

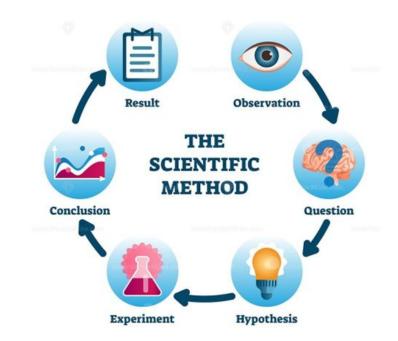
# The reproducibility crisis

02476 Machine Learning Operations Nicki Skafte Detlefsen



### What is reproducibility?

- Being able to reproduce other peoples experimental results is an essential part of the scientific method
- Well known problem throughout most fields (physics, chemistry, biology and computer science)
- With the rise of deep learning, the problem has only been made worse due to competition





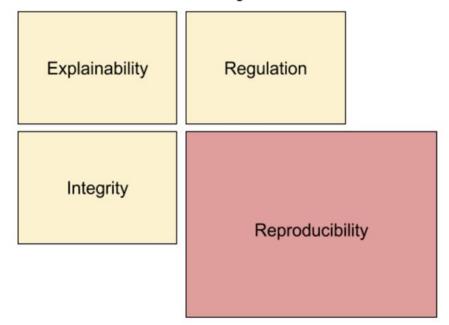
## Why to we need reproducibility?

Reproducibility is a key component in *Trustworthy ML* 

#### Case:

Imaging an AI agent used for diagnostics. Without reproducibility two persons with the exact same symptoms could get different diagnosis

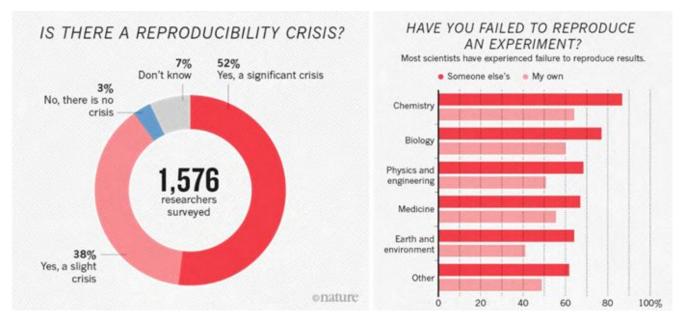
#### Trustworthy ML





#### How bad is it?

Wow this is bad...



Machine Learning around 22%



## A closer look at machine learning

Re-Implementation of 255 paper. Hypothesis testing on what "paper features" have an effect on reproducibility.

#### A Step Toward Quantifying Independently Reproducible Machine Learning Research

#### **Edward Raff**

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#### Abstract

What makes a paper independently reproducible? Debates on reproducibility center around intuition or assumptions but lack empirical results. Our field focuses on releasing code, which is important, but is not sufficient for determining reproducibility. We take the first step toward a quantifiable answer by manually attempting to implement 255 papers published from 1984 until 2017, recording features of each paper, and performing statistical analysis of the results. For each paper, we did not look at the authors code, if released, in order to prevent bias toward discrepancies between code and paper.

Table 1: Significance test of which paper properties impact reproducibility. Results significant at  $\alpha \le 0.05$  marked with"\*".

Feature	p-value
Year Published	0.964
Year First Attempted	0.674
Venue Type	0.631
Rigor vs Empirical*	$1.55 \times 10^{-9}$
Has Appendix	0.330
Looks Intimidating	0.829
Readability*	$9.68 \times 10^{-25}$
Algorithm Difficulty*	$2.94 \times 10^{-5}$
Pseudo Code*	$2.31\times10^{-4}$
Primary Topic*	$7.039 \times 10^{-4}$
Exemplar Problem	0.720
Compute Specified	0.257
Hyperparameters Specified*	$8.45 \times 10^{-6}$
Compute Needed*	$8.75 \times 10^{-5}$
Authors Reply*	$6.01 \times 10^{-8}$
Code Available	0.213
Pages	0.364
Publication Venue	0.342
Number of References	0.740
Number Equations*	0.004
Number Proofs	0.130
Number Tables*	0.010
Number Graphs/Plots	0.139
Number Other Figures	0.217
Conceptualization Figures	0.365
Number of Authors	0.497



## What are the field trying to do?

#### https://paperswithcode.com/



#### Neurips checklist:

- 3. If you ran experiments...
  - (a) Did you include the code, data, and instructions needed to reproduce the main experimental results (either in the supplemental material or as a URL)?
    - . The instructions should contain the exact command and environment needed to run to reproduce the results.
    - Please see the NeurlPS code and data submission guidelines for more details.
    - Main experimental results include your new method and baselines. You should try to capture as many of the minor
      experiments in the paper as possible. If a subset of experiments are reproducible, you should state which ones are.
    - While we encourage release of code and data, we understand that this might not be possible, so "no because the code is proprietary" is an acceptable answer.
    - · At submission time, to preserve anonymity, remember to release anonymized versions.
  - (b) Did you specify all the training details (e.g., data splits, hyperparameters, how they were chosen)?
    - The full details can be provided with the code, but the important details should be in the main paper.
  - (c) Did you report error bars (e.g., with respect to the random seed after running experiments multiple times)?
    - Answer "yes" if you report error bars, confidence intervals, or statistical significance tests for your main experiments.
  - (d) Did you include the amount of compute and the type of resources used (e.g., type of GPUs, internal cluster, or cloud provider)?
    - Ideally, you would provide the compute required for each of the individual experimental runs as well as the total compute.
    - Note that your full research project might have required more compute than the experiments reported in the paper (e.g., preliminary or failed experiments that didn't make it into the paper). The total compute used may be harder to characterize, but if you can do that, that would be even better.
    - You are also encouraged to use a CO2 emissions tracker and provide that information. See, for example, the
      experiment impact tracker (Henderson et al.), the ML CO2 impact calculator (Lacoste et al.), and CodeCarbon.



### Reproducibility are not a fixed concept

Reproducibility of model

Reproducibility of results

Dataset	<b>Model Architecture</b>	Random Init	Transfer	Parameters	IMAGENET Top5
RETINA	Resnet-50	96.4% ± 0.05	96.7% ± 0.04	23570408	92.% ± 0.06
RETINA	Inception-v3	$96.6\% \pm 0.13$	96.7% ± 0.05	22881424	93.9%
RETINA	CBR-LargeT	$96.2\% \pm 0.04$	$96.2\% \pm 0.04$	8532480	$77.5\% \pm 0.03$
RETINA	CBR-LargeW	$95.8\% \pm 0.04$	95.8% ± 0.05	8432128	$75.1\% \pm 0.3$
RETINA	CBR-Small	$95.7\% \pm 0.04$	$95.8\% \pm 0.01$	2108672	$67.6\% \pm 0.3$
RETINA	CBR-Tiny	$95.8\% \pm 0.03$	$95.8\% \pm 0.01$	1076480	$73.5\% \pm 0.05$

What is the difference?



### What can you do about it?

Make sure to document everything about your experiments

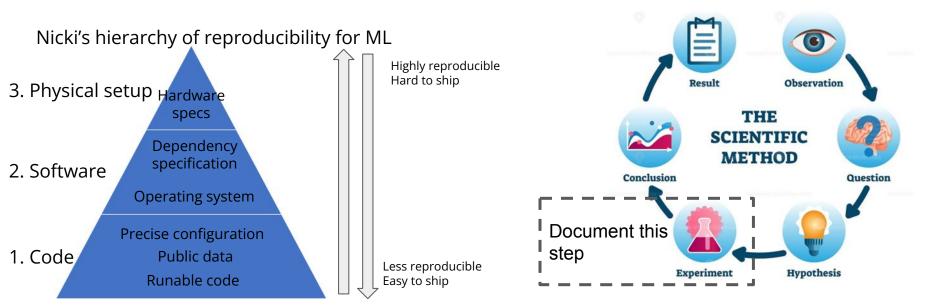


Figure credit: https://www.australianenvironmentaleducation.com.au/education-resources/what-is-the-scientific-method/



#### A closer look

Use software to abstract away problem

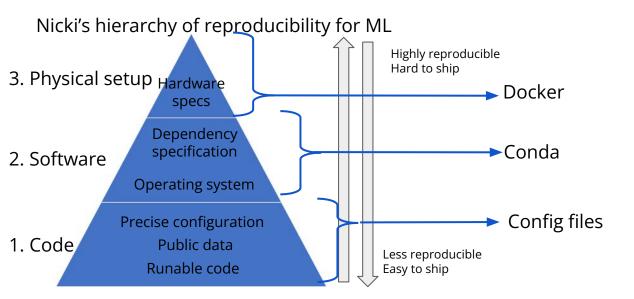


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# **Reproducibility 1: Code level**

There is a lot of subjective choices that we do when running experiments, most notable the hyperparameters.

Parameter in script	Argparser	Config files
<pre>class hparams:     lr = 0.1     batch_size = 16     num_layers = 5</pre>	python my_script.py \lr 0.1 \batch_size 16 \num_layers 5	experiment1.yaml  lr: 0.001  batch_size: 16  num_layers: 5  python my_script.py \  config=experiment1.yaml
<ul> <li>No easy configuration</li> <li>Experiments may be lost if not careful</li> </ul>	<ul> <li>Easy to configure new experiment</li> <li>Falls on user to save the config</li> </ul>	<ul> <li>Moderate level of configurability</li> <li>Parameters are systematically saved with experiments</li> </ul>



## Reproducibility 1: Code level

Hydra is a framework for elegantly configuring complex (ML) applications

my\_app()

https://github.com/facebookresearch/hydra





```
conf
    config.yaml
    dataset
      cifar10.yaml
      imagenet.yaml
my_app.py
```

#### Other options

- https://github.com/IDSIA/sacred https://mlflow.org/



## **Reproducibility 2: Software level**

For python: Just use a package management system

#### Examples:

- Conda (what I like)
- Pipenv
- venv
- pyenv

```
(lightning) C:\Users\nsde\Documents\metrics>conda env list
# conda environments:
                         C:\Users\nsde\Anaconda3
ensemble
                         C:\Users\nsde\Anaconda3\envs\ensemble
laplace
                        C:\Users\nsde\Anaconda3\envs\laplace
                      * C:\Users\nsde\Anaconda3\envs\lightning
lightning
mixerensemble
                         C:\Users\nsde\Anaconda3\envs\mixerensemble
                        C:\Users\nsde\Anaconda3\envs\mlops
mloos
                        C:\Users\nsde\Anaconda3\envs\protein
protein
                        C:\Users\nsde\Anaconda3\envs\pvae
ovae
stochman
                         C:\Users\nsde\Anaconda3\envs\stochman
(lightning) C:\Users\nsde\Documents\metrics>
```

(lightning) C:\Users\r			
# packages in environm	ent at C:\Users\n	sde\Anaconda3\envs\1	ightning:
#			
# Name	Version		Channel
absl-py	1.2.0	pypi_0	рурі
aiohttp	3.8.3	pypi_0	рурі
aiosignal	1.2.0	pypi_0	рурі
alabaster	0.7.12	pypi_0	pypi
asttokens	2.0.5	pyhd3eb1b0_0	
async-timeout	4.0.2	pypi_0	рурі
atomicwrites	1.4.1	pypi_0	рурі
attrs	22.1.0	pypi_0	pypi
babel	2.10.3	pypi_0	pypi
backcall	0.2.0	pyhd3eb1b0_0	
beautifulsoup4	4.11.1	pypi_0	рурі
black	22.8.0	pypi_0	pypi
blas	2.116	mkl	conda-forge
blas-devel	3.9.0	16_win64_mkl	conda-forge
bleach	5.0.1	pypi_0	pypi
brotlipy	0.7.0	py38h294d835_1004	conda-forg
build	0.8.0	pypi_0	pypi
ca-certificates	2022.07.19	haa95532_0	
cachetools	5.2.0	pypi_0	pypi
certifi	2022.9.14	py38haa95532_0	
cffi	1.15.1	py38hd8c33c5_0	conda-forge
cfgv	3.3.1	pypi_0	pypi
charset-normalizer	2.1.1	pyhd8ed1ab_0	conda-forge
check-manifest	0.48	pypi_0	pypi
click	8.1.3	pypi_0	pypi
cloudpickle	2.2.0	pypi_0	рурі
colorama	0.4.5	py38haa95532_0	
commonmark	0.9.1	pypi_0	pypi
contourpy	1.0.5	pypi_0	pypi
coverage	6.4.4	pypi_0	рурі
cryptography	37.0.4	py38hb7941b4_0	conda-forge
cudatoolkit	11.6.0	hc9ea762_10	conda-forge
cycler	0.11.0	pypi 0	pypi
decorator	5.1.1	pyhd3eb1b0 0	
defusedxml	0.7.1	pypi_0	рурі
distlib	0.3.6	pypi_0	pypi
docutils	0.17.1	pypi_0	pypi
dython	0.7.2	pypi 0	pypi



### **Reproducibility 3: Hardware level**

The easiest would be to hand over your machine. In practise not that possible.

Instead lets hand over a "virtual machine".

Virtual machines takes hardware from the host and creates virtual CPU, RAM, storage for each virtual machine. The VMs are completely independent from the host.







The core advantage of a VM is that it in principal can run on any host without changes, because it is independent.

Docker can be seen as an lightweight version of full VMs.

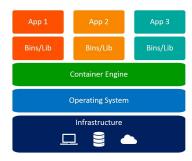
VM is isolation of machines, while Containers is isolation of processes







Virtual machine by stable diffusion



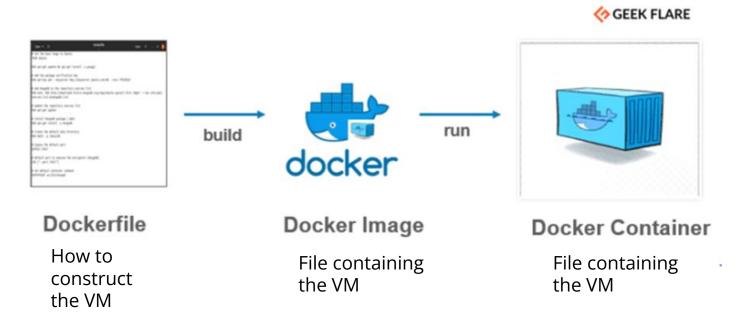
Containers





### **Reproducibility 3: Hardware level**

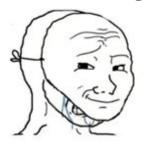
A way to create containerized applications = specialized VMs





### Meme of the day

#### **Programmers**



This code is unreadable and your dataset is flawed. No one will be able to reproduce your results!



It's not my fault the legacy environment is messed up! We still have 97.3% unit test coverage.

#### Scientist



This code is unreadable and your dataset is flawed. No one will be able to reproduce your results!



I know:)