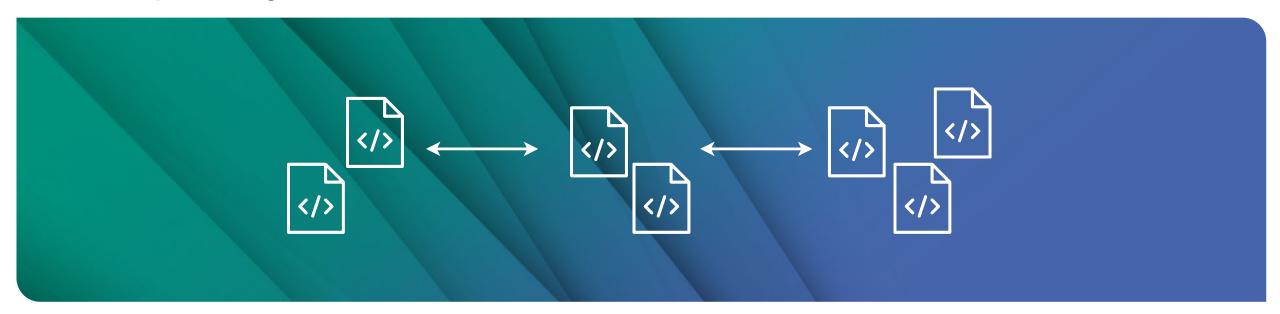




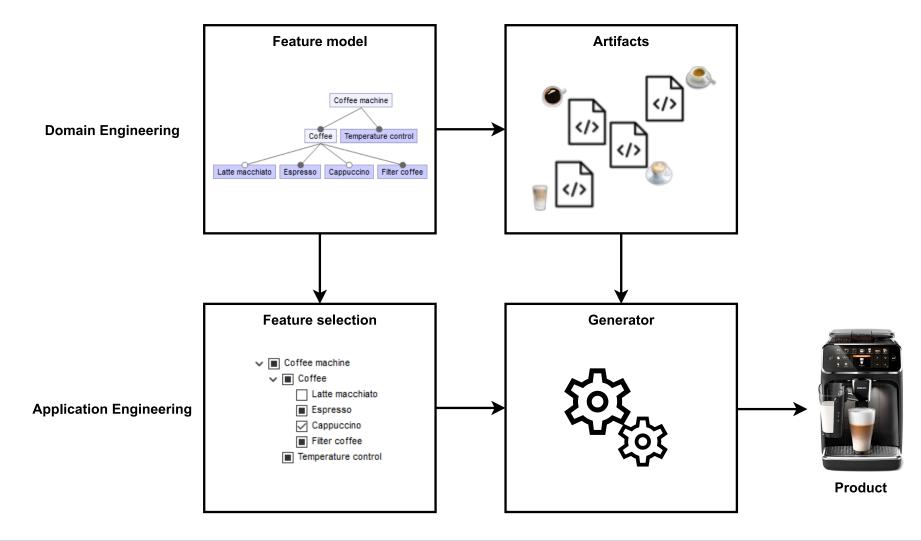
Synchronizing software variants: A two-dimensional approach

Christoph König, Kamil Rosiak, Lukas Linsbauer, Ina Schaefer



Software Product Lines





2

Horizontal cloning



Basic



- + Filter coffee
- + Espresso
- + Temperature control

Pro 100



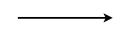
- + Filter coffee
- + Espresso
- + Cappuccino
- + Temperature control

Pro 250

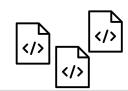


- + Filter coffee
- + Espresso
- + Latte Macchiato
- + Cappuccino
- + Temperature control









Vertical cloning











Latte Macchiato



- Heat milk
- Grind coffee
- Brew espresso
- Foam milk
- 6.



Heat water

- Heat milk
- Grind coffee
- Foam milk
- 5. Brew espresso
- 6.

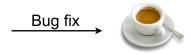


Synchronizing software variants



Basic







Pro 100









Pro 250





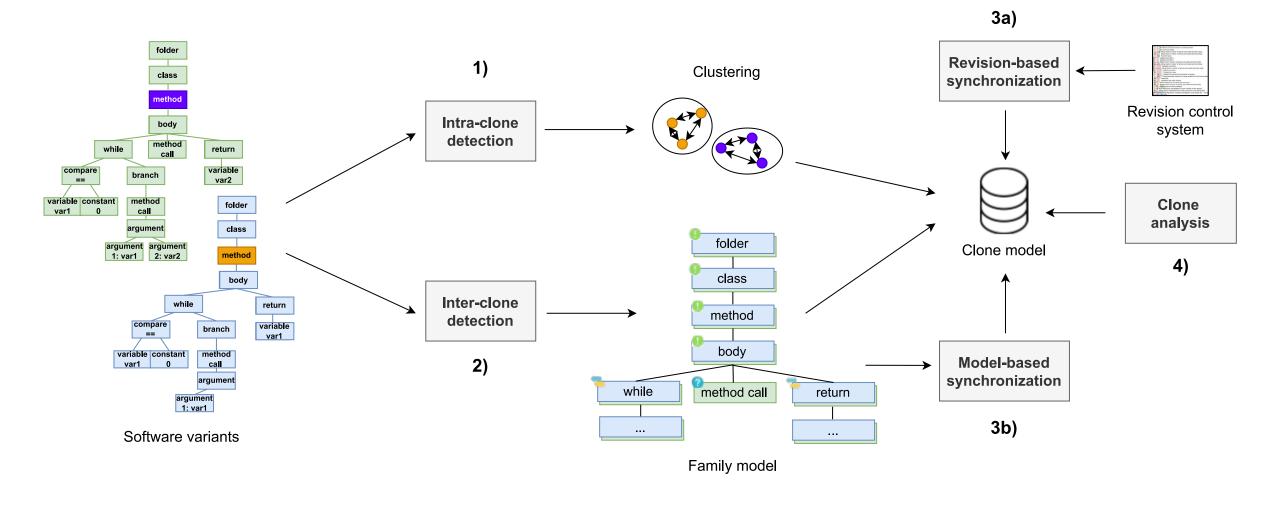






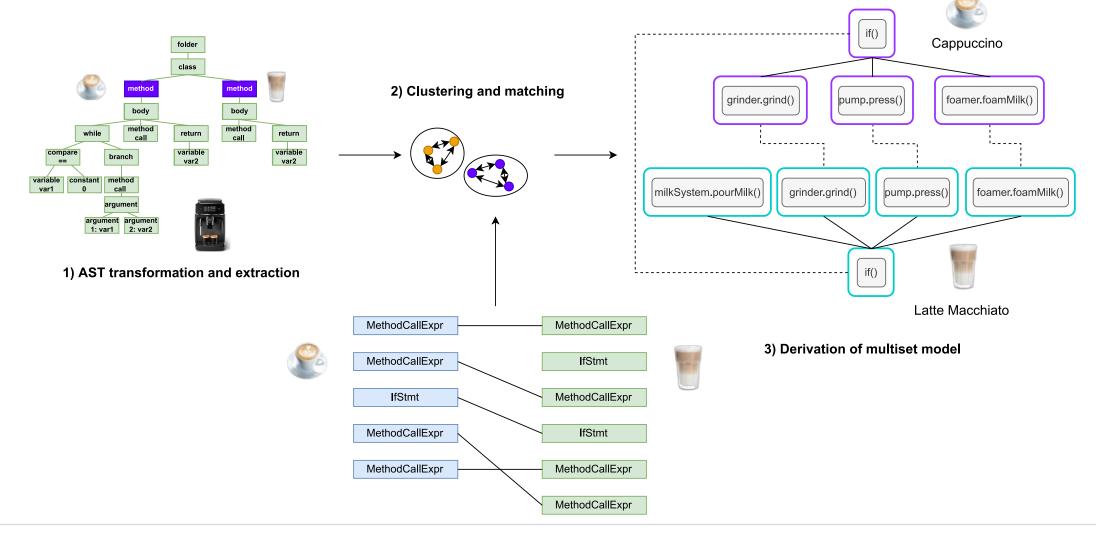
Overview





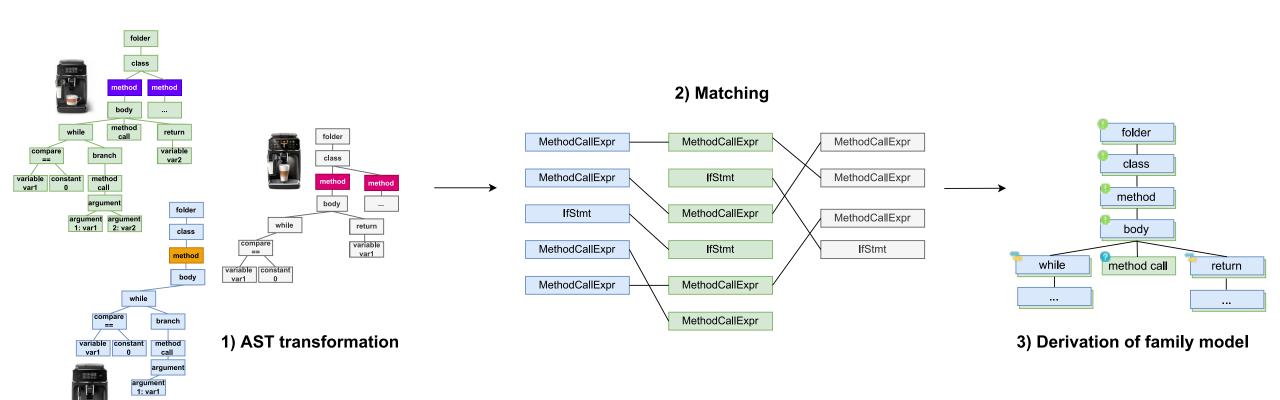
Intra-clone detection





Inter-clone detection

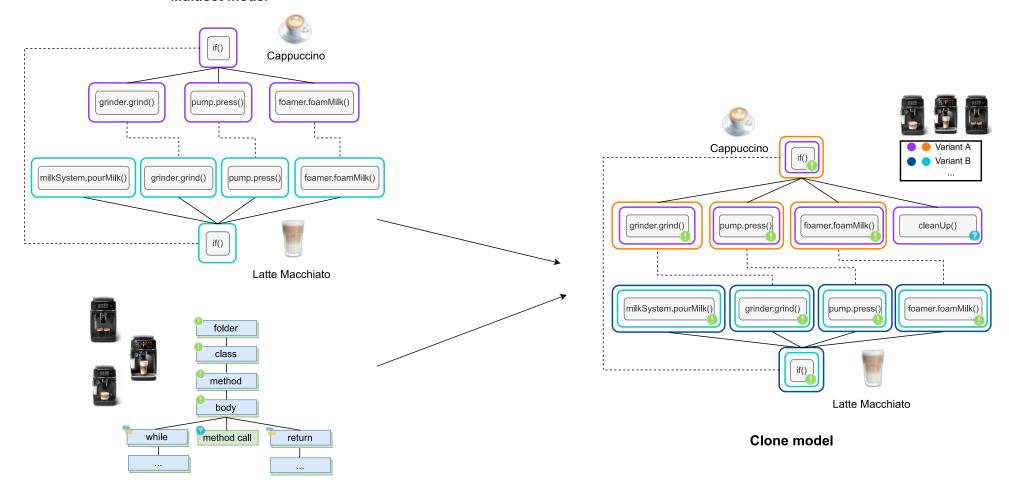




Clone model



Multiset model



Family model



Evaluation



- RQ1: How relevant is two-dimensional clone synchronization?
- RQ2: How correctly can we synchronize intra- and inter-clones?
- RQ3: How does synchronization scale in time and space with increasing number of variants?

Evaluation



Experimental subject: Five revisions of MobileMedia SPL¹

Metrics

- Co-occurrence: Relationship between two inter-clones c₁ and c₂, where c₁ has at least one intra-clone c_{1a}
- Subsequent revisions R_i and R_i (i < j)</p>
 - True positives (TP): Clones manually synchronized in R_i and two-dimensionally in R_i
 - False negatives (FN): Clones manually synchronized in R_i but not two-dimensionally in R_i
 - False positives (FP): Clones not manually synchronized in R_i but two-dimensionally in R_i
 - Precision (PR): $\frac{TP}{TP + FP}$
 - Recall (RC): $\frac{TP}{TP + FN}$

¹ Eduardo Figueiredo, Nelio Cacho, Claudio Sant'Anna, Mario Monteiro, Uira Kulesza, Alessandro Garcia, Sergio Soares, Fabiano Ferrari, Safoora Khan, Fernando Castor Filho, and Francisco Dantas. Evolving software product lines with aspects

RQ1: Relevance



		#Intra-clones			#Inter-clones					
Revision	#Variants	T_{I}	T_{II}	T_{III}	T_I	T_{II}	T_{III}	∅ Intra-group size	∅ Inter-group size	#Co-occurrence
R_1	1	1	1	1	0	0	0	4.40	0	0
R_2	2	2	4	2	86	0	5	4.00	2.00	24
R_3	4	4	8	4	88	2	6	4.00	3.92	48
R_4	8	8	8	52	104	2	7	3.19	7.61	172
R_5	16	16	44	100	134	2	10	3.30	13.62	424

The number of intra- and inter-clones per clone type, the average group size and the number of co-occurrences in each revision of MobileMedia

RQ2: Correctness



		Int	ra-clo	nes		Inter-clones				
	TP	FP	FN	PR	RC	TP	FP	FN	PR	RC
R_1	92	3	6	0.97	0.93	94	0	0	1.00	1.00
R_2	146	5	26	0.97	0.85	211	87	0	0.71	1.00
R_3	419	31	153	0.93	0.73	1308	352	4	0.78	0.99
R_4	404	111	28	0.79	0.94	2040	804	0	0.72	1.00

Precision and recall of variant synchronization between subsequent revisions of MobileMedia

Threats to validity



- Bias of matching algorithm
- MobileMedia as synthetic clone-and-own experimental subject
- Java source code only

Summary



- Approach: Synchronizing intra- and inter-clones simultaneously
- High precision and recall in change propagation
- Relevance of two-dimensional software variant synchronization