Supplementary Table 3. Summary of number, minor allele frequency (MAF) and density of markers used for spot blotch GWAS in 294 wheat genotypes of hard winter wheat association mapping panel (HWWAMP).

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **Chromosomes** | **No. of markers** | **Chromosome length (cM)** | **Minor allele frequency** | **Average number of marker/cM** | **Average distance between** **markers (cM)** |
| Genome A1A-7A | 1 | 1036 | 161.335 | 0.2419 | 6.4 | 0.16 |
| 2 | 956 | 179.116 | 0.1998 | 5.3 | 0.19 |
| 3 | 773 | 188.378 | 0.2442 | 4.1 | 0.24 |
| 4 | 744 | 164.13 | 0.1889 | 4.5 | 0.22 |
| 5 | 790 | 148.304 | 0.2333 | 5.3 | 0.19 |
| 6 | 958 | 164.089 | 0.2093 | 5.8 | 0.17 |
| 7 | 954 | 244.155 | 0.2417 | 3.9 | 0.26 |
| - | 887**†** | 178.501**†** | 0.2227† | 5.0**†** | 0.20**†** |
| Genome B1B-7B | 1 | 1254 | 173.624 | 0.2415 | 7.2 | 0.14 |
| 2 | 1162 | 185.666 | 0.2485 | 6.3 | 0.16 |
| 3 | 1117 | 149.634 | 0.2727 | 7.5 | 0.13 |
| 4 | 451 | 119.446 | 0.2591 | 3.8 | 0.26 |
| 5 | 1554 | 219.773 | 0.2100 | 7.1 | 0.14 |
| 6 | 1237 | 127.049 | 0.2620 | 9.7 | 0.10 |
| 7 | 855 | 178.856 | 0.2259 | 4.8 | 0.21 |
| - | 1090 **†** | 164.864**†** | 0.2457† | 6.6**†** | 0.16**†** |
| Genome D1D-7D | 1 | 438 | 192.274 | 0.1939 | 2.3 | 0.44 |
| 2 | 581 | 138.737 | 0.1688 | 4.2 | 0.24 |
| 3 | 240 | 164.672 | 0.2176 | 1.5 | 0.69 |
| 4 | 52 | 170.428 | 0.2205 | 0.3 | 3.28 |
| 5 | 159 | 207.329 | 0.2580 | 0.8 | 1.30 |
| 6 | 146 | 160.498 | 0.2517 | 0.9 | 1.10 |
| 7 | 133 | 223.524 | 0.2662 | 0.6 | 1.68 |
| - | 250**†** | 179.637**†** | 0.2252† | 1.5**†** | 1.25**†** |
| Total | - | 742 | 174.334 | 0.2312 | 4.4 | 0.54 |

† corresponding mean values of genomes A, B, and D