

## Breseq Protocol

1. Upload your data to `data/` folder.

Your fastq files needs to be directly in the folder you upload to `data/`.

DO NOT change the name of your fastq files.

### Example folder structure:

Here we have two samples `test` and `test2` in the `data/` folder.

```
data/  
  test  
    test_R1.fastq.gz  
    test_R2.fastq.gz  
  test2  
    test2_R1.fastq.gz  
    test2_R2.fastq.gz
```

2. Open terminal in the breseq folder.
3. Run the following command in the terminal Replace `data/test` with the path to your data folder.

```
sbatch main_slurm.sh data/test
```