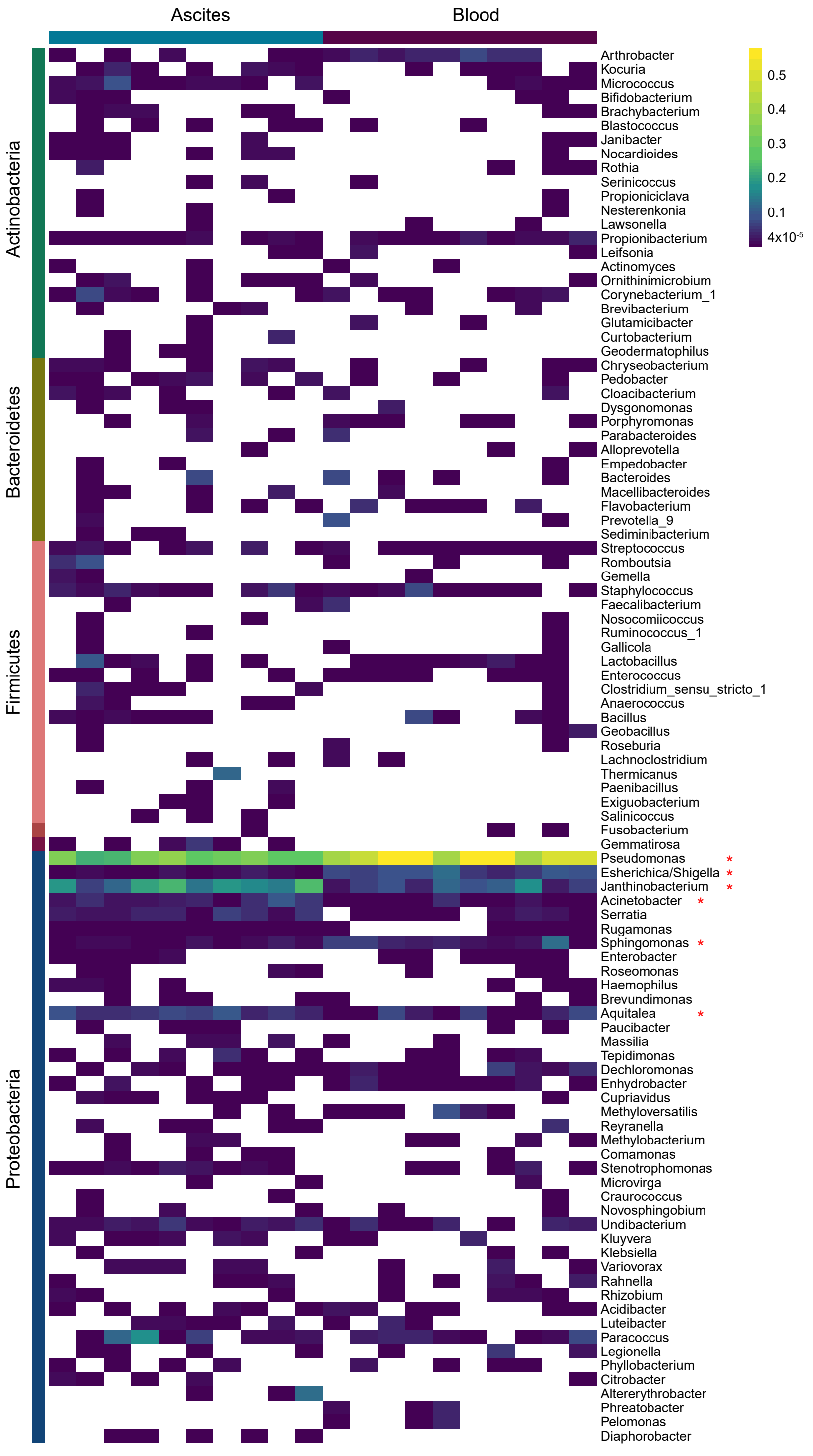

Figure S2. Microbiome composition in our cohort of 10 patients. Heatmap shows the relative abundance of different genera. Phylum affiliation of genera are marked to the left. Six core genera detected in all samples in both compartments are marked with red asterisks.