

Parameterized Algorithms and Complexity for Function Merging with Branch Reordering

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Code size optimization

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Applications: mobile apps, embedded systems.

Function merging

Function Merging

Input: Two functions F_1 and F_2 .

Output: Merged function $F_\star$ such that:

- $F_\star$ captures both F_1 and F_2 , and
- $F_\star$ is as small as possible.

Function merging (ex.)

Simplifying assumption: size = # lines.

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21 def F2(x):
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Function merging (ex.)

23 lines $\rightarrow$ 22 lines!

Function merging via sequence alignment

SOTA approach: **sequence alignment** [CGO'19, PLDI'20].

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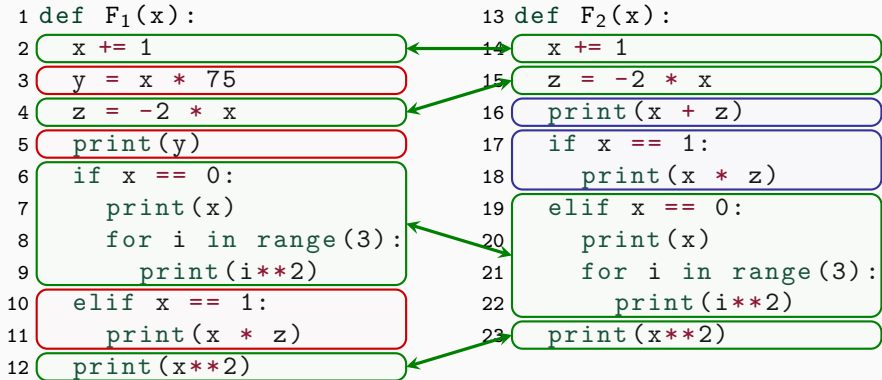
More aligned instructions $\Rightarrow$ smaller $F_\star$.

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Input: Two strings S_1 and S_2 .

Output: Optimal sequence alignment of S_1 and S_2 .

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Textbook dynamic programming for Sequence Alignment
 $\Rightarrow \mathcal{O}(|F_1| \cdot |F_2|)$ -time algorithm.

Sequence alignment + branch reordering

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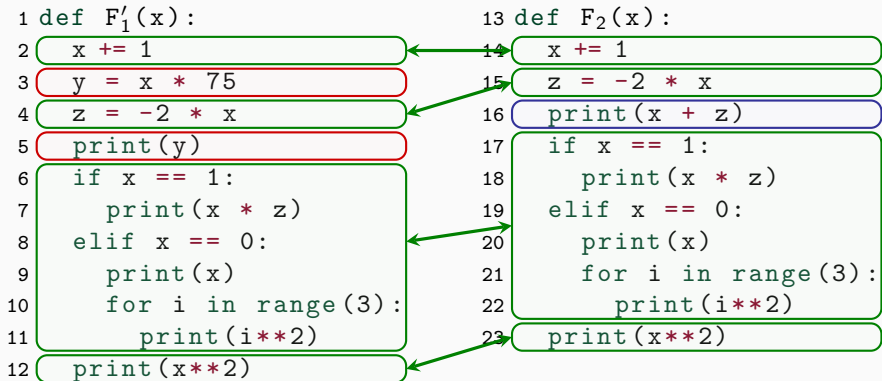
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23 lines $\rightarrow$ 22 lines $\rightarrow$ 20 lines!

Problem statement

Function Merging with Branch Reordering

Input: Two functions F_1 and F_2 .

Output: Compute

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We study **algorithms and complexity** of this problem.

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Function Merging with Branch Reordering is NP-hard.

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Identify some **parameters** of the input such that:

- (1) On inputs of interest, these parameters are **restricted**.
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Satisfying (1) and (2) $\Rightarrow$ profit!

Parameters: branching factor and nesting depth

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Hypothesis: These parameters are typically small.

Two parameterized algorithms

Four parameters:

- branching factor of F_1/F_2 (b_1/b_2),
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Function Merging with Branch Reordering can be solved in

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Function Merging with Branch Reordering can be solved in $2^{\mathcal{O}(b_1 d_1 + b_2)} \cdot (|F_1| \cdot |F_2|)^4$ time.

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Open: what if b_1, b_2 are restricted, but d_1, d_2 are unbounded?

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dp algorithm for $\text{SequenceAlignment}(S_1, S_2)$

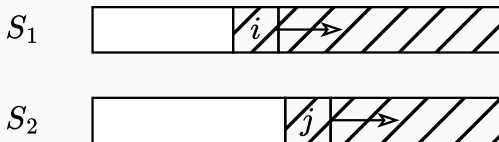
$\approx$

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$$dp[i, j] = \text{SequenceAlignment}(S_1[i \dots], S_2[j \dots])$$



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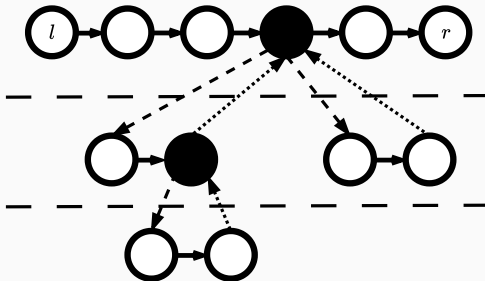
incremental, simultaneous exploration of $\text{Reorder}(F_1)$ and $\text{Reorder}(F_2)$; compute sequence alignment **on the fly**.

Programs to branching graphs

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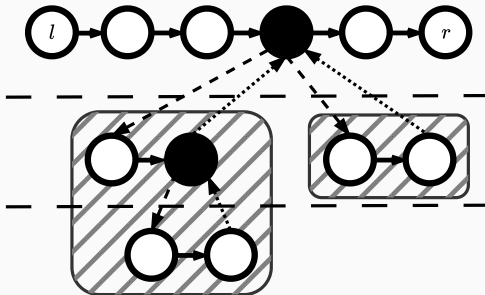
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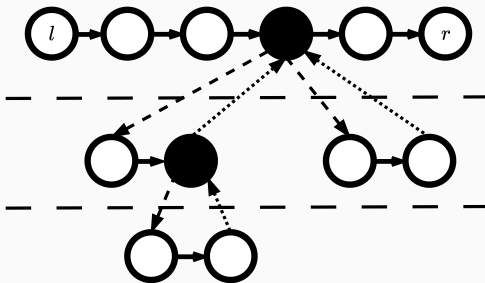


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Exploring $\text{Reorder}(F)$



$\text{EXPLORE}(\ell, \emptyset, \varepsilon)$.

procedure $\text{EXPLORE}(u, X, S)$

if X contains all nodes: **print** S ; **return**.

$X \leftarrow X \cup \{u\}$.

if u is $\bigcirc$: $\text{EXPLORE}(\text{next}(u), X, S \cdot \text{instr}(u))$.

if u is $\bullet$: **forall** $v \in \text{branches}(u) \setminus X$: $\text{EXPLORE}(v, X, S)$.

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Lemma (Compactness)

The number of distinct (u, X) encountered in EXPLORE is at most

$$2^{\mathcal{O}(bd)} \cdot |F|.$$

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EXPLORE($u, X, \dots$)



FUNCTIONMERGING(u_1, X_1, u_2, X_2)

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FUNCTIONMERGING(u_1, X_1, u_2, X_2)

dynamic programming states: $2^{\mathcal{O}(b_1 d_1 + b_2 d_2)} \cdot |F_1| \cdot |F_2|$.

Work per state: $\text{poly}(b_1, b_2, d_1, d_2) \cdot (|F_1| + |F_2|)$.

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Optimized $X_1, X_2 \Rightarrow \text{poly}(b_1, b_2, d_1, d_2)$ work per state.

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Theorem (Restricting b_1, b_2, d_1, d_2)

Function Merging with Branch Reordering can be solved in

$2^{\mathcal{O}(b_1 d_1 + b_2 d_2)} \cdot |F_1| \cdot |F_2|$ *time.*

Overview of the second algorithm

Theorem (Restricting b_1, b_2, d_1)

Function Merging with Branch Reordering can be solved in

$$2^{\mathcal{O}(b_1 d_1 + b_2)} \cdot (|F_1| \cdot |F_2|)^4 \text{ time.}$$

View first algorithm as

weighted reachability on a finite graph

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Overview of the second algorithm

Theorem (Restricting b_1, b_2, d_1)

Function Merging with Branch Reordering can be solved in

$$2^{\mathcal{O}(b_1 d_1 + b_2)} \cdot (|F_1| \cdot |F_2|)^4 \text{ time.}$$

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Use classical reachability algorithm [SAS'03].