

Microbial Association with Remission	At Baseline	6 weeks post UST treatment
<i>Escherichia/Shigella</i> (OTU1)	lower relative abundance	–
<i>Faecalibacterium</i> (OTU7)	higher relative abundance	higher relative abundance
<i>Roseburia</i> (OTU12)	–	higher relative abundance
<i>Bacteroides</i> (OTU19)	higher relative abundance	–
<i>Ruminococcus</i> (OTU35)	higher relative abundance	–
<i>Ruminococcaceae</i> (OTU53)	–	higher relative abundance
<i>Clostridium XIVa</i> (OTU73)	–	higher relative abundance
<i>Blautia</i> (OTU124)	–	higher relative abundance
α – Diversity	higher	–