



G A G G A 0 1 1 0 1  
G A A T T 0 0 1 1 0  
A C A A G 0 0 1 0 0  
T A C C A 0 0 1 0 0  
T T A C A 0 1 0 0 0  
A C C T C 0 0 0 0 1  
A A G G A 0 0 0 0 0  
G A T G A 0 0 1 1 1  
T A G A T 0 0 1 0 0  
G A T G A 1 0 0 1 1  
T G T A 0 1 0 0 0  
T A G T A 0 0 0 0 0  
G A T A 0 0 0 0 0  
G A G T T 0 0 0 0 0  
G A G T T 0 0 0 0 0  
T G A T A 0 0 0 0 0  
A T T A G G H T  
T G G H T  
G T R  
R G  
A C

**EMENTA:**

- PPT Introduction
- Get Starting
- View a DNA Sequence
- Side Panel Settings
- GenBank Search and Download
- View a DNA Sequence
- BLAST: Locate a Protein Sequence on the Chromosome
- RNA Folding and Structure Prediction
- Working with Annotations
- Align Sequences
- Primer Design
- Assembly
- Cloning
- Detect Restriction Sites
- Proteolytic Cleavage
- Information Along Sequences
- Phylogenetic trees
- HPC

Material : CLC Main Workbench CLC Cell e CUBE

Recursos: Datashow.