Table S1. Study sites were located in two mixed maritime forests in the early stages of succession. These two forests, which we called Forest 1 and Forest 2, are approximately 9 km apart and are located near Bellingham, Washington, USA. Altitude and soil organic matter content of the studied forests are presented as means \pm standard errors. Study sites 1 through 4 were located in Forest 1, while 5 through 8 were located in Forest 2.

| Location ID | n | Altitude (m) | Organic matter (%) | Study Sites |
|-------------|---|--------------|--------------------|----------------|
| Forest 1 | 4 | 174 | 14.4 ± 3.8 | 1-4, composite |
| Forest 2 | 4 | 471 | 24.5 ± 5.0 | 5 - 8 |

Table S3. Arbsucular mycorrhizal fungal taxa found in litter, soil, or both litter and soil after resampling to 114 sequences/sample.

| Where found | Virtual taxa | Genus |
|------------------------------|--------------|-----------------|
| Only in leaf litter | | |
| | VTX00067 | Glomus |
| | VTX00108 | Glomus |
| | VTX00113 | Glomus |
| | VTX00115 | Glomus |
| | VTX00143 | Glomus |
| | VTX00165 | Glomus |
| | VTX00197 | Glomus |
| | VTX00159 | Glomus |
| | VTX00239 | Paraglomus |
| In both leaf litter and soil | | |
| | VTX00004 | Archaeospora |
| | VTX00057 | Claroideoglomus |
| | VTX00225 | Claroideoglomus |
| | VTX00357 | Claroideoglomus |
| | VTX00160 | Glomus |
| | VTX00281 | Paraglomus |
| | VTX00308 | Paraglomus |
| Only in soil | | |
| • | VTX00030 | Acaulospora |
| | VTX00056 | Claroideoglomus |
| | VTX00193 | Claroideoglomus |
| | VTX00340 | Claroideoglomus |
| | VTX00061 | Diversispora |
| | VTX00062 | Diversispora |
| | VTX00072 | Glomus |
| | VTX00074 | Glomus |
| | VTX00084 | Glomus |
| | VTX00088 | Glomus |
| | VTX00122 | Glomus |
| | VTX00129 | Glomus |
| | VTX00153 | Glomus |
| | VTX00199 | Glomus |
| | VTX00214 | Glomus |
| | VTX00216 | Glomus |
| | VTX00219 | Glomus |

Table S4. Results of a permutational multivariate analysis of variance of AM fungal communities considering two factors, site and source of mycorrhizal community (*i.e.* soil or litter). This analysis used the same data that are displayed in Fig. 1c. Note that the factor 'site' included a separate category for composite samples.

| Source | df | SS | R^2 | F | p |
|-------------|----|-------|-------|------|-------|
| site | 8 | 39418 | 0.698 | 2.86 | 0.006 |
| soil/litter | 1 | 10127 | 0.179 | 5.88 | 0.004 |
| residual | 4 | 6891 | 0.122 | | |
| Total | 13 | 56436 | | | |