

# Fall Update

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## nf-core modules

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- A repository of reusable processes for common software
- Nextflow DSL 2 Released in v20.07.1

```
$ nf-core modules install . fastqc
```



```
nf-core/tools version 1.10.2
```

```
INFO      Installing fastqc
```

```
INFO      Downloaded 3 files to ./modules/nf-core/software/fastqc
```

```
#!/usr/bin/env nextflow
```

```
nextflow.enable.dsl = 2
```

```
include { FASTQC } from './modules/nf-core/software/fastqc'
```

```
workflow {
```

```
    FASTQC(SRA_DOWNLOAD.out.raw_reads)
```

```
}
```

# nf-core rnaseq

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nf-core groseq rewrite

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# Europa

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