### INFRASTRUCTURE AS CODE

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# WHAT IS IAC?

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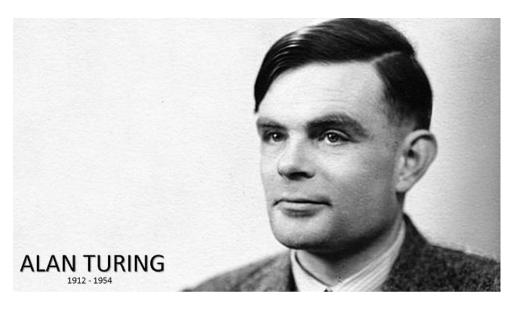
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### BEFORE THAT, SOME BASIC CONCEPTS

- Can the system perform logic?
- It is capable of iterate?
- And to expand memory?

#### SOME (UNINTENTIONAL) TURING COMPLETE SYSTEMS

- Microsoft excell
- C++ templates
- Power point
- Minesweeper
- Minecraft
- Magic: The Gathering



#### BEFORE THAT, SOME BASIC CONCEPTS

process { executor = 'sge' 2 3 mapping.clusterOptions = "-P project -pe omp 16 -l mem\_total=94G" 4 counting.clusterOptions = "-P project -pe omp 8 -l mem\_total=16G" 5 6 indir = "/Users/pipeliner/pipelines/toy\_data/rna-seq" 7 outdir = "/Users/pipeliner/pipelines/rna-seg-results" 8 fasta = "\${params.indir}/genome\_reference.fa" 9 10 qtf = "\${params.indir}/genome\_annotation.gtf" 11 12 // General pipeline parameters 13 paired = true 14 aligner = "hisat" 15 quantifier = "featurecounts" 16 17 index.use\_existing = true 18 index.path = "\${params.indir}/alignment\_indices/hisat\_index/index/part" 19 // Process-specific parameters 20 21 feature\_counts.cpus = 8 22 feature\_counts.type = "exon" 23 feature\_counts.id = "gene\_id" 24 ..... feature\_counts.xargs = = "" 25 feature\_counts.ainj 26



#### BEFORE THAT, SOME BASIC CONCEPTS

Do we, humans, treat the same way config files and code files?

Are config files different from programs?

### BEFORE THAT, SOME BASIC CONCEPTS

### Software does everything code does... and more!



#### Asserting libraries

- Integration testing
- Cross compiling
- Debugging
- Logging
- Versioning
- Refactoring
- Upgrading (Downgrading!)
- Flow control
- Linting
- Syntax control
- Observability
- Hinting, User experience
- Repeatability
- Compatibility
- Composability
- Inheritance
- Polymorphism



#### executor = 'sge'

mapping.clusterOptions = "-P project -pe omp 16 -l mem\_total=94G" counting.clusterOptions = "-P project -pe omp 8 -l mem\_total=166"

indir = "/Users/pipeliner/pipelines/toy\_data/rna-seq"
outdir = "/Users/pipeliner/pipelines/rna-seq-results"
fasta = "\${params.indir}/genome\_reference.fa"

gtf = "\${params.indir}/genome\_annotation.gtf"

// General pipeline parameters

- paired = true
  aligner = "hisat"
- aligner = "hisat" quantifier = "featurecounts"

quantifier = reactifect

index.use\_existing = true index.path = "\${params.indir}/alignment\_indices/hisat\_index/index/part"

// Process-specific parameters
feature\_counts.cpus = 8
feature\_counts.type = "exon"
feature\_counts.id = "gene\_id"

- feature\_counts.id = "g
  feature\_counts.xargs = ""
- 24 teature\_counts.xargs = ""
  25 feature\_counts.ainj = ""
- }



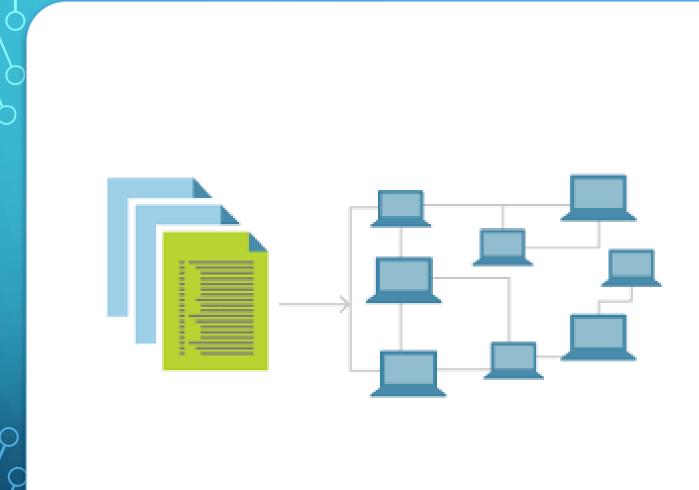
### ANALYSIS OF BASIC CONCEPTS

Example: imagine you have to configure a server and you are in a Turing Complete use case



### CONFIGURATE INFRASTRUCTURE

- Nowadays we are already using config files to manage infrastructures
- Config file: hard coding
- Use software: advantages everywhere



#### SO, WHAT IS IAC OR IAS?

Avoid the use of config files to manage the configuration of infrastructure and instead use code or software

# WHAT ARE THE COMMON USES OF IAC?

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### DECENTRALIZED COMPANIES

- Very common: companies with several hq
- $\diamond$  Hq in different regions == compatibility issues
  - IaC unifies system configurations.

#### SERVER DEPLOYMENT



- Need of contuous services
- Possible scenario: machine break down
- IaC allows for faster deployment

### CONTINUOUS INTEGRATION

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- Deploying manually and periodically = time consuming
- IaC = solution



- IaC template incident
- Vulnerability propagation
- Solution: Compliance as code



# HAVE WE USED IAC AS STUDENTS?

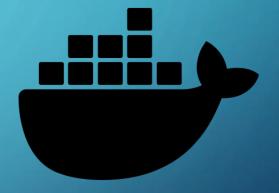
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VM





Docker



# WHEN DO WE USE IAC?

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# SHOULD I USE IAC?

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# SIMPLIC



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### ABILITY

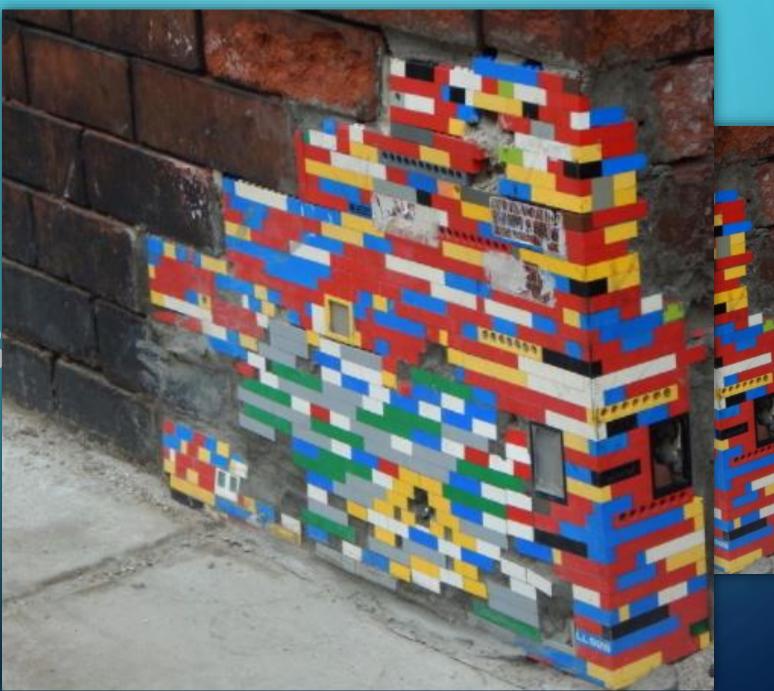
# SHAREABILITY

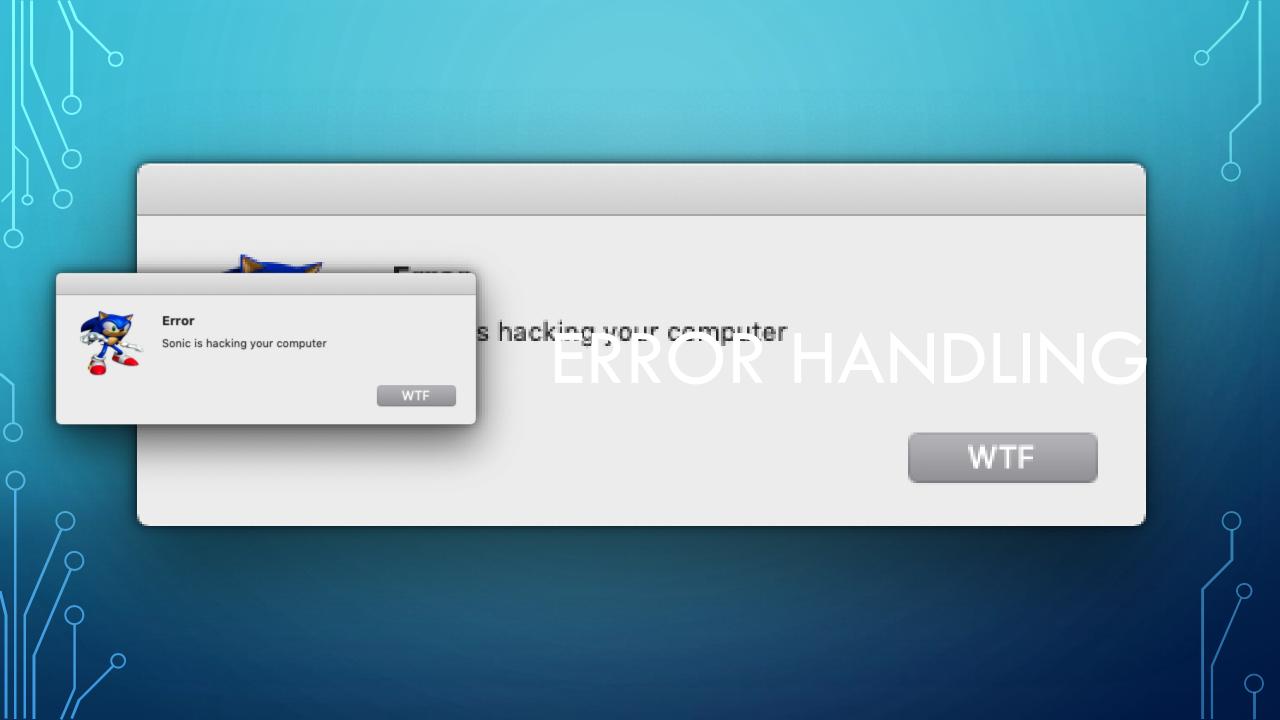


### STAB

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### SCALA

My IaC code with different configurations

with different

My IaC code



# SIMPLICITY





#### UNDERSTANDABILITY



#### LEARNABILITY



#### ERROR HANDLING







SCALABILITY



MAINTAINABILITY