An Investigation of Compression Techniques to Speed up Mutation Testing

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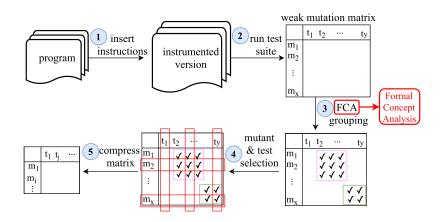
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our approach: ComMT

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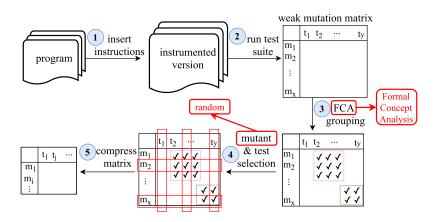
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Assumption: mutants which have high similarity in weak mutation are very likely to have the same outcome in strong mutation.



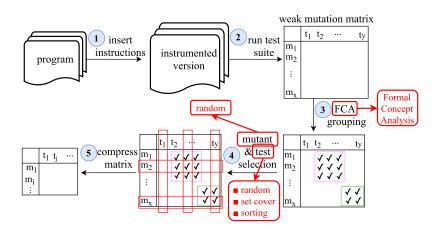
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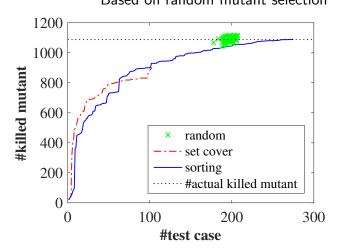


Experimental study

- 6 open-source projects
- automatically generated test suites (by Evosuite)
- comparison of test case selection random, set cover, sorting
- ComMT vs. other optimisations coverage-based, infection-based

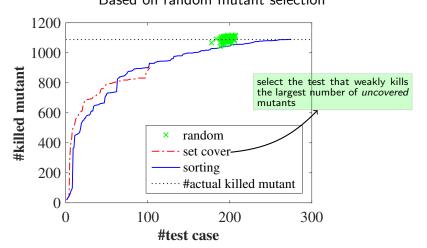
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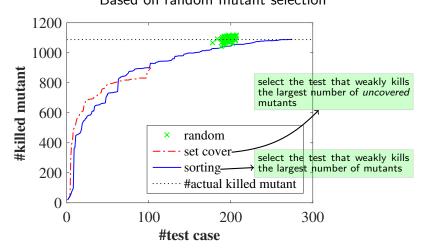
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ComMT vs. other optimisations

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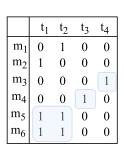
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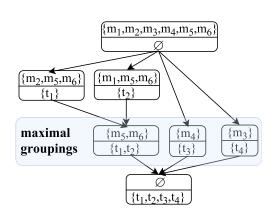
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- Implement on top of an existing mutation tool

FCA grouping





• Trade-offs between execution time reduction and error rate (%):

| Project | Cov. based | Inf. based | ComMT | | | | | | | |
|---------------------------------------|---------------|---------------|-------|---------|-----------|--------|--------|-------|---------|-------|
| · · · · · · · · · · · · · · · · · · · | | | No se | lection | Set cover | | Random | | Sorting | |
| baseline < | Exec. | Red. | Red. | Err. | Red. | Err. | Red. | Err. | Red. | Err. |
| jsecurity | 1.39 min | 16.78 | 87.71 | 0.13 | 90.27 | -17.42 | 88.55 | -2.46 | 87.31 | -0.04 |
| summa | 1.54 min | 13.57 | 90.97 | 0.74 | 93.58 | -14.1 | 92.18 | -2.25 | 90.87 | 0.69 |
| db-everywhere | 0.02 min | 3.65 | 59.29 | -0.26 | 80.81 | -18.83 | 70.43 | -6.22 | 63.95 | -0.26 |
| noen | 1.58 min | 6.09 | 88.52 | 0.30 | 91.08 | -30.69 | 89.52 | -7.29 | 87.98 | 0.14 |
| jtailgui | 0.31 min | 18.7 | 87.66 | 0.07 | 91.33 | -13.11 | 89.86 | -3.42 | 87.89 | 0.07 |
| caloriecount | 19.21 mir | 9.39 | 89.44 | 0.04 | 91.83 | -21.99 | 90.5 | -6.89 | 89.06 | -0.38 |
| Mean | - | 11.37 | 83.93 | 0.257 | 89.82 | -19.36 | 86.84 | -4.76 | 84.51 | 0.262 |