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Search Based Software Engineering for Variability Management

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Project Context



- FWF Lise Meitner Fellowship
 - Named after distinguished austrian nuclear fission scientist
 - Sponsored by the Austrian Science Fund (FWF)
- Fellowship duration: 2 years
 Start date: August 2012
- Personnel involved
 - Research fellow
 - Academic host: Alexander Egyed





- Give an overview of the project
 - ➤ Goals
 - Novel contributions
- Brief description of ...
 - Early results
 - Ongoing and upcoming work

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Background – Feature Models

Feature models

de facto standard to model variability

> denote sets of "valid" feature combinations





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Background – Software Evolution



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Reverse Engineering

Process of analyzing a software system to identify its components and their relationships with the goal of creating a higher level abstraction of them

Software Evolution

Process of progressive changes to the software artifacts or their properties

Background – Consistency Checking





Consistency checking

- Verifies that artifacts adhere to consistency rules that describe the semantic relationships among elements
- Example. UML consistency rule

Message action must be defined as an operation in receiver's class.













Reverse Engineering 2 **VARIABIL** Consistency Software Checking **Evolution**













Problems Addressed

Problem 1



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Fixing inconsistencies in the presence of variability



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Products

Problem 1 – Example



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Reverse Engineering of Variability

Most common scenario from products variants to a SPL



Search Based Variability Mining

Problem 3



Variability Evolution

Adding new features, new members of the product family, modifications

Feature Model





Search Based Feature Oriented Refactoring





Early Results

Reverse Engineering Feature Models from Product

Configurations (SBSE 2012)

Collaboration with University of Seville

David Benavides, Jose Galindo, Sergio Segura, Jose Parejo



Problem Pictorial View



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Feature Sets



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ETHOM Structural Encoding Example



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ETHOM CTC Encoding Example

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Crossover — One point



Crossover — One point (2) Result



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E,3,6	R,2,6
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Experimental Setting







Evaluation





Case studies

59 feature models from SPLOT repository
No. products 1...896
No. features 9 ... 27

Executions

- Initial populations for each feature model were the same for the fitness functions being analysed
- 10 runs for each feature model for each fitness function
- > 16 cores at 2.40 GHz, 25GB RAM, Cent OS, Java 1.6





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Relaxed Fitness Function – maximized FFRelaxed(sfs, fm) = | {fs : sfs | validFor(fs,fm) } |

- sfs = set of desired feature setsfm = feature model to evaluate fs = a feature set
- Auxiliary function validFor
 checks if a feature set is valid in a FM
 computed with FAMA using propositional logic
- Maximizes containment of desired feature sets

FFRelaxed Results (1)



SFA



feature sets



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maximum = cardinality of the desired feature sets



FFRelaxed Results (3)



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Surplus(sfs,fm) = $\frac{\text{#products(fm) - |sfs|}}{|sfs|} \times 100$







- Analyzing other fitness functions
 Finer comparison granularity
- Comparison with local search approaches
 Extension to HeuristicLab platform
- Studying variability-aware chromosome operators
 Crossover and mutation
- Extensions to feature model encodings
 Based on genetic programming









http://www.sea.uni-linz.ac.at/sbse4vm/



Acknowledgements





Consistency and Composition for Managing Variability in Multi-View Systems





