

Run CloVR-Comparative

1. Upload data

1. On Data sets tab: **Add**

2. Paste URL:

http://cb2.igs.umaryland.edu/neisseria_comparative.tgz

3. **File Type:** "GenBank"

4. **Name:** e.g. "neisseria_5genomes_test"

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2. Configure Pipeline

1. Select from applications panel: **Comparative**
2. **Input Genbank Tags:** "neisseria_5genomes_test"
3. **Account:** "local"
4. **Pipeline Description:** e.g. "neisseria_test"
5. **Submit**