

Test Intelligence

Lücken schließen und überflüssige Tests entfernen,
um mehr Fehler in weniger Zeit zu finden

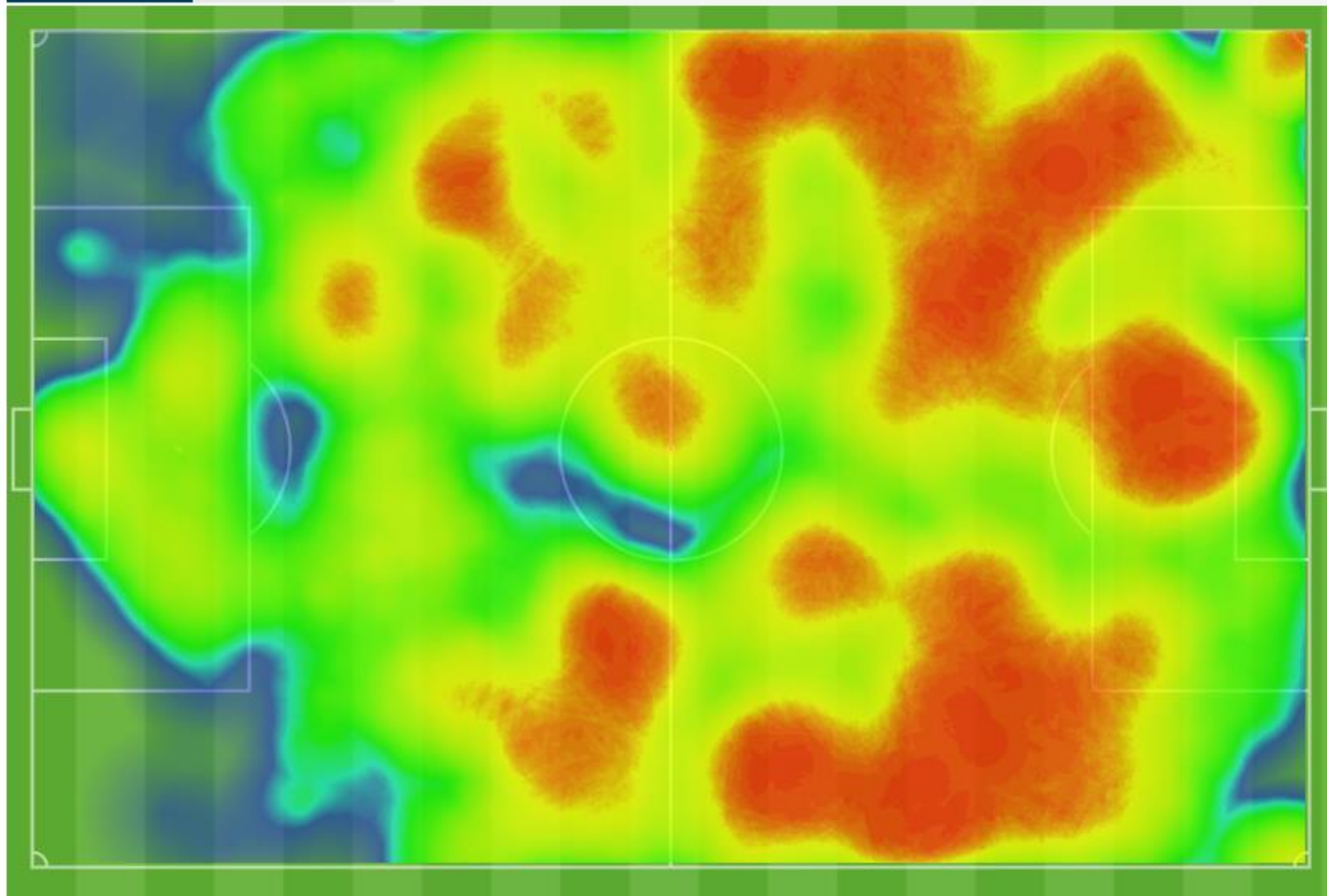
Dr. Elmar Juergens



FC Bayern München

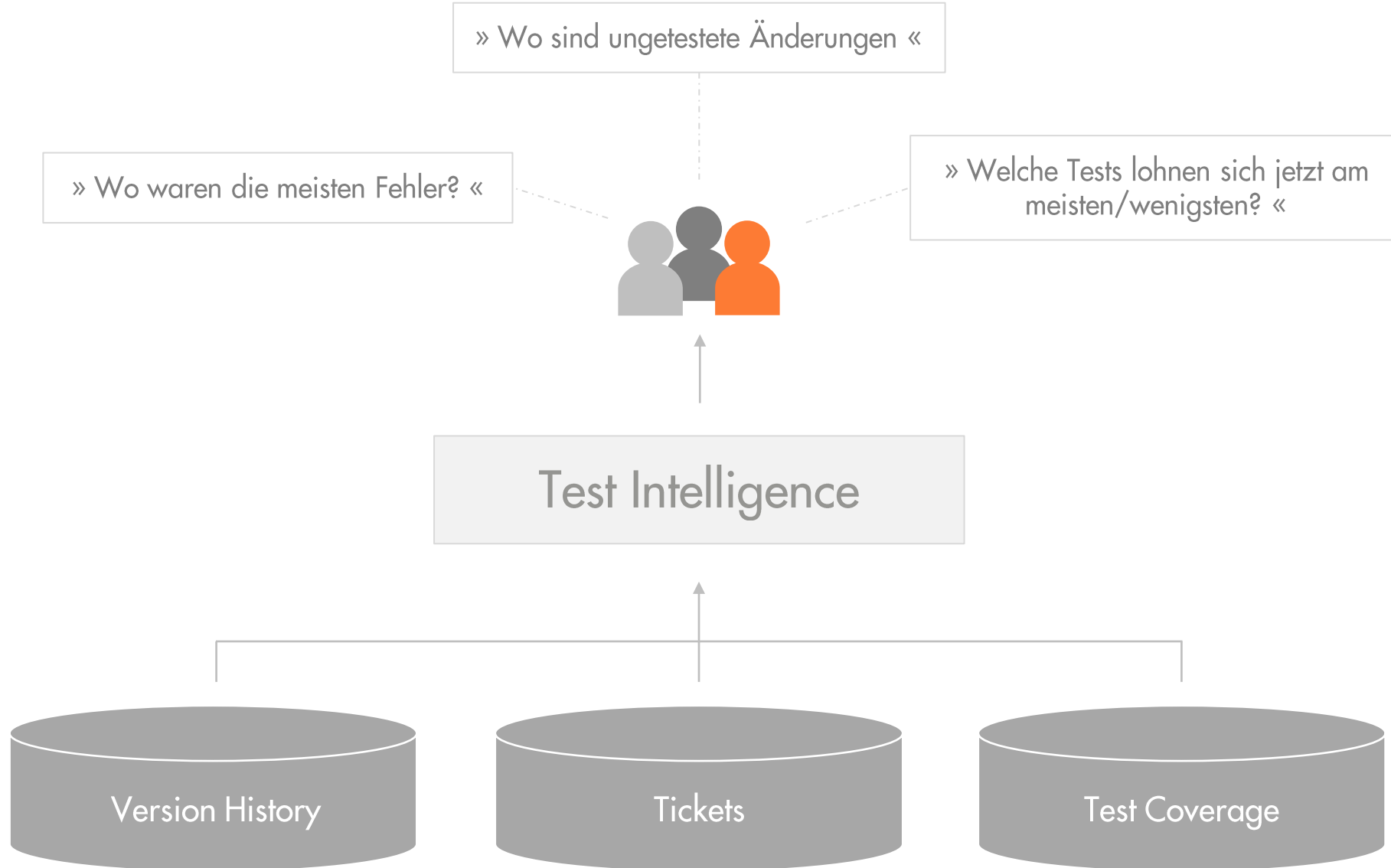
- FC Bayern München
- TW 1 Manuel Neuer
- AW 21 Lucas Hernández
- AW 5 Benjamin Pavard
- AW 4 Niklas Süle
- AW 27 David Alaba
- MF 29 Kingsley Coman
- MF 10 Leroy Sané
- MF 25 Thomas Müller
- MF 6 Joshua Kimmich
- MF 18 Leon Goretzka
- ST 13 Eric Maxim Choupo-Moting
- Einwechselfspieler
- 17 Jérôme Boateng
- 19 Alphonso Davies

Heatmap Touchmap



Paris Saint-Germain

- Paris Saint-Germain
- TW 1 Keylor Navas
- AW 5 Marquinhos
- AW 31 Colin Dagba
- AW 22 Abdou Diallo
- AW 3 Presnel Kimpembe
- MF 23 Julian Draxler
- MF 27 Idrissa Gueye
- MF 10 Neymar
- MF 15 Danilo Pereira
- MF 11 Ángel Di María
- ST 7 Kylian Mbappé
- Einwechselfspieler
- 25 Mitchel Bakker
- 18 Moise Kean
- 21 Ander Herrera
- 12 Rafinha

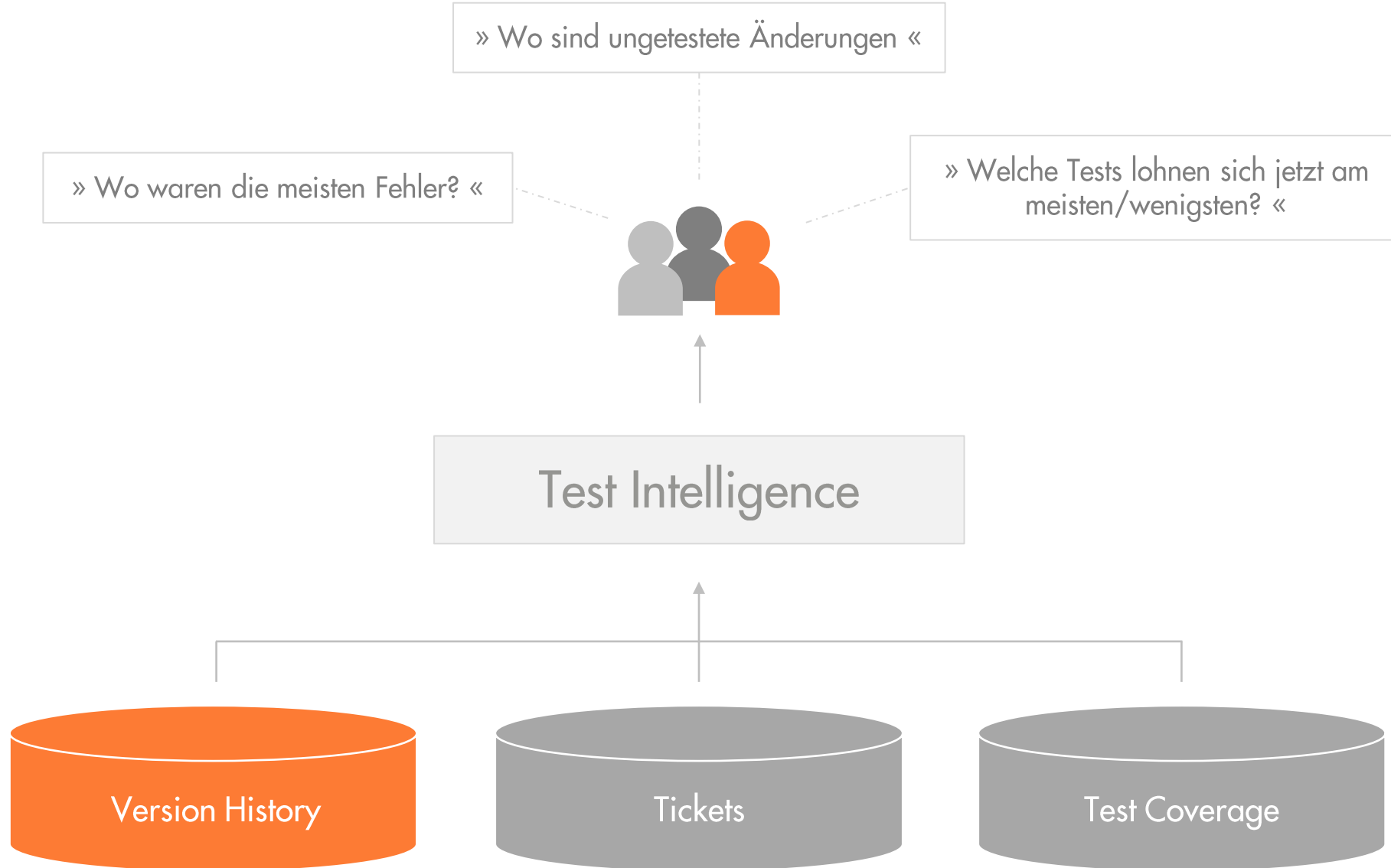




TUM



CQSE



```

23
24
25
26 /* package */class NamespaceRenames {
27
28     /** Maps from old name fragment to new name fragment */
29     public Set<ImmutablePair<String, String>> namespaceRenames = new HashSet<
30
31     /**
32      * Computes a rename rule from an old and a new namespace name based on a
33      * type name correspondence. <code>
34      * For name1 = a.oldnamespace.b.c.D and name2 = a.newnamespace.b.c.D, res
35      * </code>
36      * */
37     public static ImmutablePair<String, String> computeRenameRule(
38         String fqTypeName1, String fqTypeName2, String separator) {
39
40         if (fqTypeName1.equals(fqTypeName2)) {
41             return null;
42         }
43
44         String commonSuffix = StringUtils.longestCommonSuffix(fqTypeName1,
45             fqTypeName2);
46         if (StringUtils.isEmpty(commonSuffix)) {
47             return null;
48         }
49
50         int separatorPosition = commonSuffix.indexOf(separator);
51         if (separatorPosition != -1) {
52             commonSuffix = commonSuffix.substring(separatorPosition);
53         }
54
55         String from = StringUtils.stripSuffix(commonSuffix, fqTypeName1);
56         String to = StringUtils.stripSuffix(commonSuffix, fqTypeName2);
57
58         return new ImmutablePair<String, String>(from, to);
59     }
60
61
62
63
64

```

```

24
25
26 /* package */class NamespaceRenames {
27
28     /** Maps from old name fragment to new name fragment */
29     public Set<ImmutablePair<String, String>> namespaceRenames = new Has
30
31     // TODO (LH) Doc vs. method name : 'Compute' vs. 'find'
32     /**
33      * Computes a rename rule from an old and a new namespace name based
34      * type name correspondence. <code>
35      * For name1 = a.oldnamespace.b.c.D and name2 = a.newnamespace.b.c.D
36      * </code>
37      * */
38     // TODO (LH) Please reflect in identifier names that they refer to
39     // names
40     public static ImmutablePair<String, String> findRenameRule(String na
41         String name2, String separator) {
42
43         if (name1.equals(name2)) {
44             return null;
45         }
46
47         // TODO (LH) Looks like StringUtils is missing a 'longestCommon
48         String commonSuffix = reverse(StringUtils.longestCommonPrefix(
49             reverse(name1), reverse(name2)));
50         if (StringUtils.isEmpty(commonSuffix)) {
51             return null;
52         }
53
54         int separatorPosition = commonSuffix.indexOf(separator);
55         if (separatorPosition != -1) {
56             commonSuffix = commonSuffix.substring(separatorPosition);
57         }
58
59         // TODO (LH) Please reflect in identifier names that these are
60         // namespaces
61         String from = StringUtils.stripSuffix(commonSuffix, name1);
62         String to = StringUtils.stripSuffix(commonSuffix, name2);
63
64         return new ImmutablePair<String, String>(from, to);
65

```

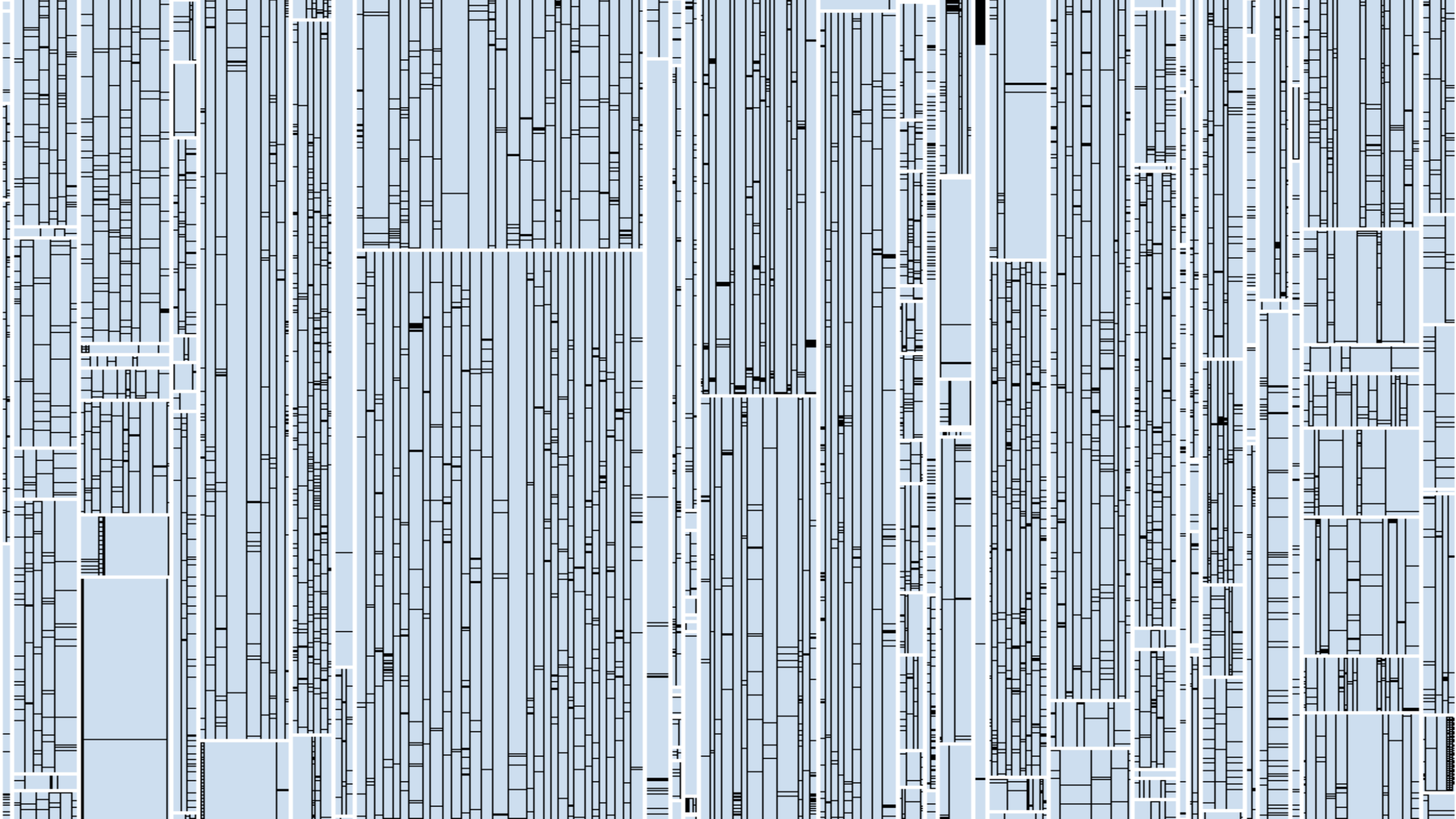

GUI.Dialogs

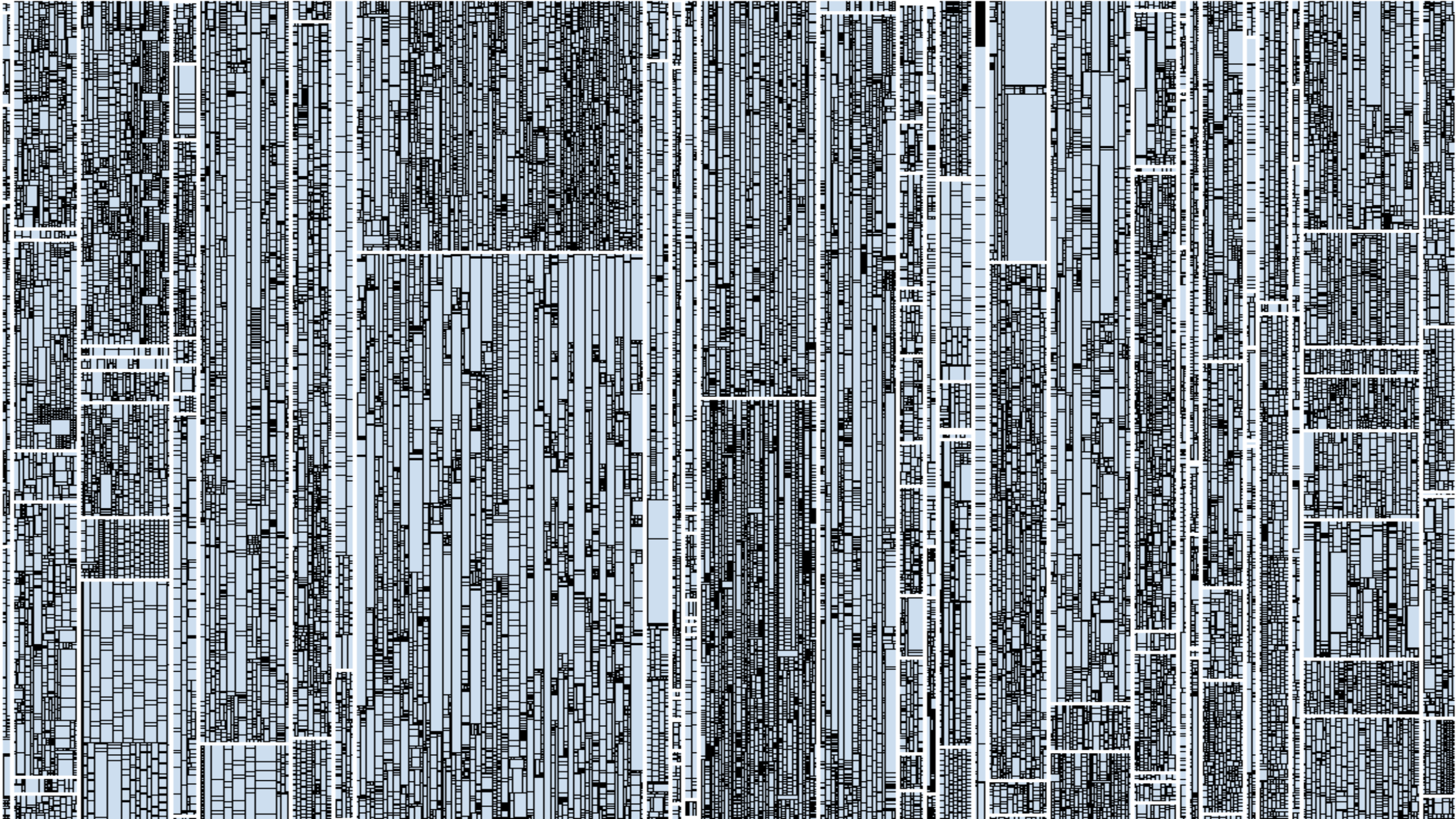
UI Controls

GUI.Base

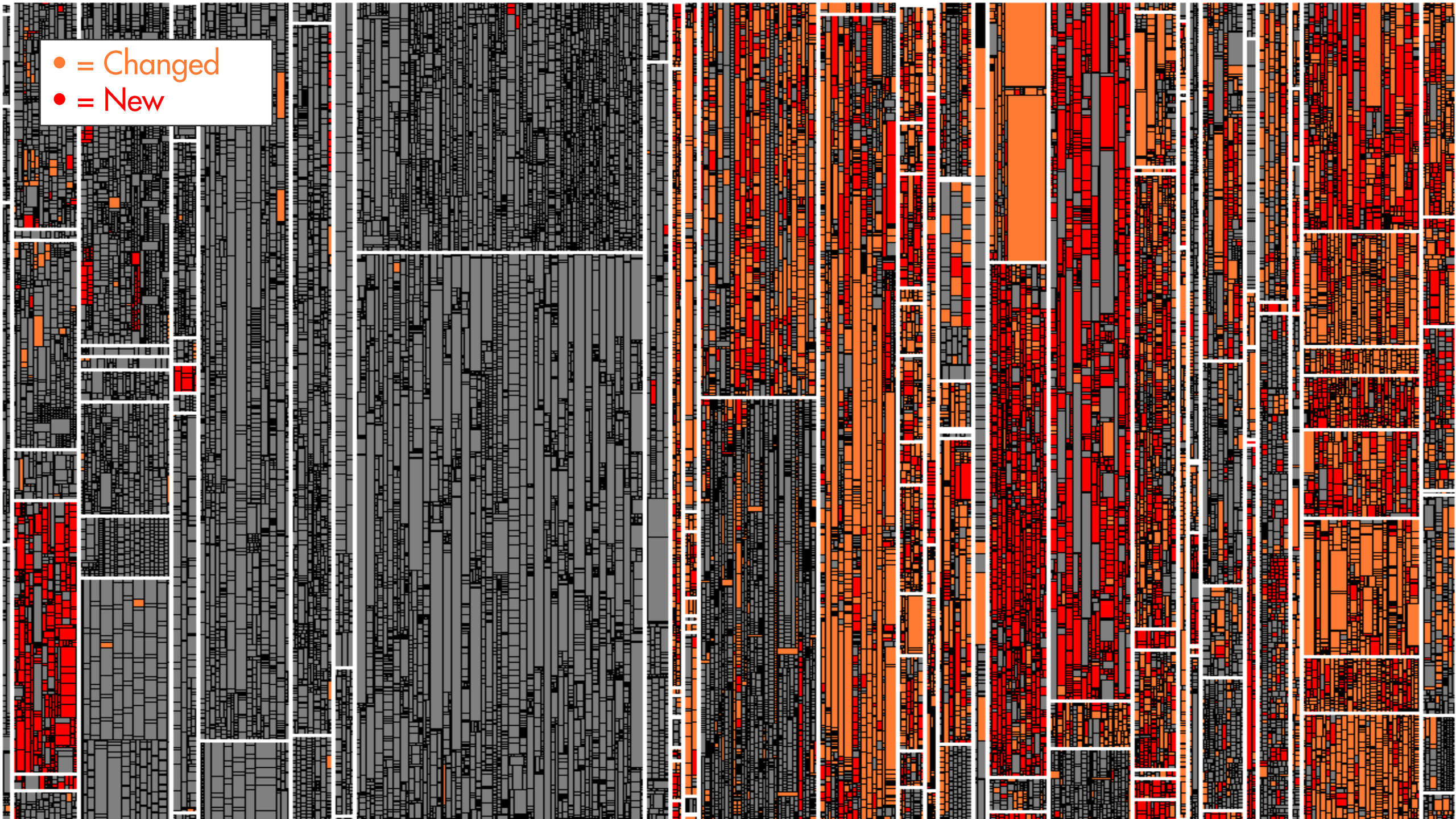
Authentication

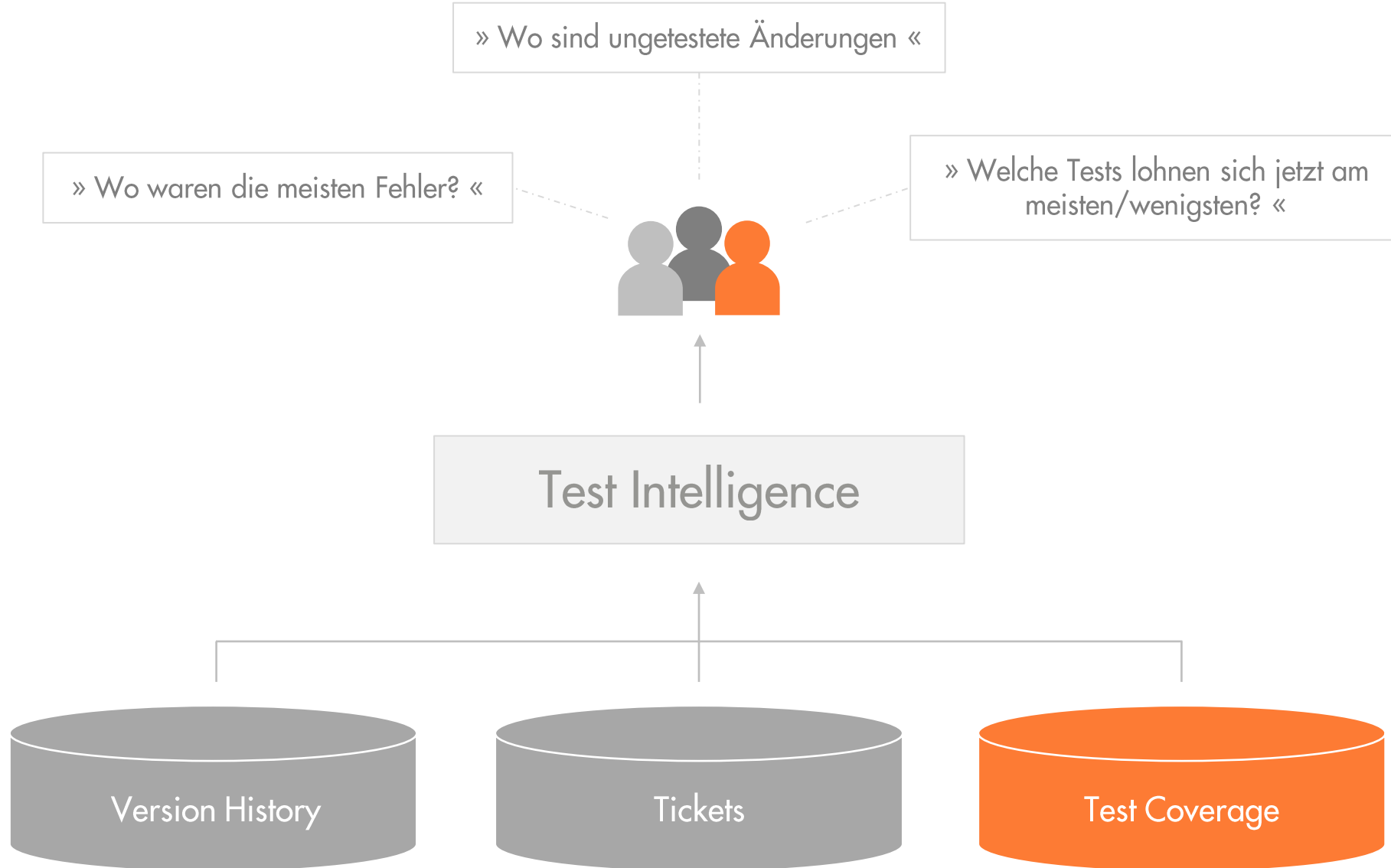
Data Validation





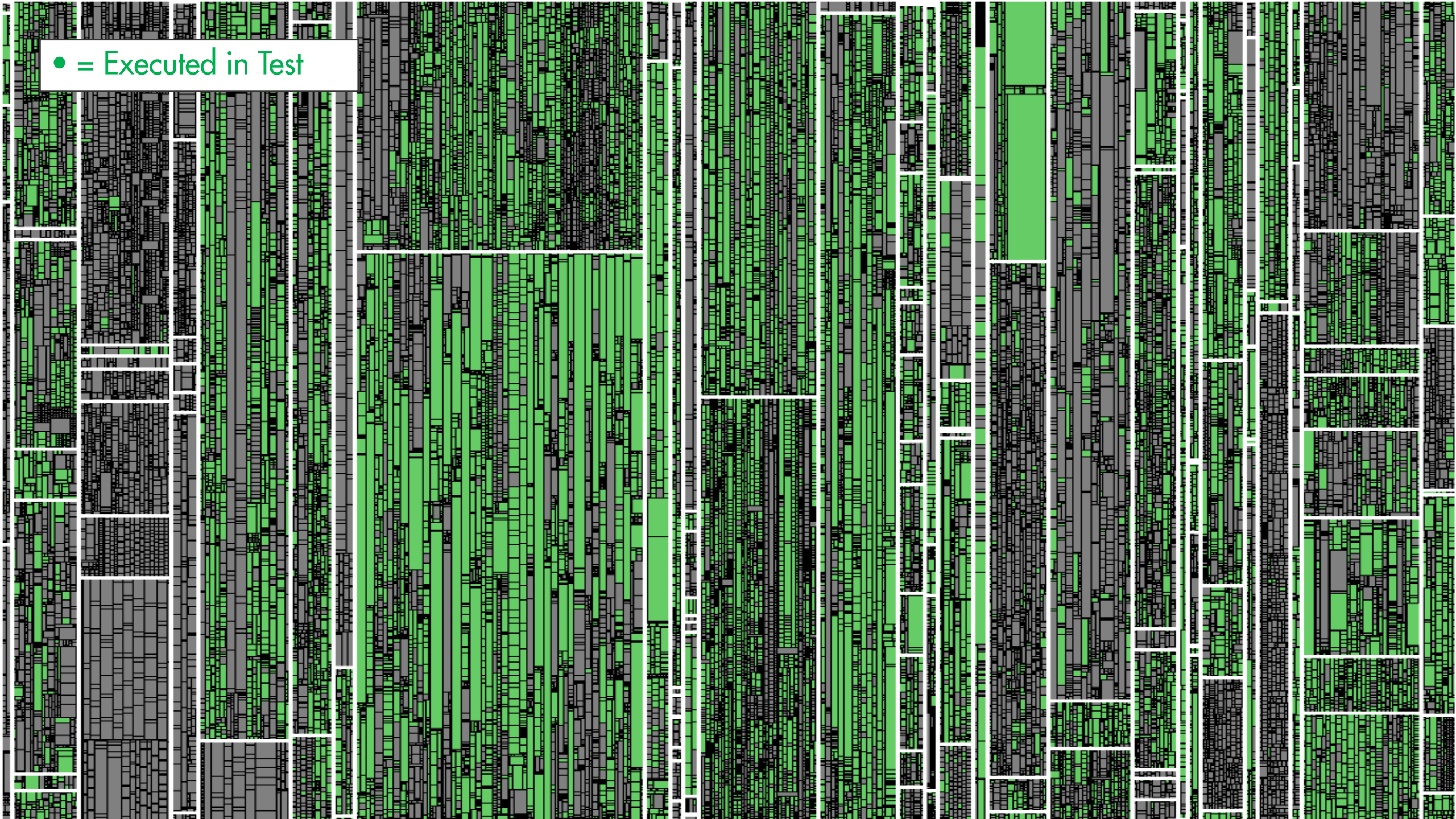
- = Changed
- = New

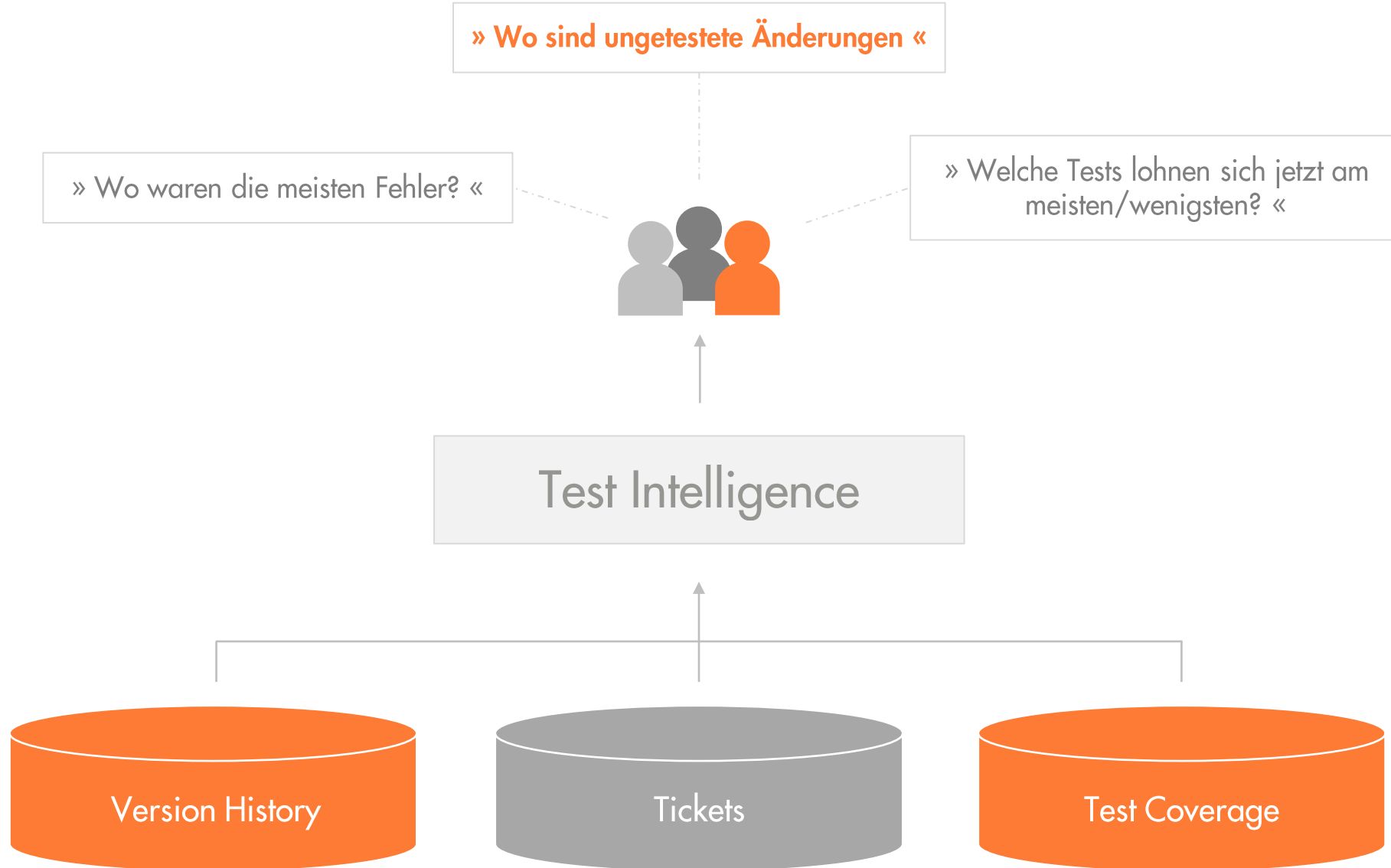




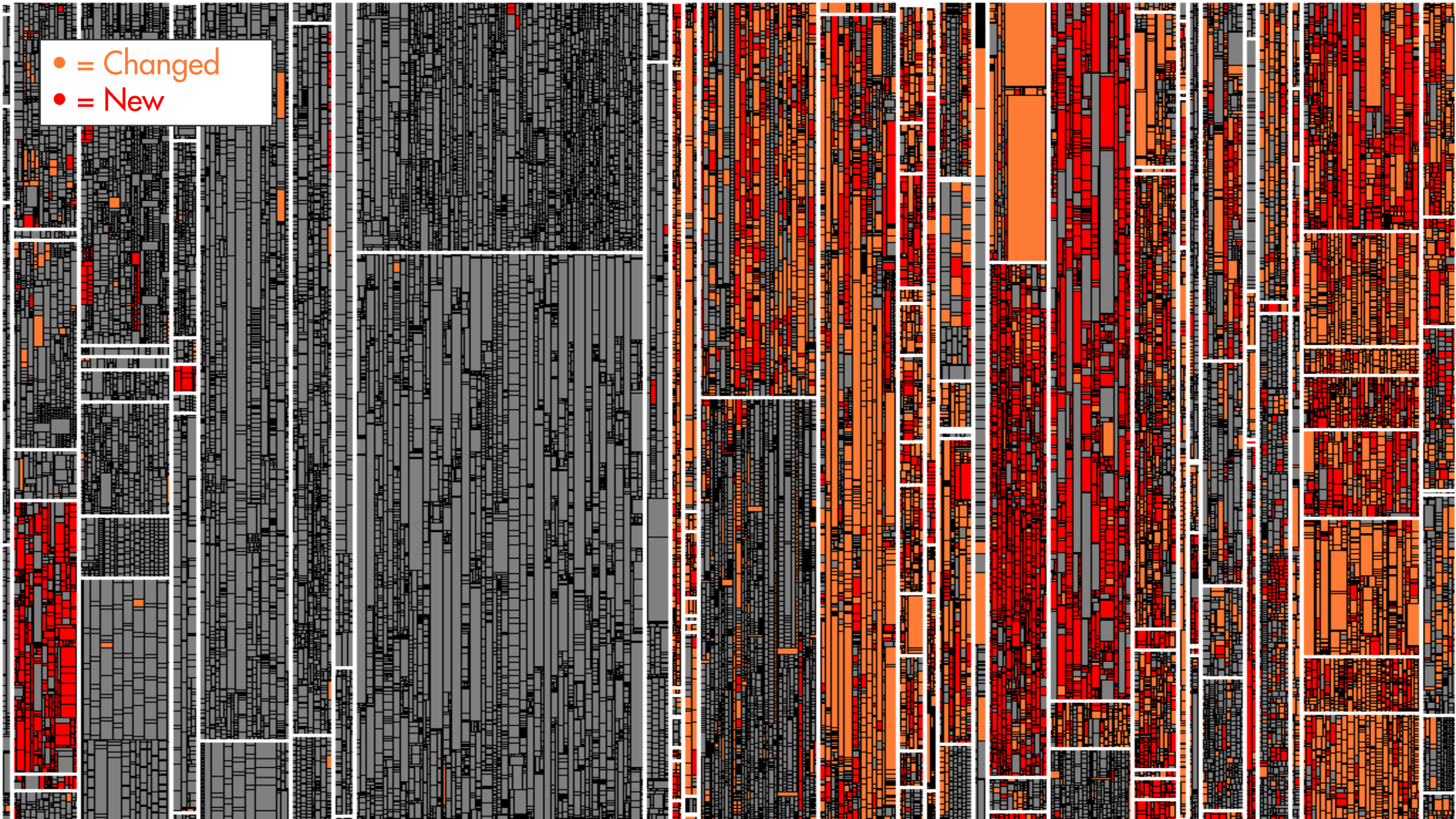
```
1  protected void calculateIndirectAmbiguities() {
2      Map<NucleotideCompound, List<NucleotideCompound>> equivalentMap = new HashMap<>>();
3
4      List<NucleotideCompound> ambiguousCompounds = new ArrayList<NucleotideCompound>();
5      for (NucleotideCompound compound : getAllCompounds()) {
6          if (!compound.isAmbiguous()) {
7              continue;
8          }
9          ambiguousCompounds.add(compound);
10     }
11
12     for (NucleotideCompound sourceCompound : ambiguousCompounds) {
13         Set<NucleotideCompound> sourceConstituents = sourceCompound.getConstituents();
14         for (NucleotideCompound targetCompound : ambiguousCompounds) {
15             Set<NucleotideCompound> targetConstituents = targetCompound.getConstituents();
16             if (targetConstituents.containsAll(sourceConstituents)) {
17                 NucleotideCompound lcSourceCompound = toLowerCase(sourceCompound);
18                 NucleotideCompound lcTargetCompound = toLowerCase(targetCompound);
19                 checkAdd(equivalentMap, sourceCompound, targetCompound);
20                 checkAdd(equivalentMap, sourceCompound, lcTargetCompound);
21                 checkAdd(equivalentMap, targetCompound, sourceCompound);
22                 checkAdd(equivalentMap, lcTargetCompound, sourceCompound);
23                 checkAdd(equivalentMap, lcSourceCompound, targetCompound);
24                 checkAdd(equivalentMap, lcSourceCompound, lcTargetCompound);
25             }
26         }
27     }
28
29     for (NucleotideCompound key : equivalentMap.keySet()) {
30         List<NucleotideCompound> vals = equivalentMap.get(key);
31         for (NucleotideCompound value : vals) {
32             addEquivalent((C) key, (C) value);
33             addEquivalent((C) value, (C) key);
34         }
35     }
36 }
```


● = Executed in Test



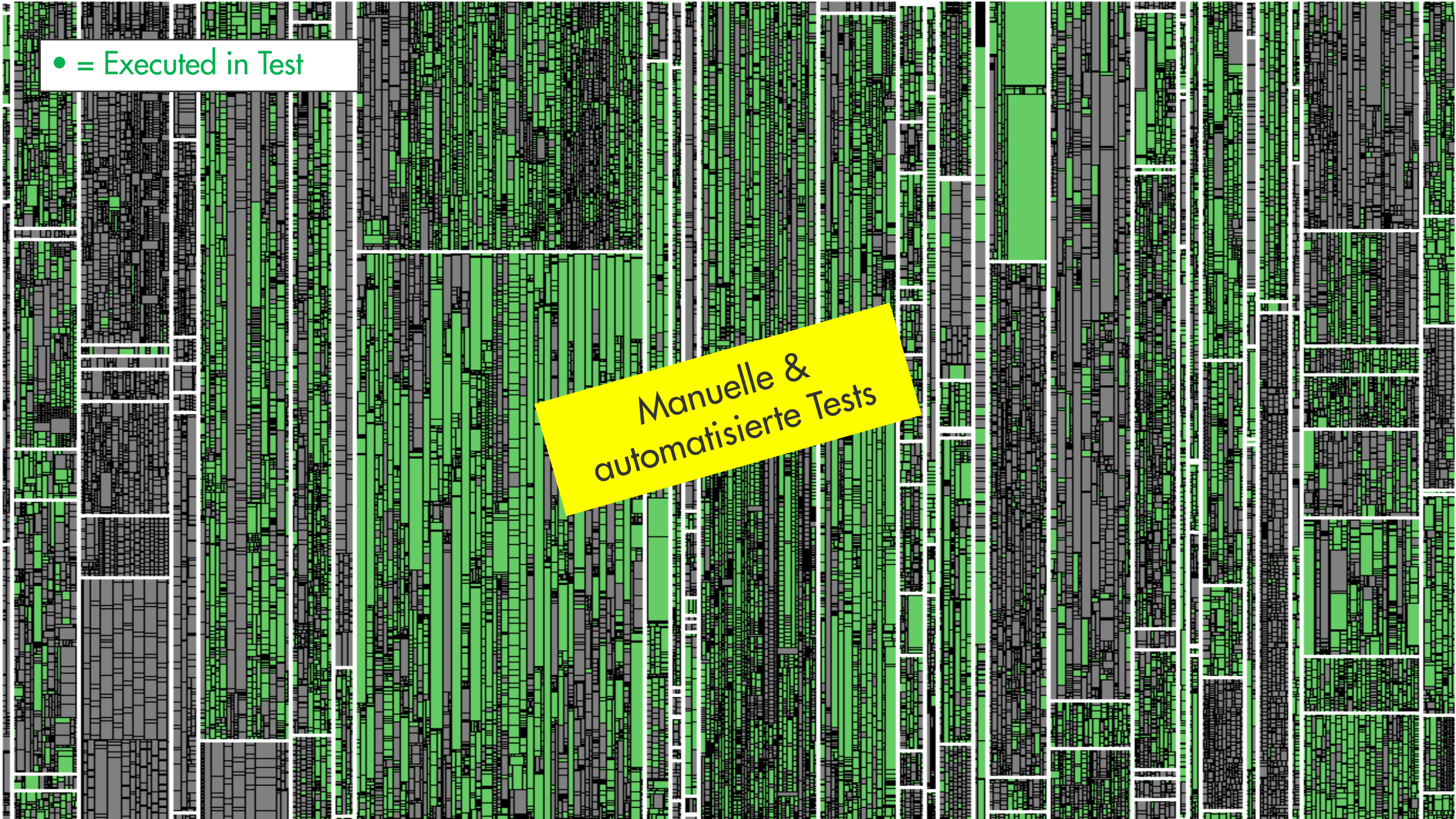


- = Changed
- = New

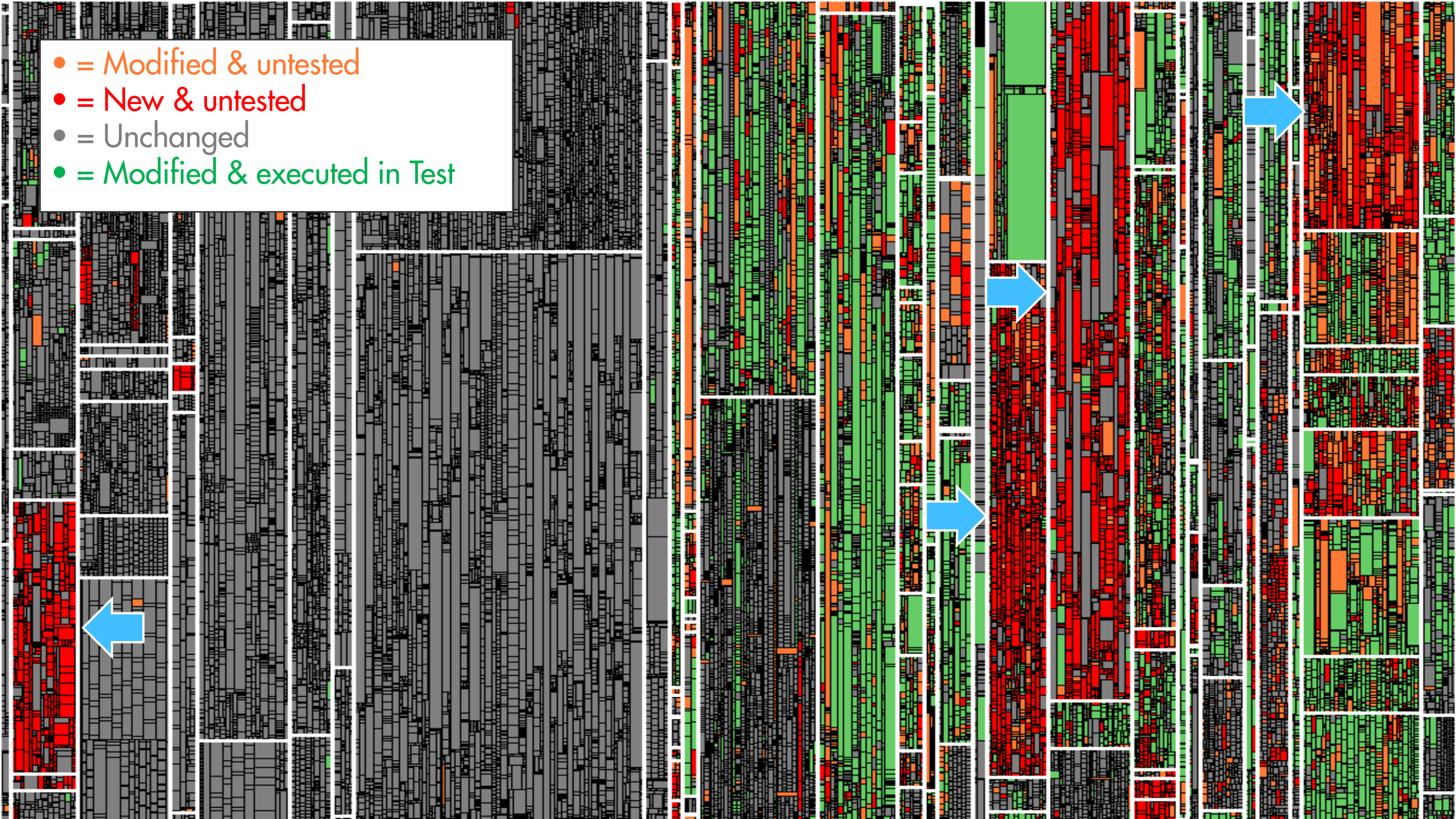


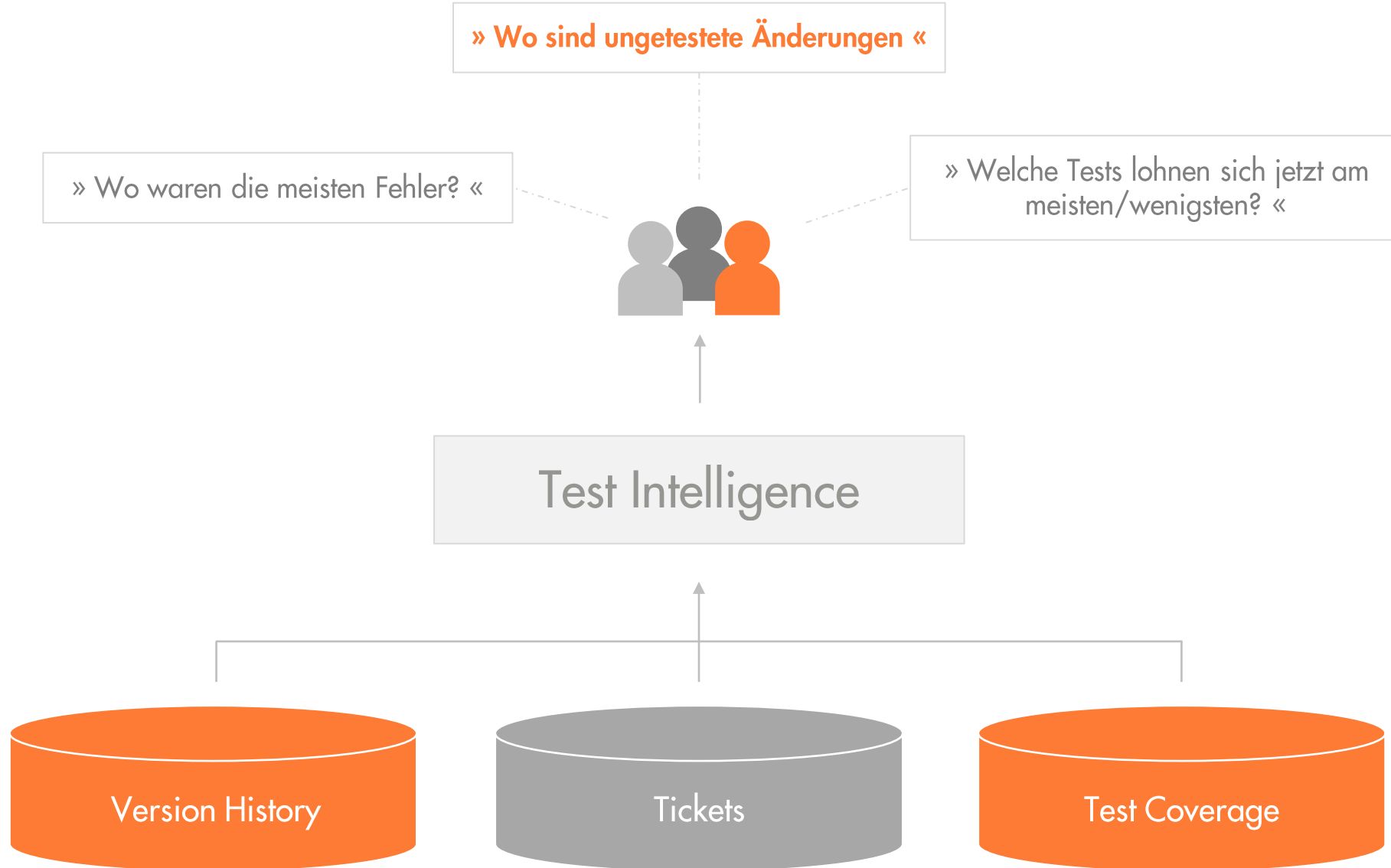
● = Executed in Test

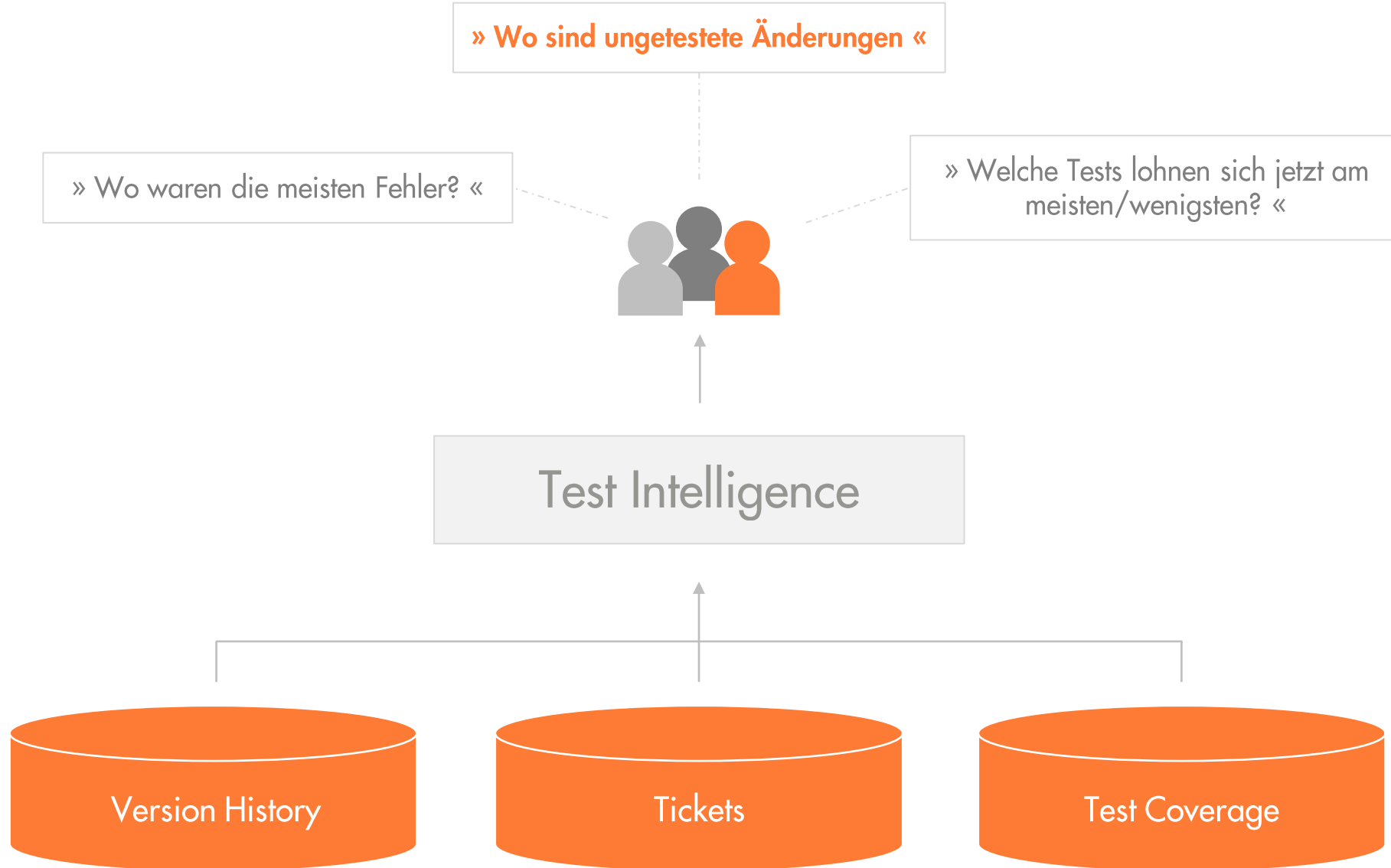
Manuelle &
automatisierte Tests
























- = Modified & untested
- = New & untested
- = Unchanged
- = Modified & executed in Test









Issue # ▼	Subject	Done		Test Gap
↗ TS-10549	Undo/Redo for web-based architecture editor	Done		0% 
↗ TS-10784	Fix long method finding in TaintAnalysisRunner	Done		0% 
↗ TS-10923	Implement metric 'Nesting Depth' for Simulink	Done		29% 
↗ TS-11364	External findings are not registered during first upload	Done		14% 
↗ TS-11942	Manual test coverage upload during development	Done		43% 
↗ TS-12050	Tool for transferring findings blacklists and tasks	Done		50% 
↗ TS-12262	Cannot set or alter alias without reanalysis	Done		0% 
↗ TS-13151	Fetch parent relationship of TFS work items	Done		0% 

Issue # ▾	Subject		Test Gap
TS-14421	Get rid of TestGapSynchronizer block	Done 	0% 
TS-14733	Remove Dataflow blocks	Done 	22% 

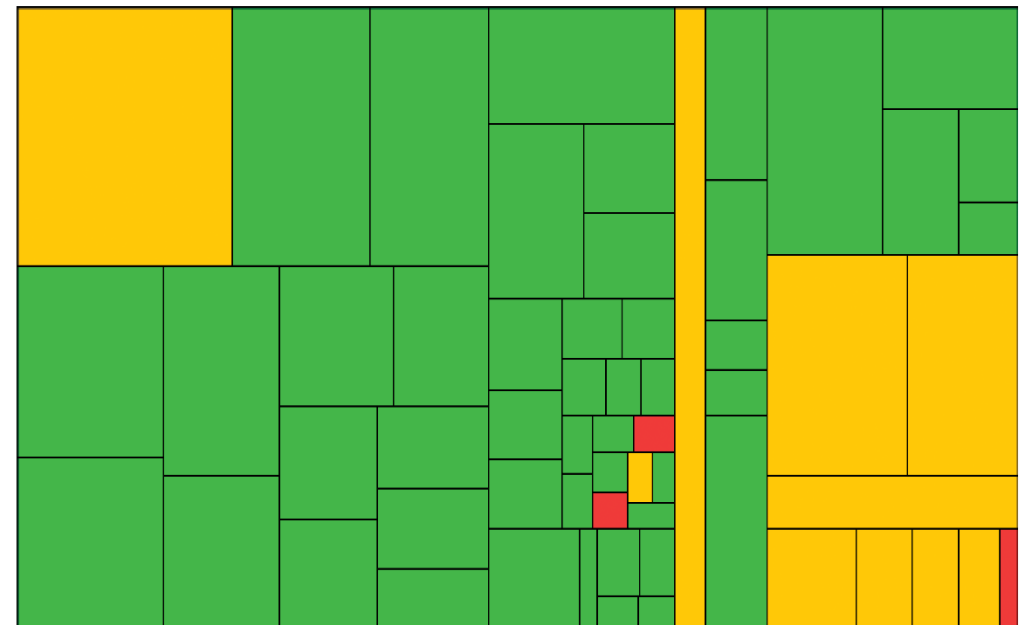
Done Issue TS-14733 - Remove Dataflow blocks

Creator:  (on Apr 06 2018 19:44) Last update: Aug 24 2018 09:32

Assignee: 

Project	Type	Priority	Resolution	Fix Version
TS	Maintenance	Normal	Green	Teamscale 4.5
Component	Labels	Affected Version	Customer	Customer Issue
Backend	Performance			
Epic Name	Freshdesk URL	Merge Request		
		https://git.cqse.eu/cqse/teamscale/3621		

Aug 15 2018 12:37–Now | Test Gap: 22%



Virtueller Workshop

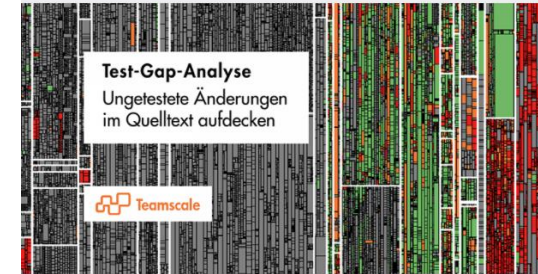
21. April 21, 10:30 – 12:00 Uhr

Anmeldung:

<http://cqse.eu/tga-workshop-gtb>

eventbrite

Müssen Sie einige Updates vornehmen? [Event bearbeiten](#)



Registrieren

Ablauf

- Test-Gap-Analyse (TGA) in Forschung und Praxis
- Live-Demonstration
- Nutzen-Analyse zum Einsatz bei der Munich Re

Vortragende

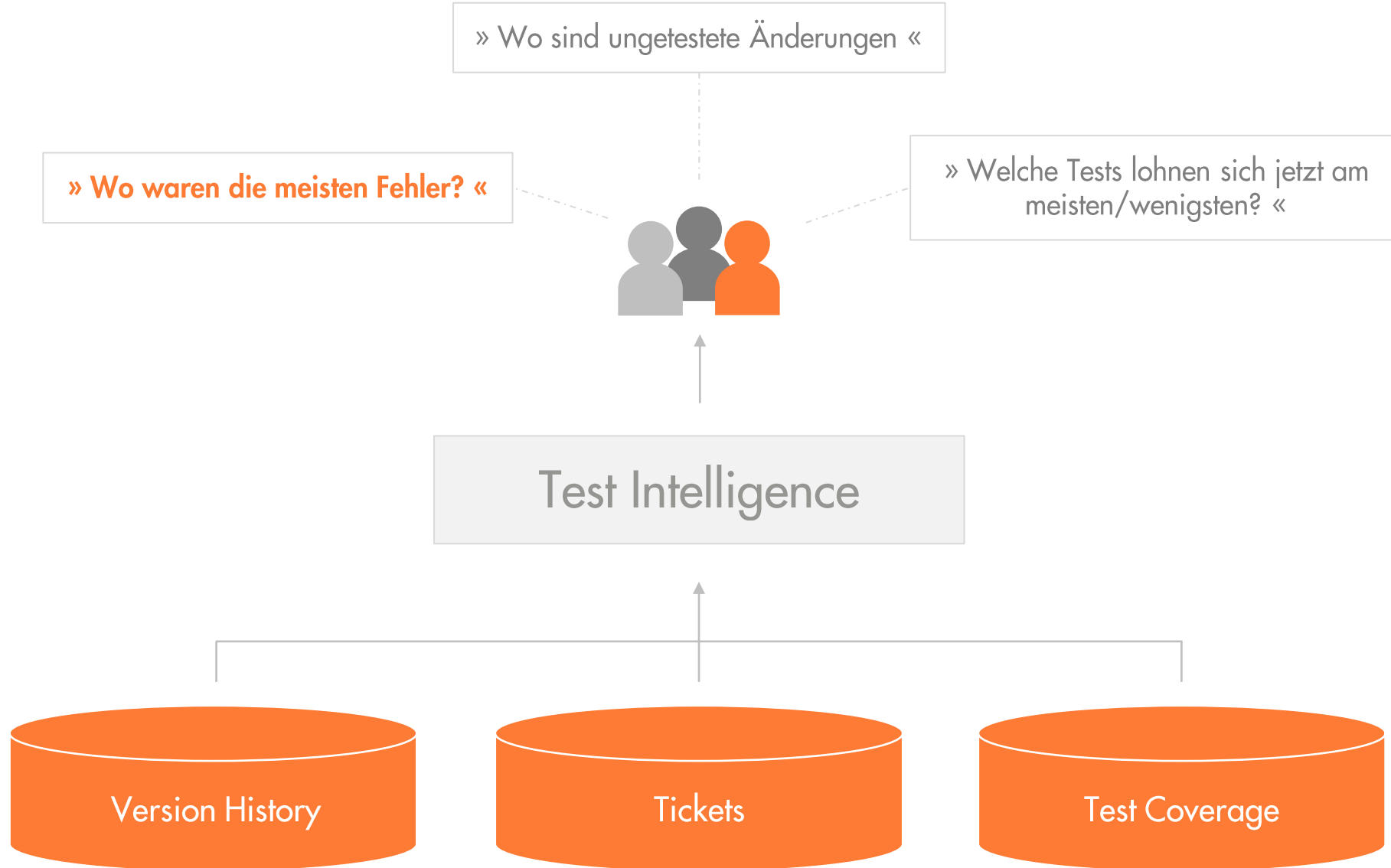
Dr. Elmar Jürgens. Mitgründer der CQSE und beschäftigt sich sein ganzes Berufsleben mit Qualitätsanalysen von Software.

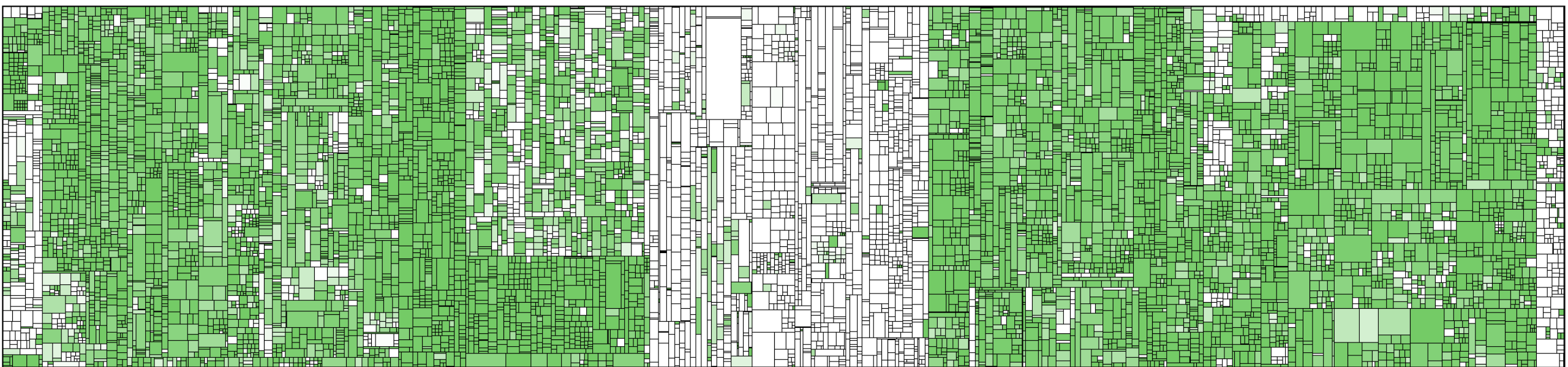
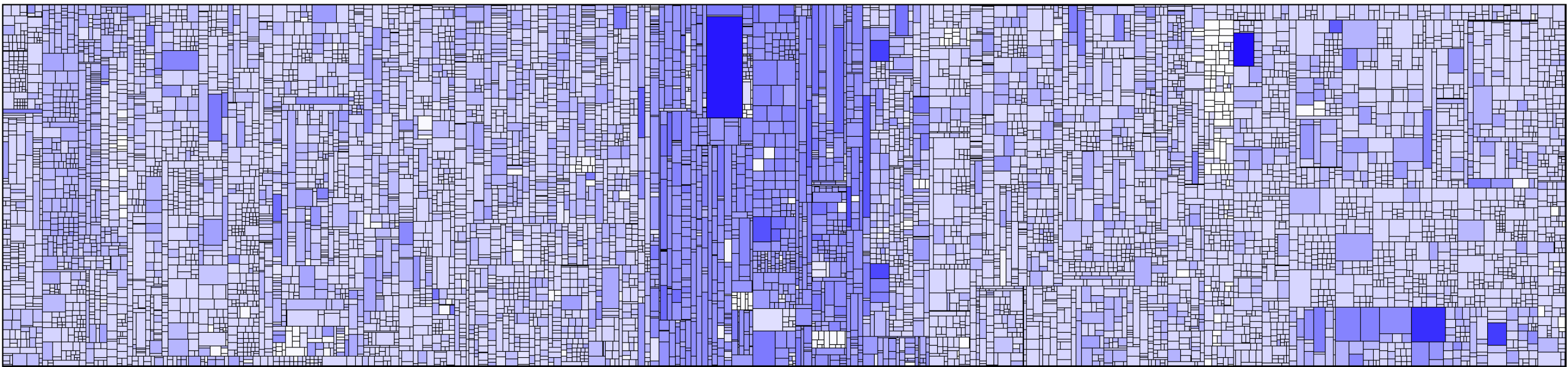
Fabian Streitl. Hat als Leiter des Piloten-Teams die TGA bei vielen Kunden erfolgreich eingeführt.

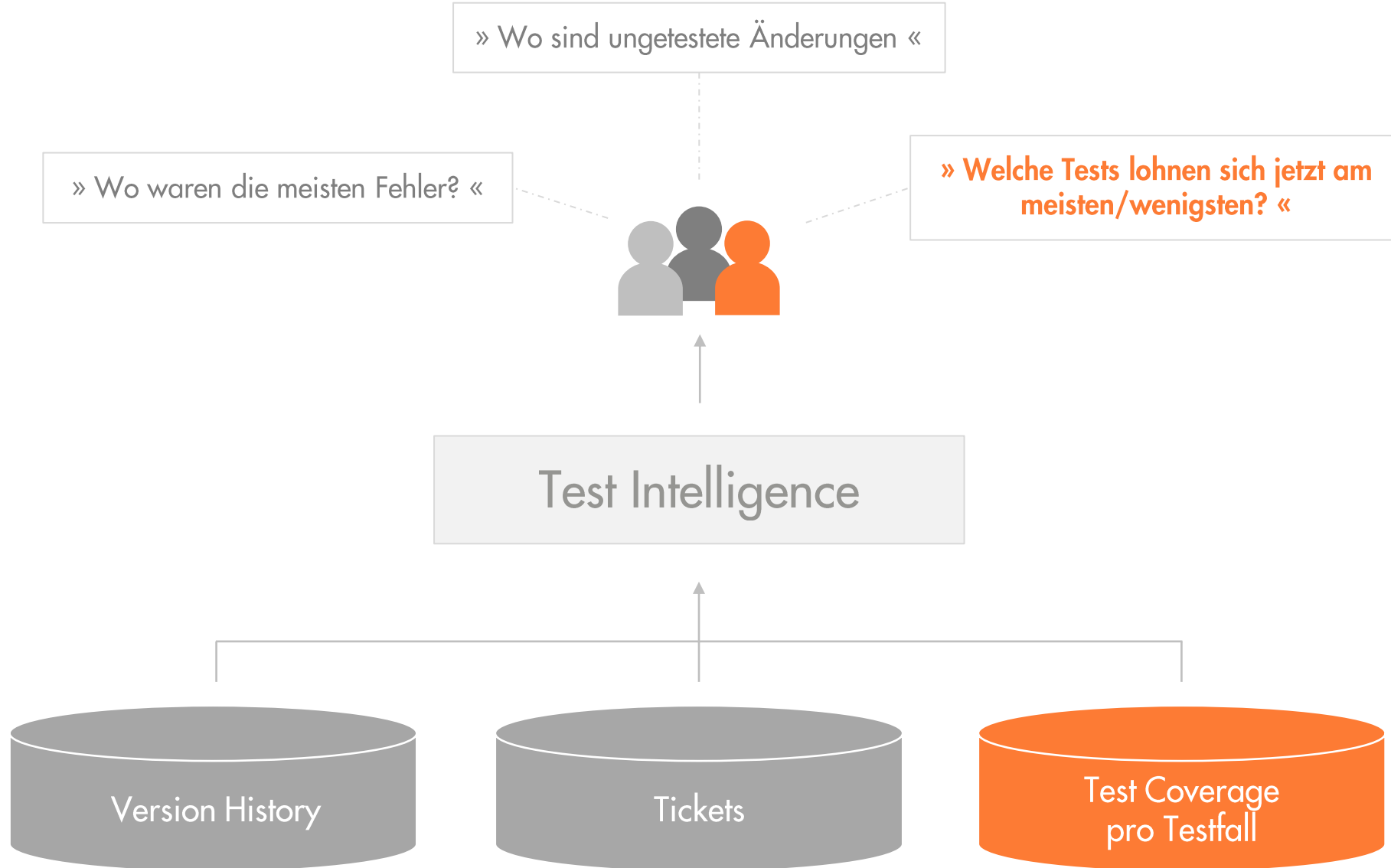
Jakob Rott. Setzt die TGA im Kundenumfeld täglich ein und hat viele Ausreden gehört, warum Sachen nicht getestet wurden.

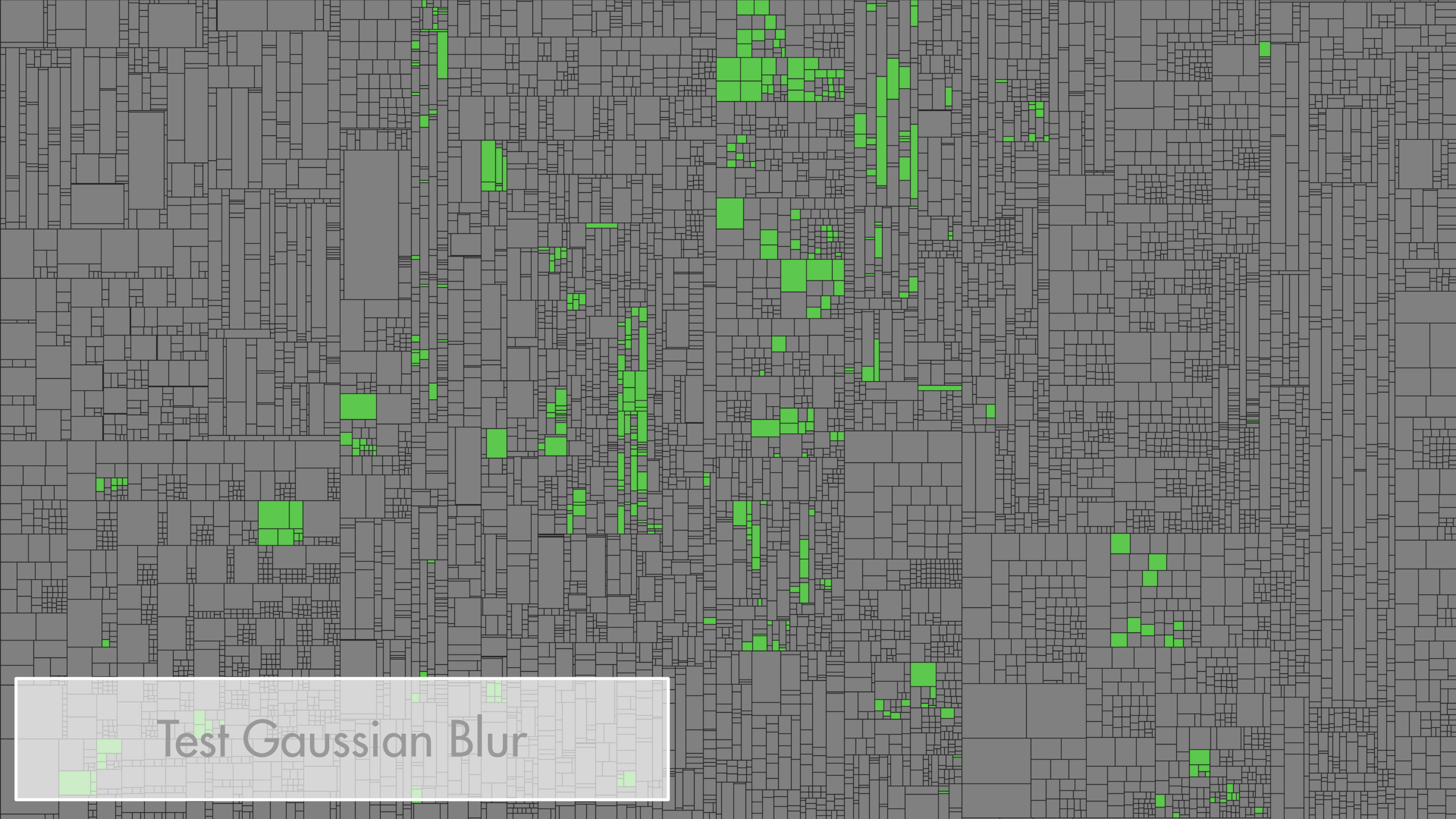
Einordnung

Der Workshop fasst mehrere Vorträge zusammen, die in den letzten Jahren auf dem *German Testing Day*, den *Software Quality Days* und dem *QS-Tag* mit Best Presentation Awards ausgezeichnet wurden. Anders als diese Einzelvorträge stellt er aber keine neuen Analysen oder neue Forschungsarbeiten vor, sondern gibt einen konsolidierten Überblick für diejenigen, die sich für das Thema *Test-Gap-Analyse* neu interessieren.





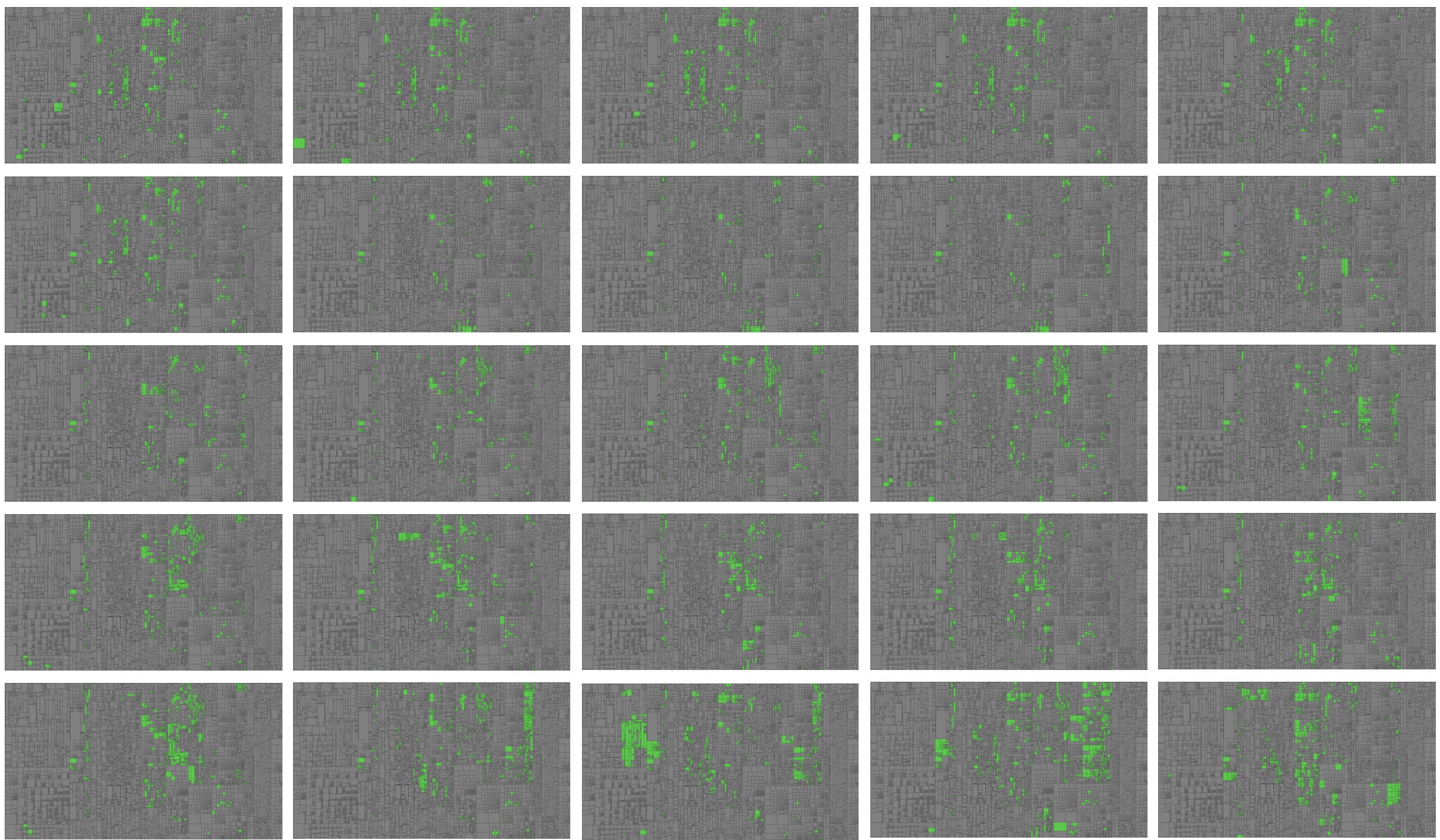




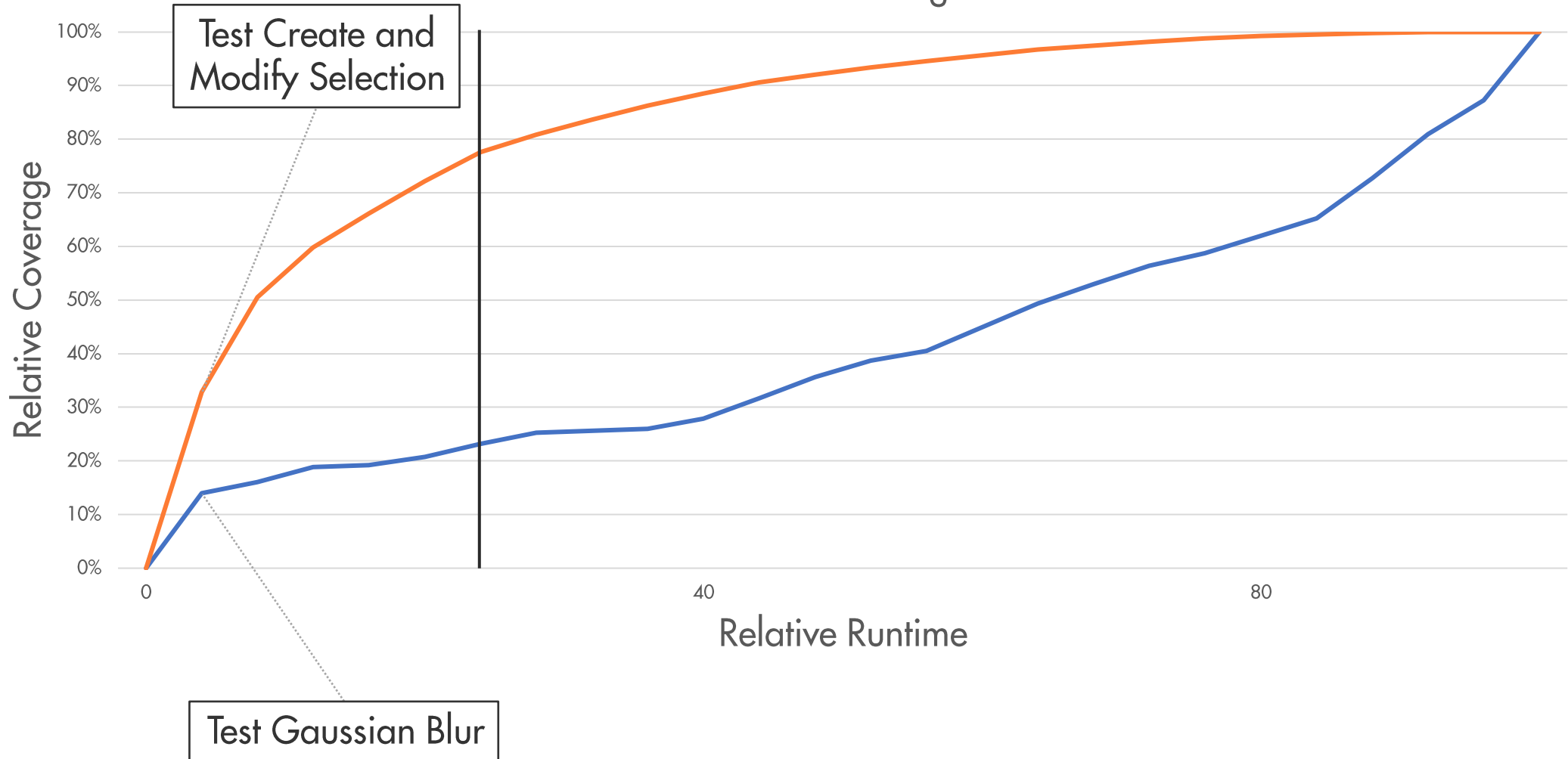
Test Gaussian Blur

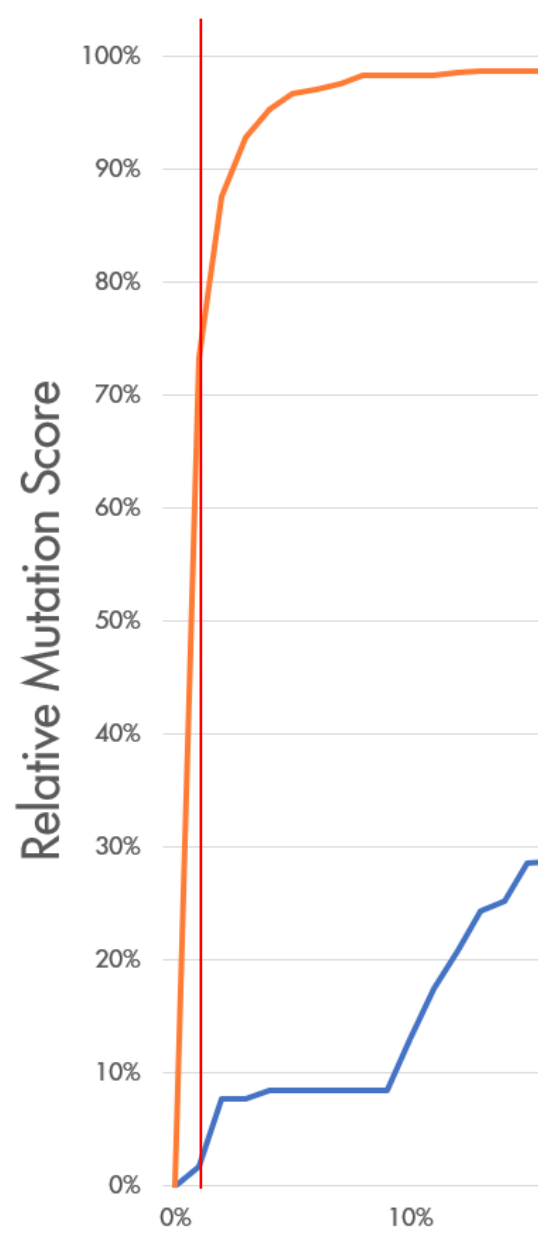


Test Create and Modify
Selection

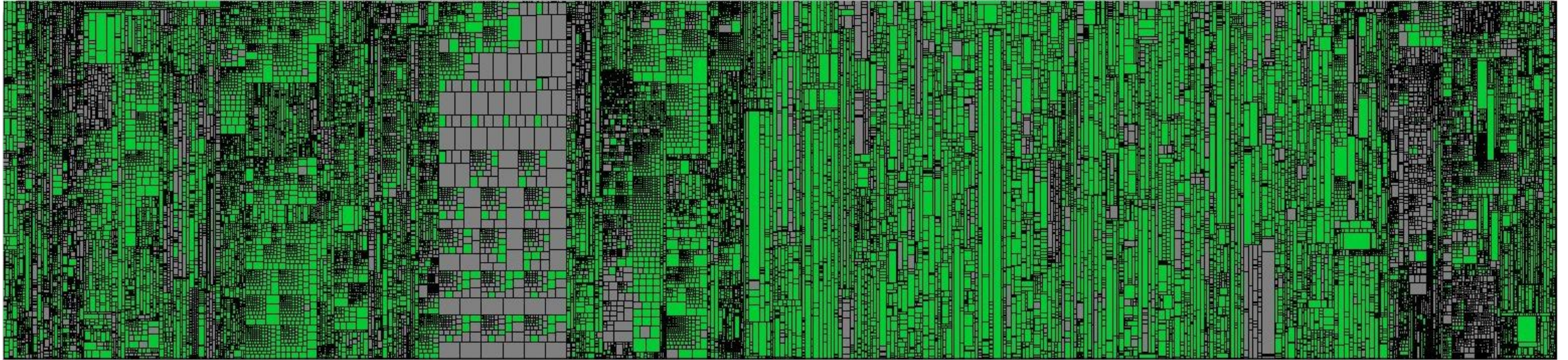


Time vs Code Coverage

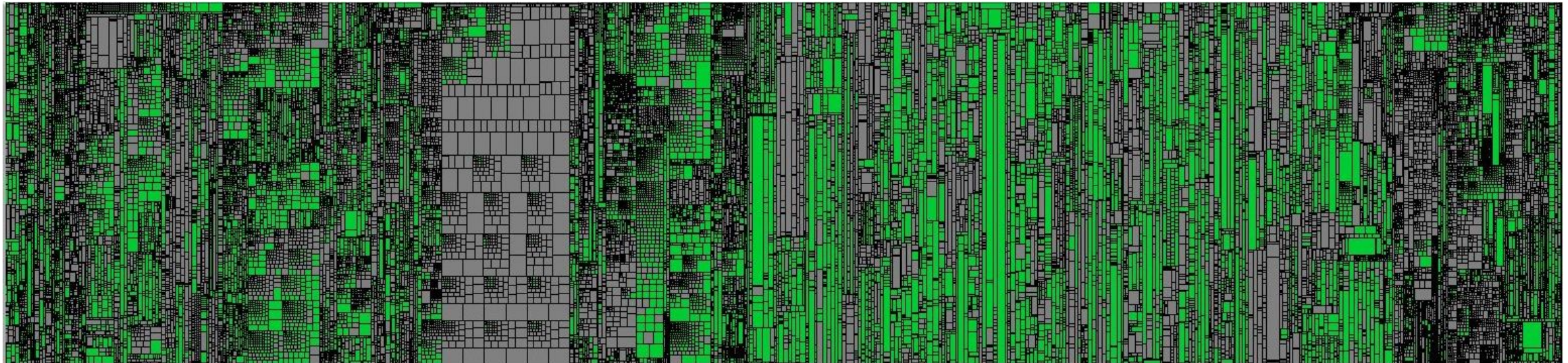




400+ Stunden Testausführung



1 Stunde Testausführung: >50% Coverage

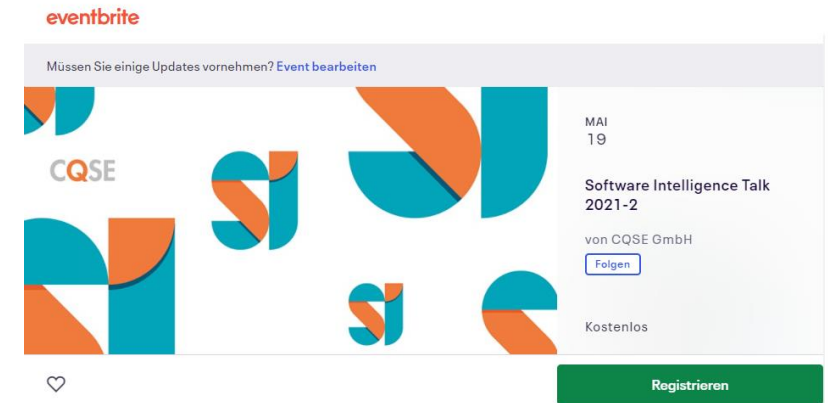


Virtueller Workshop

19. Mai 21, 10:30 – 12:00 Uhr

Anmeldung:

<http://cqse.eu/si-talks/2gtb>



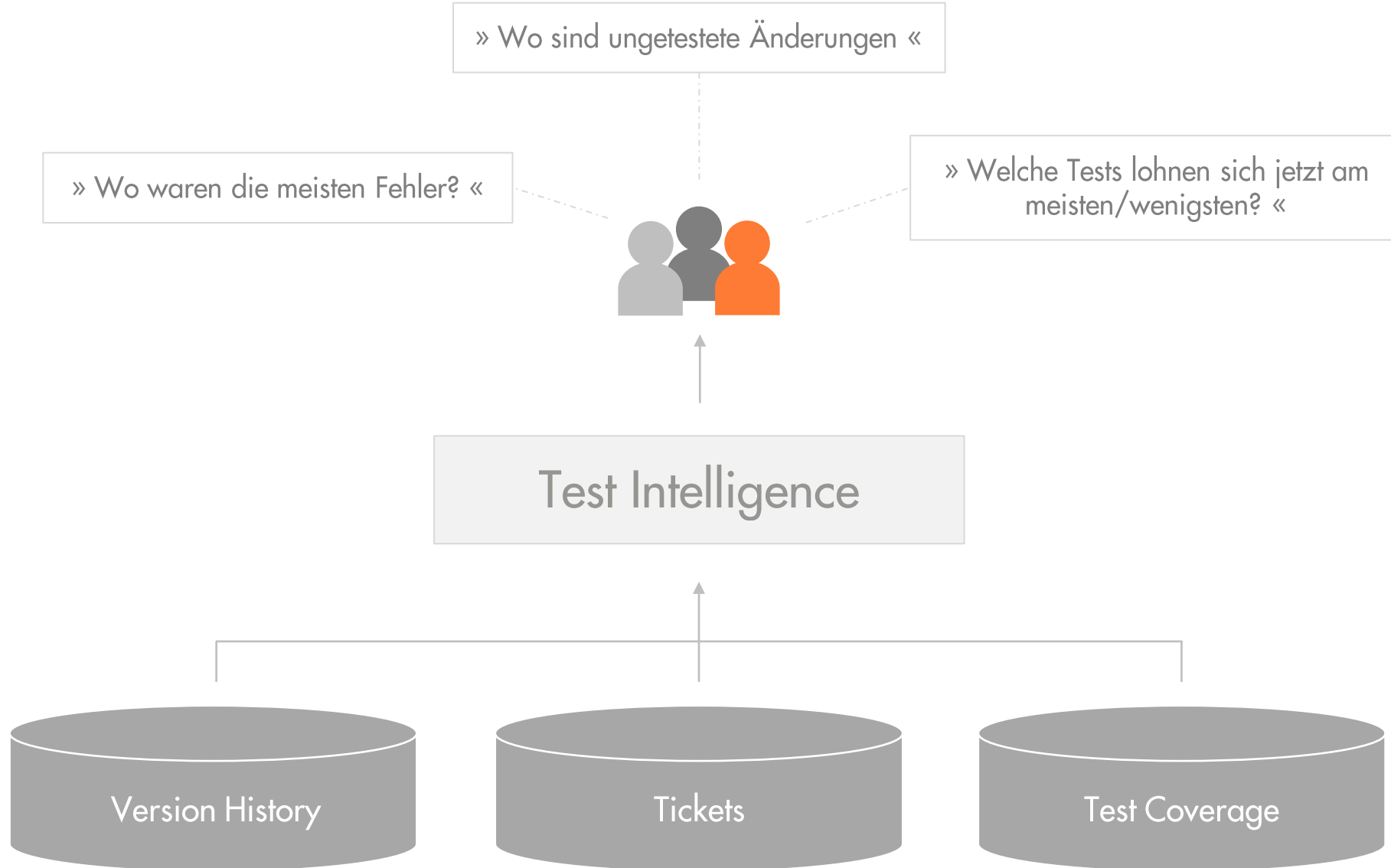
Zu den Sprechern

Jeannette Wernicke Jeannette Wernicke ist Innovation Managerin im Innovation Lab der Bayerische Versorgungskammer. Sie beschäftigt sich dort damit neue Technologien und Trends auf ihre Anwendbarkeit in der BVK zu untersuchen. Davor war sie als Software-Entwicklerin und -Architektin mit Schwerpunkt Software-Qualität tätig.

Raphael Nömmel hat seinen Master an der Technischen Universität München mit einem Fokus auf Software-Engineering und Software-Qualität abgeschlossen. Er hat seine Masterarbeit zu Test-Suite-Minimierung und Pareto-Testing erstellt und promoviert bei der CQSE in diesem Themenbereich.

Fabian Streitl hat als Leiter des CQSE Piloten-Teams die Test-Analysen bei vielen Kunden erfolgreich eingeführt.

Dr. Elmar Jürgens hat die CQSE mitgegründet und beschäftigt sich sein ganzes Berufsleben mit Qualitätsanalysen von Software.



Kontakt – Ich freue mich auf Fragen 😊



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