

# Bioconductor

<https://www.bioconductor.org/>

Svaki korisnik lokalno istalira R pakete.

Upute za instalaciju paketa:

Pokrenite R

```
R
```

Instalacije glavnog paketa:

```
Sys.setenv(R_INSTALL_STAGED = FALSE)
if (!requireNamespace("BiocManager", quietly = TRUE))
  install.packages("BiocManager")
BiocManager::install()
```

Instalaciju dodatnih bioconductor paketa (npr. GenomicFeatures):

```
Sys.setenv(R_INSTALL_STAGED = FALSE)
BiocManager::install("GenomicFeatures")
```

Pokretanje aplikacije:

**bioconductor.sge**

```
#!/bin/bash
#$ -N test
#$ -cwd

Rscript imeSkripte.R
```

```
qsub bioconductor.sge
```