

Supplementary Table 2: Differentially Expressed Genes (DEGs) between DEX and CTR groups (DEX vs. CTR).

#	Agilent Array ID	Symbol	Corresponding Gene	Gene Full Name	ENTREZ ID	Fold change	q Value (FDR%)
1	A_73_100146	LOC509808	LIPG	lipase, endothelial	509808	-14.76	0.00
2	A_73_107233	CYP1A1	CYP1A1	cytochrome P450, subfamily I (aromatic compound-inducible), polypeptide 1	282870	-10.41	2.75
3	A_73_117912	LOC535704	FKBP5	FK506 binding protein 5	535704	-8.65	1.87
4	A_73_118642	LOC505329	CRISPLD2	cysteine-rich secretory protein LCCL domain containing 2	505329	-7.07	2.75
5	A_73_120575	MGP	MGP	matrix Gla protein	282660	-6.66	2.75
6	A_73_108526	LOC510967	CTSW	cathepsin W	510967	-6.35	1.87
7	A_73_108314	LOC540417	IRX3	iroquois homeobox 3	540417	-5.82	5.04
8	A_73_108050	XLKD1	LYVE1	lymphatic vessel endothelial hyaluronan receptor 1	404179	-5.74	1.87
9	A_73_119473	LOC539430	OSR1	odd-skipped related 1 (Drosophila)	539430	-5.3	3.30
10	A_73_111228	LOC540817	MOB3B	MOB kinase activator 3B	540817	-5.17	4.55
11	A_73_114373	LOC617258	CCL24	chemokine (C-C motif) ligand 24	617258	-5.14	1.87
12	A_73_106522	PGF	PGF	placental growth factor	280894	-5.11	3.30
13	A_73_108953	AGTR1	AGTR1	angiotensin II receptor, type 1	281607	-4.98	0.00
14	A_73_117872	SLAM	SLAMF1	signaling lymphocytic activation molecule family member 1	281489	-4.93	4.48
15	A_73_114230	PGDH	HPGD	hydroxyprostaglandin dehydrogenase 15-(NAD)	512259	-4.88	3.86
16	A_73_111455	LOC533844	CD163	CD163 molecule	533844	-4.8	2.75
17	A_73_114133	LOC507449	RASD1	RAS, dexamethasone-induced 1	507449	-4.75	0.00
18	A_73_112275	CPXM	CPXM1	carboxypeptidase X (M14 family), member 1	532219	-4.71	2.75
19	A_73_110765	LOC617799	CST7	cystatin F (leukocystatin)	617799	-4.7	3.39
20	A_73_116484	LOC504445	DKK1	dickkopf homolog 1 (Xenopus laevis)	504445	-4.59	2.75
21	A_73_114811	LOC504909	ABCA10	ATP-binding cassette, sub-family A (ABC1), member 10	504909	-4.54	2.55
22	A_73_114060	LOC535300	TRGC6	T-cell receptor gamma chain TRGC6	535300	-4.51	2.75
23	A_73_100352	MYOC	MYOC	myocilin, trabecular meshwork inducible glucocorticoid response	281342	-4.44	0.00
24	A_73_112354	LOC523285	ARRDC2	arrestin domain containing 2	523285	-4.4	2.75
25	A_73_108054	LOC534928	PFKFB4	6-phosphofructo-2-kinase/fructose-2,6-biphosphatase 4	534928	-4.37	2.78
26	A_73_116172	EIF4B	EIF4B	eukaryotic translation initiation factor 4B	505850	-4.34	1.87
27	A_73_102507	NFKBIA	NFKBIA	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, alpha	282291	-4.26	1.87

28	A_73_110879	LOC505824	NTRK2	neurotrophic tyrosine kinase, receptor, type 2	505824	-4.04	4.55
29	A_73_106942	LOC540812	ARMC12	armadillo repeat containing 12	540812	-3.99	3.30
30	A_73_109217	LOC516318	PER1	period circadian clock 1	516318	-3.95	1.87
31	A_73_111837	PTGES	PTGES	prostaglandin E synthase	282019	-3.79	0.00
32	A_73_106842	LOC509968	C1QC	complement component 1, q subcomponent, C chain	509968	-3.66	5.04
33	A_73_118275	C1QA	C1QA	complement component 1, q subcomponent, A chain	534961	-3.6	3.86
34	A_73_118933	LOC504278	ABCA9	ATP-binding cassette, sub-family A (ABC1), member 9	504278	-3.49	0.00
35	A_73_104727	CD3E	CD3E	CD3e molecule, epsilon (CD3-TCR complex)	281054	-3.42	0.00
36	A_73_121418	SLC11A1	SLC11A1	solute carrier family 11 (proton-coupled divalent metal ion transporters), member 1	282470	-3.42	3.39
37	A_73_104525	LOC514285	ANGPTL5	angiopoietin-like 5	514285	-3.41	0.00
38	A_73_114363	PRF1	PRF1	perforin 1 (pore forming protein)	369025	-3.31	4.48
39	A_73_102645	CD69	CD69	CD69 molecule	281058	-3.31	3.30
40	A_73_119464	LOC510187	MEDAG	mesenteric estrogen-dependent adipogenesis	510187	-3.28	2.75
41	A_73_119745	LOC511774	C1QTNF1	C1q and tumor necrosis factor related protein 1	511774	-3.17	1.87
42	A_73_115704	LOC511711	FGL2	fibrinogen-like 2	511711	-3.16	1.87
43	A_73_107180	LOC535075	ALDH1A2	aldehyde dehydrogenase 1 family, member A2	535075	-3.14	4.48
44	A_73_118870	LOC514557	OAF	OAF homolog (Drosophila)	514557	-3.09	1.87
45	A_73_107724	LRAT	LRAT	lecithin retinol acyltransferase (phosphatidylcholine--retinol O-acyltransferase)	281285	-3.05	2.75
46	A_73_119307	LOC534389	FAM105A	family with sequence similarity 105, member A	534389	-3.04	2.55
47	A_73_104367	LOC617024	E2F2	E2F transcription factor 2	617024	-3.02	2.75
48	A_73_103736	LOC536989	PLXDC1	plexin domain containing 1	NA	-2.99	2.75
49	A_73_100779	LOC615535	ENPP1	ectonucleotide pyrophosphatase/phosphodiesterase 1	615535	-2.97	0.00
50	A_73_121104	LOC404176	TRD@	T-cell receptor delta chain	407199	-2.94	2.75
51	A_73_104591	LOC511286	LOC511286	polyadenylate-binding protein 5-like	511286	-2.92	3.86
52	A_73_118155	SNAI2	SNAI2	snail homolog 2 (Drosophila)	520631	-2.85	2.11
53	A_73_100851	LOC533166	SORL1	sortilin-related receptor, L(DLR class) A repeats containing	NA	-2.81	2.78
54	A_73_108986	LOC523616	C1QTNF9	C1q and tumor necrosis factor related protein 9	NA	-2.78	1.87
55	A_73_117589	ICAM1	ICAM1	intercellular adhesion molecule 1	281839	-2.78	1.87
56	A_73_115458	LOC517232	RECK	reversion-inducing-cysteine-rich protein with kazal motifs	517232	-2.77	2.55
57	A_73_111558	LOC512705	TDRD6	tudor domain containing 6	512705	-2.74	3.30
58	A_73_115499	LOC513053	OLFML2B	olfactomedin-like 2B	513053	-2.7	2.75
59	A_73_111072	NP	PNP	purine nucleoside phosphorylase	493724	-2.69	2.55

60	A_73_109372	AOX1	AOX1	aldehyde oxidase 1	338074	-2.69	2.75
61	A_73_117551	LOC529661	TSKU	tsukushi small leucine rich proteoglycan homolog (Xenopus laevis)	NA	-2.66	1.87
62	A_73_117868	LOC509044	LOC509044	E2F transcription factor 6 pseudogene	509044	-2.64	2.75
63	A_73_114013	MMP2	MMP2	matrix metalloproteinase 2 (gelatinase A, 72kDa gelatinase, 72kDa type IV collagenase)	282872	-2.58	0.00
64	A_73_110199	LOC514062	EHMT2	euchromatic histone-lysine N-methyltransferase 2	514062	-2.55	3.86
65	A_73_101341	LOC532674	GPR116	G protein-coupled receptor 116	532674	-2.55	1.87
66	A_73_100213	FZD1	FZD1	frizzled family receptor 1	445417	-2.54	1.87
67	A_73_105025	SULT1A1	SULT1A1	sulfotransferase family, cytosolic, 1A, phenol-preferring, member 1	282485	-2.54	0.00
68	A_73_110571	LOC507339	C7	complement component 7	507339	-2.54	4.48
69	A_73_104593	LOC538691	HS3ST1	heparan sulfate (glucosamine) 3-O-sulfotransferase 1	538691	-2.53	2.55
70	A_73_117114	LOC515235	CCDC80	coiled-coil domain containing 80	515235	-2.53	3.30
71	A_73_115034	LOC539139	KLF9	Kruppel-like factor 9	539139	-2.52	1.87
72	A_73_113155	LOC510973	NUMBL	numb homolog (Drosophila)-like	510973	-2.48	4.48
73	A_73_116772	LOC523226	PCDHA13	protocadherin alpha 13	523226	-2.48	2.75
74	A_73_110273	LOC513870	HPCAL1	hippocalcin-like 1	513870	-2.48	0.00
75	A_73_118201	GCNT1	GCNT1	glucosaminyl (N-acetyl) transferase 1, core 2	281778	-2.47	2.75
76	A_73_103439	LOC523294	GLT8D2	glycosyltransferase 8 domain containing 2	523294	-2.46	2.75
77	A_73_105124	LOC530657	COL6A3	collagen, type VI, alpha 3	530657	-2.43	1.87
78	A_73_108287	MGC128119	GPC3	glypican 3	615239	-2.42	2.75
79	A_73_116856	LOC518469	GPCPD1	glycerophosphocholine phosphodiesterase GDE1 homolog (S. cerevisiae)	518469	-2.38	0.00
80	A_73_104301	FZD4	FZD4	frizzled class receptor 4	445416	-2.37	0.00
81	A_73_110545	LOC513468	CD244	CD244 molecule, natural killer cell receptor 2B4	513468	-2.36	2.75
82	A_73_120343	LOC515988	STAT4	signal transducer and activator of transcription 4	515988	-2.35	3.86
83	A_73_104758	DCN	DCN	decorin	280760	-2.35	2.11
84	A_73_102351	LOC529235	DDIT4	DNA-damage-inducible transcript 4	529235	-2.35	5.04
85	A_73_112805	LOC534859	B3GNT8	UDP-GlcNAc:betaGal beta-1,3-N-acetylglucosaminyltransferase 8	534859	-2.32	3.30
86	A_73_118691	MFAP2	MFAP2	microfibrillar-associated protein 2	NA	-2.32	1.87
87	A_73_109561	LOC510055	GIMAP7	GTPase, IMAP family member 7	100125415	-2.29	5.04
88	A_73_105084	LOC614109	LOC614109	sodium/bile acid cotransporter 7-like	614109	-2.27	1.87

89	A_73_120962	MGC127421	GSN	gelsolin	535077	-2.26	3.86
90	A_73_104305	LOC504741	COLEC12	collectin sub-family member 12	504741	-2.25	2.78
91	A_73_119299	DMBT1	DMBT1	deleted in malignant brain tumors 1	404104	-2.24	3.30
92	A_73_104396	LOC526800	ANKRD44	ankyrin repeat domain 44	526800	-2.24	2.75
93	A_73_112690	LOXL4	LOXL4	lysyl oxidase-like 4	281904	-2.22	3.86
94	A_73_103413	LOC510173	SH2D3A	SH2 domain containing 3A	510173	-2.21	2.55
95	A_73_105509	PIK3R1	PIK3R1	phosphoinositide-3-kinase, regulatory subunit 1 (alpha)	282307	-2.2	2.75
96	A_73_105627	LOC538419	EPS8	epidermal growth factor receptor pathway substrate 8	538419	-2.19	0.00
97	A_73_102713	LOC536815	ROBO1	roundabout, axon guidance receptor, homolog 1 (Drosophila)	536815	-2.18	1.87
98	A_73_118731	CCND3	CCND3	cyclin D3	540547	-2.17	0.00
99	A_73_120257	LOC510988	GIMAP7	GTPase, IMAP family member 7	NA	-2.17	1.87
100	A_73_103818	LOC511883	ANKRD13A	ankyrin repeat domain 13A	511883	-2.17	2.55
101	A_73_112008	LOC540224	RND3	rho-related GTP-binding protein RhoE	540224	-2.17	0.00
102	A_73_116912	PTPN13	PTPN13	protein tyrosine phosphatase, non-receptor type 13 (APO-1/CD95 (Fas)-associated phosphatase)	282333	-2.16	2.75
103	A_73_120676	LOC508029	TRPM2	transient receptor potential cation channel, subfamily M, member 2	508029	-2.14	4.48
104	A_73_118934	LOC507012	TACC1	transforming, acidic coiled-coil containing protein 1	507012	-2.13	1.87
105	A_73_103264	MYO10	MYO10	myosin X	281935	-2.13	4.55
106	A_73_110964	LOC534512	HMCN2	hemicentin 2	534512	-2.13	1.87
107	A_73_109408	SERPING1	SERPING1	serpin peptidase inhibitor, clade G (C1 inhibitor), member 1	281035	-2.13	1.87
108	A_73_120175	LOC616702	LOC616702	polyhomeotic-like protein 2-like	616702	-2.12	3.39
109	A_73_110662	ms4A8B	MS4A8B	membrane-spanning 4-domains, subfamily A, member 8B	415111	-2.12	1.87
110	A_73_104788	LOC539762	ZFHX4	zinc finger homeobox 4	539762	-2.12	3.39
111	A_73_110856	LOC538590	COL5A2	collagen, type V, alpha 2	538590	-2.11	4.48
112	A_73_106577	LOC518833	CYFIP2	cytoplasmic FMR1 interacting protein 2	518833	-2.1	4.48
113	A_73_101912	EGFR	EGFR	epidermal growth factor receptor	407217	-2.09	5.04
114	A_73_114788	LOC505220	PTPN9	protein tyrosine phosphatase, non-receptor type 9	505220	-2.09	2.78
115	A_73_118971	LOC527934	CORO7	coronin 7	527934	-2.09	1.87
116	A_73_108023	LOC539216	KRT26	keratin 26	539216	-2.08	4.48
117	A_73_109510	LOC505067	ZNF404	zinc finger protein 404	505067	-2.08	2.75
118	A_73_108520	LOC617566	BMP6	bone morphogenetic protein 6	617566	-2.07	2.75

119	A_73_111335	LOC507890	PRDM8	PR domain containing 8	507890	-2.06	2.78
120	A_73_100615	LOC533523	DDR2	discoidin domain receptor tyrosine kinase 2	533523	-2.05	0.00
121	A_73_102824	MGC127307	KLF6	Kruppel-like factor 6	505884	-2.04	2.78
122	A_73_105540	GABARAPL1	GABARAPL1	GABA(A) receptor-associated protein like 1	338472	-2.03	2.75
123	A_73_107041	LOC534787	ITIH5	inter-alpha-trypsin inhibitor heavy chain family, member 5	534787	-2.03	0.00
124	A_73_117183	LGALS4	LGALS4	lectin, galactoside-binding, soluble, 4	614804	-2	2.75
125	A_73_117734	EE898361	EST	B13281A FFB Bos taurus cDNA clone B1328 3-, mRNA sequence	NA	-6.47	3.385321
126	A_73_121153	BI774772	EST	466988 MARC 1BOV Bos taurus cDNA 5-, mRNA sequence	NA	-5.51	0
127	A_73_102870	BE757006	EST	211333 MARC 2BOV Bos taurus cDNA 5-, mRNA sequence	NA	-5.28	2.754722
128	A_73_116074	CB420054	EST	592969 MARC 6BOV Bos taurus cDNA 3-, mRNA sequence	NA	-5.11	2.754722
129	A_73_115556	BF602046	EST	267182 MARC 3BOV Bos taurus cDNA 5-, mRNA sequence	NA	-5.07	1.873211
130	A_73_104896	BM251639	EST	BOTL0100011_E09 Normalized Bovine Total Leukocyte cDNA Library, mRNA sequence	NA	-4.76	2.754722
131	A_73_101432	AV612448	EST	AV612448 Bos taurus lung fetus Bos taurus cDNA clone E1LU045F02 3-, mRNA sequence	NA	-4.57	3.29848
132	A_73_105610	CB457253	EST	714591 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-4.38	1.873211
133	A_73_101119	BM929188	EST	LM24HW00494 Bos taurus LM-24-HW cDNA library Bos taurus cDNA clone	NA	-4.05	1.873211
134	A_73_107216	CB468423	EST	734243 MARC 6BOV Bos taurus cDNA 3-, mRNA sequence	NA	-4.02	2.754722
135	A_73_105437	BE590089	EST	197038 BARC 5BOV Bos taurus cDNA 5-, mRNA sequence	NA	-3.94	2.107362
136	A_73_104653	EE911626	EST	A46231A FFB Bos taurus cDNA clone A4623 3-, mRNA sequence	NA	-3.73	1.873211
137	A_73_115308	BF076063	EST	225461 MARC 2BOV Bos taurus cDNA 5-, mRNA sequence	NA	-3.71	3.860816

138	A_73_101427	EE902681	EST	B38100A FFB Bos taurus cDNA clone B3810 5-, mRNA sequence	NA	-3.51	1.873211
139	A_73_104046	BE668539	EST	158369 MARC 4BOV Bos taurus cDNA 5-, mRNA sequence	NA	-3.5	0
140	A_73_118346	CB530263	EST	737046 MARC 6BOV Bos taurus cDNA 3-, mRNA sequence	NA	-3.34	2.107362
141	A_73_111525	EE906844	EST	A19370A FFB Bos taurus cDNA clone A1937 5-, mRNA sequence	NA	-3.24	1.873211
142	A_73_121057	EE912709	EST	A52341A FFB Bos taurus cDNA clone A5234 3-, mRNA sequence	NA	-3.23	0
143	A_73_100263	hmm89873	EST	Bos taurus similar to Homo sapiens SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 2 (SMARCD2)	NA	-3.22	5.036463
144	A_73_117300	CB448153	EST	702214 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-3.2	1.873211
145	A_73_120831	AV613356	EST	AV613356 Bos taurus adipocyte cell line Bos taurus cDNA clone E0AD003F06 3-, mRNA sequence	NA	-3.18	4.483749
146	A_73_104923	BM251291	EST	BOTL0100004XE05R Normalized Bovine Total Leukocyte cDNA Library, mRNA sequence	NA	-3.18	2.754722
147	A_73_117696	AV664629	EST	AV664629 Bos taurus brain fetus Bos taurus cDNA clone E1BR043E08 5-, mRNA sequence	NA	-3.15	2.107362
148	A_73_107406	CA035529	EST	4001394 BARC 5BOV Bos taurus cDNA clone 5BOV_78I15 3-, mRNA sequence	NA	-3.13	3.29848
149	A_73_119914	CB463539	EST	723921 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-3.05	3.385321
150	A_73_105393	CB441503	EST	691988 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.99	3.860816
151	A_73_115493	AU277966	EST	AU277966 Cloned bovine placenta cDNA Bos taurus cDNA clone placenta0009 3-, mRNA sequence	NA	-2.97	1.873211
152	A_73_120047	CB444768	EST	695972 MARC 6BOV Bos taurus cDNA 3-, mRNA sequence	NA	-2.96	2.754722
153	A_73_106449	CB459252	EST	718851 MARC 6BOV Bos taurus cDNA 3-, mRNA sequence	NA	-2.94	0

154	A_73_117645	CB437117	EST	684554 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.86	2.754722
155	A_73_120589	CB455280	EST	712414 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.76	4.545291
156	A_73_103725	EE890884	EST	A71560A FFB Bos taurus cDNA clone A7156 5-, mRNA sequence	NA	-2.72	2.754722
157	A_73_100461	CB442787	EST	693596 MARC 6BOV Bos taurus cDNA 3-, mRNA sequence	NA	-2.68	2.783309
158	A_73_106056	AW314086	EST	9897 MARC 4BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.67	3.29848
159	A_73_102653	CB463123	EST	723483 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.66	2.554378
160	A_73_119171	CB454202	EST	710930 MARC 6BOV Bos taurus cDNA 3-, mRNA sequence	NA	-2.66	0
161	A_73_113276	AV605678	EST	AV605678 Bos taurus kidney fetus Bos taurus cDNA clone E1KI030B12 5-, mRNA sequence	NA	-2.65	2.754722
162	A_73_115578	BE751862	EST	204100 MARC 2BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.63	2.754722
163	A_73_119001	AU233205	EST	AU233205 Cloned bovine fetus cDNA Bos taurus cDNA clone Cln2639 3-, mRNA sequence	NA	-2.61	3.29848
164	A_73_108977	BE808713	EST	213977 MARC 2BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.6	1.873211
165	A_73_103649	EE897589	EST	B09160A FFB Bos taurus cDNA clone B0916 5-, mRNA sequence	NA	-2.52	0
166	A_73_102260	CB461540	EST	721519 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.51	2.107362
167	A_73_108130	CB421364	EST	594380 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.49	2.754722
168	A_73_110101	EE927967	EST	C50790A FLM Bos taurus cDNA clone C5079 5-, mRNA sequence	NA	-2.49	2.783309
169	A_73_119452	AU234098	EST	AU234098 Bovine placenta cDNA Bos taurus cDNA clone Cln689 3-, mRNA sequence	NA	-2.48	1.873211
170	A_73_119380	BE756602	EST	210787 MARC 2BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.48	0

171	A_73_107480	CB531076	EST	742326 MARC 6BOV Bos taurus cDNA 3-, mRNA sequence	NA	-2.44	2.554378
172	A_73_115732	CB443118	EST	693981 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.39	1.873211
173	A_73_116546	AV591119	EST	AV591119 Bos taurus brain fetus Bos taurus cDNA clone E1BR016G01 3-, mRNA sequence	NA	-2.39	4.483749
174	A_73_103623	CB434062	EST	610645 MARC 6BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.37	1.873211
175	A_73_116770	EE904623	EST	B49130A FFB Bos taurus cDNA clone B4913 5-, mRNA sequence	NA	-2.35	2.754722
176	A_73_101194	AU278667	EST	AU278667 Cloned bovine placenta cDNA Bos taurus cDNA clone placenta0908 3-, mRNA sequence	NA	-2.34	1.873211
177	A_73_118075	BE664117	EST	148327 MARC 4BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.32	2.783309
178	A_73_121311	EE909038	EST	A03950A FFB Bos taurus cDNA clone A0395 5-, mRNA sequence	NA	-2.32	1.873211
179	A_73_115999	AU233009	EST	AU233009 Cloned bovine fetus cDNA Bos taurus cDNA clone Cln2358 3-, mRNA sequence	NA	-2.3	2.754722
180	A_73_118549	EE891345	EST	A01741A FFB Bos taurus cDNA clone A0174 3-, mRNA sequence	NA	-2.29	2.754722
181	A_73_120769	BM087935	EST	501266 MARC 2BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.29	3.29848
182	A_73_104104	AW670011	EST	113707 MARC 1BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.27	1.873211
183	A_73_112368	BI540081	EST	453023 MARC 1BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.26	2.107362
184	A_73_109549	BM087067	EST	499723 MARC 2BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.22	1.873211
185	A_73_110980	AW417127	EST	53100 MARC 1BOV Bos taurus cDNA 5-, mRNA sequence	NA	-2.22	0
186	A_73_106175	EE904189	EST	B46500A FFB Bos taurus cDNA clone B4650 5-, mRNA sequence	NA	-2.22	2.783309
187	A_73_115949	LOC509513	uncharacterized LOC509513	////	NA	-2.11	1.873211
188	A_73_113881	LOC518677	Withdrawn	////	NA	-2.1	1.873211
189	A_73_113483	LOC615516	Withdrawn	////	NA	-2.1	5.036463

190	A_73_104118	LOC520492	Withdrawn	////	NA	-2.09	4.545291
191	A_73_114289	LOC540066	Withdrawn	////	NA	-2.08	5.036463
192	A_73_118629	LOC530791	Withdrawn	////	NA	-2.07	3.29848
193	A_73_111205	LOC537532	Withdrawn	////	NA	-2.04	2.754722
194	A_73_105716	LOC537902	Withdrawn	////	NA	-2.03	3.860816
195	A_73_100558	LOC514145	Withdrawn	////	NA	-2.02	1.873211
196	A_73_108800	LOC617970	Withdrawn	////	NA	-2.01	0
197	A_73_116241	LOC617194	Withdrawn	////	NA	-2.01	2.754722
198	A_73_116068	LOC520357	Withdrawn	////	NA	-2	4.545291

NOTES:*EST: Estimated Sequence Tag**NA: Not Available**Withdrawn: Not available anymore in the genebank database (*