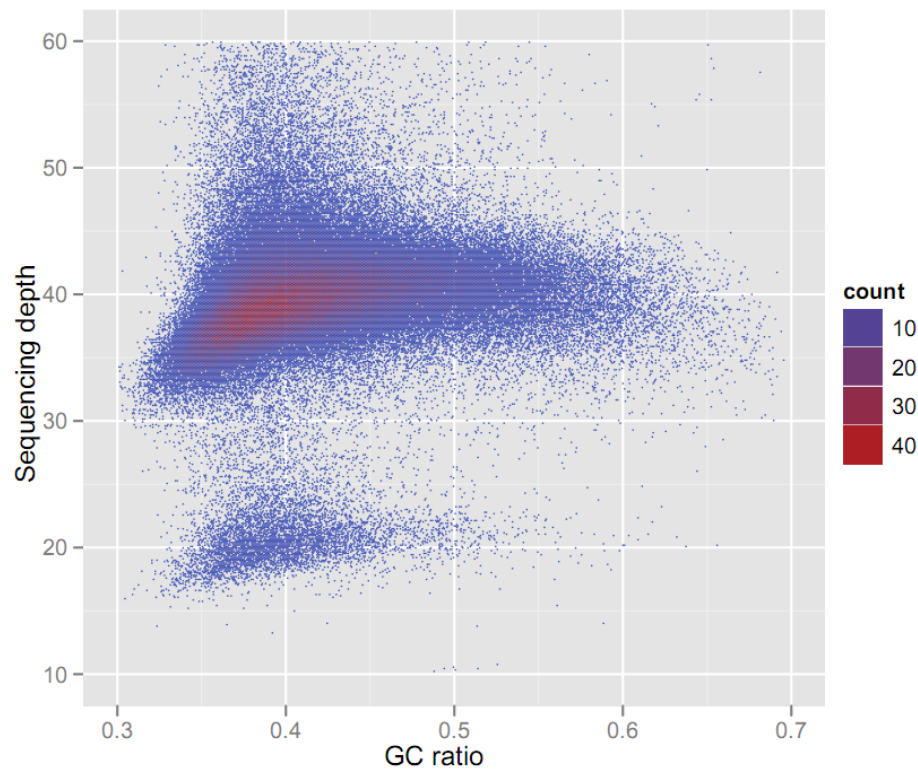




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Supplementary Figure 4: GC content and average sequencing depth values for the *S. caffer* genome. The x and y-axes represent the GC content and average depth values respectively. The dominant GC content value lies within 35-40%. The subgroup (the lower cloud with depth 18-25 and lower GC content) is a representation of the sex chromosomes.

