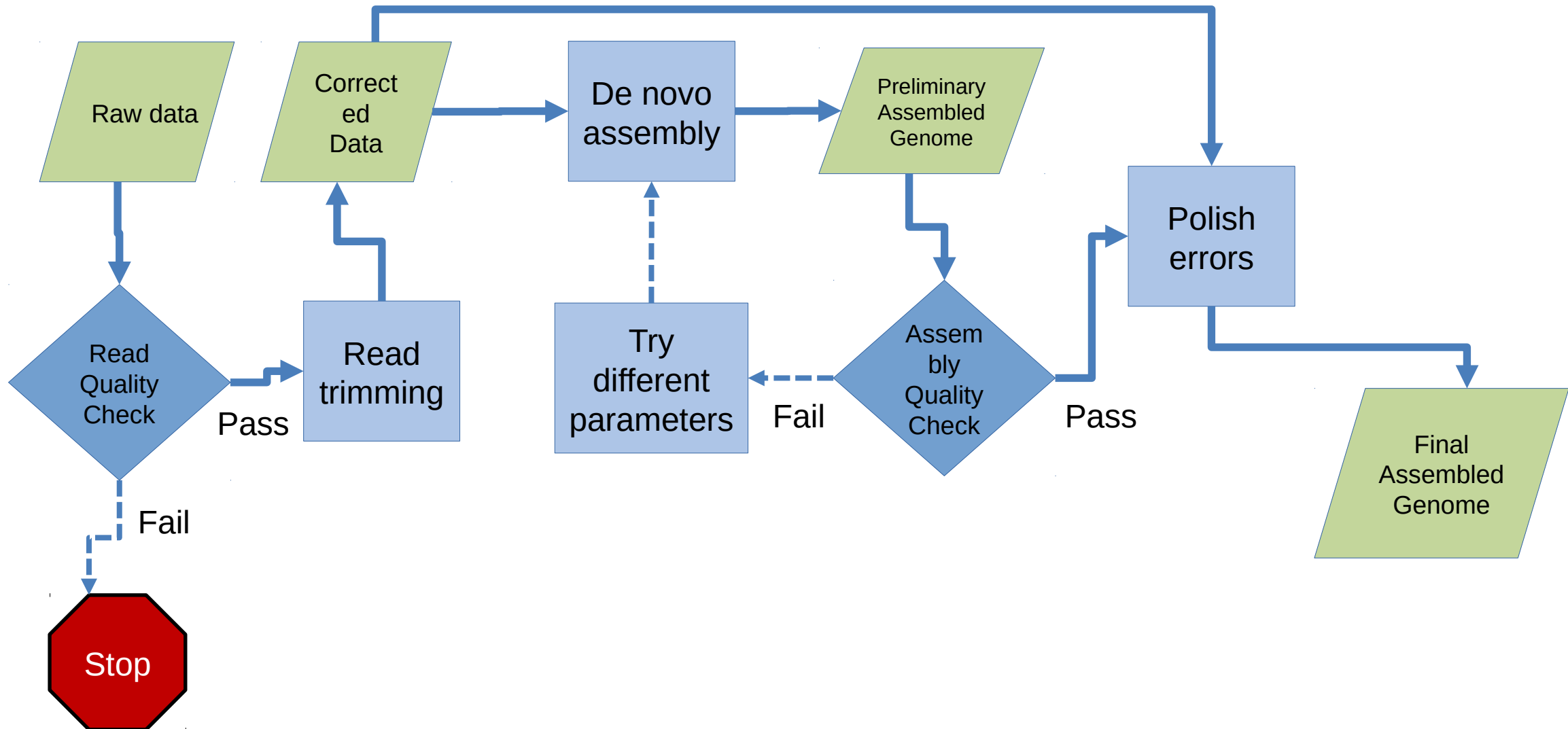
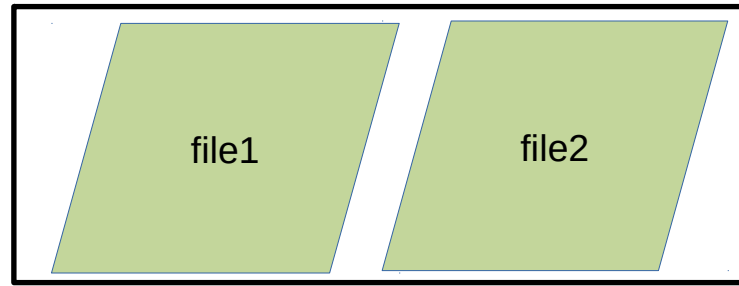


Assembly overview

Assemble the
viral genome
based on
sequences



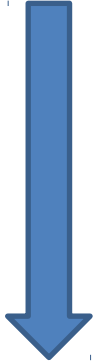
The files necessary to complete this step



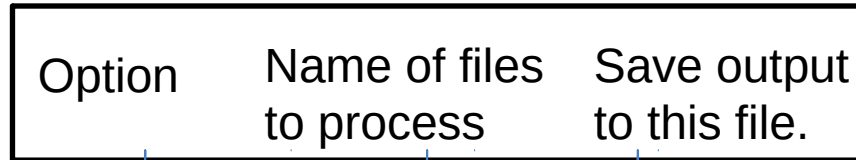
Plain English summary of what the command does

- Start with file1 and file2.
- Run command_name
- End up with file3.

Input:



Breakdown of commands.

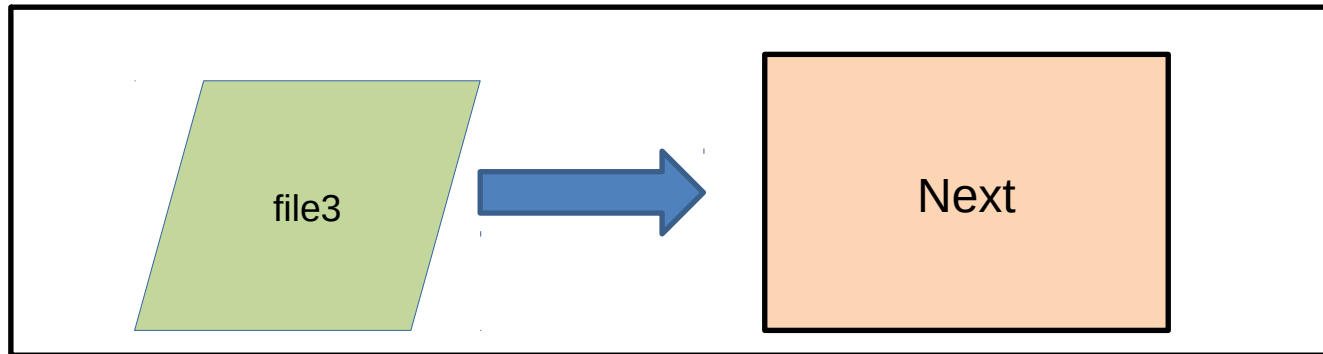


Command Line:

```
command_name --option file1 file2 > file3
```

What to enter on the command line

Output:



Files and reports generated, and the next steps in the analysis.