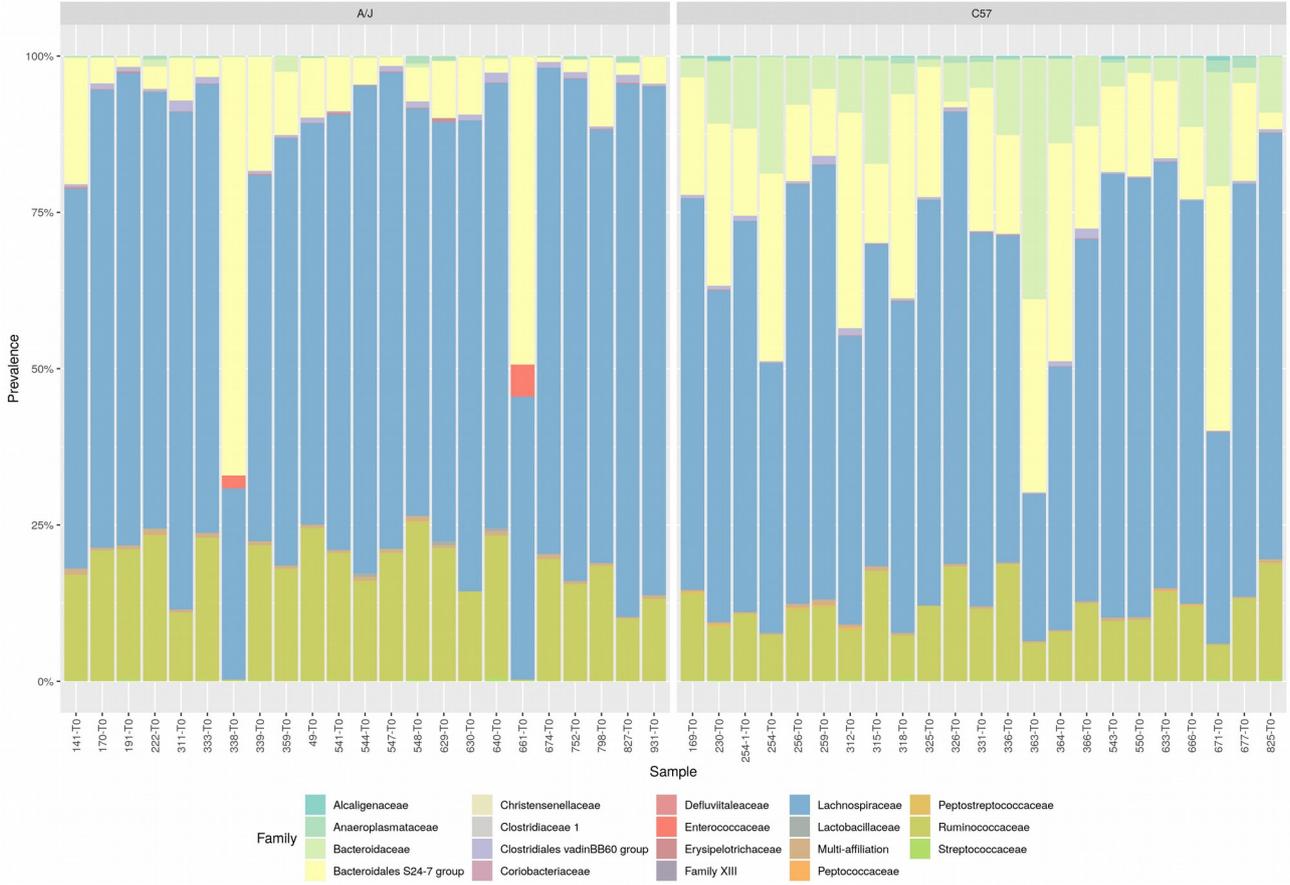
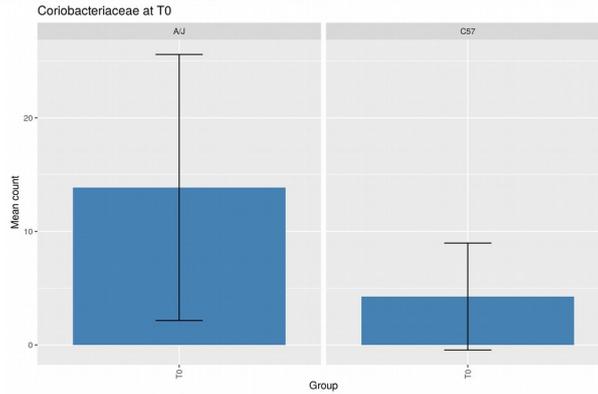
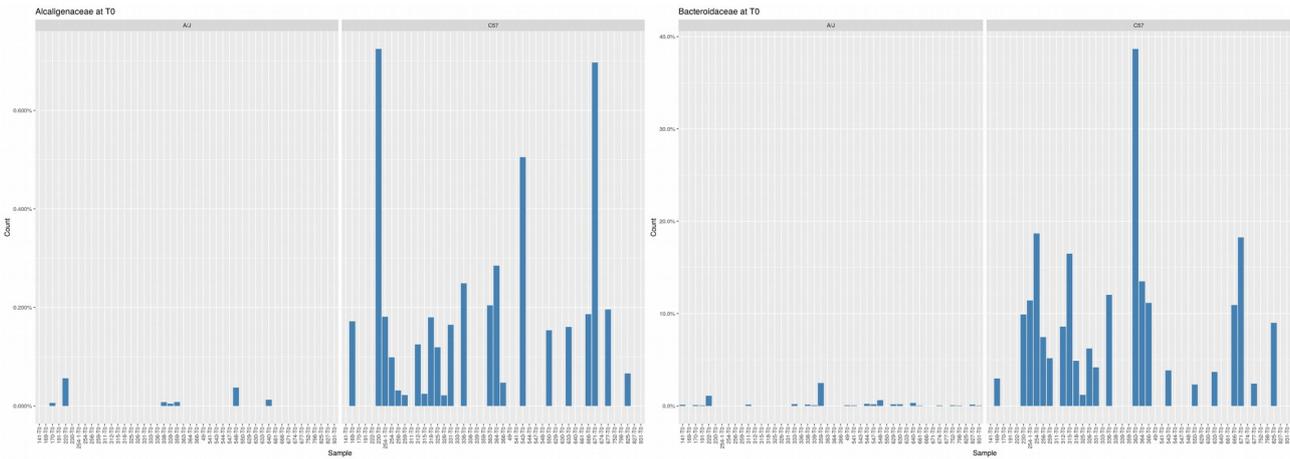


Family composition of all treatment groups at T0



Different microbiome composition in the family level between AJ and C57 strain.



The mean count of different bacteria in AJ and C57 strain before treatment.