

CORRIGENDUM

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An integrated map of chromosome 18 CAG trinucleotide repeat loci

In Table 2 of this paper some of the data is factually incorrect. The correct Table 2 is given below.

Table 2 Database homology and map location of chromosome 18 CAG repeat loci

<i>STS</i>	<i>Database homology</i>	<i>Physical position (FLpter)</i>	<i>Sub-band localisation</i>	<i>Linked marker on radiation hybrid map^a</i>	<i>Distance from linked marker (centiRays)</i>
LL18CAG10	T06032	pter	18p11.32	D18S1132	6.43
LL18CAG12	Foetal Brain EST U55976, H09960, R25492, R59809, H15991, R19422, R42000 ESTs	0.18-0.23	18p11.1-p11.21	D18S1228 D18S852	11.69 14.80
LL18CAG17	-	0.15-0.18	18p11.21	D18S1226	63.70
LL18CAG19	-	0.57-0.72	18q12.3-q21.2	D18S1311	33.52
LL18CAG20	Z78337 Foetal Brain EST	0.65-0.74	18q21.2-q21.31	D18S1311	31.40
LL18CAG25	-	0.55-0.66	18q12.3-q21.2	D18S1212	14.84
LL18CAG30	-	0.91-0.95	18q22.3	D18S823	17.35
LL18CAG31	-	failed	-	D18S1282	4.11
LL18CAG39	-	0.16-0.19	18p11.21	WI-5607	6.83 ^c
LL18CAG48	-	0.35-0.52	18q11.2-q12.1	D18S869	15.55 ^c
LL18CAG50	-	0.9-1.0	18q22.3-q23	D18S1161	41.36
LL18CAG61	L34155 LAMA3	failed	18q11.2 ^b	D18S869 D18S1067	27.10 41.10
LL18CAG73	-	pter	18p11.32	D18S1231	3.1
LL18CAG218	-	failed	-	D18S60	25.97
GCT3E06	-	-	-	D18S1161	31.20
GCT3G01	-	-	-	D18S828	0 ^d
GCT6F12	-	-	-	D18S58	0 ^d
GCT6G01	-	-	-	D18S542	24.09
GCT7G01	-	-	-	D18S71	31.58
GCT5D07	U55976	-	-	D18S573	30.73
GCT13D05	-	-	-	D18S869	35.89
GCT3A09	-	-	-	D15S148	33.45

^aMarkers were typed on the StanfordG3 Radiation Hybrid panel unless indicated otherwise.

^bFISH failed, however sequence identity to LAMA3 which is already mapped to 18q11.2.

^cThese markers were typed on the Genebridge4 hybrid panel.

^dThese markers are present on the Stanford G3 hybrid map.