; expression cor: -0.00; max zscore: 5.31 [RC]
2 1		HLH: M05979_2.00 Nvec_vc1.1_XM_001636622.3
0 []]	gene score cor	HLH: M06004_2.00 Nvec_vc1.1_XM_001640990.3 .HG1.77:like:NEUROD1/NEUROD2/ r: 0.29; expression cor: 0.46; max zscore: 19.43; gene score cor: 0.27; expression cor: 0.42; max zscore: 42.35
1 d		HLH: M08734_2.00 Nvec_vc1.1_XM_032363017.2 Lyl1/Tal2/Tal1 : 0.61; expression cor: -0.16; max zscore: 31.17 8; gene score cor: 0.32; expression cor: -0.02; max zscore: 5.36 [RC]
2 1 0		HLH: M04097_2.00 Nvec_vc1.1_XM_032364139.2 Sohlh2
2 1 0		: -0.10; expression cor: 0.10; max zscore: 20.47; gene score cor: 0.56; expression cor: -0.07; max zscore: 42.35 [RC] HLH: M08734_2.00 Nvec_vc1.1_XM_032364348.2
2 1 0	gene score cor	H.HG1.34:like:ASCL1/ASCL2/ASCL r: 0.44; expression cor: 0.01; max zscore: 31.17 lene score cor: 0.37; expression cor: 0.13; max zscore: 6.81 [RC]
2 1-		HLH: M05971_2.00 Nvec_vc1.1_XM_032374977.2 Tcf23/Tcf24 -0.14; expression cor: -0.19; max zscore: 41.26 3; gene score cor: 0.27; expression cor: -0.25; max zscore: 5.36 [RC]
2		HLH: M05974_2.00 Nvec_vc1.1_XM_032375280.2 Pasd1 : 0.05; expression cor: -0.09; max zscore: 16.65; gene score cor: -0.08; expression cor: 0.37; max zscore: 42.35
1- 0		CACCTCE HLH: M05984_2.00 Nvec_vc1.1_XM_032381908.2 HLH.HG1.66:like:HAND1/HAND2 r: 0.52; expression cor: 0.43; max zscore: 13.35
2 1 0		CASATGGC HLH: M08731_2.00 Nvec_vc1.1_XM_032382158.2 Tcf3/Tcf12/Tcf4 r: 0.25: expression cor: 0.43: max zscore: 36.65
2 1 0	ARCH867_ZEB1PL0011.1_hlh	r: 0.25; expression cor: 0.43; max zscore: 36.65 ; gene score cor: 0.25; expression cor: 0.37; max zscore: 42.35 CACCTG HLH: M06000_2.00 Nvec_vc1.1_XM_032384474.2
2 1 0	gene score cor	Nvec_vc1.1_XM_032384474.2
2 1		
2		HLH: M06003_2.00 Nvec_vc1.1_XM_048733302.1 Tfe3/Tfec/Tfeb/Mitf r: 0.02; expression cor: 0.03; max zscore: 23.04 score cor: 0.17; expression cor: -0.01; max zscore: 5.31 [RC]
1- 0		CACGTGA HMGbox_Sox: M00206_2.00 Nvec_vc1.1_XM_001632568.3 HMGbox_Sox.HG1.34:SRY r: 0.38; expression cor: 0.11; max zscore: 18.56 ; gene score cor: 0.44; expression cor: 0.10; max zscore: 11.92 [RC]
2 1 0	ARCH1174_/Sox4/Sox11_Sox13/	; gene score cor: 0.44; expression cor: 0.10; max zscore: 11.92 [RC] HMGbox_Sox: M02487_2.00 Nvec_vc1.1_XM_032364225.2 MGbox_Sox.HG1.64:like:HMG20B
2 1 0	gene score cor ARCH1159_/Lef1/Tcf7l1/Tcf7;	r: -0.05; expression cor: 0.23; max zscore: 9.98; gene score cor: -0.09; expression cor: 0.34; max zscore: 7.72 ATATATATATATATATATATATATATATATATATATA
2 1 0	gene score cor: ARCH924_/Lef1/Tcf7I1/Tcf7;	HMGbox_Sox.HG1.65:HMG20B: -0.02; expression cor: -0.05; max zscore: 9.98 gene score cor: 0.23; expression cor: -0.14; max zscore: 4.56
2 1	gene score cor ARCH1162_/Sox4/Sox11_Sox13/;	HMGbox_Sox: M03983_2.00 Nvec_vc1.1_XM_032386296.2 Gbox_Sox.HG1.18:like:SOX1/SOX2 r: 0.18; expression cor: -0.18; max zscore: 5.21 gene score cor: 0.31; expression cor: -0.35; max zscore: 21.37 [RC]
0 []]	HMG gene score cor:	HMGbox_Sox: M02487_2.00 Nvec_vc1.1_XM_048725617.1 Bbox_Sox.HG1.57:like:HMG20A/HM : -0.01; expression cor: -0.04; max zscore: 9.98 S; gene score cor: -0.09; expression cor: 0.36; max zscore: 3.36
1-0		HMGbox_Sox: M11336_2.00 Nvec_vc1.1_XM_048732776.1 Mael -0.03; expression cor: -0.09; max zscore: 21.07 ox4; gene score cor: -0.04; expression cor: 0.05; max zscore: 6.05
2 1 0	HSF	HSF: M05519_2.00 Nvec_vc1.1_XM_001625118.3 F.HG1.0:like:HSF1/HSF2/HSF4/HS
2 1 0	ARCH729_Hsf1/Hsf2/Hsf4/Hsf3; g	: 0.34; expression cor: -0.11; max zscore: 12.27 lene score cor: 0.04; expression cor: -0.04; max zscore: 16.61 [RC] TAGAACG HSF: M05519_2.00 Nvec_vc1.1_XM_001633014.3
2 1 0		Hsf5 -0.30; expression cor: -0.31; max zscore: 12.27 [RC]
pits 1		Homeodomains: M06530_2.00 Nvec_vc1.1_XM_001623084.3 Six4/Six5 r: 0.18; expression cor: 0.14; max zscore: 14.72 ; gene score cor: 0.66; expression cor: 0.09; max zscore: 18.25 [RC]
0 []]		Homeodomains: M02082_2.00 Nvec_vc1.1_XM_001623946.3 Hesx1 : 0.06; expression cor: -0.03; max zscore: 18.04; gene score cor: 0.25; expression cor: -0.09; max zscore: 5.49
0		Homeodomains: M02082_2.00 Nvec_vc1.1_XM_001634768.3 eodomains.HG2.11:DUX4/DUXA/DUXB r: 0.54; expression cor: 0.34; max zscore: 18.04
1-		Homoodomoino, M02002, 200
o		Homeodomains: M02082_2.00 Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a2/Rhox3a/Rhox3c/Rhox3e/Rhox7a/Rhox7b
0 2 1 0 0	gene score cor:	Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a2/Rhox3a/Rhox3c/Rhox3e/Rhox7a/Rhox7b: -0.06; expression cor: 0.02; max zscore: 18.04; gene score cor: 0.17; expression cor: -0.17; max zscore: 10.94 Homeodomains: M03808_2.00
1-	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor	Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a/Rhox3a/Rhox3c/Rhox3e/Rhox7a/Rhox7b: -0.06; expression cor: 0.02; max zscore: 18.04; gene score cor: 0.17; expression cor: -0.17; max zscore: 10.94 Homeodomains: M03808_2.00 Nvec_vc1.1_XM_001647471.3
1- 0	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Home	Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a2/Rhox3a/Rhox3c/Rhox3e/Rhox7a/Rhox7b : -0.06; expression cor: 0.02; max zscore: 18.04; gene score cor: 0.17; expression cor: -0.17; max zscore: 10.94 Homeodomains: M03808_2.00 Nvec_vc1.1_XM_001647471.3
1- 0 2 1- 0	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger	Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a2/Rhox3a/Rhox3c/Rhox3e/Rhox7a/Rhox7b: -0.06; expression cor: 0.02; max zscore: 18.04; gene score cor: 0.17; expression cor: -0.17; max zscore: 10.94 Homeodomains: M03808_2.00 Nvec_vc1.1_XM_001647471.3 Otx1/Otx2 r: 0.40; expression cor: 0.10; max zscore: 25.40 gene score cor: 0.32; expression cor: 0.40; max zscore: 12.43 [RC] GGATTA Homeodomains: M03126_2.00 Nvec_vc1.1_XM_032362396.2 eodomains.HG1.69:HOXA3/HOXB3/0.13; expression cor: -0.17; max zscore: 26.92 ne score cor: 0.46; expression cor: -0.29; max zscore: 24.10 [RC]
1- 0 2 1- 0	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx; Home gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx;	Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a2/Rhox3a/Rhox3c/Rhox7a/Rhox7b: -0.06; expression cor: 0.02; max zscore: 18.04; gene score cor: 0.17; expression cor: -0.17; max zscore: 10.94 Homeodomains: M03808_2.00 Nvec_vc1.1_XM_001647471.3 Otx1/Otx2 r: 0.40; expression cor: 0.10; max zscore: 25.40 gene score cor: 0.32; expression cor: 0.40; max zscore: 12.43 [RC] Homeodomains: M03126_2.00 Nvec_vc1.1_XM_032362396.2 eedomains.HG1.69:HOXA3/HOXB3/0.13; expression cor: -0.17; max zscore: 26.92 ne score cor: 0.46; expression cor: -0.29; max zscore: 24.10 [RC] Homeodomains: M06272_2.00 Nvec_vc1.1_XM_032362713.2 is11/Is12 r: 0.19; expression cor: 0.36; max zscore: 23.92 gene score cor: 0.40; expression cor: 0.29; max zscore: 10.84 [RC] Homeodomains: M06267_2.00 Nvec_vc1.1_XM_03236673.2 neodomains.HG1.68:AC004080.3/A r: 0.09; expression cor: 0.42; max zscore: 18.59
1- 0 2 1- 0 2 1- 0	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx; Home gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx;	Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a2/Rhox3a/Rhox3c/Rhox7a/Rhox7b: -0.06; expression cor: 0.02; max zscore: 18.04; gene score cor: 0.17; expression cor: -0.17; max zscore: 10.94 Homeodomains: M03808_2.00 Nvec_vc1.1_XM_001647471.3
1- 0 2 1- 0 2 1- 0	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx; Home gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx;	Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a/Rhox3a/Rhox3c/Rhox7a/Rhox7b:-0.06; expression cor: 0.02; max zscore: 18.04; gene score cor: 0.17; expression cor: -0.17; max zscore: 10.94 Homeodomains: M03808_2.00 Nvec_vc1.1_XM_001647471.3 Otx1/Otx2 r: 0.40; expression cor: 0.10; max zscore: 25.40 gene score cor: 0.32; expression cor: 0.40; max zscore: 12.43 [RC] GGATTA Homeodomains: M03126_2.00 Nvec_vc1.1_XM_032362396.2 eodomains: HG1.69:HOXA3/HOXB3/0.13; expression cor: -0.17; max zscore: 26.92 ne score cor: 0.46; expression cor: -0.29; max zscore: 24.10 [RC] MATTA Homeodomains: M06272_2.00 Nvec_vc1.1_XM_032362713.2 Is1/Isl2 r: 0.19; expression cor: 0.36; max zscore: 23.92 gene score cor: 0.40; expression cor: 0.29; max zscore: 10.84 [RC] Homeodomains: M06267_2.00 Nvec_vc1.1_XM_032366733.2 neodomains.HG1.68:AC004080.3/A r: 0.09; expression cor: 0.42; max zscore: 18.59 gene score cor: 0.47; expression cor: 0.37; max zscore: 10.88 [RC] Homeodomains: M06354_2.00 Nvec_vc1.1_XM_032366836.2 Independent in the control of
1- 0 2 1- 0 2 1- 0 2 1- 0	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx; Home gene score cor: ARCH1028_Alx3/Alx4/Alx1_Bar; H gene score cor: ARCH1552_Pbx4/Pbx2/Pbx1; ge	Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a/Rhox3a/Rhox3c/Rhox3e/Rhox7a/Rhox7b:0.06; expression cor: 0.02; max zscore: 18.04; gene score cor: 0.17; expression cor: -0.17; max zscore: 10.94 Homeodomains: M03808_2.00 Nvec_vc1.1_XM_001647471.3 Otx1/Otx2 r: 0.40; expression cor: 0.10; max zscore: 25.40 gene score cor: 0.32; expression cor: 0.40; max zscore: 12.43 [RC] GGATTA Homeodomains: M03126_2.00 Nvec_vc1.1_XM_032362396.2 eodomains: M06164747.3 -0.13, expression cor: -0.17; max zscore: 26.92 eodomains: M06272_2.00 Nvec_vc1.1_XM_032362713.2 is11/is12 r: 0.19; expression cor: 0.36; max zscore: 23.92 gene score cor: 0.40; expression cor: 0.29; max zscore: 10.84 [RC] FOR STATA Homeodomains: M06272_2.00 Nvec_vc1.1_XM_032366733.2 eodomains: M06267_2.00 Nvec_vc1.1_XM_032366733.2 reodomains: M06267_2.00 Nvec_vc1.1_XM_032366733.2 reodomains: M06267_2.00 Nvec_vc1.1_XM_032366730.2 eodomains: M06267_2.00 Nvec_vc1.1_XM_032366362 domeodomains: M0634_2.00 Nvec_vc1.1_XM_032366362 domeodomains: M06354_2.00 Nvec_vc1.1_XM_032375635.2 domeodomains: M03808_2.00 Nvec_vc1.1_XM_032375635.2 Otx1/Otx2 Otx1/Otx2 otx1/Otx2 r: 0.52; expression cor: 0.15; max zscore: 16.48 ene score cor: -0.14; expression cor: 0.44; max zscore: 25.39 [RC] GGATTA Homeodomains: M03808_2.00 Nvec_vc1.1_XM_032375635.2 Otx1/Otx2 otx1/Otx2 r: 0.52; expression cor: 0.10; max zscore: 25.40 ene score cor: 0.53; expression cor: 0.04; max zscore: 9.96 [RC]
1-0 21-0 21-0 21-0 21-0 21-0	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx; Home gene score cor: ARCH1028_Alx3/Alx4/Alx1_Bar; ### gene score cor: ARCH1552_Pbx4/Pbx2/Pbx1; ger Gene score cor: ARCH1539_Nkx2-2/Nkx2-9; ger #### gene score cor: ARCH1539_Nkx2-2/Nkx2-9; ger ###################################	Nvec_vc1.1_XM_001636859.3 Rhox3g/Rhox3a2/Rhox3a2/Rhox3a/Rhox3c/Rhox7a/Rhox7b:-0.06; expression cor: 0.02; max zscore: 18.04; gene score cor: 0.17; expression cor: -0.17; max zscore: 10.94 Homeodomains: M03808_2.00 Nvec_vc1.1_XM_001647471.3 Otx1/Otx2 r: 0.40; expression cor: 0.10; max zscore: 25.40 gene score cor: 0.32; expression cor: 0.40; max zscore: 12.43 [RC] GGATTA Homeodomains: M03126_2.00 Nvec_vc1.1_XM_032362396.2 eodomains.HG1.69:HOXA3/HOXB3/0.13; expression cor: -0.17; max zscore: 26.92 ne score cor: 0.46; expression cor: -0.29; max zscore: 24.10 [RC] Homeodomains: M06272_2.00 Nvec_vc1.1_XM_032366731.2 Isl1/Isl2 r: 0.19; expression cor: 0.36; max zscore: 23.92 gene score cor: 0.40; expression cor: 0.29; max zscore: 10.84 [RC] Homeodomains: M06267_2.00 Nvec_vc1.1_XM_032366733.2 pendomains: HG1.68:AC0004080.3/A r: 0.09; expression cor: 0.42; max zscore: 18.59 gene score cor: 0.47; expression cor: 0.37; max zscore: 10.88 [RC] Homeodomains: M06354_2.00 Nvec_vc1.1_XM_032366836.2 lomeodomains.HG1.54:Ilke:NOTO -0.10; expression cor: -0.15; max zscore: 16.48 ne score cor: -0.14; expression cor: 0.44; max zscore: 25.39 [RC] TATA Homeodomains: M03808_2.00 Nvec_vc1.1_XM_032375635.2 Otx1/Otx2 r: 0.52; expression cor: 0.10; max zscore: 25.40
1-0 2 1-0 2	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Hom gene score cor: ARCH1362_Six1/_Six4/Six5; ger Gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx; Hom gene score cor: ARCH1028_Alx3/Alx4/Alx1_Bar; gene score cor: ARCH1552_Pbx4/Pbx2/Pbx1; ge Gene score cor: ARCH1539_Nkx2-2/Nkx2-9; ger Gene score cor: ARCH1362_Six1/_Six4/Six5; ger	Nvec. vol.1. XM_001636859.3 Rhox3g/Rhox3a2/Rho
1-0 21-0 21-0 21-0 21-0 21-0	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH335_Alx3/Alx4/Alx1_Barx; Home gene score cor: ARCH1028_Alx3/Alx4/Alx1_Bar; He gene score cor: ARCH1552_Pbx4/Pbx2/Pbx1; ger Gene score cor: ARCH1539_Nkx2-2/Nkx2-9; gr ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Gene score cor: ARCH1362_Six1/_Six4/Six5; ger Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger	Nvec. vol.1. XM_001636859.3 RNo3g/Rhox3a2/Rhox
1-0 2 1-0 2	gene score cor: ARCH233_Alx3/Alx4/Alx1_Barx gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH335_Alx3/Alx4/Alx1_Barx; Home gene score cor: ARCH1028_Alx3/Alx4/Alx1_Bar; gene score cor: ARCH1552_Pbx4/Pbx2/Pbx1; ger ARCH1539_Nkx2-2/Nkx2-9; gr He gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Home gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger	Nec. vc1.1_XM_001636859.3 Rhox3g/Rh
1-0 2 1-0 2	gene score cor ARCH181_Barx2/Barx1_Dlx4/Dl; Hom gene score cor: ARCH1362_Six1/_Six4/Six5; ger Gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx; Hom gene score cor ARCH335_Alx3/Alx4/Alx1_Barx; Hom gene score cor ARCH1028_Alx3/Alx4/Alx1_Bar; gene score cor: ARCH1552_Pbx4/Pbx2/Pbx1; ge Gene score cor ARCH1539_Nkx2-2/Nkx2-9; gr Gene score cor ARCH1362_Six1/_Six4/Six5; ger Hom gene score cor ARCH1362_Six1/_Six4/Six5; ger Hor gene score cor ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Hor gene score cor ARCH1362_Six1/_Six4/Six5; ger	Nee vol.1, XM, 001636859.3, 20 RNox3g/Rhox3g/Rhox3g/Rhox3g/Rhox3g/Rhox7a/Rhox7b 0.06; expression cor: 0.02; max zscore: 18.04
1-0 2 1-0 2 1-0 2 1-0 2 1-0 2 1-0 2 1-0	gene score cor ARCH181_Barx2/Barx1_Dlx4/Dl; Hom gene score cor: ARCH1362_Six1/_Six4/Six5; ger RCH1362_Alx3/Alx4/Alx1_Barx; Hom gene score cor ARCH335_Alx3/Alx4/Alx1_Barx; Hom gene score cor ARCH1028_Alx3/Alx4/Alx1_Bar; gene score cor: ARCH1552_Pbx4/Pbx2/Pbx1; ge RCH1539_Nkx2-2/Nkx2-9; ge Gene score cor ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Hom gene score cor ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Cdc5lrt1/Cdc5lrt10/Cdc Cdc5lrt1/Cdc5lrt10/Cdc	Nece vc1.1 xM 001636859.3 Rhox3g/Rhox
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$ \begin{bmatrix} 1 \\ 0 \end{bmatrix} \begin{bmatrix} 2 \\ 1 \\ \end{bmatrix} \begin{bmatrix} 2 \\ 1 \\ \end{bmatrix} \begin{bmatrix} 2 \\ \end{bmatrix} \begin{bmatrix} $	gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Homm gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH335_Alx3/Alx4/Alx1_Barx; Homm gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx; Homm gene score cor: ARCH1028_Alx3/Alx4/Alx1_Bar; #### Gene score cor: ARCH1552_Pbx4/Pbx2/Pbx1; ger Gene score cor: ARCH1362_Six1/_Six4/Six5; ger #### Gene score cor: ARCH1379_MYB17; gene scor	Nee_vc1.1_XM_00163689.3 Nee_vc1.1_XM_00163689.3 Nehoxig/Rhoxig/Rhoxig/Rhoxig/Rhox/gr/Rhox/7e/Rhox/7e 0.06_expression cor: 0.02: max zscore: 18.04 Homeodomains: M03808_2.00 Nvec_vc1.1_XM_001647471.3 Oxiv(Vict2) 0.04; expression cor: 0.10; max zscore: 12.43 [RC] GGATTA Homeodomains: M03126_2.00 Nvec_vc1.1_XM_03286363.2 0.13; expression cor: 0.10; max zscore: 25.40 gene score cor: 0.32; expression cor: 0.40; max zscore: 12.43 [RC] GGATTA Homeodomains: M03126_2.00 Nvec_vc1.1_XM_032862366.2 0.13; expression cor: -0.17; max zscore: 26.92 0.8 score cor: 0.46; expression cor: -0.29; max zscore: 24.10 [RC]
$ \begin{bmatrix} 1 \\ 0 \end{bmatrix} \begin{bmatrix} 2 \\ 1 \end{bmatrix} \begin{bmatrix} 1 \\ 0 \end{bmatrix} \begin{bmatrix} 2 \\ 1 \end{bmatrix} \begin{bmatrix} 1 \\ 0 \end{bmatrix} \begin{bmatrix} 2 \\ 1 \end{bmatrix} \begin{bmatrix} 1 \\ 0 \end{bmatrix} \begin{bmatrix} 1 \\ \end{bmatrix} \begin{bmatrix} $	gene score cor ARCH181_Barx2/Barx1_Dlx4/Dl; Hom gene score cor: ARCH181_Barx2/Barx1_Dlx4/Dl; Hom gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH335_Alx3/Alx4/Alx1_Barx; Hom gene score cor: ARCH335_Alx3/Alx4/Alx1_Barx; Hom gene score cor: ARCH1028_Alx3/Alx4/Alx1_Barx; Gene score cor: ARCH1552_Pbx4/Pbx2/Pbx1; ge Gene score cor: ARCH1362_Six1/_Six4/Six5; ger Hom gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Cdc5lrt1/Cdc5lrt10/Cdc gene score cor: ARCH1362_Meis3/Meis2/Meis1; Gene score cor: ARCH1362_Meis3/Meis2/Meis1; Gene score cor: ARCH1362_Six1/_Six4/Six5; ger Gene score cor: ARCH1362_Six1/_Six4/Six5; ger ARCH1362_Six1/_Six4/Six5; ger Gene score cor: AR	Neez_er1_XM_00163869.3 Rhoxdg/Rhoxdg/Rhoxdg/Rhoxde/Rhox/a/Rhox/b -0.00: oppresion cor: 0.02: max zscore: 18.04 -1.00: oppresion cor: 0.02: max zscore: 18.04 -1.00: oppresion cor: 0.02: max zscore: 18.04 -1.00: oppresion cor: 0.07: oppresion cor: 0.07: max zscore: 25.40 -1.00: oppresion cor: 0.01: max zscore: 26.92 -1.01: oppresion cor: 0.01: max zscore: 26.92 -1.01: oppresion cor: 0.01: max zscore: 26.92 -1.01: oppresion cor: 0.01: max zscore: 28.92 -1.01: oppresion cor: 0.01: max zscore: 28.92 -1.01: oppresion cor: 0.01: max zscore: 28.92 -1.01: oppresion cor: 0.02: max zscore: 28.92 -1.01: oppresion cor: 0.02: max zscore: 18.59 -1.01: oppresion cor: 0.01: oppresion cor: 0.03: max zscore: 25.39 [RC] -1.01: oppresion cor: 0.01: oppresion cor: 0.04: max zscore: 25.39 [RC] -1.01: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.01: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.01: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.02: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.02: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.02: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.03: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.03: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.03: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.03: oppresion cor: 0.01: oppresion cor: 0.01: max zscore: 25.40 -1.03: oppresion cor: 0.01: oppresion cor: 0.
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	gene score con ARCH1362_Six1/_Six4/Six5; gene ARCH1363_ARCH1	Nome, and 1, 244 Col (16/88) and 2, 200 Col (
	gene score con ARCH233_Alx3/Alx4/Alx1_Barx. ARCH181_Barx2/Barx1_Dlx4/Dl; gene score con ARCH1362_Six1/_Six4/Six6; gen ARCH335_Alx3/Alx4/Alx1_Barx; gene score con ARCH1028_Alx3/Alx4/Alx1_Barx; gene score con ARCH1552_Pbx4/Pbx2/Pbx1; gen ARCH1539_Nlx2-2/Pbx4/Six6; gen ARCH1362_Six1/_Six4/Six6; gen ARCH1364_Meis3/Meis2/Meis1; Cdc5/rt1/Cdc5/rt10/Cdc gene score con ARCH1379_MYB17: gene ARCH1379_MYB17: gene ARCH1379_MYB17: gene ARCH1379_MYB17: gene ARCH1389_Rtx1/Rtx3/Rtx2_Rtx6; ARCH1359_Rtx1/Rtx3/Rtx2_Rtx6;	More with 13th 6016/36/2013 gard source out 5.17, expression cont -0.17, max zacone 10.94 gard source out 5.17, expression cont -0.17, max zacone 10.94 Horraccomanic MASS 202 Note _vit 13th 601644 21.3 gard source out 5.25 expression cont -0.17, max zacone 10.40 [RC] gard source out 5.25 expression cont -0.17, max zacone 10.40 [RC] gard source out 5.25 expression cont -0.47 max zacone 12.40 [RC] gard source out 5.25 expression cont -0.47 max zacone 12.40 [RC] gard source out 5.25 expression cont -0.47 max zacone 2.40 [RC] gard source out 5.25 expression cont -0.47 max zacone 2.40 [RC] gard source out 5.40 captesion cont -0.49 max zacone 2.40 [RC] gard source out 5.40 captesion cont -0.49 max zacone 2.40 [RC] gard source out 5.40 captesion cont -0.49 max zacone 10.44 [RC] gard source out 5.40 captesion cont -0.49 max zacone 10.49 [RC] gard source out 5.40 captesion cont -0.49 max zacone 10.49 [RC] gard source out 5.40 captesion cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 2.40 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.49 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.40 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.40 [RC] gard source out 5.47 expression cont -0.47 max zacone 10.40 [RC] gard source out
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	gene score con ARCH233_Alx3/Alx4/Alx1_Barx gene score con ARCH362_Six17_Six4/Six5, gen ARCH336_Alx3/Alx4/Alx1_Barx; gene score con ARCH336_Alx3/Alx4/Alx1_Barx; gene score con ARCH1028_Alx3/Alx4/Alx1_Barx; gene score con ARCH1539_Nlo/2_2/Nlo/2_9; gene ARCH1540_Six17_Six4/Six5; gene ARCH1362_Six17_Six4/Six5; gene ARCH1364_Meis3/Meis2/Meis1; Cdc5irt1/Cdc5irt10/Cdc gene score con ARCH1379_MYB17; gene s ARCH1379_MYB17; gene s ARCH1379_MYB17; gene s ARCH1379_Arx5/Pax8/Pax8/Pax8/Pax8/Pax8/Pax8/Pax8/Pax8	New 241 1, 204 (2016) 258 (2016) 277, with south through the house of the processor of the 102 men and the 2016 (2016) 278, with south through the 2016 (2017) 278, with south through the 2017 (2017) 278, with south through through through the 2017 (2017) 278, with south through
$\begin{bmatrix} 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 & 1 $	ARCH1362_Six1/_Six4/Six5; gen ARCH1362_Six1/_Six4/Six5; gen ARCH1362_Six1/_Six4/Six5; gen ARCH1362_Alx3/Alx4/Alx1_Barx; ARCH1362_Alx3/Alx4/Alx1_Barx; ARCH1362_Alx3/Alx4/Alx1_Barx; ARCH1362_Alx3/Alx4/Alx1_Barx; ARCH1362_Six1/_Six4/Six5; gen ARCH1362_Six1/_Six4/Six5; gen ARCH1362_Six1/_Six4/Six5; gen ARCH1362_Six1/_Six4/Six5; gen ARCH1362_Six1/_Six4/Six5; gen ARCH1362_Alx3/Alx4/Alx1_Barx; Cdc5ii+11/Cdc5i+110/Cdc gene score cor ARCH1362_Alx3/Alx4/Alx1_Barx; Cdc5ii+11/Cdc5i+110/Cdc gene score cor ARCH1362_Alx3/Alx4/Alx1_Barx; ARCH1379_Alx171_gene score cor ARCH1379_Alx171_gene score cor ARCH1379_Alx171_gene score cor ARCH1379_Alx171_Figene score cor ARCH1389_Rix1_Rix3_Rix2_Rix6; ARCH1389_Rix1_Rix3_Rix1_Rix5_Rix6; ARCH1389_Rix1_Rix3_Rix1_Rix5_Rix6_Rix6_Rix6_Rix6_Rix6_Rix6_Rix6_Rix6	ANSALOT LANG ADDITIONAL STATES AND ADDITIONA
$\begin{bmatrix} 1 \\ 0 \\ 2 \\ 1 \\ 0 \end{bmatrix}$ $\begin{bmatrix} 2 \\ 1 \\ 0 \\ 2 \\ 1 \\ 0 \end{bmatrix}$ $\begin{bmatrix} 2 \\ 1 \\ 0 \\ 2 \\ 1 \\ 0 \end{bmatrix}$ $\begin{bmatrix} 2 \\ 1 \\ 0 \\ 2 \\ 1 \\ 0 \end{bmatrix}$ $\begin{bmatrix} 2 \\ 1 \\ 0 \\ 0 \end{bmatrix}$ $\begin{bmatrix} 2 \\ 1 \\ 0 \\ 0 \end{bmatrix}$ $\begin{bmatrix} 2 \\ 1 \\ 0 \\ 0 \end{bmatrix}$ $\begin{bmatrix} 2 \\ 1 \\ 0 \end{bmatrix}$	ARCH1342 Alx3/Alx4/Alx1 Barx Gene score con ARCH181_Ban2/Bah1_Dlx4/Dl Hom ARCH1362_Six1/_Six4/Six6; gene ARCH1362_Alx3/Alx4/Alx1_Barx; Gene score con ARCH1028_Alx3/Alx4/Alx1_Barx; Gene score con ARCH1529_Alx3/Alx4/Alx1_Barx; Gene score con ARCH1539_Nav2-2Nbx2-9; gene ARCH1539_Nav2-2Nbx2-9; gene ARCH1362_Six1/_Six4/Six6; gene ARCH1362_Six1/_Six4/Six6; gene ARCH1362_Six1/_Six4/Six6; gene ARCH1364_Meis3/Me	News, 2011 J. 200, 2019 September 101, 14 (1) prints process 101, 14 (1) pr
$\begin{bmatrix} 2 \\ 1 \\ 0 \end{bmatrix}$	ARCH1342 Alx3/Alx4/Alx1 Barz. Gene score con ARCH181 Barz2/Barx1_Black/DL.; gene score con ARCH1362_Stx1_Stx4/Stx6; gen Horn ARCH1362_Alx3/Alx4/Alx1 Barz.; Gene score con ARCH1028_Alx3/Alx4/Alx1 Barz.; Gene score con ARCH1028_Alx3/Alx4/Alx1 Barz.; Gene score con ARCH1532_Nax2-ZnNax2-9; gene ARCH1532_Nax2-ZnNax2-9; gene ARCH1362_Stx1_Stx4/Stx5; gen ARCH1362_Stx1_Stx4/Stx5; gen ARCH1362_Stx1_Stx4/Stx5; gene ARCH1362_Stx1_Stx4/Stx5; gene ARCH1362_Stx1_Stx4/Stx5; gene ARCH1362_Stx1_Stx4/Stx5; gene ARCH1362_Stx1_Stx4/Stx5; gene ARCH1362_Stx1_Stx4/Stx6; gene ARCH1362_Ttx1_Stx4/Stx6; gene ARCH1362_Ttx1_Stx4/Stx6, gene ARCH1362_Ttx1_Stx4/Stx6, gene ARCH1362_Ttx1_Stx4/Stx6, gene ARCH1362_Ttx1_Stx4/Stx6, gene ARCH1362_Ttx1_Stx4/Stx4/Stx4/Stx4/Stx4/Stx4/Stx4/Stx4/	The common of th
	ARCH1342_Masanta4/Mypop_Terfap ARCH1342_Masanta4/Mypop_Terfap ARCH1362_Masanta4/Mypop_Terfap ARCH1362_Masanta4/Mypop_Terfap ARCH1362_Masanta4/Mypop_Terfap ARCH1362_Masanta4/Mypop_Terfap ARCH1363_Masanta4/Mypop_Terfap ARCH1364_Masanta4/Mypop_Terfap ARCH1379_MYPO17; gene score con ARCH1389_Refut/Refus	March 1,044, 5015, 508569
	ARCH1342, Means score con ARCH1342, Means score con ARCH1342, Abd/Akk4Ak1, Barx Hom ARCH1362, Six1, Six4/Six5; gen ARCH1362, Six1, Six4/Six5; gen ARCH1362, Ak3/Ak4/Ak1, Barx ARCH1362, Ak3/Ak4/Ak1, Barx ARCH1362, Ak3/Ak4/Ak1, Barx ARCH1362, Ak3/Ak4/Ak1, Barx ARCH1362, Six1, Six4/Six5; gen ARCH1363, Pax5/Rax8, Rax2; gen ARCH1363, Pax5/Rax8, Rax2; gen ARCH1364, Meantox4/Rypop, Tortzip ARCH1369, Rix1, Rix3, Rix2, Rix5; gen ARCH1369, Rix1, Rix3, Rix1, Rix3, Rix2, Rix5; gen ARCH1369, Rix1, Rix3, Rix1, R	New york and Challegood and Challegood Collego
	ARCH1812_Ranz Gene score con ARCH1812_Alx3/Alx4/Alx1_Barx Horn ARCH182_Alx3/Alx4/Alx1_Barx Gene score con ARCH1835_Alx3/Alx4/Alx1_Barx ARCH1828_Alx3/Alx4/Alx1_Barx Gene score con ARCH1838_Alx3/Alx4/Alx1_Barx ARCH1828_Alx3/Alx4/Alx1_Barx Gene score con ARCH1838_Nsc2-Alx3/Alx4/Alx1_Barx ARCH1838_Nsc2-Alx3/Alx4/Alx1_Barx Gene score con ARCH1838_Nsc2-Alx3/Alx4/Alx1_Barx ARCH1838_Nsc2-Alx3/Alx4/Alx1_Barx Gene score con ARCH1838_Nsc2-Alx3/Alx4/Alx1_Barx ARCH1838_Nsc2-Alx3/Alx4/Alx1_Barx ARCH1848_Alx3/Alx4/Alx3/Alx4/Alx1_Barx ARCH1848_Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx3/Alx4/Alx4/Alx3/Alx4/Alx4/Alx3/Alx4/Alx4/Alx4/Alx4/Alx4/Alx4/Alx4/Alx4	Name and 1 and Conference
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	gene score cor ARCH1315_AKCANAMAN_BAT_BATA. ARCH1818_BATA_BATA. ARCH1818_BATA_BATA. ARCH182_SIAT_SIAASIAG_SIG_ ARCH1315_AKCANAMAN_BATA. ARCH1928_ANCANAMAN_BATA. ARCH1928_ANCANAMAN_BATA. ARCH1928_ANCANAMAN_BATA. ARCH1938_NINCZ_ZINACZ_B. Gene score cor ARCH1938_NINCZ_ZINACZ_B. ARCH1938_ANCANAMAN_BATA. ARCH1938_ANCANAMAN. ARCH1938_ANCANAMAN.	Note 1, 11 (1. 10. 00) CONTROL
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