|  |  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- | --- |
| **Sibship1** | **N** | **Clusters** | **Nmax** | **Nmin** | **Mothers** | **Fathers** | **Families** | **Full sib** | **Full sib p** |
| **GOM** | 48 ± 5 | 7 ± 1 | 22 ± 6 | 1 ± 1 | 22 ± 1 | 24 ± 2 | 45 ± 4 | 10 ± 5 | 0.174 ± 0.105 |
| **MED** | 50 ± 0 | 7 ± 2 | 17 ± 3 | 1 ± 1 | 22 ± 3 | 24 ± 1 | 47 ± 1 | 8 ± 4 | 0.228 ± 0.135 |

**1**Replicate data consisted of triplicate multilocus genotypes at 8 microsatellite loci, and for 7 and 6 loci (excluding Tth16-2 and also Ttho1 due to null allele presence).