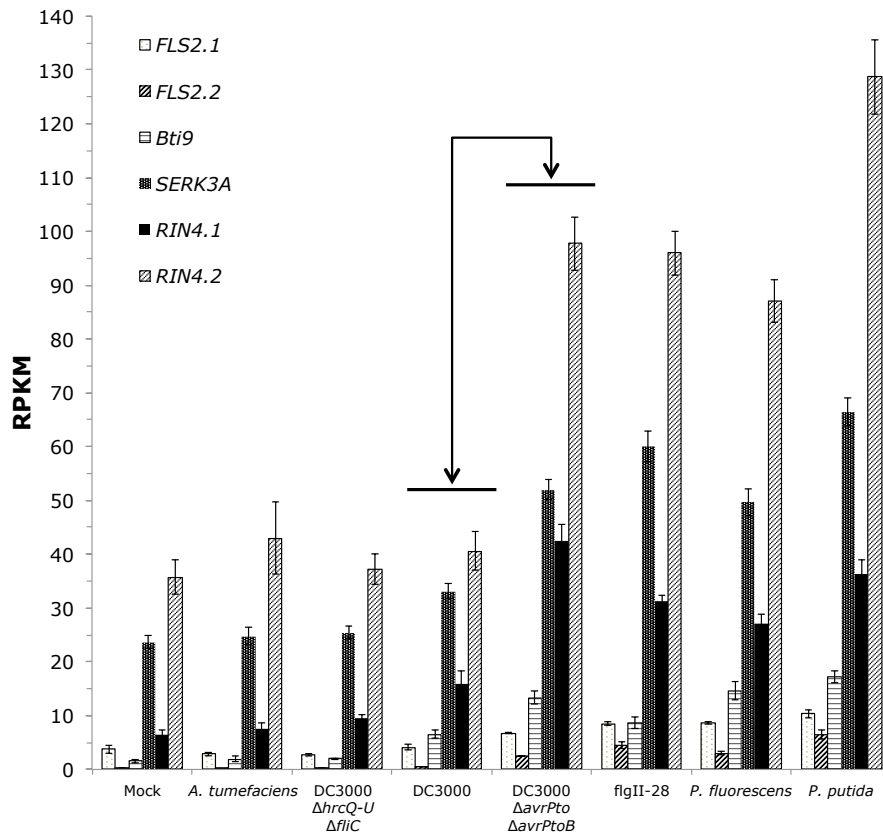




Additional file 1: Figure S1. Expression level of selected immunity-related genes used for establishing a minimum RPKM cut off (see Methods). Bars represent the average of three biological replicates with the corresponding standard error. Arrows highlight transcript abundance differences between wild type DC3000 and DC3000 $\Delta avrPto\Delta avrPtoB$. *FLS2.1* Solyc02g070890, *FLS2.2* Solyc02g070910, *Bti9* Solyc02g079600, *SERK3A* Solyc10g047140, *RIN4.1* Solyc09g059430 and *RIN4.2* Solyc06g083390.