**S3 Table.** **S-D-R simplex analysis in the pH-manipulation experiment.**

|  |  |  |  |
| --- | --- | --- | --- |
| Data format | Turnover | Richness agreement | Nestedness |
| Presence/absence | 49.5a | 66.2 | 84.3 |
| Abundance | 65.0 | 69.7 | 65.4 |

a Percentage relative contributions of phylotype turnover, richness agreement, and nestedness to the distribution pattern of AM fungi along the pH gradient are indicated. The same data matrices for the NODF/WNODF analysis were used.