Making Data Fun Again

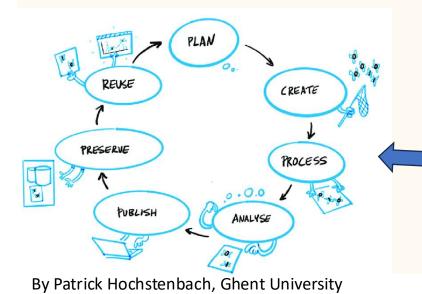
Extending EESSI to improve Research Data Management

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What is Research Data (Management)?

Research data is any data (consumed or produced) in research.



Research Data Management comprises all

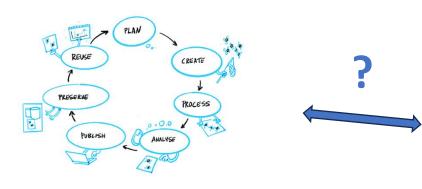
kinds of activities to "organise" research data

with the goal to enable its reuse.



Dilemma

Structure



workflows

DMPs

Creative Chaos (the fun part)





The actual problem with Research Data Management

- not acknowledged in hiring processes and proposal evaluations
- underfunded (varying support from organizations)
- costs time
- → researchers do the minimum to satisfy (funders') requirements

• Plus, RDM is just one issue for researchers using IT...



What in IT do researchers struggle with?

- many IT systems: laptop, cloud, HPC
- use many software packages
- develop code for increasingly heterogeneous architectures

- manage their own virtual data infrastructure
 - low-level (data) management operations to move data between systems *STER*
 - manage storage spaces with different performance and quota



How EESSI helps already and what could be next? many IT systems: laptop, cloud, HPC **EESSI** easy access to many software packages develop code for increasingly heterogeneous architectures manage their own virtual data infrastructure

- low-level (data) management operations to move data between systems
- manage storage spaces with different performance and quota



"Extending" EESSI to make data handling easier



Don't limit creativity

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Avoid (low-level) data management operations



"extend" EESSI

everyone should be already using EESSI EESSI provides a good basis



EESSI in a nutshell

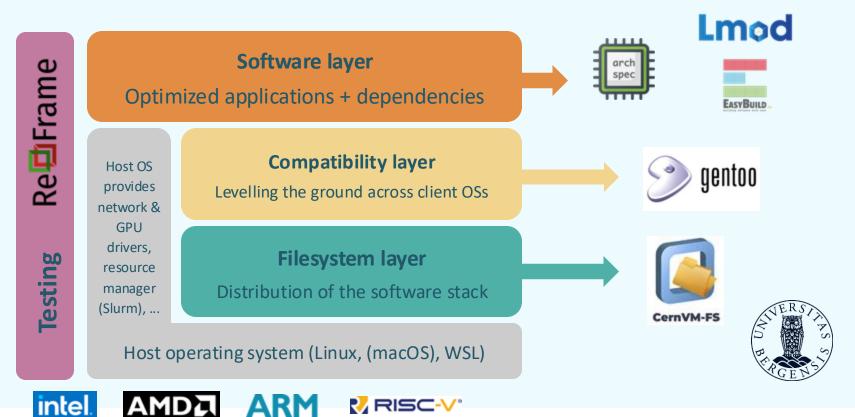
- **On-demand streaming** of **optimized** scientific software installations
- Works on any Linux distribution thanks to EESSI compat layer
- Uniform software stack across various systems: laptop, HPC, cloud, ...
- Community-oriented: let's tackle the challenges we see together!



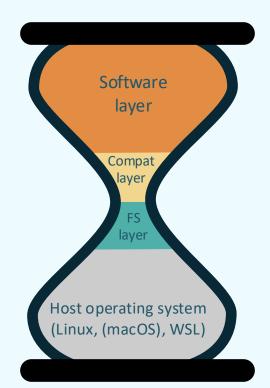




EESSI's layered architecture



How can we "extend" EESSI?



- don't want to rebuild every software installation
- better: modify or augment a core component
- assumption: most data accesses pass through compat layer
- key data access functions:
 - *open*:open, fopen, openat, fopenat, open64, fopen64, freopen
 - read, write
 - close, fclose
- all part of GLIBC: changes there apply to all/most software



How can we change GLIBC functions?

- Directly change GLIBC
 - pros: transparent for users, no change needed
 - cons: is always on whether needed/wanted or not; may have undesired consequences; need to change a shipped installation
- Wrap functions and use \$LD_PRELOAD
 - pros: keep default GLIBC; easily switched on/off by users
 - cons: need to adjust "every" run for full coverage; may create conflicts if set globally
- For prototype: wrap functions and use \$LD_PRELOAD



What could we do with wrapping GLIBC functions?



log information about certain calls (open, exec, ...)

post-process logs to create data flow graphs



enable using remote data "directly"

no manual download of data before processing it



define a virtual data infrastructure

declare which data is needed

runtime ensures that data is available



• Example: open ()

int open(const char* pathname, int flags, mode_t mode) {
 printf("vdi_log: call %s with '%s', %d, %d\n",
 __func__, pathname, flags, mode);
 int (*actual_open)() = dlsym(RTDL_NEXT, __func__);
 return actual_open(pathname, flags, mode);

- gcc -fPIC -shared -o libvdi.so vdi.c -ldl
- LD_PRELOAD=libvdi.so cat /etc/os-release



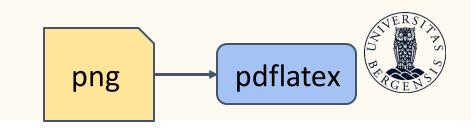
• Log lines

gnuplot

timestamp hostname user \$HOME pid ppid pgid \$PWD program argv starttime elapsed call call_args

- analyse logs to construct data flow graph
- each line produces

png



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timestamp hostname user \$HOME pid ppid pgid \$PWD program argv starttime elapsed call call_args

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• Use cases

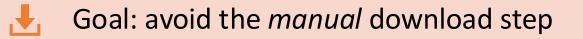
- automatically describe how results (data/figs) were produced
- automatically generate "workflow" descriptions
- detect if some (input) files were not used
- create a timeline of the work (travel back finally possible)
- improve program start-up times (e.g., <u>Spindle</u>)
- EESSI: ensure that build process used the correct data/libs
- EESSI: determine files to pre-load caches



Idea 2: enable access to remote data "directly"



Pattern: a file/dataset is downloaded, then it is used





makes explicit which data source has been used if data has a PID -> can get metadata for dataset



• Example: open()

int open(const char *pathname, int flags, ...) {
 if (is_url(pathname)) {
 if (download(pathname, &local_path) == 0) {
 return actual_open(local_path, flags);
 }
}



- Example: open()
- gcc -fPIC -shared -o libvdi.so vdi.c -ldl lcurl
- (optional) export VDI_DOWNLOAD_BASE=/project/vdi_example
 - **default**:/tmp/\$USER/vdi/downloads
- vdi run wc -l https://zenodo.org/records/13830932/files/ru thalicia_longipes_spades_01.fasta.3.fasta?do wnload=1

"bit" long to write - could support alternatives: doi://10.5281/zenodo.13830932 or zenodo://13830932 or zenodo://13830932?ruthalicia longipes spades 01.fasta.3.fasta

 vdi run wc -l https://zenodo.org/records/13830932/files/ru thalicia_longipes_spades_01.fasta.3.fasta?do wnload=1

- works surprisingly well though not with all commands such as tar
- only for *read* access
- *write* could work too:
 - requires some kind of API at receiver (zenodo provides an API)
 - upload when file closed
- should be possible to optimize
 - only download once or remove download when file closed
 - download a dataset early to prefetch it
- using PIDs (DOI, etc) allows to obtain metadata



- What if all data is accessed via some prefix/namespace?
- vdi://namespace/path_or_label/file[?params]
- A researcher would have to...
 - give the virtual data infrastructure a name (namespace)
 - register data/files/URLs with it



- Runtime could then
 - figure out where the file is stored (locally / remotely),
 - redirect or load the file,
 - $\circ~$ upload files to a sync server or service,
 - create copies,

 \bigcirc

. . .

 \circ give access to others,



- First step: create namespace, add files, obtain URL, ...
- Commands added

vdi view create <name>

vdi view list

vdi view upload <name> <file>

vdi view files <name>

vdi view geturl <name> <file>

View list	Welcome Dat	in View	
iew name Create View	Files in view S	akemake	
A'25 ₪	genome.fa		? 🗊
	genome.fa.am	6	? 🗊
sSo 會 make 會	denome.ta.anr	6	? 🗊
	genome.fa.bw	6	? 🗊
	genome.fa.fai	0	? 🗊
	genome.fa.pad	0	? 🗊
	genome.fa.sa	0	? 💼
	A.fastq	0	? 🗊
	B.fastq	0	? 💼
	C.fastq	0	? 🗊
	Dr	p files here or click to upload files	

Manage namespaces via GUI

Snakemak

• Example remote data access

vdi run head https://vdi.nessi.no/download/Snakemake/genome.fa

• Basic proof of concept



Status

- prototype implementation
- vdi cli command and wrapper library
 - open source, builds and installs in less than 1 min
 - could be shipped with EESSI
- vdi "server" (frontend, API, backend/storage)
 - supports easy creation of data flow graphs
 - supports basic virtual data infrastructure



Outlook

- data flow graph
 - exports: Nextflow, snakemake, LaTeX, ...
 - intercept more functions: *exec*
 - improve performance
- virtual data infrastructure
 - enable use of backends: iRODs, *Cloud, ...
 - runtime that actively manages data in the background



Summary

- Working with research data is not so easy
- Two tools
 - logging data accesses and create data flow graphs
 - effortlessly accessing remote data
- Early prototype: vdi client, wrapper library, frontend & backend
- ➔ Researchers can focus on working with the data ... tools take care of the data management

Resources

- Original report describing the ideas: <u>https://doi.org/10.5281/zenodo.14788711</u>
- Follow-up case study with iRODS: <u>https://doi.org/10.7494/csci.2012.13.4.21</u>
- <u>EESSI European Environment for Scientific Software Installations</u>
 - <u>docs</u> <u>Join EESSI Slack</u> <u>Paper (open access)</u> <u>github</u>
- <u>MultiXscale EuroHPC Centre of Excellence</u> (main development of EESSI)
- Code developed for this talk: <u>https://github.com/virtual-data-infrastructure</u>