|  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- |
| **Table x**. Beta-dispersion for fungal microbial communities associated with above ground plant tissues (shoots) from T0 to T1 for three inoculation treatments and an uninoculated control. | | | | | | | |
| *Inoculum Treatment* | *Effect* | *df* | *SS* | *MS* | *F* | *Permutations* | *p-value* | |
| Dry Inoculum | T0 to T1 | 1 | 0.223 | 0.223 | 10.804 | 999 | **0.001** | |
|  | Residuals | 50 | 1.034 | 0.021 |  |  |  | |
|  |  |  |  |  |  |  |  | |
| Moderate Inoculum | T0 to T1 | 1 | 1.045 | 1.045 | 121.280 | 999 | **0.001** | |
|  | Residuals | 52 | 0.448 | 0.009 |  |  |  | |
|  |  |  |  |  |  |  |  | |
| Wet Inoculum | T0 to T1 | 1 | 0.403 | 0.403 | 47.538 | 999 | **0.001** | |
|  | Residuals | 44 | 0.373 | 0.008 |  |  |  | |
|  |  |  |  |  |  |  |  | |
| Control, Uninoculated | T0 to T1 | 1 | 0.191 | 0.191 | 13.414 | 999 | **0.001** | |
|  | Residuals | 49 | 0.698 | 0.014 |  |  |  | |
| *df* = degrees of freedom; *SS* = sum of squares; *MS* = mean squares. Significant effects (*p*<0.05) are in bold. | | | | | | | |