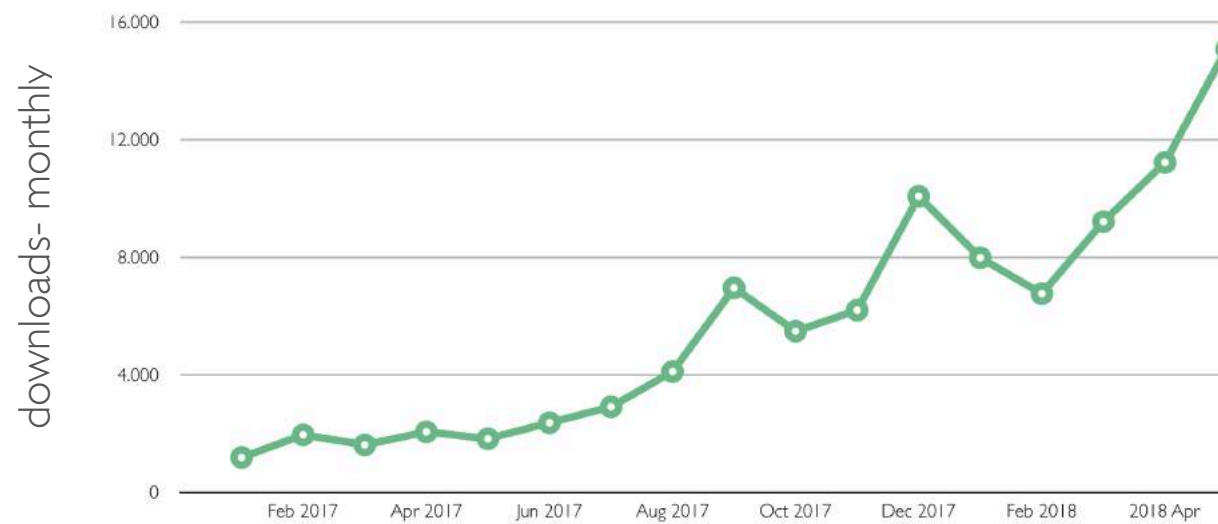




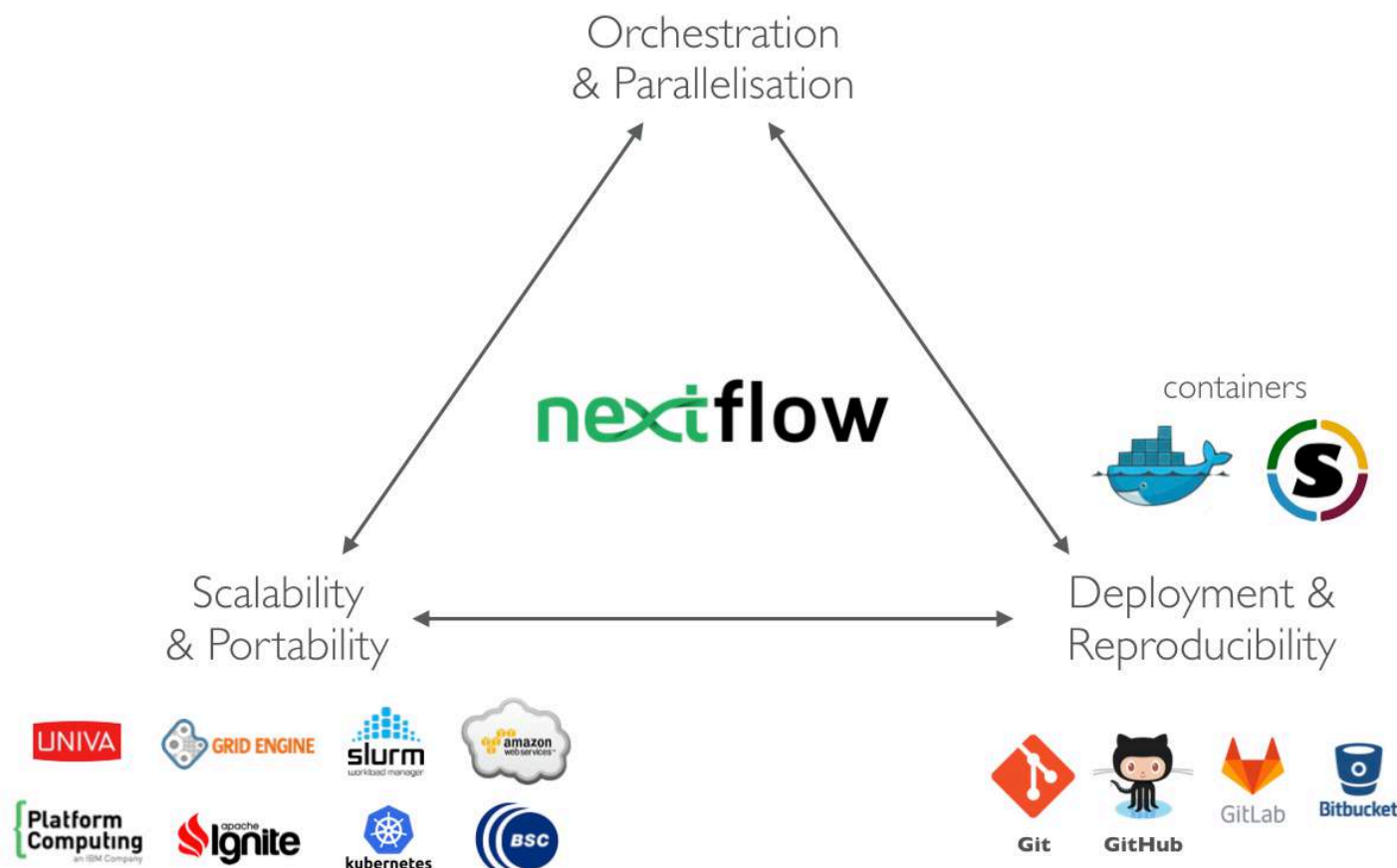
 seqeralabs

Monthly downloads of Nextflow (previous 18 months)

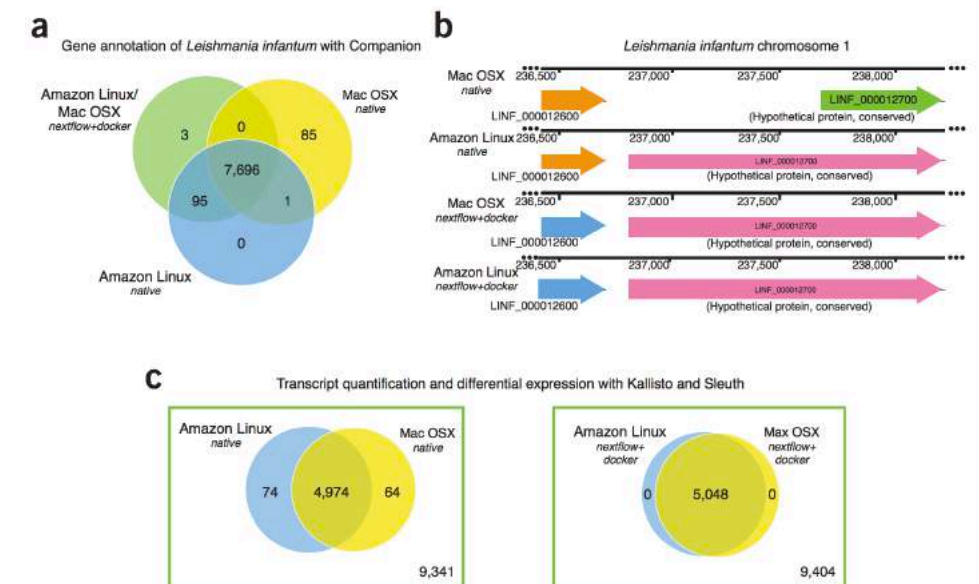


nextflow

<https://github.com/nextflow-io/nextflow>



Nextflow enables reproducible computational workflows



Di Tommaso et. al., *Nextflow enables reproducible computational workflows. Nature Biotechnology* (2017)

Pharma

Clinics

Biotech

Facilities



Our users are also **enterprises** looking for **professional solutions**



Weill Cornell Medical College

UNIVERSITEIT VAN PRETORIA
UNIVERSITY OF PRETORIA
YUNIBESITHI YA PRETORIA





our **mission**

oss project **sustainability**

the 3 c's

contributors

collaborations

community

first-class support

inspirational training events

innovative solutions



First-class support

- Maintenance releases
- Bug fixes (up to 2 years from stable)
- Support Incidents
- Private Issue tracking
- Private support channel
- Phone/Video support
- Application code review
- SLA: same business day
- Enterprise extensions package



Training

- Language syntax & datastructures
- Dataflow variables & channel semantics
- Nextflow operators
- Nextflow processes
- Basic pipeline implementation
- Pipeline sharing & best practices



Training

- Nextflow configuration & profiles
- Managing dependencies with containers
- Executors & deployment scenarios
- Error recovery & strategies
- Workflow / runtime metadata and events
- Common implementation patterns

solutions for **sensitive** data

solutions for **critical** applications

solutions for **massive scale** users



the **vision**

~~YAP (yet another platform)~~

enabling technology

the future is open source

 **seqeralabs**





Paolo Di Tommaso



Evan Floden



Powering the next generation
of big data analysis applications

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