

# Stateless Model Checking Concurrent/Distributed Programs

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Michalis Kokologiannakis    Viktor Vafeiadis  
POPL 2025

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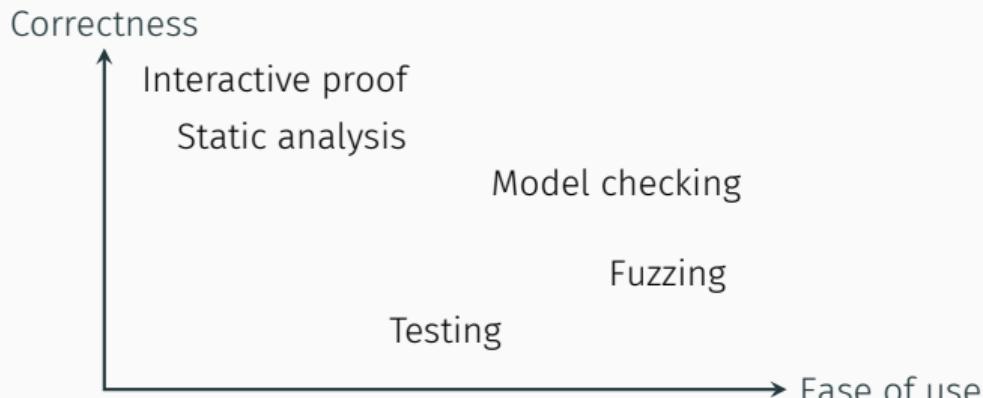
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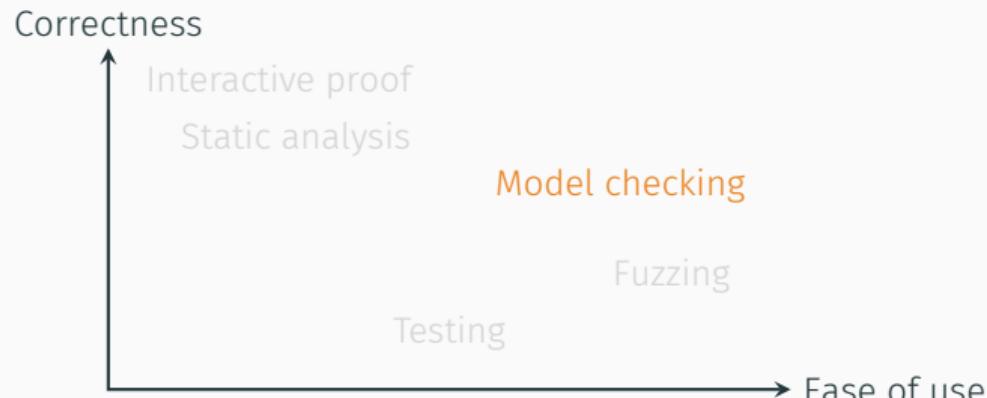
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# Model-checking approaches

## Stateful:

- Visit program states **while recording** visited states
- Assumes program has **bounded state-space**
- **High** memory usage

## Stateless:

- Visit program states **without recording** visited states
- Assumes program **always terminates**
- **Low** memory usage

## SMT-based:

- Encode program and specification as an **SMT query**
- Assumes program **always terminates**
- **High** memory usage

# Our weapon of choice

[PLDI'19] Model checking weakly-consistent libraries

[POPL'22] Truly stateless, optimal dynamic partial order reduction

**GENMC**: state-of-the-art **stateless** model checker

- Correct, optimal, highly-parallelizable
- Works with almost any memory model
- Small memory footprint



[plv.mpi-sws.org/genmc](http://plv.mpi-sws.org/genmc)

Two papers in POPL'25 (Thu @ 15:00):

- Automatically checking linearizability under weak memory consistency
- Model checking C/C++ with mixed-size accesses

# Outline

How does GENMC work?

- SMC basics
- Execution graphs
- Exploration algorithm

How to apply GENMC to our code?

- State-space reductions
- Estimating state-space size
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Each part will be followed by a demo

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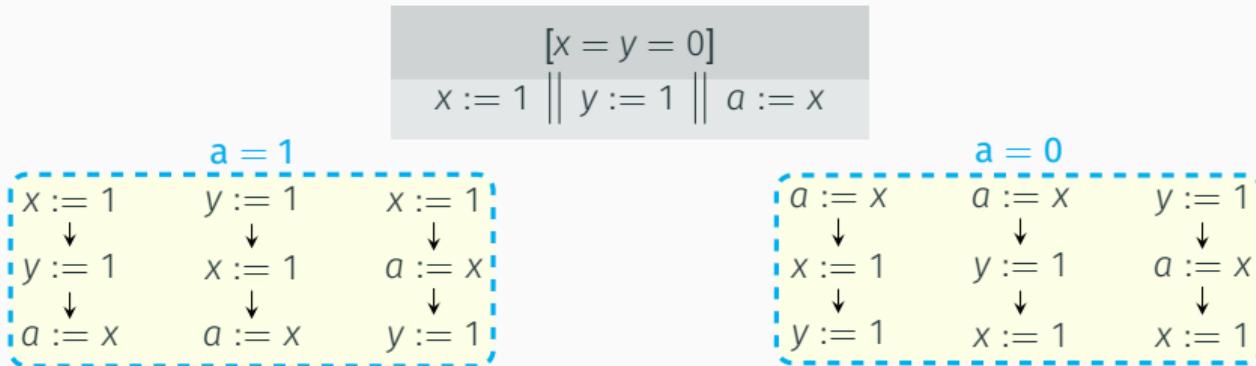
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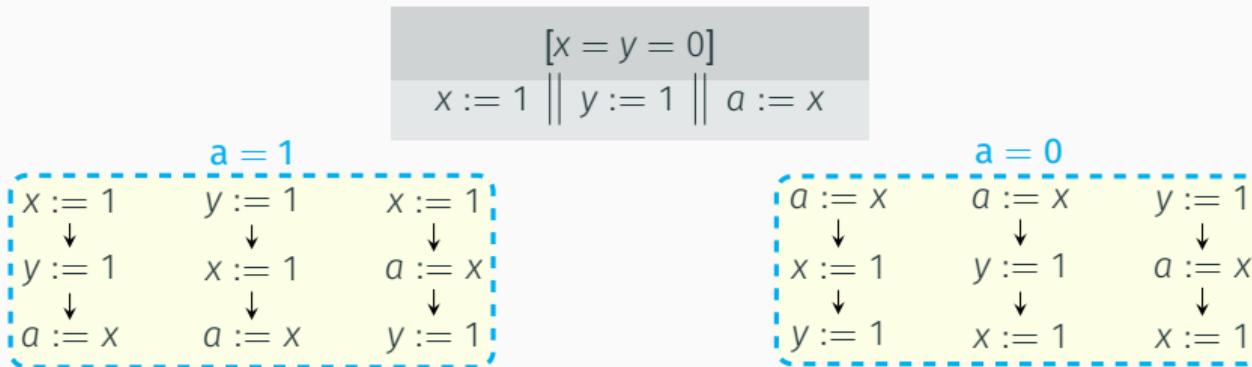
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How to enumerate one interleaving per equivalence class?  
(Partial order reduction)

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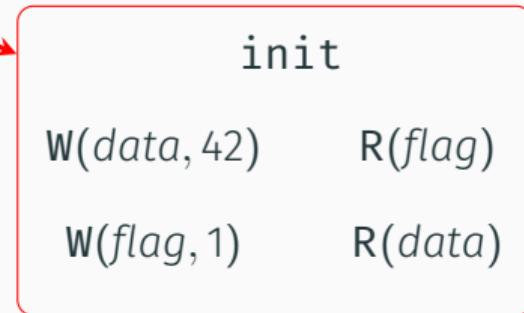
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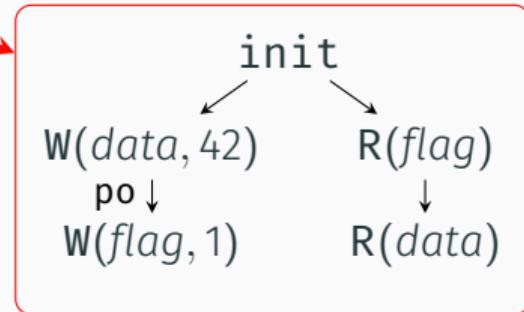
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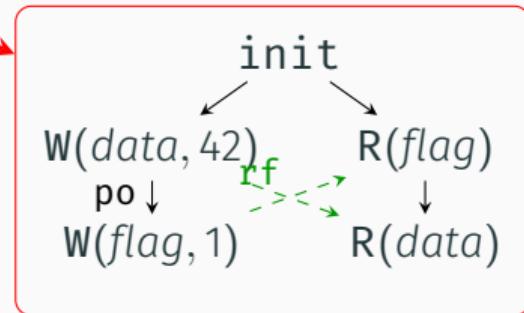
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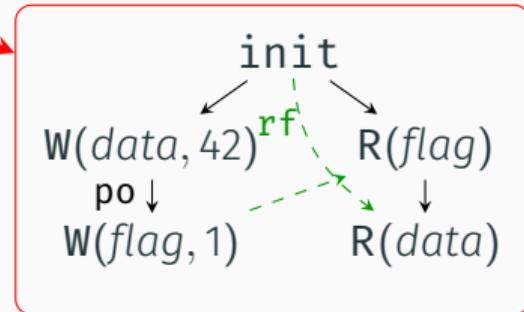
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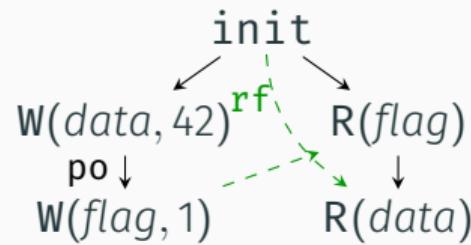
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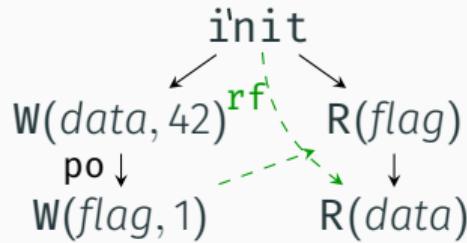


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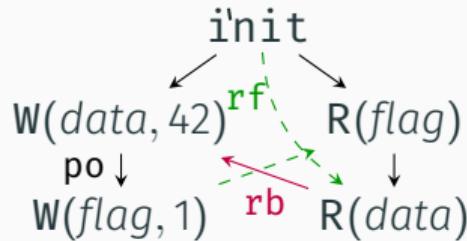


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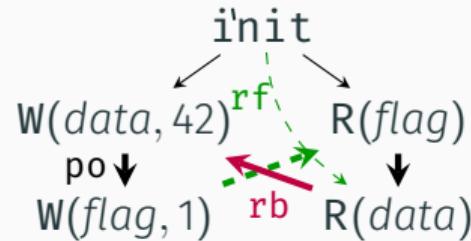


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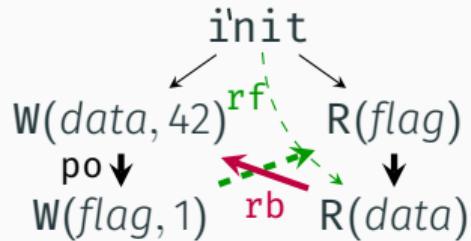


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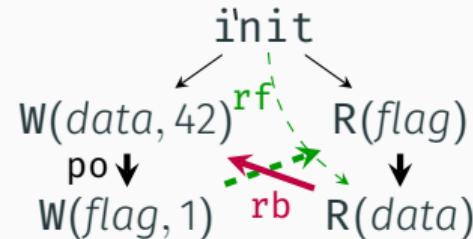
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Real model SC: irreflexive $((\text{po} \cup \text{rf} \cup \text{co} \cup \text{rb})^+)$

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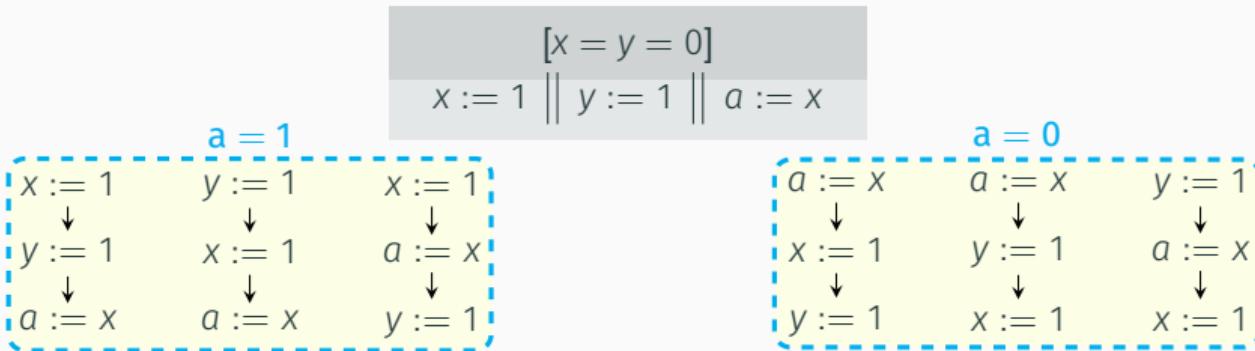
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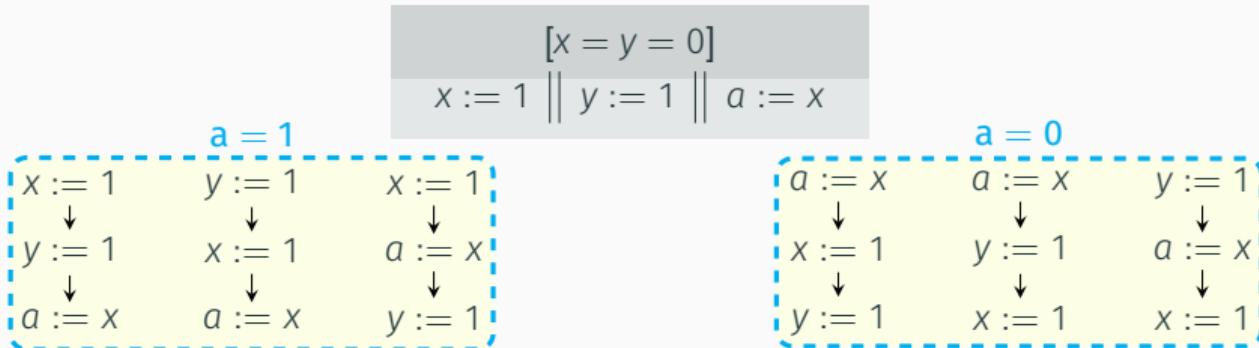
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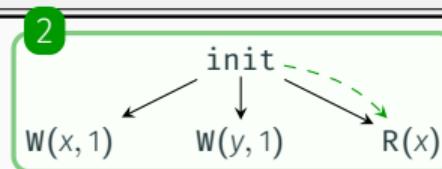
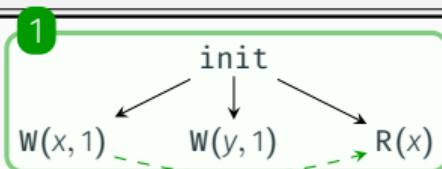
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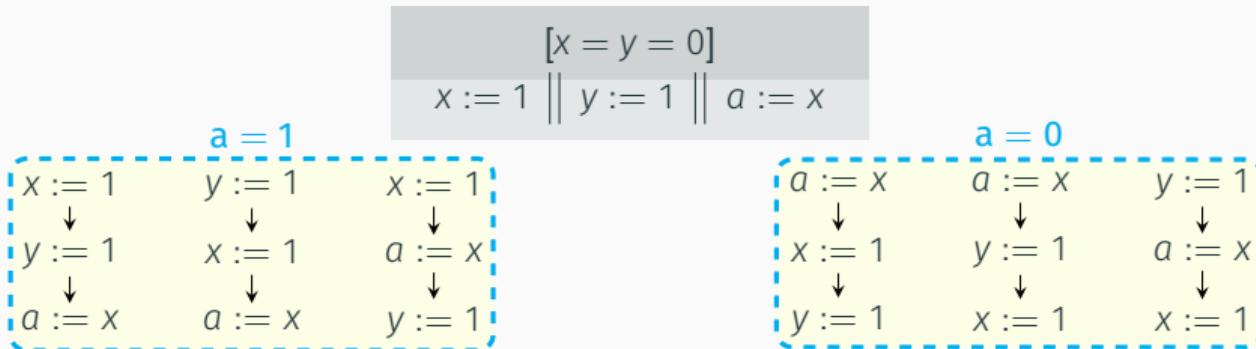


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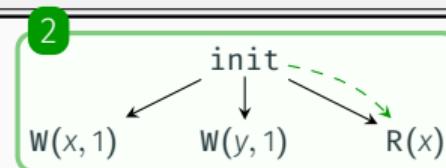
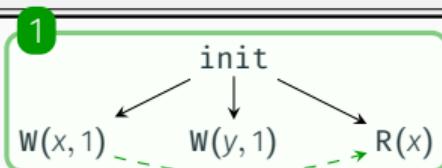


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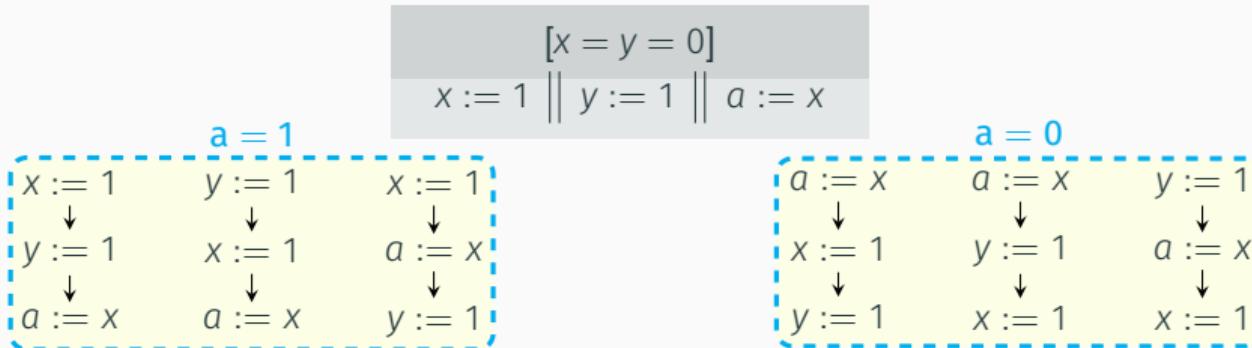
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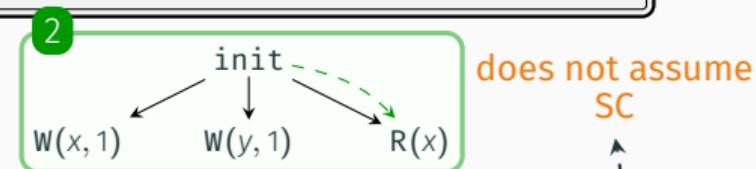
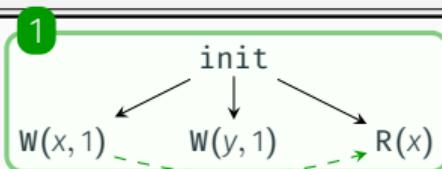
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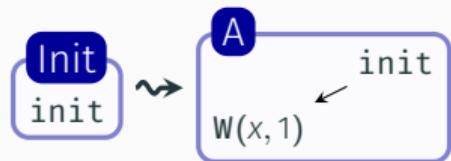
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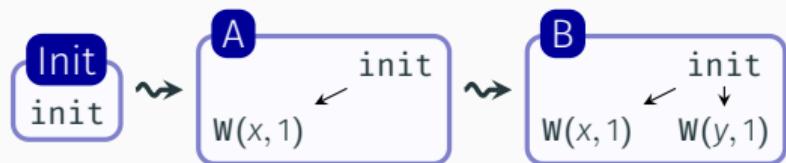
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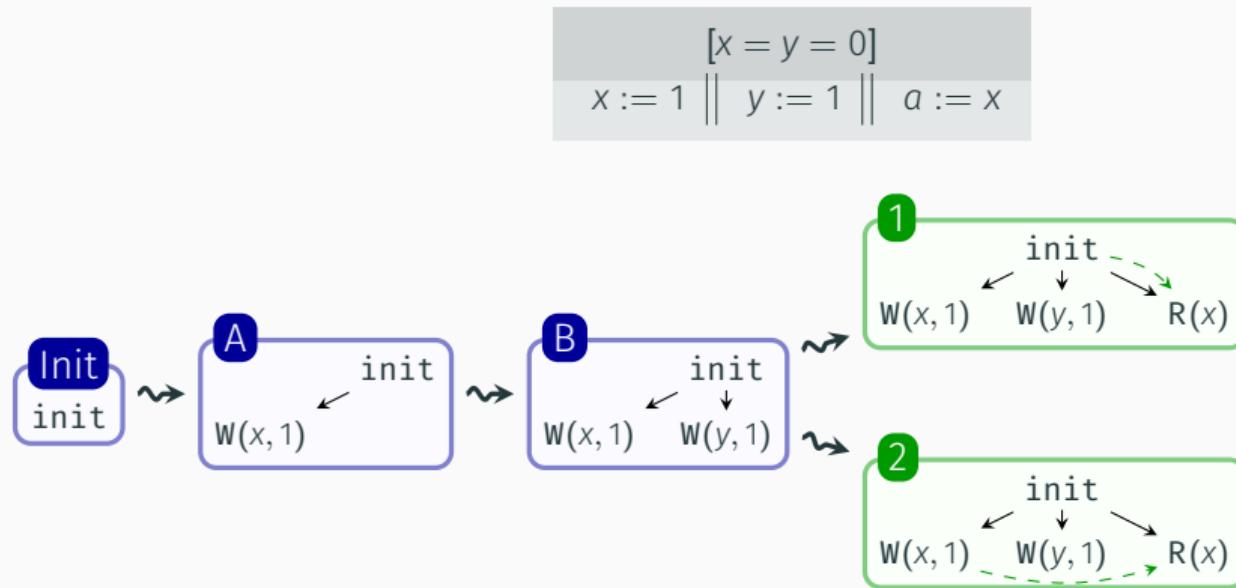
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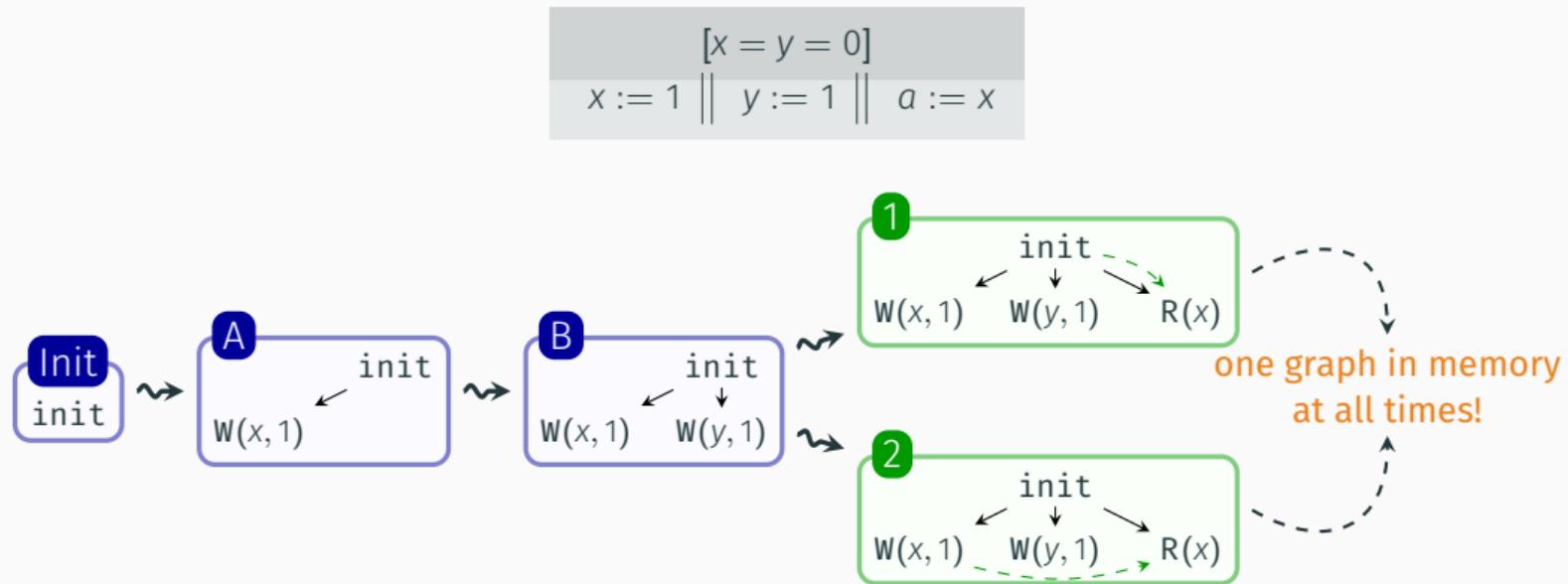
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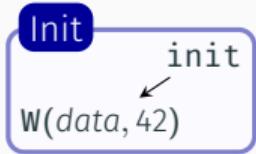


## GENMC's algorithm: Example #2

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[data = flag = 0]
data := 42  ||  if (flag = 1)
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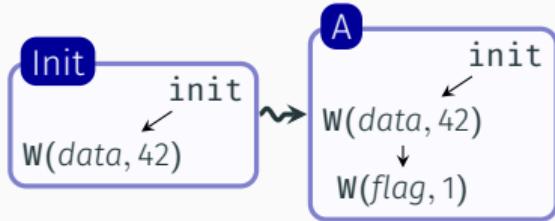
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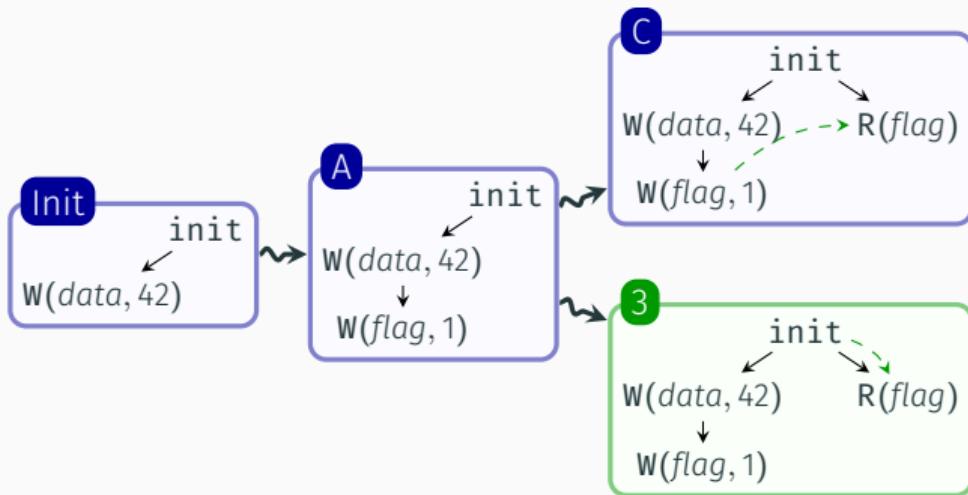
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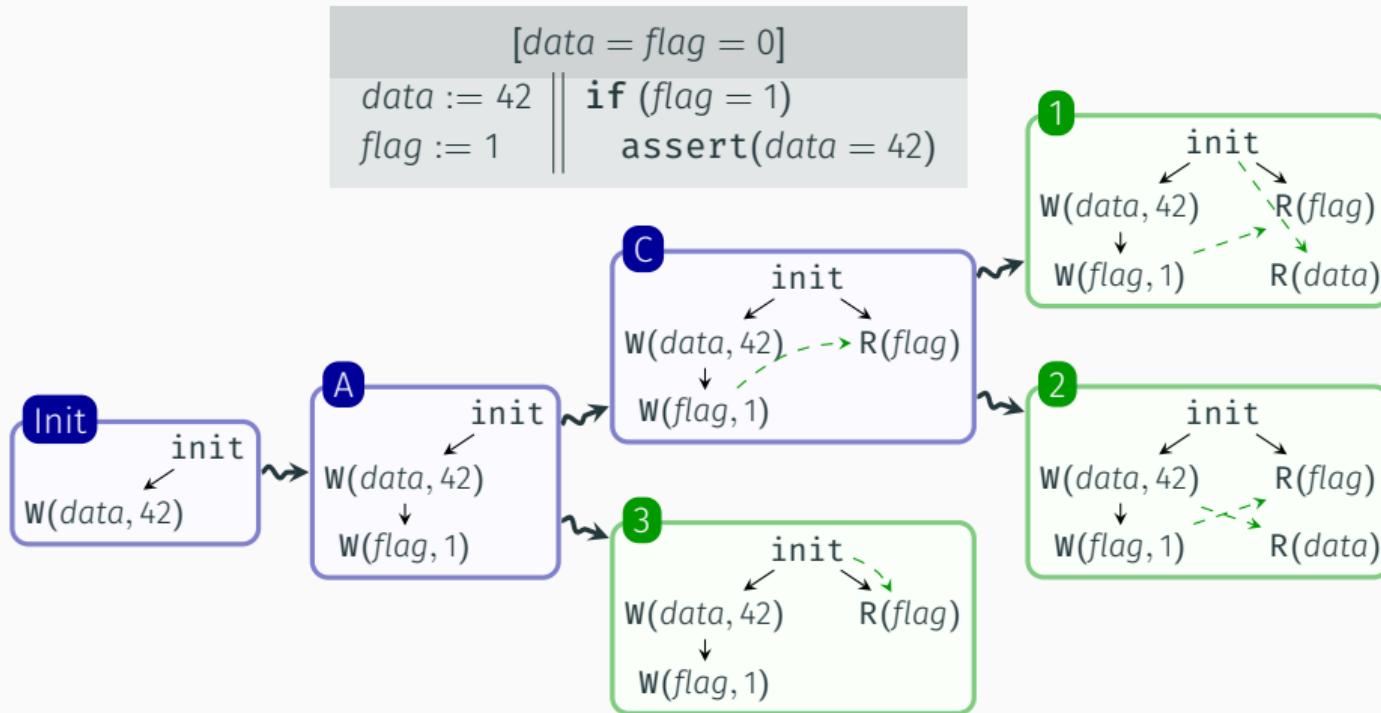


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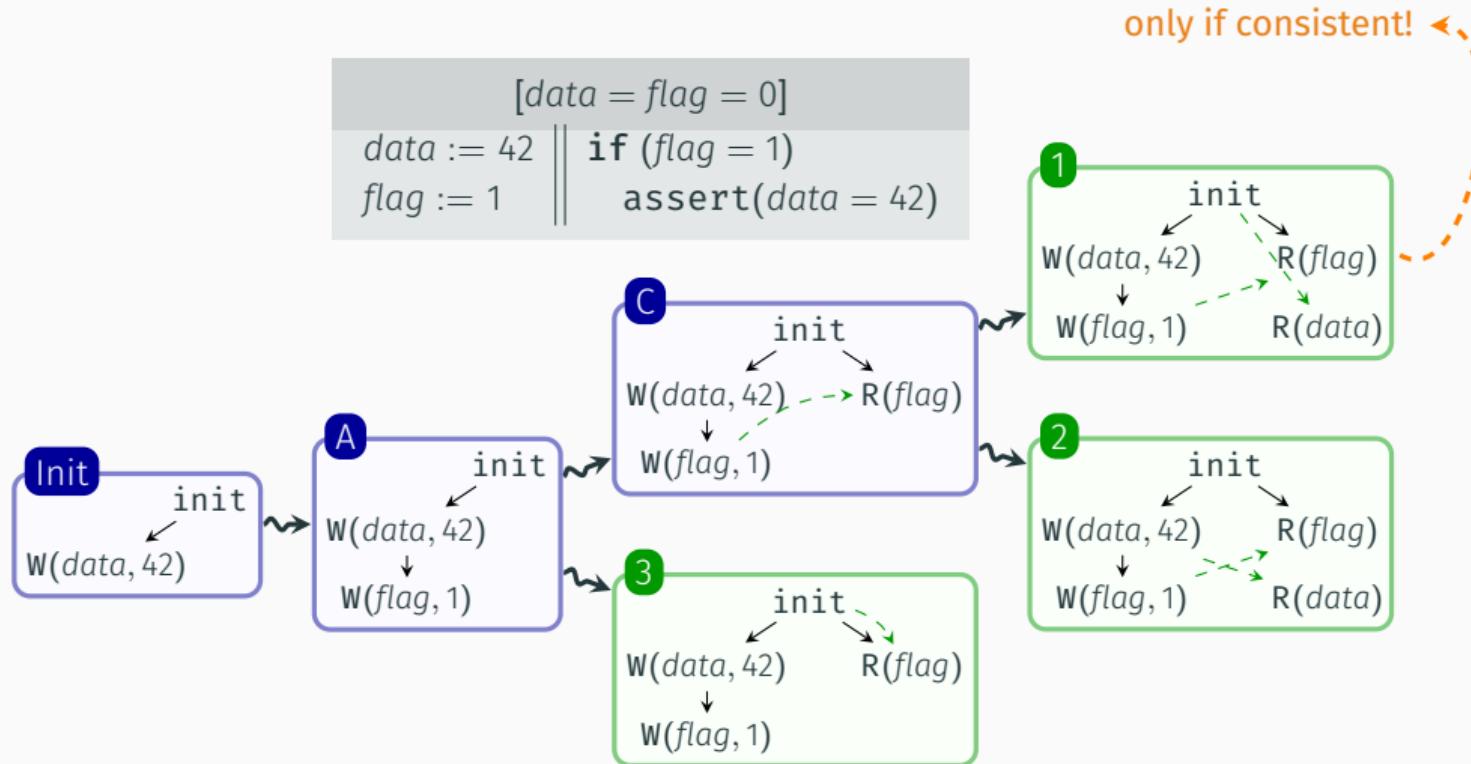
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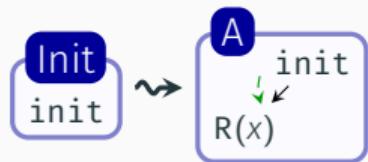
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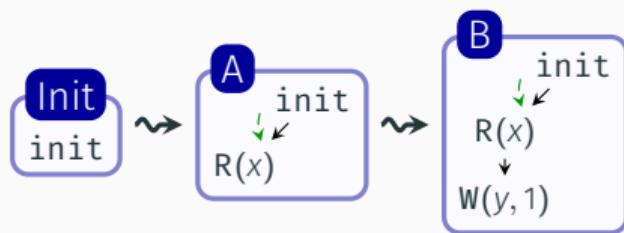
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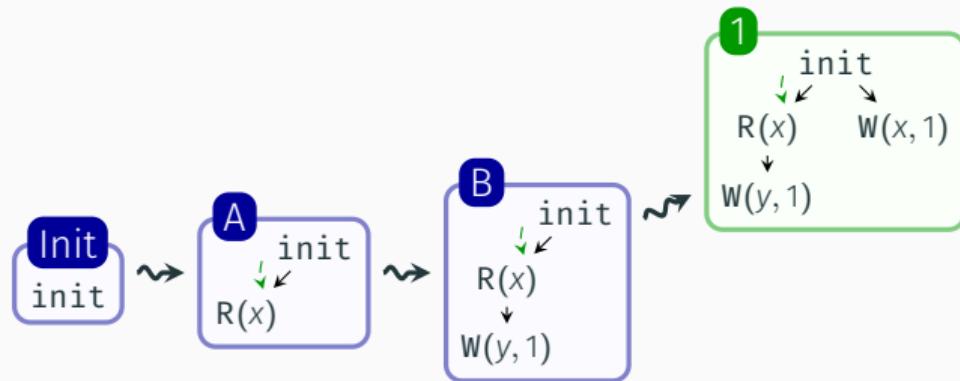
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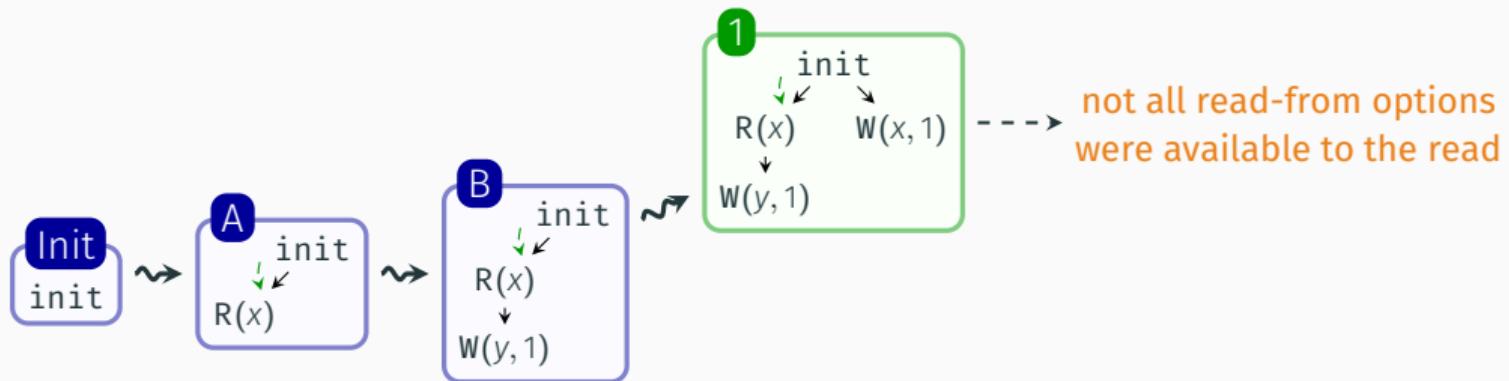
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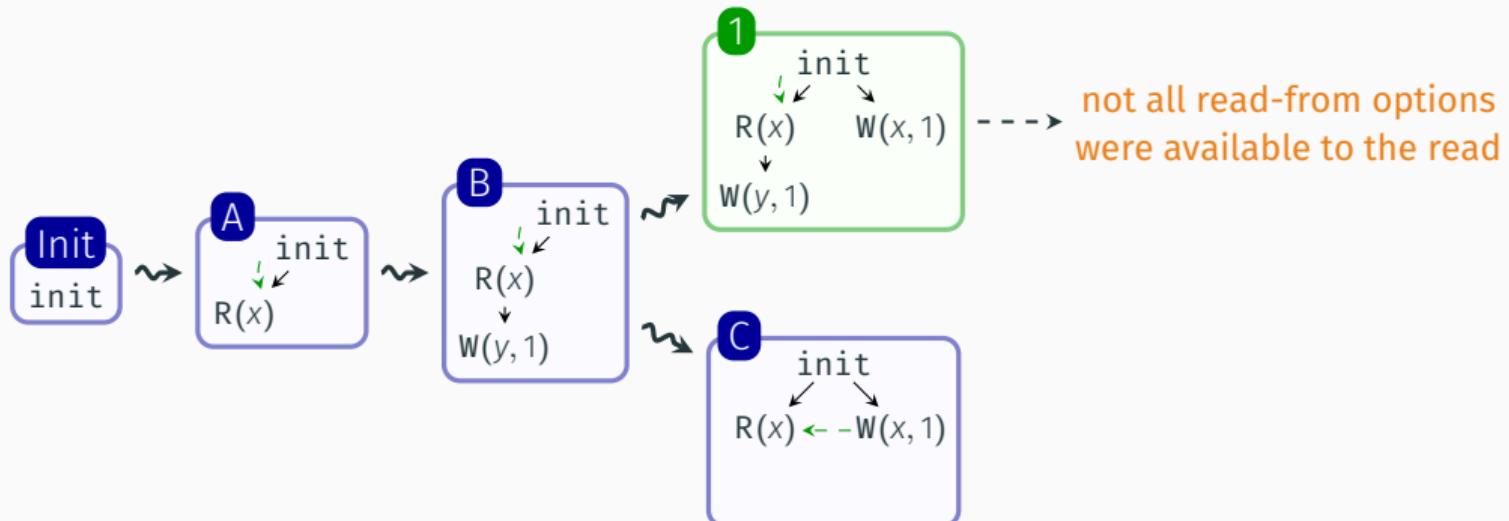
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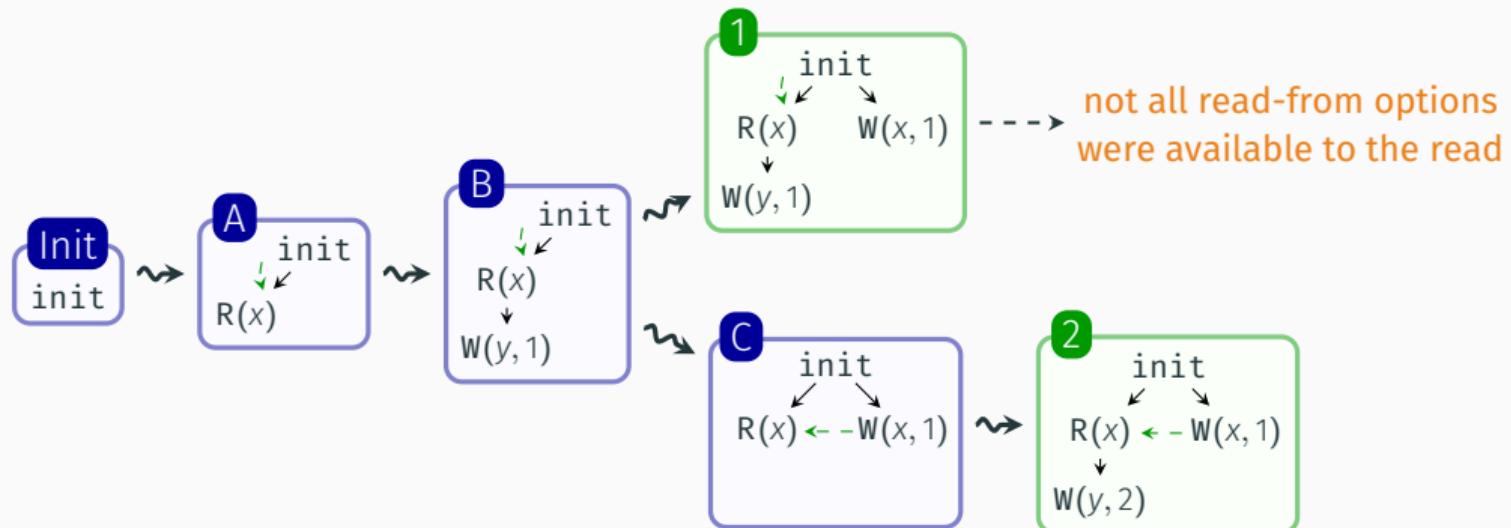


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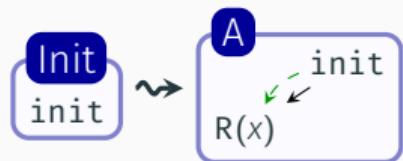
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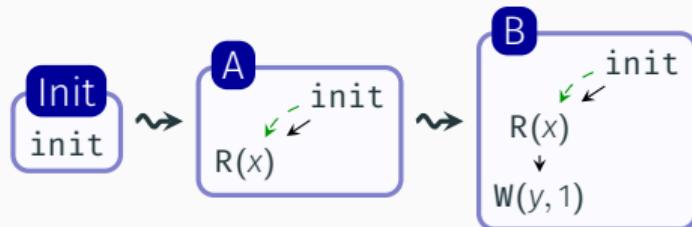
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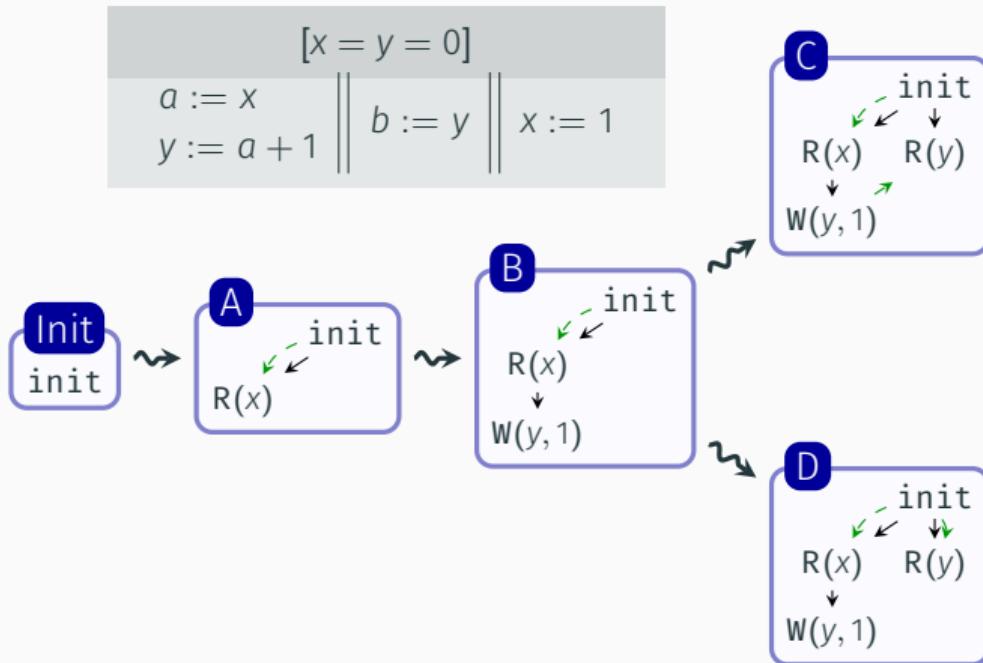


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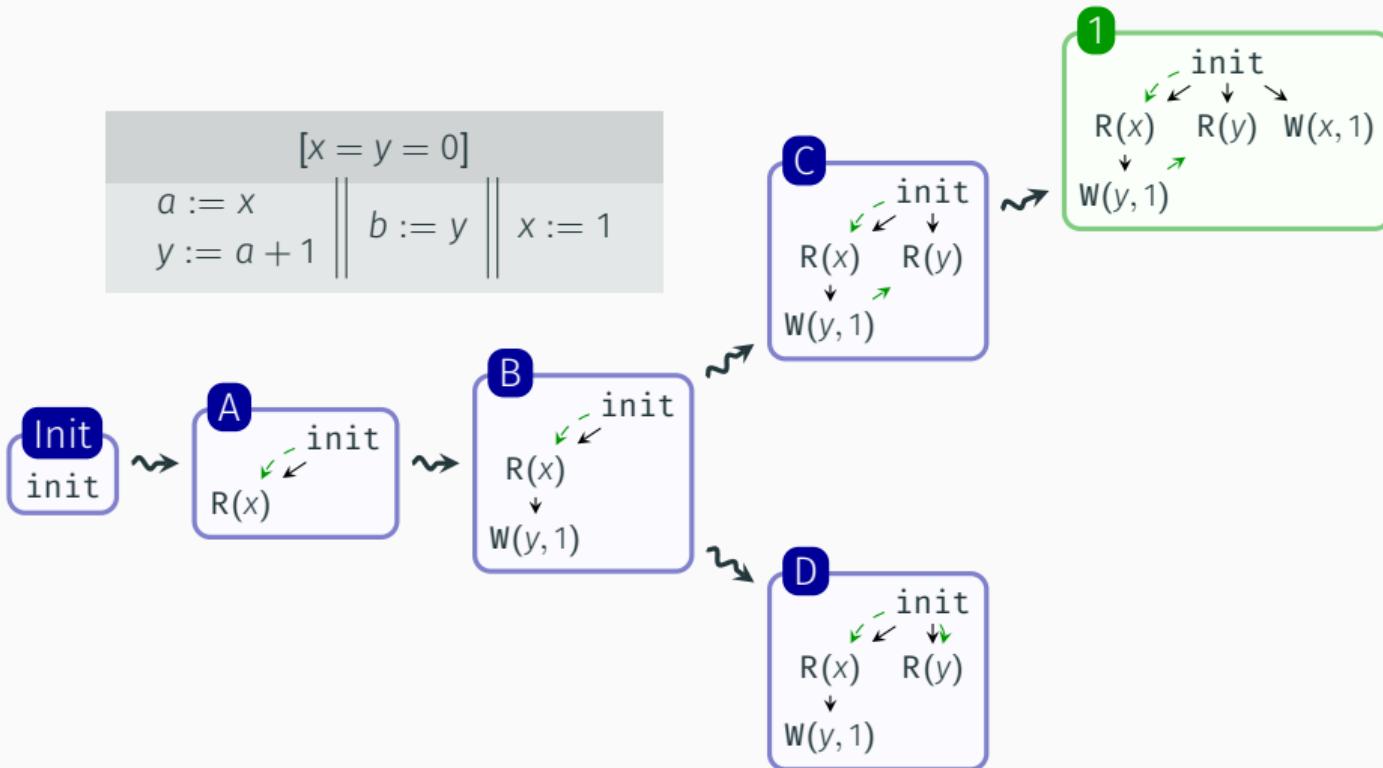
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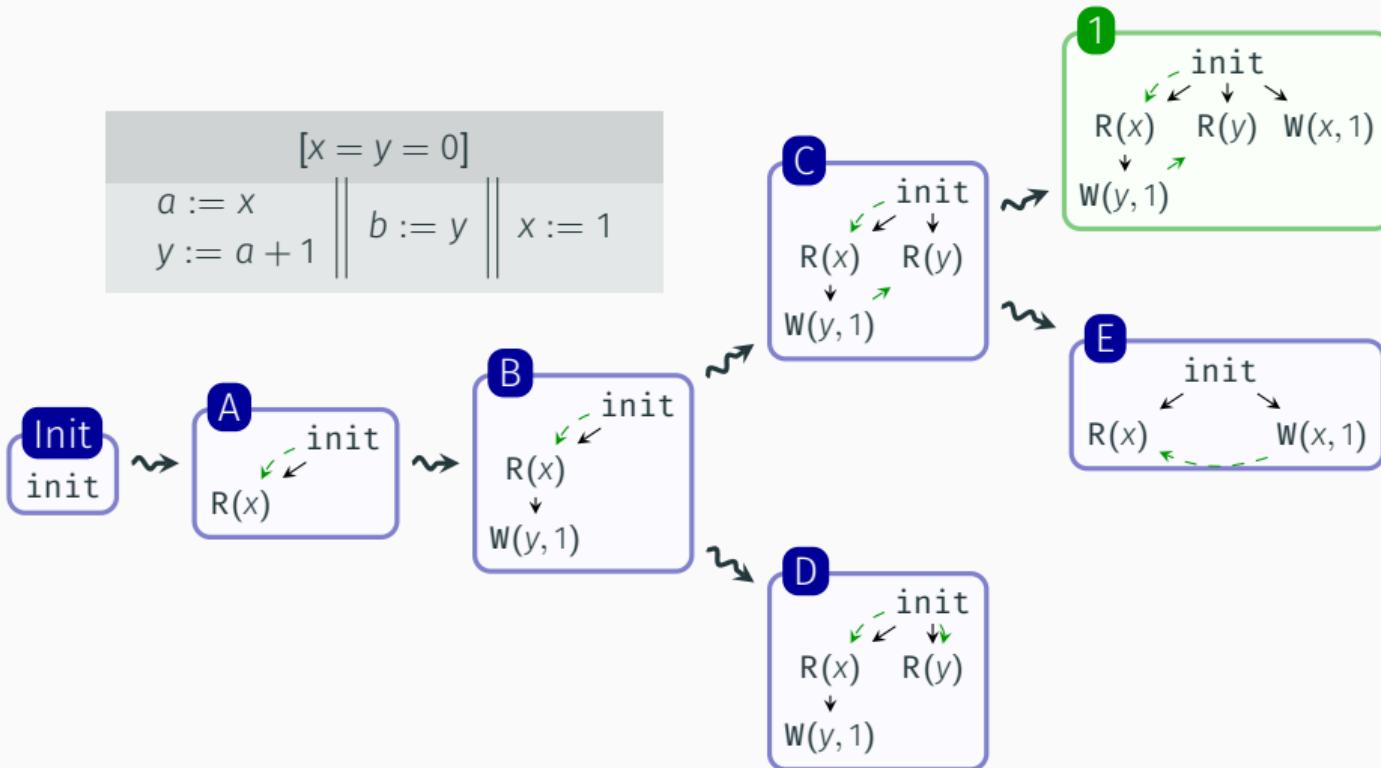
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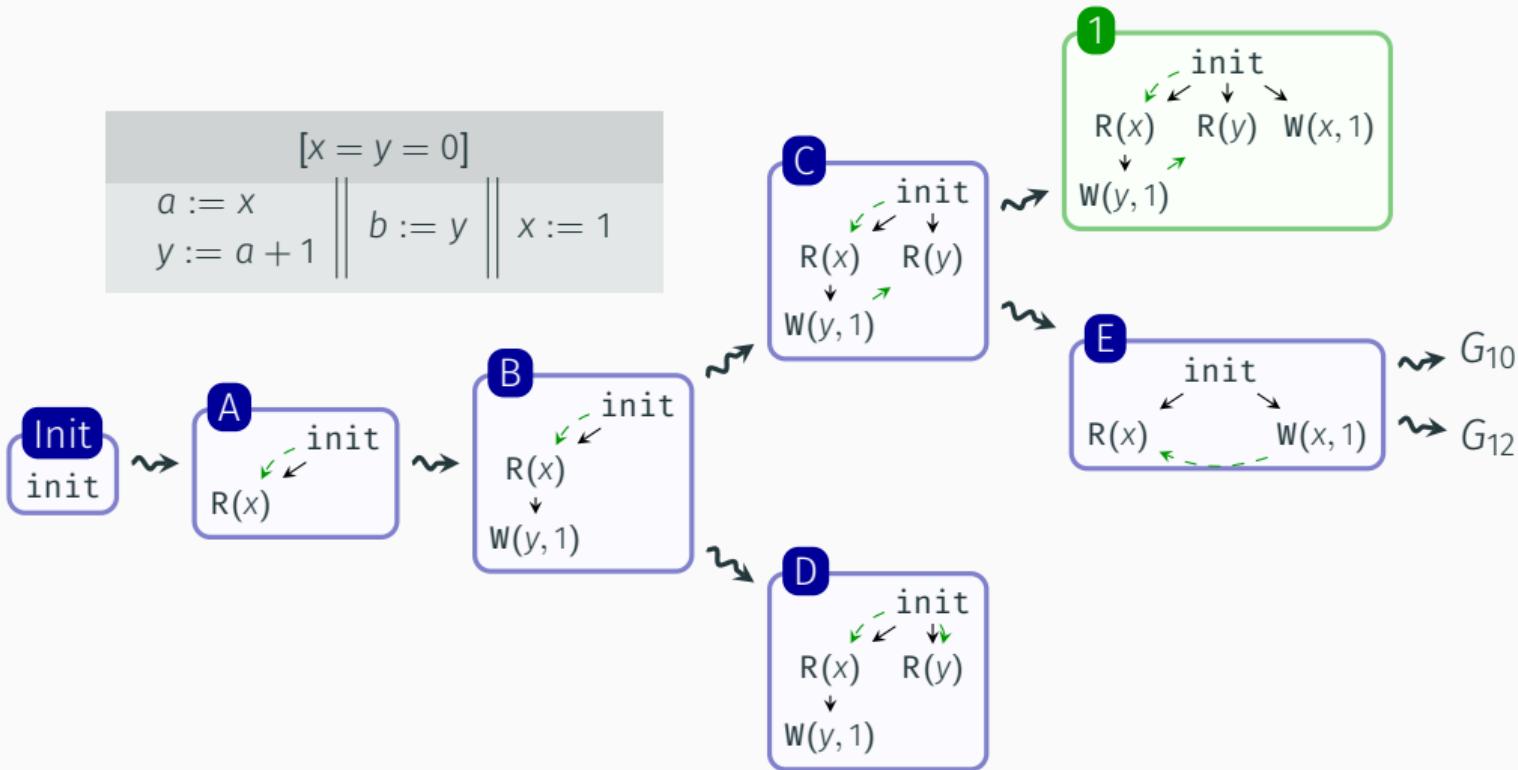
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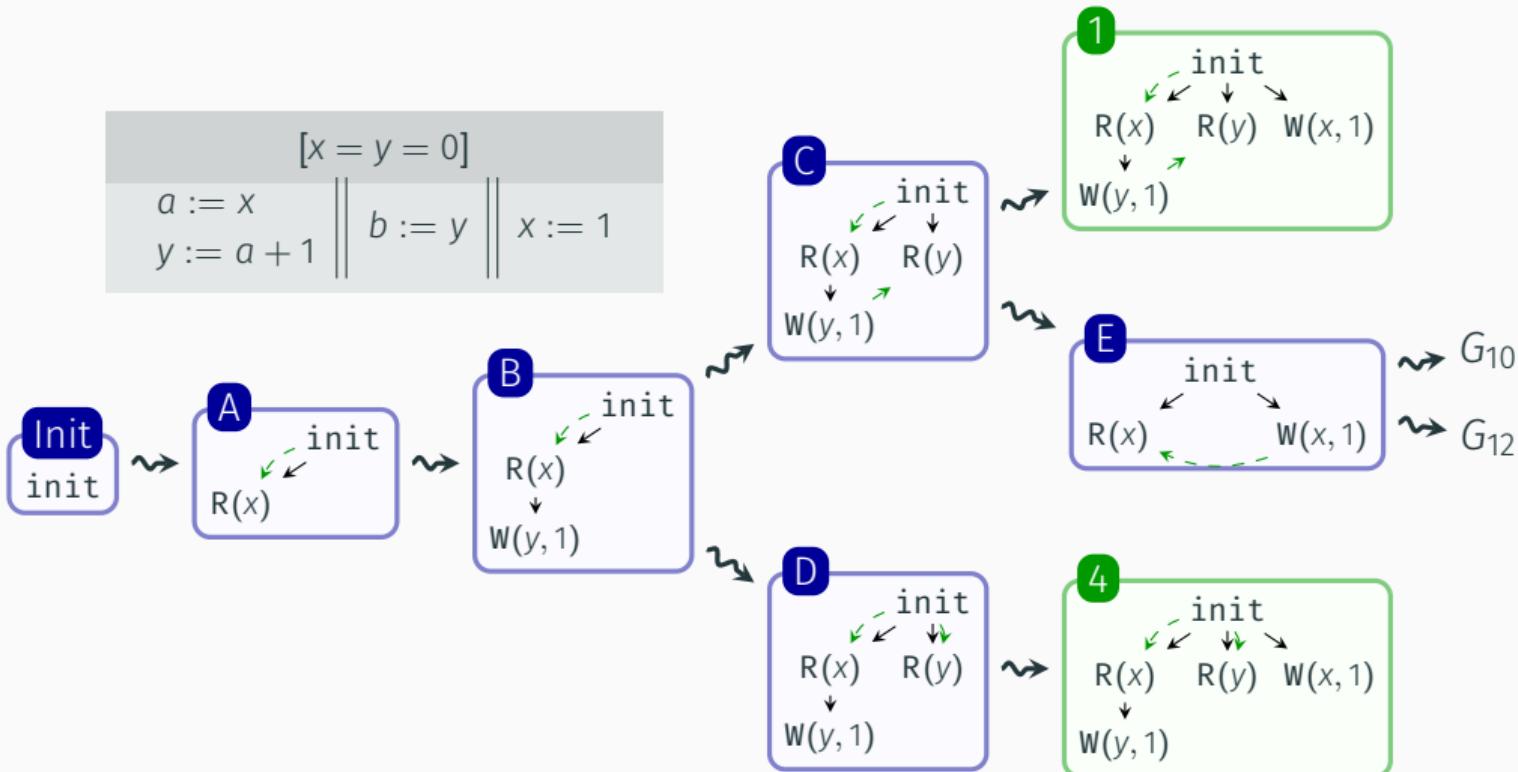
# GENMC's algorithm: Example #4



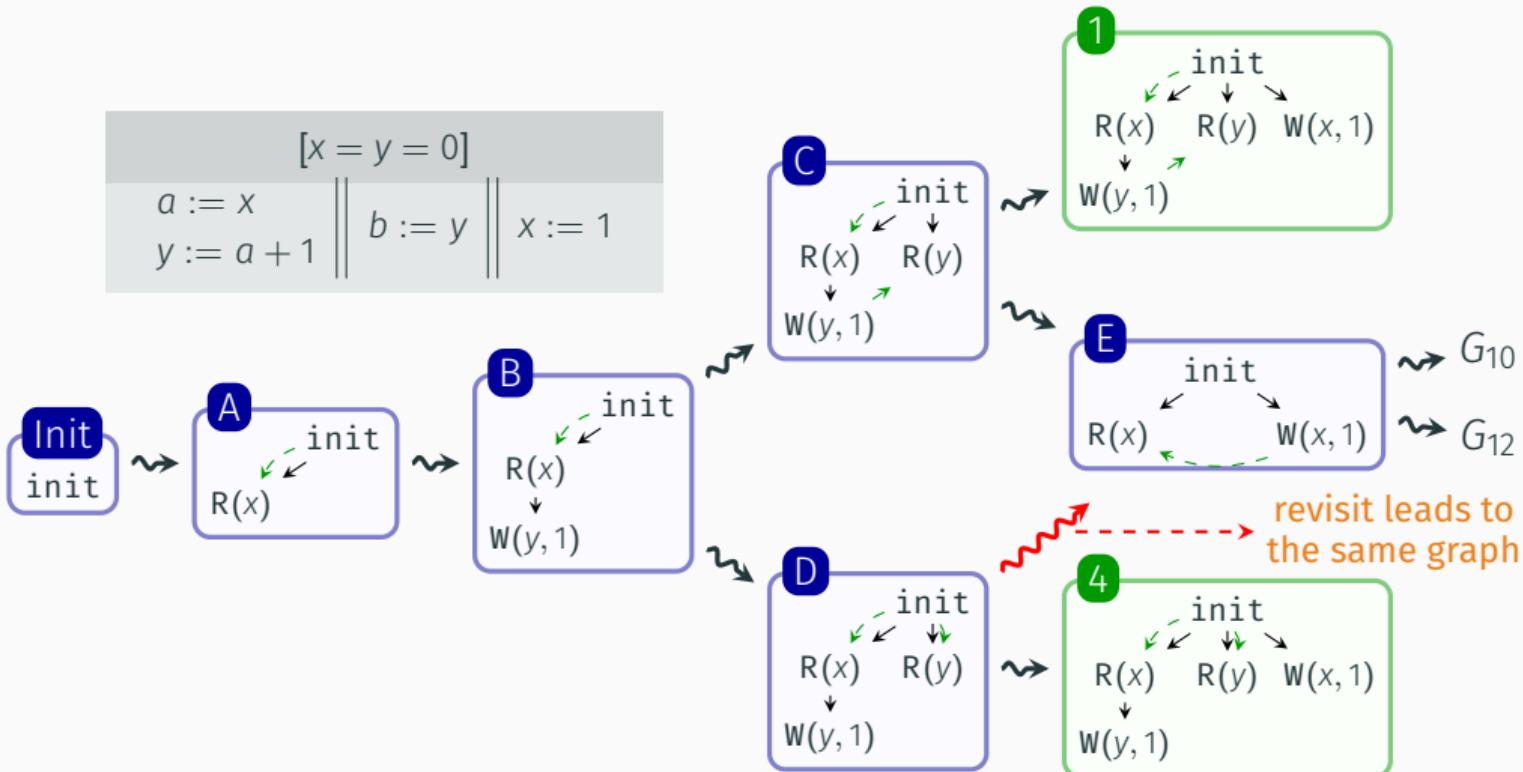
# GENMC's algorithm: Example #4



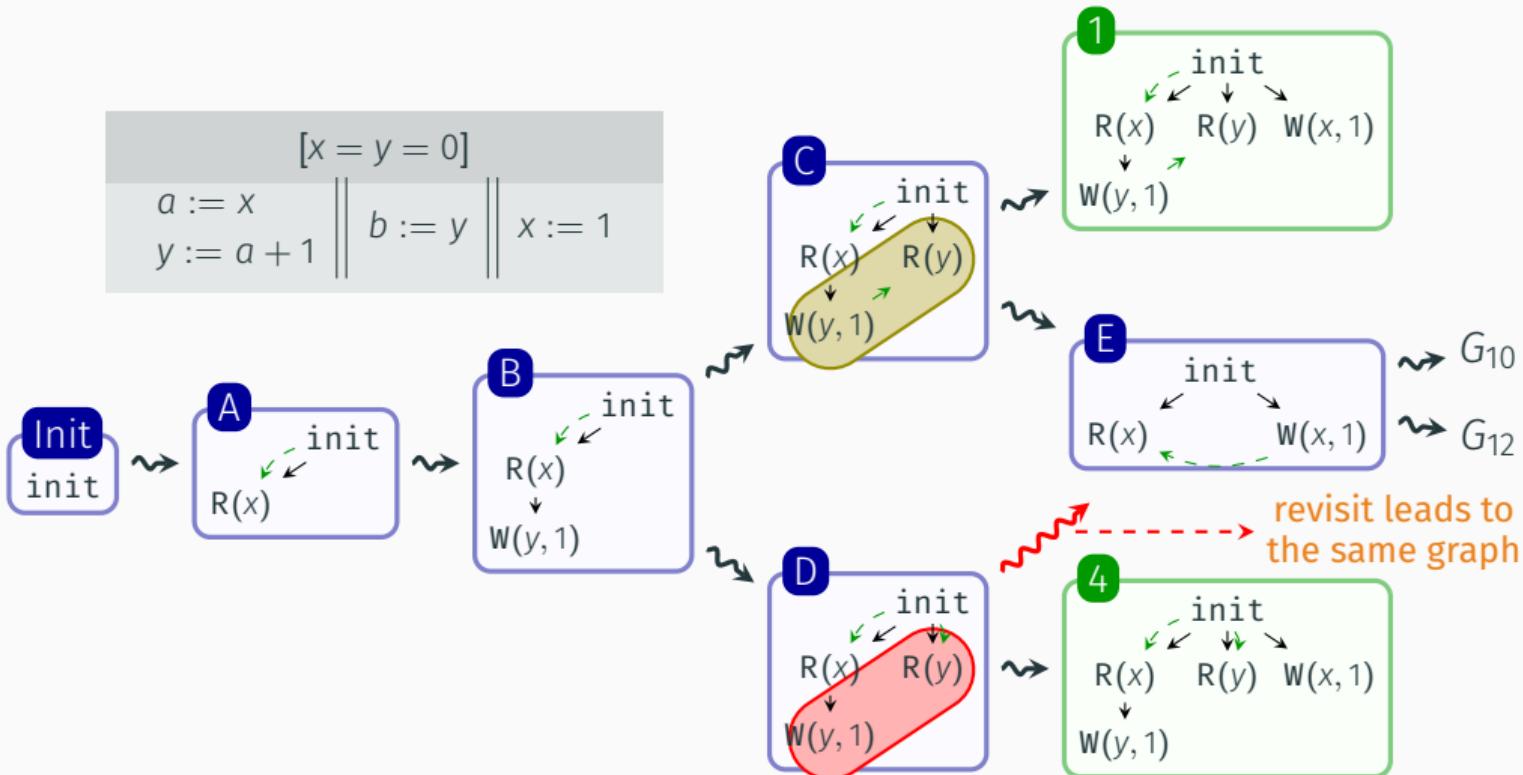
# GENMC's algorithm: Example #4



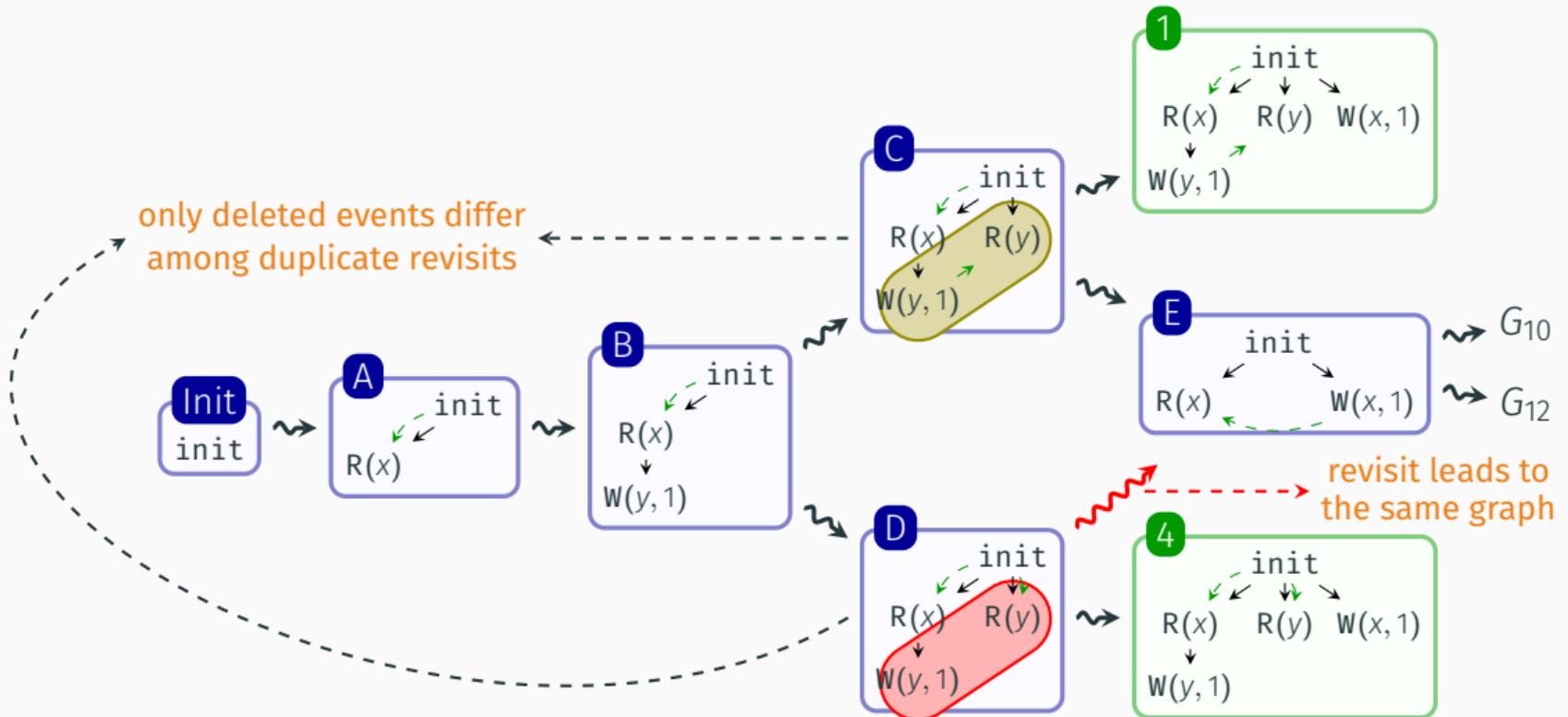
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## Memory-model conditions

GENMC enumerates all consistent execution graphs for **any** memory model  $M$ , if

- $\text{cons}_M(\cdot)$  implies irreflexive $((\text{po} \cup \text{rf})^+)$
- $\text{cons}_M(\cdot)$  is **prefix-closed**
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GENMC enumerates all consistent execution graphs for **any** memory model  $M$ , if

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- $\text{cons}_M(\cdot)$  is **prefix-closed**
- $\text{cons}_M(\cdot)$  is **maximally extensible**

These conditions hold for SC, TSO, PSO, RC11  
(can be relaxed for POWER, ARM, IMM, LKMM)

# Outline

How does GENMC work?

- SMC basics
- Execution graphs
- Exploration algorithm

How to apply GENMC to our code?

- State-space reductions
- Estimating state-space size
- Exploration bounding

Each part will be followed by a demo

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Each part will be followed by a **demo**

# Demo #1

## 1. Install Docker:

Debian/Ubuntu:

```
apt install docker.io
```

MacOS:

```
brew install --cask docker
```

Windows:

```
wsl --install
```

```
apt install docker.io
```

## 2. Run GENMC container:

```
docker pull genmc/genmc
```

```
docker run -it genmc/genmc:latest
```

## 3. Download tutorial material:

```
wget https://plv.mpi-sws.org/genmc/popl2025/examples.tar.gz
```

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# Approaches for state-space explosion

Partial order reduction (POR):

Avoid ordering **independent actions**

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$$a_1 = \dots = a_N = 0$$

$$a_1 := 1 \parallel \dots \parallel a_N := N$$

$$\# \text{ of executions} \left\{ \begin{array}{l} \text{SMC : } N! \\ \text{POR : } 1 \\ \vdots \end{array} \right.$$

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Symmetry reduction (SR):

Avoid ordering **symmetric threads**

$$x = 0$$

$$\text{fetch\_add}(x, 1) \parallel \dots \parallel \text{fetch\_add}(x, 1)$$

$$\# \text{ of executions} \left\{ \begin{array}{l} \text{SMC : } N! \\ \text{POR : } N! \\ \text{SR : } 1 \end{array} \right.$$

# Combining SR and POR

[PLDI'24] SPORE: Combining symmetry and partial order reduction

$$x = a[1] = \dots = a[N] = 0$$

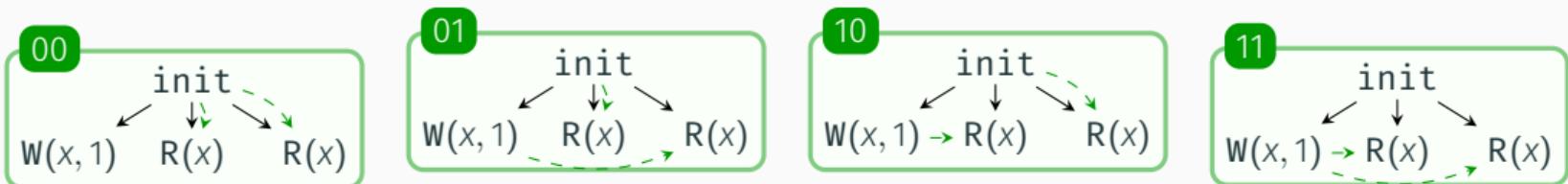
$i := \text{fetch\_add}(x, 1)$  || ... ||  $i := \text{fetch\_add}(x, 1)$   
 $a[i] := i$                    || ... ||  $a[i] := i$

SMC	:	$(2N)!/(N \cdot 2!)$
POR	:	$N!$
SR	:	$(2N - 1)!!$
POR + SR	:	1

# Combining SR and POR

[PLDI'24] SPORE: Combining symmetry and partial order reduction

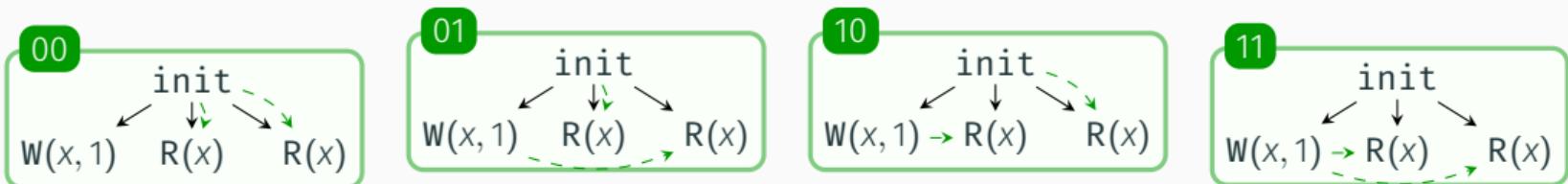
$$\begin{array}{c} [x = 0] \\ \top_1: x := 1 \parallel \top_2: r := x \parallel \top_3: r := x \end{array}$$



# Combining SR and POR

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$$[x = 0] \\ T_1: x := 1 \parallel T_2: r := x \parallel T_3: r := x$$

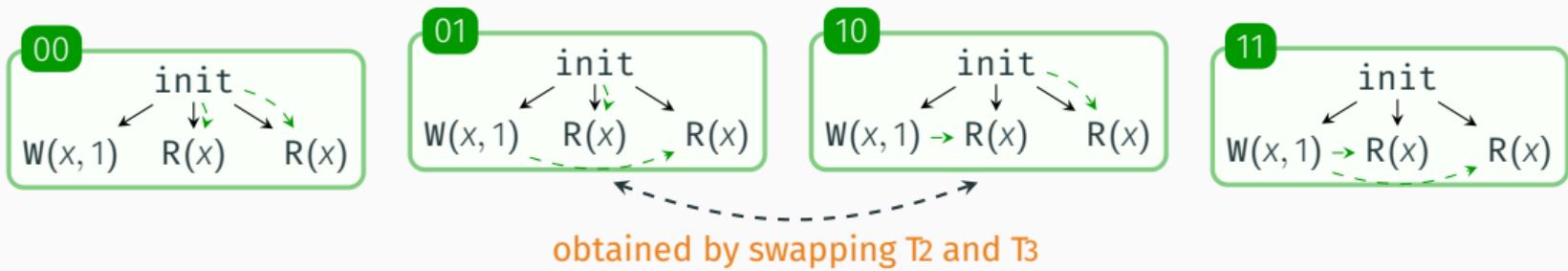


# Combining SR and POR

[PLDI'24] SPORE: Combining symmetry and partial order reduction

[ $x = 0$ ]

**Key Idea:** Identify symmetries on the execution graphs

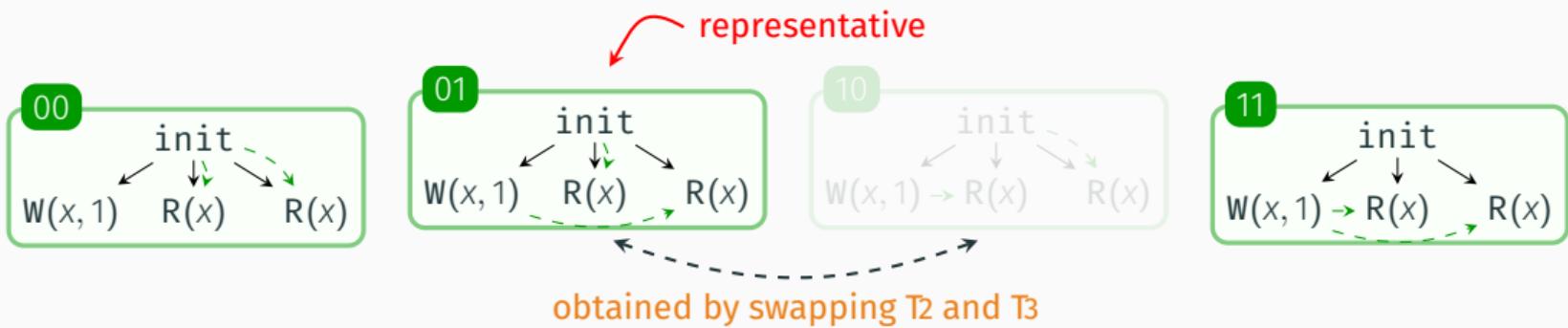


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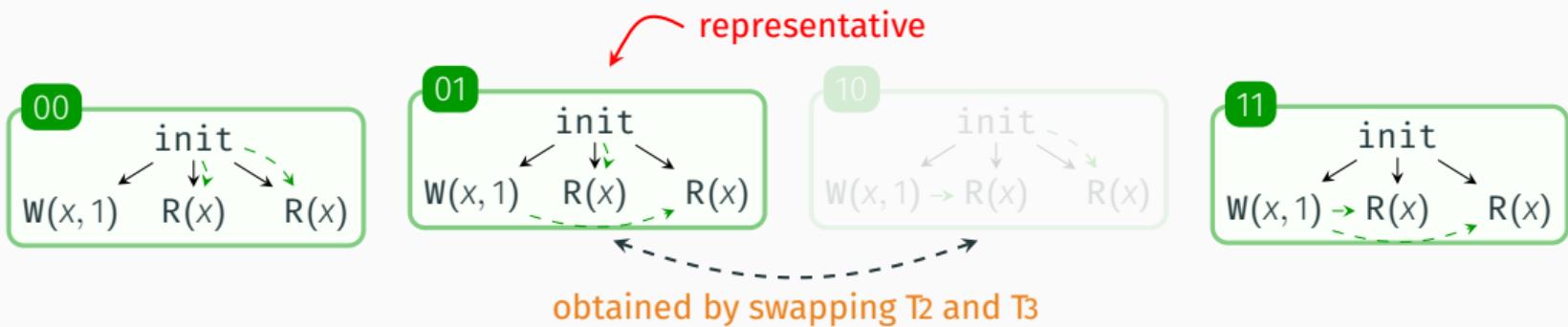


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**Key Idea:** Identify symmetries on the execution graphs  
Only generate **representative** graphs

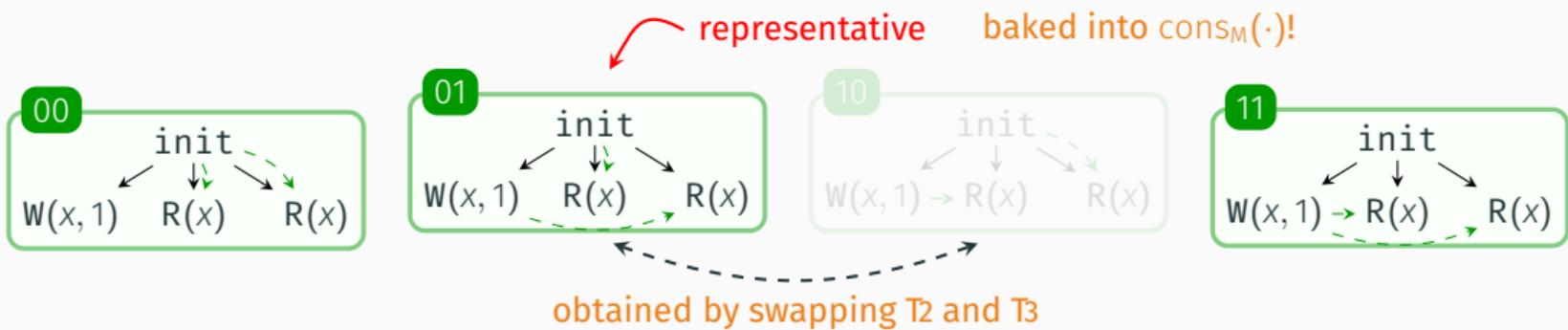


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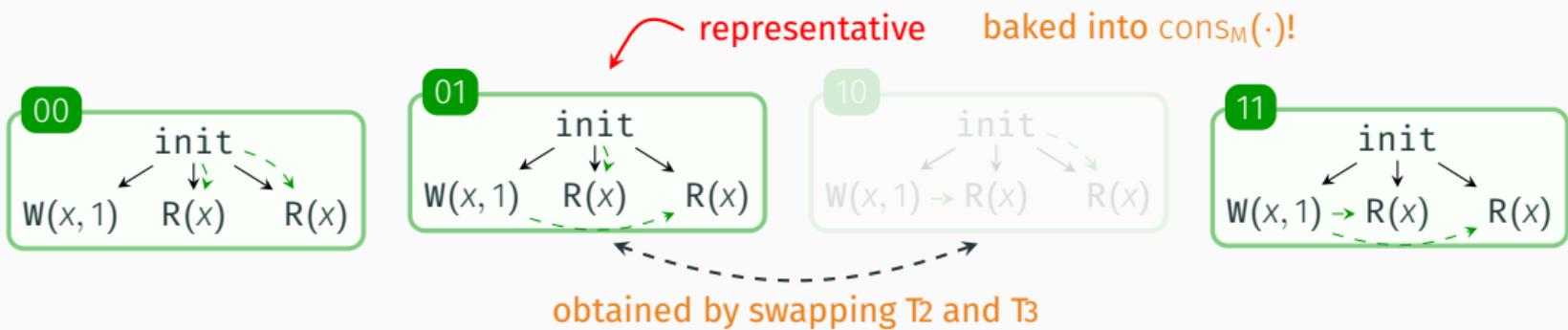


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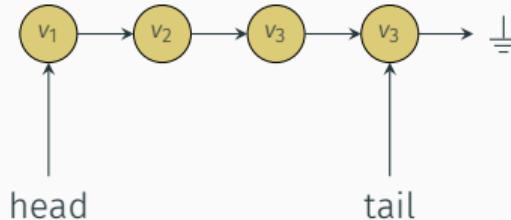
There are many types of symmetries  
~~ these can be incorporated into  $\text{cons}_M(\cdot)$

# Internal symmetries example: Michael-Scott queue

```
enqueue(v)  $\triangleq$ 
  node := malloc(...)
  node.value := v
  node.next := NULL
  do
    t := tail
    next := t.next
    if (t  $\neq$  tail) continue
    if (next  $\neq$  NULL)
      CAS(tail, t, next)
      continue
  while ( $\neg$ CAS(t.next, next, node))
  CAS(tail, t, node)
```

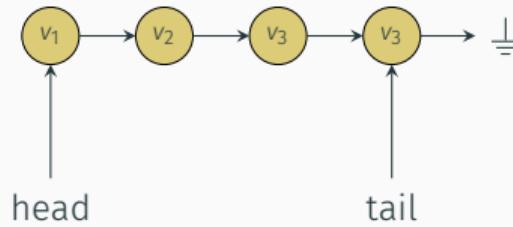
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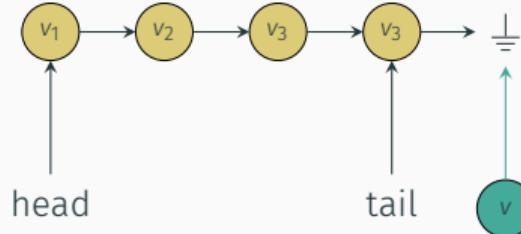
# Internal symmetries example: Michael-Scott queue

$\text{enqueue}(v) \triangleq$   
 $\xrightarrow{T_1}$  *node* := malloc(...)  
*node.value* := *v*  
*node.next* := NULL  
do  
  *t* := tail  
  *next* := *t.next*  
  **if** (*t*  $\neq$  tail) **continue**  
  **if** (*next*  $\neq$  NULL)  
    CAS(tail, *t*, *next*)  
    **continue**  
**while** ( $\neg$ CAS(*t.next*, *next*, *node*))  
  CAS(tail, *t*, *node*)



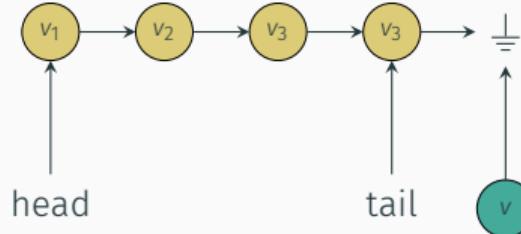
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 $\textcolor{teal}{\rightarrow} \text{ } T_1$      $\text{node := malloc(...)}$   
           $\text{node.value := } v$   
           $\text{node.next := NULL}$   
**do**  
     $t := \underline{\text{tail}}$   
     $\text{next := } t.\text{next}$   
    **if** ( $t \neq \underline{\text{tail}}$ ) **continue**  
    **if** ( $\text{next} \neq \text{NULL}$ )  
       $\text{CAS}(\underline{\text{tail}}, t, \text{next})$   
      **continue**  
**while** ( $\neg \text{CAS}(t.\text{next}, \text{next}, \text{node}))$   
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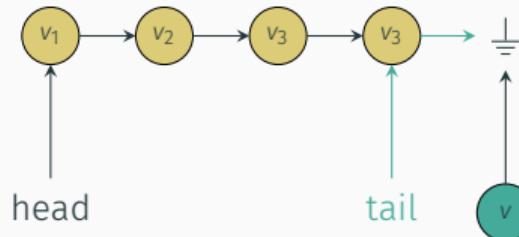
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$\text{enqueue}(v) \triangleq$   
 $\text{node} := \text{malloc}(\dots)$   
 $\text{node.value} := v$   
 $\text{node.next} := \text{NULL}$

**do**

T<sub>1</sub>

```
t := tail
next := t.next
if (t ≠ tail) continue
if (next ≠ NULL)
    CAS(tail, t, next)
    continue
while (¬CAS(t.next, next, node))
    CAS(tail, t, node)
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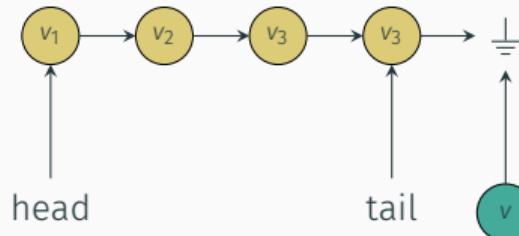
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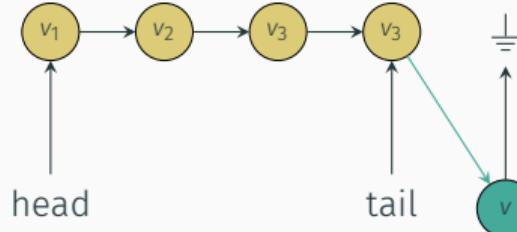
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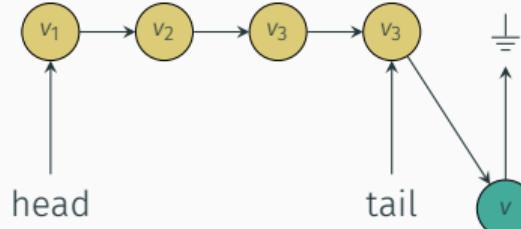
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# Internal symmetries example: Michael-Scott queue

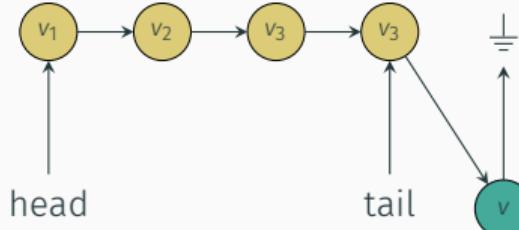
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T<sub>1</sub>

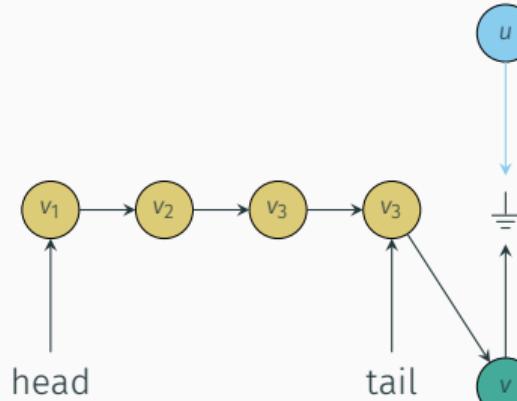
# Internal symmetries example: Michael-Scott queue

enqueue( $v$ )  $\triangleq$   
T2 →  $node := \text{malloc}(\dots)$   
 $node.value := v$   
 $node.next := \text{NULL}$   
do  
   $t := \underline{tail}$   
   $next := t.next$   
  **if** ( $t \neq \underline{tail}$ ) **continue**  
  **if** ( $next \neq \text{NULL}$ )  
    CAS( $\underline{tail}, t, next$ )  
    **continue**  
  **while** ( $\neg \text{CAS}(t.next, next, node)$ )  
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T1 →



# Internal symmetries example: Michael-Scott queue

enqueue( $v$ )  $\triangleq$   
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    **continue**  
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# Internal symmetries example: Michael-Scott queue

`enqueue(v)  $\triangleq$   
node := malloc(...)`

`node.value := v  
node.next := NULL`

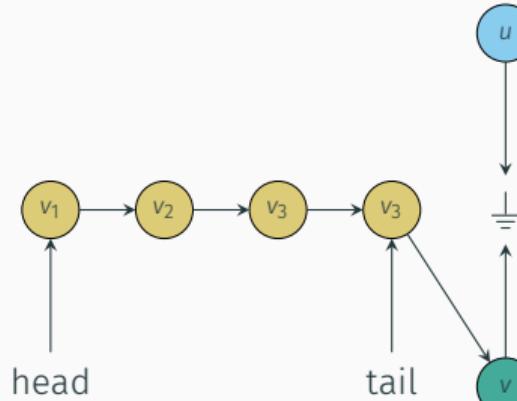
`do`

T<sub>2</sub>

`t := tail  
    next := t.next  
    if (t  $\neq$  tail) continue  
    if (next  $\neq$  NULL)  
        CAS(tail, t, next)  
    continue`

`while ( $\neg$ CAS(t.next, next, node))  
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T<sub>1</sub>



# Internal symmetries example: Michael-Scott queue

$\text{enqueue}(v) \triangleq$   
 $\text{node} := \text{malloc}(\dots)$

$\text{node.value} := v$

$\text{node.next} := \text{NULL}$

**do**

T<sub>2</sub>

$t := \underline{\text{tail}}$

$\text{next} := t.\text{next}$

**if** ( $t \neq \underline{\text{tail}}$ ) **continue**

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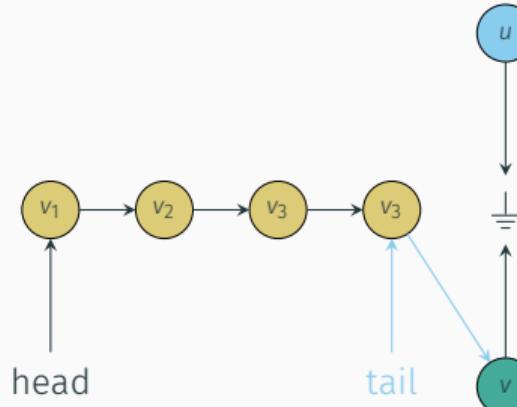
$\text{CAS}(\underline{\text{tail}}, t, \text{next})$

**continue**

**while** ( $\neg \text{CAS}(t.\text{next}, \text{next}, \text{node})$ )

$\text{CAS}(\underline{\text{tail}}, t, \text{node})$

T<sub>1</sub>



# Internal symmetries example: Michael-Scott queue

`enqueue(v)  $\triangleq$`

`node := malloc(...)`

`node.value := v`

`node.next := NULL`

`do`

`t := tail`

`next := t.next`

`if (t  $\neq$  tail) continue`

`if (next  $\neq$  NULL)`

`CAS(tail, t, next)`

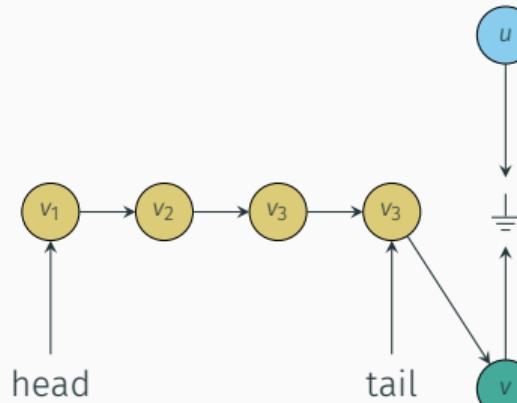
`continue`

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T<sub>2</sub>

T<sub>1</sub>



# Internal symmetries example: Michael-Scott queue

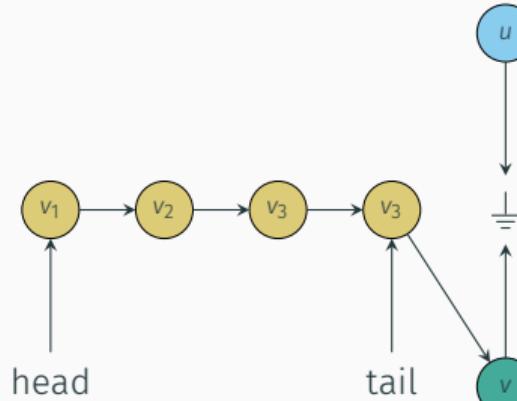
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      CAS(tail, t, node)
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T<sub>2</sub>

T<sub>1</sub>



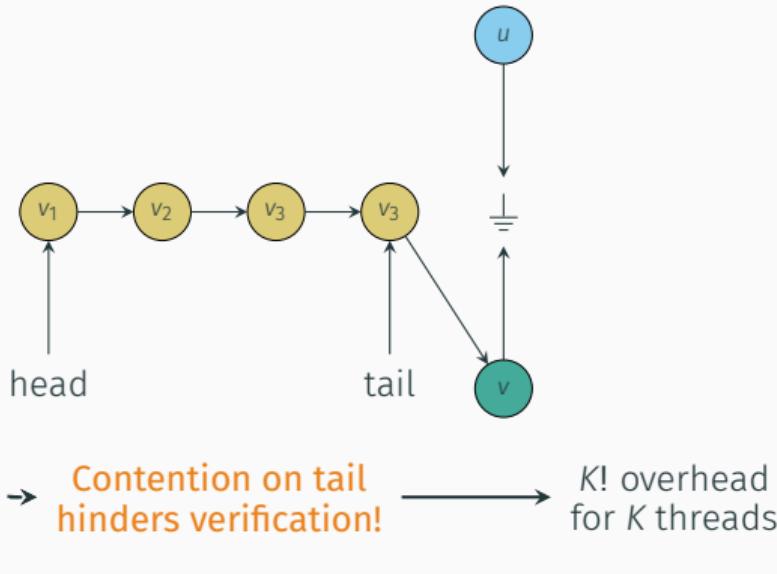
Contention on tail  
hinders verification!

# Internal symmetries example: Michael-Scott queue

```
enqueue(v) ≡  
node := malloc(...)  
node.value := v  
node.next := NULL  
do
```

```
    t := tail  
    next := t.next  
    if (t ≠ tail) continue  
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        CAS(tail, t, next)  
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```
T2 →  
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# Internal symmetries example: Michael-Scott queue

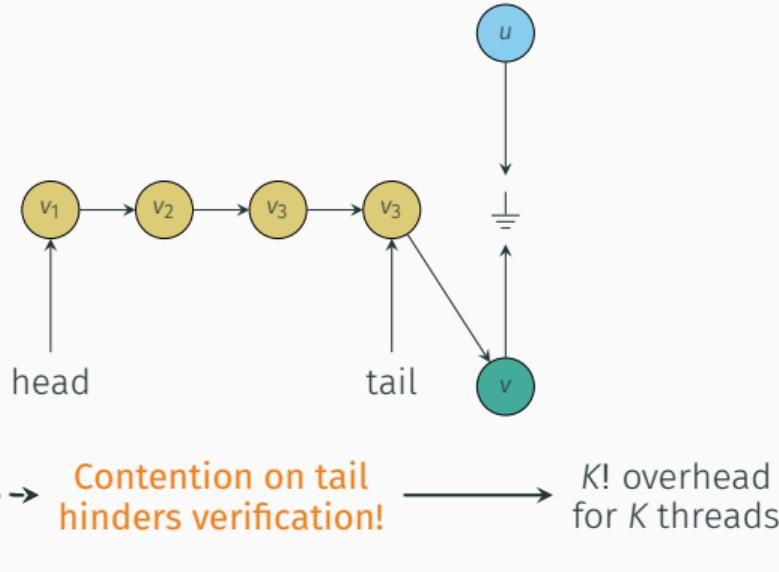
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```
while (¬CAS(t.next, next, node))  
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T2

T1



Observe: The conflicting operations are **identical** and **idempotent**

GENMC leverages idempotent operations by splitting them to `main` and `helping`  
~~ this requires **annotating** the input program

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**Key Idea:** only explore executions where `main` succeeds

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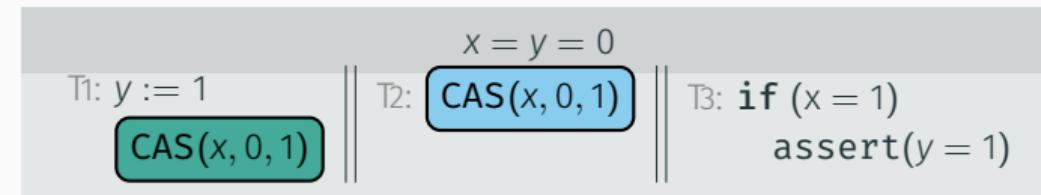
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Is this sound?

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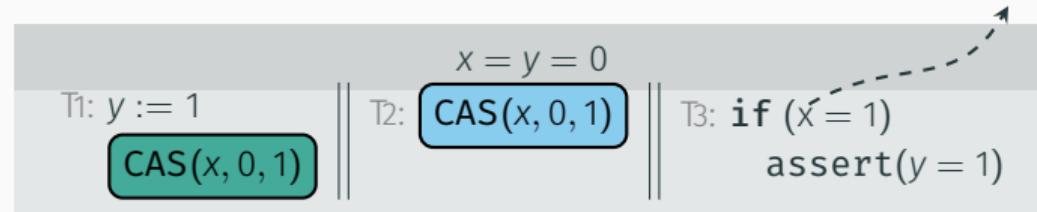


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Is this sound?

reading only from `main`  
misses the bug!

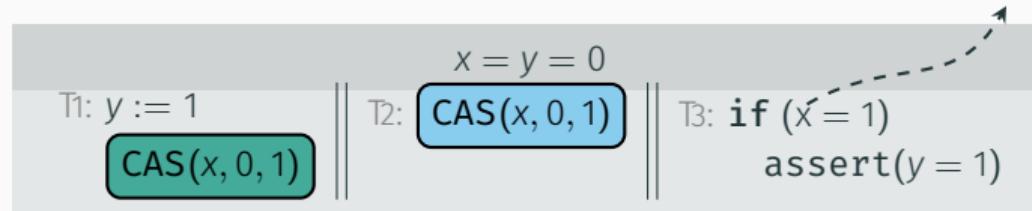


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**Key Idea:** only explore executions where `main` succeeds

reading only from `main`  
misses the bug!

Is this sound?



GENMC presents **sufficient conditions** for leveraging idempotent operations

~~ `main` and `helping` can be functions ...

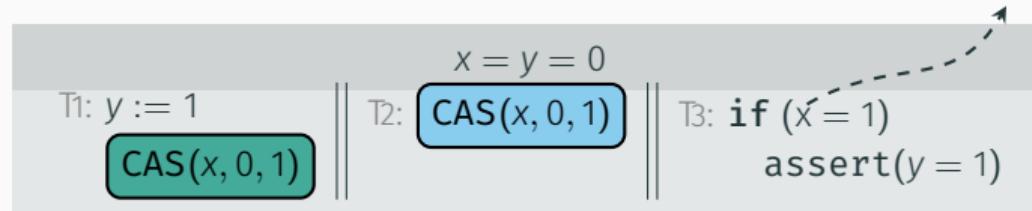
~~ but they have to satisfy certain conditions (e.g., induce same synchronization)

GENMC leverages idempotent operations by splitting them to `main` and `helping`  
~~ this requires **annotating** the input program

**Key Idea:** only explore executions where `main` succeeds

reading only from `main`  
misses the bug!

Is this sound?



GENMC presents **sufficient conditions** for leveraging idempotent operations

~~ `main` and `helping` can be functions ...

~~ but they have to satisfy certain conditions (e.g., induce same synchronization)

**Internal symmetries can also be leveraged in non-symmetric programs!**

# Outline

How does GENMC work?

- SMC basics
- Execution graphs
- Exploration algorithm

How to apply GENMC to our code?

- State-space reductions
- Estimating state-space size
- Exploration bounding

Each part will be followed by a demo

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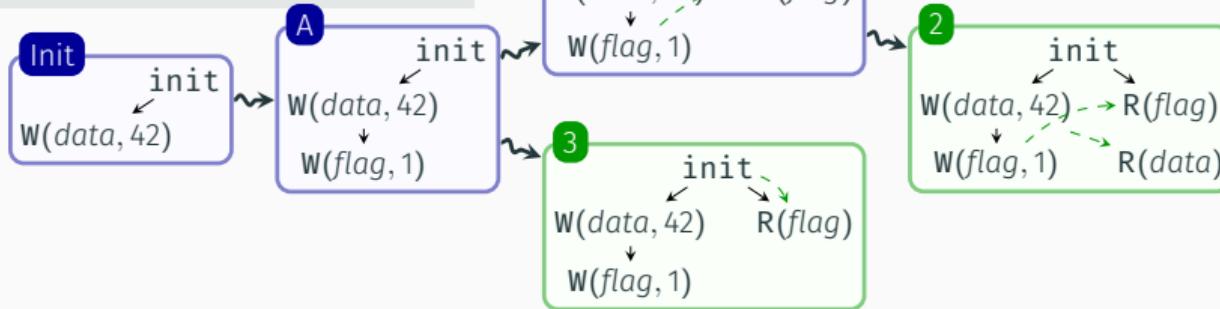
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# How to estimate the state-space size?

```
[data = flag = 0]
data := 42  ||  if (flag = 1)
flag := 1    ||  assert(data = 42)
```

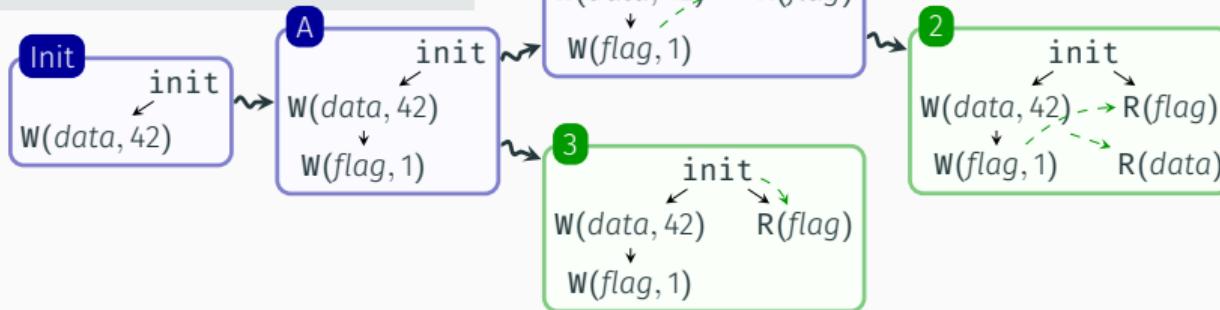
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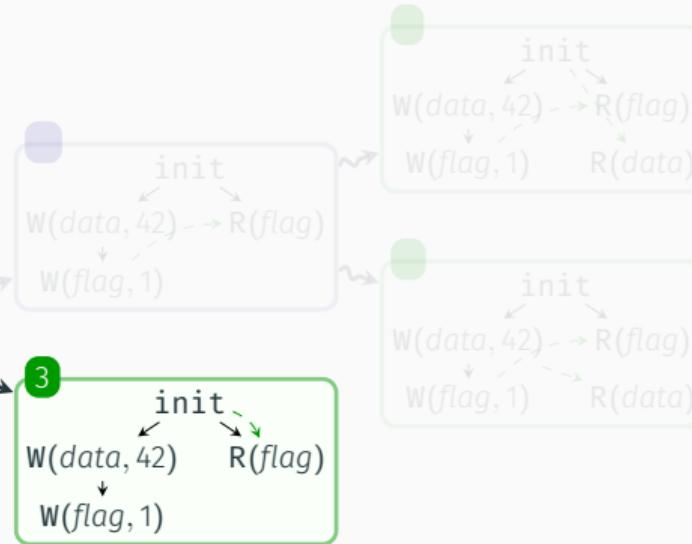
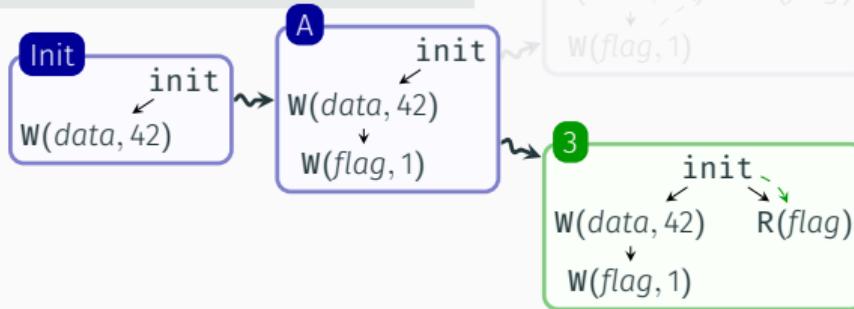


Naive “solution”:

- Assume symmetric state space
- Estimate based on explored space

# How to estimate the state-space size?

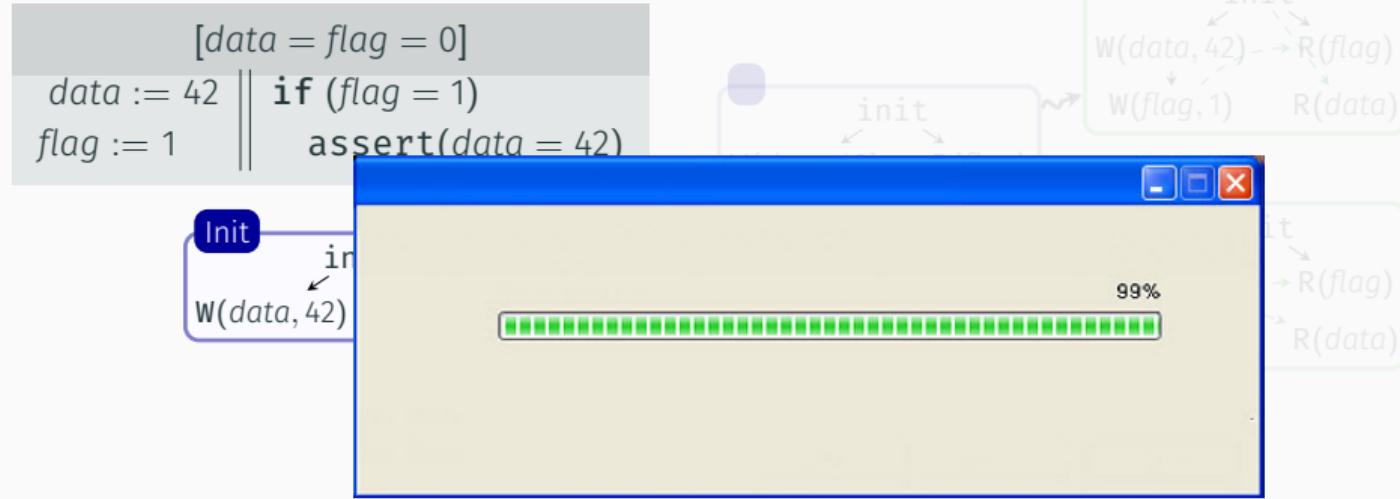
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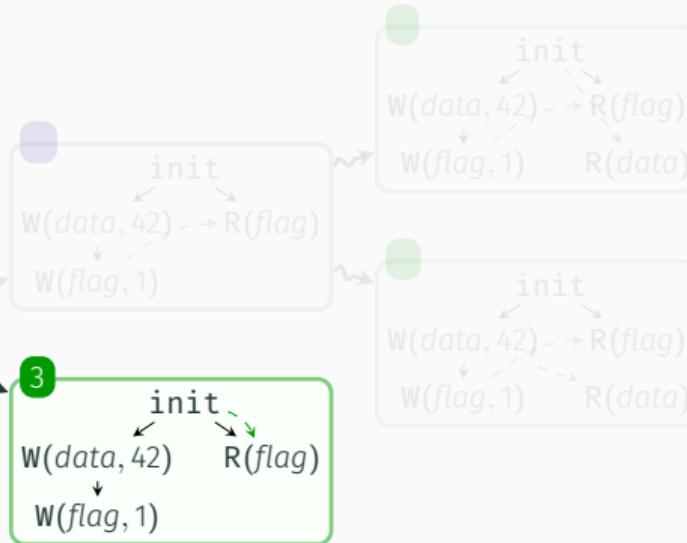
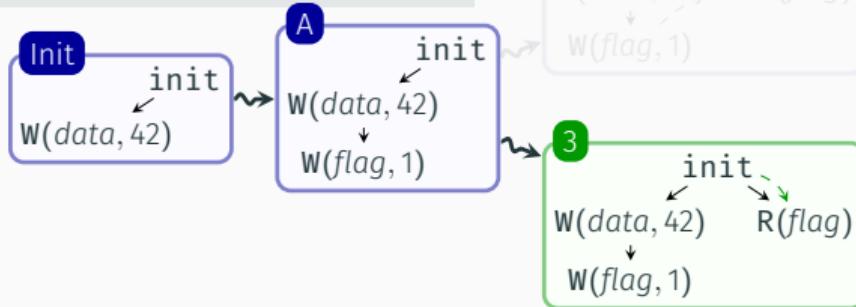


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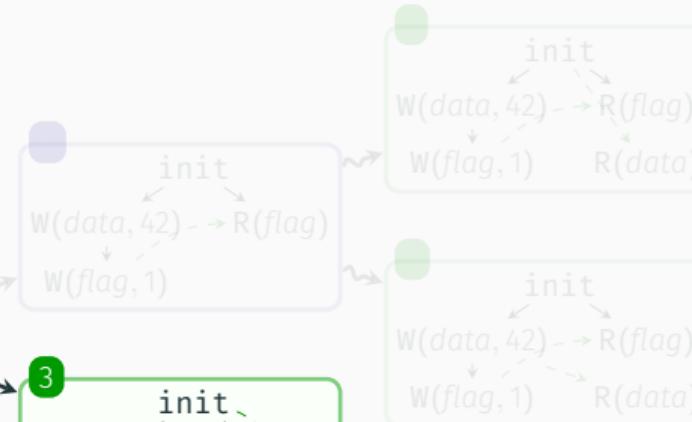
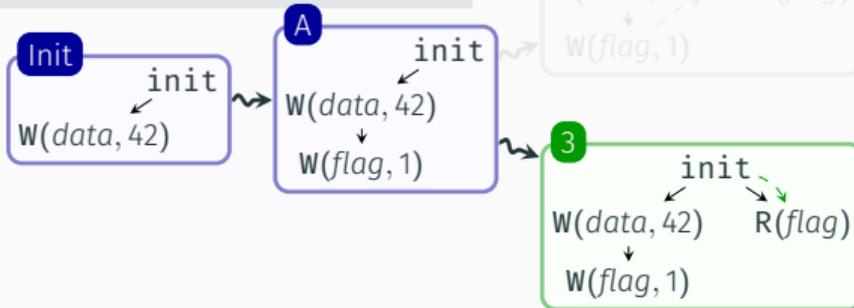
- Assume symmetric state space
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**Our Idea: Monte Carlo Simulation before verification**

- Take random samples (assuming symmetric space)
- Law of large numbers guarantees accuracy (if unbiased)

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**Our Idea:** Monte Carlo Simulation **before** verification

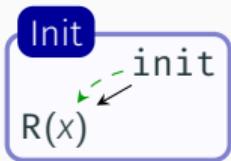
- Take random samples (assuming symmetric space)
- Law of large numbers guarantees accuracy (**if unbiased**)

# Reducing bias in estimation

```
[x = 0]  
a := x  
if (a > 0) b := x || x := 1 || x := 2
```

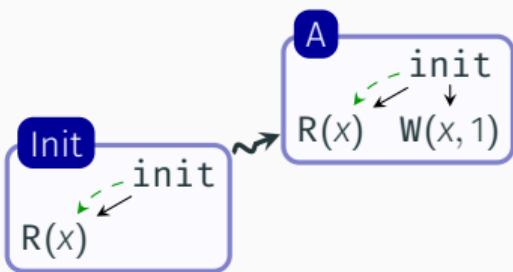
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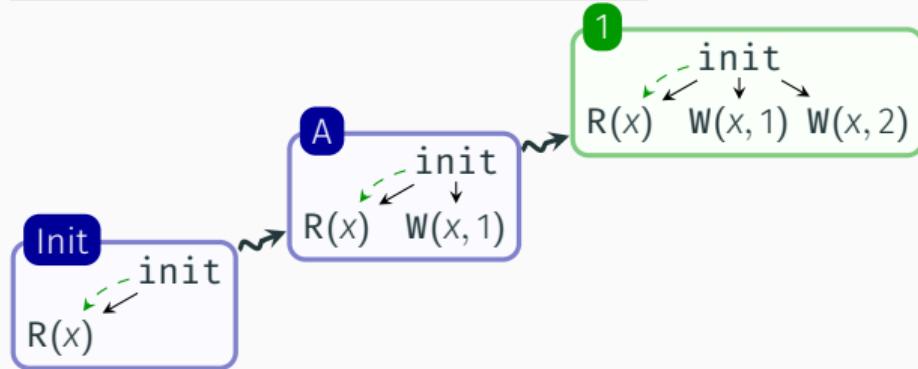
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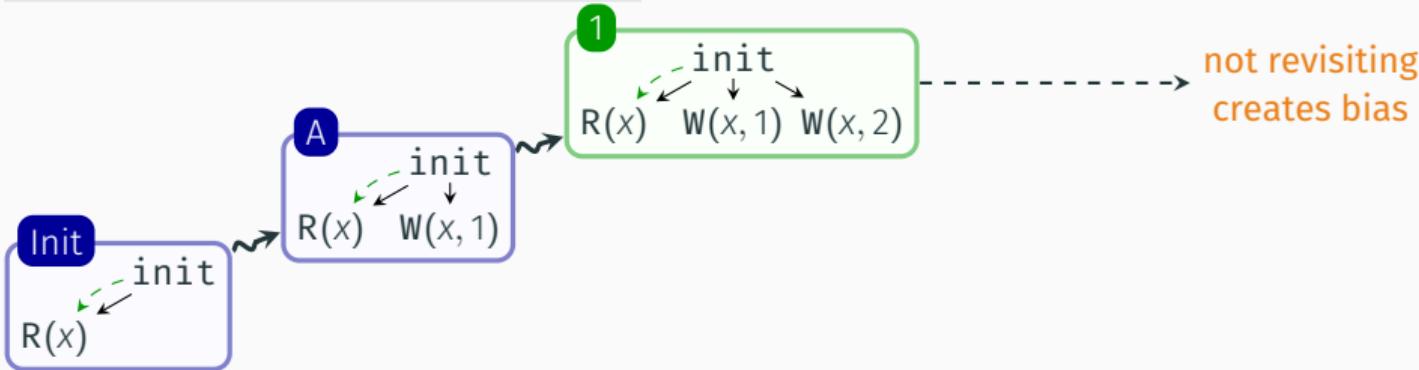
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if (a > 0) b := x ||| x := 1 ||| x := 2
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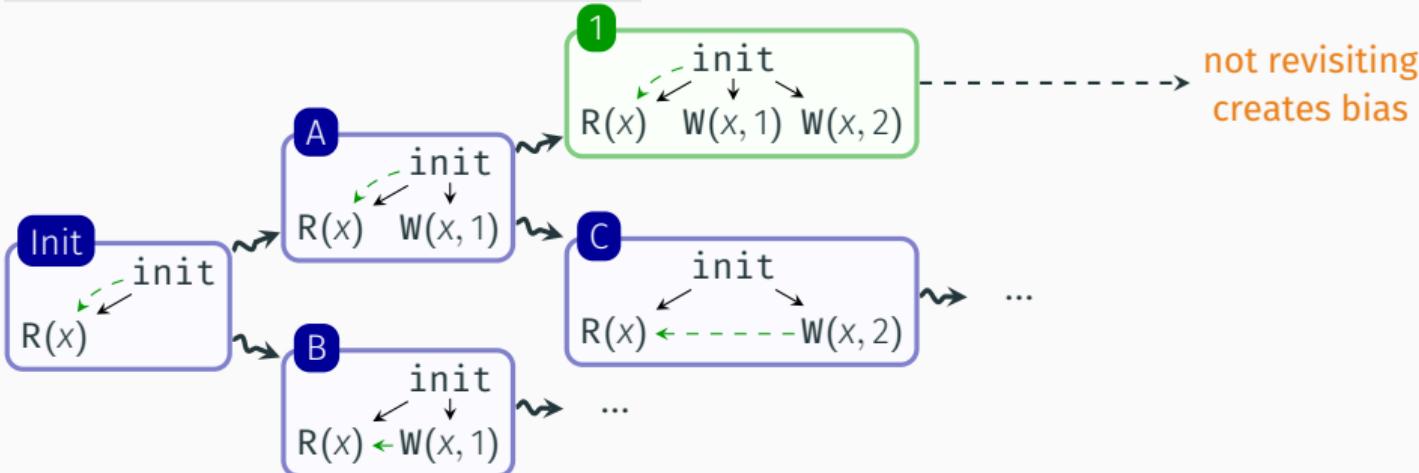
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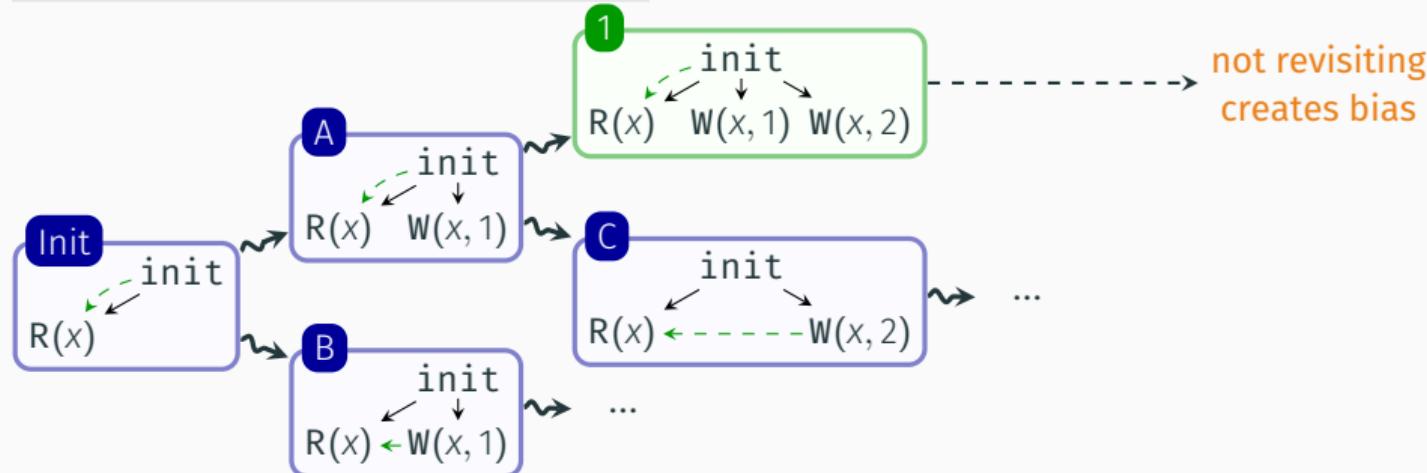
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# Reducing bias in estimation

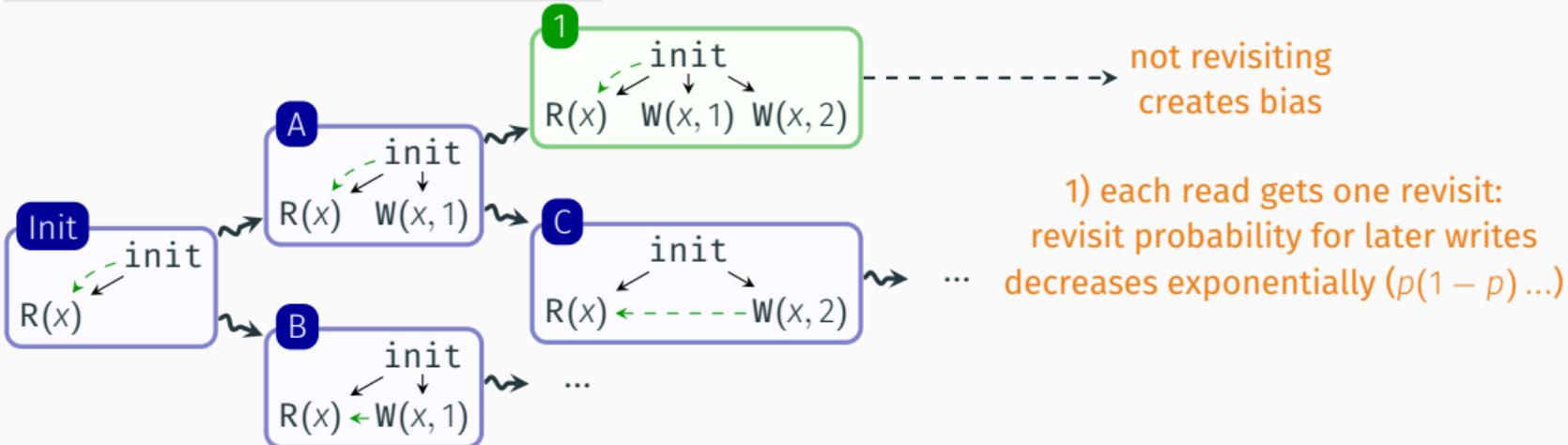
```
[x = 0]  
a := x  
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```



**Problem:** When to perform a revisit?

# Reducing bias in estimation

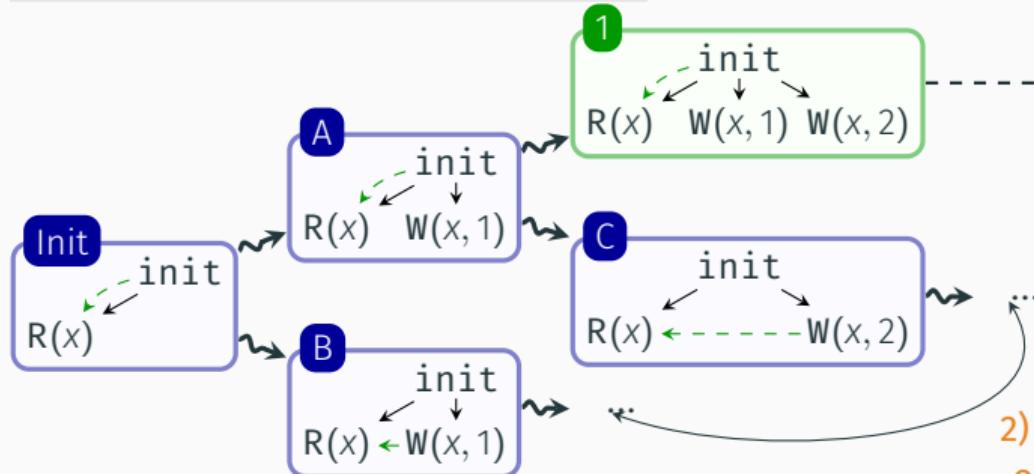
```
[x = 0]  
a := x  
if (a > 0) b := x ||| x := 1 ||| x := 2
```



**Problem:** When to perform a revisit?

# Reducing bias in estimation

```
[x = 0]  
a := x  
if (a > 0) b := x ||| x := 1 ||| x := 2
```



not revisiting creates bias

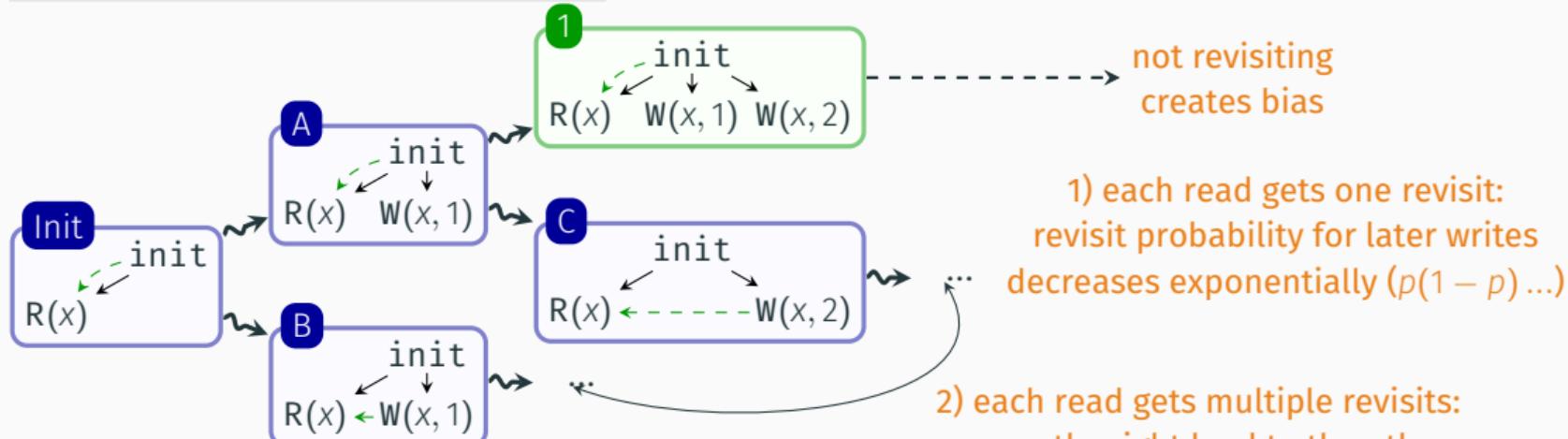
1) each read gets one revisit:  
revisit probability for later writes  
decreases exponentially ( $p(1 - p) \dots$ )

2) each read gets multiple revisits:  
one path might lead to the other  
~~ long estimation time

Problem: When to perform a revisit?

# Reducing bias in estimation

```
[x = 0]  
a := x  
if (a > 0) b := x ||| x := 1 ||| x := 2
```



**Problem:** When to perform a revisit?

**Our Solution:** No revisits — random scheduler that prioritizes writes over reads

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# Bounded exploration and POR

$[x = y = 0]$
$a := x \parallel y := 1$
$b := y \parallel y := 2$

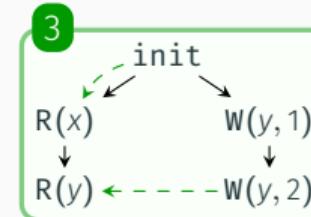
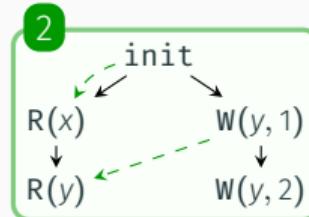
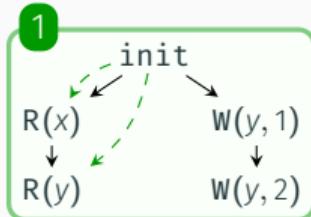
# Bounded exploration and POR

$$\begin{array}{c} [x = y = 0] \\ a := x \parallel y := 1 \\ b := y \parallel y := 2 \end{array}$$

**b = 0**  
 $a := x$   
↓  
 $b := y$   
↓  
 $y := 1$   
↓  
 $y := 2$

**b = 1**  
 $a := x$        $y := 1$   
↓                  ↓  
 $y := 1$        $a := x$   
↓                  ↓  
 $b := y$        $b := y$   
↓                  ↓  
 $y := 2$        $y := 2$

**b = 2**  
 $a := x$        $y := 1$        $y := 1$   
↓                  ↓                  ↓  
 $y := 1$        $a := x$        $y := 2$   
↓                  ↓                  ↓  
 $y := 2$        $y := 2$        $a := x$   
↓                  ↓                  ↓  
 $b := y$        $b := y$        $b := y$



# Bounded exploration and POR

$$\begin{array}{c} [x = y = 0] \\ a := x \parallel y := 1 \\ b := y \parallel y := 2 \end{array}$$

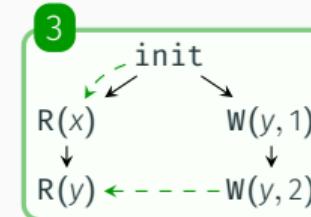
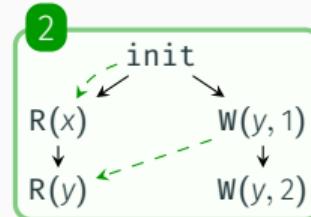
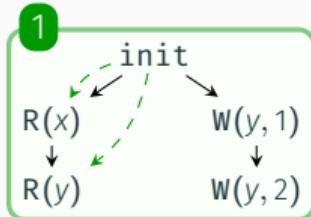
$b = 0$

$$\begin{array}{l} a := x \\ \downarrow \\ b := y \\ \downarrow \\ y := 1 \\ \downarrow \\ y := 2 \end{array}$$

$b = 1$

$$\begin{array}{ll} a := x & y := 1 \\ \downarrow & \downarrow \\ y := 1 & a := x \\ \downarrow & \downarrow \\ b := y & b := y \\ \downarrow & \downarrow \\ y := 2 & y := 2 \end{array}$$

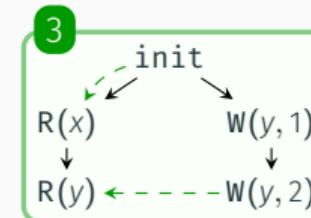
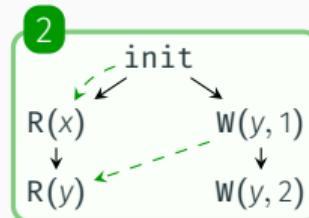
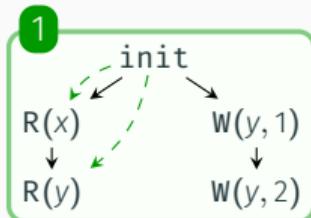
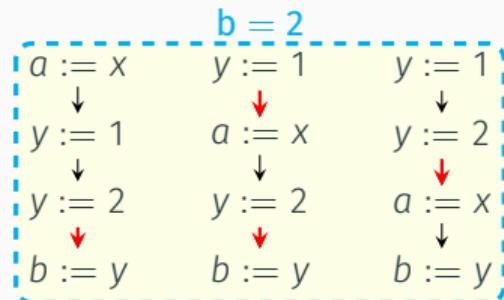
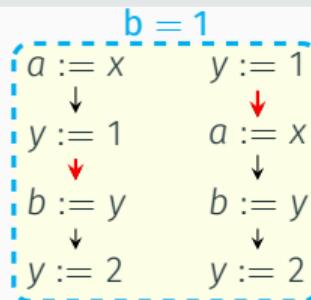
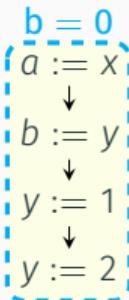
$b = 2$

$$\begin{array}{lll} a := x & y := 1 & y := 1 \\ \downarrow & \downarrow & \downarrow \\ y := 1 & a := x & y := 2 \\ \downarrow & \downarrow & \downarrow \\ y := 2 & y := 2 & a := x \\ \downarrow & \downarrow & \downarrow \\ b := y & b := y & b := y \end{array}$$


Goal: Only explore graphs  $G \in \llbracket P \rrbracket$  where exists  $t \in \text{trace}(G)$  s.t.  $B(t) \leq K$

# Bounded exploration and POR

$$\begin{array}{c} [x = y = 0] \\ a := x \parallel y := 1 \\ b := y \parallel y := 2 \end{array}$$



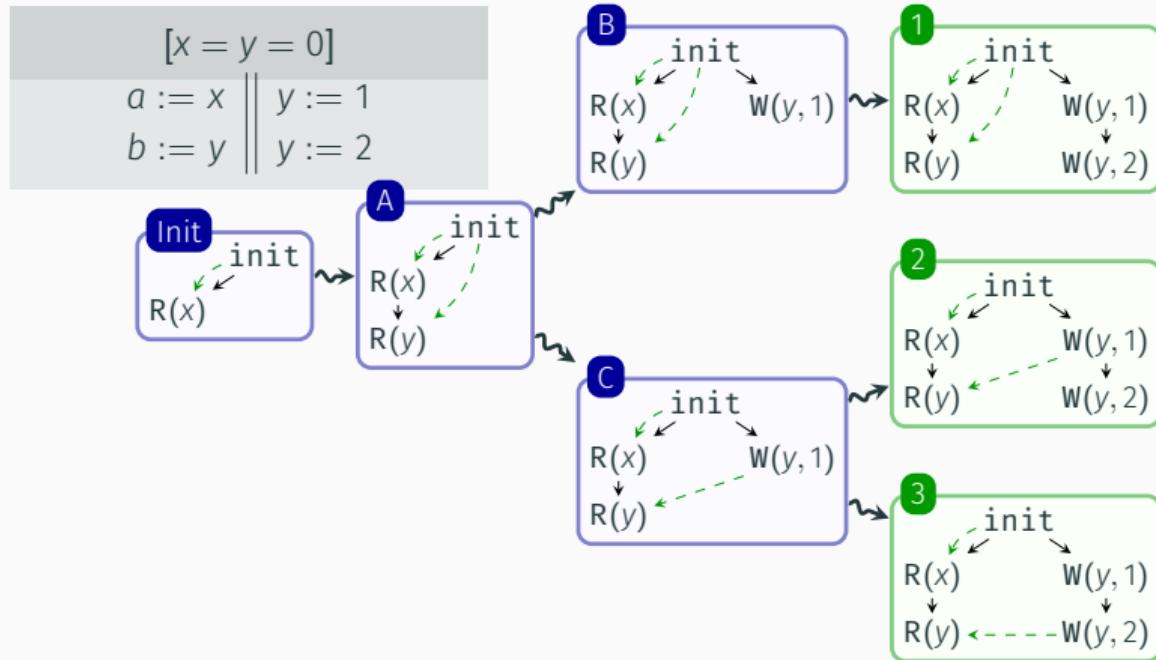
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here: round-robin rounds

# Bounded POR

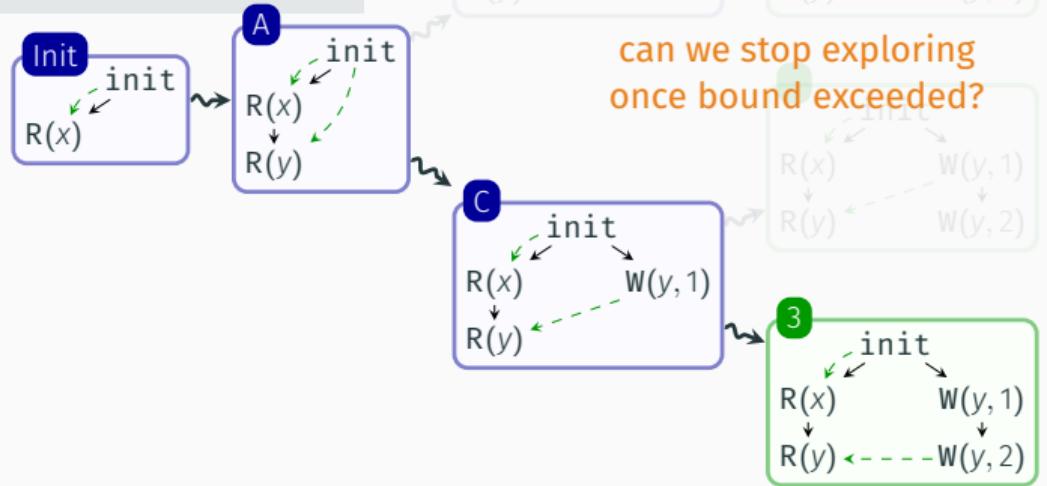
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# Bounded POR



# Bounded POR

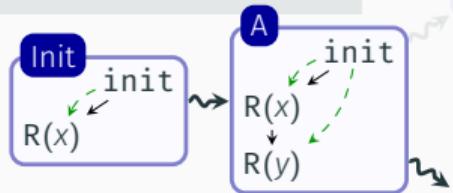
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# Bounded POR

$[x = y = 0]$

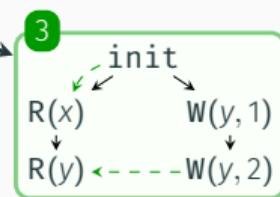
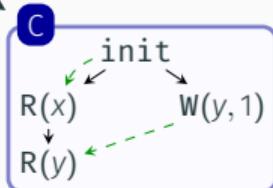
$$a := x \parallel y := 1$$

$$b := y \parallel y := 2$$


can we stop exploring  
once bound exceeded?

$a := x \quad y := 1$   
 $\downarrow \quad \downarrow$   
 $y := 1 \quad a := x$   
 $\downarrow \quad \downarrow$   
 $b := y \quad b := y$

$\min(B) = 1$

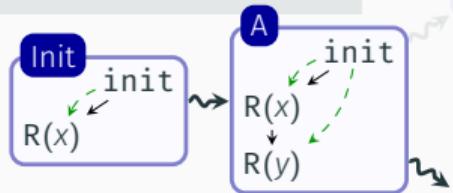


$a := x \quad y := 1 \quad y := 1$   
 $\downarrow \quad \downarrow \quad \downarrow$   
 $y := 1 \quad a := x \quad y := 2$   
 $\downarrow \quad \downarrow \quad \downarrow$   
 $y := 2 \quad y := 2 \quad a := x$   
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# Bounded POR

$$[x = y = 0] \\ \begin{array}{c|c} a := x & y := 1 \\ b := y & y := 2 \end{array}$$

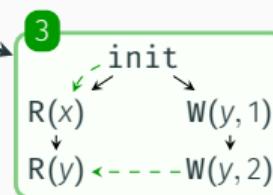
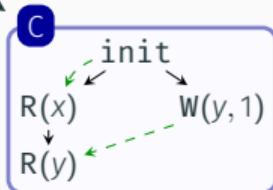


$$\begin{array}{ll} a := x & y := 1 \\ \downarrow & \downarrow \\ y := 1 & a := x \\ \downarrow & \downarrow \\ b := y & b := y \end{array}$$

$\min(B) = 1$



can we stop exploring  
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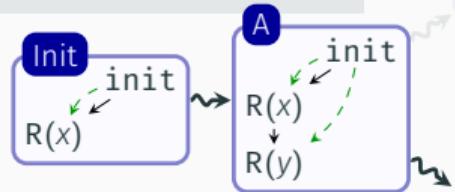
Yes, if bound is **monotone** across exploration (e.g., **round-robin**)

$$\begin{array}{llll} a := x & y := 1 & y := 1 & \\ \downarrow & \downarrow & \downarrow & \\ y := 1 & a := x & y := 2 & \\ \downarrow & \downarrow & \downarrow & \\ y := 2 & y := 2 & a := x & \\ \downarrow & \downarrow & \downarrow & \\ b := y & b := y & b := y & \end{array}$$

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# Bounded POR

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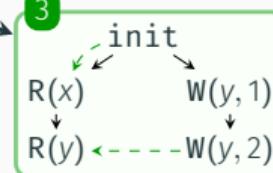
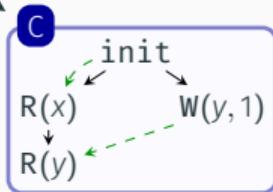


$$\begin{array}{ll} a := x & y := 1 \\ \downarrow & \downarrow \\ y := 1 & a := x \\ \downarrow & \downarrow \\ b := y & b := y \end{array}$$

$\min(B) = 1$



can we stop exploring  
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$$\begin{array}{llll} a := x & y := 1 & y := 1 & y := 1 \\ \downarrow & \downarrow & \downarrow & \downarrow \\ y := 1 & a := x & y := 2 & a := x \\ \downarrow & \downarrow & \downarrow & \downarrow \\ y := 2 & y := 2 & a := x & a := x \\ \downarrow & \downarrow & \downarrow & \downarrow \\ b := y & b := y & b := y & b := y \end{array}$$

$\min(B) = 0$

Yes, if bound is **monotone** across exploration (e.g., **round-robin**)

~~**context bounding**: intermediate graphs might have bound  $\leq T + K - 2$

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