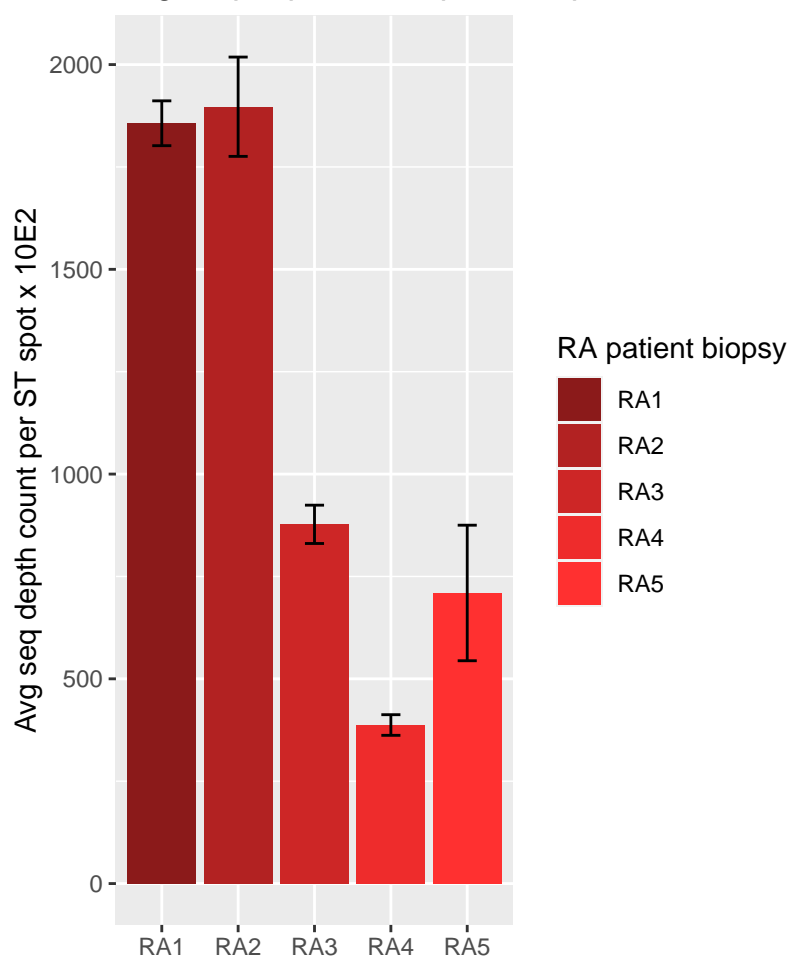
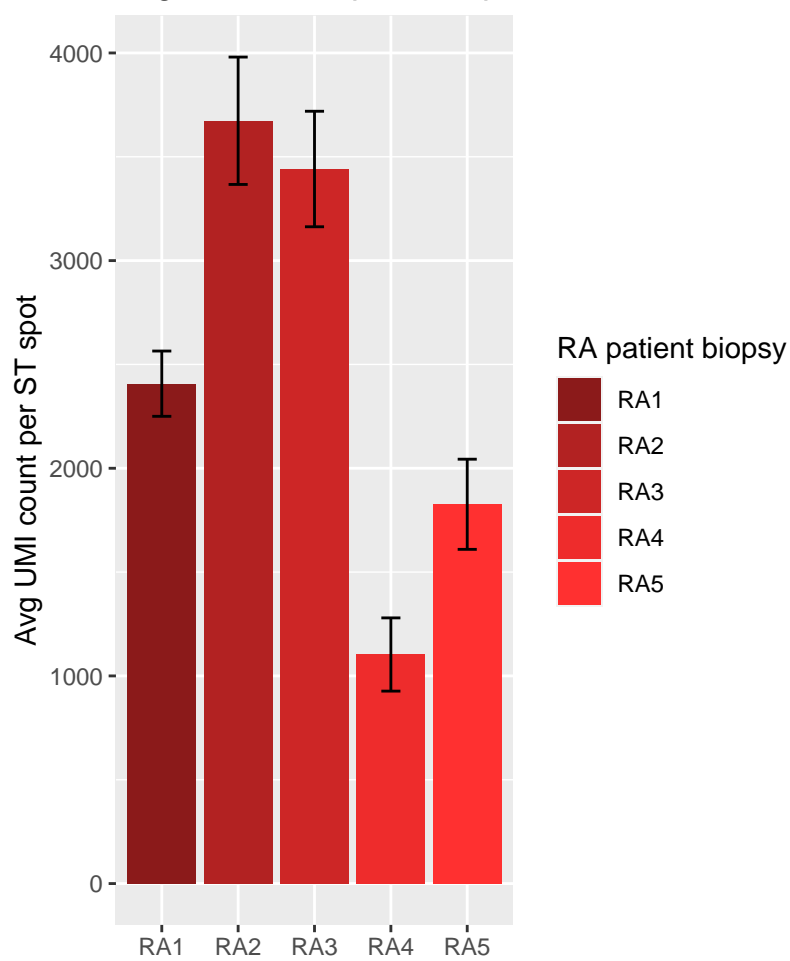


Avg seq depth count per ST spot x 10E2



Avg UMI count per ST spot



Avg gene count per ST spot

