

**Table 3**

Fold changes of sample pairs as determined by QRT-PCR (Fp) and cDNA microarray (Fa(cDNA))

Gene	E3/C3				E4/C4				E5/C5			
	PCR	cDNA array			PCR	cDNA array			PCR	cDNA array		
LRG-21	50.8	13.6	12.3	11.4	208.3	34.5	30.0	42.6	102.1	56.6	62.8	45.1
egr1	406.9	6.2	6.4	7.7	277.8	16.7	15.3	16.4	227.8	21.1	18.7	19.3
Nr4a1 (nur77)	46.7	4.0	4.2	4.5	30.6	4.6	5.7	5.6	51.6	10.0	9.9	10.3
ler2 (Pip92)	20.3	3.9	4.3	4.8	14.1	3.6	4.5	4.1	15.0	10.0	9.4	6.1
RGS2	7.4	2.4	2.7	2.7	7.8	3.9	3.4	3.3	7.1	6.7	7.1	12.4
	7.4	0.8	2.1	0.7	7.8	3.6	6.5	3.9	7.1	5.3	4.6	5.1
c-jun	6.0	3.1	3.0	2.6	6.4	3.9	3.7	3.5	4.8	6.0	7.7	6.7
TSC22	2.3	2.6	2.5	2.8	3.4	4.0	4.0	4.1	3.8	4.0	4.0	4.1
	2.3	2.2	2.4	2.4	3.4	5.2	5.3	3.1	3.8	3.9	3.9	4.0
gamma-actin	2.8	2.3	2.1	2.1	4.3	2.6	2.6	2.7	5.0	3.8	3.6	3.9
beta-actin	1.4	1.1	0.9	1.0	2.2	1.3	1.3	1.1	1.9	2.2	2.2	2.1
	1.4	1.4	1.4	1.3	2.2	3.0	3.1	2.9	1.9	2.4	2.2	2.5
KLF4	3.1	1.2	1.4	1.3	3.2	1.6	1.8	1.9	2.3	1.9	1.6	2.2
Period1	3.9	1.8	1.7	1.8	3.8	2.2	2.2	2.3	3.1	1.9	2.1	2.1
Ptp4a1	1.8	1.0	0.9	0.9	2.2	1.4	1.6	1.7	2.3	2.8	2.7	2.7
	1.8	1.3	1.1	1.4	2.2	2.4	3.0	2.6	2.3	2.8	2.7	2.6
gly96	3.7	1.4	1.7	1.2	3.1	0.8	1.3	1.3	4.2	2.5	4.3	2.4
I-kappa B	2.5	1.5	1.6	1.6	2.5	1.5	1.6	1.7	1.5	1.8	1.6	1.6
Gem	3.1	1.2	1.2	1.5	4.0	2.2	2.0	1.5	3.9	1.8	1.7	2.2
egr2	288.3	1.1	1.4	1.4	167.8	1.3	1.6	1.3	372.4	2.5	2.5	2.8
ptpn16 (MKP1)	3.6	1.4	1.4	1.6	2.4	1.0	1.3	1.2	2.6	2.1	1.4	1.7
SCL	1.9	0.8	0.8	0.9	2.3	0.9	1.0	0.7	2.4	1.4	1.4	1.4
glucose transport protein	1.2	0.7	0.8	0.7	2.8	1.0	1.0	0.7	2.9	2.4	2.5	2.4
junD	1.3	1.5	1.7	1.4	1.9	1.4	1.5	1.5	1.1	2.8	2.4	2.7
mthfd2	1.1	1.0	0.9	0.9	1.4	1.7	1.8	1.9	1.6	1.5	1.6	1.5
Stat3B	1.3	1.3	1.1	0.9	1.1	1.6	1.6	1.8	1.6	1.4	1.4	1.2
Nrf2	1.1	0.9	1.1	1.2	1.1	1.6	1.8	1.7	1.5	1.3	1.3	1.4
	1.1	1.2	1.0	1.2	1.1	1.5	1.5	1.5	1.5	1.4	1.3	1.5
Gata2	1.0	0.8	0.7	0.8	0.5	0.6	0.5	0.6	1.0	0.9	1.0	0.9
Gata6	0.7	1.0	1.1	1.0	0.6	0.8	0.7	0.7	0.5	0.8	0.8	0.6
chloride channel 3	1.0	1.2	1.2	1.2	1.2	1.4	1.3	1.5	1.5	1.3	1.3	1.4
Cox8a	1.1	1.0	1.1	1.2	0.9	0.6	0.7	0.7	0.9	1.1	1.1	1.1
elongation factor 2	3.4	1.0	1.1	1.0	1.1	1.3	1.2	1.3	0.5	1.1	1.2	1.1
fatty acid synthase	2.0	1.3	1.3	1.5	0.9	1.2	1.2	1.1	0.7	1.3	1.5	1.4
G-binding protein	0.9	1.1	1.2	1.1	0.9	1.1	1.1	1.1	1.1	1.1	1.1	1.1
growth factor-induced	0.8	0.8	0.9	0.8	0.8	0.6	0.6	0.6	1.0	1.4	1.3	1.4
Hdac1	0.9	1.0	1.0	1.0	0.9	1.1	1.5	1.6	1.1	0.9	0.9	0.9
	1.4	1.1	1.1	1.1	0.9	1.0	1.0	1.0	1.2	1.1	1.2	1.1
HSP86-1	1.4	1.0	0.8	0.8	0.9	1.3	1.3	0.8	1.2	1.1	1.0	1.1
Na,K-ATPase	0.9	0.7	0.6	0.9	1.4	1.1	1.4	1.1	1.3	1.4	1.5	1.4
26S proteasome	1.0	0.9	1.0	0.9	0.9	1.1	1.0	1.0	1.0	0.9	0.8	0.8
PRL2	1.0	1.0	1.0	1.0	1.5	0.8	0.8	0.8	1.2	1.1	1.2	1.2
small GTP-binding protein	1.1	1.0	1.0	1.0	1.2	0.8	0.8	0.8	1.2	1.0	1.2	1.0
ribosomal protein L37	0.9	0.8	0.7	0.8	0.9	1.0	1.0	0.8	1.1	1.0	0.9	1.0
ribosomal protein L7	1.1	0.9	0.9	0.9	1.2	1.0	1.2	1.0	1.3	0.9	0.9	0.9
ribosomal protein S11	0.8	1.1	1.1	1.0	0.8	1.1	1.0	1.0	1.1	0.9	1.0	0.9
ribosomal protein S3	1.2	0.8	0.9	0.8	1.1	1.1	1.2	1.0	1.1	0.9	1.0	0.9
alpha-tubulin	0.9	1.2	1.2	1.2	0.9	0.9	1.0	1.0	1.0	1.2	1.2	1.2
HSP40	1.5	1.2	1.1	1.3	1.2	1.2	1.2	1.3	1.3	2.0	1.6	1.7
MARCKS-related protein	1.1	1.1	1.1	1.2	1.0	1.0	1.1	0.9	1.1	1.1	1.1	1.1
STY Kinase	1.6	1.2	1.3	1.2	1.3	1.4	1.3	1.3	1.4	1.2	1.2	1.4
hsr.1	0.9	1.0	1.0	1.1	1.1	0.8	0.8	0.8	1.1	1.2	1.1	1.2
malate dehydrogenase	1.0	1.0	1.0	1.1	0.9	1.0	1.0	1.1	1.2	0.8	0.9	0.8