

Concurrent Join Calculus in Scala

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Parallelism vs. Asynchrony vs. Concurrency

Parallelism

Parallelism means...

- ...to use multithreading to speed up a *sequential* computation
 - ▶ main problem: to “parallelize” a computation efficiently
- examples: parallel collections, map/reduce, Spark, ...

Typical task: count words in 10,000 text files

Concurrency vs. Parallelism vs. Asynchrony

Asynchrony

Asynchrony means...

- ...to optimize *sequential* computations that may have long wait times
 - ▶ main problem: to interleave wait times on a single-thread runloop
- examples: futures/promises, async/await, streams, FRP, coroutines, ...

Typical task: implement interactive Excel tables with auto-updating cells

Concurrency vs. Parallelism vs. Asynchrony

Concurrency

Concurrency means...

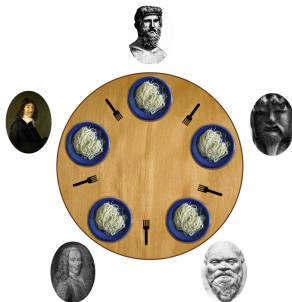
- ...mutually interacting computations, running in unknown order
 - ▶ main problem: to decide when to start a new process (or to wait)
- examples: Thread, synchronized, semaphore, ...

Typical task: simulate “dining philosophers”

Dining philosophers

The exemplary problem of concurrency

Five philosophers sit at a round table, taking turns eating and thinking for random time intervals



Problem: run the process, avoiding deadlock

Problems with concurrency

Imperative concurrency is difficult:

- callbacks, threads, semaphores, mutexes, shared mutable state...
- testing is hard – non-deterministic runtime behavior!
 - ▶ race conditions, deadlocks, livelocks

We try to *avoid* concurrency whenever possible!

How I learned to forget deadlocks and to love concurrency

In this talk:

- Introduction to the “**join calculus**” style of concurrency
- **JoinRun** -- a new Scala implementation
- Examples and demos

Join Calculus: The new hope

...and some new hype

Join Calculus is ...

- ...a declarative language for general-purpose concurrency
- “What if actors were stateless, auto-started, and type-safe”
- No threads/semaphores/locks/mutexes/forks, no shared mutable state
- Concurrency is data-driven, not scheduled
- Easier to use than anything I’ve seen so far!

Metaphors for join calculus:

- “chemical reactions”
- “concurrent pure functions” applied to “concurrent data”

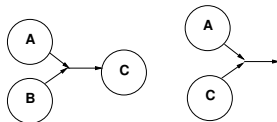
Join Calculus: The genesis

a.k.a. the “Reflexive Chemical Abstract Machine” [Fournet & Gonthier 1996]

Abstract chemistry:

- Chemical “soup” contains many “molecules”
- A combination of certain molecules starts a “chemical reaction”

“Chemical laws”:



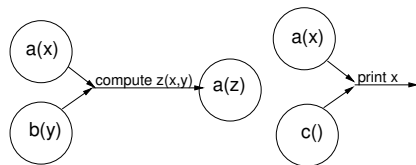
- Define molecules a , b , c , ... and arbitrary chemical laws
- Inject some molecules into the “soup”
- The runtime system evolves the soup *concurrently*

Join Calculus in a nutshell

“Better concurrency through chemistry”

Translating the “chemical metaphor” into practice:

- Each molecule carries a **value**
- Each reaction computes a “molecule-valued” **expression** that depends on input values
- Resulting molecules are injected back into the soup
- Reactions start concurrently if input molecules are available



```
join(  
  run { case a(x) + b(y) =>  
    val z = compute_z(x,y); a(z) },  
  run { case a(x) + c() =>  
    println(x) } )
```

When a reaction starts: input molecules disappear, expression is computed, output molecules are injected

Using JoinRun: basic features

Molecule injectors, reaction definitions

Define some **molecule injectors**:

```
val a = new JA[Int]
val b = new JA[Option[Int]]
```

Declare some **reactions** using the known molecules:

```
val r0 = run { case a(0) => println("finished") }
val r1 = run { case a(x) + b(None) => a(x-1) }
val r2 = run { case a(x) + b(Some(y)) => a(x+y) }
```

Activate a “**join definition**” and inject some molecules:

```
join(r0, r1, r2)
a(10) // non-blocking side-effect
b(Some(5)) // ditto; now we have a(15)
b(None) // now we have a(14)
```

- A molecule – e.g. `a(14)` – is a side-effect, not a value

Using JoinRun: more features

Blocking vs. non-blocking molecules

Blocking molecule:

- implicitly carries a “reply” injector
- the “reply” value will be returned to caller

```
val getN = new JS[Unit, Int]
// revise the join definition, appending this reaction:
... val r3 = run { case r(x) + getN(_, reply) => reply(x) }
join(r0, r1, r2, r3)
// inject non-blocking molecules...
// now inject the blocking molecule:
val x = getN() // blocking call, returns Int
```

JoinRun: Examples I

First benchmark: Counting to zero

Non-blocking counter:

```
val c = ja[Int]("counter")
val g = js[Unit,Int]("getValue")
val d = ja[Unit]("decr")
val f = js[LocalDateTime, Long]("finished")

join(
  run { case c(0) + f(t, reply) =>
    val elapsed = t.until(LocalDateTime.now, ChronoUnit.MILLIS)
    reply(elapsed) },
  run { case g(_,reply) + c(n) => c(n) + reply(n) },
  run { case c(n) + d(_) if n > 0 => c(n-1) }
)
val initialTime = LocalDateTime.now
c(1000)
(1 to 1000).foreach{ _ => d() }
val result = f(initialTime)
```

JoinRun: Examples II

Options, Futures, and Map/Reduce

Future with blocking poll (“`get`”):

```
{ case fut( (f,x) ) => finished( f(x) ) }  
{ case get(_, r) + finished(fx) => r(fx) }
```

Map/Reduce:

```
{ case res(list) + c(s) => res(s::list) }  
{ case get(_, reply) + res(list) => reply(list) }  
res( Nil )
```

```
Seq(1,2,3).foreach(x => c(x*2))  
get() // this returned Seq(4,6,2) in one test
```

JoinRun: Examples III

Five Dining Philosophers

Philosophers 1, 2, 3, 4, 5; forks f12, f23, f34, f45, f51.

```
// ... some declarations omitted for brevity
join (
  run{ case t1(_) => wait(); hA() },
  run{ case t2(_) => wait(); hB() },
  run{ case t3(_) => wait(); hC() },
  run{ case t4(_) => wait(); hD() },
  run{ case t5(_) => wait(); hE() },
  run{ case h1(_) + f12(_) + f51(_) => wait(); t1() + f12() + f51() },
  run{ case h2(_) + f23(_) + f12(_) => wait(); t2() + f23() + f12() },
  run{ case h3(_) + f34(_) + f23(_) => wait(); t3() + f34() + f23() },
  run{ case h4(_) + f45(_) + f34(_) => wait(); t4() + f45() + f34() },
  run{ case h5(_) + f51(_) + f45(_) => wait(); t5() + f51() + f45() }
)
t1() + t2() + t3() + t4() + t5()
f12() + f23() + f34() + f45() + f51()
```

JoinRun: Examples IV

Concurrent merge-sort

The `mergesort` molecule is “recursive”:

- receives the upper-level “`sortedResult`” molecule
- defines its own “`sorted`” molecules in *local scope*
- emits upper-level “`sortedResult`” when done

```
val mergesort = new JA[(Array[T], JA[Array[T]])]  
join(  
  run { case mergesort((arr, sortedResult)) =>  
    if (arr.length <= 1) sortedResult(arr)  
    else {  
      val sorted1 = new JA[Array[T]]  
      val sorted2 = new JA[Array[T]]  
      join(  
        run { case sorted1(x) + sorted2(y) => sortedResult(arrayMerge(x,y))  
      }  
      val (part1, part2) = arr.splitAt(arr.length/2)  
      // inject lower-level mergesort  
      mergesort(part1, sorted1) + mergesort(part2, sorted2)  
    }  
  })
```


Additional features of JoinRun

More bells and whistles

- Per-reaction thread pools:

```
val tp1 = new JProcessPool(threads = 1)
val tp8 = new JProcessPool(threads = 8)
join(
  run { case a(x) + b(y) => ... } onThreads tp1,
  run { case a(x) + c(y) => ... } onThreads tp8
)
```

- Auto-resume failed reactions:

```
run { case a(x) + b(y) => if (bad) throw new Exception(); ... }
```

JoinRun will reinject input molecules if exception is thrown

Roadmap for JoinRun

Need still more bells and whistles

- Graceful global shutdown
- Resilience to failure in join definitions
- Runtime diagnostics, health monitoring
- Run on a cluster (“Distributed Join Calculus”)
- What else is needed for industry-readiness?

Other approaches to concurrency

- STM
- Erlang's message-passing \approx Akka's "actors"
- CSP, π -calculus, **join calculus** (academic so far)

Join Calculus in the wild

- Previous implementations:
 - ▶ Funnel [M. Odersky et al., 2000]
 - ▶ JOCaml (jocaml.inria.fr) [Fournet et al. 2003]
 - ▶ “Join in Scala” compiler patch [V. Cremet 2003]
 - ▶ Joinads (F#, Haskell) [Petricek and Syme 2011]
 - ▶ ScalaJoins [P. Haller 2008]
 - ▶ Scala Join [J. He 2011]
 - ▶ CocoaJoin (iOS), AndroJoin (Android) [S.W. 2013]
- JoinRun -- a new JC prototype in Scala (this talk)
 - ▶ Better syntax, more checks of code sanity
 - ▶ (Some) automatic fault tolerance
 - ▶ Fair scheduling of reactions
 - ▶ Can use thread pools or Akka event-driven actor pools

Comparison: Join Calculus vs. Actor model

Reaction \approx actor

Injected molecule \approx message to actor

Actors:

- need to be created and managed explicitly
- will process one message at a time
- typically hold mutable state

Reactions:

- autostart when required input molecules are available
- many reactions can start at once, automatically concurrent
- immutable, stateless, and type-safe
- all reactions are defined statically, but locally scoped

And I thought “actors” were easy...

Akka's **documentation for the Actor class**:

Actor, ActorSystem, Props (but note the 4 edge cases and 2 warnings)

Actor's companion object; ActorRef

inbox, self, sender, context, supervisorStrategy, watch

Actor lifecycle, Actor selection

send, receive, receive timeout, forward

Future, pipeTo

exceptions, exceptions vs. Future callbacks, andThen

stop, gracefulStop, PoisonPill, Kill

become, unbecome, upgrade, stash

This was item 1 in the Actors documentation.

There are 14 further items...

Everything you need to know about JC...

Most descriptions of JC use the “message/channel” metaphor...

“Chemistry”	JC terminology	JoinRun
molecule	message on channel	<code>a(123) // side effect</code>
injector	channel (port) name	<code>val a : JA[Int]</code>
blocking injector	blocking channel	<code>val q : JS[Int]</code>
reaction	process	<code>run { case a(x) + ... }</code>
injecting a molecule	sending a message	<code>a(123) // side effect</code>
join definition	join definition	<code>◀ ◻ join(r1, r2, ...) ≡</code>

Conclusions and outlook

- Join Calculus = declarative, purely functional concurrency
- Similar to “Actors”, but far easier and “more purely functional”
- Very little known, and very little used in practice
- Existing literature is not suitable as introduction to practical use
- A new Scala implementation, JoinRun, is in the works