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 **201** to **300**

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	Gen Bank Id	Clone Id (Sequence)	Vector
201	4691	-1.3	**-1.6**	1605	13.3	61	2499	2100	28.3	61	A	10	0210AGN2	tropomyosin 2 (beta)	Passed	AA283746 Entrez UniGene	3176845	pINCY
202	3208	-1.4	**-1.6**	1977	14.1	53	3262	2741	29.3	53	C	8	021ZAGLC	minichromosome maintenance deficient (S. cerevisiae) 7	Passed	BG683250 Entrez UniGene	986752	pSport1

203	4196	-1.3	**-1.6**	2088	16.8	50	3243	2725	39.9	50	E	4	0214AGMH	neuropeptide Y	Passed	K01911 Entrez UniGene	2494284	pINCY
204	7009	-1.3	**-1.6**	3116	25.3	86	4963	4171	59.7	86	F	1	0219AGMS	CD44 antigen (homing function and Indian blood group system)	Passed	BG822701 Entrez UniGene	549196	pSport1
205	542	-1.3	**-1.6**	2439	17.9	50	3901	3278	40.5	50	E	3	021NAGL7	FK506-binding protein 1A (12kD)	Passed	AL539248 Entrez UniGene	2204204	pINCY
206	4094	-1.3	**-1.6**	2284	18.8	93	3628	3049	42.2	93	C	4	021CAGMD	dual specificity phosphatase 4	Passed	NM_001394 Entrez UniGene	740878	pSport1
207	6084	-1.4	**-1.6**	126	2.0	86†	205	172	3.2	86	H	11	021IAGLP	sperm associated antigen 1	Passed	NM_003114 Entrez UniGene	2962332	pINCY
208	4080	-1.3	**-1.5**	192	2.5	93	287	241	4.4	93	E	12	0215AGMC	uridine phosphorylase	Passed	BG492119 Entrez UniGene	1806435	pINCY
209	2448	-1.3	**-1.5**	118	1.9	65†	179	150	2.8	65	G	11	021PAGNE	KIAA0222 gene product	Passed	AW510659 Entrez UniGene	1527755	pINCY

210	1535	-1.2	**-1.5**	240	2.5	74	355	298	4.8	74	G	9	0215AGMC	cytochrome P450, subfamily IIC (mephenytoin 4-hydroxylase), polypeptide 18	Passed	M61853 Entrez UniGene	2595728	pINCY
211	289	-1.3	**-1.5**	256	2.6	71	383	322	5.4	71	A	1	021WAGKX	glutamine-fructose-6-phosphate transaminase 2	Passed	AK001242 Entrez UniGene	1904696	pINCY
212	679	-1.2	**-1.5**	259	2.8	88	383	322	5.5	88	C	1	0216AGLD	B-cell CLL/lymphoma 6 (zinc finger protein 51)	Passed	NM_001706 Entrez UniGene	1920672	pSport1
213	8351	-1.2	**-1.5**	288	3.0	73	426	358	5.7	73	B	10	021DAGLE	B-cell CLL/lymphoma 1	Passed	Z23022 Entrez UniGene	1994739	pSport1
214	2949	-1.2	**-1.5**	290	3.1	77	424	356	5.9	77	E	6	021HAGL1	paraoxonase 2	Passed	BG699928 Entrez UniGene	2134968	pINCY
215	2492	-1.3	**-1.5**	295	3.1	58	439	369	5.4	58	G	3	0213AGNG	Incyte EST	Passed		87242	pBlue
216	9552	-1.3	**-1.5**	300	3.3	81	462	388	6.1	81	B	12	0219AGMS	gamma-glutamyl hydrolase (conjugase, folylpolygammaglutamyl hydrolase)	Passed	BG717516 Entrez UniGene	1997967	pSport1

217	7716	-1.2	**-1.5**	301	3.3	73	441	371	6.9	73	F	12	021YAGKN	aldo-keto reductase family 1, member B1 (aldose reductase)	Passed	AV703256 Entrez UniGene	1901073	pINCY
218	2495	-1.3	**-1.5**	327	3.1	49	496	417	5.3	49	G	9	0213AGNG	ubiquitin carrier protein	Passed	AI571293 Entrez UniGene	2057823	pSport1
219	5872	-1.3	**-1.5**	334	3.8	85	516	434	8.0	85	B	7	021YAGLH	CDK2-associated protein 1	Passed	NM_004642 Entrez UniGene	902381	pSport1
220	1044	-1.2	**-1.5**	347	3.2	71	505	424	5.9	71	C	11	0213AGLS	early growth response 3	Passed	NM_004430 Entrez UniGene	2633001	pINCY
221	2540	-1.3	**-1.5**	360	3.8	70	537	451	7.8	70	G	3	021MAKNL	Control: Complex Target (Homo sapiens)				
222	5682	-1.3	**-1.5**	380	4.1	76	577	485	8.6	76	B	11	0211AGL9	calcium and integrin binding protein (DNA-dependent protein kinase interacting protein)	Passed	BF685744 Entrez UniGene	4626895	pINCY

223	2082	-1.2	**-1.5**	394	4.2	82	578	486	8.8	82	E	11	021MAGMZ	ZW10 interactor	Passed	AW409765 Entrez UniGene	1576329	pINCY
224	10178	-1.3	**-1.5**	430	5.2	96	644	541	10.3	96	D	4	021MAKNL	Control: Sensitivity 20pg				
225	455	-1.3	**-1.5**	436	4.2	75	657	552	9.5	75	G	9	021VAGL3	myotubularin related protein 4	No Amplification	BG012919 Entrez UniGene	2890336	pINCY
226	10179	-1.3	**-1.5**	438	5.1	91	652	548	10.4	91	D	6	021MAKNL	Control: Sensitivity 200pg				
227	3097	-1.3	**-1.5**	489	4.8	68	732	615	10.5	68	G	2	021NAGL7	agrin	Passed	AF016903 Entrez UniGene	4610962	pINCY
228	9607	-1.3	**-1.5**	506	4.6	72	763	641	8.1	72	F	2	021NAGMU	replication protein A2 (32kD)	Passed	BG333934 Entrez UniGene	1729876	pINCY
229	6538	-1.2	**-1.5**	537	5.5	97	784	659	10.8	97	H	7	0210AGM8	KIAA0399 protein	Passed	AB007859 Entrez UniGene	2347845	pINCY
230	2529	-1.2	**-1.5**	568	5.3	84	844	709	11.6	84	C	5	021MAKNL	Control: Sensitivity 200pg				

231	6166	-1.3	**-1.5**	613	6.0	100	939	789	13.0	100	D	7	021AAGLT	cell division cycle 2, G1 to S and G2 to M	Passed	BF968871 Entrez UniGene	1525795	pINCY
232	1046	-1.2	**-1.5**	661	6.0	71	979	823	12.0	71	E	3	0213AGLS	glucosidase I	Passed	NM_006302 Entrez UniGene	1555701	pINCY
233	2346	-1.3	**-1.5**	688	5.2	58	1038	872	10.7	58	E	11	021XAGNA	HS1 binding protein	Passed	AL523082 Entrez UniGene	155194	pBlue
234	4669	-1.3	**-1.5**	743	6.9	65	1109	932	12.2	65	C	2	021TAGN1	pim-1 oncogene	Passed	NM_002648 Entrez UniGene	2679117	pINCY
235	7892	-1.3	**-1.5**	746	5.8	58	1135	954	11.4	58	B	4	021IAGKV	decidual protein induced by progesterone	Passed	AK000219 1 Entrez UniGene	1960889	pSport1
236	341	-1.2	**-1.5**	773	7.0	77	1128	948	14.0	77	A	9	021AAGKZ	ras homolog gene family, member	Passed	AI167227 Entrez UniGene	1910324	pINCY
237	6355	-1.3	**-1.5**	779	7.0	100	1180	992	16.3	100	D	1	021NAGM1	interleukin 4 receptor	Multiple Bands	NM_000418 Entrez UniGene	1808529	pINCY

238	4633	-1.3	**-1.5**	799	8.9	67	1201	1009	17.5	67	G	2	021MAGMZ	SWI/SNF related, matrix associated, actin dependent regulator of chromatin, subfamily d, member 1	Multiple Bands	BC009368 Entrez UniGene	1658083	pINCY
239	7021	-1.3	**-1.5**	898	8.2	100	1382	1161	17.1	100	B	1	021GAGMT	protein translocation complex beta	Passed	BG029603 Entrez UniGene	1966933	pSport1
240	8845	-1.3	**-1.5**	952	9.4	65	1454	1222	21.7	65	H	2	0219AGLY	small nuclear ribonucleoprotein polypeptide G	Passed	AV766263 Entrez UniGene	2449837	pINCY
241	2530	-1.3	**-1.5**	975	8.7	68	1454	1222	19.8	68	C	7	021MAKNL	Control: Sensitivity 2000pg				
242	582	-1.3	**-1.5**	978	8.2	100	1468	1234	16.2	100	A	11	0211AGL9	hypocretin (orexin) neuropeptide precursor	Passed	BG194545 Entrez UniGene	4588486	pINCY
243	1092	-1.3	**-1.5**	1020	7.9	62	1568	1318	17.9	62	C	11	021HAGLU	autoimmune regulator (autoimmune polyendocrinopathy candidiasis ectodermal dystrophy)	Passed	AJ009610 Entrez UniGene	2914128	pINCY

244	5028	-1.2	**-1.5**	1056	10.1	67	1566	1316	20.4	67	A	12	0213AGNG	acyl-Coenzyme A oxidase 2, branched chain	Passed	BG545036 Entrez UniGene	943744	pSport1
245	6619	-1.2	**-1.5**	1081	9.3	100	1595	1340	19.1	100	D	1	0215AGMC	activity-regulated cytoskeleton-associated protein	Passed	AF193421 Entrez UniGene	1382579	pINCY
246	3638	-1.2	**-1.5**	1135	9.6	100	1654	1390	20.6	100	C	4	021HAGLU	phosphorylase, glycogen; muscle (McArdle syndrome, glycogen storage disease type V)	Passed	NM_005609 Entrez UniGene	2635943	pINCY
247	10107	-1.2	**-1.5**	1204	9.0	62	1752	1472	16.8	62	D	6	021WAGNF	Nef-associated factor 1	Passed	NM_006058 Entrez UniGene	1985586	pSport1
248	792	-1.3	**-1.5**	1212	8.7	59	1830	1538	18.3	59	G	11	021YAGLH	interferon-induced, hepatitis C-associated microtubular aggregate protein (44kD)	Passed	NM_006417 Entrez UniGene	1922658	pSport1
249	2963	-1.2	**-1.5**	1216	9.9	100	1799	1512	20.6	100	A	10	021OAGL2	thyroid hormone receptor interactor 10	Passed	NM_004240 Entrez UniGene	2814551	pINCY

250	5597	-1.2	**-1.5**	1382	12.4	59	2043	1717	30.1	59	F	9	0219AGL5	thyroid hormone receptor interactor 10	Passed	NM_004240 Entrez UniGene	1655192	pINCY
251	3646	-1.2	**-1.5**	1389	11.8	72	2043	1717	25.2	72	E	8	021HAGLU	non-metastatic cells 3, protein expressed in	Passed	BC000250 Entrez UniGene	2455170	pINCY
252	2069	-1.3	**-1.5**	1439	13.1	65	2150	1807	31.1	65	A	9	021MAGMZ	cell membrane glycoprotein, 110000M(r) (surface antigen)	Passed	BF033463 Entrez UniGene	1720946	pINCY
253	5122	-1.2	**-1.5**	1459	10.5	51	2134	1793	24.6	51	H	7	0216AKON	Internal_Control_Z				
254	6682	-1.2	**-1.5**	1459	12.6	100	2123	1784	28.0	100	H	7	021JAGME	minichromosome maintenance deficient (S. cerevisiae) 5 (cell division cycle 46)	Passed	AU149507 Entrez UniGene	1746529	pINCY
255	72	-1.2	**-1.5**	1573	12.2	75	2284	1919	25.3	75	G	11	021YAGKN	ferredoxin reductase	Passed	NM_004110 Entrez UniGene	1819763	pINCY
256	2355	-1.2	**-1.5**	1725	12.3	62	2541	2135	27.4	62	A	5	0214AGNB	HMT1 (hnRNP methyltransferase, S. cerevisiae)-like 2	Passed	BG167159 Entrez UniGene	2888814	pINCY

257	9137	-1.3	**-1.5**	1744	17.0	100	2616	2198	35.3	100	H	10	021RAGMA	calcium channel, voltage-dependent, gamma subunit 1	Passed	NM_000727 Entrez UniGene	4014318	pINCY
258	5782	-1.3	**-1.5**	1871	15.7	65	2783	2339	34.9	65	D	7	0216AGLD	cyclin D1 (PRAD1: parathyroid adenomatosis 1)	Passed	X59798 Entrez UniGene	2057653	pSport1
259	3808	-1.3	**-1.5**	1938	17.3	62	2936	2467	33.5	62	C	8	021NAGM1	cathepsin C	Passed	NM_001814 Entrez UniGene	1822716	pINCY
260	1034	-1.3	**-1.5**	1939	18.3	62	2992	2514	43.6	62	A	3	0213AGLS	ras homolog gene family, member G (rho G)	Passed	BG338917 Entrez UniGene	1342744	pINCY
261	5600	-1.3	**-1.5**	1996	14.6	45	3052	2565	31.9	45	H	3	0219AGL5	delta-like homolog (Drosophila)	Passed	BF969929 Entrez UniGene	3288394	pINCY
262	7388	-1.3	**-1.5**	2222	18.6	49	3397	2855	38.6	49	D	3	0216AGN8	protein tyrosine phosphatase, receptor type, U	Passed	AL049570 Entrez UniGene	2941607	pINCY
263	2862	-1.3	**-1.5**	2523	21.5	59	3763	3162	46.8	59	G	12	021WAGKX	selenophosphate synthetase 2	Passed	BG674956 Entrez UniGene	1687542	pINCY

264	7993	-1.2	**-1.5**	2648	24.9	47	3892	3271	54.6	47	D	2	021AAGKZ	amphiregulin (schwannoma-derived growth factor)	Passed	AL546917 Entrez UniGene	2350594	pINCY
265	2486	-1.3	**-1.5**	3108	28.3	56	4658	3914	64.1	56	E	3	0213AGNG	neurogranin (protein kinase C substrate, RC3)	Multiple Bands	AW117600 Entrez UniGene	1943863	pBlue
266	7741	-1.3	**-1.5**	3676	29.5	47	5535	4651	59.1	47	H	2	0215AGKO	hypothetical protein FLJ20030	Passed	AI589003 Entrez UniGene	3027978	pINCY
267	2009	-1.2	**-1.5**	4770	37.1	98	7003	5885	86.3	98	E	9	0211AGMW	calreticulin	Passed	BG675100 Entrez UniGene	2970280	pINCY
268	113	-1.2	**-1.5**	4897	30.3	62	7233	6078	68.5	62	E	9	021CAGKP	double C2-like domains, alpha	Passed	BE646046 Entrez UniGene	1797982	pINCY
269	3041	-1.3	**-1.5**	5768	47.9	59	8870	7454	104.5	59	C	10	0219AGL5	calcitonin/ calcitonin-related polypeptide, alpha	Passed	X03662 Entrez UniGene	2498725	pINCY
270	4860	-1.2	**-1.5**	151	2.1	79†	223	187	3.3	79	A	12	021DAGN9	SH3-domain binding protein 1	Passed	AL157480 Entrez UniGene	2132470	pSport1

271	89	-1.1	**-1.4**	30400	202.3	50	41185	34609	444.7	50	E	9	0215AGKO	actin, gamma 1	Passed	BG422944 Entrez UniGene	3225977	pINCY
272	1030	-1.2	**-1.4**	217	2.6	95	306	257	5.0	95	G	7	021WAGLR	hypothetical protein MGC2495	Passed	AA972572 Entrez UniGene	958633	pSport1
273	4504	-1.2	**-1.4**	7057	47.7	48	10036	8434	90.7	48	C	8	021NAGMU	lysosomal-associated membrane protein 2	Passed	BG719002 Entrez UniGene	2503017	pINCY
274	9289	-1.1	**-1.4**	6364	49.2	52	8644	7264	104.8	52	D	2	0214AGMH	major histocompatibility complex, class I, B	Multiple Bands	BG754309 Entrez UniGene	2859033	pINCY
275	3018	-1.2	**-1.4**	4521	34.2	92	6248	5250	63.5	92	C	12	0212AGL4	vinexin beta (SH3-containing adaptor molecule-1)	Passed	BG252405 Entrez UniGene	1957809	pINCY
276	28	-1.2	**-1.4**	3815	22.7	49	5424	4558	50.0	49	A	7	021RAGKM	tubulin, alpha 1 (testis specific)	Passed	BE742772 Entrez UniGene	1872936	pINCY
277	8014	-1.2	**-1.4**	3581	32.4	52	5042	4237	67.2	52	B	8	021AAGL0	mitogen-activated protein kinase kinase kinase 11	Passed	NM_002419 Entrez UniGene	1992626	pINCY

278	1689	-1.1	**-1.4**	3433	24.0	49	4676	3929	51.4	49	C	5	021IAGMJ	tyrosine 3-monooxygenase/ tryptophan 5-monooxygenase activation protein, theta polypeptide	Passed	NM_006826 Entrez UniGene	4169223	pINCY
279	2989	-1.1	**-1.4**	3360	29.0	89	4595	3861	60.4	89	C	2	021VAGL3	Homo sapiens clone 23608 mRNA sequence	Passed	BC008032 Entrez UniGene	2642865	pINCY
280	7756	-1.2	**-1.4**	2910	25.7	45	4130	3471	53.6	45	D	8	021CAGKP	KIAA0544 protein	Passed	BF801696 Entrez UniGene	4174315	pINCY
281	69	-1.2	**-1.4**	2745	19.4	52	3826	3215	40.2	52	G	5	021YAGKN	splicing factor, arginine/ serine-rich 3	Passed	BG287081 Entrez UniGene	2474214	pINCY
282	9609	-1.2	**-1.4**	2629	23.7	49	3739	3142	45.9	49	F	6	021NAGMU	ets variant gene 2	Passed	AF000671 Entrez UniGene	1320685	pINCY
283	7037	-1.1	**-1.4**	2590	22.8	64	3525	2962	48.0	64	F	9	021GAGMT	RAN, member RAS oncogene family	Passed	BG111605 Entrez UniGene	552654	pSport1

284	8324	-1.1	**-1.4**	2533	24.7	58	3452	2901	46.3	58	B	4	0216AGLD	vesicle-associated membrane protein 3 (cellubrevin)	Passed	BG400142 3 Entrez UniGene	1988078	pSport1
285	393	-1.1	**-1.4**	2244	13.7	44	3062	2573	25.7	44	C	5	021HAGL1	ATPase, H+ transporting, lysosomal (vacuolar proton pump), member D	Passed	BF311825 Entrez UniGene	2923189	pINCY
286	7629	-1.1	**-1.4**	2022	17.2	44	2755	2315	34.1	44	D	5	021MAKNL	Control: Sensitivity 200pg				
287	2636	-1.2	**-1.4**	2002	15.9	49	2741	2303	29.8	49	E	4	0215AGKO	H2B histone family, member Q	Passed	BF794197 Entrez UniGene	3040858	pINCY
288	3822	-1.1	**-1.4**	210	2.8	97	287	241	4.6	97	G	12	021NAGM1	M-phase phosphoprotein 6	Passed	BI088194 Entrez UniGene	1806540	pINCY
289	2121	-1.2	**-1.4**	1879	14.0	64	2628	2208	29.7	64	C	5	021TAGN1	SH3-domain GRB2-like 1	Passed	BE563866 Entrez UniGene	2346978	pINCY
290	2835	-1.2	**-1.4**	1589	13.7	70	2225	1870	28.9	70	G	6	021PAGKW	plexin B2	Passed	BC004542 Entrez UniGene	1831287	pINCY

291	903	-1.2	**-1.4**	1430	10.2	93	2028	1704	24.7	93	E	5	021XAGLM	COBW-like protein	Passed	BG700904 Entrez UniGene	2220048	pINCY
292	2153	-1.2	**-1.4**	1401	10.5	67	2004	1684	25.1	67	E	9	0210AGN2	glutathione peroxidase 1	Passed	BG764734 Entrez UniGene	3137033	pINCY
293	4983	-1.1	**-1.4**	1350	12.2	62	1846	1551	23.5	62	C	6	021PAGNE	splicing factor, arginine/serine-rich (transformer 2 Drosophila homolog) 10	Passed	AL560128 Entrez UniGene	2457759	pINCY
294	2298	-1.2	**-1.4**	1343	8.9	48	1868	1570	18.4	48	E	11	0216AGN8	Finkel-Biskis-Reilly murine sarcoma virus (FBR-MuSV) ubiquitously expressed (fox derived); ribosomal protein S30	Passed	AA316067 Entrez UniGene	2734906	pINCY
295	1532	-1.2	**-1.4**	1275	12.9	73	1756	1476	26.4	73	G	3	0215AGMC	Homo sapiens, clone MGC:18203 IMAGE:4155840, mRNA, complete cds	Passed	AW248283 Entrez UniGene	2135596	pINCY
296	1866	-1.2	**-1.4**	1264	10.2	100	1743	1465	23.2	100	E	11	021VAGMQ	keratin 7	Passed	AA307373 Entrez UniGene	1649959	pINCY

297	2641	-1.1	**-1.4**	1239	10.4	100	1677	1409	20.3	100	G	2	0215 AGKO	crystallin, alpha B	Passed	BF727296 Entrez UniGene	3016305	pINCY
298	1348	-1.2	**-1.4**	1185	9.5	66	1627	1367	20.0	66	A	7	021F AGM5	albumin	Passed	N24732 Entrez UniGene	3354436	pINCY
299	1555	-1.2	**-1.4**	1157	9.2	100	1584	1331	19.7	100	G	1	021C AGMD	serine/ threonine kinase 12	Passed	BG616089 Entrez UniGene	161207	pBlue
300	7956	-1.2	**-1.4**	177	2.3	82 †	245	206	3.5	82	F	12	021 WAGKX	era (E. coli G-protein homolog)-like 1	Passed	BE871890 Entrez UniGene	1686892	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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