

**eTable 1: Study Specimens Included in Analyses**

<b>Study Measure</b>	<b>Controls</b>	<b>Sepsis (Day 1-2)</b>	<b>Sepsis (Day 3-6)</b>	<b>Sepsis (Day 8-14)</b>
Number eligible	44	43	43	43
Stool, microbiome	44 (100)	17 (40)	12 (28)	8 (19)
Stool, SCFAs	0	17 (40)	15 (35)	12 (28)
Plasma, SCFAs	0	39 (91)	35 (810)	23 (53)
PBMC, mitochondrial respiration	0	36 (84)	29 (67)	16 (37)
LPS-stimulated TNF- $\alpha$	0	36 (84)	29 (67)	16 (37)

Data are presented as n (%)

SCFAs, short-chain fatty acids; PBMC, peripheral blood mononuclear cells; LPS, lipopolysaccharide; TNF, tumor necrosis factor