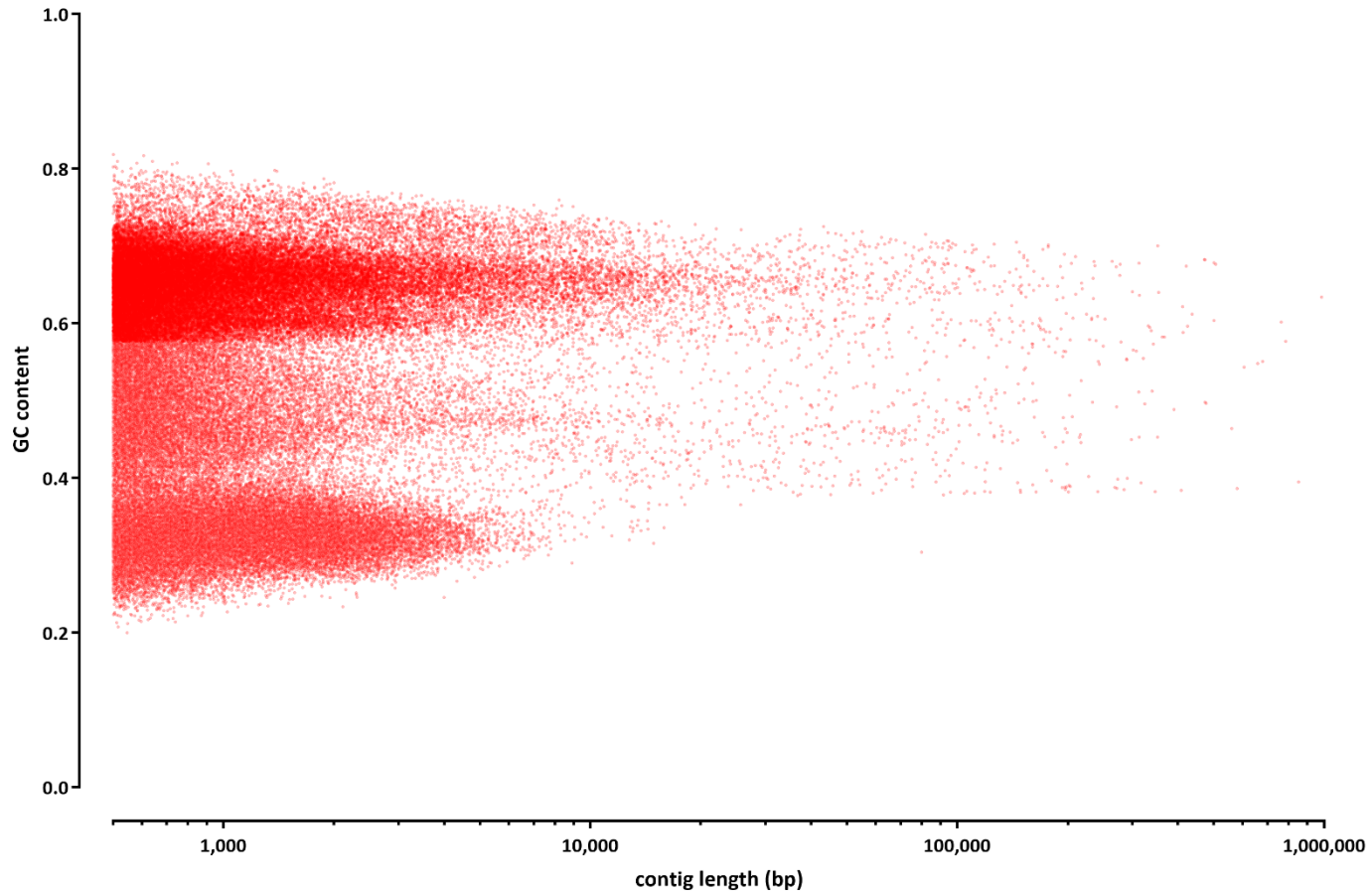
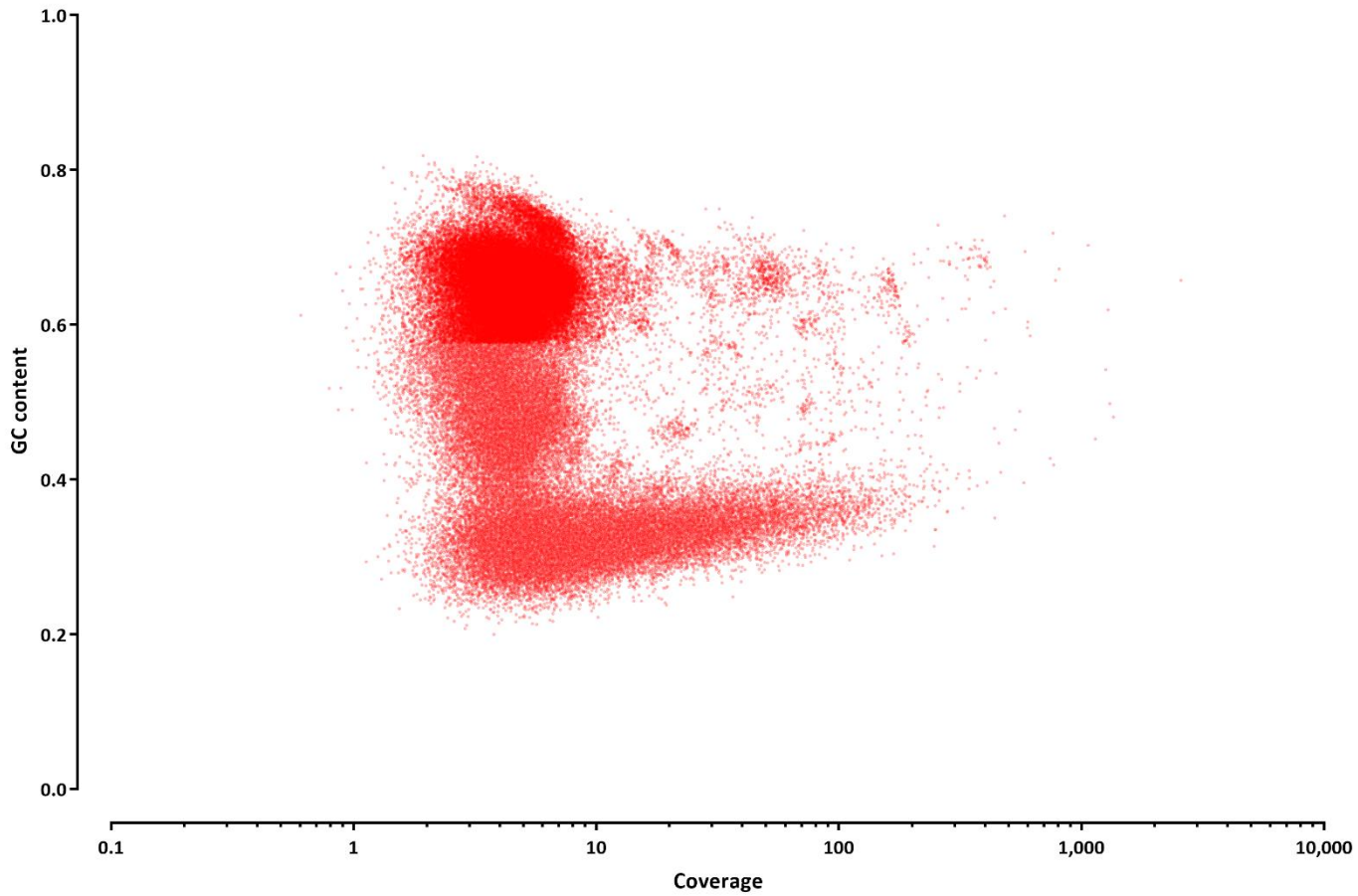


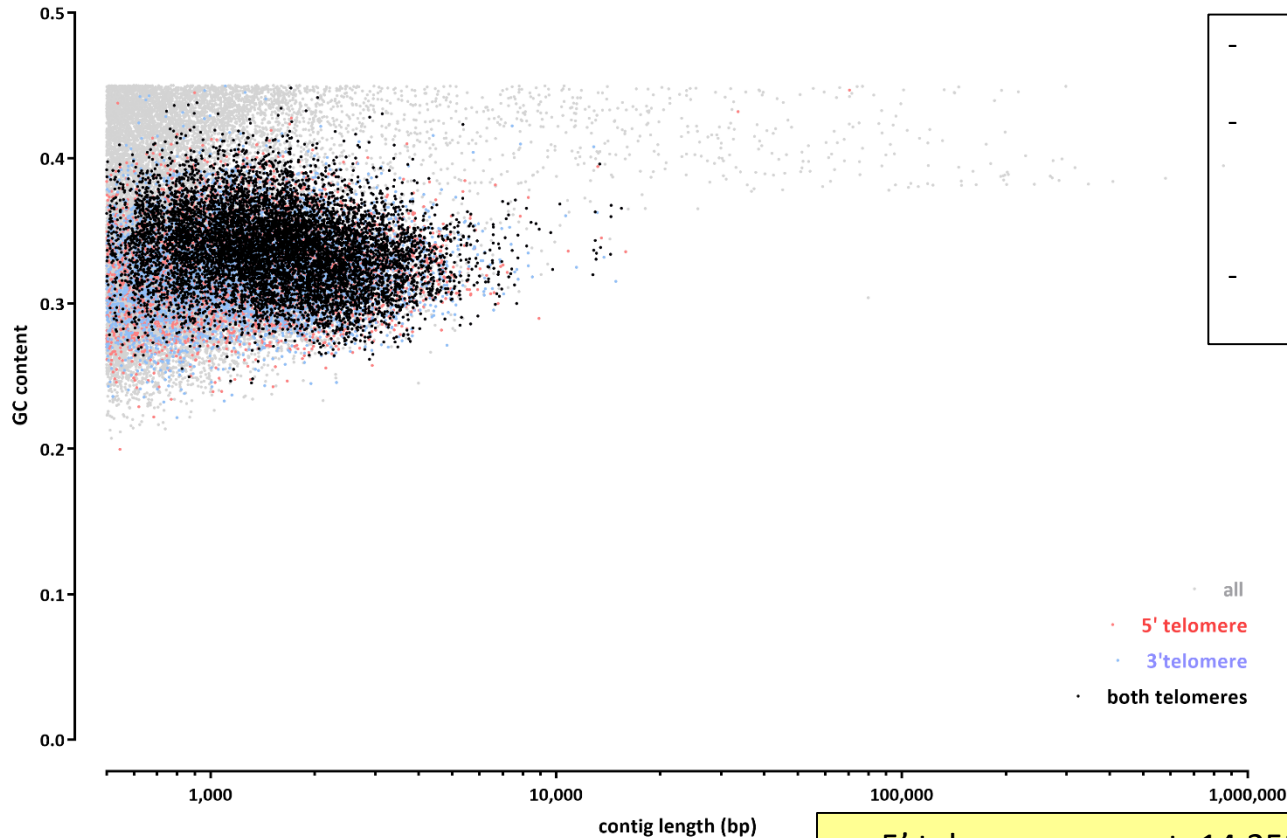
Euplotes gDNA



Euplotes gDNA coverage vs GC



Euplotes gDNA



- Split contigs at GC% = 45
- Those lower than that were assumed to be *Euplotes*, those higher to be *Symbiodinium*
- Looked for 5'-ccccaaaa-3' as the telomeric sequence in Euplotes contigs

- 5' telomere present: 14,252 contigs
- 3' telomere present: 14,290 contigs
- telomeres present on both ends: 8,521 contigs