

# Efficient Answer Set Counting with `aspmc`

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funded by FWF project W1255-N23

20<sup>th</sup> of September 2021

**FWF**

Der Wissenschaftsfonds.

**logics**  LOGICAL METHODS IN  
COMPUTER SCIENCE

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- ▶ Preferential Reasoning [Brewka *et al.*, 2015]
- ▶ Algebraic Answer Set Counting (AASC) [Eiter and Kiesel, 2020], [Kimmig *et al.*, 2011]

## Solving AASC I

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    - ↔ only feasible for “few” answer sets
  2. dynamic programming on a tree decomposition
    - ↔ only feasible for very low treewidth
  3. compilation into a *tractable circuit representation* like d-DNNF or SDD
    - ↔ compilers like c2d [Darwiche, 2004] work on CNFs

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- ▶ We favor compilation  
↔ translate ASP to CNF via *cycle-breaking*
- ▶ Compilation has performance guarantees based on *trewidth*  
↔ find *trewidth-aware* cycle-breaking

## Example Program

Non-ground smokers program

0.3 :: stress( $X$ )  $\leftarrow$  person( $X$ )

smokes( $X$ )  $\leftarrow$  stress( $X$ )

0.2 :: inf( $X$ ,  $Y$ )  $\leftarrow$  friend( $X$ ,  $Y$ )

smokes( $Y$ )  $\leftarrow$  smokes( $X$ ), inf( $X$ ,  $Y$ )

## Example Program

Non-ground smokers program

```

0.3 :: stress(X) ← person(X)
      smokes(X) ← stress(X)
0.2 :: inf(X, Y) ← friend(X, Y)
      smokes(Y) ← smokes(X), inf(X, Y)
    
```

And input data

|              |              |              |
|--------------|--------------|--------------|
| person(1)    | person(2)    | person(3)    |
| friend(1, 2) | friend(2, 3) | friend(3, 1) |

## Example cont.

Putting the input data and the non-ground program together results in

$$\begin{array}{ll}
 0.3 :: \text{stress}(x) \leftarrow & \text{for } x = 1, 2, 3 \\
 \text{smokes}(x) \leftarrow \text{stress}(x) & \text{for } x = 1, 2, 3 \\
 0.2 :: \text{inf}(x, y) \leftarrow & \text{for } x + 1 \equiv y \pmod{3} \\
 \text{smokes}(y) \leftarrow \text{smokes}(x), \text{inf}(x, y) & \text{for } x + 1 \equiv y \pmod{3}
 \end{array}$$

## (Positive) Dependency Graph

The (positive) *dependency graph* of a program  $\Pi$  is the digraph  $\text{DEP}(\Pi) = (V, E)$ , where

- ▶  $V = \mathcal{A}(\Pi)$  is the set of propositional variables that occur in  $\Pi$
- ▶  $(b, a) \in E$  if there is a rule  $a \leftarrow b_1, \dots, b_n, \text{not } c_1, \dots, \text{not } c_m$  in  $\Pi$  with  $b = b_i$

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### Theorem ([Fages, 1994])

If  $\text{DEP}(\Pi)$  is acyclic, then  $\text{Clark}(\Pi)$ , the Clark-completion of  $\Pi$ , is a propositional formula whose models are the answer sets of  $\Pi$ .



## Example cont.

The ground program...

|   |                               |
|---|-------------------------------|
| 0.3 :: stress( $x$ ) $\leftarrow$                       | for $x = 1, 2, 3$             |
| smokes( $x$ ) $\leftarrow$ stress( $x$ )                | for $x = 1, 2, 3$             |
| 0.2 :: inf( $x, y$ ) $\leftarrow$                       | for $x + 1 \equiv y \pmod{3}$ |
| smokes( $y$ ) $\leftarrow$ smokes( $x$ ), inf( $x, y$ ) | for $x + 1 \equiv y \pmod{3}$ |

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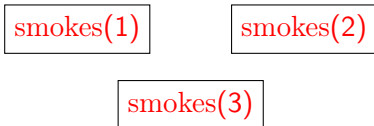
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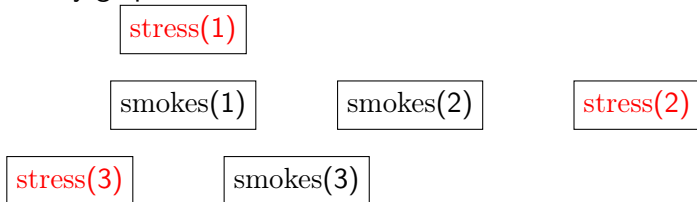


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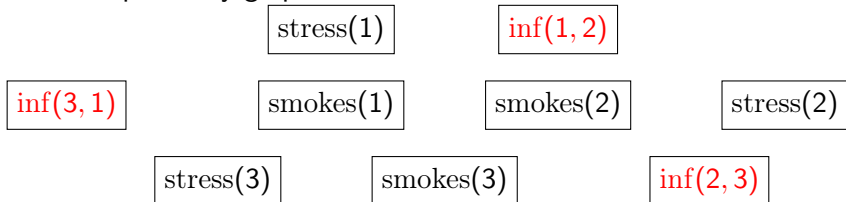
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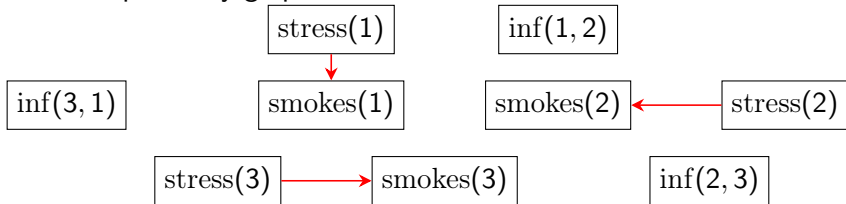
0.3 :: stress(x) ← for x = 1, 2, 3

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... has dependency graph

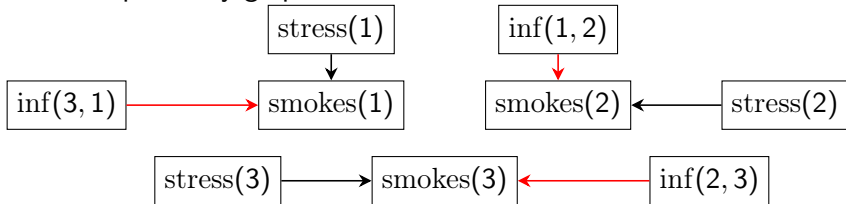


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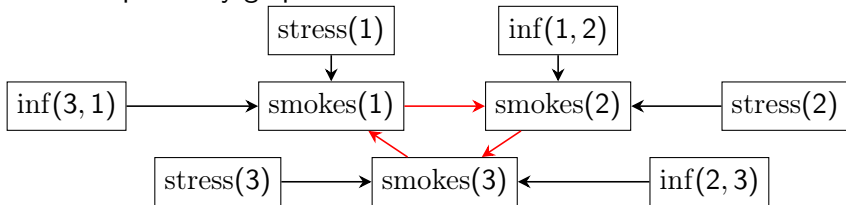
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  - ▶ only use derivations with probabilistic facts or already introduced copies
- ▶ Do this iteratively until a (semantic) fixed point is reached
- ▶ **Insight:** the order in which atoms are considered is crucial!

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$$0.3 :: \text{stress}(x) \leftarrow \text{for } x = 1, 2, 3 \quad (1)$$

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$$0.2 :: \text{inf}(x, y) \leftarrow \text{for } x + 1 \equiv y \pmod{3} \quad (2)$$

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$$\text{smokes}(1)^1 \leftarrow \text{stress}(1) \quad \text{smokes}(1)^1 \leftarrow \text{inf}(3, 1), \perp$$

no copy of  $\text{smokes}(3) \leftrightarrow$  use falsum

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---

**Algorithm 1**  $T_{\mathcal{P}}$ -Unfold( $\Pi, s$ )

---

**Input** A program  $\Pi$  and an unfolding sequence  $s \in \mathcal{A}(\Pi)^*$ .

**Output** An acyclic program  $\Pi'$ .

```
1: last = {a ↦ ⊥ | a ∈  $\mathcal{A}(\Pi)$ }
2: cnt = {a ↦ 0 | a ∈  $\mathcal{A}(\Pi)$ }
3: for  $i = 1, \dots, \text{len}(s)$  do
4:   if isLastOccurrence( $s_i, i, s$ ) then
5:     head =  $s_i$ 
6:   else
7:     head =  $s_i^{\text{cnt}(s_i)+1}$ 
8:   for  $s_i \leftarrow b_1, \dots, b_n, \text{not } c_1, \dots, \text{not } c_m \in \Pi$  do
9:     Add head  $\leftarrow \text{last}(b_1), \dots, \text{last}(b_n), \text{not } c_1, \dots, \text{not } c_m$ 
10:  last( $s_i$ ) = head
11:  cnt( $s_i$ ) = cnt( $s_i$ ) + 1
```

---

---

## Algorithm 2 $T_{\mathcal{P}}$ -Unfold( $\Pi, s$ )

---

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1: last = {a ↦ ⊥ | a ∈ A(Π)}
2: cnt = {a ↦ 0 | a ∈ A(Π)}
3: for i = 1, ..., len(s) do
4:   if isLastOccurrence(si, i, s) then
5:     head = si
6:   else
7:     head = sicnt(si)+1
8:   for si ← b1, ..., bn, not c1, ..., not cm ∈ Π do
9:     Add head ← last(b1), ..., last(bn), not c1, ..., not cm
10:  last(si) = head
11:  cnt(si) = cnt(si) + 1

```

---

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### Theorem

✓, if for every simple directed path  $\pi = (a_1, \dots, a_n)$  in  $\text{DEP}(\Pi)$  there is a directed path  $\pi_c = (a_1^{c_1}, \dots, a_n^{c_n})$  in  $\text{DEP}(T_{\mathcal{P}}\text{-Unfold}(\Pi, s))$ .

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### Lemma

*Let  $\Pi$  be an answer set program with treewidth  $k$  and  $s \in \mathcal{A}(\Pi)^*$  be an unfolding sequence. If every variable occurs at most  $m$  times in  $s$ , then the treewidth of  $\text{T}_{\mathcal{P}}\text{-Unfold}(\Pi, s)$  is less or equal to  $k \cdot m$ .*

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- ▶ We are interested in path preserving  $m$ -unfolding sequences with small  $m$

## Component-Boosted Backdoor Size

### Definition ( $\text{cbs}(G)$ )

Let  $G$  be a digraph. Then  $\text{cbs}(G)$ , the component-boosted backdoor size of  $G$ , is

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- ▶  $\max\{\text{cbs}(C) \mid C \in \text{SCC}(G)\}$ , if  $G$  is cyclic but not strongly connected

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Intuitively,  $\text{cbs}(G)$  measures the *cyclicity* of  $G$  by decomposition into “easy to solve” subgraphs

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*For every digraph  $G$  there exists a path preserving  $\text{cbs}(G)$ -unfolding sequence.*

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### Theorem

*For every digraph  $G$  there exists a path preserving  $\text{cbs}(G)$ -unfolding sequence.*

For the original point of interest this means that:

### Theorem

*For every answer set program  $\Pi$ , there exists an unfolding sequence  $s \in \mathcal{A}(\Pi)^*$  such that*

- 1. the answer sets are preserved bijectively*
- 2. the treewidth of  $\mathbb{T}_{\mathcal{P}\text{-Unfold}}(\Pi, s)$  is less or equal to  $k \cdot \text{cbs}(\text{DEP}(\Pi))$ , where  $k$  is the treewidth of  $\Pi$ .*

## Scenarios

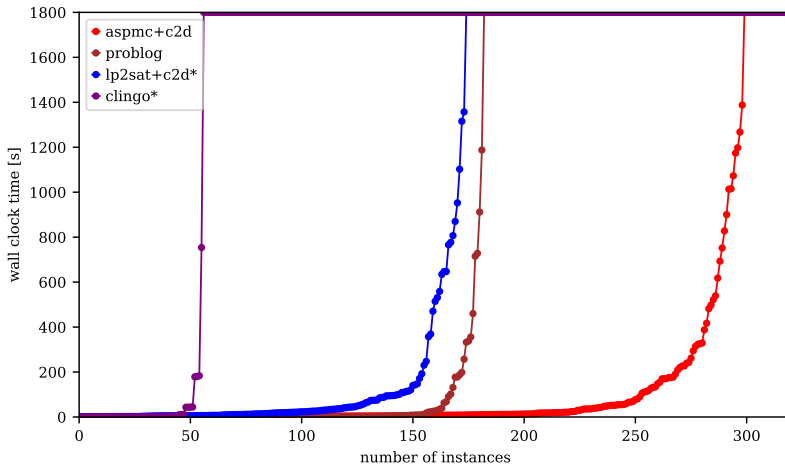
- S1 Probabilistic reasoning: Computing probabilities for atoms of Problog programs
- S2 Counting (small number of solutions on average): Counting the number of different paths between stations in public transport networks
- S3 Counting (many solutions on average): Counting conflict-free extensions in abstract argumentation

## Solvers

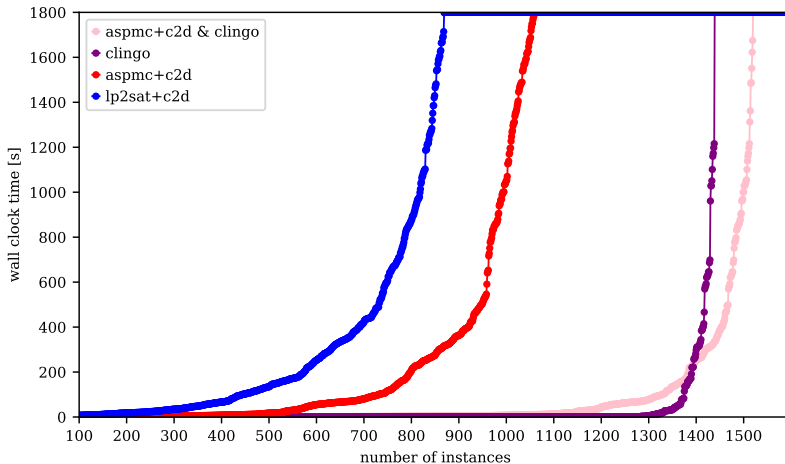
- ▶ Problog, version 2.1.0.42, run with arguments “-k sdd”
- ▶ clingo, version 5.4.0, run with arguments “-q -n 0”
- ▶ lp2sat+c2d: cycle breaking due to [Bomanson, 2017] followed by compilation using c2d [Darwiche, 2004]
- ▶ aspmc+c2d: our cycle breaking followed by compilation using c2d [Darwiche, 2004]



## Results S1



## Results S2



## Results S3

| solver<br>configuration | $\Sigma$   | tw ranges  |           |           | unique    | time[h]      |
|-------------------------|------------|------------|-----------|-----------|-----------|--------------|
|                         |            | 0-300      | 300-600   | >600      |           |              |
| aspmc+c2d               | <b>241</b> | <b>185</b> | <b>26</b> | <b>30</b> | <b>12</b> | <b>45.16</b> |
| lp2sat+c2d              | 182        | 182        | 0         | 0         | 0         | 73.85        |
| clingo                  | 144        | 97         | 21        | 26        | 2         | 94.78        |

## Conclusions





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- ▶  $T_{\mathcal{P}}$ -unfolding allows treewidth-aware cycle-breaking
- ▶ Our prototypical implementation partially outperforms other solvers

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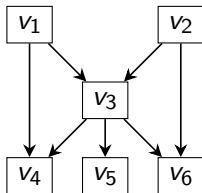
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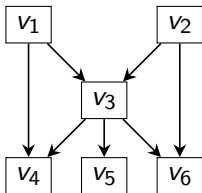
## Case 1: DAG

- ▶ We can take any unfolding sequence  $s$  that is in topological order



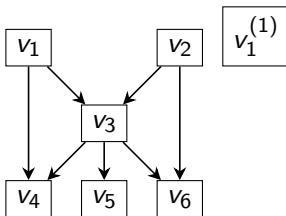
## Case 1: DAG

- ▶ We can take any unfolding sequence  $s$  that is in topological order
- ▶ Such an  $s$  can be constructed by iteratively removing a vertex  $a$  without ancestors from  $G$  and appending it to  $s$



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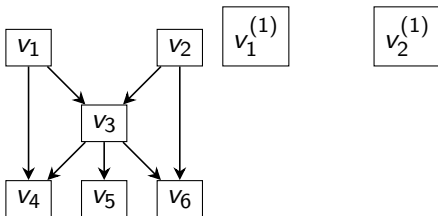
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Use unfolding sequence  $s = v_1$  .

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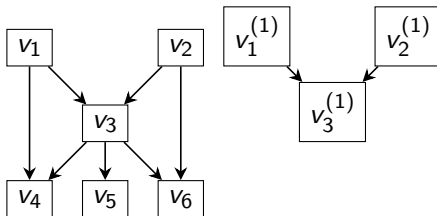
- ▶ We can take any unfolding sequence  $s$  that is in topological order
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Use unfolding sequence  $s = v_1 v_2$  .

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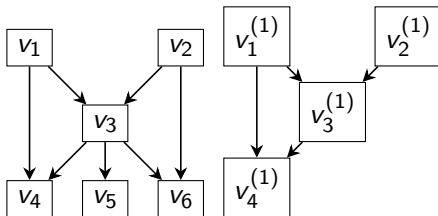
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Use unfolding sequence  $s = v_1 v_2 v_3$  .

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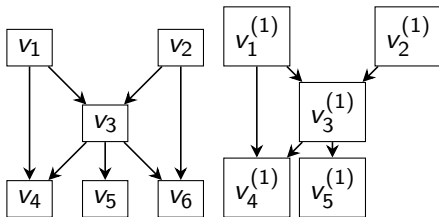
- ▶ We can take any unfolding sequence  $s$  that is in topological order
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Use unfolding sequence  $s = v_1 v_2 v_3 v_4$  .

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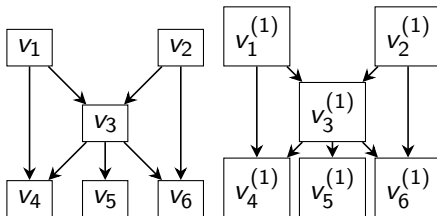


Use unfolding sequence  $s = v_1 v_2 v_3 v_4 v_5$  .



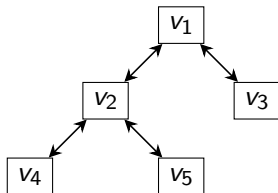
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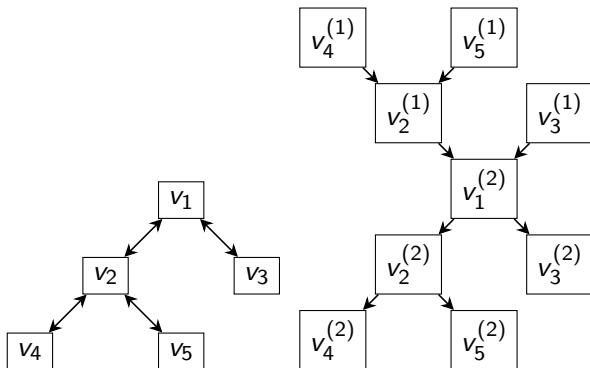
Use unfolding sequence  $s = v_1 v_2 v_3 v_4 v_5 v_6$ .

## Case 2: Polytree



Use unfolding sequence  $s = s_{post}s_{pre}$ , where  
 $s_{post} = v_4v_5v_2v_3v_1$ ,  $s_{pre} = v_3v_2v_5v_4$ .

## Case 2: Polytree

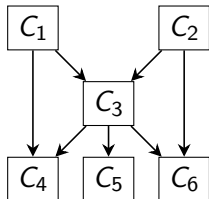


Use unfolding sequence  $s = s_{post}s_{pre}$ , where

$s_{post} = v_4v_5v_2v_3v_1, s_{pre} = v_3v_2v_5v_4$ .

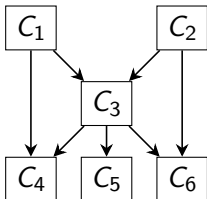
## Case 3: Cyclic but not Strongly Connected

- ▶ We can handle each strongly connected component  $C_i, i = 1, \dots, k$  of  $G$  separately and combine them



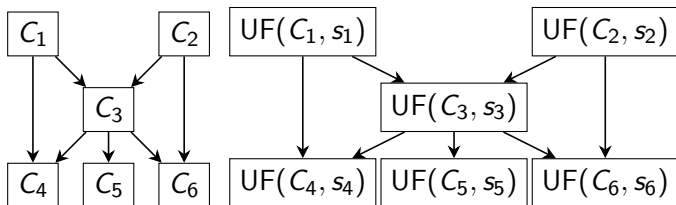
## Case 3: Cyclic but not Strongly Connected

- ▶ We can handle each strongly connected component  $C_i, i = 1, \dots, k$  of  $G$  separately and combine them
- ▶ Take  $s = s_{i_1} \dots s_{i_k}$  (in topological order of the SCCs), where  $s_{i_j}$  is a path-preserving  $\text{cbs}(C_{i_j})$ -unfolding sequence for  $C_{i_j}$



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- ▶  $G$  is strongly connected but not a polytree

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