# Declarative distributed concurrency in Scala

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### Talk summary

How I learned to forget semaphores and to love concurrency

Chymyst = an implementation of the Chemical Machine (CM) paradigm

- ullet CM pprox Actors made purely functional and auto-parallelized
- Intuitions about why CM works better than other concurrency models
  - ► Comparison with related work: ING Baker, BPMN (workflow)
- New extension for distributed programming: DCM
- Code examples

Not in this talk: academic theory

- ullet Petri nets,  $\pi$ -calculus, join calculus, joinads, mobile agent calculus...
- DCM formulated within a theory of distributed programming?

# Concurrent & parallel programming: How we cope

*Imperative* concurrency & parallelism is difficult to reason about:

- low-level API: callbacks, threads, semaphores, mutex locks
- hard to reason about mutable state and running processes
- hard to test non-deterministic runtime behavior!
- race conditions, deadlocks, livelocks
  Known declarative approaches to avoid these problems:

Kind of concurrency	Typeclass	Scala implementation
synchronous parallelism	applicative functor	Spark, .par.map()
asynchronous streaming	monadic functor	Future, async/await,
DAG		RxJava, Akka Streams
unrestricted streaming	recursive monad+	Flink, fs2, ZIO
unrestricted concurrency	?	Akka, Chymyst

For distributed computing: challenges remain

- coordination and consensus, persistence and fault tolerance
- cluster configuration and discovery
  - distributed coordination as a service: Apache ZooKeeper, etcd

# "Dining philosophers"

The paradigmatic example of concurrency, parallelism and resource contention

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



Problem: simulate the process, avoiding deadlock and starvation Solutions in various programming languages: see Rosetta Code

- Can this be implemented via (effectful) streams? (I think not.)
- The Chemical Machine code is purely declarative

#### What is the Chemical Machine

#### The Chemical Machine paradigm:

- A declarative language for concurrent and parallel computations
  - largely unknown and unused by the software engineering community
  - ► Chymyst an open-source library & embedded DSL for Scala
  - presented in my SBTB talks in 2016 and 2017
- Implement anything in 10 lines of code
- Derive programs quickly from specifications

### Chemical Machine vs. Amazon's AWS Lambda

#### How AWS $\lambda$ works:

- wait for an event that signals arrival of input data
- run a computation whenever input data becomes available
- the computation is automatically parallelized, data-driven
- writing the output data will create a new event
- Modify the AWS $\lambda$  execution model by adding new requirements:
  - a Lambda should be able to wait for several *unrelated* events
  - several Lambdas may contend *atomically* on shared input events ith these new requirements. AWS  $\lambda$  becomes "AWS $\pi$ " a model of
- With these new requirements, AWS $\lambda$  becomes "AWS $\pi$ " a model of "unrestricted concurrency"
  - (Implementation on AWS will be tricky)

#### The Chemical Machine vs. the Actor model

Modify the Actor execution model by adding new requirements:

- when messages arrive, actors are auto-created, maybe in parallel
- actors may wait atomically for messages in *several* different mailboxes It follows from these requirements that...
  - Auto-created actor instances are stateless and invisible to user
  - User code defines mailboxes and computations that consume messages
  - Repeated messages may be consumed in parallel
  - Messages are sent to mailboxes, not to specific actor instances:

```
// Akka // Chymyst val a: ActorRef = ... receive(x) \Rightarrow ... go { case a(x) \Rightarrow ... } val b: ActorRef = ... receive(y) \Rightarrow ... go { case b(y) + c(z) \Rightarrow ... } a ! 100 a(100) b ! 1; b ! 2; b ! 3 b ! 3; c("hello");
```

- All data resides on messages in mailboxes, is consumed automatically
- Mailboxes and computations are *values*, can be sent on messages

Any Actor program can be straightforwardly translated into CM

# Understanding the CM via the chemical metaphor From real to abstract chemistry

Real chemistry is "asynchronous", "concurrent", and "distributed"

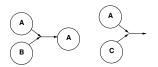
$$HCI + NaOH \rightarrow NaCI + H_2O$$

Want to run *computations* similarly to how chemical reactions run! Begin by formulating the execution model of "abstract chemistry":

- Abstract "molecules" float around in a "chemical reaction site"
- Certain sorts of molecules may combine to start a "reaction":

#### Abstract chemical laws:

$$a + b \rightarrow a$$
  
 $a + c \rightarrow \emptyset$ 



- Program code defines molecules a, b, c, ... and chemical laws
- At initial time, the code emits some molecules into the site
- The runtime system evolves the molecules in parallel
  - ▶ A simulator for this is easy to implement in any language

### Chemical Machine in a nutshell

From abstract chemistry to computation

Translating the chemical metaphor into a model of computation:

- Each molecule carries a value ("concurrent data")
- Each reaction computes new values from its input values
- Some molecules with new values may be emitted back into the reaction site

```
(a(x))
compute z(x,y)
(a(x))
c(y)
c(y)
```

```
site(
go { case a(x) + b(y) \Rightarrow
val z = f(x, y); a(z) \},
go { case a(x) + c(\_) \Rightarrow
println(x) }
```

When a reaction starts: input molecules disappear, new values are computed, output molecules are emitted

Reactions are functions from input values to output values

Need to learn how to "think in molecules"

### Chemical Machine vs. Actor model

- $\bullet$  reaction  $\approx$  function body for an (auto-started) actor
- ullet emitted molecule with value pprox message with value, in a mailbox
- molecule emitters  $\approx$  mailbox references

#### Programming with actors:

- user code creates and manages explicit actor instances
- actors typically hold mutable state and/or mutate "behavior"
  - ▶ reasoning is about running processes *and* the data sent on messages

#### Programming with the Chemical Machine:

- processes auto-start when the needed input molecules are available
- many reactions may start at once, with automatic parallelism
  - user code does not manipulate references to processes
    - ★ no state, no supervision, no lifecycle, no "dead letters", no routers
  - reasoning is only about the data currently available on molecules
    - ★ no reasoning about running processes having state

Chymyst code is typically 2x - 3x shorter than equivalent Akka code

### Example: throttling

Throttle emitting molecules s(x) with minimum allowed delay of delta ms

```
def throttle[X](s: M[X], delta: Long): M[X] = {
 val r = m[X]
 val enable = m[Unit]
// This molecule is confined to the local scope.
 site(
  go { case r(x) + enable(_) \Rightarrow
        s(x)
        Thread.sleep(delta)
        enable()
 enable() // Enable emitting 's' initially.
 r // Outside scope will be able to emit 'r'.
```

- No threads/semaphores/locks, no mutable state
- External code may emit r(x) at will, and s(x) is then throttled

Implementations in Akka, in Monix, and ZIO: > 50 LOC each

# Example: map/reduce

#### A simple map/reduce implementation:

```
val c = m[A] // Initial values have type 'A'.
val d = m[(Int, B)] // 'B' is a commutative monoid.
val res = m[B] // Final result of type 'B'.
val fetch = b[Unit, B] // Blocking emitter.
site(
 // "map"
  go { case c(x) \Rightarrow d((1, long\_computation(x))) },
 // "reduce"
  go { case d((n1, b1)) + d((n2, b2)) \Rightarrow
  val (newN, newB) = (n1 + n2, b1 | + | b2)
   if (newN == total) res(newB) else d((newN, newB))
  1.
  go { case fetch(\_, reply) + res(b) \Rightarrow reply(b) }
(1 to 100).foreach(x \Rightarrow c(x))
fetch() // Blocking call will return the final result.
```

Compare with the Akka implementation here (100+ LOC)

# Example: parallel merge-sort

#### Chymyst code: MergeSortSpec.scala

```
val mergesort = m[(Array[T], M[Array[T]])]
site(
  go { case mergesort((arr, sortedResult)) ⇒
    if (arr.length <= 1) sortedResult(arr)</pre>
      else {
        val sorted1 = m[Array[T]]
        val sorted2 = m[Array[T]]
        site(
          go { case sorted1(x) + sorted2(y) \Rightarrow
             sortedResult(arrayMerge(x,y))
        val (part1, part2) = arr.splitAt(arr.length/2)
        // Emit lower-level mergesort molecules:
        mergesort(part1, sorted1) + mergesort(part2, sorted2)
1)
```

Implementation in Akka: 30 LOC for the same functionality

# Example: Dining philosophers

Five Dining Philosophers

```
Philosophers 1, 2, 3, 4, 5 and forks f12, f23, f34, f45, f51
     // ... definitions of emitters, think(), eat() omitted for brevity
     site (
        go { case t1(_) \Rightarrow think(1); h1() },
        go { case t2(_) \Rightarrow think(2); h2() },
        go { case t3(_) \Rightarrow think(3); h3() },
        go { case t4(_) \Rightarrow think(4); h4() },
        go { case t5(\_) \Rightarrow think(5); h5() },
        go { case h1() + f12() + f51() \Rightarrow eat(1); t1() + f12() + f51() },
        go { case h2(_) + f23(_) + f12(_) \Rightarrow eat(2); t2() + f23() + f12() },
        go { case h3(_) + f34(_) + f23(_) \Rightarrow eat(3); t3() + f34() + f23() },
        go { case h4(_) + f45(_) + f34(_) \Rightarrow eat(4); t4() + f45() + f34() },
        go { case h5(_) + f51(_) + f45(_) \Rightarrow eat(5); t5() + f51() + f45() }
     t1() + t2() + t3() + t4() + t5()
     f12() + f23() + f34() + f45() + f51()
```

Source code: DiningPhilosophers.scala

For more examples, see the code repository (first-of, barriers, rendezvous, critical sections, readers/writers, Game of Life, elevators, etc.)

# Reasoning about code in the Chemical Machine paradigm

#### Reasoning about concurrent data:

- ullet Emit molecule with value pprox lift data into the "concurrent world"
- Define reaction ≈ lift a function into the "concurrent world"
- ullet Reaction site pprox container for concurrent functions and data
- ullet Reaction consumes molecules pprox function consumes input values
- ullet Reaction emits molecules pprox function returns result values

#### Reasoning about code:

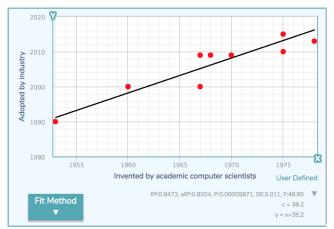
- What data do we need to handle concurrently? (Put it on molecules.)
- What computations consume this data? (Define reactions.)

#### Guarantees:

- Molecule emitters and reactions are immutable values in local scopes
- Reaction sites are immutable once activated; can refactor to libraries
- Multiple input molecules are consumed atomically by reactions

# Chemical Machine paradigm to become mainstream in 2033

- The Chemical Machine paradigm was invented in 1995
- The gap from academic invention to industry adoption is 38.2 years (declarative math, map/reduce, continuations, OOP, CSP, Actors, constraint programming, DAG dataflow,  $\lambda$ -functions, Hindley-Milner)



# Current features of Chymyst

- Blocking molecules with timeouts and back-signalling
- Automatic pipelining of molecules (ordered mailboxes)
- "Static" molecules with read-only access (similar to Akka "agents")
- Compile-time and early run-time DSL error reporting
- Logging, debugging, unit-testing facilities
- Thread pools with thread priority control

### Related frameworks: Petri nets

Workflow management: an approach based on Petri nets

- ING Baker a DSL for workflow management
- Process modeling and control ("elevator system" etc.)
- Business process management (BPM) systems

Chymyst implements a rich version of Petri nets:

- Transitions admit arbitrary guard conditions and error recovery
- Transitions carry values, reactions are values, can be nested
- Nondeterministic, asynchronous, parallel execution

A Petri net model is straightforwardly translated into a CM program

### Distributed Chemical Machine

Run concurrent code on a cluster with no code changes

- Declare some molecules as "distributed", of type DM[T]
- No other new language constructions are necessary!
  - early prototype in progress, as extension of Chymyst

A simple implementation of map/reduce in DCM: 15 LOC

```
implicit val cluster = ClusterConfig(???)
val c = dm[Int] ; val d = dm[Int] // distributed
val res = m[(Int, List[Int])] // local
val fetch = b[Unit, List[Int]]
site(
  go { case c(x) \Rightarrow d(x * 2) }, // "map" on cluster,
// "reduce" on the driver node only.
  go { case res((n, list)) + d(x) \Rightarrow res((n-1, s::list)) },
// fetch results
  go { case fetch(\_, reply) + res((0, list)) \Rightarrow reply(list) }
if (isDriver) { // 'true' only on the driver node.
  Seq(1, 2, 3).foreach(x \Rightarrow c(x))
  res((3, Nil)); fetch() // Returns the result.
```

Comparison: Akka implementation of distributed map/reduce (400+ LOC)

#### Distributed cache in 10 lines of code

Mutable Map[String, String] with operations: put, get, delete

```
implicit val cluster = ClusterConfig(???)
val data = dm[mutable.Map[String, String]]
val put = dm[(String, String)]
val get = dm[(String, M[Option[String]]]
val delete = dm[String]
site(
   go { case data(dict) + put((k, v)) ⇒ data(dict.updated(k, v)) },
   go { case data(dict) + get((k, r)) ⇒ data(dict); r(dict.get(k)) },
   go { case data(dict) + delete(k) ⇒ dict.remove(k); data(dict) }
)
if (isDriver) data(mutable.Map[String, String]())
```

• Comparison: Distributed cache in 100 lines of Akka

# Distributed peer-to-peer chat in 15 LOC

- Register user names in chat room
- Fetch list of users
- Send and receive text messages

```
implicit val cluster = ClusterConfig(???)
val users = dm[List[DM[String]]] // List of users' message emitters.
val carrier = dm[DM[String]] // Carries this node's message emitter.
val fetch = b[Unit, List[DM[String]]]
site(go { case users(es) + carrier(e) \Rightarrow users(e :: es) }
, go { case users(es) + fetch(\_, r) \Rightarrow r(es) } )
val peerName = ??? // Read from config on node.
val sender = new DM[String] (peerName) // Assign unique molecule name.
site(go { case sender(x) \Rightarrow println(s"Peer $peerName reads $x")})
carrier(sender)
if (isDriver) users(Nil)
// Fetch list of users and send a message to Sergei if present.
fetch()
  .find(_.name == "Sergei")
  .foreach(sender ⇒ sender("hello"))
```

• Comparison: Distributed chat in > 100 lines of Akka

# Reasoning in the Distributed Chemical Machine

#### Distributed computing is made declarative

- Determine which data needs to be distributed and/or concurrent
- Determine which computations will need to consume that data
- Emit initial molecules and let the DCM run

#### Pure peer-to-peer architecture

- Distributed molecules may be consumed by any DCM peer
- All DCM peers operate in the same way (no master/worker)
- All DCM peers need to define the same distributed reaction sites
  - ► To designate a DCM peer as a "driver", use config files

### Examples (see documentation)

- Broadcast (DCM peers see it exactly once upon connecting)
- Distributed peer-to-peer chat

# Chemical Machine: implementation details

- Each reaction site has a scheduler thread and a worker thread pool
- Each molecule is "bound" to a unique reaction site
- Each emitted molecule is stored in a multi-set at its reaction site
- Each emitted molecule triggers a search for possible reactions
  - Reaction search proceeds concurrently for different reaction sites
- Reactions are scheduled on the worker thread pool
  - ▶ The thread pool can be configured per-reaction or per-site
- Scala macros are used for static analysis and optimizations
  - Automatically pipelined molecules
  - Simplify and analyze Boolean conditions
- Error analysis is also performed at early run time
  - Reaction site with errors remain inactive

# Distributed Chemical Machine: implementation details

- Each distributed molecule (DM) is bound to a unique reaction site
- Emitted DM data goes into the ZK instance
- Each DCM peer listens to ZK messages and checks for its DMs
  - Once a DM is found, its data is downloaded and deserialized
- On a DCM peer, each DM is identified with a unique local RS
  - ▶ Downloaded molecules are emitted into the local RS to run reactions
  - ► All DCM peers must run identical reaction code for DMs
- Each DCM peer acquires a distributed lock on its DMs
  - ▶ Lock is released once reaction scheduling is complete
- If a node goes down or network fails, molecules will be unconsumed
  - Another DCM peer will pick up these molecules later

### Conclusions and outlook

- Chemical Machine = declarative, purely functional concurrency
  - ► Similar to "Actors", but easier to use and "more purely functional"
  - Significantly shorter code, easier to reason about
- An open-source Scala implementation: Chymyst
  - ► Static DSL code analysis (with Scala macros)
  - Industry-strength features (thread priority control, pipelining, fault tolerance, unit testing and debugging APIs)
  - Extensive documentation: tutorial book
- Promising applications:
  - Workflow management
  - Distributed peer-to-peer systems
  - Process modeling, GUIs, BPM
- Distributed Chemical Machine in the works