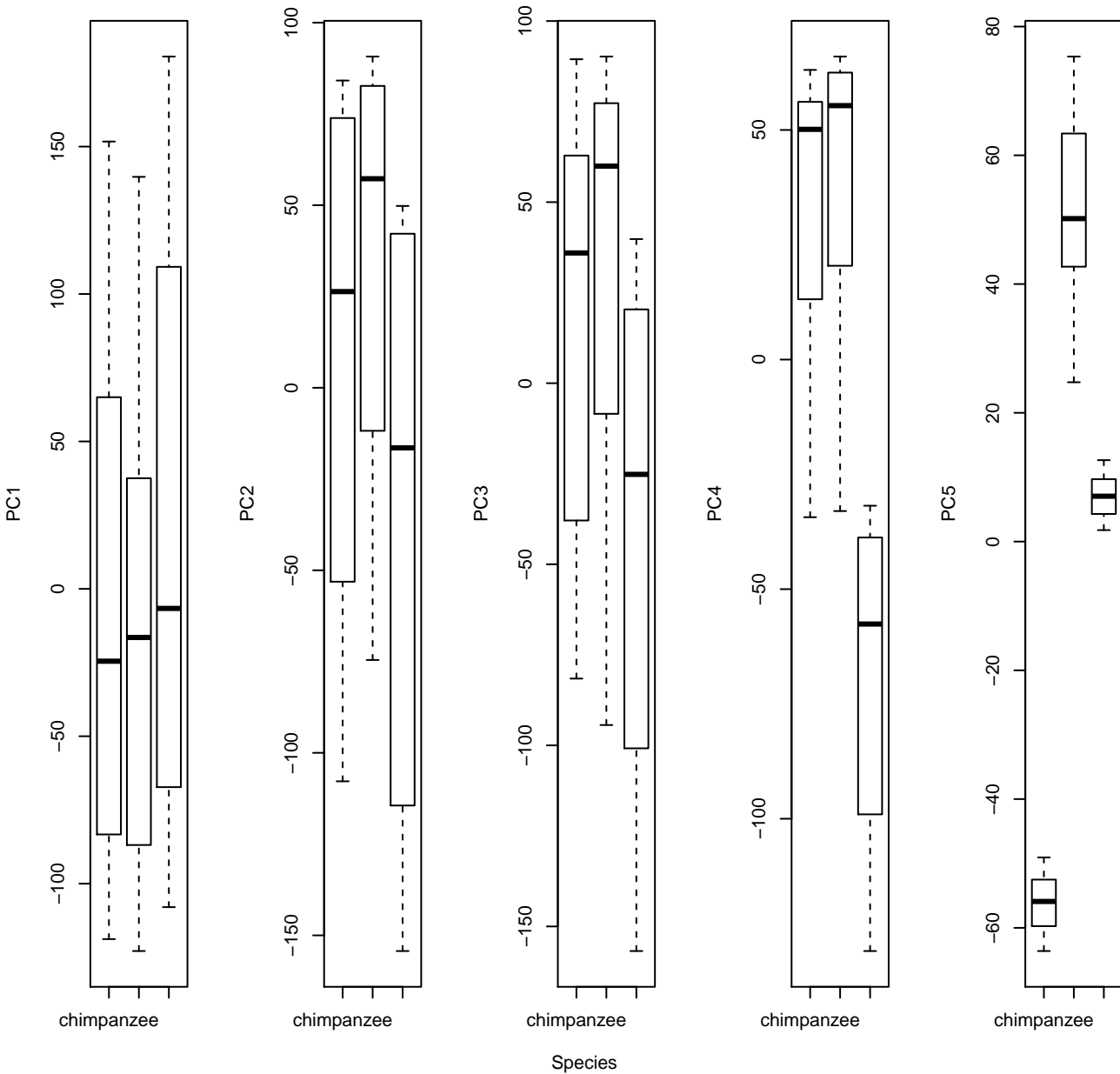
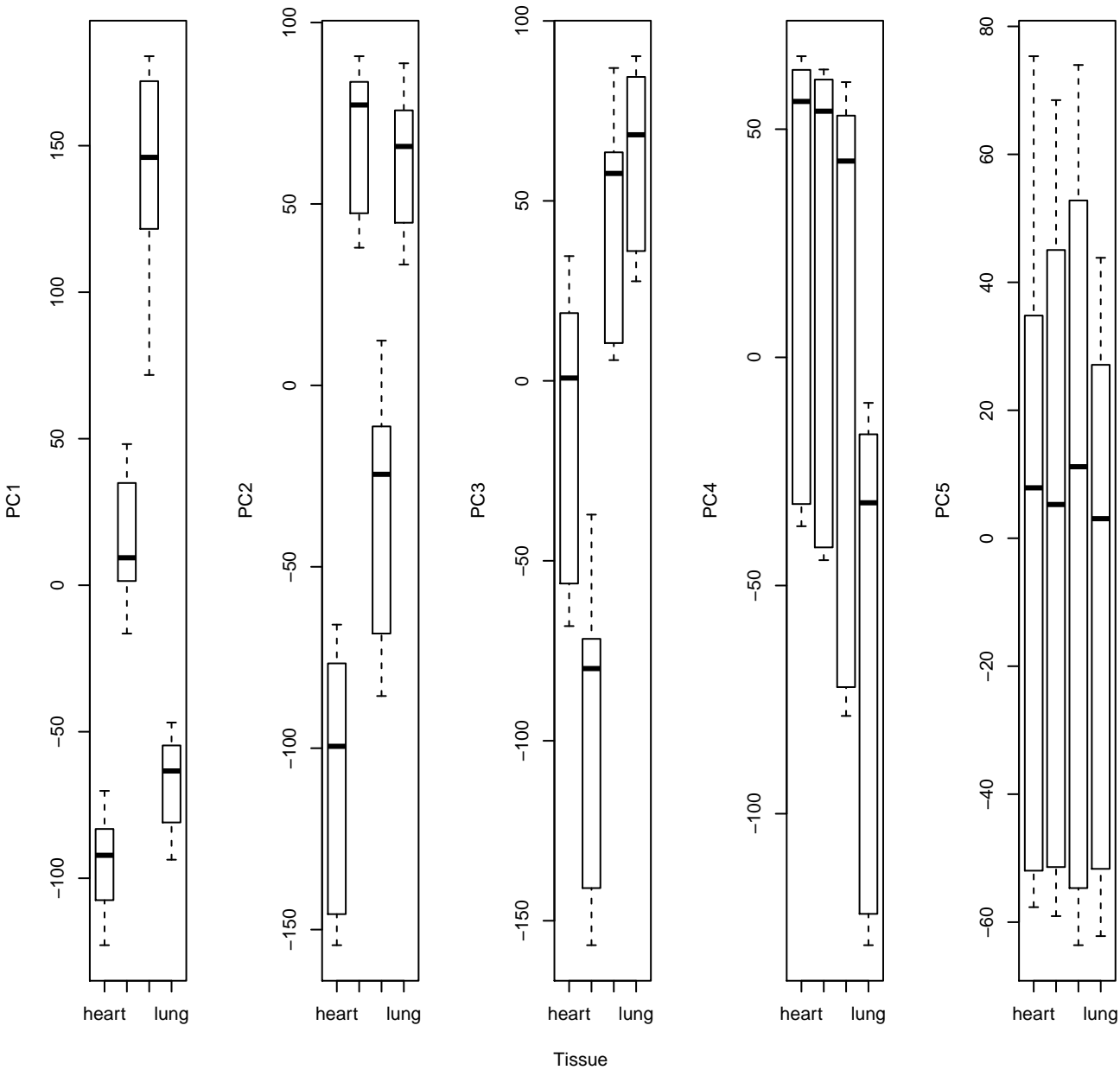


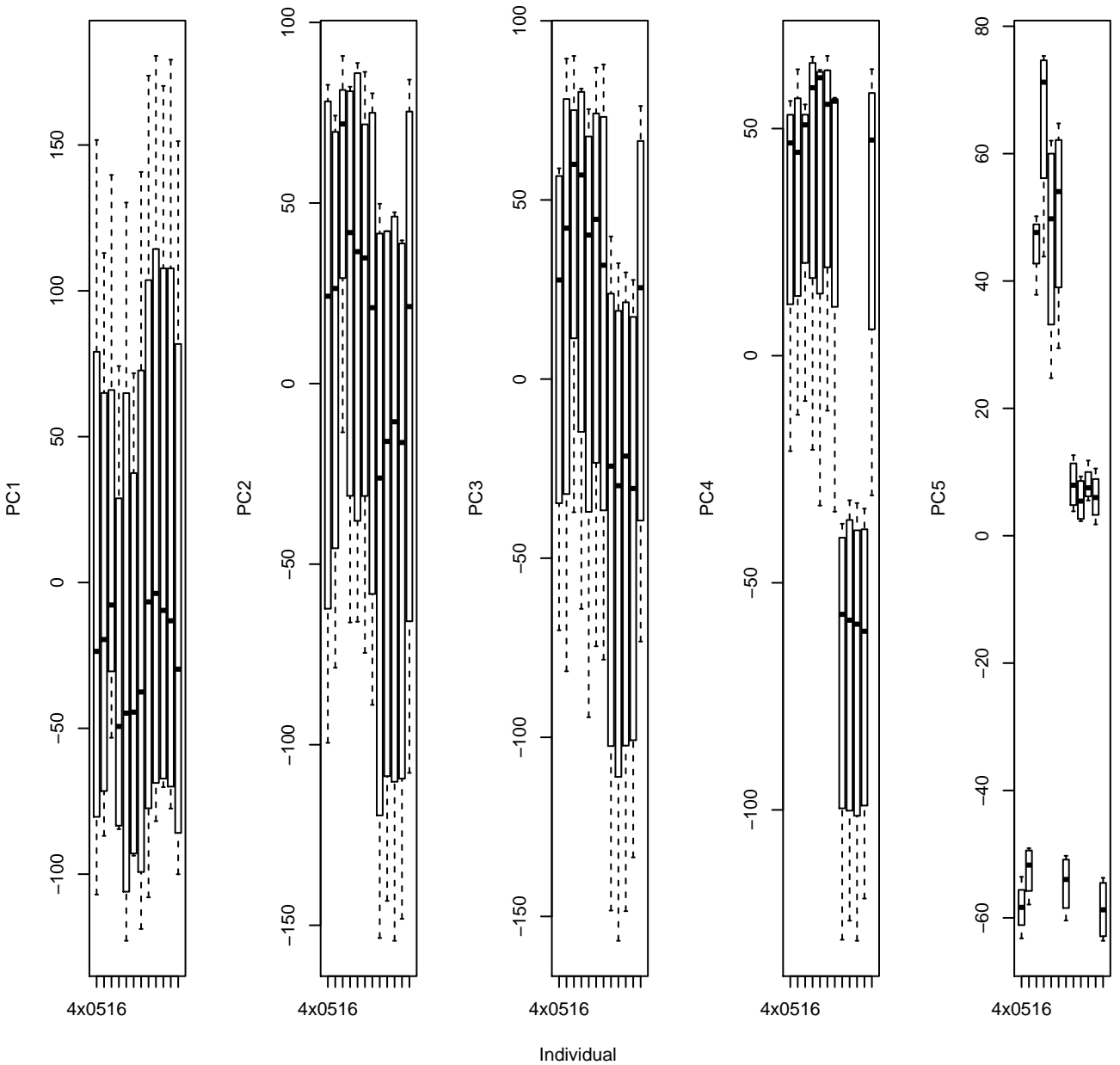
PCs vs. Species



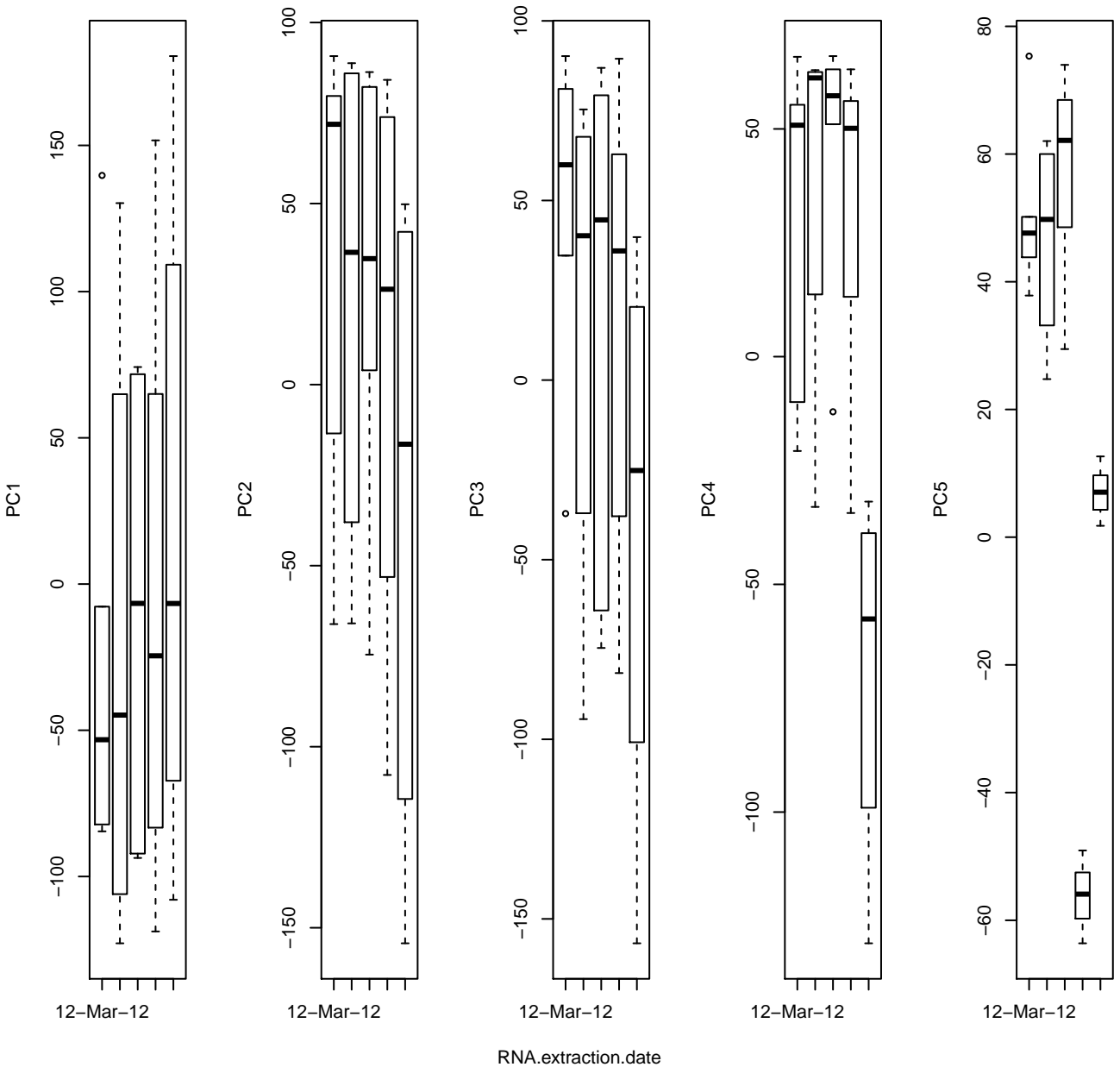
PCs vs. Tissue



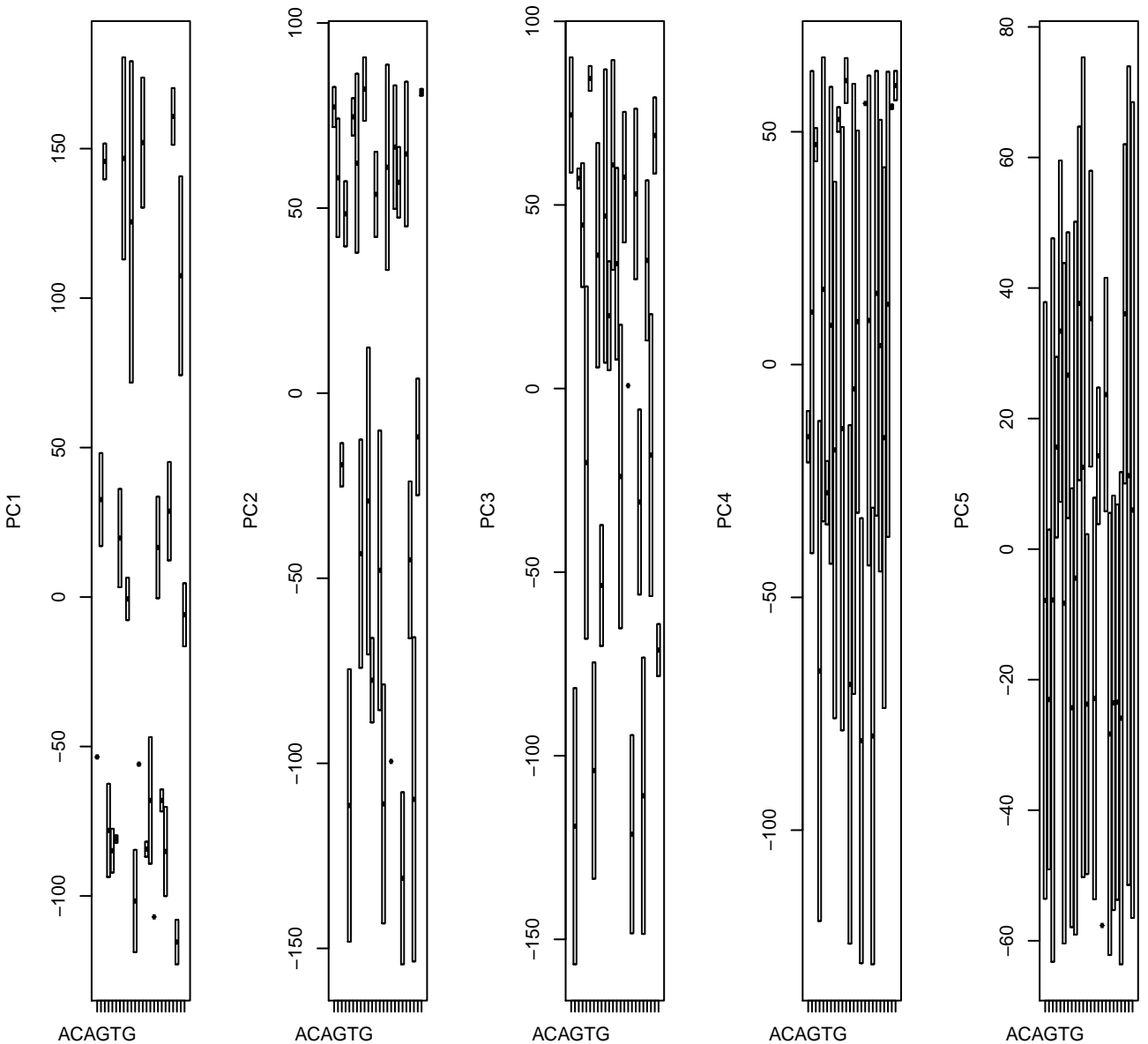
PCs vs. Individual



PCs vs. RNA.extraction.date

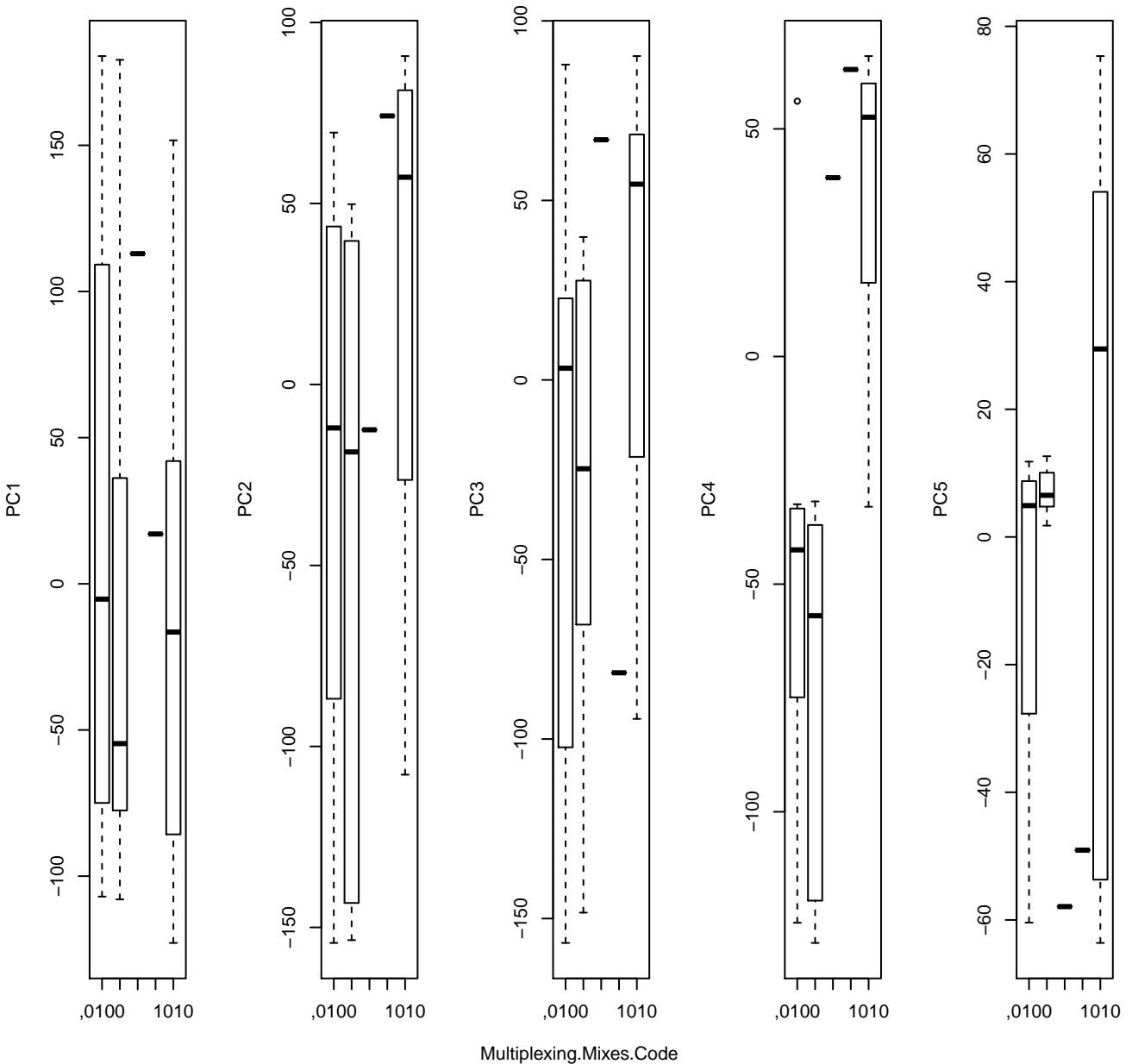


PCs vs. Multiplexing.index.sequence

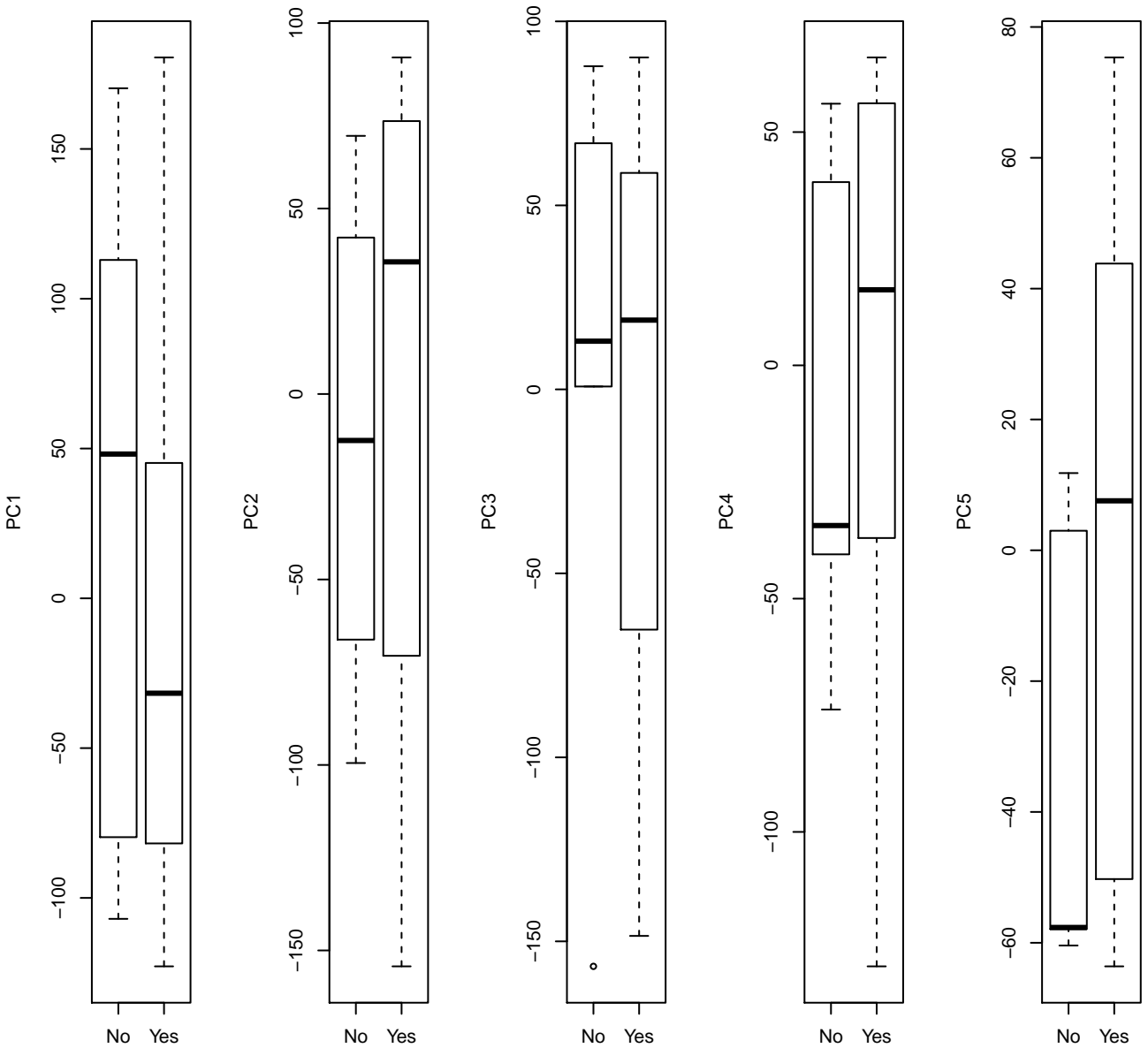


Multiplexing.index.sequence

PCs vs. Multiplexing.Mixes.Code

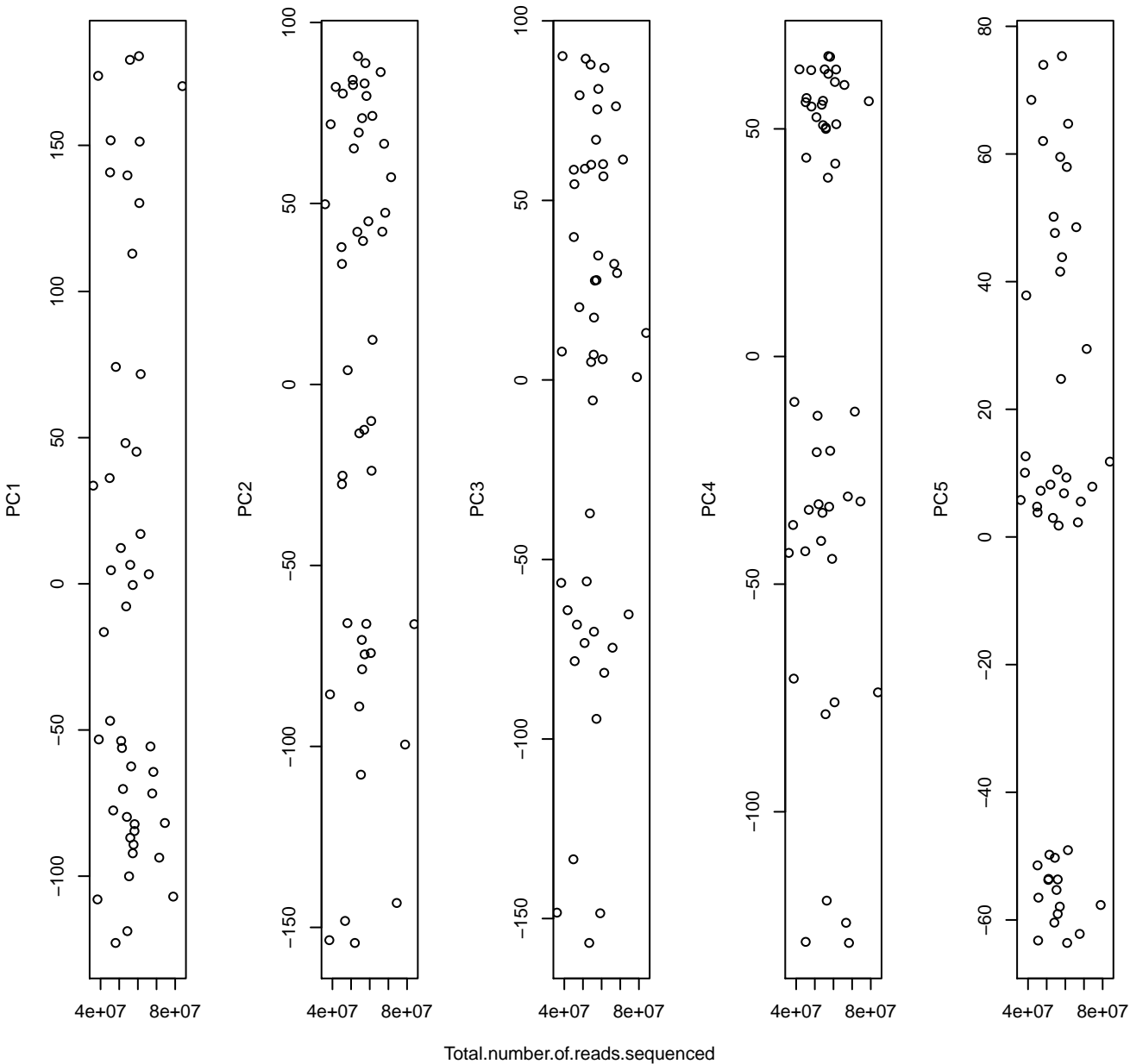


PCs vs. Sequencing.location..at.UCGF

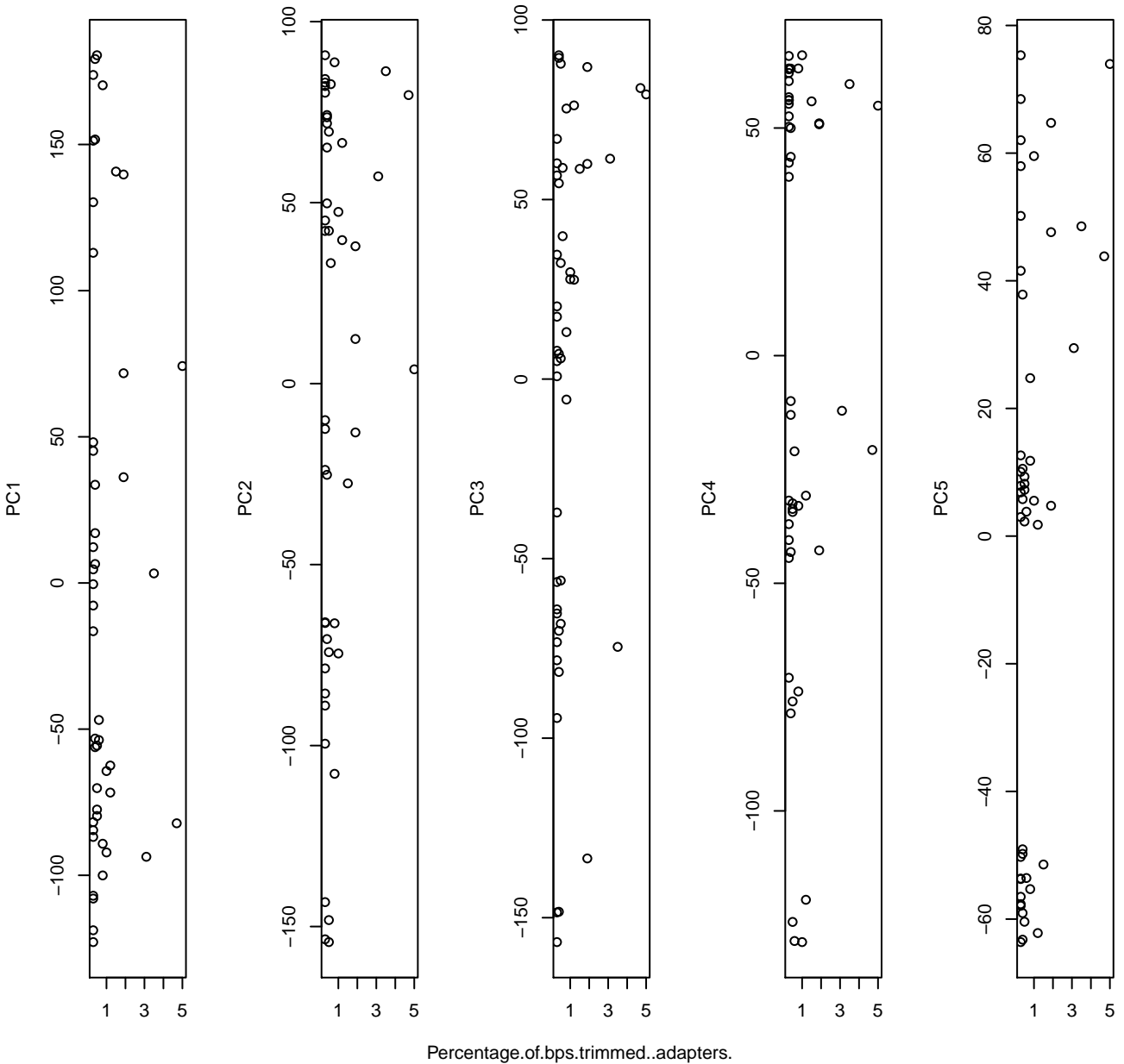


Sequencing.location..at.UCGF

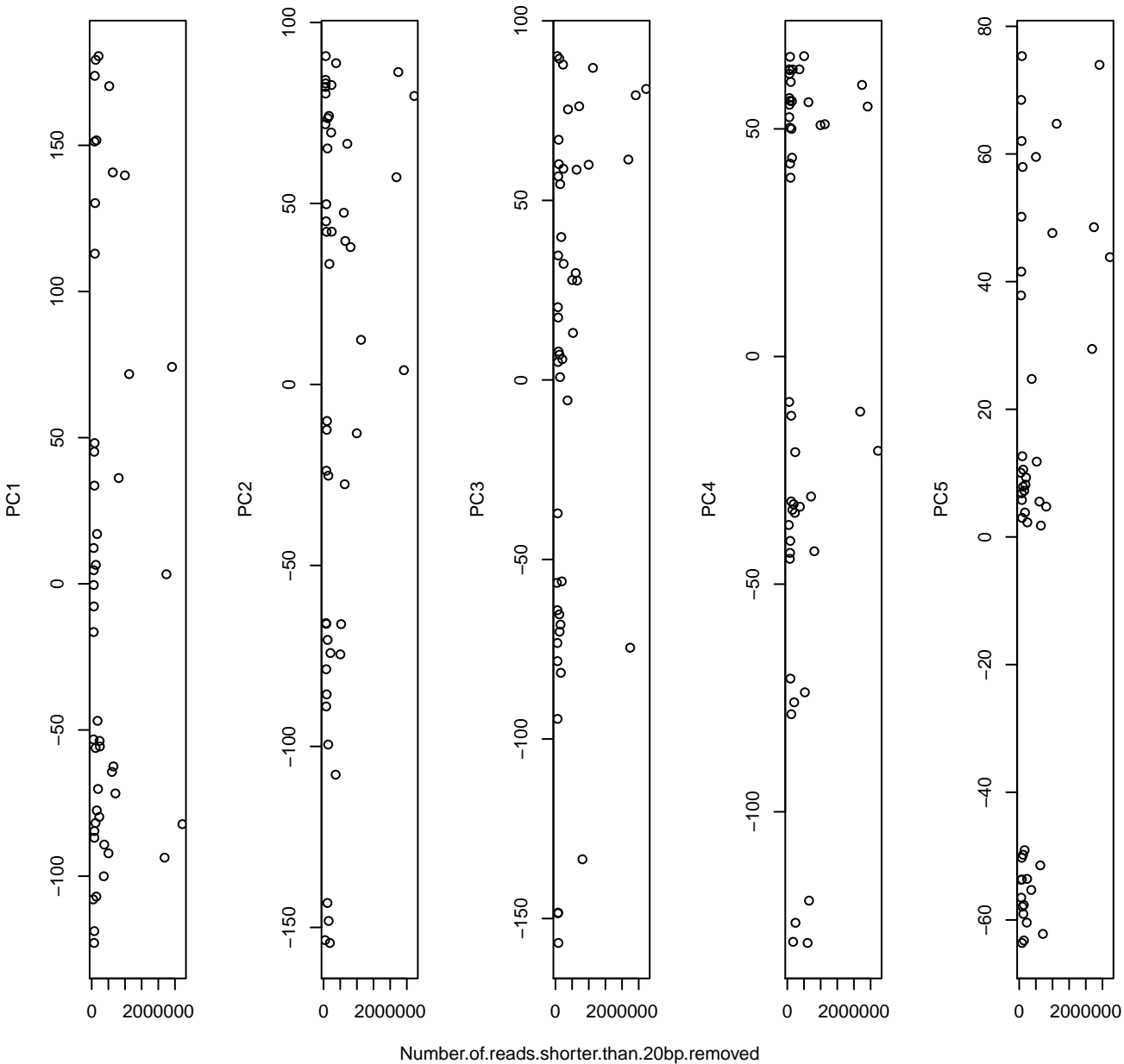
PCs vs. Total.number.of.reads.sequenced



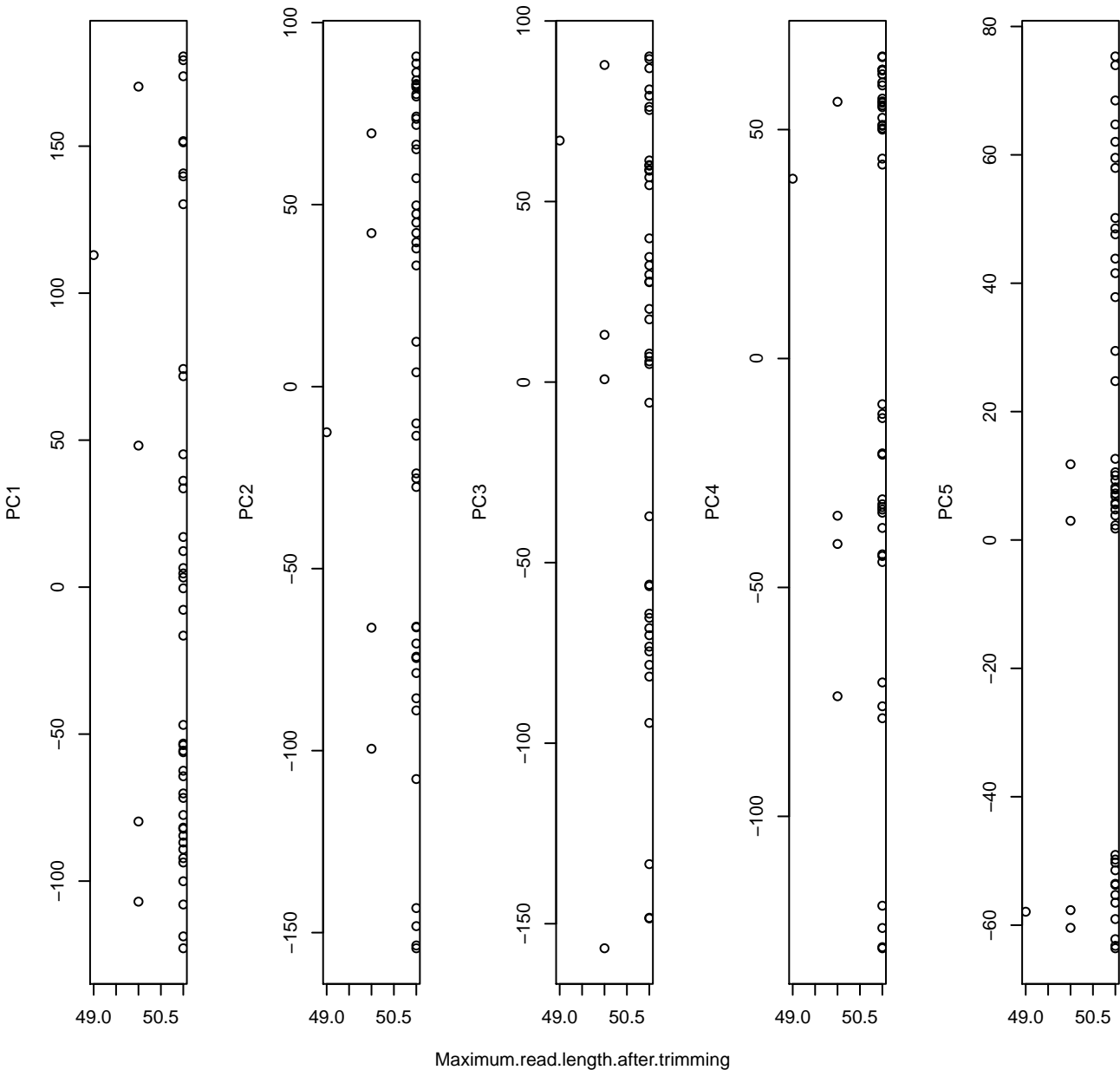
PCs vs. Percentage.of.bps.trimmed..adapters.



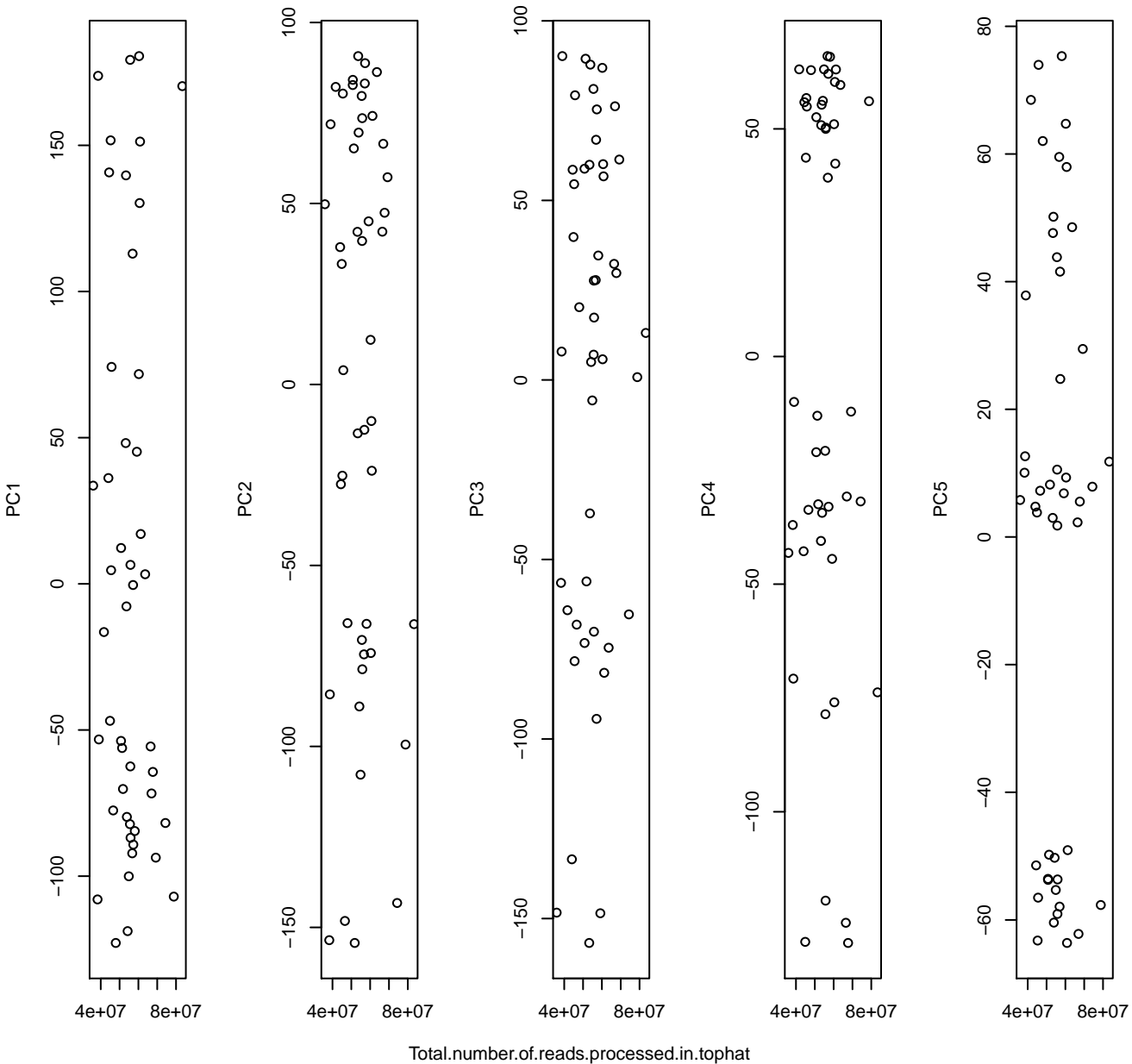
PCs vs. Number.of.reads.shorter.than.20bp.removed



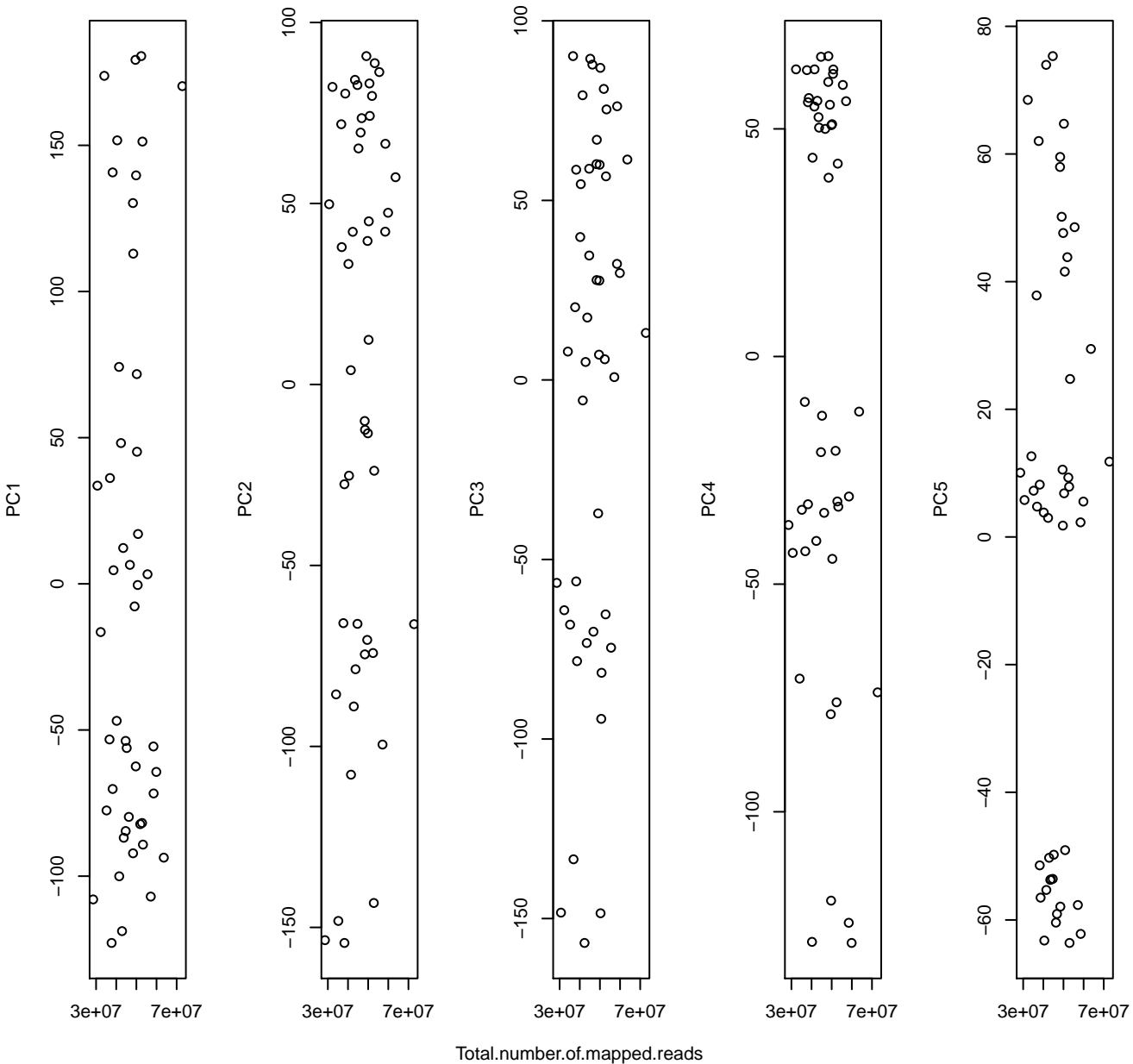
PCs vs. Maximum.read.length.after.trimming



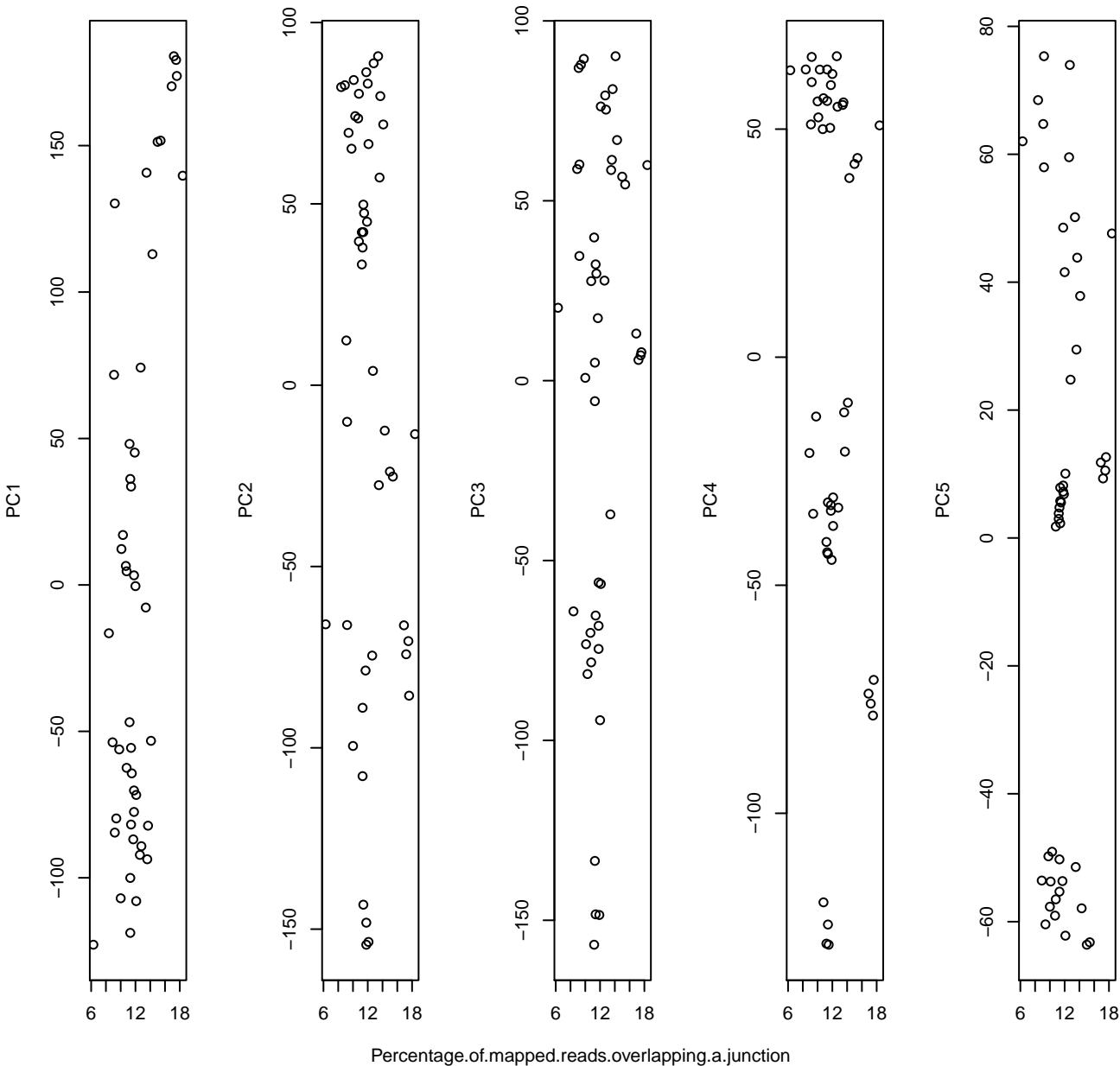
PCs vs. Total.number.of.reads.processed.in.tophat



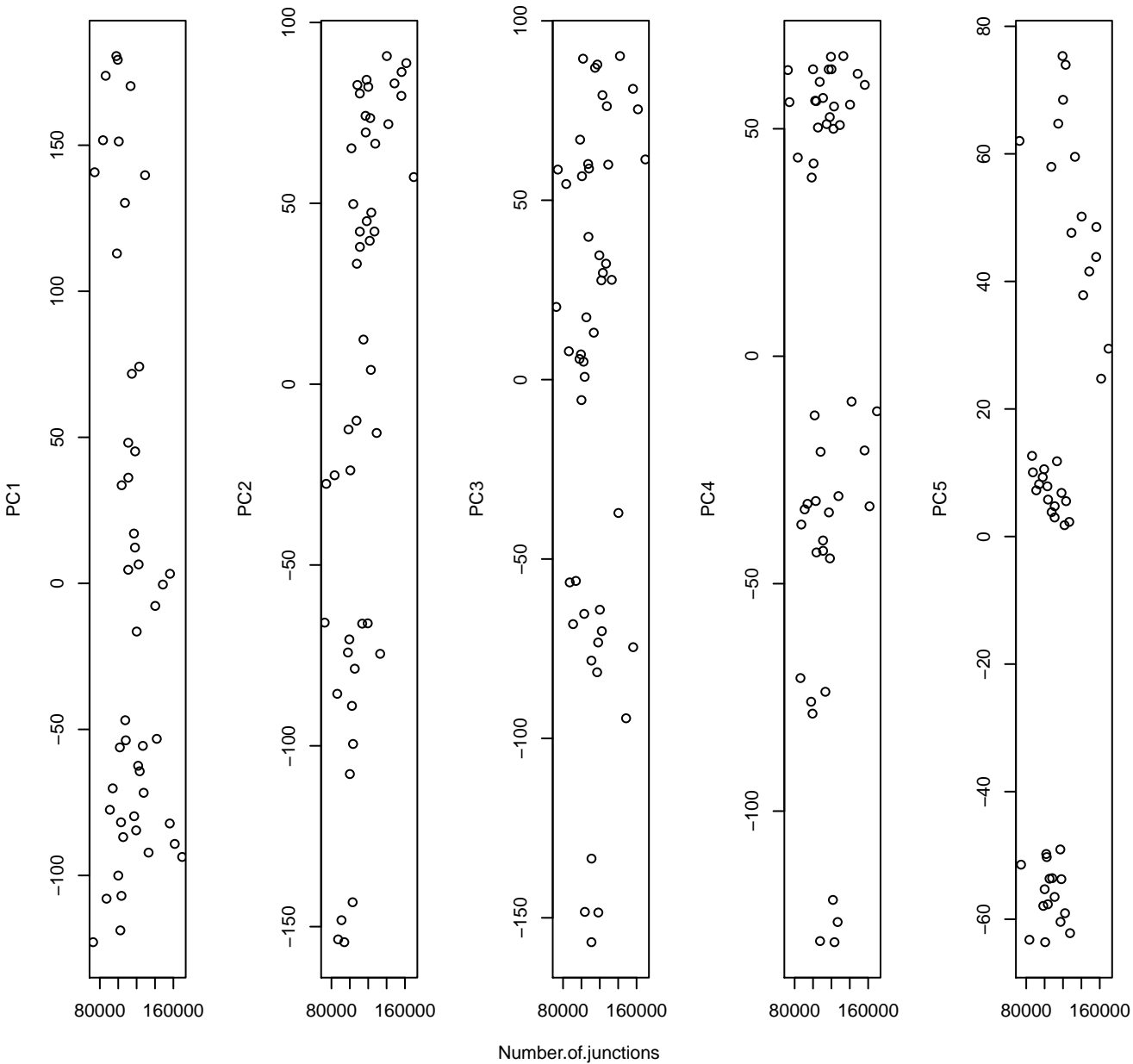
PCs vs. Total.number.of.mapped.reads



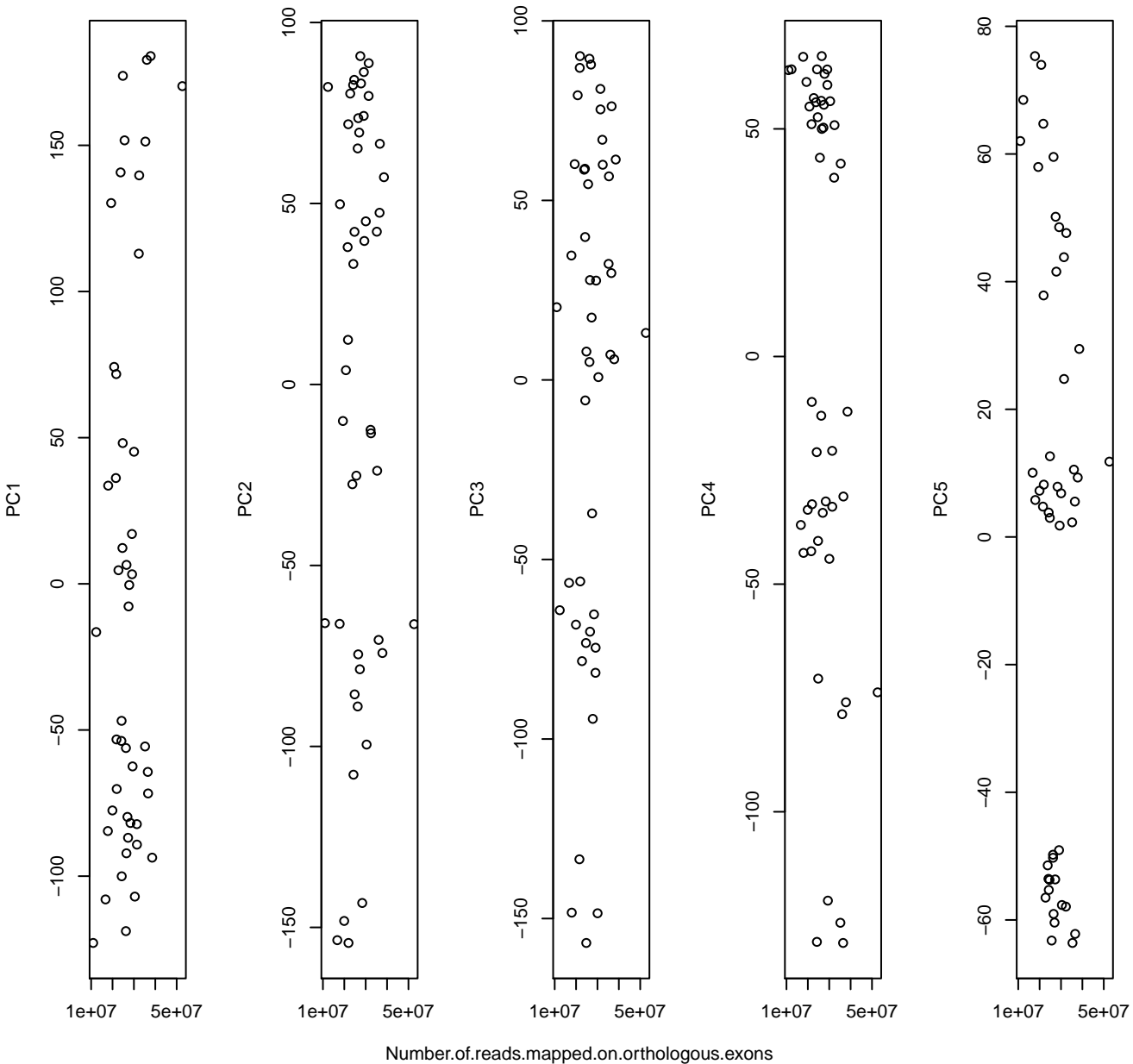
PCs vs. Percentage.of.mapped.reads.overlapping.a.junction



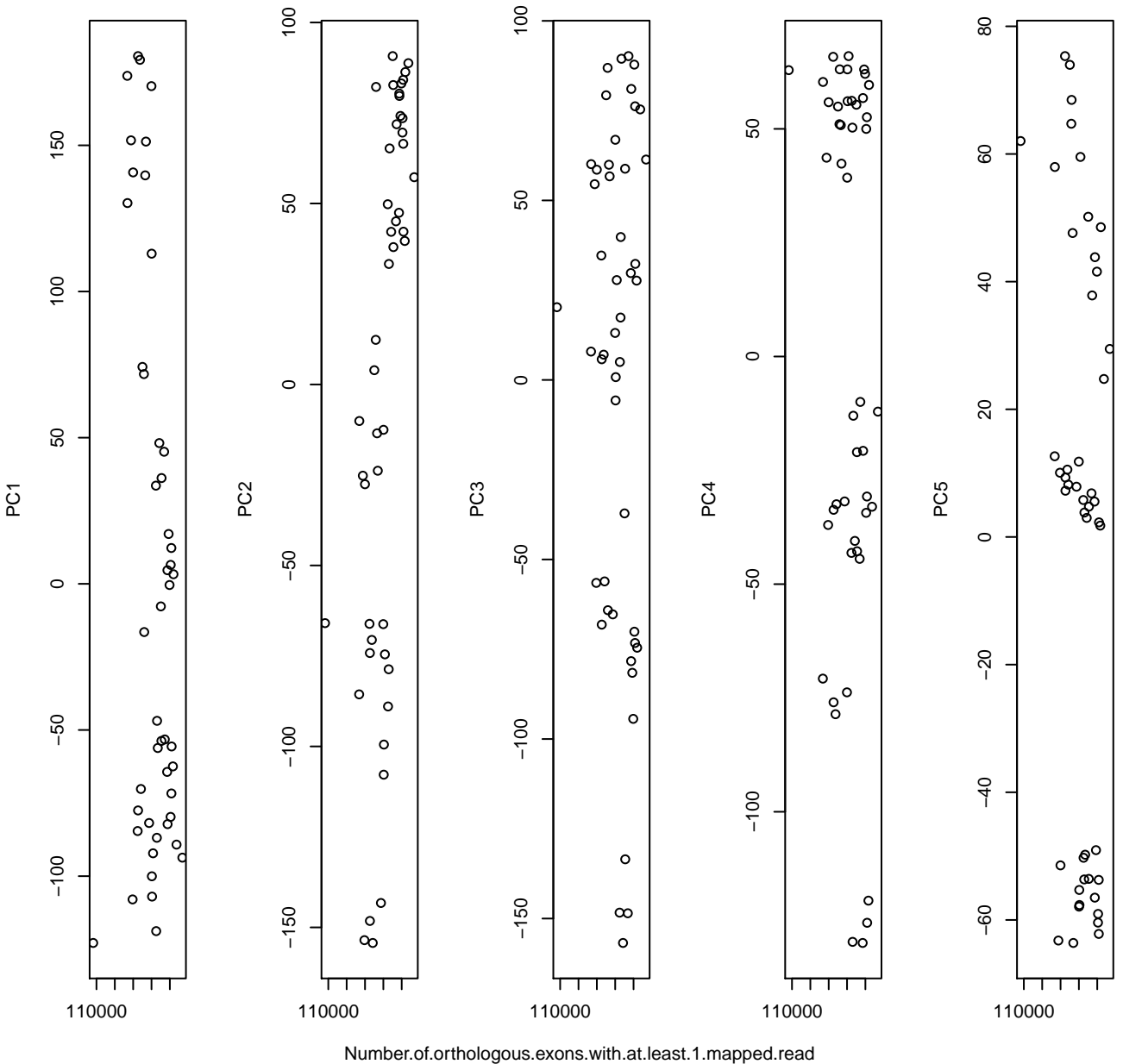
PCs vs. Number.of.junctions



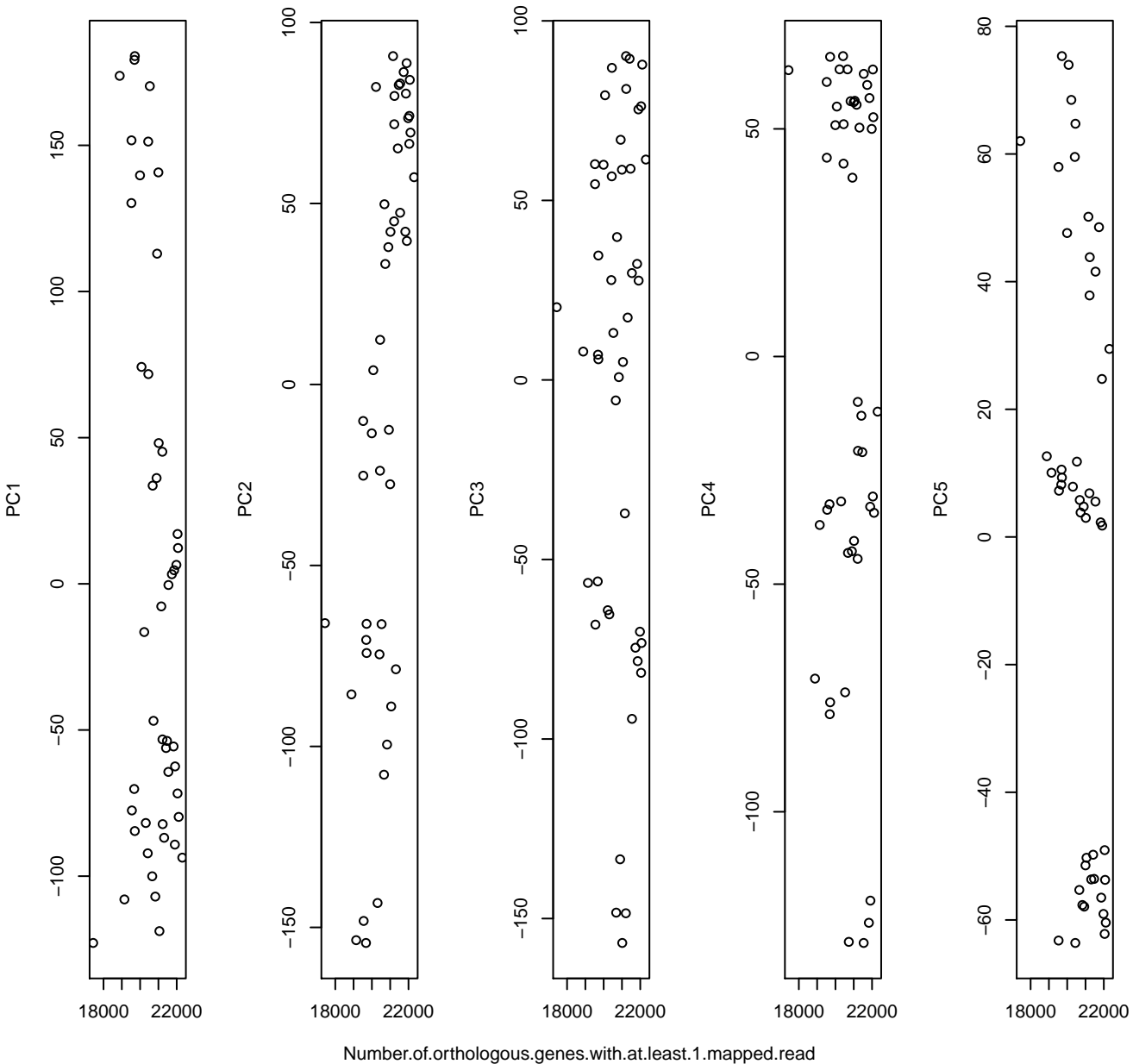
PCs vs. Number.of.reads.mapped.on.orthologous.exons



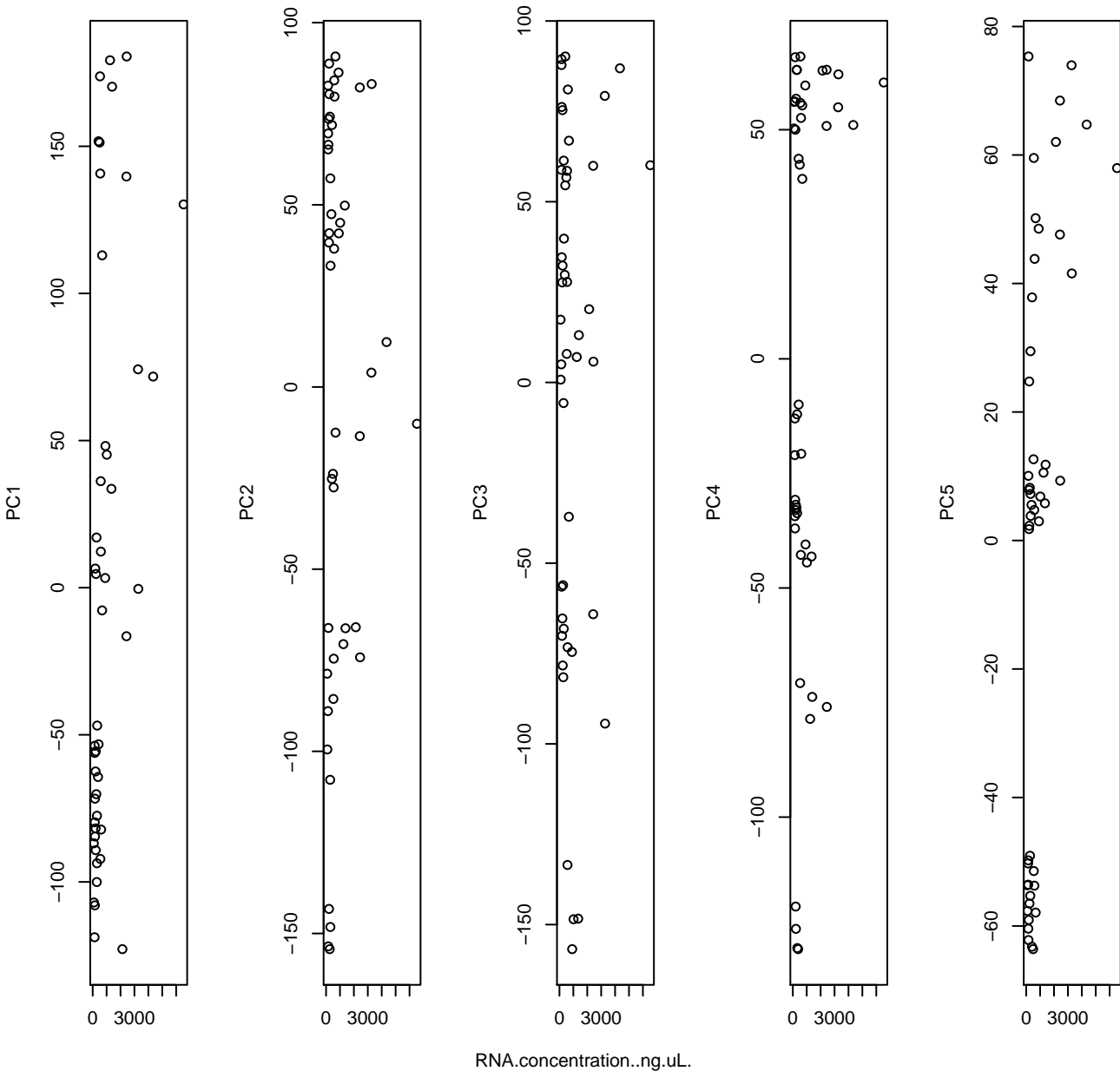
PCs vs. Number.of.orthologous.exons.with.at.least.1.mapped.read



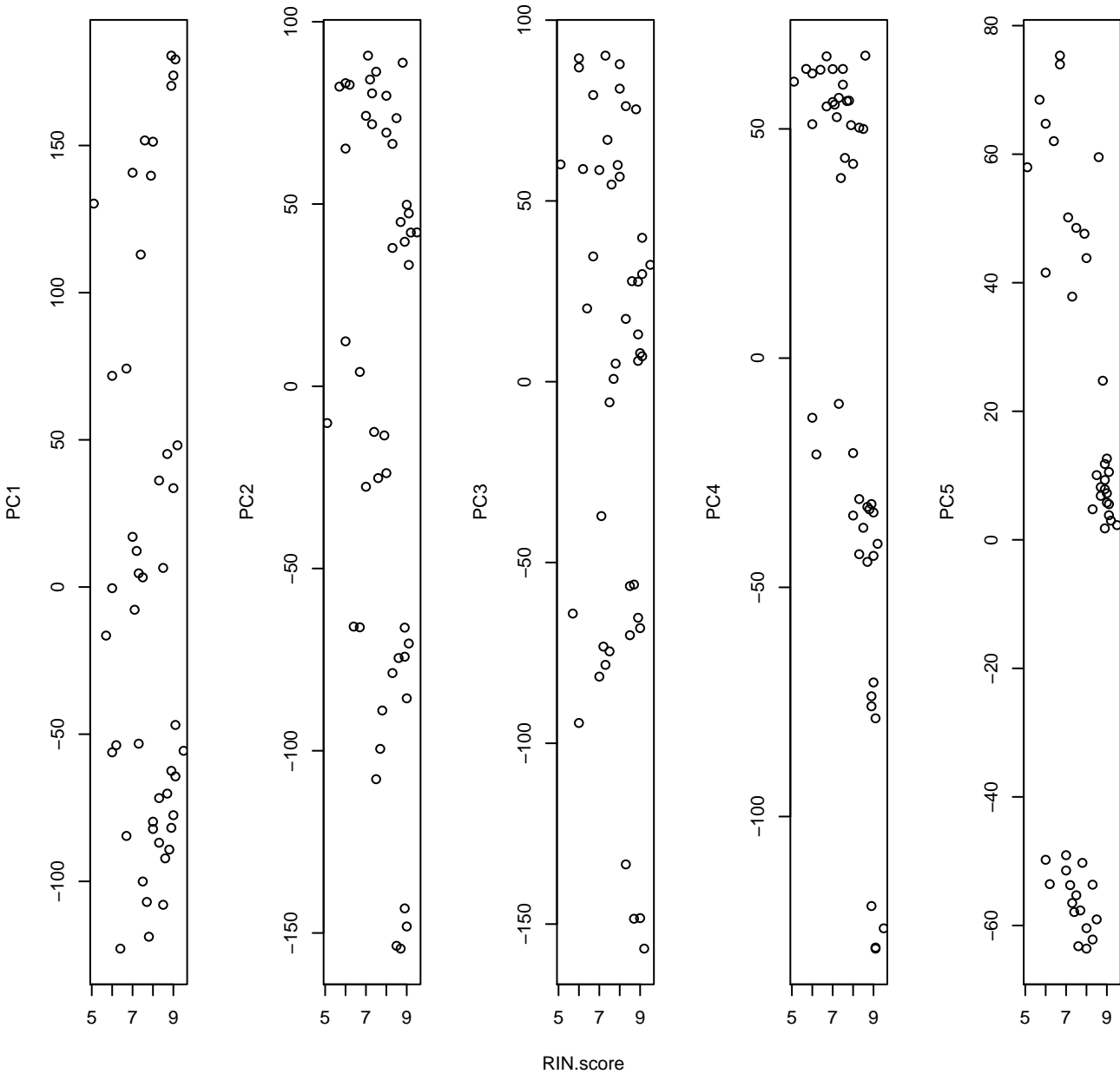
PCs vs. Number.of.orthologous.genes.with.at.least.1.mapped.read



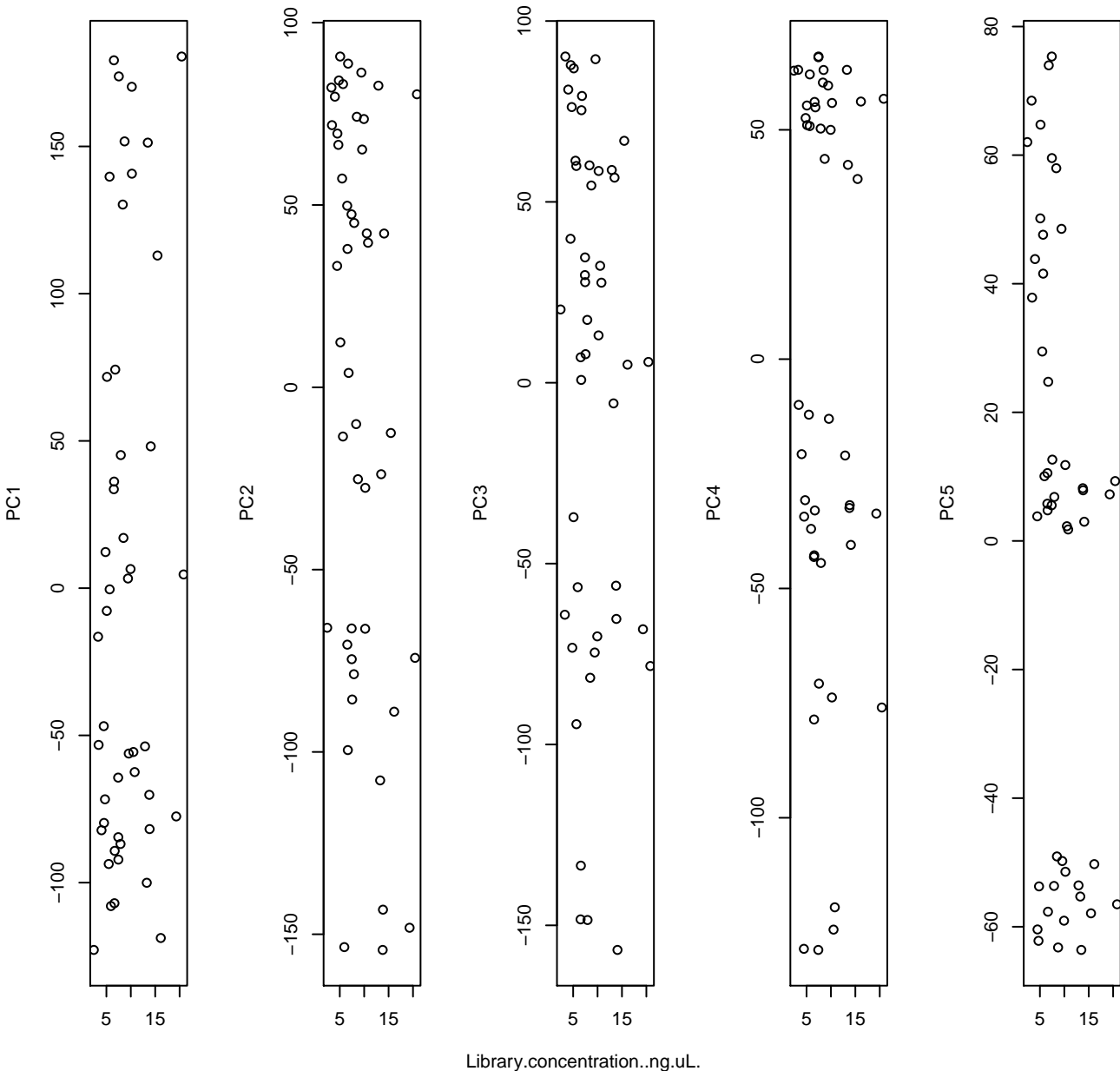
PCs vs. RNA.concentration..ng.uL.



PCs vs. RIN.score



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



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

