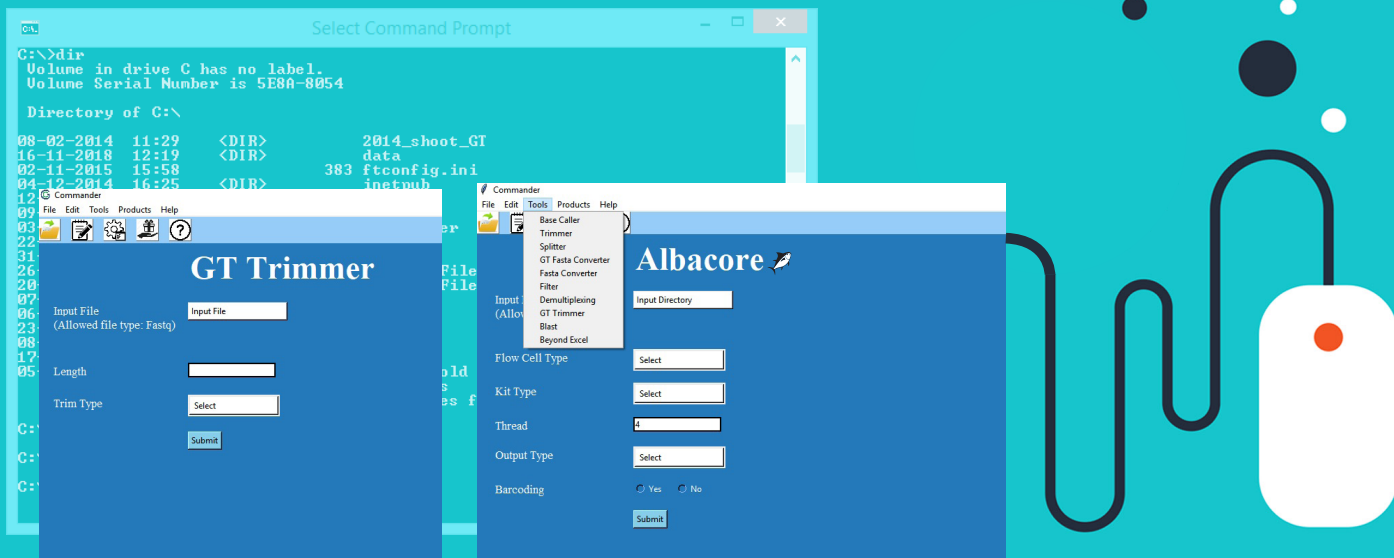


BIOINFORMATICS WITHOUT COMMAND LINE SCRIPTING



COMMANDER (Early access version) is a novel GUI based analysis interface that can **bypass command line scripting**, enabling NGS Data Analysis just at a click of a button. Beyond Excel feature and other tools that are integral to NGS analysis, Commander comes handy for bioinformatics and many other applications.

Commander Early Access Version:

- Base calling of Nanopore FAST5 files
- Conversion of FASTQ to FASTA format
- BCL2 to FASTQ demultiplexing
- QC of NGS raw data
- Trimming of the reads
- Filtering the reads based on length, quality or sequence
- Sequence Search
- Customized analysis and reports*
- Set up and run local blast
- Beyond Excel

* Conditions apply

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