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## Microarray genelists

**1. - Genes upregulated by iFGFR1**

Affymetrix probe set	GenBank Accession	Gene Symbol	Fold change
XI.736.1.S1_at	AF310008	TA-2	19.12
XI.23988.1.S1_at	BJ044287	Rexp44 mRNA	14.34
XI.4965.1.S1_at	AJ278067	irg1	6.99
XI.24294.1.S1_at	BJ098811	Cico01 mRNA	6.61
XI.24793.1.S1_at	CB563927	tspan1	6.47
XI.24337.1.A1_at	CB564601	Predicted NADPH oxidase 1-like	4.39
XI.8124.1.S1_at	BJ045090	MGC115642	4.32
XI.637.1.A1_at	AF250345	egr1-a	3.81
XI.9671.1.S1_at	BC042349	capn8-a	3.64
XI.14397.1.S2_at	BC044326	nek6	2.78
XI.16457.1.A1_at	CB563787	junb	2.73
XI.4789.1.S1_at	BC043748	MGC52875	2.72
XI.10415.1.A1_at	BF024850	MGC80142	2.66
XI.24218.1.S1_at	CB943601	dynll1-a	2.60
XI.23897.1.S1_at	BQ386549	cnfn-a	2.58
XI.9549.1.S1_at	AF374473	lmo2	2.42
XI.708.1.S1_at	AF283562	lefty-a	2.41
XI.880.1.S1_at	L07538	wnt3a	2.37
XI.5082.1.A1_at	BF072347	MGC68521	2.34
XI.1044.1.S1_at	AF283563	lefty-b	2.32
XI.12130.1.S1_at	M24752	hoxa7	2.31
XI.7720.1.A1_at	BF615090	Meso05 mRNA	2.20
XI.10684.1.A1_at	BE505501	Unknown	2.17
XI.16649.1.A1_at	BJ077367	dlgap4	2.15
XI.22857.1.A1_at	BJ088428	Predicted Plekho1	2.15
XI.24572.1.S1_a_at	AB003078	tnnc2	2.14
XI.24608.1.A1_at	CB561182	lap2omega/tmpo	2.14
XI.11965.1.S1_at	AF331825	LOC398232	2.14
XI.1465.1.S1_at	BC046253	ido1	2.14
XI.216.2.S1_a_at	Y08734	mst1	2.12
XI.18073.1.A1_at	BG885063	Unknown	2.11
XI.24205.1.S1_at	BJ086130	Predicted itga6-like	2.03
XI.23718.1.A1_at	BJ075987	mmp14	2.03
XI.21639.1.S1_at	AJ319749	hoxa10	2.03
XI.1358.1.S1_at	U04302	cdx1	2.02
XI.2665.2.A1_at	BM172523	des.1-b	2.00
XI.14334.1.S1_at	BI444166	Predicted phldb1	2.00
XI.21634.1.S1_at	CA792675	msx2	2.00
XI.12011.1.A1_at	BG812177	Predicted fbrsl1	1.97
XI.15701.1.S1_at	AY150813	kremen2	1.96
XI.1154.1.S1_at	S74933	crabp2	1.94

XI.6263.1.S1_at	CB563516	mylpf	1.94
XI.49.1.S1_at	X57234	wnt8a	1.93
XI.37.1.S1_at	X98454	ventx2.2	1.92
XI.5846.1.S1_at	BC046739	MGC53823	1.91
XI.11964.1.S2_at	AF369901	spry2	1.91
XI.16206.1.A1_at	BM172525	pnf	1.90
XI.21864.1.S1_at	AY167741	hoxc10	1.90
XI.2936.2.S1_x_at	BQ735779	c8orf42	1.89
XI.15374.1.A1_at	BJ077463	Rexp91 mRNA	1.89
XI.558.1.S1_at	AF149307	ventx3.2	1.89
XI.1056.2.S1_a_at	AJ009308	myl1	1.88
XI.22960.2.A1_at	BJ085655	creb3	1.87
XI.13835.1.S1_at	BC043838	greb1	1.87
XI.6592.1.A1_at	BE491206	Predicted laminin subunit	1.87
XI.1035.1.S1_at	U17598	elavl3	1.84
XI.802.1.S1_at	BG016128	LOC397753	1.83
XI.14402.1.A1_at	BJ082817	LOC733321	1.83
XI.4725.1.S1_at	AB072909	cldn4L2	1.82
XI.1055.1.S1_at	CB562934	MGC53335	1.81
XI.15887.1.S1_x_at	BJ088835	Predicted mpc2	1.81
XI.7618.2.A1_a_at	BG346681	plscr1	1.81
XI.1002.1.S1_at	U43223	dusp6	1.81
XI.23957.1.S1_at	BI313816	cfp	1.80
XI.13621.1.A1_at	BJ087999	Predicted HoxB8	1.78
XI.10874.1.A1_at	BU907171	gprc5c	1.78
XI.16519.1.A1_at	BI446890	hoxd3	1.77
XI.4880.2.S1_a_at	CB563123	sat1-a	1.76
XI.8315.1.A1_at	AW147996	nuak2	1.75
XI.13086.1.A1_at	BF615358	fam46a	1.75
XI.9286.1.S1_at	BC049827	add1	1.74
XI.868.1.S1_at	L19716	wnt5a	1.73
XI.11165.1.A1_at	AW767805	Predicted LOC101734664	1.73
XI.9871.1.A1_s_at	BJ053647	thbs4	1.73
XI.1210.1.S1_at	X06592	hoxb7-b	1.73
XI.26209.1.S1_at	BJ056944	Predicted LOC419301	1.73
XI.2955.1.A1_at	BJ091076	Predicted fam181a	1.71
XI.209.1.S1_at	AW164925	Predicted hes6.1	1.71
XI.1073.1.S1_at	L23542	wnt11b	1.70
XI.15165.1.A1_at	BJ077194	MGC131003	1.69
XI.25941.1.A1_at	BM180508	tpbg	1.69
XI.2466.1.A1_at	BG020193	traf4b	1.68
XI.8010.1.A1_at	AW782315	Unknown	1.68
XI.13344.1.A1_at	BJ050847	Unknown	1.68
XI.10269.1.S1_at	U02034	cdx4	1.67
XI.25587.1.A1_at	BQ387107	tmcc1	1.66

XI.23739.1.A1_at	CB564190	cdx1	1.66
XI.1202.1.S1_at	CA986862	gs17	1.66
XI.2244.1.A1_at	BG439336	Predicted MGC82906	1.65
XI.13343.1.A1_at	BJ089629	Predicted HoxD9	1.65
XI.1039.1.S1_at	AB037688	ptch2-b	1.65
XI.2235.1.A1_at	BJ082132	Unknown	1.65
XI.15436.1.A1_at	BJ082303	Predicted Irrfip1	1.65
XI.61113.1.A1_at	AW767973	Predicted NEDD4-binding protein 3-like	1.65
XI.262.1.S1_at	D10259	dlx6-b	1.64
XI.3540.1.S1_at	AF131890	LOC398134	1.63
XI.13987.1.S1_x_at	BQ731171	schip1	1.62
XI.644.1.S1_at	AF223426	foxd5b	1.62
XI.25397.1.A1_at	BE026782	Unknown	1.62
XI.11117.1.S1_at	BI445771	gsto1	1.62
XI.21869.1.S1_at	BC045041	LOC398437	1.61
XI.23670.1.A1_at	CB563705	Predicted tspan4	1.61
XI.6155.1.A1_at	BJ052899	tfap2b	1.61
XI.26380.1.A1_at	BJ050459	cdc42se2-a	1.61
XI.15872.2.S1_at	BJ086614	fbn3	1.60
XI.11057.1.A1_at	AW764507	itgb1bp2	1.60
XI.642.1.S1_at	AF162782	foxd4l1.1-a	1.60
XI.280.1.S1_at	AF061727	serpind1	1.60
XI.1265.1.S1_at	BG022051	pou2f1	1.59
XI.16273.1.A1_at	BJ045836	hoxc8	1.59
XI.15623.1.A1_at	CB756273	pfkfb3	1.59
XI.1209.1.S1_at	X12499	hoxc6	1.59
XI.1491.1.S1_at	D50552	sox13	1.59
XI.6247.1.S1_at	BC047252	hyal2-b	1.58
XI.2061.1.A1_at	BI443086	tmem169	1.58
XI.26119.1.A1_at	BJ043617	gprc5c	1.58
XI.17496.1.S1_at	BG347936	wnt5b	1.58
XI.15990.1.A1_at	BJ043686	tmem69	1.57
XI.803.1.S1_at	M23916	hoxb7-a	1.57
XI.16181.1.A1_at	BJ045364	fbxl22	1.57
XI.1082.1.S1_at	S93559	foxa4-b	1.56
XI.2507.1.A1_at	BJ078949	Unknown	1.56
XI.862.1.S1_at	L28111	dio3	1.56
XI.16131.1.S1_at	BJ045564	nexn	1.56
XI.26220.1.A1_at	BJ085819	MGC84721	1.55
XI.514.1.S1_at	M77243	t-a	1.55
XI.22996.1.S1_at	BC046259	hapln3	1.55
XI.457.1.S1_at	AF065135	gdf3	1.55
XI.15270.1.A1_at	BI447679	Unknown	1.55
XI.16035.1.A1_at	BJ081350	Unknown	1.54

XI.9439.1.S1_at	BC041731	hoxa3	1.54
XI.1873.1.S1_at	BM179250	cdh2-a	1.54
XI.688.1.S1_at	AF281080	nkx2-1	1.54
XI.1289.1.S1_at	U40457	tdgf1	1.53
XI.8165.1.S1_at	CB207508	tcf12	1.53
XI.25895.1.A1_at	BJ075592	LOC494785	1.53
XI.1817.1.A1_at	BJ049297	smyd1	1.53
XI.14231.1.A1_at	BM191311	st3Gal-II	1.52
XI.24158.1.A1_at	CB561731	c4orf31	1.52
XI.17342.1.A1_at	BJ080764	Predicted prtg	1.52
XI.7766.1.A1_at	BI443123	Unknown	1.52
XI.18908.1.A1_at	BI447750	stard4	1.52
XI.14733.1.A1_at	BG552346	insm1	1.52
XI.13019.1.S1_at	BC041294	rasl11b	1.51
XI.16321.1.A1_at	BJ047359	hes5.2-b	1.51
XI.5454.1.S1_at	AF394111	xmc	1.51
XI.6074.1.A1_at	AW767695	lox1	1.51
XI.15054.1.S1_at	BC043745	fbln1	1.51
XI.11405.1.S1_a_at	BC044063	atp2a2	1.50
XI.7620.1.S1_at	BC045059	arl5b-b	1.50
XI.5940.1.A1_at	AW766423	Predicted KIF26A-like	1.50

## 2 Genes downregulated by iGFR1

Affymetrix probe set	GenBank Accession	Gene name	Fold Change
XI.25847.1.S1_at	BC052102	agr2	-16.43
XI.6266.1.S1_at	AB105372	itln1	-8.74
XI.2924.1.S1_at	BG023525	MGC53311	-8.63
XI.15702.1.A1_at	BJ076178	ATPase 13A4-like	-7.36
XI.1685.1.S1_at	AF314056	LOC398260	-7.32
XI.847.1.S1_s_at	L10987	otog	-7.14
XI.7213.2.A1_at	BJ047679	klhl24	-6.88
XI.18858.1.A1_at	BI446930	Unknown	-6.37
XI.20089.1.S1_at	BC042303	foxi1	-5.88
XI.909.1.S1_at	AB018694	xepsin	-5.37
XI.15894.1.S1_at	CD099356	ppl	-4.99
XI.6048.1.S1_at	BJ044577	fuclectin	-4.72
XI.10583.1.S1_at	BC042234	slc3a2	-4.71
XI.5324.1.S1_at	BJ043563	otog	-4.66
XI.24565.1.A1_at	BG485946	Unknown	-4.58
XI.9836.1.A1_at	BJ075817	liph-a	-4.33
XI.1589.1.S1_at	AF025474	agr3	-4.27
XI.1616.1.A1_at	BJ083887	fam115a	-3.94
XI.16262.1.A1_at	BJ051781	Unknown	-3.86
XI.1683.1.S1_at	BJ044640	MGC68910	-3.80
XI.16436.1.A1_at	BJ081297	Predicted G protein receptor	-3.75
XI.16543.1.S1_at	AF513854	Ras dva	-3.68
XI.7354.1.A1_at	AW766695	sytl2	-3.60
XI.26141.1.S1_at	BC044313	MGC80993	-3.60
XI.15156.1.S1_at	CB562846	vill	-3.53
XI.13575.1.A1_at	BU913085	b3gnt1	-3.50
XI.12727.1.A1_at	BM192846	fa2h	-3.49
XI.24199.1.A1_at	CB756654	Unknown	-3.41
XI.10362.1.A1_at	BF072092	ca12	-3.31
XI.16466.1.A1_at	BJ082483	Predicted lectin	-3.31
XI.21983.1.S1_at	M11940	xk81a1	-3.26
XI.11136.1.A1_at	BJ045099	LOC100037100	-3.25
XI.5912.1.A1_at	BG020669	eppk1	-3.24
XI.1479.1.A1_at	BG038587	Unknown	-3.21
XI.8098.1.A1_at	BJ090592	LOC494641	-3.20
XI.522.1.S1_at	AF217647	pitx1	-3.17
XI.13893.1.A1_at	BJ083655	fam3d	-3.10
XI.22874.1.A1_at	BJ090165	ubp1	-3.06
XI.15089.1.A1_at	BJ056659	lgals9	-3.04
XI.5930.1.A1_at	AW766360	aldh11	-2.99
XI.11187.1.A1_at	AW782510	Predicted adenovirus receptor homolog	-2.97
XI.13767.1.A1_at	BJ075935	MGC85058	-2.96

XI.10855.1.A1_at	BE575595	grhl3	-2.96
XI.5486.1.A1_at	BJ091754	Unknown	-2.95
XI.16487.1.S1_at	BC045029	ehd4	-2.84
XI.7307.1.S1_at	BJ084368	LOC443682	-2.83
XI.2610.1.S1_at	M60768	anxa2-a	-2.82
XI.7848.1.A1_at	AW148259	Unknown	-2.79
XI.23326.1.S1_at	BC045031	krt-b	-2.77
XI.16658.3.A1_at	BJ083088	kcnc3	-2.74
XI.16672.1.S1_at	BC044108	kitlg	-2.74
XI.15841.1.A1_at	BJ049236	Predicted anion transporter	-2.68
XI.9961.1.A1_at	BG812624	syt1	-2.67
XI.11145.1.A1_at	AW766955	Unknown	-2.66
XI.14452.1.A1_at	BG811297	bcat1	-2.62
XI.13681.1.A1_at	BJ092589	znf750-b	-2.61
XI.25460.1.A1_at	AW148116	grhl1	-2.60
XI.9076.1.S1_at	BC047968	nkx3-1-a	-2.58
XI.16589.1.A1_at	BJ080730	Predicted sulfotransferase	-2.58
XI.13862.1.A1_at	BJ078064	Unknown	-2.57
XI.1450.1.A1_at	BG579799	Possible ATPase	-2.55
XI.7017.1.S1_at	BF615728	krt12	-2.49
XI.6104.1.A1_at	BJ076947	rab27a	-2.49
XI.2418.1.A1_at	BG023326	ccno	-2.48
XI.7213.3.S1_a_at	BQ737049	LOC100158288	-2.48
XI.11935.1.A1_at	BJ088440	Possible syt1	-2.47
XI.9959.1.A1_at	BJ045735	degs3	-2.45
XI.25518.1.S1_at	BJ053813	LOC443659	-2.44
XI.7756.1.S1_at	BC043737	znf750-a	-2.42
XI.23708.1.A1_at	AI031433	MGC81939	-2.37
XI.7213.1.S1_at	BC042338	cmahp	-2.36
XI.13291.1.A1_at	BJ078773	Predicted FAM55c-like	-2.36
XI.16504.1.A1_at	BJ090565	fut6	-2.36
XI.16435.1.A1_at	BJ089860	gdpd1	-2.35
XI.793.1.A1_at	M76565	gata3	-2.33
XI.21868.1.S1_at	BC044973	elf-1	-2.33
XI.10200.1.A1_at	BJ084274	tmem181	-2.32
XI.25344.1.A1_at	BE678810	Unknown	-2.31
XI.1605.1.S1_at	BJ080966	Predicted envoplakin	-2.30
XI.9656.1.S1_at	BC046858	glb112	-2.29
XI.4183.2.A1_at	BJ051393	LOC100158277	-2.29
XI.23835.1.A1_s_at	BJ081027	Predicted prr4	-2.29
XI.16564.1.A1_s_at	BJ054555	fam3a	-2.26
XI.1076.1.S1_at	BC046838	ag1	-2.26
XI.16096.1.A1_at	BJ046407	tmem45b	-2.25
XI.16320.1.S1_at	BC046669	anxa9	-2.25
XI.104.1.S1_at	AJ005787	pitx2-a	-2.24



XI.5296.1.A1_at	BJ080084	Unknown	-2.23
XI.21239.1.A1_at	BU908560	EIG121L	-2.23
XI.12647.1.A1_at	BJ088990	Predicted paqr5	-2.19
XI.23267.1.S1_at	BC044298	dnajb14	-2.19
XI.1380.1.S1_at	BC051601	gale	-2.18
XI.1003.1.S1_at	U28370	hesx1-a	-2.18
XI.2852.1.A1_at	BG555933	Predicted slc16a3	-2.18
XI.9651.1.A1_at	BG161001	Unknown	-2.18
XI.21677.2.A1_at	BJ057218	Lapl04 mRNA	-2.17
XI.16144.1.A1_at	CB592728	cd81-a	-2.16
XI.15347.1.A1_at	BJ052833	eps8l1	-2.15
XI.9058.1.A1_at	BG347632	mmp3	-2.14
XI.2659.1.S1_at	BC045045	atp12a-b	-2.13
XI.8004.1.S1_at	AW199480	Predicted mob1b	-2.13
XI.16249.1.A1_at	BJ091874	Predicted mucin-4-like	-2.12
XI.1466.1.S1_at	CB560706	slc35a3.2	-2.12
XI.12819.2.A1_at	BJ056722	gmpr2	-2.09
XI.24099.1.A1_at	CB562751	dvl3	-2.07
XI.9576.1.S1_at	CB560639	ca2	-2.06
XI.10887.1.A1_at	BM172535	LOC496380	-2.06
XI.973.1.S1_at	L47990	gbx2.2	-2.06
XI.12147.1.A1_at	BJ051202	Unknown	-2.06
XI.15019.1.A1_at	BQ384301	Predicted LOC100127290	-2.05
XI.6308.1.A1_at	AW632863	mtus1	-2.05
XI.14521.1.S1_at	BC041533	LOC398688	-2.04
XI.15893.1.A1_at	BJ056712	Predicted ankyrin repeat domain 43	-2.04
XI.1242.1.S1_at	BC043635	arg1	-2.03
XI.8929.1.A1_at	BG554539	Predicted ATPase subunit C2-like	-2.03
XI.26274.1.A1_at	BE189559	MGC82269	-2.02
XI.509.1.S1_at	BC046269	atp1b2	-2.02
XI.15540.1.A1_at	BJ054782	atf5.2	-2.02
XI.22091.1.A1_at	BE188962	Predicted superoxide dismutase	-2.02
XI.2854.1.S1_at	BC043826	fam3a	-2.02
XI.16658.2.A1_at	BJ090409	Predicted FAM55D-like	-2.01
XI.1604.1.A1_at	BJ057329	MGC78986	-2.01
XI.4812.1.A1_at	BG555240	Unknown	-2.00
XI.3804.1.A1_at	BJ080125	rab25	-2.00
XI.17407.1.S1_at	BM928994	sgsm3	-2.00
XI.792.1.S1_at	M76564	gata2	-1.99
XI.24155.1.A1_at	BJ090126	ada	-1.98
XI.2256.1.S1_at	BJ056357	hal.1	-1.97
XI.23515.1.S1_at	BC042293	cldn1	-1.97
XI.656.1.S1_at	AF163313	cml	-1.97
XI.15637.1.A1_at	BJ067363	capn9	-1.96
XI.3793.1.S1_at	AB031348	xenf	-1.95

XI.15310.1.A1_at	BJ045220	gmppb-b	-1.95
XI.15698.1.A1_at	BG020321	Predicted tmem63A	-1.95
XI.21384.1.S1_at	AB085630	cat1	-1.95
XI.23980.1.A1_at	AW764542	Predicted LOC100494996	-1.94
XI.15057.1.A1_at	BJ081522	kiaa1324l	-1.94
XI.30.1.S1_at	Z48770	kit-a	-1.93
XI.15857.1.A1_at	BJ099826	tmem38b-b	-1.92
XI.21926.1.S1_at	BC042252	prp5	-1.92
XI.460.1.S1_at	BG021592	fzd4	-1.92
XI.14093.2.A1_at	BJ079222	gna14-a	-1.91
XI.14093.2.A1_at	BJ079222	gna14-a	-1.91
XI.9247.1.A1_at	BG438869	elf1	-1.91
XI.23255.1.S1_at	AF176665	mtus1	-1.91
XI.18645.1.A1_at	BI443285	Unknown	-1.90
XI.1241.1.S1_at	D83649	sox7	-1.89
XI.5792.1.A1_at	BJ053583	glo1	-1.89
XI.18670.1.A1_at	BI443571	MGC115605	-1.88
XI.23845.2.S1_at	CB756807	capns2	-1.88
XI.10620.1.S1_at	BJ047148	slc1a5	-1.88
XI.15658.1.A1_at	BJ047928	scel	-1.88
XI.15475.1.A1_at	BJ083397	U8 snoRNA	-1.87
XI.7111.1.S1_at	BJ086019	efhd1	-1.86
XI.7483.1.S2_at	L29057	cdh1	-1.86
XI.24073.1.A1_at	CB561420	fut1	-1.86
XI.14584.1.A1_at	BJ091208	sbk1	-1.86
XI.16031.1.A1_at	CD326985	Predicted uaca	-1.84
XI.8104.1.A1_at	BJ056584	LOC446305	-1.84
XI.21920.1.S1_at	BC043746	slc19a3	-1.84
XI.7751.1.S1_at	AF536830	menf.1	-1.83
XI.5611.1.S1_at	BJ045338	elovl7	-1.83
XI.16165.1.A1_at	BJ051817	Predicted nasek-a	-1.83
XI.3829.1.S1_at	BC041743	osbp2	-1.82
XI.3898.1.A1_at	BF232494	Predicted vangl1	-1.82
XI.16562.1.A1_at	BJ084204	Predicted gcm1	-1.82
XI.627.1.S1_at	AF183571	six3-a	-1.81
XI.17336.1.A1_at	BJ084049	rassf10	-1.81
XI.14862.1.A1_at	BG885914	Predicted rnf149	-1.81
XI.6133.1.S1_at	M11032	krt-a	-1.81
XI.10182.1.A1_at	BJ043711	Predicted LOC100487499 isoform 1	-1.80
XI.13377.1.A1_at	BJ051589	kctd15	-1.80
XI.8717.1.S1_at	BC041217	cd81-a	-1.79
XI.8076.1.A1_at	BF428354	MGC82342	-1.79
XI.5611.2.A1_at	BJ043782	MGC80262	-1.78
XI.12539.1.A1_at	BJ087662	Predicted pmm2	-1.78
XI.14870.1.A1_at	BJ077643	hk2	-1.78

XI.11134.1.S1_at	AF465789	LOC398485	-1.77
XI.6421.1.A1_at	AW633197	tob1	-1.77
XI.15247.1.A1_at	BI443131	rnft1	-1.77
XI.19422.1.A1_at	BI444791	c1orf21	-1.76
XI.3818.1.S1_at	AF368041	snai2-a	-1.76
XI.16875.1.A1_at	AW199209	LOC100137681	-1.76
XI.5854.1.A1_at	BJ091200	angpt4	-1.76
XI.24090.1.A1_at	CB561571	kit-b	-1.74
XI.9325.1.A1_at	BJ080496	Unknown	-1.74
XI.511.1.S1_at	AW164934	crx-b	-1.74
XI.24391.1.A1_at	BJ092498	Predicted hexokinase -2-like	-1.74
XI.6160.1.S1_at	BC041187	MGC52585	-1.73
XI.25380.1.A1_at	BJ044089	Predicted LOC100101281	-1.73
XI.6291.1.A1_at	BJ043812	cldn4	-1.73
XI.15574.1.A1_at	BJ083528	tfcp2l1-b	-1.72
XI.14620.1.A1_at	BJ056766	ctn4	-1.72
XI.16197.1.A1_at	BJ044460	fezf2	-1.72
XI.139.1.S1_at	X74316	foxi4.2-b	-1.72
XI.1451.1.A1_at	BG579801	tbc1d24.2	-1.72
XI.3776.1.S1_at	CB563309	capn1	-1.72
XI.841.1.S1_a_at	M23238	pdgfa	-1.71
XI.25780.1.S1_at	BG016938	gdf1	-1.71
XI.1525.1.A1_at	BJ056088	Predicted LOC100485898	-1.71
XI.14738.1.A1_at	BM179208	cab39l	-1.71
XI.25758.1.S1_at	CD325687	MGC68655	-1.69
XI.15706.1.A1_at	BJ092390	nek8	-1.69
XI.5424.1.A1_at	BI444735	dapk2	-1.69
XI.1654.1.A1_at	BG019624	Predicted zfp729-like	-1.69
XI.10638.1.A1_at	BE505190	tnfrsf1a	-1.68
XI.15970.1.A1_at	BJ051939	MGC80468	-1.68
XI.6005.1.S1_at	CB943930	Unknown	-1.68
XI.2098.1.A1_at	BG486824	LOC503673	-1.67
XI.9829.1.S1_at	BG018208	foxn4	-1.67
XI.10967.1.S1_at	BE026444	Predicted LOC100036671	-1.67
XI.16033.1.A1_at	CD362297	Predicted LOC100158300	-1.67
XI.8108.1.A1_at	BJ047639	LOC100036828	-1.66
XI.24323.1.A1_at	CB564657	tesc	-1.66
XI.15597.1.A1_at	BJ080111	Predicted slc35d1	-1.66
XI.7078.2.S1_a_at	BJ078851	amd1	-1.65
XI.8089.1.A1_at	BQ384705	hs3st1	-1.65
XI.8089.1.A1_at	BQ384705	hs3st1	-1.65
XI.131.1.S1_at	X60099	hesx1-b	-1.65
XI.15963.1.S1_at	BC044687	sfrp2	-1.64
XI.142.1.S1_at	X52691	c-ets-1b	-1.64
XI.11114.1.A1_at	BJ051736	LOC100337617	-1.64

XI.4128.1.S1_at	BJ047451	LOC100037197	-1.63
XI.24966.1.S1_at	BC049292	MGC53924	-1.63
XI.15272.1.A1_at	BJ051205	st3gal4	-1.62
XI.15265.1.A1_at	BJ047004	Unknown	-1.61
XI.4752.1.A1_at	BJ057383	Predicted tbc1	-1.61
XI.7919.1.S1_at	BJ090309	MGC81684	-1.60
XI.2986.1.S1_x_at	BJ099184	Unknown	-1.60
XI.14049.2.A1_at	BJ083561	Predicted LOC100497155	-1.60
XI.18686.1.A1_at	BJ045936	Predicted ankrd9	-1.60
XI.4275.1.S1_at	BC043822	nek2	-1.60
XI.15572.1.A1_at	BJ088128	hes7.1	-1.60
XI.3204.1.S1_at	U62807	hspa5	-1.60
XI.13505.1.A1_at	AW767880	Predicted hexd-like	-1.59
XI.5679.1.S1_at	BE026676	nipal4	-1.59
XI.6436.1.S1_at	BQ398712	Predicted LOC101730357	-1.59
XI.4391.1.A1_at	BE678263	mreg	-1.59
XI.4460.1.A1_at	BM262138	MGC84886	-1.59
XI.10784.1.A1_at	BE508497	Unknown	-1.59
XI.6002.1.A1_at	AW766813	fam154b	-1.58
XI.1395.1.A1_x_at	BG022417	LOC496316	-1.58
XI.16025.1.A1_at	BJ043667	gnb3	-1.58
XI.8559.1.A1_at	BJ043709	Predicted pcdh10 gene product	-1.58
XI.11931.1.A1_at	BG893379	Predicted syt2	-1.58
XI.15997.2.A1_at	BJ050776	sytl1	-1.58
XI.16087.1.A1_x_at	BJ046353	MGC81163	-1.57
XI.13379.1.A1_at	BJ085119	Predicted trpm4	-1.57
XI.1424.1.S1_a_at	BJ043534	LOC100127277	-1.57
XI.16451.1.A1_at	BJ049278	LOC495248	-1.57
XI.22394.1.S1_at	AW633564	gmppa-b	-1.56
XI.14426.1.A1_at	BJ081435	Predicted krt8.2	-1.56
XI.551.1.S1_at	BI449397	pacsin2	-1.56
XI.13161.1.S1_at	BU901413	b3gnt2	-1.56
XI.1016.1.S1_at	AB007037	fgfr4-b	-1.56
XI.18621.1.A1_s_at	BI443159	Unknown	-1.56
XI.12881.2.A1_at	BJ084140	MGC81002	-1.56
XI.9387.1.A1_at	BJ052978	Unknown	-1.55
XI.693.1.S1_at	AF233632	camk2g	-1.55
XI.11329.1.S1_at	BF072268	Unknown	-1.55
XI.23696.1.S1_at	BC048022	tsga10	-1.55
XI.25710.1.S1_at	BF048877	hist1h2am	-1.55
XI.2143.1.S1_at	M59455	tfap2a-a	-1.55
XI.21648.1.S1_at	BJ062907	kit-b	-1.55
XI.6195.1.A1_at	AW766846	Predicted LOC100486791	-1.54
XI.9628.1.S1_x_at	BQ398502	ap1ar	-1.54
XI.15267.1.A1_at	BJ053669	gata4-a	-1.54

XI.2478.1.S1_at	BJ088660	gpd1	-1.54
XI.11740.1.A1_at	BJ045587	Predicted Dmx1	-1.54
XI.7235.1.A1_at	BG885754	Predicted hic2	-1.53
XI.6577.1.S1_at	BE506347	LOC100137643	-1.53
XI.5614.1.S1_at	BE026933	meig1	-1.53
XI.14513.1.A1_at	BE491698	dlx3-b	-1.53
XI.2520.1.S1_at	BC041506	eif1	-1.53
XI.15843.1.A1_at	BJ052021	LOC100127337	-1.53
XI.5439.1.A1_at	BE575910	Predicted rtn1-a	-1.53
XI.9703.1.A1_at	BG161470	Predicted slc10a3	-1.53
XI.10349.1.A1_at	BF072039	oaf	-1.52
XI.21813.1.S1_at	BC041253	vtcn1	-1.52
XI.6054.1.A1_at	BG555273	gata2	-1.52
XI.11581.1.S1_at	BG016863	serinc2	-1.52
XI.12043.1.A1_at	BJ092311	LOC100037144	-1.52
XI.10209.1.S1_at	BJ083595	esrra	-1.51
XI.14515.1.S1_at	BC044002	esrp1	-1.51
XI.15772.1.A1_at	BJ083484	Unknown	-1.51
XI.3326.2.S1_a_at	X63427	bmp7.2	-1.51
XI.315.1.S1_at	AF043643	pcdh7	-1.51
XI.21645.1.S1_at	X54678	pou2f3	-1.51
XI.6072.1.A1_at	BG731664	Predicted Hand1	-1.50
XI.13666.1.A1_x_at	BJ091634	slc25a24-a	-1.50
XI.24384.2.S1_at	BI314483	cab39l	-1.50
XI.186.1.S1_at	AF017273	rax-a	-1.50

### 3. Genes upregulated by iFGFR2

Affymetrix Probe Set	Genbank Accession	Gene Symbol	Expression Fold change
XI.736.1.S1_at	AF310008	T-A	23.07
XI.23988.1.S1_at	BJ044287	Rexp44	11.14
XI.4965.1.S1_at	AJ278067	irg1	8.06
XI.16457.1.A1_at	CB563787	junb	6.98
XI.24294.1.S1_at	BJ098811	Cico01 RNA	6.20
XI.8124.1.S1_at	BJ045090	MGC115642	6.04
XI.24793.1.S1_at	CB563927	tspan1	5.52
XI.23897.1.S1_at	BQ386549	cnfn-a	4.76
XI.2213.1.A1_at	BG022871	socs3	4.35
XI.637.1.A1_at	AF250345	egr1-a	4.26
XI.9671.1.S1_at	BC042349	capn8-a	3.92
XI.5082.1.A1_at	BF072347	MGC68521	3.06
XI.4789.1.S1_at	BC043748	MGC52875	2.64
XI.14397.1.S2_at	BC044326	nek6	2.62
XI.24094.1.A1_at	BJ083532	kcnc	2.57
XI.8010.1.A1_at	AW782315	Unknown	2.44
XI.15701.1.S1_at	AY150813	kremen2	2.42
XI.49.1.S1_at	X57234	wnt8a	2.36

XI.24337.1.A1_at	CB564601	Predicted NADPH oxidase 1-like	2.35
XI.24218.1.S1_at	CB943601	dynll1-a	2.30
XI.708.1.S1_at	AF283562	lefty-a	2.25
XI.11964.1.S2_at	AF369901	spry2	2.25
XI.7618.2.A1_a_at	BG346681	plscr1	2.22
XI.558.1.S1_at	AF149307	ventx3.2	2.22
XI.12130.1.S1_at	M24752	hoxa7	2.16
XI.21515.1.S1_at	CB197658	cfos-A	2.15
XI.880.1.S1_at	L07538	wnt3a	2.15
XI.8976.1.S1_at	BG555629	Unknown	2.13
XI.23900.1.A1_at	BF232275	cfos-A	2.11
XI.8315.1.A1_at	AW147996	nuak2	2.09
XI.15623.1.A1_at	CB756273	pkfkb3	2.07
XI.1265.1.S1_at	BG022051	pou2f1	2.07
XI.18073.1.A1_at	BG885063	Unknown	2.07
XI.1082.1.S1_at	S93559	foxa4-b	2.01
XI.23957.1.S1_at	BI313816	cfp	2.00
XI.12011.1.A1_at	BG812177	Predicted fbrsl1	2.00
XI.4880.2.S1_a_at	CB563123	sat1-a	1.97
XI.14334.1.S1_at	BI444166	Predicted phldb1	1.96
XI.6155.1.A1_at	BJ052899	tfap2b	1.96
XI.4725.1.S1_at	AB072909	cldn4L2	1.95
XI.37.1.S1_at	X98454	ventx2.2	1.95
XI.2466.1.A1_at	BG020193	traf4b	1.94
XI.22960.2.A1_at	BJ085655	Unknown	1.93
XI.7252.1.S1_at	AY172320	LOC398520	1.93
XI.1465.1.S1_at	BC046253	ido1	1.89
XI.23938.2.S1_at	CB563963	laptm4a	1.89
XI.23963.1.A1_at	BM181061	LOC100137667	1.88
XI.19778.1.A1_at	BQ398893	bhlhe40	1.87
XI.26209.1.S1_at	BJ056944	Unknown	1.87
XI.1044.1.S1_at	AF283563	lefty-b	1.87
XI.13904.1.A1_at	BJ077369	Predicted LOC100216133	1.85
XI.13019.1.S1_at	BC041294	rasl11b	1.84
XI.16519.1.A1_at	BI446890	hoxd3	1.83
XI.7889.1.A1_at	AW198901	cmip	1.83
XI.11965.1.S1_at	AF331825	LOC398232	1.83
XI.10874.1.A1_at	BU907171	gprc5c	1.83
XI.25397.1.A1_at	BE026782	Unknown	1.81
XI.10415.1.A1_at	BF024850	MGC80142	1.79
XI.25941.1.A1_at	BM180508	tpbg	1.79
XI.5350.1.S1_a_at	BF612893	LOC494681	1.78
XI.12012.2.A1_at	BM191810	btrc	1.77
XI.25587.1.A1_at	BQ387107	tmcc1	1.77
XI.24205.1.S1_at	BJ086130	Predicted itga6-like	1.76
XI.2061.1.A1_at	BI443086	tmem169	1.75
XI.23247.1.S1_at	BQ400459	ddr1	1.74
XI.16396.1.S1_at	BC048018	MGC52622	1.74
XI.4107.1.A1_at	BF025212	Predicted MGC82982	1.73
XI.24824.1.A1_at	BG555750	kng1	1.72
XI.7620.1.S1_at	BC045059	arl5b-b	1.72
XI.801.1.S1_at	BJ056432	gsc-b	1.72
XI.11148.1.A1_at	AW766989	chn2	1.72
XI.15887.1.S1_x_at	BJ088835	mpc2	1.72
XI.21869.1.S1_at	BC045041	LOC398437	1.71
XI.24158.1.A1_at	CB561731	c4orf31	1.70
XI.18712.1.A1_s_at	BI444259	kalm	1.70
XI.19998.1.S1_at	M55054	wnt3	1.70

Xl.10607.1.A1_at	BE491057	Unknown	1.70
Xl.457.1.S1_at	AF065135	gdf3	1.69
Xl.644.1.S1_at	AF223426	foxd5b	1.69
Xl.14206.1.A1_at	BM190943	ripk1	1.68
Xl.15374.1.A1_at	BJ077463	rexp91	1.68
Xl.700.1.S1_at	AF294272	myd88-a	1.67
Xl.7683.1.S1_at	BG816867	fam174	1.66
Xl.19790.1.S1_at	BQ399032	MGC82544	1.66
Xl.3618.1.A1_at	BG020413	cebpb-b	1.66
Xl.1073.1.S1_at	L23542	wnt11b	1.66
Xl.15131.1.A1_at	BJ048969	plekhn1	1.65
Xl.203.1.S1_at	AF012925	ventx1.1-a	1.65
Xl.23718.1.A1_at	BJ075987	mmp14	1.65
Xl.16273.1.A1_at	BJ045836	hoxc8	1.65
Xl.10269.1.S1_at	U02034	cdx4	1.65
Xl.23286.1.S1_at	BC046843	MGC52924	1.65
Xl.24197.1.S1_at	CD327596	scin	1.64
Xl.1002.1.S1_at	U43223	dusp6	1.64
Xl.4819.1.S1_at	AF461119	robo1	1.64
Xl.18364.1.A1_at	BI315001	Predicted ATPase	1.63
Xl.15037.1.S1_at	BJ078533	cxc7	1.63
Xl.15165.1.A1_at	BJ077194	MGC131003	1.63
Xl.4290.1.S1_at	BC046657	ptafr	1.63
Xl.12622.1.A1_at	BJ088645	Predicted aim1l	1.63
Xl.23976.1.A1_at	CB562187	jund	1.62
Xl.23739.1.A1_at	CB564190	cdx1	1.62
Xl.21639.1.S1_at	AJ319749	hoxa10	1.62
Xl.8351.1.A1_s_at	AW148150	prdm1	1.62
Xl.21562.1.S1_at	CB756418	rasd1	1.62
Xl.7556.1.S1_at	AF387815	prickle1-a	1.61
Xl.5940.1.A1_at	AW766423	Predicted KIF26a	1.61
Xl.1358.1.S1_at	U04302	cdx1	1.61
Xl.216.2.S1_a_at	Y08734	mst1	1.60
Xl.9662.1.A1_at	BJ082037	exoc3	1.60
Xl.13054.1.S1_at	AF521098	lin28a	1.60
Xl.13275.1.A1_at	BJ090323	ca14	1.60
Xl.866.1.S2_at	L39213	shh	1.60
Xl.824.1.S1_at	M27063	mix1-a	1.60
Xl.14572.1.A1_at	BJ052988	Predicted LOC100487322	1.60
Xl.9439.1.S1_at	BC041731	hoxa3	1.59
Xl.642.1.S1_at	AF162782	foxd411.1-a	1.59
Xl.6113.1.A1_at	AW767973	LOC100487322	1.59
Xl.24136.1.A1_at	BJ087326	slc43a1	1.58
Xl.15736.1.A1_at	BJ045157	Unknown	1.57
Xl.3713.1.A1_at	BF428167	cir1	1.57
Xl.15811.1.A1_at	BJ049061	txn	1.57
Xl.3294.1.S1_at	AF509738	mark2	1.57
Xl.11438.1.A1_at	BM172656	gxylt1	1.56
Xl.1209.1.S1_at	X12499	hoxc6	1.56
Xl.11188.1.S1_at	AW782515	eps8l3	1.56
Xl.21864.1.S1_at	AY167741	hoxc10	1.55
Xl.9549.1.S1_at	AF374473	lmo2	1.55
Xl.802.1.S1_at	BG016128	LOC397753	1.55
Xl.10766.1.A1_at	BJ053815	kng1	1.55
Xl.16140.1.A1_at	BJ081438	LOC100036879	1.55
Xl.13997.2.A1_at	BJ078256	LOC443660	1.55
Xl.415.1.S1_at	AJ010497	il1b	1.55
Xl.2208.1.A1_at	BJ089383	pkp3	1.54



XI.514.1.S1_at	M77243	t-a	1.54
XI.14496.1.A1_at	BF025525	LOC100191024	1.54
XI.19198.1.S1_at	BQ384108	Predicted LOC100490392	1.54
XI.4792.1.A1_at	BG022307	MGC68744	1.53
XI.15716.1.A1_at	BJ055126	smo	1.53
XI.22636.1.A1_at	BJ087054	pfkfb3	1.53
XI.6868.1.A1_at	BG555913	pvr1	1.53
XI.3085.1.S1_at	BC046743	sgk223	1.53
XI.20394.1.S1_at	BJ089309	myo6	1.53
XI.15284.1.A1_at	BJ080869	akap13	1.53
XI.3540.1.S1_at	AF131890	LOC398134	1.52
XI.16035.1.A1_at	BJ081350	Unknown	1.52
XI.8165.1.S1_at	CB207508	tcf12	1.52
XI.2757.1.S1_at	BC044125	Unknown	1.52
XI.14601.1.A1_at	BJ078962	Unknown	1.51
XI.12606.1.S1_at	BU908750	egln3	1.51
XI.15754.1.A1_at	BG021580	cpne3	1.51
XI.2351.1.A1_at	BJ047002	tfe3	1.50
XI.12180.1.S1_at	AB095361	higd1a-a	1.50
XI.12568.1.A1_at	BJ087957	phf6	1.50
XI.22272.1.A1_at	CD328921	krt16	1.50
XI.23306.1.S1_at	AJ009313	cldn6.1	1.50
XI.16644.1.S1_at	BJ088493	gpx3-b	1.50
XI.18908.1.A1_at	BI447750	stard4	1.50
XI.12111.1.S1_at	AY055473	prickle1-b	1.50

#### 4. Genes downregulated by iFGFR2

Affymetrix Probe Set	GenBank Accession	Gene Symbol	Fold Change
XI.25847.1.S1_at	BC052102	agr2	-6.21
XI.18858.1.A1_at	BI446930	Unknown	-3.50
XI.16543.1.S1_at	AF513854	Ras dva	-3.18
XI.1685.1.S1_at	AF314056	Ag2a	-3.15
XI.24199.1.A1_at	CB756654	Unknown	-3.08
XI.20089.1.S1_at	BC042303	foxi1	-3.03
XI.6266.1.S1_at	AB105372	itln1	-2.98
XI.13575.1.A1_at	BU913085	b3gnt1	-2.85
XI.15702.1.A1_at	BJ076178	ATPase 13a4-like	-2.80
XI.2924.1.S1_at	BG023525	MGC53311	-2.71
XI.26141.1.S1_at	BC044313	MGC80993	-2.52
XI.10583.1.S1_at	BC042234	slc3a2	-2.48
XI.22874.1.A1_at	BJ090165	ubp1	-2.40
XI.11145.1.A1_at	AW766955	Unknown	-2.38
XI.909.1.S1_at	AB018694	xepsin	-2.38
XI.24565.1.A1_at	BG485946	Predicted LOC100049150	-2.34
XI.2659.1.S1_at	BC045045	atp12a-b	-2.30
XI.16262.1.A1_at	BJ051781	Predicted LOC100497338	-2.28
XI.12727.1.A1_at	BM192846	fa2h	-2.26
XI.841.3.S1_a_at	X17545	pdgfa	-2.25
XI.186.1.S1_at	AF017273	rax-a	-2.25
XI.1589.1.S1_at	AF025474	agr3	-2.22
XI.14452.1.A1_at	BG811297	bcat1	-2.22
XI.1683.1.S1_at	BJ044640	MGC68910	-2.21
XI.460.1.S1_at	BG021592	fzd4	-2.17
XI.15475.1.A1_at	BJ083397	U8 snoRNA	-2.16
XI.16466.1.A1_at	BJ082483	Predicted Igalsl-a	-2.14
XI.3789.1.S1_at	BC041714	cebpa	-2.13
XI.104.1.S1_at	AJ005787	pitx2-a	-2.11



XI.10362.1.A1_at	BF072092	ca12	-2.09
XI.11114.1.A1_at	BJ051736	LOC100337617	-2.09
XI.522.1.S1_at	AF217647	pitx1	-2.09
XI.5296.1.A1_at	BJ080084	Unknown	-2.07
XI.5324.1.S1_at	BJ043563	otog	-2.06
XI.628.1.S1_at	M84163	pparg	-2.06
XI.9576.1.S1_at	CB560639	ca2	-2.04
XI.5912.1.A1_at	BG020669	eppk1	-2.03
XI.16589.1.A1_at	BJ080730	Predicted sulfotransferase	-2.01
XI.1616.1.A1_at	BJ083887	fam115a	-2.01
XI.7213.3.S1_a_at	BQ737049	LOC100158288	-2.01
XI.16144.1.A1_at	CB592728	cd81-a	-1.99
XI.1479.1.A1_at	BG038587	Unknown	-1.98
XI.9836.1.A1_at	BJ075817	liph-a	-1.96
XI.5149.1.S1_at	CB592843	sbk1	-1.95
XI.13291.1.A1_at	BJ078773	Predicted Fam55c-like	-1.94
XI.26274.1.A1_at	BE189559	MGC82269	-1.93
XI.24932.1.A1_at	BG811878	Predicted mafb	-1.91
XI.15156.1.S1_at	CB562846	vill	-1.90
XI.2610.1.S1_at	M60768	anxa2-a	-1.88
XI.11136.1.A1_at	BJ045099	LOC100037100	-1.87
XI.16564.1.A1_s_at	BJ054555	fam3a	-1.87
XI.15841.1.A1_at	BJ049236	Predicted slco2b1	-1.87
XI.21868.1.S1_at	BC044973	elf-1	-1.86
XI.1003.1.S1_at	U28370	hesx1-a	-1.86
XI.793.1.A1_at	M76565	gata3	-1.85
XI.6308.1.A1_at	AW632863	mtus1	-1.84
XI.1654.1.A1_at	BG019624	Predicted zfp729-like	-1.84
XI.10855.1.A1_at	BE575595	grhl3	-1.84
XI.7213.1.S1_at	BC042338	cmahp	-1.84
XI.5930.1.A1_at	AW766360	aldh1l1	-1.83
XI.5792.1.A1_at	BJ053583	glo1	-1.81
XI.14532.1.S1_at	BC044975	nr2f2	-1.81
XI.847.1.S1_s_at	L10987	otog	-1.79
XI.7751.1.S1_at	AF536830	menf.1	-1.78
XI.7751.1.S1_at	AF536830	menf.1	-1.78
XI.18395.1.S1_at	BC043753	MGC52890	-1.77
XI.21430.1.S1_at	AF175342	six3	-1.77
XI.509.1.S1_at	BC046269	atp1b2	-1.76
XI.627.1.S1_at	AF183571	six3-a	-1.76
XI.21926.1.S1_at	BC042252	prp5	-1.76
XI.25344.1.A1_at	BE678810	Unknown	-1.76
XI.847.1.S1_at	L10987	otog	-1.75
XI.3804.1.A1_at	BJ080125	rab25	-1.75
XI.21986.1.S1_at	BC043898	tmem188	-1.75
XI.6421.1.A1_at	AW633197	tob1	-1.74
XI.7354.1.A1_at	AW766695	sytl2	-1.74
XI.7354.1.A1_at	AW766695	sytl2	-1.74
XI.4631.1.S1_at	U75681	slbp	-1.74
XI.21983.1.S1_at	M11940	xk81a1	-1.74
XI.16435.1.A1_at	BJ089860	gdpd1	-1.73
XI.16562.1.A1_at	BJ084204	Predicted gmc1	-1.73
XI.24073.1.A1_at	CB561420	fut1	-1.71
XI.15637.1.A1_at	BJ067363	capn9	-1.70
XI.7848.1.A1_at	AW148259	Unknown	-1.70
XI.23835.1.A1_s_at	BJ081027	Predicted prr4	-1.70
XI.13767.1.A1_at	BJ075935	MGC85058	-1.70
XI.16320.1.S1_at	BC046669	anxa9	-1.69

XI.12647.1.A1_at	BJ088990	Predicted paqr5	-1.69
XI.15019.1.A1_at	BQ384301	Predicted sgsm3	-1.68
XI.30.1.S1_at	Z48770	kit-a	-1.68
XI.16658.3.A1_at	BJ083088	kcnc3	-1.67
XI.16672.1.S1_at	BC044108	kitlg	-1.67
XI.2418.1.A1_at	BG023326	ccno	-1.67
XI.25460.1.A1_at	AW148116	grhl1	-1.65
XI.24099.1.A1_at	CB562751	dvl3	-1.65
XI.15893.1.A1_at	BJ056712	Predicted ankyrin repeat domain 43	-1.64
XI.7307.1.S1_at	BJ084368	LOC443682	-1.64
XI.7307.1.S1_at	BJ084368	LOC443682	-1.64
XI.16197.1.A1_at	BJ044460	fezf2	-1.64
XI.26465.1.A1_at	BJ089263	rarres1	-1.64
XI.13893.1.A1_at	BJ083655	fam3d	-1.64
XI.9961.1.A1_at	BG812624	syt1	-1.64
XI.13862.1.A1_at	BJ078064	Predicted LOC100036636	-1.64
XI.973.1.S1_at	L47990	gbx2.2	-1.62
XI.16096.1.A1_at	BJ046407	tmem45b	-1.62
XI.1424.1.S1_a_at	BJ043534	LOC100127277	-1.62
XI.5486.1.A1_at	BJ091754	Unknown	-1.61
XI.2143.1.S1_at	M59455	tfap2a-a	-1.60
XI.647.3.S1_x_at	U67887	pax6-a	-1.60
XI.16436.1.A1_at	BJ081297	Predicted G protein receptor	-1.60
XI.16451.1.A1_at	BJ049278	LOC495248	-1.60
XI.14584.1.A1_at	BJ091208	sbk1	-1.60
XI.10620.1.S1_at	BJ047148	slc1a5	-1.60
XI.633.2.S1_at	AF114151	fzd7	-1.59
XI.647.4.S1_x_at	AF154553	pax6-b	-1.59
XI.4183.2.A1_at	BJ051393	LOC100158277	-1.59
XI.142.1.S1_at	X52691	c-ets-1b	-1.59
XI.10209.1.S1_at	BJ083595	esrra	-1.58
XI.159.1.S1_s_at	X87366	rxrb-a	-1.58
XI.23136.1.S1_at	M77013	ctnnb1-b	-1.57
XI.792.1.S1_at	M76564	gata2	-1.57
XI.25758.1.S1_at	CD325687	MGC68655	-1.57
XI.12819.2.A1_at	BJ056722	gmpr2	-1.57
XI.1351.1.A1_at	BG439162	MGC68638	-1.57
XI.15347.1.A1_at	BJ052833	eps81	-1.57
XI.17667.1.A1_at	BG811141	LOC100037144	-1.57
XI.25518.1.S1_at	BJ053813	LOC443659	-1.57
XI.7078.2.S1_a_at	BJ078851	amd1	-1.57
XI.1604.1.A1_at	BJ057329	MGC78986	-1.56
XI.7756.1.S1_at	BC043737	znf750-a	-1.56
XI.7756.1.S1_at	BC043737	znf750-a	-1.56
XI.17407.1.S1_at	BM928994	sgsm3	-1.56
XI.6104.1.A1_at	BJ076947	rab27a	-1.56
XI.5679.1.S1_at	BE026676	nipal4	-1.55
XI.16518.1.S1_at	BC042248	dbp	-1.55
XI.8098.1.A1_at	BJ090592	LOC494641	-1.55
XI.20084.1.S1_at	BJ056997	Predicted tcp1112	-1.55
XI.21751.1.S1_at	BG347993	LOC100189575	-1.55
XI.16487.1.S1_at	BC045029	ehd4	-1.55
XI.7523.1.S1_at	D82960	LOC398141	-1.54
XI.377.2.S1_a_at	AY130967	nr6a1	-1.54
XI.9247.1.A1_at	BG438869	elf1	-1.54
XI.23267.1.S1_at	BC044298	dnajb14	-1.54
XI.22292.1.S1_at	BG017113	LOC100158394	-1.53
XI.15772.1.A1_at	BJ083484	Unknown	-1.53

XI.8004.1.S1_at	AW199480	Predicted mob1b	-1.53
XI.2256.1.S1_at	BJ056357	hal.1	-1.53
XI.2698.1.S1_at	BC041749	MGC53587	-1.53
XI.15057.1.A1_at	BJ081522	kiaa1324l	-1.53
XI.13666.1.A1_x_at	BJ091634	Predicted slc25a24-a	-1.53
XI.1011.1.S1_at	AB038496	prss27	-1.53
XI.917.1.S1_at	AB034702	crx-a	-1.53
XI.10887.1.A1_at	BM172535	LOC496380	-1.52
XI.21920.1.S1_at	BC043746	slc19a3	-1.51
XI.24391.1.A1_at	BJ092498	Predicted hexokinase -2-like	-1.51
XI.12783.1.A1_at	BJ055803	Unknown	-1.51
XI.2715.1.S1_at	BG555028	atp6v0a4	-1.51
XI.10182.1.A1_at	BJ043711	Predicted LOC100487499 isoform 1	-1.51
XI.21239.1.A1_at	BU908560	EIG121L	-1.50
XI.7919.1.S1_at	BJ090309	MGC81684	-1.50

### 5. Genes upregulated by iGFR3

Affymetrix ProbeSet	Genbank Accession	Gene Symbol	Fold change
XI.736.1.S1_at	AF310008	TA-2	17.95
XI.23988.1.S1_at	BJ044287	Rexp44 mRNA	14.88
XI.4965.1.S1_at	AJ278067	irg1	14.42
XI.8124.1.S1_at	BJ045090	MGC115642	6.96
XI.2213.1.A1_at	BG022871	socs3	5.09
XI.24294.1.S1_at	BJ098811	Cico01 mRNA	4.52
XI.16457.1.A1_at	CB563787	junb	4.35
XI.4290.1.S1_at	BC046657	ptafr	2.88
XI.23897.1.S1_at	BQ386549	cnfn-a	2.22
XI.909.1.S1_at	AB018694	xepsin	2.02
XI.12535.1.A1_at	BJ087645	Predicted clcn6	1.91
XI.16555.1.S1_at	BC043762	rassf6	1.82
XI.8976.1.S1_at	BG555629	Unknown	1.82
XI.20140.1.A1_at	BI315434	Predicted muc19-like	1.75
XI.16035.1.A1_at	BJ081350	Unknown	1.74
XI.1235.1.S1_at	AF085280	ras	1.70
XI.22152.1.S1_at	BJ045156	Predicted angpt4 precursor	1.69
XI.7618.2.A1_a_at	BG346681	plscr1	1.69
XI.12642.1.A1_at	BJ091176	Predicted ChacC	1.68
XI.15811.1.A1_at	BJ049061	txn	1.68
XI.9867.1.A1_at	BG019737	Predicted tiparp	1.68
XI.6841.1.A1_at	BG552502	slc38a5	1.63
XI.2098.1.A1_at	BG486824	LOC503673	1.63
XI.24634.1.A1_at	CB562143	serpina1	1.59
XI.1683.1.S1_at	BJ044640	MGC68910	1.59
XI.13793.1.A1_at	BJ078301	Predicted map1b	1.58
XI.16396.1.S1_at	BC048018	MGC52622	1.57
XI.9671.1.S1_at	BC042349	capn8-a	1.57
XI.22911.1.A1_at	BJ097463	cecr1	1.54
XI.19790.1.S1_at	BQ399032	MGC82544	1.54
XI.9610.1.A1_at	BG038543	ext1	1.53
XI.23587.1.A1_at	BU911087	yap1	1.52
XI.1094.1.S1_at	S69801	fst	1.52
XI.1491.1.S1_at	D50552	sox13	1.52
XI.14402.2.A1_at	BJ092913	flnc	1.51
XI.4440.1.S1_at	BC046372	gpr160	1.50
XI.17323.1.A1_at	BG023008	bag2	1.50
XI.13904.1.A1_at	BJ077369	Predicted LOC100216133	1.50

### 6. Genes downregulated by iFGFR3

Affymetrix ProbeSet	Genbank Accession	Gene Symbol	Fold change
XI.1239.1.S1_at	AW198919	LOC397911	-1.81
XI.4420.1.A1_at	BJ083126	pkp2	-1.78
XI.1616.1.A1_at	BJ083887	fam115a	-1.69
XI.25780.1.S1_at	BG016938	gdf1	-1.67
XI.15645.1.S1_at	BJ061898	dsc3	-1.65
XI.15051.1.A1_at	BJ046997	cinp	-1.63
XI.847.1.S1_at	L10987	otog	-1.62
XI.13379.1.A1_at	BJ085119	Predicted trpm4-like	-1.60
XI.14584.1.A1_at	BJ091208	sbk1	-1.60
XI.7246.1.S1_at	BG438768	Predicted phactr2	-1.59
XI.21734.1.A1_at	BJ077150	thumpd1	-1.59
XI.12041.1.A1_at	BG370976	Predicted zcchc6	-1.59
XI.1082.1.S1_at	S93559	foxa4-b	-1.58
XI.12938.1.A1_at	BJ092938	arhgap1-a	-1.54
XI.16658.2.A1_at	BJ090409	Predicted FAM55D-like	-1.54
XI.6302.1.A1_at	BJ052581	tmbim4	-1.53
XI.7571.1.A1_at	BQ387361	hsph1-a	-1.53
XI.5859.1.S1_at	BJ081030	gamt-b	-1.51
XI.24877.1.S1_at	BG346297	rps6kb1	-1.50
XI.4836.1.A1_s_at	BJ044149	aldh4a1	-1.50

### 7. Genes upregulated by iFGFR4

Affymetrix ProbeSet	GenBank Accession	Symbol	Fold Change
XI.4965.1.S1_at	AJ278067	irg1	2.75
XI.736.1.S1_at	AF310008	LOC398207	2.28
XI.8124.1.S1_at	BJ045090	MGC115642	2.02
XI.9671.1.S1_at	BC042349	capn8-a	2.00
XI.11188.1.S1_at	AW782515	eps8l3	1.93
XI.23988.1.S1_at	BJ044287	Rexp44 mRNA	1.91
XI.24294.1.S1_at	BJ098811	Cico01 mRNA	1.86
XI.4492.1.A1_at	BE678826	atg16l1	1.82
XI.269.1.S2_x_at	AF055980	igf1r	1.76
XI.22272.1.A1_at	CD328921	krt16	1.74
XI.22275.1.S1_at	BJ041762	api5-a	1.71
XI.4290.1.S1_at	BC046657	ptafr	1.71
XI.11602.1.A1_at	AW460740	Predicted slc12a3	1.63
XI.20140.1.A1_at	BI315434	Predicted muc19-like	1.63
XI.4175.2.A1_at	BJ081456	Unknown	1.62
XI.15329.2.A1_at	BJ075982	ndfip2	1.61
XI.7720.1.A1_at	BF615090	Meso05 mRNA	1.61
XI.6097.1.S1_at	BJ091445	Unknown	1.60
XI.16231.1.S1_at	BJ046324	tim10-a	1.59
XI.25982.1.S1_at	BJ056119	Predicted Mov10l1	1.59
XI.10251.1.A1_at	BG811095	Predicted atp2b4	1.58
XI.15070.1.S1_at	BQ386483	MGC80314	1.58
XI.15701.1.S1_at	AY150813	kremen2	1.58
XI.656.1.S1_at	AF163313	cml	1.57

XI.15089.1.A1_at	BJ056659	lgals9	1.57
XI.3661.1.S1_at	U44050	mcm6	1.56
XI.2206.1.S1_at	AF353715	klf17	1.55
XI.4091.1.A1_at	BF025171	fh13	1.55
XI.16035.1.A1_at	BJ081350	Unknown	1.54
XI.16655.1.S1_at	BC047967	slc16a6	1.53
XI.16690.1.A1_at	BM172616	Predicted arhgap12	1.53
XI.18899.1.A1_at	BI447680	Predicted torsin1a interacting protein	1.52
XI.880.1.S1_at	L07538	wnt3a	1.52
XI.6134.1.A1_at	AW782303	Predicted wdr12	1.52
XI.21562.1.S1_at	CB756418	rasd1	1.51
XI.23938.2.S1_at	CB563963	laptm4a	1.50

### 8. Genes downregulated by iFGFR4

Affymetrix ProbeSet	GenBank Accession	Symbol	Fold change
XI.8976.1.S1_at	BG555629	Unknown	-1.89
XI.2659.1.S1_at	BC045045	atp12a-b	-1.83
XI.10583.1.S1_at	BC042234	slc3a2	-1.76
XI.15997.1.A1_at	BJ043804	sytl1	-1.75
XI.3789.1.S1_at	BC041714	cebpa	-1.75
XI.8821.1.A1_at	BJ077414	Predicted psph	-1.74
XI.24572.1.S1_a_at	AB003078	tnnc2	-1.70
XI.5846.1.S1_at	BC046739	MGC53823	-1.68
XI.909.1.S1_at	AB018694	xepsin	-1.65
XI.9990.1.A1_at	BM180289	Predicted Dnase	-1.64
XI.24769.1.S1_at	BQ387765	rpp30	-1.64
XI.19182.1.S1_at	AF368235	hmx3	-1.63
XI.19213.1.S1_at	BU912806	mlst8	-1.62
XI.18146.1.S1_at	CD303354	hmp19	-1.60
XI.25085.1.A1_at	CB756377	rab4b	-1.59
XI.1616.1.A1_at	BJ083887	fam115a	-1.59
XI.9829.1.S1_at	BG018208	foxn4	-1.58
XI.12281.1.S1_at	BG162103	yif1b-a	-1.58
XI.2665.2.A1_at	BM172523	des.1-b	-1.57
XI.19970.1.S1_at	BG409836	extl2	-1.57
XI.17442.1.A1_at	BJ090152	pcolce2	-1.56
XI.12647.1.A1_at	BJ088990	Predicted paqr5	-1.56
XI.26350.1.A1_at	BJ084015	timp2	-1.55
XI.1424.1.S1_a_at	BJ043534	LOC100127277	-1.55
XI.9722.1.S1_a_at	CD326266	Unknown	-1.55
XI.16436.1.A1_at	BJ081297	Predicted G protein receptor	-1.54
XI.22874.1.A1_at	BJ090165	ubp1	-1.52
XI.5232.1.S1_at	AF309688	parn	-1.52
XI.23712.1.S1_at	BG555088	cnfn-b	-1.52
XI.8717.1.S1_at	BC041217	cd81-a	-1.52
XI.5860.1.S1_x_at	CB560495	ckm	-1.51
XI.1165.1.A1_at	X72902	drd2-b	-1.51
XI.9058.1.A1_at	BG347632	mmp3	-1.51

#### 4. Overlapping genes found by the microarray genelist

Overlapping Categories	Number of genes	Genes					
iFGFR1 iFGFR2 iFGFR3 iFGFR4	7	cnfn MGC115642	capn8	fam115a	xepsin	Unknown	irg1
iFGFR1 iFGFR2 iFGFR3	6	plscr1	MGC68910	junb	otog	sbk1	foxa4
iFGFR1 iFGFR2 iFGFR4	7	cd81 G protein receptor	wnt3a	kremen2	slc3a2	LOC100127277	ubp1
iFGFR1 iFGFR3 iFGFR4	2	Rexp44 mRNA	Cico01 mRNA				
iFGFR1 iFGFR2	148	lefty kcnc3 fut1  mst1 tob1 agr3 hoxc10 ido1 LOC496380 xk81a1 ventx2.2  MGC68521 esrra fezf2 dynll1 cdx4 dvl3 loc398404 LOC443659 nek6	syt1 elf1 MGC85058  dnajb14 wnt8a slc1a5 stard4 aldh11 LOC397753 rasl11b LOC100158288  tfap2b hoxd3 LOC100037100 LOC100158277 MGC82269 LOC494641 anxa9 anxa2 wnt11b	pitx2 rab27a sat1  hal.1 hoxc6 foxd4l1.1 tmem45b tcf12 arl5b cldn4L2 znf750  grhl3 liph tspan1 hesx1 MGC52875 capn9 Ras dva six3 fam3a	tfap2a lmo2 foxi1  ehd4 amd1 U8 snoRNA tpbg rax fa2h agr2 hoxa7 hexokinase -2- like MGC78986 LOC100037144 hoxa3 ventx3.2 EIG121L egr1 tmcc1 LOC100487499	pdgfa MGC53311 b3gnt1  gata3 gprc5c cmahp eppk1 atp1b2 eps8l1 LOC398232 gdpd1  pou2f1 tmem169 kiaa1324l sgsm3 bcat1 hoxc8 MGC81684 LOC100337617 MGC131003	nipal4 gdf3 fam3d ankyrin repeat domain 43 c4orf31 sulfotransferase gbx2.2 nuak2 pfkfb3 pitx1 slc19a3  prp5 kit mmp14 sytl2 dusp6 MGC80142 foxd5b crx kitlg

		slc25a24 cdx1 mtus1 vill LOC398134	spry2 gmpr2 LOC495248 rab25 itln1	ca12 cfd fzd4 LOC443682 MGC80993	zfp729 glo1 traf4b gata2 t	LOC398437 MGC68655 ca2 menf.1 hoxa10	grhl1 mpc2 ccno
iFGFR1 iFGFR3	4	sox13	TA-2	LOC503673	gdf1		
iFGFR1 iFGFR4	9	atp12a tnnc2	MGC53823 cml	mmp3 des.1	foxn4	lgals9	sytl1
iFGFR2 iFGFR3	5	ptafr	MGC82544	MGC52622	socs3	txn	
iFGFR2 iFGFR4	5	laptm4a	krt16	eps8l3	rasd1	paqr5	
iFGFR3 iFGFR4	1	muc19-like					
iFGFR1	207	gsto1 ptch2  pcdh7 arg1  fucolectin krt12 ada tbcx capns2 cetn4 tmem69 sfrp2  xmc LOC100037197 LOC100137681 meig1 lrrfp1 hes5.2 slc35d1 hk2	wnt5a hes6.1  krt prp4 ATPase subunit C2- like MGC84721 hic2 trpm4 gcm1 c1orf21 FAM55D-like foxi4.2  tmpo nkx3-1 cab39l smyd1 slc16a3 serinc2 slc35a3.2 vtcn1	pacsin2 ap1ar NEDD4-binding protein 3-like cldn1  tmem38b hes7.1 insm1 synt2 tspan4 schip1 nexn cdh2  Possible ATPase scel fgfr4 LOC100494996 MGC53924 LOC733321 LOC446305 tsga10	pou2f3 dapk2  atp2a2 pnp  c-ets-1 cat1 fam154b MGC81163 LOC100485898 nek8 ATPase 13A4 LOC100497155  elovl7 ag1 mreg vangl1 camk2g cdc42se2 klhl24 kctd15	nkx2 LOC100137643  LOC398688 dio3  bmp7.2 creb3 LOC398260 hyal2 krt8.2 xenf elf-1 rassf10  LOC100486791 hs3st1 hspa5 oaf Possible syt1 tfcp2l1 loxl1	nek2 paqr5  prtg uaca  cdh1 fhn3 ankrd9 tnfrsf1a gnb3 LOC494785 LOC100036671 lectin  MGC80468 hexd itgb1bp2 glb1l2 envoplakin tdgf1 tmem63A Dmx1



		MGC52585 rnft1 anion transporter LOC101734664 tesc fbxl22 st3Gal-II angpt4 wnt5b LOC100101281 MGC82342 add1 osbpl2 HoxB8	LOC100036828 Rexp91 mRNA gpd1 FAM55c-like efhd1 gata4 greb1 fam46a hoxb7 msx2 MGC53335 HoxD9 fut6 adenovirus homolog	LOC100127337 phldb1 b3gnt2 MGC81002 LOC101730357 tbc1d24.2 fam181a dlgap4 rtn1 MGC84886 tmem181 c8orf42 ppl gna14	hist1h2am gmppa st3gal4 mylpf eif1 MGC82906 LOC419301 degs3 MGC115605 Plekho1 dlx6 cldn4 pcdh10 esrp1	LOC100127290 sox7 fbln1 Meso05 mRNA gmppb elavl3 Hand1 atf5.2 dlx3 slc10a3 thbs4 LOC496316 crabp2 rnf149 MGC81939	Lapl04 mRNA rnasek superoxide dismutase LOC100158300 mucin-4-like gs17 MGC80262 hapln3 myl1 itga6-like gale pmm2 LOC398485 serpind1
iFGFR2	107	MGC68638 LOC100487322 wnt3 scin ATPase mix1 shh LOC100036636 KIF26a MGC52924 exoc3 Ag2a pkp3 ventx1.1 ripk1  ctnnb1 lgalsl	LOC100497338 mafb itga6-like elf cfos sgsm3 Rexp44 mob1b plekhn1 LOC100049150 sgk223 myd8a rxrb slco2b1  cir1 kng1	prickle1 atp6v0a4 il1b prss27 LOC100158394 robo1 ATPase 13a4-like c-ets kalrn slc43a1 cmip LOC100137667 tcp11l2 smo jund  fzd7 atp12	mark2 cxcr7 Fam55c-like bhlhe40 pparg egln3 phf6 LOC100490392 aim1l cebpa lin28a MGC52890 MGC53587 LOC100216133 nr2f2  myo6 LOC100487322	LOC443660 cldn6.1 higd1a cebpb prdm1 ddr1 LOC100036879 pax6 akap13 LOC494681 btrc MGC82982 cpne3 LOC100189575 nr6a1 NADPH oxidase 1-like fam174	pvr1 gxylt1 LOC398520 tmem188 gpx3 T-A slbp Cico01 RNA phldb1 LOC398141 dbp gmc1 rarres1 ca14 tfe3  kcnc LOC100191024

		prp4	chn2	gsc	MGC68744		
iFGFR3	32	bag2 serpina1 cecr1 rassf6 thumpd1 slc38a5	ext1 zcchc6 ras tmbim4 tiparp clcn6	cinp phactr2 hsph1 map1b ChacC	dsc3 fst trpm4-like FAM55D-like LOC100216133	gpr160 rps6kb1 arhgap1 aldh4a1 yap1	pkp2 LOC397911 gamt flnc angpt4 precursor
iFGFR4	32	arhgap12 parn ndfip2 slc16a6 hmp19 wdr12	atp2b4 klf17 pcolce2 mcm6 Meso05Unknown timp2	ckm extl2 rab4b hmx3 api5	fh13 ptafrslc12a3 cebapasp LOC398207 timm10	Mov10l1 Dnase MGC80314 yif1b mlst8	igf1r torsin1a interacting drd2 atg16l1 rpp30

## RNA-seq Genelists

**10 - Genes upregulated by iFGFR1**

Gene	Aligns to Source	Genome region	Expression fold-change
FOXA4-B	c.Audic201207_X015973	JGIv7b.000031469_915655-919113+	166.5
FOXA4-B	c.Audic201207_X039184	JGIv7b.000146669_874246-877928+	55.0
POU4F2	c.Ueno201210eye_X001811	JGIv7b.000190940_1080000-1081640+	52.2
FOXA4-B	c.Chang2013_X041679	NIGv2.S00005482_210567-214238+	38.6
FOXA4-B	c.Taira201203st10_X004823	NIGv2.S00000078_44808-48405-	35.9
RNF223	c.Taira201203stomach_X001511	JGIv7b.000066475_1968499-1980832-	26.1
LEFTY-A	c.Audic201207_X034596	JGIv7b.000102974_5086276-5089519+	19.6
IKZF2	c.Ueno201210st09_X000205	JGIv7b.000020345_864299-915085+	15.5
WNT11B	c.Ueno2012102cells_X000642	JGIv7b.000036991_2149589-2155869-	14.1
UNNAMED	c.Chang2013_X002796	JGIv7b.000006875_1358573-1361506-	13.2
DUSP5	c.Quigley201212_X053012	NIGv2.S00001026_193780-200530+	10.6
OSBP	c.Audic201207_X051959	NIGv2.S00000286_2585-35077+	10.3
DUSP5	c.Quigley201212_X012016	JGIv7b.000020641_3538718-3561871+	9.5
OSBP	c.TXGP201107_X010742	NIGv2.S00004591_55245-117384-	9.3
NOT-B	c.Quigley201212_X014344	JGIv7b.000027038_130890-133875+	8.1
SPRY1	c.Taira201203brain_X010913	JGIv7b.000088765_2315207-2319681-	8.0
UNNAMED	c.JGIL6RMv1_XeXenL6RMv10032768m	JGIv7b.000007555_1690594-1693172-	7.4
FOXD4L1.2	c.Chang2013_X037957	JGIv7b.000399468_514029-542898+	7.3
SPRY1	c.Taira201203st09_X003581	JGIv7b.000112610_1220629-1226046-	7.2
OSBP	c.TXGP201107_X001151	JGIv7b.000011978_364042-432008+	6.3
FOXB1	c.UniGene_XI_S13590653	JGIv7b.000038656_1756198-1758738+	6.0
ADMP	c.Chang2013_X033894	JGIv7b.000231526_1125358-1130767+	5.7
UNNAMED	c.JGIL6RMv1_XeXenL6RMv10025054m	JGIv7b.000016863_2508114-2510618-	5.6
FOXD4L1.1-B	c.Chang2013_X039478	NIGv2.S00001139_12051-40798-	5.2
ZXDC	c.Taira201203eye_X014955	NIGv2.S00000309_453048-462189-	5.2
SGK1	c.Taira201203ovary_X003188	JGIv7b.000039762_1596878-1614597+	5.0

SPRY1	c.Amin201106_X029644	NIGv2.S00003069_305986-309195+	5.0
LOC398207	c.Quigley201212_X033084	JGIv7b.000094770_18082-25147+	4.8
ARL5C	c.XGI_TC426294	JGIv7b.000075417_5337831-5344493-	4.7
BTG2	c.Quigley201112_X018699	JGIv7b.000196263_531292-533437-	4.5
SP5L	c.JGIL6RMv1_XeXenL6RMv10032472m	JGIv7b.000070222_2093542-2103684-	4.3
FOXD4L1.1-B	c.Chang2013_X038965	NIGv2.S00000686_540615-542511+	4.2
TSPAN1	c.Audic201207_X053503	NIGv2.S00001300_769342-778120+	4.1
ZSWIM4	c.Ueno2012102cells_X001536	JGIv7b.000175714_234323-250264-	4.1
UNNAMED	c.JGIL6RMv1_XeXenL6RMv10028945m	JGIv7b.000043061_1325599-1369985-	4.1
FOXD4L1.1-B	c.TeperekTkacz201202_X000807	JGIv7b.000106782_2609184-2610932+	4.0
SP5L	c.Chang2013_X040311	NIGv2.S00002193_122413-127676-	4.0
ZSWIM4	c.TXGP201107_X010004	NIGv2.S00000140_899537-914561+	3.9
TSPAN1	c.Ueno201210kidney_X001802	JGIv7b.000276272_818115-826957+	3.9
CTDSPL	c.Park201106_X027161	NIGv2.S00002443_4948-59702-	3.9
UNNAMED	c.Quigley201212_X053887	NIGv2.S00001621_263619-264537+	3.4
ID3	c.Quigley201212_X005126	JGIv7b.000008834_232083-234500+	3.4
PNP	c.TXGP201107_X009208	JGIv7b.000272406_956346-987977+	3.4
PGBD4	c.Park201106_X015855	JGIv7b.000086967_109082-110412-	3.4
Oct-25	c.Chang2013_X013645	JGIv7b.000037448_494399-508009-	3.3
ID3	c.Audic201207_X054642	NIGv2.S00002590_1022943-1025176-	3.3
MT-ND4L	c.JGIL6RMv1_XeXenL6RMv10002896m	JGIv7b.000323037_5513-7181-	3.3
PFKFB3	c.Taira201203st10_X000227	JGIv7b.000005375_625926-678516+	3.2
PTGS2	c.Ueno201210heart_X000956	JGIv7b.000078978_2024793-2032785+	3.1
QSER1	c.Quigley201212_X028931	JGIv7b.000074352_3829490-3877907-	3.1
IRG1	c.Quigley201212_X033094	JGIv7b.000094770_38299-44565+	3.0
ZFP36L1	c.Audic201207_X044942	JGIv7b.000218195_958966-963341-	3.0
FOXD4L1.1-A	c.TeperekTkacz201202_X001018	JGIv7b.000214452_17356-19212-	2.9
SETD2	c.JGIL6RMv1_XeXenL6RMv10033051m	NIGv2.S00003637_448414-489792-	2.9
MRRF	c.TeperekTkacz201206_X001915	JGIv7b.000034020_290760-308242+	2.9
CDC25B-B	c.Quigley201212_X056550	NIGv2.S00007616_27531-41541+	2.9

HES1	c.JGIL6RMv1_XeXenL6RMv1003 8392m	JGIv7b.000030987_2402981- 2405745+	2.9
RHOB	c.Ueno201210heart_X001032	JGIv7b.000093416_2010188- 2013106-	2.9
ZNF629	c.Taira201203st15_X003375	NIGv2.S00007957_89052-92237+	2.8
CDC25B-A	c.Quigley201212_X014347	JGIv7b.000027038_1265417- 1280660+	2.8
PRICKLE1	c.Taira201203egg_X004705	JGIv7b.000074339_3116341- 3188715-	2.8
IYD	c.Taira201203intestine_X003967	JGIv7b.000045784_2485606- 2494783-	2.8
ATP6V0C	c.JGIL6RMv1_XeXenL6RMv1002 3514m	JGIv7b.000023403_4435801- 4447982-	2.8
POU3F4	c.Quigley201212_X020593	JGIv7b.000046073_2406892- 2453403+	2.8
ATP6V0C	c.XGI_TC428762	JGIv7b.000000939_2385517- 2396566+	2.8
PDGFB	c.Quigley201212_X030415	JGIv7b.000078978_3403347- 3420648-	2.8
UNNAMED	c.Quigley201112_X001095	JGIv7b.000003737_943400- 944398-	2.7
CDC25B-A	c.Chang2013_X041859	NIGv2.S00006568_301501- 315199-	2.7
RNF111	c.Quigley201212_X026207	JGIv7b.000059883_2180394- 2206352-	2.7
ZFP36L2	c.Chang2013_X026204	JGIv7b.000102974_14415127- 14420720-	2.7
CDC25B-B	c.Taira201203kidney_X000277	JGIv7b.000001168_7102490- 7117401-	2.7
ARID1A	c.Quigley201212_X054694	NIGv2.S00002590_373032- 409653+	2.7
YAP1	c.UniGene_XI_S55517271	JGIv7b.000006290_698493- 749864-	2.7
PTGS2	c.Chang2013_X004724	JGIv7b.000012423_153604- 158988+	2.7
TPBG	c.Chang2013_X029203	JGIv7b.000146478_52070-57563+	2.7
RPS3	c.JGIL6RMv1_XeXenL6RMv1000 3218m	JGIv7b.000050406_350421- 370416-	2.6
GPX4	c.Park201106_X000258	JGIv7a.000101906_462283- 468199-	2.6
MT-ND4L	c.Taira201203st12_X004364	NIGv2.C23850859_23-547-	2.6
ID3	c.JGIL6RMv1_XeXenL6RMv1002 7462m	JGIv7b.000098463_1551637- 1553867+	2.6
UNNAMED	c.mgEST_1013119916	JGIv7b.000217632_684-2008-	2.6
MXI1	c.TeperekTkacz201206_X001389	JGIv7b.000020641_3480668- 3492185+	2.6
MCMBP	c.Amin201106_X029621	NIGv2.S00002933_255375- 265588-	2.6
PRICKLE1	c.TeperekTkacz201206_X005778	NIGv2.S00000920_346369- 410352-	2.6
ZFP36L2	c.Park201106_X000114	JGIv7a.000017698_6247896- 6253061+	2.6
IRG1	c.Quigley201212_X017592	JGIv7b.000036991_115611- 122292+	2.6
FOXK2	c.Quigley201212_X040617	JGIv7b.000163806_1576700- 1627098+	2.6

SALL1	c.TeperekTkacz201205_X000856	JGIv7b.000033905_3475663-3497233+	2.6
SALL1	c.Taira201203st12_X000017	JGIv7a.000020499_457812-477054-	2.5
POU3F4	c.Ismailoglu201203_X013681	NIGv2.S00002741_201687-204164-	2.5
TRIB2	c.Audic201207_X032976	JGIv7b.000093416_6021763-6045133-	2.5
WHSC2	c.Quigley201212_X013424	JGIv7b.000024597_1013702-1040465+	2.5
ARID1A	c.TeperekTkacz201205_X002770	NIGv2.S00003103_73152-114642-	2.5
TRIB2	c.Quigley201212_X052794	NIGv2.S00000877_812197-831768+	2.5
UBQLN4	c.Chang2013_X010550	JGIv7b.000026819_2815543-2828727-	2.5
UNNAMED	c.Ueno201210st12_X000337	JGIv7b.000074488_887627-894002-	2.5
UNNAMED	c.Taira201203kidney_X001646	JGIv7b.000011405_882624-883983+	2.5
PTCH2-A	c.Ueno201210eye_X002089	JGIv7b.000325141_1096289-1126340+	2.5
PFKFB3	c.TXGP201107_X010066	NIGv2.S00000358_1134122-1185330-	2.4
OCT -91	c.Chang2013_X041872	NIGv2.S00006607_4387-6427-	2.4
LMBRD2	c.Quigley201212_X023738	JGIv7b.000052441_5992095-6032818+	2.4
UNNAMED	c.JGIL6RMv1_XeXenL6RMv10055418m	JGIv7b.000075417_5512673-5517835-	2.4
UNNAMED	c.Quigley201212_X054313	NIGv2.S00002168_444164-451470-	2.4
FOXK2	c.Ueno201210ovary_X000574	JGIv7b.000081941_2354504-2409334-	2.4
UNNAMED	c.Taira201203heart_X000951	JGIv7b.000009528_1568246-1645156+	2.4
ZFP36L1	c.Amin201106_X000043	JGIv7a.000007480_2568139-2572061+	2.4
DROSHA	c.Taira201203kidney_X006805	JGIv7b.000057180_2913941-3058638-	2.4
CDC25B-A	c.Quigley201212_X000025	JGIv7a.000001422_178764-193695-	2.4
SETD2	c.Taira201203lung_X005942	JGIv7b.000093635_127204-196840-	2.4
FAM126B	c.Ueno201210brain_X000726	JGIv7b.000020345_3471385-3524477+	2.3
ZNF629	c.Quigley201212_X012224	JGIv7b.000021594_76004-85345+	2.3
ZNF214	c.Taira201203eye_X016061	NIGv2.S00005637_215-139265+	2.3
WASL	c.Chang2013_X038876	NIGv2.S00000626_1526987-1553647+	2.3
STC1	c.Chang2013_X017838	JGIv7b.000051940_55199-70780+	2.3
DYNLL1-A	c.Audic201207_X053894	NIGv2.S00001717_2085-4762-	2.3
PTGS2	c.Taira201203heart_X008958	NIGv2.S00003847_492214-500068-	2.3
UBE2D4	c.Quigley201212_X053841	NIGv2.S00001574_1842836-1872122+	2.3

POU3F1	c.Chang2013_X039997	NIGv2.S00001621_254496-259081+	2.3
SALL1	c.XenBase_288557289	JGIv7b.000062355_1954057-1972637-	2.3
CHMP2A	c.Taira201203brain_X018099	NIGv2.S00002094_484041-490219+	2.3
CECR2	c.Quigley201212_X047146	JGIv7b.000256647_18135-55076-	2.3
UBE2D4	c.Taira201203eye_X008955	JGIv7b.000079772_2548726-2578727-	2.2
HNRNPA3	c.TeperekTkacz201205_X002644	NIGv2.S00000679_862568-872644+	2.2
RBM42	c.Chang2013_X038988	NIGv2.S00000706_368369-371959-	2.2
BCL9	c.TeperekTkacz201206_X005606	JGIv7b.000399803_808628-838129-	2.2
UNNAMED	c.Audic201207_X054489	NIGv2.S00002441_336305-342411-	2.2
AGMAT	c.Taira201203kidney_X014957	NIGv2.S00003382_450731-459756+	2.2
ARL5C	c.Chang2013_X038725	NIGv2.S00000489_675387-681630+	2.2
CCDC160	c.Audic201207_X023147	JGIv7b.000050694_3511837-3515127+	2.2
GPRC5C	c.Taira201203kidney_X013921	JGIv7b.000402746_1345231-1435883+	2.2
ABHD15	c.Taira201203spleen_X004590	JGIv7b.000159212_666954-684116+	2.2
CPEB2	c.Taira201203st08_X004644	JGIv7b.000146311_365282-431607-	2.2
IRX1-B	c.Quigley201112_X010920	JGIv7b.000057559_47731-52835-	2.2
UNNAMED	c.Chang2013_X039319	NIGv2.S00001027_700145-703404+	2.1
CA14	c.Audic201207_X014301	JGIv7b.000026819_2924131-2951166+	2.1
BCL9	c.Quigley201207_X014499	NIGv2.S00001884_287121-355017+	2.1
HNRNPA3	c.TeperekTkacz201205_X000128	JGIv7b.000004321_5551952-5566829+	2.1
FBXL14	c.Quigley201112_X020416	JGIv7b.000256647_503440-506888+	2.1
UBE2R2	c.XGI_TC417190	JGIv7b.000169907_3865426-3906484+	2.1
CREBBP	c.JGIL6RMv1_XeXenL6RMv10023475m	JGIv7b.000023403_3292998-3363583+	2.1
C17ORF63	c.Taira201203st25_X001759	JGIv7b.000036364_3720727-3756693+	2.1
CAMSAP3	c.Taira201203brain_X013982	JGIv7b.000171831_601330-667752-	2.1
C5ORF15	c.Audic201207_X011351	JGIv7b.000017836_1329301-1339385-	2.1
HNRNPA0	c.Quigley201207_X013950	NIGv2.S00000330_13760-15405-	2.1
CDC42EP4	c.Audic201207_X000266	JGIv7a.000026529_1333726-1360157-	2.1
YY1	c.Amin201106_X024715	JGIv7b.000236382_2132334-2149616+	2.1
FCN2	c.Taira201203kidney_X014553	NIGv2.S00001621_95963-100340+	2.1



NOTCH3	c.Park201106_X020666	JGIv7b.000171831_680633-733284-	2.1
ETV3	c.Chang2013_X014268	JGIv7b.000041091_3521026-3559652-	2.1
ACOX2	c.Taira201203kidney_X014616	NIGv2.S00001847_315152-339798+	2.1
CD276	c.Audic201207_X052520	NIGv2.S00000663_600421-616317+	2.1
DUSP1	c.UniGene_XI_S13684642	JGIv7b.000018892_2914011-2917473+	2.1
GPRC5C	c.Chang2013_X038888	NIGv2.S00000640_561649-567108-	2.1
MORN2	c.Chung201110_X007967	NIGv2.S00000914_155417-172394+	2.1
UNNAMED	c.Taira201203kidney_X009137	JGIv7b.000096766_2029557-2056237+	2.1
CRABP2	c.Taira201203egg_X003235	JGIv7b.000041091_3369386-3397974-	2.1
GPRC5C	c.Taira201203eye_X001105	JGIv7b.000007000_226270-261891-	2.1
SMURF2	c.Quigley201212_X040698	JGIv7b.000163806_4832005-4880003+	2.0
CST3	c.Chang2013_X041680	NIGv2.S00005502_148526-151519-	2.0
HP1BP3	c.Quigley201112_X018100	JGIv7b.000175822_18694-28456-	2.0
DYRK1B	c.Taira201203stomach_X001921	JGIv7b.000102557_219148-291907+	2.0
SCARB1	c.Audic201207_X041239	JGIv7b.000167265_511996-570757-	2.0
U2AF1	c.XGI_TC433037	JGIv7b.000347078_702320-709716+	2.0
A2M	c.Ueno201210st10_X000172	JGIv7b.000007045_3704835-3735363+	2.0
ZFP36L2	c.Audic201207_X056261	NIGv2.S00006220_476386-483244-	2.0
KHSRP	c.Taira201203st10_X003887	JGIv7b.000175714_268495-294223-	2.0
UNNAMED	c.Park201106_X017595	JGIv7b.000108888_148572-173804-	2.0

#### 11 – Genes downregulated by iGFR1

Gene	Aligns to source:	Genome region	Expression Fold Change
KRT12	c.XenBase_27696404	JGIv7b.000013265_3627584-3644405-	-5.3
SLC12A3	c.TeperekTkacz201206_X004009	JGIv7b.000111469_66839-126398+	-5.0
KRT12	c.Ueno201210st35_X000016	NIGv2.S00002938_233278-248330+	-4.1
UNNAMED	c.Amin201106_X030176	NIGv2.S00005487_252198-256251+	-3.6
HESX1	c.JGIL6RMv1_XeXenL6RMv10033507m	JGIv7b.000033876_482101-485204-	-3.2
SNRPG	c.Quigley201212_X009407	JGIv7b.000014870_225925-236198+	-3.2

HESX1	c.UniGene_XI_S13589749	JGlv7b.000177844_181426-184575+	-3.0
IFT172	c.Chung201110_X004537	JGlv7b.000080529_201168-231666+	-3.0
SNRPG	c.Park201106_X000036	JGlv7a.000004727_1139488-1149380-	-3.0
RBM8A	c.JGIL6RMv1_XeXenL6RMv10010425m	JGlv7b.000012462_1741658-1749852+	-2.9
HIST2H3A	c.TeperekTkacz201206_X004021	JGlv7b.000111824_6775263-6775607-	-2.8
SNAI1	c.Amin201106_X029102	NIGv2.S00001881_253774-258061+	-2.8
PITPNB	c.Ismailoglu201203_X007692	JGlv7b.000077809_130001-140882+	-2.8
PTMA-A	c.Chang2013_X036024	JGlv7b.000293841_203483-206579+	-2.7
RBM8A	c.Ueno2012106cells_X002372	NIGv2.S00002347_114737-122574+	-2.7
PPIL1	c.Chang2013_X039410	NIGv2.S00001083_232616-234252-	-2.6
APPL1	c.TeperekTkacz201206_X006093	NIGv2.S00005117_156006-183356-	-2.6
OTUD6B	c.JGIL6RMv1_XeXenL6RMv10002547m	JGlv7b.000022127_9001-21497-	-2.6
TOMM22	c.Chang2013_X040115	NIGv2.S00001865_407984-416289-	-2.5
BRP44L-B	c.Quigley201212_X056153	NIGv2.S00005505_85388-93227-	-2.4
SRP9	c.Amin201106_X027761	NIGv2.S00000024_616994-624649-	-2.4
CWC25	c.Quigley201112_X000013	JGlv7a.000002298_2384297-2389993+	-2.4
HES3.1	c.Chang2013_X039721	NIGv2.S00001322_2013951-2016303+	-2.4
HNRPDL	c.Taira201203st15_X003372	NIGv2.S00007280_40360-44246-	-2.4
FAM55D	c.Ismailoglu201203_X007979	JGlv7b.000083106_152889-161353+	-2.4
SGK494	c.Taira201203eye_X014978	NIGv2.S00000348_1620323-1629386+	-2.4
RPL21	c.Chung201110_X008136	NIGv2.S00001639_1135756-1148673-	-2.4
TPMT	c.Audic201207_X053187	NIGv2.S00001069_373040-389472+	-2.3
RUVBL2	c.Ueno201210egg_X001199	JGlv7b.000139674_1787321-1801883+	-2.3
SNRPD1	c.Park201106_X000689	JGlv7b.000001268_1685868-1701614+	-2.3
RP5-977B1.10	c.Chang2013_X000311	JGlv7a.000078524_134965-145190+	-2.3
HNRPDL	c.JGIL6RMv1_XeXenL6RMv10001854m	JGlv7b.000058878_2007697-2013395-	-2.3
PPIE	c.Ueno2012102cells_X001385	JGlv7b.000137317_2000210-2012876-	-2.3
HNRNPH1-B	c.Taira201203egg_X006162	JGlv7b.000140825_732234-746785+	-2.3
UNNAMED	c.Chang2013_X033037	JGlv7b.000208071_3368708-3370756+	-2.3
LDLRAP1	c.Taira201203st30_X000390	JGlv7b.000007281_1927654-1960093+	-2.3

UNNAMED	c.Taira201203st08_X004257	JGlv7b.000107078_130406-136077+	-2.3
UNNAMED	c.Quigley201207_X005310	JGlv7b.000044780_1693944-1705020-	-2.2
UNNAMED	c.Amin201106_X019023	JGlv7b.000107347_2455544-2457909+	-2.2
RPL39	c.Quigley201212_X010080	JGlv7b.000015436_5400196-5404618-	-2.2
SFSWAP	c.Taira201203intestine_X010650	NIGv2.S00002754_425243-477210+	-2.2
RPL27A	c.Chang2013_X035887	JGlv7b.000287959_146467-152199+	-2.2
MAK16	c.Taira201203st15_X002384	JGlv7b.000139741_1526513-1535556-	-2.2
NDUFB6	c.Quigley201207_X008771	JGlv7b.000090041_5628093-5636504+	-2.2
TUBA1A-B	c.mgEST_1013155827	JGlv7b.000127187_1109513-1113080-	-2.2
COQ4	c.Quigley201212_X017740	JGlv7b.000037448_288099-305397+	-2.2
EIF4A3	c.Taira201203brain_X010967	JGlv7b.000089475_854935-897659-	-2.2
MFAP1	c.Chang2013_X000051	JGlv7a.000006679_5732770-5737706+	-2.2
LSM5	c.Audic201207_X052452	NIGv2.S00000589_73817-77516-	-2.2
C12ORF45	c.Ueno201210lung_X000864	NIGv2.S00004457_446048-448697+	-2.2
CALCOCO1	c.Ueno201210st10_X002327	NIGv2.S00002501_240558-264815-	-2.2
TAF15	c.Taira201203stomach_X000009	JGlv7a.000016056_140368-162530+	-2.2
MDK	c.Chung201110_X002883	JGlv7b.000043483_642289-654640+	-2.2
C12ORF45	c.JGIL6RMv1_XeXenL6RMv1003 1748m	JGlv7b.000005925_7519623-7523842-	-2.2
C11ORF58	c.Chang2013_X038804	NIGv2.S00000548_1067259-1071887+	-2.2
HNRNP11-A	c.Audic201207_X022527	JGlv7b.000048253_786566-811362+	-2.2
ARG1	c.Audic201207_X034514	JGlv7b.000102277_1378103-1389903+	-2.1
ELOF1	c.Chang2013_X000313	JGlv7a.000080025_251822-256641+	-2.1
PKDCC.2	c.Ueno201210brain_X000869	JGlv7b.000027036_317796-362184+	-2.1
MED9	c.Chang2013_X022043	JGlv7b.000074548_369381-382000+	-2.1
EIF3J	c.Chang2013_X000153	JGlv7a.000021089_540005-554241+	-2.1
ANP32A	c.Quigley201212_X053357	NIGv2.S00001260_397655-405232-	-2.1
RBM8A-A	c.Quigley201112_X023672	NIGv2.S00002567_199325-202713-	-2.1
ZNF622	c.Quigley201112_X002635	JGlv7b.000011199_1862234-1874227+	-2.1
C8ORF40	c.Amin201106_X030036	NIGv2.S00004598_559626-561767-	-2.1
MRT04	c.Chung201110_X000032	JGlv7a.000015632_608771-614811-	-2.1

PCBD1	c.Chang2013_X040476	NIGv2.S00002416_67527-72806-	-2.1
RPF1	c.Quigley201207_X003442	JGlv7b.000024235_1715202-1728596-	-2.1
PPAN-B	c.XGI_TC418418	NIGv2.S00006828_10160-16544-	-2.1
H2AFJ	c.Ueno2012104cells_X000824	JGlv7b.000074488_1286181-1298618-	-2.1
HNRNPA1	c.Taira201203brain_X012675	JGlv7b.000133382_406820-446822+	-2.1
PITPNB	c.Taira201203egg_X008593	NIGv2.S00003591_314926-322588-	-2.1
GLRX	c.JGIL6RMv1_XeXenL6RMv10028215m	JGlv7b.000001187_4083787-4090024+	-2.1
EIF3B	c.Quigley201112_X000049	JGlv7a.000009276_5375917-5396714+	-2.1
CWC25	c.Quigley201212_X029323	JGlv7b.000075417_5226786-5234433+	-2.1
H3F3A	c.Quigley201212_X051737	NIGv2.S0000233_993527-1000649+	-2.0
SNRNP70	c.Amin201106_X027634	JGlv7b.000402746_367479-382531+	-2.0
MRPS26	c.Quigley201212_X000390	JGlv7a.000106822_27276-34467+	-2.0
HNRNPH1-A	c.Chang2013_X000118	JGlv7a.000015444_3784852-3807177-	-2.0
RPL10A	c.Chang2013_X000250	JGlv7a.000044917_626050-629118+	-2.0
HNRNPK	c.Taira201203st40_X001865	JGlv7b.000086871_2140635-2152595-	-2.0
POLR2L.1	c.XGI_TC463547	JGlv7b.000021980_2317439-2319477-	-2.0
SRP9	c.XenBase_148234311	JGlv7b.000102974_5193679-5201441-	-2.0
GNB3	c.Quigley201112_X013169	JGlv7b.000079772_5446576-5470832+	-2.0
HNRNPH1-B	c.Amin201106_X029551	NIGv2.S00002804_295426-309569+	-2.0
RBM8A-A	c.Taira201203ovary_X003602	JGlv7b.000046492_203712-208136-	-2.0
UNNAMED	c.Ismailoglu201203_X004665	JGlv7b.000039723_1326545-1329420+	-2.0
CIRBP	c.mgEST_1013253913	JGlv7b.000039437_1967966-1973134+	-2.0
ANP32A	c.Taira201203liver_X003877	JGlv7b.000325448_345352-354714-	-2.0
HIST2H2AB	c.Chang2013_X035898	JGlv7b.000287959_3113019-3114049+	-2.0
PIN4	c.Chang2013_X013351	JGlv7b.000036864_3174743-3179026+	-2.0

## 12. Genes upregulated by iFGFR4

Gene	Aligns to Source:	Genome region	Expression fold-change
ACLY	c.Park201106_X026788	NIGv2.S00001609_104960-120704-	44.4
PADI2	c.JGIL6RMv1_XeXenL6RMv10033213m	JGlv7b.000036295_1002602-1023208-	30.5

MRRF	c.TeperekTkacz201206_X001915	JGlv7b.000034020_290760-308242+	28.6
HBA1	c.Taira201203st30_X003398	JGlv7b.000120240_1982132-1984062-	24.5
MARK4	c.Taira201203st40_X003255	NIGv2.S00005156_627695-652388+	24.4
SLC25A4	c.Taira201203eye_X000097	JGlv7a.000093157_254175-263290-	22.7
PADI2	c.TeperekTkacz201206_X006137	NIGv2.S00007245_10058-34147+	15.7
APPL1	c.TeperekTkacz201206_X006093	NIGv2.S00005117_156006-183356-	15.0
CARHSP1	c.Amin201106_X007374	JGlv7b.000026364_2644128-2662567+	9.7
IFT172	c.Chung201110_X004537	JGlv7b.000080529_201168-231666+	6.5
NQO1	c.Chung201110_X008036	NIGv2.S00001215_83829-96206+	6.4
CCNB2	c.Taira201203kidney_X007160	JGlv7b.000059883_2168606-2176741-	6.3
MCMBP	c.Amin201106_X029621	NIGv2.S00002933_255375-265588-	6.1
SLC12A3	c.TeperekTkacz201206_X004009	JGlv7b.000111469_66839-126398+	5.4
CXORF56	c.Amin201106_X019451	JGlv7b.000118020_87186-102070-	5.2
LOC73372 8	c.TXGP201107_X003099	JGlv7b.000032511_552358-563142-	5.1
KRTCAP3	c.Amin201106_X030131	NIGv2.S00005238_57878-69325-	5.0
FBXW4	c.Chang2013_X020551	JGlv7b.000061874_688873-700001+	4.7
FRS3	c.Chang2013_X004216	JGlv7b.000009994_1616589-1623506-	4.6
NEDD9	c.Taira201203heart_X003857	JGlv7b.000051654_212436-234627-	4.3
C13ORF15	c.Taira201203st08_X000568	JGlv7b.000008129_2441581-2449217+	4.1
SPSB1	c.Quigley201212_X027248	JGlv7b.000065414_48610-67548+	4.0
OAT.1	c.Audic201207_X000047	JGlv7a.000002565_100020-118676-	3.9
SPSB1	c.Audic201207_X052320	NIGv2.S00000512_649491-655558+	3.9
IER5	c.Chung201110_X003630	JGlv7b.000053263_7195493-7198049-	3.7
DDX39B	c.Amin201106_X015857	JGlv7b.000075398_1100580-1119081-	3.7
SSX2IP	c.XenBase_148236336	JGlv7b.000024235_1612955-1629872+	3.7
PATL2	c.TXGP201107_X000601	JGlv7b.000006590_5196886-5213377+	3.6
MORN2	c.Chung201110_X007967	NIGv2.S00000914_155417-172394+	3.5
NDUFB8	c.Chung201110_X008334	NIGv2.S00003140_77811-90953-	3.5
MOB3C	c.Taira201203lung_X009424	NIGv2.S00003101_168454-201204+	3.5
SGK1	c.Taira201203ovary_X003188	JGlv7b.000039762_1596878-1614597+	3.4

KLF2	c.Taira201203brain_X016955	JGIv7b.000357348_471871-475471+	3.4
SPSB1	c.Audic201207_X027549	JGIv7b.000066475_149868-163046-	3.4
PDIA3	c.Amin201106_X018916	JGIv7b.000106789_1878320-1894173-	3.3
MAP1LC3C	c.Chang2013_X020661	JGIv7b.000062432_227149-233998+	3.3
CDCA7	c.Chang2013_X016983	JGIv7b.000049342_728889-735053+	3.3
RGS4	c.Taira201203st10_X003952	JGIv7b.000181903_1352400-1362094-	3.3
VANGL2	c.Ueno201210egg_X000601	JGIv7b.000041091_1975712-2030640+	3.3
CCNA1	c.Quigley201212_X032721	JGIv7b.000091366_786361-795530+	3.2
RNF138	c.Chang2013_X040712	NIGv2.S00002774_218588-228009-	3.2
SPSB1	c.Taira201203eye_X015081	NIGv2.S00000590_555846-572578+	3.2
CCNA1	c.Chang2013_X041006	NIGv2.S00003350_160219-168990-	3.2
SIVA1	c.Ueno201210st10_X001965	JGIv7b.000230550_3128515-3134801-	3.2
OCT -60	c.Ueno201210st12_X000686	NIGv2.S00001621_270628-275793+	3.2
SPDYC-B	c.mgEST_1013156593	JGIv7b.000021980_494497-503231+	3.2
CSNK1E	c.Audic201207_X054625	NIGv2.S00002589_664278-702891-	3.1
DNAJB14	c.Park201106_X027317	NIGv2.S00002782_21805-25261+	3.1
CCNB1	c.JGIL6RMv1_XeXenL6RMv10035758m	JGIv7b.000220499_647410-652378+	3.1
TNFSF11	c.XGI_TC415993	JGIv7b.000014978_7271141-7282475-	3.1
TBP	c.TXGP201107_X008044	JGIv7b.000176005_572262-579759+	3.1
BTG4	c.UniGene_XI_S14220550	JGIv7b.000078584_1125227-1127450-	3.1
CCNA1	c.Quigley201212_X043616	JGIv7b.000200825_2199521-2208550+	3.1
REEP6	c.Quigley201212_X056701	NIGv2.S00013727_36150-40518+	3.0
ROMO1	c.Ueno2012106cells_X002149	JGIv7b.000345631_1246002-1250872-	3.0
CCNB1	c.Ueno201210st09_X001000	NIGv2.S00000097_65075-71908-	2.9
UNNAMED	c.TXGP201107_X003472	JGIv7b.000037448_486318-492292-	2.9
CITED2	c.TXGP201107_X010770	NIGv2.S00004977_24236-61333+	2.9
TBP	c.Quigley201207_X000083	JGIv7a.000042089_691873-698741+	2.9
HES2	c.Audic201207_X032154	JGIv7b.000087017_2605107-2607011+	2.8
CITED2	c.Taira201203lung_X000289	JGIv7b.000003036_19-1129-	2.8
PIM3	c.Audic201207_X035421	JGIv7b.000108224_328683-333812+	2.8

CSNK1E	c.Chang2013_X036098	JGIv7b.000298452_154108-193571+	2.8
ZAR1L	c.JGIL6RMv1_XeXenL6RMv10004417m	JGIv7b.000200825_707245-714280-	2.8
CCNB1	c.Chang2013_X038225	NIGv2.S00000078_323966-328917+	2.8
TMEM57	c.XenBase_59897110	JGIv7b.000167390_21266-49398-	2.7
UAP1	c.TeperekTkacz201206_X004667	JGIv7b.000181903_1492576-1508305-	2.7
RAD51D	c.XGI_TC421430	JGIv7b.000267344_1554170-1569938+	2.7
RAD51D	c.XGI_TC421430	JGIv7b.000267344_1554170-1569938+ 12.084153834 33.0660071545 2.73631133869	2.7
SSX2IP	c.XGI_TC451478	JGIv7b.000137809_1838967-1862244+	2.7
BTG4	c.TeperekTkacz201202_X000972	JGIv7b.000185843_399483-403249+	2.7
RASSF3	c.Taira201203spleen_X000358	JGIv7b.000005925_322282-417925+	2.7
RBM24	c.Ueno2012104cells_X001046	JGIv7b.000121479_384572-396468+	2.7
CHMP1A	c.Quigley201212_X051081	JGIv7b.000398601_1827-11556-	2.7
TTC5	c.Taira201203lung_X008868	NIGv2.S00000077_302823-308542-	2.7
ZFAND2A	c.TXGP201107_X001035	JGIv7b.000009994_8681051-8693755+	2.7
PARP15	c.Chang2013_X009064	JGIv7b.000021603_9359704-9378965+	2.7
HBA1	c.Ueno201210st12_X000456	JGIv7b.000120240_2034592-2044970-	2.6
SNX10	c.Taira201203st30_X005028	NIGv2.S00001230_520830-547530-	2.6
CCNB1	c.Chang2013_X012184	JGIv7b.000032212_7697205-7707852-	2.6
CDC6	c.Ueno2012104cells_X000219	JGIv7b.000013265_3282967-3290855+	2.6
UNNAMED	c.Audic201207_X036836	JGIv7b.000125077_505053-507505+	2.6
COX5B.2	c.mgEST_1013088791	JGIv7b.000203187_564639-572568-	2.6
RIF1	c.UniGene_XI_S60885257	JGIv7b.000004321_14052078-14104109-	2.6
DUSP22	c.UniGene_XI_S20755577	JGIv7b.000274508_1364859-1406902-	2.6
ALDH2	c.JGIL6RMv1_XeXenL6RMv10040201m	JGIv7b.000025254_4358924-4376397+	2.6
DEPDC4	c.Audic201207_X034415	JGIv7b.000100253_3192496-3205330-	2.6
LMBRD2	c.Quigley201212_X023738	JGIv7b.000052441_5992095-6032818+	2.5
UNNAMED	c.Taira201203st30_X000113	JGIv7b.000002049_1721239-1731925+	2.5
PRDX2	c.XGI_TC424241	JGIv7b.000055171_763313-771105+	2.5
FABP4	c.JGIL6RMv1_XeXenL6RMv10008819m	JGIv7b.000022127_2889081-2897540+	2.5



SNX7	c.mgEST_1013252172	JGlv7b.000120545_1486943-1527899+	2.5
ZNF337	c.Audic201207_X007264	JGlv7b.000012518_12603145-12616647+	2.5
PRRG1	c.Chang2013_X038519	NIGv2.S00000267_3222777-3223253+	2.5
DNAJB14	c.TXGP201107_X007950	JGlv7b.000167800_1369425-1389999-	2.5
CHP	c.XenBase_148236366	JGlv7b.000218195_2686150-2715482-	2.5
TMEM57	c.Ismailoglu201203_X005490	JGlv7b.000047026_493573-514727-	2.5
KLF2	c.Chang2013_X039910	NIGv2.S00001484_754702-758538-	2.5
ZNF300	c.Chang2013_X033558	JGlv7b.000223728_307126-310308+	2.5
CCDC117	c.Taira201203brain_X018472	NIGv2.S00003454_691237-706773+	2.5
TMEM57	c.Audic201207_X054695	NIGv2.S00002689_26954-54534-	2.4
TBP	c.Taira201203st08_X002042	JGlv7b.000034423_1781424-1792792-	2.4
PPARG	c.XenBase_38014780	NIGv2.S00003119_62006-98298-	2.4
RPAIN	c.TeperekTkacz201202_X001249	NIGv2.S00003419_1262929-1270325+	2.4
GREM1			2.4
GOLT1B	c.Amin201106_X030315	NIGv2.S00006552_75449-79526+	2.4
SNX10	c.Taira201203st10_X003152	JGlv7b.000099185_1240707-1267816-	2.4
C13ORF15	c.Taira201203st25_X000983	JGlv7b.000014978_7745525-7756248-	2.4
UNNAMED	c.Taira201203pancreas_X002475	JGlv7b.000162663_1276806-1284586+	2.4
LIPH	c.Taira201203st08_X000725	JGlv7b.000011316_5046664-5065548+	2.4
CEBPG	c.TXGP201107_X010361	NIGv2.S00001380_2151666-2162683+	2.4
KIAA0355	c.Ueno201210egg_X001390	JGlv7b.000249035_666016-723445+	2.4
TRIM14	c.UniGene_Xl_S14219872	JGlv7b.000074488_1497018-1508582+	2.4
CHMP7	c.Taira201203kidney_X003977	JGlv7b.000027313_20393-37906-	2.4
UNNAMED	c.Taira201203brain_X018123	NIGv2.S00002155_161459-164407-	2.4
GREM1	c.JGIL6RMv1_XeXenL6RMv1004425	JGlv7b.000049557_885127-888412+	2.4
ISM1	c.TeperekTkacz201202_X000139	JGlv7b.000012020_11809576-11845053-	2.4
DNAJC9	c.Amin201106_X030268	NIGv2.S00006259_56963-63694+	2.4
PIF1	c.JGIL6RMv1_XeXenL6RMv10012147m	JGlv7b.000067962_340725-351534-	2.4
TCEB2	c.Quigley201212_X053864	NIGv2.S00001595_114830-125047-	2.4
SLC18A2	c.XGI_TC416830	JGlv7b.000139741_1438356-1457109+	2.4



CCDC68	c.Chang2013_X041113	NIGv2.S00003474_201149-229730-	2.4
ACO2	c.Amin201106_X028808	NIGv2.S00001315_1221169-1241309+	2.4
PIM3	c.Quigley201212_X039731	JGlv7b.000152894_17327-23415-	2.3
FAM177A1	c.Amin201106_X025404	JGlv7b.000265107_1397790-1409483+	2.3
RNF8	c.Ueno2012102cells_X001705	JGlv7b.000256136_750566-757156+	2.3
C19ORF44	c.Ismailoglu201203_X008318	JGlv7b.000090041_5824040-5831999-	2.3
DEPDC7	c.Quigley201112_X010106	JGlv7b.000051988_3615759-3631460+	2.3
GRAMD3	c.Taira201203eye_X014301	JGlv7b.000332973_1550687-1602350-	2.3
B9D2	c.Park201106_X008957	JGlv7b.000039723_9341434-9344109+	2.3
C14ORF109	c.Taira201203liver_X001461	JGlv7b.000039723_1331764-1339011-	2.3
BFAR	c.XGI_TC418564	JGlv7b.000009994_1592416-1612079-	2.3
GPKOW	c.Audic201207_X050801	JGlv7b.000376231_14717-32725-	2.3
ATL2	c.Quigley201212_X042339	JGlv7b.000180104_1646919-1703974-	2.3
CRY1	c.Quigley201212_X021579	JGlv7b.000047533_5654488-5685330+	2.3
INSM1	c.Taira201203egg_X002601	JGlv7b.000030080_2706723-2717676+	2.3
BIRC5.1-B	c.Taira201203egg_X005990	JGlv7b.000133382_1856472-1859157-	2.3
RAB21	c.UniGene_XI_S17527732	JGlv7b.000237412_340926-357427-	2.3
ARHGAP18	c.XGI_TC418315	NIGv2.S00001447_15280-42007+	2.3
EIF1	c.Quigley201112_X023222	NIGv2.S00001391_645171-647700-	2.3
UAP1	c.Quigley201112_X009177	JGlv7b.000047606_5100243-5116395+	2.3
TMEM18	c.Ismailoglu201203_X004936	JGlv7b.000043061_3754291-3763196-	2.3
RAB21	c.Ismailoglu201203_X006060	JGlv7b.000052352_755272-767082-	2.3
FRYL	c.Taira201203st25_X001618	JGlv7b.000032657_3068830-3109249+	2.3
CBR4	c.Audic201207_X009465	JGlv7b.000014692_9422410-9434441+	2.3
BTG3	c.Amin201106_X021722	JGlv7b.000160841_3387681-3397411-	2.2
GNG4	c.Chang2013_X010110	JGlv7b.000025196_277278-317283-	2.2
GLMN	c.JGIL6RMv1_XeXenL6RMv10052860m	JGlv7b.000047606_1433123-1455862+	2.2
FAM46B	c.TXGP201107_X006481	JGlv7b.000098463_2092359-2103126+	2.2
GMNN	c.TeperekTkacz201206_X005993	NIGv2.S00002971_104644-109494+	2.2
ZNF300	c.Taira201203st09_X000193	JGlv7b.000003552_26743-27560+	2.2

FGFR1OP2	c.JGIL6RMv1_XeXenL6RMv1005070 8m	JGIv7b.000070246_641260- 647706+	2.2
CSNK1E	c.Taira201203spleen_X003556	JGIv7b.000078978_5282261- 5322709-	2.2
GRAMD3	c.Taira201203intestine_X001420	JGIv7b.000012879_564891- 614168-	2.2
CHMP1A	c.JGIL6RMv1_XeXenL6RMv1004108 3m	JGIv7b.000010177_1212605- 1225218-	2.2
CPSF4	c.Quigley201112_X022573	NIGv2.S00000371_1893358- 1895886+	2.2
MCM3	c.TXGP201107_X008579	JGIv7b.000215439_1169508- 1191663-	2.2
NOG	c.Taira201203brain_X011562	JGIv7b.000100342_354700- 356309-	2.2
UNNAMED	c.UniGene_XI_S16181855	JGIv7b.000019169_2231616- 2244772-	2.2
FOPNL	c.Chang2013_X004220	JGIv7b.000009994_3745254- 3755831-	2.2
TMEM111. 2	c.Amin201106_X029979	NIGv2.S00004312_858139- 867545+	2.2
VLDLR	c.Taira201203st10_X003481	JGIv7b.000126823_118223- 160899-	2.2
KLHL13	c.Taira201203brain_X007899	JGIv7b.000050694_2072891- 2126536-	2.2
ATAT1	c.XenBase_58403331	JGIv7b.000134683_522316- 556855-	2.2
PPP1CC	c.XGI_TC418335	JGIv7b.000025254_4420786- 4434616+	2.2
CDC42SE2	c.XGI_TC417024	JGIv7b.000009266_3529229- 3550496+	2.2
NDE1	c.Chang2013_X004144	JGIv7b.000009994_3653920- 3668883+	2.2
RNF4	c.Taira201203st12_X000133	JGIv7b.000003467_4873946- 4889341+	2.2
MTHFR	c.Taira201203eye_X010382	JGIv7b.000108577_106317- 135153-	2.2
STX5	c.Amin201106_X008522	JGIv7b.000031469_1757845- 1766955-	2.2
ACOT13	c.JGIL6RMv1_XeXenL6RMv1005520 9m	JGIv7b.000133814_5456-9847-	2.2
TAPT1	c.Taira201203eye_X011378	JGIv7b.000146311_74713-112330+	2.2
LYRM4	c.XGI_TC429950	JGIv7b.000046891_32557-76294-	2.2
UNNAMED	c.XGI_TC453389	JGIv7b.000035086_72578-76147-	2.2
PAPD5	c.Taira201203st35_X001255	JGIv7b.000032008_585419- 611266-	2.2
ATG4C	c.XGI_TC462242	JGIv7b.000053042_1264305- 1295314-	2.2
TMEM169	c.Chang2013_X009003	JGIv7b.000021603_174379- 181180+	2.2
ITPKC	c.XenBase_83405250	JGIv7b.000039723_10057060- 10086843-	2.1
H1FOO	c.Chang2013_X034727	JGIv7b.000255257_262055- 268442+	2.1
ARL2BP	c.XGI_TC422181	JGIv7b.000033905_3874492- 3885266+	2.1
ALDOC	c.Audic201207_X017272	JGIv7b.000035361_3467404- 3489396+	2.1

UNNAMED	c.UniGene_XI_S25791702	JGIv7b.000121300_9688-33844+	2.1
SNX10	c.Taira201203intestine_X008223	JGIv7b.000178713_2590081-2614536-	2.1
SIRT3.2	c.Audic201207_X038188	JGIv7b.000137507_1853230-1871470+	2.1
ZNF3	c.Taira201203brain_X004212	JGIv7b.000021594_704284-724209-	2.1
UNNAMED	c.Chang2013_X005252	JGIv7b.000012518_8294195-8344255-	2.1
POC5	c.Chung201110_X008544	NIGv2.S00010585_13475-41861+	2.1
UAP1	c.Quigley201212_X052410	NIGv2.S00000616_398194-413500+	2.1
DEPDC7	c.Quigley201212_X054003	NIGv2.S00001820_537040-557127+	2.1
ACO2	c.Ueno201210heart_X000191	JGIv7b.000012423_2228713-2250819+	2.1
ATG4C	c.Audic201207_X040002	JGIv7b.000155039_2303502-2327858-	2.1
LARP6	c.Taira201203egg_X002993	JGIv7b.000035716_1966319-1980564-	2.1
PPHLN1	c.Quigley201212_X015301	JGIv7b.000030470_3514159-3571642+	2.1
RBM24	c.XGI_TC417312	JGIv7b.000011136_1872989-1883587-	2.1
GMNN	c.Park201106_X027971	NIGv2.S00005321_186838-188634-	2.1
PGK1	c.Amin201106_X029109	NIGv2.S00001898_865570-878802-	2.1
GADD45G	c.XGI_TC421689	JGIv7b.000013576_4101295-4106965+	2.1
ACO2	c.Audic201207_X030539	JGIv7b.000078978_2882729-2904948-	2.1
HN1	c.Chang2013_X038637	NIGv2.S00000387_384492-395632-	2.1
GNG10	c.JGIL6RMv1_XeXenL6RMv10011906m	JGIv7b.000046215_225505-236096-	2.1
KIAA0889	c.Taira201203ovary_X004788	JGIv7b.000071264_2754863-2817946+	2.1
NOG	c.Taira201203st35_X003850	JGIv7b.000373158_1037035-1038876+	2.1
SOX13	c.XenBase_46250057	JGIv7b.000151578_693353-785783+	2.1
PDCD10	c.Amin201106_X014159	JGIv7b.000058517_2924605-2943478-	2.1
PMCH	c.Taira201203brain_X003935	JGIv7b.000018184_5024474-5028781+	2.1
HRAS	c.TeperekTkacz201205_X001797	JGIv7b.000109526_292263-327631-	2.1
ZNF484	c.Taira201203egg_X006088	JGIv7b.000136952_451253-459505+	2.1
DAPK1	c.Taira201203ovary_X001492	JGIv7b.000013576_1230568-1320502+	2.1
OAT.2	c.Taira201203st10_X000021	JGIv7a.000044240_1462478-1479658-	2.1
ZNF639	c.TXGP201107_X005028	JGIv7b.000058994_8462-20315-	2.1
UNNAMED	c.TXGP201107_X008557	JGIv7b.000215439_1104601-1114534+	2.0

FBXO33	c.Park201106_X026736	NIGv2.S00001505_834374-839817+	2.0
DUS4L	c.Chang2013_X005142	JGIv7b.000012518_18164100-18178627+	2.0
STK35	c.Chang2013_X002898	JGIv7b.000007103_501795-506878+	2.0
ATP5D	c.mgEST_1013092630	JGIv7b.000054274_2468502-2472064-	2.0
NDNL2	c.UniGene_Xl_S18078526	JGIv7b.000043648_238817-252510-	2.0
PDCD10	c.Amin201106_X025510	JGIv7b.000267727_608606-626750-	2.0
CLNS1A	c.JGIL6RMv1_XeXenL6RMv10021444m	JGIv7b.000262000_1405538-1422069+	2.0
DEPDC7	c.Taira201203intestine_X005584	JGIv7b.000074352_3780008-3800394-	2.0
GALNT4	c.XenBase_147907289	JGIv7b.000070754_794116-833753+	2.0
LSM2	c.Amin201106_X029261	NIGv2.S00002210_37773-42176-	2.0
CEP85	c.Quigley201207_X015122	NIGv2.S00007337_87917-102249+	2.0
SERTAD2	c.Chang2013_X033798	JGIv7b.000230550_2789873-2797899+	2.0
RND3	c.JGIL6RMv1_XeXenL6RMv10031510m	JGIv7b.000021603_9128421-9149344-	2.0
SLC20A1	c.Ismailoglu201203_X013158	NIGv2.S00000368_64191-89886-	2.0
CBX2	c.Taira201203st40_X002816	JGIv7b.000277717_37187-63068-	2.0
NUAK2	c.Taira201203eye_X015865	NIGv2.S00003839_264501-280393-	2.0
SOD1	c.Taira201203st08_X000445	JGIv7b.000006590_6414248-6420299-	2.0
BRF1	c.Ismailoglu201203_X011522	JGIv7b.000230550_1877377-2079767+	2.0
DEPDC7	c.Quigley201212_X051448	NIGv2.S00000039_416616-434040+	2.0
HIST1H2A A	c.Taira201203egg_X001018	JGIv7b.000011316_6703374-6705050+	2.0

### 13 Genes downregulated by iFGFR4

Gene	Aligns to Source	Genome region	Expression fold-change
ZEB2	c.XenBase_8925961	JGIv7b.000004321_15892491-15987808+	-18.1
ZEB2	c.Taira201203st20_X001347	JGIv7b.000021603_7456409-7551033-	-12.3
LRAT	c.Audic201207_X055311	NIGv2.S00003495_975186-978844+	-12.2
ZEB2	c.Taira201203kidney_X015323	NIGv2.S00007837_359787-458457+	-11.8
ZEB2	c.Taira201203kidney_X000017	JGIv7a.000002575_192291-280172-	-10.3
CRABP2	c.Audic201207_X014291	JGIv7b.000026819_2001871-2017104+	-9.6

AG1-A	c.JGIL6RMv1_XeXenL6RMv100 37629m	JGIv7b.000013787_5208193- 5215643+	-8.7
HESX1	c.UniGene_XI_S13589749	JGIv7b.000177844_181426- 184575+	-7.8
DYNLL1-A	c.Audic201207_X053894	NIGv2.S00001717_2085-4762-	-5.3
MDK	c.Chung201110_X002883	JGIv7b.000043483_642289- 654640+	-5.3
CYP26A1	c.Chang2013_X000110	JGIv7a.000014833_4137270- 4142183-	-5.0
HESX1	c.JGIL6RMv1_XeXenL6RMv100 33507m	JGIv7b.000033876_482101-485204-	-4.8
PKDCC.2	c.Ueno201210brain_X000869	JGIv7b.000027036_317796- 362184+	-4.8
PKDCC.2	c.Taira201203heart_X005631	JGIv7b.000102974_12998538- 13043290+	-4.8
NR6A1	c.Quigley201112_X000037	JGIv7a.000006263_166884- 203776+	-4.7
CYP26A1	c.Chang2013_X039292	NIGv2.S00000972_68480-73355+	-4.4
ARG1	c.Audic201207_X034514	JGIv7b.000102277_1378103- 1389903+	-4.4
CFH	c.Taira201203kidney_X012568	JGIv7b.000245044_3631680- 3797857-	-3.9
ADAP1	c.Audic201207_X030499	JGIv7b.000078978_5596836- 5611399+	-3.9
CYP26A1	c.Ueno201210st10_X000202	JGIv7b.000007555_3441089- 3446814+	-3.9
NR6A1	c.Quigley201112_X023210	NIGv2.S00001385_714320-753033+	-3.9
TRIM29	c.Park201106_X024075	JGIv7b.000287959_1656133- 1688834+	-3.7
SLC27A3	c.JGIL6RMv1_XeXenL6RMv100 13872m	JGIv7b.000012462_395592-403463-	-3.6
ROR2	c.Ismailoglu201203_X002182	JGIv7b.000013576_4963312- 5056011-	-3.5
STK40	c.JGIL6RMv1_XeXenL6RMv100 15764m	JGIv7b.000298574_255290- 267525+	-3.4
KRT5.7	c.Taira201203brain_X001103	JGIv7b.000005732_8831252- 8837956-	-3.4
EPPK1	c.Quigley201212_X055874	NIGv2.S00004660_124529-182304-	-3.3
CNN2	c.XGI_TC417055	JGIv7b.000054274_2567215- 2577179-	-3.3
TRIM29	c.Quigley201212_X054528	NIGv2.S00002350_692632-725034+	-3.3
CNN1	c.XenBase_3746796	JGIv7b.000039437_1774002- 1782237+	-3.2
FAM55D	c.Ismailoglu201203_X007979	JGIv7b.000083106_152889- 161353+	-3.2
LRP2	c.Taira201203testis_X001603	JGIv7b.000043242_7285838- 7393816+	-3.2
SLC30A8	c.Quigley201112_X017380	JGIv7b.000160942_6433391- 6462650+	-3.2
UNNAMED	c.Chang2013_X041601	NIGv2.S00004944_40501-42272+	-3.2
GNB3	c.Quigley201112_X013169	JGIv7b.000079772_5446576- 5470832+	-3.1
KIAA1324L	c.Quigley201212_X006996	JGIv7b.000012518_13813324- 13861697+	-3.0

LIN28A	c.Taira201203st25_X000544	JGIv7b.000008834_1121721-1140275-	-3.0
C9	c.Ismailoglu201203_X006109	JGIv7b.000052441_7261050-7287758-	-3.0
STARD13	c.Quigley201212_X015911	JGIv7b.000031941_1881436-1986425-	-3.0
VIM	c.Quigley201212_X026614	JGIv7b.000061124_70427-86528-	-3.0
CYP26A1	c.TeperekTkacz201205_X002611	NIGv2.S00000318_464174-468759+	-2.9
VIM	c.Ismailoglu201203_X013259	NIGv2.S00000766_2696693-2712306+	-2.9
KIT	c.Taira201203lung_X002929	JGIv7b.000032657_1734171-1767962-	-2.9
CYP26A1	c.TeperekTkacz201205_X000583	JGIv7b.000016863_3785916-3790655+	-2.9
OTX2	c.TeperekTkacz201205_X001673	JGIv7b.000091950_411122-418355+	-2.9
UPK3B	c.Audic201207_X047411	JGIv7b.000267344_1483270-1496800-	-2.9
UNNAMED	c.Ismailoglu201203_X003280	JGIv7b.000024242_674171-676991+	-2.9
SHROOM3	c.Taira201203heart_X004291	JGIv7b.000058878_5502767-5562019-	-2.8
CNN1	c.Quigley201207_X013837	NIGv2.S00000082_492864-496632-	-2.8
NR6A1	c.Ueno201210st20_X000895	NIGv2.S00000673_537275-562026+	-2.8
ANGPTL3	c.Quigley201212_X052087	NIGv2.S00000405_1066114-1069514-	-2.8
CKAP4	c.Quigley201112_X023483	NIGv2.S00002102_42661-49168+	-2.8
UNNAMED	c.Taira201203heart_X003470	JGIv7b.000045784_2039071-2113089-	-2.7
UNNAMED	c.Quigley201212_X038159	JGIv7b.000135348_2833347-2910067-	-2.7
MCM6.2-B	c.UniGene_XI_S13831231	JGIv7b.000060608_495830-515096-	-2.7
WFDC2	c.Quigley201112_X018501	JGIv7b.000187321_2337925-2345788-	-2.7
CKAP4	c.Quigley201112_X016484	JGIv7b.000137507_2068193-2074541-	-2.7
MDK	c.Quigley201112_X012346	JGIv7b.000074352_63252-76586-	-2.7
CYGB	c.Quigley201212_X036348	JGIv7b.000120240_2355646-2375117-	-2.6
RAX	c.Audic201207_X024176	JGIv7b.000052441_2501506-2508259-	-2.6
PARP3	c.Audic201207_X043154	JGIv7b.000187321_2587167-2606620-	-2.6
PKDCC.1	c.JGIL6RMv1_XeXenL6RMv10013085m	JGIv7b.000236382_1908213-1923883+	-2.6
CCND1	c.XGI_TC422275	JGIv7b.000074352_1060302-1076913+	-2.6
ZNF608	c.Taira201203brain_X002631	JGIv7b.000012879_1052370-1165078+	-2.6
KIAA1324L	c.JGIL6RMv1_XeXenL6RMv10009242m	JGIv7b.000033104_5396372-5433588-	-2.6
ZNF740	c.XenBase_147898513	JGIv7b.000005732_8194028-8209521-	-2.5

AEN	c.Amin201106_X010297	JGIv7b.000041091_3468250-3477129-	-2.5
UNNAMED	c.Taira201203st40_X003121	NIGv2.S00001319_1814861-1822370+	-2.5
LRAT	c.Taira201203eye_X007824	JGIv7b.000061741_1096071-1112641-	-2.5
DYNLL1	c.JGIL6RMv1_XeXenL6RMv10042969m	JGIv7b.000113816_508606-510472-	-2.5
DYNLL1	c.Quigley201212_X055124	NIGv2.S00003088_565653-567846-	-2.5
OTX2	c.Quigley201212_X051915	NIGv2.S00000300_108072-110208-	-2.5
CYP2J2	c.Chang2013_X033182	JGIv7b.000214452_9948-52224+	-2.4
UNNAMED	c.Amin201106_X025116	JGIv7b.000248633_526499-535913+	-2.4
ANGPTL3	c.mgEST_1013086036	JGIv7b.000005925_1870076-1909413+	-2.4
LIN28A	c.Park201106_X028195	NIGv2.S00007337_220421-239976+	-2.4
ORAOV1	c.Taira201203st12_X001800	JGIv7b.000043483_1012989-1020298-	-2.4
XEPSIN	c.Taira201203skin_X003526	JGIv7b.000132609_46128-75477+	-2.4
LIG3	c.JGIL6RMv1_XeXenL6RMv10001947m	JGIv7b.000150750_933402-974025-	-2.4
TPSG1	c.Audic201207_X037375	JGIv7b.000132609_88539-98813+	-2.4
PTRH2	c.Chung201110_X001263	JGIv7b.000013523_3440014-3450087-	-2.4
CNN2	c.Quigley201212_X052425	NIGv2.S00000641_860802-870505-	-2.4
AGR2	c.Chung201110_X003762	JGIv7b.000057216_2613250-2620205-	-2.3
FXYD3	c.mgEST_1013251433	JGIv7b.000019916_18854-33417+	-2.3
CKAP4	c.JGIL6RMv1_XeXenL6RMv10016662m	JGIv7b.000054336_24051-31518-	-2.3
PARP1	c.Ismailoglu201203_X008816	JGIv7b.000102974_4858525-4894562+	-2.3
FAM55B	c.Park201106_X027114	NIGv2.S00002314_904499-956201-	-2.3
PRPF39.2	c.Taira201203st12_X001103	JGIv7b.000021980_1925557-1957291-	-2.3
RBM34	c.Taira201203lung_X005379	JGIv7b.000075417_5233396-5251942-	-2.3
CINP	c.Audic201207_X046200	JGIv7b.000236382_2970847-2980970-	-2.3
UNNAMED	c.Quigley201212_X006696	JGIv7b.000012518_76668-87512+	-2.3
HNRNPR	c.Quigley201207_X004040	JGIv7b.000030711_1549550-1574750+	-2.3
DYNLL1	c.Quigley201207_X004595	JGIv7b.000036586_112311-114604-	-2.3
PTAFR	c.XenBase_148225481	JGIv7b.000013204_932024-944528+	-2.3
UNNAMED	c.JGIL6RMv1_XeXenL6RMv10045325m	JGIv7b.000337760_15901-58446+	-2.3
PKDCC.1	c.XGI_TC417985	NIGv2.S00000053_637642-653407+	-2.2



ARL6IP1	c.Amin201106_X029507	NIGv2.S00002733_149987-163003-	-2.2
FTH1	c.Chang2013_X041004	NIGv2.S00003324_455821-456877-	-2.2
RPS21	c.XGI_TC456641	JGIv7b.000014557_962191-970899-	-2.2
KRT5.7	c.Taira201203kidney_X011862	JGIv7b.000200825_6634405-6640781+	-2.2
CASP6	c.XGI_TC452334	JGIv7b.000051940_545160-548822+	-2.2
LIN28A	c.TeperekTkacz201205_X002748	NIGv2.S00002590_129600-146832+	-2.2
C5ORF30	c.XenBase_148234248	JGIv7b.000001187_2165049-2174001-	-2.2
UNNAMED	c.Chang2013_X040571	NIGv2.S00002584_1679464-1680608+	-2.2
SESN1	c.TXGP201107_X000908	JGIv7b.000008355_4543017-4556237-	-2.2
UTP18	c.Chang2013_X011369	JGIv7b.000030353_49970-67598+	-2.1
SRP19	c.TeperekTkacz201206_X000865	JGIv7b.000012879_5175567-5182374-	-2.1
TDG	c.XGI_TC417017	JGIv7b.000005925_7934169-7954927-	-2.1
UNNAMED	c.Chang2013_X033037	JGIv7b.000208071_3368708-3370756+	-2.1
GREB1L	c.Quigley201112_X013620	JGIv7b.000085591_509475-593911+	-2.1
UNNAMED	c.JGIL6RMv1_XeXenL6RMv10008609m	JGIv7b.000103160_83520-85320-	-2.1
SHISA2	c.JGIL6RMv1_XeXenL6RMv10028709m	JGIv7b.000137879_671290-673328-	-2.1
ASB3	c.Amin201106_X011834	JGIv7b.000047457_291467-304807-	-2.1
CRX-B	c.JGIL6RMv1_XeXenL6RMv10025952m	JGIv7b.000039723_9163473-9172981+	-2.1
FTH1	c.JGIL6RMv1_XeXenL6RMv10053072m	JGIv7b.000139674_1673507-1677285+	-2.0
CRX-A	c.UniGene_XI_S22245695	JGIv7b.000050079_2769673-2781629-	-2.0
RPL27A	c.Chang2013_X035887	JGIv7b.000287959_146467-152199+	-2.0
HPGD	c.Ismailoglu201203_X012202	JGIv7b.000272351_96769-138279-	-2.0
TUBA1A-B	c.mgEST_1013155827	JGIv7b.000127187_1109513-1113080-	-2.0
WDR12	c.JGIL6RMv1_XeXenL6RMv10042759m	JGIv7b.000139113_1098311-1114621-	-2.0
CDH26	c.Quigley201212_X024302	JGIv7b.000053223_277006-322333+	-2.0
POLR2K	c.JGIL6RMv1_XeXenL6RMv10019863m	JGIv7b.000034503_3942828-3945719-	-2.0
UNNAMED	c.Taira201203st08_X004257	JGIv7b.000107078_130406-136077+	-2.0
SRP19	c.mgEST_1013086260	NIGv2.S00002506_35793-42413-	-2.0
ZFP36L1	c.Ismailoglu201203_X013721	NIGv2.S00003031_949948-957707-	-2.0
HAS-RS	c.Ismailoglu201203_X013818	NIGv2.S00003642_116689-121596+	-2.0



