S4 Fig. Raw gel images for the BOX-PCR fingerprints of diazotrophic/N-scavenging bacteria isolated from different sources and from soils under different management conditions used. The resulting fingerprints were analyzed using BioNumerics v. 4.6 to produce a dendrogram based on the Jaccard coefficient from the distance matrix (2% tolerance in terms of band size) and the unweighted pair-group method with arithmetic mean (UPGMA) as shown in Fig 2. 
M: Indicate the lanes containing the molecular weight marker (1kb DNA ladder, LGC biotecnologia, Brazil). X: Indicate the lanes not included in the final Fig 2.