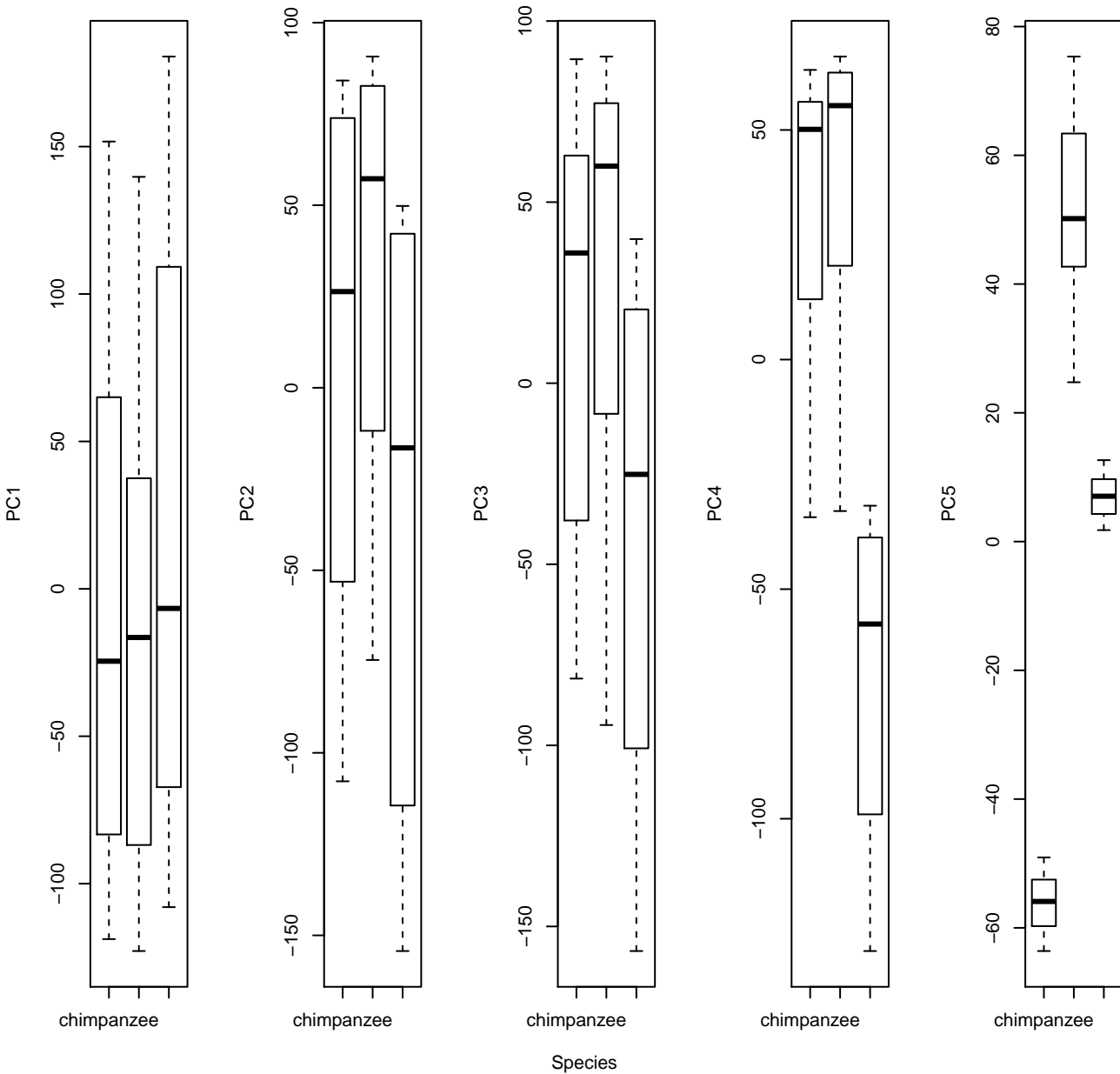
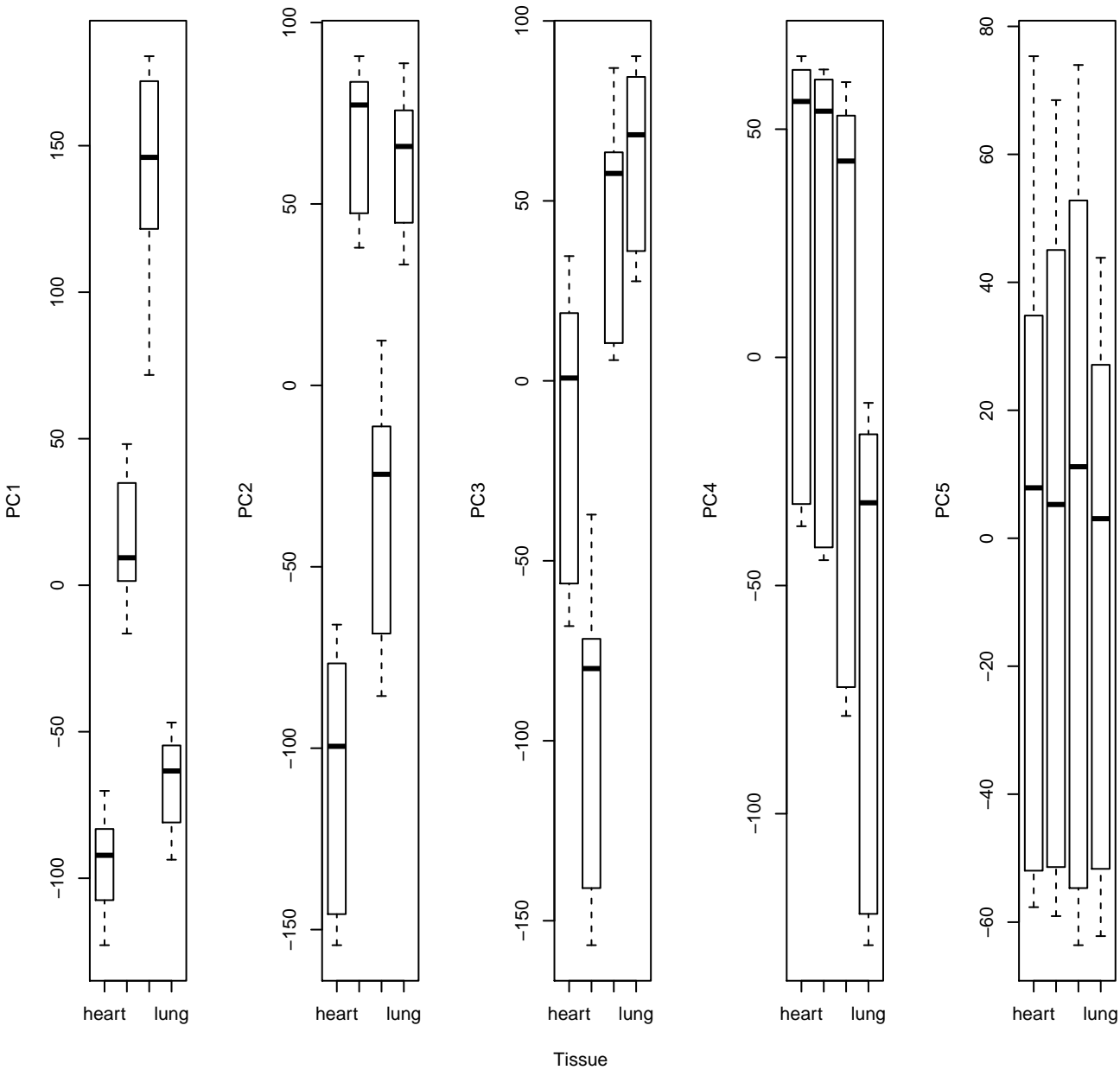


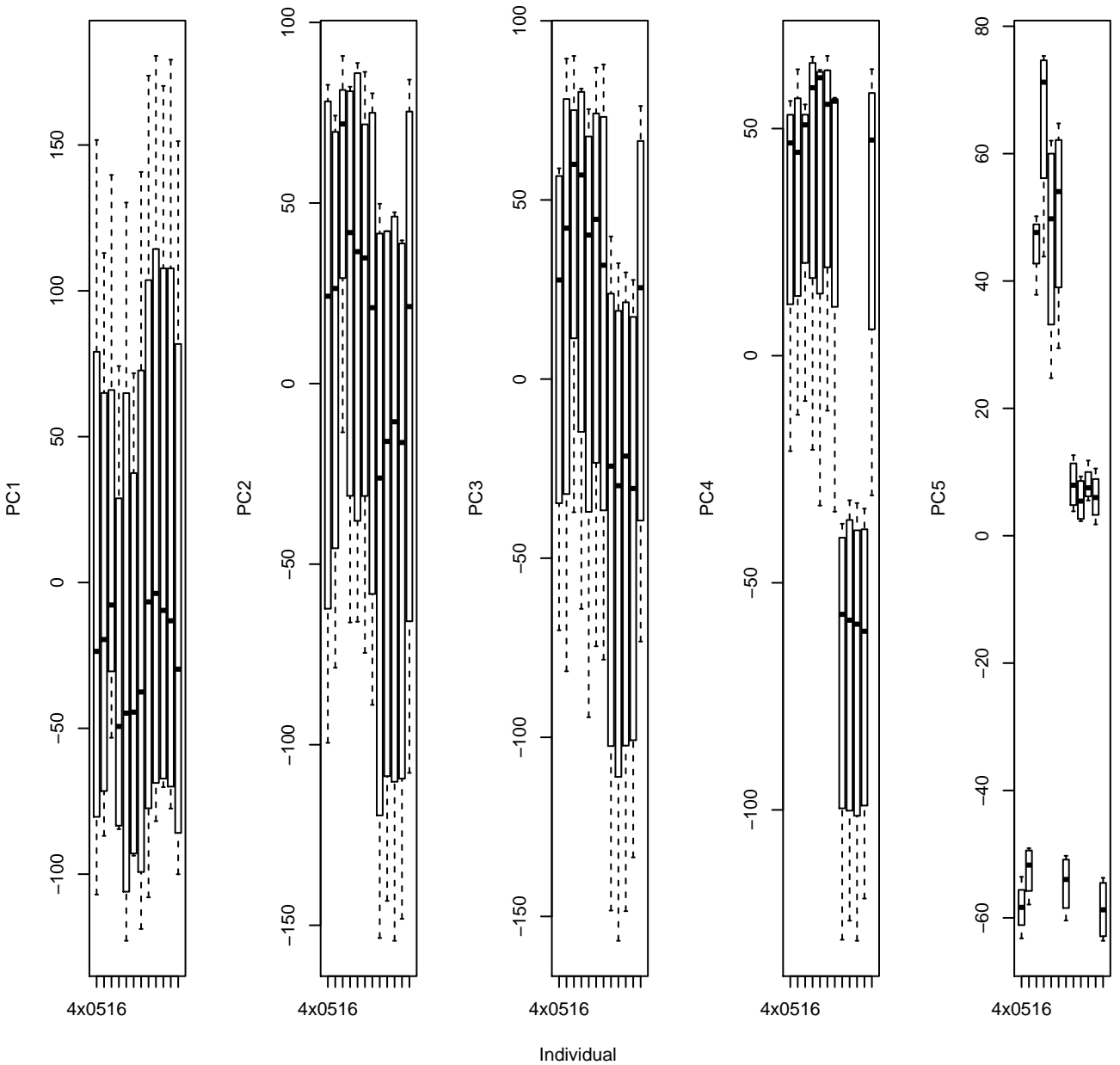
# PCs vs. Species



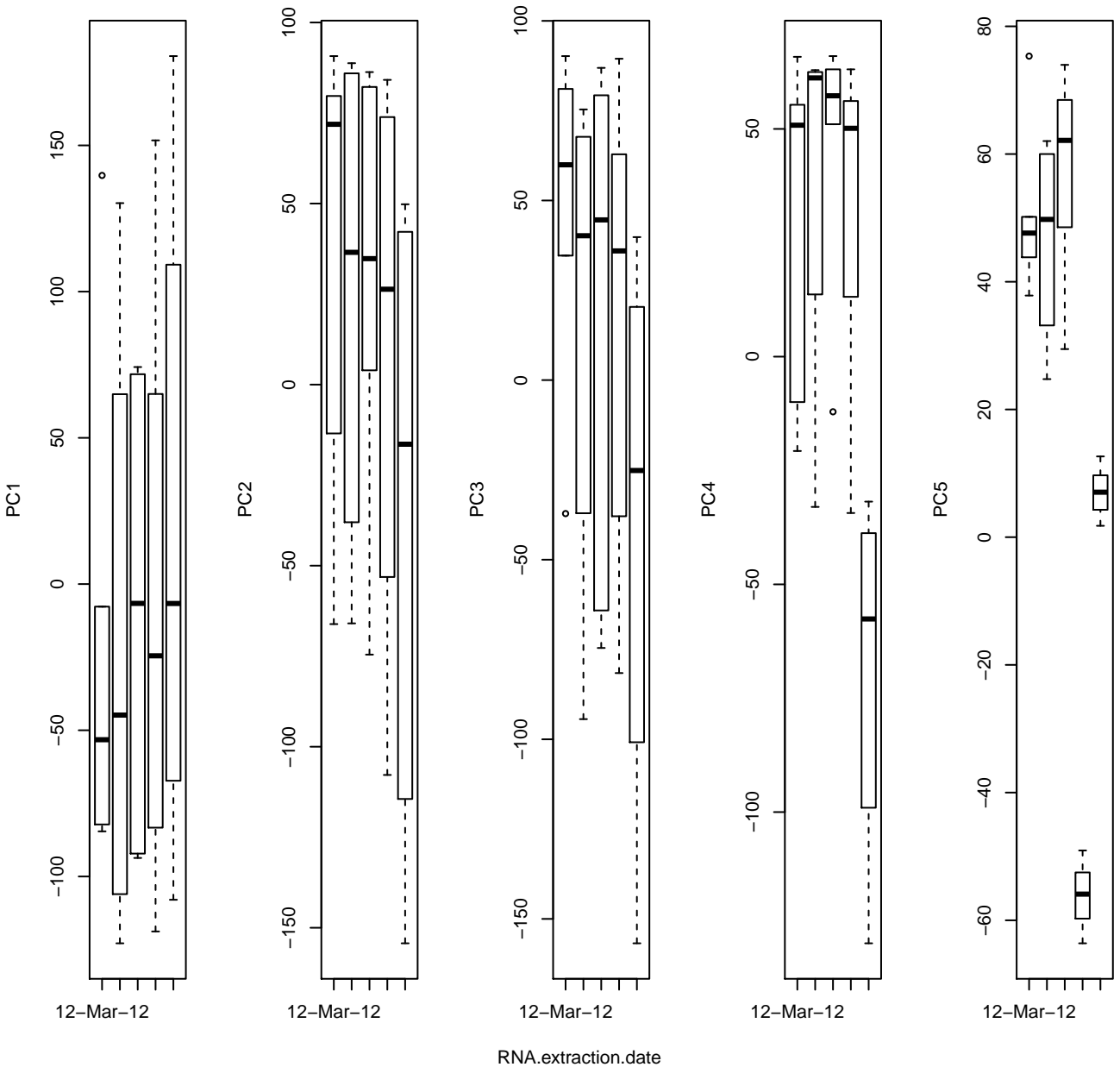
PCs vs. Tissue



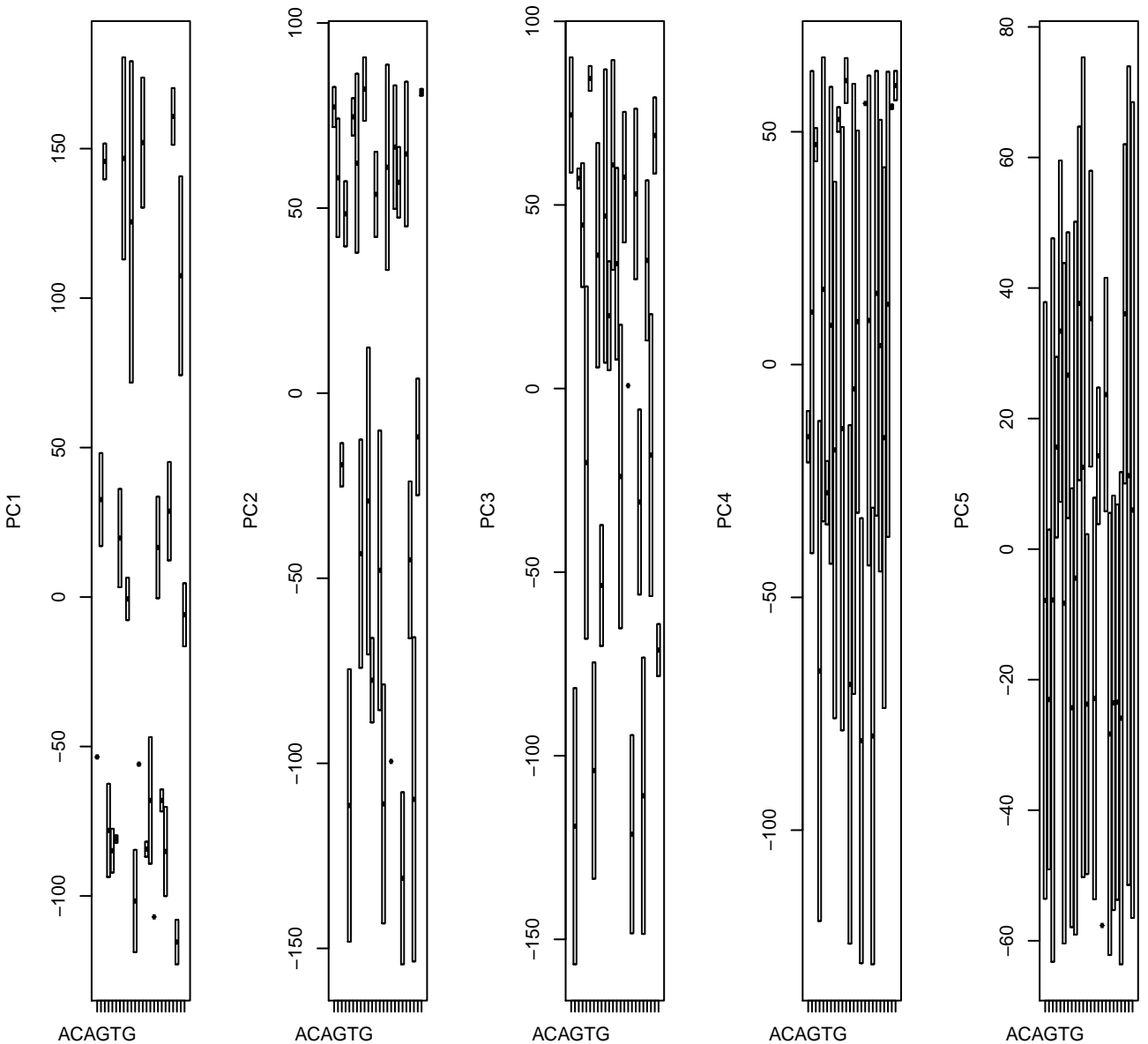
# PCs vs. Individual



PCs vs. RNA.extraction.date

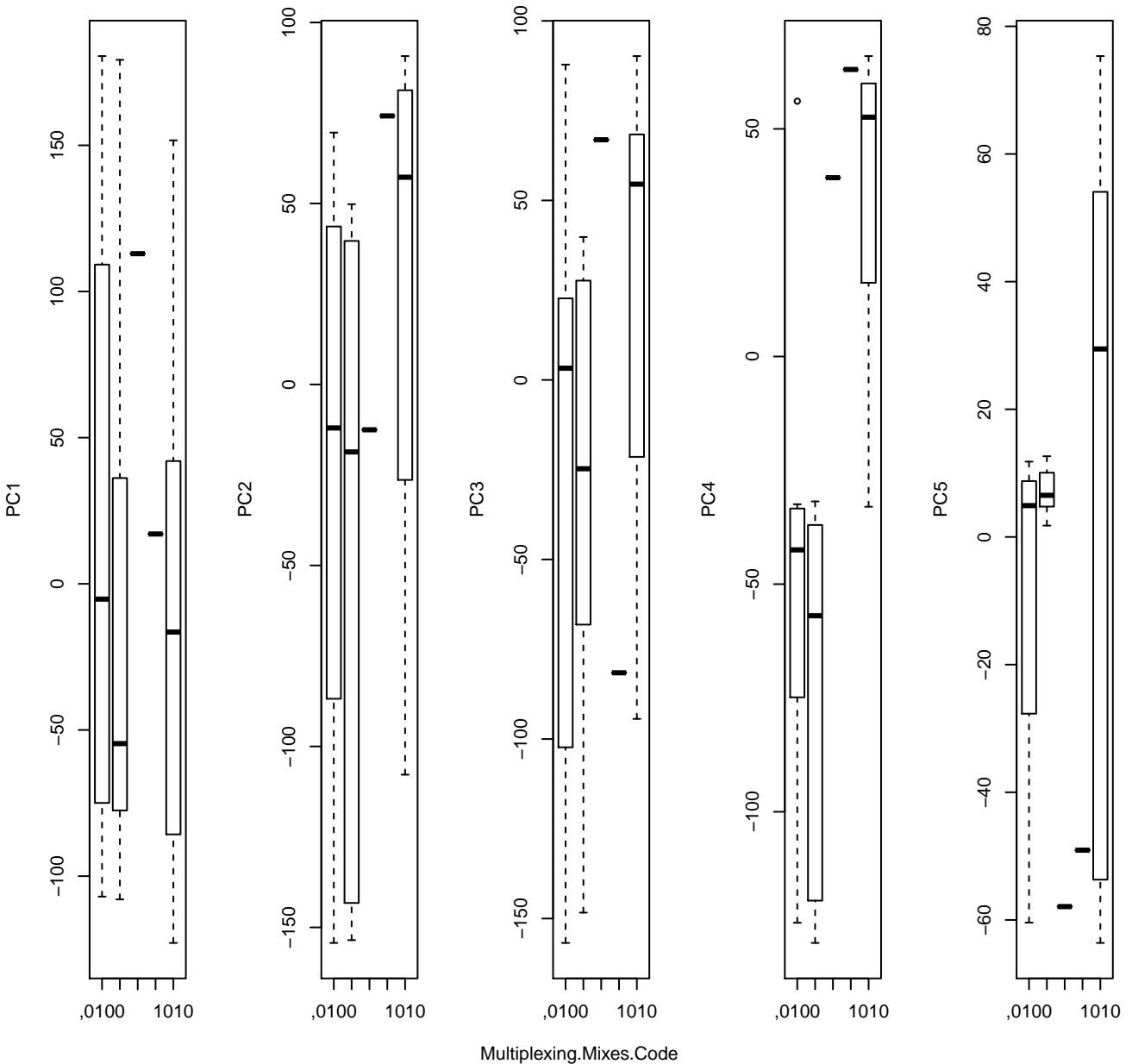


# PCs vs. Multiplexing.index.sequence

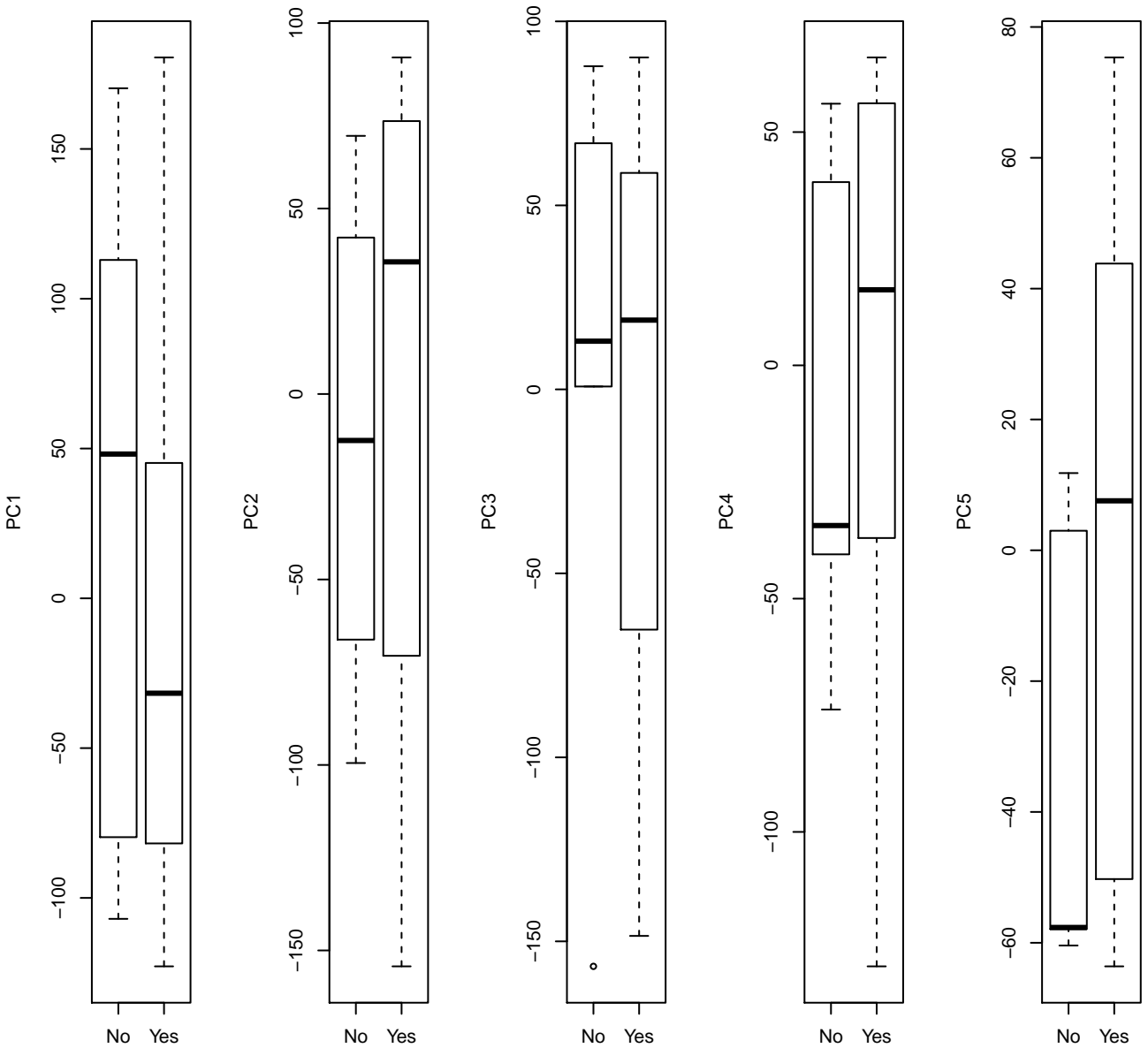


Multiplexing.index.sequence

PCs vs. Multiplexing.Mixes.Code

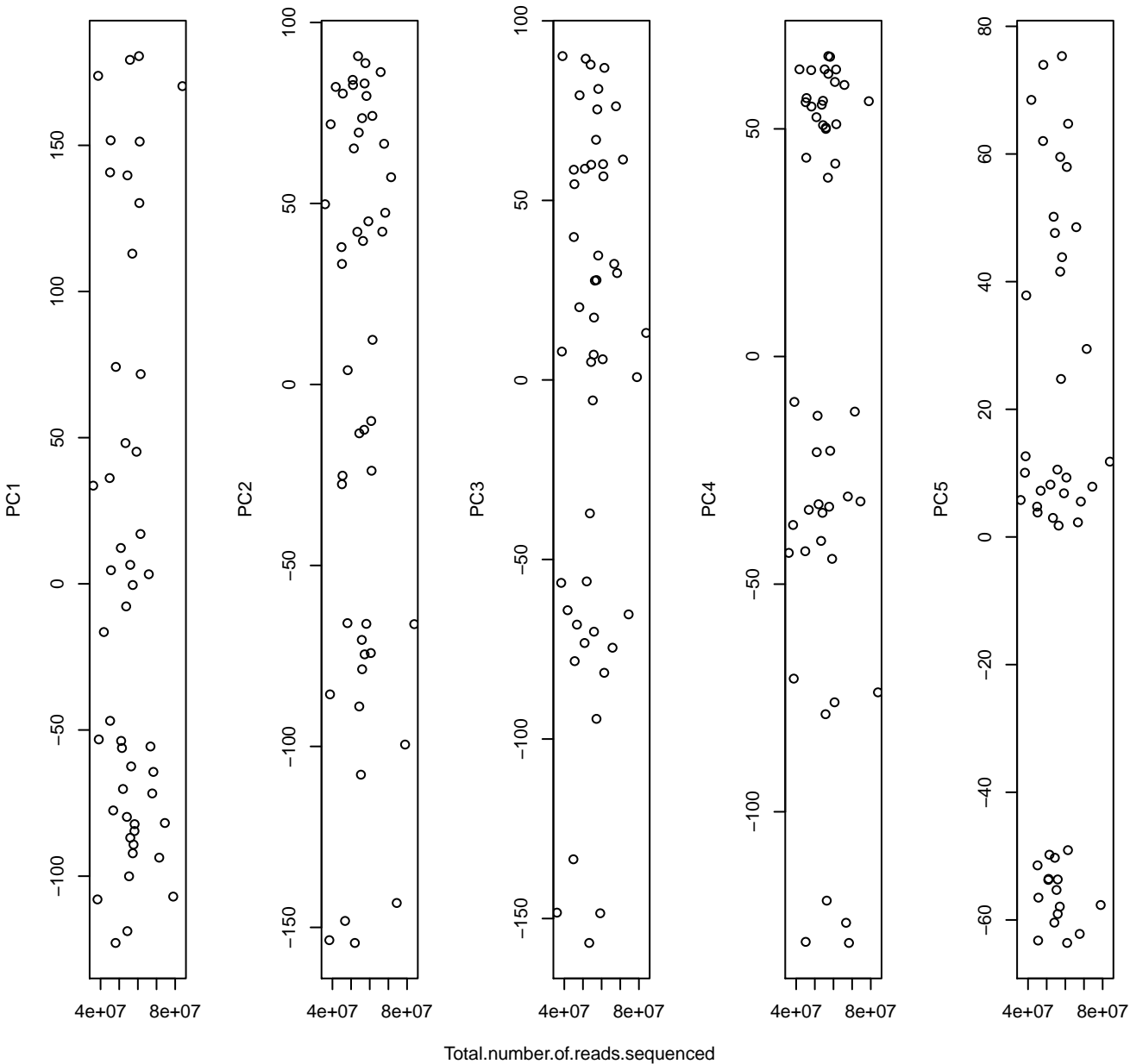


PCs vs. Sequencing.location..at.UCGF



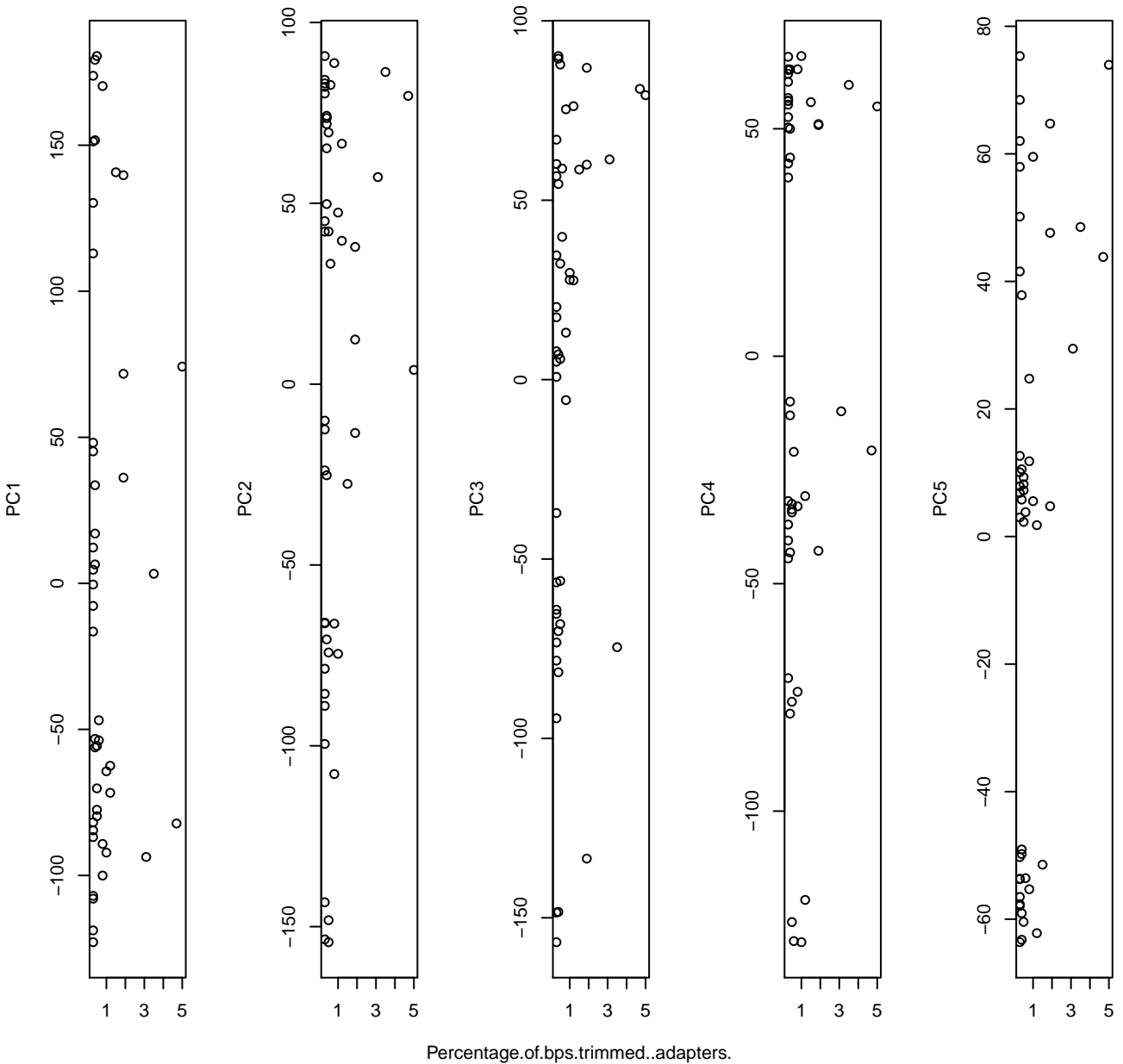
Sequencing.location..at.UCGF

PCs vs. Total.number.of.reads.sequenced

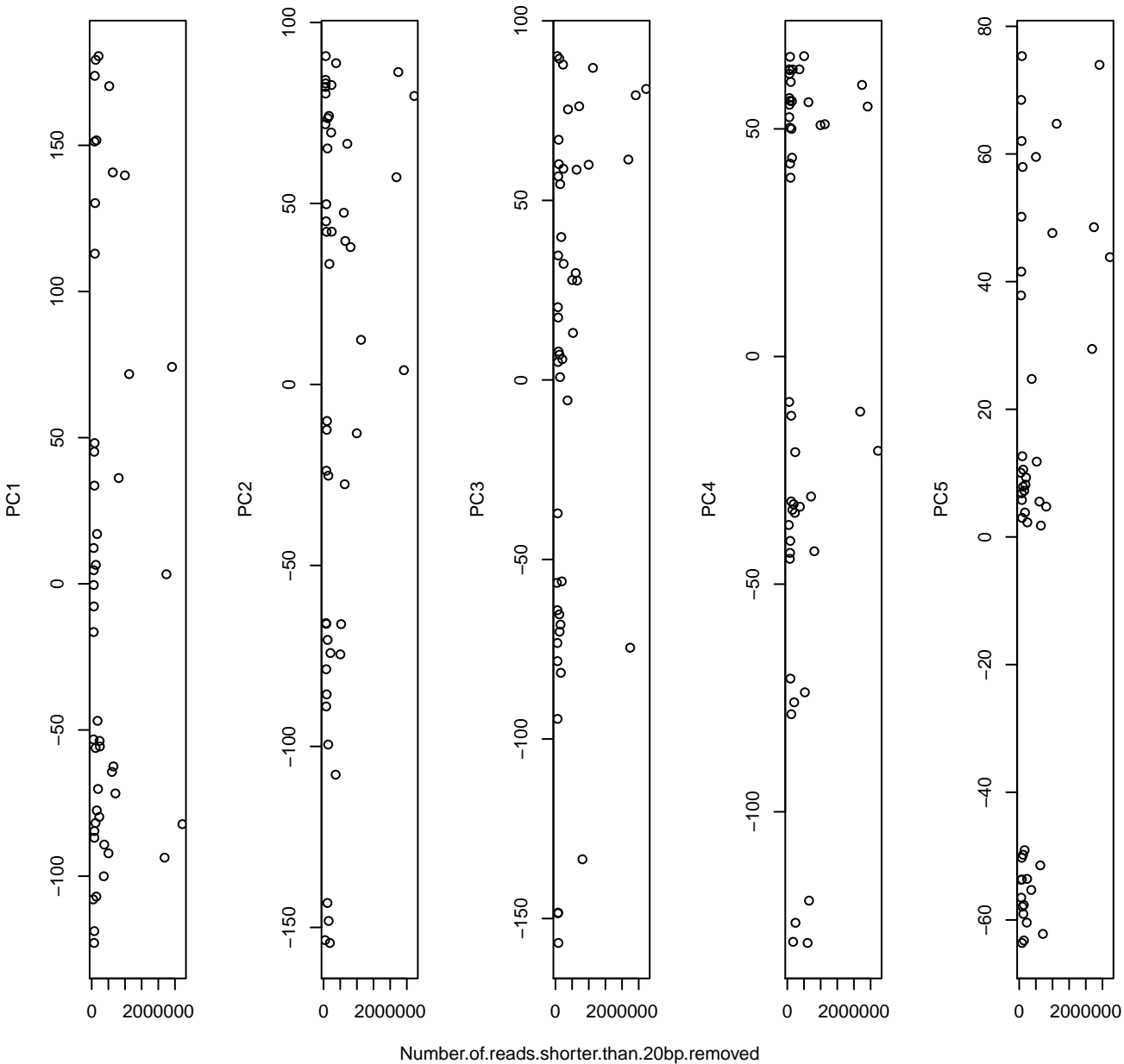




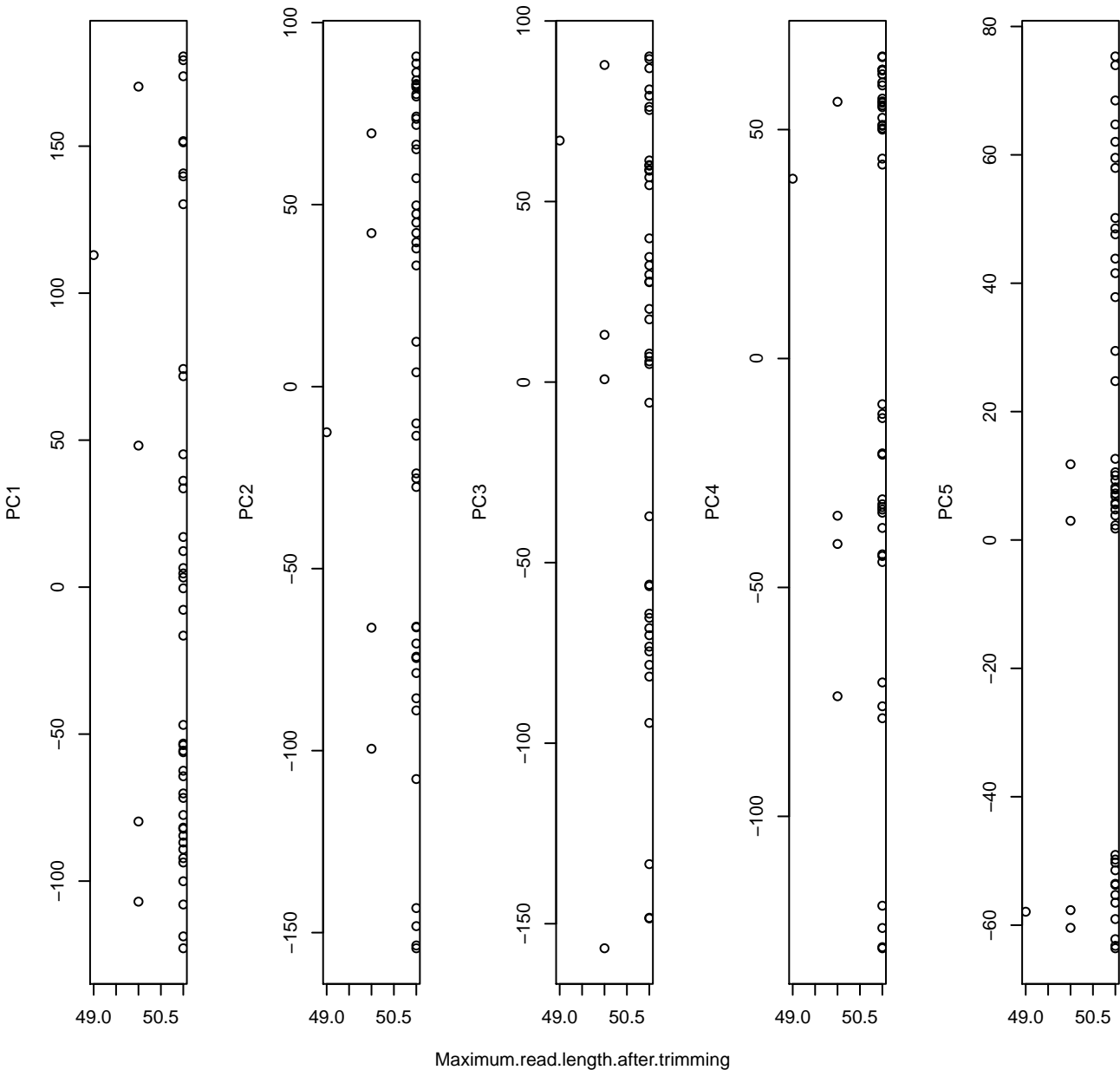
PCs vs. Percentage.of.bps.trimmed..adapters.



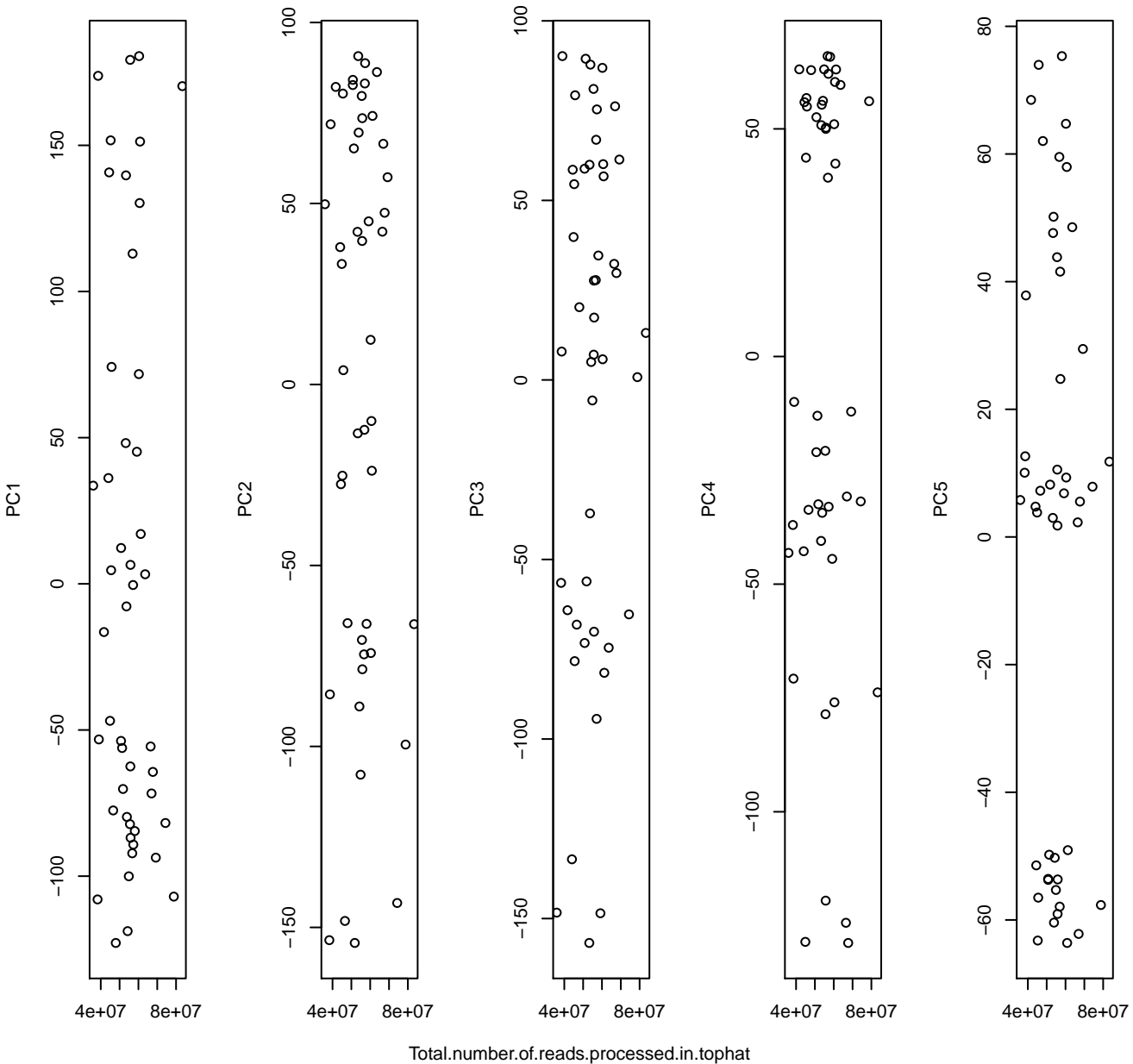
PCs vs. Number.of.reads.shorter.than.20bp.removed



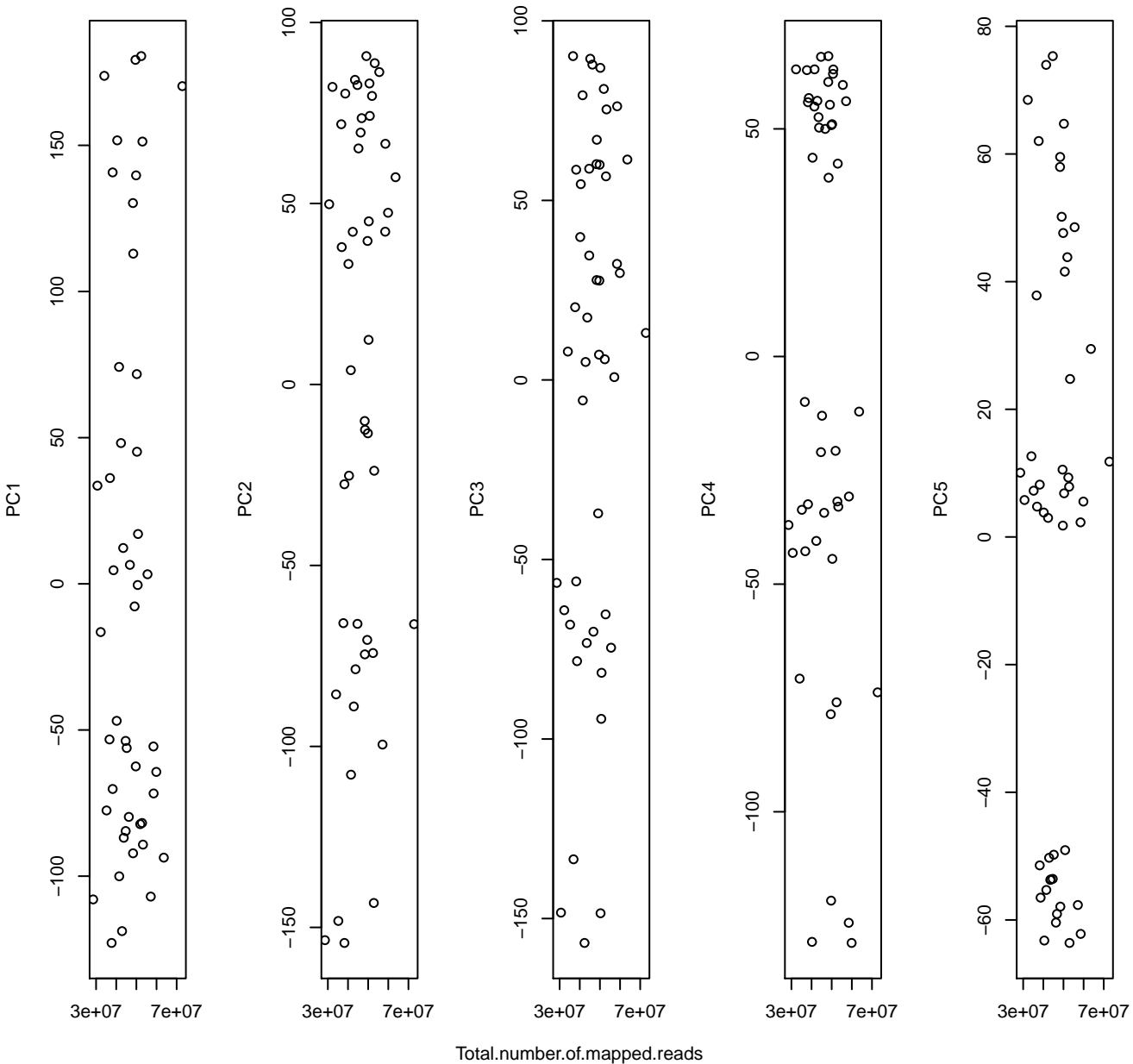
PCs vs. Maximum.read.length.after.trimming



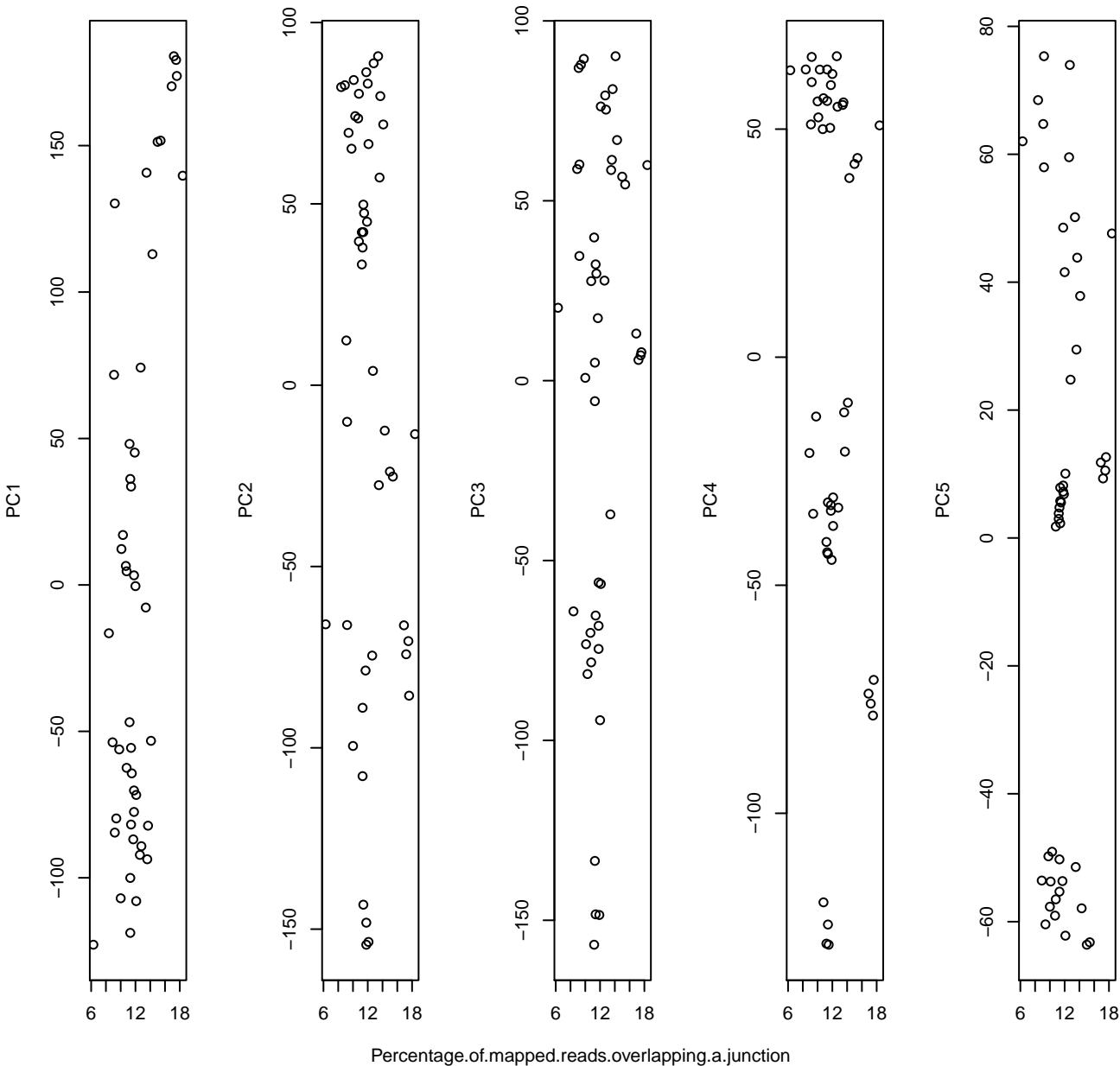
PCs vs. Total.number.of.reads.processed.in.tophat



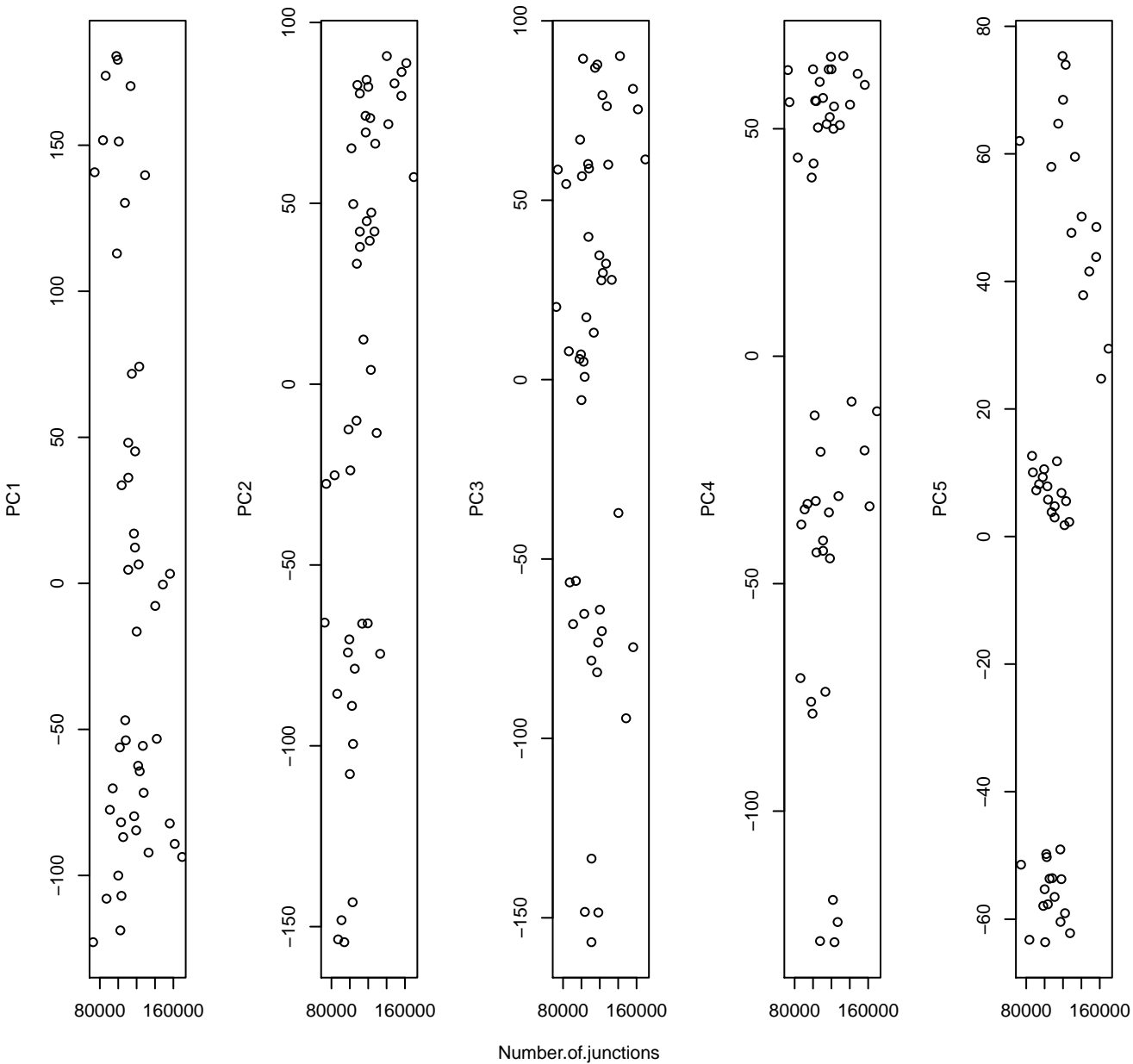
PCs vs. Total.number.of.mapped.reads



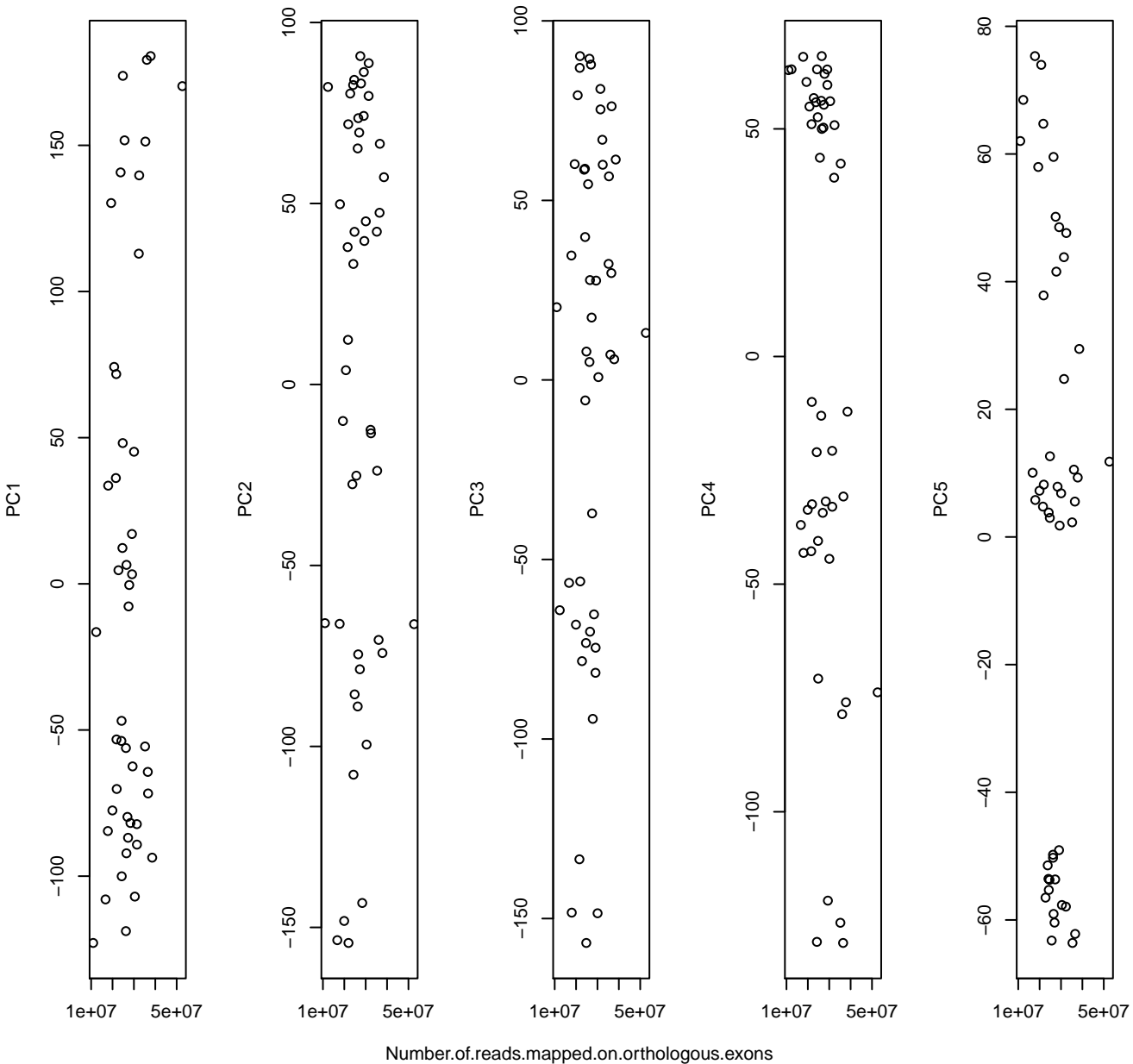
PCs vs. Percentage.of.mapped.reads.overlapping.a.junction



PCs vs. Number.of.junctions

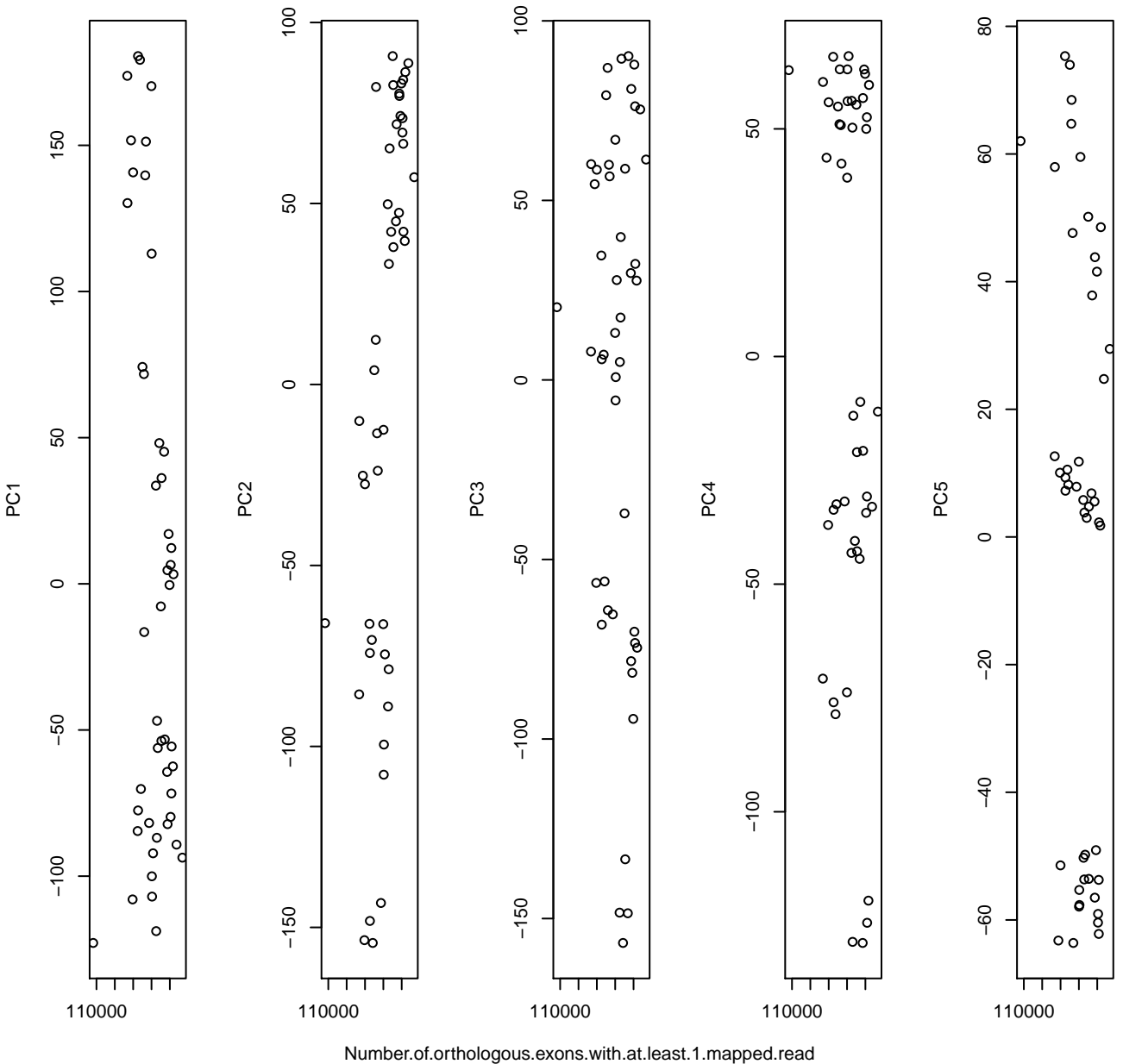


PCs vs. Number.of.reads.mapped.on.orthologous.exons

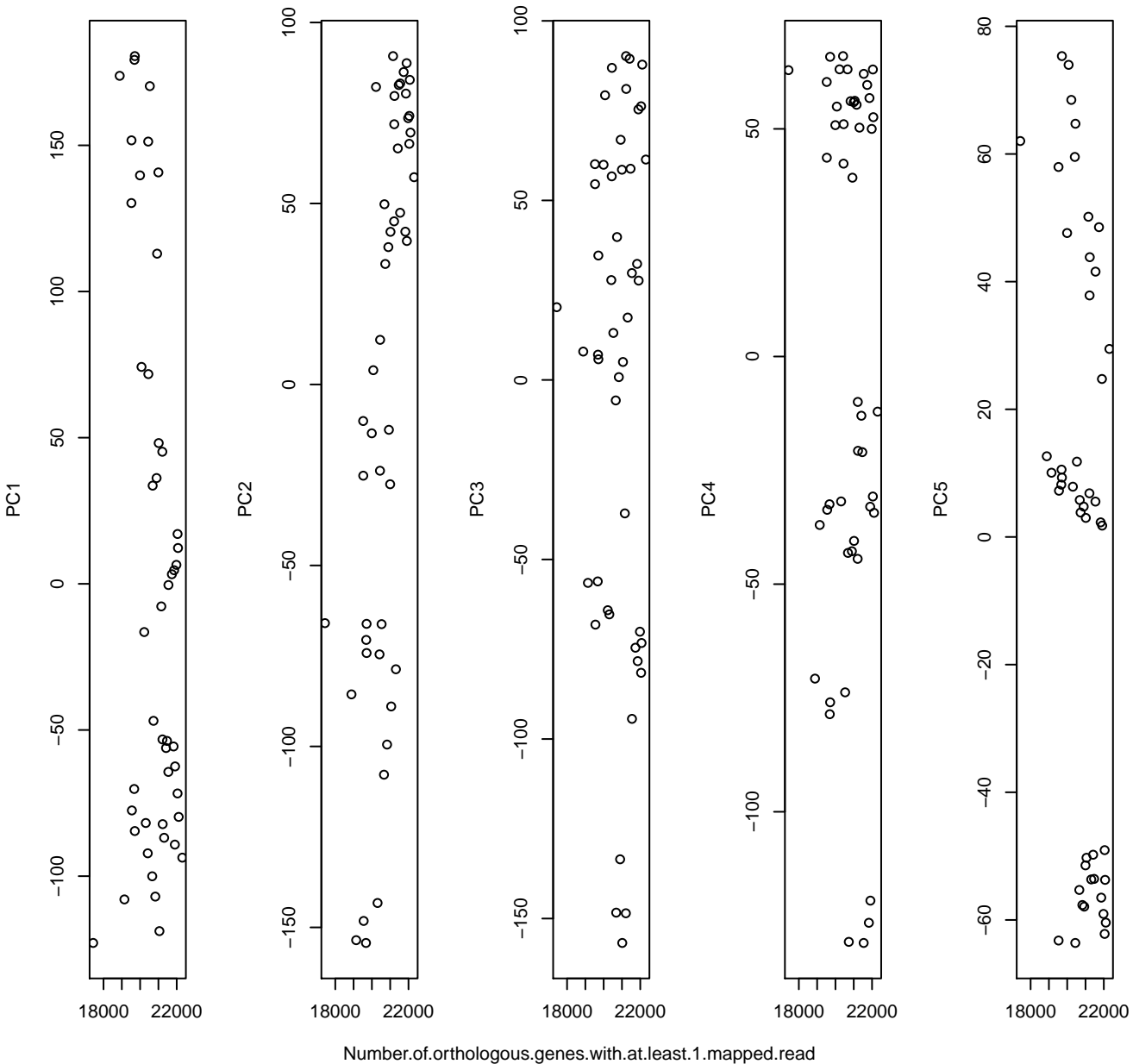




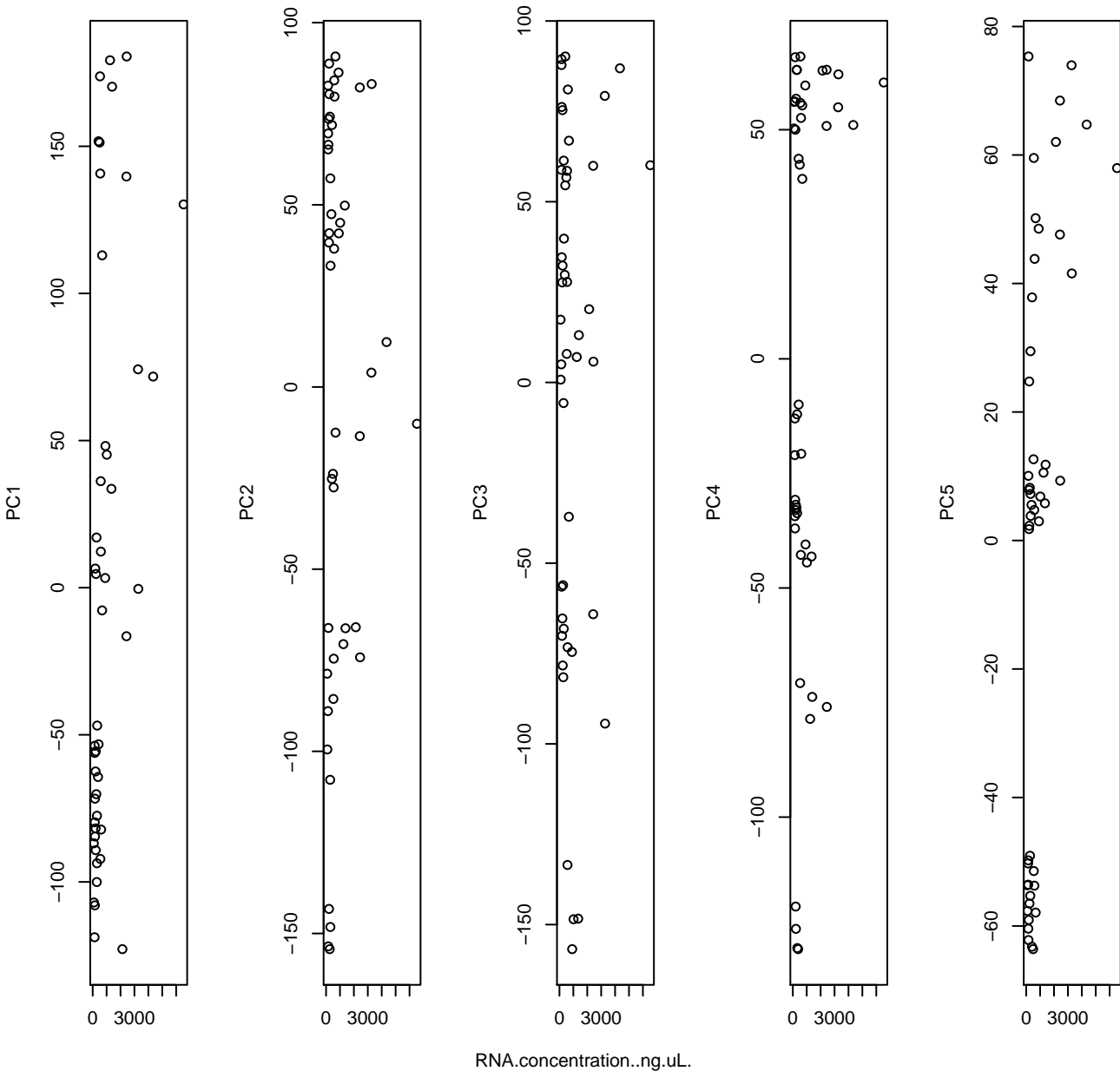
PCs vs. Number.of.orthologous.exons.with.at.least.1.mapped.read



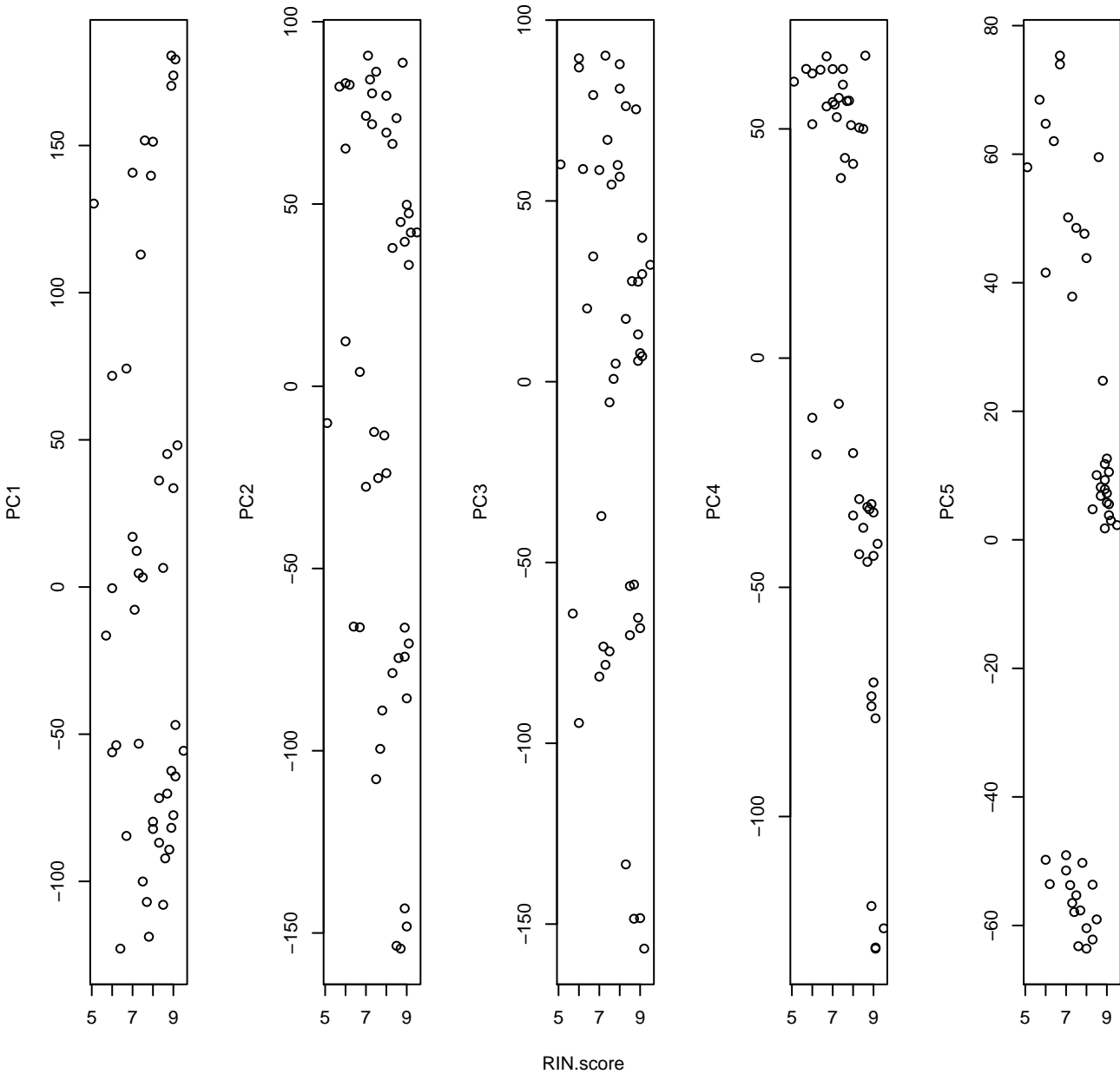
PCs vs. Number.of.orthologous.genes.with.at.least.1.mapped.read



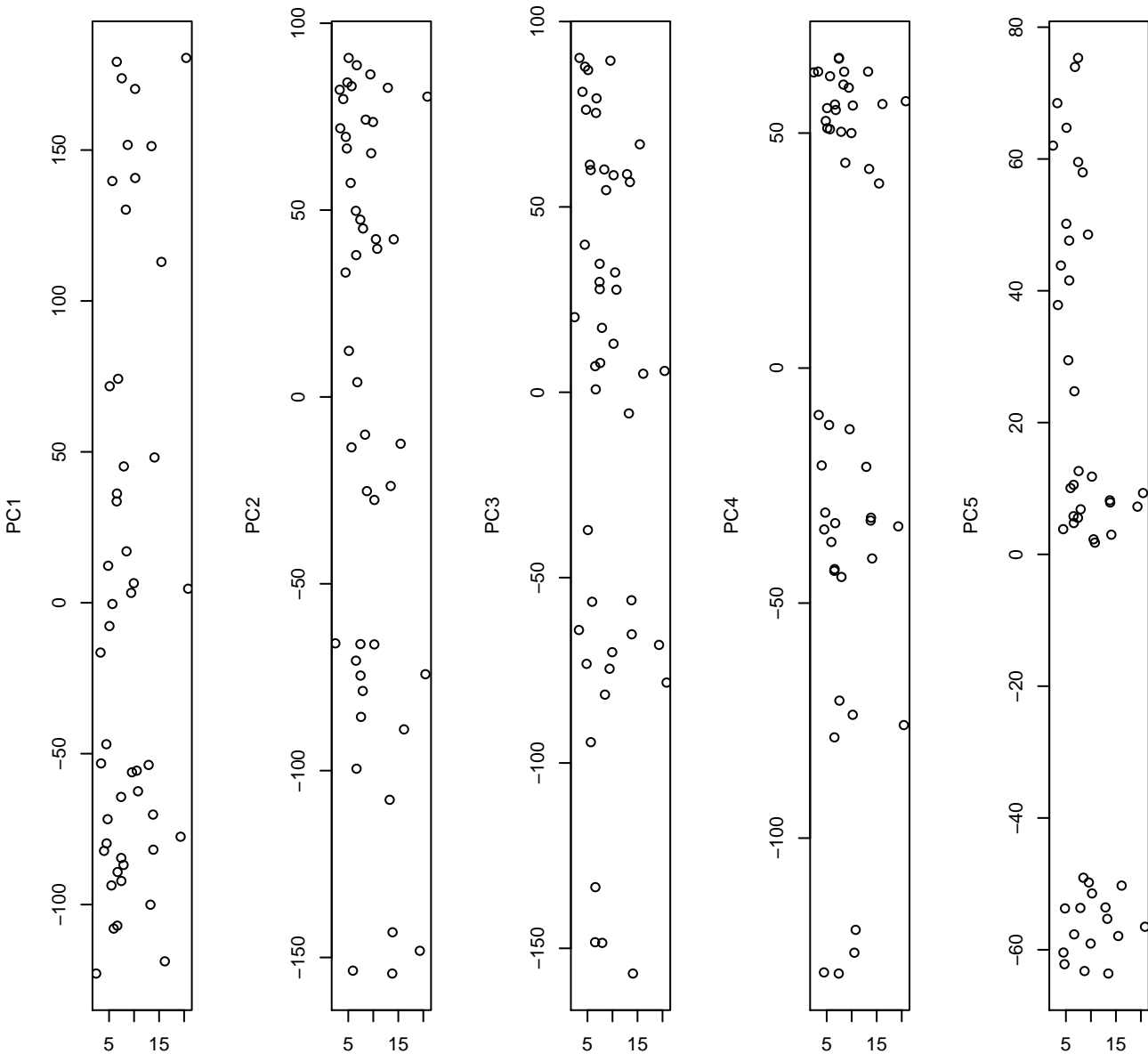
PCs vs. RNA.concentration..ng.uL.



# PCs vs. RIN.score



PCs vs. Library.concentration..ng.uL.



Library.concentration..ng.uL.

PCs vs. Library.fragments.size..bp.

