



Fig. S4. Taxa showing differences in longitudinal pattern between patient and control skin. (a–d) Mean relative abundances and standard error of the mean for different time-points for *Corynebacterium*, *Mucilaginibacter*, *Tsukamurella tyrosinosolvens* and *Janibacter*, showing the differential longitudinal pattern between patient and control skin (raw $p < 0.05$). F: family; G: genus; S: species.