# **Bioinformatics** session

A two-day workshop for bioinformaticians and molecular biologists with focus on the TSO500 pipeline in InPreD



#### **Overview**

- 1. Setup
- 2. Development & Collaboration
- 3. Nextflow
- 4. tso500\_nxf\_workflow
- 5. Python

# 1. Setup

• go to https://github.com/ and click on Sign up



• enter your email

```
Welcome to GitHub!

Let's begin the adventure

Enter your email*

→ coder@inpred.no

Continue
```

set a password

```
Welcome to GitHub!
Let's begin the adventure
Enter your email*

√ coder@inpred.no

Create a password*
                                                          Continue
```

• choose a username

```
Welcome to GitHub!
Let's begin the adventure
Enter your email*

√ coder@inpred.no

Create a password*
  •••••
Enter a username*
→ inpredder
                                                       Continue
```

• choose email preferences



• solve the puzzle





• create your account



find the activation code in the email you received



Here's your GitHub launch code, @inpredder!



select the desired options

This will help us guide you to the tools that are best suited for your projects. How many team members will be working with you? Just me O 2-5 5-10 0 10-20 20-50 50+ Are you a student or teacher? O N/A Student Teacher Continue

# What specific features are you interested in using?

Select all that apply so we can point you to the right GitHub plan.

Collaborative coding

Codespaces, Pull requests, Notifications, Code review, Code review assignments, Code owners, Draft pull requests, Protected branches, and more.

Automation and CI/CD

Actions, Packages, APIs, GitHub Pages, GitHub Marketplace, Webhooks, Hosted runners, Self-hosted runners, Secrets management, and more.

Security

Private repos, 2FA, Required reviews, Required status checks, Code scanning, Secret scanning, Dependency graph, Dependabot alerts, and more.

• choose the free plan



### 1. Setup

Be added to InPreD organisation at GitHub

#### 1. Resources

• Getting started with your GitHub account

#### 2. Development & Collaboration

#### Short git introduction

- distributed version control system
- tracks history of changes committed by different contributors
- every developer has full copy of project and its history

#### Short git introduction

#### Basic git commands

```
git init : initialises new git repository
git clone <repository url> : creates local copy of remote repository
git add <file/s> : stage new or changed files (anything that should be committed to the repository)
```

git commit -m "feat: my new feature" : commit changes to the repository

#### Basic git commands

#### commit message conventions

```
<type>[optional scope]: <description>
```

- feat : new feature
- fix : patching bug
- refactor : code change that neither is neither feat nor fix
- build : build system related changes
- perf: improving performance

#### commit message conventions

```
<type>[optional scope]: <description>
```

- chore : code unrelated changes, e.g. dependencies
- style : code change that does not change meaning
- test : changes to tests
- docs : adding/updating documentation
- ci : continuous integration, e.g. github actions

#### Basic git commands

git status: overview over untracked, modified and staged changes

git branch: show local branches

git merge: merge branches

git pull: load changes from remote counterpart

git push: upload changes to remote counterpart

**Branching strategy - Gitflow** 

- github actions (linting, testing, building)
- pull requests (best practice)
- release and semantic versioning
- licensing

#### 2. Resources

- About Git
- Gitflow workflow

#### 3. Nextflow

- general (install, best practice)
- nf-core template
- stubbing

# 4. tso500\_nxf\_workflow

- status
- demonstration

# 5. Python

- general (best practice, cli)
- unit testing (pytest)

#### Resources

- github actions
- nf-core
- pytest unittesting
- semantic versioning