

**Table 1. Microsatellite markers for first round of the genome screen with logarithm of odds (lod) >1 (SIBPAIR) (1)**

Microsatellite marker	Cytogenetic location	lod screen1	<i>P</i> value screen1	lod screen2	<i>P</i> value screen2
D1S238	1q31.1	<b>1.16</b>	0.010	0.09	0.251
D1S425	1q32.3	<b>1.20</b>	0.009	0.21	0.125
D14S292	14q32.3	<b>1.10</b>	0.010	0.00	0.500
D16S503	16q21	<b>1.16</b>	0.010	0.00	0.500
D17S1852	17p11.2	<b>1.00</b>	0.016	0.00	0.500
D19S220	19q13.1	<b>1.00</b>	0.016	0.00	0.500
D21S1252	21q22.12	<b>1.58</b>	0.003	<b>1.00</b>	0.016

Results from the second round are shown in bold.

1. Satsangi, J., Parkes, M., Louis, E., Hashimoto, L., Kato, N., Welsh, K., Terwilliger, J. D., Lathrop, G. M., Bell, J. I. & Jewell, D. P. (1996) *Nat. Genet.* **14**, 199–202.