|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| LG | Map length (cM) | Number of markers | Unique map positions | Avg. genetic dist. between unique markers in cM (sd) |
| 1 | 93.9 | 969 | 335 | 0.28 (0.33) |
| 2 | 71.37 | 893 | 253 | 0.28 (0.88) |
| 3 | 76.3 | 912 | 256 | 0.30 (0.65) |
| 4 | 70.2 | 851 | 257 | 0.28 (0.41) |
| 5 | 78.7 | 778 | 295 | 0.27 (0.29) |
| 6 | 69.1 | 741 | 247 | 0.28 (0.88) |
| 7 | 59.8 | 738 | 234 | 0.26 (0.33) |
| 8 | 65.6 | 674 | 246 | 0.27 (0.41) |
| 9 | 68.6 | 623 | 182 | 0.37 (0.58) |
| 10 | 71.1 | 410 | 150 | 0.48 (0.96) |
| Avg. | 72.74 (8.69) | 758.90 (155.75) | 245.50 (49.04) | 0.31 (0.06) |