



**Figure S2.** Relative abundance of the predicted KEGG Orthologous groups (KOs) at baseline (error lines indicate standard deviation) in faeces (A) and in saliva (B) from the two centers – KI and HP and the most abundant significantly differently predicted functional groups (L3) in the faecal (C) and salivary (D) baseline samples from the two study sites, HP (N=37) and KI (N=29).