**Table S2.** Two‐way analysis of variance (ANOVA) of seedling rot dose-response assay at the V2 stage. DF = degree of freedom, Sum Sq= sum of squares, Mean sq= mean squares, \*Significantly different at 0.05 probability level.

|  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- |
| **Source of variance** | **Df** | **Sum Sq** | **Mean Sq** | **F value** | **Pr(>F)** |
| **Severity** |  |  |  |  |  |
| Genotype | 1 | 10054 | 10054 | 127.365 | <2e-16\* |
| Doses | 3 | 32338 | 10779 | 136.560 | <2e-16\* |
| Genotype:Doses | 2 | 1163 | 582 | 7.369 | 0.00107\* |
| Residuals | 93 | 7341 | 79 |  |  |
| **Height** |  |  |  |  |  |
| Genotype | 1 | 19893 | 19893 | 61.12 | 1.40e-11\* |
| Doses | 3 | 16541 | 5514 | 16.94 | 1.08e-08\* |
| Genotype:Doses | 3 | 9954 | 3318 | 10.20 | 8.53e-06\* |
| Residuals | 84 | 27340 | 325 |  |  |
| **Root fresh weight** |  |  |  |  |  |
| Genotype | 1 | 16405 | 16405 | 68.367 | 1.14e-12\* |
| Doses | 3 | 25827 | 8609 | 35.878 | 2.41e-15\* |
| Genotype:Doses | 3 | 6893 | 2298 | 9.576 | 1.50e-05\* |
| Residuals | 90 | 21596 | 240 |  |  |
| **Aerial fresh weight** |  |  |  |  |  |
| Genotype | 1 | 6687 | 6687 | 36.752 | 2.63e-08\* |
| Doses | 3 | 29232 | 9744 | 53.550 | < 2e-16\* |
| Genotype:Doses | 3 | 3580 | 1193 | 6.559 | 0.000442\* |
| Residuals | 96 | 17468 | 182 |  |  |