

# Pure Functional Epidemics

## An Agent-Based Approach

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### ABSTRACT

Agent-Based Simulation (ABS) is a methodology in which a system is simulated in a bottom-up approach by modelling the micro interactions of its constituting parts, called agents, out of which the global system behaviour emerges.

So far mainly object-oriented techniques and languages have been used in ABS. Using the SIR model of epidemiology, which allows to simulate the spreading of an infectious disease through a population, we show how to use pure Functional Reactive Programming to implement ABS. With our approach we can guarantee the reproducibility of the simulation already at compile time and rule out specific classes of run-time bugs, something that is not possible with traditional object-oriented languages. Also, we found that the representation in a purely functional format is conceptually quite elegant and opens the way to formally reason about ABS.

### KEYWORDS

Functional Reactive Programming, Monadic Stream Functions, Agent-Based Simulation

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## 1 INTRODUCTION

The traditional approach to Agent-Based Simulation (ABS) has so far always been object-oriented techniques, due to the influence of the seminal work of Epstein et al [9] in which the authors claim "[...] object-oriented programming to be a particularly natural development environment for Sugarscape specifically and artificial societies generally [...]" (p. 179). This work established the metaphor in the ABS community, that *agents map naturally to objects* [22] which still holds up today.

In this paper we challenge this metaphor and explore ways of approaching ABS in a pure (lack of implicit side-effects) functional way using Haskell. By doing this we expect to leverage the benefits of pure functional programming [12]: higher expressivity through

declarative code, being polymorph and explicit about side-effects through monads, more robust and less susceptible for bugs due to explicit data flow and lack of implicit side-effects.

As use case we introduce the SIR model of epidemiology with which one can simulate epidemics, that is the spreading of an infectious disease through a population, in a realistic way.

Over the course of four steps, we derive all necessary concepts required for a full agent-based implementation. We start from a very simple solution running in the Random Monad which has all general concepts already there and then refine it in various ways, making the transition to Functional Reactive Programming (FRP) [36] and to Monadic Stream Functions (MSF) [26].

The aim of this paper is to show how ABS can be done in *pure* Haskell and what the benefits and drawbacks are. By doing this we give the reader a good understanding of what ABS is, what the challenges are when implementing it and how we solve these in our approach.

The contributions of this paper are:

- We present an approach to agent-based simulation using *declarative* analysis with FRP in which we systematically introduce the concepts of ABS to *pure* functional programming in a step-by-step approach. Also this work presents a new field of application to FRP as to the best of our knowledge the application of FRP to ABS (on a technical level) has not been addressed before. The result of using FRP allows expressing continuous time-semantics in a very clear, compositional and declarative way, abstracting away the low-level details of time-stepping and progress of time within an agent.
- Our approach can guarantee reproducibility already at compile time, which means that repeated runs of the simulation with same initial conditions will always result in the same dynamics, something highly desirable in simulation in general. This can only be achieved through purity, which guarantees the absence of implicit side-effects which allows to rule out non-deterministic influences at compile time through the strong static type system. This becomes only possible in pure functional programming where we can control the side-effects and can program side-effect polymorph, something not possible with traditional object-oriented approaches. Further, through purity and the strong static type system we can rule out important classes of run-time bugs e.g. related to dynamic typing, and the lack of implicit data-dependencies

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which are common in traditional imperative object-oriented approaches.

In Section 2 we define agent-based simulation, introduce functional reactive programming, arrowized programming and monadic stream functions, because our approach builds heavily on these concepts. In Section 3 we introduce the SIR model of epidemiology as an example model to explain the concepts of ABS. The heart of the paper is Section 4 in which we derive the concepts of a pure functional approach to ABS in four steps, using the SIR model. Section 5 discusses related work. Finally, we draw conclusions and discuss issues in Section 6 and point to further research in Section 7.

## 2 BACKGROUND

### 2.1 Agent-Based Simulation

Agent-Based Simulation is a methodology to model and simulate a system where the global behaviour may be unknown but the behaviour and interactions of the parts making up the system is known. Those parts, called agents, are modelled and simulated, out of which then the aggregate global behaviour of the whole system emerges.

So, the central aspect of ABS is the concept of an agent which can be understood as a metaphor for a pro-active unit, situated in an environment, able to spawn new agents and interacting with other agents in some neighbourhood by exchange of messages.

We informally assume the following about our agents [18, 30, 37]:

- They are uniquely addressable entities with some internal state over which they have full, exclusive control.
- They are pro-active which means they can initiate actions on their own e.g. change their internal state, send messages, create new agents, terminate themselves.
- They are situated in an environment and can interact with it.
- They can interact with other agents situated in the same environment by means of messaging.

Epstein [8] identifies ABS to be especially applicable for analysing "spatially distributed systems of heterogeneous autonomous actors with bounded information and computing capacity". It exhibits the following properties:

- Linearity & Non-Linearity - actions of agents can lead to non-linear behaviour of the system.
- Time - agents act over time which is also the source of their pro-activity.
- States - agents encapsulate some state which can be accessed and changed during the simulation.
- Feedback-Loops - because agents act continuously and their actions influence each other and themselves in subsequent time-steps, feedback-loops are the norm in ABS.
- Heterogeneity - although agents can have same properties like height, sex,... the actual values can vary arbitrarily between agents.
- Interactions - agents can be modelled after interactions with an environment or other agents.

- Spatiality & Networks - agents can be situated within e.g. a spatial (discrete 2D, continuous 3D,...) or complex network environment.

### 2.2 Functional Reactive Programming

Functional Reactive Programming is a way to implement systems with continuous and discrete time-semantics in pure functional languages. There are many different approaches and implementations but in our approach we use *Arrowized FRP* [13, 14] as implemented in the library Yampa [4, 11, 21].

The central concept in Arrowized FRP is the Signal Function (SF) which can be understood as a *process over time* which maps an input- to an output-signal. A signal can be understood as a value which varies over time. Thus, signal functions have an awareness of the passing of time by having access to  $\Delta t$  which are positive time-steps with which the system is sampled.

$$\text{Signal } \alpha \approx \text{Time} \rightarrow \alpha$$

$$\text{SF } \alpha \beta \approx \text{Signal } \alpha \rightarrow \text{Signal } \beta$$

Yampa provides a number of combinators for expressing time-semantics, events and state-changes of the system. They allow to change system behaviour in case of events, run signal functions and generate stochastic events and random-number streams. We shortly discuss the relevant combinators and concepts we use throughout the paper. For a more in-depth discussion we refer to [4, 11, 21].

*Event.* An event in FRP is an occurrence at a specific point in time which has no duration e.g. the recovery of an infected agent. Yampa represents events through the *Event* type which is programmatically equivalent to the *Maybe* type.

*Dynamic behaviour.* To change the behaviour of a signal function at an occurrence of an event during run-time, the combinator *switch* ::  $\text{SF } a (b, \text{Event } c) \rightarrow (c \rightarrow \text{SF } a \ b) \rightarrow \text{SF } a \ b$  is provided. It takes a signal function which is run until it generates an event. When this event occurs, the function in the second argument is evaluated, which receives the data of the event and has to return the new signal function which will then replace the previous one.

*Randomness.* In ABS one often needs to generate stochastic events which occur based on e.g. an exponential distribution. Yampa provides the combinator *occasionally* ::  $\text{RandomGen } g \Rightarrow g \rightarrow \text{Time} \rightarrow b \rightarrow \text{SF } a (\text{Event } b)$  for this. It takes a random-number generator, a rate and a value the stochastic event will carry. It generates events on average with the given rate. Note that at most one event will be generated and no 'backlog' is kept. This means that when this function is not sampled with a sufficiently high frequency, depending on the rate, it will lose events.

Yampa also provides the combinator *noise* ::  $(\text{RandomGen } g, \text{Random } b) \Rightarrow g \rightarrow \text{SF } a \ b$  which generates a stream of noise by returning a random number in the default range for the type *b*.

*Running signal functions.* To *purely* run a signal function Yampa provides the function *embed* ::  $\text{SF } a \ b \rightarrow (a, [(DTime, \text{Maybe } a)]) \rightarrow [b]$  which allows to run an SF for a given number of steps where in each step one provides the  $\Delta t$  and an input *a*. The function then returns the output of the signal function for each step. Note that the

input is optional, indicated by *Maybe*. In the first step at  $t = 0$ , the initial  $a$  is applied and whenever the input is *Nothing* in subsequent steps, the last  $a$  which was not *Nothing* is re-used.

### 2.3 Arrowized programming

Yampa's signal functions are arrows, requiring us to program with arrows. Arrows are a generalisation of monads which, in addition to the already familiar parameterisation over the output type, allow parameterisation over their input type as well [13, 14].

In general, arrows can be understood to be computations that represent processes, which have an input of a specific type, process it and output a new type. This is the reason why Yampa is using arrows to represent their signal functions: the concept of processes, which signal functions are, maps naturally to arrows.

There exists a number of arrow combinators which allow arrowized programming in a point-free style but due to lack of space we will not discuss them here. Instead we make use of Paterson's do-notation for arrows [23] which makes code more readable as it allows us to program with points.

To show how arrowized programming works, we implement a simple signal function, which calculates the acceleration of a falling mass on its vertical axis as an example [27].

```
fallingMass :: Double -> Double -> SF () Double
fallingMass p0 v0 = proc _ -> do
  v <- arr (+v0) <<< integral -< (-9.8)
  p <- arr (+p0) <<< integral -< v
  returnA -< p
```

To create an arrow, the *proc* keyword is used, which binds a variable after which then the *do* of Patersons do-notation [23] follows. Using the signal function *integral :: SF v v* of Yampa which integrates the input value over time using the rectangle rule, we calculate the current velocity and the position based on the initial position  $p0$  and velocity  $v0$ . The *<<<* is one of the arrow combinators which composes two arrow computations and *arr* simply lifts a pure function into an arrow. To pass an input to an arrow, *-<* is used and *<-* to bind the result of an arrow computation to a variable. Finally to return a value from an arrow, *returnA* is used.

### 2.4 Monadic Stream Functions

Monadic Stream Functions (MSF) are a generalisation of Yampa's signal functions with additional combinators to control and stack side effects. An MSF is a polymorphic type and an evaluation function which applies an MSF to an input and returns an output and a continuation, both in a monadic context [25, 26]:

```
newtype MSF m a b =
  MSF { unMSF :: MSF m a b -> a -> m (b, MSF m a b) }
```

MSFs are also arrows which means we can apply arrowized programming with Patersons do-notation as well. MSFs are implemented in Dunai, which is available on Hackage. Dunai allows us to apply monadic transformations to every sample by means of combinators like *arrM :: Monad m => (a -> m b) -> MSF m a b* and *arrM\_ :: Monad m => m b -> MSF m a b*.

## 3 THE SIR MODEL

To explain the concepts of ABS and of our pure functional approach to it, we introduce the SIR model as a motivating example and



Figure 1: States and transitions in the SIR compartment model.

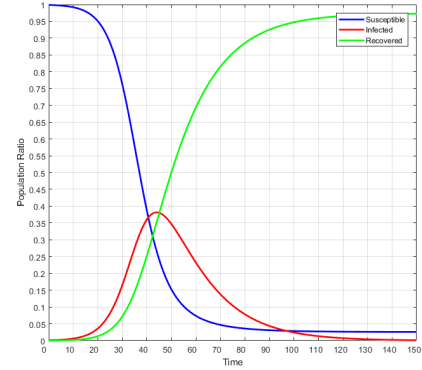


Figure 2: Dynamics of the SIR compartment model using the System Dynamics approach. Population Size  $N = 1,000$ , contact rate  $\beta = \frac{1}{5}$ , infection probability  $\gamma = 0.05$ , illness duration  $\delta = 15$  with initially 1 infected agent. Simulation run for 150 time-steps.

use-case for our implementation. It is a very well studied and understood compartment model from epidemiology [16] which allows to simulate the dynamics of an infectious disease like influenza, tuberculosis, chicken pox, rubella and measles spreading through a population [7].

In this model, people in a population of size  $N$  can be in either one of three states *Susceptible*, *Infected* or *Recovered* at a particular time, where it is assumed that initially there is at least one infected person in the population. People interact *on average* with a given rate of  $\beta$  other people per time-unit and become infected with a given probability  $\gamma$  when interacting with an infected person. When infected, a person recovers *on average* after  $\delta$  time-units and is then immune to further infections. An interaction between infected persons does not lead to re-infection, thus these interactions are ignored in this model. This definition gives rise to three compartments with the transitions seen in Figure 1.

This model was also formalized using System Dynamics (SD) [28]. In SD one models a system through differential equations, allowing to conveniently express continuous systems which change over time, solving them by numerically integrating over time which gives then rise to the dynamics. We won't go into detail here and provide the dynamics of such a solution for reference purposes, shown in Figure 2.

### An Agent-Based approach

The approach of mapping the SIR model to an ABS is to discretize the population and model each person in the population as an individual agent. The transitions between the states are happening due to discrete events caused both by interactions amongst the

agents and time-outs. The major advantage of ABS is that it allows to incorporate spatiality as shown in Section 4.4 and simulate heterogeneity of population e.g. different sex, age,... something not possible with other simulation methods e.g. SD or Discrete Event Simulation (DES).

According to the model, every agent makes *on average* contact with  $\beta$  random other agents per time unit. In ABS we can only contact discrete agents thus we model this by generating a random event on average every  $\frac{1}{\beta}$  time units. We need to sample from an exponential distribution because the rate is proportional to the size of the population [2]. Note that an agent does not know the other agents' state when making contact with it, thus we need a mechanism in which agents reveal their state in which they are in *at the moment of making contact*. This mechanism is an implementation detail which we will derive in our implementation steps. For now we only assume that agents can make contact with each other somehow.

This results in the following agent behaviour:

- *Susceptible*: A susceptible agent makes contact *on average* with  $\beta$  other random agents. For every *infected* agent it gets into contact with, it becomes infected with a probability of  $\gamma$ . If an infection happens, it makes the transition to the *Infected* state.
- *Infected*: An infected agent recovers *on average* after  $\delta$  time units. This is implemented by drawing the duration from an exponential distribution [2] with  $\lambda = \frac{1}{\delta}$  and making the transition to the *Recovered* state after this duration.
- *Recovered*: These agents do nothing because this state is a consuming state from which there is no escape: recovered agents stay immune and can not get infected again in this model.

## 4 DERIVING A PURE FUNCTIONAL APPROACH

We presented a high-level agent-based approach to the SIR model in the previous section, which focused only on the states and the transitions, but we haven't talked about technical implementation.

In [33] two fundamental problems of implementing an agent-based simulation from a programming-language agnostic point of view is discussed. The first problem is how agents can be proactive and the second how interactions and communication between agents can happen. For agents to be pro-active, they must be able to perceive the passing of time, which means there must be a concept of an agent-process which executes over time. Interactions between agents can be reduced to the problem of how an agent can expose information about its internal state which can be perceived by other agents. Further the authors have shown the influence of different deterministic and non-deterministic elements in agent-based simulation on the dynamics and how the influence of non-determinism can completely break them down or result in different dynamics despite same initial conditions. This means that we want to rule out any potential source of non-determinism.

In this section we will derive a pure functional approach for an agent-based simulation of the SIR model in which we will pose solutions to the previously mentioned problems. We will start out with a very naive approach and show its limitations which we

overcome by adding FRP. Then in further steps we will add more concepts and generalisations, ending up at the final approach which utilises Monadic Stream Functions, a generalisation of FRP.

Of paramount importance is to keep our implementations pure which rules out the use of the IO Monad and thus any potential source of non-determinism under all circumstances because we would lose all compile time guarantees about reproducibility. Still we will make use of the Random and State Monad which indeed allow side-effects but the crucial point here is that we restrict side-effects only to these types in a controlled way without allowing general unrestricted effects <sup>1</sup>.

### 4.1 Naive beginnings

We start by modelling the states of the agents. Infected agents are ill for some duration, meaning we need to keep track when an infected agent recovers. Also a simulation is stepped in discrete or continuous time-steps thus we introduce a notion of *time* and  $\Delta t$  by defining:

```
type Time      = Double
type TimeDelta = Double
```

```
data SIRState = Susceptible | Infected TimeDelta | Recovered
```

Now we can represent every agent simply as its SIR state. We hold all our agents in a list:

```
type SIRAgent = SIRState
type Agents   = [SIRAgent]
```

Next we need to think about how to actually step our simulation. For this we define a function which advances our simulation with a fixed  $\Delta t$  until a given time  $t$  where in each step the agents are processed and the output is fed back into the next step. This is the source of pro-activity as agents are executed in every time step and can thus initiate actions based on the passing of time. Note that we step the simulation with a hard-coded  $\Delta t = 1.0$  for reasons which become apparent when implementing the *susceptible* behaviour. As already mentioned, the agent-based implementation of the SIR model is inherently stochastic which means we need access to a random-number generator. We decided to use the Random Monad at this point as threading a generator through the simulation and the agents would be very cumbersome. Thus our simulation stepping runs in the Random Monad:

```
runSimulation :: RandomGen g => Time -> Agents -> Rand g [Agents]
runSimulation tEnd as = runSimulationAux 0 as []
  where
    runSimulationAux :: RandomGen g
      => Time -> Agents -> [Agents] -> Rand g [Agents]
    runSimulationAux t as acc
      | t >= tEnd = return (reverse (as : acc))
      | otherwise = do
        as' <- stepSimulation as
        runSimulationAux (t + 1.0) as' (as : acc)

stepSimulation :: RandomGen g => Agents -> Rand g Agents
stepSimulation as = mapM (runAgent as) as
```

Now we can implement the behaviour of an individual agent. First we need to distinguish between the agents SIR states:

```
processAgent :: RandomGen g => Agents -> SIRAgent -> Rand g SIRAgent
processAgent as Susceptible = susceptibleAgent as
processAgent _ (Infected dur) = return (infectedAgent dur)
processAgent _ Recovered     = return Recovered
```

<sup>1</sup>The code of all steps can be accessed freely through the following URL: <https://github.com/thalerjonathan/phd/tree/master/public/purefunctionalepidemics/code>



An agent gets fed the states of all agents in the system from the previous time-step so it can draw random contacts - this is one, very naive way of implementing the interactions between agents.

From our implementation it becomes apparent that only the behaviour of a susceptible agent involves randomness and that a recovered agent is simply a sink - it does nothing and stays constant.

Lets look how we can implement the behaviour of a susceptible agent. It simply makes contact on average with a number of other agents and gets infected with a given probability if an agent it has contact with is infected. Here it becomes apparent that we implicitly assume that the simulation is stepped with a  $\Delta t = 1.0$ . If we would use a different  $\Delta t$  then we need to adjust the contact rate accordingly because it is defined *per time-unit*. This would amount to multiplying with the  $\Delta t$  which in combination with the discretisation using *floor* would lead to too few contacts being made, ultimately resulting in completely wrong dynamics.

When the agent gets infected, it calculates also its time of recovery by drawing a random number from the exponential distribution, meaning it is ill on average for *illnessDuration*.

```
susceptibleAgent :: RandomGen g => Agents -> Rand g SIRAgent
susceptibleAgent as = do
  -- draws from exponential distribution
  rc <- randomExpM (1 / contactRate)
  cs <- replicateM (floor rc) (makeContact as)
  if or cs
  then infect
  else return Susceptible
where
  makeContact :: RandomGen g => Agents -> Rand g Bool
  makeContact as = do
    randContact <- randomElem as
    case randContact of
      -- returns True with given probability
      (Infected _) -> randomBoolM infectivity
      _              -> return False

  infect :: RandomGen g => Rand g SIRAgent
  infect = randomExpM (1 / illnessDuration)
  >>= \rd -> return (Infected rd)
```

The infected agent is trivial. It simply recovers after the given illness duration which is implemented as follows:

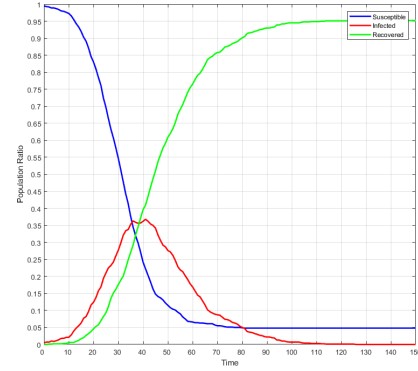
```
infectedAgent :: TimeDelta -> TimeDelta -> SIRAgent
infectedAgent dur
  | dur' <= 0 = Recovered
  | otherwise = Infected dur'
where
  dur' = dur - 1.0
```

Again note the hard-coded  $\Delta t = 1.0$ .

**4.1.1 Results.** When running our naive implementation with a population size of 1,000 we get the dynamics as seen in Figure 3. When comparing it to the dynamics of the reference in Figure 2, the agent-based dynamics are not as smooth which stems from the fact that the agent-based approach is inherently discrete and stochastic [17].

**4.1.2 Discussion.** Reflecting on our first naive approach we can conclude that it already introduced most of the fundamental concepts of ABS

- Time - the simulation occurs over virtual time which is modelled explicitly divided into *fixed*  $\Delta t$  where at each step all agents are executed.
- Agents - we implement each agent as an individual, with the behaviour depending on its state.



**Figure 3: Naive simulation of SIR using the agent-based approach. Population of 1,000, contact rate  $\beta = \frac{1}{5}$ , infection probability  $\gamma = 0.05$ , illness duration  $\delta = 15$  with initially 1 infected agent. Simulation run for 150 time-steps with fixed  $\Delta t = 1.0$ .**

- Feedback - the output state of the agent in the current time-step  $t$  is the input state for the next time-step  $t + \Delta t$ .
- Environment - as environment we implicitly assume a fully-connected network (complete graph) where every agent 'knows' every other agent, including itself and thus can make contact with all of them.
- Stochasticity - it is an inherently stochastic simulation, which is indicated by the Random Monad type and the usage of *randomBoolM* and *randomExpM*.
- Deterministic - repeated runs with the same initial random-number generator result in same dynamics. This may not come as a surprise but in Haskell we can guarantee that property statically already at compile time because our simulation runs in the Random Monad and *not* in the IO Monad. This guarantees that no external, uncontrollable sources of non-determinism can interfere with the simulation.

Nonetheless our approach has also weaknesses and dangers:

- (1) We are using a hard-coded  $\Delta t = 1.0$  because of the way our susceptible behaviour is implemented. Also  $\Delta t$  is dealt with explicitly in the infected behaviour where it is hard-coded to  $\Delta t = 1.0$ . This is not very modular and elegant and a potential source of errors - can we do better and find a more elegant solution?
- (2) The way our agents are represented is not very modular. The state of the agent is explicitly encoded in an ADT and when processing the agent, the behavioural function always needs to distinguish between the states. Can we express it in a more modular way e.g. continuations?

We now move on to the next section in which we will address these points.

## 4.2 Adding Functional Reactive Programming

As shown in the first step, the need to handle  $\Delta t$  explicitly can be quite messy, is inelegant and a potential source of errors, also the explicit handling of the state of an agent and its behavioural function is not very modular. We can solve both these weaknesses

by switching to the Functional Reactive Programming paradigm, because it allows to express systems with discrete and continuous time-semantics.

In this step we are focusing on Arrowized FRP [13] using the library Yampa [11]. In it, time is handled implicitly, meaning it cannot be messed with, which is achieved by building the whole system on the concept of signal functions (SF). An SF can be understood as a process over time and is technically a continuation which allows to capture state using closures. Both these fundamental features allow us to tackle the weaknesses of our first step and push our approach further towards a truly elegant functional approach.

**4.2.1 Implementation.** We start by re-defining the SIR states, whereas now the illness duration is not needed any more. Also we re-define an agent to be an SF which receives the SIR states of all agents as input and outputs the SIR state of the agent:

```
data SIRState = Susceptible | Infected | Recovered
```

```
type SIRAgent = SF [SIRState] SIRState
```

Now we can define the behaviour of an agent to be the following:

```
sirAgent :: RandomGen g => g -> SIRState -> SIRAgent
sirAgent g Susceptible = susceptibleAgent g
sirAgent g Infected    = infectedAgent g
sirAgent g Recovered   = recoveredAgent
```

Depending on the initial state we return the corresponding behaviour. Most notably is the difference that we are now passing a random-number generator instead of running in the Random Monad because signal functions as implemented in Yampa are not capable of being monadic. We see that the recovered agent ignores the random-number generator which is in accordance with the implementation in the previous step where it acts as a sink which returns constantly the same state:

```
recoveredAgent :: SIRAgent
recoveredAgent = arr (const Recovered)
```

When an event occurs we can change the behaviour of an agent using the Yampa combinator *switch*, which is much more elegant and expressive than the initial approach as it makes the change of behaviour at the occurrence of an event explicit. Thus a susceptible agent behaves as susceptible until it becomes infected. Upon infection an *Event* is returned which results in switching into the *infectedAgent* SF, which causes the agent to behave as an infected agent from that moment on. Instead of randomly drawing the number of contacts to make, we now follow a fundamentally different approach by using Yampas *occasionally* function. This requires us to carefully select the right  $\Delta t$  for sampling the system as will be shown in results.

```
susceptibleAgent :: RandomGen g => g -> SIRAgent
susceptibleAgent g =
  switch (susceptible g) (const (infectedAgent g))
  where
    susceptible :: RandomGen g
    => g -> SF [SIRState] (SIRState, Event ())
    susceptible g = proc as -> do
      makeContact <- occasionally g (1 / contactRate) () -< ()
      if isEvent makeContact
      then do
        a <- drawRandomElemSF g -< as
        case a of
          Infected -> do
            i <- randomBoolSF g infectivity -< ()
            if i
            then returnA -< (Infected, Event ())
            else returnA -< (Susceptible, NoEvent)
```

```
        -> returnA -< (Susceptible, NoEvent))
    else returnA -< (Susceptible, NoEvent)
```

We deal with randomness different now and implement signal functions built on the *noiseR* function provided by Yampa. This is an example for the stream character and statefulness of a signal function as it needs to keep track of the changed random-number generator internally through the use of continuations and closures. Here we provide the implementation of *randomBoolSF*. *drawRandomElemSF* works similar but takes a list as input and returns a randomly chosen element from it:

```
randomBoolSF :: RandomGen g => g -> Double -> SF () Bool
randomBoolSF g p = proc _ -> do
  r <- noiseR ((0, 1) :: (Double, Double)) g -< ()
  returnA -< (r <= p)
```

The infected agent behaves as infected until it recovers on average after the illness duration after which it behaves as a recovered agent by switching into *recoveredAgent*. As in the case of the susceptible agent, we use the *occasionally* function to generate the event when the agent recovers. Note that the infected agent ignores the states of the other agents as its behaviour is completely independent of them.

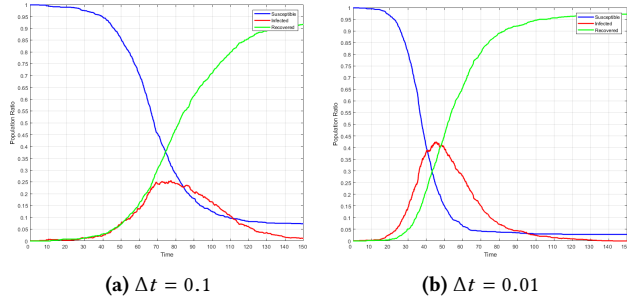
```
infectedAgent :: RandomGen g => g -> SIRAgent
infectedAgent g = switch infected (const recoveredAgent)
  where
    infected :: SF [SIRState] (SIRState, Event ())
    infected = proc _ -> do
      recEvt <- occasionally g illnessDuration () -< ()
      let a = event Infected (const Recovered) recEvt
      returnA -< (a, recEvt)
```

Running and stepping the simulation works now a bit differently, using Yampas function *embed*:

```
runSimulation :: RandomGen g
=> g -> Time -> DTime -> [SIRState] -> [[SIRState]]
runSimulation g t dt as
  = embed (stepSimulation sfs as) ((), dts)
  where
    steps      = floor (t / dt)
    dts        = replicate steps (dt, Nothing)
    n          = length as
    (rngs, _)   = rngSplits g n [] -- unique rngs for each agent
    sfs        = zipWith sirAgent rngs as
```

What we need to implement next is a closed feedback-loop - the heart of every agent-based simulation. Fortunately, [4, 21] discusses implementing this in Yampa. The function *stepSimulation* is an implementation of such a closed feedback-loop. It takes the current signal functions and states of all agents, runs them all in parallel and returns the new agent states of this step. Note the use of *notYet* which is required because in Yampa switching occurs immediately at  $t = 0$ . If we don't delay the switching at  $t = 0$  until the next step, we would enter an infinite switching loop - *notYet* simply delays the first switching until the next time-step.

```
stepSimulation :: [SIRAgent] -> [SIRState] -> SF () [SIRState]
stepSimulation sfs as =
  dpSwitch
    -- feeding the agent states to each SF
    (\_ sfs' -> (map (\sf -> (as, sf)) sfs'))
    -- the signal functions
    sfs
    -- switching event, ignored at t = 0
    (switchingEvt >>> notYet)
    -- recursively switch back into stepSimulation
    stepSimulation
  where
    switchingEvt :: SF ((), [SIRState]) (Event [SIRState])
    switchingEvt = arr (\ (_, newAs) -> Event newAs)
```



**Figure 4: FRP simulation of agent-based SIR showing the influence of different  $\Delta t$ .** Population size of 1,000 with contact rate  $\beta = \frac{1}{5}$ , infection probability  $\gamma = 0.05$ , illness duration  $\delta = 15$  with initially 1 infected agent. Simulation run for 150 time-steps with respective  $\Delta t$ .

Yampa provides the `dpSwitch` combinator for running signal functions in parallel, which has the following type-signature:

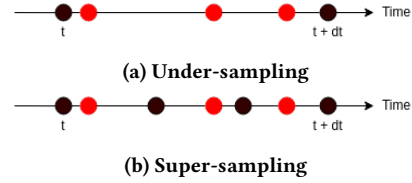
```
dpSwitch :: Functor col
  -- routing function
  => (forall sf. a -> col sf -> col (b, sf))
  -- SF collection
  -> col (SF b c)
  -- SF generating switching event
  -> SF (a, col c) (Event d)
  -- continuation to invoke upon event
  -> (col (SF b c) -> d -> SF a (col c))
  -> SF a (col c)
```

Its first argument is the pairing-function which pairs up the input to the signal functions - it has to preserve the structure of the signal function collection. The second argument is the collection of signal functions to run. The third argument is a signal function generating the switching event. The last argument is a function which generates the continuation after the switching event has occurred. `dpSwitch` returns a new signal function which runs all the signal functions in parallel and switches into the continuation when the switching event occurs. The `d` in `dpSwitch` stands for decoupled which guarantees that it delays the switching until the next time-step: the function into which we switch is only applied in the next step, which prevents an infinite loop if we switch into a recursive continuation.

Conceptually, `dpSwitch` allows us to recursively switch back into the `stepSimulation` with the continuations and new states of all the agents after they were run in parallel.

**4.2.2 Results.** The dynamics generated by this step can be seen in Figure 4.

In this step we followed the FRP approach which is fundamentally different from the previous step in Section 4.1 because in FRP we assume a continuous flow of time as opposed to discrete time-steps of  $\Delta t = 1.0$  previously. This means that we need to select a *correct*  $\Delta t$  otherwise we would end up with wrong dynamics. The selection of a correct  $\Delta t$  depends in our case on *occasionally* in the *susceptible* behaviour, which randomly generates an event on average with *contact rate* following the exponential distribution. To arrive at the correct dynamics, this requires us to sample *occasionally*, and thus the whole system, with small enough  $\Delta t$



**Figure 5: A visual explanation of under-sampling and super-sampling.** The black dots represent the time-steps of the simulation. The red dots represent virtual events which occur at specific points in continuous time. In the case of under-sampling, 3 events occur in between the two time steps but *occasionally* only captures the first one. By increasing the sampling frequency either through a smaller  $\Delta t$  or super-sampling all 3 events can be captured.

which matches the frequency of events generated by *contact rate*. If we choose a too large  $\Delta t$ , we lose events which will result in wrong dynamics as can be seen in Figure 4a. This issue is known as under-sampling and is described in Figure 5.

For tackling this issue we have two options. The first one is to use a smaller  $\Delta t$  as can be seen 4b, which results in the whole system being sampled more often, thus reducing performance. The other option is to implement super-sampling and apply it to *occasionally* which would allow us to run the whole simulation with  $\Delta t = 1.0$  and only sample the *occasionally* function with a much higher frequency.

An approach to super-sampling would be to introduce a new combinator to Yampa which allows to super-sample other signal functions.

```
superSampling :: Int -> SF a b -> SF a [b]
```

It evaluates the *SF* argument for  $n$  times, each with  $\Delta t = \frac{\Delta t}{n}$  and the same input argument  $a$  for all  $n$  evaluations. At time 0 no super-sampling is performed and just a single output of the *SF* argument is calculated. A list of  $b$  is returned with length of  $n$  containing the result of the  $n$  evaluations of the *SF* argument. If 0 or less super samples are requested exactly one is calculated. We could then wrap the *occasionally* function which would then generate a list of events. We have investigated super-sampling more in-depth but have to omit this due to lack of space.

**4.2.3 Discussion.** By moving on to FRP using Yampa we made a huge improvement in clarity, expressivity and robustness of our implementation. State is now implicitly encoded, depending on which signal function is active. Also by using explicit time-semantics with *occasionally* we can achieve extremely fine grained stochastics. Compared to drawing a random number of events we create only a single event or none at all. This requires to sample the system with a much smaller  $\Delta t$  than in the previous step: we are treating it as a truly continuous agent-based system.

A very severe problem, very hard to find with testing but detectable with in-depth validation analysis, is the fact that in the *susceptible* agent the same random-number generator is used in *occasionally*, `drawRandomElemSF` and `randomBoolSF`. This means that all three stochastic functions, which should be independent from each other, are inherently correlated. This is something one

wants to prevent under all circumstances in a simulation, as it can invalidate the dynamics on a very subtle level, and indeed we have tested the influence of the correlation in this example and it has an impact. We left this severe bug in for explanatory reason, as it shows an example where functional programming actually encourages very subtle bugs if one is not careful. A possible solution would be to simply split the initial random-number generator in *sirAgent* three times (using one of the splitted generators for the next split) and pass three random-number generators to *susceptible*. Note that this is not an issue in Sections 4.1 and 4.3 as we are using the Random Monad, which never uses the same random-number generator twice thus resulting in guaranteed uncorrelated stochastics.

So far we have an acceptable implementation of an agent-based SIR approach. What we are lacking at the moment is a general treatment of an environment. To conveniently introduce it we want to make use of monads which is not possible using Yampa. In the next step we make the transition to Monadic Stream Functions as introduced in Dunai [26] which allows FRP within a monadic context.

### 4.3 Generalising to Monadic Stream Functions

A part of the library Dunai is BearRiver, a wrapper which re-implements Yampa on top of Dunai, which should allow us to easily replace Yampa with MSFs. This will enable us to run arbitrary monadic computations in a signal function, which we will need in the next step when adding an environment.

**4.3.1 Identity Monad.** We start by making the transition to BearRiver by simply replacing Yampas signal function by BearRivers' which is the same but takes an additional type parameter *m* indicating the monadic context. If we replace this type-parameter with the Identity Monad we should be able to keep the code exactly the same, except from a few type-declarations, because BearRiver re-implements all necessary functions we are using from Yampa. We simply re-define our agent signal function, introducing the monad stack our SIR implementation runs in:

```
type SIRMonad = Identity
type SIRAgent = SF SIRMonad [SIRState] SIRState
```

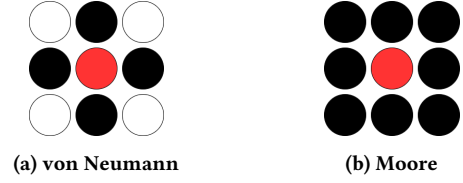
**4.3.2 Random Monad.** Using the Identity Monad does not gain us anything but it is a first step towards a more general solution. Our next step is to replace the Identity Monad by the Random Monad which will allow us to get rid of the RandomGen arguments to our functions and run the whole simulation within the Random Monad *again* just as we started but now with the full features functional reactive programming. We start by re-defining the SIRMonad and SIRAgent:

```
type SIRMonad g = Rand g
type SIRAgent g = SF (SIRMonad g) [SIRState] SIRState
```

The question is now how to access this Random Monad functionality within the MSF context. For the function *occasionally*, there exists a monadic pendant *occasionallyM* which requires a MonadRandom type-class. Because we are now running within a MonadRandom instance we simply replace *occasionally* with *occasionallyM*.

```
occasionallyM :: MonadRandom m => Time -> b -> SF m a (Event b)
```

**4.3.3 Discussion.** So far making the transition to MSFs does not seem as compelling as making the move from the Random Monad



**Figure 6: Common neighbourhoods in discrete 2D environments of Agent-Based Simulation.**

to FRP in the beginning. Running in the Random Monad within FRP is convenient but we could achieve the same with passing RandomGen around as we already demonstrated. Still it guarantees us that we won't have correlated stochastics as discussed in the previous section. In the next step we introduce the concept of a read/write environment which we realise using a StateT monad. This will show the real benefit of the transition to MSFs.

### 4.4 Adding an environment

In this step we will add an environment in which the agents exist and through which they interact with each other. This is a fundamental different approach to agent interaction but is as valid as the approach in the previous steps.

In ABS agents are often situated within a discrete 2D environment [9] which is simply a finite  $N \times M$  grid with either a Moore or von Neumann neighbourhood (Figure 6). Agents are either static or can move freely around with cells allowing either single or multiple occupants.

We can directly map the SIR model to a discrete 2D environment by placing the agents on a corresponding 2D grid with an unrestricted neighbourhood. The behaviour of the agents is the same but they select their interactions directly from the environment. Also instead of feeding back the states of all agents as inputs, agents now communicate through the environment by revealing their current state to their neighbours by placing it on their cell. Agents can read the states of all their neighbours which tells them if a neighbour is infected or not. For purposes of a more interesting approach, we restrict the neighbourhood to Moore (Figure 6b).

**4.4.1 Implementation.** We start by defining our discrete 2D environment for which we use an indexed two dimensional array. In each cell the agents will store their current state, thus we use the *SIRState* as type for our array data:

```
type Disc2dCoord = (Int, Int)
type SIREnv = Array Disc2dCoord SIRState
```

Next we redefine our monad stack and agent signal function. We use a StateT transformer on top of our Random Monad from the previous step with *SIREnv* as type for the state. Our agent signal function now has unit input and output type, which indicates that the actions of the agents are only visible through side-effects in the monad stack they are running in.

```
type SIRMonad g = StateT SIREnv (Rand g)
type SIRAgent g = SF (SIRMonad g) () ()
```

The implementation of a susceptible agent is now a bit different. The agent directly queries the environment for its neighbours and randomly selects one of them. The remaining behaviour is similar:



```

929 susceptibleAgent :: RandomGen g => Disc2dCoord -> SIRAgent g
930 susceptibleAgent coord
931   = switch susceptible (const (infectedAgent coord))
932   where
933     susceptible :: RandomGen g
934     => SF (SIRMonad g) () ((), Event ())
935     susceptible = proc _ -> do
936       makeContact <- occasionallyM (1 / contactRate) () <- ()
937       if not (isEvent makeContact)
938       then returnA <-< ((), NoEvent)
939       else do
940         env <- arrM_ (lift get) <- ()
941         let ns = neighbours env coord agentGridSize moore
942         s <- drawRandomElemS <-< ns
943         case s of
944           Infected -> do
945             infected <- arrM_
946               (lift $ lift $ randomBoolM infectivity) <- ()
947             if infected
948             then do
949               arrM (put . changeCell coord Infected) <- env
950               returnA <-< ((), Event ())
951             else returnA <-< ((), NoEvent)
952           _ -> returnA <-< ((), NoEvent)
953
954 neighbours :: SIREnv -> Disc2dCoord -> Disc2dCoord
955           -> [Disc2dCoord] -> [SIRState]
956
957 moore :: [Disc2dCoord]
958 moore = [ topLeftDelta, topDelta, topRightDelta,
959           leftDelta, rightDelta,
960           bottomLeftDelta, bottomDelta, bottomRightDelta ]
961
962 topLeftDelta :: Disc2dCoord
963 topLeftDelta = (-1, -1)
964 topDelta :: Disc2dCoord
965 topDelta = (0, -1)
966 ...

```

Querying the neighbourhood is done using the *neighbours* function. It takes the environment, the coordinate for which to query the neighbours for, the dimensions of the 2D grid and the neighbourhood information and returns the data of all neighbours it could find. Note that on the edge of the environment, it could be the case that fewer neighbours than provided in the neighbourhood information will be found due to clipping.

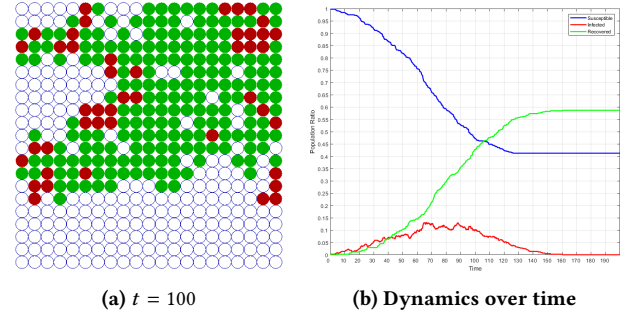
The behaviour of an infected agent is nearly the same as in the previous step, with the difference that upon recovery the infected agent updates its state in the environment from Infected to Recovered.

Running the simulation with MSFs works slightly different. The function *embed* we used before is not provided by BearRiver but by Dunai which has important implications. Dunai does not know about time in MSFs, which is exactly what BearRiver builds on top of MSFs. It does so by adding a ReaderT Double which carries the  $\Delta t$ . This is the reason why we need lifts e.g. in case of getting the environment. Thus *embed* returns a computation in the ReaderT Double Monad which we need to peel away using *runReaderT*. This then results in a StateT computation which we evaluate by using *evalStateT* and an initial environment as initial state. This then results in another monadic computation of the Random Monad type which we evaluate using *evalRand* which delivers the final result. Note that instead of returning agent states we simply return a list of environments, one for each step. The agent states can then be extracted from each environment.

```

984 runSimulation :: RandomGen g => g -> Time -> DTime
985 -> SIREnv -> [(Disc2dCoord, SIRState)] -> [SIREnv]
986 runSimulation g t dt env as = evalRand esRand g

```



**Figure 7: Simulating the agent-based SIR model on a 21x21 2D grid with Moore neighbourhood (Figure 6b), a single infected agent at the center and same SIR parameters as in Figure 2. Simulation run until  $t = 200$  with fixed  $\Delta t = 0.1$ . Last infected agent recovers shortly after  $t = 160$ . The susceptible agents are rendered as blue hollow circles for better contrast.**

```

where
  steps = floor (t / dt)
  dts   = replicate steps ()
  -- initial SFs of all agents
  sfs   = map (uncurry sirAgent) as
  -- running the simulation
  esReader = embed (stepSimulation sfs) dts
  esState = runReaderT esReader dt
  esRand = evalStateT esState env

```

Due to the different approach of returning the SIREnv in every step, we implemented our own MSF:

```

stepSimulation :: RandomGen g
=> [SIRAgent g] -> SF (SIRMonad g) () SIREnv
stepSimulation sfs = MSF (\_ -> do
  -- running all SFs with unit input
  res <- mapM ('unMSF' ()) sfs
  -- extracting continuations, ignore output
  let sfs' = fmap snd res
  -- getting environment of current step
  env <- get
  -- recursive continuation
  let ct = stepSimulation sfs'
  return (env, ct))

```

**4.4.2 Results.** We implemented rendering of the environments using the gloss library which allows us to cycle arbitrarily through the steps and inspect the spreading of the disease over time visually as seen in Figure 7.

Note that the dynamics of the spatial SIR simulation which are seen in Figure 7b look quite different from the reference dynamics of Figure 2. This is due to a much more restricted neighbourhood which results in far fewer infected agents at a time and a lower number of recovered agents at the end of the epidemic, meaning that fewer agents got infected overall.

**4.4.3 Discussion.** At first the environment approach might seem a bit overcomplicated and one might ask what we have gained by using an unrestricted neighbourhood where all agents can contact all others. The real win is that we can introduce arbitrary restrictions on the neighbourhood as shown with the Moore neighbourhood.

Of course an environment is not restricted to be a discrete 2D grid and can be anything from a continuous N-dimensional space

to a complex network - one only needs to change the type of the StateT monad and provide corresponding neighbourhood querying functions. The ability to place the heterogeneous agents in a generic environment is also the fundamental advantage of an agent-based over other simulation approaches and allows to simulate much more realistic scenarios.

#### 4.5 Additional Steps

ABS involves a few more advanced concepts which we didn't fully explore in this paper due to lack of space. Instead we give a short overview and discuss them without presenting code or going into technical details.

**4.5.1 Agent-Transactions.** Agent-transactions are necessary when an arbitrary number of interactions between two agents need to happen instantaneously without time-lag. The use-case for this are price negotiations between multiple agents where each pair of agents needs to come to an agreement in the same time-step [9]. In object-oriented programming, the concept of synchronous communication between agents is implemented directly with method calls.

We have implemented synchronous interactions, which we termed agent-transactions in an additional step. We solved it pure functionally by running the signal functions of the transacting agent pair as often as their protocol requires but with  $\Delta t = 0$ , which indicates the instantaneous character of agent-transactions.

**4.5.2 Event Scheduling.** Our approach is inherently time-driven where the system is sampled with fixed  $\Delta t$ . The other fundamental way to implement an ABS in general, is to follow an event-driven approach [20] which is based on the theory of Discrete Event Simulation [38]. In such an approach the system is not sampled in fixed  $\Delta t$  but advanced as events occur where the system stays constant in between. Depending on the model, in an event-driven approach it may be more natural to express the requirements of the model.

In an additional step we have implemented a rudimentary event-driven approach which allows the scheduling of events but had to omit it due to lack of space. Using the flexibility of MSFs we added a State transformer to the monad stack which allows enqueueing of events into a priority queue. The simulation is advanced by processing the next event at the top of the queue which means running the MSF of the agent which receives the event. The simulation terminates if there are either no more events in the queue or after a given number of events or if the simulation time has advanced to some limit. Having made the transition to MSFs, implementing this feature was quite straight forward which shows the power and strength of the generalised approach to FRP using MSFs.

**4.5.3 Dynamic Agent creation.** In the SIR model, the agent population stays constant - agents don't die and no agents are created during simulation - but some simulations [9] require dynamic agent creation and destruction. We can easily add and remove agents signal functions in the recursive switch after each time-step. The only problem is that creating new agents requires unique agent ids but with the transition to MSFs we can add a monadic context which allows agents to draw the next unique agent id when they create a new agent.

## 5 RELATED WORK

The amount of research on using pure functional programming with Haskell in the field of ABS has been moderate so far. Most of the papers are related to the field of Multi Agent Systems and look into how agents can be specified using the belief-desire-intention paradigm [6, 15, 32].

A library for DES and SD in Haskell called *Aivika 3* is described in the technical report [31]. It is not pure, as it uses the IO Monad under the hood and comes only with very basic features for event-driven ABS, which allows to specify simple state-based agents with timed transitions.

Using functional programming for DES was discussed in [15] where the authors explicitly mention the paradigm of FRP to be very suitable to DES.

A domain-specific language for developing functional reactive agent-based simulations was presented in [35]. This language called FRABJOUS is human readable and easily understandable by domain-experts. It is not directly implemented in FRP/Haskell but is compiled to Yampa code which they claim is also readable. This supports that FRP is a suitable approach to implement ABS in Haskell. Unfortunately, the authors do not discuss their mapping of ABS to FRP on a technical level, which would be of most interest to functional programmers.

Object-oriented programming and simulation have a long history together as the former one emerged out of Simula 67 [5] which was created for simulation purposes. Simula 67 already supported Discrete Event Simulation and was highly influential for today's object-oriented languages. Although the language was important and influential, in our research we look into different approaches, orthogonal to the existing object-oriented concepts.

Lustre is a formally defined, declarative and synchronous dataflow programming language for programming reactive systems [10]. While it has solved some issues related to implementing ABS in Haskell it still lacks a few important features necessary for ABS. We don't see any way of implementing an environment in Lustre as we do in our approach in Section 4.4. Also the language seems not to come with stochastic functions, which are but the very building blocks of ABS. Finally, Lustre does only support static networks, which is clearly a drawback in ABS in general where agents can be created and terminated dynamically during simulation.

## 6 CONCLUSIONS

Our approach is radically different from traditional approaches in the ABS community. First it builds on the already quite powerful FRP paradigm. Second, due to our continuous time approach, it forces one to think properly of time-semantics of the model and how small  $\Delta t$  should be. Third it requires to think about agent interactions in a new way instead of being just method-calls.

Because no part of the simulation runs in the IO Monad and we do not use `unsafePerformIO` we can rule out a serious class of bugs caused by implicit data-dependencies and side-effects which can occur in traditional imperative implementations.

Also we can statically guarantee the reproducibility of the simulation, which means that repeated runs with the same initial conditions are guaranteed to result in the same dynamics. Although we allow side-effects within agents, we restrict them to only the

Random and State Monad in a controlled, deterministic way and never use the IO Monad which guarantees the absence of non-deterministic side effects within the agents and other parts of the simulation.

Determinism is also ensured by fixing the  $\Delta t$  and not making it dependent on the performance of e.g. a rendering-loop or other system-dependent sources of non-determinism as described by [27]. Also by using FRP we gain all the benefits from it and can use research on testing, debugging and exploring FRP systems [24, 27].

## Issues

Currently, the performance of the system is not comparable to imperative implementations but our research was not focusing on this aspect. We leave the investigation and optimization of the performance aspect of our approach for further research.

Despite the strengths and benefits we get by leveraging on FRP, there are errors that are not raised at compile time, e.g. we can still have infinite loops and run-time errors. This was for example investigated in [29] where the authors use dependent types to avoid some run-time errors in FRP. We suggest that one could go further and develop a domain specific type system for FRP that makes the FRP based ABS more predictable and that would support further mathematical analysis of its properties. Furthermore, moving to dependent types would pose a unique benefit over the traditional object-oriented approach and should allow us to express and guarantee even more properties at compile time. We leave this for further research.

In our pure functional approach, agent identity is not as clear as in traditional object-oriented programming, where an agent can be hidden behind a polymorphic interface which is much more abstract than in our approach. Also the identity of an agent is much clearer in object-oriented programming due to the concept of object-identity and the encapsulation of data and methods.

We can conclude that the main difficulty of a pure functional approach evolves around the communication and interaction between agents, which is a direct consequence of the issue with agent identity. Agent interaction is straight-forward in object-oriented programming, where it is achieved using method-calls mutating the internal state of the agent, but that comes at the cost of a new class of bugs due to implicit data flow. In pure functional programming these data flows are explicit but our current approach of feeding back the states of all agents as inputs is not very general and we have added further mechanisms of agent interaction which we had to omit due to lack of space.

## 7 FURTHER RESEARCH

We see this paper as an intermediary and necessary step towards dependent types for which we first needed to understand the potential and limitations of a non-dependently typed pure functional approach in Haskell. Dependent types are extremely promising in functional programming as they allow us to express stronger guarantees about the correctness of programs and go as far as allowing to formulate programs and types as constructive proofs which must be total by definition [1, 19, 34].

So far no research using dependent types in agent-based simulation exists at all. In our next paper we want to explore this for

the first time and ask more specifically how we can add dependent types to our pure functional approach, which conceptual implications this has for ABS and what we gain from doing so. We plan on using Idris [3] as the language of choice as it is very close to Haskell with focus on real-world application and running programs as opposed to other languages with dependent types e.g. Agda and Coq which serve primarily as proof assistants.

We hypothesize that dependent types could help ruling out even more classes of bugs already at compile time and encode invariants and model specifications on the type level, which implies that we don't need to test them using e.g. property-testing with QuickCheck. This would allow the ABS community to reason about a model directly in code:

- Accessing the environment in section 4.4 involves indexed array access which is always potentially dangerous as the indices have to be checked at run-time. Using dependent types it should be possible to encode the environment dimensions into the types. In combination with suitable data types for coordinates one should be able to ensure already at compile time that access happens only within the bounds of the environment.
- In the SIR implementation one could make wrong state-transitions e.g. when an infected agent should recover, nothing prevents one from making the transition back to susceptible. Using dependent types it might be possible to encode invariants and state-machines on the type level which can prevent such invalid transitions already at compile time. This would be a huge benefit for ABS because many agent-based models define their agents in terms of state-machines.
- An infected agent recovers after a given time - the transition of infected to recovered is a timed transition. Nothing prevents us from *never* doing the transition at all. With dependent types we might be able to encode the passing of time in the types and guarantee on a type level that an infected agent has to recover after a finite number of time steps.
- In more sophisticated models agents interact in more complex ways with each other e.g. through message exchange using agent IDs to identify target agents. The existence of an agent is not guaranteed and depends on the simulation time because agents can be created or terminated at any point during simulation. Dependent types could be used to implement agent IDs as a proof that an agent with the given id exists *at the current time-step*. This also implies that such a proof cannot be used in the future, which is prevented by the type system as it is not safe to assume that the agent will still exist in the next step.
- In our implementation, we terminate the SIR model always after a fixed number of time-steps. We can informally reason that restricting the simulation to a fixed number of time-steps is not necessary because the SIR model *has to* reach a steady state after a finite number of steps. This means that at that point the dynamics won't change any more, thus one can safely terminate the simulation. Informally speaking,



the reason for that is that eventually the system will run out of infected agents, which are the drivers of the dynamic. We know that all infected agents will recover after a finite number of time-steps *and* that there is only a finite source for infected agents which is monotonously decreasing. Using dependent types it might be possible to encode this in the types, resulting in a total simulation, creating a correspondence between the equilibrium of a simulation and the totality of its implementation. Of course this is only possible for models in which we know about their equilibria a priori or in which we can reason somehow that an equilibrium exists.

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