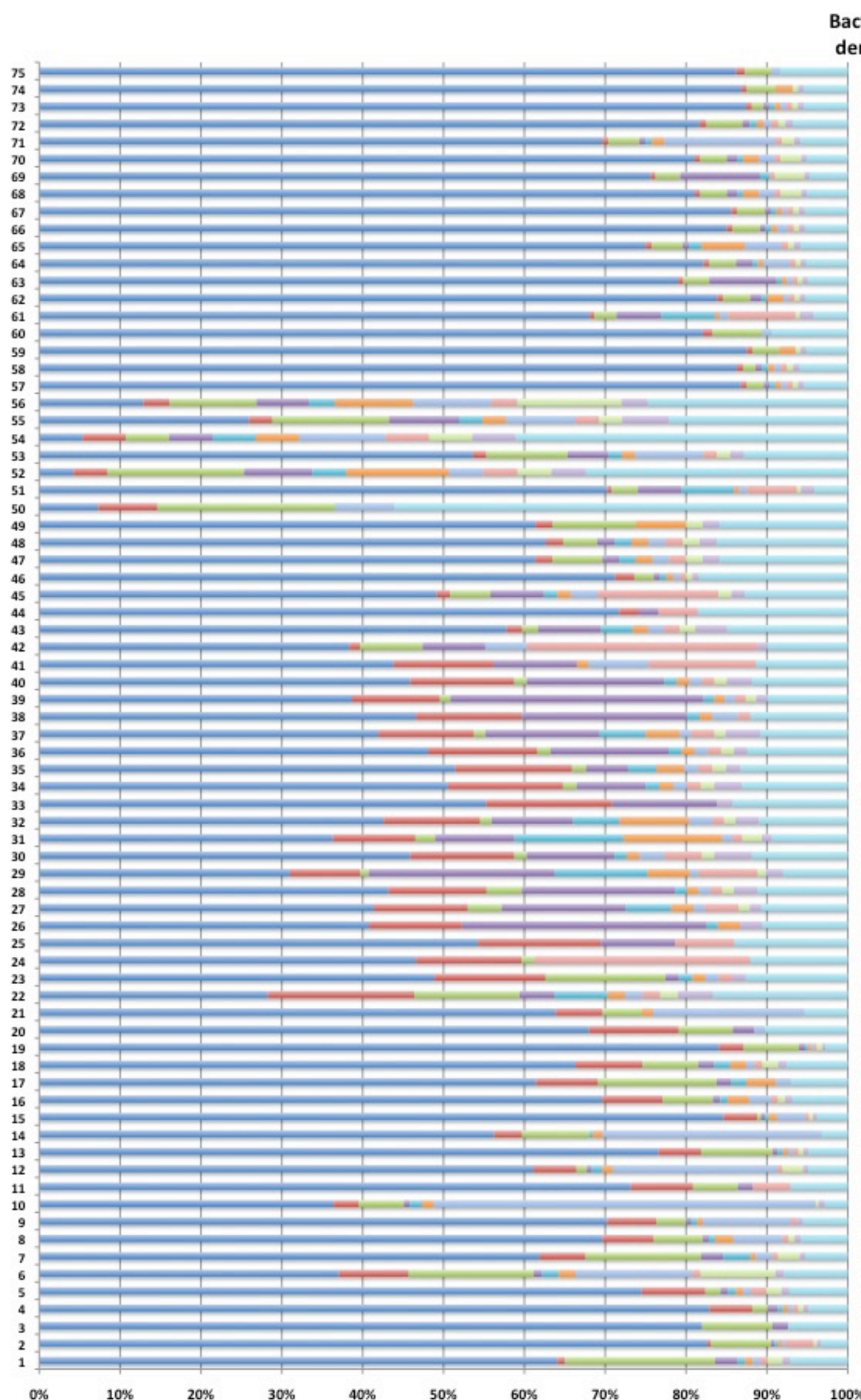
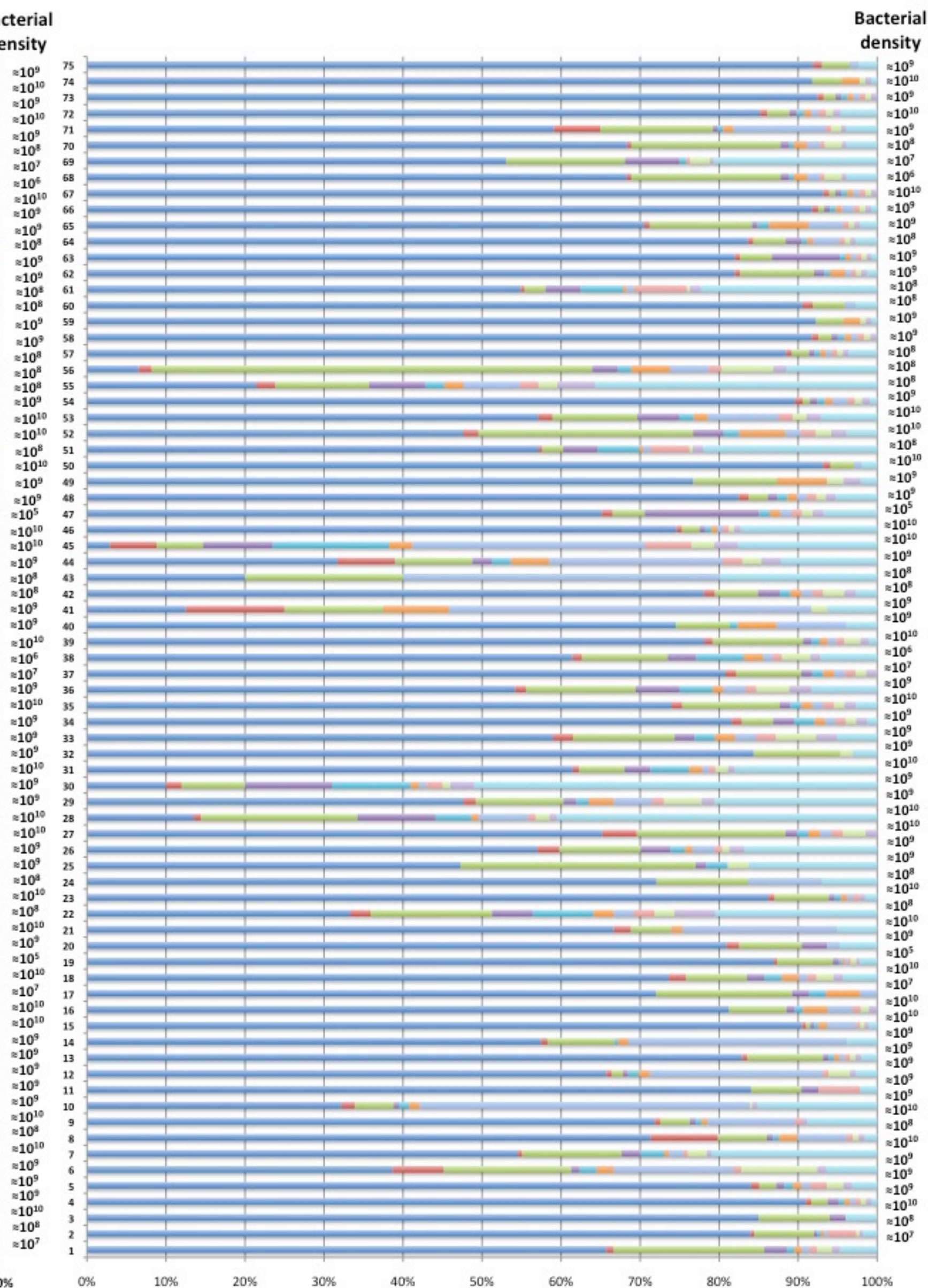
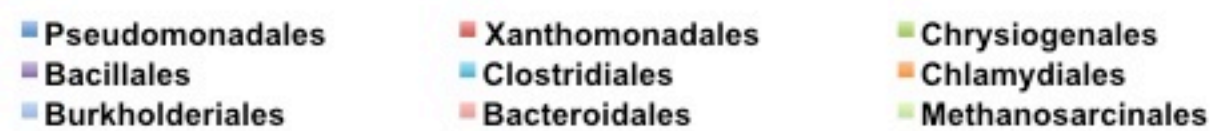


Title	Microbiota and metabolite profiling reveal specific alterations in bacterial community structure and environment in the cystic fibrosis airway during exacerbation
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% Relative abundance of OTUs from DNA



% Relative abundance of OTUs from RNA

