



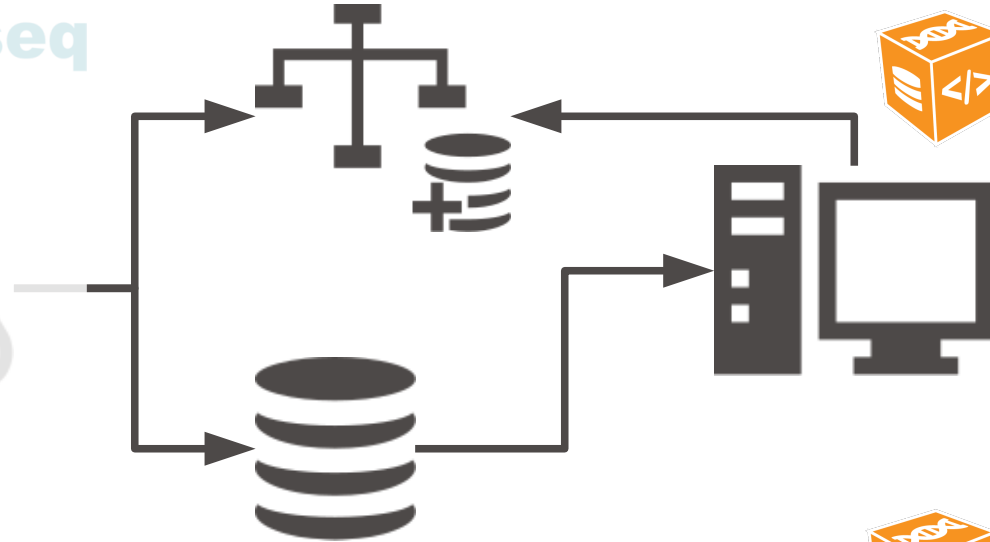
Data Storage Platforms



Korbinian Bösl
Data management coordinator
ELIXIR Norway/Digital Life Norway
10th December 2020

NeLS

Norwegian e-Infrastructure for Life Sciences



sensitive
data



NeLS



Feide ola@uib.no



My Projects



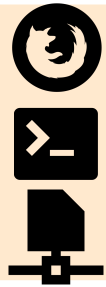
NeLS



Norwegian e-Infrastructure for Life Sciences

Data curation

NeLS



 **Feide** ola@uib.no
My Projects

NeLS

Norwegian e-Infrastructure for Life Sciences

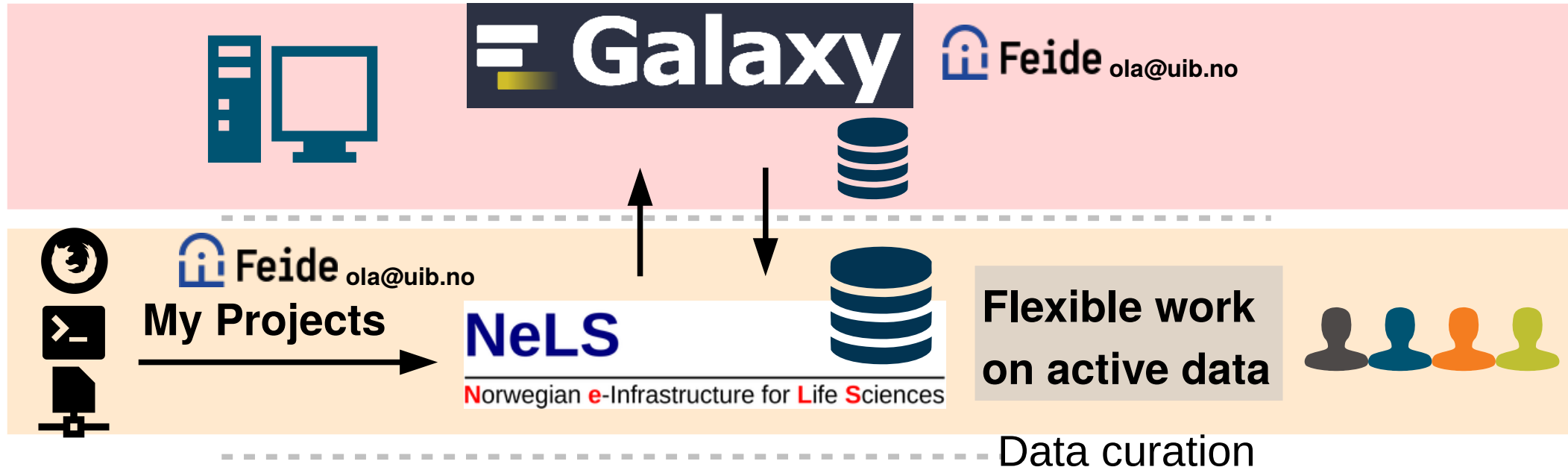


**Flexible work
on active data**

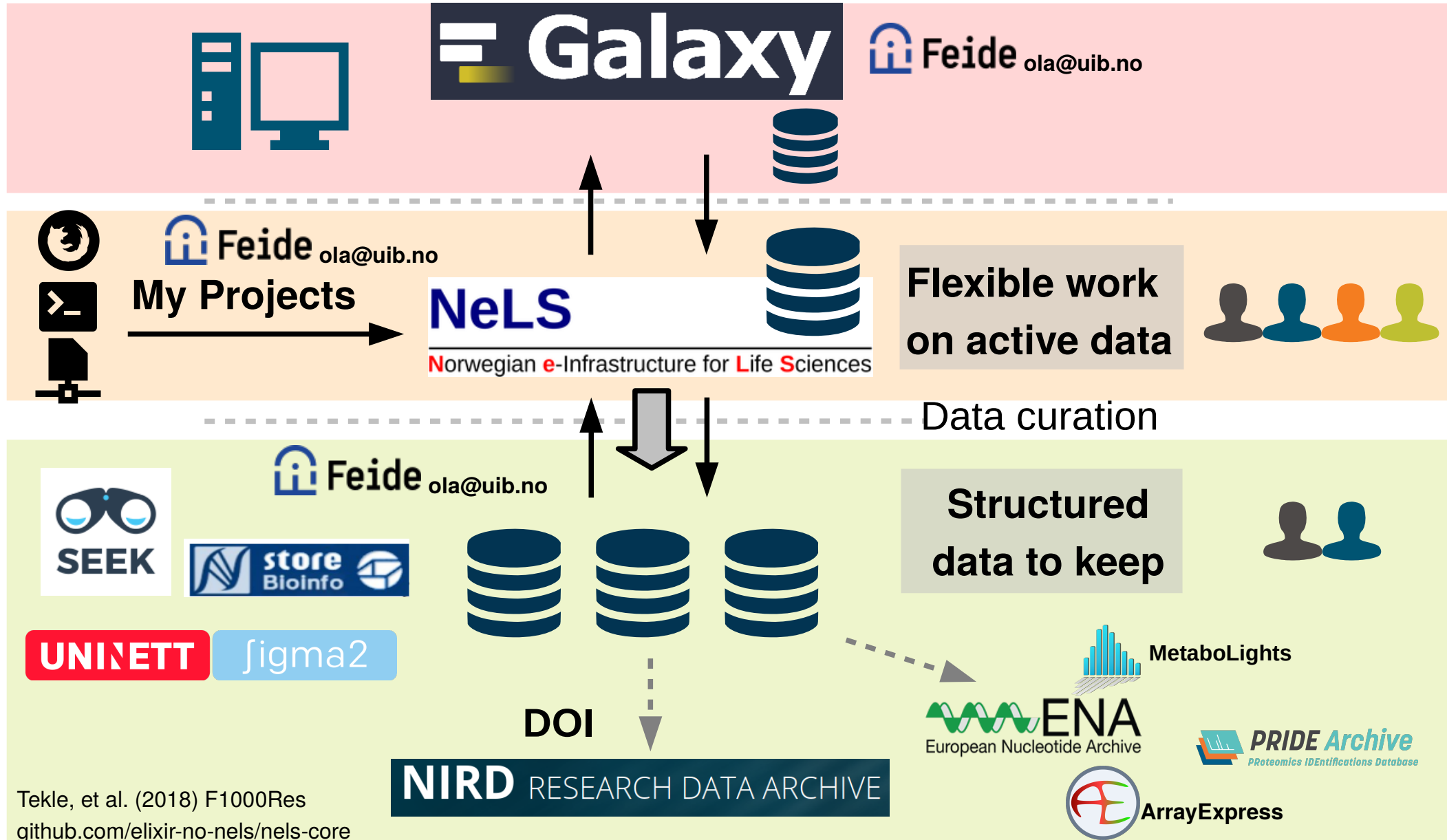


Data curation

NeLS



NeLS



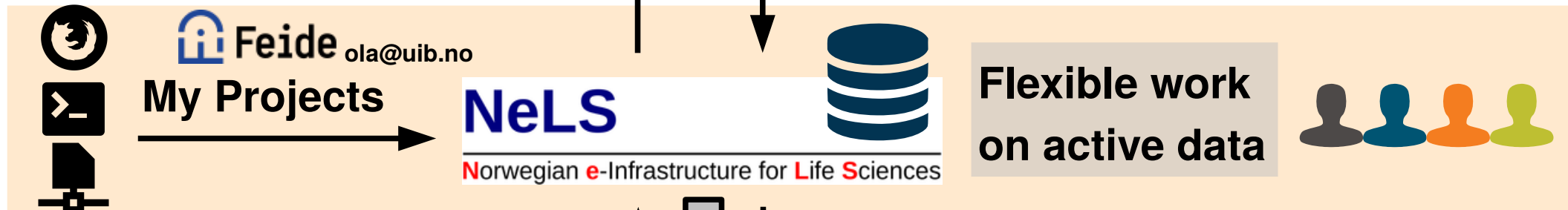
Tekle, et al. (2018) F1000Res
github.com/elixir-no-nels/nels-core

NeLS

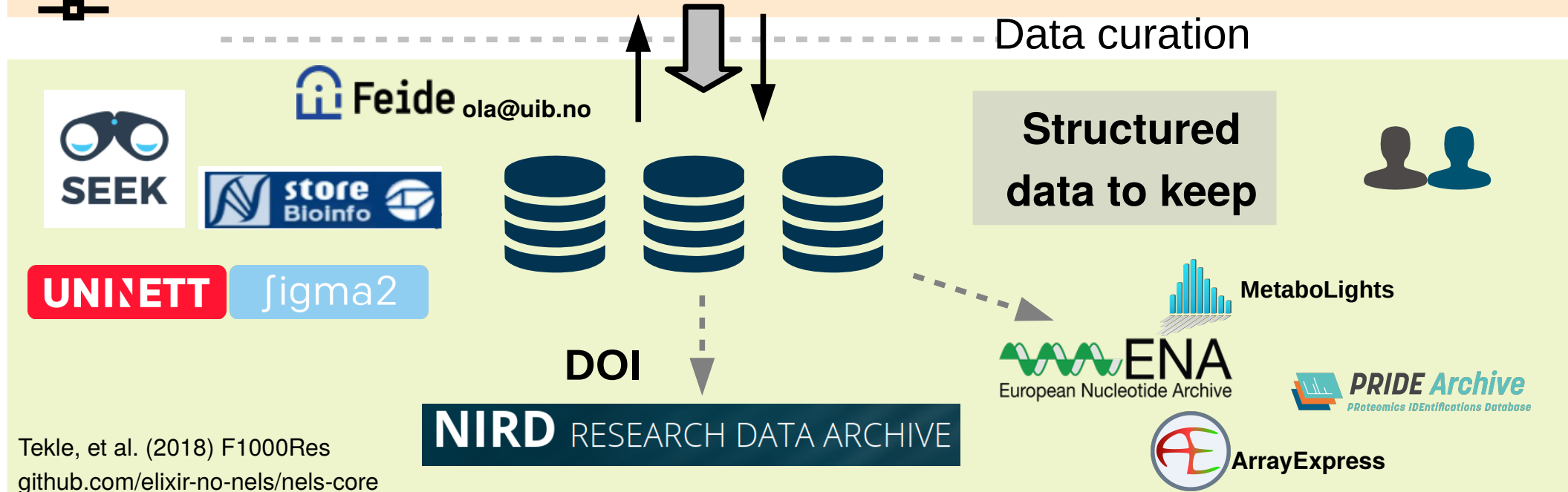
Days - weeks



 **nor seq**
Months



Years



Decades

Tekle, et al. (2018) F1000Res
github.com/elixir-no-nels/nels-core

<https://nels.bioinfo.no/>

NeLS - Accessibility

small datasets



ola@uib.no

NeLS - Accessibility

small datasets



ola@uib.no

larger datasets



ssh-key



git



sFTP

Data Analysis in NeLS

Name	Annotation	Owner	Community Rating	Community Tags	Last Updated↓
16S Workflow with Mothur program ▾		kjetil-klepper	★★★★★		Oct 17, 2020
NGS Pipeline for Paired End Reads (R1 and R2) ▾		kjetil-klepper	★★★★★		Oct 17, 2020
miRNA differential expression (miRBase, hg38) ▾		kjetil-klepper	★★★★★	nels	Oct 15, 2020
miRNA differential expression (MirGeneDB, hg38) ▾		kjetil-klepper	★★★★★	nels	Oct 14, 2020
COVID-19: PE Variation ▾		kjetil-klepper	★★★★★	nels	Oct 13, 2020
Pre-process COVID-19 PE collections ▾		kjetil-klepper	★★★★★	nels	Oct 13, 2020
Pre-process COVID-19 PE single sample ▾		kjetil-klepper	★★★★★	nels	Oct 13, 2020

Data Analysis in NeLS

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Pre-process COVID-19 PE single sample ▾		kjetil-klepper	★★★★★	nels	Oct 13, 2020

Courses Q1 2021

NeLS - Getting Access

NeLS

Norwegian e-Infrastructure for Life Sciences

Typically > 1TB data

Free < 10TB & <15 mill. NOK

Allocation through Helpdesk: contact@bioinfo.no

Direct upload through Nourseq facilities

<https://nels.bioinfo.no/pages/user-terms.xhtml>



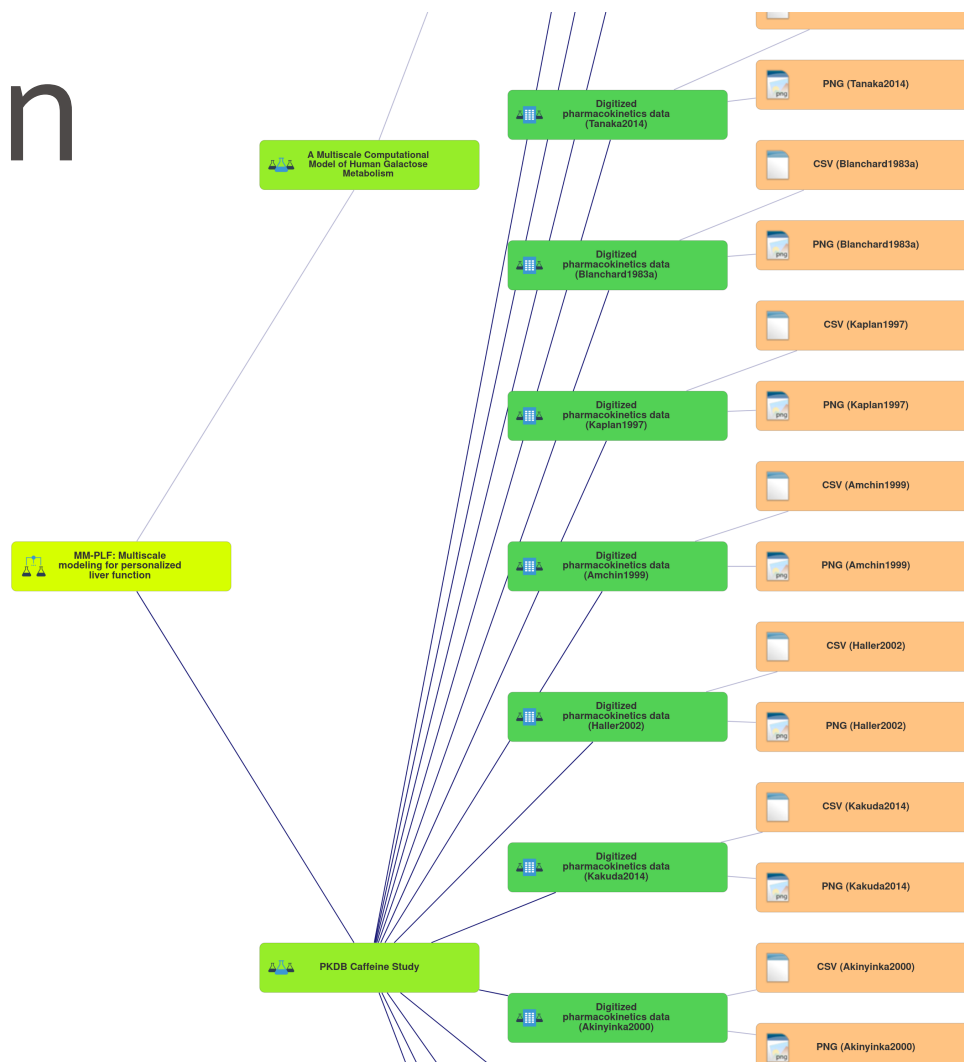
SEEK





SEEK

Investigation Study Assay





RightField



File Edit Sheet Help

	A	B	C	D	E	F
1	# This is an excel template...					
2	# Use this template for ...					
3	# Click the Metadata Ex...					
4	# Field names (in blu...					
5	# CLICK HERE for the F...					
6						
7	SERIES					
8	# This section describes ...					
9						
10	title					
11	summary					
12	summary					
13	overall design					
14	contributor					
15	contributor (SEEK ID)					
16	SEEK Project	Project				
17	Experiment Class (a...	transcriptomics				
18	Experiment Design t...	ExperimentDesignT...				
19	Technology type	microarray				
20	quality control type	QualityControlDesc...				
21						
22	SAMPLES					
23	# The Sample name...					
24	# CLICK HERE to find t...					
25						
26	Sample name	title	CEL file	source name	organism	characteristics:...
27	SAMPLE 1				organism	
28	SAMPLE 2				organism	
29	SAMPLE 3				organism	
30	SAMPLE 4				organism	
31	SAMPLE 5				organism	
32	SAMPLE 6				organism	
33	SAMPLE 7				organism	
34	SAMPLE 8				organism	
35	SAMPLE 9				organism	
36	SAMPLE X				organism	
37						
38						
39	PROTOCOLS					
40	# This section includes pr...					
41	# Protocols which are ap...					
42						
43	growth protocol					
44	treatment protocol					
45	extract protocol					
46	label protocol					

Selected cells: B17:B17

ONTOLOGY HIERARCHIES

MGEDOntology.owl x

JERMOntology x

- ExperimentClassType
 - fluxomics
 - genomics
 - interactomics
 - metabolomics
 - proteomics
 - reactomics
 - single_cell
 - transcriptomics
- InformaticsAnalysisType
 - informaticsAnalysisType
- ModelAnalysisType
 - ModelAnalysisType
- CultureGrowth
 - CultureGrowth
- FactorsStudied
 - concentration
 - expression

TYPE OF ALLOWED VALUES

☐ Free text

☐ Direct subclasses

☒ Subclasses

☐ Instances

☐ Direct instances

ALLOWED VALUES

- Comparative genomic hybridization
- RNAi
- gene expression profiling
- methylation profiling
- microRNA profiling
- tiling path

Apply

Metadata Template Matrix Template Metadata Example Matrix Example 1 Matrix Example 2

NeLS

Norwegian e-Infrastructure for Life Sciences



Integration



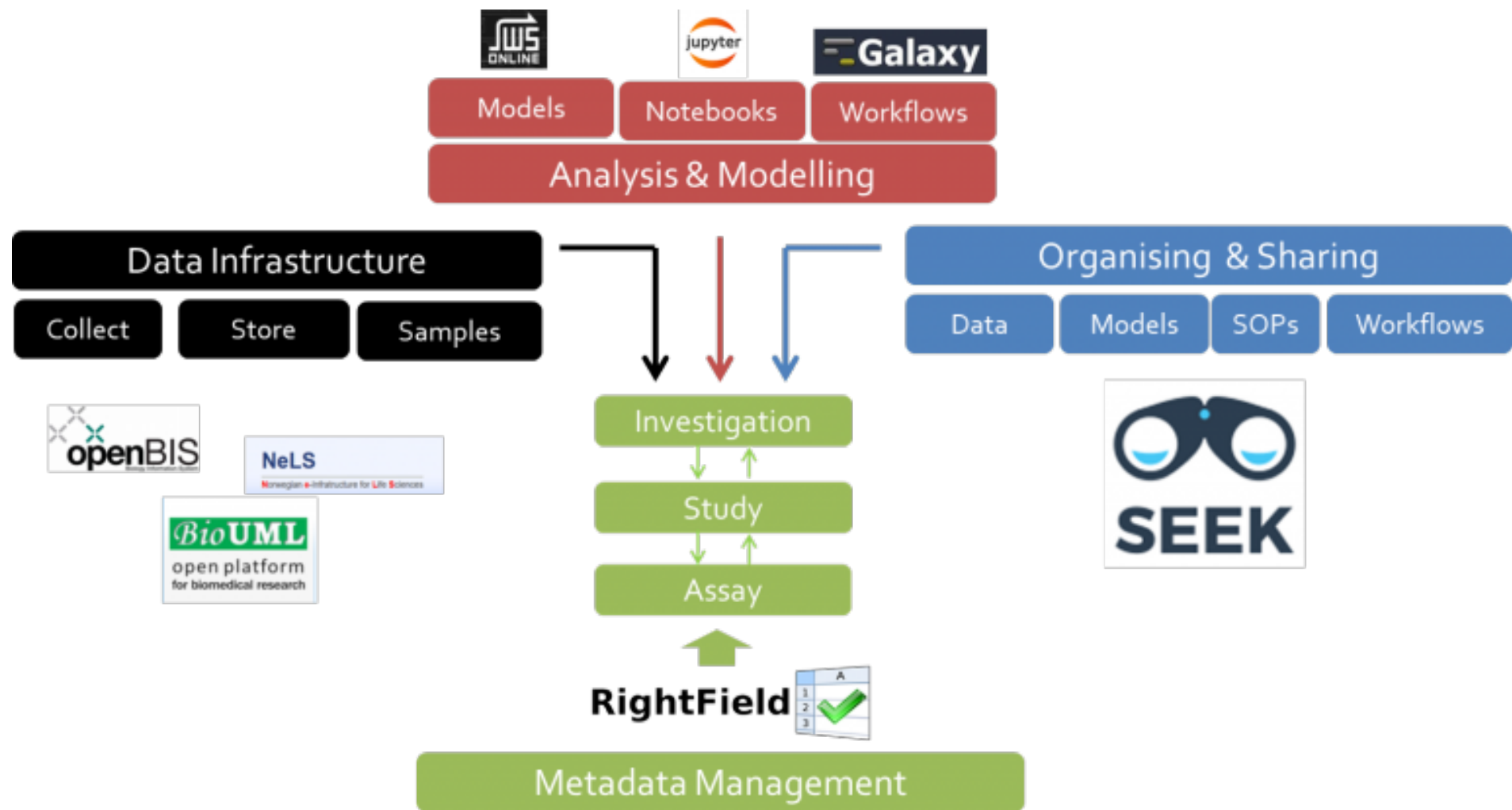
Reference data in NeLS in SEEK



Access data in NeLS through SEEK



Save Metadata to SEEK through NeLS



National HPC infrastructure

UNINETT

sigma2



Advanced Storage Needs

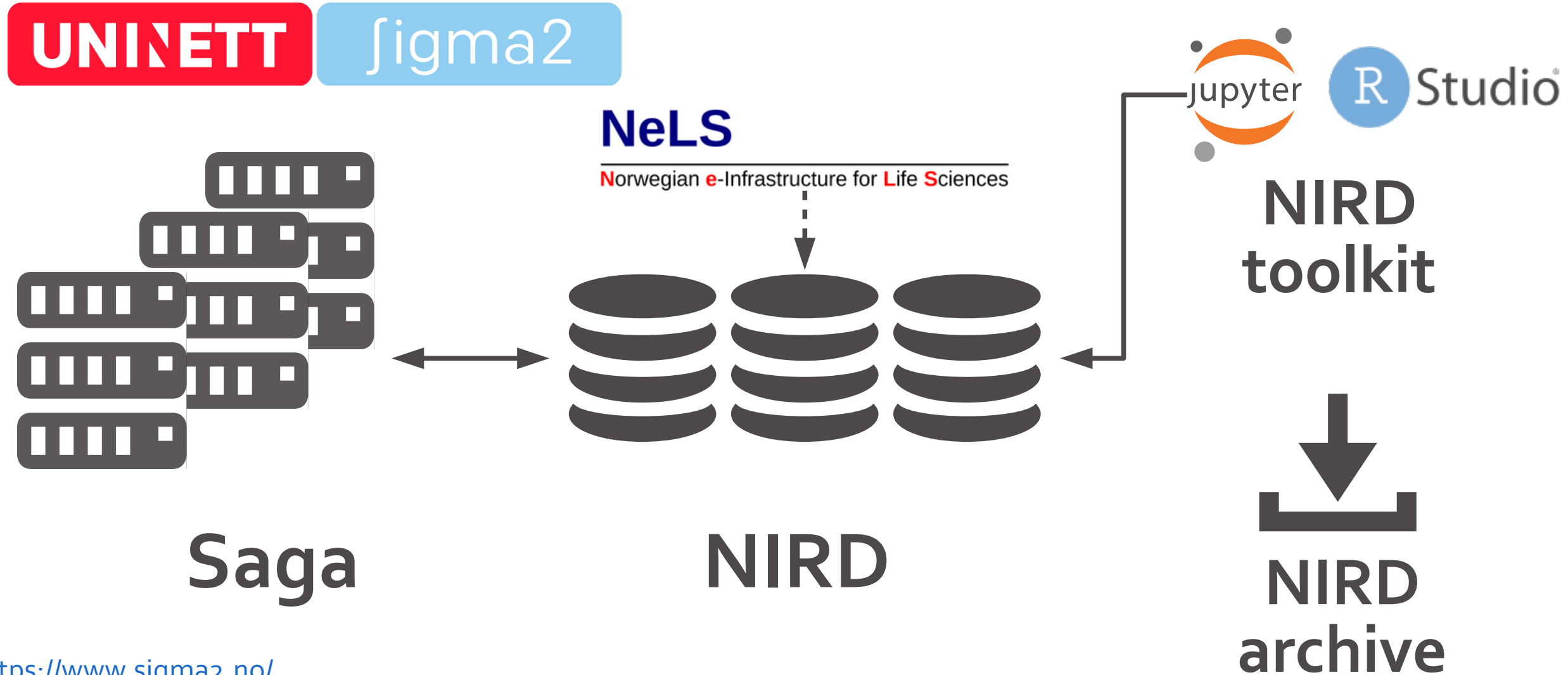


Advanced Computing Resources



Free Analysis Toolkit (Rstudio, Jupyternotebook)

National HPC infrastructure



HPC infrastructure

UNINETT

sigma2

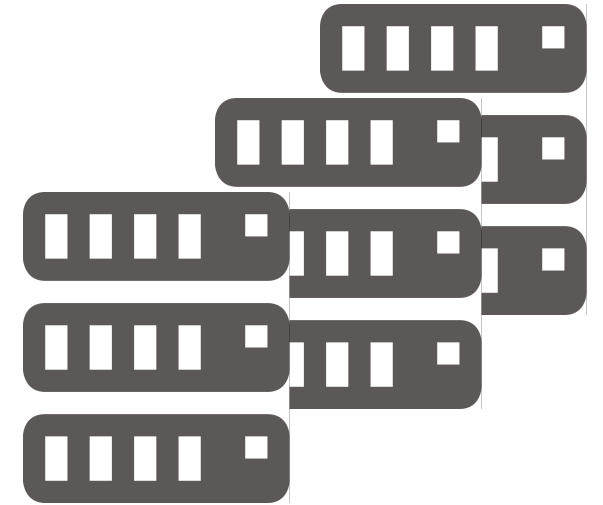
Saga

244 nodes – 9824 cores – 32 GPUs

Total 75TB RAM – 186-3066GB/node

89TB + 60TB NVMe+SSD

1PB file system



Storage infrastructure - NIRD

UNINETT

sigma2

Geo replicated

Nightly snapshots

12 PB (→ 36 PB next 3 yrs)



Sigma2 - Getting Access

UNINETT

sigma2

Bi-annual calls (January & July) – Reapplication (!)

User contribution model (projects > 15 mill. NOK)

Application for small grants continually