

		microarray		qPCR					
		MANF ^{mzΔ96} vs w		MANF ^{mzΔ96} vs w		69B>MANF ¹³³			
	GeneName	logFC	P.Value	log2	C(TT)	log2	C(TT)	gene ID	Description
1	CG10420	1.98	0.00010	5.15	9.69E-07	-3.29	1.74588E-05	CG10420	Nucleotide exchange factor SIL1 precursor
2	CG14879	-4.03	7.27E-11	-3.46	0.000367			CG14879	Concanavalin A like lectin homology
3	CG5810	-1.92	1.79E-06	-3.78	1.54E-07			CG5810	leucine rich repeat
4	Ddc	1.62	4.59E-05	2.69	0.000257	-4.99	1.5571E-05	CG10697	DOPA decarboxylase
5	DnaJ-H	1.10	4.12E-05	2.56	2.32E-05			CG9828	DnaJ homolog
6	Hrs	1.56	0.00012	1.38	0.046314			CG2903	Hepatocyte growth factor-regulated tyrosine kinase substrate
7	Hsp83	2.46	9.71E-05	4.57	3.17E-06			CG1242	Heat shock protein 83
8	InR	1.21	0.00069	1.87	3.42E-05			CG18402	Insulin-like receptor precursor
9	Pak3	-1.96	0.00027	-1.14	0.000316			CG14895	Pak , serine threonine kinase
10	path	2.59	6.62E-09	2.66	0.056037			CG3424	Pathetic, amino acid transmembrane transporter
11	Pi3K92E	1.11	0.00030	2.34	0.02905			CG4141	phosphatidylinositol-4-phosphate 3-kinase
12	pipe	-1.53	4.24E-05	-0.25	0.000121			CG9614	heparan sulfate 2-O-sulfotransferase
13	pale	2.83	3.35E-07	4.83	1.06E-06			CG10118	TH, Tyrosine 3-hydroxylase
14	punch	1.97	4.28E-06	8.41	0.000109	0.69	0.02420	CG9441	Punch, GTP cyclohydrolase I
15	Rala	1.48	1.71E-07	0.63	2.12E-05			CG2849	Ras-related protein
16	sulfateless	4.21	1.76E-09	1.62	4.28E-05			CG8339	heparan sulfate glucosaminyl N-deacetylase/N-sulfotransferase
17	Rop	3.33	4.15E-10	0.82	4.28E-05			CG15811	Ras-opposite
18	usnp	-0.43	4.36E-05	-1.18	0.001099			CG11173	Ubisnap, SNAP29 homolog
19	Manf	-1.37	0.000617	-5.39	6.31E-06	1.82	1.8427E-06	CG7013	Manf, known previously as ARMET

Table 2.