

Accession #	Chromosome	Name	Comments	Location
AC000016.1	chr 04	2U1	same as AC025090	25,142,143,145k
AC000120.1	chr 07	51		5,8,10,14,15,17,18,19,20,23,25,26k
AC004523.1	chr 19	4F12	missing N-term	3,5,7,8,10,21,22k
AC004597.1	chr 19	4F24P		35,36k
AC004609.1	chr 19	4F9P		13,24,33k
AC004790.1	chr 19	4F9P	continues on AC004609	Ok
AC004791.1	chr 19	4F pseudo	lone C-term exon	36,37k
AC004914.1	chr 07	5A1	middle	21,26,28,30,47,72,108,111k
AC004961.2	chr 07	5A1	C-term	31,33,35,44k
AC004977.1	chr 07?	3A7 frag		106k
AC005020.4	chr 07	3A5		18,22,24,26,31,32,33,34,35,38,45,46,48,50k
AC005020.4	chr 07	3A5 pseudo	exons 2,8,13	2,3,13k
AC005336.1	chr 19	4F11 like	C-term 2 diffs	35,36k
AC005336.1	chr 19	4F2	alt splice exon 2	1,2,8,11,12,14,15,16,17,19k
AC006021.2	chr 07	5A1	N-term	10k
AC007002.2	chr 02	26B1		19,22,31,32,33,34k
AC008104.3	chr 18	39A1	also chr 6? incomplete	42,50,89,92,94,95k
AC008537.2	chr 19	2A6	not in AC008537.3 or .4	17,18,19,20,21,22,104,105k
AC008537.4	chr 19	2G1P	missing 3 exons	82,83,85,89,91k
AC008537.4	chr 19	2A7		66,67,68,69,70,71,72,73k
AC008537.4	chr 19	2F1P	missing N-term	41,44,45,48,49k
AC008537.4	chr 19	2A18P	C-term only	99,100,101k
AC008537.4	chr 19	2T2P		31,32,33,34,35k
AC008537.4	chr 19	2B7P1	missing C-term exon	115,127,130,132,133,135,136k
AC008539.2	chr 05	2B7P3	exon 1 only	3k missing from later versions .3 and .4
AC008962.6	chr 19	2A7PT/2A7PC		2,3,4,5k
AC008962.6	chr 19	2G2P		28,30,33,34,35,37,39,40k
AC008962.6	chr 19	2A13	7 diffs, 5 in exon 8	66,67,68,69,70,71,72,73,74k
AC008962.6	chr 19	2F1 99%	3 diffs	94,98,99,100,101,102,103,105,106k
AC008962.6	chr 19	2T3P		112,113,114,115k
AC009144.4	chr 16	17	really chr 10?	66,67,70,71,73k
AC009229.3	chr 02?	1B1		185,186,189,190k
AC009528.7	chr 04	4V2	same as AC012525	9k
AC009678.4	chr 08	2J2	7 diffs 2 stops	47,51,64,67,69,71,78,79k missing C-term
AC009927.3	chr 08?	7A1		127,128,130,131,160,162,163k
AC009974.6	chr 02	27A1	N-term frameshifted	134,144,146,147,148,149k
AC010192.8	chr 03	8B1	2 diffs	26,27,28k
AC011074.5	chr 02?	1B1		72,76,77k
AC011492.5	chr 19	4F22		19,23,31,32,34,37,38,42,43,44,45k
AC011492.5	chr 19	4F23P		58,123,124,125,130,131,132,134,136k
AC011510.6	chr 19	2S1		96,98,99,100,102,103,104,107,108k
AC011517.5	chr 19	4F11	N-term	1,6,7,8k small dupl. At 6 and 7
AC011537.6	chr 19	4F24P	missing last exon	22,26,29,31,32,33,41,42k
AC011541.5	chr 19	2B7P2	exon 1 only	2k
AC011801.3	chr 20?	24		72k
AC011904.3	chr 07	3A43	complete gene	67,69,71,73,76,77,83,84,85,88,93,96,104k
AC011904.3	chr 07	3A43	alt exon 1	113k
AC011904.3	chr 07	3A43	exon 4 fragment	63k
AC012169.6	chr 15	19	missing 2 parts	111,112,115,116,122,123,128k
AC012525.6	chr 04	4V2	same as AC046186	206,210,214,218k
AC013318.4	chr 02	2C8 66%	pseudogene frag.	22k
AC013593.3	chr 01?	4B1	partial	26k
AC015742.3	chr 20?	24 96%	missing exon 7	20,22,51,52,53,160,161,162,164k
AC016689.2	chr 02	1B1		22,23,26,27k
AC018795.5	chr 11	2R1	missing N-term 1-112	105,110,111,112,113k
AC018804.5	chr 02	4F27P	62% to AC018949	36k
AC018864.4	chr 08	7B1 92%	3 diffs	15k
AC018872.5	chr 02	2C8 70%	same as AC013318	116k
AC018949.4	chr 15	4F25P	New pseudo	95k
AC019337.3	chr 19?	2T2P		47,48,122,125,151,152k
AC020705.4	chr 15	1A1		98,99,100,101k
AC020705.4	chr 15	1A2		139,141,142,143,144k
AC020782.4	chr 08	7A1		42,44,45,47,48,49,50k
AC020950.5	chr 19	4F2	N-term	37,39k
AC020950.5	chr 19	4F11	4 diffs	2,5,7,10,11,12,13,20,21k
AC021155.2	chr 06	2F1 fragment3	3 diffs	162k
AC022028.2	chr 15?	4F25P	same as AC018949	81k
AC022097.3	chr 05	1B1	chr 02?	135,136,140k
AC022650.3		2C9 41%	same as AC023380	0,86,87,88,92,93
AC023172.1	chr 19	2A7PT		37,38,39,40k
AC023172.1	chr 19	2B6		4,16,17,19,22,25,29k
AC023300.3	chr 15?	11A1		15,16,29,30,46k
AC023380.2		2C9 41%	same as AC022650	11k
AC024226.9	chr 05	4F pseudo	34% to 4F3, 4F11	64k
AC025090.2	chr 04	2U1	Hs.172564	76,77,104,105,107,109k
AC025165.11	chr 12q	27B1		67,68,69,70,183,184k
AC025730.6	chr 11	2R1	chr 12? Missing N-term	25,26,27,28,33k
AC025748.2	chr 12	2R1	partial chr 11?	56,183,184k
AC025769.3	chr 05?	2A6 like		30,31,32,65,66,67k
AC025769.3	chr 05?	2F1P	dup of chr 19 AC008537?	36,36,101,102k
AC025769.3	chr 05?	2B7P1	dup of chr 19 AC008537?	61,63,89,92,114k
AC025769.3	chr 05?	2A7	dup of chr 19 AC008537?	169,170,171,172,173,174,175k
AC025769.3	chr 05?	2T2P	dup of chr 19 AC008537?	200,201,202,203k
AC025769.3	chr 05?	2A18P	dup of chr 19 AC008537?	212,213,214k
AC025769.3	chr 05?	2G1P	dup of chr 19 AC008537?	222,224,228,230,231k
AC025807.2	chr 03	51P1	U36926	83,84k
AC026267.1	chr 16	24	really chr 20?	12,13,14,19,20,29k
AC026296.5	chr 20	8A1	partial	29,57,65,66,69,87,90,115k
AC026340.12	chr 12q	27B1	complete gene	72,73,74,75,76k
AC026847.3	chr 20	8A1	complete gene	21,73,74,114,117,119,137,139,143,147k
AC026935.2	chr 01	4X1		0,6,12,37,38,45,111,143,144k
AC026935.2	chr 01	4A20		11,21,45,46,47,48,133,140,161,162,163,175,176k
AC026938.3	chr 01	46	8 diffs+stop	73k
AC027105.3	chr 17	8A1	really chr 20?	C-term exon 6,7k
AC027142.1		27C1	NEW	39,41,43,85,107,108,110k
AC027416.2	chr 02?	27A1	complete gene	25,26,39,40,148,149,150,151k
AC027800.2	chr 18	4F pseudo	85% to 4F25P	81k
AC036165.2	chr 15	4F25P	same as AC018949	68k

AC046186.2	chr 04		4V2	same as AC012525	157,158k
AC053538.4	chr 18		4F pseudo	76% to 4F27P	37k
AC058798.1	chr 19		2S1		6,71,72,74,118,119,122,123,124k
AC058798.1	chr 19		2F1		39,127,131,132,133,134,135,136k
AC058798.1	chr 19		2G2P		40,42,105,106,107,110,112k
AC058798.1	chr 19		2T3P		62,63,64k
AC058798.1	chr 19		2A13		139,141,142,143,144,145k
AC058798.1	chr 19		2A	N-term	157,158,159k
AC060810.1	chr 02		4F pseudo	94% to 4F27P	51,52k
AC064854.2	chr 10		2C9 like	N-term	6,7k
AC068165.3	chr 02		4F pseudo	88% to 4F25P	115k
AC068202.2	chr 01		2J2		6,7,36,37,105,128,135,139,142,144k
AC068522.1	chr 08		7A1		100,119k
AC068644.5	chr 03		2D6-like	50% to 2D6	94,103k
AC068845.3	chr 19		4F3	alt exon 2 at 151k	7,8,148,149,153,154,156,157,159,160k
AC068845.3	chr 19		4F8	complete gene	174,175,179,180,181,183,184,185,187k
AC068845.3	chr 19		4F10P	partial	140,141,143,144k
AC068845.3	chr 19	N-term is predicted	4F12	missing 327-371 off end	10,11,103,104,107,109,136,137k
AC068845.3	chr 19		4F22	complete gene	6,7,29,30,32,68,69,73,80,81,83,84,87k
AC068845.3	chr 19		4F23P	complete gene	43,44,45,115,125,126,128,129k
AC069020.4	chr 10		2C pseudo	57% to 2C18	174,175k
AC069294.4	chr 07		3A4	missing exons 7-9	15,40,41,43,55,57,91,93,94,95,96k
AC069294.4	chr 07		3A5	missing exons 5-13	23,27,28,30k
AC069294.4	chr 07		3A7	missing exon 6	17,19,22,34,36,37,38,39,48,49,50,122,126k
AC069294.4	chr 07		3A43		4,6,100,101,102,105,110,113,151,153,156,157,185,186k
AC069294.4	chr 07		3A fragments	exons 1,2,13	16,62,69,73,141,177k
AC073128.2	chr 02		27A1	complete gene	47,48,49,50,52,53,150k
AC073385.1	chr 08		11B1	partial	4,5,15,16,26(dup of 16),27(dup of 15),32,33,60k
AC073464.2	chr 02	same as AC068165	4F pseudo	88% to 4F25P	66k
AC073888.2	chr 06		39A1	heme fragment	146k
AC073957.3	chr 07	New Human Sequence	2W1	40% to 2F1 complete	19,20,21,67,68,69,71k
AD000091.1	chr 19		4F23P	missing N-term	1,3,4,5,11,12k
AD000685.1	chr 19		4F3		26,30,32,34,35,37,38,43,44k
AD000685.1	chr 19		4F8		0,1,4,5,7,8,9,13,14k
AD000685.1	chr 19		4F10P		49,51,52k
AF215845.2	chr 08		71B1	missing 1-40	92,93,100,101,111,112,120k
AF228661.2	chr 08		7B1	missing 1-86	8,9,20,27,28k
AL021878.1	chr 22		2D6*5	gene deleted	
AL021878.1	chr 22		2D7AP		46,47,48,49,50k
AL021878.1	chr 22		2D8P		55,56,57,58,59,60k
AL033378.12	chr 06	same as AL357567.2	51 pseudo	48% to 51A1	58,59k
AL035670.29	chr 06		39A1	also chr 18	5,9,43,51,80,86,91,93,94k
AL357567.2	chr 06	same as AL033378.12	51 pseudo	48% to 51A1	33,34k
AL359997.4	chr 09		1 pseudo	45% to Xenopus 1A	27,100,106,107,109k
AL390073.3	chr 01		4A11		91k
AL049547.10	chr 06		21A2		83,84,85,86k
AL078614.2	chr 21	really chr 15?	4F25P		59k
AL109748.2	chr 21		4F28P		3,4k
AL133513.5	chr 10		2C9/2C19 hybrid		40,43,49,50,52,67,70,88k
AL133513.5	chr 10	71% to 2C9	2C N-term fra	pseudogene	114k
AL133513.5	chr 10	87% to 2C18	2C18 like	N-term frag	19k
AL135960.1	chr 01		4A11		129,131,132,133,134,135,136,137,140k
AL136000.2	chr 14		46	missing 1-118	0,6,7,15,16,18,21,22,25,26k
AL138805.8	chr 20		24	complete gene	59,60,61,64,65,66,67,71,73,75k
AL138845.6	chr 01	really chr 8?	2J2	missing 335-397	62,69,109,110,113,114,115,116,117,118k
AL138921.9	chr 10		2C pseudo	partial	7,18,37,41,136,182k
AL138924.2	chr 10	not in AL138924.19	2C19	C-term exon	107k
AL139008.10	chr 09		4F26P	77% to AC018949	108k
AL139427.3	chr 01		46 94%	C-term fragments	77k
AL157834.5	chr 10		2C8 99%	2 diffs N-term only	1,7,8,10,176k
AL157835.3	chr 10		2C18	complete gene on 2 contigs	79,80,89,93,107,137,141,148k
AL158168.13	chr 10	not in AL158168.15	2C pseudo frags	same as AL138921	8,16,57,179k
AL161645.5	chr 10		CYP2E1	N-term half only	111,112,115,116,117k
AL161895.7	chr 13		51P2	U40053	102k
AL163204.2	chr 21		4F pseudo	96% to AC027800	205,206k
AL353727.9	chr 10		17	complete gene	142,143,144,145,146,147,148k
AL356289.4	chr 01	46 is on chr 14	46 pseudo	C-term only	28k
AL356489.6	chr 09		4F26P		242,243k
AL356489.6	chr 09		4F pseudo	82% to 4F26P	19k
AL356793.10	chr 01		4B1	missing 143-166	43,44,45,47,48,49,51,63k
AL356793.10	chr 01		4A20 pseudo	96% to 4A20	1,2,16,17,18,85,95,100k
AL358613.4	chr 10		26A1	missing 1-151	80,81,82,83k
AL358613.4	chr 10	not in AL358613.6	26C1	partial	103,121,122,148,153,154k
AL358790.7	chr 10		17	complete gene	36,37,38,39,40,41,43k
AL359454.3	chr 13		51P2		3,4k
AL359672.10	chr 10		2C8	complete gene	83,94,97,101,102,103,116,118,139k
AL359672.10	chr 10		2C9	complete gene	26,35,39,40,42,128,134,135k
AL359672.10	chr 10		2C-like fragm		73,75,119k
AL359741.3	chr 13		51P2		144k
AL390726.4	chr 09		4F pseudo	92% to 4F26P	108k
AL391382.1	chr 13	2 diffs with AC027800	4F pseudo	79% to 4F25P	97,98k
AP002766.1	chr 11?	is this really chr 13?	51P2		73,74k
NT_001551	chr 07		5A1	200k gene	joins AC006021, AC004914, AC004961

Last modified Sept. 26, 2000 David Nelson

180 entries