JgiID	symbol	Entrez.GeneID	UnigeneID_XT	gi	accession	gene.name	gene.function	gene.synonyms	XB.GenpageID	XB.GeneID	UnigeneID_XL	MGI_Symbol	EntrezGeneID.MM	Seo2007_Ngnr1_NeuroD	Seo2007_Ngnr1	Seo2007_NeuroD	Masui2010	Thompson2012	baseMean	baseMean_Cc_6h_Dex	baseMean_Ptf1a-GR_6h_Dex	FC_Ptf1a-GR_6h_Dex_vs_Cc_6h_Dex	log2FC_Ptf1a-GR_6h_Dex_vs_Cc_6h_Dex	FDR
Xetrov72000709	prdm13	100490238		301605112	XM_002932160	NA	NA	NA	XB-GENEPAGE-960154	XB-GENE-960155		Prdm13	230025	-	-	-	-	-	1010,86	1,33	2020,38	865,70	9,76	0,00%
Xl.41853	-	-	-	-	-		BJ070460 NIBB Mochii normalized Xenopus tailbud library Xenopus laevis cDNA clone XL056k03 5', mRNA sequence [BJ070460]	-	-	-	Xl.41853	-	-	-	-	-	-	-	47,86	0,81	94,92	52,95	5,73	0,00%
Xetrov72027391	hes3.3	100101782	Str.51064	197246528,197246257,301608655,134024390	BC169157,BC168837,XM_002933848,BC135685	hairy and enhancer of split 3, gene 3	helix-loop-helix transcription factor/transcriptional repressor	esr2	XB-GENEPAGE-480449	XB-GENE-480450	Xl.12067,Xl.34898	Hes3	15207	-	Xl.34898	-	-	Hes3	125,09	3,72	246,45	52,46	5,71	0,00%
Xetrov72030513	pou3f1	549258	Str.25752	67867505,77626825,67972640	BC098088,CR760792,NM_001016504	POU class 3 homeobox 1	transcription factor containing POU and HOX domains	pou50|oct-6|oct6|scip|nrl-22	XB-GENEPAGE-853353	XB-GENE-853354	Xl.21449,Xl.1274	Pou3f1	18991	-	-	Xl.21449	-	-	912,30	33,35	1791,24	52,18	5,71	0,00%
Xetrov72035391		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	-	-	-	-	-	243,10	9,48	476,72	45,60	5,51	0,00%
Xetrov72017800	mxra7	594877	Str.29637	301612981,60551293,301612979	XM_002935926,BC091054,XM_002935925	NA	NA	NA	XB-GENEPAGE-953445	XB-GENE-953446	Xl.48885,Xl.15502	Mxra7	67622	Xl.15502	-	-	-	-	50,43	1,33	99,52	43,05	5,43	0,00%
Xetrov72023347	aldh1b1	100498540		301621943	XM_002940263	NA	NA	NA	XB-GENEPAGE-1013739	XB-GENE-1013740	NA	Aldh1b1	72535	-	-	-	Aldh1b1	-	35,74	0,81	70,67	39,57	5,31	0,00%
Xetrov72032576	zc3h12c	100490495		301617158	XM_002937961	NA	NA	NA	XB-GENEPAGE-6045327	XB-GENE-6045328	Xl.69989	Zc3h12c	244871	-	-	-	-	-	490,67	25,26	956,08	36,45	5,19	0,00%
Xetrov72027607	hes8	100271766	Str.15848	156914851,301608657	BC152703,XM_002933849	hairy and enhancer of split 8	transcriptional repressor		XB-GENEPAGE-5952809	XB-GENE-5952810	Xl.53198	NA	NA	-	-	-	-	-	26,31	0,52	52,09	34,85	5,12	0,03%
Xetrov72035516	id4	448114	Str.5460	52345581,49257765,77626910	NM_001004839,BC074645,CR760882	inhibitor of DNA binding 4, dominant negative helix-loop-helix protein	transcriptional regulator	idb4|bhlhb27	XB-GENEPAGE-876359	XB-GENE-876360	Xl.8624,Xl.85614	Id4	15904	-	-	-	-	-	294,46	15,81	573,11	34,15	5,09	0,00%
Xetrov72035284		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	-	-	-	-	-	1375,96	77,43	2674,50	34,11	5,09	0,00%
Xl.76807	-	-	-	-	-	NA	NA	-	-	-	Xl.76807	-	-	-	-	-	-	-	1513,91	87,11	2940,70	33,39	5,06	0,00%
Xetrov72034908	snai2	395065	Str.2	49522371,45361692,14133800,77623613	BC075376,NM_204093,AF368039,CR926188	snail homolog 2	zinc finger transcriptional repressor	xSlu|xslug|slug|Snail2|XSnail2	XB-GENEPAGE-487370	XB-GENE-487371	Xl.11972,Xl.3818	Snai2	20583	-	-	-	-	-	174,18	9,63	338,74	31,95	5,00	0,00%
Xetrov72031563	fam101b	493421	Str.18500	56118888,51704084	NM_001008058,BC080945	NA	NA	NA	XB-GENEPAGE-962745	XB-GENE-962746	Xl.56544,Xl.75079	Fam101b	76566	-	-	-	-	-	76,36	3,72	148,99	31,80	4,99	0,00%
Xetrov72028287		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	-	-	-	-	-	2398,74	156,84	4640,64	29,41	4,88	0,00%
Xetrov72002407		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	-	-	-	-	-	1521,54	101,67	2941,42	28,66	4,84	0,00%
Xetrov72016721	myt1	100490628	Str.57565	301626514	XM_002942382	myelin transcription factor 1	transcription factor containing C2HC type zinc finger	X-MyT1|xmyt1	XB-GENEPAGE-1009974	XB-GENE-1009975	Xl.57163,Xl.31433,Xl.85112	Myt1	17932	Xl.57163	-	-	-	-	3048,87	207,44	5890,31	28,26	4,82	0,00%
Xetrov72027234	hes5.1	733463	Str.11260	77621232,138519880,113205535,49523069	CR760038,BC135757,NM_001044415,BC075552	hairy and enhancer of split 5, gene 1	helix-loop-helix transcription factor/transcriptional repressor	esr-1|esr1	XB-GENEPAGE-481134	XB-GENE-481135	Xl.48575,Xl.8440	Hes5	15208	-	-	Xl.8440	-	-	618,01	41,72	1194,30	27,98	4,81	0,00%
Xl.55672	-	-	-	-	-	NA	NA	-	-	-	Xl.55672	-	-	-	-	-	-	-	40,38	1,86	78,91	27,96	4,81	0,01%
Xetrov72027170	fgf16	100492309		301604465	XM_002931813	fibroblast growth factor 16	heparin-binding growth factor		XB-GENEPAGE-483588	XB-GENE-483589	NA	Fgf16	80903	-	-	-	-	-	27,53	1,05	54,02	26,88	4,75	0,11%
Xetrov72020979	post	100494594		301631582,301605880	XM_002944831,XM_002932507	posterior protein	zinc finger, CCHC type	Xpo	XB-GENEPAGE-5867591	XB-GENE-5867592	Xl.16656,Xl.85047	NA	NA	-	-	-	-	-	4622,84	335,15	8910,53	26,51	4,73	0,00%
Xetrov72005788	hoxd1	549432	Str.10073	77625937,77682137,213627235,213627233,169642307	CR760220,NM_001016678,BC170965,BC170961,BC160395	homeobox D1	homeodomain transcription factor	Xhox.lab|xhox.lab1|mgc131028|xlab|xhoxd1	XB-GENEPAGE-482730	XB-GENE-482731	Xl.53491,Xl.11612,Xl.76294	Hoxd1	15429	-	-	-	-	-	70,28	4,82	135,74	23,51	4,56	0,00%
Xetrov72008860		NA	NA	NA	NA	NA	NA	NA	NA	NA	Xl.83186	NA	NA	-	-	-	-	-	3323,55	278,66	6368,45	22,78	4,51	0,00%
Xetrov72012722	thbs1	100170153	Str.61138	189571714,301615579	BC167555,XM_002937199	thrombospondin 1	adhesive glycoprotein	tsp|thbs|tsp1|tsp-1|thbs-1|thrombospondin 1	XB-GENEPAGE-486188	XB-GENE-486189	Xl.23965,Xl.21621	Thbs1	21825	-	-	-	-	-	237,15	19,27	455,03	22,50	4,49	0,00%
Xetrov72016283	elavl3	594912	Str.46013	51261570,134026263,77620853,77682340	BC080177,BC136134,CR761704,NM_001030326	ELAV (embryonic lethal, abnormal vision)-like 3 (Hu antigen C)	RNA binding	elrC	XB-GENEPAGE-490863	XB-GENE-490864	Xl.1035,Xl.35785	Elavl3	15571	-	-	Xl.1035	-	-	331,34	29,74	632,94	20,63	4,37	0,00%
Xl.14633	-	-	-	-	-		BJ056854 NIBB Mochii normalized Xenopus neurula library Xenopus laevis cDNA clone XL033d16 3', mRNA sequence [BJ056854]	-	-	-	Xl.14633	-	-	-	-	-	-	-	81,43	6,91	155,95	19,84	4,31	0,00%
Xetrov72023106		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	-	-	-	-	-	108,87	9,82	207,93	19,32	4,27	0,00%
Xetrov72027609	hes5.2	733755	Str.24311	134254268,77621406,113205883	BC135288,CR761792,NM_001044509	hairy and enhancer of split 5, gene 2	helix-loop-helix transcription factor/transcriptional repressor	bHLHb38|esr7|esr-7	XB-GENEPAGE-876634	XB-GENE-876635	Xl.586,Xl.68431	Hes5	15208	-	-	Xl.586	-	-	119,13	10,97	227,30	19,08	4,25	0,00%