

Figure S5. Transcription profile of the *P. megakarya* (Pm1) and *P. palmivora* (Pp2) RxLRs detected *in planta* and distribution of the RxLR families based on their expression pattern. For the relative transcription profiles, *Pmeg* (A) and *Ppal* (B) RxLRs detected in at least three of the four biological replicates (with \geq 1 paired read) of 15hpi *in planta* samples were normalized and Log₁₀ transformed. (C) Number of RxLRs in each of expression categories of the RxLR families. The heat map was generated using CIMminer (http://discover.nci.nih.gov/cimminer).