University of Zurich[UZH]

s.e.a.l.
software evolution & architecture lab

SANER'17

Klagenfurt, Austria

Reducing Redundancies in Multi-Revision Code Analysis

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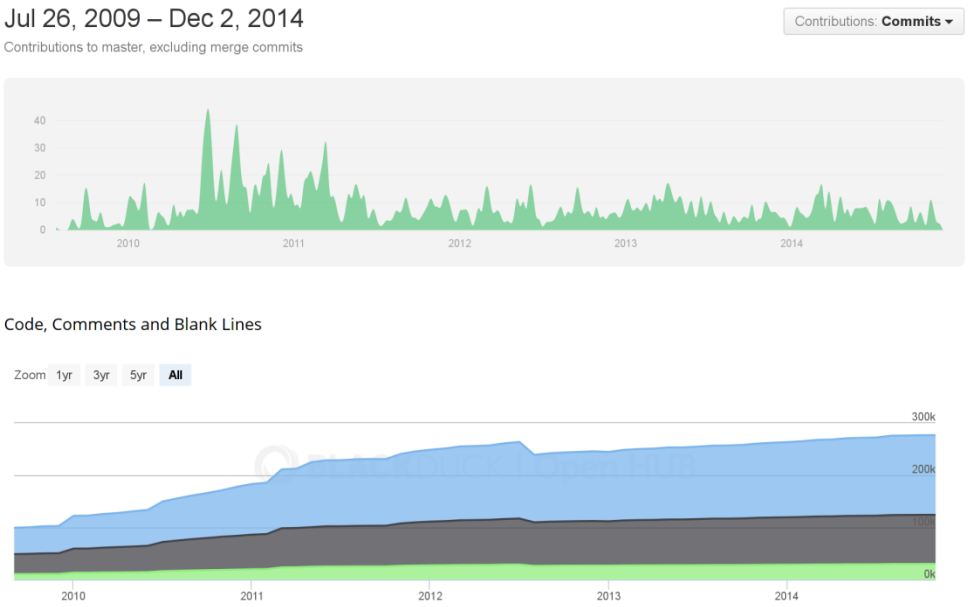
22.02.2017

The Problem Domain

- Static analysis (e.g. #Attr., McCabe, coupling...)



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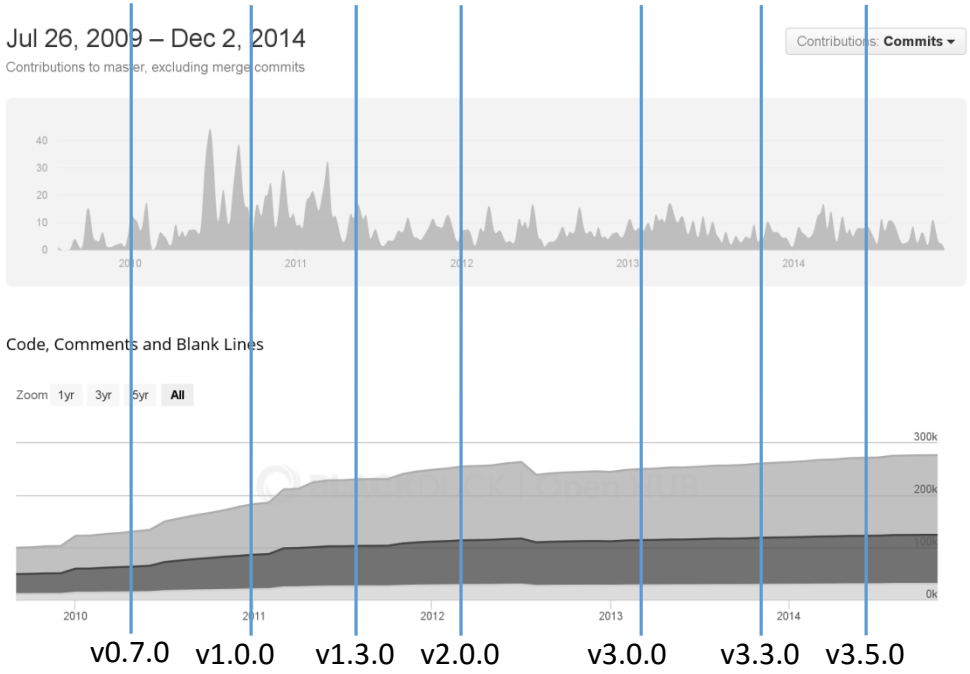
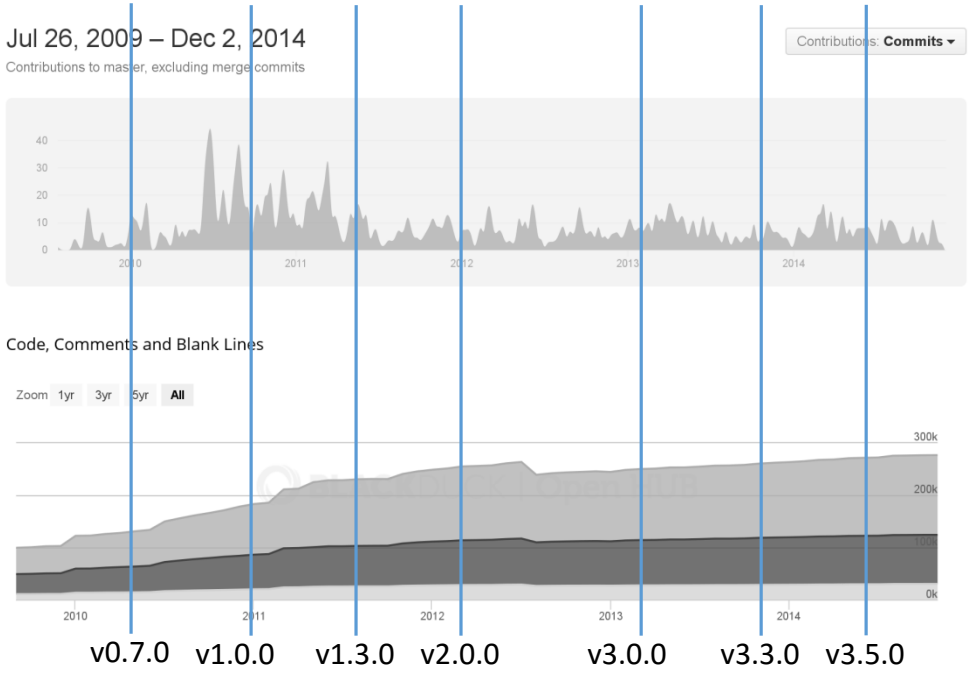
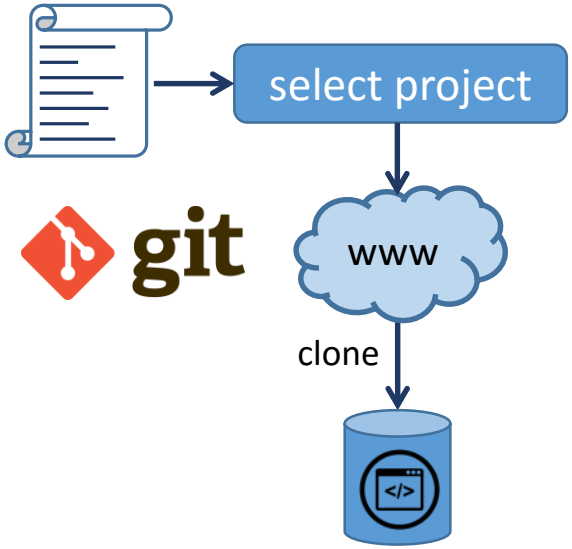


The Problem Domain

- Static analysis (e.g. #Attr., McCabe, coupling...)
- Many revisions, fine-grained historical data



A Typical Analysis Process



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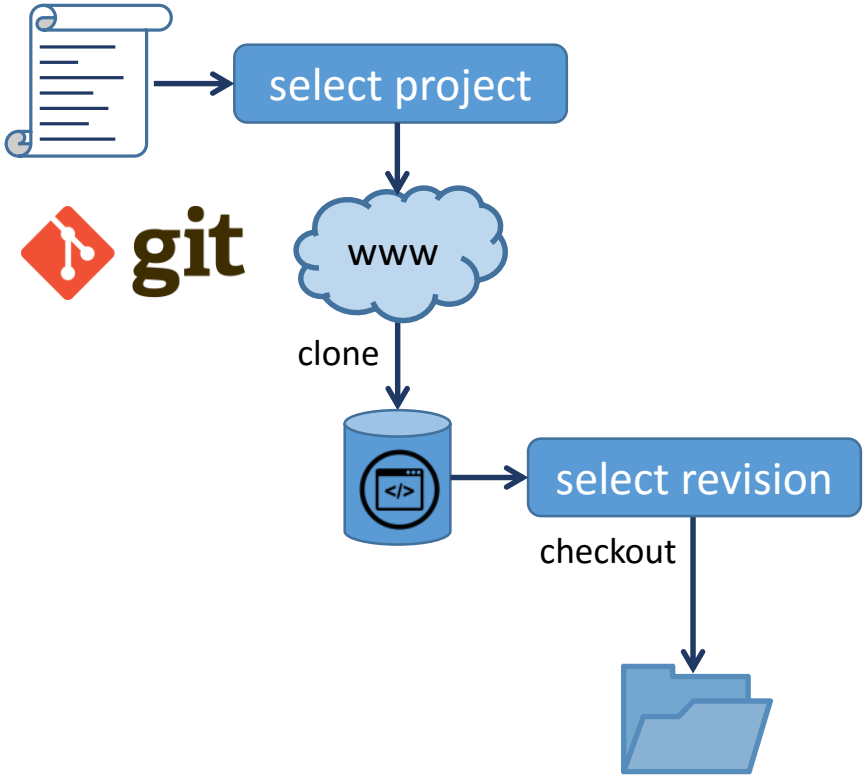
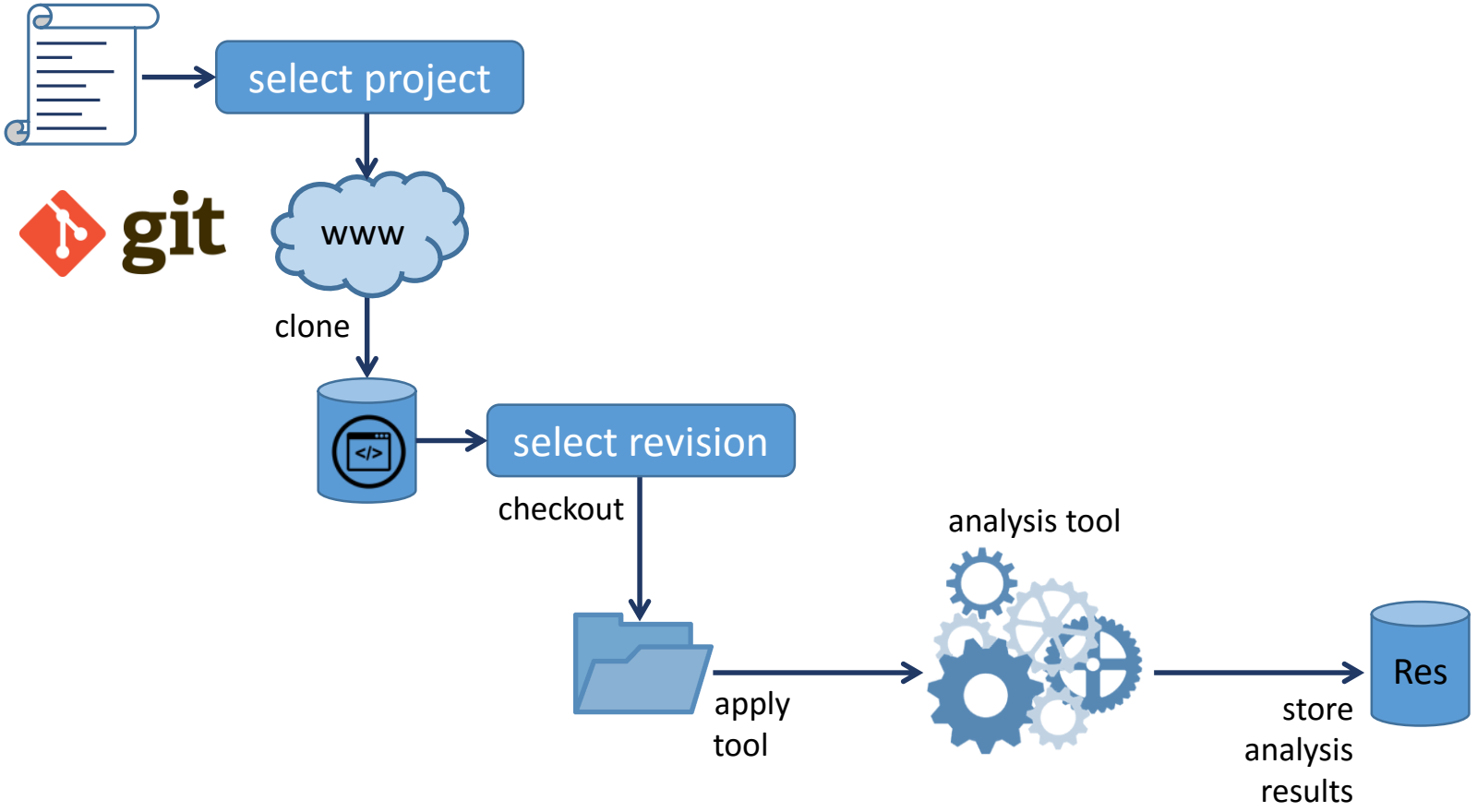
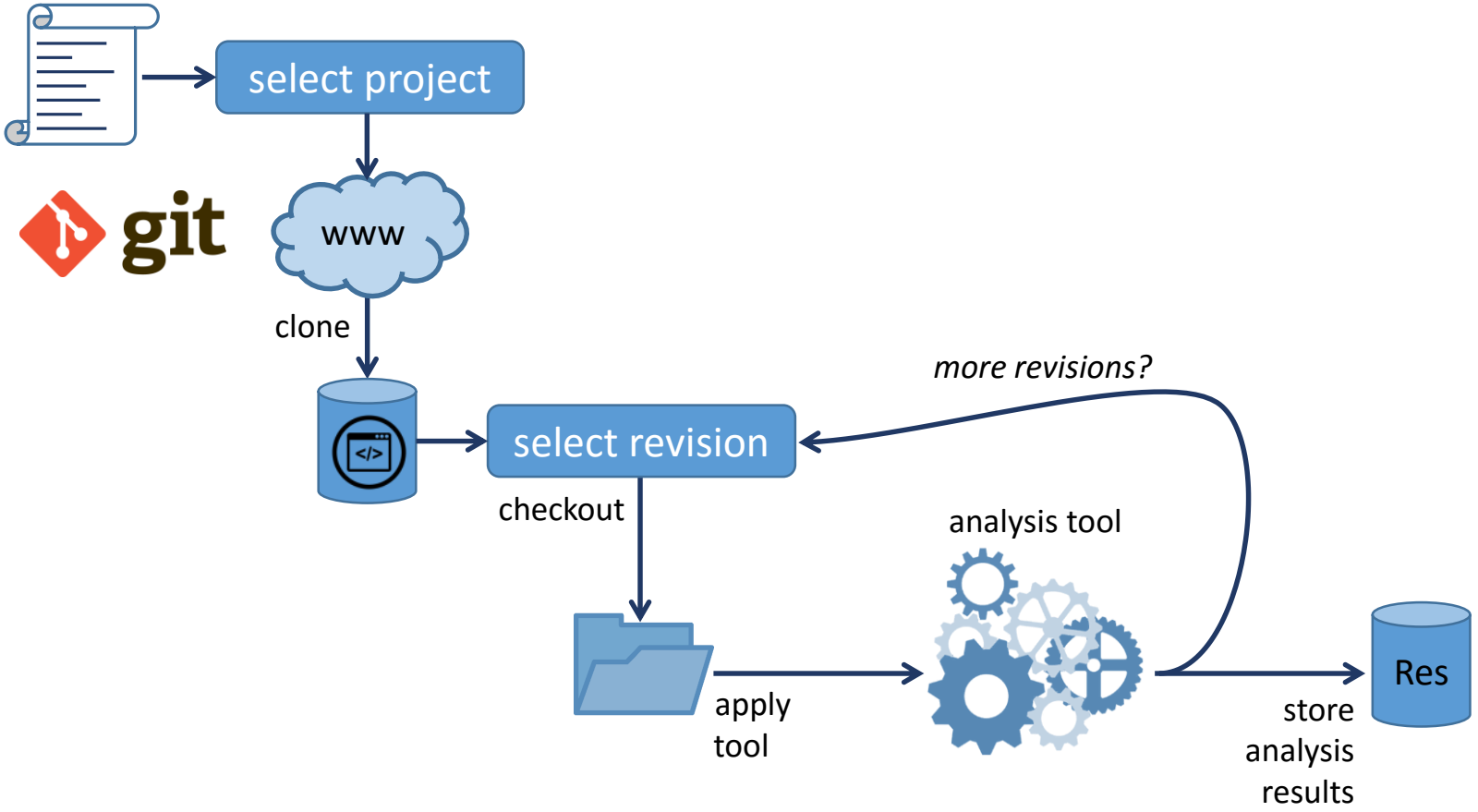


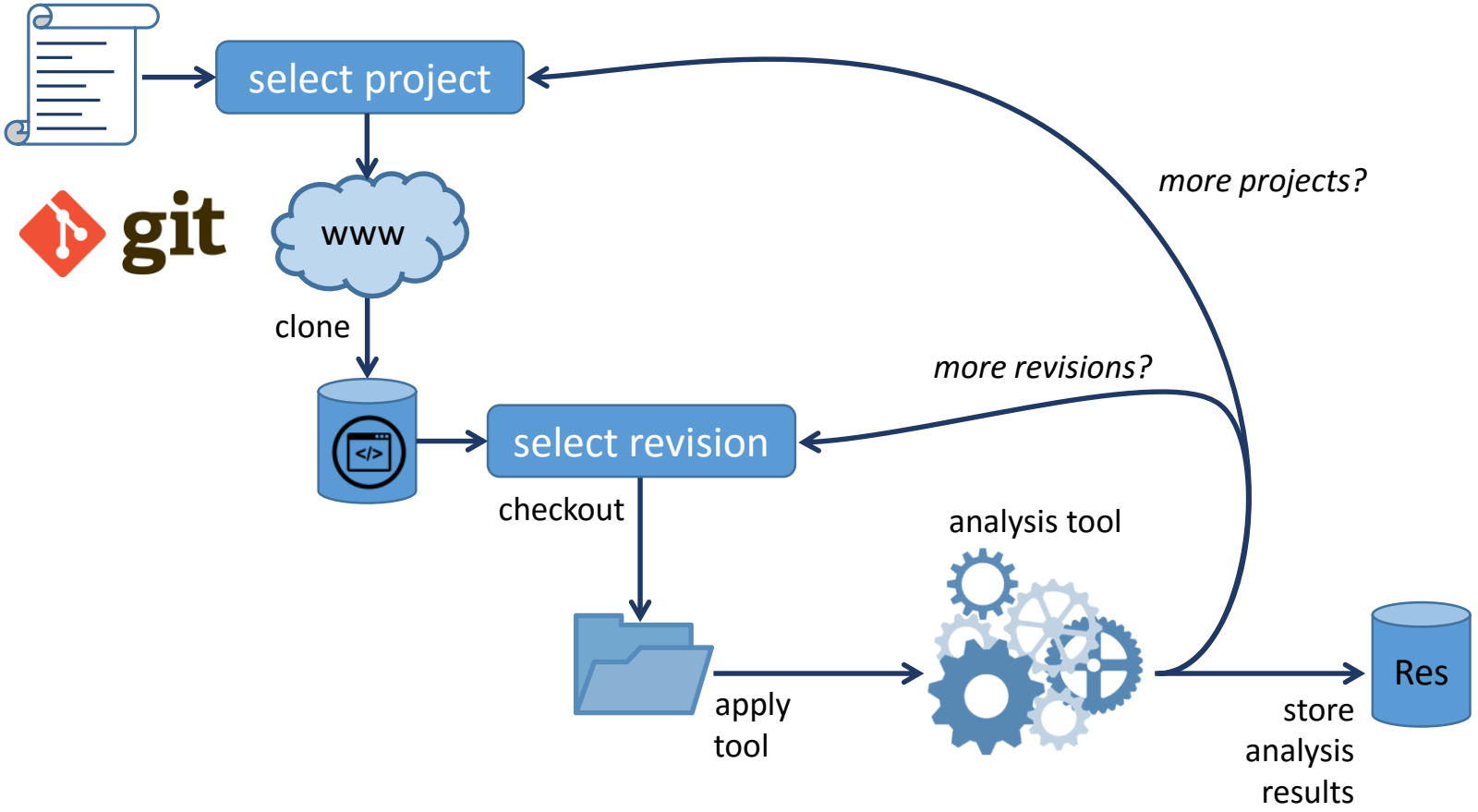
A Typical Analysis Process



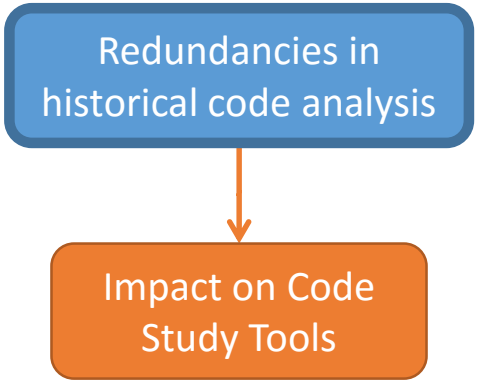
A Typical Analysis Process



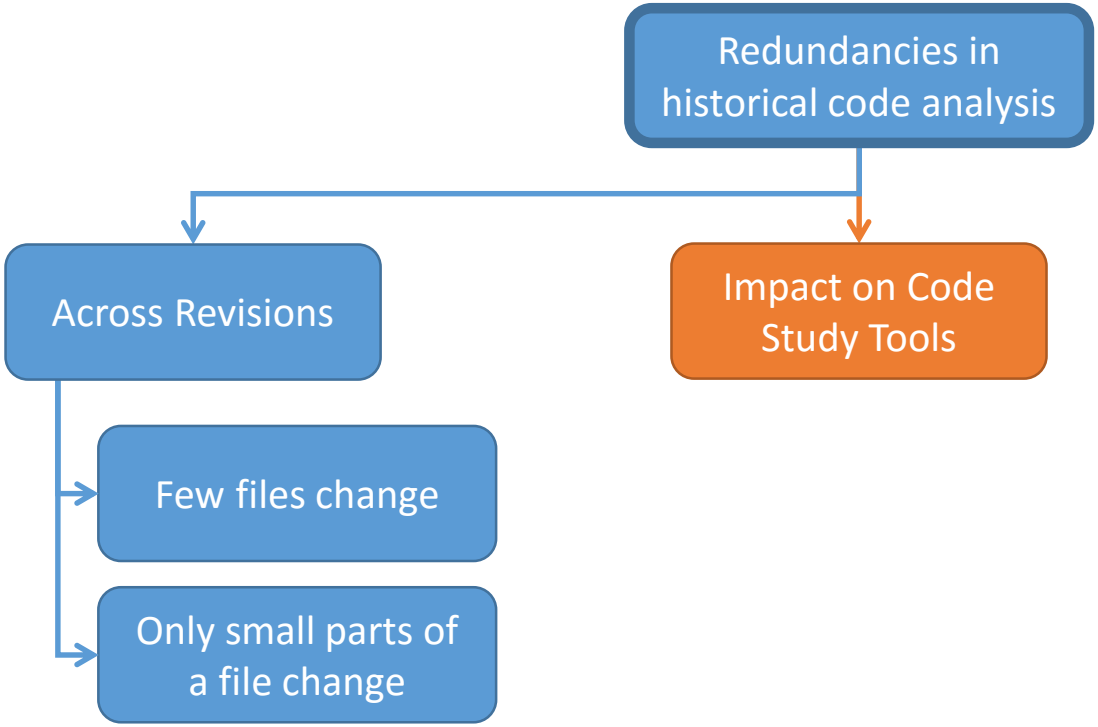
A Typical Analysis Process



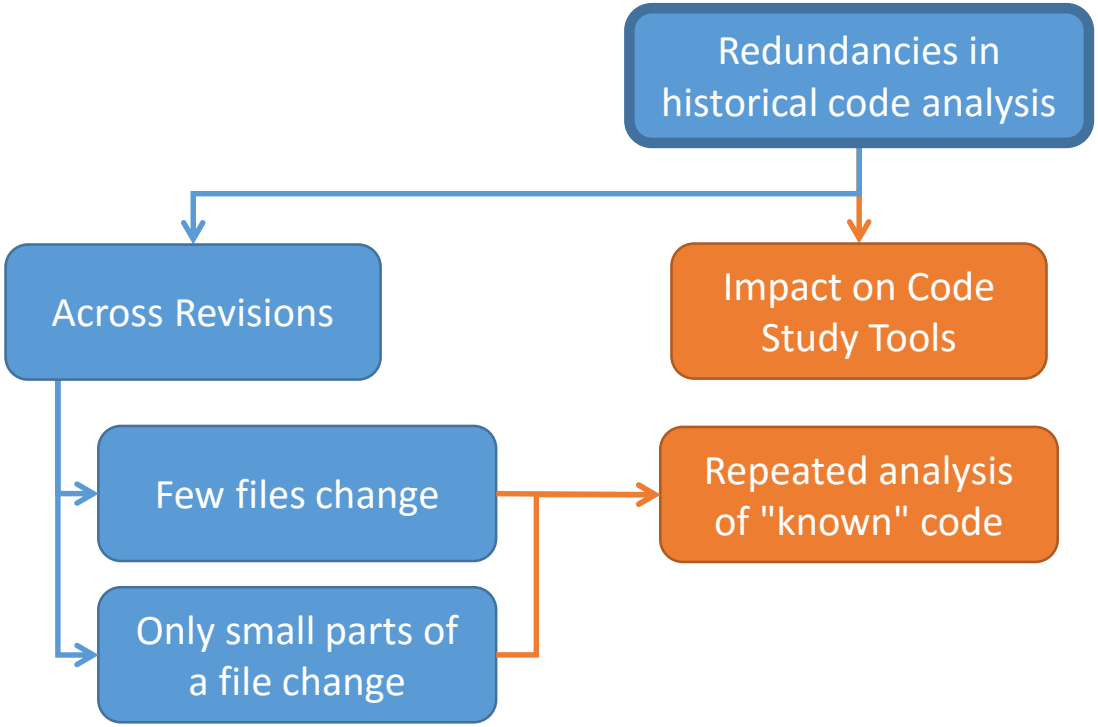
Redundancies all over...



Redundancies all over...



Redundancies all over...



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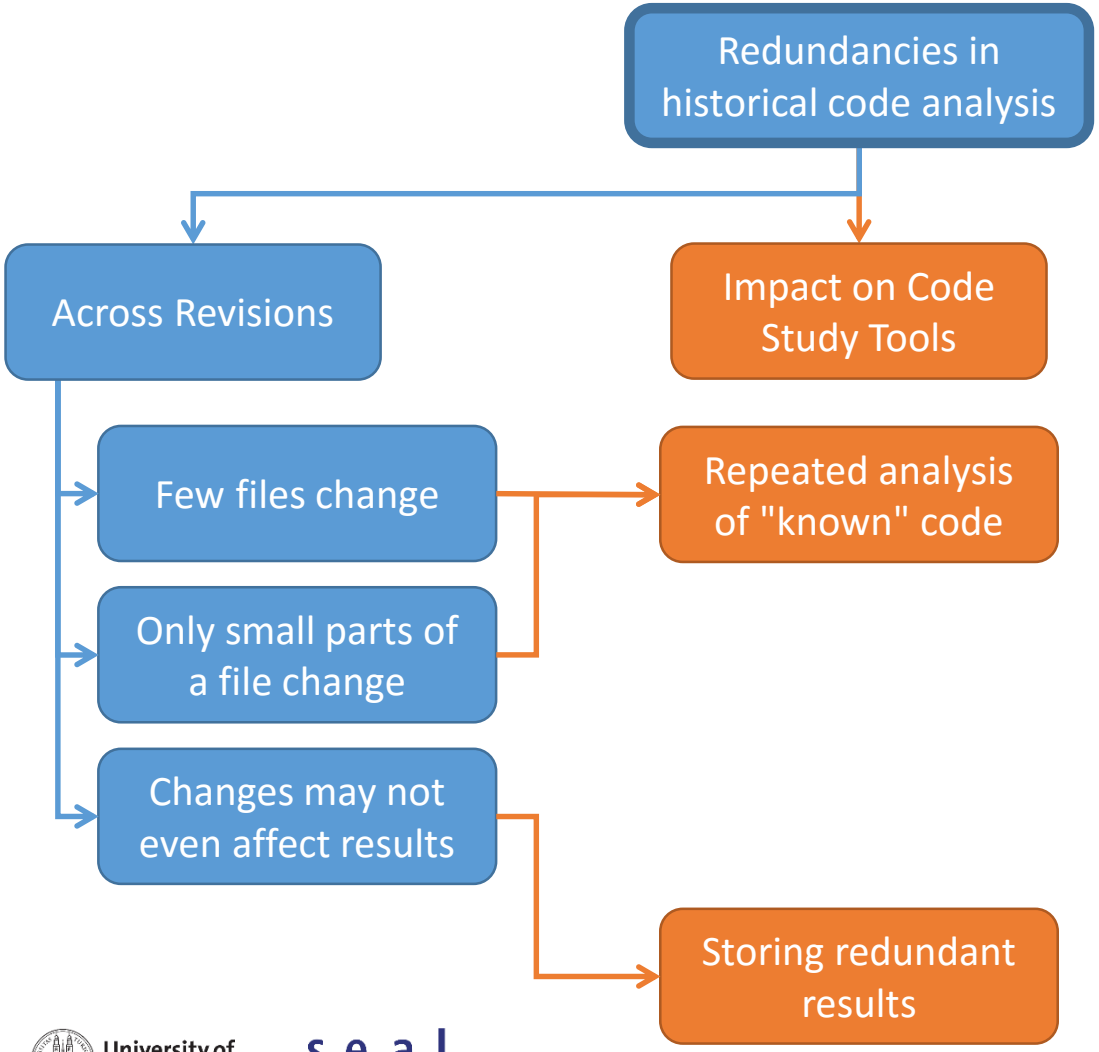
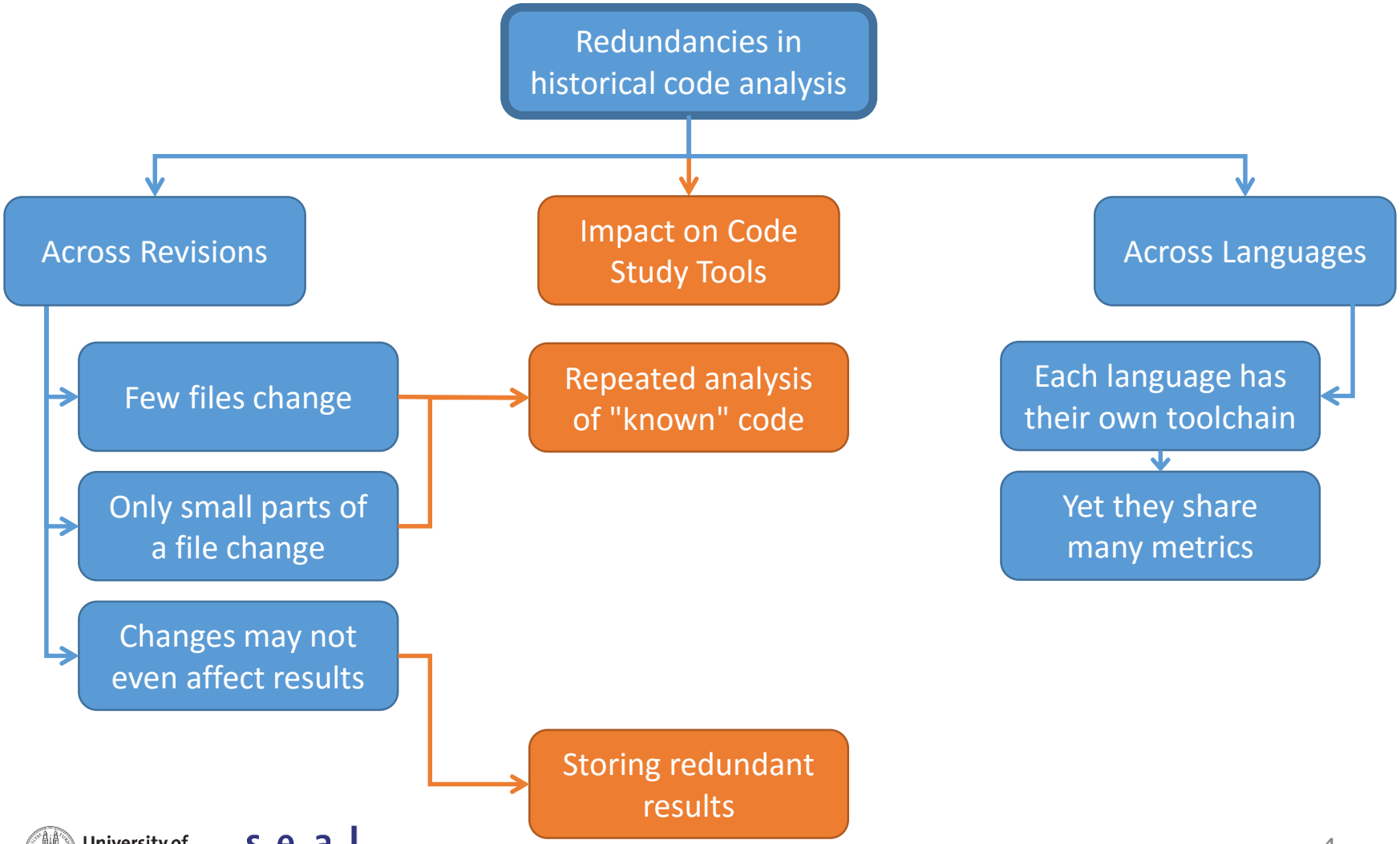
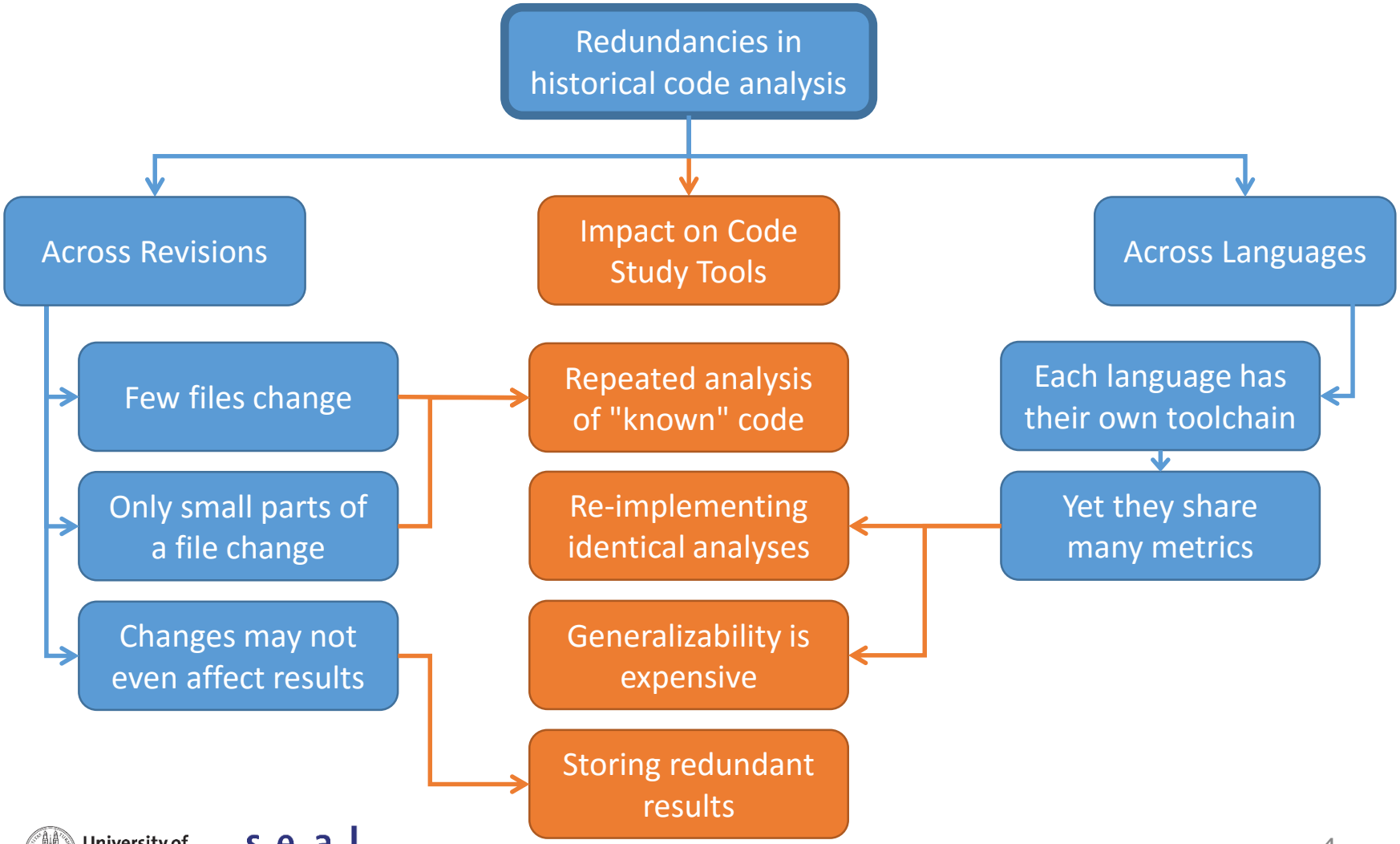


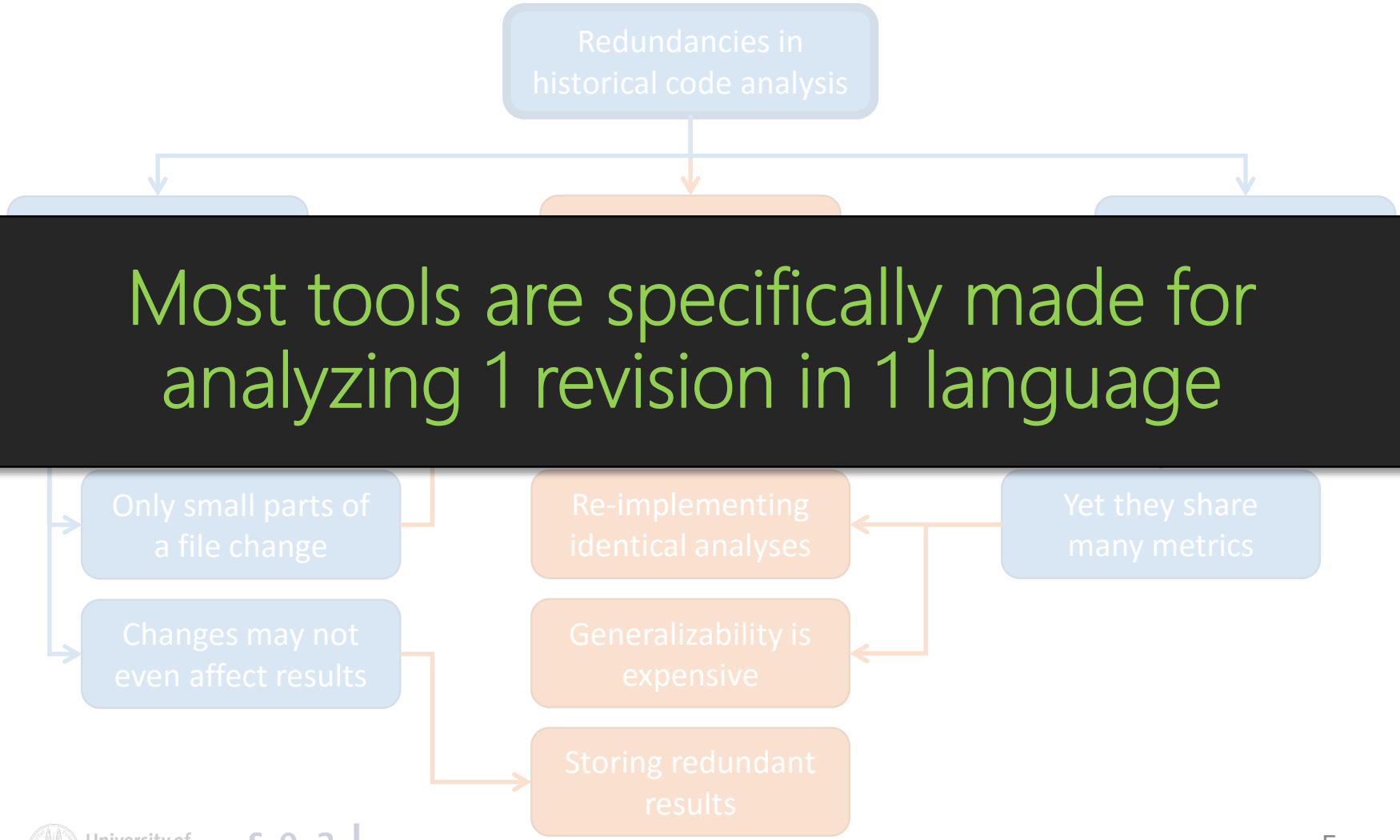
Redundancies all over...



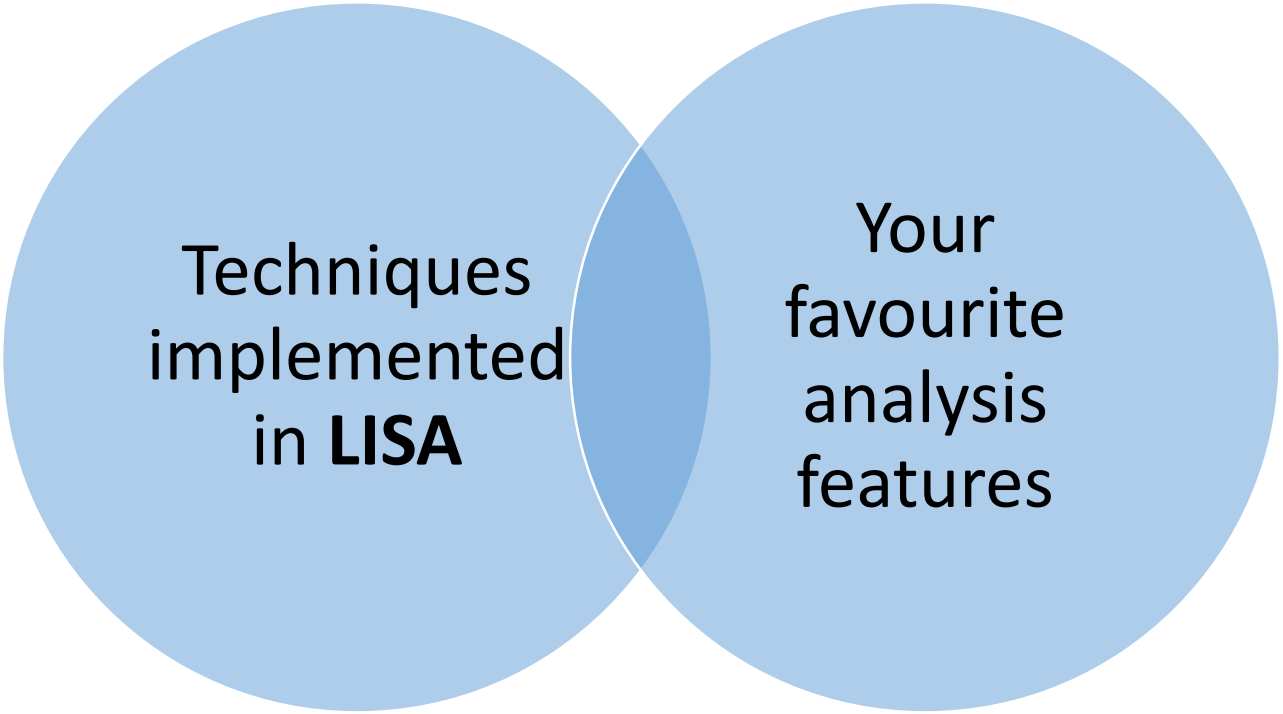
Redundancies all over...



Redundancies all over…



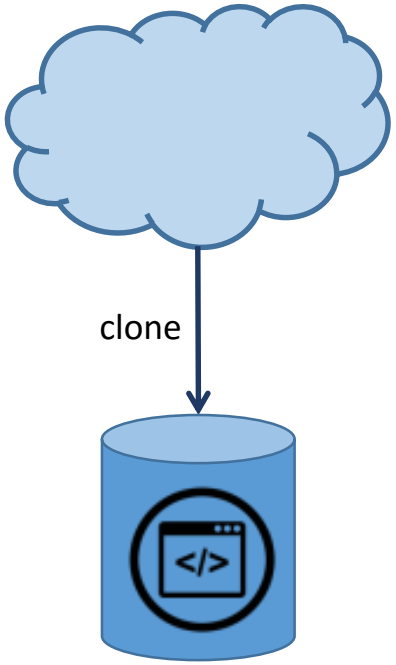
Important!



Pick what you like!

#1: Avoid Checkouts

Avoid checkouts



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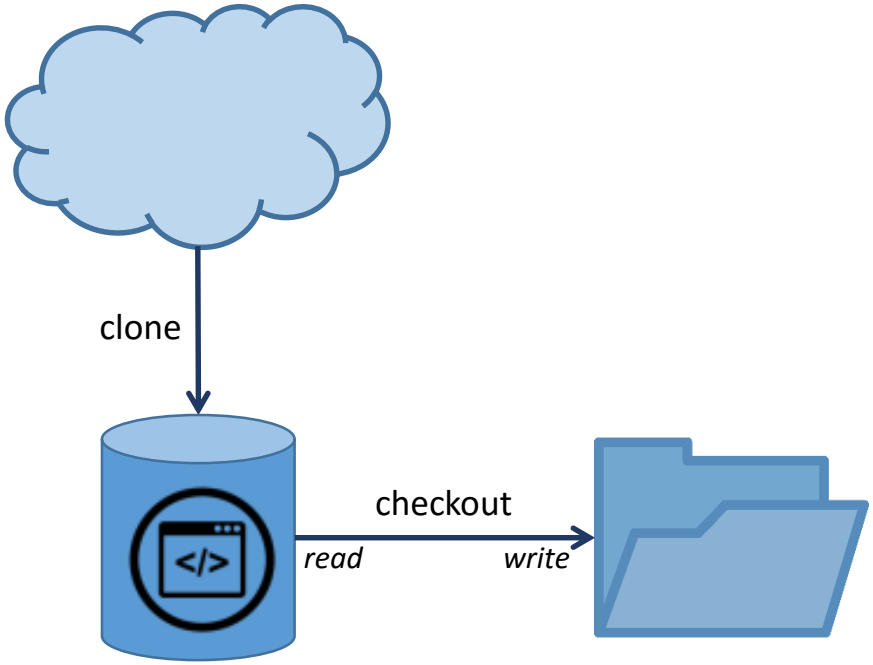

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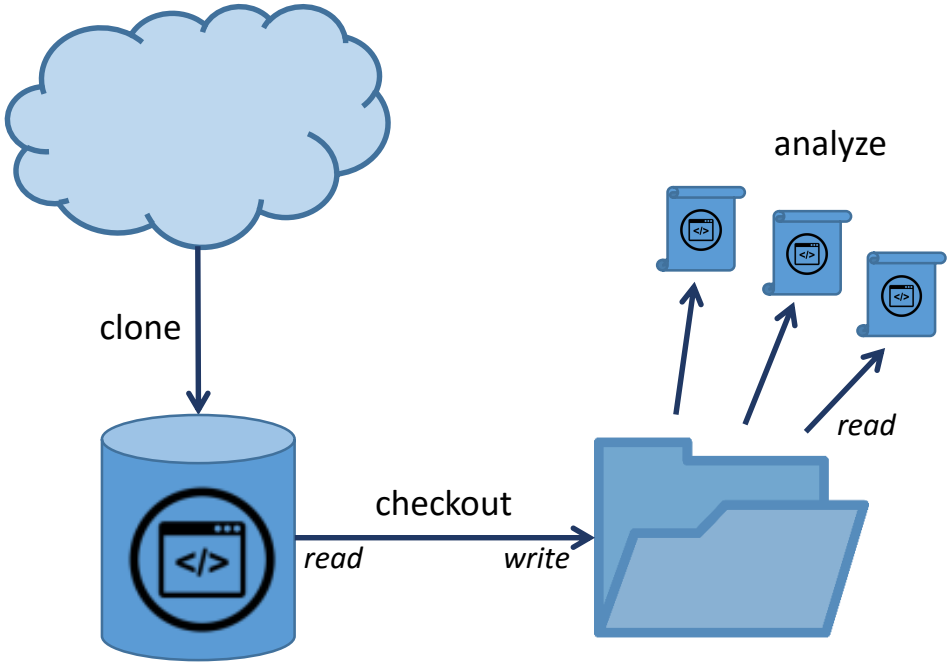
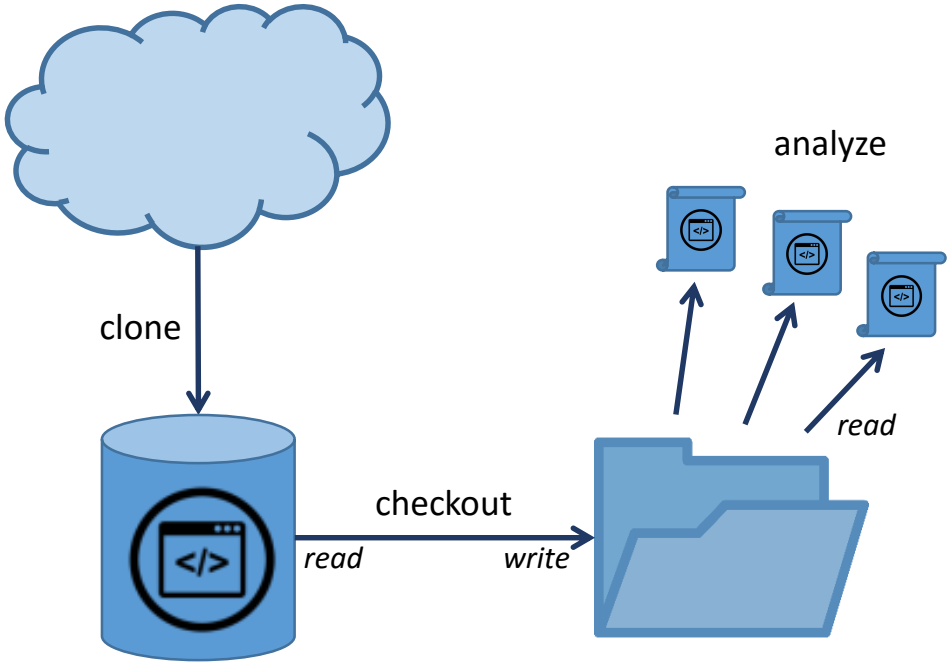
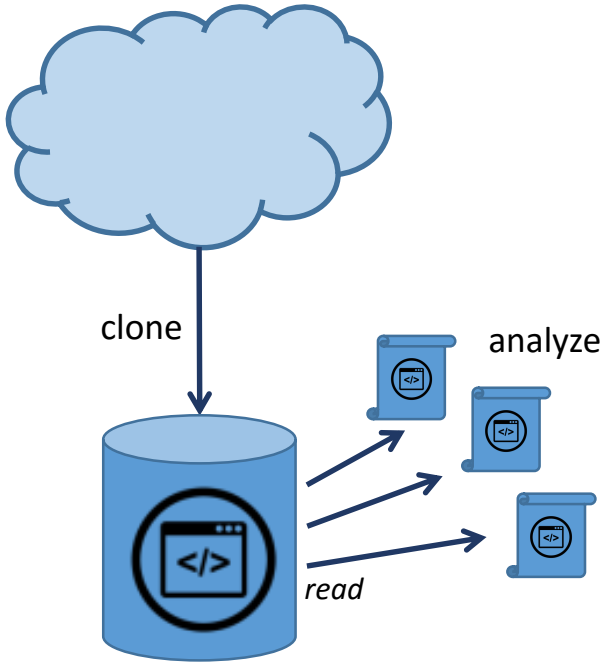


Avoid checkouts

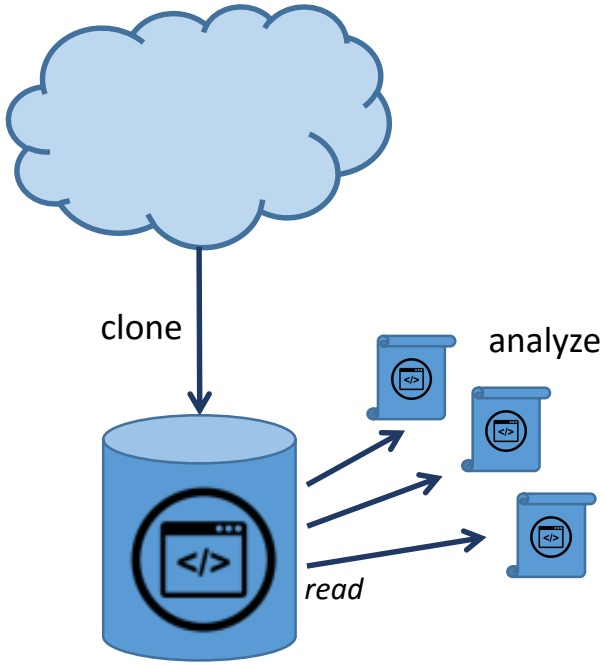


For every file: 2 read ops + 1 write op
Checkout includes irrelevant files
Need 1 CWD for every revision to be analyzed in parallel

Avoid checkouts

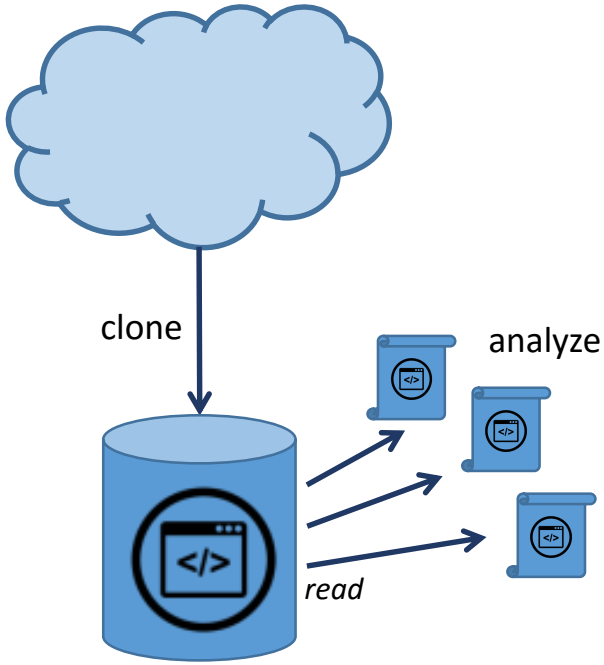


Avoid checkouts



Only read relevant files in a single read op
No write ops
No overhead for parallization

Avoid checkouts



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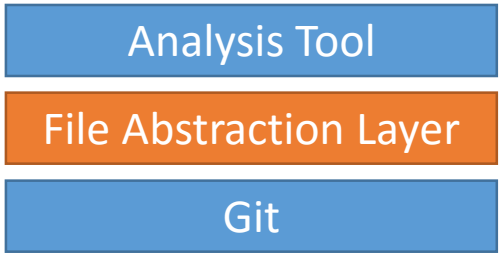
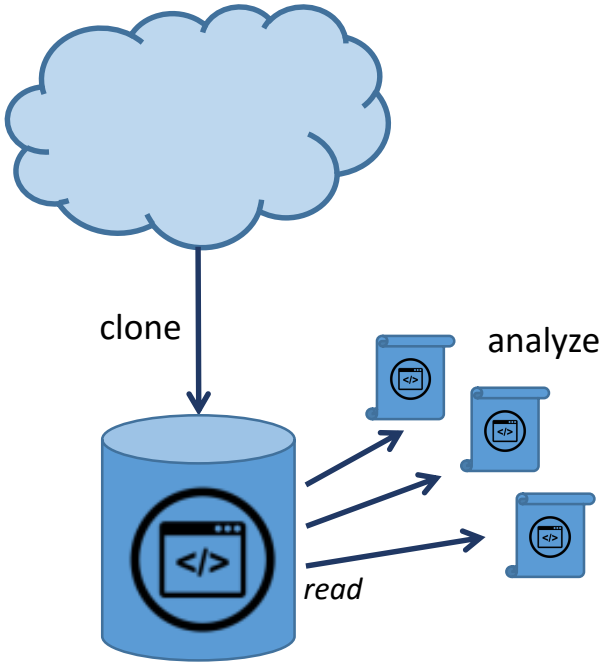


Avoid checkouts



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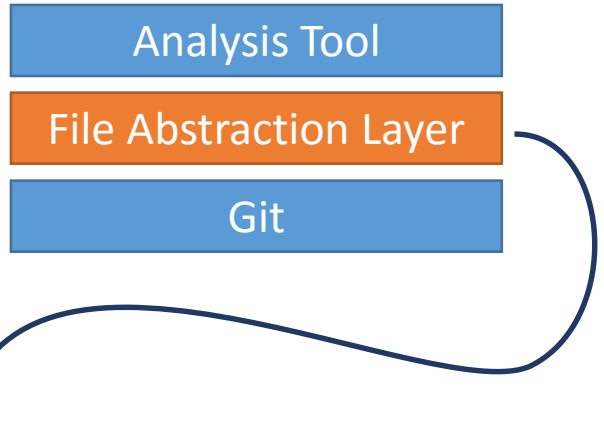


E.g. for the JDK Compiler:

```
class JavaSourceFromCharArray(name: String, val code: CharBuffer)
extends SimpleJavaFileObject(URI.create("string:///" + name), Kind.SOURCE) {
  override def getCharContent(): CharSequence = code
}
```

Avoid checkouts

Only read relevant files in a single read op
No write ops
No overhead for parallization

The simplest time-saver:
If you can - operate directly on bare Git

E.g. for the JDK Compiler:

```
class JavaSourceFromCharArray(name: String, val code: CharBuffer)
extends SimpleJavaFileObject(URI.create("string:///" + name), Kind.SOURCE) {
  override def getCharContent(): CharSequence = code
}
```

#2: Use a multi-revision representation of your sources

Merge ASTs

rev. 1 

rev. 2 

rev. 3 

rev. 4 

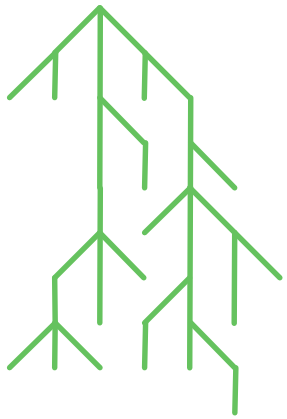
Merge ASTs

rev. 1

rev. 2

rev. 3

rev. 4



rev. 1

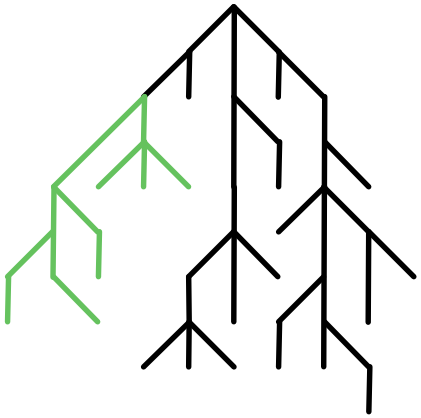
Merge ASTs

rev. 1

rev. 2

rev. 3

rev. 4



rev. 2

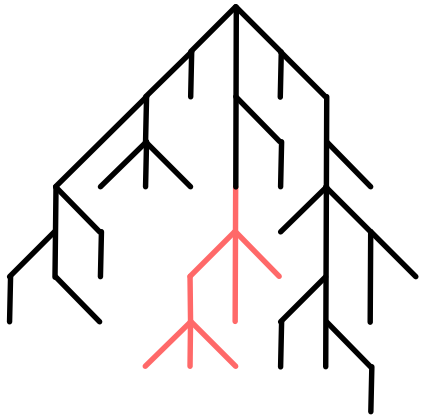
Merge ASTs

rev. 1

rev. 2

rev. 3

rev. 4



rev. 3

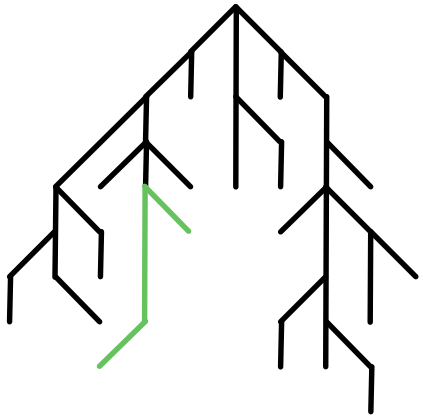
Merge ASTs

rev. 1

rev. 2

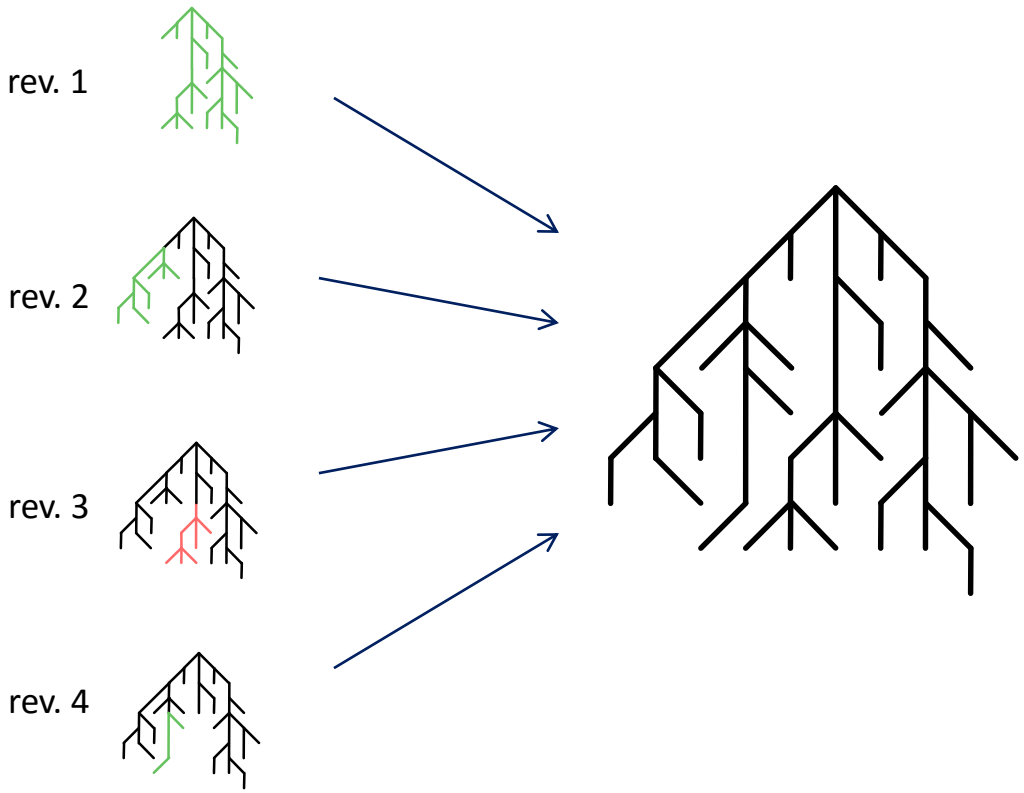
rev. 3

rev. 4

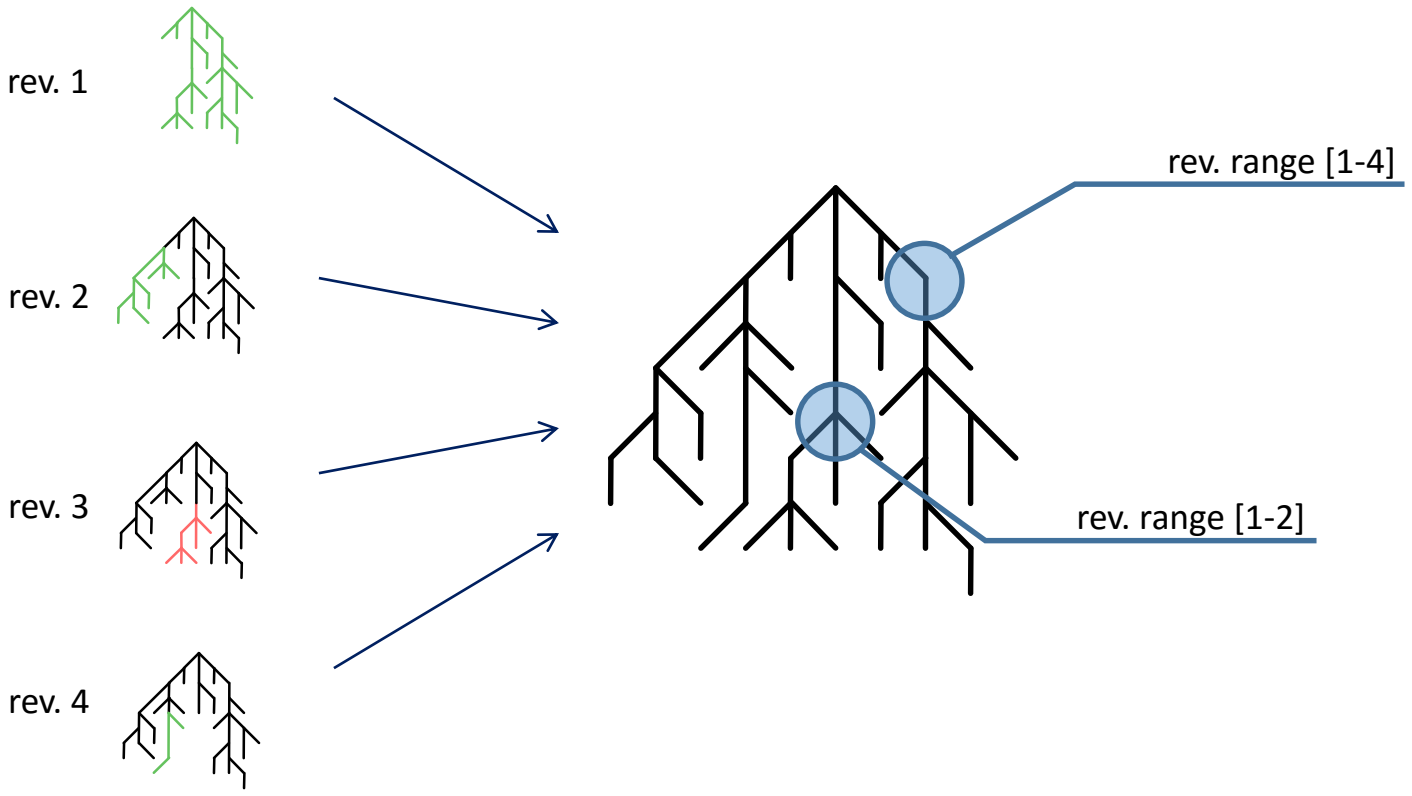


rev. 4

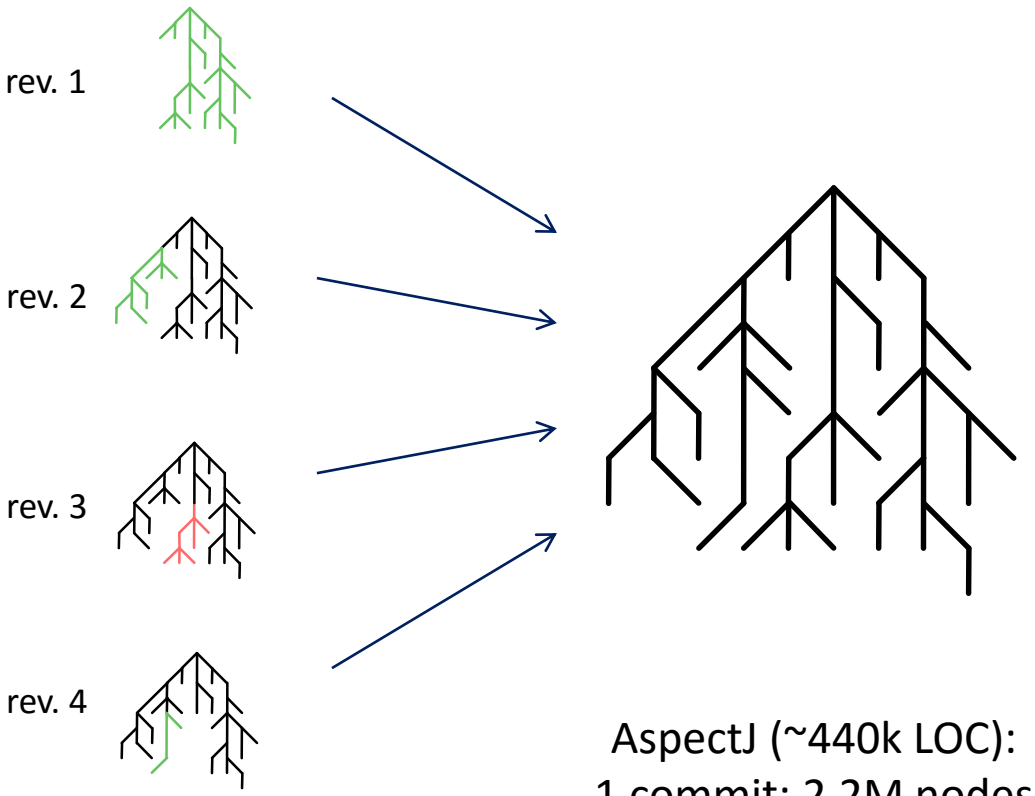
Merge ASTs



Merge ASTs



Merge ASTs



AspectJ (~440k LOC):
1 commit: 2.2M nodes
All >7000 commits: 6.5M nodes

Merge ASTs

rev. 1

rev. 4

Merging ASTs brings exponential space and time savings

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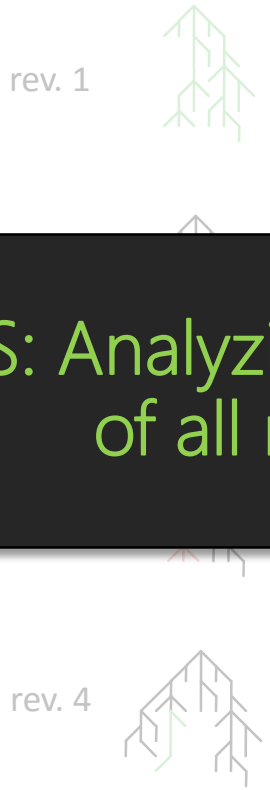
1 commit: 2.2M nodes
All >7000 commits: 6.5M nodes

Merge ASTs



PS: Analyzing multiple revisions implies building a graph of all revisions *first*, and analyzing it *afterwards*

AspectJ (~440k LOC):
1 commit: 2.2M nodes
All >7000 commits: 6.5M nodes

#3: Store AST nodes only if they're needed for analysis

```
public class Demo {
  public void run() {
    for (int i = 1; i< 100; i++) {
      if (i % 3 == 0 || i % 5 == 0) {
         System.out.println(i)
      }
  }
}
```

What's the complexity (1+#forks) and name for each method and class?

```
public class Demo {
  public void run() {
    for (int i = 1; i< 100; i++) {
      if (i % 3 == 0 || i % 5 == 0) {
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  }
}
```

parse

140 AST nodes
(using ANTLR)

What's the complexity (1+#forks) and name for each method and class?

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public class Demo {
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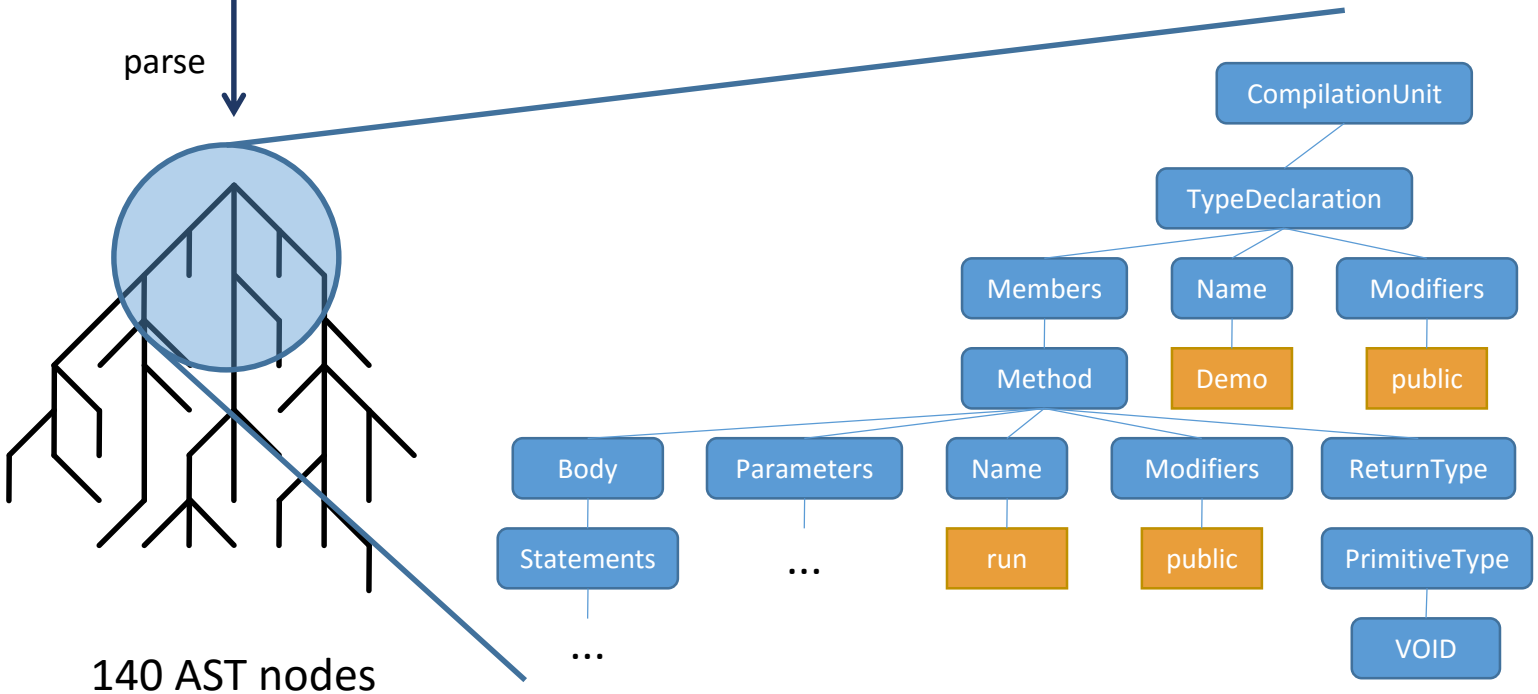
parse



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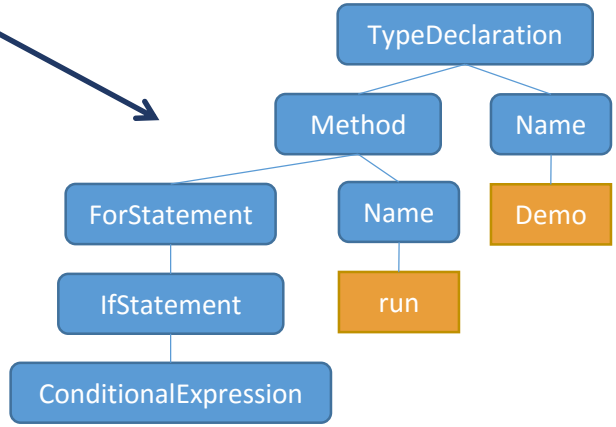
  public void run() {
    for (int i = 1; i< 100; i++) {
      if (i % 3 == 0 || i % 5 == 0) {
         System.out.println(i)
      }
  }
}
```

What's the complexity (1+#forks) and name for each method and class?

parse

filtered parse

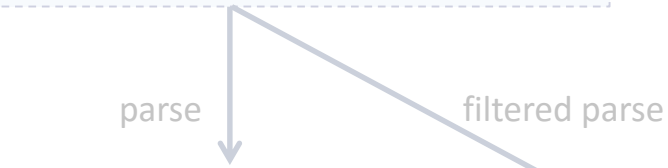


140 AST nodes
(using ANTLR)

7 AST nodes
(using ANTLR)

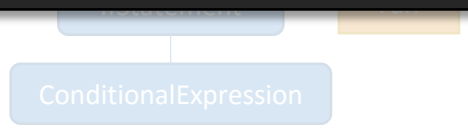
```
public class Demo {
  public void run() {
    for (int i = 1; i< 100; i++) {
      if (i % 3 == 0 || i % 5 == 0) {
         System.out.println(i)
      }
  }
}
```

What's the complexity (1+#forks) and name for eachmethod and class?



Storing only needed AST nodes applies a manyfold reduction in needed space

140 AST nodes
(using ANTLR)



7 AST nodes
(using ANTLR)

```
public class Demo {
  public void run() {
    for (int i = 1; i< 100; i++) {
      if (i % 3 == 0 || i % 5 == 0) {
         System.out.println(i)
      }
  }
}
```

What's the complexity (1+#forks) and name for eachmethod and class?

parse

filtered parse

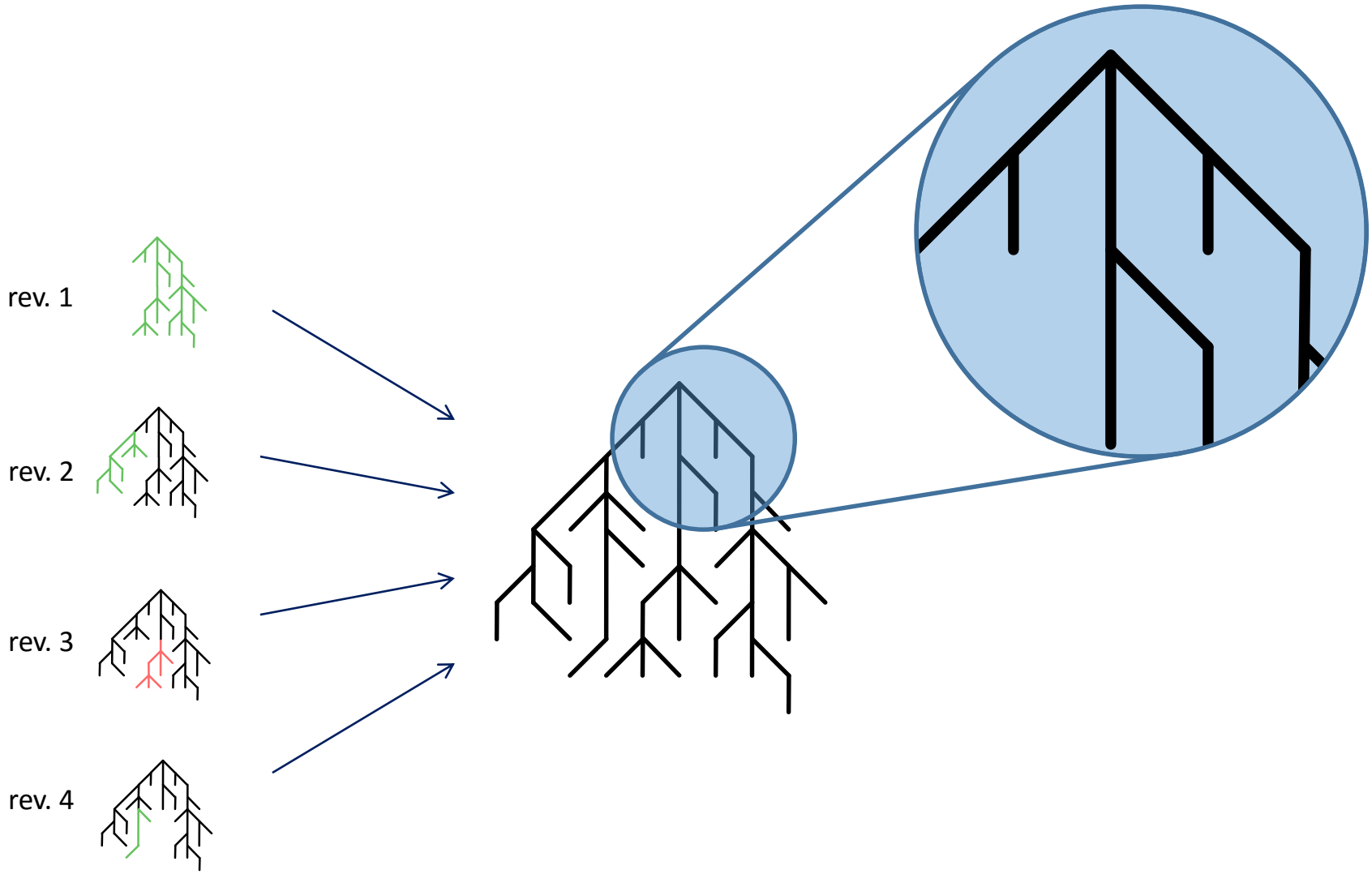
PS: Which AST nodes to load into the graph **depends on** the **analysis**

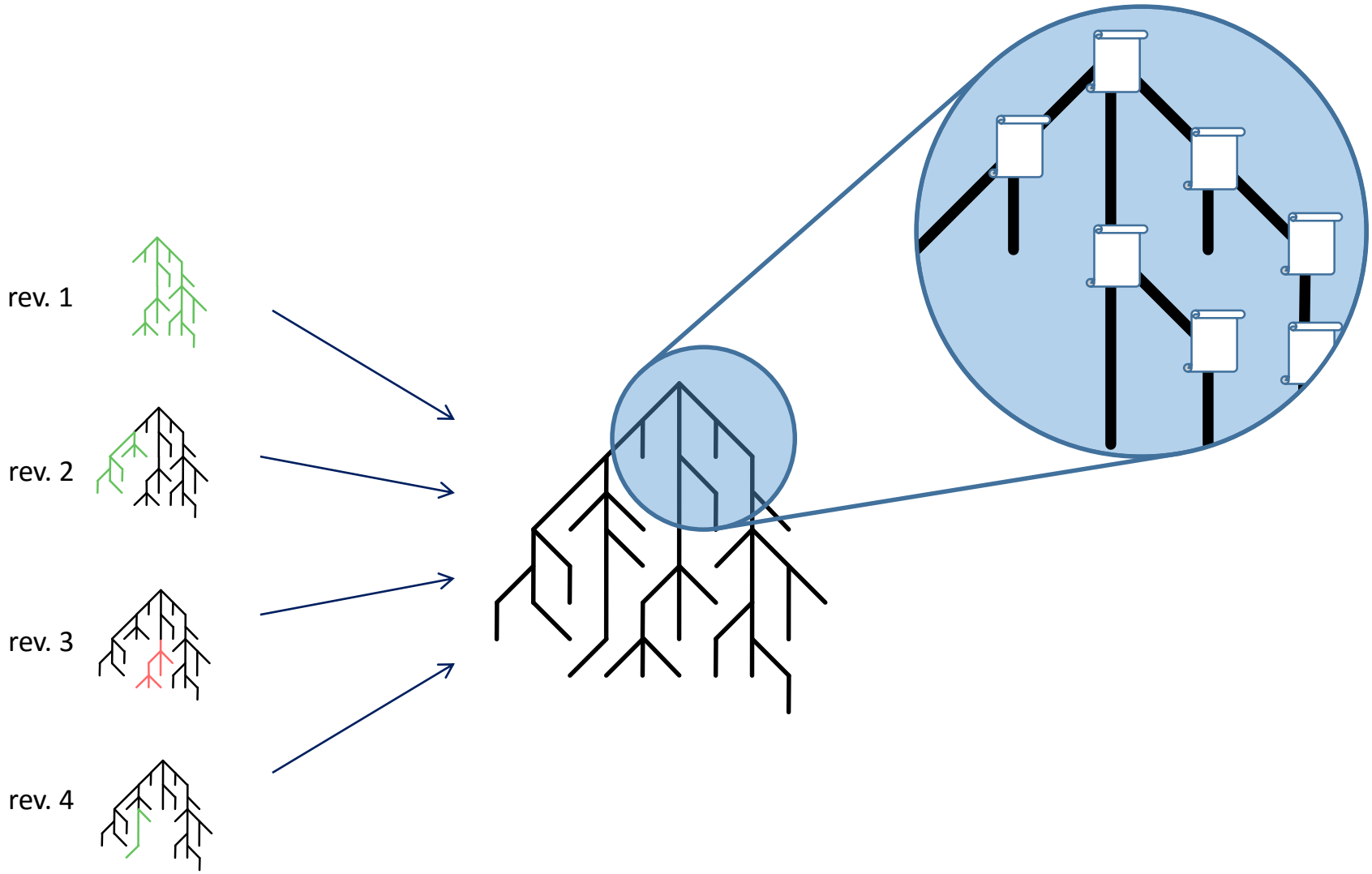
ConditionalExpression

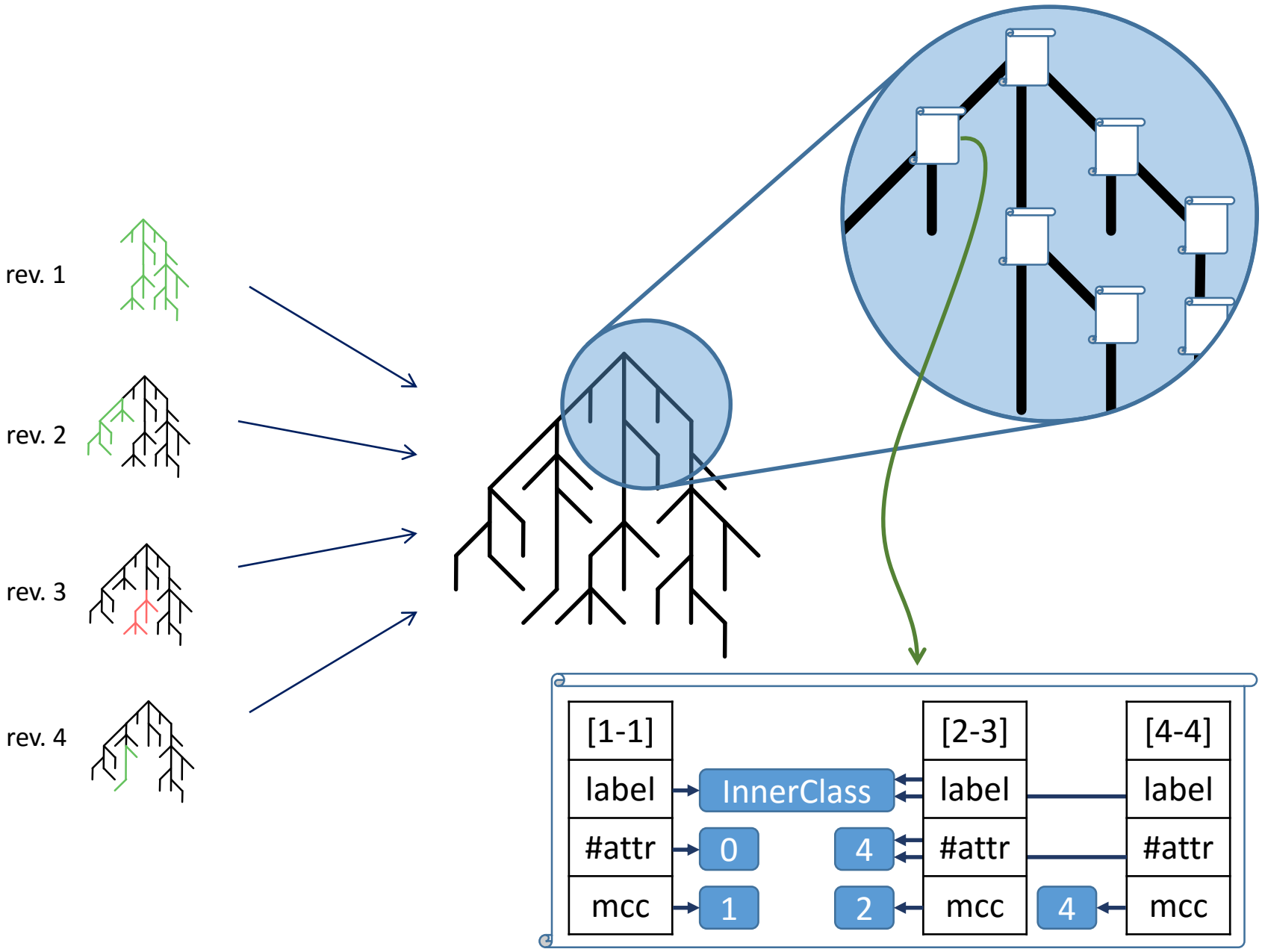
140 AST nodes
(using ANTLR)

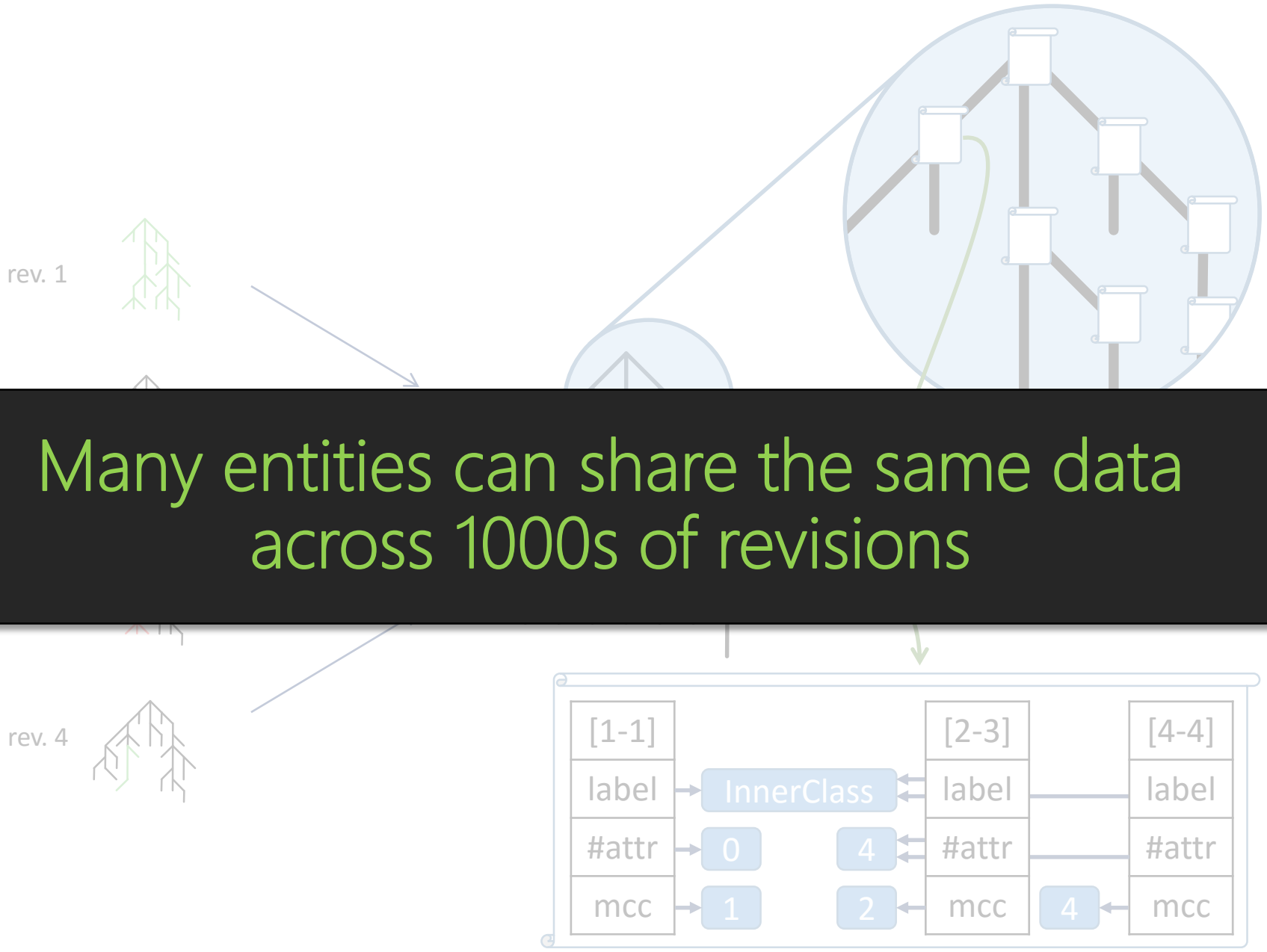
7 AST nodes
(using ANTLR)

#4: Use non-duplicative data structures to store your results


rev. 1
rev. 2
rev. 3
rev. 4


rev. 1
rev. 2
rev. 3
rev. 4


rev. 1
rev. 2
rev. 3
rev. 4
[1-1]
label
#attr
mcc
InnerClass
0
1
[2-3]
label
#attr
mcc
4
2
[4-4]
label
#attr
mcc
4



Many entities can share the same data across 1000s of revisions

LISA also does:

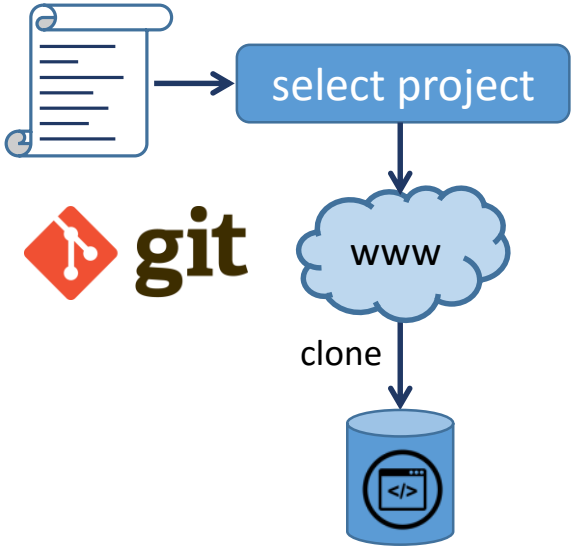
#5: Parallel Parsing

#6: Asynchronous graph computation

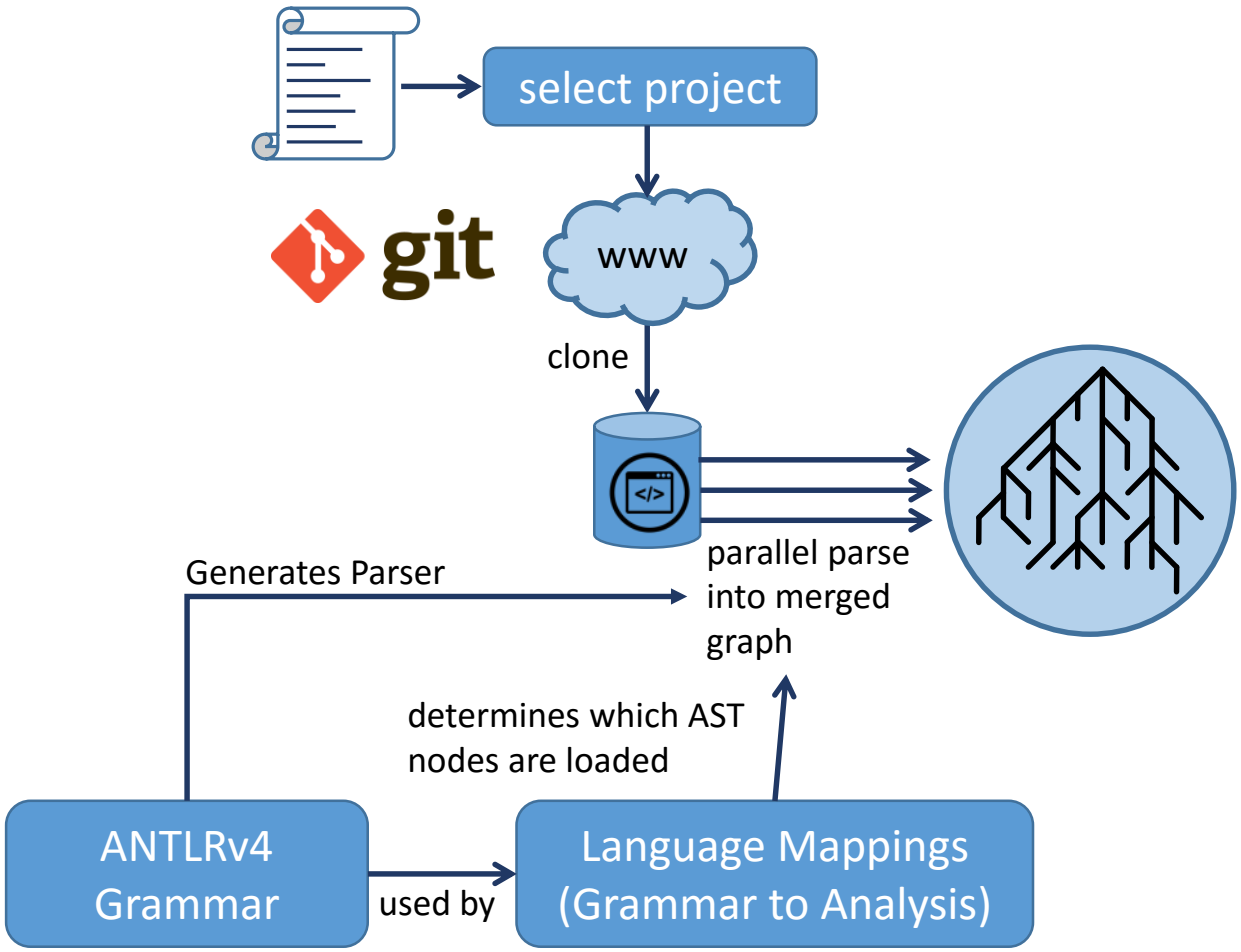
#7: **Generic graph computations** applying to ASTs from **compatible languages**

To Summarize...

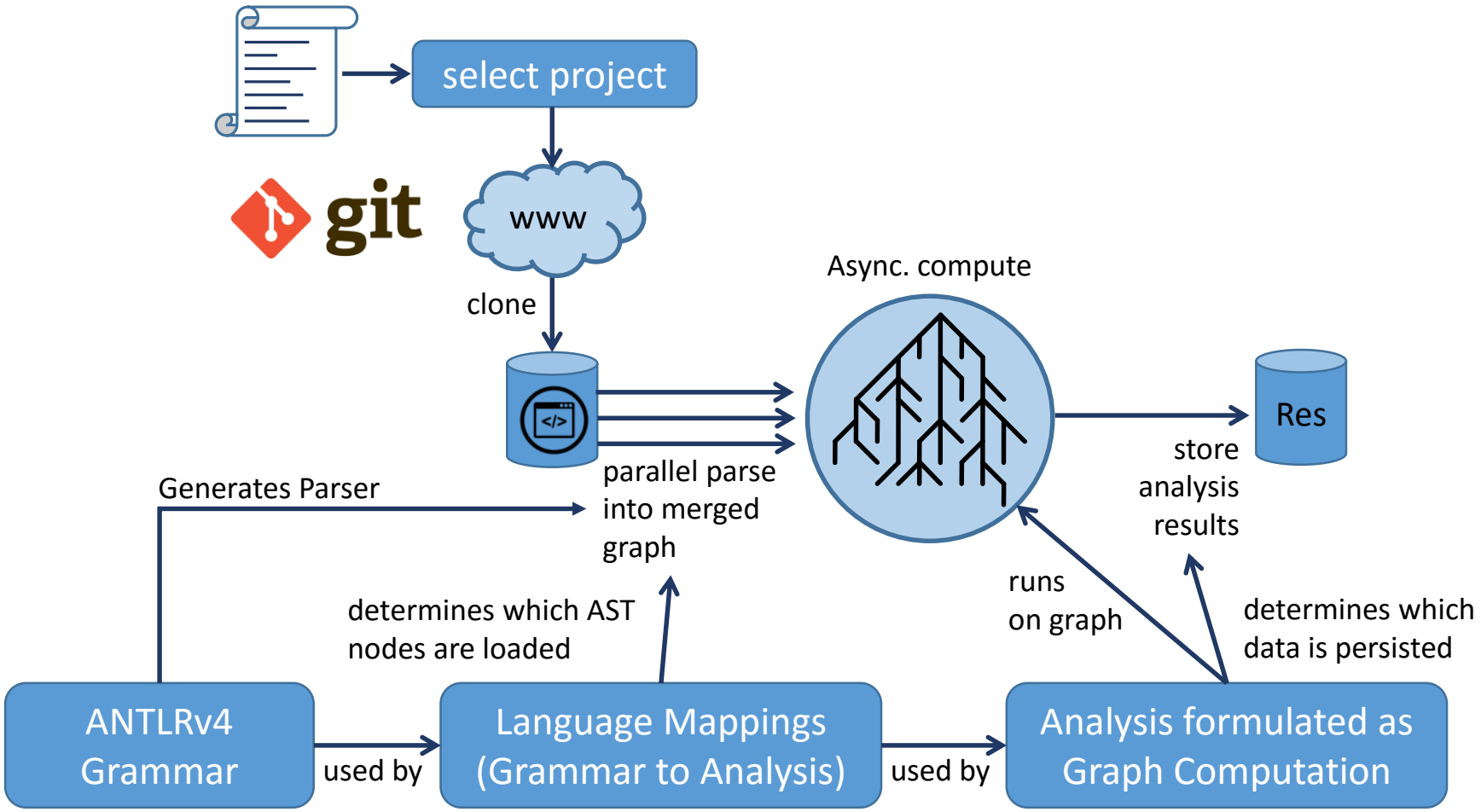
The LISA Analysis Process



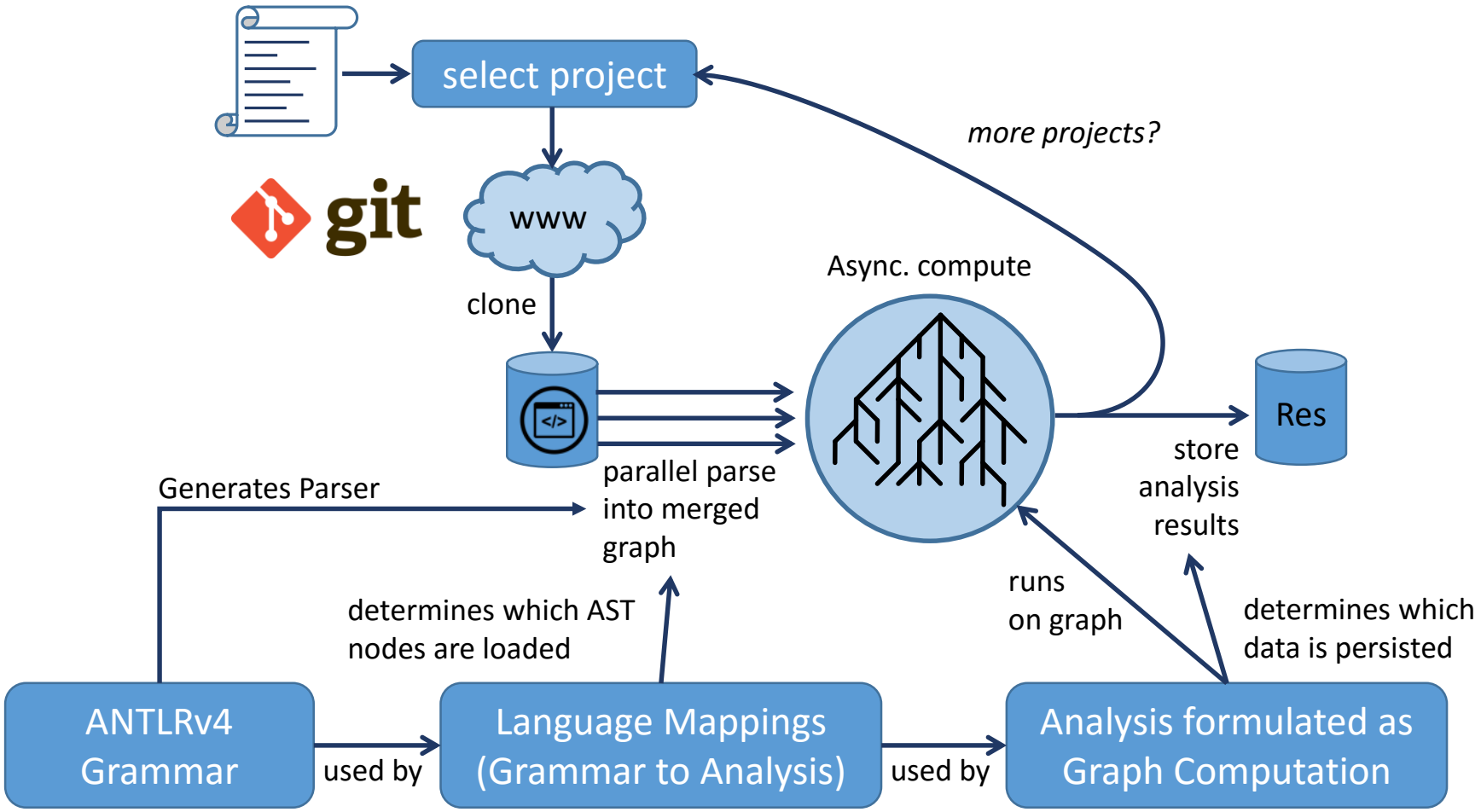
The LISA Analysis Process



The LISA Analysis Process

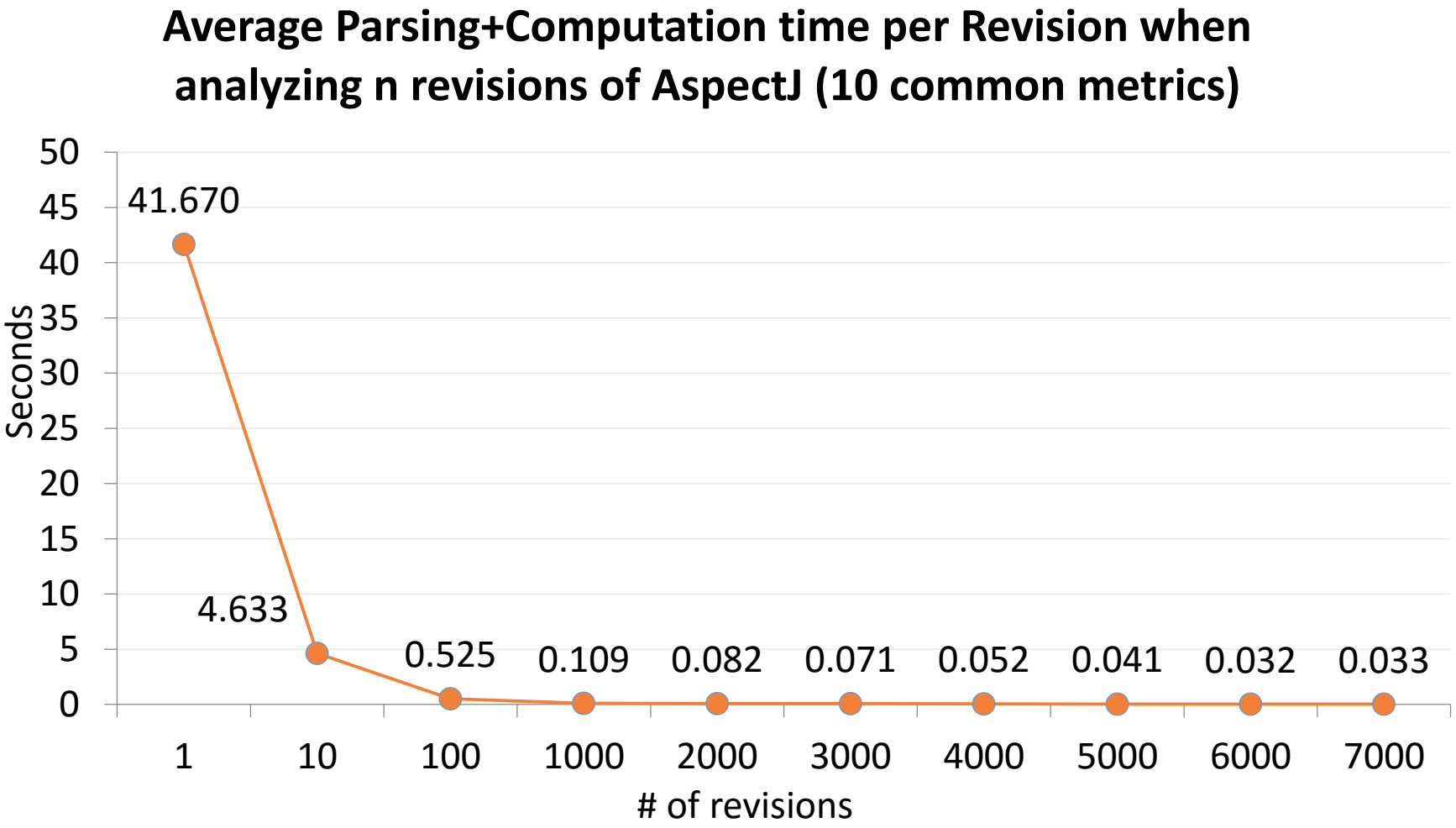


The LISA Analysis Process



How well does it work, then?

Marginal cost for +1 revision



Overall Performance Stats

Language	Java	C#	JavScript
#Projects	100	100	100
#Revisions	646'261	489'764	204'301
#Files (parsed!)	3'235'852	3'234'178	507'612
#Lines (parsed!)	1'370'998'072	961'974'773	194'758'719
Total Runtime (RT)[1]	18:43h	52:12h	29:09h
Median RT[1]	2:15min	4:54min	3:43min
Tot. Avg. RT per Rev.[2]	84ms	401ms	531ms
Med. Avg. RT per Rev.[2]	**30ms**	**116ms**	**166ms**

[1] Including cloning and persisting results
[2] Excluding cloning and persisting results

What's the catch?

(There are a few...)

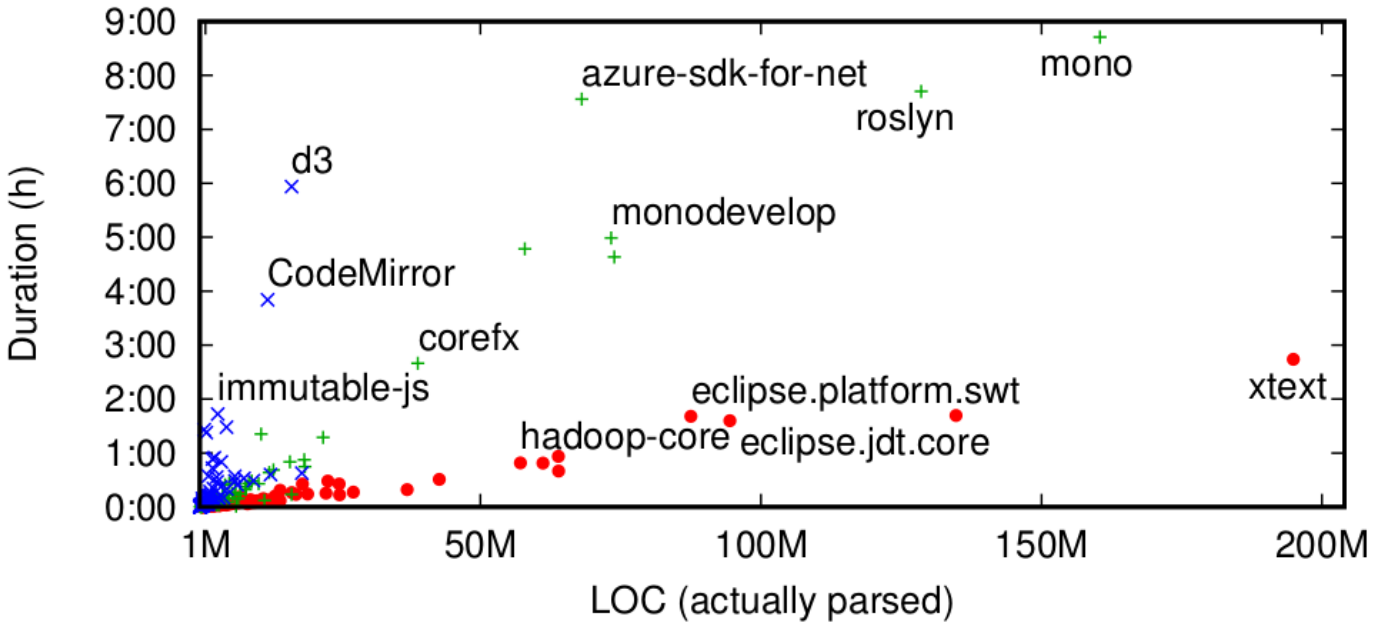
The (not so) minor stuff

- Must implement analyses from scratch
  - No help from a compiler
  - Non-file-local analyses need some effort

The (not so) minor stuff

- Must implement analyses from scratch
  - No help from a compiler
  - Non-file-local analyses need some effort
- Moved files/methods etc. add overhead
  - Uniquely identifying files/entities is hard
  - (No impact on results, though)

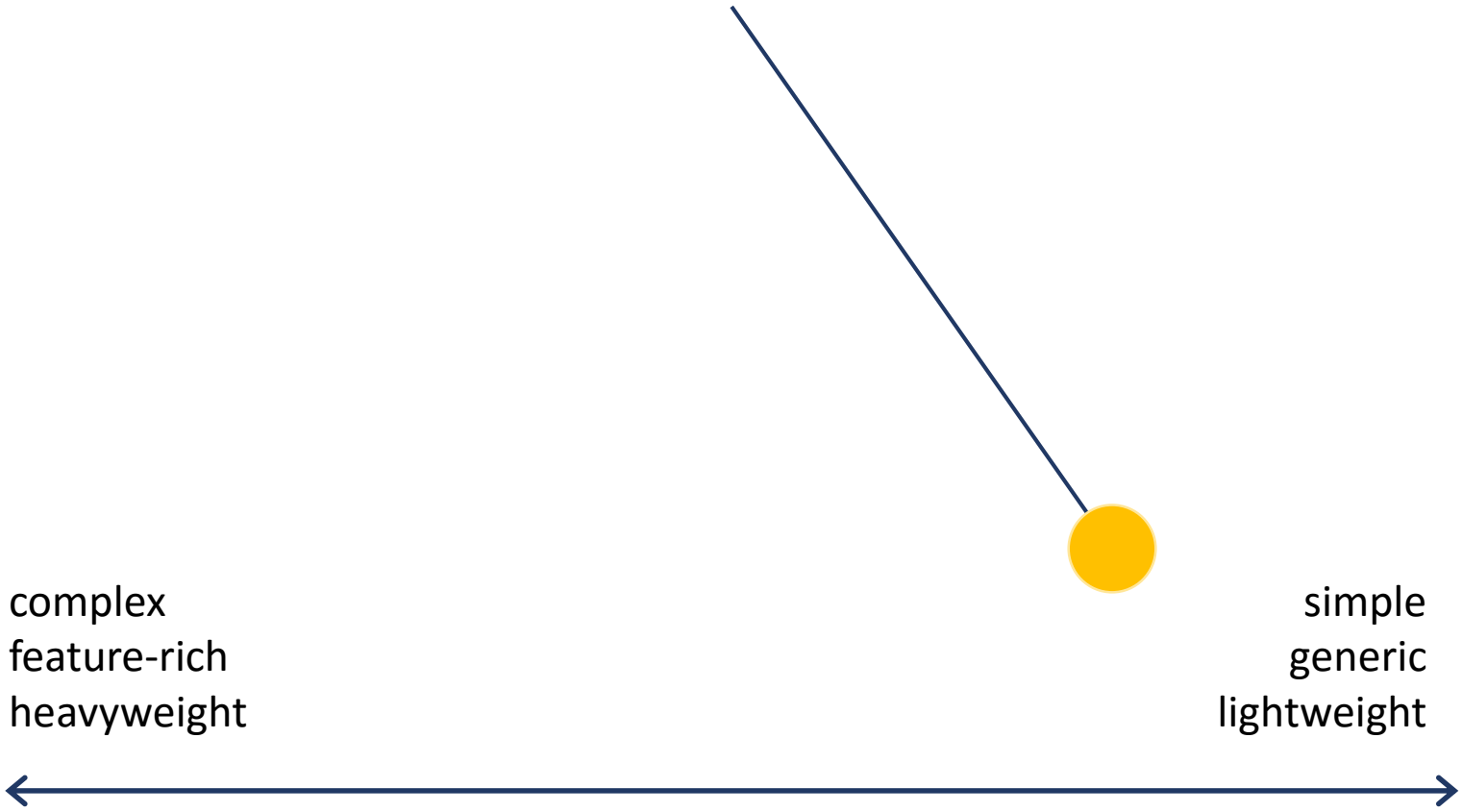
Language matters



E.g.: Javascript takes longer because:

- Larger files, less modularization
- Slower parser (automatic semicolon-insertion)

LISA is EXTREME



University of Zurich[UZH]

s.e.a.l.
software evolution & architecture lab

Thank you for your attention

Read the paper: http://t.uzh.ch/Fj

Try the tool: http://t.uzh.ch/Fk

Get the slides: http://t.uzh.ch/Fm

Contact me: alexandru@ifi.uzh.ch

Parallelize Parsing

Single Git tree traversal

src/Main.java	{1: 1251a4}, {3: fc2452}, {4: 251929}
src/Settings.java	{2: fa255a}
src/Foo.java	{1: 512fc2}, {4: 791c2a}, {5: bcb215}
src/Bar.java	{4: 8a23b2}, {5: b2399f}

Obtain *sequence* of Git blob ids for old versions of each unique path

Parallelize Parsing

Single Git tree traversal

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Obtain *sequence* of Git blob ids for old versions of each unique path

Parse files with different paths in parallel
Some files will have more revisions, taking longer to parse in total
→ Parsing only takes roughly as long as required for the file with the most revisions

Parallelize Parsing

Parallel Parsing from Git is easy and has no overhead

src/Foo.java	{1: 512fc2}, {4: 791c2a}, {5: bcb215}
src/Bar.java	{4: 8a23b2}, {5: b2399f}

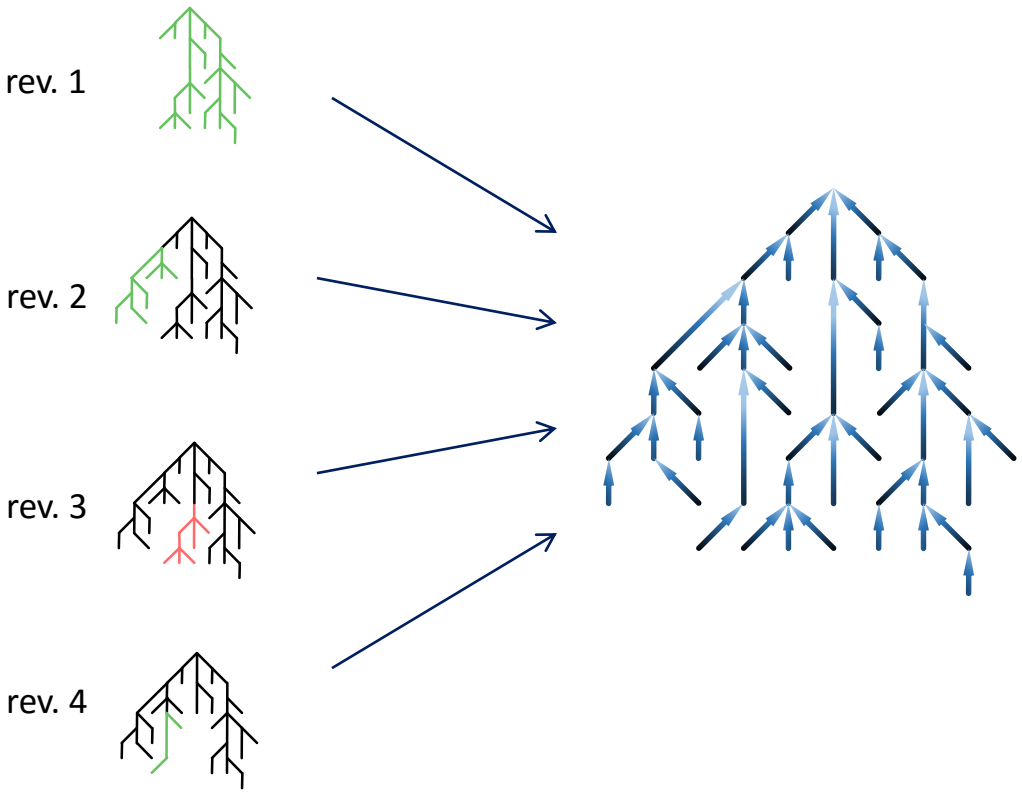
→ Parsing only takes roughly as long as required for the file with the most revisions

Obtain *sequence* of Git blob ids for old versions of each unique path

"Speed-up factor" for each technique

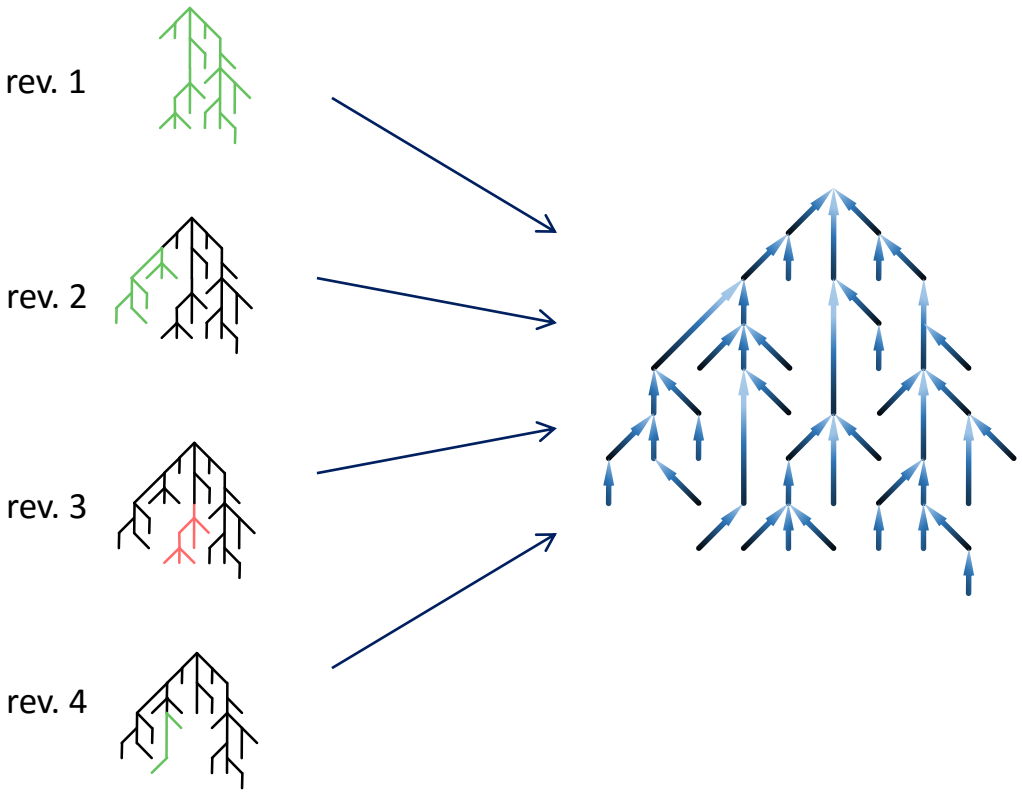
- Parallel parsing: Roughly 2
- Merged ASTs: >1000 for many revisions
- Filtered parsing: >10 during computation, depends on how much is filtered
- All depends on file sizes / parser speed

Asynchronous Graph Analysis



Depending on the node type:
- Signal specfic data

Asynchronous Graph Analysis



Depending on the node type:
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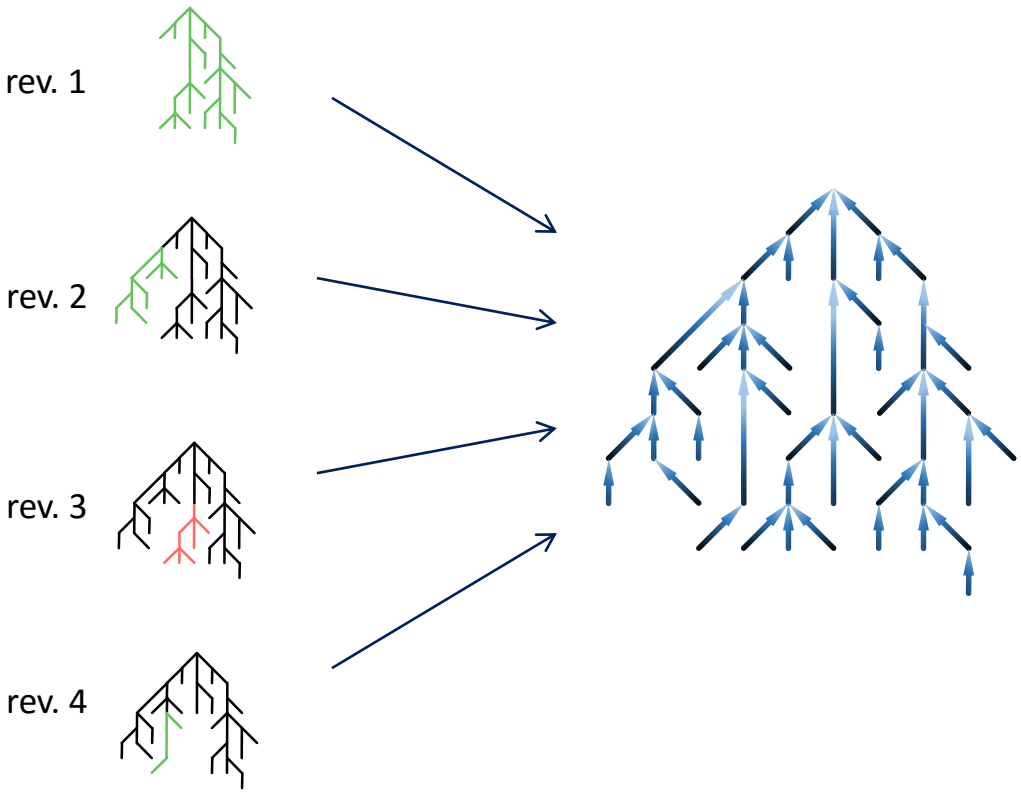
- Collect specific data

Asynchronous Graph Analysis



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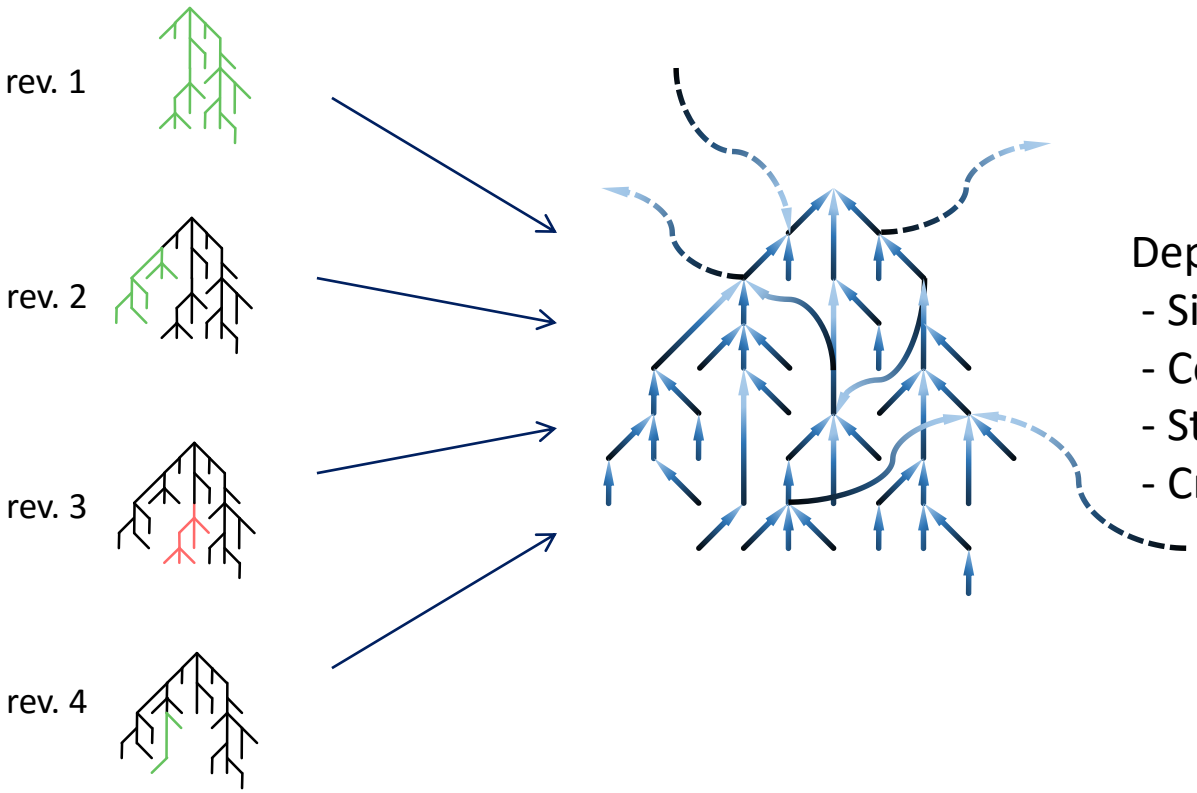

Depending on the node type:
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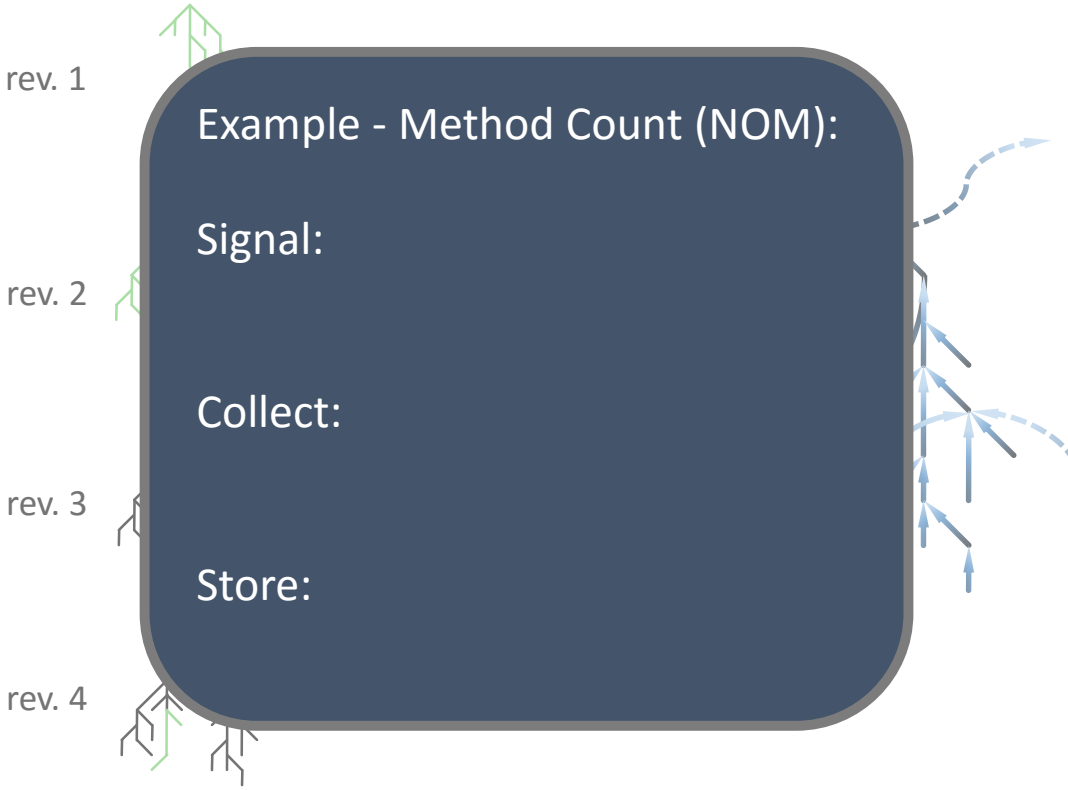
- Create new nodes & edges

Asynchronous Graph Analysis



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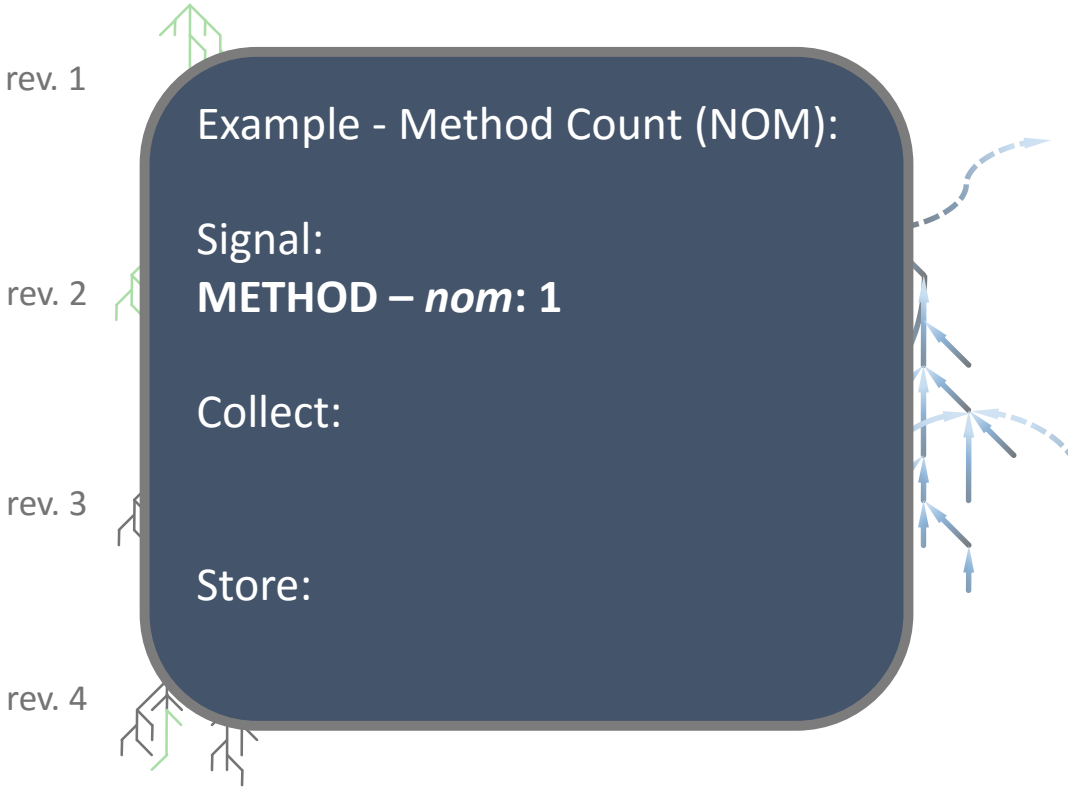

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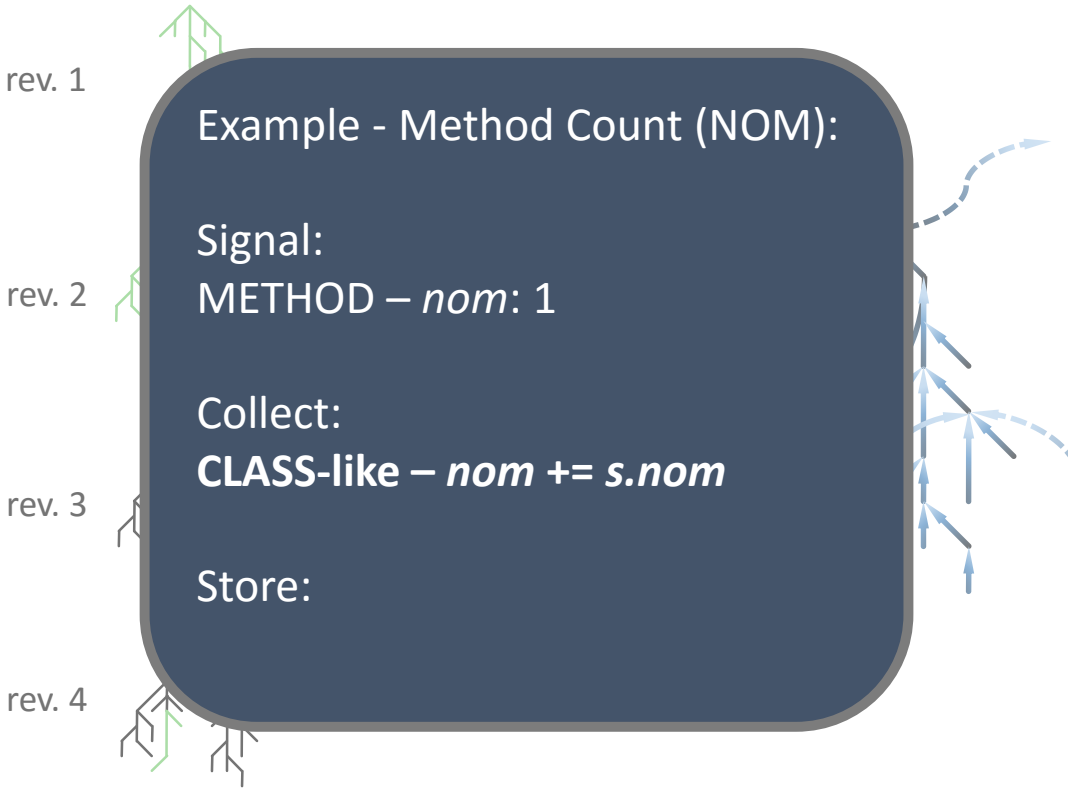

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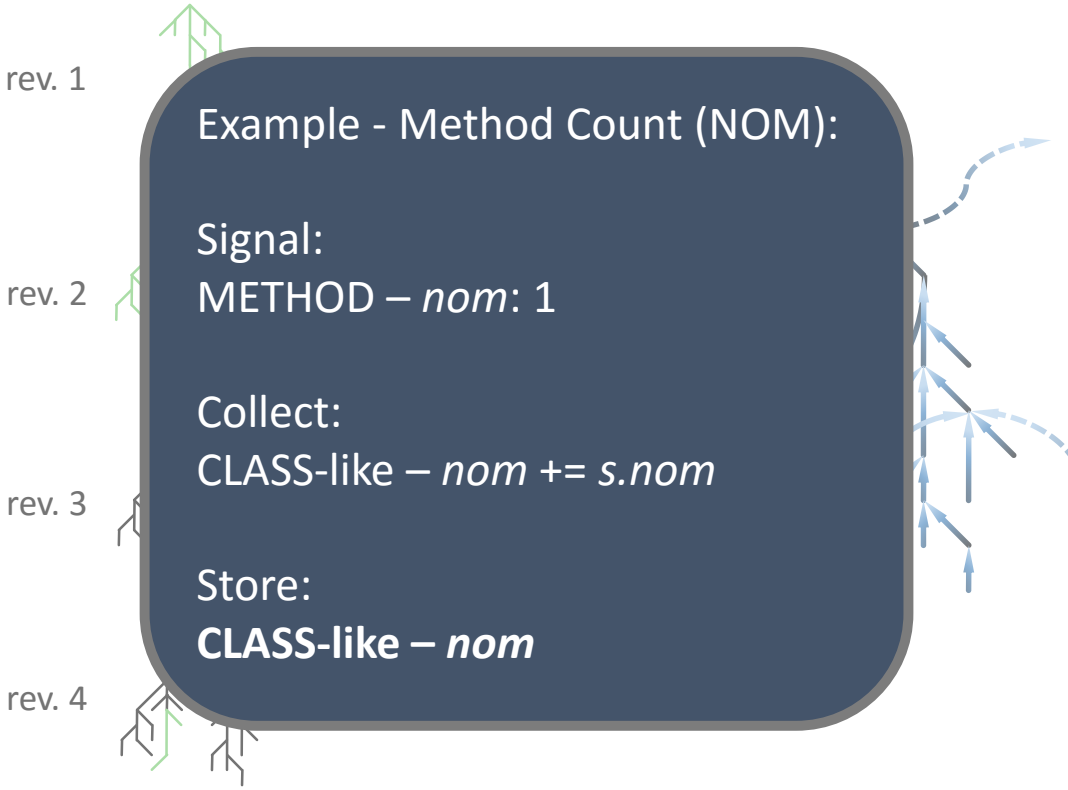

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Static source code analysis?

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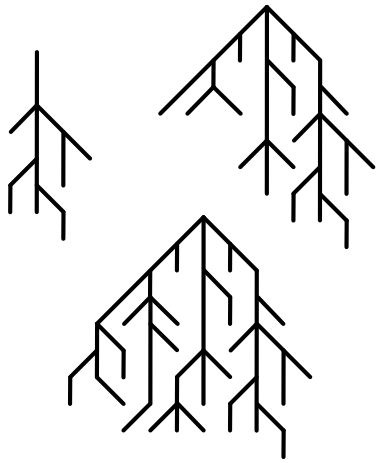
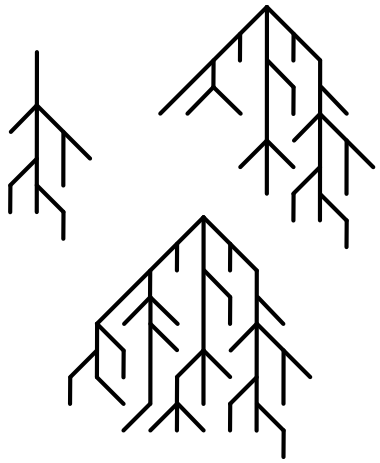


Static source code analysis?

Simple Code Metrics
(NOC, NOM, WMC, Complexity, ...)



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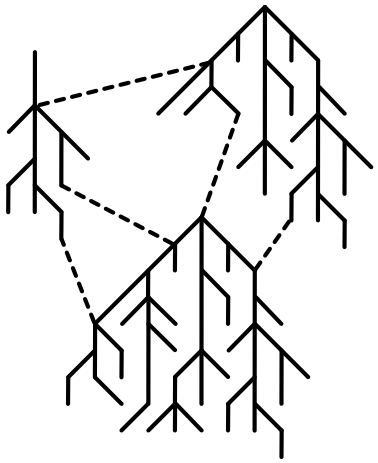
(Coupling, Inheritance, ...)



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AMW	5.45
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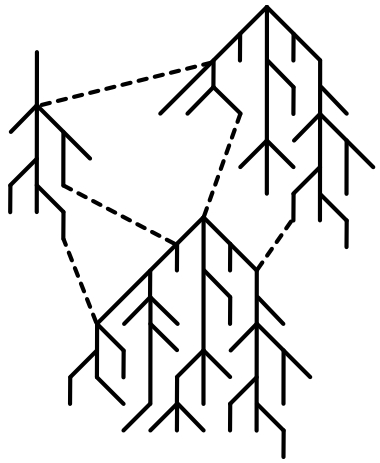

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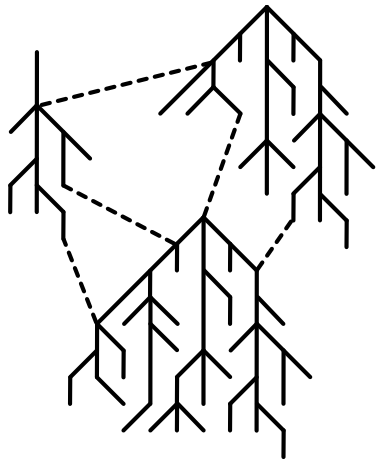

Practice
code smells
refactoring advice
hot-spot detection
bug prediction

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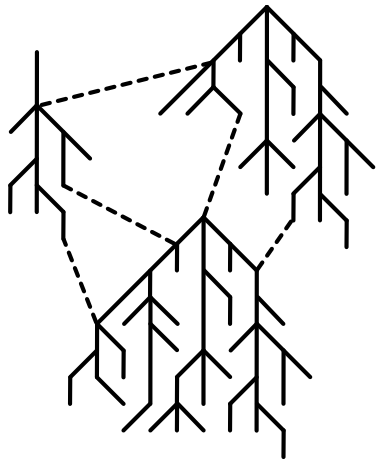

Practice	**Research**
code smells	understanding software evolution
refactoring advice	identifying patterns &
hot-spot detection	anti-patterns
bug prediction	code quality assessment techniques
	...
	→ code studies

(Many) existing studies

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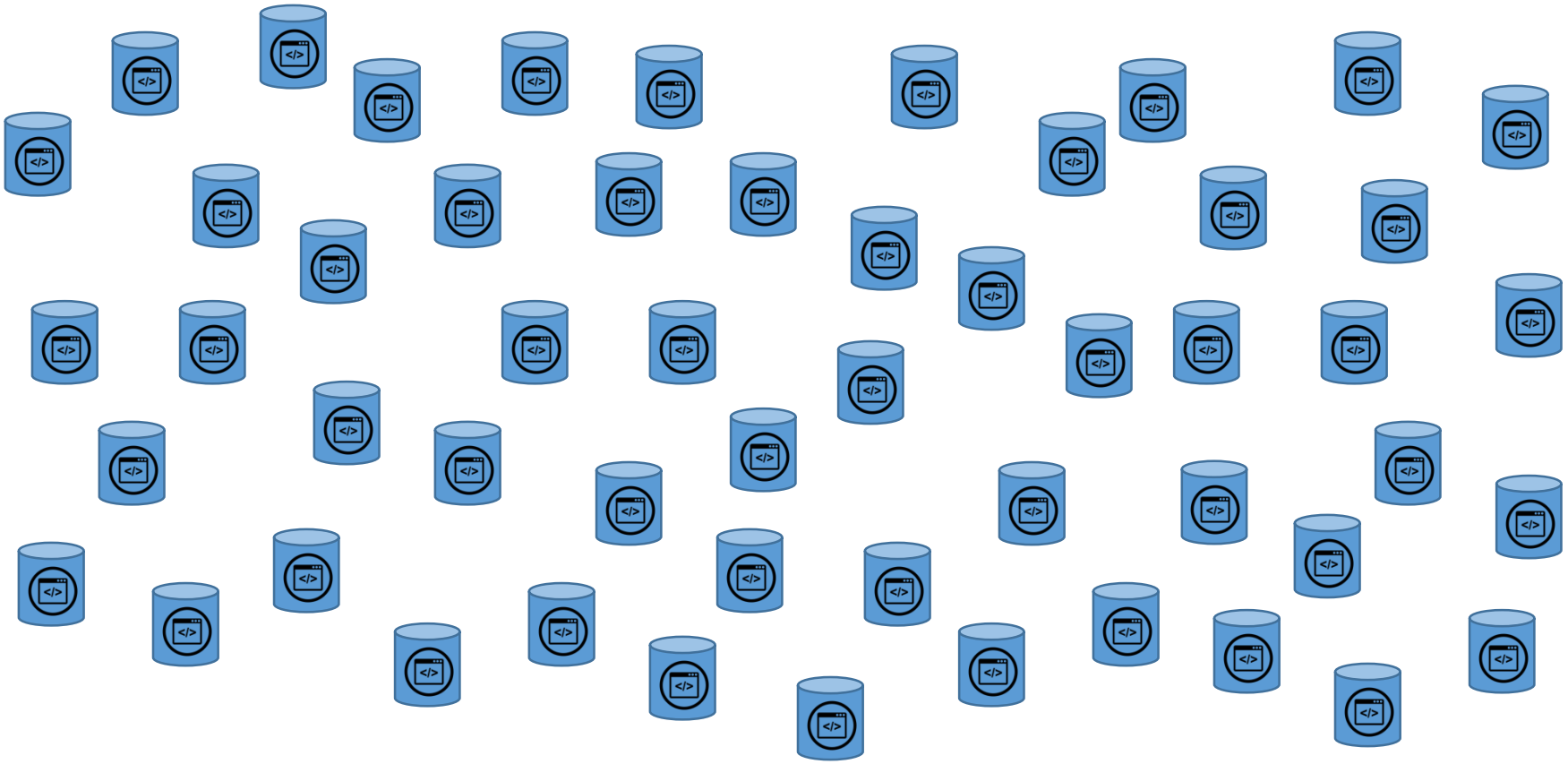
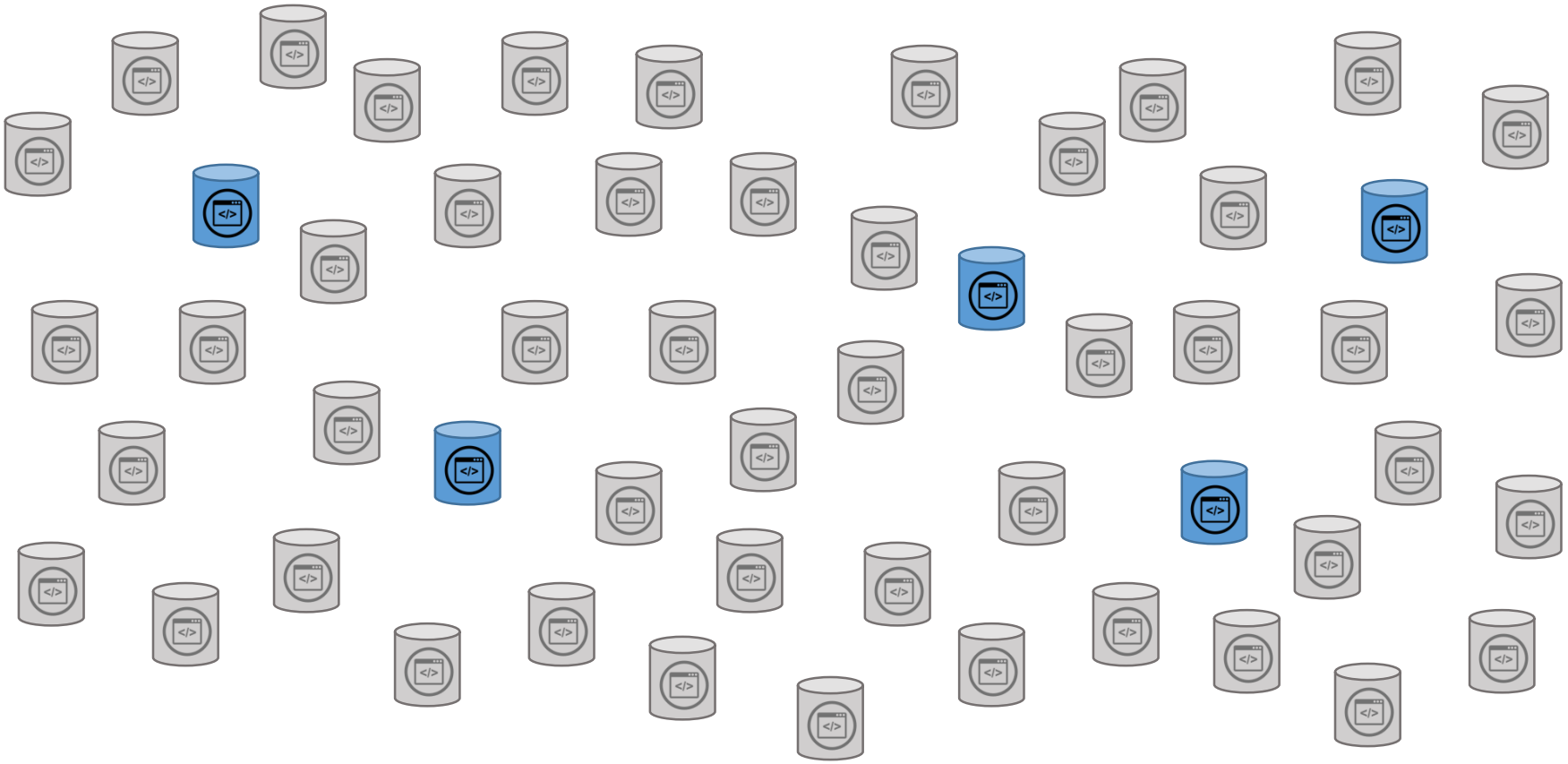


(Many) existing studies

- investigate a small number of projects



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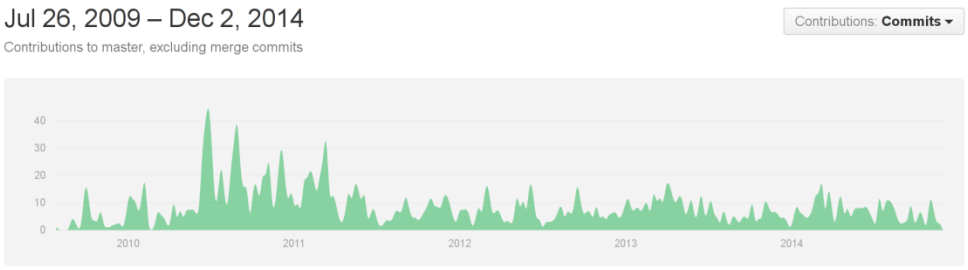
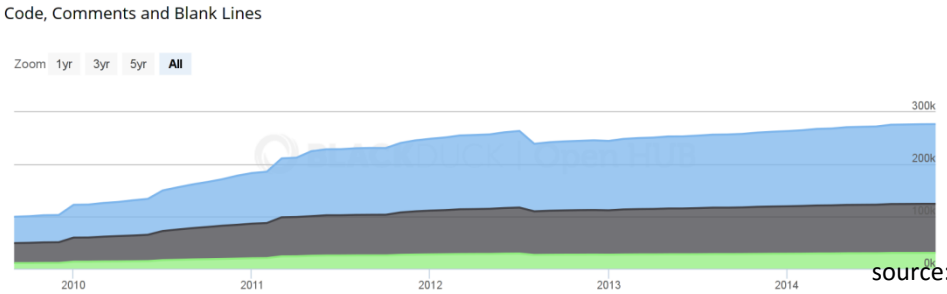


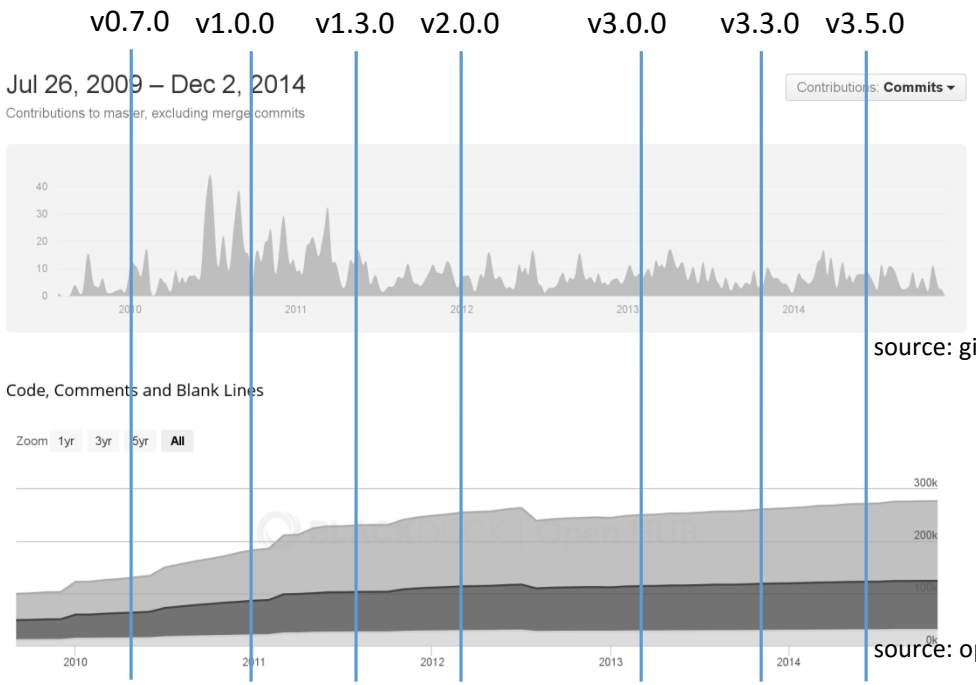
source: github.com



source: openhub.net

(Many) existing studies

- investigate a small number of projects
- analyze a few snapshots of multi-year projects



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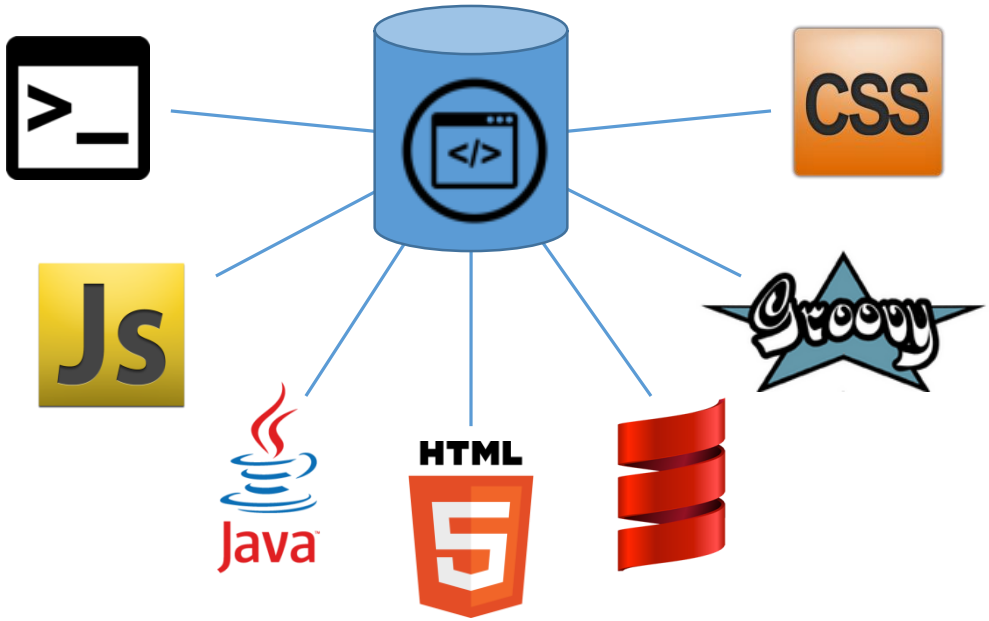
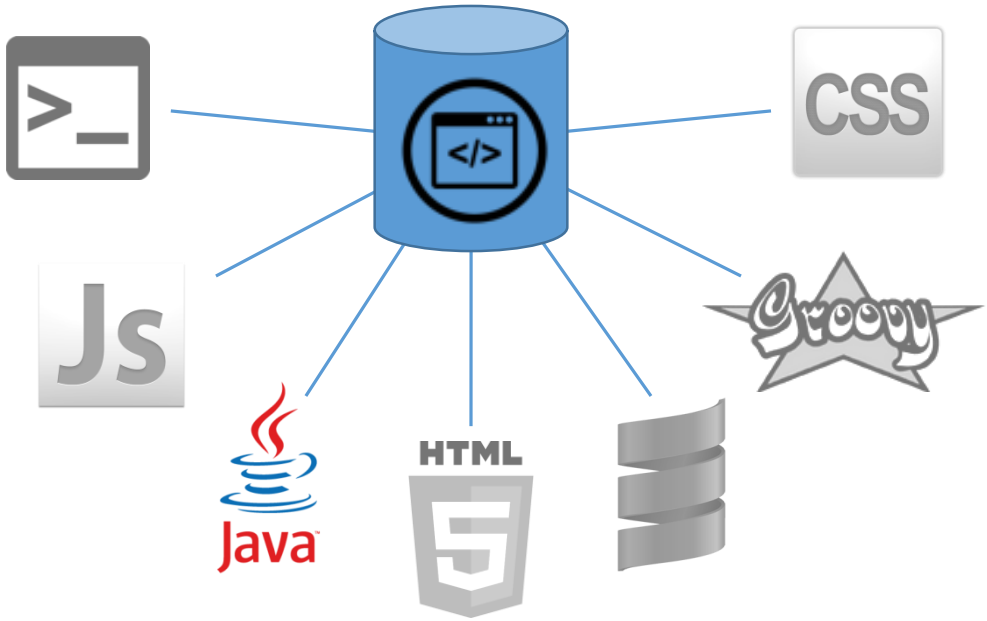


(Many) existing studies

- investigate a small number of projects
- analyze a few snapshots of multi-year projects
- focus on very few programming languages



Why?

Only a few projects

Only a few snapshots

Only few Languages

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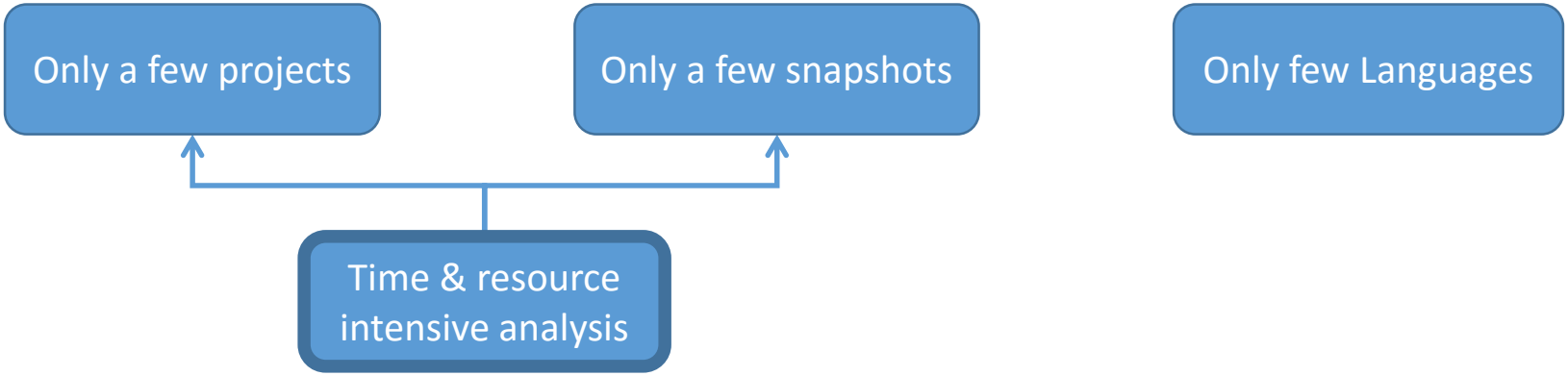
Only a few projects
Only a few snapshots
Only few Languages
Time & resource intensive analysis

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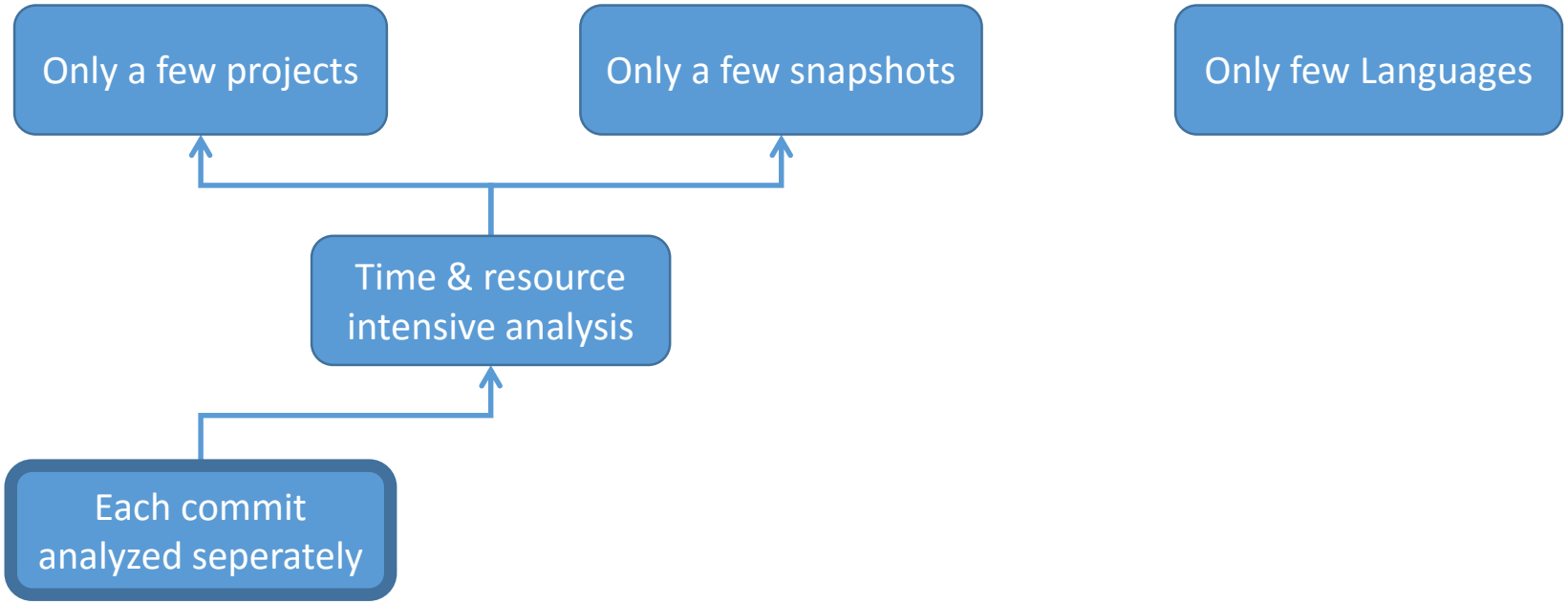
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Each commit analyzed seperately

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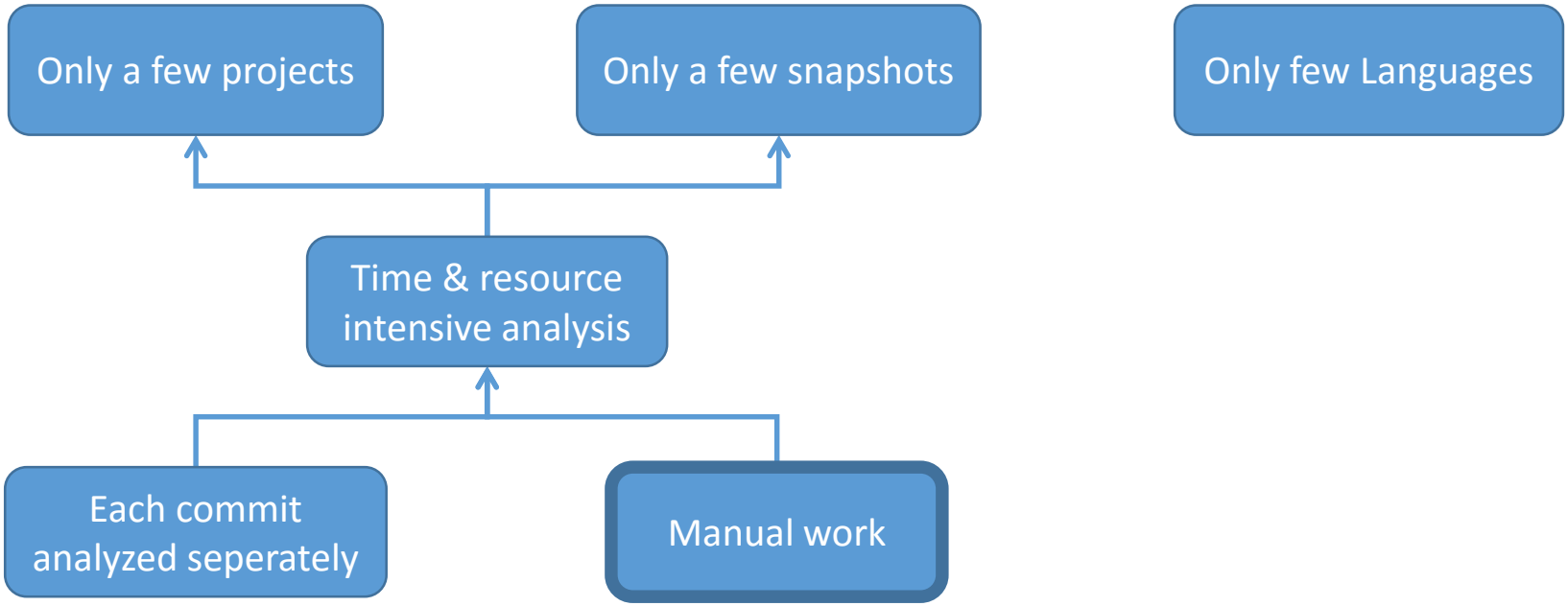
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Manual work

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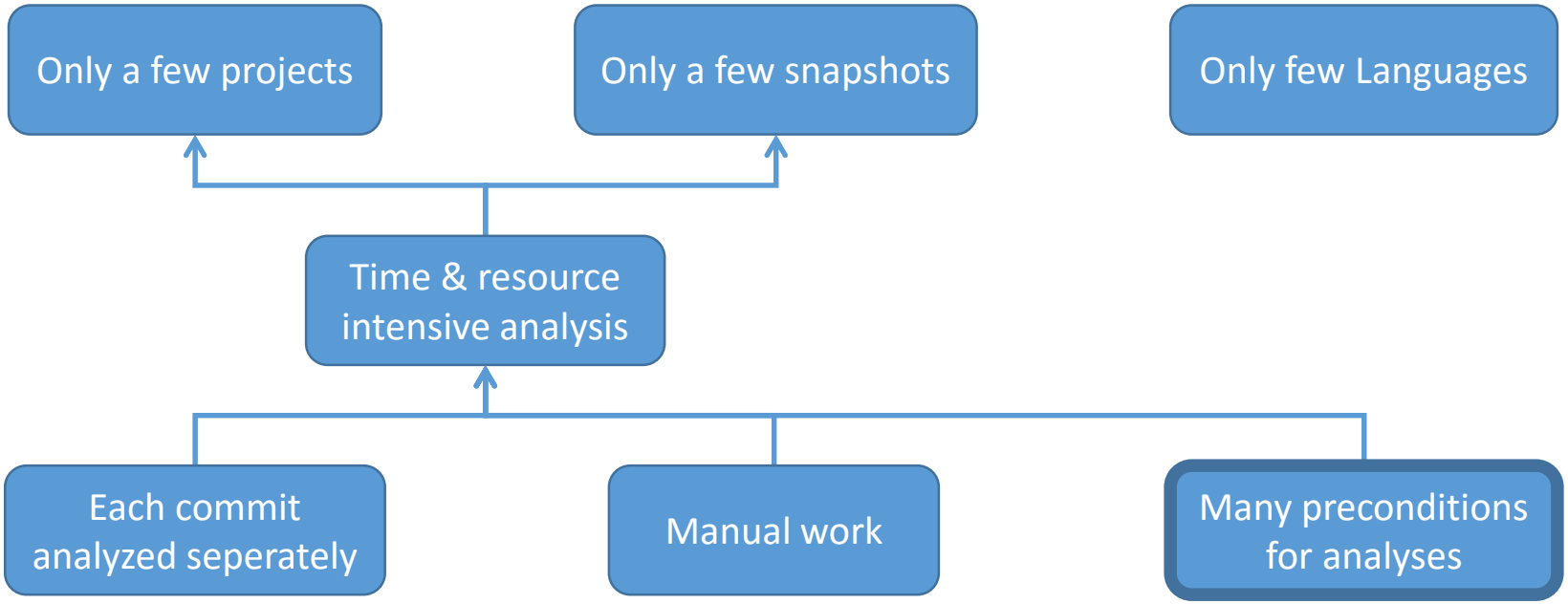
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Manual work
Many preconditions for analyses

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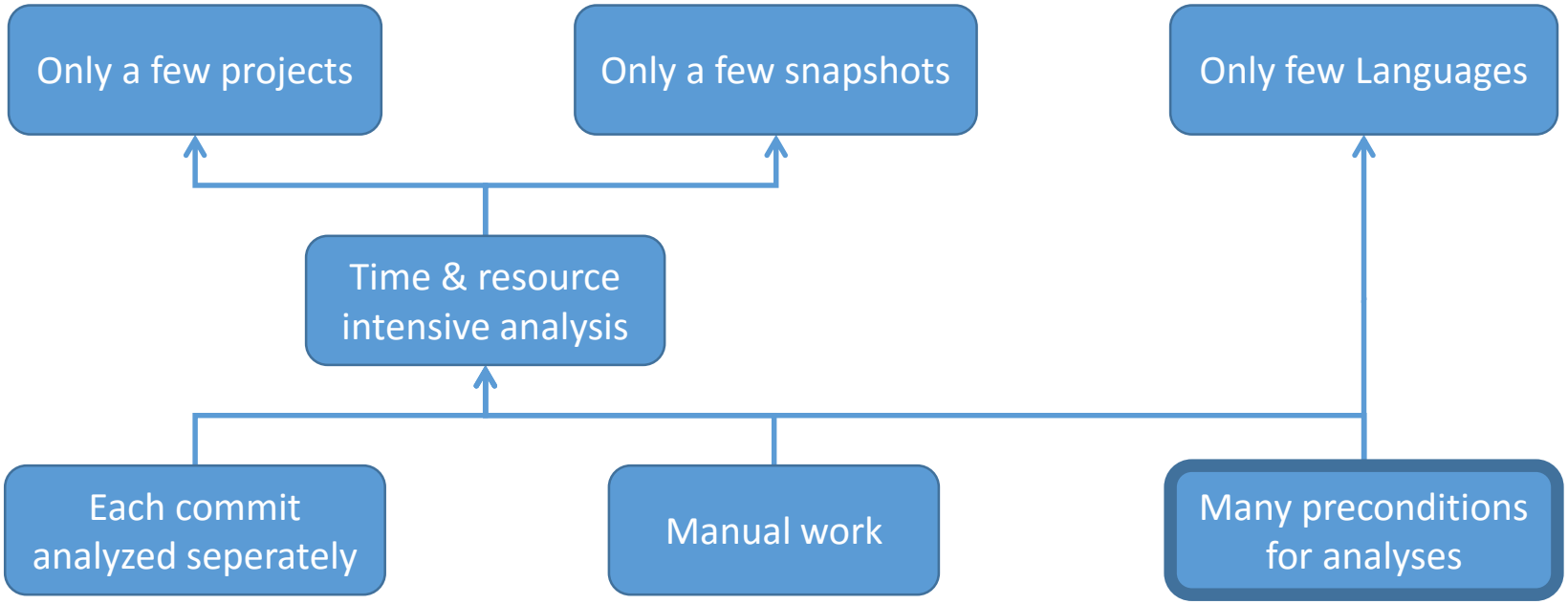
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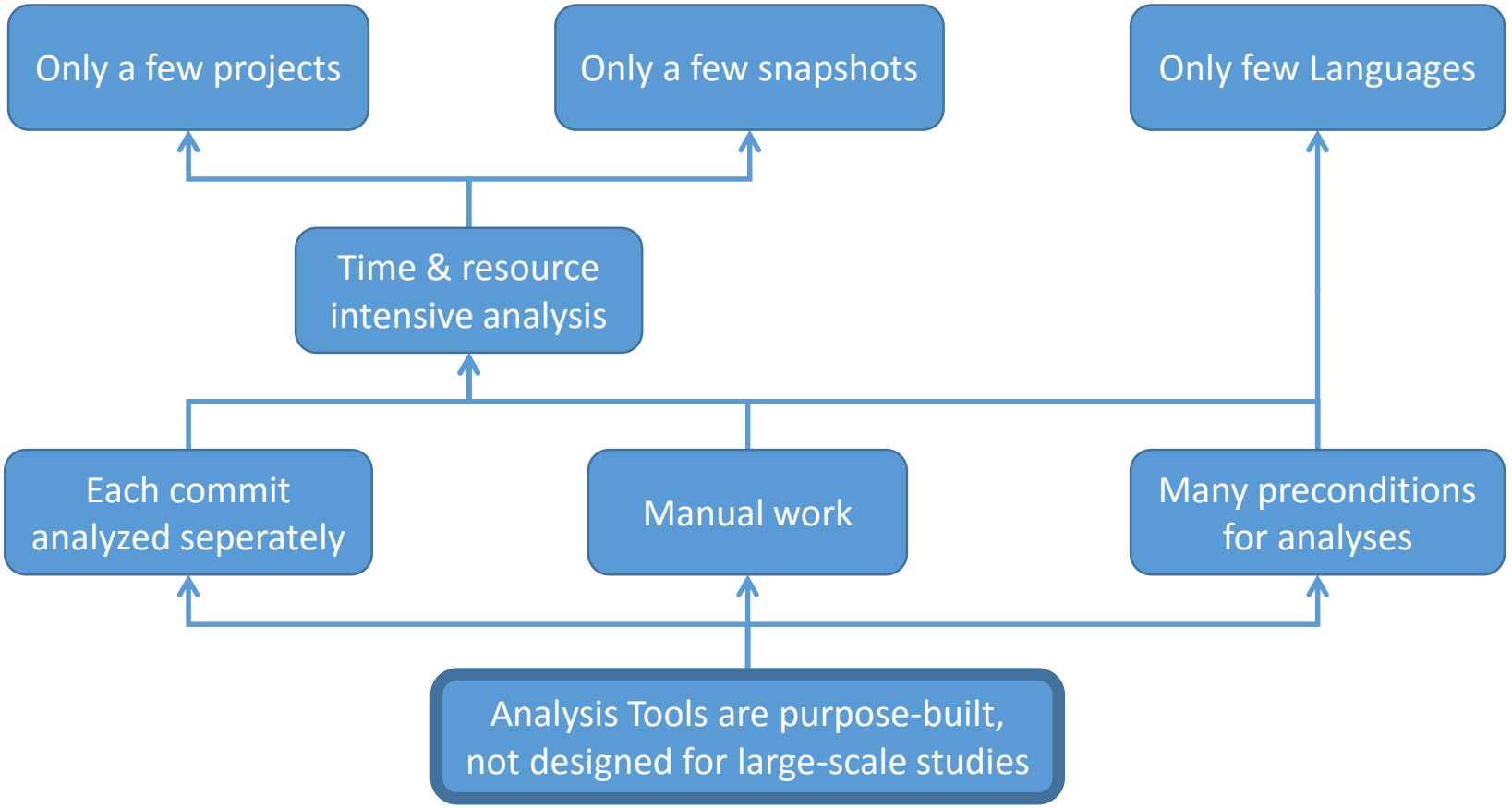
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Analysis Tools are purpose-built, not designed for large-scale studies

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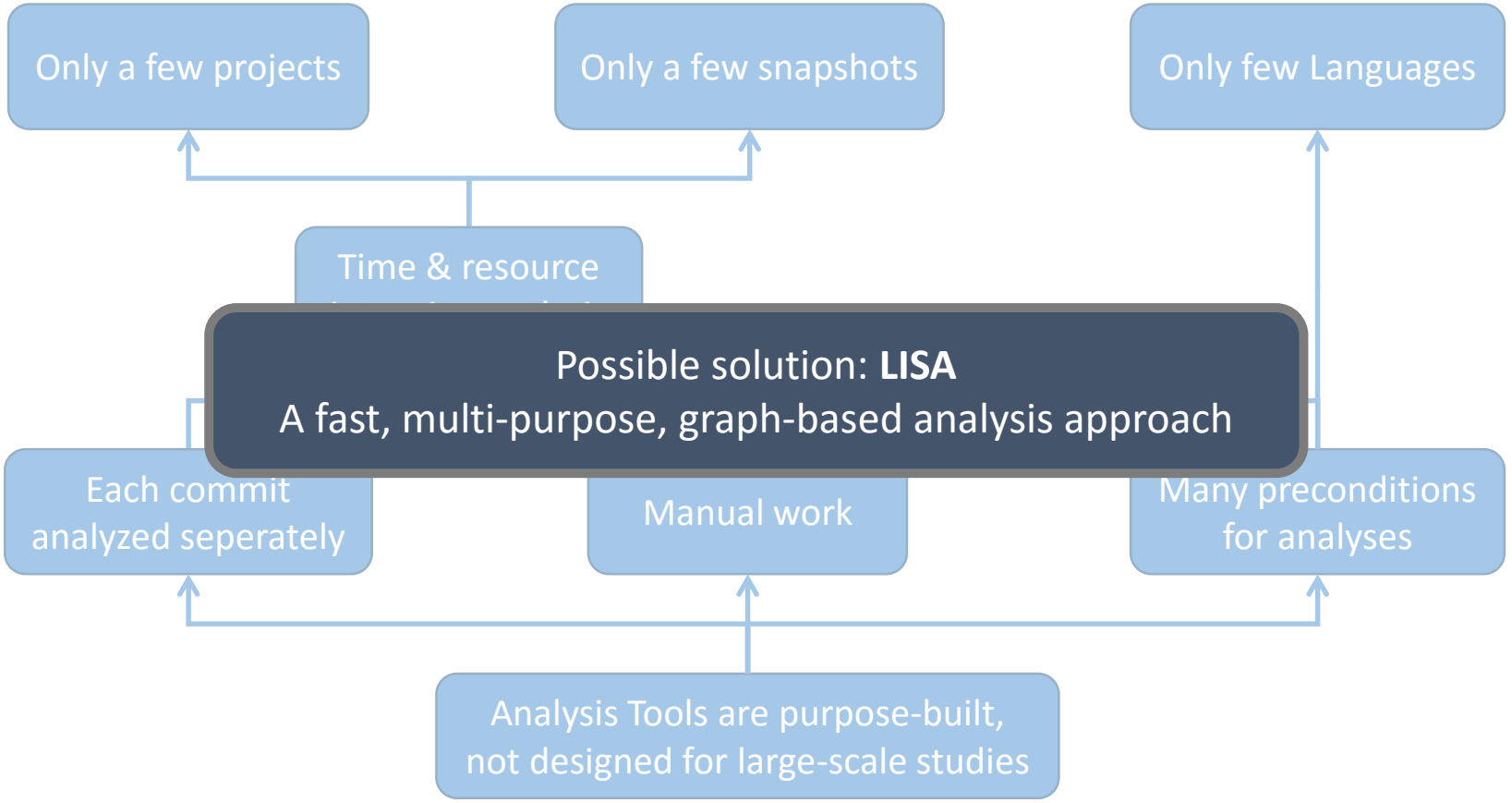
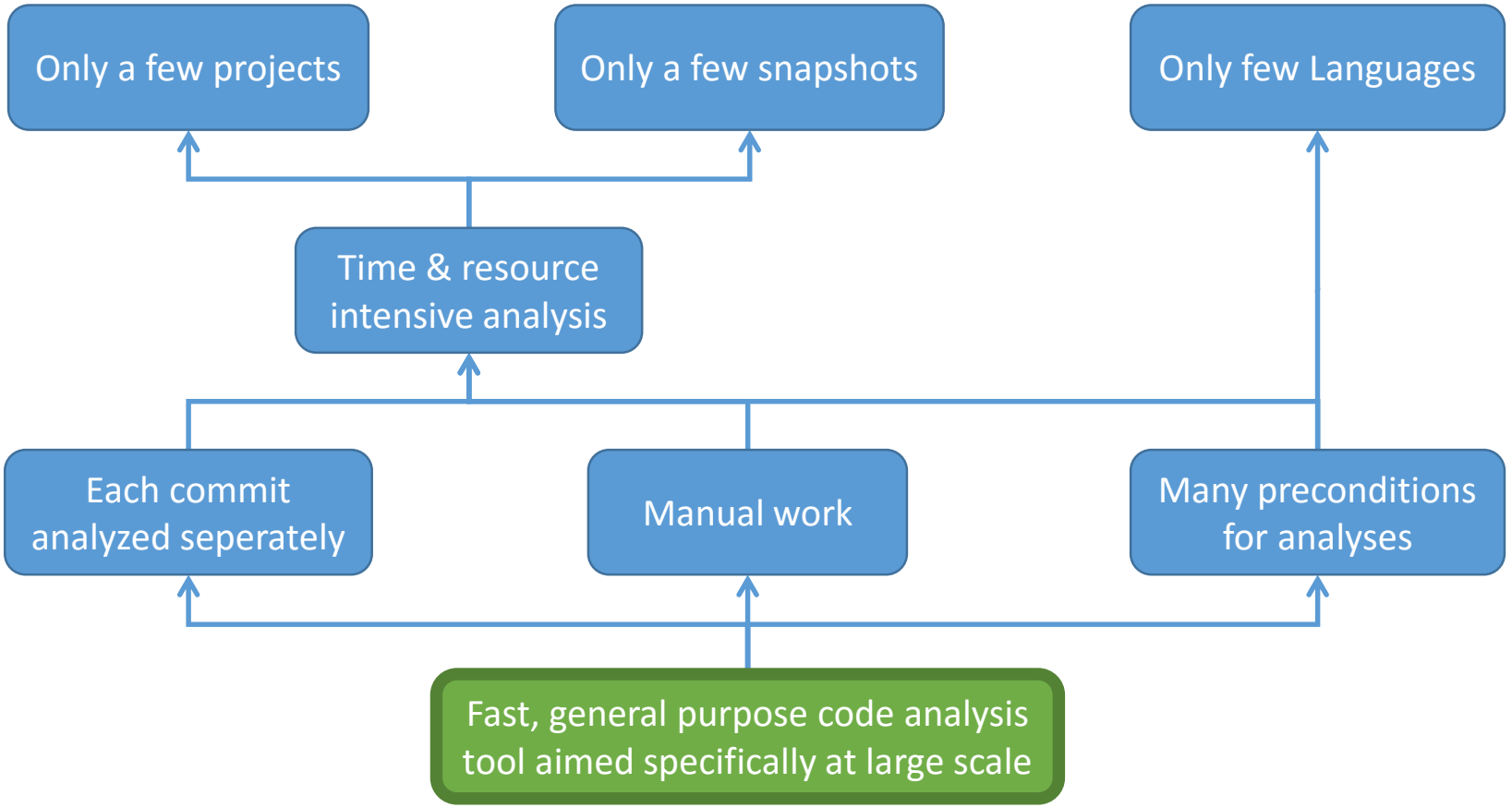

Only a few projects
Only a few snapshots
Only few Languages
Time & resource
Possible solution: **LISA**
A fast, multi-purpose, graph-based analysis approach
Each commit analyzed seperately
Manual work
Many preconditions for analyses
Analysis Tools are purpose-built, not designed for large-scale studies

Rapid Analysis using LISA



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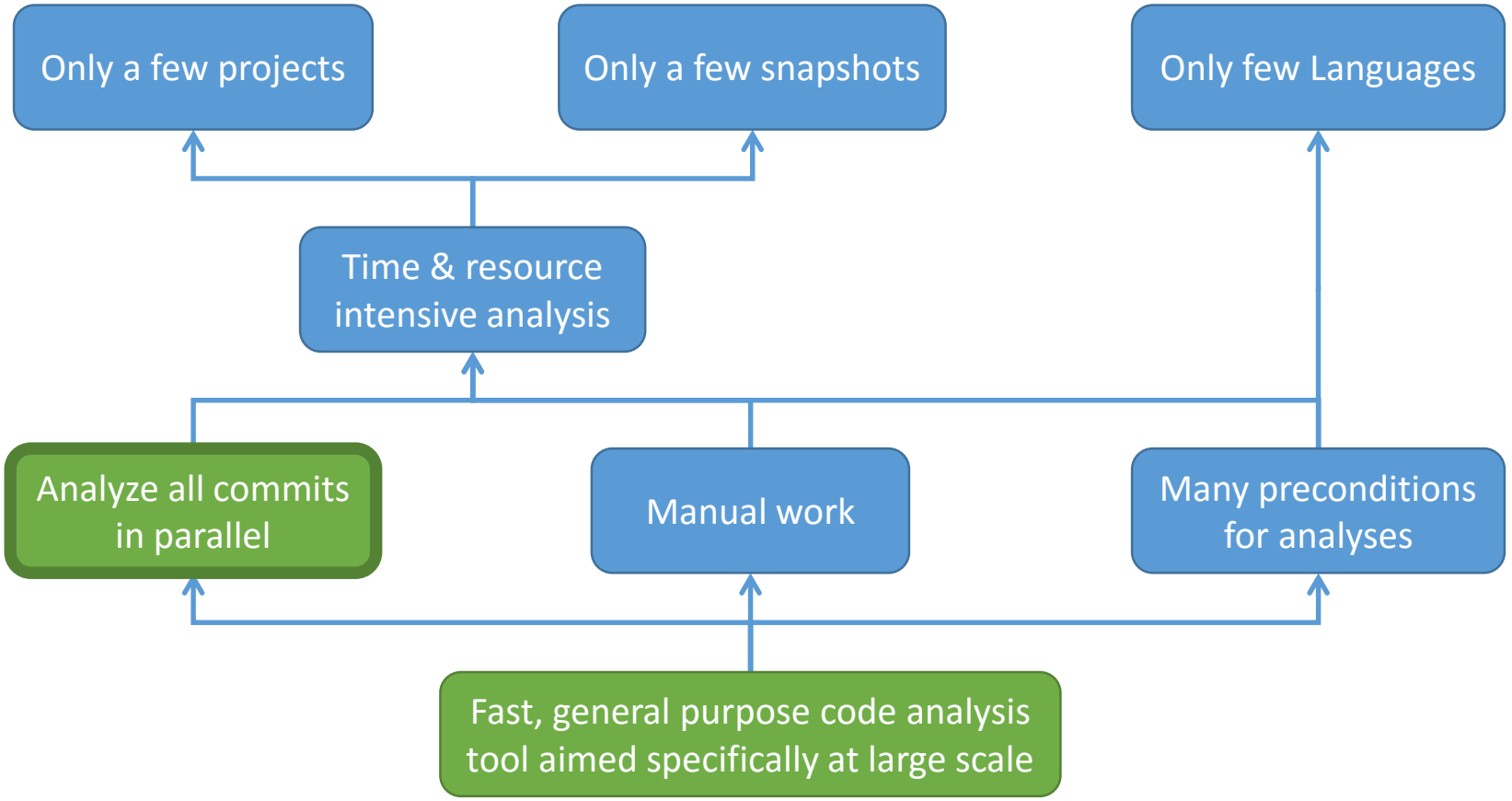

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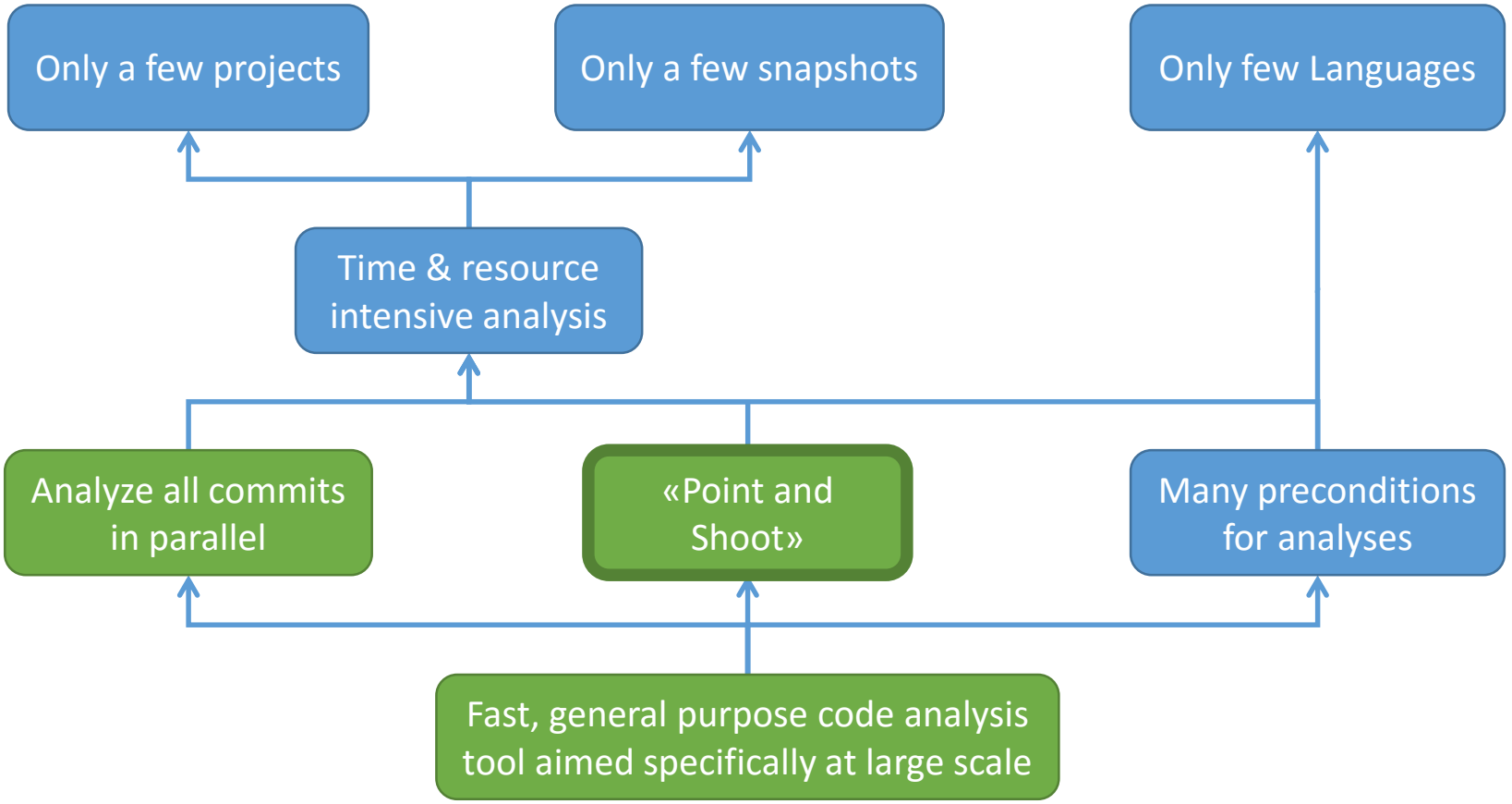

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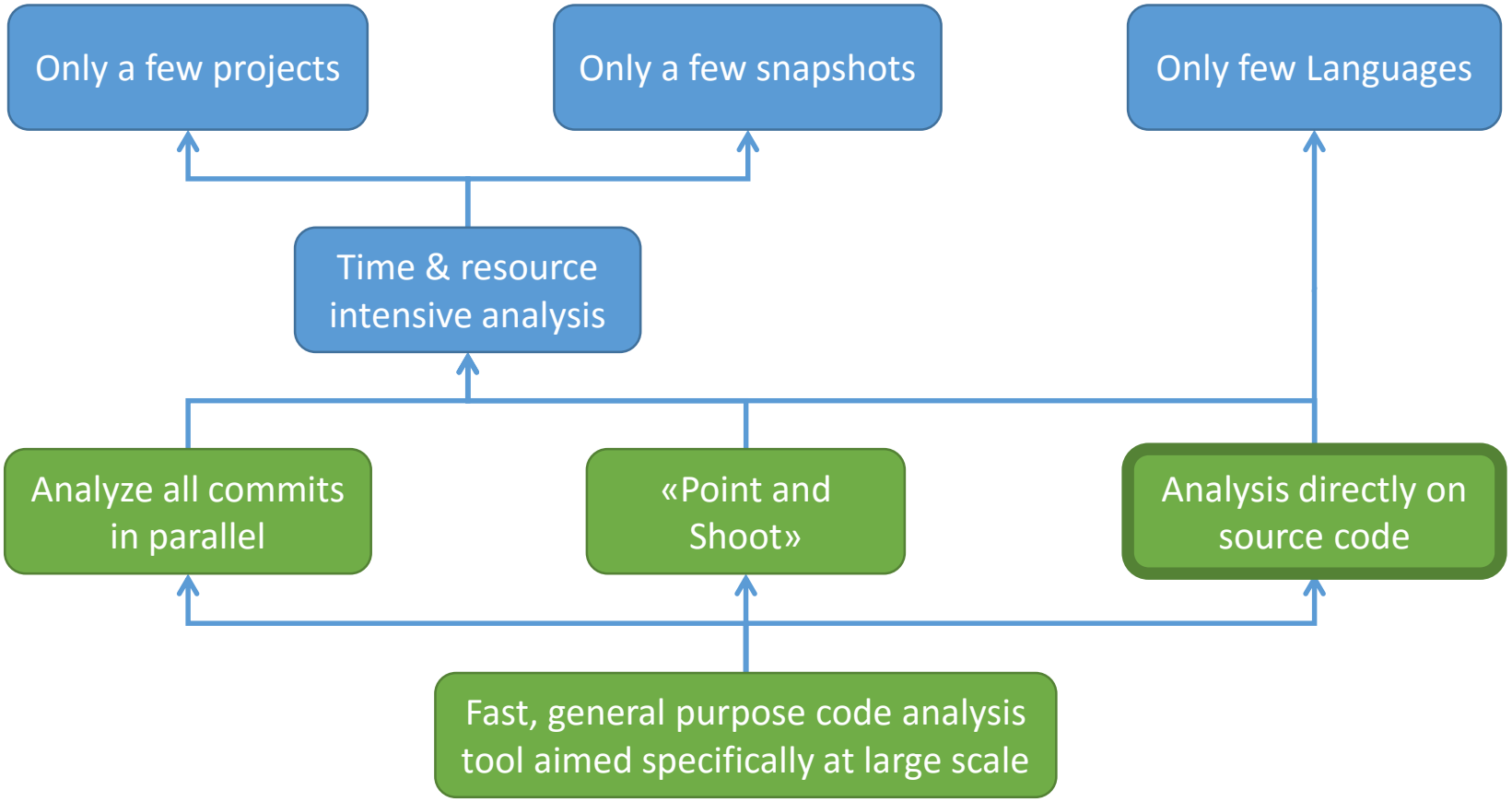

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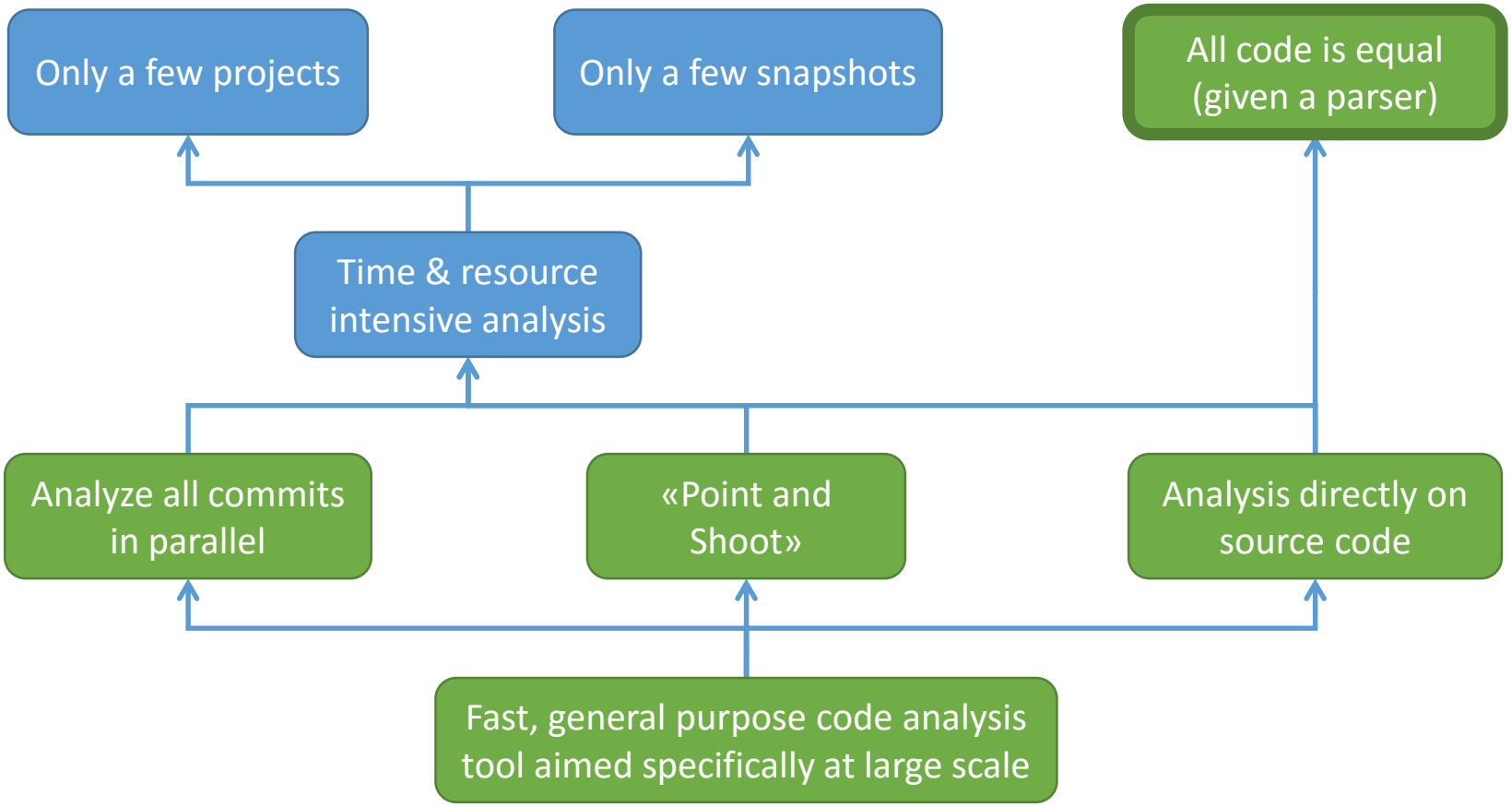

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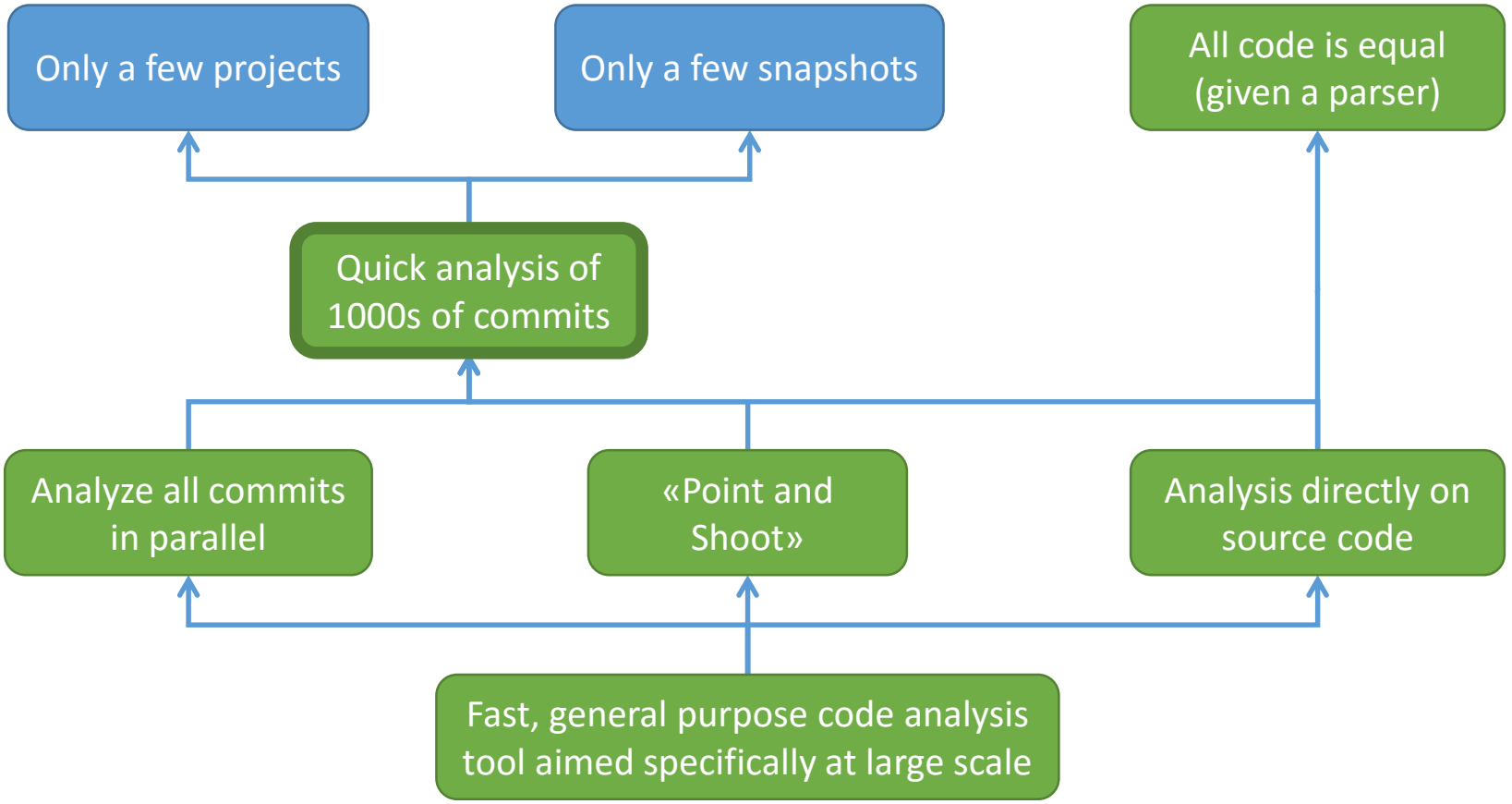

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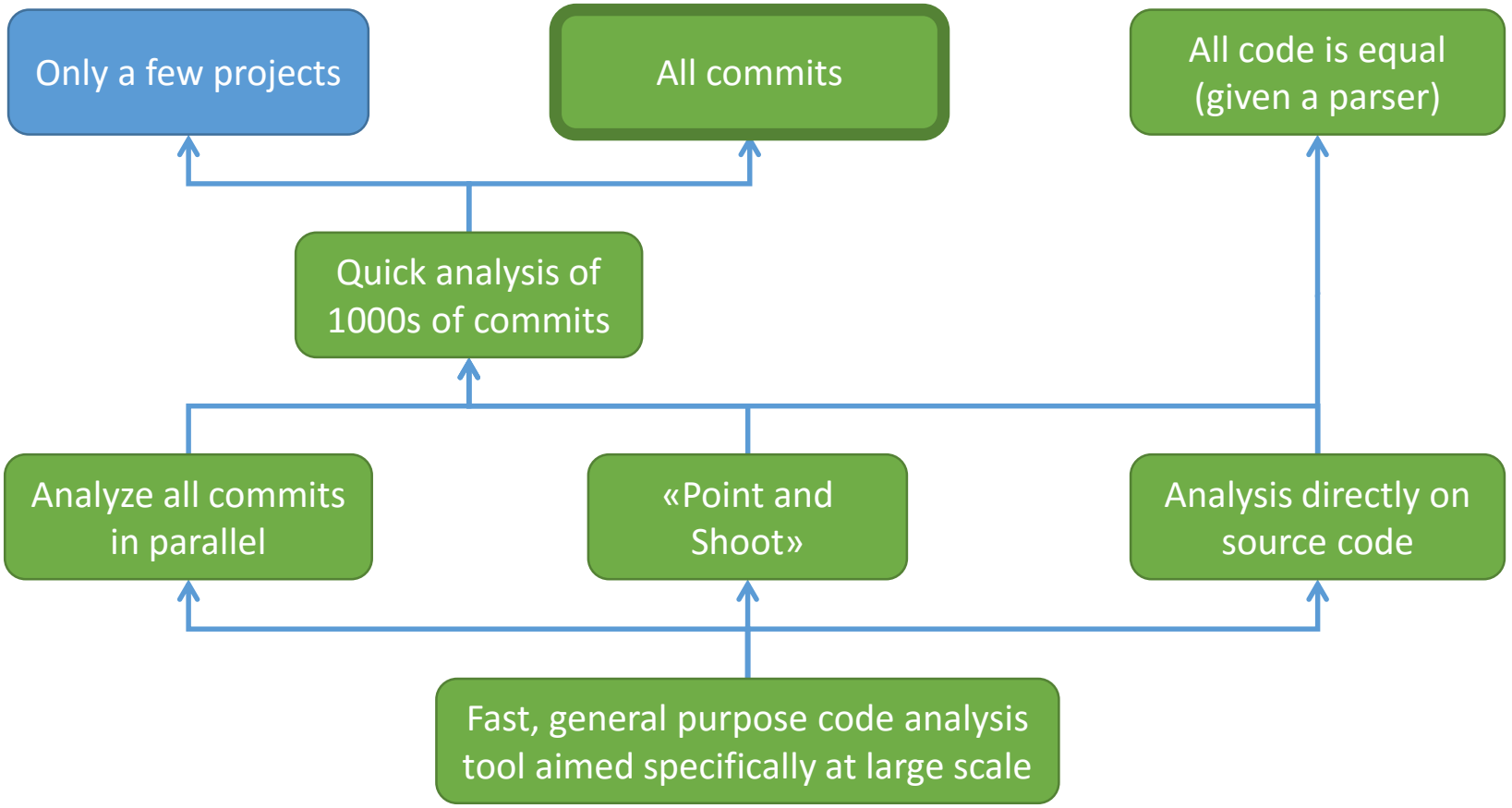

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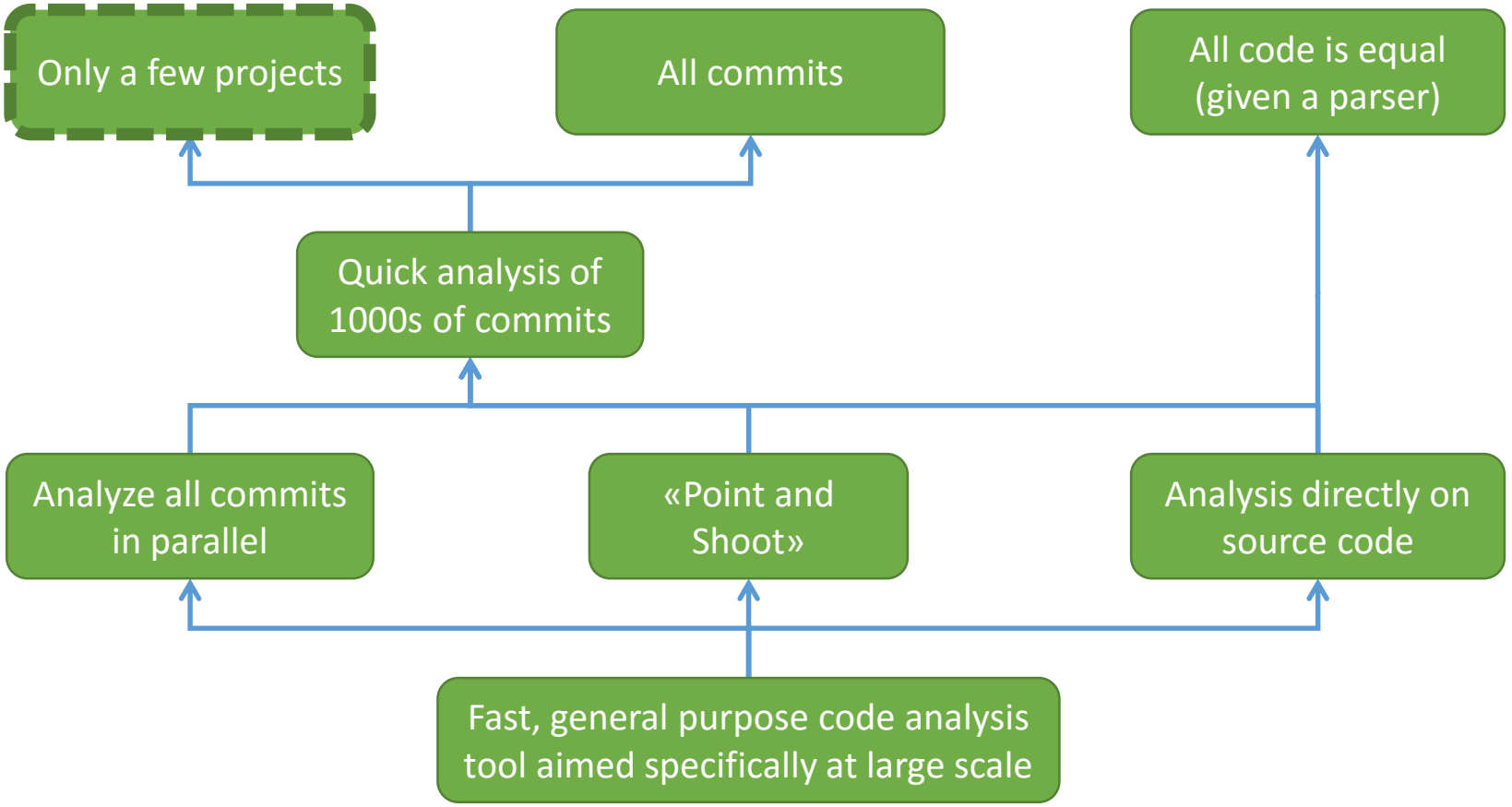
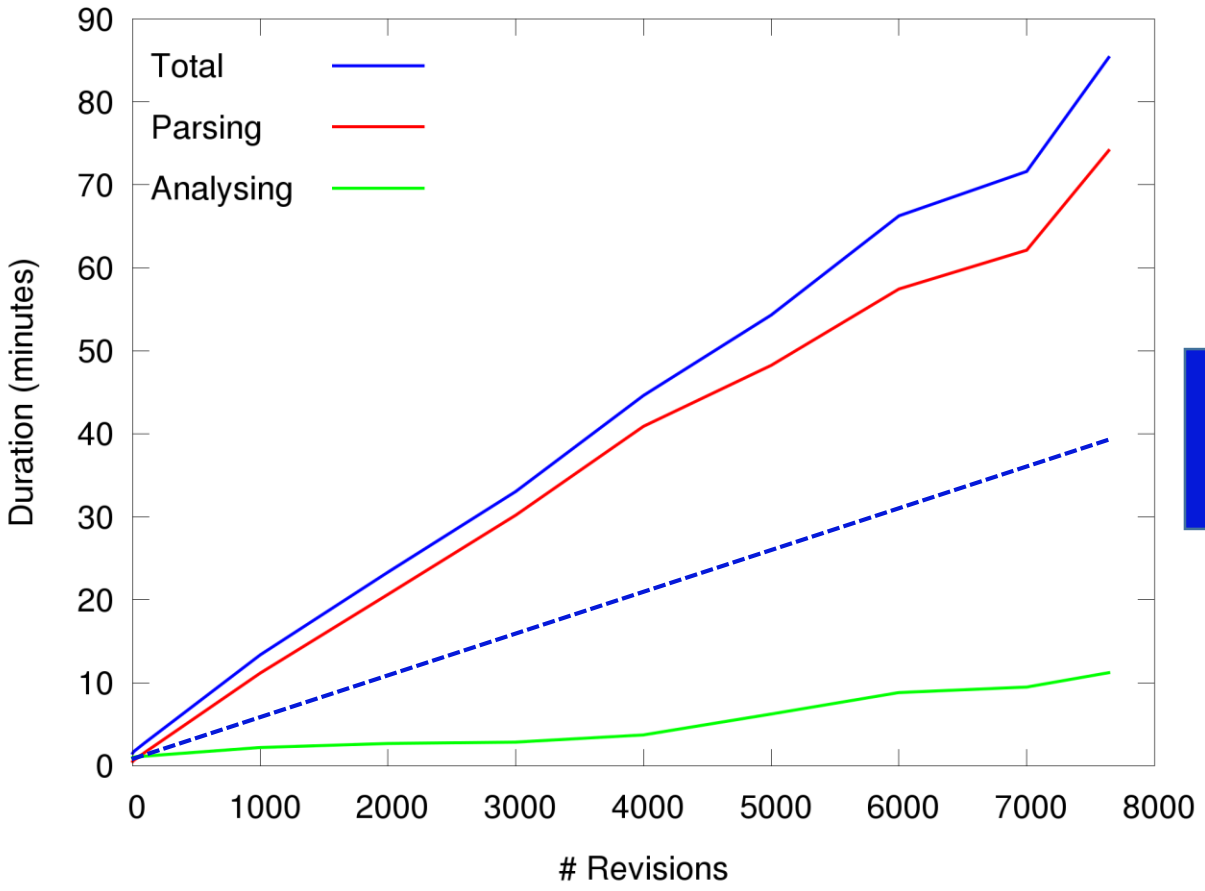


of Commits: Linear Scaling



March Update (Total)

Multi-Project Parallelization

LISA

AspectJ:
7642 commits, 440k LOC
Requirements:
20 GB memory
45 min on 4 cores

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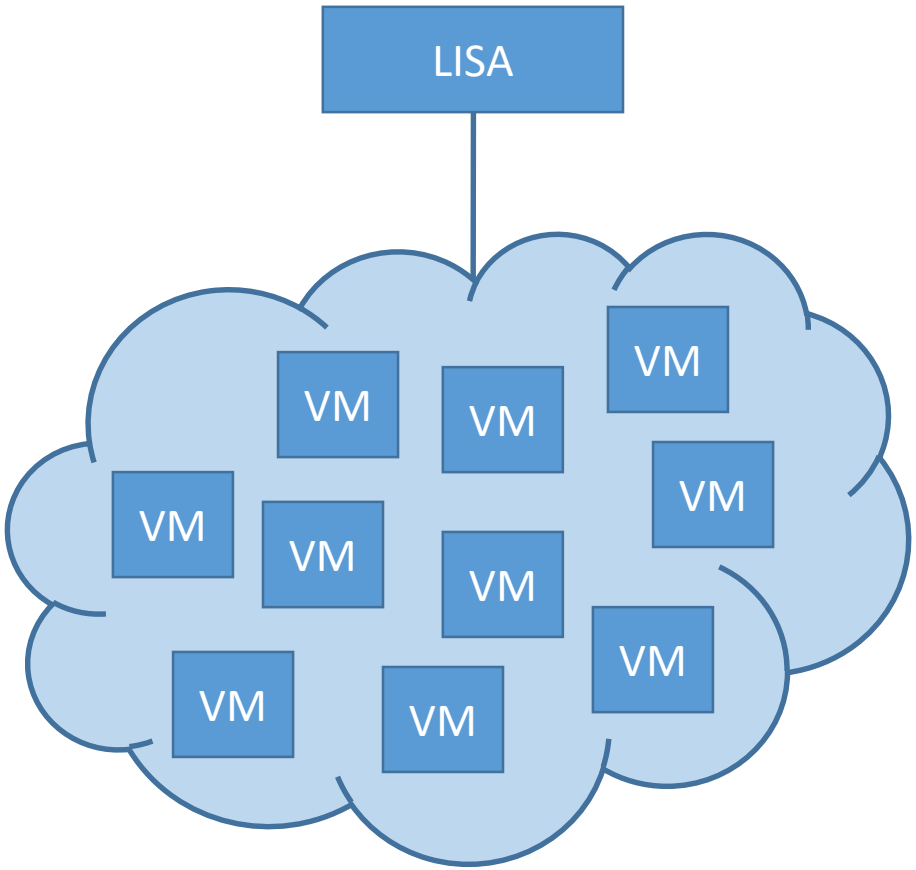

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Requirements:
20 GB memory
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Parallelization scenario:
10x Amazon EC 2 r3.8xlarge
10x 244GB memory
10x 32 cores
Potential to analyze 160 AspectJ-sized projects per hour

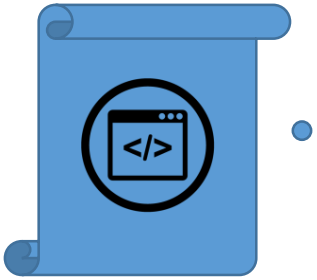
(Most projects on git-hub are much smaller)

Benchmarking

Benchmarking

Is this metric too high?

Here‘s a code smell - should I fix it?

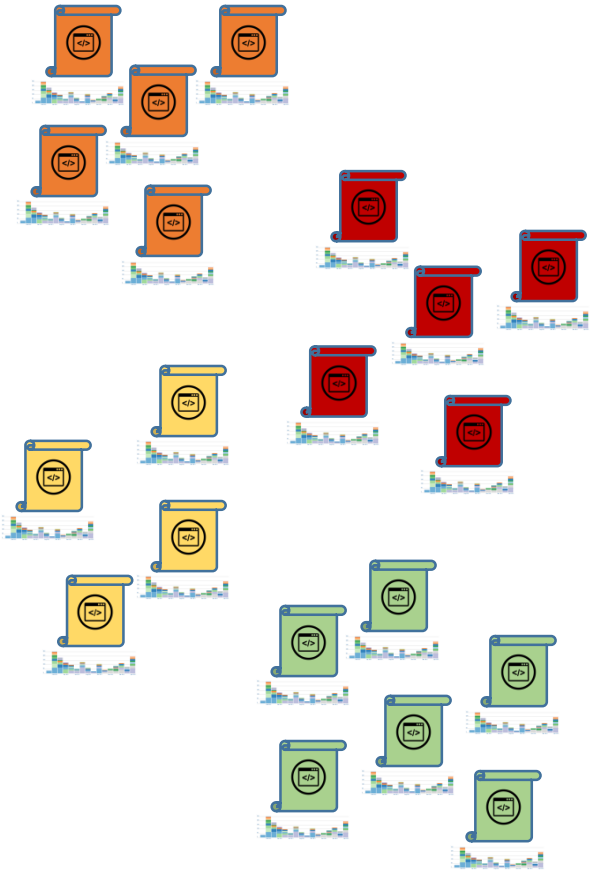


Is this good or bad? What does it even mean?

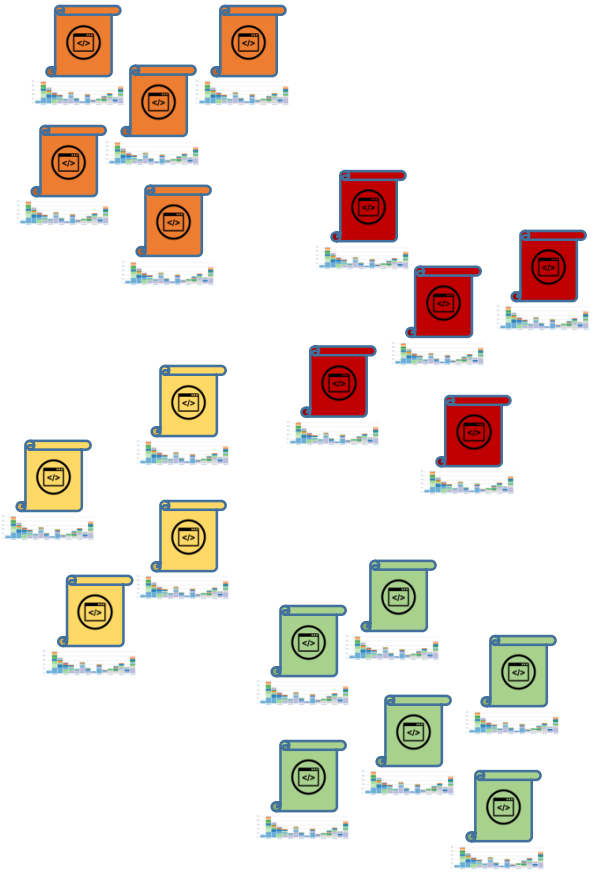
Benchmarking

Benchmarking

Cluster & Compare

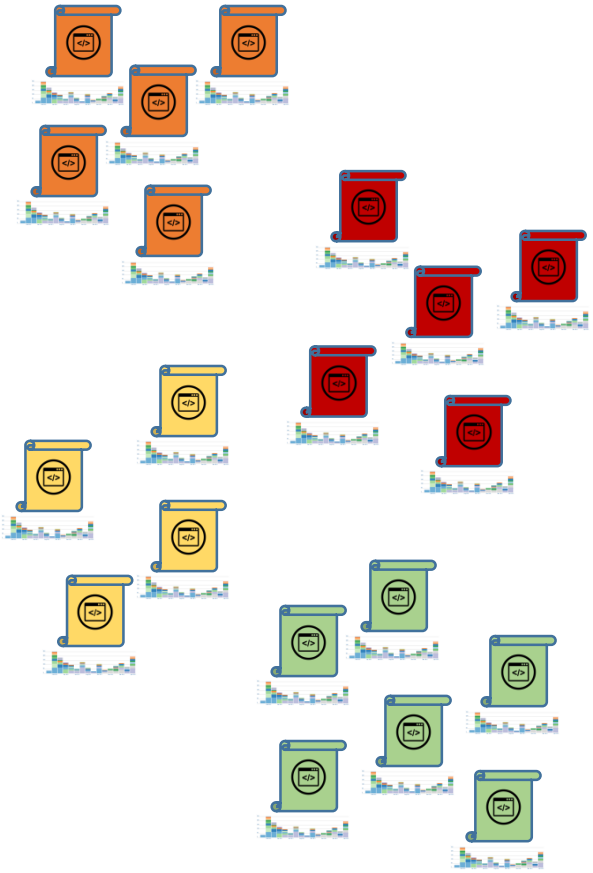


Cluster & Compare



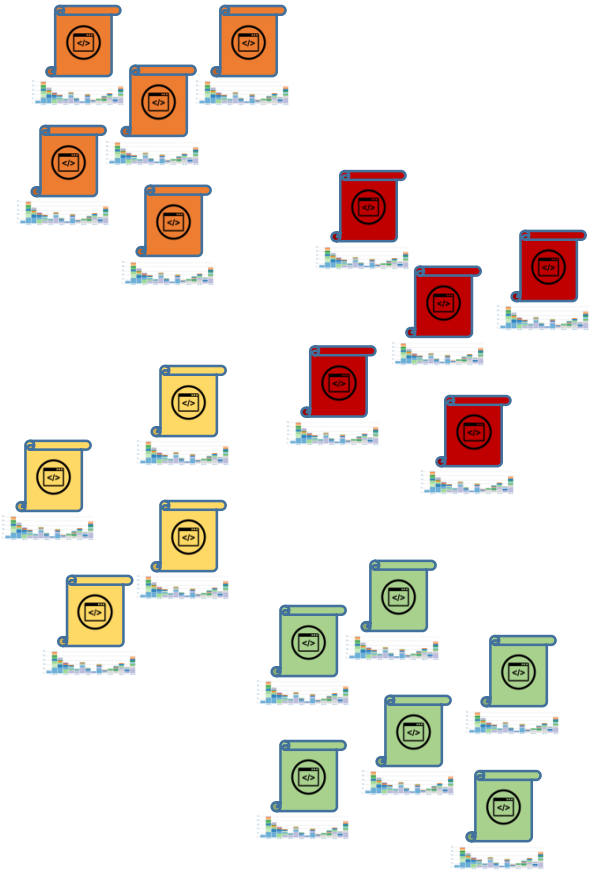
- Discover „phenotypes“
  - By metric values

Cluster & Compare



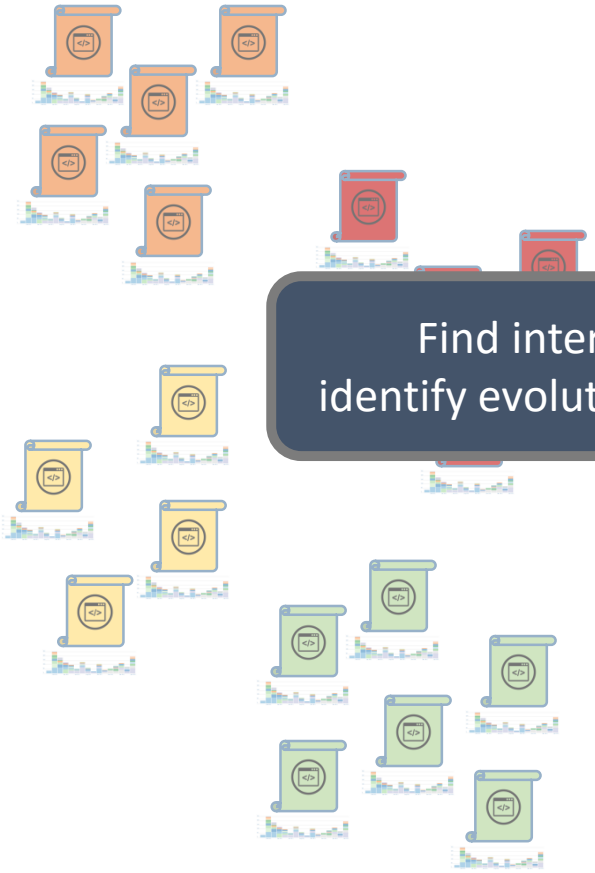
- Discover „phenotypes“
  - By metric values
  - By metric evolution over time

Cluster & Compare



- Discover „phenotypes“
  - By metric values
  - By metric evolution over time
- Compare programming languages / paradigms

Cluster & Compare

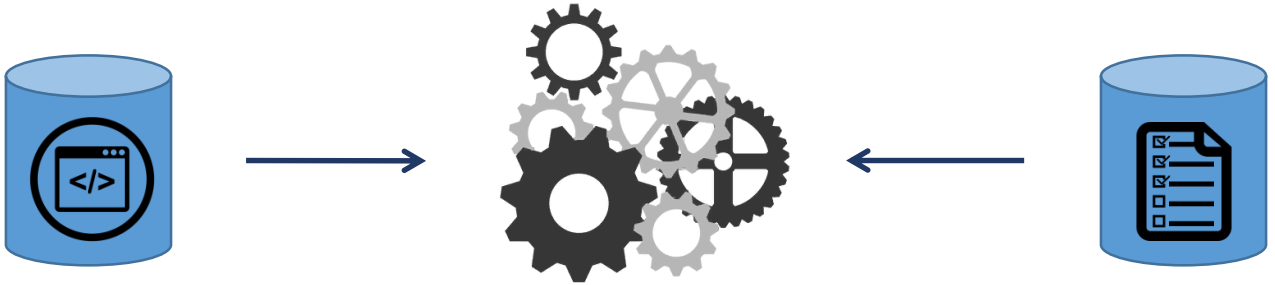


- Discover „phenotypes“
  - By metric values
  - n over

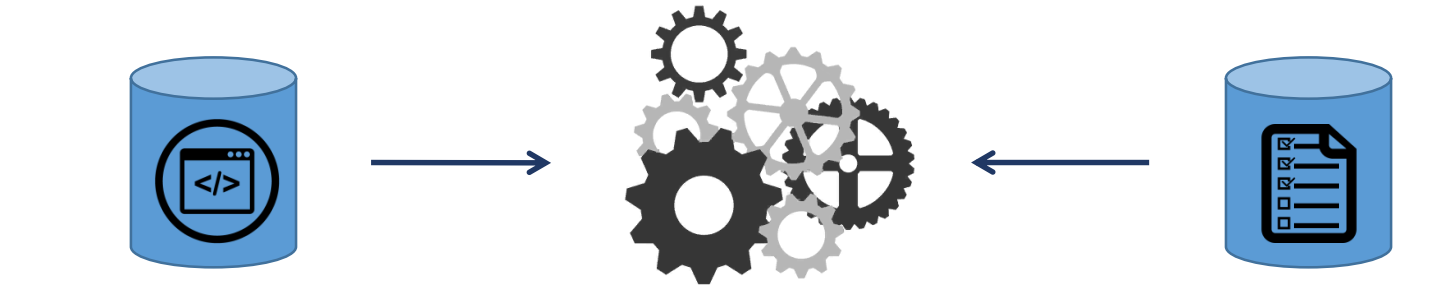
Find interesting projects for further study & identify evolution patterns to find (un-)desirable ones

- Compare programming languages / paradigms

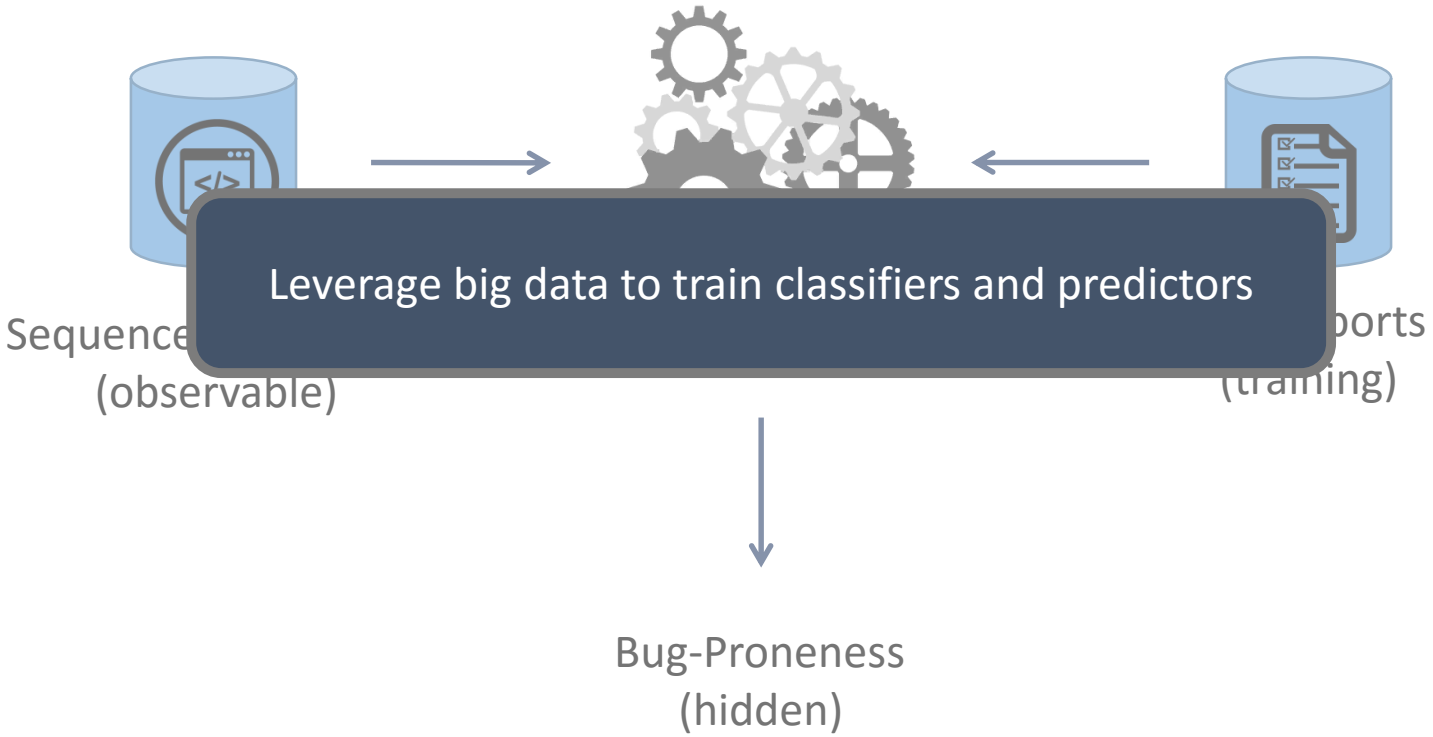
Machine Learning



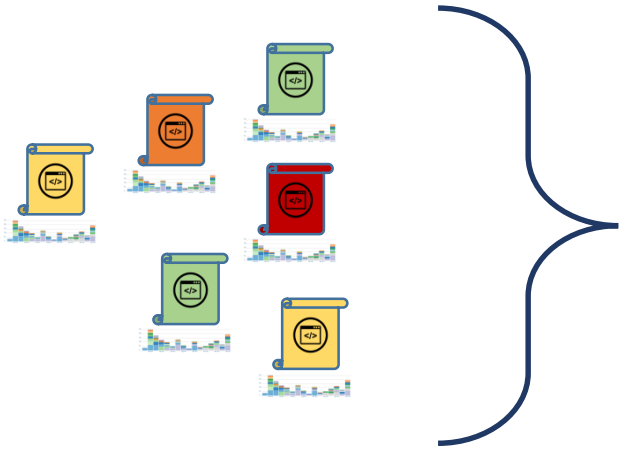
Machine Learning



Machine Learning

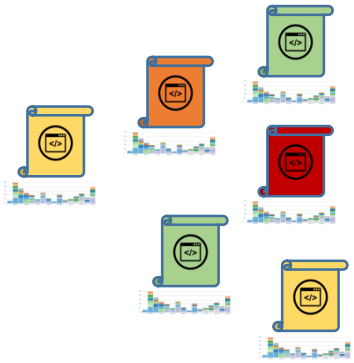


Study Replication

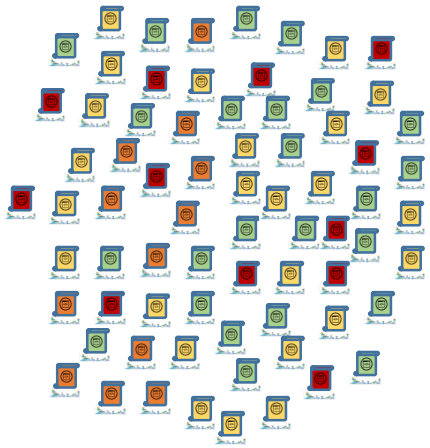


Conclusions X, Y, Z

Study Replication



Conclusions X, Y, Z

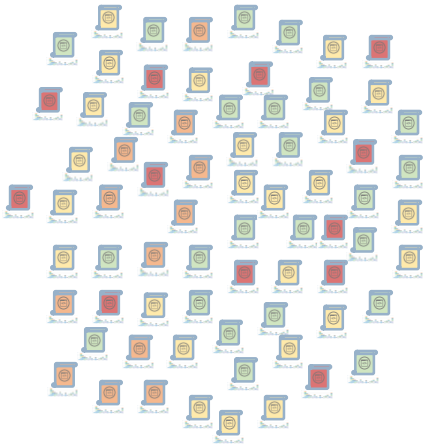


Replicable with 1000s of projects?

Study Replication

Conclusions X, Y, Z

Confirmation or rebuttal of existing assumptions



Replicable with 1000s of projects?