

Run CioVR-Comparative

1. Upload data

1. On Data sets tab: **Add**
2. Paste URL:
<http://cb2.igs.umaryland.edu/compare.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. **Pipeline Description**: e.g. “asm_comparative”
4. **Run**