

CLOSE-TO-OPTIMAL PLACEMENT AND ROUTING FOR CONTINUOUS-FLOW MICROFLUIDIC BIOCHIPS



Andreas Grimmer¹, Qin Wang², Hailong Yao², Tsung-Yi Ho³, Robert Wille¹

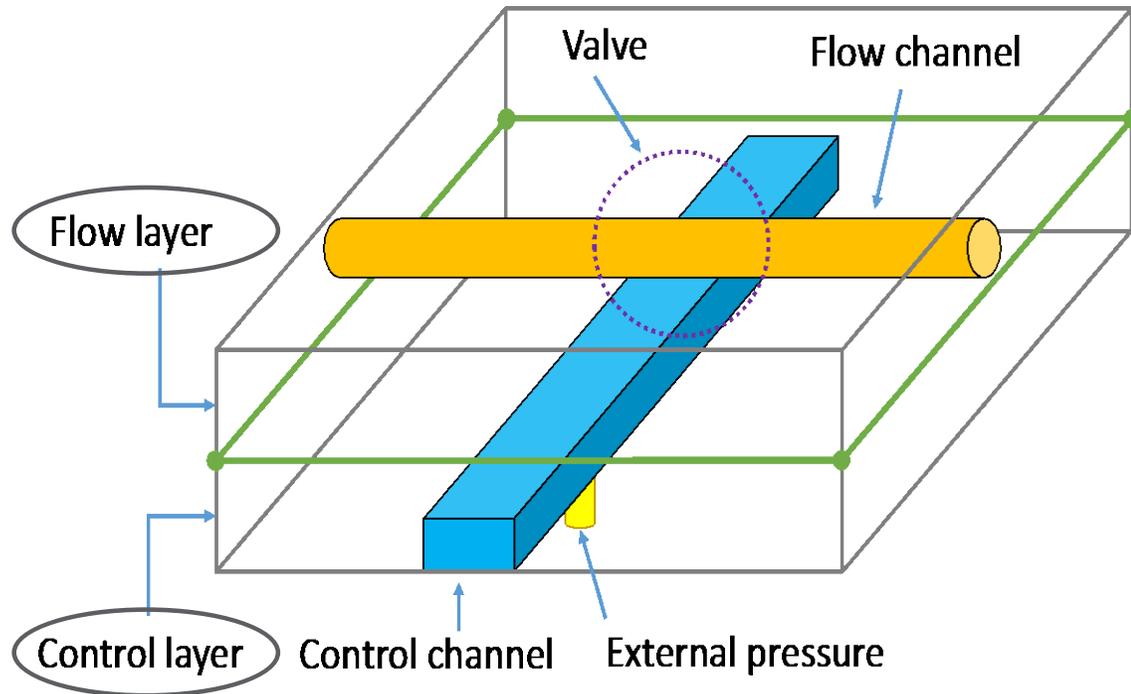
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² Tsinghua University, Beijing, China

³ National Tsing Hua University, Taiwan

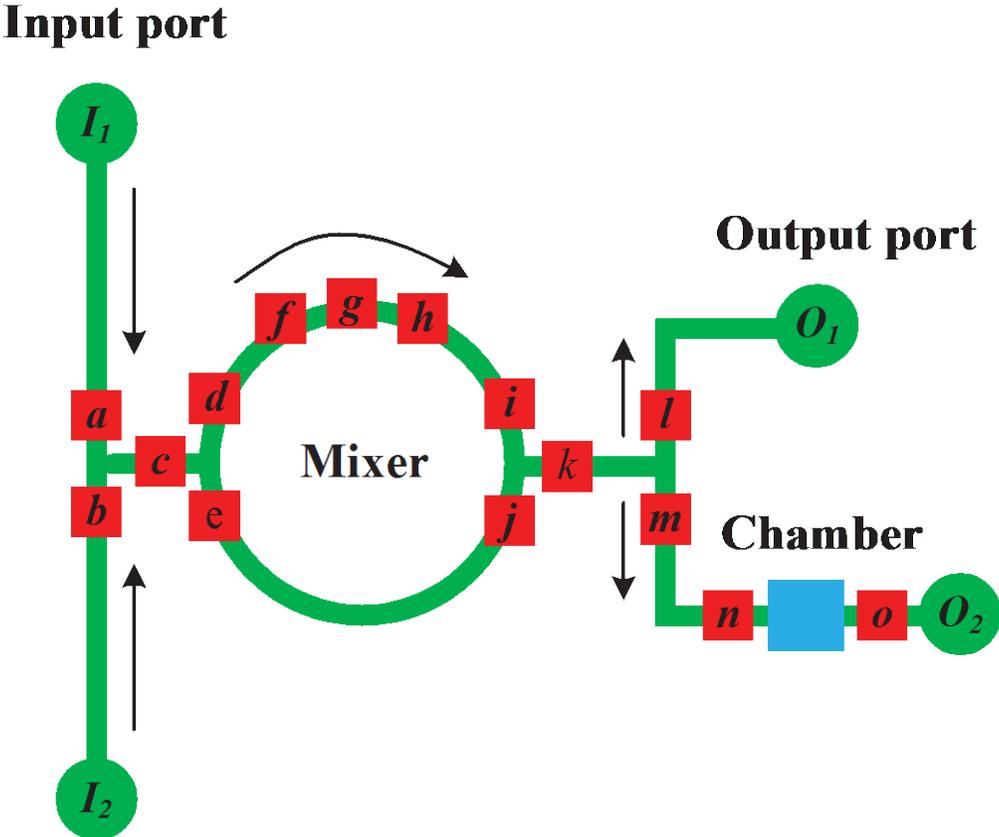
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CONTINUOUS-FLOW MICROFLUIDIC BIOCHIP



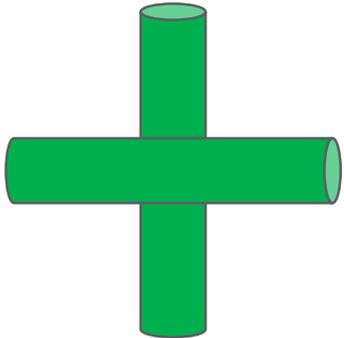
- Two-layer: control- and flow-layer
- Thousands of valves

MIXER COMPONENT



PHYSICAL DESIGN

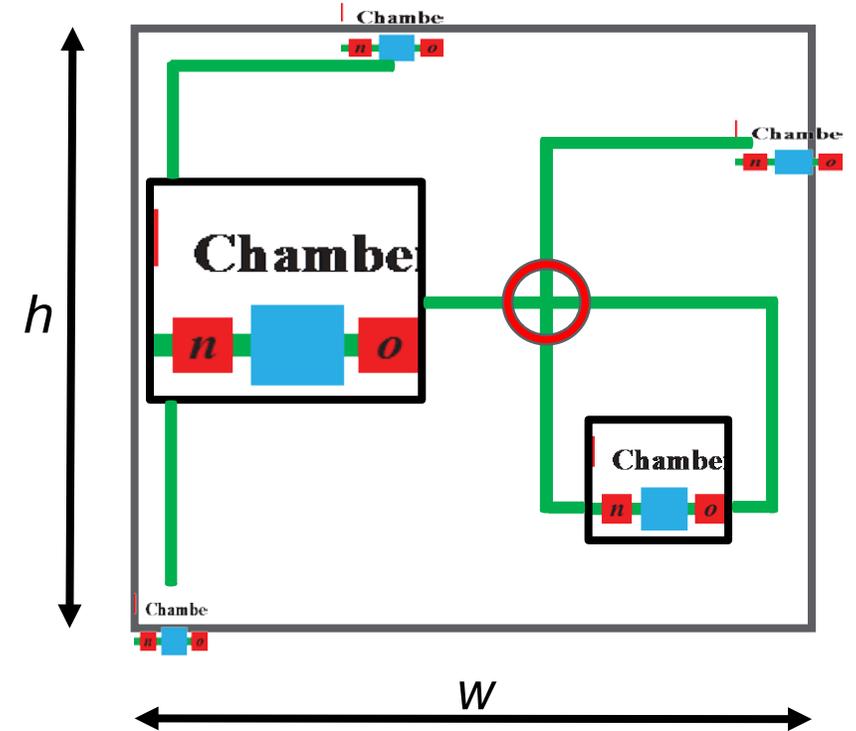
- Place components
- Route flow channels
- Quality of design:



Flow channel intersections



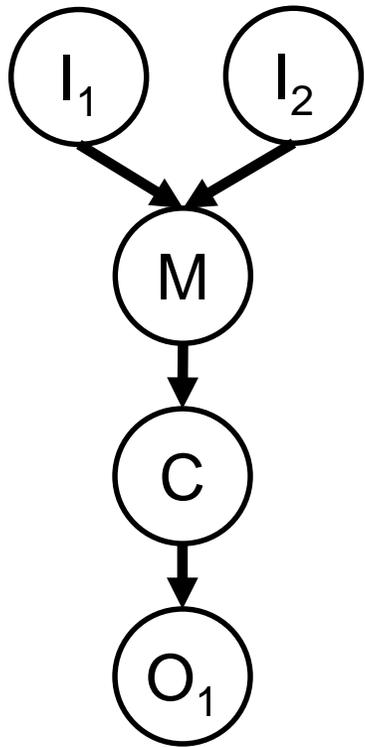
Overall flow channel length



Flow layer size

PHYSICAL DESIGN

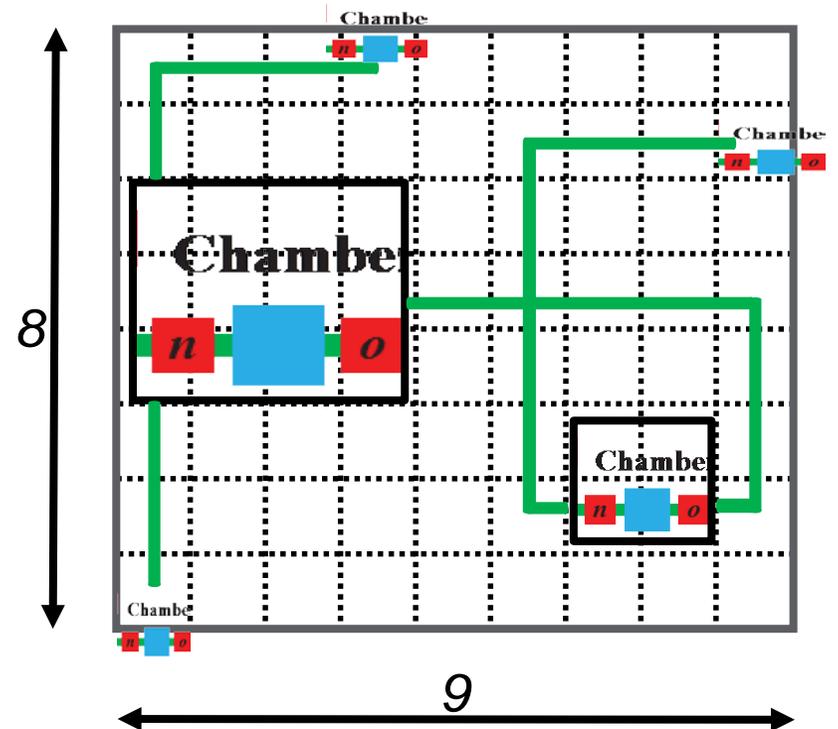
Input



Grid
9 x 8 cells

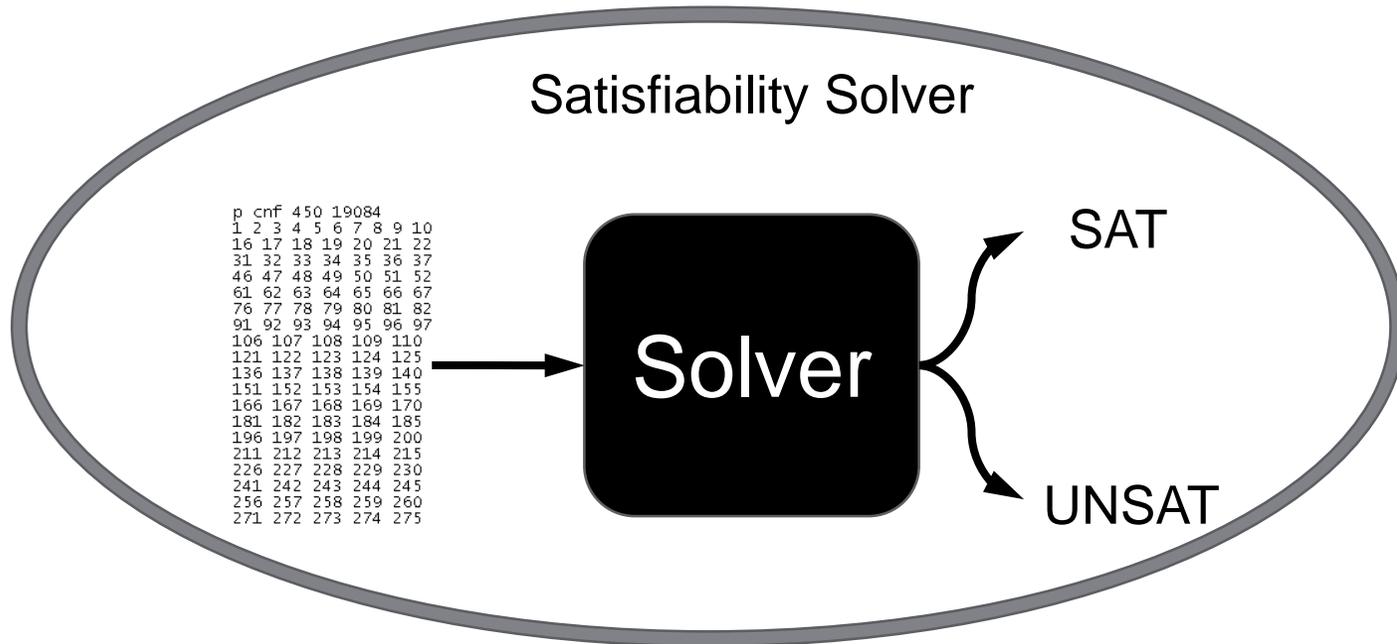
Resource	Area
Mixer	4 x 3
Chamber	2 x 2

Grid



APPROACH

- Consider placement and routing in **single** step (P&R)
→ Enormous search space



Search Space Pruning



SYMBOLIC FORMULATION FOR P&R

■ Grid: $1 \leq x \leq w$ and $1 \leq y \leq h$

■ $V = \{Mixer, Chamber, I_1, I_2, O_1\}$

■ For each component $v \in V$:

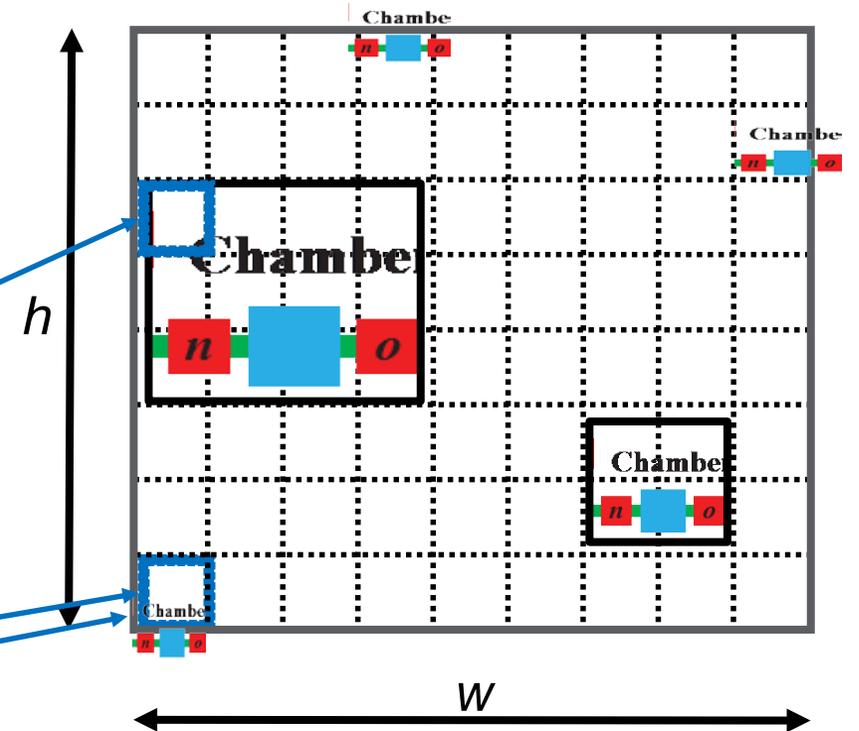
$$vp_{(x,y),v}$$

■ Example:

$$vp_{(1,6),Mixer} = 1$$

$$vp_{(1,1),Mixer} = 0$$

$$vp_{(1,1),I_2} = 1$$



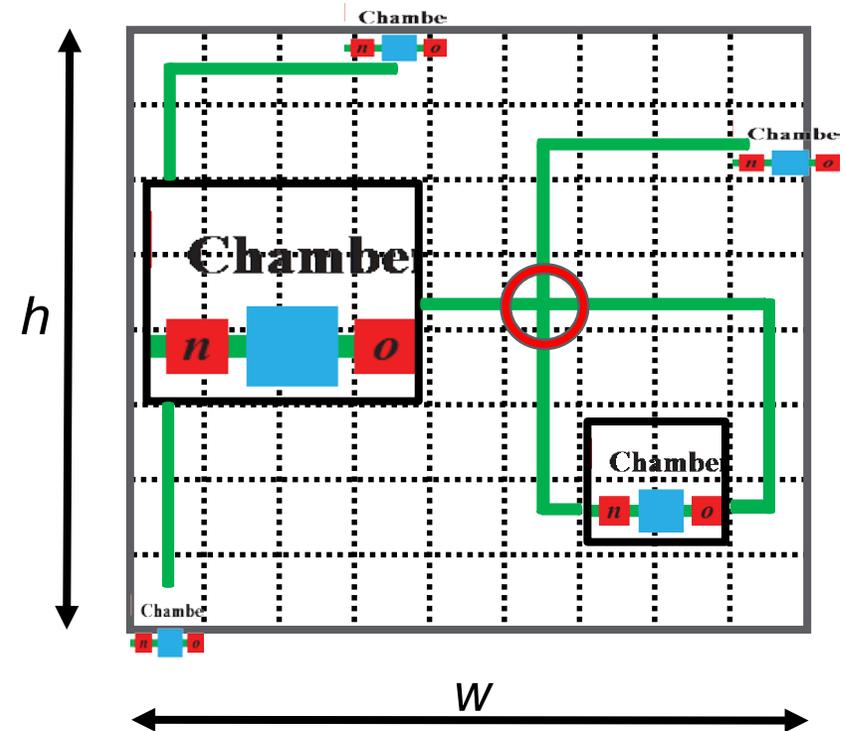
CONSTRAINTS FOR P&R

- Constraints for placing components
- Constraints for routing channels
- Enforcing the **designer's** constraints:
 - Maximal channel intersections:

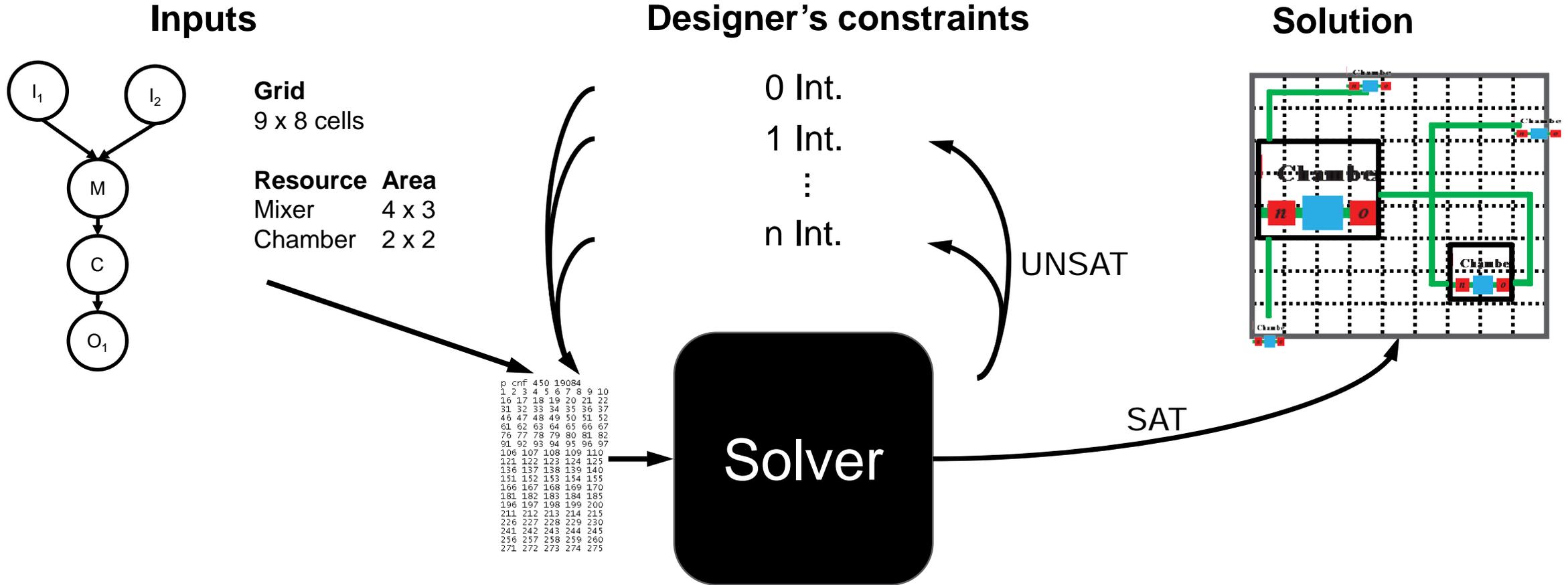
$$\sum_{\substack{1 \leq x \leq w \\ 1 \leq y \leq h}} \left(\sum_{(u,v) \in E} ep_{(x,y),(u,v)=2} \right) \leq \text{maxIntersections}.$$

- Maximal channel length:

$$\sum_{\substack{1 \leq x \leq w \\ 1 \leq y \leq h}} \left(\sum_{(u,v) \in E} ep_{(x,y),(u,v)} \right) \leq \text{maxLength}$$

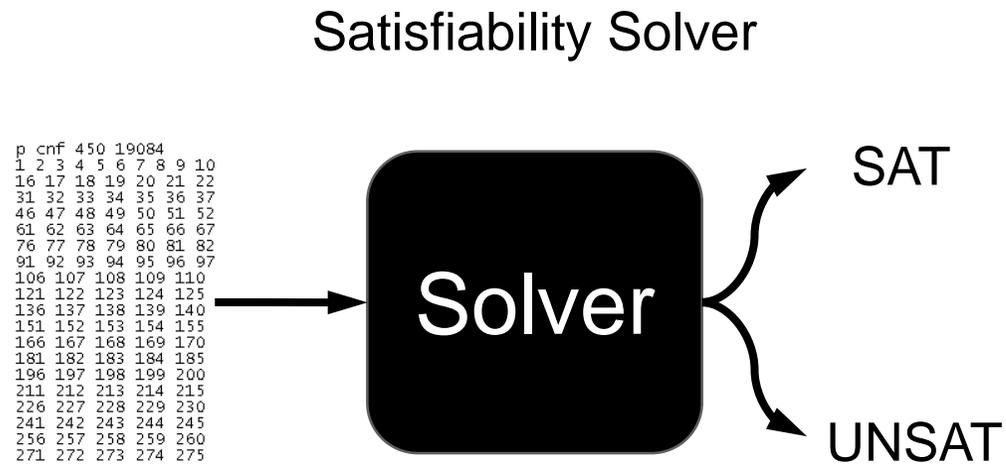


SAT-PROCEDURE FOR OPTIMAL P&R

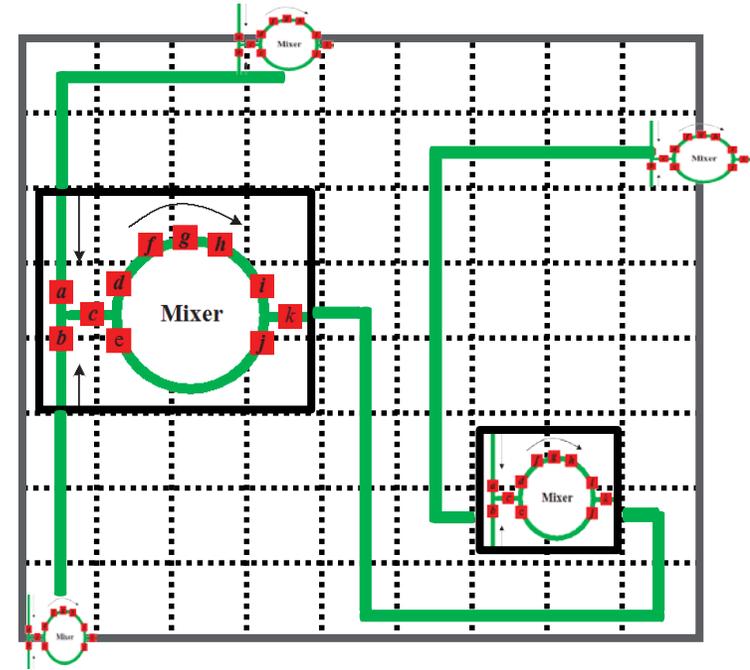
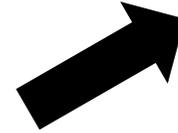
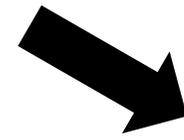
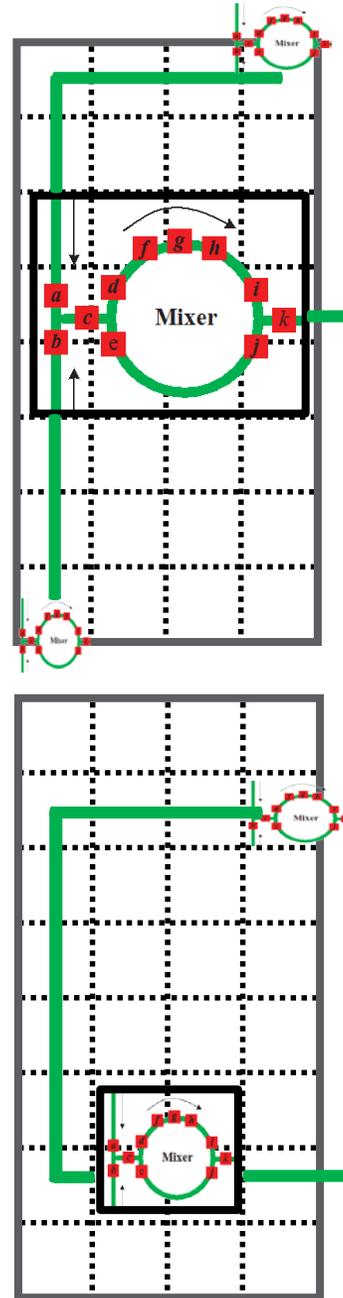
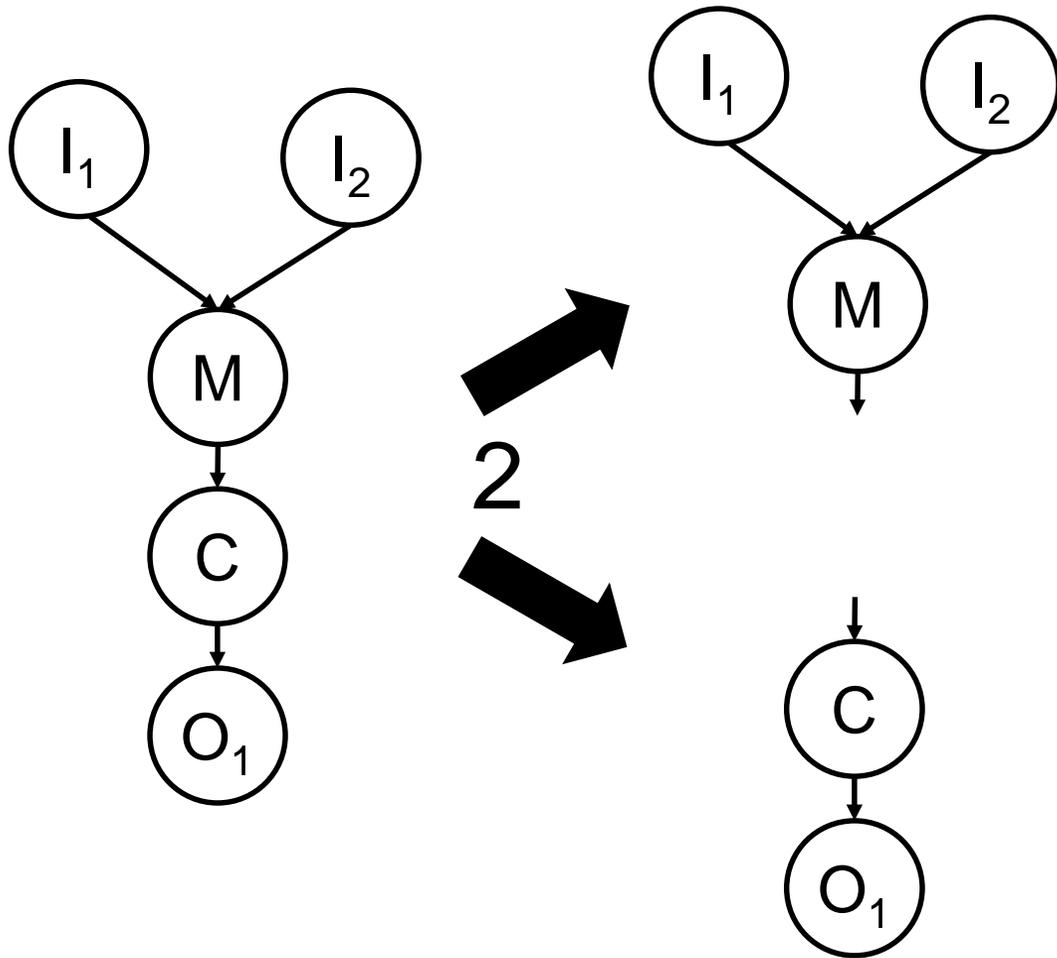


APPROACH

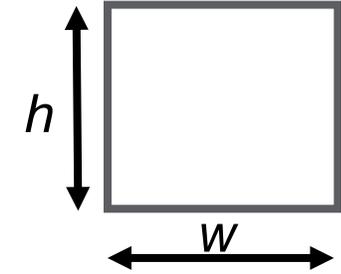
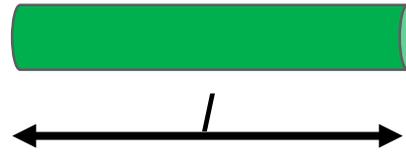
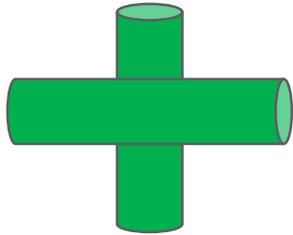
- Consider placement and routing in **single** step (P&R)
→ Enormous search space



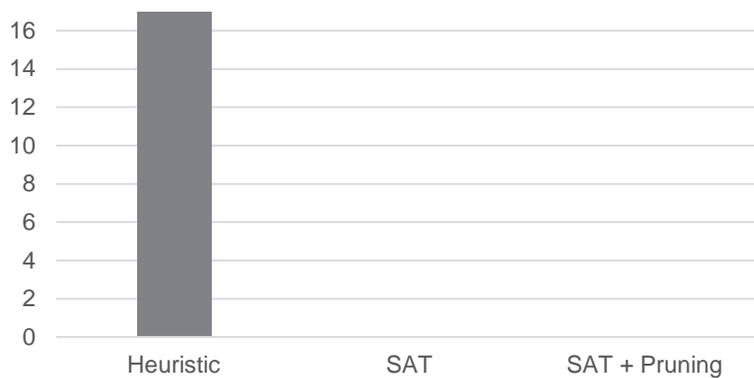
PARTITIONING



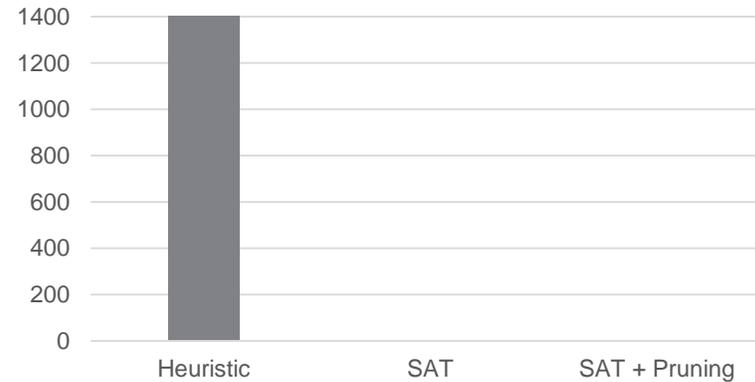
EVALUTATION - SMALL BENCHMARK



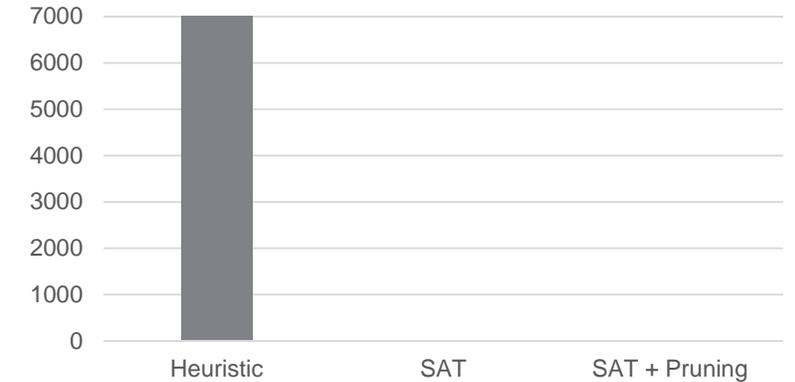
Channel Intersections



Channel Length

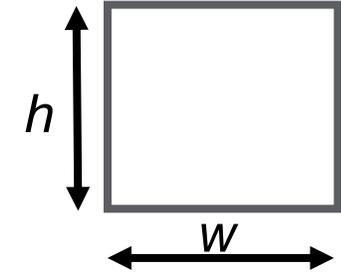
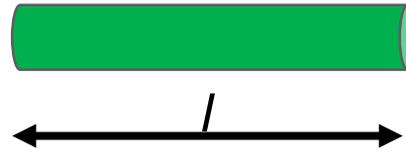
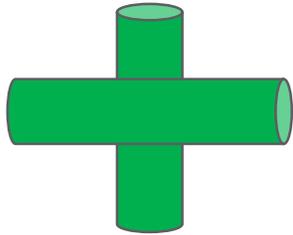


Grid Size

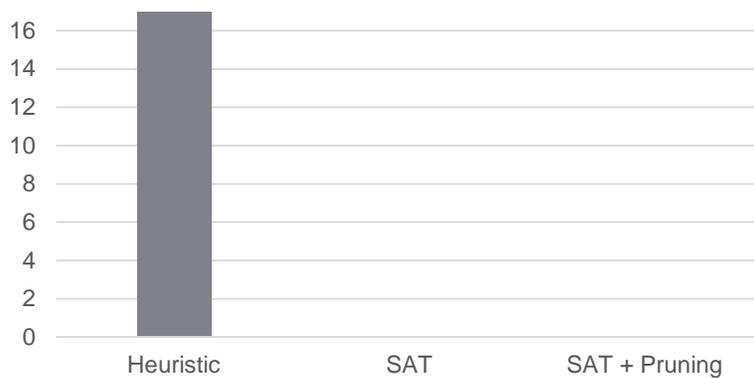


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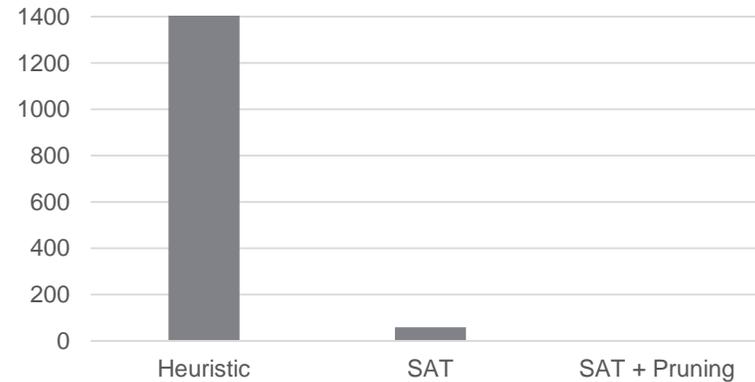
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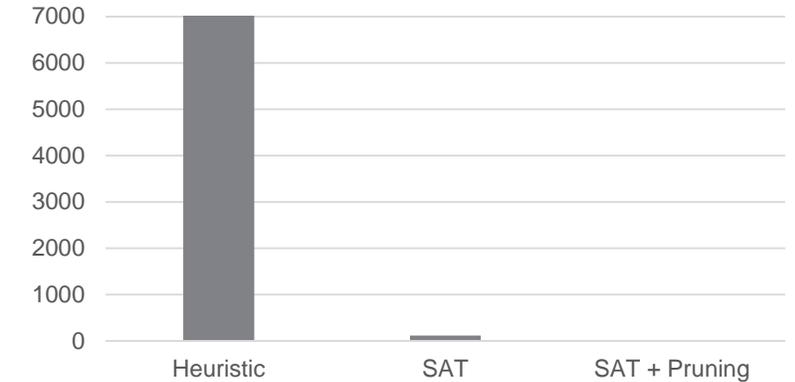
Channel Intersections



Channel Length

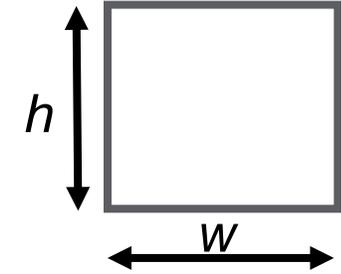
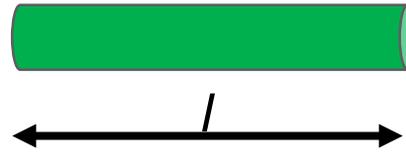
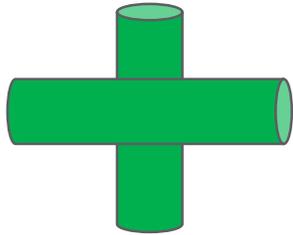


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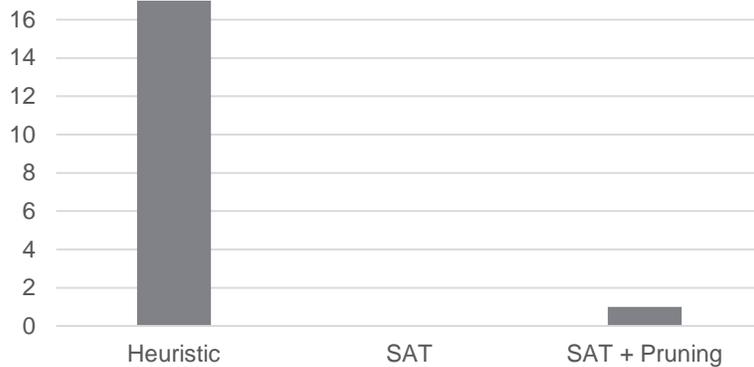


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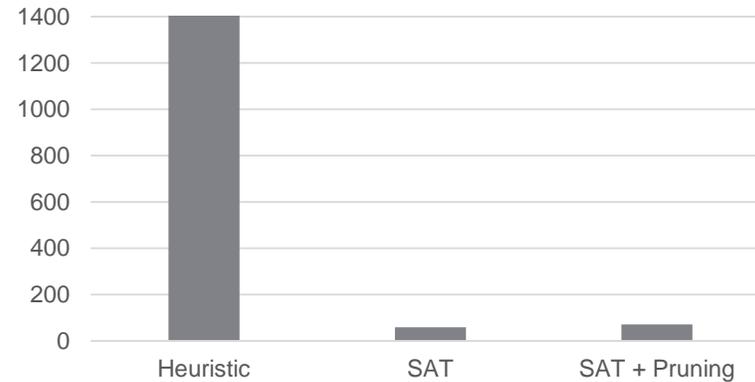
EVALUTATION - SMALL BENCHMARK



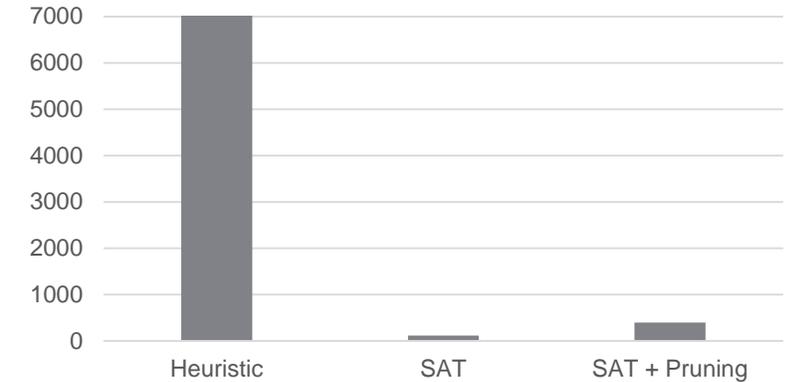
Channel Intersections



Channel Length

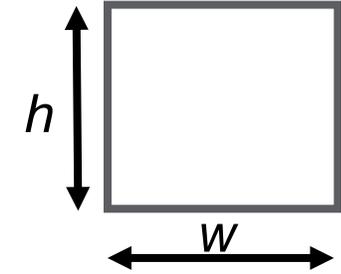
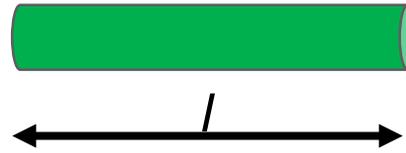
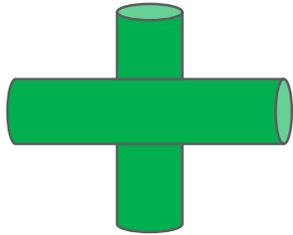


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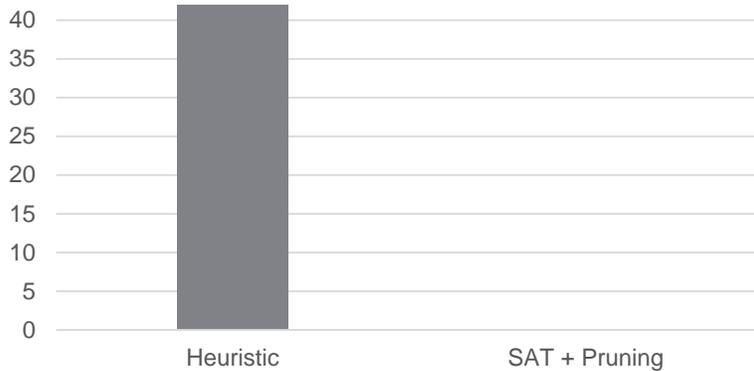


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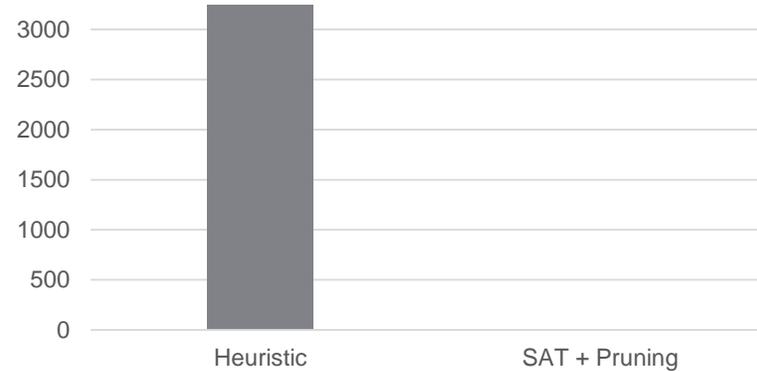
EVALUTATION - PROTEIN SPLIT



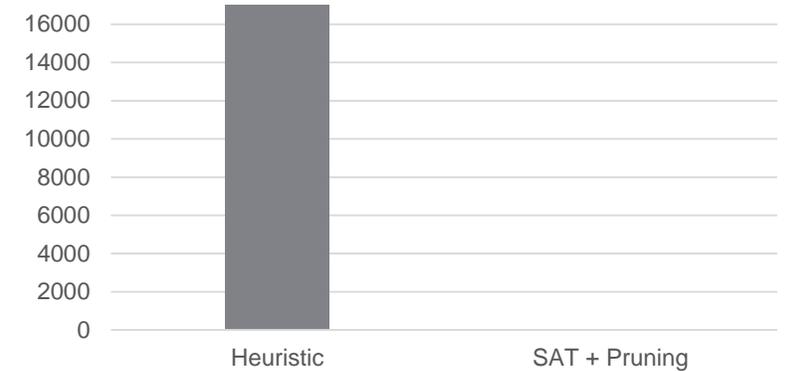
Channel Intersections



Channel Length

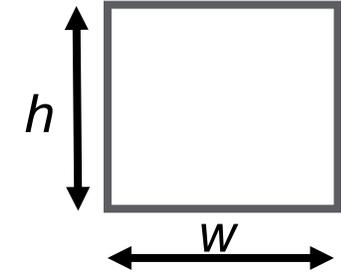
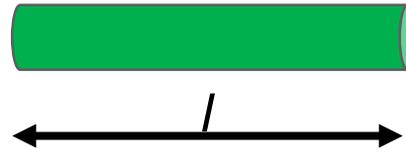
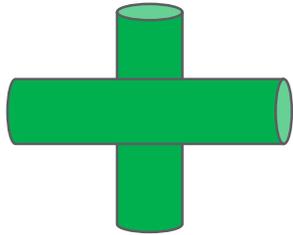


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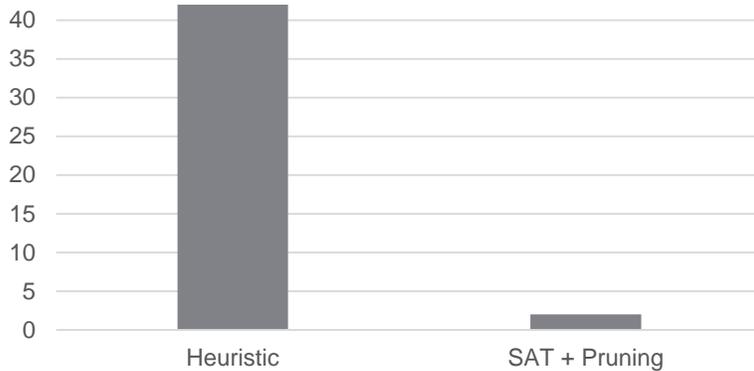


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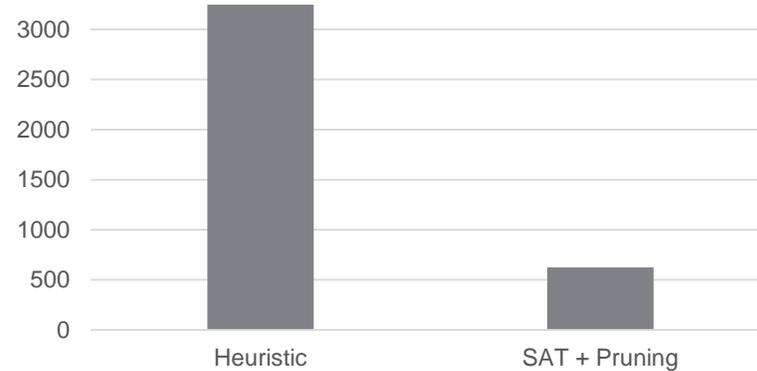
EVALUTATION - PROTEIN SPLIT



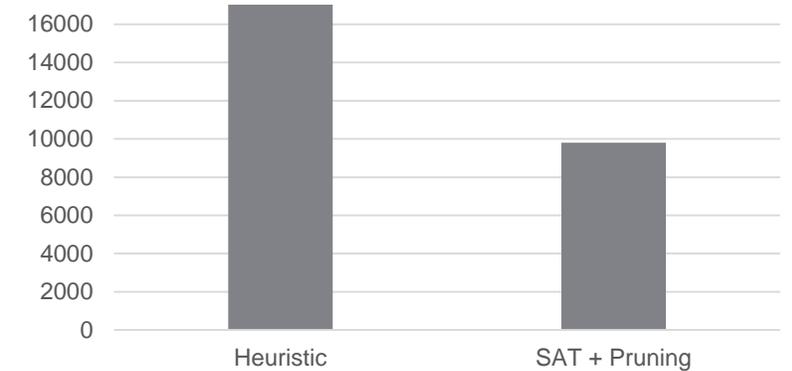
Channel Intersections



Channel Length



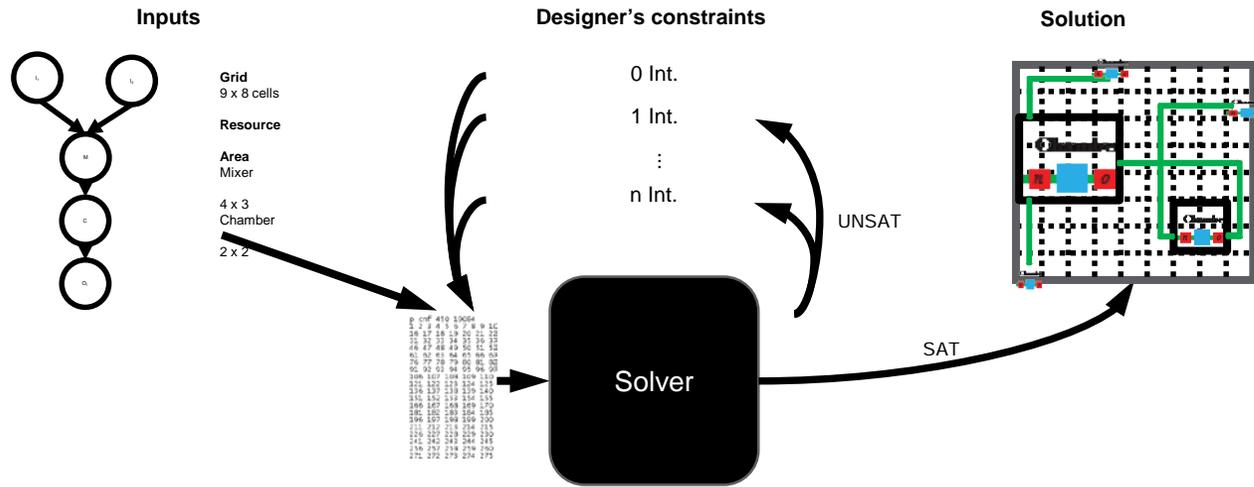
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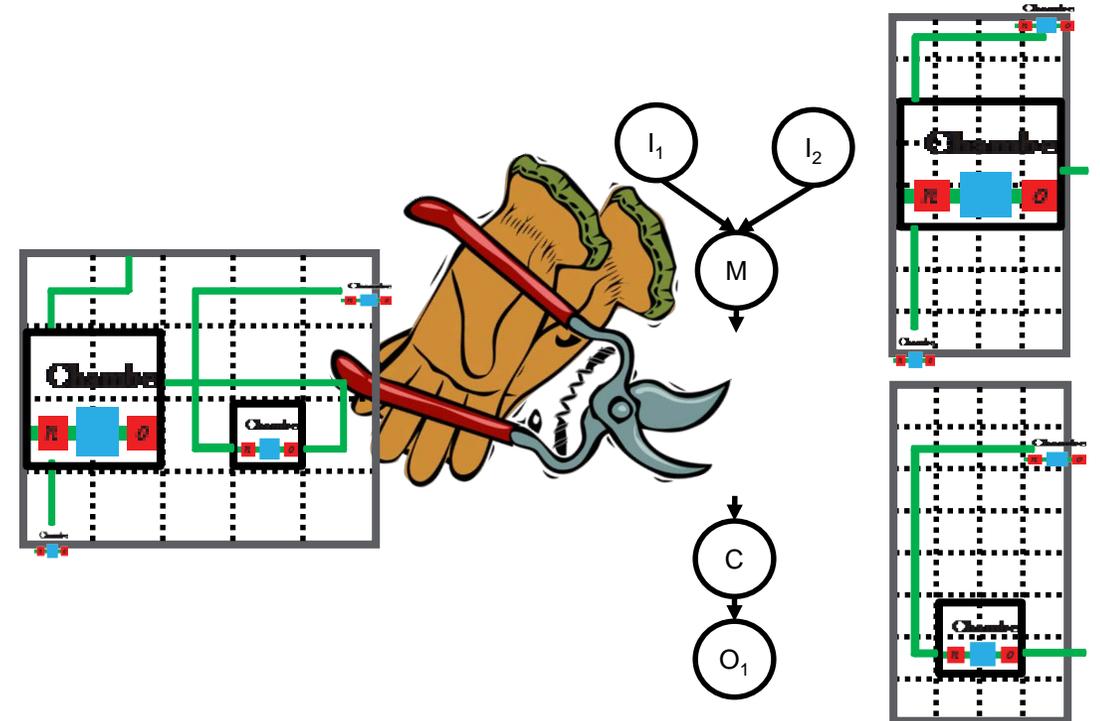
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CONCLUSION

Satisfiability Solver



Search Space Pruning



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