**Table S2. *OsPUB41* expression is induced following infection with either bacterial or fungal pathogens**

|  |  |  |  |  |  |  |  |  |
| --- | --- | --- | --- | --- | --- | --- | --- | --- |
| **Experiment** | **Fungal strains** | | | | | **Bacterial strains** | | |
| ***Magnaporthe grisea* FR13**a | | ***Magnaporthe oryzae***  **Guy11**a | | | ***PXO99A***a | ***PXO86***a | ***BXO43b*** |
| Time pointsc | 3dpi | 4dpi | 2dpi | 4dpi | 6dpi | 24hpi | 24hpi | 12hpi |
| *OsPUB41* (Fold Change)d | 7.2 | 199 | 151.7 | 19.4 | 24.2 | 7.2 | 21.3 | 5.1±0.6 |

aData in publicly available databases was analyzed. In these experiments, rice had been infected with either fungal (*Magnaporthe grisea* FR13; Rice cultivar: Nipponbare; GEO-ID: GSE7256 or *Magnaporthe oryzae* Guy11; Rice cultivar: Nipponbare; GEO-ID: GSE18361) or bacterial pathogens (Xoo strains: *PXO99A* and *PXO86*, rice cultivar: IR24; GEO-ID: GSE36272)

bBXO43: Xoo (BXO43 strain) treated 15 days old TN-1 rice leaves were harvested 12 hpi for qPCR analysis. Relative expression of *OsPUB41* in infected as compared to mock inoculated rice leaves was calculated using the 2(-ΔΔCt) method.

cdpi: days post infection, hpi: hours post infection

dRelative expression of *OsPUB41* in infected as compared to mock inoculated rice plants was calculated using Expression Console and Transcriptome Analysis Console (p ≤ 0.05).