Supplementary Figure 3: Two-dimensional heat map representing rank normalized abundances (scaled between 0 and 1) of 20 differentially abundant bacterial genera (median abundance of >0.1% in at least one group) in the gut bacterial microbiomes of controls (CT1, n = 6; CT2, n = 6; CT3, n = 6; CT4, n = 6), diabetic (DM1, n = 6; DM2, n = 6) and diabetic rats showing retinal changes (DRC1, n = 6; DRC2, n = 5). The discriminating genera were arranged along the two dimensions (axes) based on hierarchical clustering.