Lodging values of genotypes with significantly higher lodging scores than the yield checks of their respective test.

|  |  | **Lodging** | | | |
| --- | --- | --- | --- | --- | --- |
| **Genotypea** | **Test Nameb** | **Valuec** | **Rankd** | **Test Averagee** | **Check Averagef** |
| N18-1604 | Jay Test 1 | 3.35 (135.1%) | 1 | 2.63 | 2.48 |
| N18-1641 | 3.15 (127%) | 3 |
| N18-1956 | 3.05 (123%) | 5 |
| N18-1661 | Jay Test 2 | 3.04 (177.8%) | 1 | 2.32 | 1.71 |
| N18-1628 | 2.96 (173.1%) | 2 |
| N18-1577 | 2.79 (163.2%) | 3 |
| N18-1820 | 2.75 (160.8%) | 4 |
| N18-1796 | 2.71 (158.5%) | 5 |
| N18-1659 | 2.62 (153.2%) | 6 |
| N18-1579 | 2.58 (150.9%) | 7 |
| N18-1586 | 2.58 (150.9%) | 8 |
| aThe genotype name. | | | | | |
| bThe test name | | | | | |
| cThe genotype marginal mean for the phenotype (value divided by check average). | | | | | |
| dThe ranking of this genotype for the phenotype within its test. | | | | | |
| eThe average phenotype value for all genotypes in the test. | | | | | |
| fThe average value of the checks in the test. | | | | | |