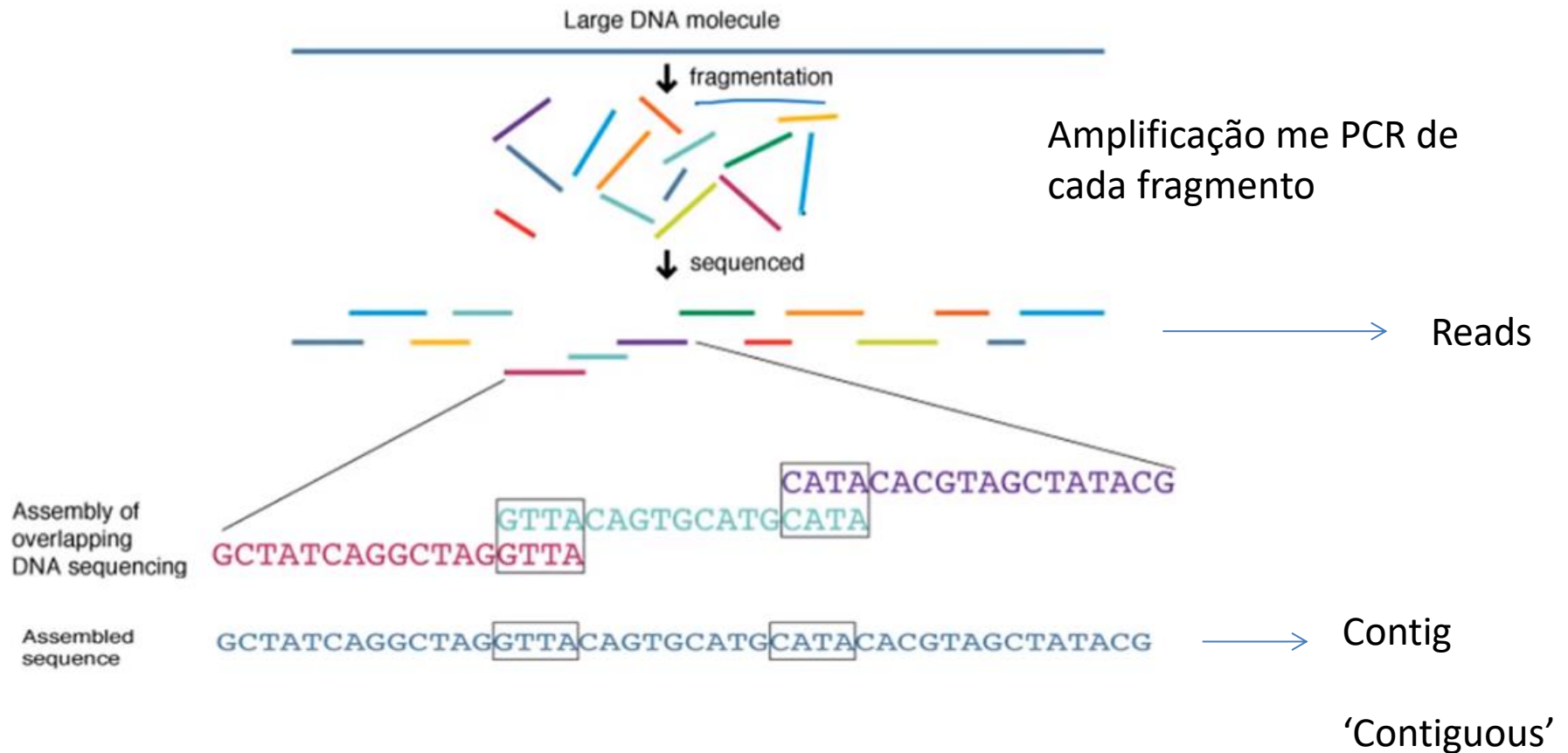


# Atividade 4 - Montagem de Genoma/Algoritmo guloso

## Sequenciamento de DNA SHOTGUN



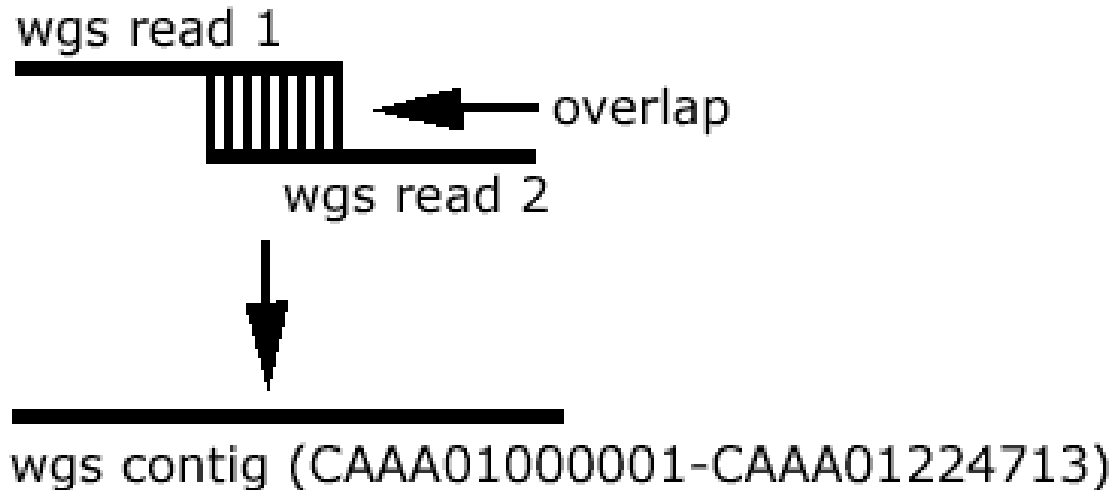


Figure 2. Production of a WGS contig. These contigs contain no gaps, although the sequence may contain 'N's due to sequence ambiguity. WGS contigs obtain accessions similar to the ones shown in the figure, with the first 4 letters representing a project code, the first two numbers representing the assembly version, and the last 6 numbers providing unique identifiers for each contig.

WGA: Whole Genome Assembly

<https://www.ncbi.nlm.nih.gov/assembly/basics/#WGA>

Reconstruct  
this

CTAGGCCCTCAATTTTT  
CTCTAGGCCCTCAATTTTT  
GGCTCTAGGCCCTCATTTTT  
CTCGGCTCTAGCCCCTCATTTT  
TATCTCGACTCTAGGCCCTCA  
TATCTCGACTCTAGGCC  
TCTATATCTCGGCTCTAGG  
GGCGTCTATATCTCG  
GGCGTCGATATCT  
GGCGTCTATATCT

From these

GGCGTCTATATCTCGGCTCTAGGCCCTCATTTTTT