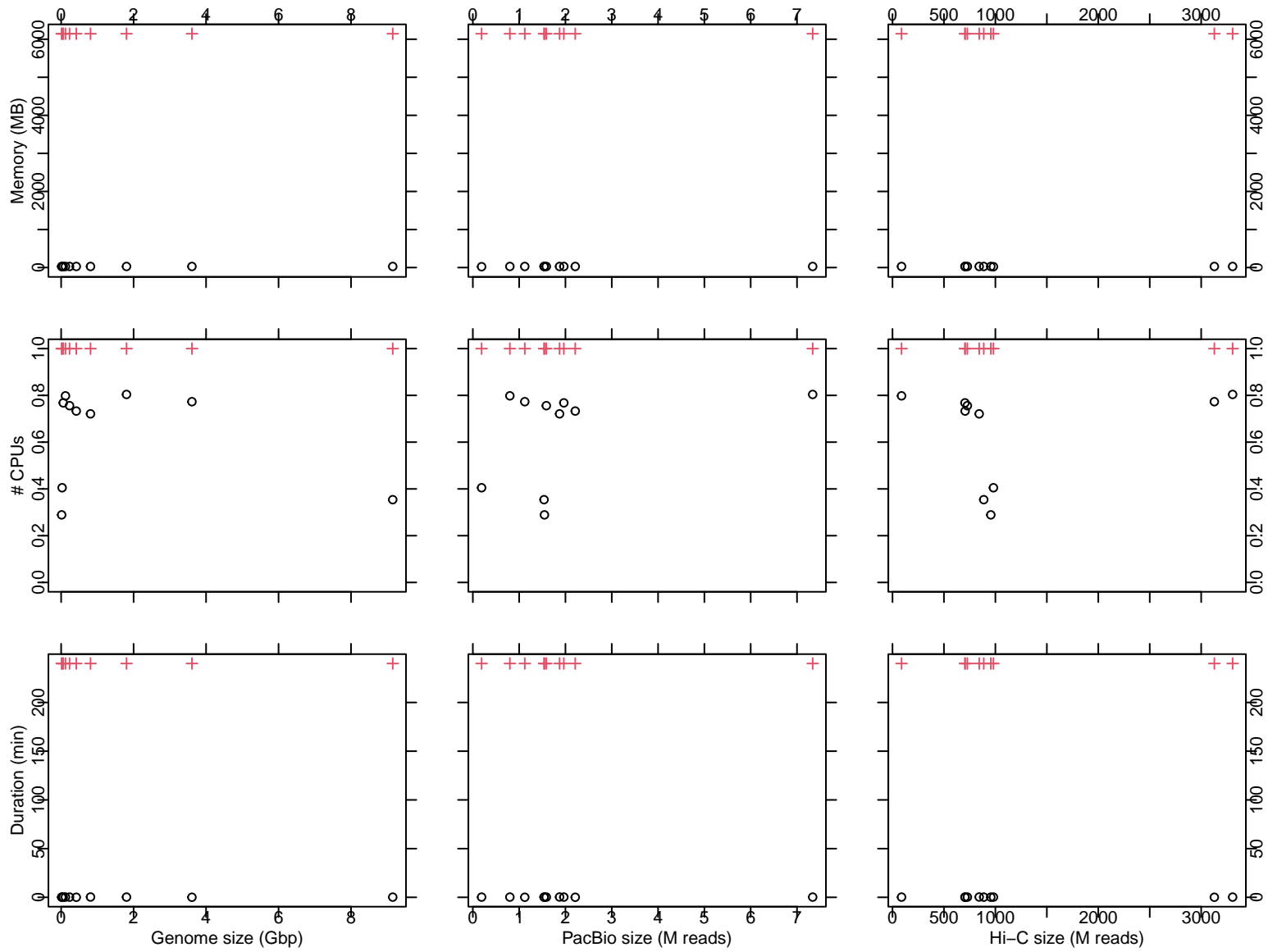
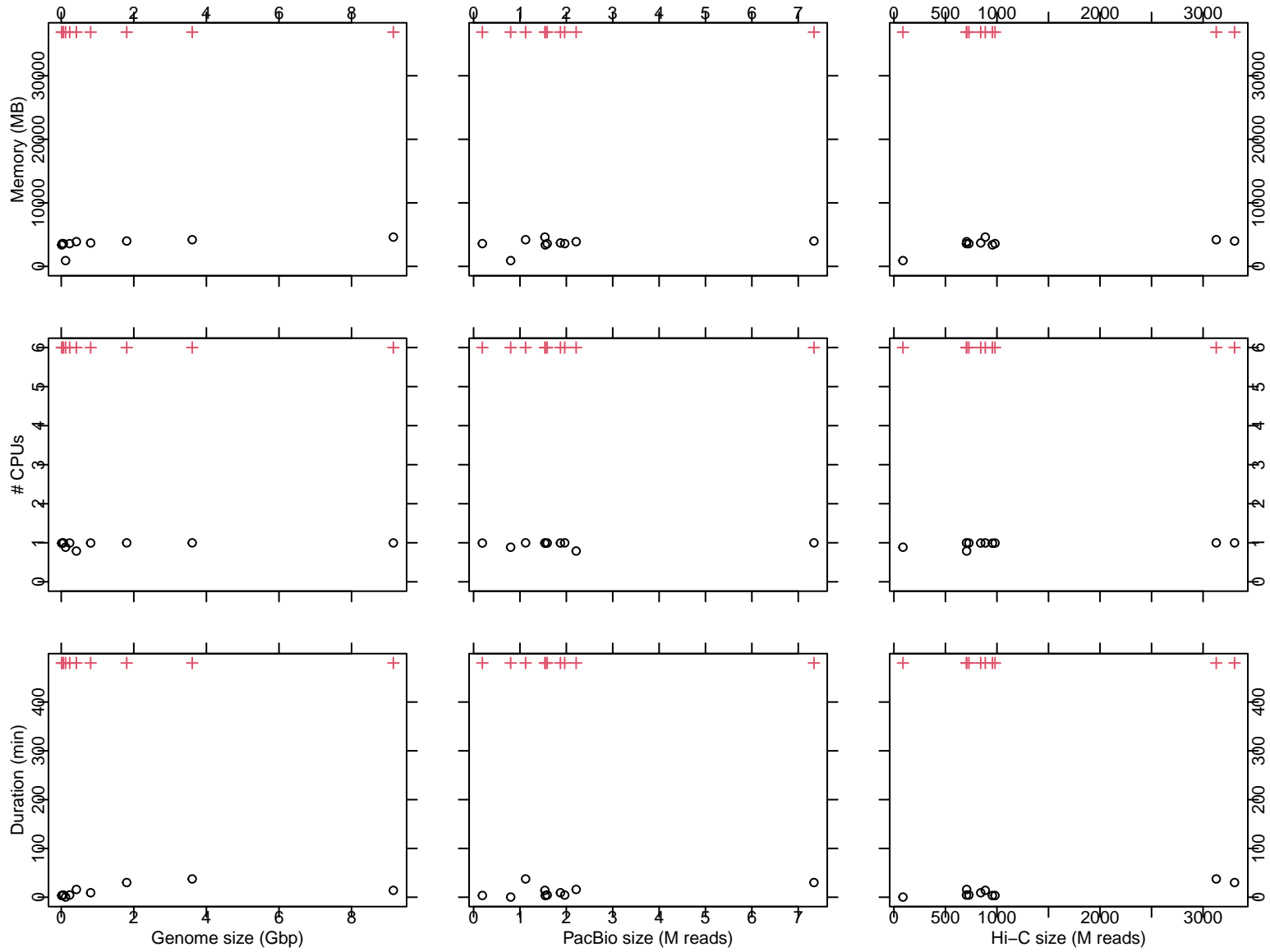


sanger-tol/genomenote - start

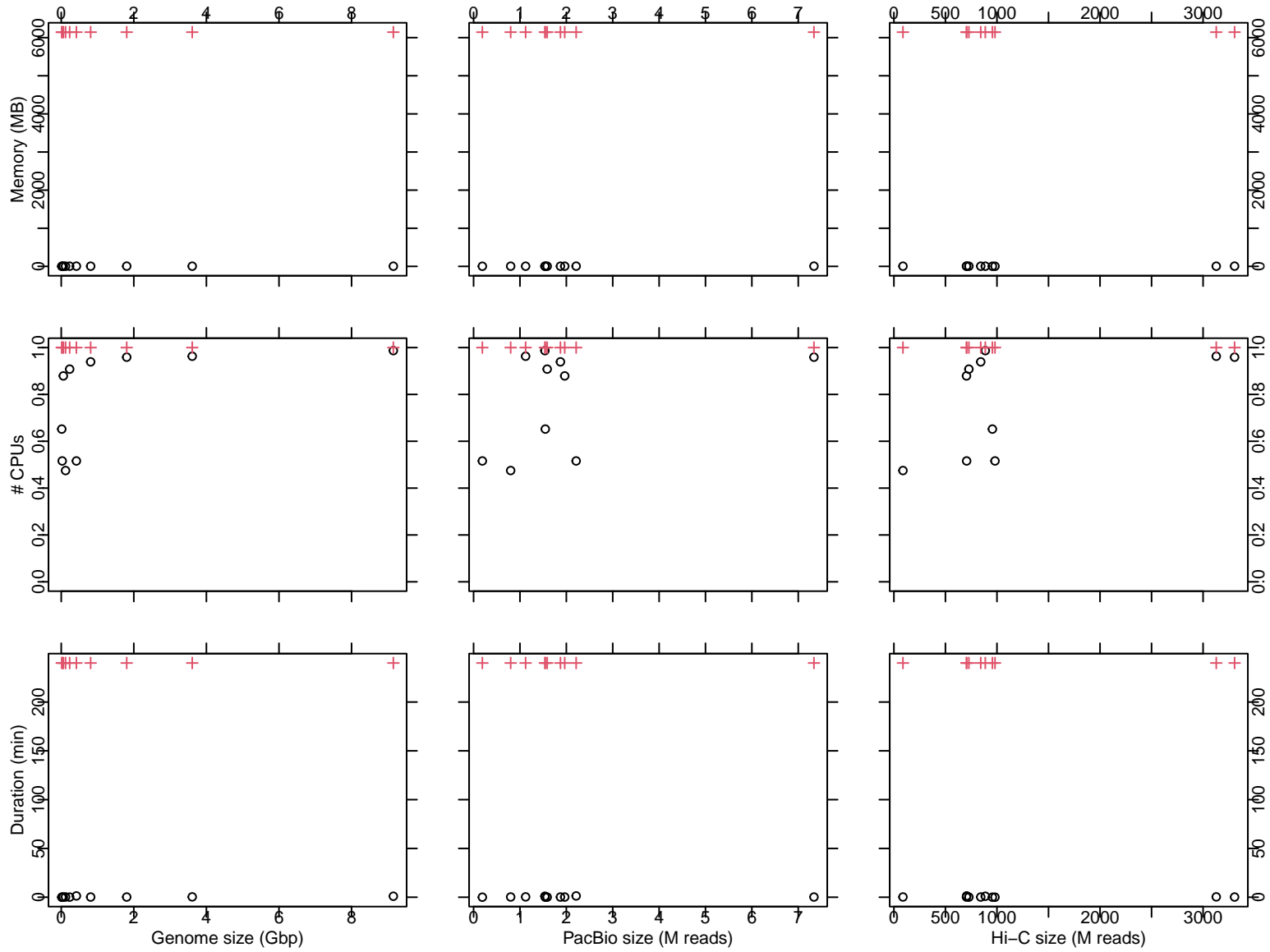
GENOME_STATISTICS:NCBI_GET_ODB



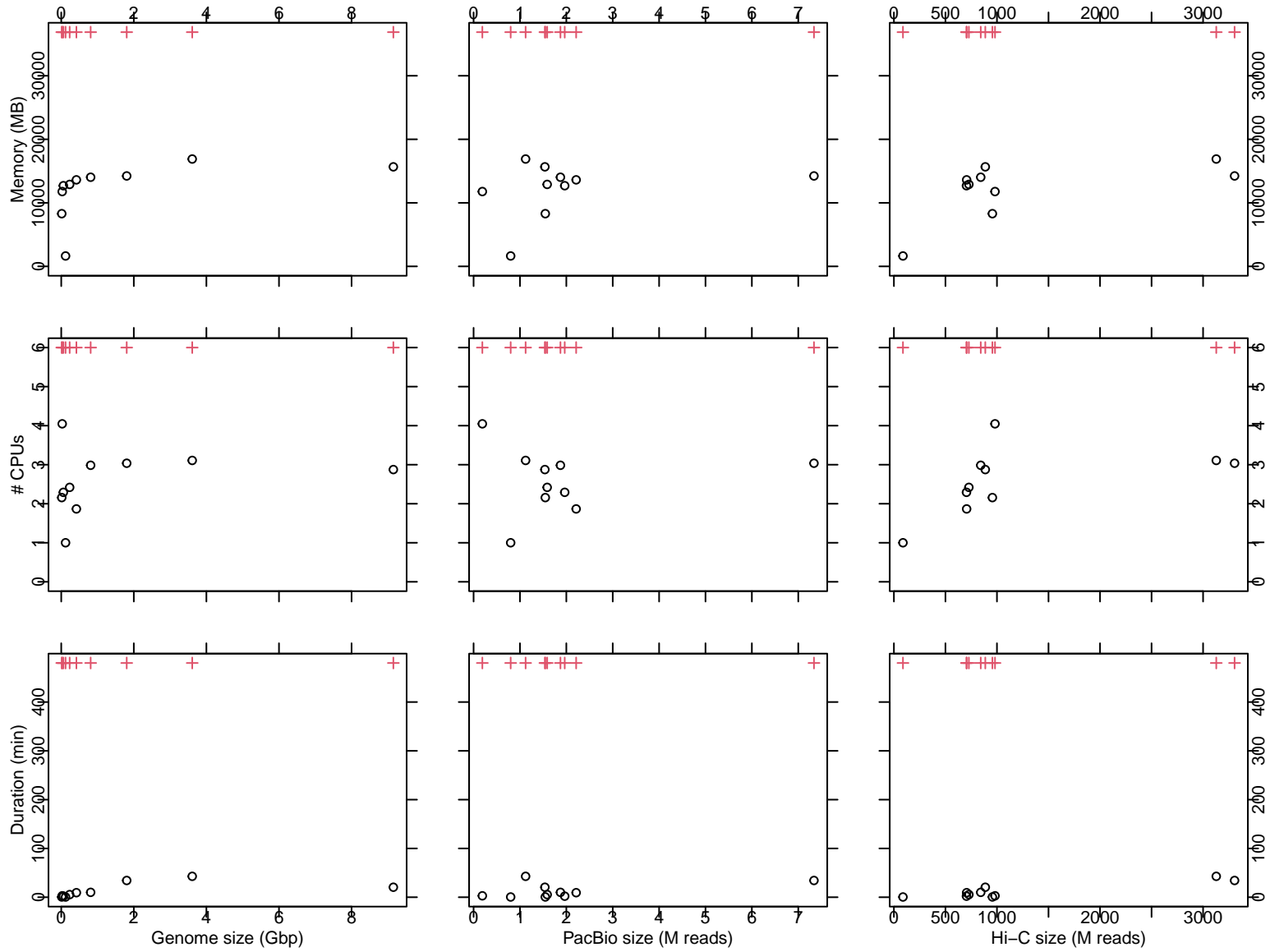
CONTACT_MAPS:COOLER_LOAD



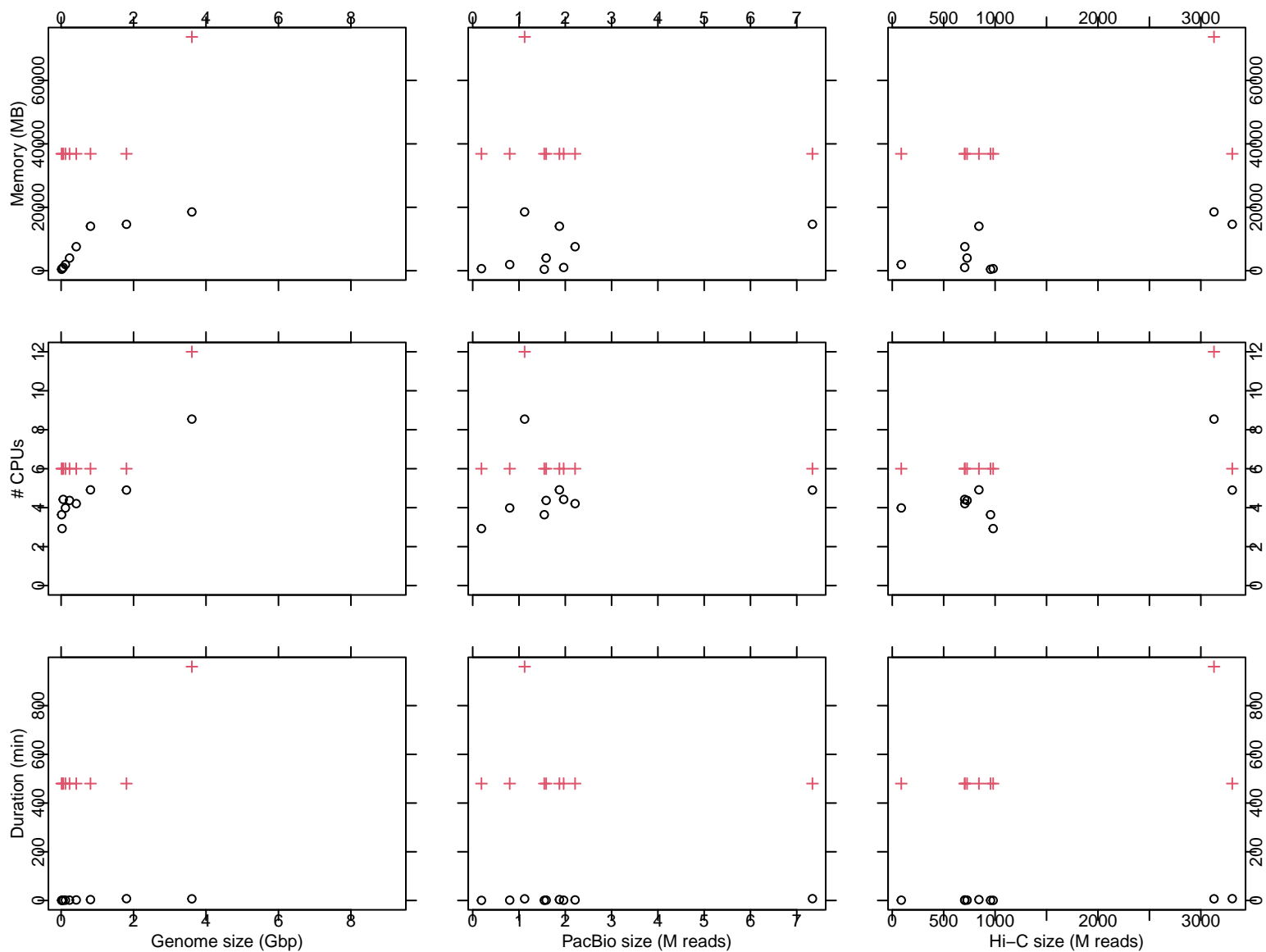
CONTACT_MAPS:SAMTOOLS_FAIDX



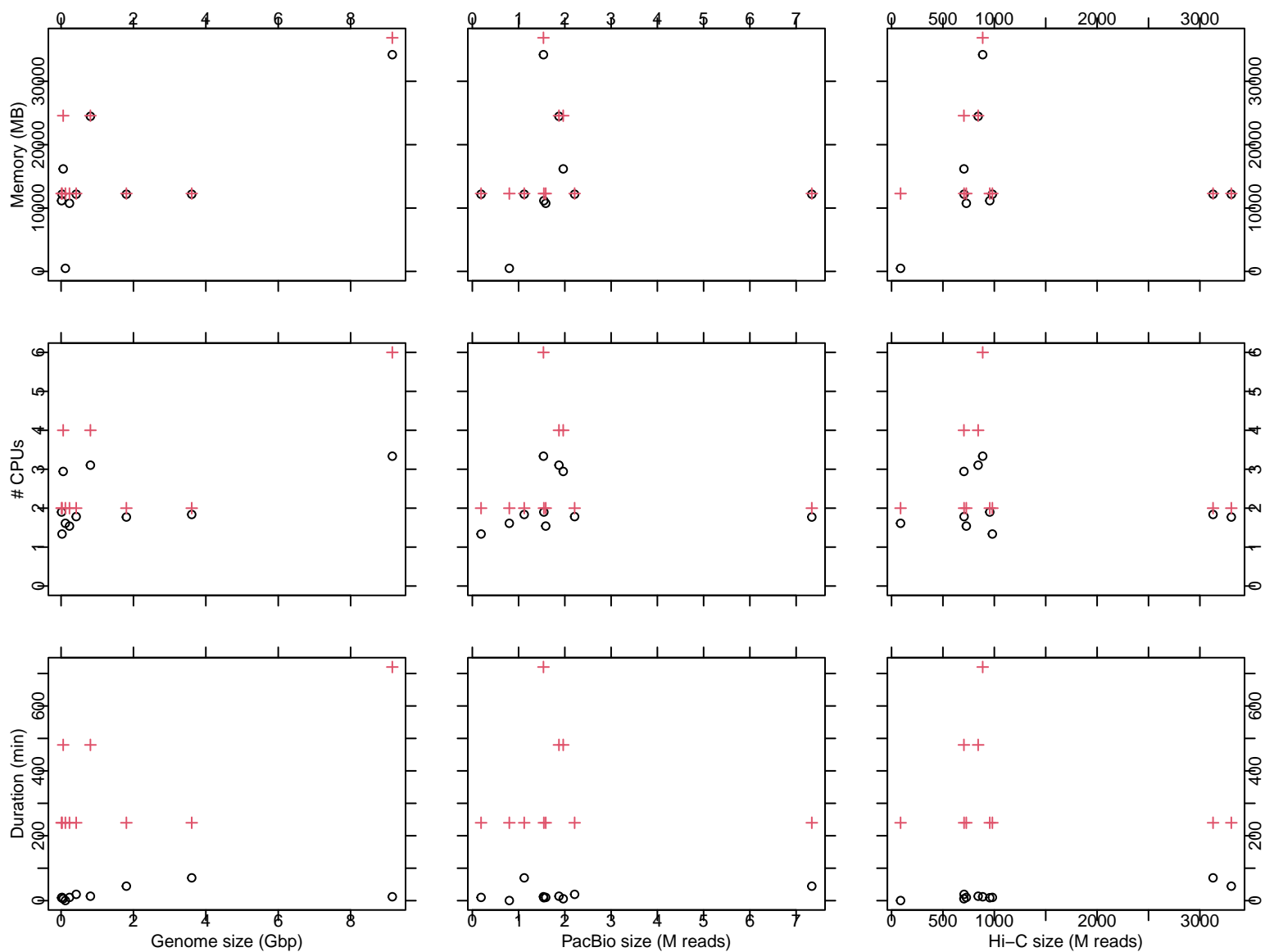
CONTACT_MAPS:COOLER_ZOOMIFY



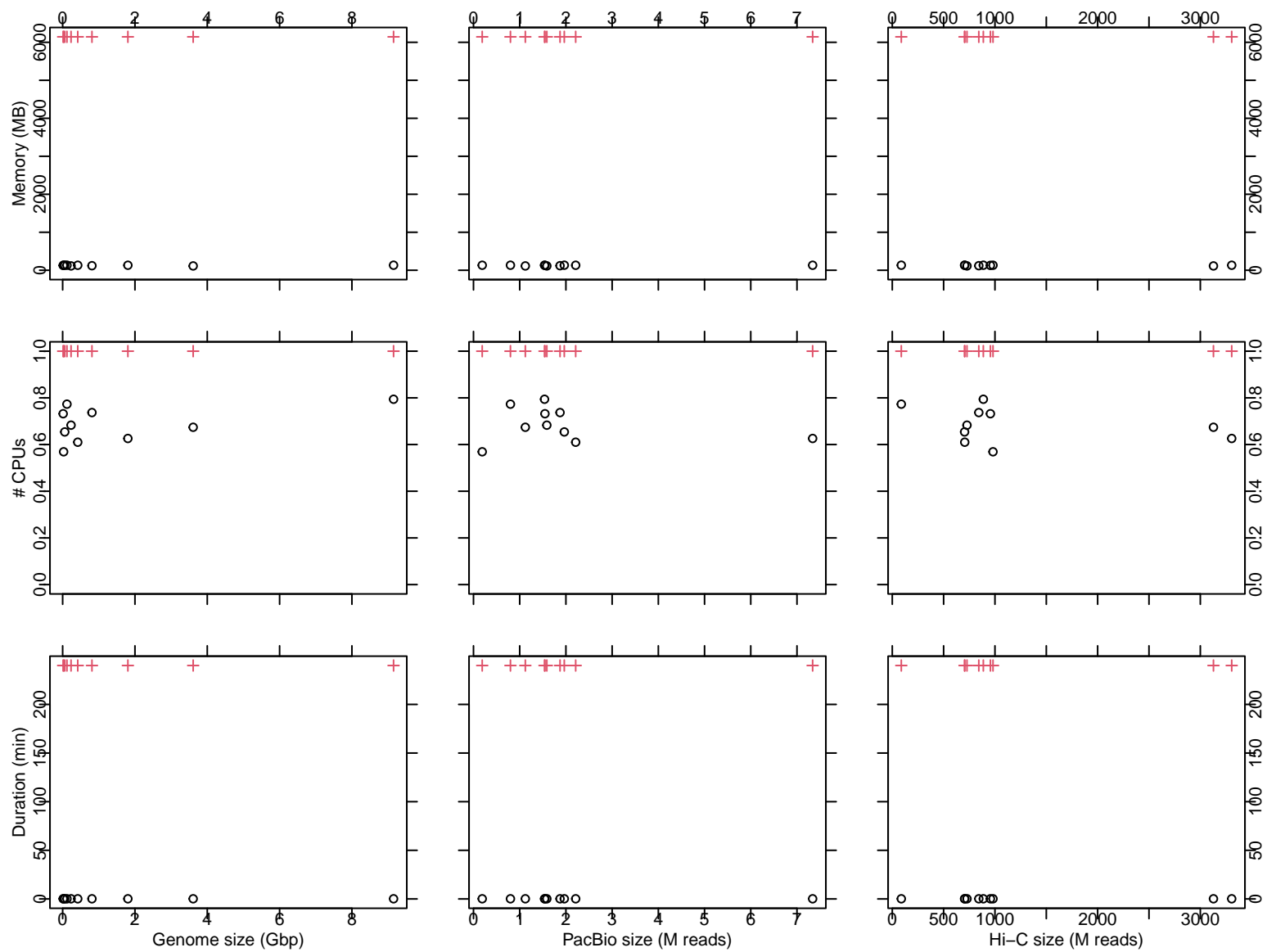
GENOME_STATISTICS:MERQURYFK_MERQURYFK



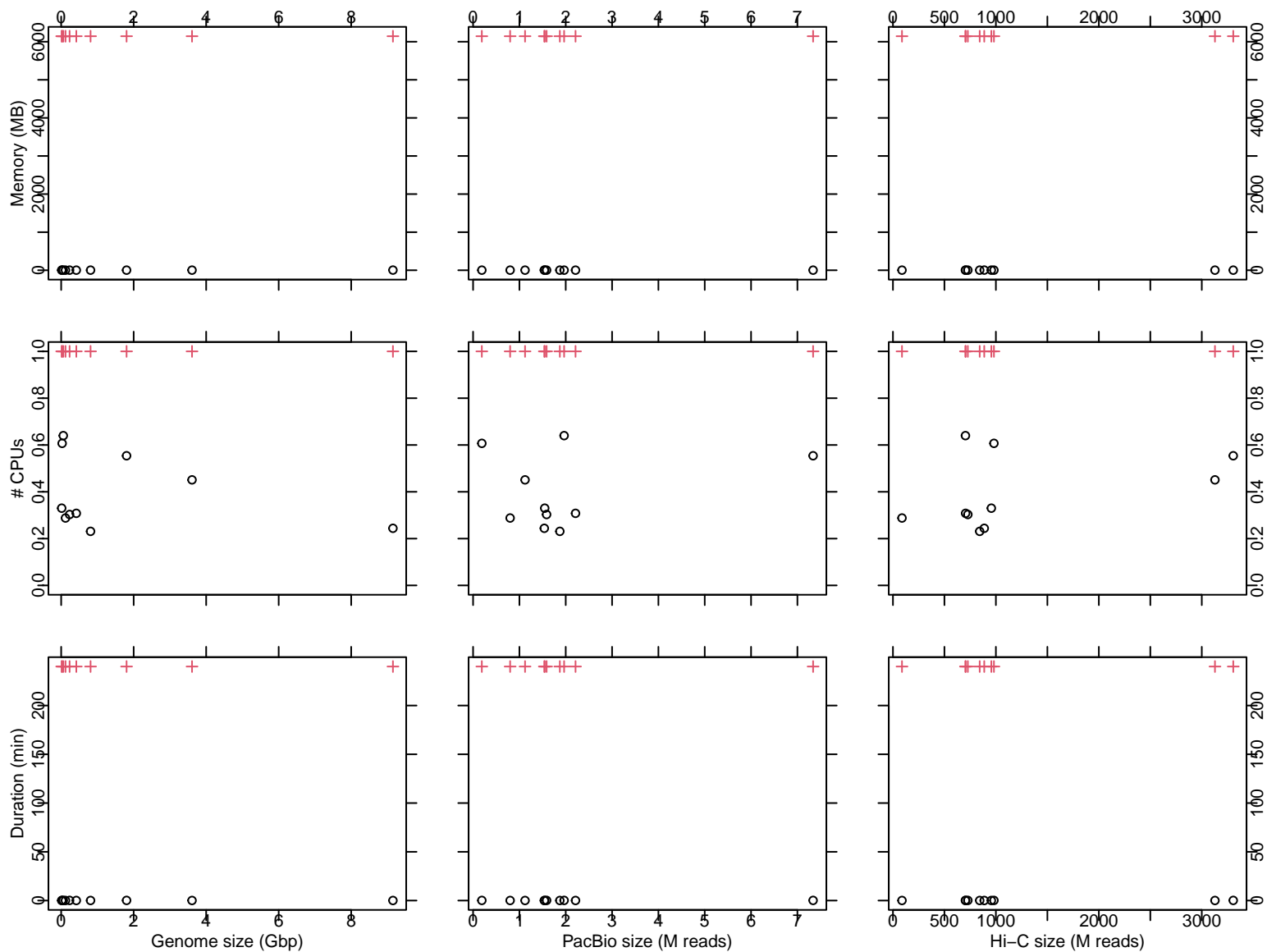
CONTACT_MAPS:FILTER_SORT



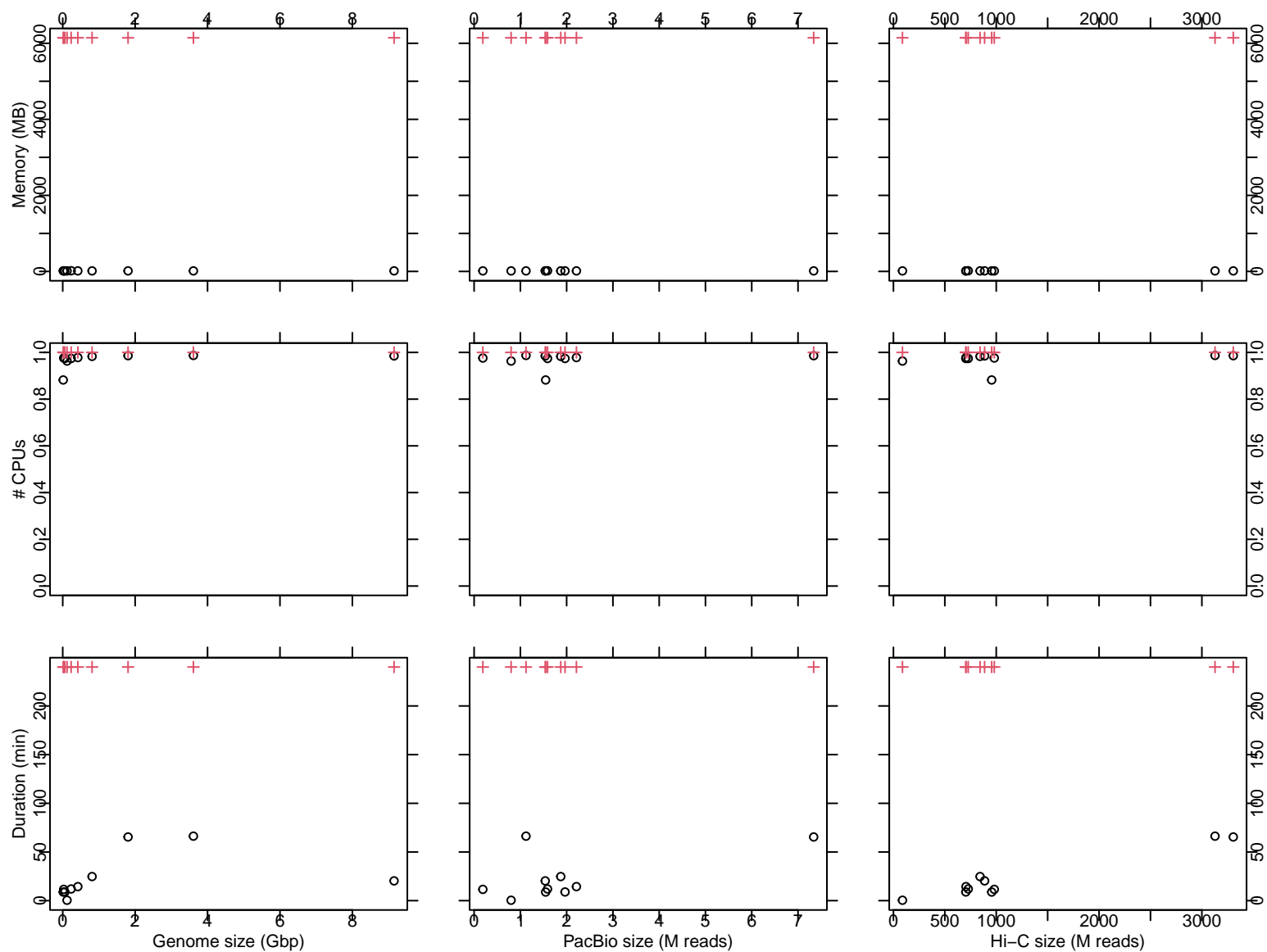
MULTIQC



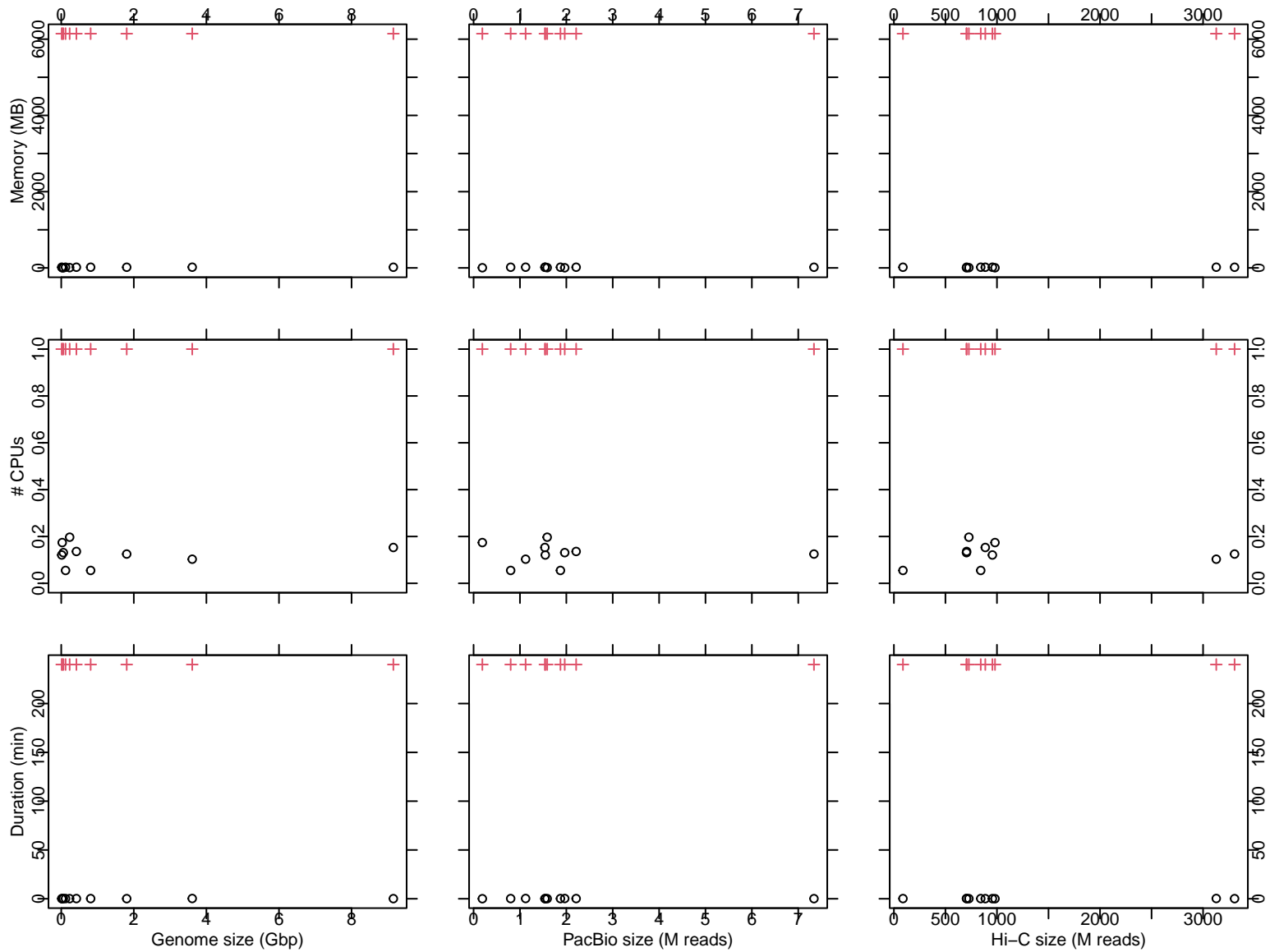
CONTACT_MAPS:FILTER_GENOME



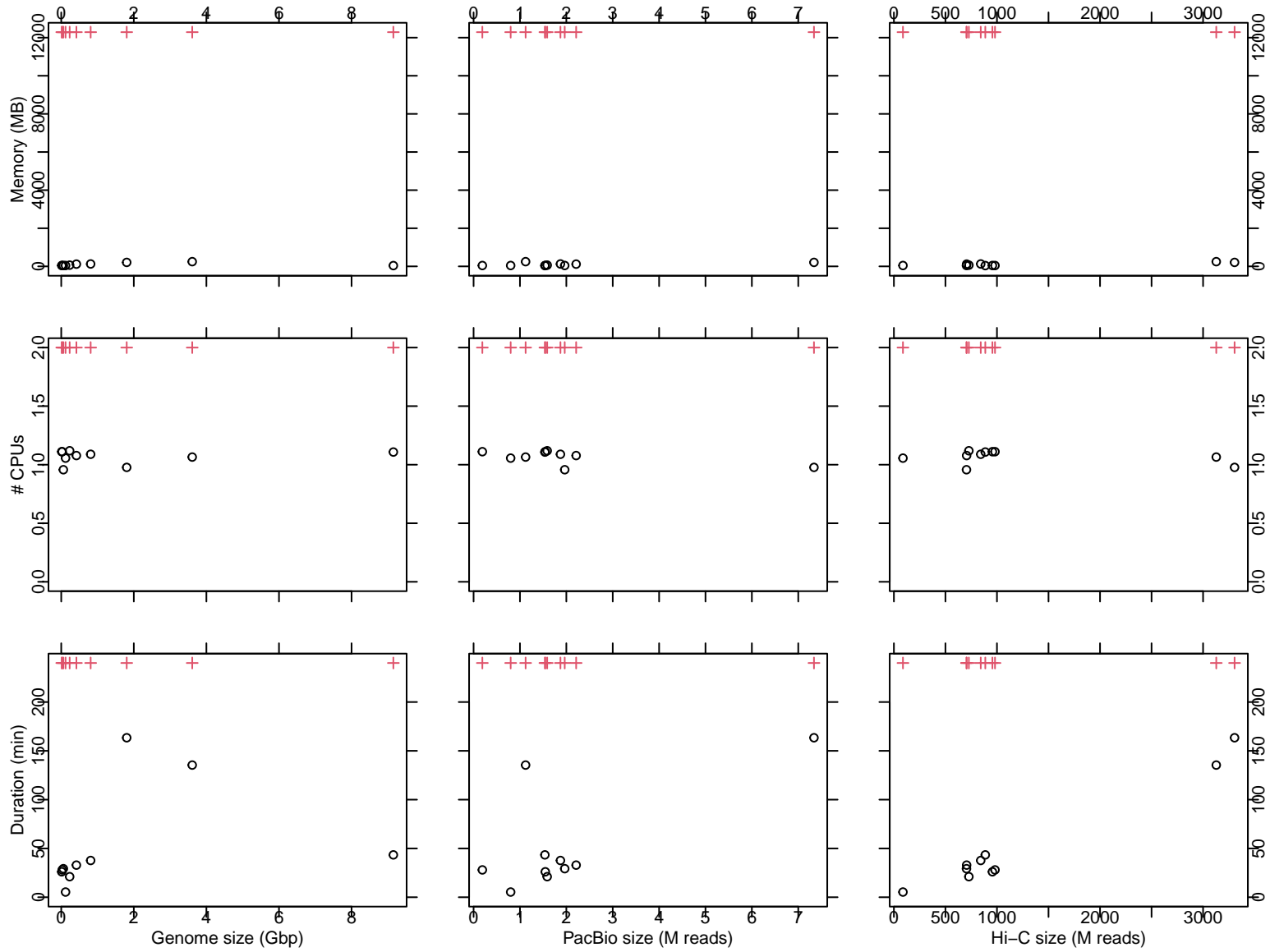
CONTACT_MAPS:FILTER_BED



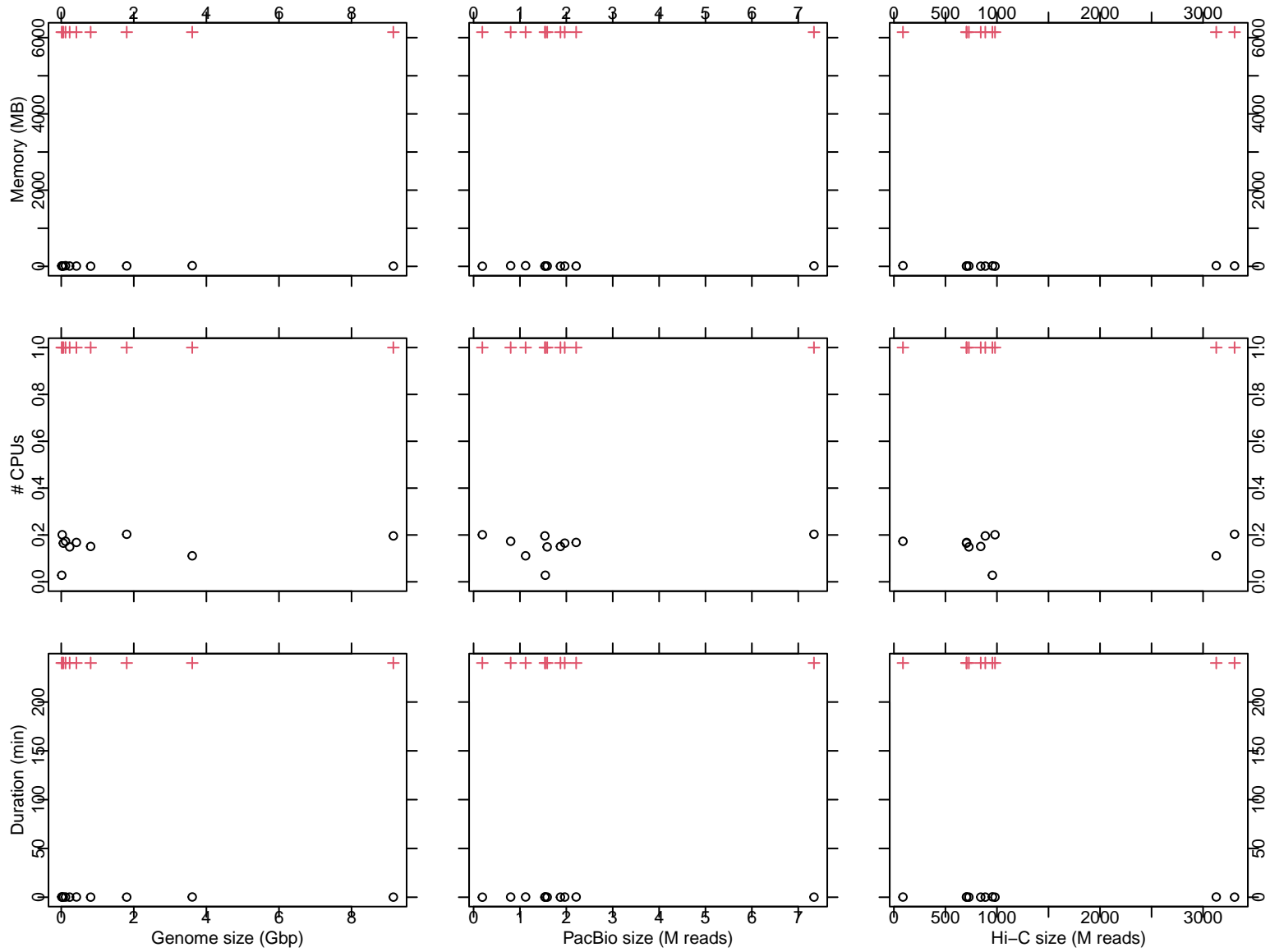
GENOME_STATISTICS:SUMMARYSEQUENCE



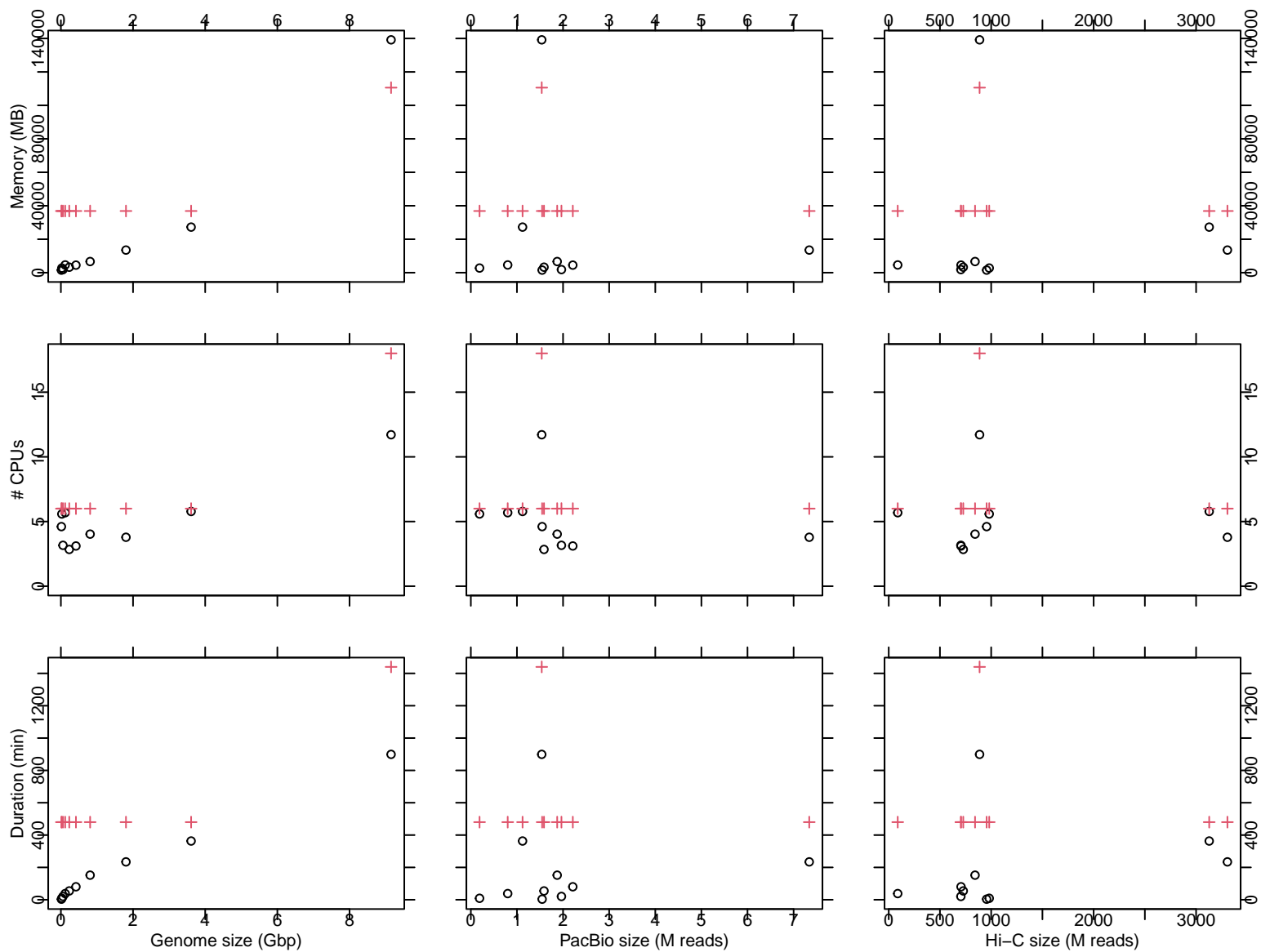
CONTACT_MAPS:SAMTOOLS_VIEW



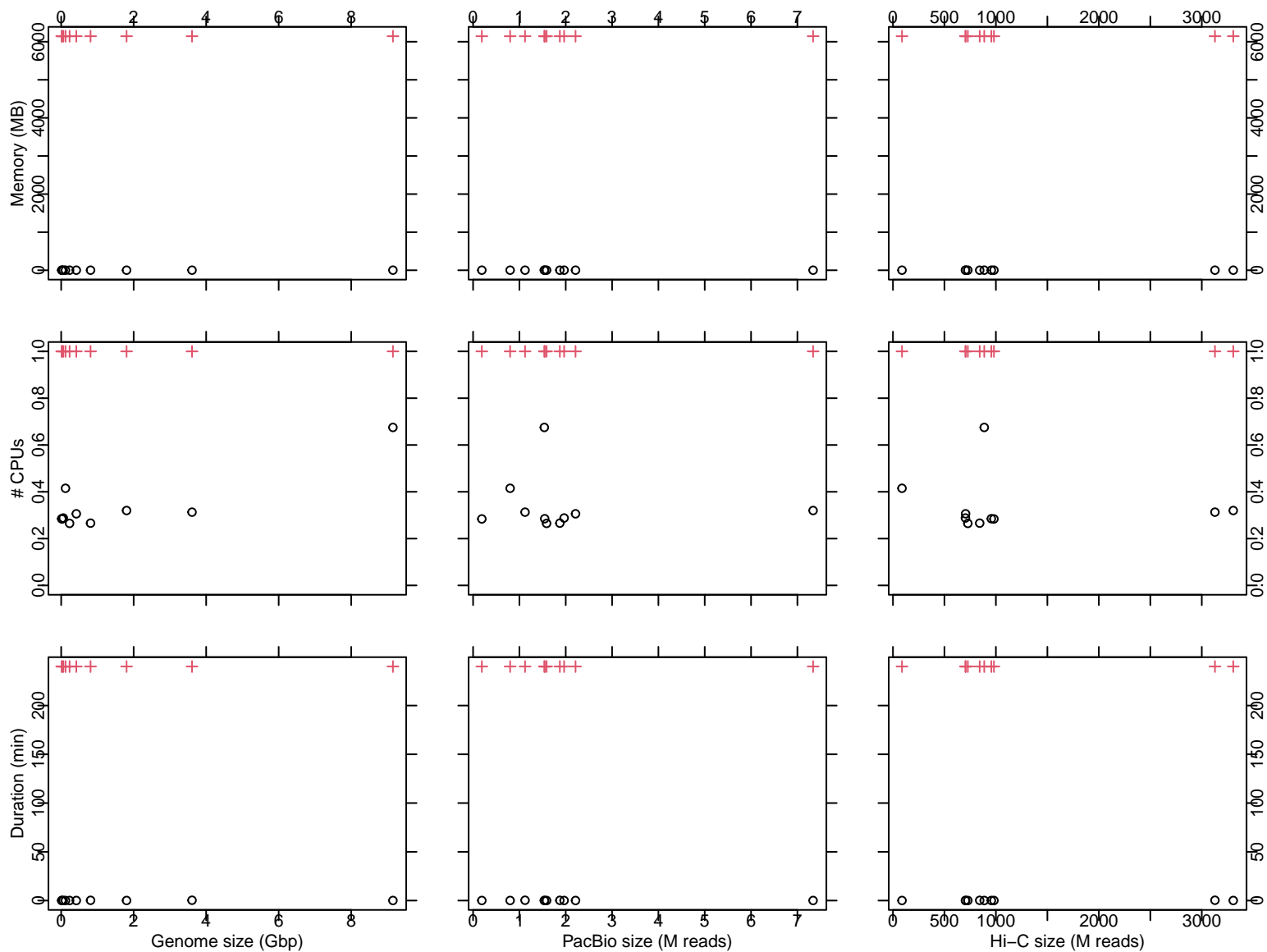
GENOME_STATISTICS:SUMMARYGENOME



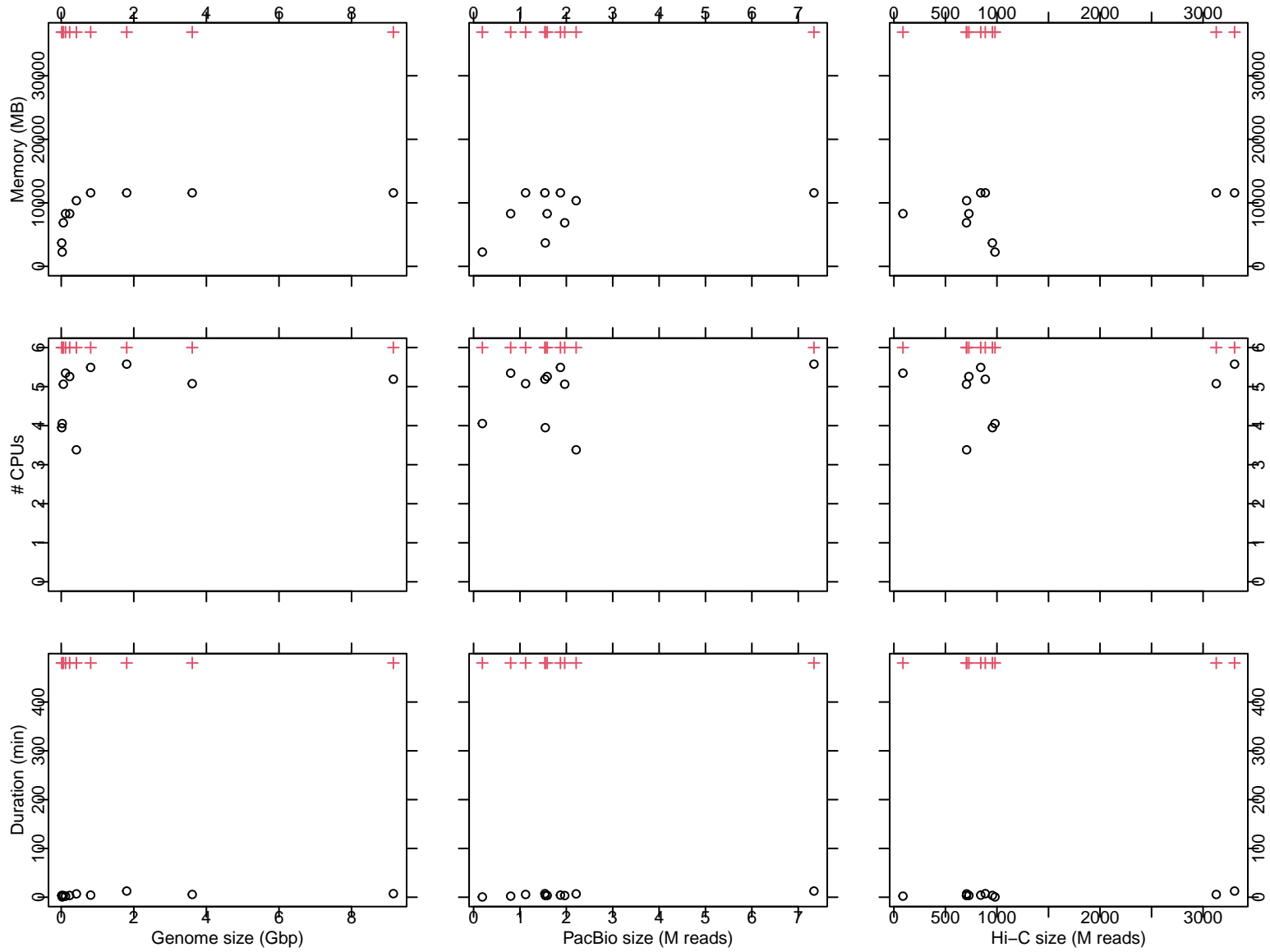
GENOME_STATISTICS:BUSCO



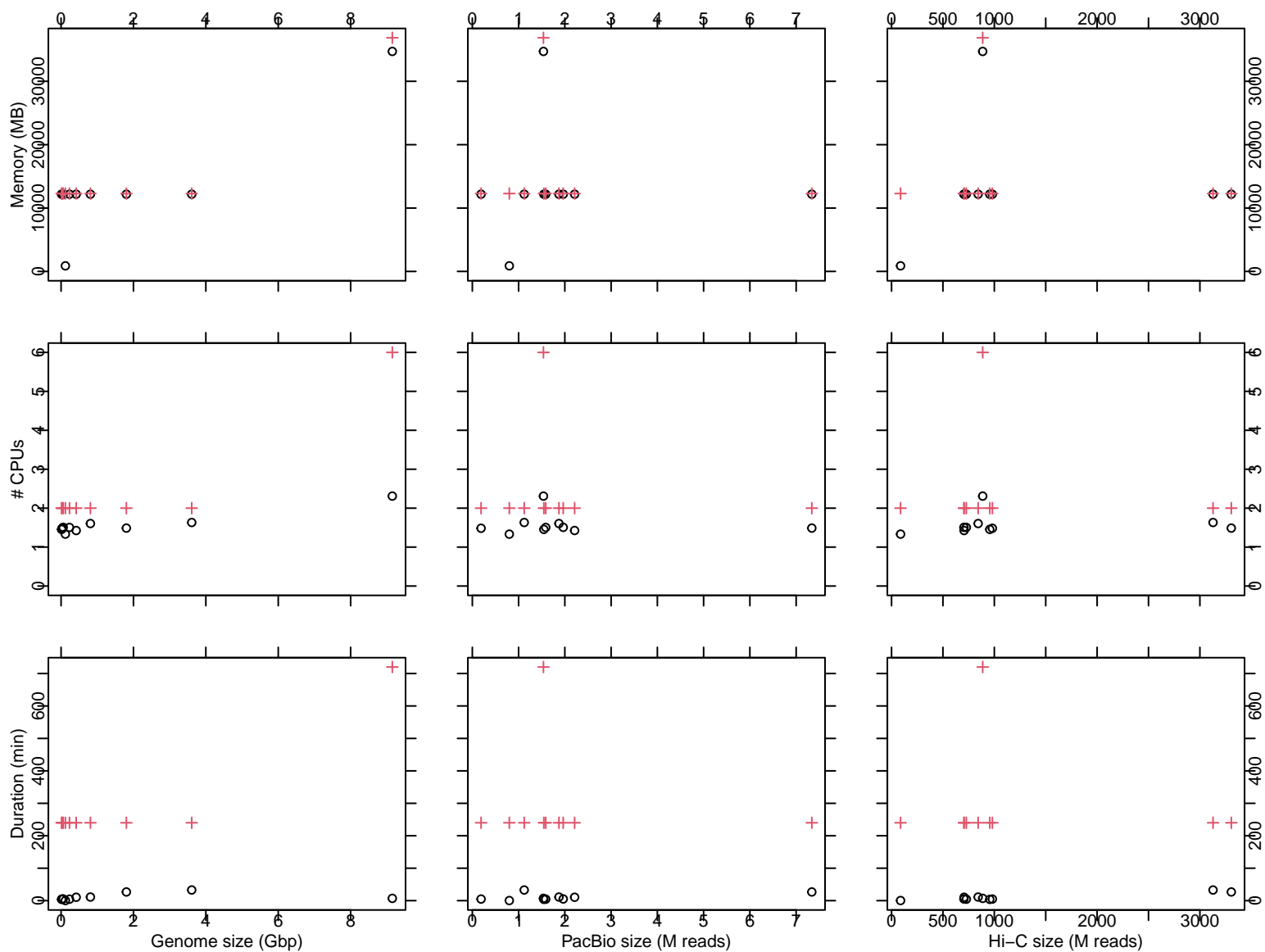
CUSTOM_DUMPSOFTWAREVERSIONS



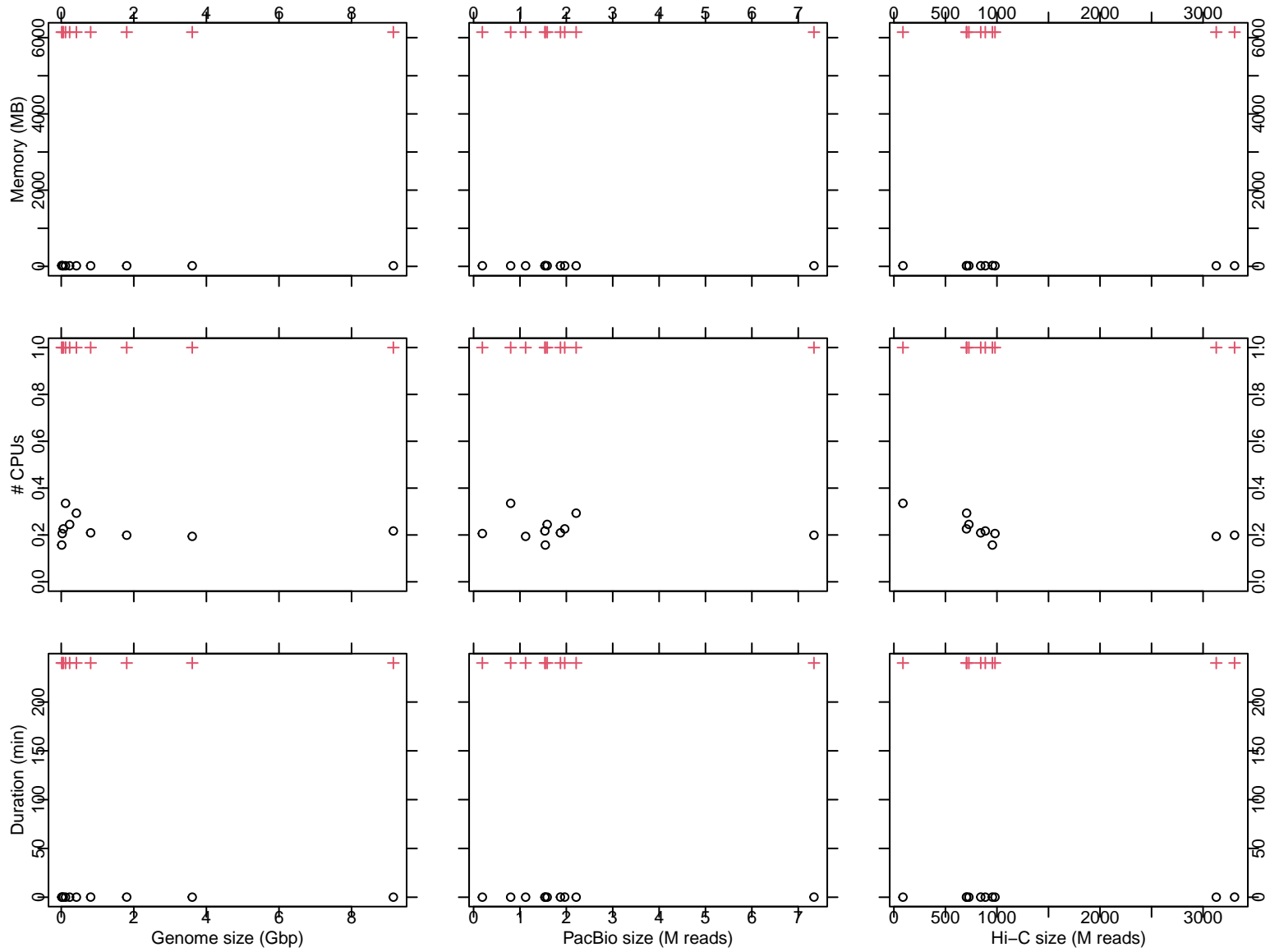
GENOME_STATISTICS:FASTK_FASTK



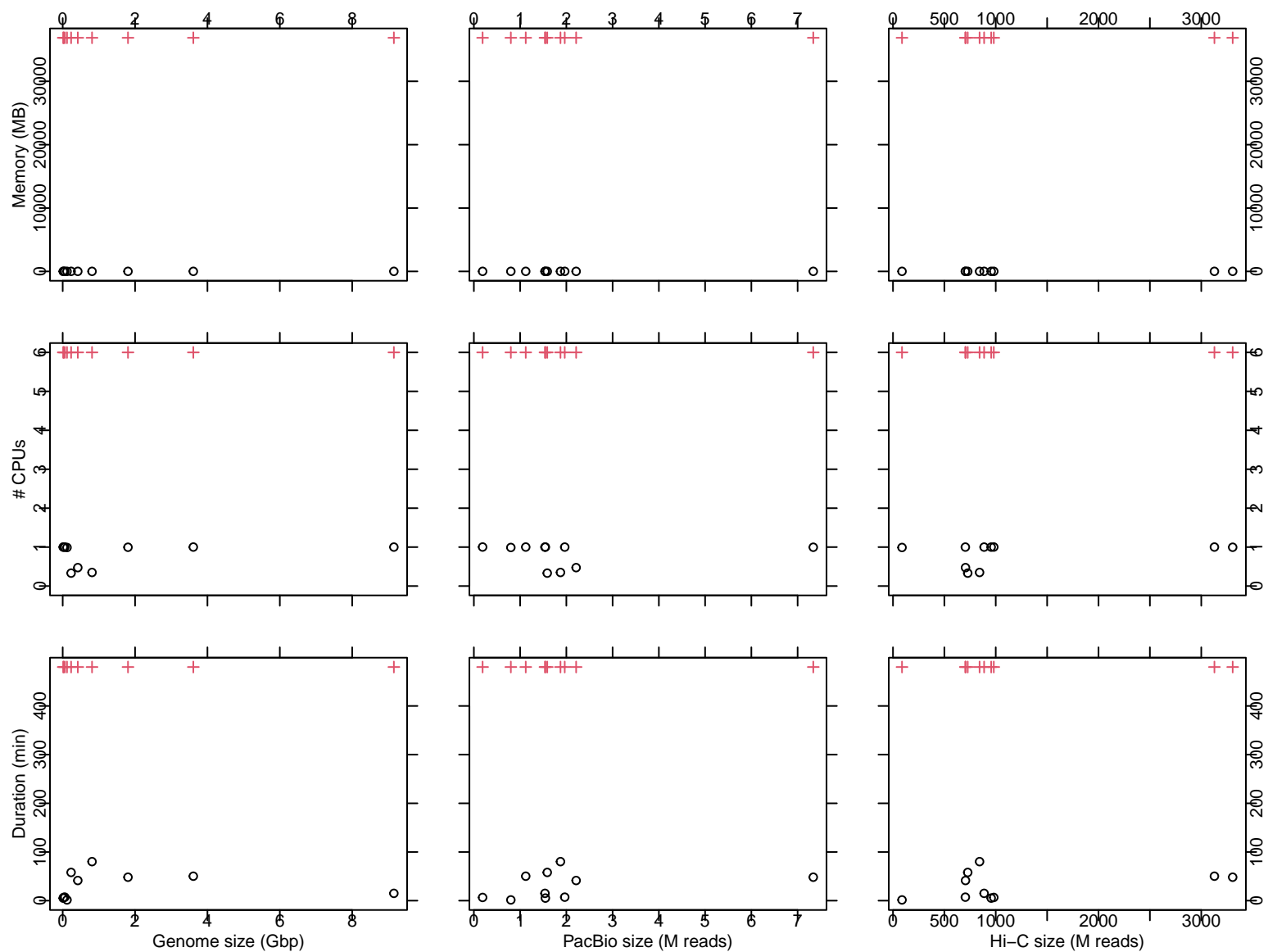
CONTACT_MAPS:BED_SORT



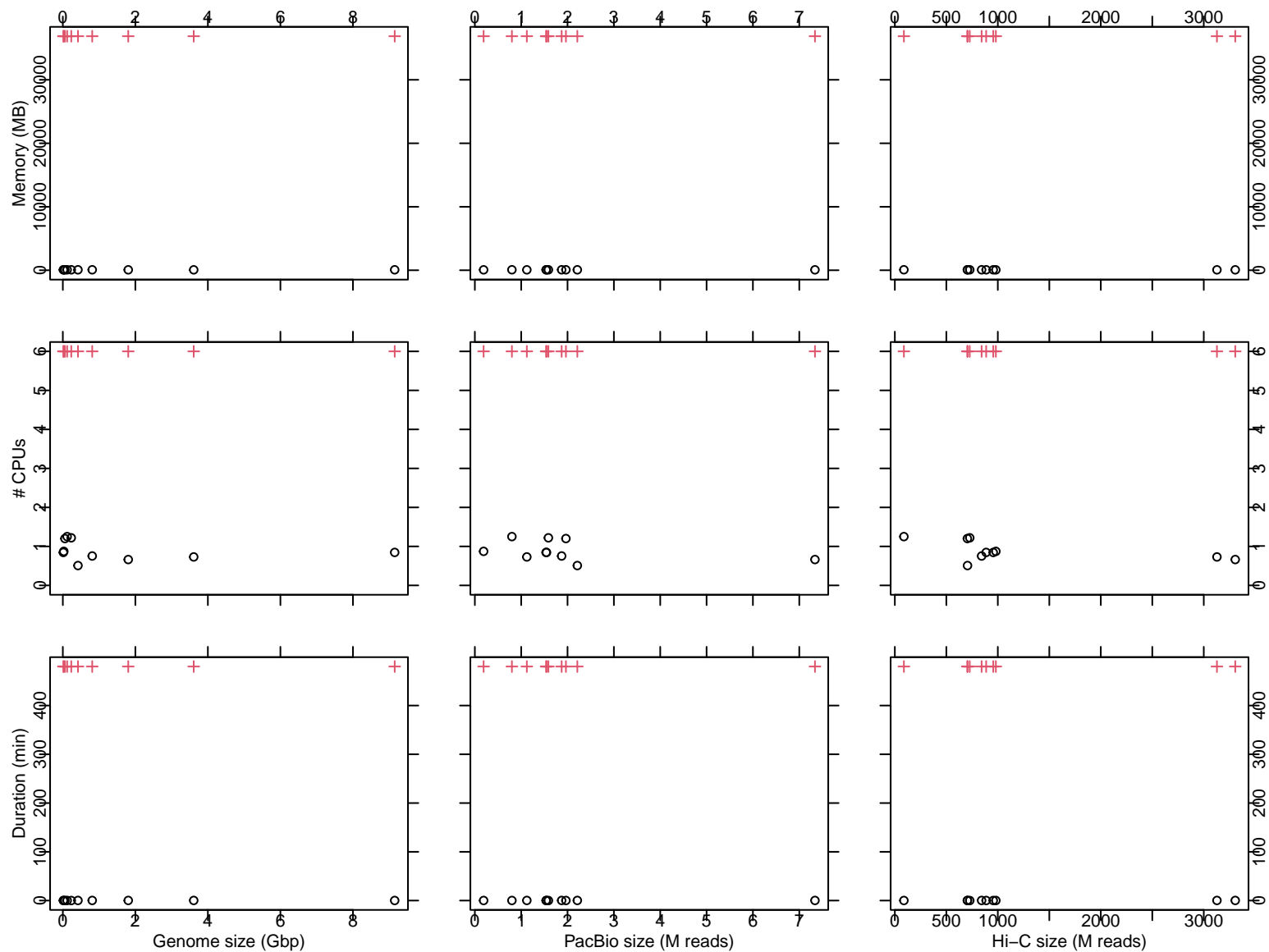
GENOME_STATISTICS:CREATETABLE



CONTACT_MAPS:BEDTOOLS_BAMTOBED



CONTACT_MAPS:COOLER_DUMP



INPUT_CHECK:SAMPLESHEET_CHECK

