

NEXTFLOW VS. SNAKEMAKE

	NEXTFLOW	SNAKEMAKE
Platform	Groovy/JVM	Python
Workflow modules	Yes	Yes
Docker support	Yes	Yes
Conda support	Yes	Yes (same author for conda project)
Build-in batch schedulers	Yes	Yes
Remote files	Yes	Yes

Ref: https://doi.org/10.1038/nbt.3820

STYLE

```
NEXTFLOW

process NAME {
    label: ...
    when: ...
    input: ...
    output: ...
    script: ...
```

```
SNAKEMAKE
rule NAME:
        version: ...
        input: ...
        output: ...
        resources: ...
        log: ...
        script: ...
        shell: ...
        notebook: ...
```

WORKFLOW

```
NEXTFLOW
process foo { ... }
process bar { ... }
workflow NAME {
       take: ...
       main:
               foo() | bar()
       emit: ...
```

```
SNAKEMAKE

subworkflow foo:
    output: pipe(...)
    ...

rule NAME:
    input: foo()
    output: ...
    shell: ...
```

GET STARTED



Requirements: Java 8 (or later)



Installation:

wget -qO- https://get.nextflow.io | bash

HELLO WORLD

```
input = Channel.of("Hello", "Hi")
workflow {
            input | p1
            p1.out.result | p2 | view
process p1{
            input: val(word)
            output: val(result), emit: result
            exec: result = "$word world"
process p2{
            input: val(data)
            output: val(res)
            exec: res = data.toUpperCase()
```

```
$ nextflow test.nf

NEXTFLOW ~ version 20.10.0

Launching `test.nf` [stoic_meucci] - revision: 565aea2060

executor > local (4)

[6a/c3ef0e] process > p1 (1) [100%] 2 of 2 

[dc/1d0443] process > p2 (2) [100%] 2 of 2 

HI WORLD

HELLO WORLD
```

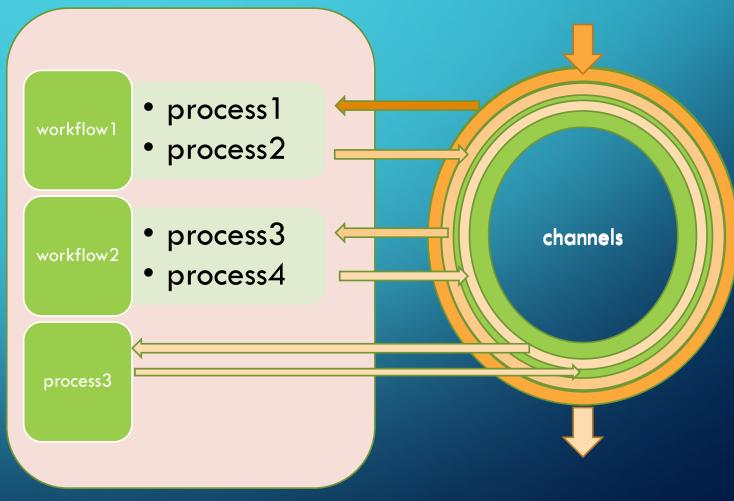


BASIC CONCEPTS

- params: an object can be access by whole pipeline
- channel: asynchronous FIFO queues
- process: a function unit
- workflow: a function unit by multiple process



UNDERSTAND THE CONCEPTS







MORE INFORMATION

- https://nextflow.io/
- https://nf-co.re/
- https://groovy-lang.org/