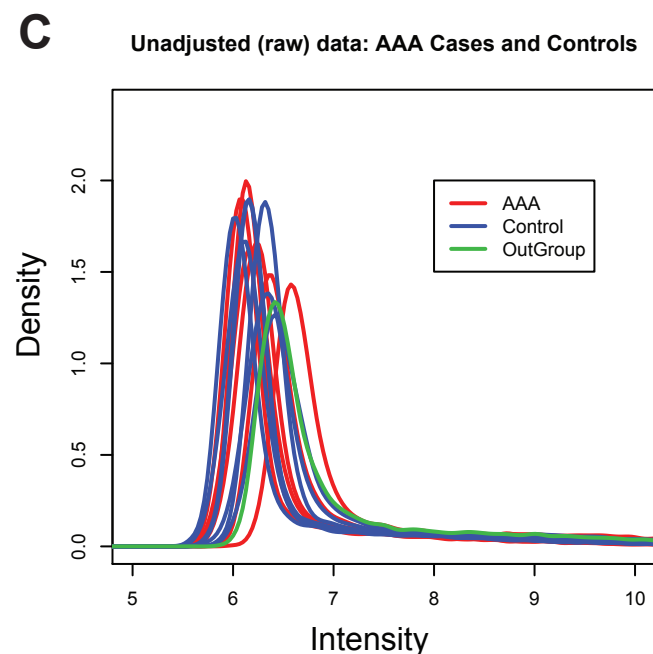
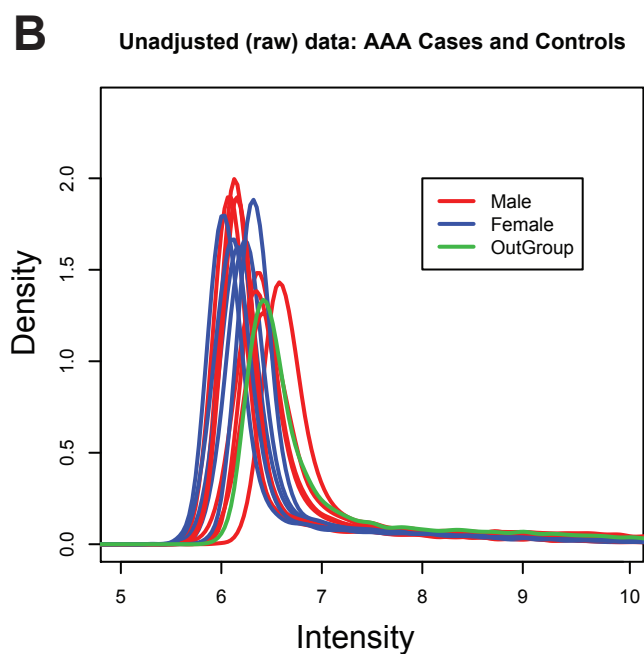
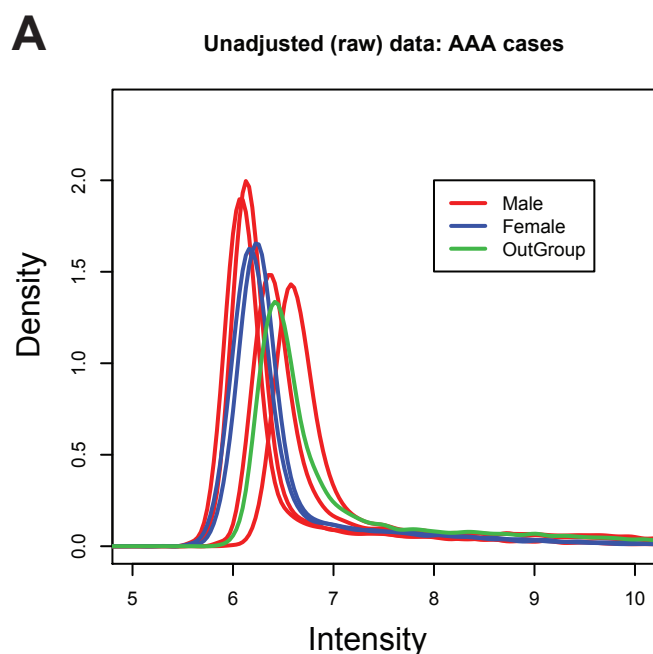




Supplemental Figure II Signal Intensity histograms. Unadjusted, i.e. raw, signals were converted into histograms using R (R: A Language and Environment for Statistical Computing; R version 2.4.1). The histograms are colored by sex (Panels **A** and **B**) and tissue type (Panel **C**; AAA vs. Control). There is no systematic bias in the observed variation, either by sex or tissue. Lack of systematic bias between AAA and control tissue suggest the difference in tissue source is not reflected in a difference in RNA signal and therefore RNA integrity at a global level.