



Fig. S1. Dynamics of the microbial structures of consortium SV79 in five different biomass substrates for 7 days, performed by DGGE, using the V3 region of the 16S rRNA gene.

Lanes 1–5: five different biomass substrates. 1-sugarcane bagasse, 2-spent mushroom substrate, 3-*Spartina anglica*, 4-*Miscanthus floridulus*, and 5-*Pennisetum sinense* Roxb.