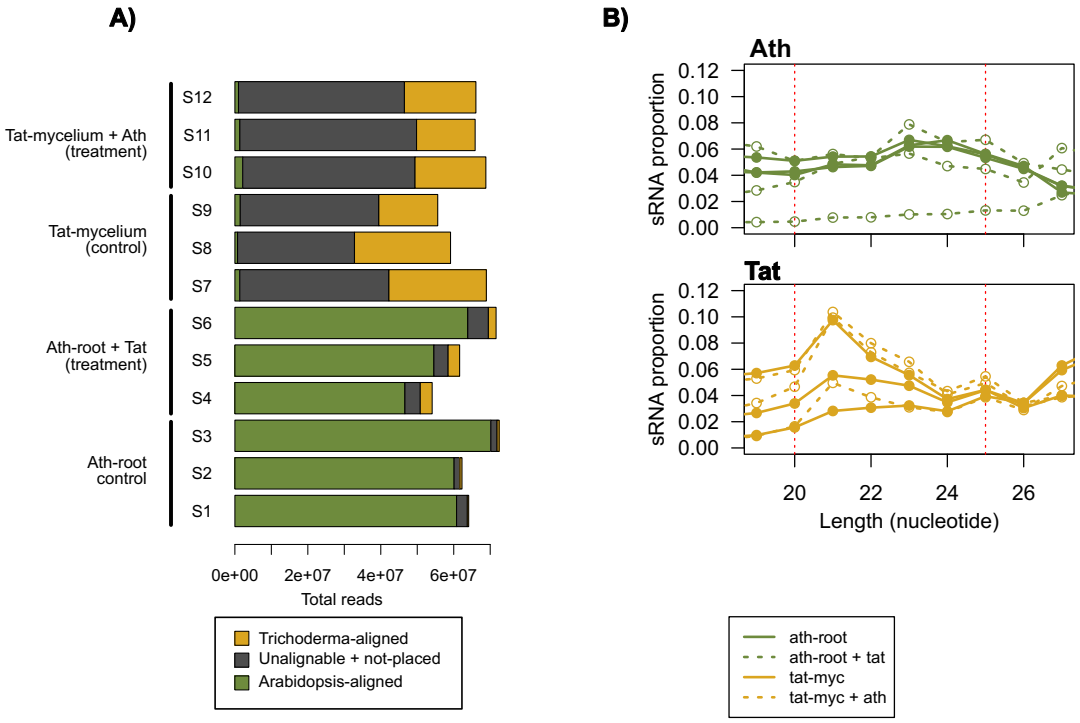


Fig. S7. Small RNA alignment details. (A) Alignment rates for sRNA libraries to merged genome. Colors indicate reads aligned to the *Arabidopsis thaliana* (olivegreen) or *Trichoderma atroviride* (goldenrod) genomes, and reads that were unalignable or not-placed due to high-count of multi-mapping sites or lack of multi-mapper weighting. (B) Size profiles of aligned reads in *Arabidopsis thaliana* and *Trichoderma atroviride*. Solid lines indicate samples with the organism alone and dotted-lines indicate interaction conditions. Y-axis is the proportioin of all aligned reads in the library.