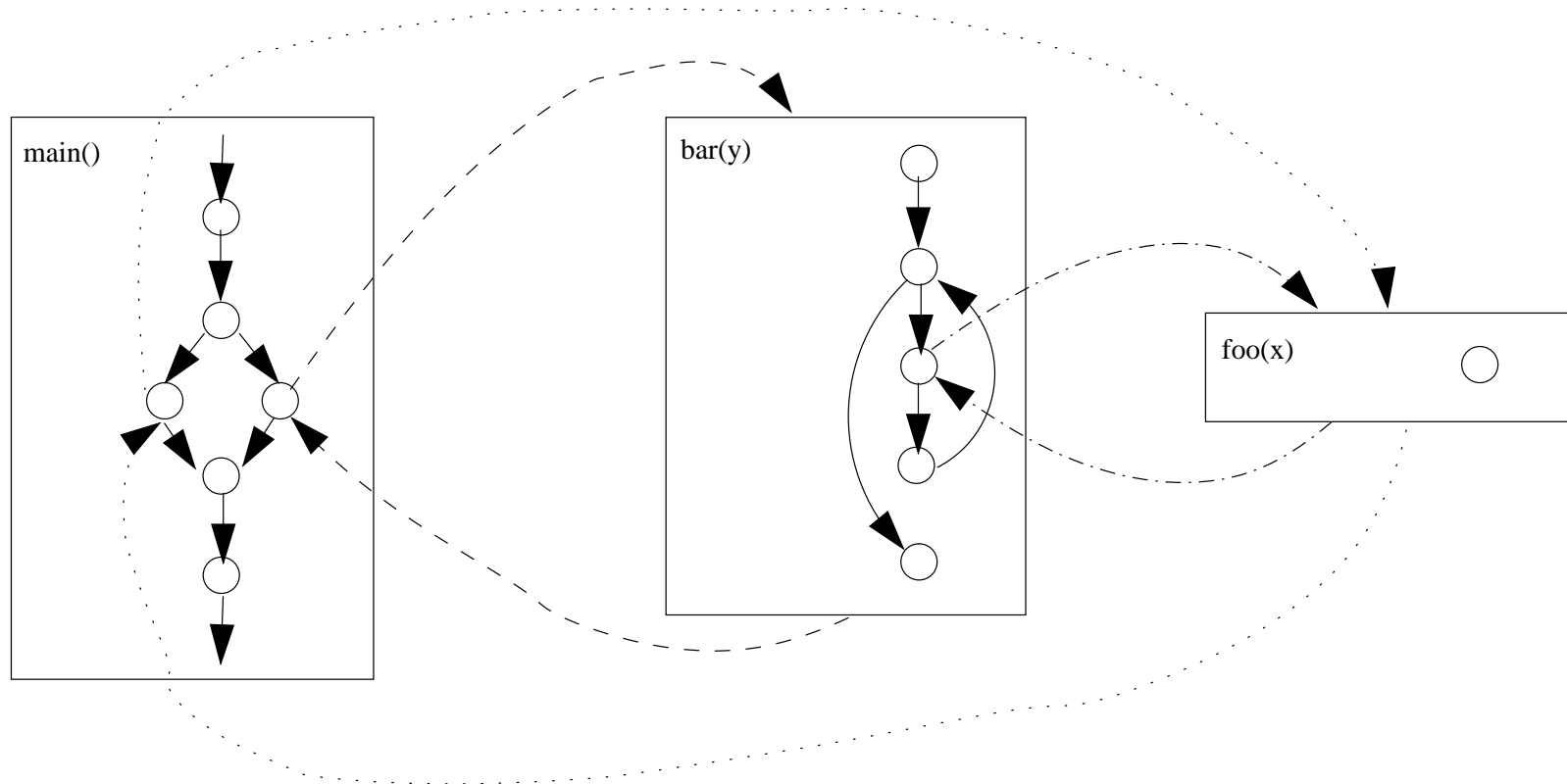


```
main() {  
    ...  
    if (x > 0) {  
        x = foo(x);  
    } else {  
        x = bar(x);  
    }  
    ...  
}  
  
int bar(int y) {  
    int t;  
    while (y > 0) {  
        t = t + foo(y);  
        t++;  
    }  
    return t;  
}  
  
int foo(int x) {  
    return x + x;  
}
```

Context-sensitive approaches attempt to match up call sites with called methods and distinguish values flowing from different call sites.

Interprocedural Analysis



Call Graph

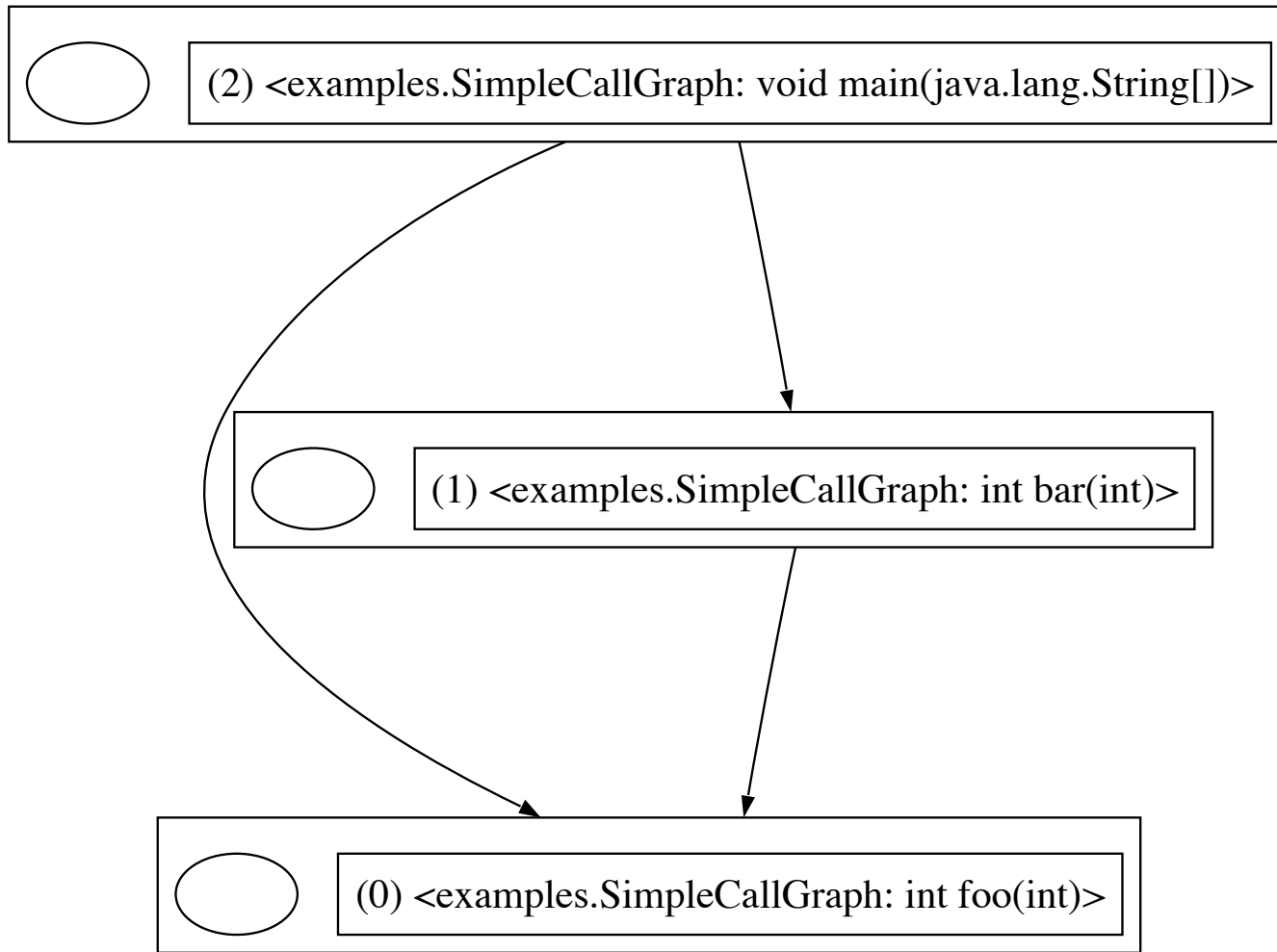
A *call graph* is an abstraction of the possible calling relationships among program methods

Just like a control flow graph a call graph *overestimates* the actual program behavior

It also leaves out a lot of detail

- no information about the call sites in a method
- no information about the number of calls in a method
- no information about the order of calls in a method

Interprocedural Analysis



SummaryCallGraph

Call Graph : Recursion

```
void main() {  
    arec(10);  
    rec(7);  
}
```

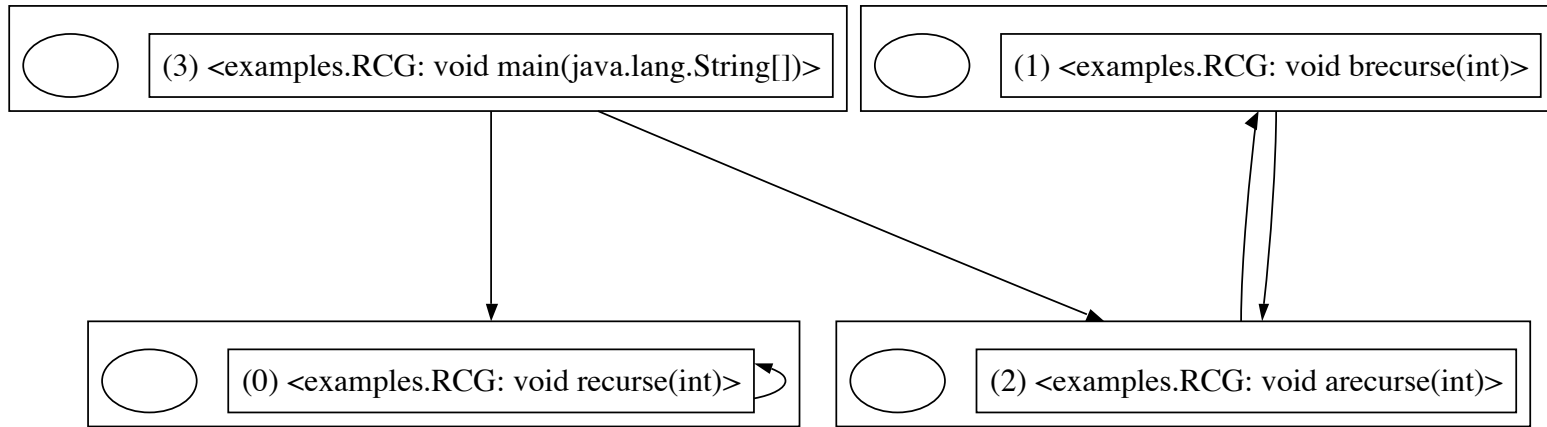
```
void rec(int x) {  
    if (x > 0) rec(x--);  
}
```

```
void arec(int x) {  
    if (x > 0) brec(x--);  
}
```

```
void brec(int x) {  
    if (x > 0) arec(x--);  
}
```

Need to be able to reflect both *direct* and *indirect* recursion.

Interprocedural Analysis



SummaryCallGraph

Call Graph : Polymorphism

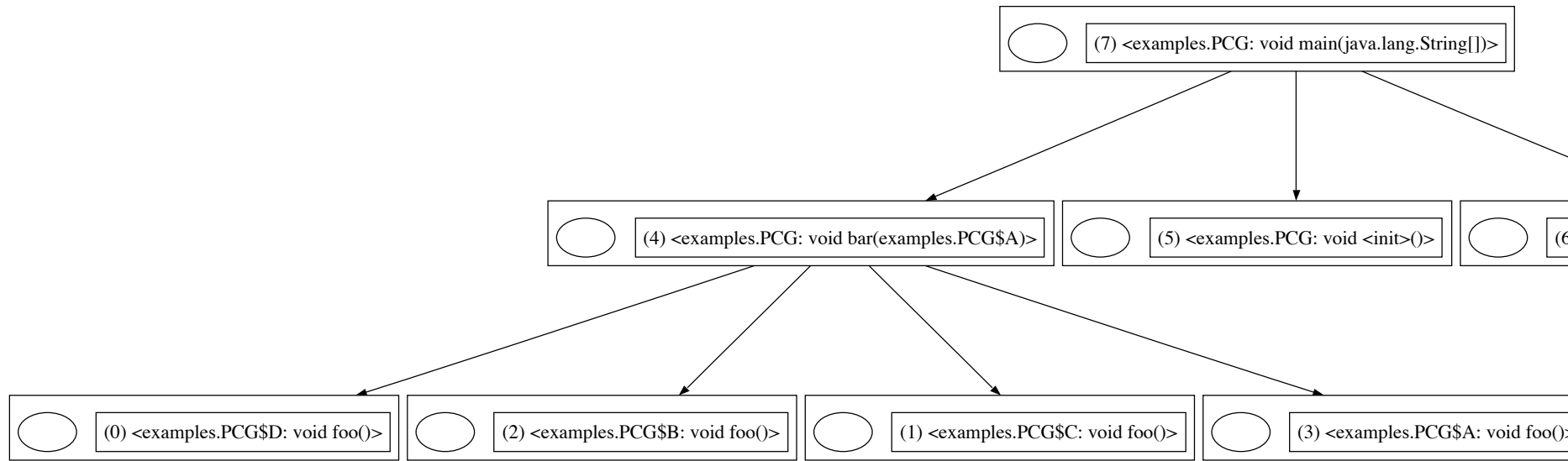
```
class PCG {  
    class A { void foo() { x = 1; } }  
    class B extends A { void foo() { x = 2; } }  
    class C extends A { void foo() { x = 3; } }  
    class D extends B { void foo() { x = 4; } }  
  
    void main() { bar(new PCG().new A()); }  
  
    static void bar(A a) { a.foo(); }  
}
```

Dynamic dispatch can lead to significant overestimation of the possible calling relationships.

This is one reason *points to* analysis is so important.

Which calls to `foo` in `bar` are possible?

Interprocedural Analysis



SummaryCallGraph

Summary-based Interprocedural Analysis

The basic idea is to perform *dependent* but separate local flow analyses for each method

Dependences between methods, i.e., `foo()` calls `bar()`, are captured by constructing and applying *method summaries*

Summaries are calculated during the analysis until a fix-point is reached

Just as in a local flow analysis order is important, so an interprocedural analysis orders the calculation of summaries according to call dependences, i.e., analyze methods in reverse order of calls.

Method Summaries

A method summary reflects the data flow values that are computed on output of a method call

i.e., the least-upper bound of out at `return` stmts

Summaries are often elements of the underlying flow analysis lattice D , but they need not be.

If a method m calls another method n , the values on exit of m may depend on the summary for n . If the summary for n changes we want force a recomputation of m 's summary.

Applying Method Summaries

When analyzing a method body we use a summary to calculate a *method call transfer function* for a call to n

- if n already has a computed summary we use it
- if n has no stored summary we use the *default* summary

Since summaries account for all of the behavior of a method, transitively through all of its possible calls, a summary can be quite imprecise.

Methods that are not subjected to analysis, e.g., library methods, are assigned a *default* summary.

Parameterized (or Context-sensitive) Method Summaries

One can construct summaries that are functions $D \rightarrow \mathcal{P}(D)$

this can be generalized to some other context Δ as the domain

Intuitively, the domain value defines a calling context and the image defines the summarized effects of the methods in that context.

To construct such a summary, for each value $d \in D$ repeat the following

1. set $\iota = d$ for the extremal node
2. run flow analysis to fix-point
3. calculate $s = \bigsqcup_{r \in \text{Returns}} A_{out}(r)$
4. install the map entry $[d \mapsto s]$ in the summary

Interprocedural State Propagation Analysis

```
void oknested() {  
    open(); coloop(); close();  
}
```

```
void coloop(int x) {  
    while (x < 10) {  
        close(); x++; open();  
    }  
}
```

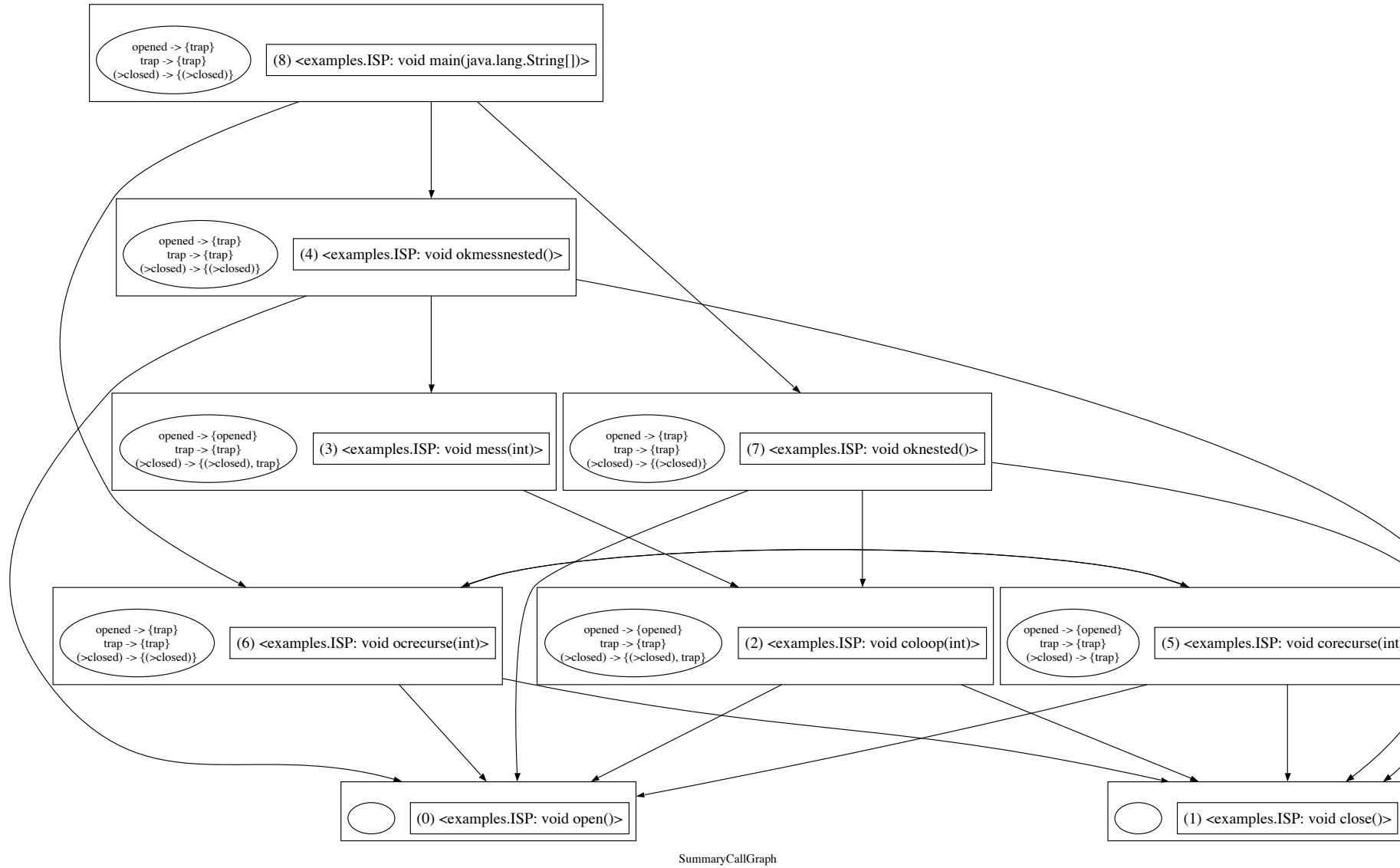
```
void okmessnested() {  
    open(); mess(); close();  
}
```

```
void ocrecurse(int x) {  
    open();  
    if (x > 0) corecurse(x--);  
}
```

```
void corecurse(int x) {  
    close();  
    if (x > 0) ocrecurse(x--);  
}
```

```
void mess(int x) {  
    if (x > 0)  
        while (x-- > 0) coloop(x);  
    else coloop(x); }
```

Interprocedural Analysis



Order of Analysis

Bottom-up in the call graph (but there is lots of freedom to break tie).

The first ones to process are the *leaves*: `open()`, `close()`

`coloop()` is called from multiple methods so we need to process this relatively early in the order.

The chain of `mess()`, `okmessnested()` are processed in reverse call order.

This cluster is *co-dependent* we actually have to reanalyze one of them:

`corecurse()`, `ocrecurse()`, `corecurse()`

`oknested()` could have come earlier, but need not

and finally `main()`