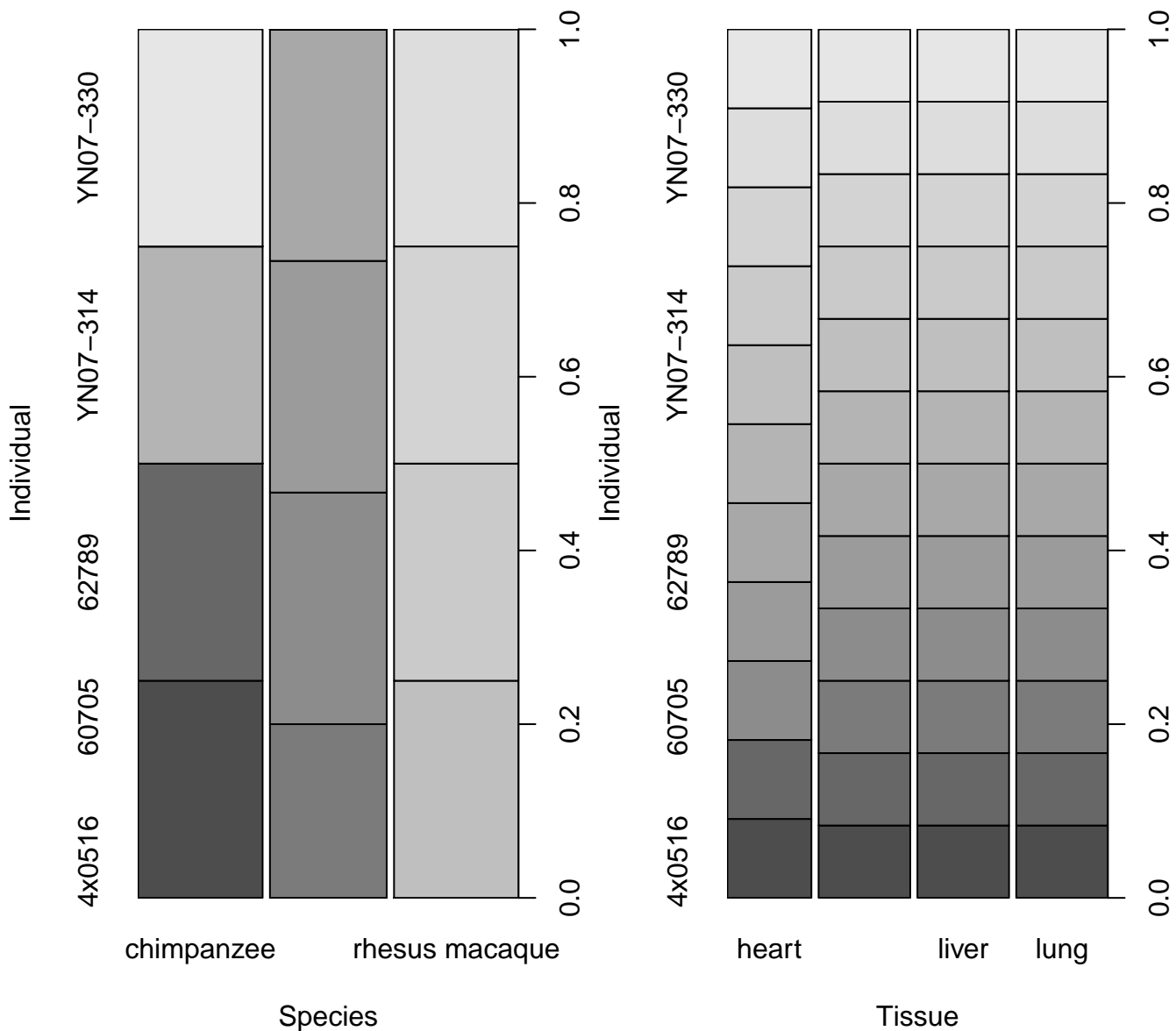
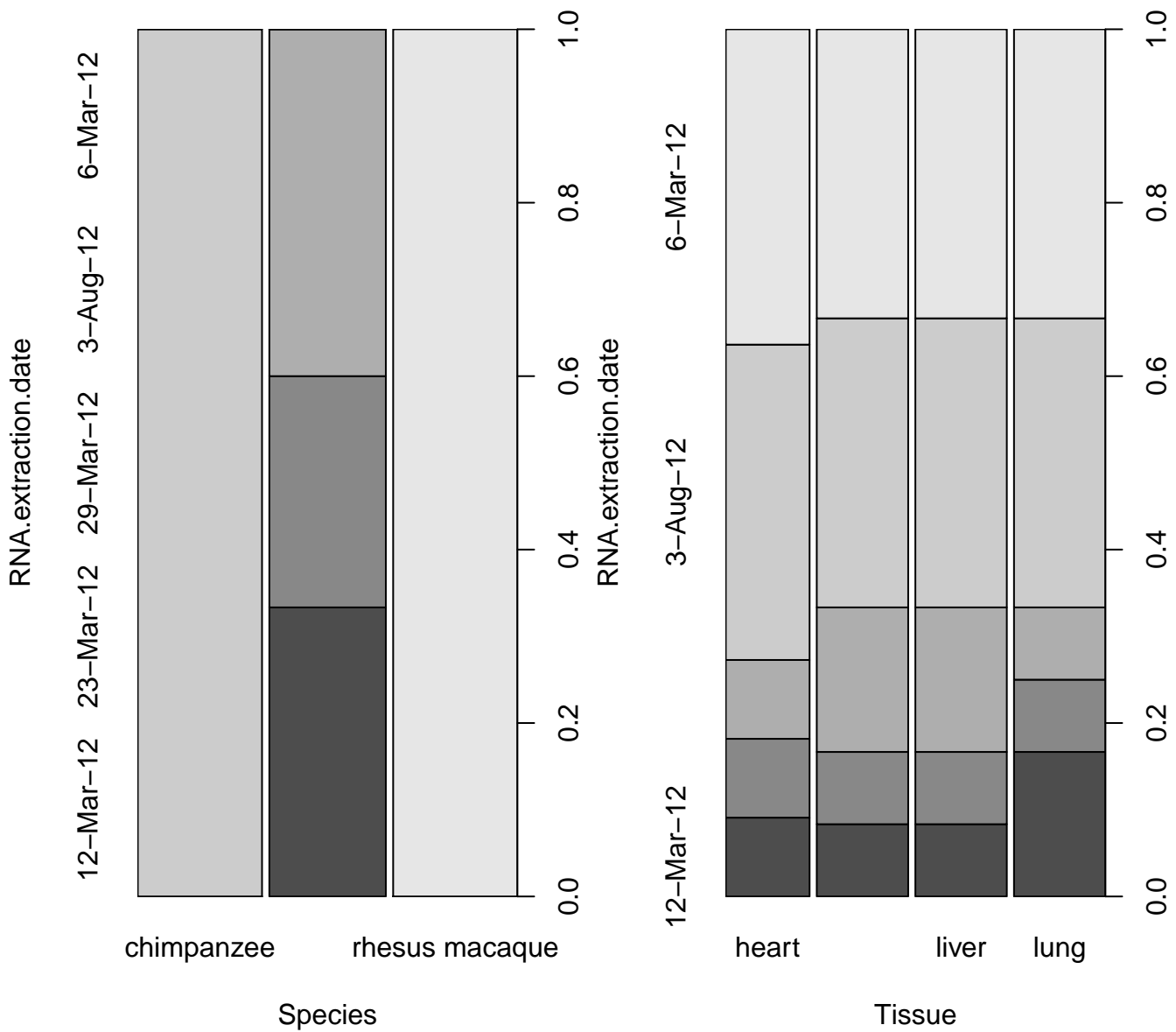


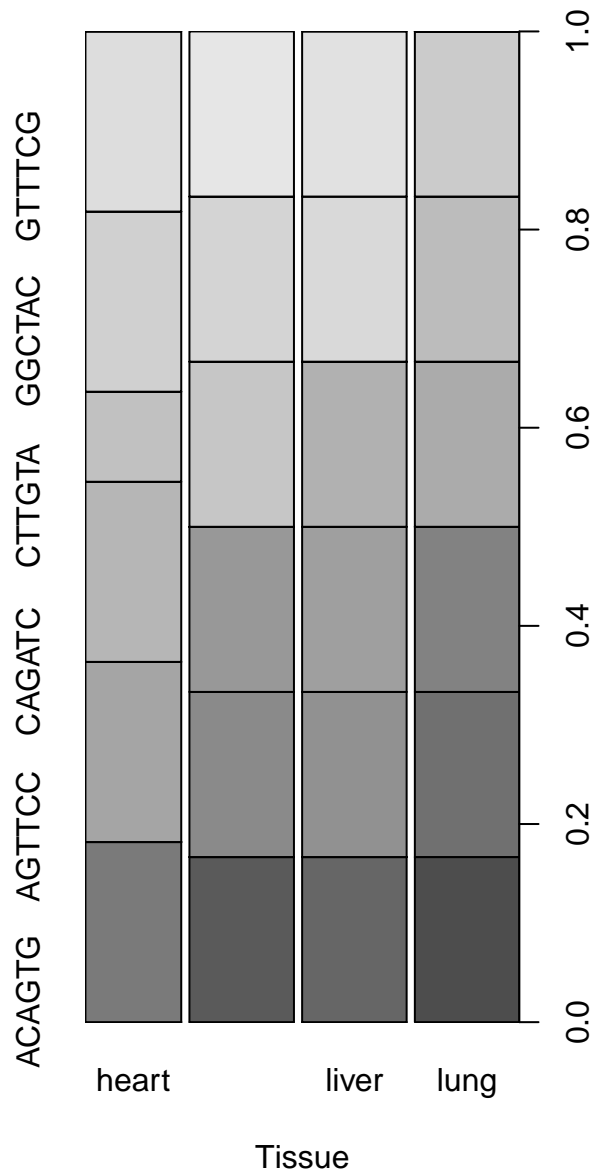
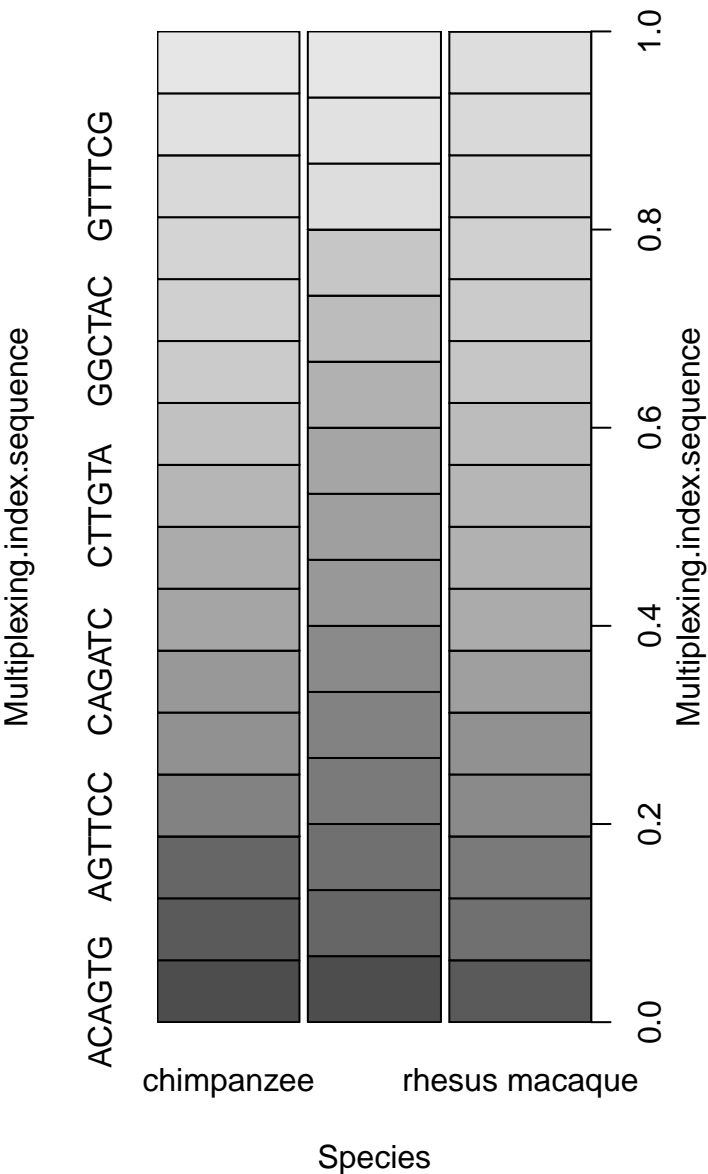
Individual vs. Species and Tissue



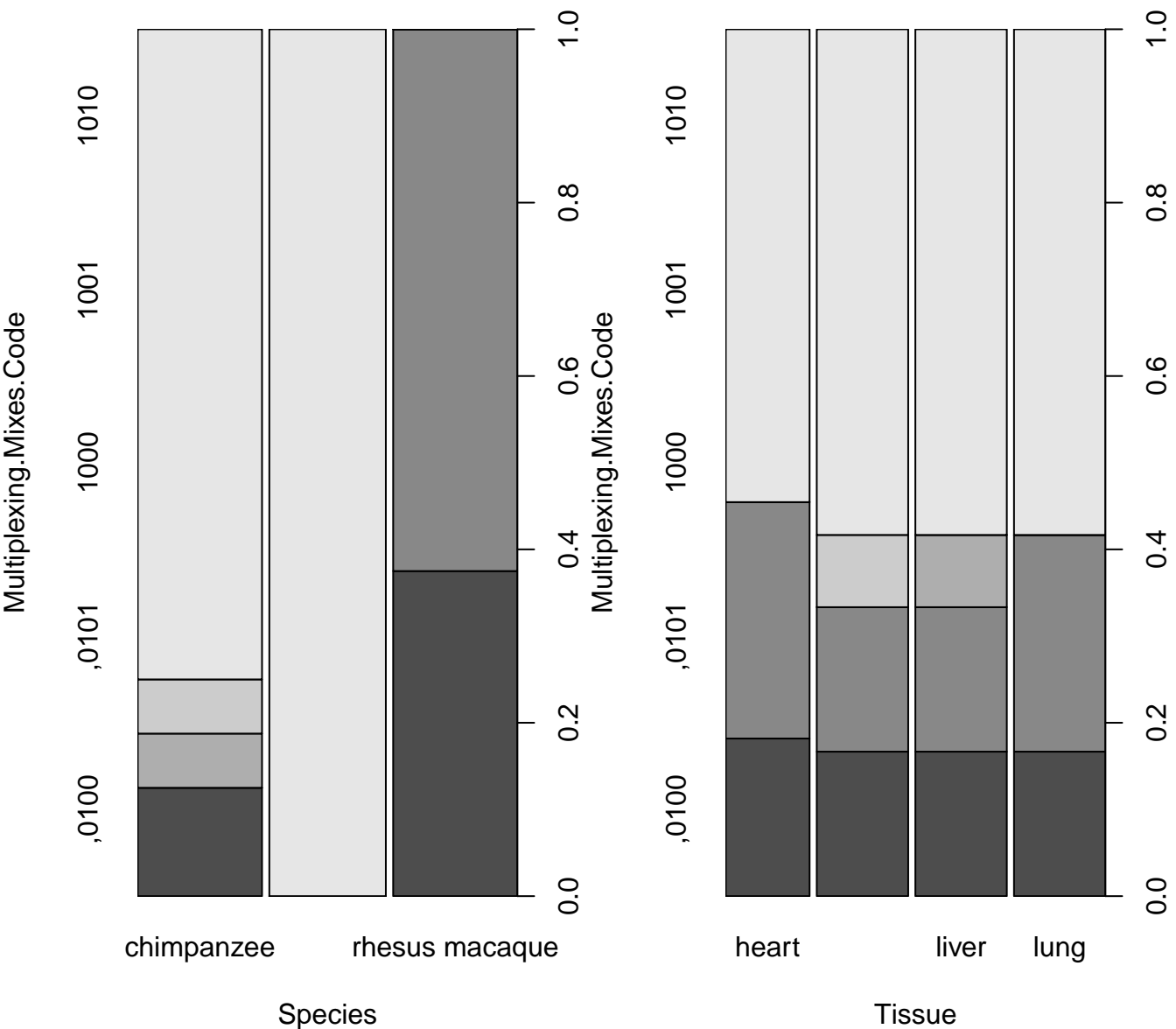
RNA.extraction.date vs. Species and Tissue



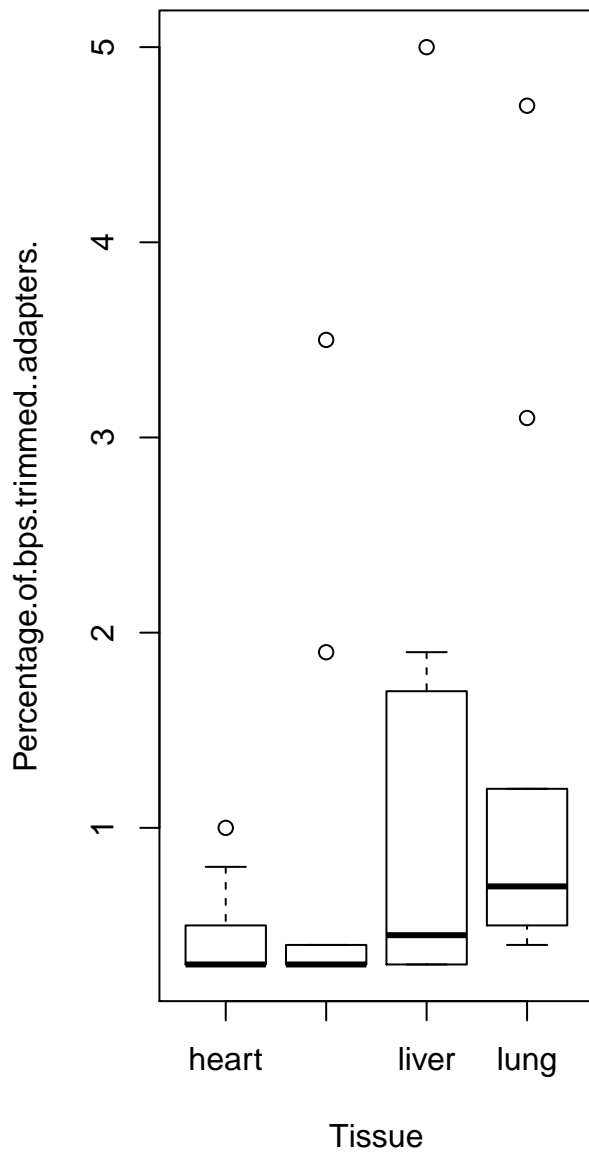
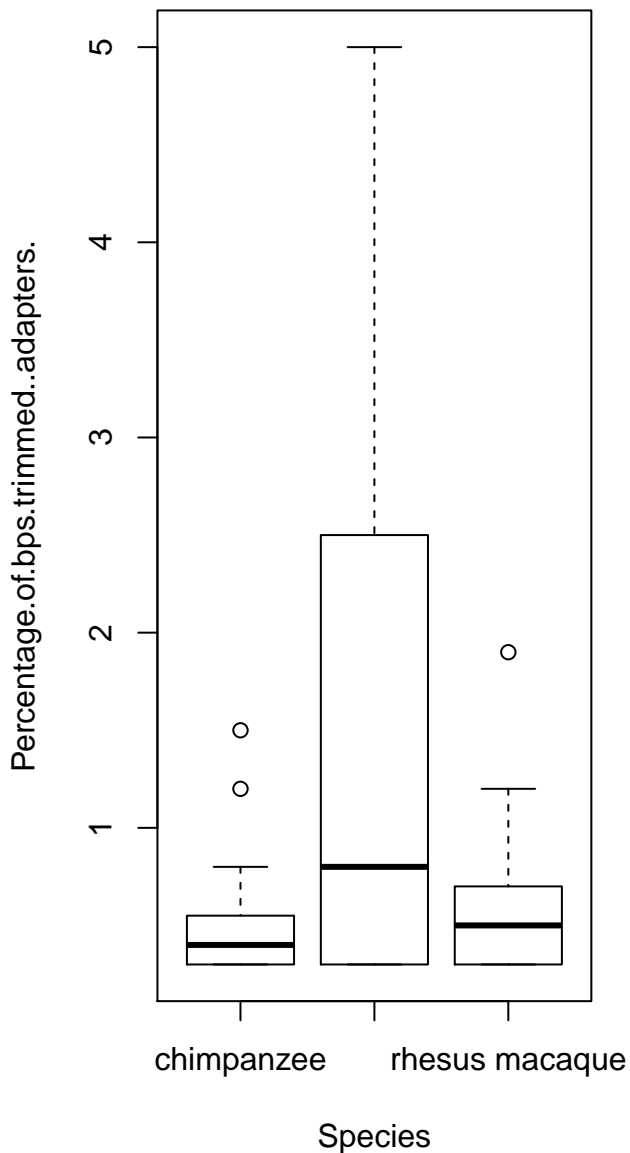
# Multiplexing.index.sequence vs. Species and Tissue



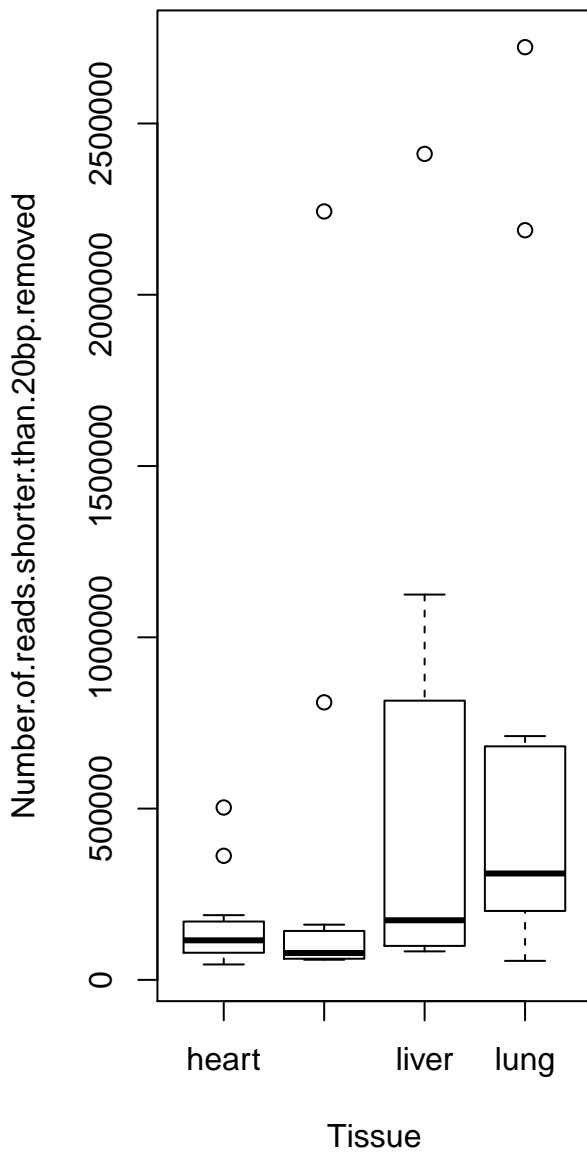
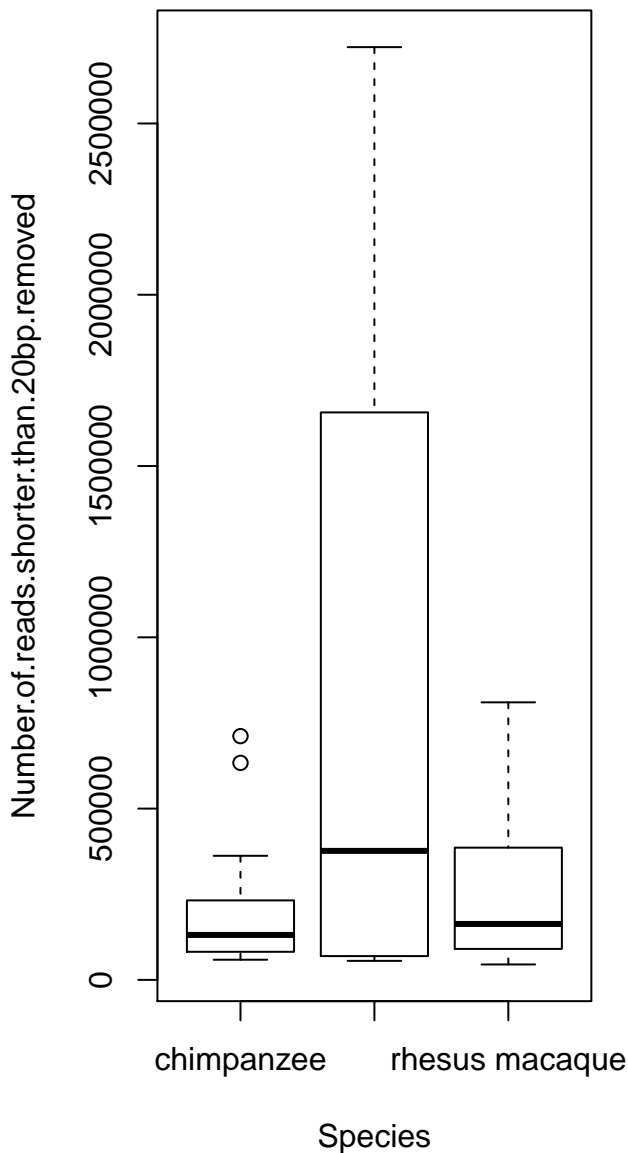
Multiplexing.Mixes.Code vs. Species and Tissue



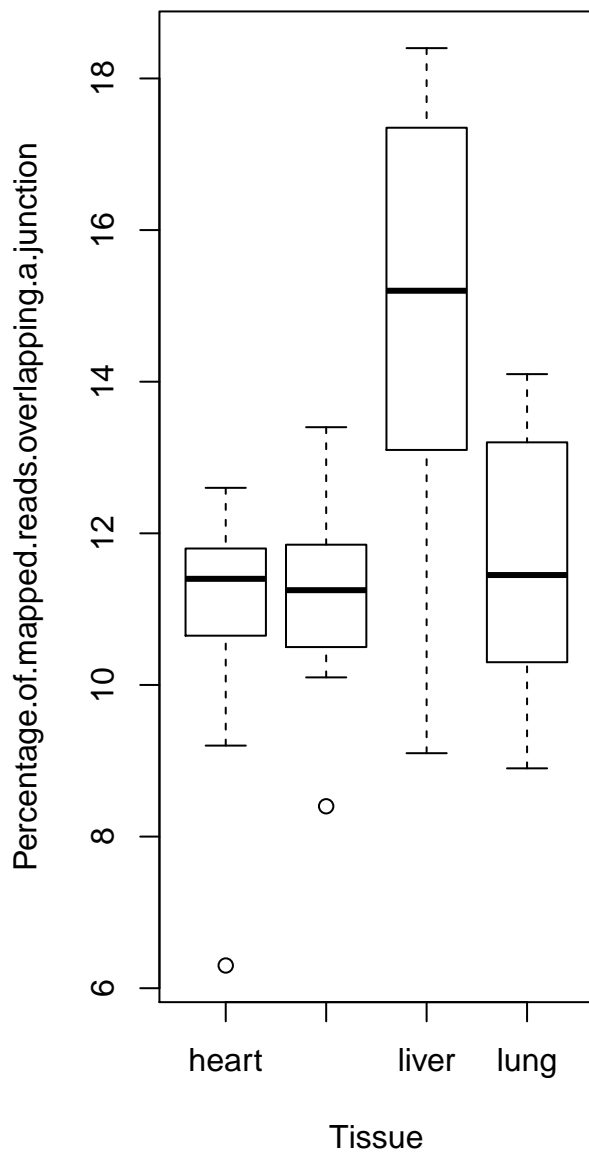
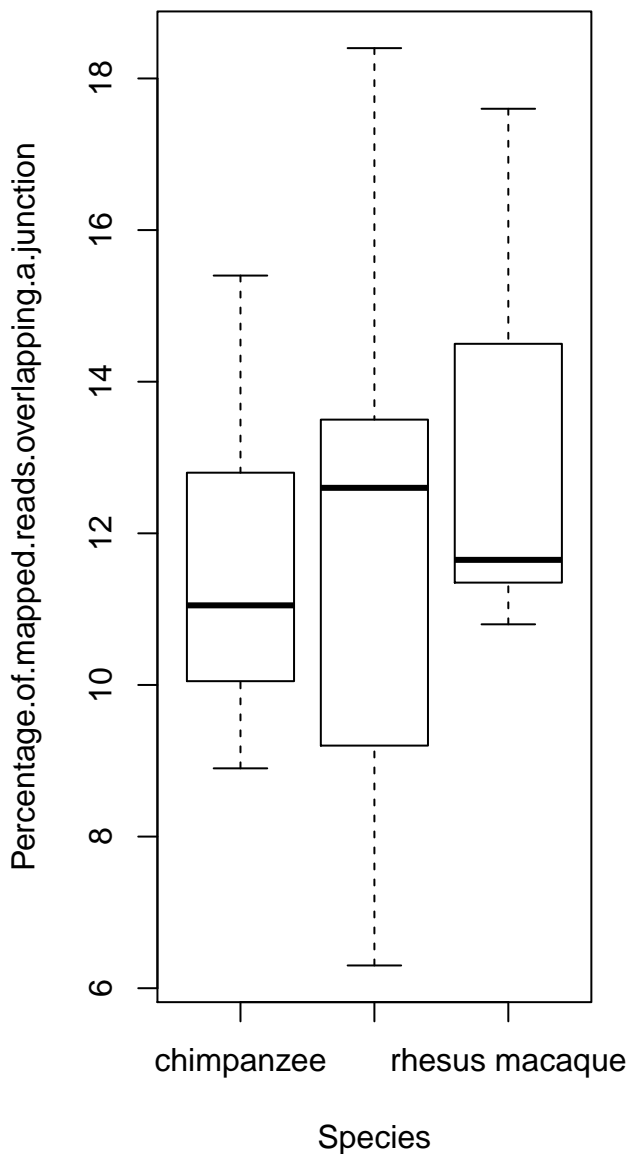
Percentage.of.bps.trimmed..adapters. vs. Species and Tissue



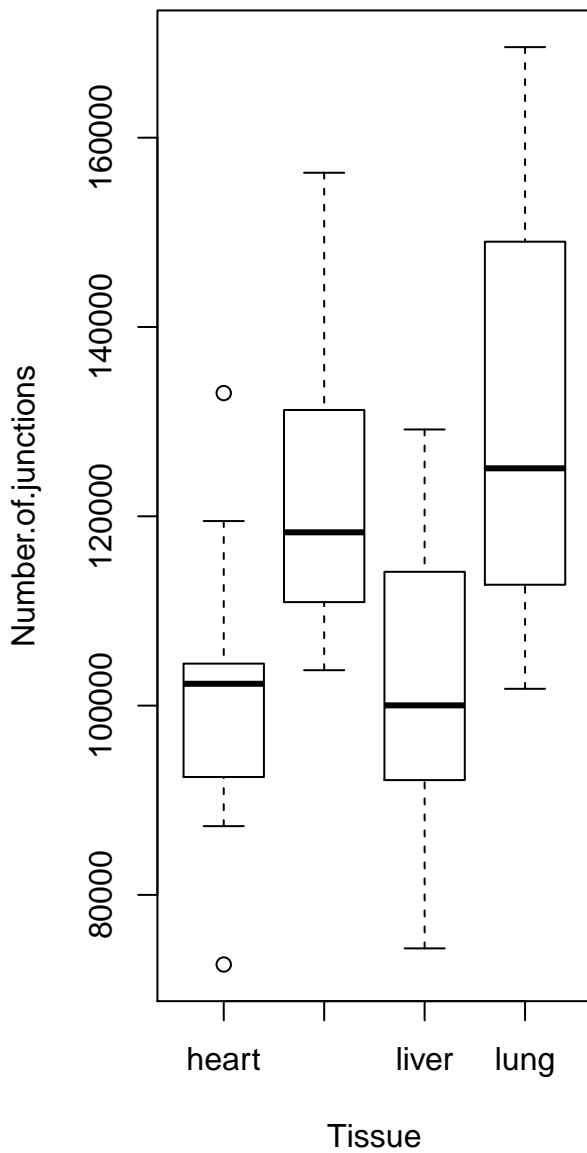
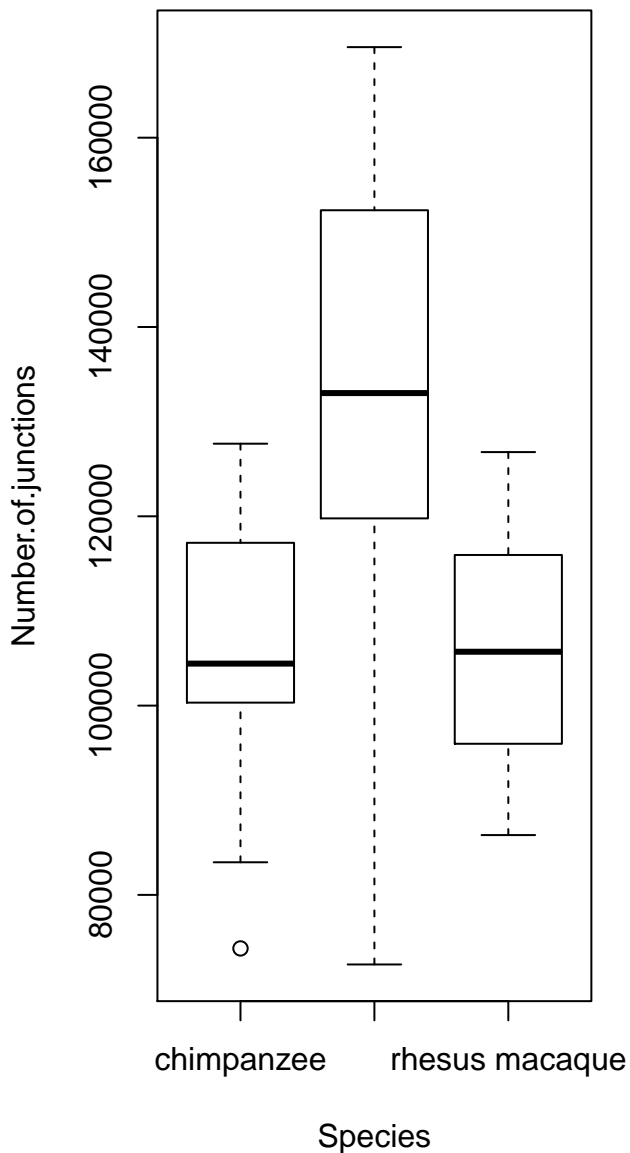
Number.of.reads.shorter.than.20bp.removed vs. Species and Tissue



Percentage.of.mapped.reads.overlapping.a.junction vs. Species and Tissue

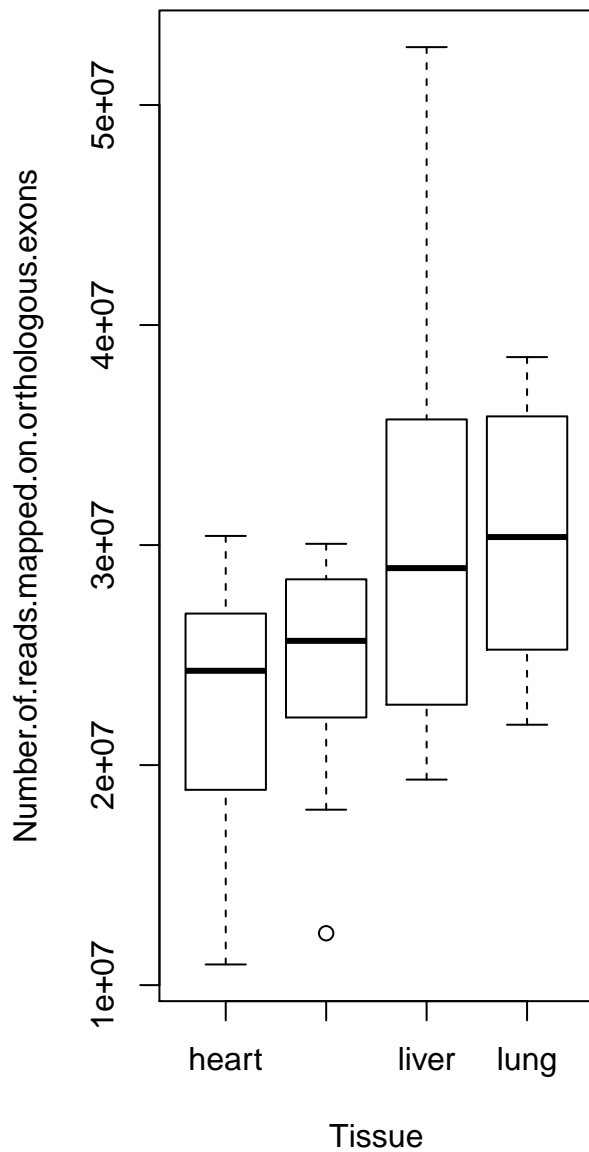
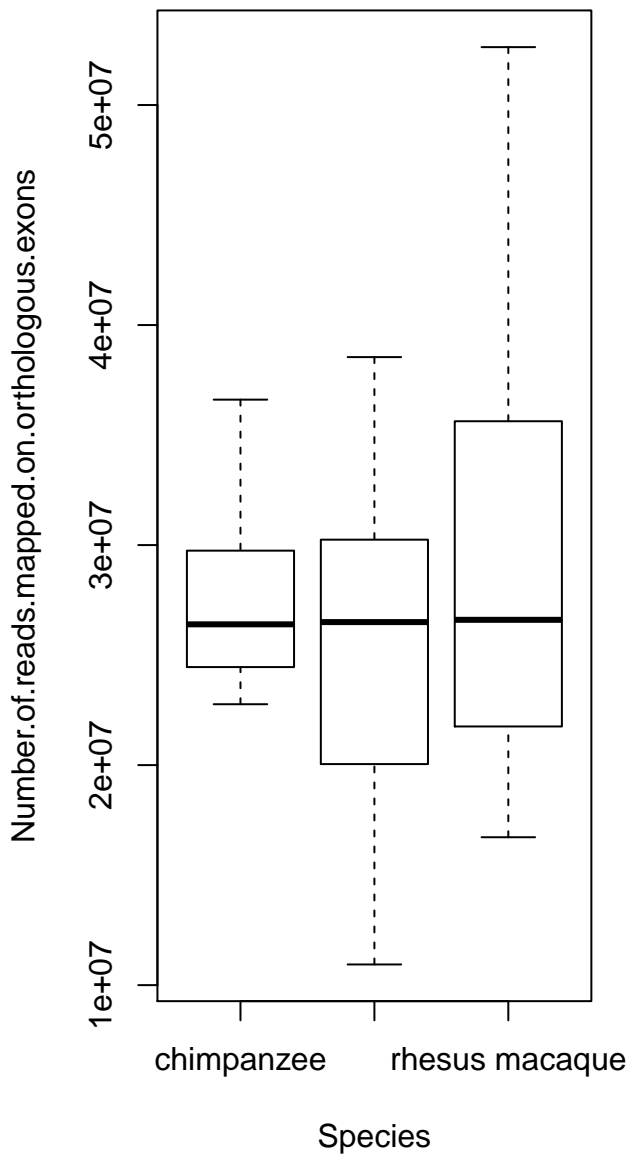


Number.of.junctions vs. Species and Tissue

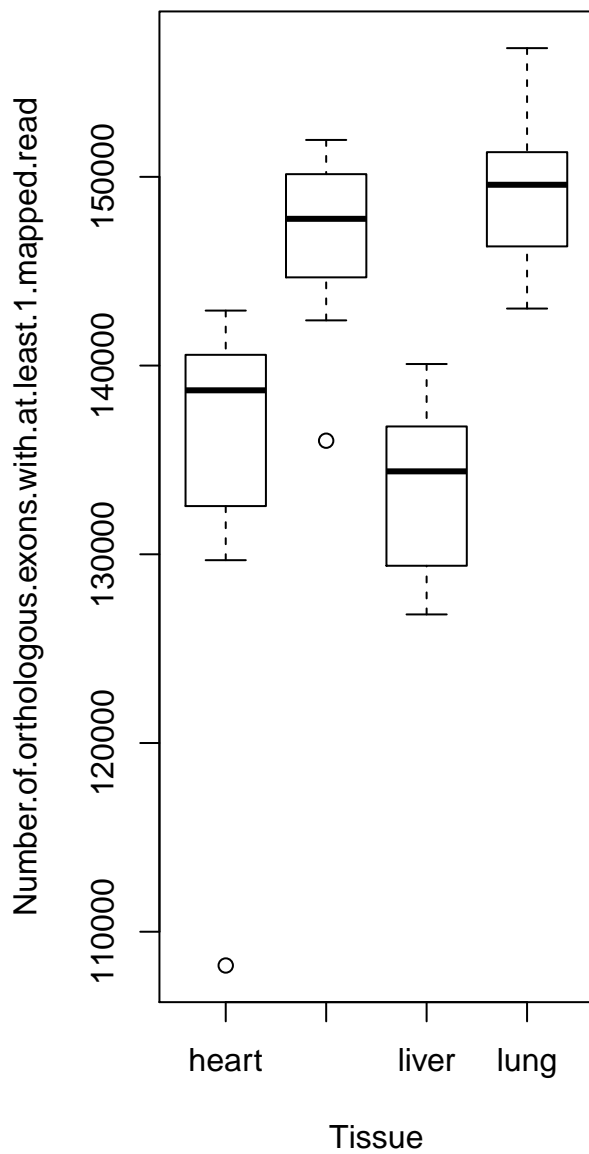
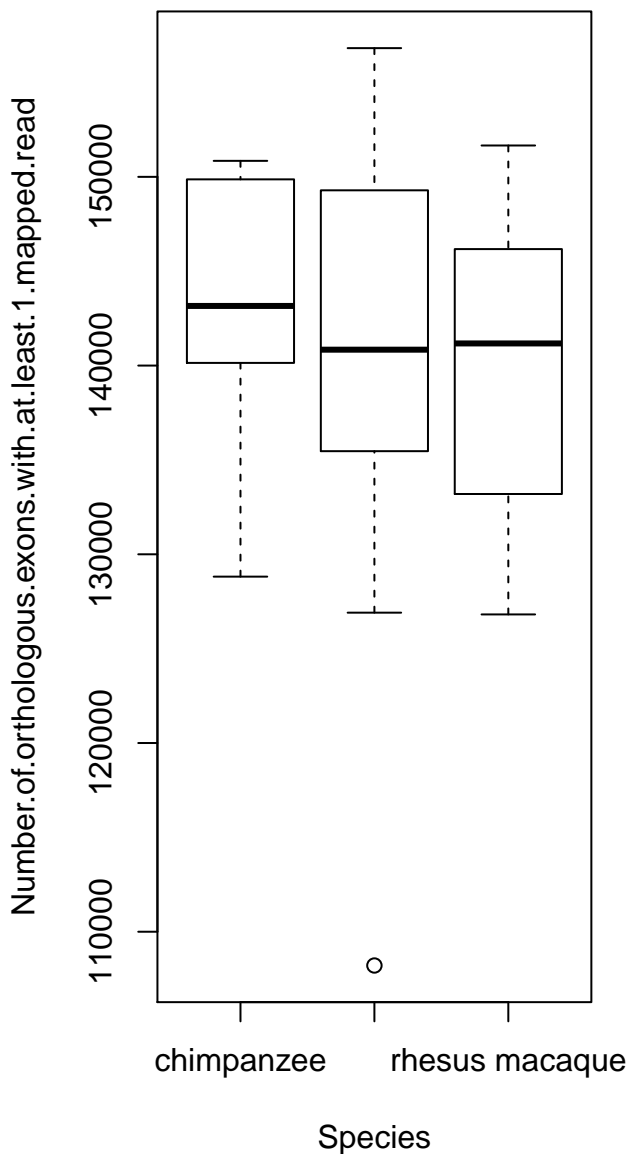




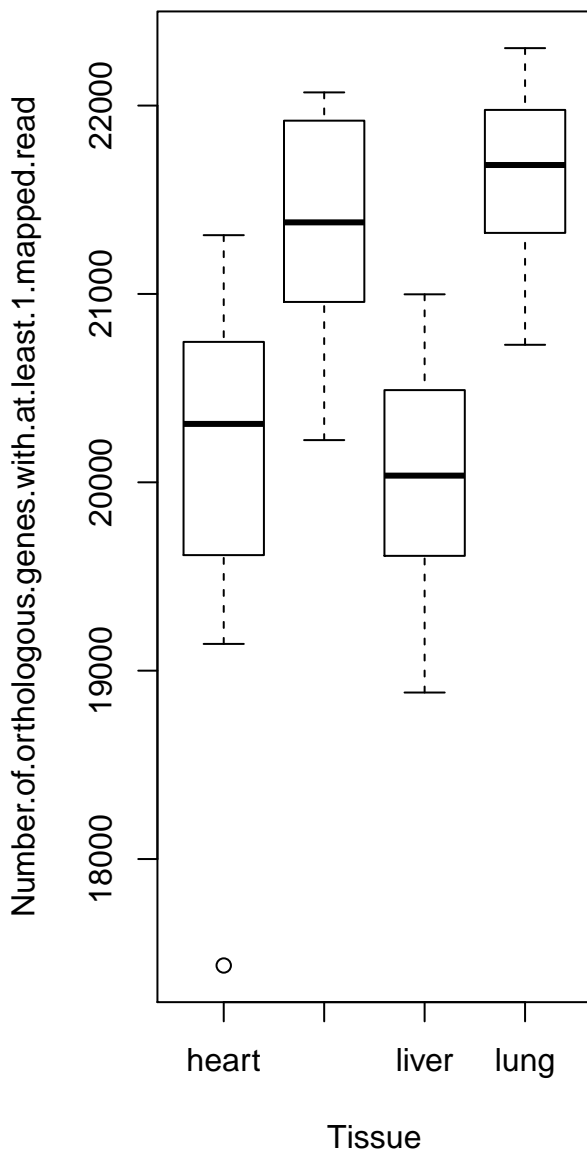
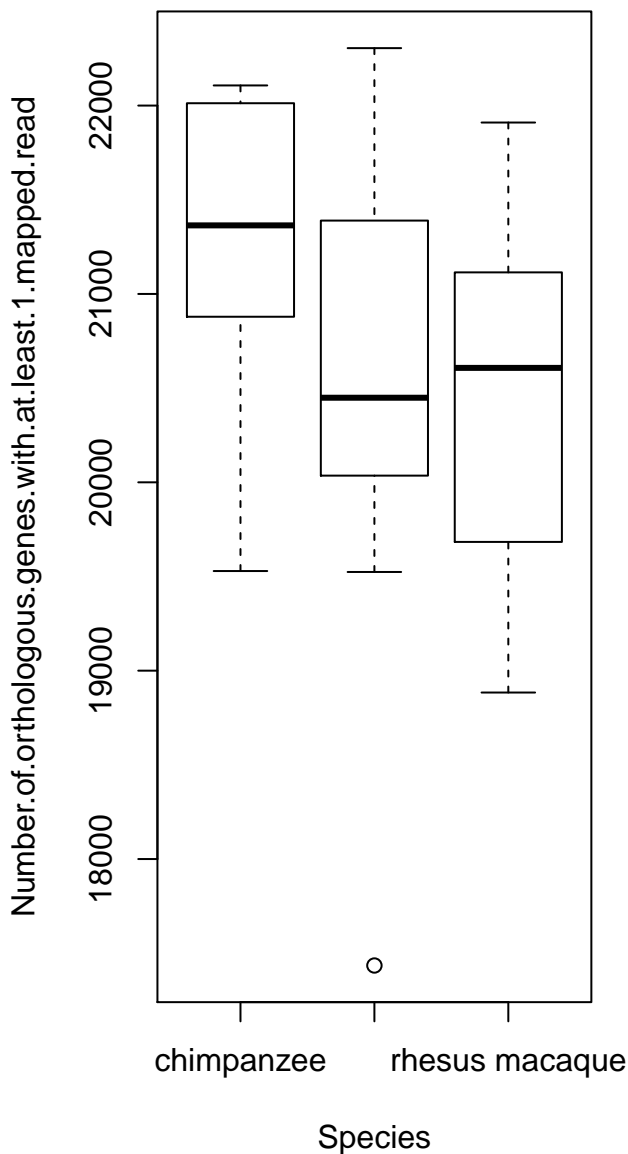
Number.of.reads.mapped.on.orthologous.exons vs. Species and Tissue



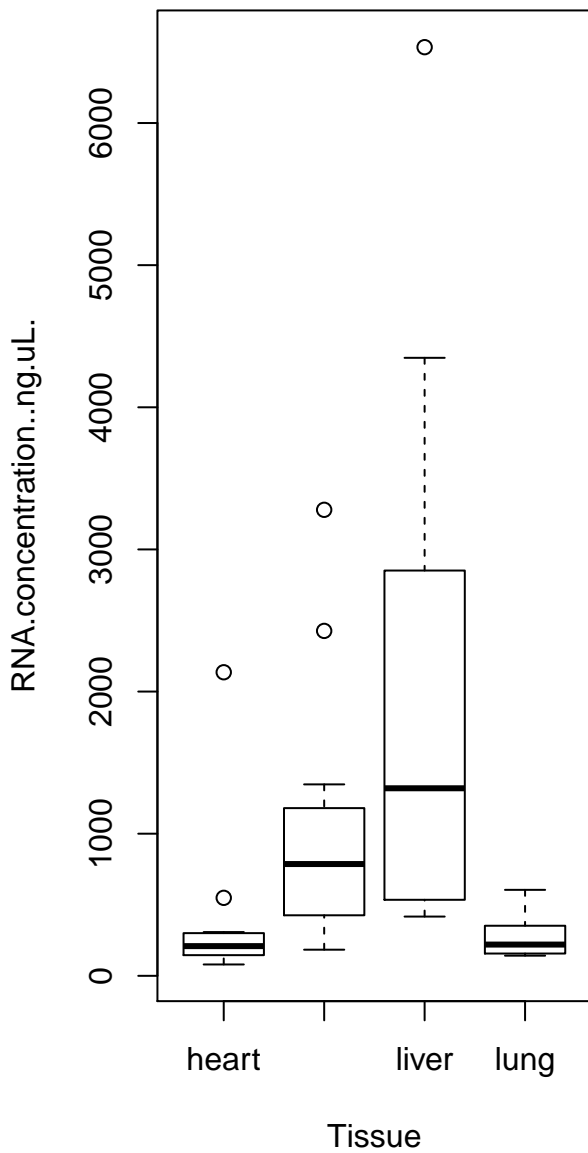
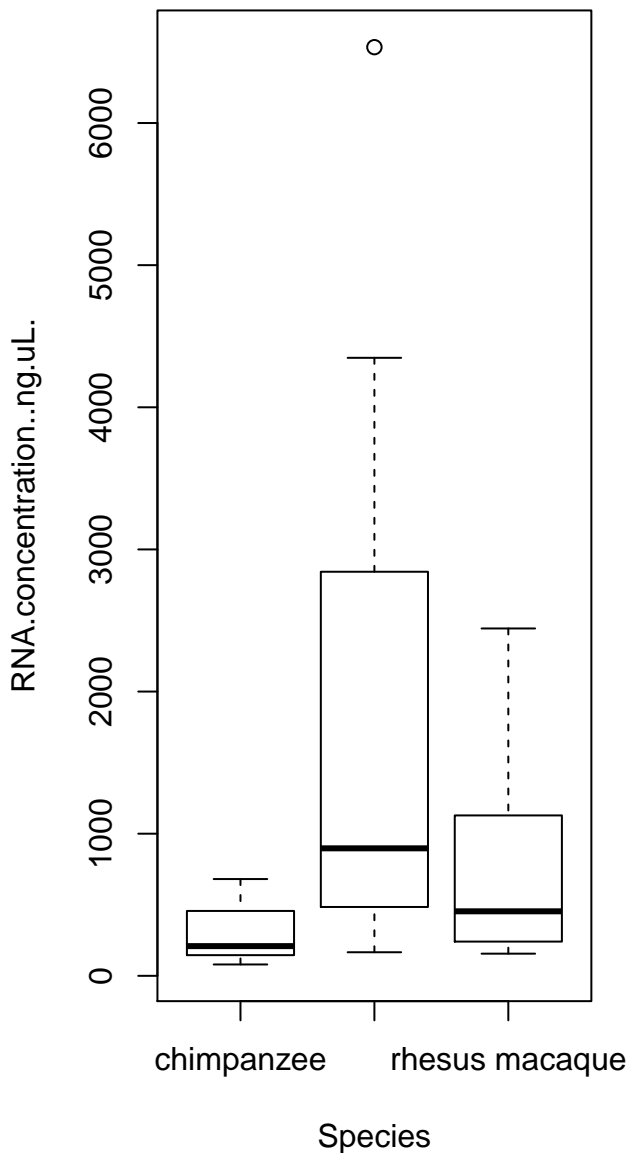
Number.of.orthologous.exons.with.at.least.1.mapped.read vs. Species and Tissue



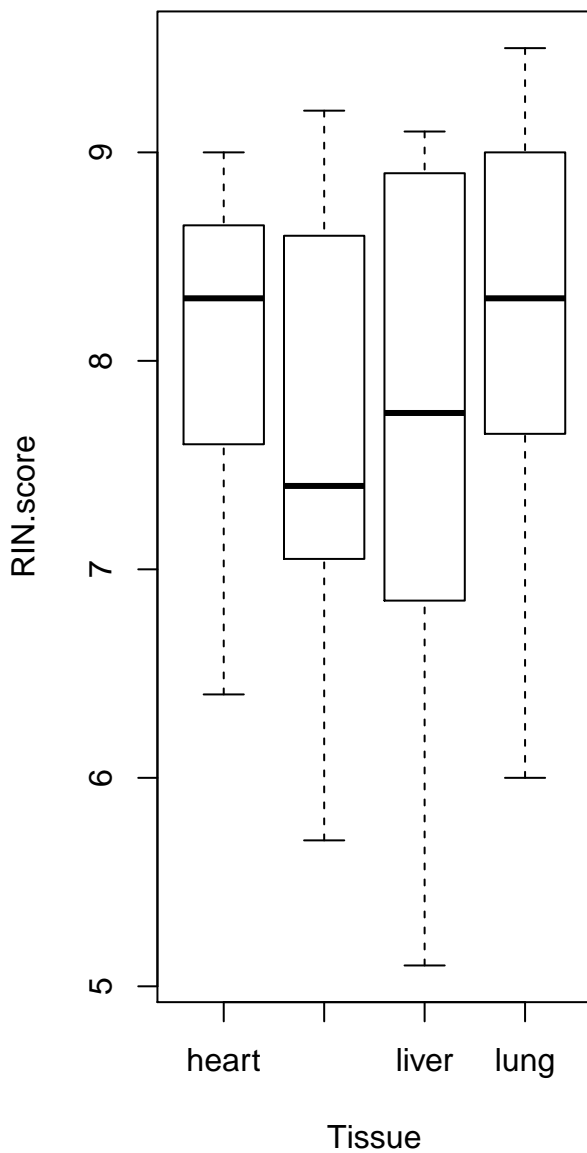
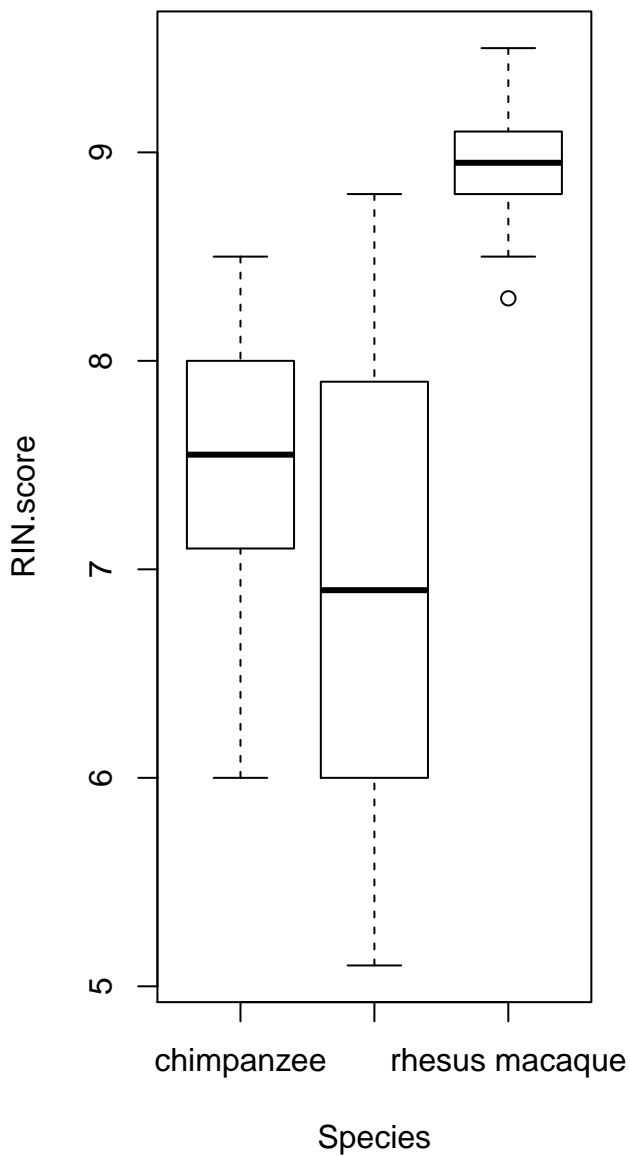
Number.of.orthologous.genes.with.at.least.1.mapped.read vs. Species and Tissue



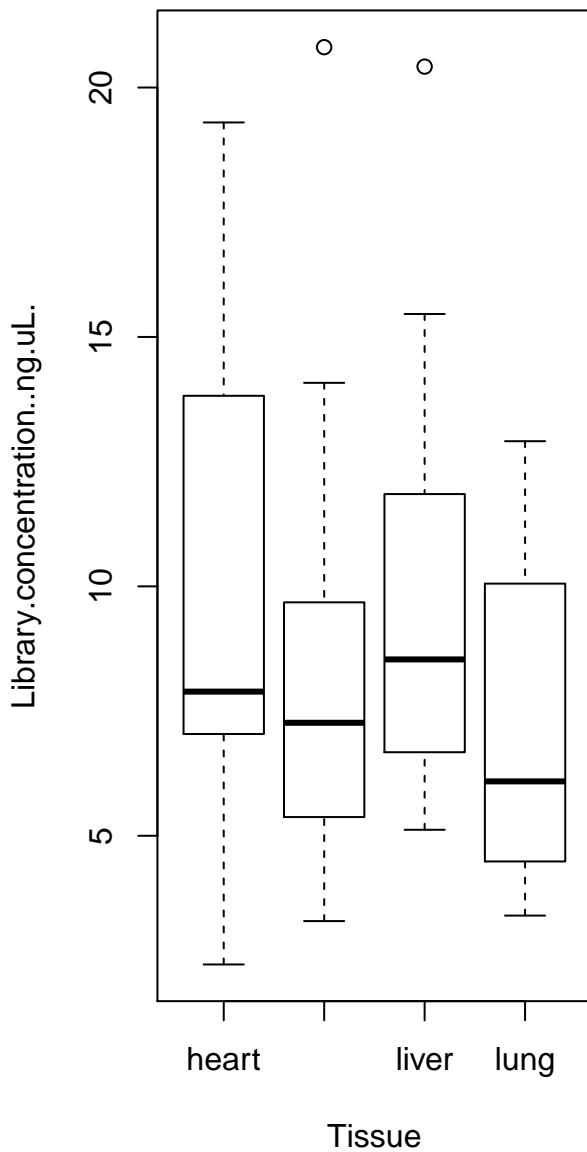
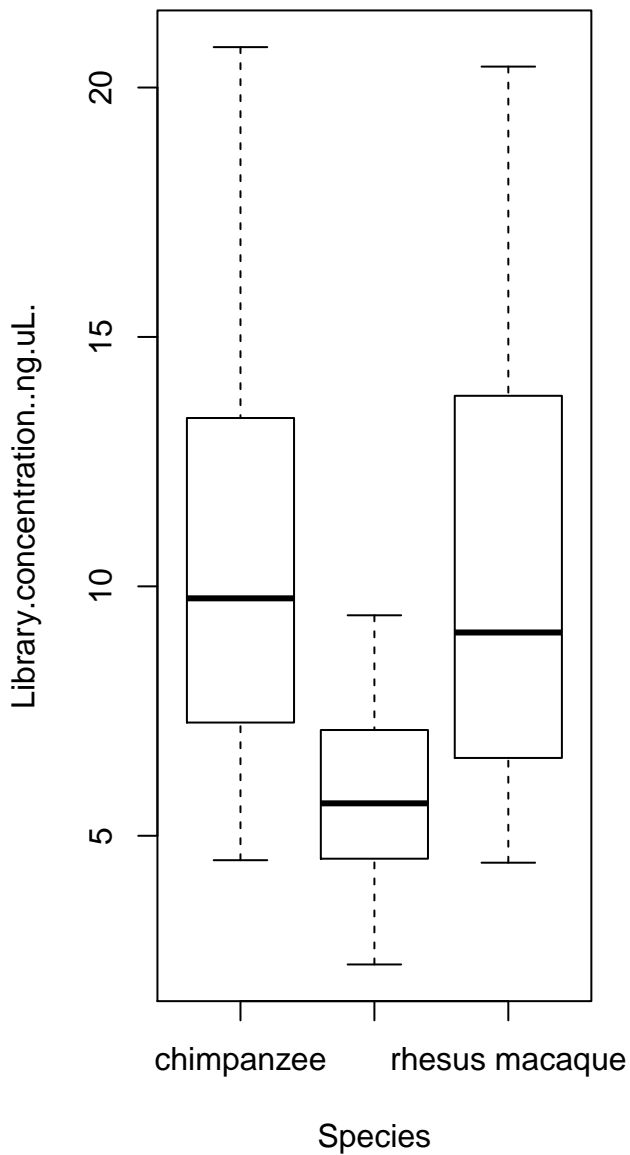
RNA.concentration..ng.uL. vs. Species and Tissue



RIN.score vs. Species and Tissue



Library.concentration..ng.uL. vs. Species and Tissue



Library.fragments.size..bp. vs. Species and Tissue

