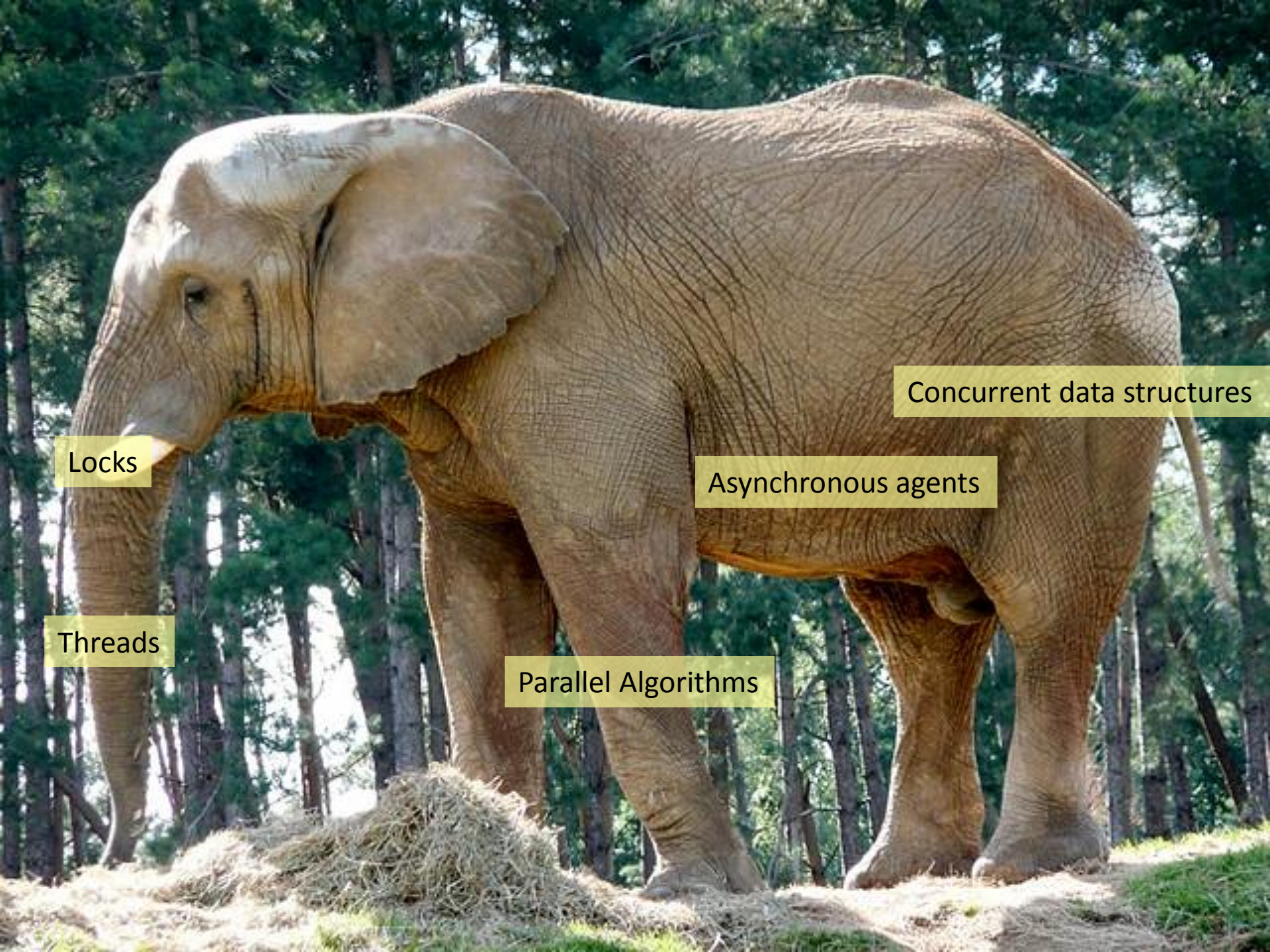


Parallel and Concurrent Haskell

Part I

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Locks

Threads

Parallel Algorithms

Asynchronous agents

Concurrent data structures

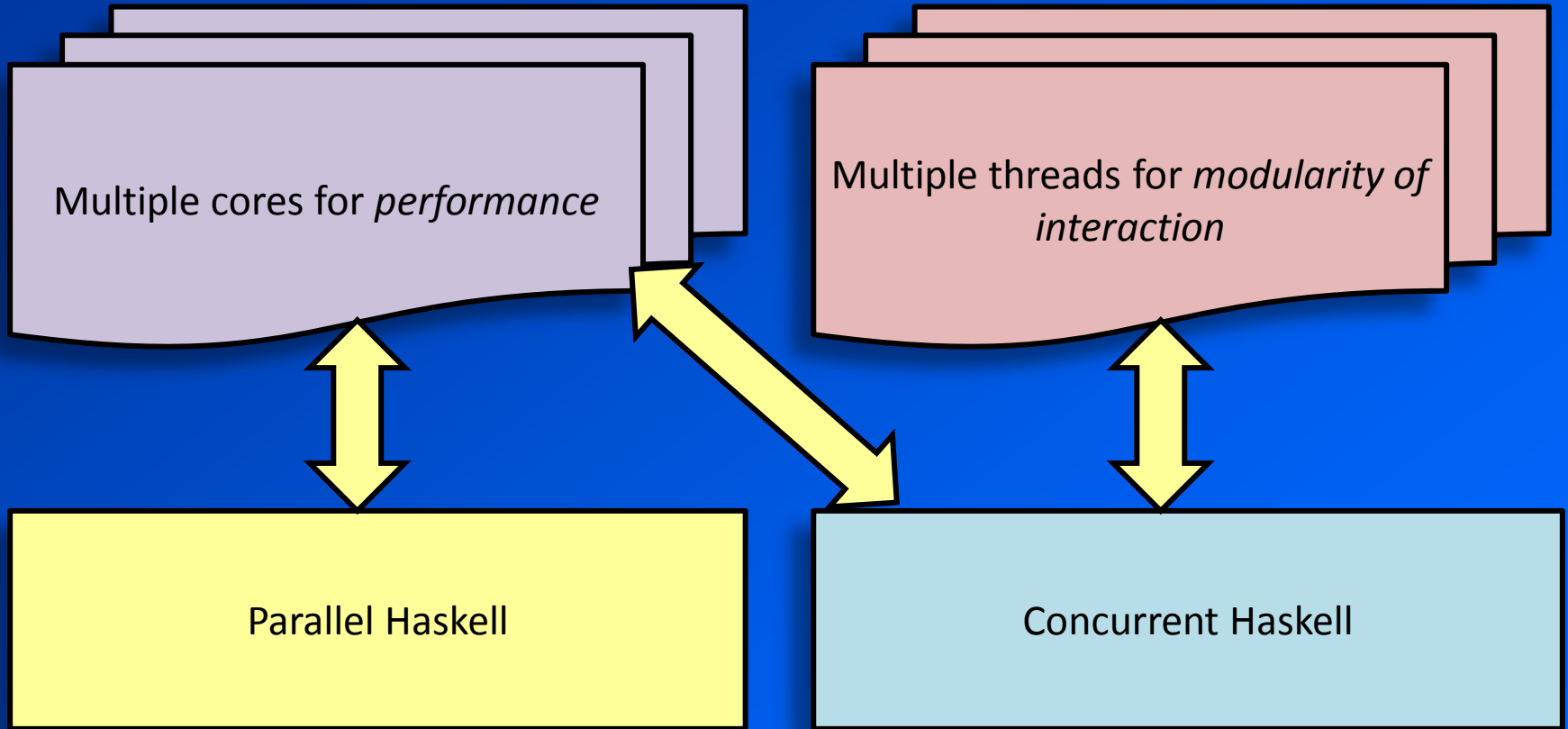
All you need is X

- Where X is actors, threads, transactional memory, futures...
- Often true, but for a given application, some X s will be much more suitable than others.
- In Haskell, our approach is to give you lots of different X s
 - “Embrace diversity (but control side effects)”
(Simon Peyton Jones)

Parallel and Concurrent Haskell ecosystem



Parallelism vs. Concurrency



Parallelism vs. Concurrency

- Primary distinguishing feature of Parallel Haskell: **determinism**
 - The program does “the same thing” regardless of how many cores are used to run it.
 - No race conditions or deadlocks
 - add parallelism without sacrificing correctness
 - Parallelism is used to speed up pure (non-IO monad) Haskell code

Parallelism vs. Concurrency

- Primary distinguishing feature of Concurrent Haskell: threads of control
 - Concurrent programming is done in the IO monad
 - because threads have *effects*
 - effects from multiple threads are interleaved **nondeterministically** at runtime.
 - Concurrent programming allows programs that interact with multiple external agents to be *modular*
 - the interaction with each agent is programmed separately
 - Allows programs to be structured as a collection of interacting agents (actors)

I. Parallel Haskell

- In this part of the course, you will learn how to:
 - Do basic parallelism:
 - compile and run a Haskell program, and measure its performance
 - parallelise a simple Haskell program (a Sudoku solver)
 - use ThreadScope to profile parallel execution
 - do dynamic partitioning
 - measure parallel speedup
 - use Amdahl's law to calculate possible speedup
 - Work with Evaluation Strategies
 - build simple Strategies
 - parallelise a data-mining problem: K-Means
 - Work with the Par Monad
 - Use the Par monad for expressing dataflow parallelism
 - Parallelise a type-inference engine

Running example: solving Sudoku

- code from the Haskell wiki (brute force search with some intelligent pruning)
- can solve all 49,000 problems in 2 mins
- input: a line of text representing a problem

```
.....2143.....6.....2.15.....637.....68..4....23.....7....  
.....241..8.....3..4..5..7....1.....3.....51.6...2...5..3..7...  
.....24...1.....8.3.7...1..1..8..5....2.....2.4..6.5..7.3.....
```

```
import Sudoku
```

```
solve :: String -> Maybe Grid
```

Solving Sudoku problems

- Sequentially:
 - divide the file into lines
 - call the solver for each line

```
import Sudoku
import Control.Exception
import System.Environment

main :: IO ()
main = do
    [f] <- getArgs
    grids <- fmap lines $ readFile f
    mapM (evaluate . solve) grids
```

`evaluate :: a -> IO a`

Compile the program...

```
$ ghc -O2 sudoku1.hs -rtsopts  
[1 of 2] Compiling Sudoku           ( sudoku.hs, sudoku.o )  
[2 of 2] Compiling Main             ( sudoku1.hs, sudoku1.o )  
Linking sudoku1 ...  
$
```

Run the program...

```
$ ./sudoku1 sudoku17.1000.txt +RTS -s
```

```
2,392,127,440 bytes allocated in the heap
```

```
36,829,592 bytes copied during GC
```

```
191,168 bytes maximum residency (11 sample(s))
```

```
82,256 bytes maximum slop
```

```
2 MB total memory in use (0 MB lost due to fragmentation)
```

```
Generation 0: 4570 collections,      0 parallel, 0.14s, 0.13s elapsed
```

```
Generation 1: 11 collections,      0 parallel, 0.00s, 0.00s elapsed
```

```
...
```

```
INIT time 0.00s ( 0.00s elapsed)
```

```
MUT time 2.92s ( 2.92s elapsed)
```

```
GC time 0.14s ( 0.14s elapsed)
```

```
EXIT time 0.00s ( 0.00s elapsed)
```

```
Total time 3.06s ( 3.06s elapsed)
```

```
...
```

Now to parallelise it...

- Doing parallel computation entails specifying coordination in some way – compute A in parallel with B
- This is a constraint on evaluation order
- But by design, Haskell *does not have a specified evaluation order*
- So we need to add something to the language to express constraints on evaluation order

The Eval monad

```
import Control.Parallel.Strategies

data Eval a
instance Monad Eval

runEval :: Eval a -> a

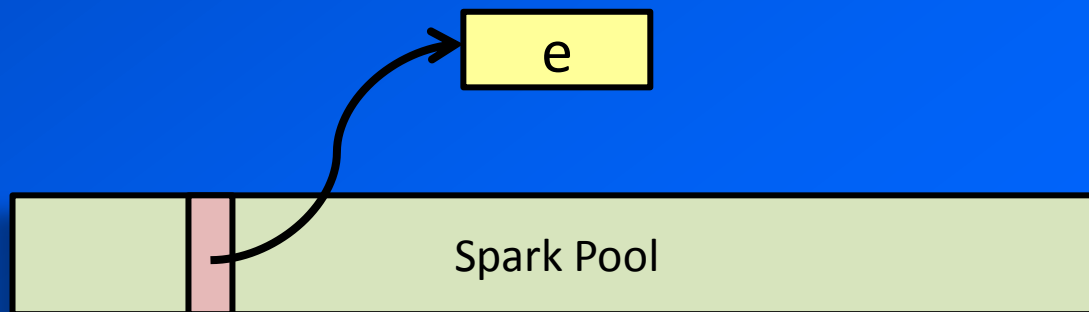
rpar :: a -> Eval a
rseq :: a -> Eval a
```

- Eval is pure
- Just for expressing sequencing between rpar/rseq – nothing more
- Compositional – larger Eval sequences can be built by composing smaller ones using monad combinators
- Internal workings of Eval are very simple (see Haskell Symposium 2010 paper)

What does rpar *actually* do?

```
x <- rpar e
```

- rpar creates a *spark* by writing an entry in the *spark pool*
 - rpar is very cheap! (not a thread)
- the spark pool is a circular buffer
- when a processor has nothing to do, it tries to remove an entry from its own spark pool, or steal an entry from another spark pool (*work stealing*)
- when a spark is found, it is evaluated
- The spark pool can be full – watch out for spark overflow!



Basic Eval patterns

- To compute a in parallel with b, and return a pair of the results:

do

```
a' <- rpar a  
b' <- rseq b  
return (a',b')
```

Start evaluating
a in the
background

Evaluate b, and
wait for the
result

- alternatively:

do

```
a' <- rpar a  
b' <- rseq b  
rseq a'  
return (a',b')
```

- what is the difference between the two?

Parallelising Sudoku

- Let's divide the work in two, so we can solve each half in parallel:

```
let (as,bs) = splitAt (length grids `div` 2) grids
```

- Now we need something like

```
runEval $ do
  as' <- rpar (map solve as)
  bs' <- rpar (map solve bs)
  rseq as'
  rseq bs'
  return ()
```

But this won't work...

```
runEval $ do
  as' <- rpar (map solve as)
  bs' <- rpar (map solve bs)
  rseq as'
  rseq bs'
  return ()
```

- rpar evaluates its argument to Weak Head Normal Form (WHNF)
- WTF is WHNF?
 - evaluates as far as the *first constructor*
 - e.g. for a list, we get either [] or (x:xs)
 - e.g. WHNF of “map solve (a:as)” would be “solve a : map solve as”
- But we want to evaluate the whole list, and the elements

We need to go deeper

```
import Control.DeepSeq
deep :: NFData a => a -> a
deep a = deepseq a a
```

- deep fully evaluates a nested data structure and returns it
 - e.g. a list: the list is fully evaluated, including the elements
- uses overloading: the argument must be an instance of NFData
 - instances for most common types are provided by the library

Ok, adding deep

```
runEval $ do
  as' <- rpar (deep (map solve as))
  bs' <- rpar (deep (map solve bs))
  rseq as'
  rseq bs'
  return ()
```

- Now we just need to evaluate this at the top level in 'main':

```
evaluate $ runEval $ do
  a <- rpar (deep (map solve as))
  ...
```

- (normally using the result would be enough to force evaluation, but we're not using the result here)

Let's try it...

- Compile sudoku2
 - (add -threaded -rtsopts)
 - run with `sudoku17.1000.txt +RTS -N2`
- Take note of the Elapsed Time

Runtime results...

```
$ ./sudoku2 sudoku17.1000.txt +RTS -N2 -s
 2,400,125,664 bytes allocated in the heap
 48,845,008 bytes copied during GC
 2,617,120 bytes maximum residency (7 sample(s))
 313,496 bytes maximum slop
    9 MB total memory in use (0 MB lost due to fragmentation)
```

```
Generation 0: 2975 collections, 2974 parallel, 1.04s, 0.15s elapsed
Generation 1:    7 collections,    7 parallel, 0.05s, 0.02s elapsed
```

```
Parallel GC work balance: 1.52 (6087267 / 3999565, ideal 2)
```

```
SPARKS: 2 (1 converted, 0 pruned)
```

```
INIT   time    0.00s  ( 0.00s elapsed)
MUT    time    2.21s  ( 1.80s elapsed)
GC     time    1.08s  ( 0.17s elapsed)
EXIT   time    0.00s  ( 0.00s elapsed)
Total  time    3.29s  ( 1.97s elapsed)
```

Calculating Speedup

- Calculating speedup with 2 processors:
 - Elapsed time (1 proc) / Elapsed Time (2 procs)
 - NB. not CPU time (2 procs) / Elapsed (2 procs)!
 - NB. compare against sequential program, not parallel program running on 1 proc
- Speedup for sudoku2: $3.06/1.97 = 1.55$
 - not great...

Why not 2?

- there are two reasons for lack of parallel speedup:
 - less than 100% utilisation (some processors idle for part of the time)
 - extra overhead in the parallel version
- Each of these has many possible causes...

A menu of ways to screw up

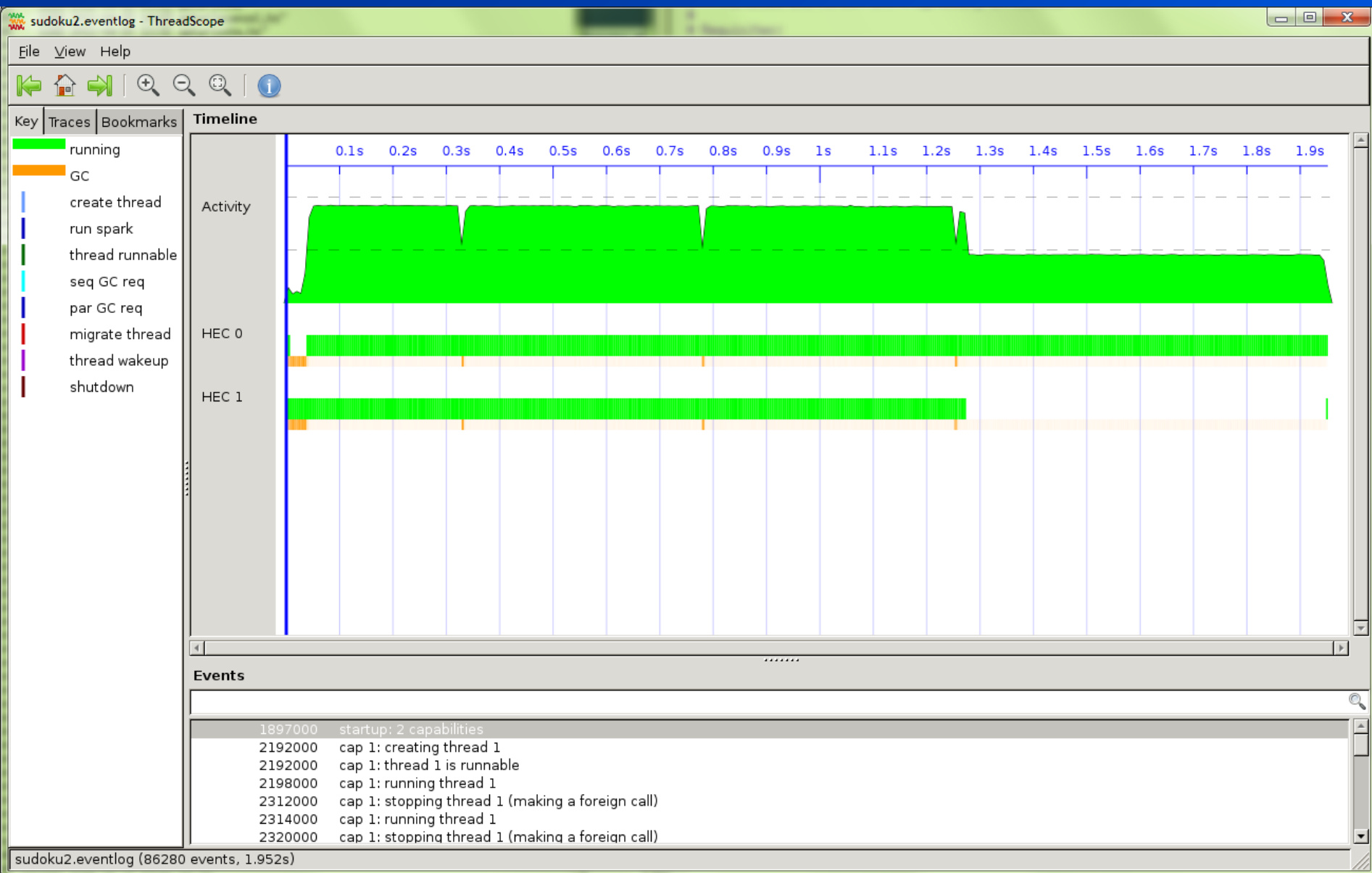
- less than 100% utilisation
 - parallelism was not created, or was discarded
 - algorithm not fully parallelised – residual sequential computation
 - uneven work loads
 - poor scheduling
 - communication latency
- extra overhead in the parallel version
 - overheads from rpar, work-stealing, deep, ...
 - lack of locality, cache effects...
 - larger memory requirements leads to GC overhead
 - GC synchronisation
 - duplicating work

So we need *tools*

- to tell us why the program isn't performing as well as it could be
- For Parallel Haskell we have ThreadScope

```
$ rm sudoku2; ghc -O2 sudoku2.hs -threaded -rtsopts -eventlog  
$ ./sudoku2 sudoku17.1000.txt +RTS -N2 -ls  
$ threadscope sudoku2.eventlog
```

- -eventlog has very little effect on runtime
 - important for profiling parallelism



Uneven workloads...

- So one of the tasks took longer than the other, leading to less than 100% utilisation

```
let (as,bs) = splitAt (length grids `div` 2) grids
```

- One of these lists contains more work than the other, even though they have the same length
 - sudoku solving is not a constant-time task: it is a searching problem, so depends on how quickly the search finds the solution

Partitioning

```
let (as,bs) = splitAt (length grids `div` 2) grids
```

- Dividing up the work along fixed pre-defined boundaries, as we did here, is called *static partitioning*
 - static partitioning is simple, but can lead to under-utilisation if the tasks can vary in size
 - static partitioning does not adapt to varying availability of processors – our solution here can use only 2 processors

Dynamic Partitioning

- Dynamic partitioning involves
 - dividing the work into smaller units
 - assigning work units to processors dynamically at runtime using a *scheduler*
 - good for irregular problems and varying number of processors
- GHC's runtime system provides spark pools to track the work units, and a work-stealing scheduler to assign them to processors
- So all we need to do is use smaller tasks and more rpars, and we get dynamic partitioning

Revisiting Sudoku...

- So previously we had this:

```
runEval $ do
  a <- rpar (deep (map solve as))
  b <- rpar (deep (map solve bs))
  ...
```

- We want to push rpar down into the map
 - each call to solve will be a separate spark

A parallel map

```
parMap :: (a -> b) -> [a] -> Eval [b]
parMap f [] = return []
parMap f (a:as) = do
  b <- rpar (f a)
  bs <- parMap f as
  return (b:bs)
```

Create a spark to
evaluate (f a) for
each element a

Return the new list

- Provided by Control.Parallel.Strategies
- Also:

```
parMap f xs = mapM (rpar . f) xs
```

Putting it together...

```
evaluate $ deep $ runEval $ parMap solve grids
```

- NB. `evaluate $ deep` to fully evaluate the result list
- Code is simpler than the static partitioning version!

Results

```
./sudoku3 sudoku17.1000.txt +RTS -s -N2 -ls
 2,401,880,544 bytes allocated in the heap
 49,256,128 bytes copied during GC
 2,144,728 bytes maximum residency (13 sample(s))
 198,944 bytes maximum slop
    7 MB total memory in use (0 MB lost due to fragmentation)
```

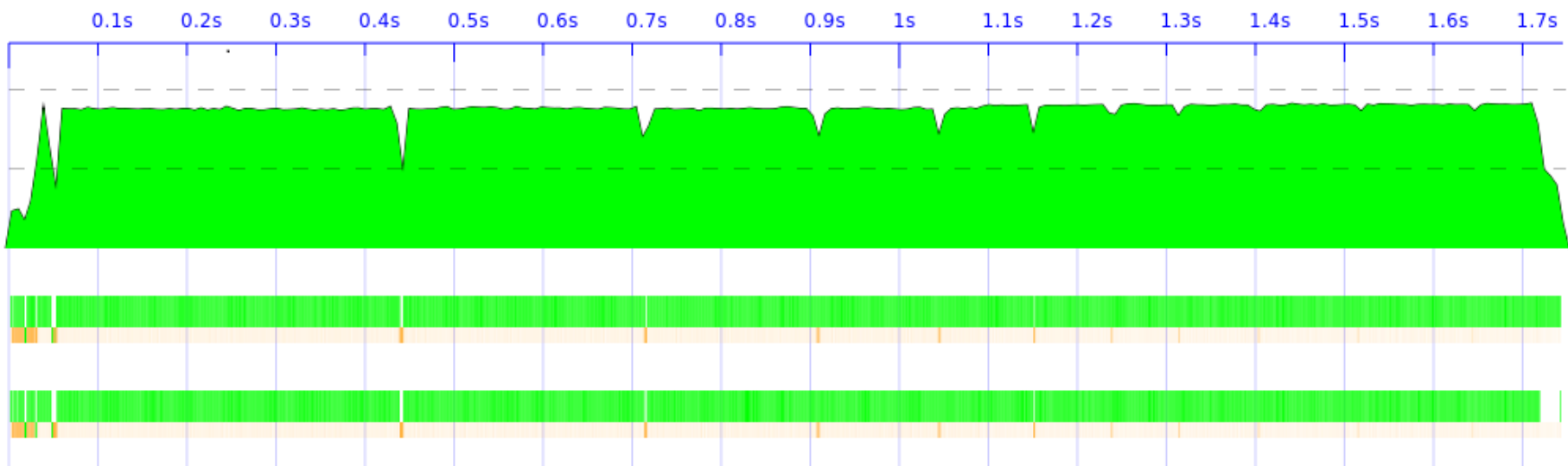
```
Generation 0: 2495 collections, 2494 parallel, 1.21s, 0.17s elapsed
Generation 1:   13 collections,   13 parallel, 0.06s, 0.02s elapsed
```

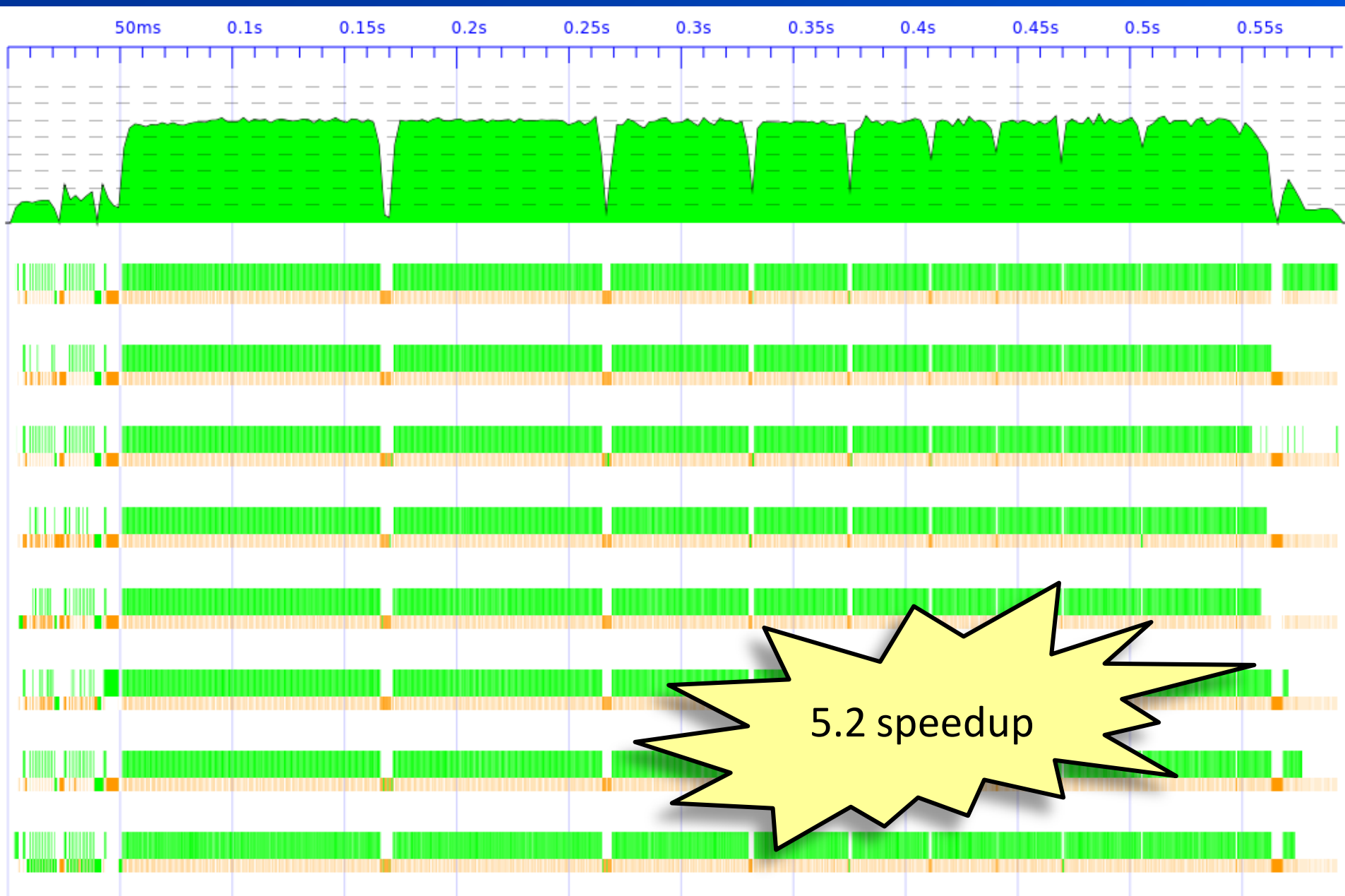
```
Parallel GC work balance: 1.64 (6139564 / 3750823, ideal 2)
```

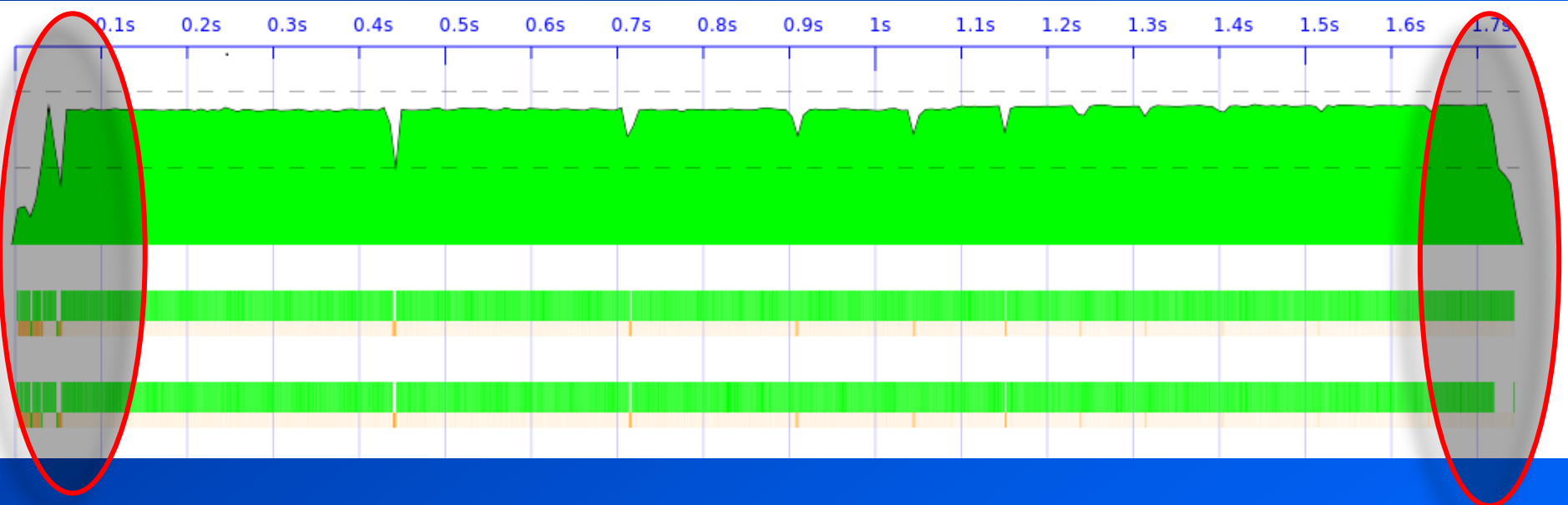
```
SPARKS: 1000 (1000 converted, 0 pruned)
```

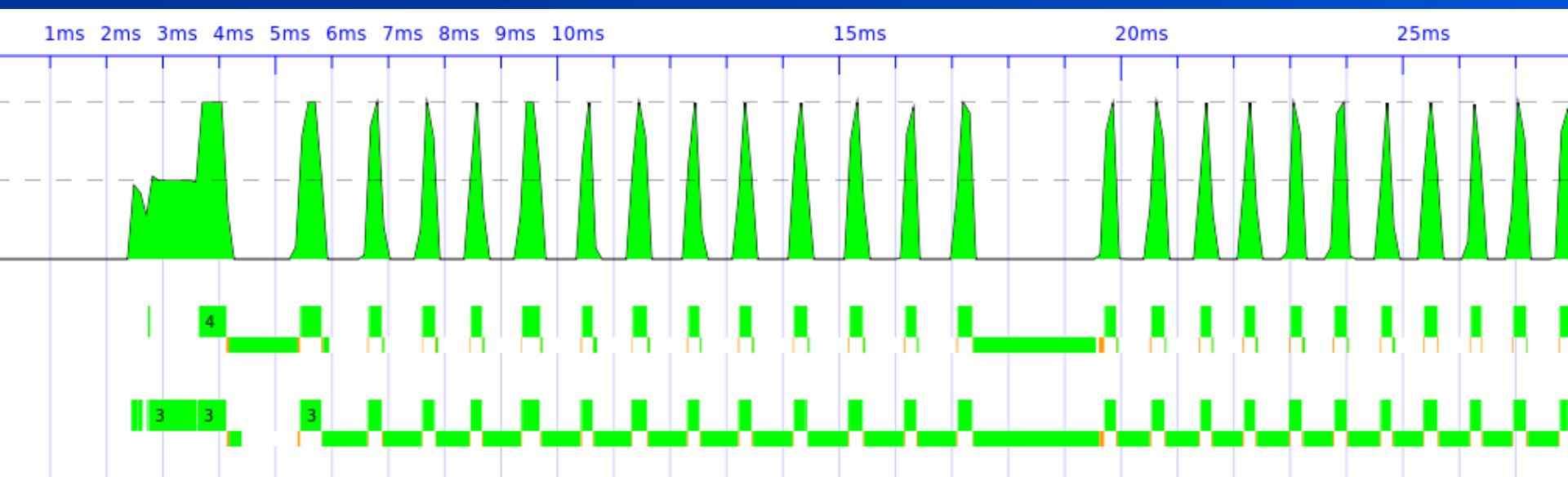
```
INIT   time    0.00s  ( 0.00s elapsed)
MUT    time    2.19s  ( 1.55s elapsed)
GC     time    1.27s  ( 0.19s elapsed)
EXIT   time    0.00s  ( 0.00s elapsed)
Total  time    3.46s  ( 1.74s elapsed)
```

Now 1.7 speedup









- Lots of GC
- One core doing all the GC work
 - indicates one core generating lots of data

```
import Sudoku
import Control.Exception
import System.Environment

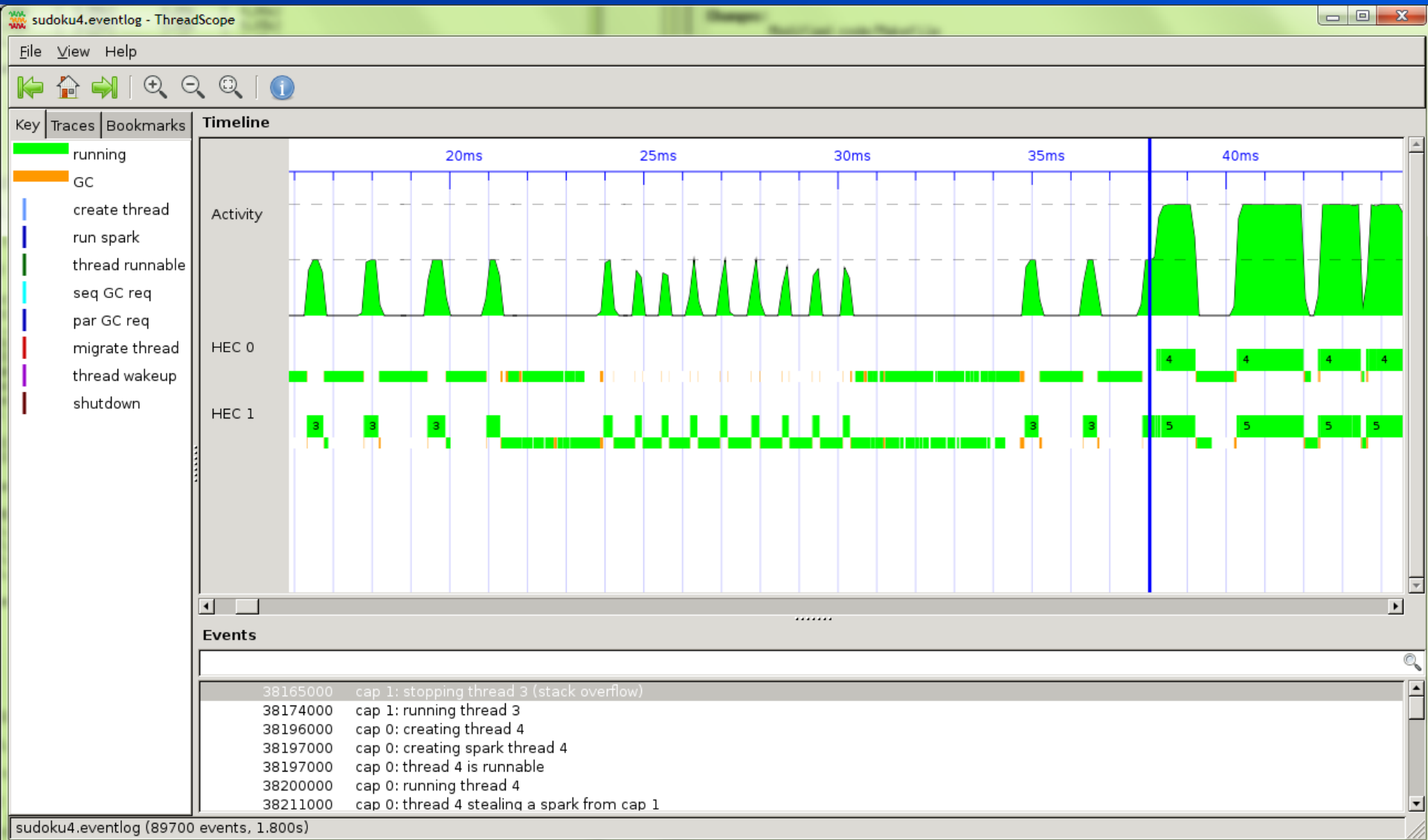
main :: IO ()
main = do
    [f] <- getArgs
    grids <- fmap lines $ readFile f
    evaluate $ deep $ runEval $ parMap solve grids
```

- Are there any sequential parts of this program?
- **readFile** and **lines** are not parallelised

- Suppose we force the sequential parts to happen first...

```
import Sudoku
import Control.Exception
import System.Environment

main :: IO ()
main = do
    [f] <- getArgs
    grids <- fmap lines $ readFile f
    evaluate (length grids)
    evaluate $ deep $ runEval $ parMap solve grids
```



Calculating possible speedup

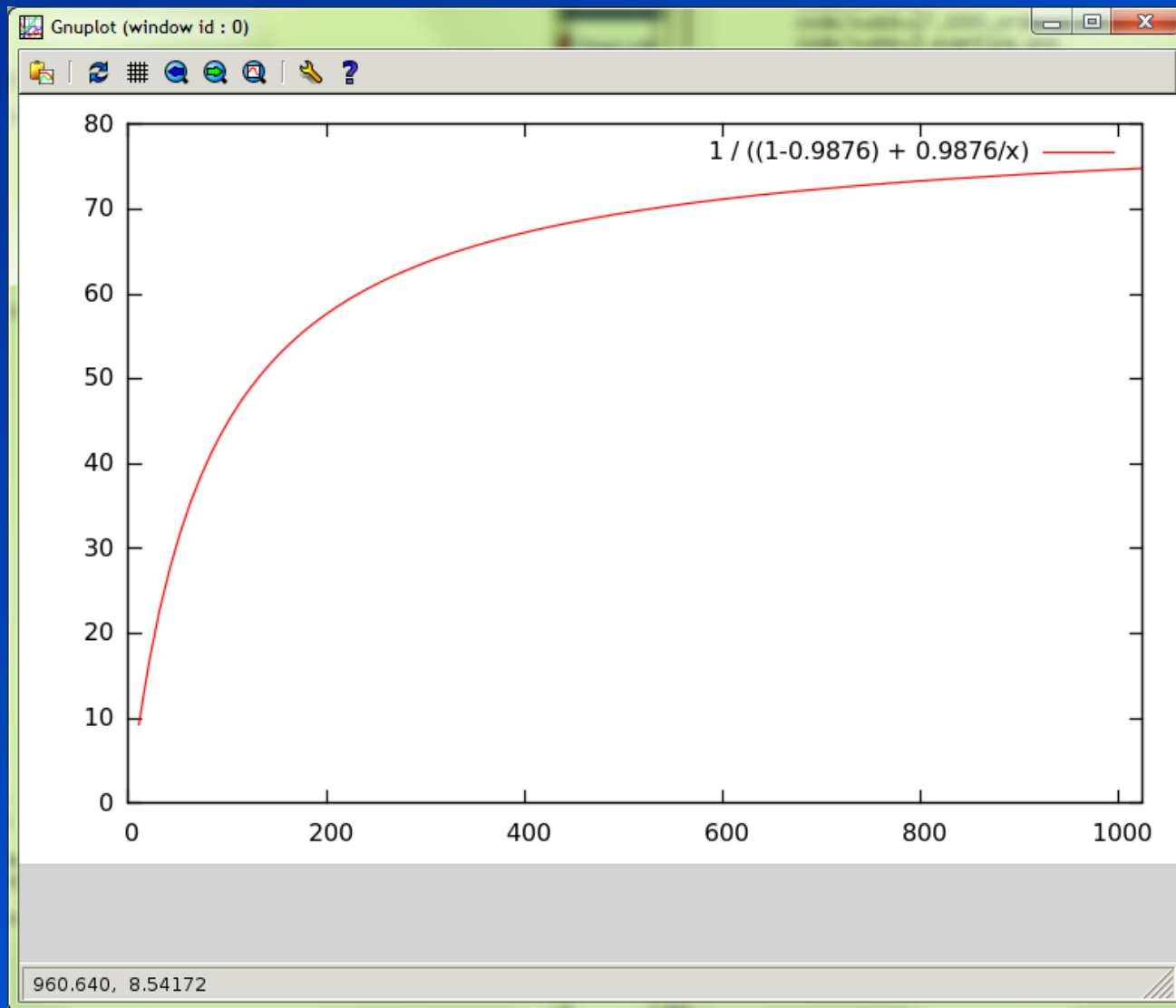
- When part of the program is sequential, Amdahl's law tells us what the maximum speedup is.

$$\frac{1}{(1 - P) + \frac{P}{N}}$$

- P = parallel portion of runtime
- N = number of processors

Applying Amdahl's law

- In our case:
 - runtime = 3.06s (NB. sequential runtime!)
 - non-parallel portion = 0.038s ($P = 0.9876$)
 - $N = 2$, max speedup = $1 / ((1 - 0.9876) + 0.9876/2)$
 - ≈ 1.98
 - on 2 processors, maximum speedup is not affected much by this sequential portion
 - $N = 64$, max speedup = 35.93
 - on 64 processors, 38ms of sequential execution has a dramatic effect on speedup



- diminishing returns...
- See “*Amdahl's Law in the Multicore Era*”, Mark Hill & Michael R. Marty

- Amdahl's law paints a bleak picture
 - speedup gets increasingly hard to achieve as we add more cores
 - returns diminish quickly when more cores are added
 - small amounts of sequential execution have a dramatic effect
 - proposed solutions include heterogeneity in the cores
 - likely to create bigger problems for programmers
- See also Gustafson's law – the situation might not be as bleak as Amdahl's law suggests:
 - with more processors, you can solve a bigger problem
 - the sequential portion is often fixed or grows slowly with problem size
- Note: in Haskell it is **hard to identify the sequential parts anyway**, due to lazy evaluation

Evaluation Strategies

- So far we have used Eval/rpar/rseq
 - these are quite low-level tools
 - but it's important to understand how the underlying mechanisms work
- Now, we will raise the level of abstraction
- Goal: encapsulate parallel idioms as re-usable components that can be composed together.

The Strategy type

```
type Strategy a = a -> Eval a
```

- A Strategy is...
 - A function that,
 - when applied to a value 'a',
 - evaluates 'a' to some degree
 - (possibly sparking evaluation of sub-components of 'a' in parallel),
 - and returns an equivalent 'a' in the Eval monad
- NB. the return value should be observably equivalent to the original
 - (why not the same? we'll come back to that...)

Example...

```
parList :: Strategy [a]
```

- A Strategy on lists that sparks each element of the list
- This is usually not sufficient – suppose we want to evaluate the elements fully (e.g. with `deep`), or do `parList` on nested lists.
- So we parameterise `parList` over the Strategy to apply to the elements:

```
parList :: Strategy a -> Strategy [a]
```

Defining parList

```
type Strategy a = a -> Eval a  
parList :: Strategy a -> Strategy [a]
```

- We have the building blocks:

```
rpar :: a -> Eval a  
      :: Strategy a
```

```
parList :: (a -> Eval a) -> [a] -> Eval [a]  
parList s [] = return []  
parList s (x:xs) = do  
    x' <- rpar (runEval (s x))  
    xs' <- parList s xs  
    return (x':xs')
```

By why *do* Strategies return a value?

```
parList (a -> Eval a) -> [a] -> Eval [a]
parList s []          = return ()
parList s (x:xs) = do
  x'  <- rpar (runEval (s x))
  xs' <- parList s xs
  return (x':xs')
```

- Spark pool points to (runEval (s x))
- If nothing else points to this expression, the runtime will discard the spark, on the grounds that it is not required
- *Always keep hold of the return value of rpar*
- (see the notes for more details on this)

Let's generalise...

- Instead of `parList` which has the sparking behaviour built-in, start with a basic traversal in the `Eval` monad:

```
evalList :: (a -> Eval a) -> [a] -> Eval [a]
evalList f []      = return ()
evalList f (x:xs) = do
  x'  <- f x
  xs' <- parList f xs
  return (x':xs')
```

- and now:

```
parList f = evalList (rpar `dot` f)
  where s1 `dot` s2 = s1 . runEval . s2
```

Generalise further...

- In fact, `evalList` already exists for arbitrary data types in the form of 'traverse'.

```
evalTraversable  
  :: Traversable t => Strategy a -> Strategy (t a)  
  
evalTraversable = traverse  
  
evalList = evalTraversable
```

- So, building Strategies for arbitrary data structures is easy, given an instance of `Traversable`.
- (not necessary to understand `Traversable` here, just be aware that many Strategies are just generic traversals in the `Eval` monad).

How do we *use* a Strategy?

```
type Strategy a = a -> Eval a
```

- We could just use runEval
- But this is better:

```
x `using` s = runEval (s x)
```

- e.g.

```
myList `using` parList rdeepseq
```

- Why better? Because we have a “law”:
 - $x \text{ `using` } s \approx x$
 - We can insert or delete “using s” without changing the semantics of the program

Is that really true?

- Well, not entirely.
- 1. It relies on Strategies returning “the same value” (*identity-safety*)
 - Strategies from the library obey this property
 - Be careful when writing your own Strategies
- 2. `x `using` s` might do more evaluation than just `x`.
 - So the program with `x `using` s` might be `_|_`, but the program with just `x` might have a value
- if identity-safety holds, adding `using` cannot make the program produce a different result (other than `_|_`)

But we wanted 'parMap'

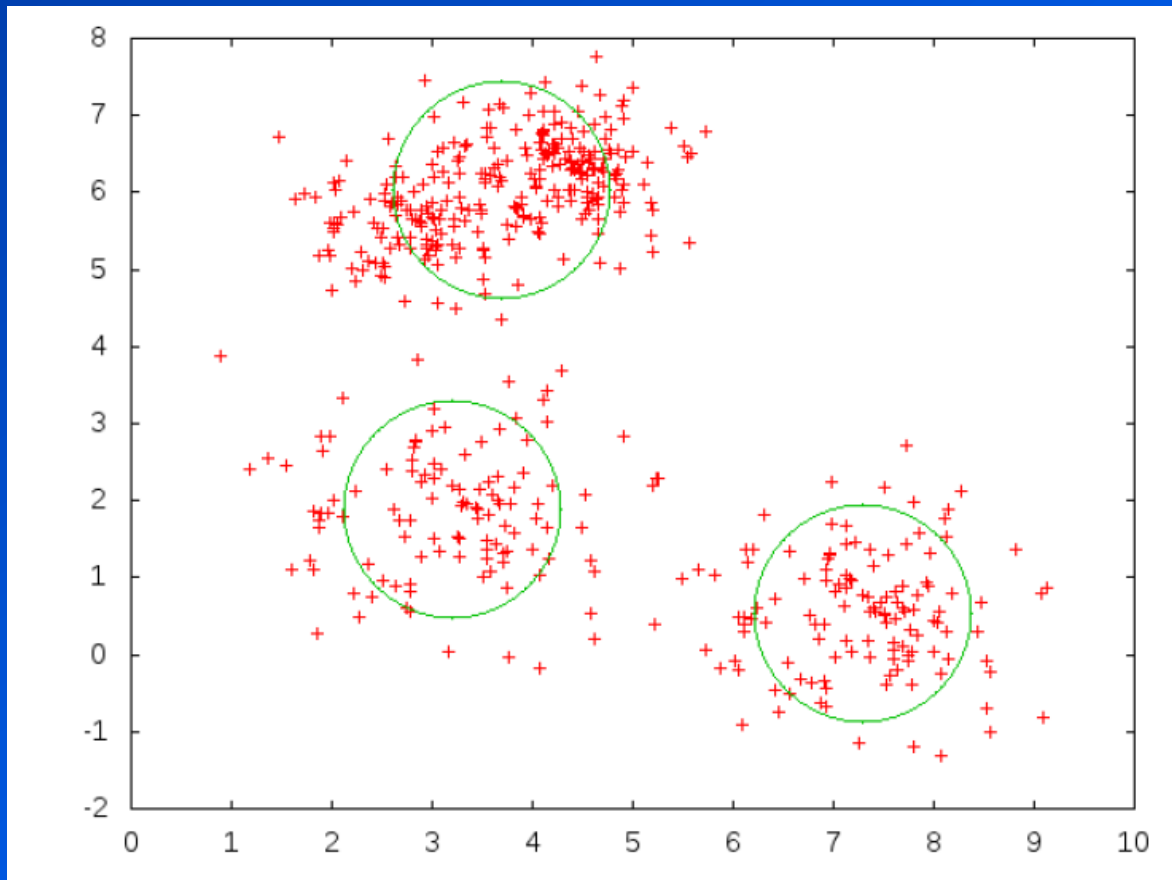
- Earlier we used parMap to parallelise Sudoku
- But parMap is a combination of two concepts:
 - The *algorithm*, 'map'
 - The *parallelism*, 'parList'

```
parMap f x = map f xs `using` parList
```

- With Strategies, the algorithm can be separated from the parallelism.
 - The algorithm produces a (lazy) result
 - A Strategy filters the result, but does not do any computation – it returns the same result.

K-Means

- A data-mining algorithm, to identify clusters in a data set.



K-Means

- We use a heuristic technique (Lloyd's algorithm), based on iterative refinement.
 1. Input: an initial guess at each cluster location
 2. Assign each data point to the cluster to which it is closest
 3. Find the *centroid* of each cluster (the average of all points)
 4. repeat 2-3 until clusters stabilise
- Making the initial guess:
 1. Input: number of clusters to find
 2. Assign each data point to a random cluster
 3. Find the centroid of each cluster
- Careful: sometimes a cluster ends up with no points!

K-Means: basics

```
data Vector = Vector Double Double

addVector :: Vector -> Vector -> Vector
addVector (Vector a b) (Vector c d) = Vector (a+c) (b+d)

data Cluster = Cluster
    {
        clId      :: !Int,
        clCount   :: !Int,
        clSum     :: !Vector,
        clCent    :: !Vector
    }

sqDistance :: Vector -> Vector -> Double
-- square of distance between vectors

makeCluster :: Int -> [Vector] -> Cluster
-- builds Cluster from a set of points
```

K-Means:

```
assign
  :: Int          -- number of clusters
  -> [Cluster]    -- clusters
  -> [Vector]     -- points
  -> Array Int [Vector] -- points assigned to clusters

makeNewClusters :: Array Int [Vector] -> [Cluster]
  -- takes result of assign, produces new clusters

step :: Int -> [Cluster] -> [Vector] -> [Cluster]
step nclusters clusters points =
  makeNewClusters (assign nclusters clusters points)
```

- assign is step 2
- makeNewClusters is step 3
- step is (2,3) – one iteration

Putting it together.. sequentially

```
kmeans_seq :: Int -> [Vector] -> [Cluster] -> IO [Cluster]
kmeans_seq nclusters points clusters = do
  let
    loop :: Int -> [Cluster] -> IO [Cluster]
    loop n clusters | n > tooMany = return clusters
    loop n clusters = do
      hPrintf stderr "iteration %d\n" n
      hPutStr stderr (unlines (map show clusters))
      let clusters' = step nclusters clusters points
      if clusters' == clusters
        then return clusters
        else loop (n+1) clusters'
  --
  loop 0 clusters
```


Parallelise makeNewClusters?

```
makeNewClusters :: Array Int [Vector] -> [Cluster]
makeNewClusters arr =
  filter ((>0) . clCount) $
    [ makeCluster i ps | (i,ps) <- assocs arr ]
```

- essentially a map over the clusters
- number of clusters is small
- not enough parallelism here – grains are too large, fan-out is too small

How to parallelise?

- Parallelise assign?

```
assign :: Int -> [Cluster] -> [Vector] -> Array Int [Vector]
assign nclusters clusters points =
  accumArray (flip (:)) [] (0, nclusters-1)
    [ (clId (nearest p), p) | p <- points ]
  where
    nearest p = ...
```

- essentially map/reduce: map nearest + accumArray
- the map parallelises, but accumArray doesn't
- could divide into chunks... but is there a better way?

Sub-divide the data

- Suppose we divided the data set in two, and called **step** on each half
- We need a way to combine the results:

```
step n cs (as ++ bs) == step n cs as `combine` step n cs bs
```

- but what is **combine**?

```
combine :: [Cluster] -> [Cluster] -> [Cluster]
```

- assuming we can match up cluster pairs, we just need a way to combine two clusters

Combining clusters

- A cluster is notionally a set of points
- Its *centroid* is the average of the points
- A Cluster is represented by its centroid:

```
data Cluster = Cluster
    {
        clId      :: !Int,
        clCount   :: !Int,      -- num of points
        clSum     :: !Vector,   -- sum of points
        clCent    :: !Vector    -- clSum / clCount
    }
```

- but note that we cached clCount and clSum
- these let us merge two clusters and recompute the centroid in $O(1)$

Combining clusters

- So using

```
combineClusters :: Cluster -> Cluster -> Cluster
```

- we can define

```
reduce :: Int -> [[Cluster]] -> [Cluster]
```

- (see notes for the code; straightforward)
- now we can express K-Means as a map/reduce

Final parallel implementation

```
kmeans_par :: Int -> Int -> [Vector] -> [Cluster] -> IO [Cluster]
kmeans_par chunks nclusters points clusters = do
  let chunks = split chunks points
  let
    loop :: Int -> [Cluster] -> IO [Cluster]
    loop n clusters | n > tooMany = return clusters
    loop n clusters = do
      hPrintf stderr "iteration %d\n" n
      hPutStr stderr (unlines (map show clusters))
      let
        new_clusterss =
          map (step nclusters clusters) chunks
          `using` parList rdeepseq

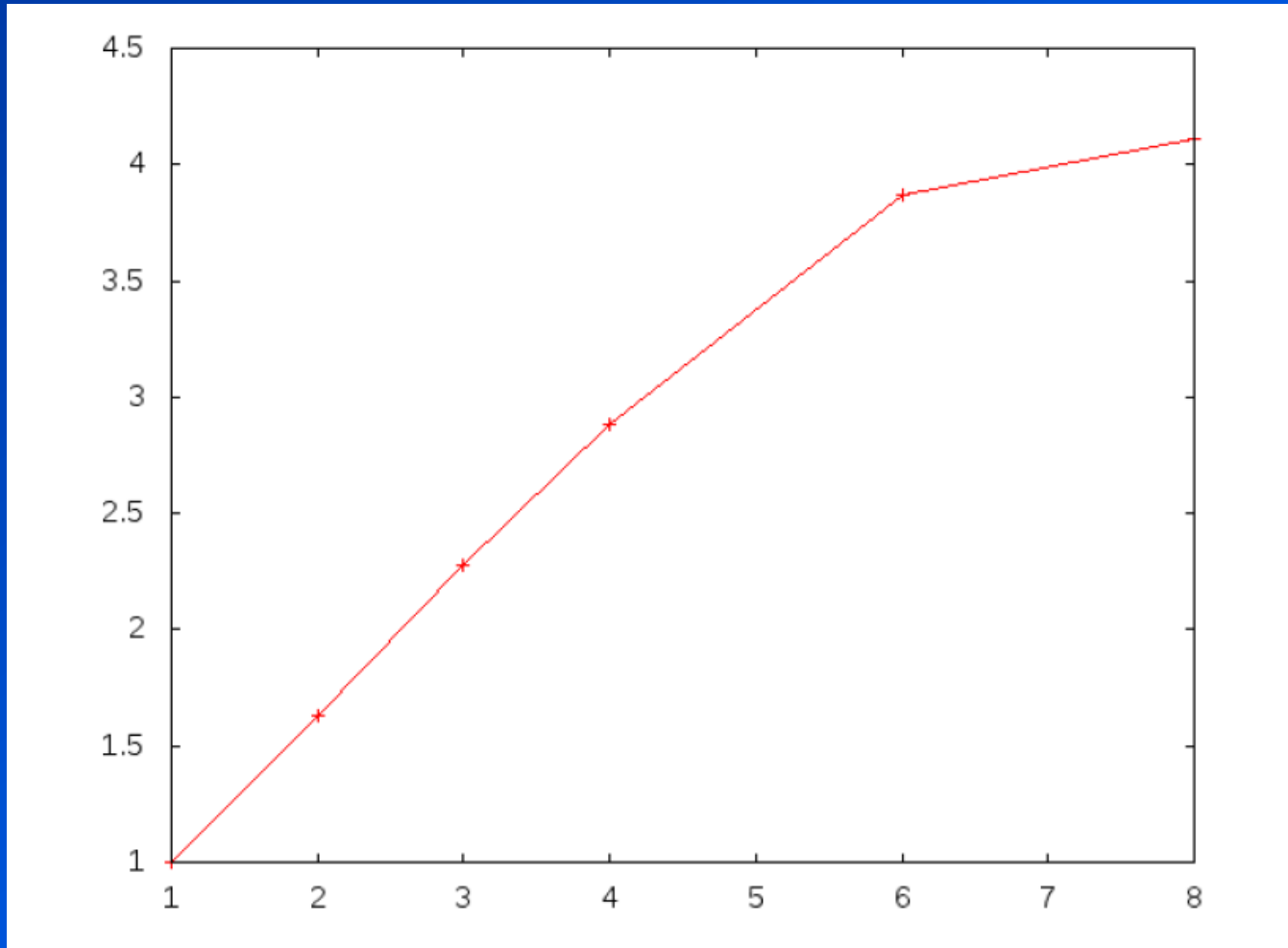
        clusters' = reduce nclusters new_clusterss

      if clusters' == clusters
      then return clusters
      else loop (n+1) clusters'
  --
  loop 0 clusters
```

What chunk size?

- Divide data by number of processors?
 - No! Static partitioning could lead to poor utilisation (see earlier)
 - there's no need to have such large chunks, the RTS will schedule smaller work items across the available cores

- Results for 170000 2-D points, 4 clusters, 1000 chunks



Further thoughts

- We had to restructure the algorithm to make the maximum amount of parallelism available
 - map/reduce
 - move the branching point to the top
 - make reduce as cheap as possible
 - a tree of reducers is also possible
- Note that the parallel algorithm is data-local – this makes it particularly suitable for distributed parallelism (indeed K-Means is commonly used as an example of distributed parallelism).
- But be careful of static partitioning

State of play

- yesterday we:
 - looked at the Eval monad, rpar and rseq, and Strategies
 - got confused about laziness
- This morning:
 - short intro to another programming model for parallelism in Haskell, the Par monad
 - Lab session (Parallel Haskell)
- This afternoon:
 - Concurrent Haskell

- Strategies, in theory:
 - *Algorithm + Strategy = Parallelism*
- Strategies, in practice (sometimes):
 - *Algorithm + Strategy = No Parallelism*
- lazy evaluation is the magic ingredient that bestows modularity, but lazy evaluation can be tricky to deal with.
- The Par monad:
 - abandon modularity via lazy evaluation
 - get a more direct programming model
 - avoid some common pitfalls
 - modularity via higher-order skeletons
 - a beautiful implementation

A menu of ways to screw up

- less than 100% utilisation
 - parallelism was not created, or was discarded
 - algorithm not fully parallelised – residual sequential computation
 - uneven work loads
 - poor scheduling
 - communication latency
- extra overhead in the parallel version
 - overheads from rpar, work-stealing, deep, ...
 - lack of locality, cache effects...
 - larger memory requirements leads to GC overhead
 - GC synchronisation
 - duplicating work

The **Par** Monad

```
data Par  
instance Monad Par
```

Par is a monad for
parallel computation

```
runPar :: Par a -> a
```

Parallel computations
are pure (and hence
deterministic)

```
fork :: Par () -> Par ()
```

forking is *explicit*

```
data IVar
```

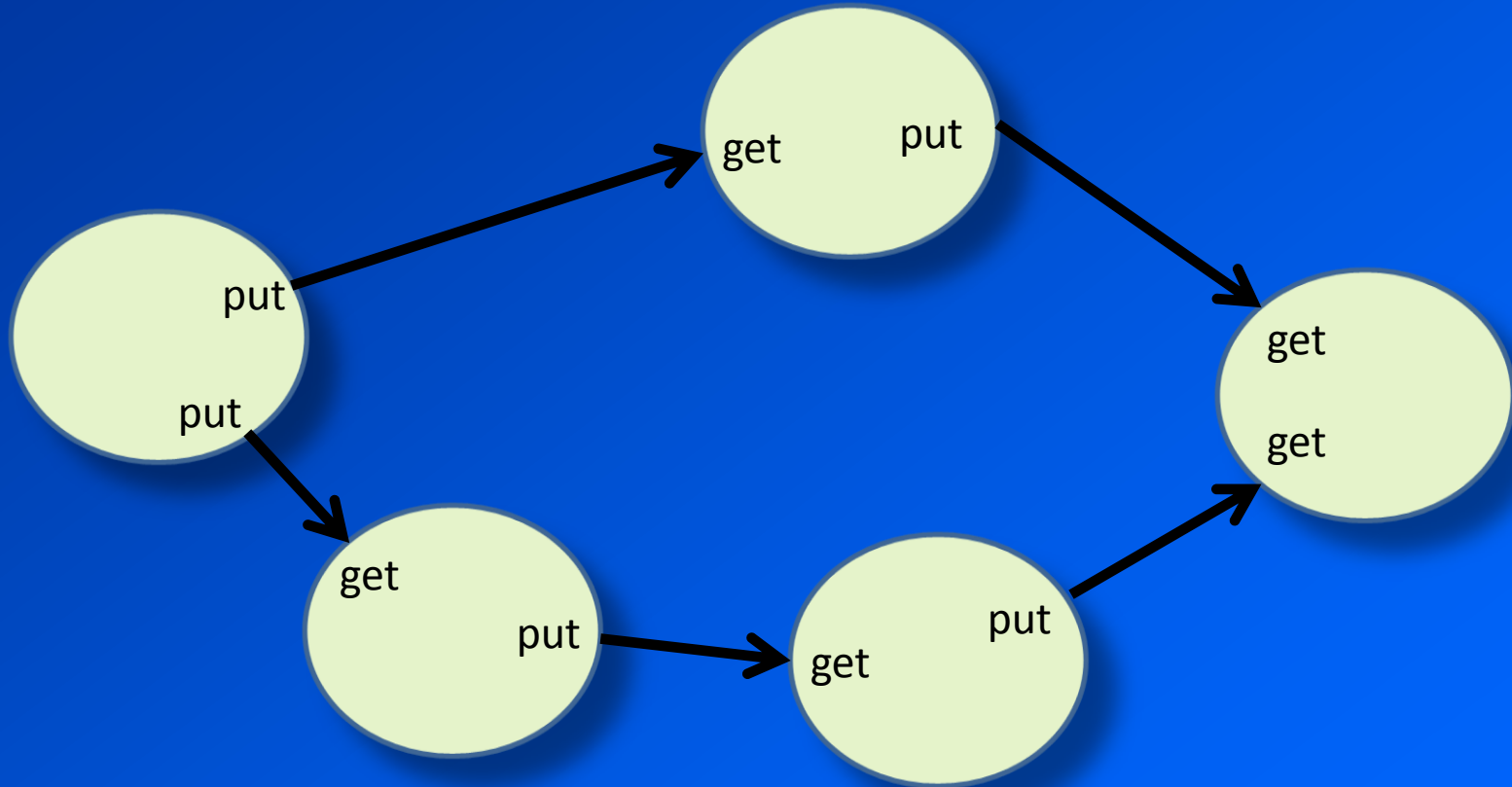
```
new :: Par (IVar a)
```

```
get :: IVar a -> Par a
```

```
put :: NFData a => IVar a -> a -> Par ()
```

results are communicated
through IVars

Par expresses dynamic dataflow



Examples

- Par can express regular parallelism, like **parMap**. First expand our vocabulary a bit:

```
spawn :: Par a -> Par (IVar a)
spawn p = do r <- new
            fork $ p >>= put r
            return r
```

- now define **parMap** (actually parMapM):

```
parMapM :: NFData b => (a -> Par b) -> [a] -> Par [b]
parMapM f as = do
  ibs <- mapM (spawn . f) as
  mapM get ibs
```

Examples

- Divide and conquer parallelism:

```
parfib :: Int -> Int -> Par Int
parfib n
  | n <= 2      = return 1
  | otherwise = do
    x <- spawn $ parfib (n-1)
    y <- spawn $ parfib (n-2)
    x' <- get x
    y' <- get y
    return (x' + y')
```

- In practice you want to use the sequential version when the grain size gets too small

How did we avoid laziness?

- put is hyperstrict.
- (by default)
- there's also a WHNF version called `put_`

Dataflow problems

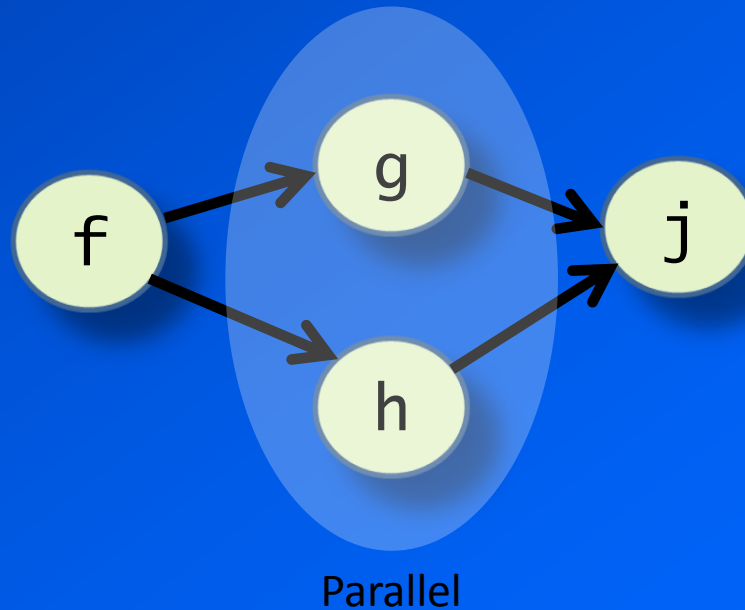
- Par really shines when the problem is easily expressed as a dataflow graph, particularly an irregular or dynamic graph (e.g. shape depends on the program input)
- Identify the nodes and edges of the graph
 - each node is created by fork
 - each edge is an IVar

Example

- Consider typechecking (or inferring types for) a set of non-recursive bindings.
- Each binding is of the form $x = e$ for variable x , expression e
- To typecheck a binding:
 - input: the types of the identifiers mentioned in e
 - output: the type of x
- So this is a dataflow graph
 - a node represents the typechecking of a binding
 - the types of identifiers flow down the edges

Example

```
f = ...  
g = ... f ...  
h = ... f ...  
j = ... g ... h ...
```



Implementation

- We parallelised an existing type checker (nofib/infer).
- Algorithm works on a single term:

```
data Term = Let VarId Term Term | ...
```

- So we parallelise checking of the top-level Let bindings.

The parallel type inferencer

- Given:

```
inferTopRhs :: Env -> Term -> PolyType  
makeEnv    :: [(VarId,Type)] -> Env
```

- We need a type environment:

```
type TopEnv = Map VarId (IVar PolyType)
```

- The top-level inferencer has the following type:

```
inferTop :: TopEnv -> Term -> Par MonoType
```

Parallel type inference

```
inferTop :: TopEnv -> Term -> Par MonoType
inferTop topenv (Let x u v) = do
  vu <- new

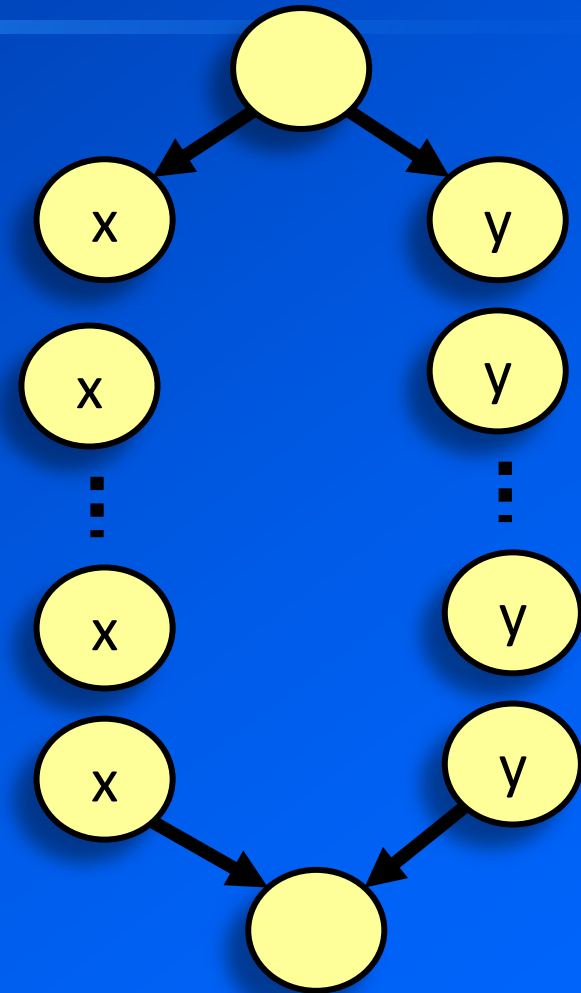
  fork $ do
    let fu = Set.toList (freeVars u)
    tfu <- mapM (get . fromJust . flip Map.lookup topenv) fu
    let aa = makeEnv (zip fu tfu)
    put vu (inferTopRhs aa u)

  inferTop (Map.insert x vu topenv) v

inferTop topenv t = do
  -- the boring case: invoke the normal sequential
  -- type inference engine
```

Results

```
let id = \x.x in
  let x = \f.f id id in
  let x = \f . f x x in
  let x = \f . f x x in
  let x = \f . f x x in
  ...
  let x = let f = x in \z . z in
  let y = \f.f id id in
  let y = \f . f y y in
  let y = \f . f y y in
  let y = \f . f y y in
  ...
  let x = let f = y in \z . z in
  \f. let g = \a. a x y in f
```



- -N1: 1.12s
- -N2: 0.60s (1.87x speedup)
- available parallelism depends on the input: these bindings only have two branches

Thoughts to take away...

- *Parallelism is not the goal*
 - Making your program faster is the goal
 - (unlike Concurrency, which is a goal in itself)
 - If you can make your program fast enough without parallelism, all well and good
 - However, designing your code with parallelism in mind should ensure that it can ride Moore's law a bit longer
 - maps and trees, not folds

Lab

- Download the sample code here:

```
http://community.haskell.org/~simonmar/par-tutorial.tar.gz
```

- or get it with git:

```
git clone https://github.com/simonmar/par-tutorial.git
```

- code is in par-tutorial/code
- lab exercises are here:

```
http://community.haskell.org/~simonmar/lab-exercises.pdf
```

- install extra packages:

```
cabal install xml utf8-string
```

Open research problems?

- How to do safe nondeterminism
- Par monad:
 - implement and compare scheduling algorithms
 - better raw performance (integrate more deeply with the RTS)
- Strategies:
 - ways to ensure identity safety
 - generic clustering