

# WESkit

GA4GH-compliant workflow execution

Sven Twardziok

Online training

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# GA4GH - WES

# Workflows & Cloud Genomics in healthcare

**Expected 60 millions of human genomes sequenced within global healthcare systems by 2025 - this will transform whole healthcare systems<sup>1</sup>**



National sequencing initiatives<sup>2</sup>

## Challenges<sup>2</sup>:

- Technical:** infrastructure and bioinformatics capacity
- Research:** data integration and interpretation
- Social:** public acceptability, and ethical and legislative issues

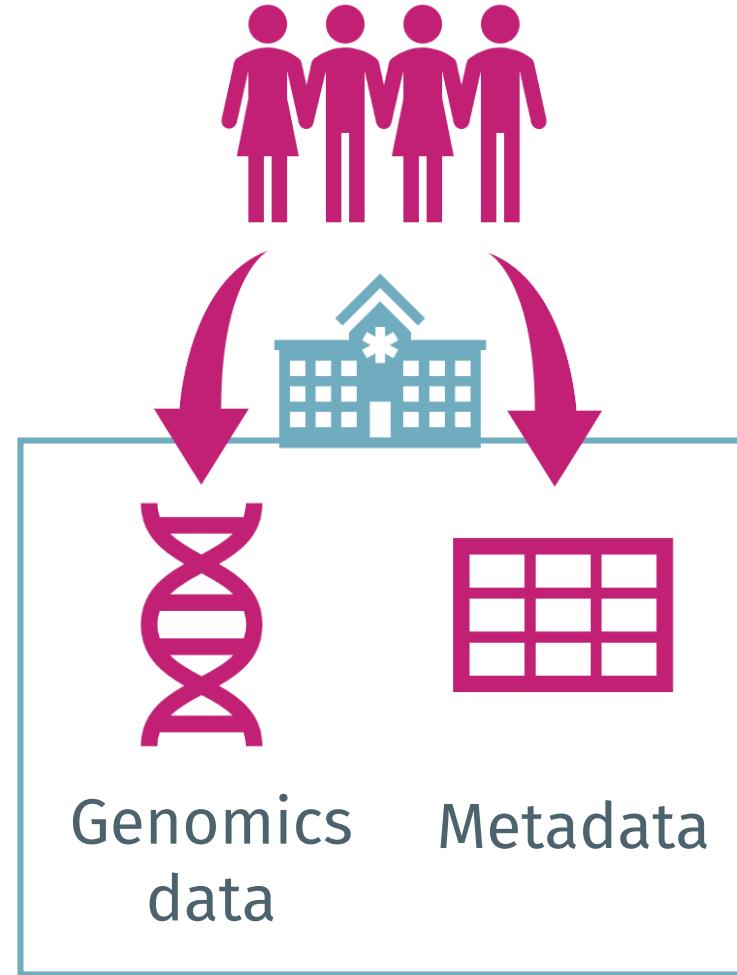
<sup>1</sup> Genomics in healthcare: GA4GH looks to 2022 (2017) | bioRxiv.,

<sup>2</sup> Integrating Genomics into Healthcare: A Global Responsibility (2019) | AJHG

# Genomics data in healthcare

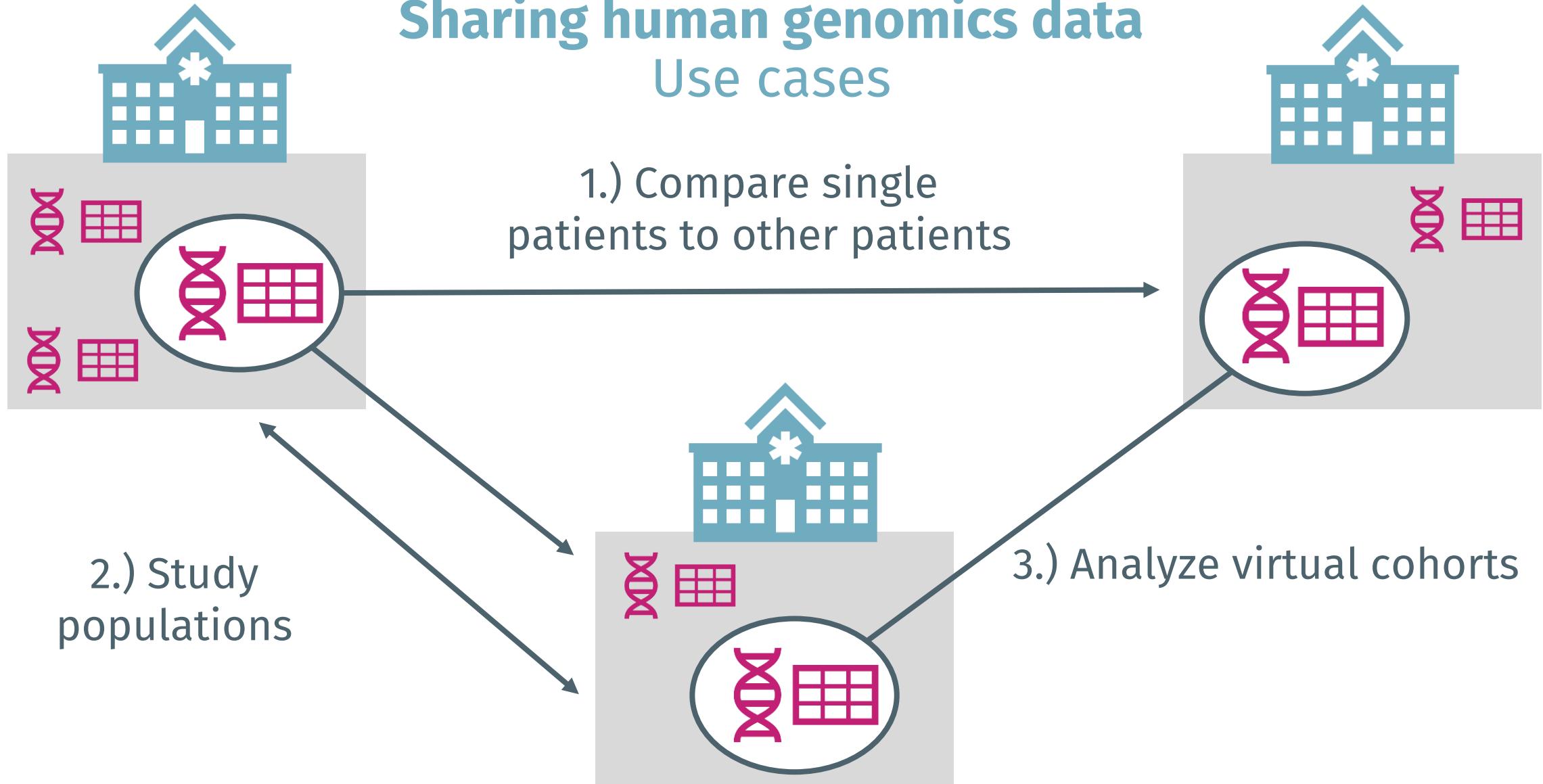
## Features

- **Big:** hundreds of gigabytes per patient
- **Sensitive:** genetic data is specifically protected by GDPR
- **Structured:** linked with many metadata such as clinical data
- **Decentralized:** located in sequencing facilities, (university) hospitals, private laboratories



# Sharing human genomics data

## Use cases



# Science Cloud

## Sending analysis workflows to data

### Requirements for standardization

- Organize a decentralized cloud infrastructure including a decentralized user access management.
- Send and process workflows in the cloud.
- Provide access to data and data discovery.
- Define good data/workflow management.

# Global Alliance for Genomics & Health (GA4GH)



Global Alliance  
for Genomics & Health  
Collaborate. Innovate. Accelerate.

Enabling genomic data sharing  
for the benefit of human health.



## Mission

The Global Alliance for Genomics and Health aims to accelerate progress in genomic research and human health by cultivating a **common framework of standards** and harmonized approaches for effective and responsible genomic and health-related data sharing.



GA4GH **Driver Projects** are real-world genomic data initiatives that provide input on the standards most needed for the international genomics community to share data. Driver Projects put finished standards and frameworks to immediate use.

## GA4GH Driver Projects

### National Initiatives



### Pan-African



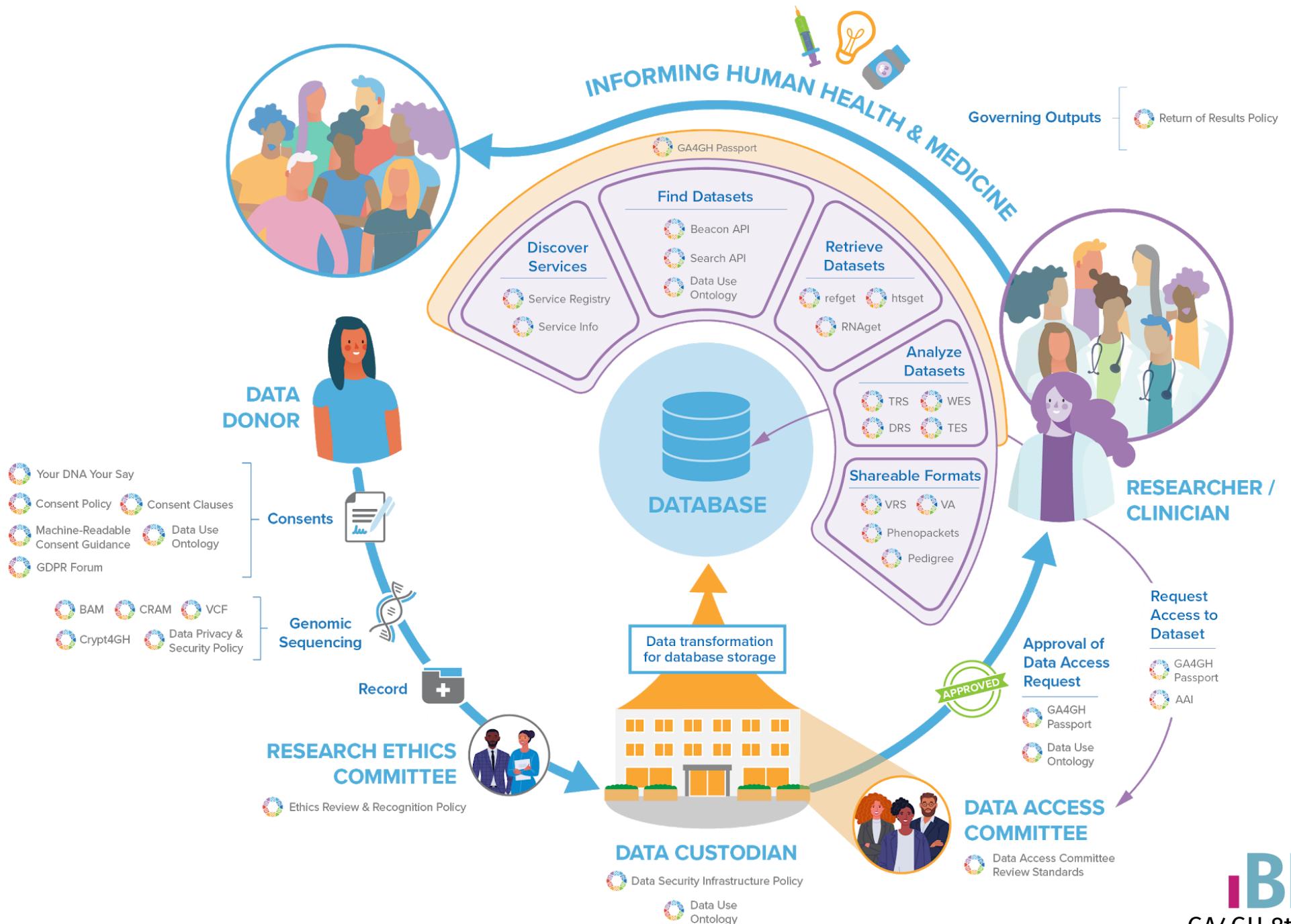
### Pan-European



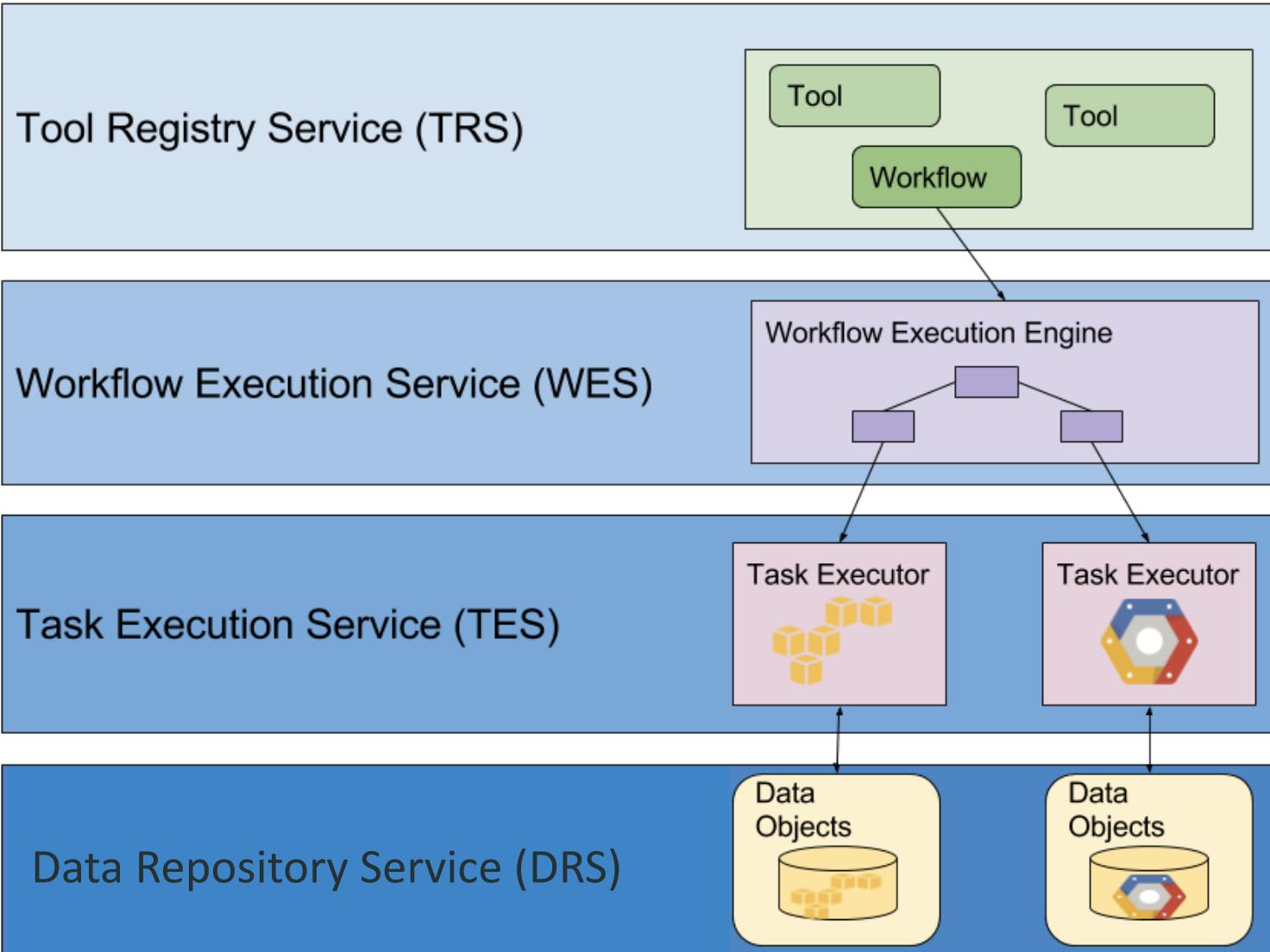
### International



<https://www.ga4gh.org/wp-content/uploads/GA4GH-onepager-v6.pdf>



# Cloud Work Stream APIs



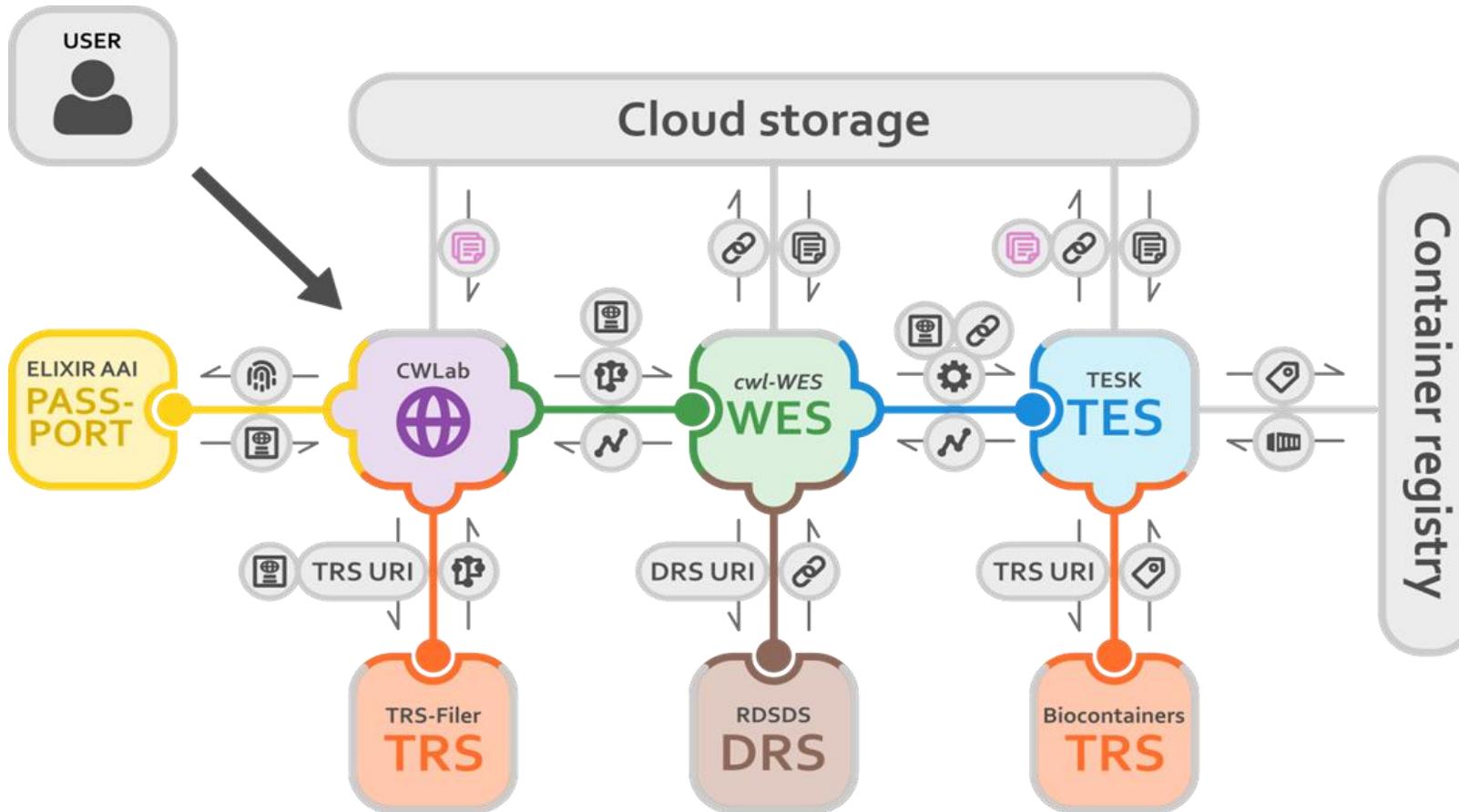
**Sharing Tools  
and Workflows**

**Executing  
Workflows**

**Executing  
Individual Tasks**

**Accessing Data**

# ELIXIR Cloud & AAI Integration of services



## Legend

- } GA4GH API client
- { GA4GH API server
- Login credentials (password / 2FA)
- Bearer token with GA4GH Passport
- Workflow
- Data access link
- Input data
- Task
- Container image name/tag
- Container image
- Logs including workflow/task state
- Output data

2020-09-30 ELIXIR Cloud & AAI @ 8th GA4GH Plenary

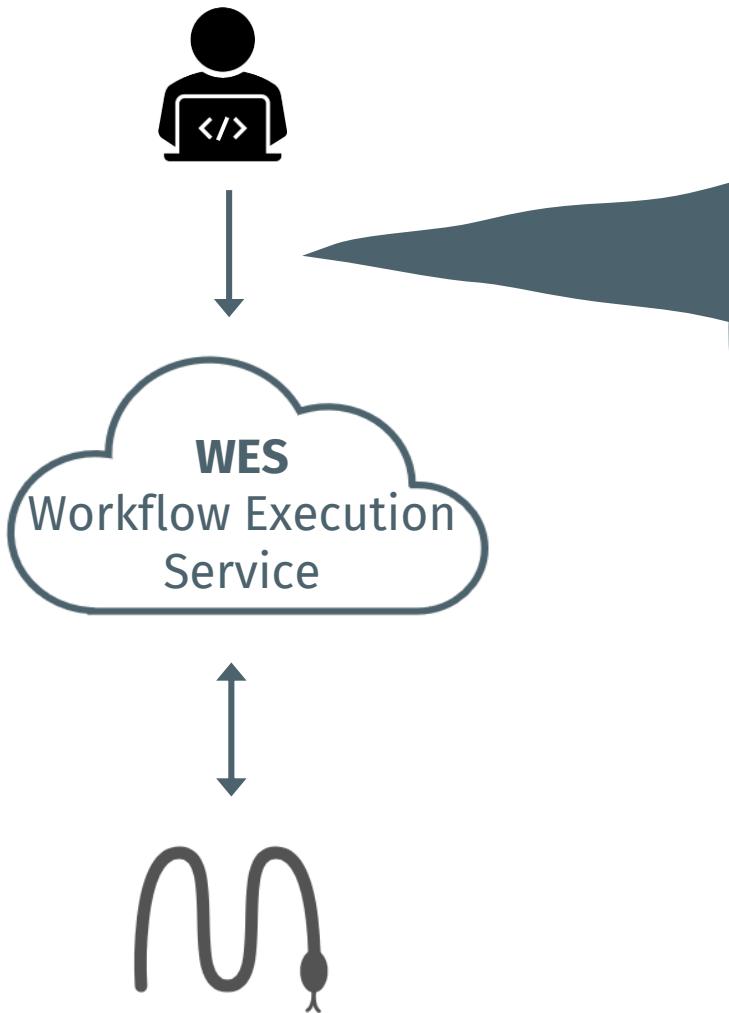
# GA4GH Workflow Execution Service (WES) Endpoints

Method	Path	Description
GET	/service-info	Get information about Workflow Execution Service.
GET	/runs	List the workflow runs.
POST	/runs	Run a workflow.
GET	/runs/{run_id}	Get detailed info about a workflow run.
GET	/runs/{run_id}/status	Get quick status info about a workflow run.
POST	/runs/{run_id}/cancel	Cancel a running workflow.

<https://github.com/ga4gh/workflow-execution-service-schemas/>

# Workflow Execution Service (WES)

## POST /run



```
requests.post(  
    url = WES_SERVER + "/ga4gh/wes/v1/runs",  
    header = {"Authorization": "Bearer " + ACCESS_TOKEN},  
    data = {  
        "workflow_params": CONFIG_JSON,  
        "workflow_type": "Snakemake",  
        "workflow_type_version": "5",  
        "workflow_url": "/workflows/wf1/Snakefile"  
    }  
}  
{"run_id": "id_123"}
```

# Workflow Execution Service (WES)

## GET /runs/{run\_id}



```
requests.get(url = WES_SERVER + "/ga4gh/wes/v1/runs/id_123")
{
    "run_id": "id_123",
    "request": {...},
    "state": "COMPLETE",
    "run_log":{
        "name": "my_workflow",
        "cmd": "snakemake ...",
        "start_time": "2021-01-12T12:00:00Z",
        "end_time": "2021-01-12T12:05:00Z",
        "stdout": "/PATH/TO/STDOUT",
        "stderr": "/PATH/TO/STDERR",
        "exit_code": 0
    },
    "task_logs": [...],
    "outputs": {...}
}
```

# WESkit



GA4GH WES implementation

<https://gitlab.com/one-touch-pipeline/weskit>

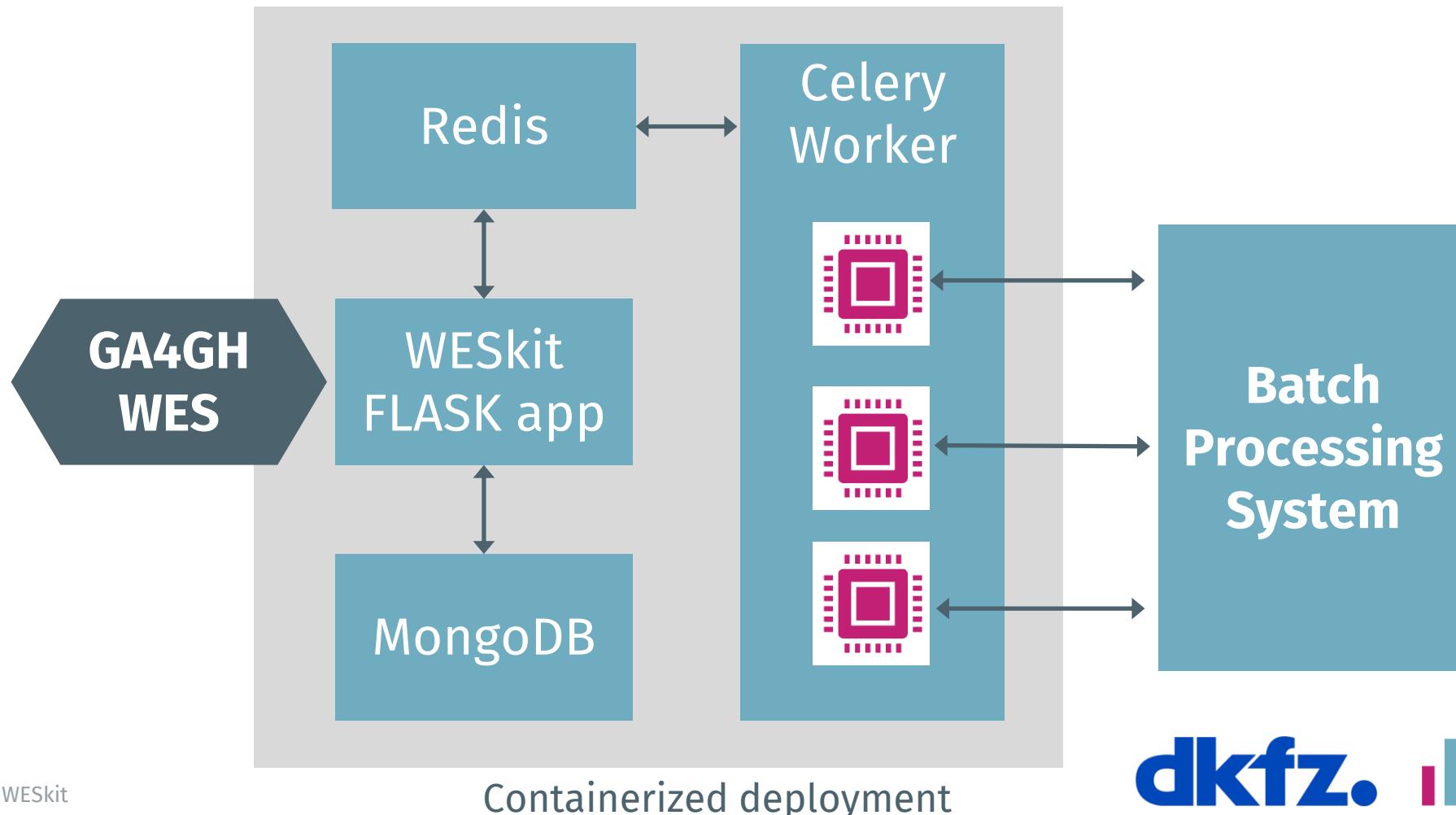


## Features

- GA4GH WES implementation supporting Snakemake and Nextflow
- Developed for stability and high data throughput at BIH and DKFZ
- HPC and cloud deployment supported
- OAuth2.0 support
- Developed in cooperation with ELIXIR compute platform



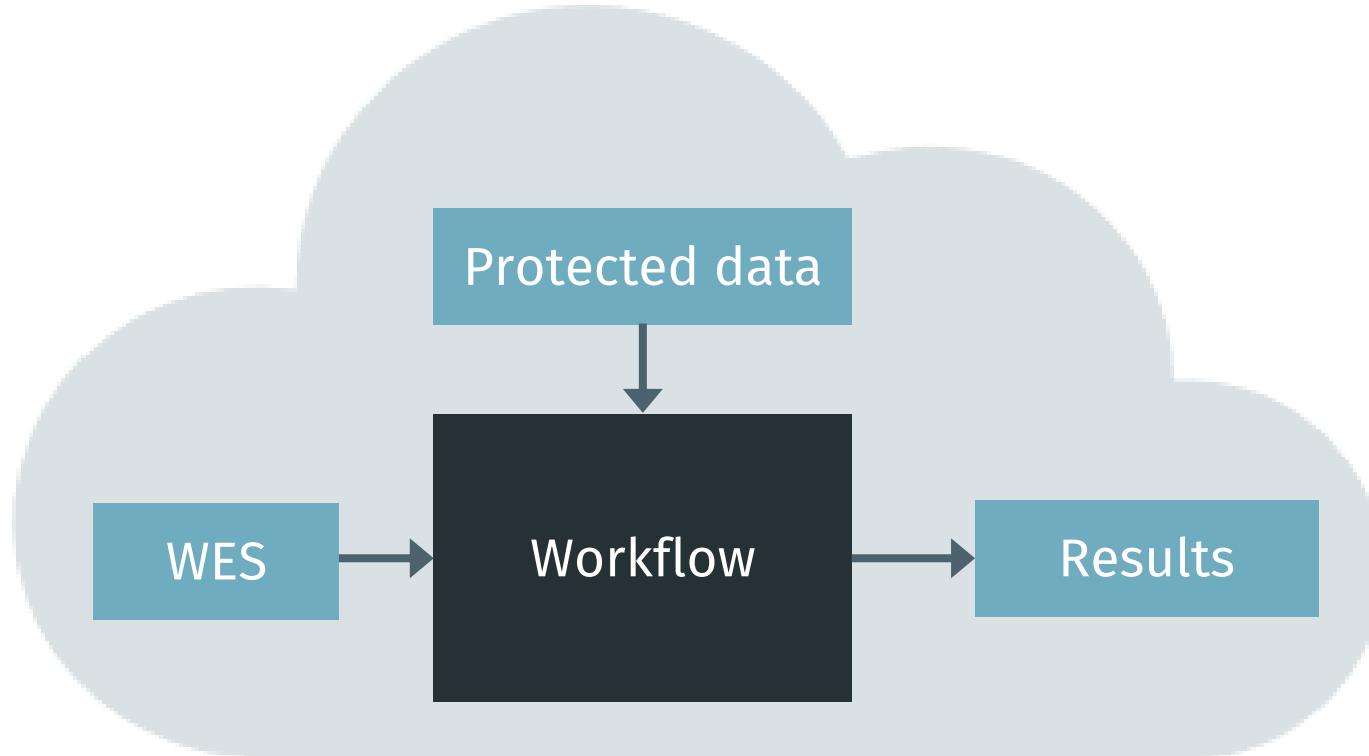
GA4GH WES implementation  
<https://gitlab.com/one-touch-pipeline/weskit>



# Demonstration

# Demo: deploying WESkit

## WESkit platform



<https://gitlab.com/one-touch-pipeline/weskit/documentation>

# Exercise

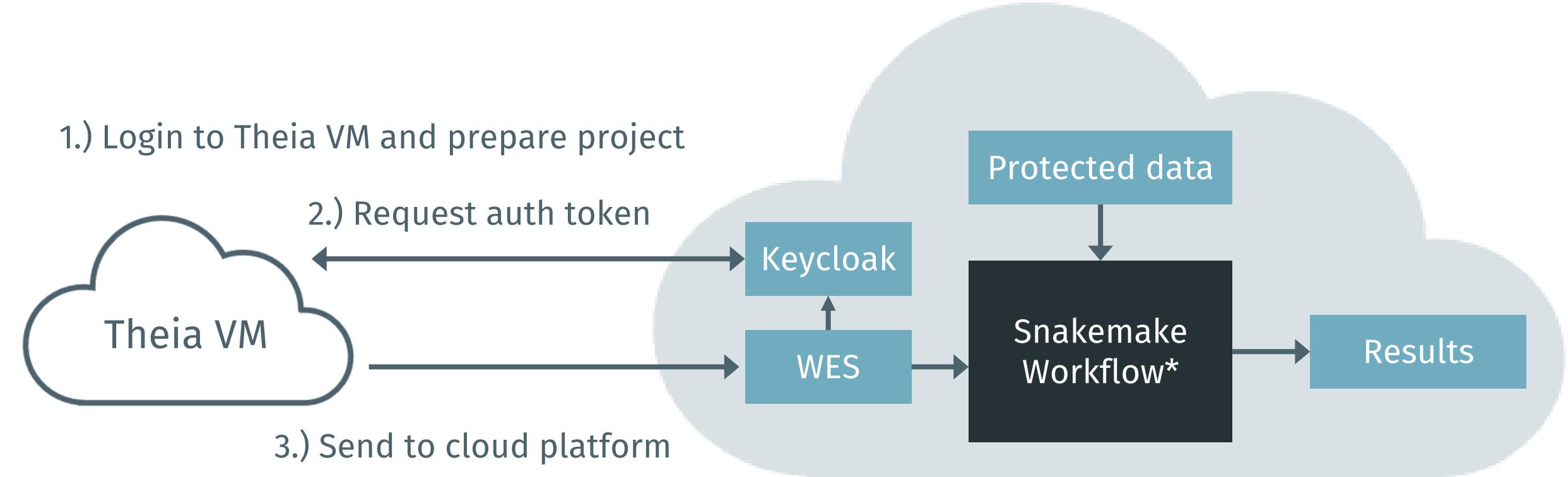
# Task: send workflow to cloud

<https://gitlab.com/twardzso/weskit-elixir-workflow-workshop-2021>

- 1.) Login to Theia VM and prepare project

- 2.) Request auth token

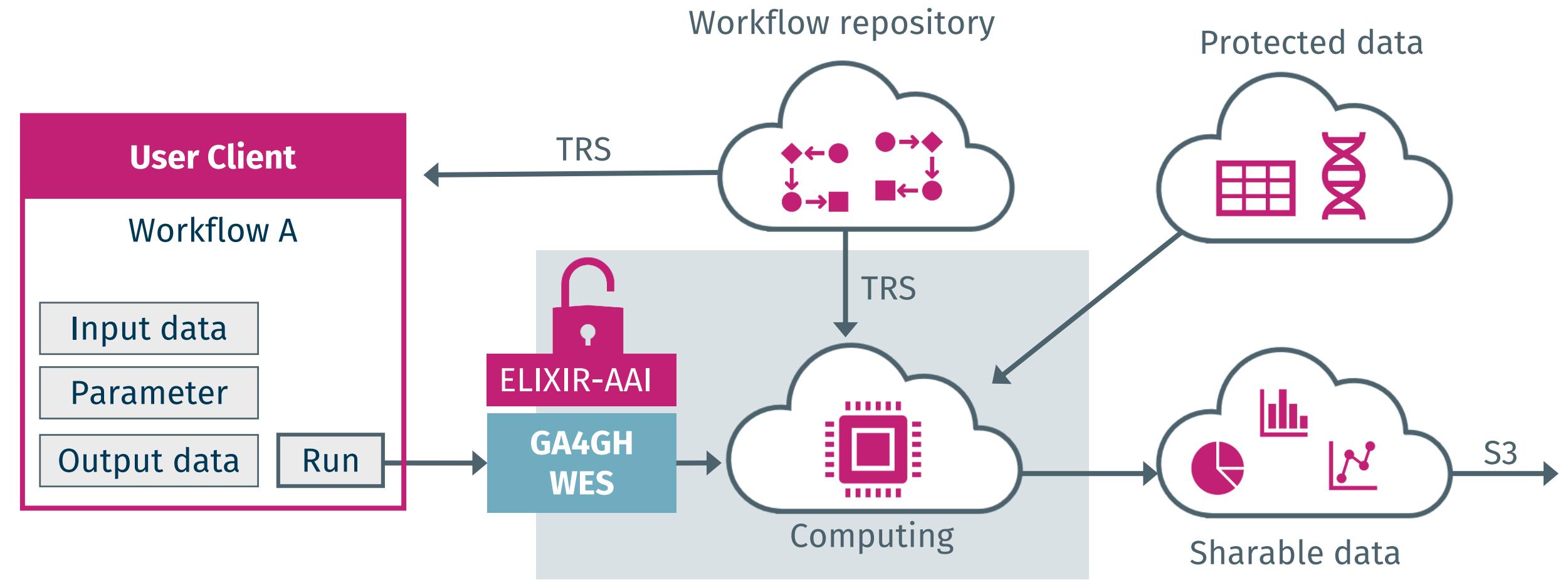
- 3.) Send to cloud platform



\*<https://snakemake.readthedocs.io/en/stable/tutorial/tutorial.html>

# WESkit use-cases

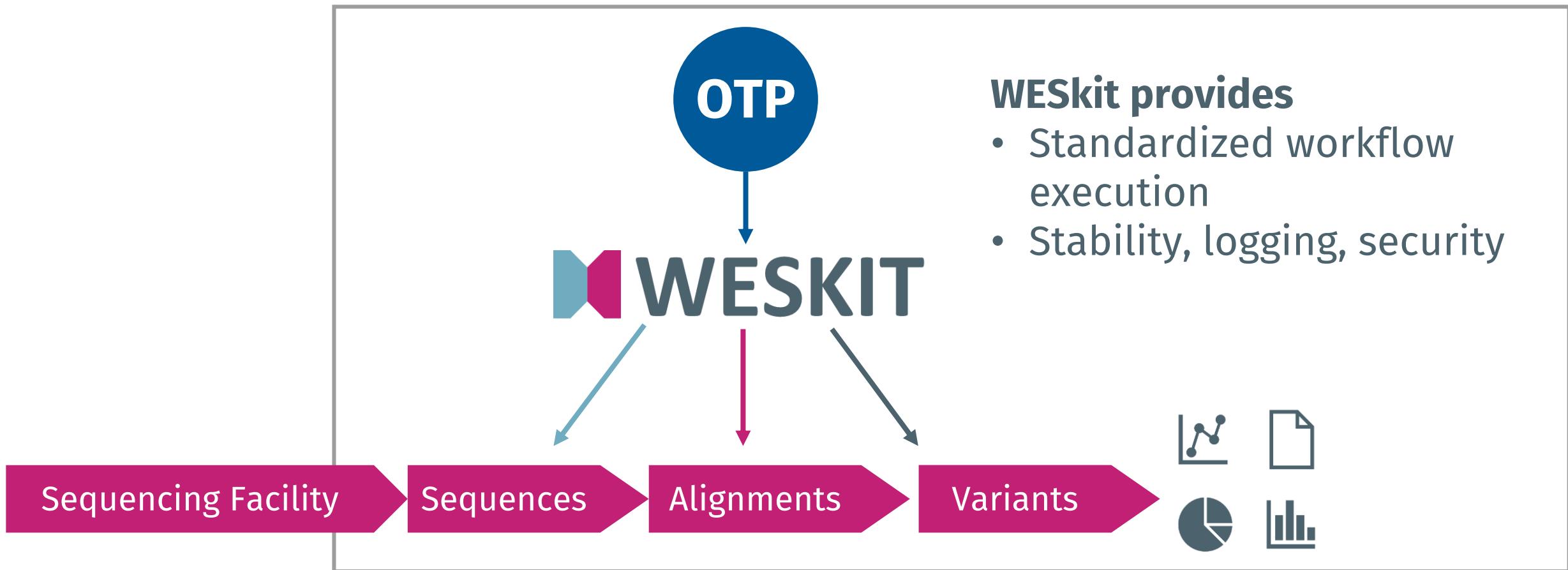
# WESkit Cloud Use Case



 **WESKIT**

# WESkit Automation Use Case

<https://gitlab.com/one-touch-pipeline>



# Summary

- GA4GH-WES: specification for sending workflows to execution environment
- WESkit: implementation of GA4GH-WES for Snakemake and Nextflow
- Cloud use case: build platform to provide access to sensitive data
- Automation use case: provide standardized interface for repetitive execution of workflows

# Thank you

## Questions:

sven.twardziok@bih-charite.de

[www.bihealth.org](http://www.bihealth.org)

**Charité - BIH**

**Roland Eils**

Andreas Kling

Valentin Schneider-  
Lunitz

Florian Kärcher

Harald Wagener

Jürgen Eils

Martin Braun

Matthias Bieg

Naveed Ishaque

Philipp Strubel

**HD-Hub**

Chris Lawerenz

Elisabeth Bürger

Ivo Buchhalter

Kersten Breuer

Pavlo Lutsik

Philip Kensche

**ELIXIR Cloud & AAI**

Alexander Kanitz

Alvaro Gonzalez

Ania Niewielska

Jonathan Tedds

Justin Clark-Casey

Kostis Zagganas

Lukas Hehtmanek

Marius Diekmann



Aus Forschung wird Gesundheit

# Questions?



Aus Forschung wird Gesundheit