

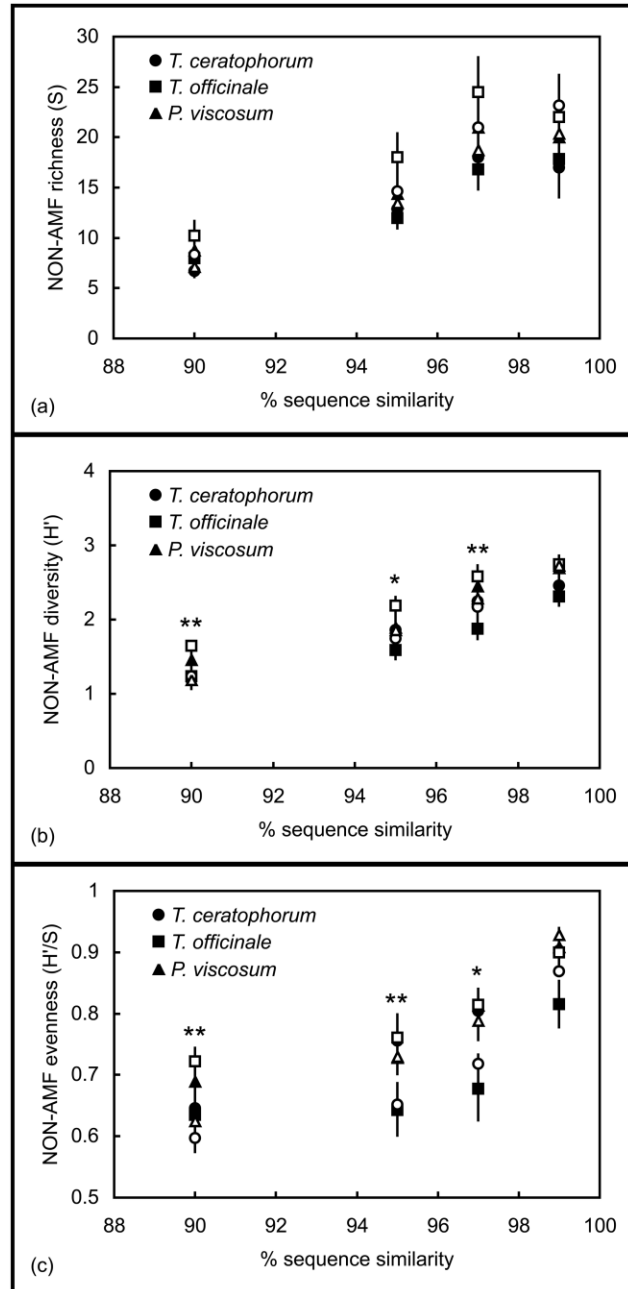


Fig. S2 Mean ($\pm$ SE) non-AMF richness, diversity, and evenness per root sample based on OTUs designated at 90%, 95%, 97%, and 99% sequence similarity. (a) OTU richness by habitat and host species; (b) Shannon's diversity index by habitat and host species; (c) OTU evenness by habitat and host species. Symbols represent samples from *T. ceratophorum* (circles), *T. officinale* (squares), and *P. viscosum* (triangles). Open and solid symbols represent samples from the open meadow and willow understory, respectively. Asterisks indicate a significant habitat by host interaction ($*P \leq 0.05$, $**P \leq 0.01$).