

# Outline

- What is *de novo* assembly?
- CGE Assembler web-tool
- Handling the output

# What is *de novo* assembly?

- It is not reference based...
- CGE Assembler
  - Velvet: De Bruijn Graph
  - Newbler: Overlap Layout Consensus (OLC)
- Remember: Assemblers are far from perfect!

# CGE Assembler

<https://cge.cbs.dtu.dk/services/Assembler/>

## Assembler 1.2

Select type of your reads

454 - single end reads



Choose type of input data

- 454 or Ion Torrent: Newbler (OLC)
- Illumina or SOLID: Velvet

Options:

Trim the reads? ☒

You can use this option if you haven't already trimmed your reads, and you want our trimmer to trim them.

 Isolate File

Name	Size	Progress	Status
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 Upload

 Remove

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## Assembler 1.2

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- 454 or Ion Torrent: Newbler (OLC)
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Trim raw data before assembly?

Isolate File

Name

Upload

Remove

Size	Progress	Status
------	----------	--------

**Min. read length:** read length / 3

**Min. base quality:** 15

**Min. avg. quality:** 20

**Min. adapter match:** 15

# Output



## Assembler-1.0 Server - Results

**Input Files:** *Styph-0808F31478\_3\_1\_sequence.txt Styph-0808F31478\_3\_2\_sequence.txt*

### CONTIGS INFO

**Technology:** *Illumina Paired Ends Reads*

**N50:** *201079*

[DOWNLOAD CONTIGS](#)

# Final note

- Most CGE tools are assembly based
- All assembly based tools use the CGE Assembler
- Optimize your jobs: Do one assembly

Bacterial analysis pipeline:

<https://cge.cbs.dtu.dk/services/cge/>