Figure S1. Rarefaction curves obtained for the HiSeq sequencing reads of operational taxonomic units (OTUs) from the V3–V4 region of bacterial 16S rDNA genes in soil samples collected at the two different forest sites. Rarefaction curves at a cutoff level of 3% were constructed at a 97% sequence similarity cutoff value in mothur (http://www.mothur.org/). RS1, RS2, RS3, RS4 represent September and December 2015 and March and June 2016 at TBDF-BTM, respectively. RS5, RS6, RS7, and RS8 represent September and December 2015 and March and June 2016 at TRF-JFL, respectively.