

cuffdiff of replicates

test_id	gene_id	gene	locus	sample_1	sample_2	status	value_1	value_2
ENSRNOG0	ENSRNOG0	NCKX1_RA	8:6917554	T	P	OK	2.31711	7.77389
ENSRNOG0	ENSRNOG0	RGD13592	10:965949	T	P	OK	2.61807	18.769
ENSRNOG0	ENSRNOG0	RGD13592	10:965949	T	N	OK	2.61807	17.3716

cuffdiff of individual genes

gene_id	gene_short	1T_FPKM	2T_FPKM	3T_FPKM	4T_FPKM	1P_FPKM	2P_FPKM	3P_FPKM
ENSRNOG0	NCKX1_RA	3.19567	1.25174	1.84031	2.30138	12.5279	13.5085	2.25365
ENSRNOG0	RGD13592	2.56691	3.12167	3.15359	1.72578	0.717251	0.852209	2.82114

log2(fold_change)	test_stat	p_value	q_value	significant
1.74631	-4.63718	3.53E-06	0.025512	yes
2.84178	-5.21674	1.82E-07	0.002302	yes
2.73016	-4.94187	7.74E-07	0.006521	yes

4P_FPKM	1N_FPKM	2N_FPKM	3N_FPKM	4N_FPKM
1.77911	2.45833	2.43647	11.3845	11.745
2.25712	1.14159	3.01218	21.64	1.54513