

Scalable N-Way Model Matching Using Multi-Dimensional Search Trees



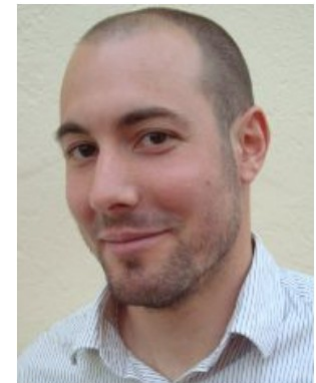
Alexander Schultheiß



Paul Maximilian Bittner



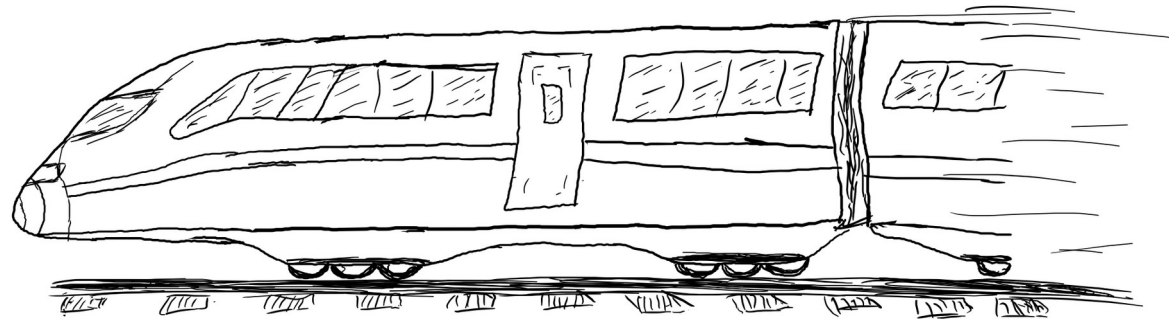
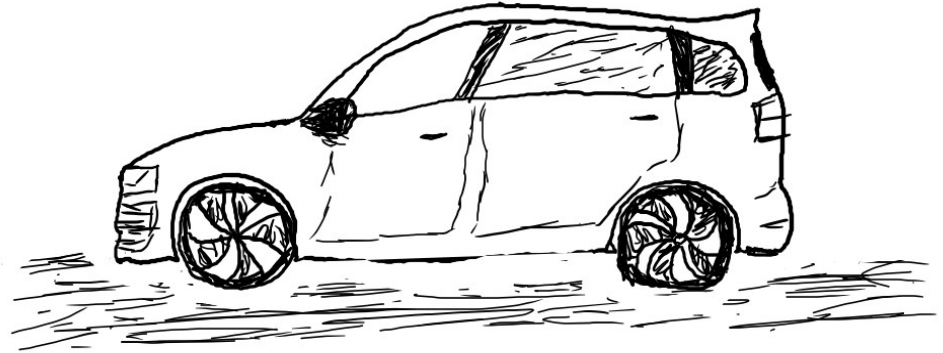
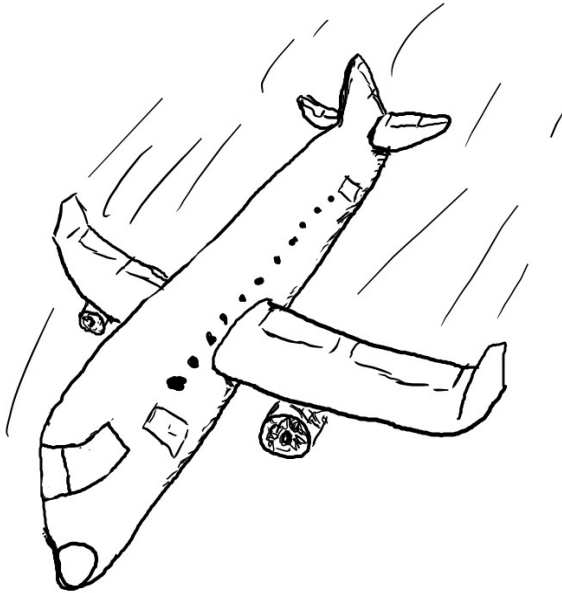
Thomas Thüm



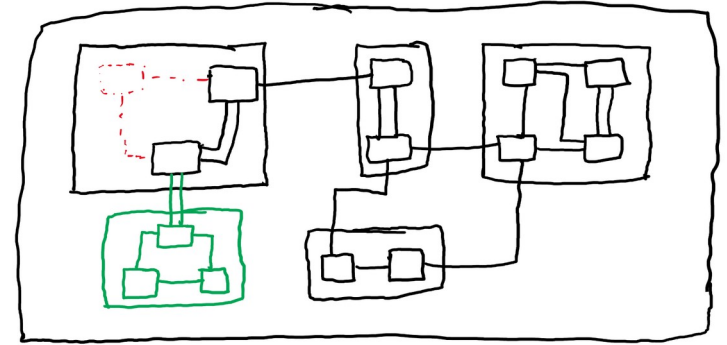
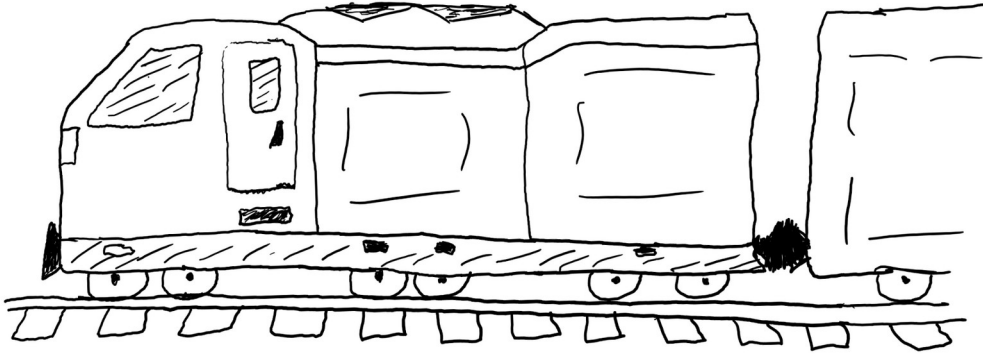
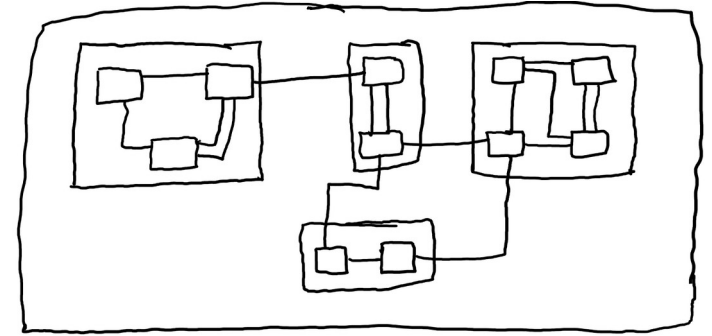
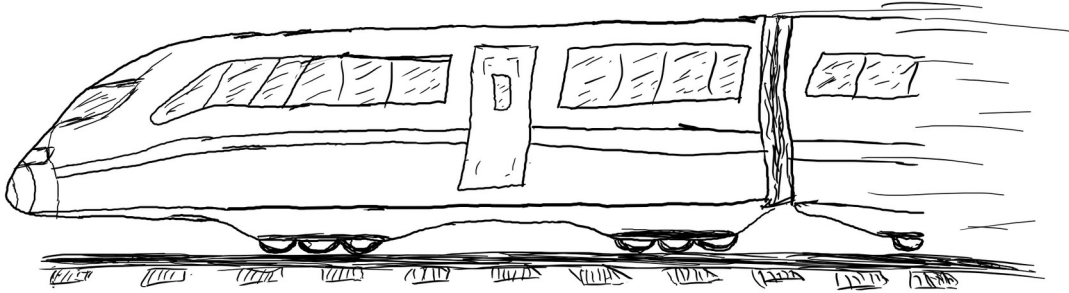
Timo Kehrer

Why do we need model matching?

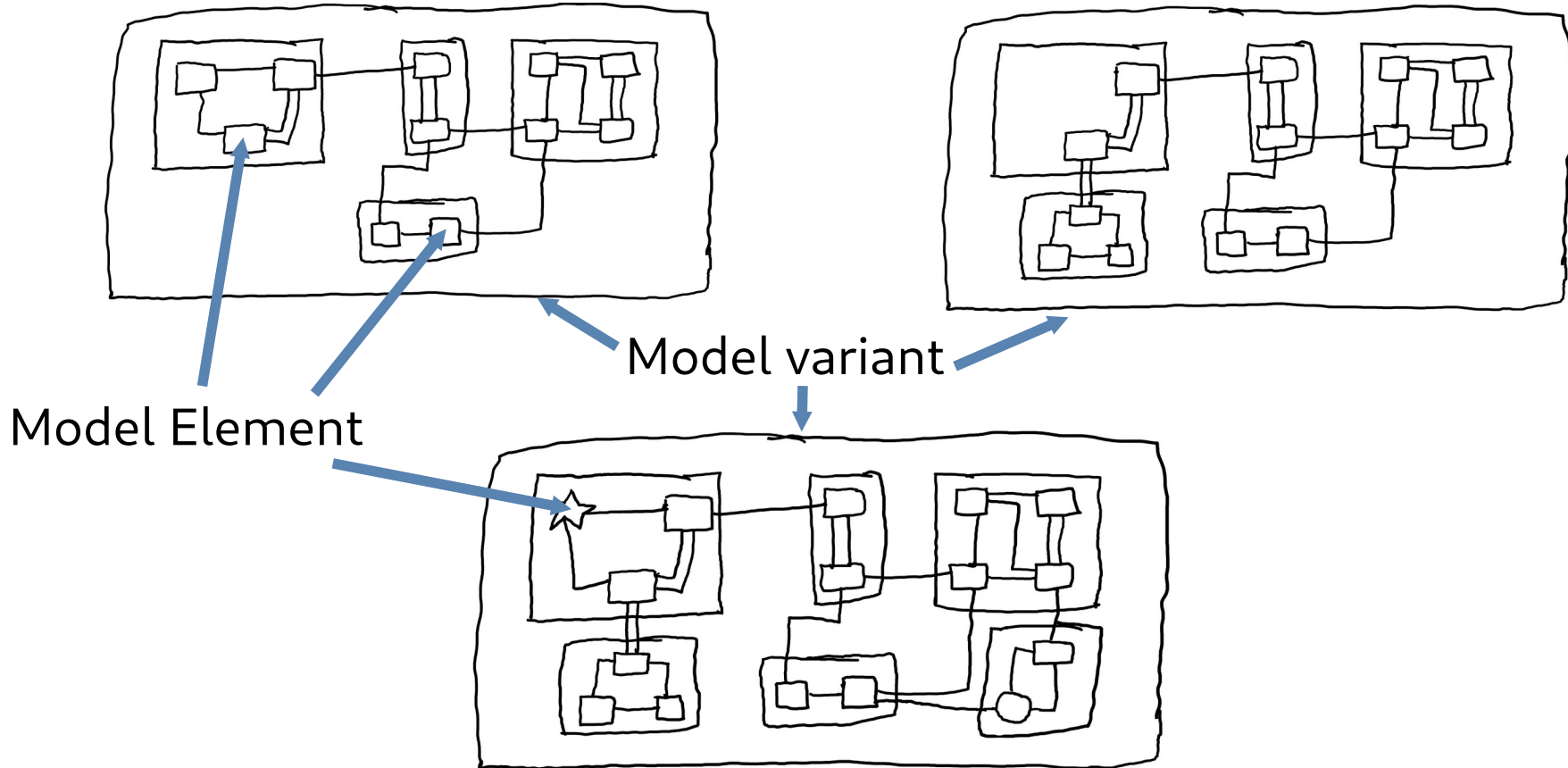
Model-driven development is crucial in certain domains



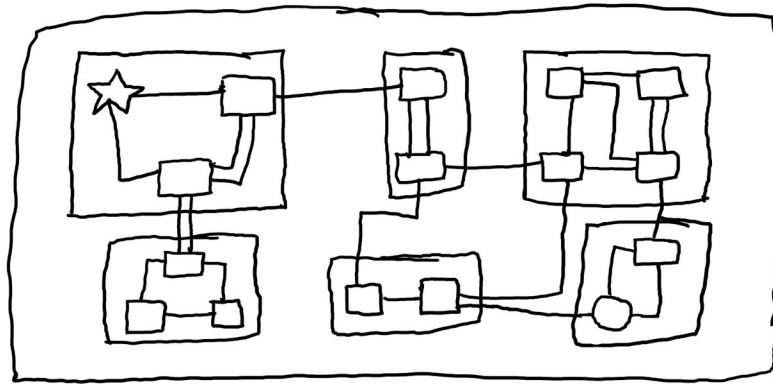
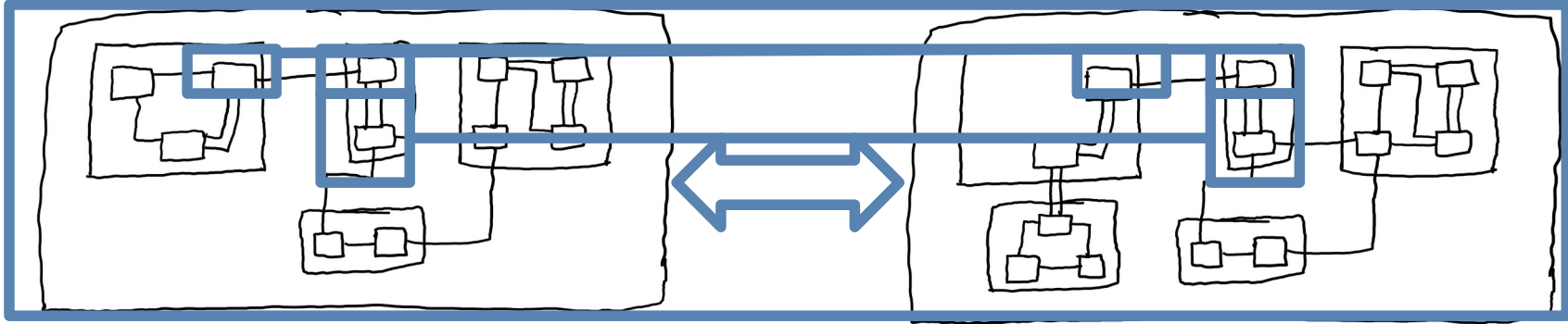
Different variants of a model are required



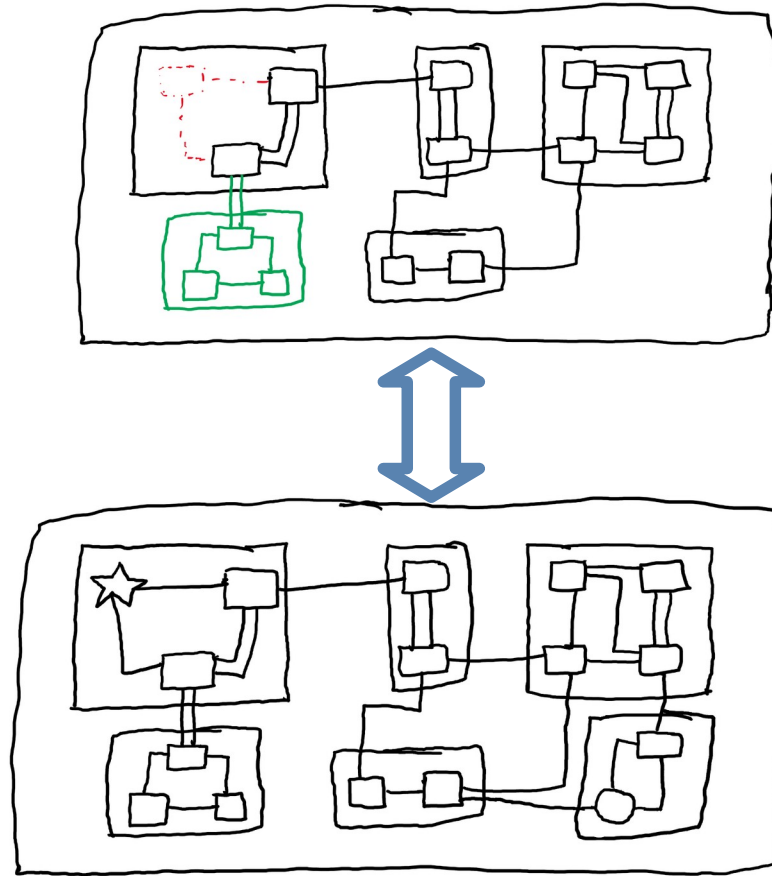
We want to match common elements



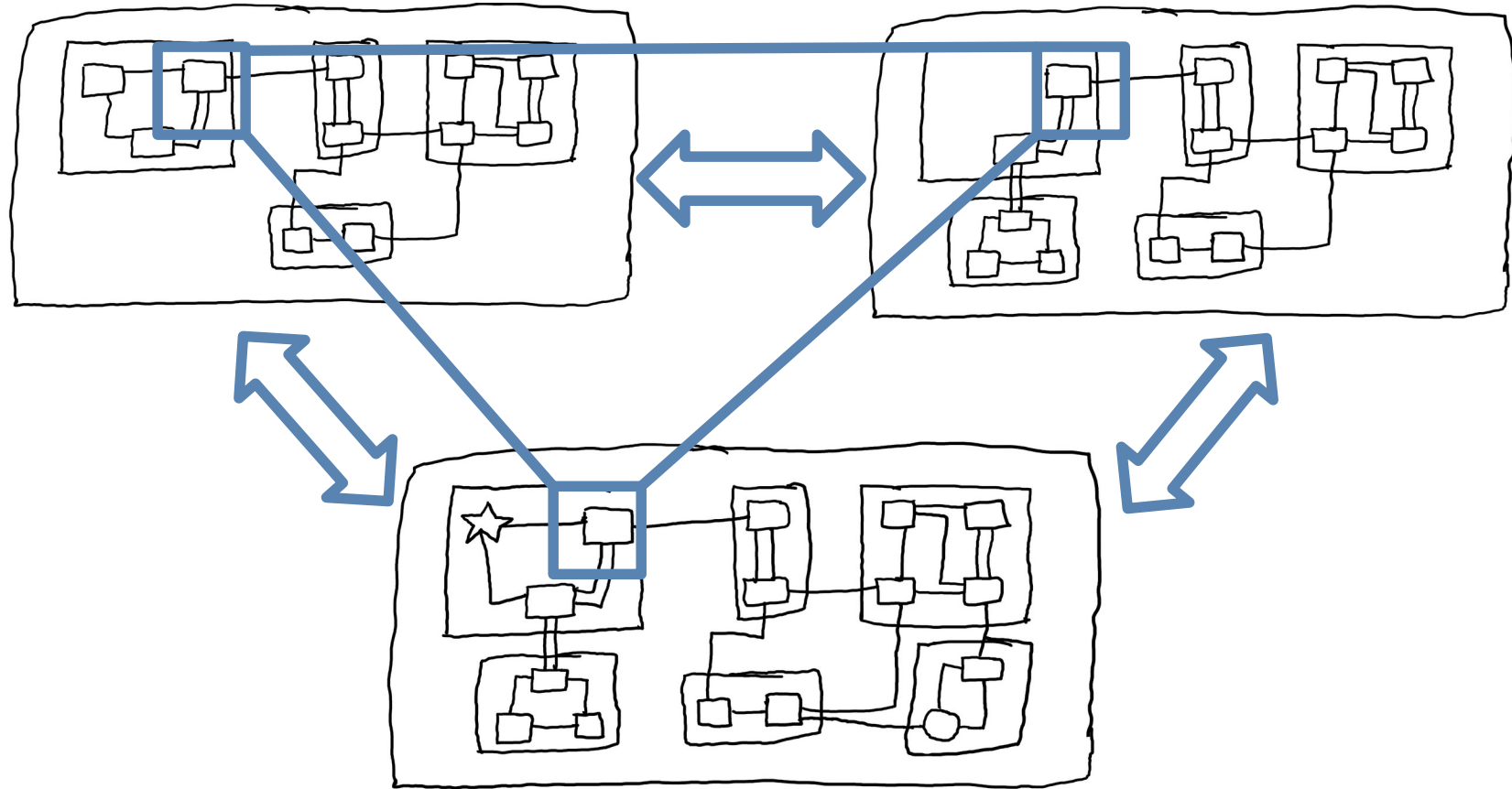
Pairwise matching considers models one-by-one



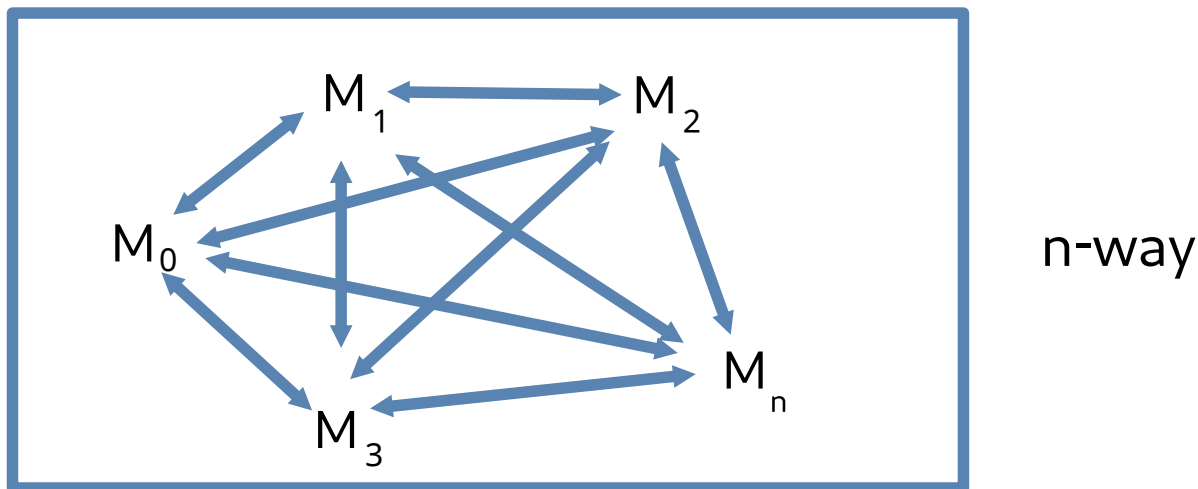
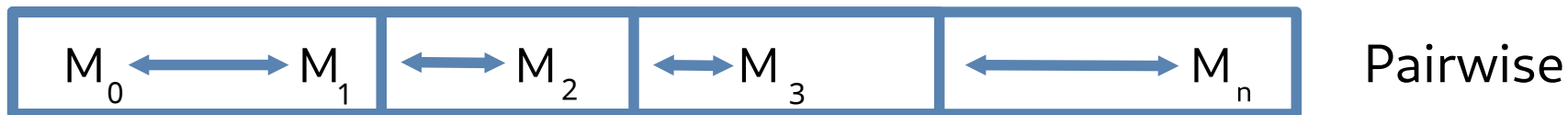
Pairwise matching considers models one-by-one



N-way matching considers all models at once

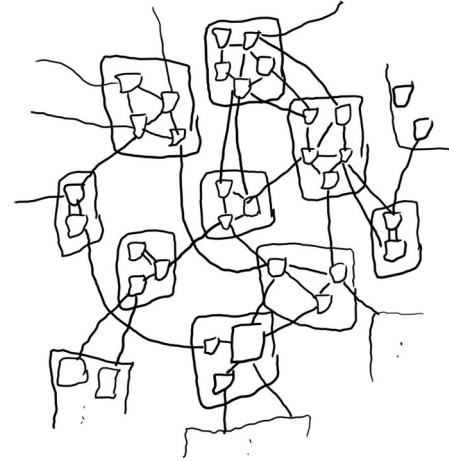
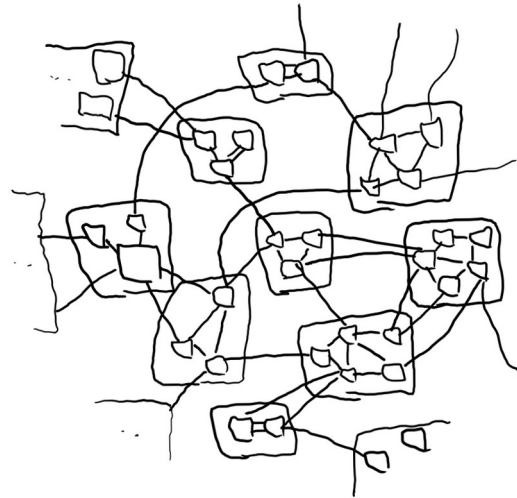
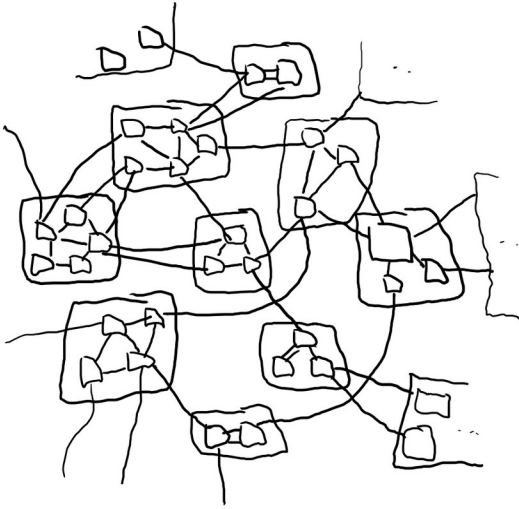


Pairwise and n-way in direct comparison



Why do we need **scalable** n-way model matching?

Models can be large



We present RaQuN

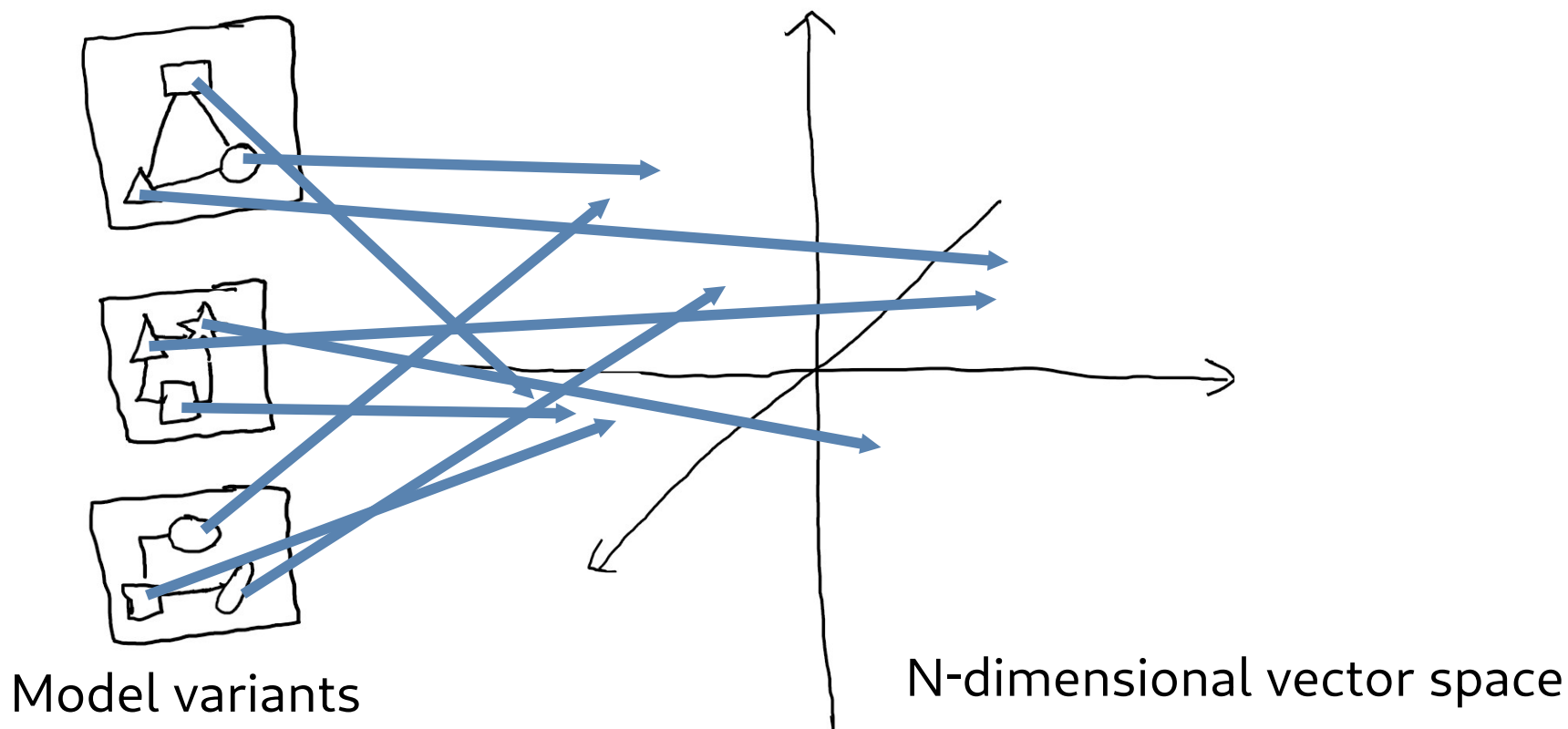
What is the idea?

Fewer element comparisons

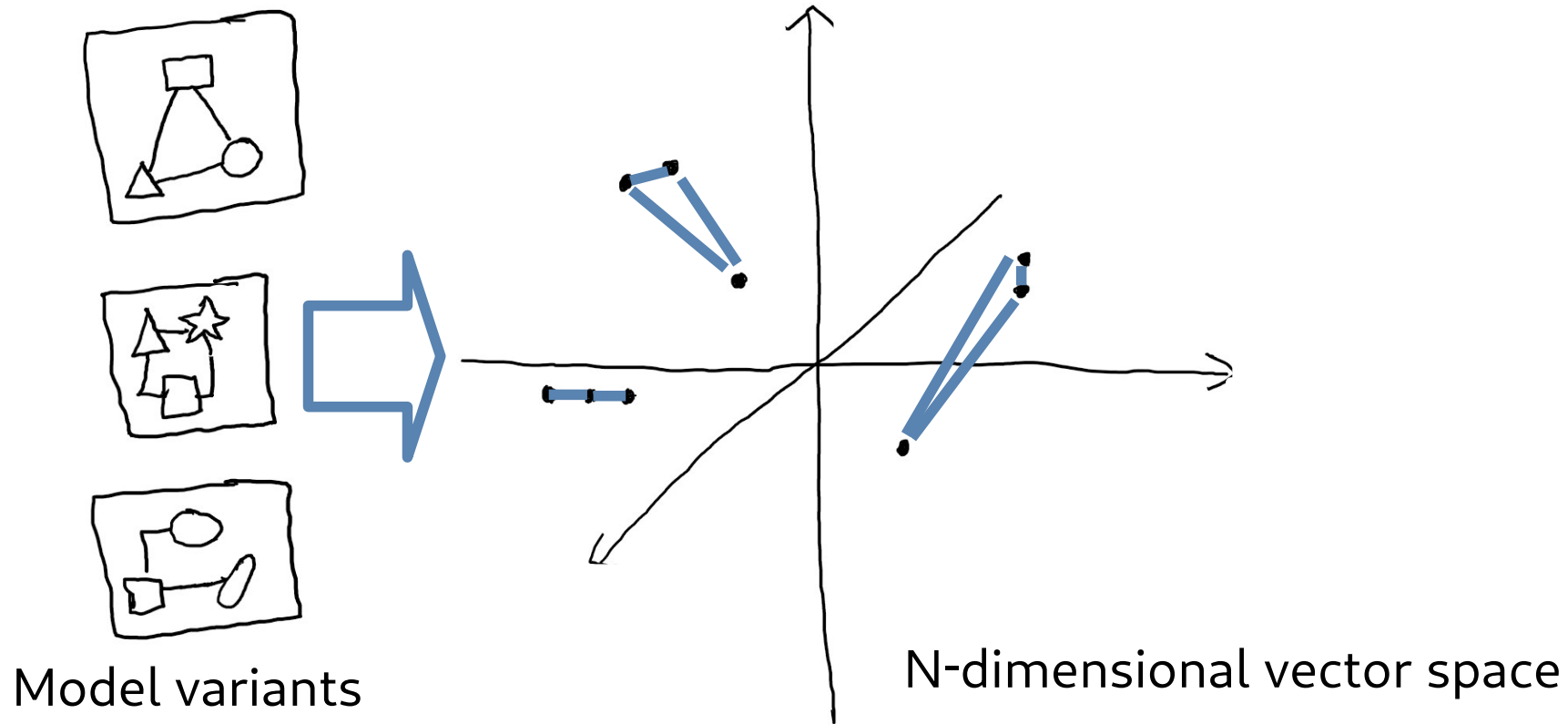


Faster matching

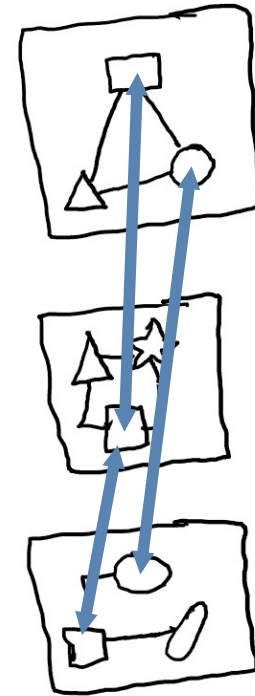
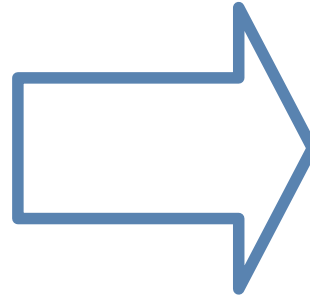
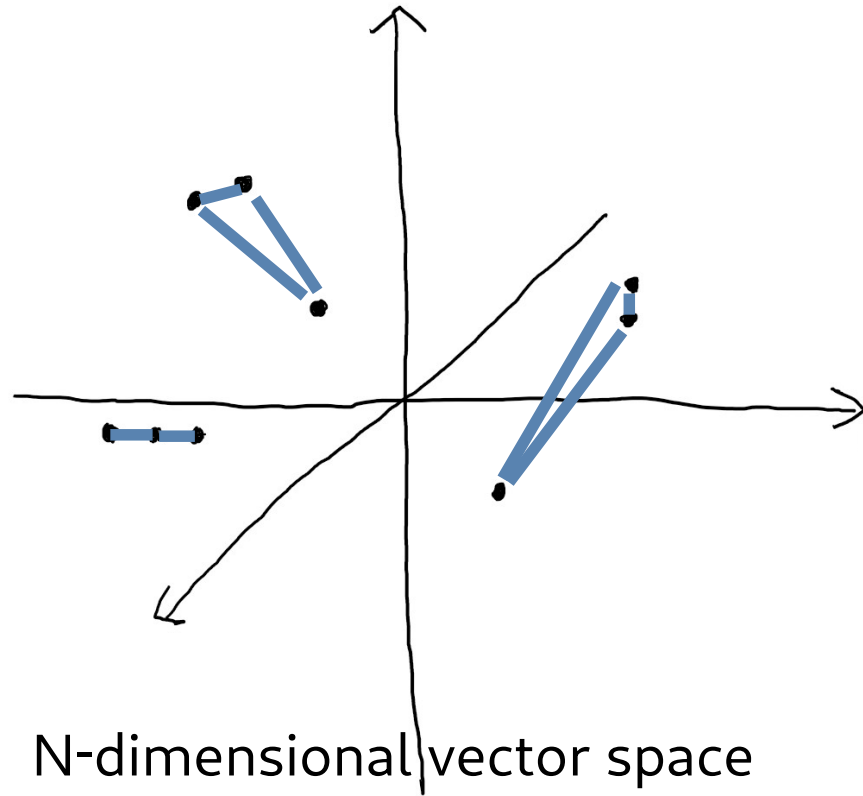
First: Map elements to points in a vector space



Second: Collect the nearest neighbors of each element



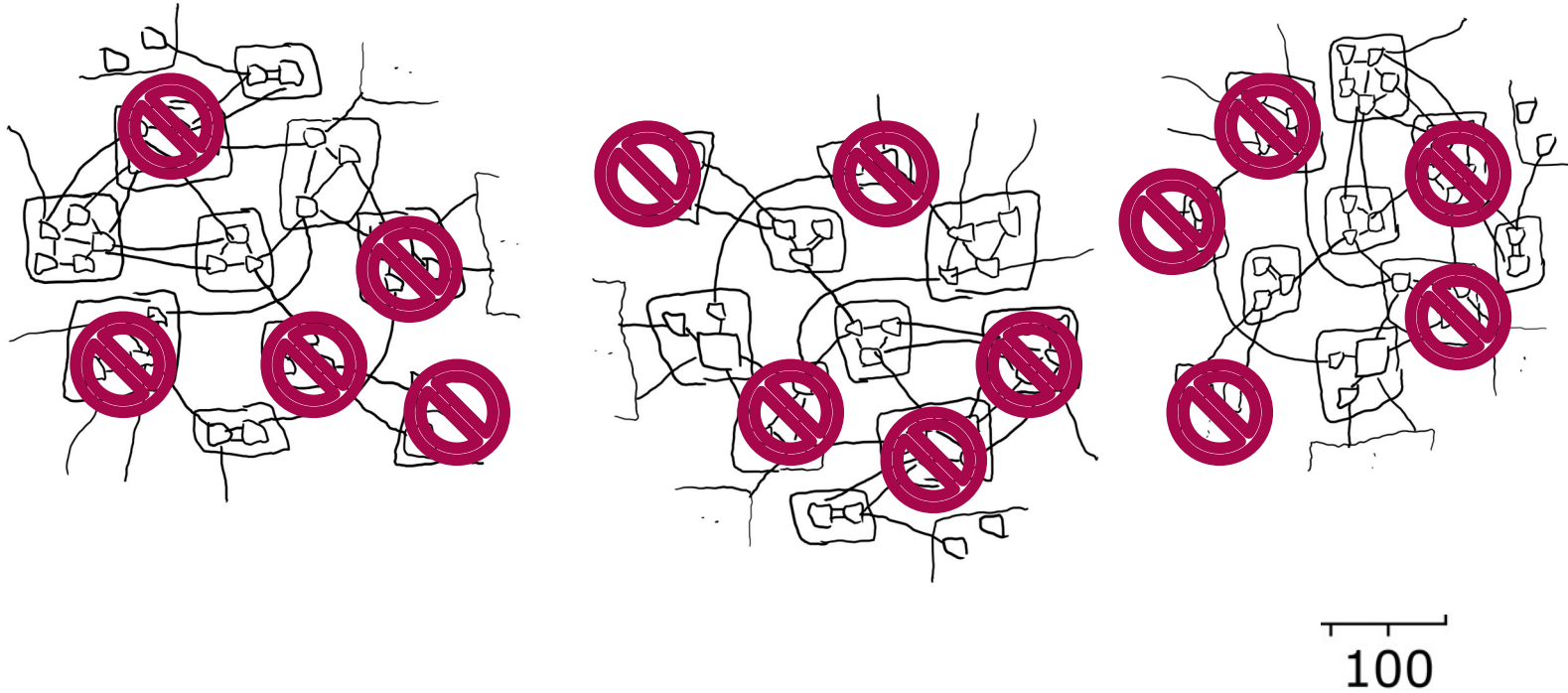
Third: Compare and match neighboring elements



Model variants

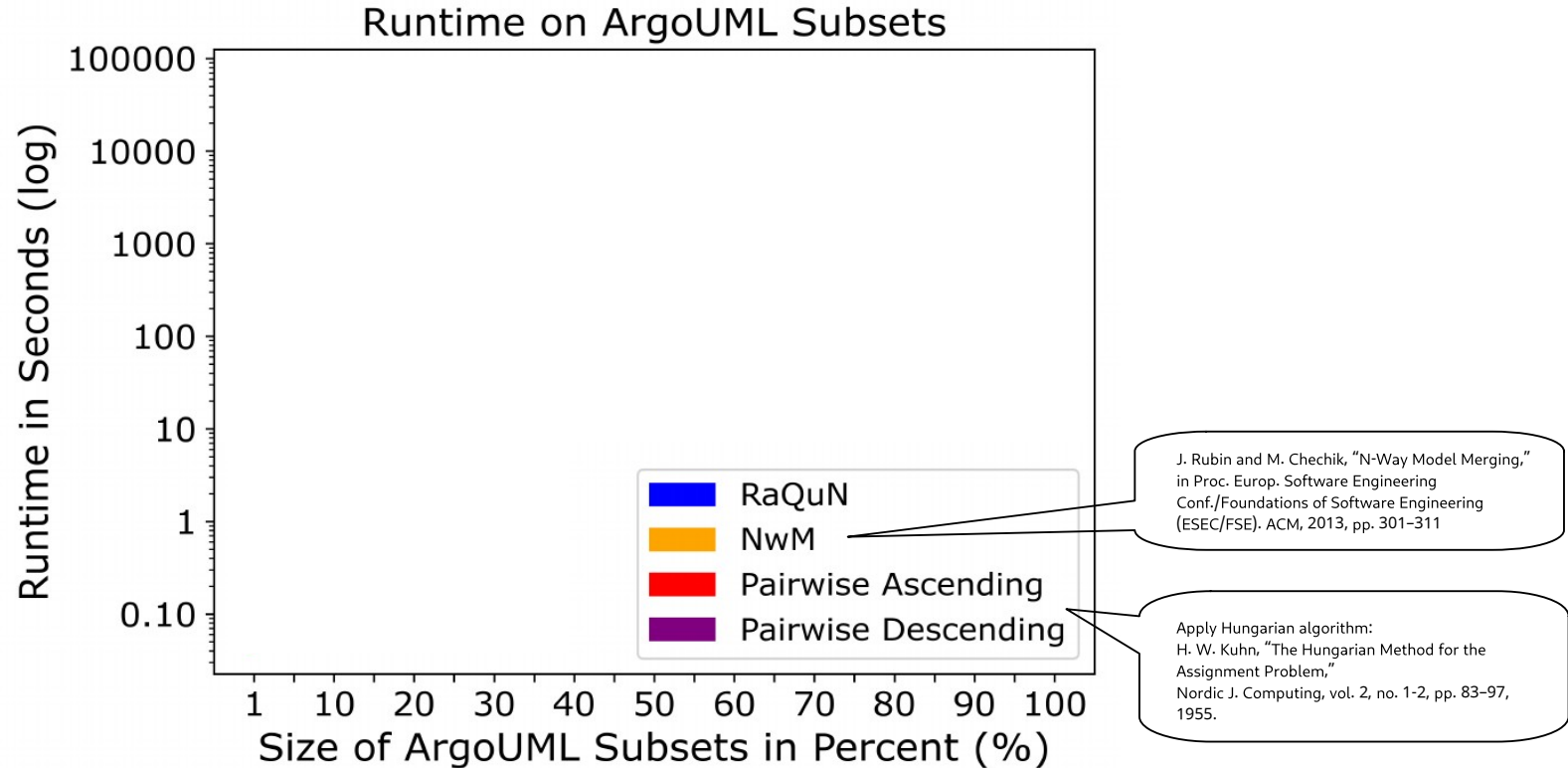
How well does RaQuN scale?

Scalability on ArgoUML

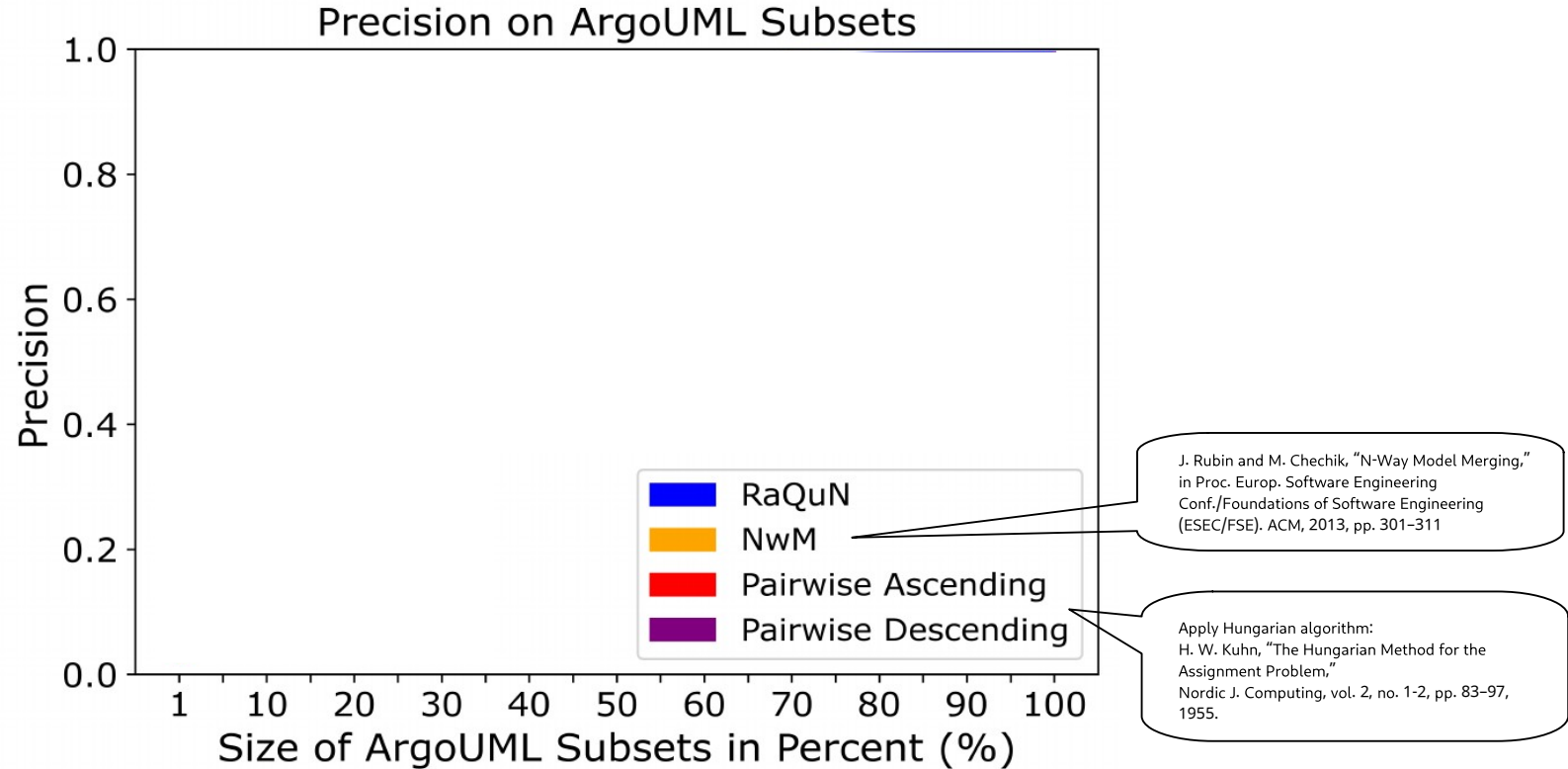


Size of ArgoUML Subsets in Percent (%)

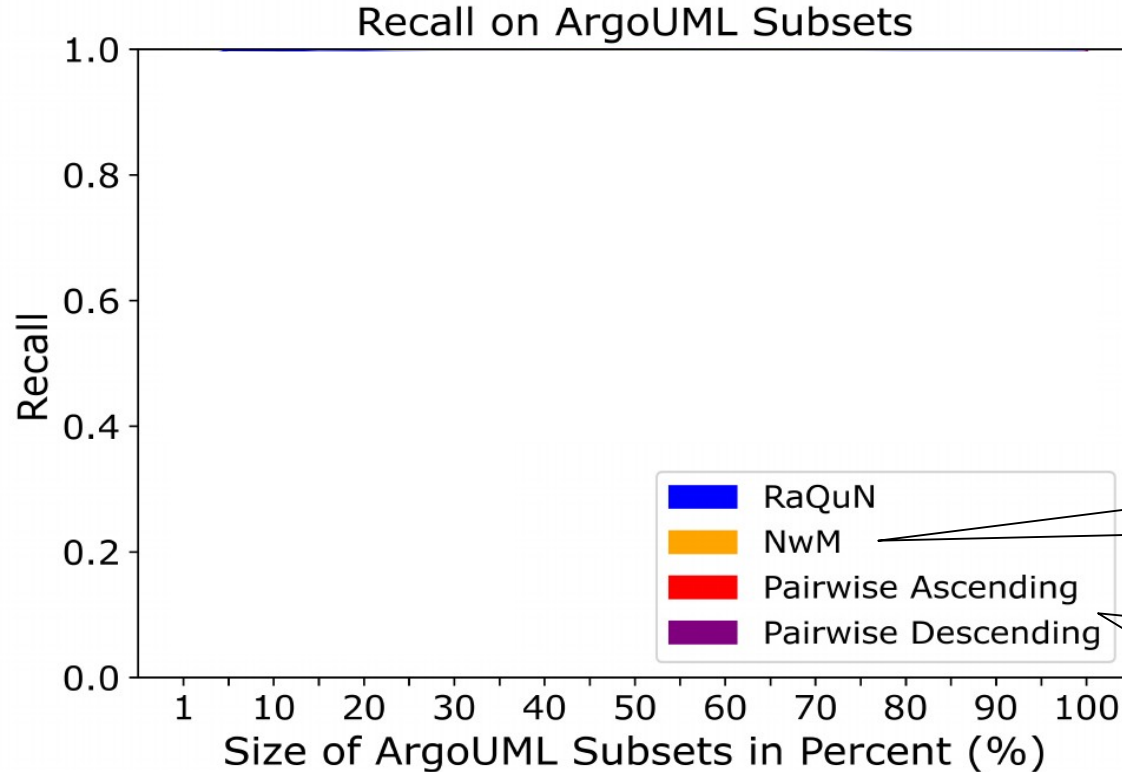
How well does RaQuN scale?



Are the formed matches correct?



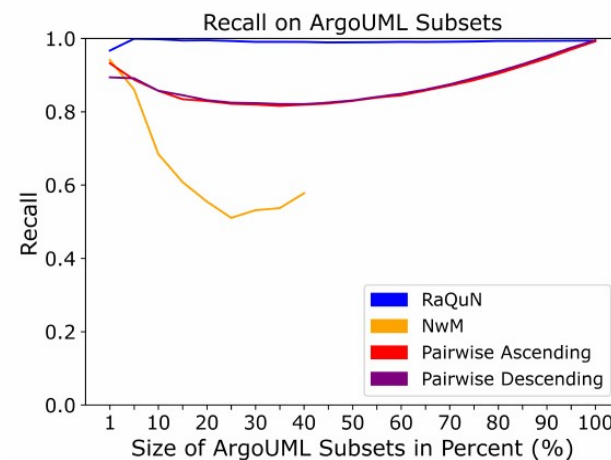
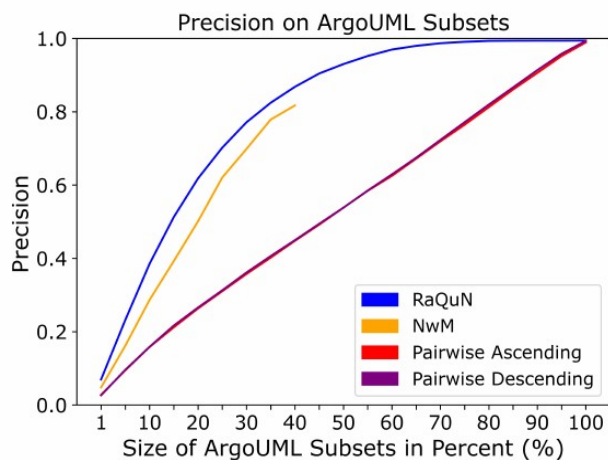
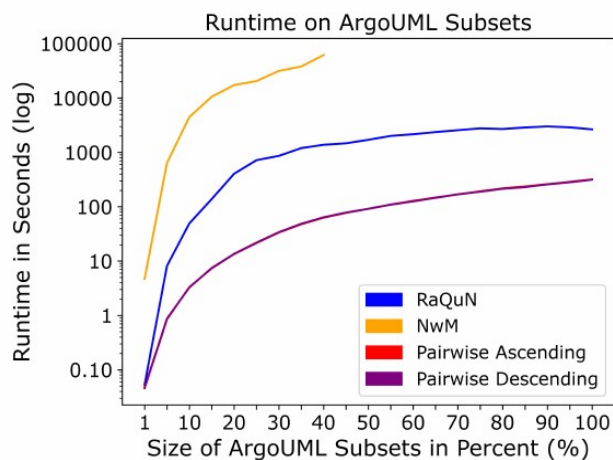
Have all relevant matches been found?



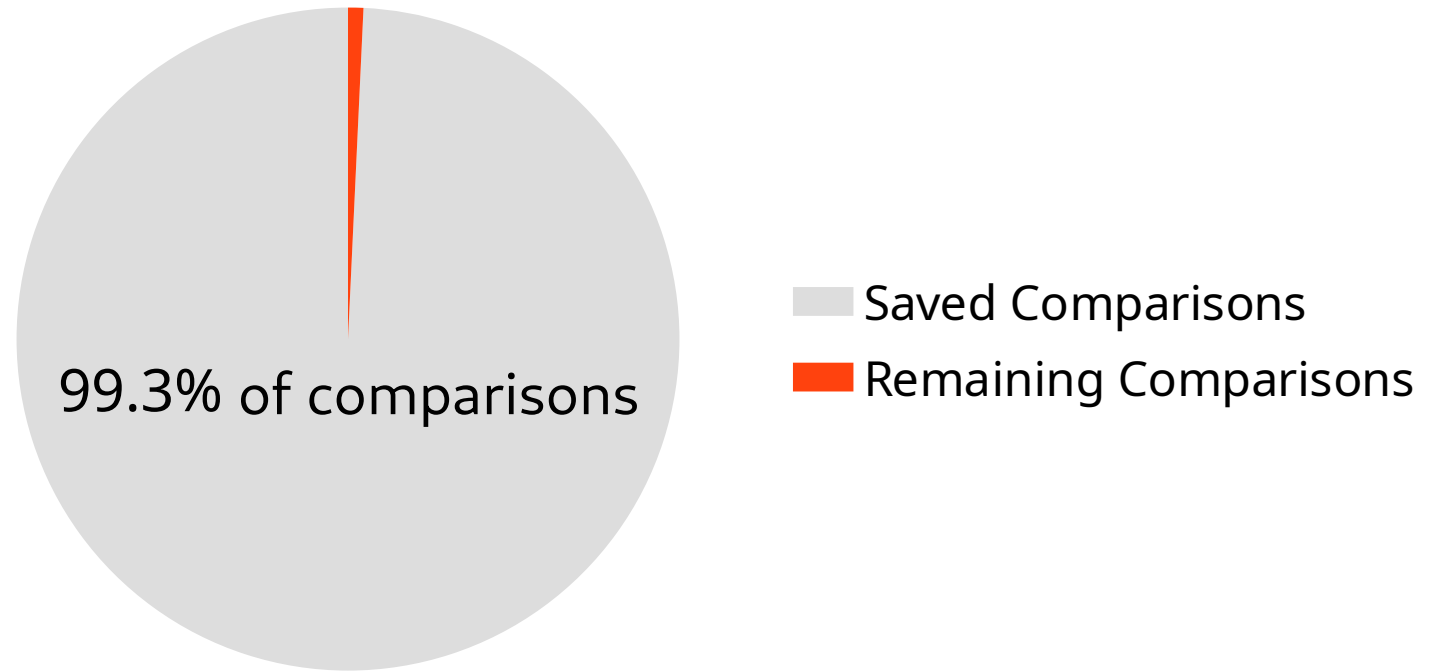
J. Rubin and M. Chechik, "N-Way Model Merging," in Proc. Europ. Software Engineering Conf./Foundations of Software Engineering (ESEC/FSE). ACM, 2013, pp. 301–311

Apply Hungarian algorithm:
H. W. Kuhn, "The Hungarian Method for the Assignment Problem," Nordic J. Computing, vol. 2, no. 1-2, pp. 83–97, 1955.

Does RaQuN achieve better runtime/quality?



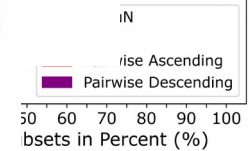
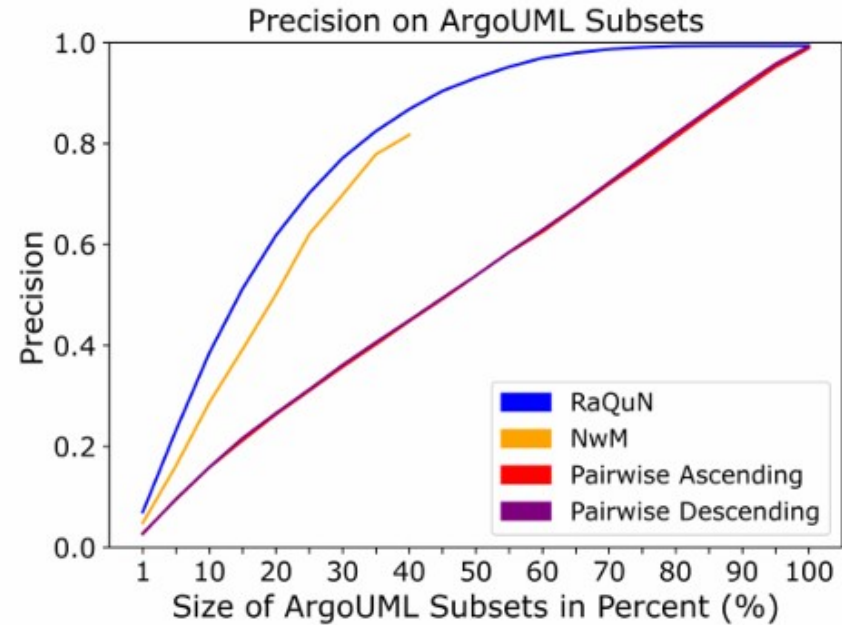
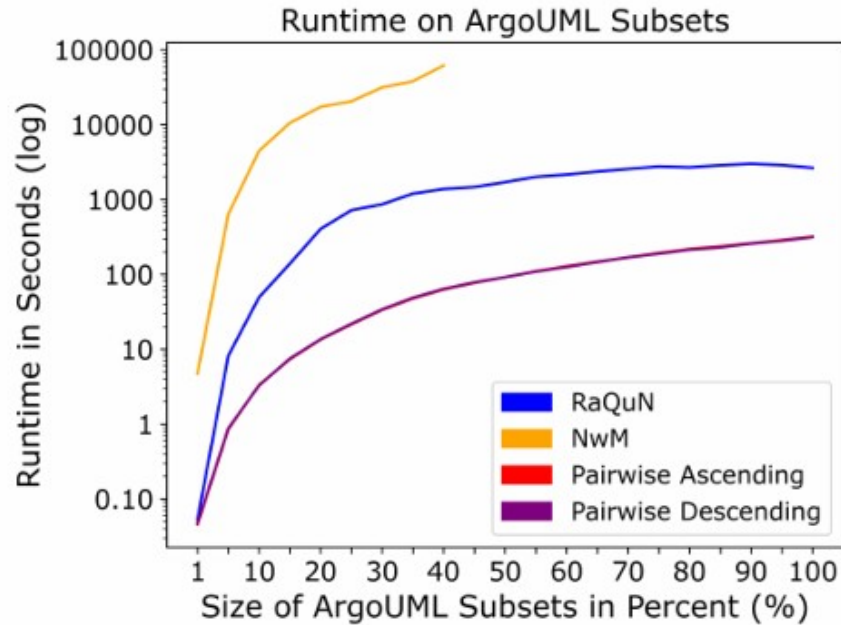
How many comparisons are saved?



Future Work

- Specialize on different domains
- Extend to n-to-m element matching
- Explore scenarios with interactive matching

Summary

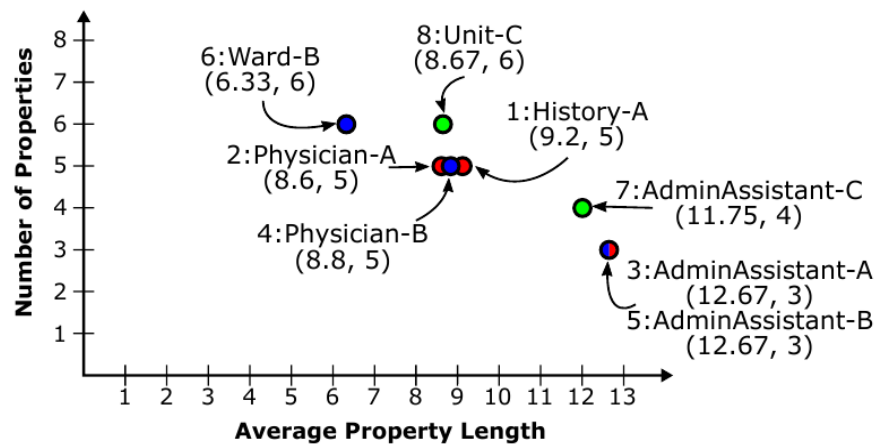
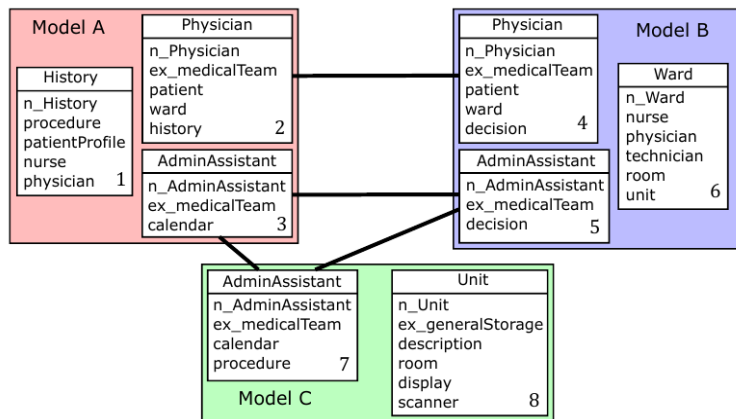


Appendix

We consider a variety of subjects

	Model Type	#Models	Elements	
			Avg.	Median
Hospital	Simple class diag.	8	27.62	26
Warehouse	Simple class diag.	16	24.25	22
Random	Synthetic	100	26.99	26
Loose	Synthetic	100	28.88	29
Tight	Synthetic	100	25.01	25
PPU Structure	SysML block diag.	13	32.15	32
PPU Behavior	UML statemachines	13	221.85	228
bCMS	UML class diag.	14	67.71	63
BCS	Component/connector	18	78.78	72
ArgoUML	UML class diag.	7	1,752.86	1,749
Apo-Games	Simple class diag.	20	63.05	60

Example vectorization



Comparisons saved

Dataset	Full N-Way Matching	RaQuN	
	#Comparisons	#Comparisons	Saved
Hospital	21,211	1,229	94.2%
Warehouse	70,037	6,408	90.9%
Random	33,600	2,499	92.6%
Loose	26,244	2,670	89.8%
Tight	25,237	2,215	91.2%
PPU Structure	80,620	41,533	48.5%
PPU Behavior	3,814,644	259,827	93.2%
bCMS	416,571	106,585	74.4%
BCS	939,346	253,862	73.0%
ArgoUML	64,521,622	481,179	99.3%
Apo-Games	750,319	31,880	95.8%

Weight metric

Given a match t , the weight is calculated as

$$w(t) = \frac{\sum_{2 \leq j \leq |t|} j^2 \cdot n_j^p}{n^2 \cdot |\pi(t)|} \quad (2)$$

where $|t|$ denotes the size of the match, n_j^p the number of properties that occur in exactly j elements of the match, and $\pi(t)$ is the set of all distinct properties of all elements in t .

Detailed Results (1)

TABLE II

COMPARISON OF ACHIEVED WEIGHTS AND RUNTIMES ACROSS ALL ALGORITHMS, AVERAGED OVER 30 RUNS FOR EACH SUBJECT.

Algorithm	Hospital		Warehouse		Random		Loose		Tight		Apo-Games	
	Weight	Time (in s)	Weight	Time (in s)	Weight	Time (in s)	Weight	Time (in s)	Weight	Time (in s)	Weight	Time (in s)
RaQuN	4.92	0.08 [0.07, 0.12]	1.63	0.32 [0.27, 0.93]	1.04	0.07 [0.05, 0.13]	1.03	0.13 [0.07, 0.33]	0.94	0.07 [0.05, 0.11]	18.27	71.12 [41.30, 104.58]
NwM	4.49	20.64 [16.18, 24.22]	1.46	74.51 [29.53, 84.87]	0.80	24.00 [1.23, 76.02]	0.79	22.40 [1.12, 70.68]	0.88	39.75 [1.63, 66.60]	17.91	5,462.91 [3,742.22, 6,597.83]
Pairwise Ascending	4.49	0.31 [0.28, 0.44]	1.11	0.36 [0.33, 0.45]	0.79	0.21 [0.10, 0.31]	0.74	0.14 [0.08, 0.22]	0.94	0.22 [0.16, 0.34]	12.96	10.42 [9.32, 12.04]
Pairwise Descending	4.72	0.16 [0.14, 0.22]	1.27	0.36 [0.32, 0.53]	0.78	0.15 [0.10, 0.23]	0.74	0.14 [0.08, 0.25]	0.93	0.22 [0.17, 0.33]	16.40	10.68 [9.27, 13.50]

Algorithm	PPU Structure		PPU Behavior		bCMS		BCS		ArgoUML	
	Weight	Time (in s)	Weight	Time (in s)	Weight	Time (in s)	Weight	Time (in s)	Weight	Time (in s)
RaQuN	28.95	0.85 [0.81, 0.90]	164.53	18.32 [16.39, 20.34]	41.53	2.84 [1.40, 3.58]	51.17	12.58 [8.10, 18.22]	1727.65	2,647.76 [1,483.48, 3,324.70]
NwM	28.64	9.27 [7.84, 10.07]	146.35	4,616.30 [3,410.86, 5,919.65]	41.25	247.31 [227.52, 275.95]	43.20	330.25 [235.74, 388.45]	- - -	timeout - - -
Pairwise Ascending	28.65	0.27 [0.22, 0.38]	145.46	11.03 [10.32, 12.61]	39.76	1.16 [1.13, 1.21]	43.83	5.69 [3.85, 7.43]	1702.91	318.26 [297.43, 379.76]
Pairwise Descending	28.89	0.26 [0.21, 0.42]	142.56	10.84 [10.20, 13.01]	38.76	1.04 [1.01, 1.07]	47.16	4.84 [3.37, 6.73]	1710.25	314.88 [302.28, 329.35]