

Towards the Absence of Bugs

An Intelligent Combination of Static and Dynamic Analysis to Identify Vulnerabilities in the Development Process

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- Static and dynamic analysis
- Interactive analysis

Verification of static analysis findings

Residual Risk Estimation

Good-Turing Estimator (GTE)

Conclusion

Acknowledgments

This work was developed within the IntelliSecTest project. The IntelliSecTest project is funded with 3.5 million euros over three years by the Fraunhofer PREPARE program.

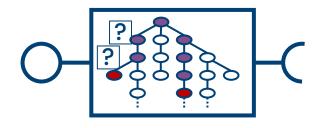


Testing of Trustworthy Systems

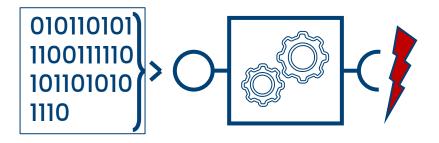
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Two major approaches

Static Analysis



Dynamic Analysis



Advantages

High path coverage
Good presentation of results

Very few false positives

 Provides input data triggering to vulnerability

Drawbacks

 High number of false positives

Random path coverage

Poor results presentation

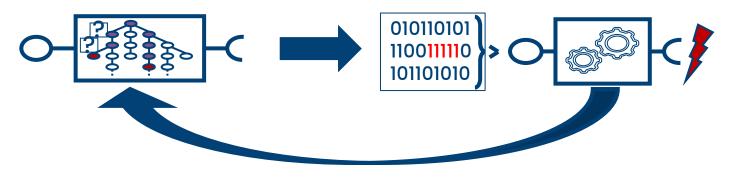






Interactive Analysis

Combination of static and dynamic Analysis





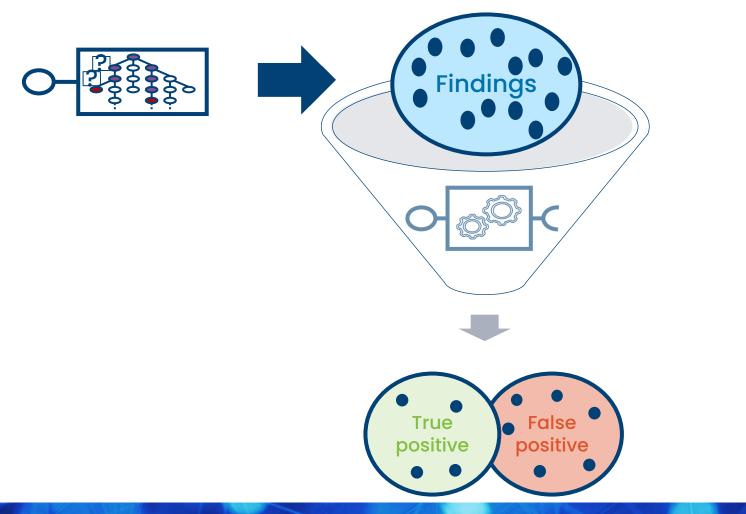
- High path coverage
- Good presentation of results
- Provides input data leading to vulnerability
- Reduce number of false positives





Verify static analysis findings





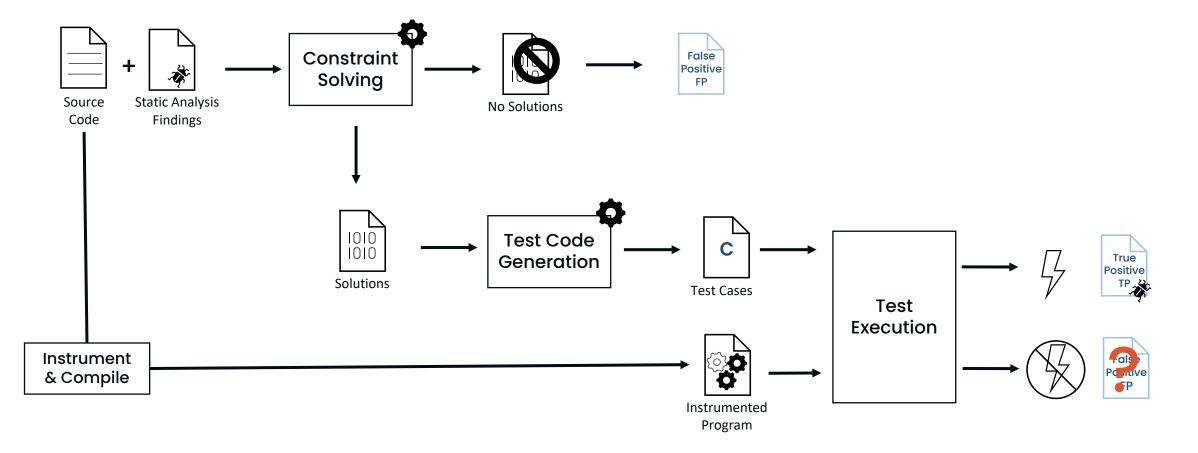




Verify static analysis findings



Create test cases using constraint solving



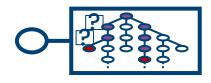


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Verify static analysis findings





Findings False positive

"Program testing can *be used* to show the presence *of* bugs, *but never* to show their absence!"

[E. Dijkstra]

→ Need a way to <u>estimate the</u> <u>residual risk</u> that there is an undiscovered vulnerability in the code







Traditionally probability calculation:

- Assumption: the ratio of every element in the set in relation to the occurrence in the sample set is universally true
- Result: no prediction for unseen elements

Good-Turing Estimation (GTE):

- Assumption: the sample data just captures a part of the set
- Consequents: probability discounting to create room for unseen elements (pseudo count)





Testing of Trustworthy Systems

Residual Risk Estimation

the sample is approximately

approximately

Missing mass estimation

"the chance that the next [...] sampled will belong to a new species is approximately"

$$P'_0 \approx \frac{n_1}{N}$$

 σ_0^\prime the probability for all unobserved species ("missing mass")

 n_r number of species that were seen exactly r times

N is the total number of counts

239

(7)

(8)

We may say that the proportion of the population represented by the sample is approximately $1 - n_1/N$, and the chance that the next animal sampled will belong to a new species is approximately n_1/N . (9) I.J.Good: THE POPULATION FREQUENCIES OF SPECIES AND THE ESTIMATION OF POPULATION PARAMETERS (1953)

I. J. GOOD

Hence also the expected total chance of all species that are represented r times or more in

 $N^{-1}\{(r+1)n_{r+1}+(r+2)n_{r+2}+\ldots\}.$

In particular, the expected total chance of all species represented at all in the sample is

 $N^{-1}(2n_2 + 3n_2 + \ldots) = 1 - n_1/N.$

elong to a new







GTE Applying to Fuzzing

"If no error has been exposed throughout the [fuzzing] campaign, the Good-Turing estimator gives an upper bound on the probability to generate a test input that exposes an error." [M. Böhme]

Residual Risk Estimation

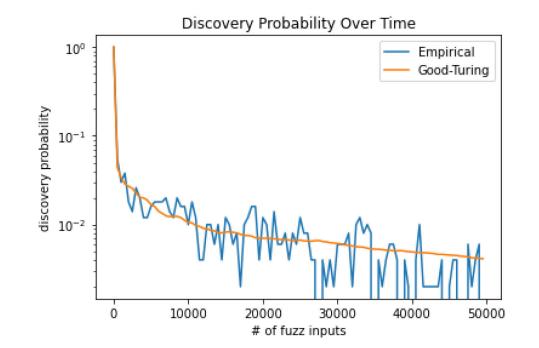
Empirical estimator

To measure the empirical probability, we execute the same population of inputs (n=50000) and measure in regular intervals (measurements=100 intervals). During each measurement, we repeat the following experiment repeats=500 times, reporting the average: If the next input yields a new trace, return 1, otherwise return 0. Note that during these repetitions, we do not record the newly discovered traces as observed. [1]

[1] https://www.fuzzingbook.org/html/WhenToStopFuzzing.html

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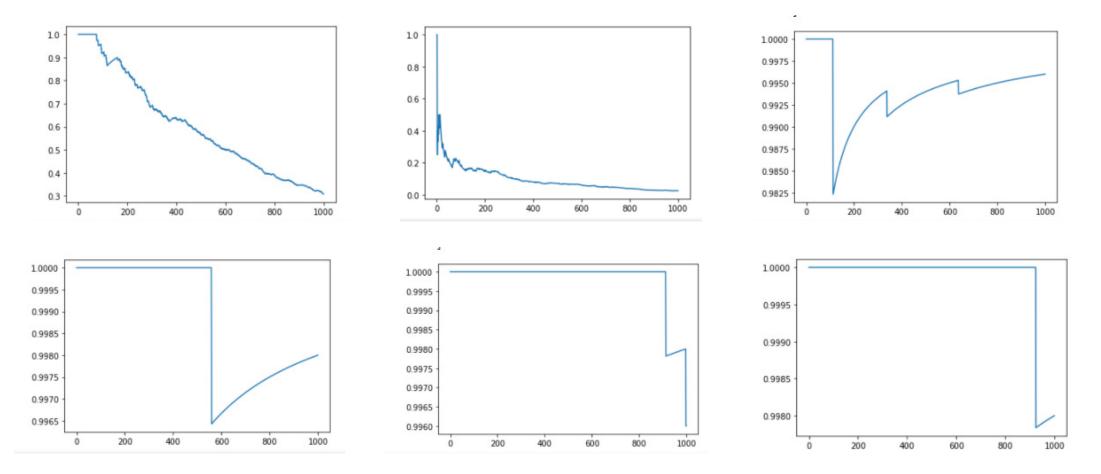






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GTE Applying to different examples

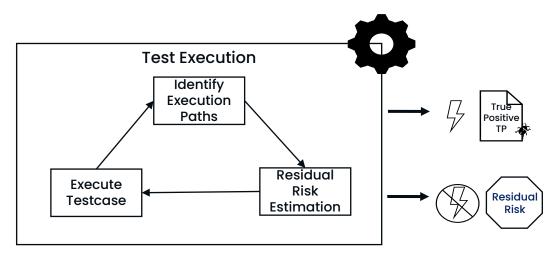




Residual Risk Estimation



Verification of Static Analysis Findings



When to stop the test execution?

- Use relative GTE Values (no absolute value)
- Consider a calibration period
- Monitoring the trend in GTE values across multiple test cases
- End test execution when no more significant changes are monitored

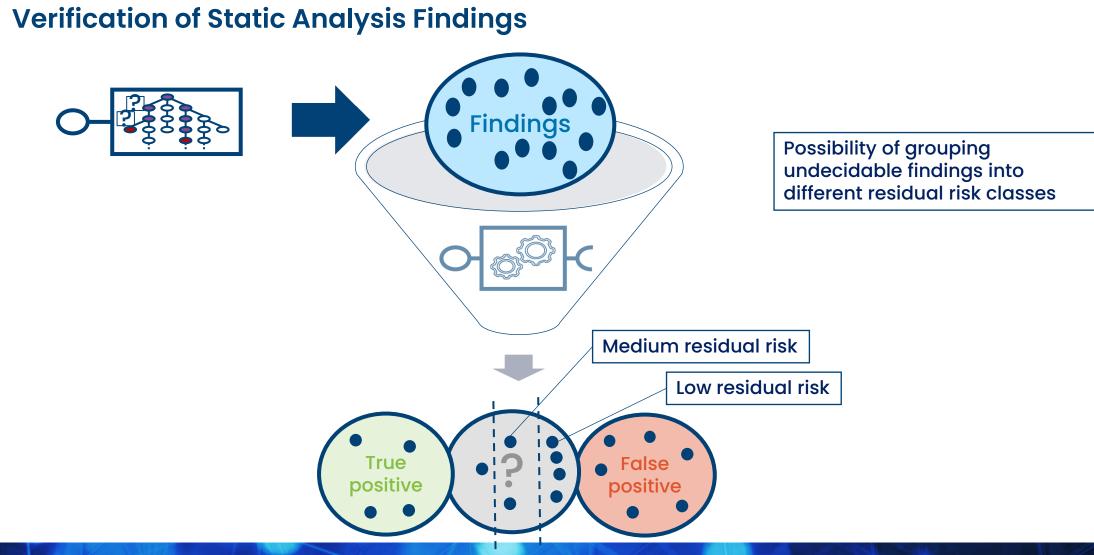




Residual Risk Estimation



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- Static and dynamic analysis can benefit from each other
- Dynamic analysis can be used to verify static analysis findings
- Good-Turing estimation can estimate the residual risk of a test campaign

Even if tests does not provide absolute evidence, a measure of evaluation can be provided to reduce the degree of uncertainty





Thank You! Any further questions?

Contact me: ramon.barakat@fokus.fraunhofer.de





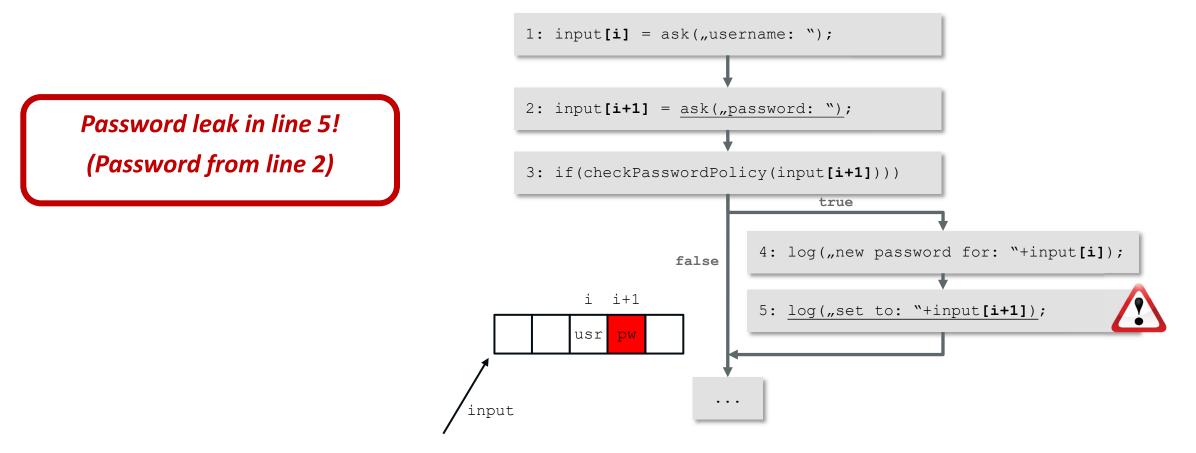


- Good, Irving J. "The population frequencies of species and the estimation of population parameters." *Biometrika* 40.3-4 (1953): 237-264.
- Böhme, Marcel. "STADS: Software testing as species discovery." ACM Transactions on Software Engineering and Methodology (TOSEM) 27.2 (2018): 1-52.
- https://www.fuzzingbook.org/html/WhenToStopFuzzing.html
- Gale, William A., and Geoffrey Sampson. "Good-turing frequency estimation without tears." *Journal of quantitative linguistics* 2.3 (1995): 217-237.
- Can Good-Turing Frequency Estimation Tell Us When to Stop Fuzzing? (Blog) <u>https://bshastry.github.io/2018/10/08/good-turing-fuzzing.html</u>









9th

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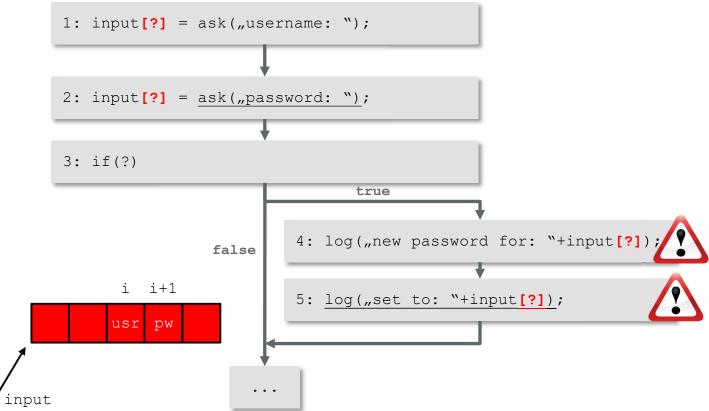


Common approximation:

Static Analysis

- Abstracting path constraints
- Abstraction of array indices





ETSI

Testing of Trustworthy Systems



9th

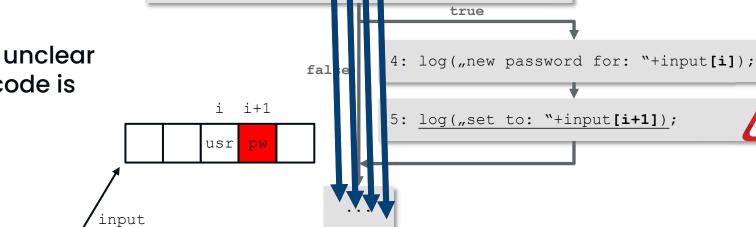
If password leak is observed, unclear • where the vulnerable line of code is

Tool-supported Security Testing today

Dynamic Analysis

- Probability to execute line 5 very low •
- Vulnerability remains undetected •

AND



3: if(checkPassworlFolicy(input[i+1])))

ssword: ");

2: input[i+1] = ask



Testing of Trustworthy Systems





9th



How does it work

GT Estimation:

$$P'_{r} = \frac{1}{N}(r+1)\frac{n_{r+1}}{n_{r}}$$

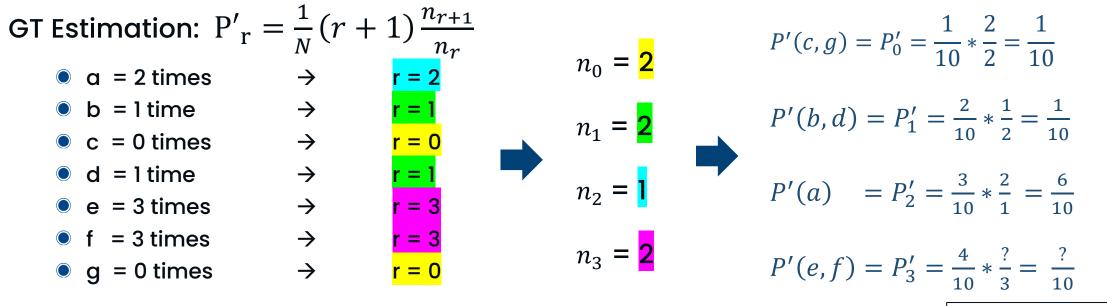
- P'_0 the probability for all unobserved species ("missing mass")
- P'_r the probability to observe r individuals for species X
- In the number of individuals that have been observed for species X
- n_r number of species that were seen exactly r times
- N is the total number of counts







How does it work Set: {a, b, c, d, e, f, g} Sample data: "aabdeeefff" (*N = 10*)



Use interpolation for higher counts or lacks



