

Plate	Position	Catalog No. of Primer	Accession No. of Gene	Symbol
QG085-01	A01	HQP018502	NM_000638	VTN
QG085-01	A02	HQP003299	NM_004385	VCAN
QG085-01	A03	HQP018155	NM_000364	TNNT2
QG085-01	A04	HQP018074	NM_000460	THPO
QG085-01	A05	HQP018068	NM_003246	THBS1
QG085-01	A06	HQP018067	NM_000361	THBD
QG085-01	A07	HQP018038	NM_001032281	TFPI
QG085-01	A08	HQP013260	NM_000934	SERPINF2
QG085-01	A09	HQP012154	NM_000602	SERPINE1
QG085-01	A10	HQP008737	NM_000185	SERPIND1
QG085-01	A11	HQP011620	NM_000488	SERPINC1
QG085-01	A12	HQP013131	NM_004155	SERPINB9
QG085-01	B01	HQP012332	NM_000624	SERPINA5
QG085-01	B02	HQP013122	NM_000295	SERPINA1
QG085-01	B03	HQP016747	NM_003006	SELPLG
QG085-01	B04	HQP022805	NM_004850	ROCK2
QG085-01	B05	HQP015059	NM_000313	PROS1
QG085-01	B06	HQP000652	NM_006404	PROCR
QG085-01	B07	HQP015041	NM_000312	PROC
QG085-01	B08	HQP013257	NM_000301	PLG
QG085-01	B09	HQP013205	NM_001005376	PLAUR
QG085-01	B10	HQP013204	NM_002658	PLAU
QG085-01	B11	HQP013201	NM_000930	PLAT
QG085-01	B12	HQP017176	NM_022788	P2RY12
QG085-01	C01	HQP011547	NM_005957	MTHFR
QG085-01	C02	HQP005585	NM_007351	MMRN1
QG085-01	C03	HQP011256	NM_004530	MMP2
QG085-01	C04	HQP011188	NM_005928	MFGE8
QG085-01	C05	HQP021938	NM_139279	MCFD2
QG085-01	C06	HQP010870	NM_002332	LRP1
QG085-01	C07	HQP010660	NM_005570	LMAN1
QG085-01	C08	HQP010114	NM_000893	KNG1
QG085-01	C09	HQP010104	NM_000892	KLKB1
QG085-01	C10	HQP009826	NM_000888	ITGB6
QG085-01	C11	HQP009818	NM_000212	ITGB3
QG085-01	C12	HQP009795	NM_000419	ITGA2B
QG085-01	D01	HQP009794	NM_002203	ITGA2
QG085-01	D02	HQP009641	NM_000576	IL1B
QG085-01	D03	HQP009083	NM_005347	HSPA5
QG085-01	D04	HQP007336	NM_153456	HS6ST3
QG085-01	D05	HQP005363	NM_153612	HS3ST5

QG085-01	D06	HQP009048	NM_000412	HRG
QG085-01	D07	HQP008908	NM_000457	HNF4A
QG085-01	D08	HQP008800	NM_000601	HGF
QG085-01	D09	HQP007183	NM_002052	GATA4
QG085-01	D10	HQP007125	NM_000820	GAS6
QG085-01	D11	HQP006022	NM_002026	FN1
QG085-01	D12	HQP000999	NM_006682	FGL2
QG085-01	E01	HQP005398	NM_005141	FGB
QG085-01	E02	HQP005396	NM_000508	FGA
QG085-01	E03	HQP005061	NM_000132	F8
QG085-01	E04	HQP005058	NM_000130	F5
QG085-01	E05	HQP005057	NM_001993	F3
QG085-01	E06	HQP021852	NM_003950	F2RL3
QG085-01	E07	HQP005056	NM_004101	F2RL2
QG085-01	E08	HQP005055	NM_005242	F2RL1
QG085-01	E09	HQP005053	NM_001992	F2R
QG085-01	E10	HQP005052	NM_000506	F2
QG085-01	E11	HQP005070	NM_001994	F13B
QG085-01	E12	HQP005069	NM_000129	F13A1
QG085-01	F01	HQP005068	NM_000505	F12
QG085-01	F02	HQP005066	NM_000128	F11
QG085-01	F03	HQP005064	NM_000504	F10
QG085-01	F04	HQP022883	NM_001776	ENTPD1
QG085-01	F05	HQP004689	NM_001972	ELA2
QG085-01	F06	HQP023069	NM_001040034	CD63
QG085-01	F07	HQP022821	NM_000072	CD36
QG085-01	F08	HQP021313	NM_001753	CAV1
QG085-01	F09	HQP019841	NM_004343	CALR
QG085-01	F10	HQP016506	NM_000710	BDKRB1
QG085-01	F11	HQP014302	NM_000706	AVPR1A
QG085-01	F12	HQP011288	NM_001181	ASGR2
QG085-01	G01	HQP008829	NM_001154	ANXA5
QG085-01	G02	HQP022625	NM_004797	ADIPOQ
QG085-01	G03	HQP004081	NM_000789	ACE
QG085-01	G04	HQP008678	NM_000014	A2M
QG085-01	G05	HQP005059	NM_000131	F7
QG085-01	G06	HQP016516	NM_000623	BDKRB2
QG085-01	G07	HQP002836	NM_001872	CPB2
QG085-01	G08	HQP005441	NM_000509	FGG
QG085-01	G09	HQP007377	NM_000821	GGCX
QG085-01	G10	HQP012463	NM_016186	SERPINA10
QG085-01	G11	HQP018130	NM_000062	SERPING1
QG085-01	G12	HQP008811	NM_001528	HGFAC

QG085-01	H01	HGDC		
QG085-01	H02	HGDC		
QG085-01	H03	HQP006940	NM_002046	GAPDH
QG085-01	H04	HQP016381	NM_001101	ACTB
QG085-01	H05	HQP015171	NM_004048	B2M
QG085-01	H06	HQP006171	NM_012423	RPL13A
QG085-01	H07	HQP009026	NM_000194	HPRT1
QG085-01	H08	HQP054253	NR_003286	RN18S1
QG085-01	H09	RT		
QG085-01	H10	RT		
QG085-01	H11	PCR		
QG085-01	H12	PCR		