CyVerse Documentation

CyVerse

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CHAPTER 1

Goal

NanoDJ is a Dockerized Jupyter Notebook for interactive Oxford Nanopore MinION sequence manipulation and genome assembly. It integrates basecalling, read trimming and quality control, simulation and plotting routines with a variety of widely used aligners and assemblers, including procedures for hybrid assembly.

- Facilitate ONT sequence analyses by integrating capabilities for data manipulation, sequence comparison and assembly in field experiments or for educational purposes.
- Interactively explore your data with beautiful visualizations that provide new perspectives.
- Easily share results with your team, even those members without NanoDJ installed.

In this quick start, we will show you how to launch NanoDJ VICE app in DE.

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CHAPTER 2

Prerequisites

2.1 Downloads, access, and services

In order to complete this tutorial you will need access to the following services/software

Prerequisite	Preparation/Notes	Link/Download	
CyVerse account	You will need a CyVerse account to complete this exercise	Register	

2.2 Platform(s)

We will use the following CyVerse platform(s):

Platform	Interface	Link	Platform Documen- tation	Learning C Docs	Center
Discovery Environ-	Web/Point-and-	Discovery Environ-	DE Manual	Guide	
ment	click	ment			

2.3 Input and example data

In order to complete this quickstart you will need to have the following inputs prepared.

Note: The example input data (which contains data and Jupyter notebooks) is already prefilled for you during app launch

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Input File(s)	Format	Preparation/Notes	Example Data
Fast5 and Jupyter note- books	Fast5Jupyter notebooks	Input data can be a list of FAST5 files from previ- ous basecalled runs (e.g. a Metrichor output) or event	data_transfer. sh (Community Data -> iplantcollabora- tive -> example_data -> nanodj and select

CHAPTER 3

Get started: Launch NanoDJ

- 1. Login to the.
- 2. Click on Apps window in the DE workspace and search for and run NanoDJ.

Tip: Alternatively you can clik this



button to directly launch the NanoDJ VICE app.

- 3. Under "Analysis Name" leave the defaults or make any desired notes.
- 4. Under "Parameters" for 'Input files', click the "+" and browse data_transfer.sh (iplantcollaborative > example_data > nanodj and select data_transfer.sh script).
- 5. Click **Launch Analysis**. You will receive couple of notifications on the bell corresponding to job submission and running with the "Access your running analysis here".
- 6. Clicking on the "Access your running analysis" will open the NanoDJ JupyterLab in another tab in the browser after a brief building phase.

Note: You will be asked to authenticate again to the JupyterLab with your CyVerse username and password

7. Open the Terminal in the main JupyterLab interface (under 'others' section), navigate to the vice folder and execute the data_transfer.sh script to initiate the test data downloading process. You will be asked to enter your CyVerse password again to initiate the downloading process.

```
cd vice
bash data_transfer.sh
Enter your current iRODS password:
```

8. Finally, once you finish analysis, navigate to the Discovery Environment tab, select the Analysis window and select the analysis, click "save and complete analysis". Upon clicking complete analysis, the analysis will be completed and all the outputs will be brought back to the analysis folder. Alternatively you can use iput -rPVT <folder name> command to transfer the data back to the data store.

Note: Currently quick starts uses a reduced test data but if you want to try the full test data, then instead of using data_transfer.sh, use data_transfer_full.sh script which is in the same folder as the other script.

3.1 Additional information, help

- Full materials for the webinar is available here
- See the original JupyterLab quick start
- See the original NanoDJ for how to run ONT data analysis

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• Contact CyVerse support by clicking the intercom button on the page.

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