Human UniGene 1 Results

Experiment Results Generated by GEMTools 2.5
Client: Genome_Systems
Export Date: Oct 12, 2001

GEM	022JC38J
Balance Coefficient	1.19
Minimum S/B	2.5
Minimum Area	40%
Probe 1	123YA1BU
P1 Description	d17/5
Probe 2	1235A1BV
P2 Description	wt

Report sorted by **Balanced Diff Expr** in **Ascending** order
Ranks from **101** to **200**

Order LifeArray clones

Legend

Rank	Location	Diff Expr	Balanced Diff Expr	P1 Signal	P1 S/B	P1 Area %	P2 Balanced Signal	P2 Signal	P2 S/B	P2 Area %	Plate Row	Plate Col	Plate ID	Gene Name	PCR Status	GenBank Id	Clone Id (Sequence)	Vector
101	8632	-1.6	**-1.9**	371	4.5	85	703	591	10.7	85	H	8	021IAGLP	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	Passed	X72889 Entrez UniGene	2944175	pINCY

102	3089	-1.6	**-1.9**	142	2.1	83†	263	221	3.9	83	C	10	021NAGL7	fascin (Strongylocentrotus purpuratus) homolog 2 (actin-bundling protein, retinal)	Passed	NM_012418 Entrez UniGene	4855125	pINCY
103	6438	-1.6	**-1.9**	480	5.2	100	934	785	13.9	100	F	11	0218AGM4	thymidine kinase 1, soluble	Passed	BE746447 Entrez UniGene	2055926	pSport1
104	629	-1.6	**-1.9**	496	4.4	100	933	784	12.4	100	A	9	021SAGLB	UDP-N-acetyl-alpha-D-galactosamine:polypeptide N-acetylgalactosaminyltransferase 2 (GalNAc-T2)	Passed	AL580930 Entrez UniGene	516735	pSport1
105	8186	-1.6	**-1.9**	514	5.8	98	997	838	14.7	98	D	4	021NAGL7	potassium intermediate/ small conductance calcium-activated channel, subfamily N, member 4	Passed	BG741215 Entrez UniGene	2793840	pINCY
106	4503	-1.6	**-1.9**	545	5.0	67	1014	852	11.7	67	C	6	021NAGMU	lamin B2	Passed	AU130032 Entrez UniGene	2414632	pINCY
107	527	-1.6	**-1.9**	859	7.0	100	1626	1366	19.0	100	G	9	021GAGL6	tumor rejection antigen (gp96) 1	Passed	AV655628 Entrez UniGene	3624631	pINCY

108	5144	-1.6	**-1.9**	974	6.2	44	1833	1540	14.9	44	H	3	021RAGKM	protein disulfide isomerase related protein (calcium-binding protein, intestinal-related)	Passed	NM_004911 Entrez UniGene	1824957	pINCY
109	63	-1.6	**-1.9**	1283	10.1	65	2395	2013	31.0	65	E	5	021YAGKN	FXYD domain-containing ion transport regulator 2	Passed	BG166035 Entrez UniGene	2380042	pINCY
110	7557	-1.6	**-1.9**	1387	12.0	65	2658	2234	34.8	65	D	5	021WAGNF	malate dehydrogenase 1, NAD (soluble)	Passed	NM_005917 Entrez UniGene	487539	pBlue
111	104	-1.6	**-1.9**	2197	13.1	55	4075	3424	35.9	55	C	3	021CAGKP	KIAA0084 protein	Passed	AL561563 Entrez UniGene	2697959	pINCY
112	415	-1.6	**-1.9**	3427	27.0	62	6562	5514	87.2	62	C	1	021OAGL2	RNA-binding protein regulatory subunit	Passed	BG715880 Entrez UniGene	1879921	pINCY
113	7301	-1.6	**-1.9**	227	2.4	79†	433	364	5.5	79	F	9	021EAGN4	hypothetical protein KIAA1165	No Amplification	R82692 Entrez UniGene	55889	pBlue

114	3622	-1.5	**-1.8**	152	2.2	100†	273	229	4.3	100	E	8	021AAGLT	arginine-rich, mutated in early stage tumors	Passed	AA974308 Entrez UniGene	1618455	pINCY
115	3787	-1.5	**-1.8**	153	2.1	92†	269	226	4.0	92	E	2	021GAGM0	KIAA0974 protein	Passed	BE669938 Entrez UniGene	4540	pBlue
116	3017	-1.5	**-1.8**	173	2.2	86†	307	258	4.4	86	C	10	0212AGL4	platelet activating receptor homolog	Passed	NM_013308 Entrez UniGene	3879095	pINCY
117	6495	-1.5	**-1.8**	154	2.1	86†	276	232	4.1	86	B	5	021TAGM7	Homo sapiens cDNA: FLJ21897 fis, clone HEP03447, highly similar to AF052178 Homo sapiens clone 24523 mRNA sequence	Passed	BE259271 Entrez UniGene	2518964	pINCY
118	6374	-1.5	**-1.8**	1481	14.2	100	2600	2185	36.2	100	B	3	021UAGM2	FYN-binding protein (FYB-120/130)	Passed	AF198052 Entrez UniGene	1614872	pINCY
119	9411	-1.5	**-1.8**	268	2.9	88	471	396	5.7	88	D	6	0213AGMM	synaptophysin-like protein	Passed	S72481 Entrez UniGene	1798283	pINCY

120	7570	-1.5	**-1.8**	287	3.3	82	512	430	7.6	82	H	7	021WAGNF	S-adenosylhomocysteine hydrolase-like 1	Passed	AL527927 Entrez UniGene	1739627	pSport1
121	7577	-1.5	**-1.8**	297	3.2	70	528	444	7.6	70	B	9	0213AGNG	CD58 antigen, (lymphocyte function-associated antigen 3)	Passed	NM_001779 Entrez UniGene	342285	pBlue
122	1322	-1.5	**-1.8**	212	2.5	96	383	322	6.0	96	A	3	0218AGM4	nuclear matrix protein p84	Passed	AV713026 Entrez UniGene	22715	pBlue
123	6786	-1.5	**-1.8**	2578	22.8	55	4618	3881	56.4	55	B	11	021IAGMJ	tubulin, beta polypeptide	Passed	BG762520 Entrez UniGene	3334367	pINCY
124	2716	-1.5	**-1.8**	1160	10.8	66	2048	1721	28.3	66	G	8	021QAGKR	diazepam binding inhibitor (GABA receptor modulator, acyl-Coenzyme A binding protein)	Passed	AA843364 Entrez UniGene	2060396	pSport1
125	4316	-1.5	**-1.8**	1114	9.6	100	2040	1714	23.9	100	E	4	0213AGMM	transporter 1, ATP-binding cassette, sub-family B (MDR/TAP)	Passed	NM_000593 Entrez UniGene	1634279	pINCY

126	488	-1.5	**-1.8**	641	5.7	68	1157	972	12.9	68	C	3	0219AGL5	TNF receptor-associated factor 2	Passed	BG677004 Entrez UniGene	3226017	pINCY
127	1926	-1.5	**-1.8**	369	4.2	85	655	550	10.6	85	A	11	021GAGMT	leptin (murine obesity homolog)	Passed	NM_000230 Entrez UniGene	1961822	pSport1
128	1273	-1.5	**-1.8**	387	3.9	93	679	571	10.2	93	A	1	021UAGM2	grancalcin, EF-hand calcium-binding protein	Passed	BG545676 Entrez UniGene	1671852	pINCY
129	1177	-1.5	**-1.8**	406	4.1	85	735	618	10.1	85	A	1	0219AGLY	early growth response 1	Passed	AL553329 Entrez UniGene	1705208	pINCY
130	5138	-1.5	**-1.8**	453	3.7	62	815	685	8.4	62	F	3	021RAGKM	ribosomal protein S6 kinase, 90kD, polypeptide 1	Passed	NM_002953 Entrez UniGene	1822236	pINCY
131	8359	-1.5	**-1.8**	462	4.3	88	833	700	9.2	88	F	2	021DAGLE	protein kinase, cAMP-dependent, regulatory, type I, beta	Passed	BF940317 Entrez UniGene	1926688	pSport1
132	7799	-1.5	**-1.8**	519	4.5	50	925	777	10.4	50	B	10	021QAGKR	spermine synthase	Passed	BG535109 Entrez UniGene	2061432	pSport1

133	6370	-1.5	**-1.8**	1054	10.6	100	1853	1557	27.0	100	H	7	021NAGM1	SEC14 (S. cerevisiae)-like 1	Passed	NM_003003 Entrez UniGene	1879956	pINCY
134	6529	-1.5	**-1.8**	540	5.6	100	988	830	14.2	100	F	1	0210AGM8	solute carrier family 29 (nucleoside transporters), member 1	Passed	BI088219 Entrez UniGene	2594080	pINCY
135	9461	-1.5	**-1.8**	135	2.1	88†	244	205	3.8	88	D	10	021HAGMO	cathepsin H	Passed	BC002479 Entrez UniGene	1749417	pINCY
136	4361	-1.5	**-1.8**	125	1.9	83†	223	187	3.4	83	C	10	021HAGMO	nuclear receptor subfamily 4, group A, member 1	Passed	NM_002135 Entrez UniGene	1958560	pINCY
137	6375	-1.5	**-1.8**	921	8.7	100	1695	1424	23.3	100	B	5	021UAGM2	v-rel avian reticuloendotheliosis viral oncogene homolog B (nuclear factor of kappa light polypeptide gene enhancer in B-cells 3)	Passed	BG748809 Entrez UniGene	1859449	pINCY
138	10180	-1.6	**-1.8**	845	9.0	62	1561	1312	24.9	62	D	8	021MAKNL	Control: Sensitivity 2000pg				
139	3168	-1.5	**-1.8**	300	3.0	87	546	459	6.4	87	E	12	021LAGLA	DNA segment on chromosome X (unique) 9879 expressed sequence	Passed	AI360458 Entrez UniGene	2585358	pINCY

140	3805	-1.5	**-1.8**	624	6.4	66	1147	964	17.1	66	C	2	021NAGM1	tetracycline transporter-like protein	Passed	L11669 Entrez UniGene	1856947	pINCY
141	1	-1.5	**-1.8**	547	3.5	55	1004	844	10.3	55	A	1	0216AKON	Internal_Control_E				
142	8522	-1.5	**-1.8**	109	2.0	97†	198	166	3.5	97	D	4	021QAGLL	hypothetical protein FLJ23231	Passed	AA640102 Entrez UniGene	2348706	pINCY
143	6701	-1.5	**-1.7**	1063	7.7	65	1858	1561	20.3	65	F	9	021QAGMF	SH2 domain protein 2A	Passed	NM_003975 Entrez UniGene	504273	pBlue
144	7630	-1.5	**-1.7**	1477	14.0	61	2567	2157	36.4	61	D	7	021MAKNL	Control: Sensitivity 2000pg				
145	7083	-1.4	**-1.7**	2448	23.3	86	4100	3445	54.0	86	F	5	021UAGMV	tumor necrosis factor receptor superfamily, member 9	Passed	BG436824 Entrez UniGene	191843	pBlue
146	1102	-1.4	**-1.7**	895	7.1	100	1482	1245	18.8	100	G	7	021HAGLU	nuclear factor of kappa light polypeptide gene enhancer in B-cells inhibitor, epsilon	Passed	BG469491 Entrez UniGene	2748942	pINCY
147	10047	-1.4	**-1.7**	910	8.5	62	1509	1268	20.2	62	H	6	021BAGNC	pericentrin 2 (kendrin)	Passed	AB007862 Entrez UniGene	1655365	pINCY

148	9493	-1.5	**-1.7**	6971	60.6	49	12108	10175	155.2	49	H	2	021OAGMP	x 006 protein	Passed	BF445491 Entrez UniGene	440017	pBlue
149	8042	-1.4	**-1.7**	19343	175.3	51	32332	27170	412.7	51	D	4	021HAGL1	nuclear receptor subfamily 1, group H, member 2	Passed	BE878950 Entrez UniGene	2581075	pINCY
150	3741	-1.4	**-1.7**	909	8.2	100	1509	1268	18.6	100	E	6	0219AGLY	tumor necrosis factor receptor superfamily, member 14 (herpesvirus entry mediator)	Multiple Bands	BC002794 Entrez UniGene	2121653	pINCY
151	9008	-1.4	**-1.7**	2975	26.9	82	4985	4189	66.5	82	F	4	021FAGM5	peroxiredoxin 1	Passed	BG612736 Entrez UniGene	3942594	pINCY
152	9528	-1.4	**-1.7**	345	4.2	92	595	500	9.3	92	B	12	0212AGMR	metallothionein 1L	Passed	BG928106 Entrez UniGene	2513883	pINCY
153	1089	-1.4	**-1.7**	170	2.1	59†	292	245	3.8	59	C	5	021HAGLU	solute carrier family 8 (sodium/ calcium exchanger), member 1	Passed	NM_021097 Entrez UniGene	2880435	pINCY
154	6796	-1.4	**-1.7**	844	7.4	62	1448	1217	17.4	62	F	7	021IAGMJ	cytochrome b-245, alpha polypeptide	Passed	BG751568 Entrez UniGene	4173205	pINCY

155	2932	-1.5	**-1.7**	757	7.3	100	1315	1105	16.8	100	G	8	021AAGL0	macrophage myristoylated alanine-rich C kinase substrate	Passed	AL534743 Entrez UniGene	2135733	pINCY
156	4287	-1.4	**-1.7**	718	7.4	100	1225	1029	17.1	100	C	6	021WAGML	tubulin, beta, 5	Passed	AL536237 Entrez UniGene	1486358	pINCY
157	1774	-1.4	**-1.7**	554	4.7	100	938	788	10.7	100	G	7	0213AGMM	growth arrest and DNA-damage-inducible, alpha	Passed	AI935984 Entrez UniGene	1702350	pINCY
158	5390	-1.4	**-1.7**	432	4.3	75	745	626	10.2	75	B	3	021WAGKX	modulator recognition factor I	Passed	M62324 Entrez UniGene	1568724	pINCY
159	117	-1.4	**-1.7**	3127	24.0	53	5178	4351	65.0	53	G	5	021CAGKP	LIM domain kinase 1	Passed	NM_002314 Entrez UniGene	3373632	pINCY
160	8372	-1.4	**-1.7**	368	4.1	92	616	518	8.8	92	B	4	021KAGLF	keratin 7	Passed	BI094014 Entrez UniGene	1962141	pSport1

161	6631	-1.4	**-1.7**	319	3.7	93	528	444	8.4	93	H	1	0215AGMC	uridine monophosphate synthetase (orotate phosphoribosyl transferase and orotidine-5'-decarboxylase)	Passed	AU133065 Entrez UniGene	1816355	pINCY
162	1887	-1.4	**-1.7**	295	3.0	73	502	422	6.4	73	E	5	0212AGMR	replication protein A3 (14kD)	Passed	BF030135 Entrez UniGene	2451279	pINCY
163	8993	-1.4	**-1.7**	264	3.3	87	449	377	7.7	87	H	10	0218AGM4	KIAA0005 gene product	Passed	NM_014670 Entrez UniGene	2059208	pSport1
164	9799	-1.4	**-1.7**	259	2.8	68	430	361	5.8	68	F	2	0210AGN2	methylenetetrahydrofolate dehydrogenase (NADP+ dependent), methenyltetrahydrofolate cyclohydrolase, formyltetrahydrofolate synthetase	Passed	BC001014 Entrez UniGene	1297179	pINCY
165	8262	-1.4	**-1.7**	226	2.7	82	383	322	5.4	82	D	12	021LAGLA	neural polypyrimidine tract binding protein	Passed	BG991819 Entrez UniGene	2578983	pINCY
166	8033	-1.5	**-1.7**	199	2.6	90	346	291	4.9	90	H	10	021AAGL0	retinoid X receptor, alpha	Passed	X52773 Entrez UniGene	2808269	pINCY

167	4534	-1.5	**-1.7**	151	1.9	76†	261	219	3.5	76	E	8	021UAGMV	E2F transcription factor 1	Passed	AL121906 Entrez UniGene	1920739	pSport1
168	5991	-1.4	**-1.7**	148	2.1	72†	253	213	3.7	72	B	5	021XAGLM	HTPAP protein	Passed	AI765129 Entrez UniGene	2626340	pINCY
169	3731	-1.4	**-1.6**	174	2.5	98	282	237	5.0	98	A	10	0219AGLY	annexin A8	Passed	BC008813 Entrez UniGene	1911622	pINCY
170	3280	-1.4	**-1.6**	165	2.4	92†	269	226	4.2	92	C	8	021KAGLF	RNA helicase-related protein	Passed	BF679543 Entrez UniGene	2757583	pSport1
171	6994	-1.3	**-1.6**	230	2.5	75	361	303	4.8	75	H	7	0212AGMR	vav 2 oncogene	Passed	AL045952 Entrez UniGene	3744592	pINCY
172	9458	-1.3	**-1.6**	280	3.1	90	445	374	5.7	90	D	4	021HAGMO	serine/ threonine kinase 4	Passed	NM_006282 Entrez UniGene	1510581	pINCY
173	2358	-1.4	**-1.6**	164	2.0	65†	268	225	3.4	65	A	11	0214AGNB	RAB9, member RAS oncogene family	Passed	BG535930 Entrez UniGene	1746329	pINCY

174	7474	-1.4	**-1.6**	335	3.9	86	543	456	8.5	86	H	7	0214AGNB	aldo-keto reductase family 1, member A1 (aldehyde reductase)	Passed	AW873466 Entrez UniGene	1634342	pINCY
175	2349	-1.3	**-1.6**	174	2.1	59†	274	230	3.8	59	G	5	021XAGNA	biliverdin reductase A	Passed	AI765830 Entrez UniGene	775330	pSport1
176	725	-1.4	**-1.6**	875	7.6	100	1443	1213	19.4	100	A	9	021KAGLF	chemokine (C-C motif) receptor 6	Passed	AL121935 Entrez UniGene	3190228	pSport1
177	7445	-1.3	**-1.6**	162	2.1	59†	256	215	3.4	59	F	9	021XAGNA	thymopoietin	Passed	BF983366 Entrez UniGene	56919	pBlue
178	4665	-1.4	**-1.6**	480	5.1	66	775	651	9.8	66	A	6	021TAGN1	insulin-like growth factor 2 (somatomedin A)	Passed	BG619049 Entrez UniGene	2955178	pINCY
179	1549	-1.3	**-1.6**	496	4.2	58	771	648	7.8	58	E	1	021CAGMD	ADP-ribosylation factor 4	Passed	BG529241 Entrez UniGene	65463	pBlue
180	6748	-1.3	**-1.6**	130	2.1	75†	207	174	3.5	75	F	7	0214AGMH	exonuclease 1	Passed	AC004783 Entrez UniGene	4385292	pINCY

181	7700	-1.4	**-1.6**	1003	7.7	58	1642	1380	19.4	58	B	4	021YAGKN	tumor necrosis factor (ligand) superfamily, member 7	Passed	BG177086 Entrez UniGene	2017463	pINCY
182	4678	-1.3	**-1.6**	528	5.5	78	833	700	11.0	78	E	8	021TAGN1	proteasome (prosome, macropain) subunit, beta type, 6	Passed	BF205294 Entrez UniGene	2989852	pINCY
183	4526	-1.4	**-1.6**	437	4.4	72	709	596	10.5	72	C	4	021UAGMV	benzodiazapine receptor (peripheral)	Passed	BE531172 Entrez UniGene	1241680	pSport1
184	4806	-1.3	**-1.6**	214	2.3	56†	342	287	4.2	56	G	12	021SAGN6	cAMP responsive element modulator	Passed	AL117336 Entrez UniGene	1921290	pSport1
185	2828	-1.4	**-1.6**	537	5.4	100	872	733	12.1	100	E	4	021PAGKW	intracellular hyaluronan-binding protein	Passed	AK025144 Entrez UniGene	1755234	pINCY
186	9466	-1.3	**-1.6**	601	6.5	100	933	784	13.4	100	F	8	021HAGMO	proteasome (prosome, macropain) subunit, alpha type, 4	Passed	BG178069 Entrez UniGene	1975642	pINCY
187	5163	-1.4	**-1.6**	1116	11.1	57	1836	1543	29.1	57	F	5	021YAGKN	general transcription factor IIA, 2 (12kD subunit)	Multiple Bands	BG431140 Entrez UniGene	2457833	pINCY

188	4999	-1.3	**-1.6**	647	6.5	89	1019	856	15.5	89	A	2	021WAGNF	cellular retinoic acid-binding protein 1	Passed	BG715841 Entrez UniGene	585432	pSport1
189	9885	-1.3	**-1.6**	1381	13.9	91	2153	1809	32.7	91	B	6	021SAGN6	transcriptional coactivator	Passed	AI088790 Entrez UniGene	1842227	pSport1
190	6897	-1.3	**-1.6**	1433	11.8	64	2280	1916	27.6	64	H	5	021AAGMN	Homo sapiens, Similar to src homology three (SH3) and cysteine rich domain, clone MGC:2793 IMAGE:2961089, mRNA, complete cds	Passed	AW410140 Entrez UniGene	2256026	pSport1
191	2434	-1.3	**-1.6**	653	5.1	64	1016	854	9.7	64	C	7	021PAGNE	KIAA0101 gene product	Passed	BG560803 Entrez UniGene	2458926	pINCY
192	3039	-1.3	**-1.6**	672	6.3	100	1066	896	12.9	100	C	6	0219AGL5	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily a, member 2	Passed	R56503 Entrez UniGene	3660322	pINCY
193	2824	-1.3	**-1.6**	8903	70.6	89	14242	11968	148.8	89	C	8	021PAGKW	protein kinase, cAMP-dependent, regulatory, type I, alpha (tissue specific extinguisher 1)	Passed	BE257854 Entrez UniGene	1627426	pINCY

194	5756	-1.4	**-1.6**	1533	12.2	100	2513	2112	28.8	100	D	3	021ZAGLC	immediate early protein	Passed	BG255669 Entrez UniGene	1217963	pSport1
195	1764	-1.3	**-1.6**	712	6.4	100	1136	955	14.1	100	C	11	0213AGMM	Sjogren's syndrome nuclear autoantigen 1	Passed	BC004118 Entrez UniGene	1710701	pINCY
196	2614	-1.3	**-1.6**	759	7.8	66	1182	993	16.8	66	E	8	021YAGKN	complement component 1, r subcomponent	Passed	M14058 Entrez UniGene	1664320	pINCY
197	373	-1.3	**-1.6**	802	7.0	83	1288	1082	18.7	83	E	1	021AAGL0	general transcription factor IIH, polypeptide 4 (52kD subunit)	Passed	AW401633 Entrez UniGene	2470646	pINCY
198	1591	-1.3	**-1.6**	464	4.1	70	734	617	8.0	70	C	1	021QAGMF	lymphocyte-activation gene 3	Passed	NM_002286 Entrez UniGene	684538	pSport1
199	9555	-1.3	**-1.6**	903	8.1	100	1415	1189	16.6	100	D	6	0219AGMS	ATX1 (antioxidant protein 1, yeast) homolog 1	No Amplification	AI276280 Entrez UniGene	2313349	pSport1
200	7470	-1.4	**-1.6**	3269	30.2	93	5365	4508	76.1	93	F	11	0214AGNB	chemokine (C-C motif) receptor 7	Passed	L08176 Entrez UniGene	2652665	pINCY

† Probe 1 did not meet selection criteria

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Previous 100 Next 100

1 **2** 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 19 20 21 22 23 24 25 26 27 28 29 30 31 32 33 34 35 36 37 38 39 40 41 42 43 44 45 46 47 48 49 50 51 52 53 54 55 56 57 58 59 60 61 62 63 64 65 66 67 68 69 70 71 72 73 74 75 76 77 78 79 80 81 82 83 84 85 86 87 88 89 90 91 92 93 94

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