

Chap. 2 Supplemental Table 1. Significant gene ontology (GO) categories for genes that showed significant strain x treatment interaction.

Gene Set	Description	Overlap/ Size	FDR	Gene Symbols
GO:0043523	regulation of neuron apoptotic process	106/242	2.82E-06	<i>Coro1a, Sigmar1, Atf4, Trp53, Nr4a2, Lig4, Mapk8, Kcnip3, Mef2c, Nr4a3, Cdc34, Kras, Fzd9, Hif1a, Ascl1, Pin1, Dlx1, Hras, Ndnf, Cited1, Set, Bdnf, Ngf, Nes, Zpr1, Cdk5r1, Hsph1, Rhoa, Bax, Bcl2, Tyro3, Nae1, Egl3, Hspd1, Ube2m, Hdac4, Hyoul, Unc5b, Atf2, Sirt1, Sod2, Grin1, Npm1, Map2k7, Bcl2l1, Aifm1, Bag1, Vstm2l, Syngap1, Egl2, Dnajc5, Gsk3a, Stxbp1, Ppp2r2b, Gabra5, Sod1, Kcnma1, Cpeb4, Chl1, Braf, Cntfr, Xiap, Trim2, Pak3, Pmaip1, Gabrb2, Ntrk2, Mcl1, Grm7, Itsn1, Apoe, Casp3, Pik3ca, Lgmn, Grid2, Smo, Mt3, Ddit3, Grik2, Casp6, Wfs1, Pawr, Tgfb2, Egr1, Lrp1, Nqo1, Ppt1, Bbc3, Cln3, Il18, F2r, Ccl12, Hmox1, Cx3cr1, Gdf5, Draxin, Ctsz, Atp7a, Foxo3, Mdk, Hrk, Mt1, Ccl3, Btg2, Bcl2l11, Jun</i>
GO:1901216	positive regulation of neuron death	54/108	2.17E-05	<i>Aimp2, Atf4, Trp53, Map3k5, Mapk8, Cdc34, Ptpn5, Ascl1, Pin1, Efnb2, Eif2s1, Cdk5r1, Rhoa, Bax, Ube2m, Hdac4, Nos1, Atf2, Map2k7, Aifm1, Ptpnz1, Gsk3a, Egl3, Ppp2r2b, Kcnma1, Capn2, Pak3, Pmaip1, Mcl1, Casp3, Adora1, Clu, Grin2b, Ddit3, Elk1, Grik2, Casp6, Pawr, Tgfb2, Egr1, Nqo1, Tlr4, Bbc3, Il18, Ctsz, Fos, Tlr6, Foxo3, Trp53bp2, Ddit4, Hrk, Ccl3, Bcl2l11, Jun</i>
GO:0008637	apoptotic mitochondrial changes	48/110	3.13E-03	<i>Trp53, Lmna, Fzd9, Timm50, Vps35, Higd1a, Ppif, Cck, Bax, Bcl2, Rhot1, Cd24a, Akt1, Atg3, Pim2, Prelid1, Hspd1, Ggct, Sh3glb1, Atf2, Sod2, Mtch2, Rhot2, Fam162a, Parl, Bcl2l1, Ndufs1, Aifm1, Slc25a5, Pam16, Mllt11, Slc25a4, Gsk3a, Ppp2r2b, Atpif1, Dnm1l, Pmaip1, Zfp13, Them4, Acaa2, Bbc3, Hgf, Nptx1, Cideb, Atp7a, Hrk, Bcl2l11, Jun</i>
GO:0008630	intrinsic apoptotic signaling pathway in response to DNA damage	46/109	8.09E-03	<i>Trp53, Aen, Myc, Dyrk2, Fnip2, Rad9a, Zfp385a, Htra2, Mlh1, Usp28, Tmem109, Bax, Cdkn1a, Bcl2, Msh6, Cdkn2d, Sirt1, Sod2, Mtch2, Mif, Bcl2l1, Bag6, Usp47, Polb, Ercc6, Pmaip1, Crip1, Mcl1, Clu, Akr3, Bcl2a1d, Tmem161a, Chek2, Hipk1, Skil, Bbc3, Cxcl12, Brca2, Bcl2a1b, Trp63, Hmox1, Pml, Xpa, Ddit4, Pik3r1, Bcl2l11</i>
GO:2001242	regulation of intrinsic apoptotic signaling pathway	64/162	8.38E-03	<i>Creb3l1, Mapk7, Trp53, Slc9a3r1, Ivns1abp, Hif1a, Ptpn1, Hdac2, Hells, Rad9a, Zfp385a, Ppif, Ybx3, Ptpn2, Bax, Bcl2, Cdkn2d, Fign1, Taf6, Src, Hyoul, Vdac2, Bag5, Sirt1, Sod2, Mtch2, Mif, Rrn3, Sfpq, Map2k1, Parl, Bcl2l1, Bclaf1, Usp47, Ndufa13, Dnm1l, Dnajal, Sod1, Bcap31, Uri1, Serinc3, Spop, Cep63, Txndc12, Mcl1, Nfe2l2, Clu, Ikbkg, Akr3, Bcl2a1d, Tmem161a, Herpud1, Pttglip, Wfs1, Skil, Bbc3, Cxcl12, Mdm2, Bcl2a1b, Nme5, Bdkrb2, Plekhf1, Eif2ak3, Bcl2l11</i>
GO:0072332	intrinsic apoptotic signaling pathway by p53 class mediator	33/74	1.21E-02	<i>Trp53, Aen, Dyrk2, Zfp385a, Mybbp1a, Usp28, Tmem109, Bax, Cdkn1a, Sirt1, Mif, Rrn3, Bag6, Hint1, Pmaip1, Cep63, Wwox, E2f2, Chek2, Pttglip, Hipk1, Bbc3, Mdm2, Brca2, Perp, Zfp385b, Jmy, Trp63, Pml, Pdk2, Bdkrb2, Trp53bp2, Ddit4</i>
GO:0030902	hindbrain development	64/173	3.62E-02	<i>Trp53, Sstr3, Cdk5r2, Gart, Atxn2, Kat2a, Sez6, Sstr2, Ascl1, Hnrnpd, Sstr1, Wnt7a, Scrib, Mafk, Cend1, Kcnc1, Cdk5r1, Bcl2, Plxna2, Flna, Arcn1, Dll1, Sec24b, Agtpbp1, Lef1, Nanos1, Myo16, Grin1,</i>

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GO:0030900	forebrain development	175/413	1.02E-08	<i>Map2k1, B4galt2, Trnp1, Ncoal, Ogdh, Mtpn, Hspa5, Nr2c2, Gnpat, Ptbp2, Zfp365, Itgb1, Zbtb18, Lrp6, Grid2, Fgf2, Atg7, Smo, Ulk1, Rere, Rpgrip11, Nfib, Prox1, Igflr, Klhl1, Rora, Wls, Egr2, Kndc1, Cbs, Rbfox2, Dixdc1, Atp7a, Smad1, Gata2, Mdk</i> <i>Crhr1, Avpr1a, Lhx6, Srf, Rara, Ephb2, Adcyap1, Arx, Sstr3, Nr4a2, Cdk5r2, Cxcr4, Crtac1, Ets1, Gart, Ttc21b, Prdm8, Ran, H2afx, Tmem108, Nr4a3, Pgap1, Ephb3, Eif2b3, Kras, Kat2a, Npy, Dlx2, Slc32a1, Dlx5, Sstr2, Ccdc85c, Hif1a, Ascl1, Hdac2, Hes5, Mboat7, Dlx1, P2ry12, Tra2b, Sstr1, Htra2, Sox1, Wnt7a, Mettl3, Jakmip1, Ptchd1, Nme1, Uba6, Kcnc1, Zfp335, Cdk5r1, Rhoa, Kif5b, Bax, Trappc9, Tyro3, Wnt4, Neurod6, Wnt7b, Dab2ip, Agtpbp1, Atat1, Gak, Rac3, Rtn4rl1, Src, Creb1, Lef1, Gsx1, Fabp7, Fat4, Rrm1, Plxna3, Sema7a, Top2b, Bcl11b, Grin1, Efna2, Cntn2, Szt2, Lrp8, Fgf13, Pou3f3, Pcm1, Ncoal, Slitrk5, Ogdh, Rtn4r, Fez1, Mettl14, Smarcc2, Gnao1, Cfl1, Rtn4, Kif3a, Srgap2, Ywhae, App, Pten, Arhgap35, Dbi, Cep120, Bcan, Lhx2, Foxp1, Tctn1, Foxg1, Mkks, Bmpr1a, Gdpd5, Ext1, Ntrk2, Tsc1, Nf2, Ndst1, Ppp1r9b, Ryk, Casp3, Tbx3, Zbtb18, Bbs1, Btbd3, Frs2, Lrp2, Lrp6, Tnr, Fgf2, Prkg1, Atg7, Smo, Dclk2, Ttc8, Rpgrip11, Notch1, Tacc2, Adgrg1, Mdga1, Nrg3, Slc1a2, Slc4a10, Nfib, Syne2, Emx2, Csf1r, Prox1, Bmp2, Lamb1, Kcna1, Cxcl12, Mas1, Bmp4, Fezf2, Slc38a2, Bbs2, Cdon, Sun2, Twsg1, Apaf1, Draxin, Myo1d, Gli3, Dixdc1, Atp7a, Fut10, Bbs4, Mfsd2a, Gata2, Ccdc141, Ndel, Mdk, Id4, Rapgef2, Slc2a1, Btg2</i>
GO:0050773	regulation of dendrite development	75/170	8.87E-05	<i>Mapk6, Foxo6, Trpc6, Rtn4ip1, Prex1, Arc, Pacsin1, Arf6, Mef2c, Bcl11a, Sez6, Ss18l1, Hdac2, Carm1, Itpka, Sarm1, Actr3, Baiap2, Asap1, Cdk5r1, Caprin1, Rhoa, Anapc2, Cyth2, Vldlr, Dab2ip, Ngef, Cask, Srcin1, Dbn1, Sh3glb1, Cdkl3, Lzts1, Camk1, Ube3a, Grin1, Lrp8, Epha4, Dhx36, Pqbp1, Ptpnz1, Nlgn3, Gsk3a, Cfl1, Arhgap33, Cyfip1, Dnm11, Pten, Ppp3ca, Cux1, Pak3, Cul7, Rapgef4, Ntrk2, Itsn1, Sdk1, Ppp1r9a, Apoe, Sdc2, Dnm3, Camk1d, Adgrb3, Fstl4, Pdlim5, Nlgn1, Mfn1, Tmem106b, Syne1, Sipal11, Kndc1, Sgk1, Lzts3, D130043K22Rik, Zmynd8, Rapgef2</i>
GO:0048814	regulation of dendrite morphogenesis	46/105	3.57E-03	<i>Trpc6, Arc, Ss18l1, Itpka, Sarm1, Actr3, Baiap2, Cdk5r1, Caprin1, Anapc2, Ngef, Cask, Srcin1, Dbn1, Sh3glb1, Cdkl3, Lzts1, Ube3a, Grin1, Lrp8, Epha4, Dhx36, Pqbp1, Ptpnz1, Nlgn3, Cfl1, Arhgap33, Dnm11, Pten, Ppp3ca, Cux1, Pak3, Cul7, Ppp1r9a, Sdc2, Dnm3, Adgrb3, Pdlim5, Nlgn1, Mfn1, Syne1, Sipal11, Kndc1, Sgk1, Lzts3, Rapgef2</i>
GO:0021766	hippocampus development	50/96	1.16E-05	<i>Srf, Rara, Cdk5r2, Ran, Tmem108, Nr4a3, Eif2b3, Dlx2, Slc32a1, Hdac2, Dlx1, Nme1, Uba6, Cdk5r1, Kif5b, Neurod6, Atat1, Lef1, Plxna3, Lrp8, Fgf13, Ncoal, Ogdh, Fez1, Cfl1, Kif3a, Ywhae, Pten, Bcan, Mkks, Tsc1, Nf2, Ppp1r9b, Casp3, Zbtb18, Bbs1, Smo, Dclk2, Emx2, Prox1, Kcna1, Mas1, Fezf2, Bbs2, Gli3, Bbs4, Mfsd2a, Mdk, Id4, Btg2</i>
GO:0021626	central nervous system maturation	6/7	1.58E-02	<i>Cdk5r2, Cend1, Cdk5r1, Arcn1, Grin1, Rere</i>
GO:0097485	neuron projection guidance	91/205	9.08E-06	<i>Etv4, Ephb2, Vegfa, Arx, Cxcr4, Unc5a, Efna3, Slit3, Nr4a3, Ephb3, Dlx5, Zswim8, Tubb3, Efnb2, Boc, Efna5, Lhx9, Sema6a, Bdnf, Efnb1, Epha8, Lmtk2, Cdk5r1, Sema6d, Etv1, Sema6c, Plxna2, Flrt2, Klif7, Efnb3, Wnt7b, Ptch1, Plxna1, Plxnd1, Vasp, Unc5b, Plxna3, Sema7a, Apbb2, Bcl11b, Efna2, Cntn2,</i>

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				<i>Epha4, Vstm2l, Dpysl5, Tubb2b, Apbb1, Cyfip1, Kif3a, Gap43, App, Arhgap35, Smad4, Mycbp2, Lhx2, Chl1, Foxp1, Foxg1, Kif5c, Ptprm, Ext1, Bmpr2, Nfasc, Ryk, Lamb2, Flrt3, Lgi1, Fzd3, Smo, Dcc, Ttc8, Ank3, Dscam, Sema3d, Sema3g, Nfib, Palld, Tgfb2, Rnfl165, Csf1r, Ephb1, Cxcl12, Fezf2, Matn2, Egr2, B3gnt2, Draxin, Gli3, Lama2, Bmpr1b, Ccdc141</i>
GO:0007611	learning or memory	110/279	3.72E-04	<i>Ldlr, Crhr1, Hmgcr, Srf, Ephb2, Hrh3, Foxo6, Adcy8, Htr7, Kcnk4, Shc3, Arc, Kit, Mef2c, Slc11a2, Kras, Kat2a, Calb1, Npix2, Fzd9, Kmt2b, Chrna7, Hif1a, Htr2a, Zfp385a, Arf4, Npas4, Grpr, Uba6, Bdnf, Sorcs3, Ngf, Cck, Abcc8, Grm4, Mapk8ip2, Adrb1, Tacr1, Cic, Creb1, Slc6a1, Cnr1, Pak7, Vdac3, Grin1, Cntn2, Fgf13, B4galt2, Syngap1, Ehmt2, Pja2, Ptprz1, Nlgn3, Gabra5, App, Pten, Dbi, Nrnx1, Gatm, Gm2a, Braf, Rag1, Prkcz, Ntrk2, Grm7, Cln8, Ppp1r9b, Apoe, Casp3, Itgb1, Tibk1, Lgmn, Btbd9, Tnr, Gmfb, Pak6, Grin2b, Ift20, Adgrb3, Atpl1a2, Vip, Pomk, Egr1, Tanc1, Lamb1, Ghssr, Ppt1, Itga3, Cln3, Rcan2, Kenk2, Cacna1c, Egr2, Cx3cr1, Paip2, Grin2a, Man2b1, Dcdc2a, Atxn11, Ctns, Fos, Rcan1, Drd5, Sgk1, Atxn1, Kenk10, Slc24a2, Mdk, Btg2, Jun</i>
GO:0001662	behavioral fear response	23/45	6.91E-03	<i>Adcyap1, Mef2c, Eif4e, Eif4g1, Bdnf, Cck, Cckbr, Bcl2, Mapk8ip2, Usp46, Vdac3, Gabra5, Dbi, Rag1, Grm7, Apoe, Grin2b, Grik2, Als2, Htr1a, Fbxl20, Gjal, Mdk</i>
GO:0007616	long-term memory	18/37	3.30E-02	<i>Ldlr, Srf, Adcy8, Arc, Kat2a, Calb1, Npas4, Tacr1, Grin1, Ehmt2, Pja2, Prkcz, Ntrk2, Apoe, Btbd9, Egr1, Ctns, Sgk1</i>
GO:0071560	cellular response to transforming growth factor beta stimulus	90/208	3.44E-05	<i>Smad2, Tgfb1l1, Mapk7, Trp53, Dnmt1, Mef2c, Tab1, Pin1, Gipcl, Men1, Akr1c18, Actr3, Usp15, Ing2, Dlx1, Stub1, Eid2, Cited1, Wnt7a, Chst11, Stk16, Nlk, Wnt4, Snx25, Ovol2, Src, Creb1, Epb4115, Dbn1, Nodal, Zfp703, Lemd3, Sirt1, Ptk2, Got1, Smurf1, Strap, Hspa5, Smad4, Yes1, Bmpr1a, Bmpr2, Itgb1, Cldn5, Adam17, Ldlrad4, Dusp15, Sox11, Wwox, Ppm1a, Pxn, Acvr1b, Itgb5, Ltbp3, Htra3, Col4a2, Tgfb2, Npnt, Smad6, Igflr, Prdm16, Skil, Bmp2, Cidea, Itga3, Dusp22, Sdcbp, Acvr11, Cd109, Tgfbr3, Cx3cr1, Pml, Twsg1, Apaf1, Adam9, Rasl11b, Gcnt2, Cflar, Fos, Fbn1, Smad1, Wfikk2, Eng, Bmpr1b, Pdgfd, Mstn, Cldn1, Htra1, Edn1, Jun</i>
GO:0032680	regulation of tumor necrosis factor production	60/145	3.52E-03	<i>Rara, Ptger4, Arhgef2, Chrna7, Hdac2, Hsf1, Dhx9, Rasgrp1, Tnfaip3, Trex1, Abcc8, Arfgef2, Cd24a, Trim27, Ifng, C1qtnf4, Hspd1, Akap8, Sirt1, Mif, Gstp1, Ube2j1, Ncl, Cyp2j6, Foxp1, Bpi, Tbc1d23, Cyba, Clu, Ptafr, Sirpa, Errfi1, Tlr1, Nod1, Lgals9, Zbtb20, Ddt, Cd300ld, Ghssr, Flt3, Tlr4, Cidea, Syt11, Ddx58, Il18, Myd88, Tspo, Cx3cr1, Ifih1, Tlr9, Mavs, Tlr6, Ly96, H2-T23, Haver2, Arg2, Ccl3, Ticam1, Pik3r1, Arid5a</i>
GO:0071354	cellular response to interleukin-6	16/31	2.66E-02	<i>Phb, Ptpn2, Fer, C1qtnf4, Ctr9, Cfl1, Rela, Smad4, Gab1, Fgf23, Il6st, Il6ra, St3gal6, Pid1, St18, Stat3</i>
GO:0071706	tumor necrosis factor superfamily cytokine production	63/150	1.75E-03	<i>Rara, Ptger4, Arhgef2, Chrna7, Hdac2, Hsf1, Dhx9, Rasgrp1, Tnfaip3, Trex1, Abcc8, Arfgef2, Cd24a, Trim27, Ifng, C1qtnf4, Hspd1, Akap8, Sirt1, Mif, Gstp1, Ube2j1, Ncl, App, Cyp2j6, Foxp1, Azi2, Bpi, Tbc1d23, Cyba, Clu, Ptafr, Sirpa, Errfi1, Tlr1, Nod1, Lgals9, Zbtb20, Ddt, Cd300ld, Ghssr, Flt3, Tlr4, Cidea, Cd274, Syt11, Ddx58, Il18, Myd88, Tspo, Cx3cr1, Ifih1, Tlr9, Mavs, Tlr6, Ly96, H2-T23, Haver2, Arg2, Ccl3, Ticam1, Pik3r1, Arid5a</i>

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GO:0002732	positive regulation of dendritic cell cytokine production	6/6	3.89E-03	<i>Ddx21, Dhx36, Ddx1, Ddx58, Mavs, Ticam1</i>
GO:0034968	histone lysine methylation	47/104	1.49E-03	<i>Setd1b, Dnmt1, Smyd2, Prdm8, Suv39h1, Lmna, Nelfe, Iws1, Kmt2b, Bcor, Setd6, Hdac2, Setd1a, Men1, Phf1, Rtf1, Cxxc1, Wdr5, Suv39h2, Zfp335, Phf19, Nelfa, Setd3, Prdm6, Rif1, Smarcb1, Ash2l, Ctr9, Ehmt1, Sirt1, Kdm4b, Pih1d1, Ehmt2, Rlf, Asxl2, Smad4, Ash1l, Kmt2e, Kdm4a, Tet3, Ogt, Kdm3a, Tet2, Setmar, Jarid2, Ezh1, Setd7</i>
GO:0031065	positive regulation of histone deacetylation	12/19	1.02E-02	<i>Vegfa, Trp53, Ing2, Srebf1, Akap8, Ctbp1, BC004004, Nipbl, Lpin1, Prkd2, Pml, Bcl6</i>
GO:0090239	regulation of histone H4 acetylation	9/13	1.52E-02	<i>Iws1, Smarcb1, Ctbp1, Sirt1, Pih1d1, Zmpste24, Spi1, Atg7, Atg5</i>
GO:0097306	cellular response to alcohol	38/84	4.87E-03	<i>Aacs, Tnc, Adcy8, Ptger4, Cdk4, Larpl, Akr1c18, Efna5, Npas4, Hmgcs1, Acaca, Prkaa2, Fancb, Akt1, Blm, Ptch1, Inhbb, Akap8, Ireb2, Sod2, Dnmt3a, Aifm1, Arpc2, Pten, Lancl2, Spi1, Mlc1, Lrp6, Smo, Iptr2, P2ry6, Cttna1, Klf4, Hdac8, Abca1, Gramd1b, Foxo3, Trp53inp1</i>
GO:0006066	alcohol metabolic process	124/298	7.45E-06	<i>Ldlr, Insig1, Mvd, Hsd17b7, Dhcr24, Hmgcr, Sqle, Fdfl1, Fdps, Cyp51, Myk, Nsdhl, Arv1, Lss, Pgp, Clcn2, Akr1b3, Thrb, Sptlc2, Sc5d, Srebf2, Akr1c18, Itpka, Rdh11, Ppip5k2, Tpi1, Hmgcs1, Prkaa2, Ces1d, Mtmr7, Wnt4, Vldlr, Ifng, Srebf1, Akr7a5, Galk1, Itpk1, Por, Plcd1, Got1, Ipmk, Nfe2l1, Gpd2, Mtmr2, App, Pten, Nudt3, Sod1, Kcnma1, Gdel, Ttc39b, Cat, Plcgl, Dgat2, Rest, Cebpa, Dpm2, Npc2, Cln8, Abhd4, Apoe, Aldh2, Scp2, Tkfc, Aldh3a2, Dkk3, Fgf2, Prkg1, Ptafr, Mt3, Rdh14, Ebp, Gdpd1, Pmp22, Akr1b10, Lrp5, Plek, Erlin2, Aldh1a1, Ip6k2, Mttp, Dpagt1, Dhhrs4, Akr1b8, Inpp5a, Acadvl, Gk5, Inpp1, Ip6k1, Soat1, Lrp1, Gpd1, Abcg1, Bmp2, Sgpp1, Acadl, Mas1, P2ry6, Lcat, Sorl1, Mbtps1, Slc37a2, Adcyap1r1, Asah2, Inpp5k, Dhhrs3, Ephx1, Lepr, Abca1, Limal, Cln6, Pecr, Rdh12, Acer2, Bmp6, Sord, Cyp39a1, Fgf1, Sult2b1, Napepld, Chka, Inpp5j, Cyp1b1, Hmgcs2</i>
GO:0034308	primary alcohol metabolic process	29/67	3.02E-02	<i>Clcn2, Akr1b3, Akr1c18, Rdh11, Wnt4, Akr7a5, Kcnma1, Gdel, Dgat2, Rest, Abhd4, Aldh2, Aldh3a2, Dkk3, Rdh14, Gdpd1, Akr1b10, Aldh1a1, Akr1b8, Gpd1, Bmp2, Slc37a2, Dhhrs3, Pecr, Rdh12, Bmp6, Napepld, Chka, Cyp1b1</i>
GO:0007548	sex differentiation	104/287	1.17E-02	<i>Vgf, Dhcr24, Ccnd1, Rara, Adcyap1, Fst, Vegfa, Nefh, Fkbp4, Slit3, Ptger4, Fgf9, Kit, Phb, Kmt2b, Adrm1, Greb1l, Ing2, Lhx9, Dach1, Eif2b4, Hmgcs1, Ybx3, Bax, Atn1, Bcl2, Flna, Tyro3, Wnt4, Asb1, Src, Zfp2, Idh1, Rrm1, Fancf, Sirt1, Ube3a, Bcl2l1, Nco1, Eif2s2, Sod1, Hspa5, Smad4, Schip1, Lgr4, Cntfr, Ermp1, Bmpr1a, Pcytlb, Nipbl, Pdgfra, Jmjd1c, Tbc1d20, Rnf38, Casp3, Tbx3, Cst3, Tiparp, Lrp2, Lrp6, Gas2, Kitl, Acvr1b, Stat5a, Fgf7, Fnac3a, Adgrg1, Dach2, Lhfp12, Prdx4, Rab13, Tgfb2, Insr, Nrip1, Plekha1, Brca2, Mas1, Bmp4, Cttna1, Trp63, Ago4, Kdm5a, Rad21l, Serpinb6a, Sox8,</i>

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				<i>Adcyap1r1, Col9a3, Tlr9, Ctsl, Map7, Wdr19, Dmrt1, Adamts1, Foxo3, Abcb1a, Bmpr1b, Bmp6, Ar, Lfng, Hmgb2, Osr1, Arid5b, Mertk, Bcl2l11</i>
