Fig6 Comparison of PCR–DGGE diagrams of 16S rRNA gene V3 region between the normal controls and the chronic gastritis group

(A) Normal group, M1-marker, M2- control, normal group samples --1, 5, 9,12,16,20

(B) Chronic gastritis group, M1-marker, M2-control, Chronic gastritis samples --11,12,13,14,15,16,17,18,19,20