

50001 - Algorithm Analysis and Design - Lecture 1

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12/11/21

Lecture Recording

Lecture recording is available here

Introduction

An algorithm is a method of computing a result for a given problem, at its core in a systematic/-mathematical means.

This course extensively uses haskell instead of pseudocode to express problems, though its lessons still apply to other languages.

Fundamentals

Insertion Problem

Given an integer x and a sorted list ys , produce a list containing $x : ys$ that is ordered.

Note that this can be solved by simply using $sort(x : ys)$ however this is considered wasteful as it does not exploit the fact that ys is already sorted.

An example algorithm would be to traverse ys until we find a suitable place for x

```
1 insert :: Int -> [Int] -> [Int]
2 insert x [] = [x]
3 insert x yss@(y:ys)
4   | x <= y    = x:yss
5   | otherwise = y : insert x ys
```

Call Steps

In order to determine the complexity of the function, we use a **cost model** and determine what steps must be taken.

For example for $insert\ 4\ [1, 3, 6, 7]$

$insert\ 4\ [1, 3, 6, 7]$

$\rightsquigarrow \{ \text{definition of } insert \}$

$1 : insert\ 4\ [3, 6, 7]$

$\rightsquigarrow \{ \text{definition of } insert \}$ Hence this requires 3 call steps.

$1 : 3 : insert\ 4\ [6, 7]$

$\rightsquigarrow \{ \text{definition of } insert \}$

$1 : 3 :: 4 : [6, 7]$

We can use recurrence relations to get a generalised formula for the worst case (maximum number of calls):

$$\begin{aligned} T_{insert} 0 &= 1 \\ T_{insert} 1 &= 1 + T_{insert}(n-1) \end{aligned}$$

We can solve the recurrence:

$$\begin{aligned}
T_{insert}n &= 1 + T_{insert}(n-1) \\
T_{insert}n &= 1 + 1 + T_{insert}(n-2) \\
T_{insert}n &= 1 + 1 + \dots + 1 + T_{insert}(n-n) \\
T_{insert}n &= n + T_{insert}(0) \\
T_{insert}n &= n + 1
\end{aligned}$$

More complex algorithms

```

1 isort :: [Int] -> [Int]
2 isort [] = []
3 isort (x:xs) = insert x (isort xs)

```

$$\begin{aligned}
T_{isort}0 &= 1 \\
T_{isort}n &= 1 + T_{insert}(n-1) + T_{isort}(n-1)
\end{aligned}$$

Hence by using our previous formula for *insert*

$$T_{isort}n = 1 + n + T_{isort}(n-1)$$

And by recurrence:

$$\begin{aligned}
T_{isort}n &= 1 + n + (1 + n - 1) + T_{isort}(n-2) \\
T_{isort}n &= 1 + n + (1 + n - 1) + (1 + n - 2) + \dots + T_{isort}(n-n) \\
T_{isort}n &= 1 + n + (1 + n - 1) + (1 + n - 2) + \dots + T_{isort}(0) \\
T_{isort}n &= n + n + (n-1) + (n-2) + \dots + (n-n) + 1 \\
T_{isort}n &= 1 + n + \sum_{i=0}^n i \\
T_{isort}n &= \sum_{i=0}^{n+1} i \\
T_{isort}n &= \frac{(n+1) \times (n+1)}{2}
\end{aligned}$$

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Evaluation & Cost Models

```
1 minimum :: [Int] -> Int
2 minimum = head . isort
```

When analysing the cost of **minimum** we must consider how the function is evaluated.

For example we could shortcut the once **isort** has determined the first element (the minimum) of the list.

Cost Model

A model to determine the time taken to execute a program.

The model assigns cost to different operations (e.g comparisons, calls, memory reads/writes)

A very generalised cost model assigns cost based on the number of **reductions** required to evaluate a program.

Small While Language

We can define a small language of expressions as follows:

$$e ::= x \mid k \mid f \ e_1 \dots e_n \mid \text{if } e \text{ then } e_1 \text{ else } e_2$$

where k means constant and x is the variable form.

Infix functions such as $+$, $-$, \times are written normally, and are also expressions as they can be used in the form $(+) \ e_1 \ e_2$.

There are also several primitive constants: *True*, *False*, $0, 1, 2, \dots$

List constants and operations are also primitive: $[], (:), \text{null}, \text{head}, \text{tail}$

Evaluation Order

- **Applicative Order** Strict evaluation

The leftmost, innermost reducible expression is evaluated first.

e.g for $fst(3 \times 2, 1 + 2)$
 $fst(3 \times 2, 1 + 2)$
 $\rightsquigarrow \{ \text{Definition of } \times \}$
 $fst(6, 1 + 2)$
 $\rightsquigarrow \{ \text{Definition of } + \}$
 $fst(6, 3)$
 $\rightsquigarrow \{ \text{Definition of } fst \}$
6

- **Normal Order** Lazy evaluation

The leftmost outer reducible is evaluated first. Effectively evaluating the function before its arguments.

e.g for $fst(0, 1 + 2)$
 $fst(3 \times 2, 1 + 2)$
 $\rightsquigarrow \{ \text{Definition of } fst \}$
 3×2
 $\rightsquigarrow \{ \text{Definition of } \times \}$
6

For a given program, if **applicative** and **normal** terminate, then they produce the same value in normal form.

However there are some programs where **normal** evaluation terminates, but **applicative** will not.

Applicative	Normal
$fst(0, \text{crazy nonsense})$	$fst(0, \text{crazy nonsense})$
$\rightsquigarrow \{ \text{By lack of definition for crazy nonsense} \}$	$\rightsquigarrow \{ \text{Definition of } fst \}$
<i>CRASH!</i>	0

The program may be syntactically correct, but have an error such as zero-division which will not be evaluated and hence not result in improper termination under **normal** order.

Applicative Terminates \Rightarrow Normal Terminates

Cost Model for Small While

We can evaluate a cost model for the small while language by creating a function T to assign cost to expressions.

Type	Function	Explanation
non-primitive function	$f \ a_1 \ \dots \ a_n = e$ $T(f) \ a_1 \ \dots \ a_n = T(e) + 1$	Given we have already computed all argument, the cost of the function is the cost of the expression it produces, and a single call.
primitive function	$T(f) \ x \ \dots \ x_n = 0$	Primitive functions are assumed to be free.
Variable	$T(x) = 0$	accessing variables is free.
Application	$T(f \ e_1 \ \dots \ e_n) = T(f) \ e_1 \ \dots \ e_n +$ $T(e_1) + \dots + T(e_n)$	When applying a function we must consider both its cost, and the cost of all argument expressions.
Conditional	$T(\text{if } p \text{ then } e_1 \text{ else } e_2) = T(p) +$ $\text{if } p \text{ then } T(e_1) \text{ else } T(e_2)$	Cost of condition and of the resulting expression.

Cost Model Example

Given the function:

$$\text{mul } m \ n = \text{if } m = 0 \text{ then } 0 \text{ else } n + \text{mul } (m - 1) \ n$$

Evaluate $T(\text{mul } 3 \ 100)$

(1)	$mul\ 3\ 100$	
(2)	$T(\text{if } 3 = 0 \text{ then } 0 \text{ else } 100 + mul\ (3 - 1)\ 100) + 1$	By Rule for non-primitive functions
(3)	$T(3 = 0) + T(100 + mul\ (3 - 1)\ 100) + 1$	By rule for conditionals
(4)	$0 + T(100 + mul\ (3 - 1)\ 100) + 1$	By primitive functions
(5)	$T(+)(100\ mul\ (3 - 1)\ 100) + T(100) + T(mul\ (3 - 1)\ 100) + 1$	By application rule
(6)	$0 + T(100) + T(mul\ (3 - 1)\ 100) + 1$	By rule for primitive functions
(7)	$0 + T(mul\ (3 - 1)\ 100) + 1$	By rule for constants
(8)	$T(mul)\ (3 - 1)\ 100 + T(3 - 1) + T(100) + 1$	By application rule
(9)	$T(mul)\ (3 - 1)\ 100 + T(-)3\ 1 + T(100) + 1$	By application rule
(10)	$T(mul)\ 2\ 100 + 1$	By application rule
(11)	$T(\text{if } 2 = 0 \text{ then } 0 \text{ else } 100 + mul\ (2 - 1)\ 100) + 1 + 1$	By Rule for non-primitive functions
(12)	$T(2 = 0) + T(100 + mul\ (2 - 1)\ 100) + 2$	By rule for conditionals
(13)	$0 + T(100 + mul\ (2 - 1)\ 100) + 2$	By primitive functions
(14)	$T(+)(100\ mul\ (2 - 1)\ 100) + T(100) + T(mul\ (2 - 1)\ 100) + 2$	By application rule
(15)	$0 + T(100) + T(mul\ (2 - 1)\ 100) + 2$	By rule for primitive functions
(16)	$0 + T(mul\ (2 - 1)\ 100) + 2$	By rule for constants
(17)	$T(mul)\ (2 - 1)\ 100 + T(2 - 1) + T(100) + 2$	By application rule
(18)	$T(mul)\ (2 - 1)\ 100 + T(-)2\ 1 + T(100) + 2$	By application rule
(19)	$T(mul)\ 1\ 100 + 2$	By application rule
(20)	$T(\text{if } 1 = 0 \text{ then } 0 \text{ else } 100 + mul\ (1 - 1)\ 100) + 2 + 1$	By Rule for non-primitive functions
(21)	$T(1 = 0) + T(100 + mul\ (1 - 1)\ 100) + 3$	By rule for conditionals
(22)	$0 + T(100 + mul\ (1 - 1)\ 100) + 3$	By primitive functions
(23)	$T(+)(100\ mul\ (1 - 1)\ 100) + T(100) + T(mul\ (1 - 1)\ 100) + 3$	By application rule
(24)	$0 + T(100) + T(mul\ (1 - 1)\ 100) + 3$	By rule for primitive functions
(25)	$0 + T(mul\ (1 - 1)\ 100) + 3$	By rule for constants
(26)	$T(mul)\ (1 - 1)\ 100 + T(1 - 1) + T(100) + 3$	By application rule
(27)	$T(mul)\ (1 - 1)\ 100 + T(-)2\ 1 + T(100) + 3$	By application rule
(28)	$T(mul)\ 0\ 100 + 3$	By application rule
(29)	$T(\text{if } 0 = 0 \text{ then } 0 \text{ else } 100 + mul\ (1 - 1)\ 100) + 3 + 1$	By Rule for non-primitive functions
(30)	$T(0 = 0) + T(0) + 4$	By rule for conditionals
(31)	$0 + T(0) + 4$	By rule for primitive functions
(32)	$0 + 4$	By rule for variables
(33)	4	

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Asymptotics

L-Function

A **Logarithmico-exponential** function f is:

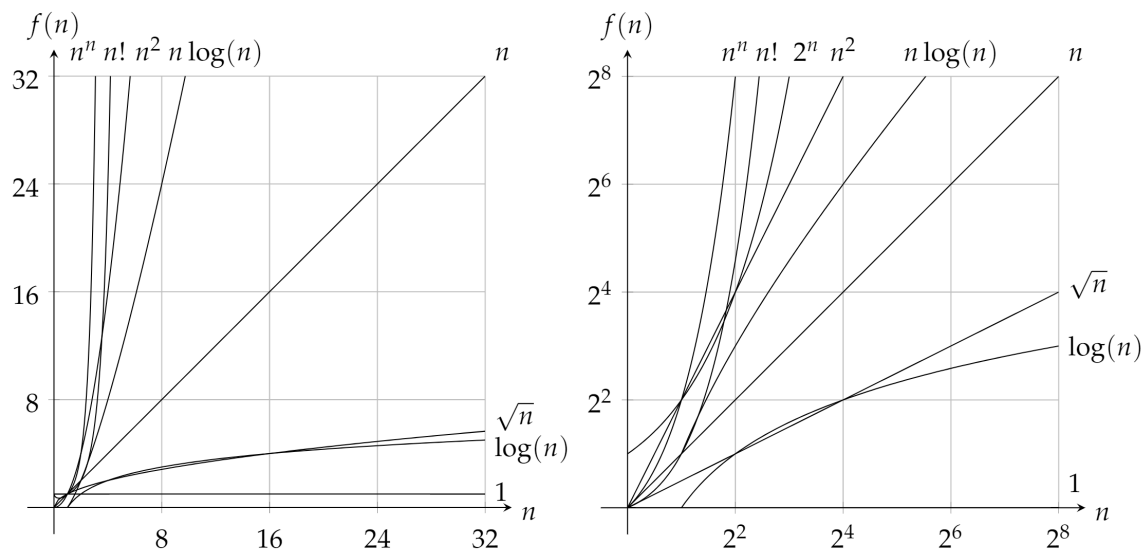
- real: $f \in X \rightarrow Y$ where $X, Y \subset \mathbb{R}$
- positive: $\forall x \in X. [f(x) \leq 0]$
- monotonic: $\forall x_1, x_2 \in X. [x_1 < x_2 \Leftrightarrow f(x_1) < f(x_2)]$ (positive monotonic) or $\forall x_1, x_2 \in X. [x_1 < x_2 \Leftrightarrow f(x_1) > f(x_2)]$ (negative monotonic)
- one valued: $\forall x \in X, y_1, y_2 \in Y. [f(x) = y_1 \wedge f(x) = y_2 \Rightarrow y_1 = y_2]$
- on a real variable defined for all values greater than some definite value: $X \equiv \{x | x > \text{definite limit} \wedge x \in \mathbb{R}\}$

L-Functions are continuous, of constant sign and as $n \rightarrow \infty$ the value $f(n)$ tends to 0, ∞ or some other positive definite limit.

Functions that aren't **L-Functions** are called **Wild Functions**.

In asymptotics we use **L-Functions** to describe the growth of time used by algorithms as the size of the input to an algorithm grows.

Common functions are shown below:



Du Bois-Reymond Theorem

Defines inequalities for the rate of increase of functions.

Where $lim = \lim_{n \rightarrow \infty} \frac{f(n)}{g(n)}$

$(<)$	$f \prec g \Leftrightarrow lim = 0$	g grows much faster than f
(\leq)	$f \preceq g \Leftrightarrow lim < \infty$	g grows much faster than f or some multiple of f
$(=)$	$f \asymp g \Leftrightarrow lim < \infty$	g grows some multiple faster than f
(\geq)	$f \succcurlyeq g \Leftrightarrow lim > 0$	f grows much faster than g or some multiple of g
$(>)$	$f \succ g \Leftrightarrow lim > \infty$	f grows much faster than g

These operators form a trichotomy such that one of the below will always hold:

$$f \prec g \quad f \asymp g \quad f \succ g$$

Further the operators \succ and \prec are converse:

$$f \succ g \Leftrightarrow g \prec f$$

And transitive:

$$\begin{aligned} f \prec g \wedge g \prec h &\Rightarrow f \prec h \\ f \preceq g \wedge g \preceq h &\Rightarrow f \preceq h \end{aligned}$$

We can place the common **L-Functions** in order:

$$1 \prec \log n \prec \sqrt{n} \prec n \prec n \log n \prec n^2 \prec n^3 \prec n! \prec n^n$$

Bachman-Landau Notation

Comparison with Bois-Reymond

Set definition

$f \in o(g(n)) \Leftrightarrow f \prec g$	$o(g(n)) = \{f \forall \delta > 0. \exists n_0 > 0. \forall n > n_0 [f(n) < \delta g(n)]\}$
$f \in O(g(n)) \Leftrightarrow f \preceq g$	$O(g(n)) = \{f \exists \delta > 0. \exists n_0 > 0. \forall n > n_0 [f(n) \leq \delta g(n)]\}$
$f \in \Theta(g(n)) \Leftrightarrow f \asymp g$	$\Theta(g(n)) = O(g(n)) \cap \Omega(g(n))$
$f \in \Omega(g(n)) \Leftrightarrow f \succcurlyeq g$	$\Omega(g(n)) = \{f \exists \delta > 0. \exists n_0 > 0. \forall n > n_0 [f(n) \geq \delta g(n)]\}$
$f \in \omega(g(n)) \Leftrightarrow f \succ g$	$\omega(g(n)) = \{f \forall \delta > 0. \exists n_0 > 0. \forall n > n_0 [f(n) > \delta g(n)]\}$

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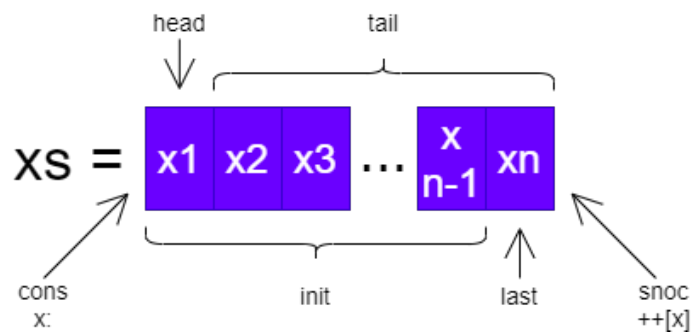
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Lists

```

1 data List [a] = [] | (:) a [a]
2
3 — or ...
4 data List a where
5   Empty :: List a
6   Cons  :: a -> List a -> List a

```



Lists in **Haskell** are a persistent data structure, meaning that when operations are applied to lists the original list is maintained (not mutated).

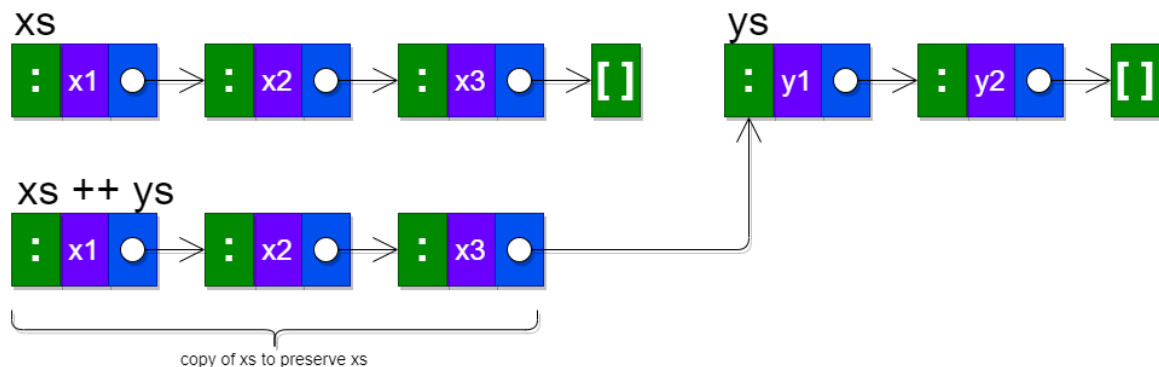
Append

We can append lists, by traversing over the first list, copying values (this ensures both argument lists are preserved).

```

1 (++) :: [a] -> [a] -> [a]
2 []    ++ ys = ys
3 (x:xs) ++ ys = x:(xs ++ ys)

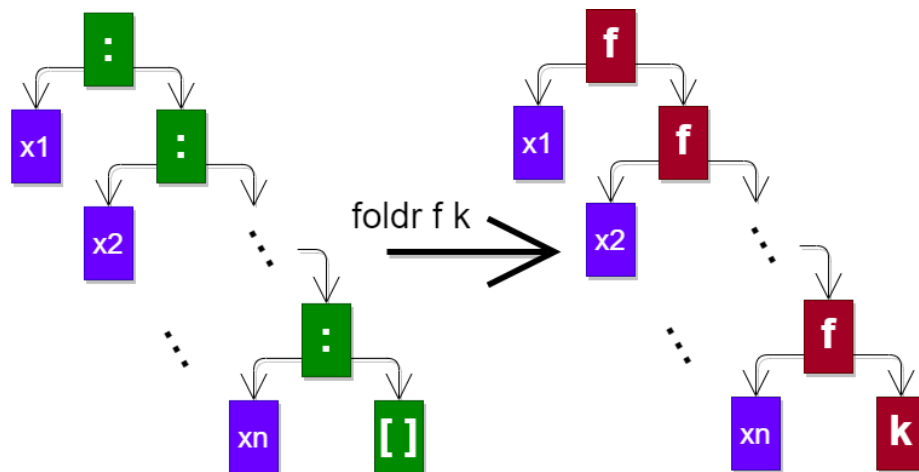
```



As the entire first list must be traversed, the cost of $xs++ys$ is $T_{(++)}(n) \in O(n)$ where $n = \text{length } xs$

Foldr

```
1 foldr :: (a -> b -> b) -> b -> [a] -> b
2 foldr f k [] = k
3 foldr f k (x:xs) = f x (foldr f k xs)
```



As you can see, $\text{foldr } (:) [] \equiv \text{id}$.

Foldr can also be expressed through bracketing

$$\text{foldr } f \ k \ [x_1, x_2, \dots, x_n] \equiv f \ x_1 \ (f \ x_2 \ (\dots (f \ x_n \ k) \dots))$$

Associativity

Associativity determines how operations are grouped in the absence of brackets.

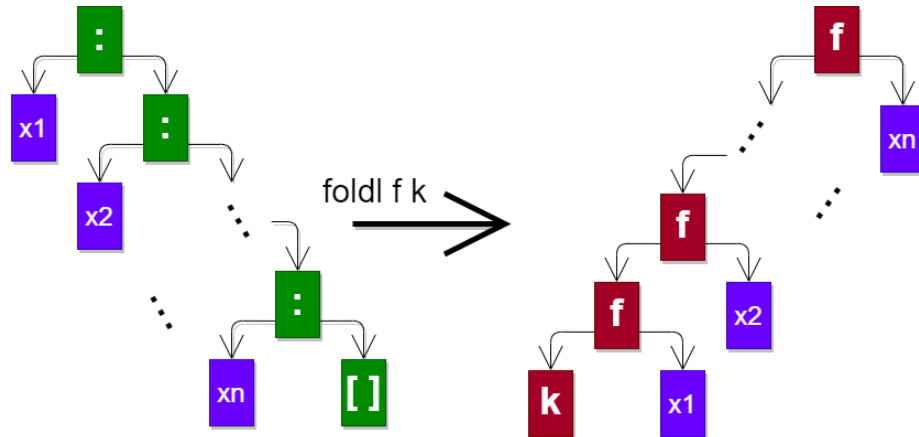
$a \spadesuit b \spadesuit c$ unbracketed statement
 $((a) \spadesuit b) \spadesuit c$ \spadesuit is left associative
 $a \spadesuit (b \spadesuit (c))$ \spadesuit is right associative

If \spadesuit is associative, then the right & left associative versions are equivalent.

foldr applies functions in a right-associative scheme.

Foldl

```
1 foldl :: (b -> a -> b) -> b -> [a] -> b
2 foldl f k [] = k
3 foldl f k (x:xs) = foldl f (f k x) xs
```



As you can see $\text{foldl} (\text{snoc}) [] \equiv \text{id}$.
Foldl can be expressed through bracketing

$$\text{foldl } f \ k \ [x_1, x_2, \dots, x_n] \equiv f (\dots (f (f \ k \ x_1) x_2) \dots x_n)$$

Monoids

Consider the case when for some \star and ϵ : $\text{foldr } \star \ \epsilon \equiv \text{foldl } \star \ \epsilon$. For this to be possible for $\star :: a \rightarrow a \rightarrow a$ and $\epsilon :: a$.

$$\begin{aligned} \star & \text{ must be associative} & x \star (y \star z) & \equiv (x \star y) \star z \\ \epsilon & \text{ must have no effect} & \epsilon \star n & = n \end{aligned}$$

These properties for a **monoid** (a, \star, ϵ) .

Other example include:

$$\begin{aligned} (\text{lists}, ++, []) & \quad (\mathbb{N}, +, 0) & (\mathbb{N}, \times, 1) & \quad (\text{bool}, \wedge, \text{true}) \\ (\text{bool}, \vee, \text{false}) & \quad (\mathbb{R}, \max, \infty) & (\mathbb{R}, \min, -\infty) & \quad (\text{Universal set}, \cup, \emptyset) \end{aligned}$$

We can also find monoids of functions:

$$(a \rightarrow a, (.), \text{id})$$

as $(\text{id} . g)x \equiv \text{id}(g \ x)$ and $((f . g) . h)x = f(g(h \ x))$

Concat

We can easily define concat recursively as:

```
1 concat :: [[a]] -> [a]
2 concat [] = []
3 concat (xs:xss) = xs ++ concat xss
```

We can also notice that $([[a]], (++) , [])$ is a monoid, so we can use **foldr** or **foldl**

```
1 concatr :: [[a]] -> [a]
2 concatr = foldr (++) []
```

```
1 concatr :: [[a]] -> [a]
2 concatr = foldl (++) []
```

as `(++)` makes a copy of the first argument (to ensure persistent data), if we apply it in a left associative bracketing scheme we will have to make larger & larger copies.

$$(\dots(((([] ++_0 xs_1) ++_m xs_2) ++_{2m} xs_3) ++_{3m} xs_4 \dots) ++_{mn} xs_n)$$

Hence where $n = \text{length } xs$ and $m = \text{length } xs_1 = \text{length } xs_2 = \dots = \text{length } xs_n$.

$$\begin{aligned} T_{concatl}(m, n) &\in O(n^2 m) \\ T_{concatr}(m, n) &\in O(nm) \end{aligned}$$

DLists

Instead of storing a list, we store a composition of functions that build up a list.

$$\begin{aligned} &xs_1 ++ xs_2 ++ xs_3 ++ \dots ++ xs_n \\ &\quad \downarrow \\ &f\ xs_1 \bullet f\ xs_2 \bullet f\ xs_3 \bullet \dots \bullet f\ xs_n \\ &\quad \downarrow \\ &(xs_1 ++) \bullet (xs_2 ++) \bullet (xs_3 ++) \bullet \dots \bullet (xs_n ++) \end{aligned}$$

We can then apply this function on the empty list `[]` to get the resulting list.

```

1  newtype DList a = DList ([a] -> [a])
2
3  instance List DList where
4      toList :: DList a -> [a]
5      toList (DList fxs) = fxs []
6
7      fromList :: [a] -> DList a
8      fromList xs = DList (xs++)
9
10     (++) :: DList a -> DList a -> DList a
11     DList fxs ++ DList fys = DList (fxs . fys)

```

We can form a **monoid** of $(DList, ++, DList\ id)$.

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DLists Continued...

Monoids (again)

A **monoid** is a triple (M, \diamond, ϵ) where \diamond is associative and of type $M \rightarrow M \rightarrow M$, and $x \diamond \epsilon \equiv x$.

```
1 class Monoid m where
2   (<>) :: m -> m -> m
3   mempty :: m
```

A haskell typeclass can then be instantiated for many other data types. For example the **monoid** $(\mathbb{Z}, +, 0)$ (note that we cannot enforce **monoid** properties through haskell, unlike languages such as **agda**).

```
1 — declaring newtype so that many monoid instance on Int do not conflict
2 newtype PlusInt = PlusInt Int
3
4 instance Monoid PlusInt where
5   (<>) :: PlusInt -> PlusInt -> PlusInt
6   (<>) = (+)
7
8   mempty :: PlusInt
9   mempty = 0
```

Likewise we can abstract Lists to a class (which we can instantiate for DLists).

List Class

```
1 class List list where
2   empty :: list a
3   single :: a -> list a
4
5   (:) :: a -> list a -> list a
6   snoc :: list a -> list a -> list a
7
8   head :: list a -> a
9   tail :: list a -> list a
10
11  last :: list a -> a
12  init :: list a -> list a
13
14  (++) :: list a -> list a -> list a
15
16  length :: list a -> Int
17
18  fromList :: [a] -> list a
19  toList :: list a -> [a]
```

$[a]$ is our abstract list type, and *lista* is our concrete type.

It is critical to ensure that $toList \bullet fromList \equiv id$

But in general $fromList \bullet toList \neq id$ (this is as the internal representation may change and much information about the internal representation cannot be preserved by `toList`, for example an unbalanced tree changed to a list maybe be balanced when converted back to a tree).

We also included $normalise :: fromList \bullet toList$ as a useful tool to reset the internal structure (for example to rebalanced the tree representation of a list)

Haskell Implementation

To prevent conflicts due to Prelude functions already being defined we can use:

```
1 import Prelude hiding(head, tail, (++), etc...)
2 import qualified Prelude
```

To help ensure correctness we can use **Quickcheck** to check properties

```
1 cabal install --lib QuickCheck
```

Then can use quickcheck to define properties we want to test:

```
1 import Test.QuickCheck
2
3 -- code to test written here ...
4
5 prop_propertyname :: InputTypes -> Bool
6 prop_propertyname = test code
7
8 -- example for normalise (takes a list type, that has equality defined for it)
9 prop_normalise (Eq a, Eq (list a), List list) => list a -> Bool
10 prop_normalise xs = (toList . fromList) xs == xs
11
12 -- Can return properties (requires show) using the triple-equals
13 prop_assoc :: (Eq (list a), Show (list a), List list)
14   => list a -> list a -> list a -> Property
15 prop_assoc xs ys zs = (xs ++ ys) ++ zs == xs ++ (ys ++ zs)
```

```
1 ghci file_to_check.hs
2 *file_to_check> quickCheck (prop_normalise :: [Int] -> Bool)
3 +++ OK, passed 100 tests.
4 *file_to_check> quickCheck (prop_normalise :: [Bool] -> Bool)
5 +++ OK, passed 100 tests.
6 *file_to_check> verboseCheck (prop_normalise :: [Bool] -> Bool)
7 Passed:
8 ...
9
10 Passed:
11 ...
12
13 etc...
14
15 +++ OK, passed 100 tests.
```

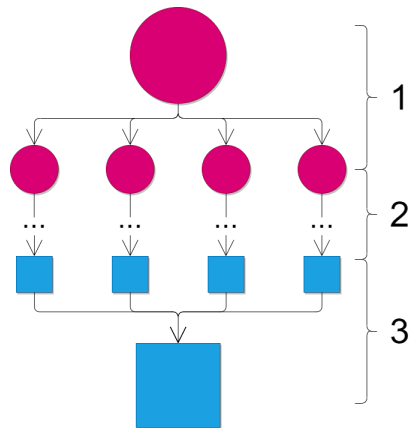
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Divide & Conquer

1. Divide a problem into subproblems
2. Divide and conquer subproblems into subsolutions
3. Conquer subsolutions into a solution



Merge Sort

```

1 msort :: Ord a => [a] -> [a]
2 msort [] = []
3 msort [x] = [x]
4 msort xs = merge (msort us) (msort vs)
5   where (us, vs) = splitAt (length xs `div` 2) xs
6
7 merge :: Ord a => [a] -> [a] -> [a]
8 merge [] ys = ys
9 merge xs [] = xs
10 merge xss@(x:xs) yss@(y:ys)
11   | x <= y    = x : merge xs yss
12   | otherwise = y : merge xss ys

```

SplitAt divides, and **merge** Conquers. We can calculate the time complexity for the recurrence relations below (based on recursive structure of **msort**):

$$T_{msort}(0) = 1$$

$$T_{msort}(1) = 1$$

$$T_{msort}(n) = T_{length}(n) + T_{splitAt}\left(\frac{n}{2}\right) + T_{merge}\left(\frac{n}{2}\right) + 2 \times T_{msort}\left(\frac{n}{2}\right)$$

We can simplify the complexity of **msort**

$$\begin{aligned}
 T_{msort}(n) &= T_{length}(n) + T_{splitAt}\left(\frac{n}{2}\right) + T_{merge}\left(\frac{n}{2}\right) + 2 \times T_{msort}\left(\frac{n}{2}\right) \\
 &= n + \frac{n}{2} + \frac{n}{2} + \frac{n}{2} + 2 \times T_{msort}\left(\frac{n}{2}\right) \\
 &= \frac{5}{2} \times n + 2 \times T_{msort}\left(\frac{n}{2}\right)
 \end{aligned}$$

Master Theorem

For an algorithm *algo* such that:

$$T_{algo}(n) = a \times T_{algo}\left(\frac{n}{b}\right) + f(n) + \text{base cases}$$

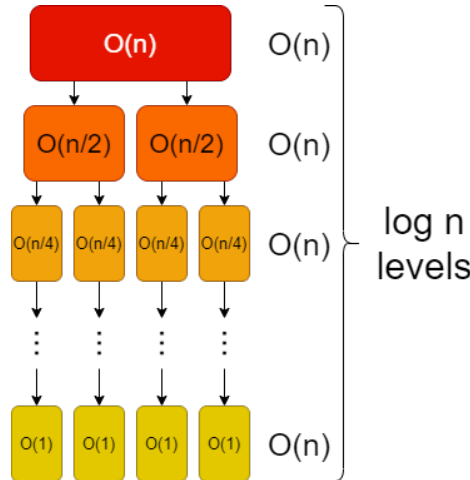
The work at recursion level $\log_b n$ is $\Theta(a^{\log_b n})$ To calculate the order of the time complexity:

1. Get the recurrence relation in the form above.
2. Get the critical exponent E by formula: $E = \log_b a = \frac{\log a}{\log b}$.
3. Given $f(n) = n^c$ we can express the work as a geometric sum $\sum_{i=0}^{\log_b n} ar^i$ where $r = \frac{a}{b^c}$.

$$r > 1 \Leftrightarrow a > b^c \Leftrightarrow \log_b a > c \Leftrightarrow E > c$$

$$\begin{aligned}
 E < c & \quad T_{algo}(n) \in \Theta(f(n)) \\
 E = c & \quad T_{algo}(n) \in \Theta(f(n) \log_b n) = \Theta(f(n) \log n) \\
 E > c & \quad T_{algo}(n) \in \Theta(a^{\log_b n}) = \Theta(n^{\log_b a}) = \Theta(n^E)
 \end{aligned}$$

By master theorem we can easily see $T_{msort}(n) \in \Theta(n \log n)$. We can also calculate it using a graph:



Quicksort

```

1 qsort :: Ord a => [a] -> [a]
2 qsort [] = []
3 qsort [x] = [x]
4 qsort (x:xs) = qsort us ++ x:qsort vs
5   where (us, vs) = partition (<x) xs
6
7 partition :: (a -> Bool) -> [a] -> ([a],[a])
8 partition p xs = (filter p xs, filter (not . p) xs)

```

Note for simplicity, we assume the lists are split into equal parts.

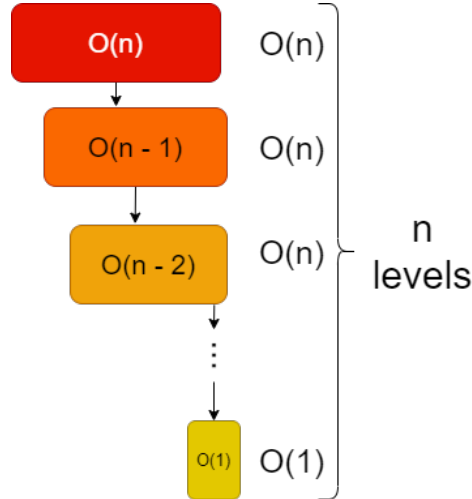
$$T_{qsort}(0) = 1$$

$$T_{qsort}(1) = 1$$

$$T_{qsort}(n) = T_{partition}(n-1) + T_{++}\left(\frac{n}{2}\right) + 2 \times T_{qsort}\left(\frac{n}{2}\right)$$

In the worst case, the partition splits xs into $(xs, [])$, we have complexity:

$$\begin{aligned}
 T_{qsort}(n) &= T_{partition}(n-1) + T_{++}(n-1) + T_{qsort}(0) + T_{qsort}(n-1) \\
 &= 2(n-1) + n + 1 + T_{qsort}(n-1)
 \end{aligned}$$



We can once again use master theorem, or a diagram such as below to see the complexity: Hence in the worst case $T_{qsort}(n) \in O(n^2)$ (same as insertion sort).

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Dynamic Programming

A technique to efficiently calculate solutions to certain recursive problems.

1. Describe an inefficient recursive algorithm.
2. Reduce inefficiency by storing intermediate shared results.

Fibonacci Sequence

Fully Recursive

```

1 fib :: Int -> Integer
2 fib 0 = 0
3 fib 1 = 1
4 fib n = fib (n-1) + fib (n-2)

```

$$T_{fib}(0) = 1$$

$$T_{fib}(1) = 1$$

$$T_{fib}(n) = 1 + T_{fib}(n-2) + T_{fib}(n-1)$$

The complexity of this algorithm is $T_{fib}(n) \in O(2^n)$.

Saving Intermediate Results

We can use a helper function which takes the remaining number of additions, and the two previous values.

```

1 fib :: Int -> Integer
2 fib n = fibHelper n 0 1
3   where
4     fibHelper :: Int -> Integer -> Integer -> Integer
5     fibHelper 0 x y = x
6     fibHelper n x y = fibHelper (n-1) y (x + y)

```

$$T_{fib}(0) = 1$$

$$T_{fib}(1) = 1$$

$$T_{fib}(n) = 1 + T_{fib}(n-1)$$

The complexity of this algorithm is $T_{fib}(n) \in O(n)$.

This way every value is calculated only once for each call. However values are not saved between calls.

Memoisation

```

1 fibs :: [Integer]
2 fibs = 0 : 1 : zipWith (+) fibs (tail fibs)
3
4 fib :: Int -> Integer
5 fib n = fibs !! n

```

This creates a large list, we must only index on the list to get the value. By using an array we can reduce the time taken to get to the n th element.

Array Based Memoisation

```

1 — Array Data Type
2 import Data.Array ( Ix(range), Array, array )
3
4 — Tabulate function
5 tabulate :: Ix i => (i,i) -> (i -> e) -> Array i e
6 tabulate (a,b) f = array (a,b) [(i,f i) | i <- range (a,b)]
7
8 — Ix (class of all indexes)
9
10 — T(!) is in O(1)
11 (!) :: Ix i => Array i e -> i -> e
12
13 — Range creates a list of indexes in a range
14 range :: Ix i => (i,i) -> [i]
15
16 — Array function creates an array from a range of indexes & values
17 array :: Ix i => (i,i) -> [(i,e)] -> Array i e

```

Hence we can make our algorithm:

```

1 import Data.Array ( Array, (!) )
2
3 fib :: Int -> Integer
4 fib n = table ! n
5 where
6     table :: Array Int Integer
7     table = tabulate (0,n) memo
8
9     memo :: Int -> Integer
10    memo 0 = 0
11    memo 1 = 1
12    memo n = table ! (n-2) + table ! (n-1)

```

Here we can do constant time lookups for values in the table. If a value is not present, it is lazily evaluated using other elements in the table.

In this way we only calculate each fibonacci number once, and only when we need it. Further it is saved for any subsequent calls to fib.

Edit-Distance

The **Edit-Distance** Problem is concerned with calculating the **Levenshtein** distance between two strings.

Levenshtein Distance

The number of insertions, deletions & updates required to convert one string into another.

$toil \rightarrow_{+1} oil \rightarrow_{+1} il \rightarrow_{+1} ill$

```
1 dist :: String -> String -> Int
2 dist xs [] = length xs
3 dist [] ys = length ys
4 dist xxs@(x:xs) yys@(y:ys)
5   = minimum
6     [dist xxs ys + 1,
7      dist xs yys + 1,
8      dist xs ys + if x == y then 0 else 1]
```

This problem becomes of order $O(3^n)$ as it recurs 3 ways for each call.

We can reuse results for two substrings through memoisation, first we make a new recursive version that uses the index we are checking in each string:

```
1 dist :: String -> String -> Int
2 dist xs ys = dist' xs ys (length xs) (length ys)
3
4 dist' :: String -> String -> Int -> Int -> Int
5 dist' xs ys i 0 = i
6 dist' xs ys 0 j = j
7 dist' xs ys i j
8   = minimum [dist' xs ys i (j-1) + 1,
9              dist' xs ys (i-1) j + 1,
10             dist' xs ys (i-1) (j-1) + if x == y then 0 else 1]
11 where
12   m = length xs
13   n = length ys
14   x = xs !! (m-i)
15   y = ys !! (n-j)
```

We can then use **tabulate** to create a memoised version.

```
1 import Data.Array ( Array, (!) )
2
3 dist :: String -> String -> Int
4 dist xs ys = table ! (m,n)
5   where
6     table = tabulate ((0,0),(m,n)) (uncurry memo)
7
8     memo :: Int -> Int -> Int
9     memo i 0 = i
10    memo 0 j = j
11    memo i j
12      = minimum [table ! (i, j - 1) + 1,
13                 table ! (i - 1, j) + 1,
14                 table ! (i - 1, j - 1) + if x == y then 0 else 1]
15    where
16      x = xs !! (m - i)
17      y = ys !! (n - j)
18
19    m = length xs
20    n = length ys
```

```
21     axs, ays :: Array Int Char
22     axs = fromList xs
23     ays = fromList ys
```

As there are at most $m \times n$ entries in the table, and each are calculated at most once, and the lookup time is constant (using arrays), the complexity is $O(mn)$.

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Amortized Analysis

So far we have studied complexity of a single, isolated run of an algorithm. **Amortized Analysis** is about understanding cost in a wider context (e.g averaged over many calls to a routine).

Dequeues

Dequeue

A Dequeue is a double ended queue. An abstract datatype that generalises a queue. Elements can be added or removed from either end.

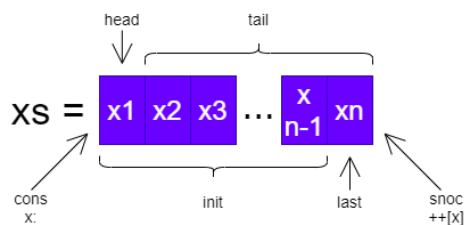
Common associated functions are:

snoc	Insert element at the back of the queue.
cons	Insert element at the front of the queue.
eject	Remove last element.
pop	remove first element.
peek	Examine but do not remove first element.

Dequeues are also called head-tail linked lists or symmetric lists.

we use a dequeue when we want to reduce the time taken to perform certain operations.

List Operation Complexity

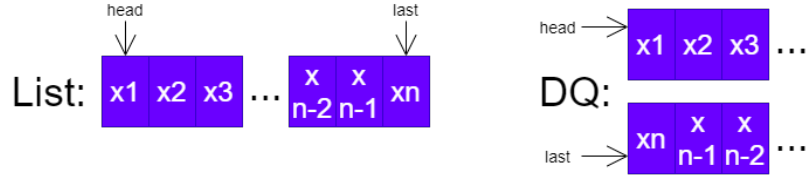


cons	$O(1)$
head	$O(1)$
tail	$O(1)$

snoc	$O(1)$
last	$O(1)$
init	$O(1)$

Dequeue Structure

To achieve $O(1)$ complexity in the **snoc**, **init** and **last** we use two lists.



One list starts contains the start of the list, and the other the end (reversed).

We keep two invariants for *Dequeue us sv*:

$$null\ us \Rightarrow null\ sv \vee single\ sv$$

$$null\ sv \Rightarrow null\ us \vee single\ us$$

In other words, if one list is empty, the other can contain at most 1 element.

An example implementation in haskell is below:

```

1  — can ignore certain patterns due to invariant
2  {-# OPTIONS.GHC -Wno-incomplete-patterns #-}
3
4  data Dequeue a = Dequeue [a] [a]
5
6  instance List Dequeue where
7    toList :: Dequeue a -> [a]
8    toList (Dequeue us sv) = us ++ reverse sv
9
10   fromList :: [a] -> Dequeue a
11   fromList xs = Dequeue us (reverse vs)
12     where (us, vs) = splitAt (length xs `div` 2) xs
13
14   — use the invariant, if [] sv then sv = [x] or []
15
16   — O(1)
17   cons :: a -> Dequeue a -> Dequeue a
18   cons x (Dequeue us []) = Dequeue [x] us
19   cons x (Dequeue us sv) = Dequeue (x:us) sv
20
21   — O(1)
22   snoc :: Dequeue a -> a -> Dequeue a
23   snoc (Dequeue [] sv) x = Dequeue sv [x]
24   snoc (Dequeue us sv) x = Dequeue us (x:sv)
25
26   — O(1)
27   last :: Dequeue a -> a
28   last (Dequeue _ (s:_)) = s
29   last (Dequeue [u] _) = u
30   last (Dequeue [] []) = error "Nothing in the dequeue"
31
32   — O(1)
33   head :: Dequeue a -> a
34   head (Dequeue (u:_ _) _) = u
35   head (Dequeue [] [v]) = v
36   head (Dequeue [] []) = error "Nothing in the dequeue"
37
38   — O(1)
39   tail :: Dequeue a -> Dequeue a

```

```

40 tail (Dequeue [] []) = error "Nothing in the dequeue"
41 tail (Dequeue [] _) = empty
42 tail (Dequeue _ []) = empty
43 tail (Dequeue [u] sv) = Dequeue (reverse su)
44   where
45     n = length sv
46     (su, sv') = splitAt (n `div` 2) sv
47     — note: could also do a fromList (reverse sv) but less efficient
48
49 tail (Dequeue (u:us) sv) = Dequeue us sv
50
51 — O(1)
52 init :: Dequeue a -> Dequeue a
53 init (Dequeue [] []) = error "Nothing in the dequeue"
54 init (Dequeue [] _) = empty
55 init (Dequeue _ []) = empty
56 init (Dequeue us [s]) = fromList us
57 init (Dequeue us (s:sv)) = Dequeue us sv
58
59 isEmpty :: Dequeue a -> Bool
60 isEmpty (Dequeue [] []) = True
61 isEmpty _ = False
62
63 empty :: Dequeue a
64 empty = Dequeue [] []

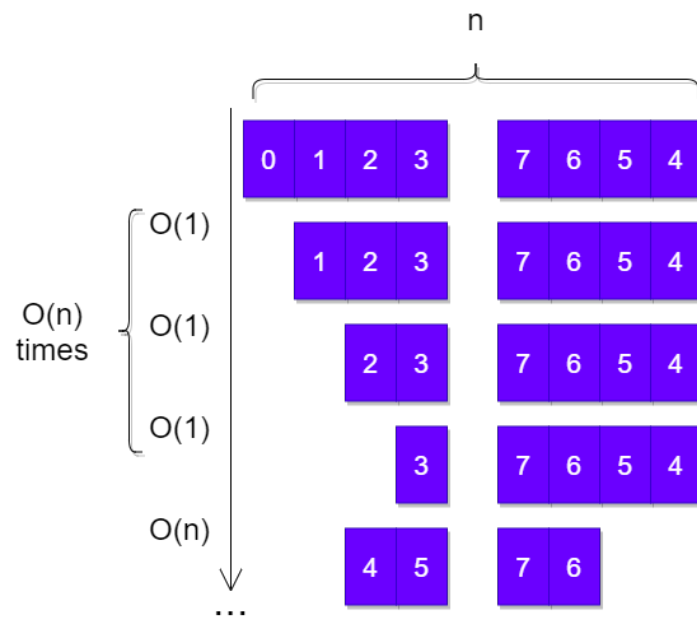
```

When considering the cost of **tail** and **init** we must consider that there are two possibilities:

High Cost	$init(Dequeue\ us[s])$ $tail(Dequeue\ [u]sv)$	This operation is $O(n)$ complexity due to the splitAt and reverse operation done on half of a list.
Low Cost	$init(Dequeue\ us(s:sv))$ $tail(Dequeue\ (u:us)sv)$	Low cost $O(1)$ operation as it requires only a pattern match on the first element.

As both of these operations rebalance the **Dequeue** to to be balanced (half the queue on each list), we these operations can have an amortized cost of $O(1)$.

We know this as the average cost is of order $O(1)$. The $O(n)$ cost is incurred every $n/2$ calls to **tail/init**.



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Amortization

The complexity of **tail** is an example of **Amortized analysis**, where operation's wider context are considered when calculating the complexity.

$$xs_0 \xrightarrow{op_0} xs_1 \xrightarrow{op_1} xs_2 \xrightarrow{op_2} xs_3 \xrightarrow{op_3} \dots \xrightarrow{op_{n-1}} xs_n$$

We defined 3 parts:

1. **Cost Function**

$C_{op_i}(xs_i)$ determines the cost of operation op_i on data xs_i . Estimating how many steps it takes for each operation to execute.

2. **Amortized Cost Function**

$A_{op_i}(xs_i)$ for each operation op_i on data xs_i .

3. **Size Function**

$S(xs)$ that calculates the size of data xs

We define these functions with the goal to show that:

$$C_{op_i}(xs_i) \leq A_{op_i}(xs_i) + S(xs_i) - S(xs_{i+1})$$

The cost of the operation is smaller than the amortized cost, plus the difference in size of the data structure before and after the operation.

Once this is shown, we can infer that:

$$\sum_{i=0}^{n-1} C_{op_i}(xs_i) \leq \sum_{i=0}^{n-1} A_{op_i}(xs_i) + S(xs_0) - S(xs_n)$$

Furthermore when $S(xs_0) = 0$ this implies:

$$\sum_{i=0}^{n-1} C_{op_i}(xs_i) \leq \sum_{i=0}^{n-1} A_{op_i}(xs_i) - S(xs_n) \Rightarrow \sum_{i=0}^{n-1} C_{op_i}(xs_i) \leq \sum_{i=0}^{n-1} A_{op_i}(xs_i)$$

This means the cost of the operations is less than the sum of the amortized costs.

For example, if $A_{op_i}(xs) = 1$ then the total cost will be bounded by $O(n)$.

Tail example

$$C_{cons}(xs) = 1 \quad C_{snoc}(xs) = 1 \quad C_{head}(xs) = 1 \quad C_{last}(xs) = 1$$

For tail we can do the following:

- | | | |
|-----|--|---|
| (1) | $C_{tail}(Dequeue\ us\ sv) = length\ sv$ | (Create a cost function of tail .) |
| (2) | $A_{op}(xs) = 2$ | (Create an arbitrary cost function.) |
| (3) | $S(Dequeue\ us\ sv) = length\ us - length\ sv $ | (Create a size function for dequeue .) |
| (4) | $Dequeue\ us\ sv$ where $length\ sv = k$ | (Worst case where us is a singleton) |
| (5) | $S(Dequeue\ us\ sv) = k - 1$
$S(Dequeue\ us'\ sv') = 1$ | (Size of the next data structure can be at most 1.) |
| (6) | $C_{tail}(Dequeue\ us\ sv) = k$ | (Calculate the worst case cost of tail .) |
| (7) | $k \leq 2 + (k - 1) - 1 = k + 2$ | (As this inequality holds, the time complexity of all n instructions is $O(n)$.) |

As the time complexity of all n instructions together is $O(n)$, the amortized cost of a single instruction is $O(1)$.

About the size function

We want to balance the size function such that:

- The size function is 0 to start with.
- The size between operations is large enough to prove the inequality.

The size function is arbitrary, if you cannot choose a size function that satisfied the goal inequality, then you're probably making a mistake

Peano Numbers

```

1 data Peano = Zero | Succ Peano
2
3 — analogous to (:) Cons
4 incr :: Peano -> Peano
5 incr = Succ
6
7 — analogous to tail
8 decr :: Peano -> Peano
9 decr (Succ n) = n
10 decr Zero = error "Cannot decrement zero"
11
12 — analogous to (++) concatenate
13 add :: Peano -> Peano -> Peano
14 add a Zero = a
15 add a (Succ b) = Succ (add a b)
16
17 — tail recursive version for extra goodness!
18 add a b = add (incr a) (decr b)

```

This shows how similar operations of similarly structured data can be.

Binary Numbers

```

1 data Bit = I | O
2
3 — [LSB,...,MSB]
4 type Binary = [Bit]
5

```

```

6 incr :: Binary -> Binary
7 incr [] = [I]
8 incr (O:bs) = I:bs
9 incr (I:bs) = O:incr bs

```

we can do amortized analysis on incr:

- (1) $C_{incr}(bs) = t + 1$ where $t = \text{length } (\text{takeWhile } (== I) \text{ } bs)$ (Create a cost function.)
- (2) $A_{incr}(bs) = 2$ (Create Amortized Cost)
- (3) $S(bs) = \text{length.filter } (== I) \text{ } bs$ (Create size function.)
- (4) Given $bs' = \text{incr } bs$ we show $C_{incr}(bs) \leq A_{incr}bs + S(bs) - S(bs')$ (Setup inequality)
- (5) $t + 1 \leq 2 + S(bs) - (S(bs) - t + 1)$ (Substitute in inequality)
- (6) $t + 1 \leq 1 + t$ (Hence inequality holds)
- (7) $S(\text{start}) = 0$ (Start size is zero.)

Hence The sum of C is smaller than the sum of A , as this is over n operations and $\sum A = 2n$, $C_{incr}(bs) \in O(1)$.

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List lookup

```

1  (!!) :: [a] -> Int -> a
2  (x:xs) !! 0 = x
3  (_:xs) !! k = xs !! (k-1)

```

As you can see `!!` costs $O(n)$ as it may traverse the entire list.

If we want this access to be faster, we can use trees:

```

1  data Tree a = Tip | Leaf a | Node Int (Tree a) (Tree a)
2
3  node :: Tree a -> Tree a -> Tree a
4  node l r = Node (size l + size r) l r
5
6  instance List Tree where
7    toList :: Tree a -> [a]
8    toList Tip = []
9    toList (Leaf n) = [n]
10   toList (Node n l r) = toList l ++ toList r
11
12   -- Invariant: size Node n a b = n = size a + size b
13   length :: Tree a -> Int
14   length Tip = 0
15   length (Leaf _) = 1;
16   length (Node n _ _) = n;
17
18   (!!) :: Tree a -> Int -> a
19   (Leaf x) !! 0 = x
20   (Node n l r) !! k
21   | k < m = l !! k
22   | otherwise = r !! (k-m)
23   where m = length l
24
25   -- case for Tip !! n or Leaf !! >0
26   _ !! _ = error "(!!): Cannot get list index"

```

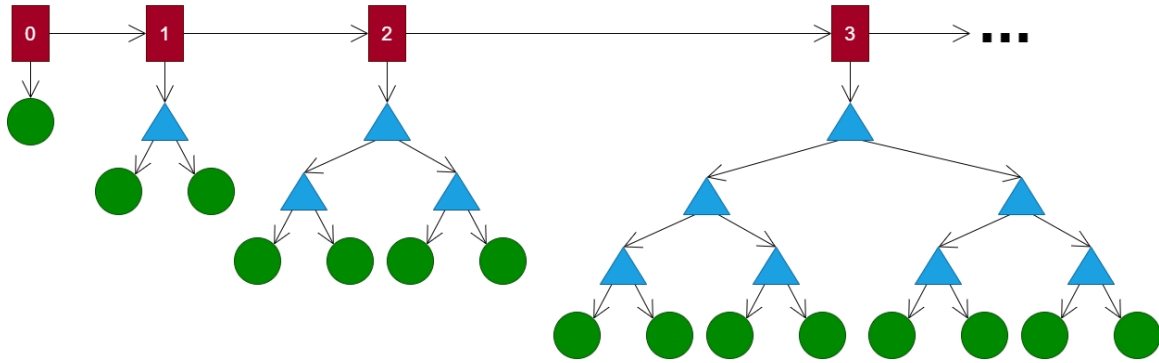
This costs $O(\log n)$ as each recursive call acts on half of the remaining list.

However we have difficulty with insertion:

Insert Quickly	$n : t = \text{node}(\text{Leaf } n)t$	Effectively becomes a linked list, $\log n$ search time ruined.
Insert Slowly	Rebalance tree (e.g. AVL tree)	Complex and no longer $O(1)$ insert.

Random Access Lists

A list containing elements that are either nothing, or a perfect tree with size the same as 2^{index} .



The empty tree can be represented by a *Tip* value (from the notes), or using type *Maybe(Tree)* (from the lecture) where $Tree\ a = Leaf\ x \mid Node\ n\ l\ r$.

When we add to a tree, we add to the first element of the **RAList**, if the invariant is breached (no longer perfect tree of size 2^0) it can be combined with the next list over (if empty, place, else combine and repeat).

This way while the worst case insert is $O(n)$, our amortized complexity is $O(1)$ much as with increment.

```

1  data Tree a = Tip | Leaf a | Node Int (Tree a) (Tree a)
2
3  type RAList a = [Tree a]
4
5  instance List RAList where
6    toList :: RAList a -> [a]
7    toList (RaList ls) = concatMap toList ls
8
9    fromList :: [a] -> RAList a
10   fromList = foldr (:) empty
11
12   empty :: RAList a
13   empty = []
14
15   (:) :: a -> RAList a -> RAList a
16   n : [] = [Leaf n]
17   n : (RAList ls) = RAList (insertTree (Leaf n) ls)
18   where
19     insertTree :: Tree a -> [Tree a] -> [Tree a]
20     insertTree t [] = [t]
21     insertTree t (tip:ls) = t:ls
22     insertTree t (t':ts) = Tip: insertTree (node t t') ts
23
24   length :: RAList a -> Int
25   length (RAList ls) = foldr ((+) . length) 0 ls
26
27   (!! ) :: RAList a -> Int -> a
28   (RAList []) !! _ = error "(!!): empty list"
29   (RAList (x:xs)) !! n
30   | n < m = x | n
31   | otherwise = (RAList xs) !! (n-m)
32   where m = length x

```


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Equality

```
1 class Eq a where
2   (==) :: a -> a -> Bool
```

Eq is the typeclass for equality, any instance of this class should ensure the equality satisfied the laws:

$$\begin{array}{ll} \text{reflexivity} & x == x \\ \text{transitivity} & x == y \wedge y == z \Rightarrow x == z \\ \text{symmetry} & x == y \Rightarrow y == x \end{array}$$

We also expect the indiscernability of identicals (Leibniz Law):

$$x == y \Rightarrow f\ x == f\ y$$

For a set-like interface we have a member function:

```
1 (in) :: Eq a => a -> Set a -> Bool
```

If we assume only that Eq holds, the complexity is $O(n)$ as we must potentially check all members of the set. To get around this, we use ordering.

Orderings

The Ord typeclass allows us to check for inequalities:

```
1 class Eq a => Ord a where
2   (<=) :: a -> a -> Bool
3   (<)  :: a -> a -> Bool
4   (>=) :: a -> a -> Bool
5   (>)  :: a -> a -> Bool
```

We must try to ensure certain properties hold, for example for to have a partial order we require:

$$\begin{array}{ll} \text{reflexivity} & x \leq x \\ \text{transitivity} & x \leq y \wedge y \leq z \Rightarrow x \leq z \\ \text{antisymmetry} & x \leq y \wedge y \leq x \Rightarrow x == y \end{array}$$

There are also total orders (all elements in the set are ordered compared to all others), for which we add the constraint:

$$\text{connexity} \quad x \leq y \vee y \leq x$$

Ordered Sets

```
1 class OrdSet ordset where
2   empty    :: ordset n
3   insert   :: Ord a => a -> ordset a -> ordset a
4   member   :: Ord a => a -> ordset a -> Bool
5   fromList :: Ord a => [a] -> ordset a
6   toList   :: Ord a => ordset a -> [a]
```

We can implement this class for Trees:

```
1 data Tree a = Tip | Node (Tree a) a (Tree a)
2
3 instance OrdSet Tree where
4   empty :: Tree n
5   empty = Tip
6
7   insert :: Ord a => a -> Tree a -> Tree a
8   insert x Tip = Node Tip x Tip
9   insert x (Node l y r)
10      | x == y    = t
11      | x < y     = Node (insert x l) y r
12      | otherwise = Node l y (insert x r)
13
14   member :: Ord a => a -> Tree a -> Bool
15   member x Tip = False
16   member x (Node l y r)
17      | x == y    = True
18      | x < y     = member x l
19      | otherwise = member x r
20
21   fromList :: Ord a => [a] -> Tree a
22   fromList = foldr insert empty
23
24   toList :: Ord a => Tree a -> [a]
25   toList Tip = []
26   toList (Node l y r) = toList l ++ y : toList r
```

However the worst case here is still $O(n)$ as we do not balance the tree as more members are inserted. If the members are added in order, the tree devolves to a linked list.

We need a way to create a tree that self balances.

Binary Search Trees (AVL Trees)

```
1 — H for height! The int stored at a node is the height of that tree.
2 data HTree a = HTip | HNode Int (HTree a) a (HTree a)
3
4 height :: HTree a -> Int
5 height HTip = 0
6 height (HNode h _ _ _) = h
7
8 hnode :: HTree a -> a -> HTree a -> HTree a
9 hnode l x r = HNode h l x r
10    where h = max (height l) (height r) + 1
```

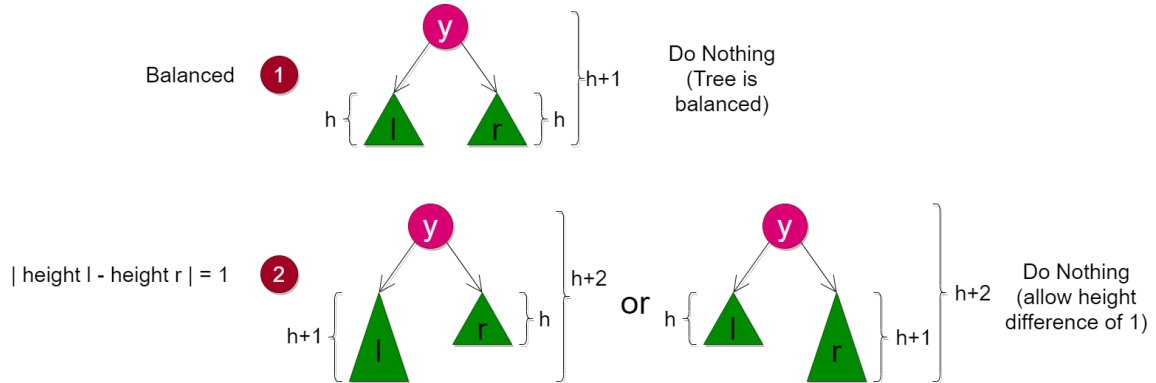
When inserting into the tree we must keep the tree balanced such that no subtree's left is more than one higher than its' right.

```

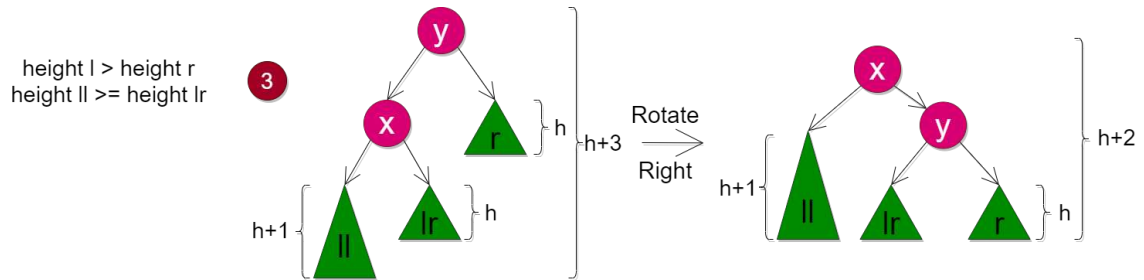
1 insert :: Ord a => a -> HTree a -> HTree a
2 insert x HTip = hnode Tip x Tip
3 insert x t@(HNode _ l y r)
4   | x == y    = t
5   | x < y     = balance (insert x l) y r
6   | otherwise = balance l y (insert x r)

```

We must rebalance the tree after insertion, this must consider the following cases:



When the tree's balanced invariant has been broken, we must follow these cases:

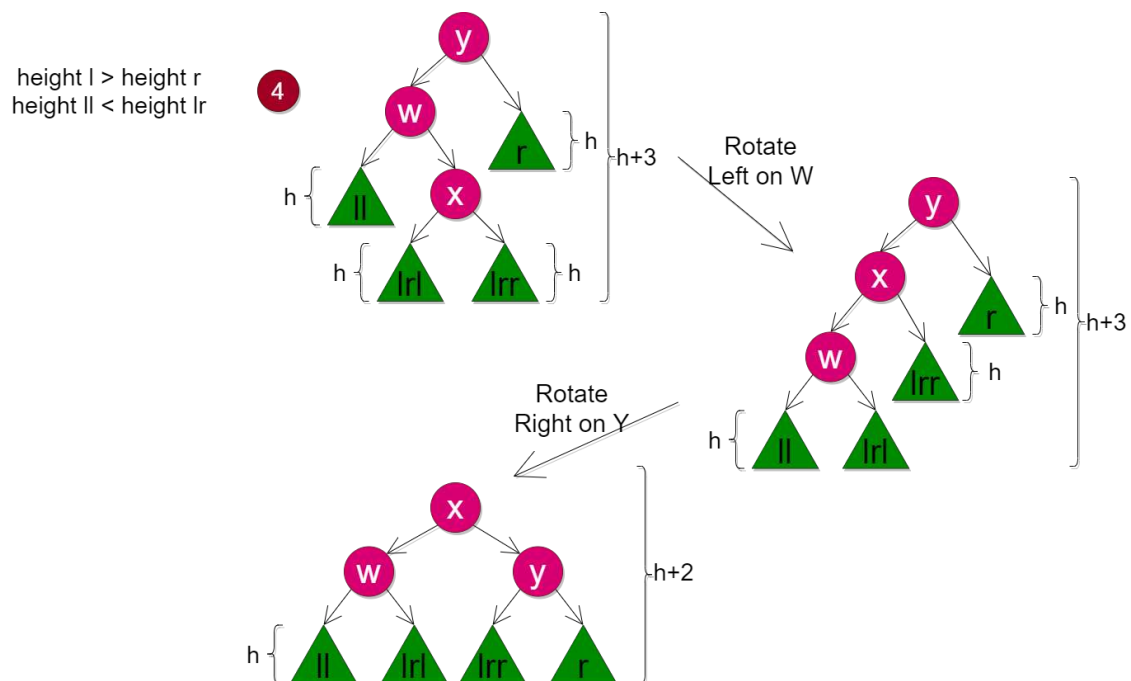


```

1 rotr :: HTree a -> HTree a
2 rotr (HNode _ (HNode _ ll x lr) y r) = hnode ll x (hnode lr y r)

```

When the right subtree's right subtree is higher:



```

1  rotl :: HTree a -> HTree a
2  rotl (HNode _ ll w (HNode _ lrl x lrr))
3    = hnode (hnode ll w lrl) x lrr
4
5  — so for the example: HNode _ wt y r -> rotr ( hnode (rotl wt) y r)

```

We can use rotate left and rotate right to for the two cases where the right subtree is 2 or more higher than the left.

```

1  balance :: Ord a => HTree a -> a -> HTree a -> HTree
2  balance l x r
3    | lh == lr || abs (lr - lrr) == 1 = t
4    | lh > lr   = rotr(hnode (if height ll < height lr then rotl l else l) x r)
5    | otherwise = rotr(hnode l x (if height rl > height rr then rotr r else r))
6  where
7    lh = height l
8    rl = height r
9    (HNode _ ll _ lrr) = l
10   (HNode _ rl _ rr) = r

```

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Red-Black Trees

AVL trees worked by storing an extra integer (height) to use in rebalancing, **red-black trees** use an extra bit to determine if a node is red or black.

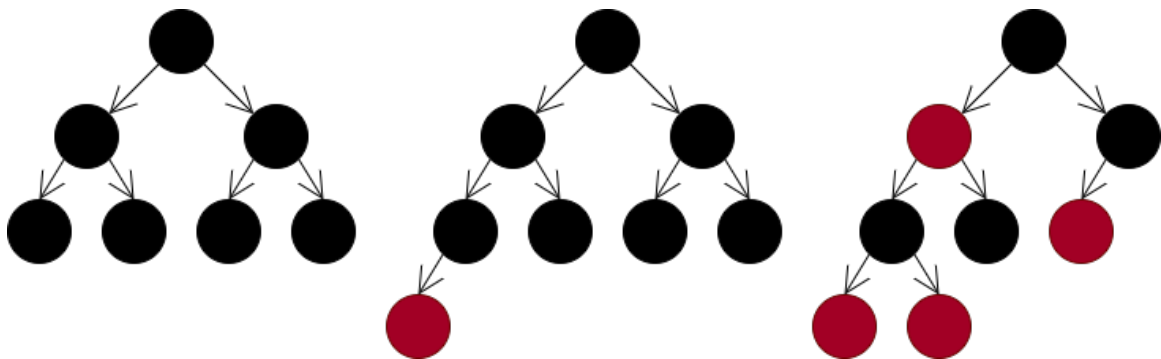
In practice they are less balanced than **AVL trees** however the insertion is faster and the data structure is a little bit smaller.

```
1 data Colour = Red | Black
2
3 data RBTREE a = Empty | Node Colour (RBTREE a) a (RBTREE a)
```

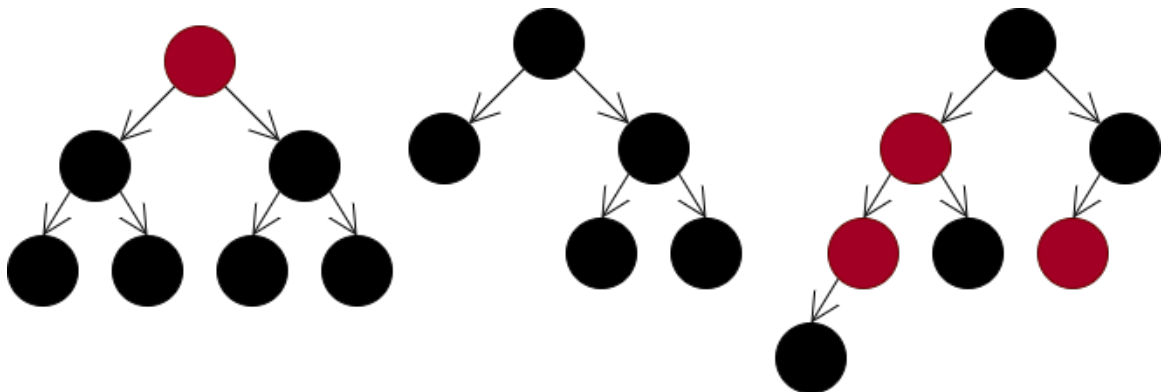
The structure relies on two invariances:

1. Every Red node must have a Black parent node.
2. Every path from the root to leaf must have the same number of black nodes.

Valid Red Black Trees



Invalid Red Black Trees

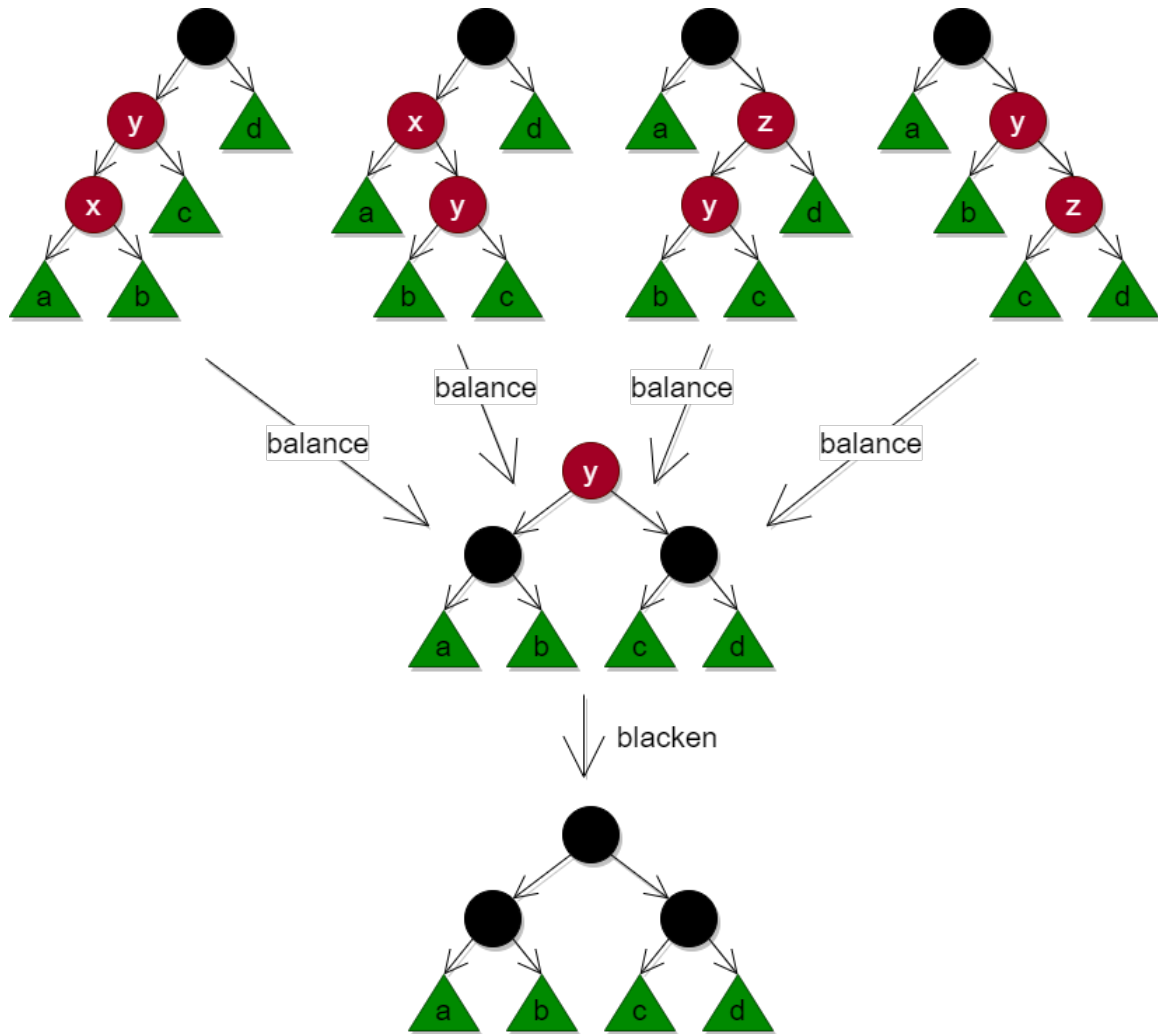


We have an insert function that needs to rebalance the tree:

```

1  blacken :: Ord a => RBTre a -> RBTre a
2  blacken (Node Red l x r) = Node Black l x r
3  blacken t                = t
4
5  balance :: Ord a => Colour -> RBTre a -> a -> RBTre a -> RBTre a
6  balance c l v r = case Node c l v r of
7    Node Black (Node Red (Node Red a x b) y c) z d -> bal x y z a b c d
8    Node Black (Node Red a x (Node Red b y c)) z d -> bal x y z a b c d
9    Node Black a x (Node Red (Node Red b y c) z d) -> bal x y z a b c d
10   Node Black a x (Node Red b y (Node Red c z d)) -> bal x y z a b c d
11   t                                                -> t
12   where
13     bal x y z a b c d = Node Red (Node Black a x b) y (Node Black c z d)
14
15  insert :: Ord a => a -> RBTre a -> RBTre a
16  insert = (blacken .) . ins
17   where
18     ins :: Ord a => a -> RBTre a -> RBTre a
19     ins x Empty = Node Red Empty x Empty
20     ins x t@(Node c l y r)
21     | x < y      = balance c (ins x l) y r
22     | x == y     = t
23     | otherwise = balance c l y (ins x r)

```

Counting

We can exploit the analogy we used with counting and trees for **RALists** here, with a difference.

Imagine a counting system that lacks zeros. We can count to 10 as:

Normal:	1	2	...	9	10	...	11	12	...	19	20	...	101	102	...	110	111
Special:	1	2	...	9	X	...	11	12	...	19	1X	...	X1	X2	...	XX	111

We can use this with the pattern of inserting elements into a red black tree (in order) to map red black trees to an incrementing number.

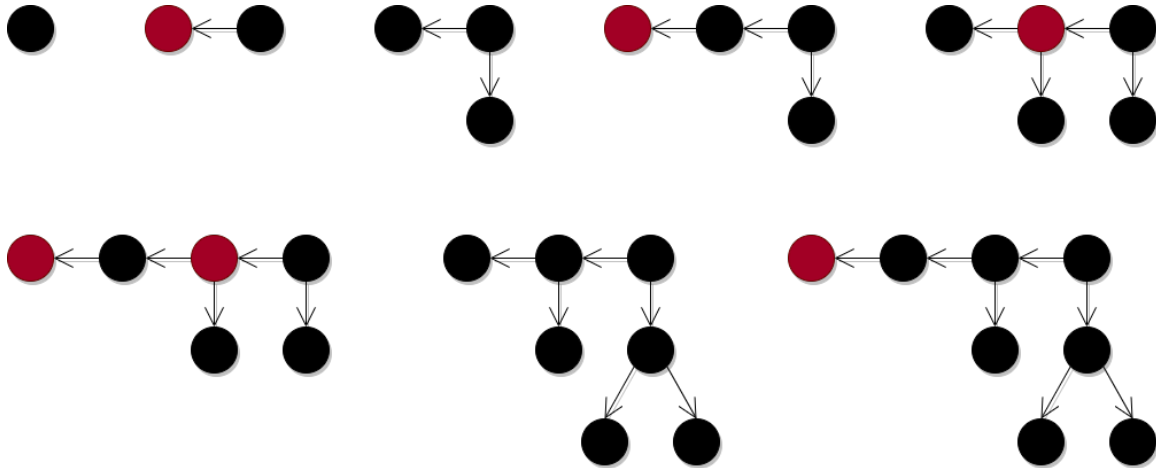
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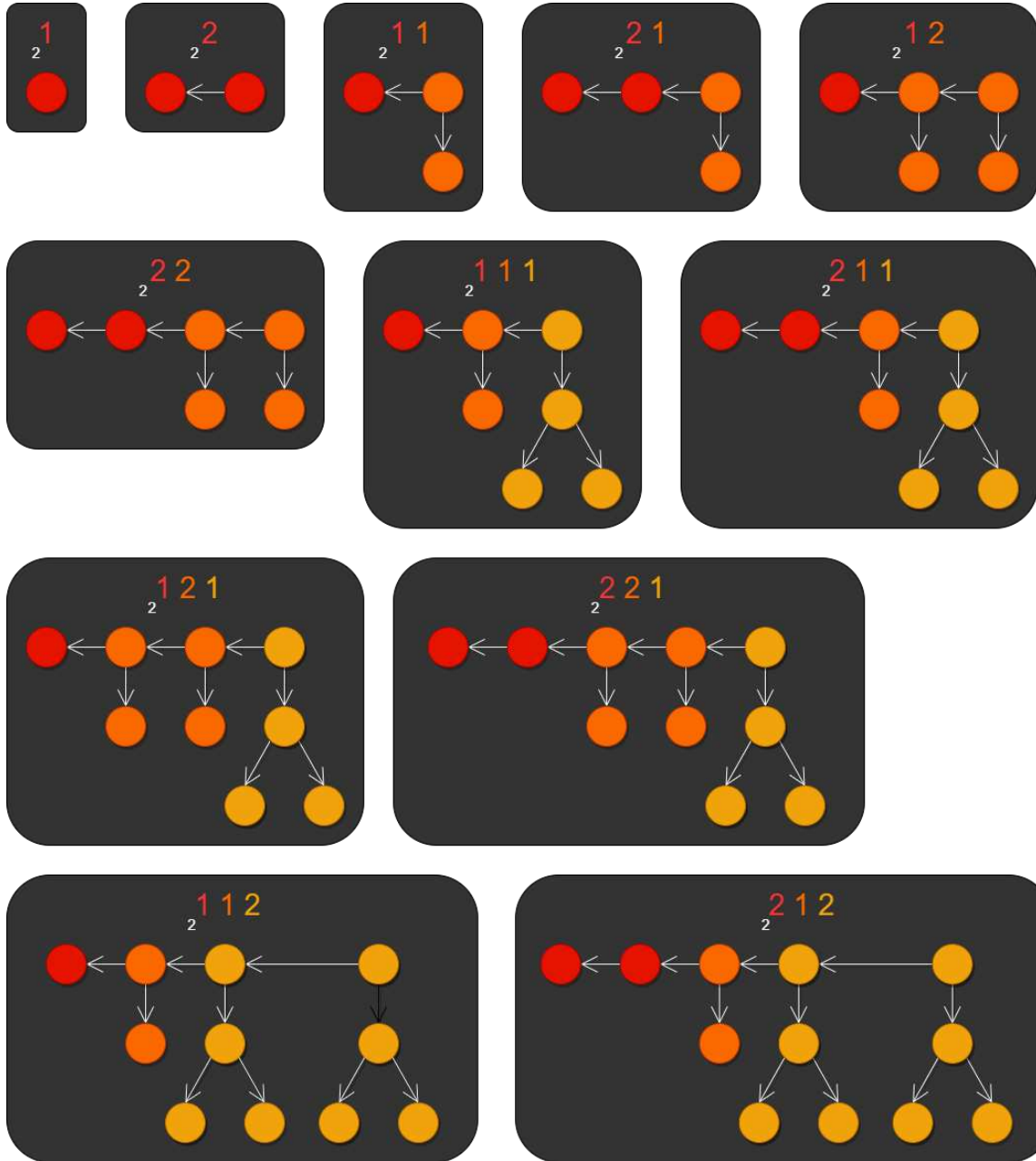
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Red Black Trees Continued

We have a pattern with inserting elements from an ordered list into the tree.



We can encode this as a special binary number system, using 1 and 2 such that the least significant bit is the number of trees of 2^0 nodes, and the n th is 2^n .



We can increment this by:

```

1 — each element is a red black tree (+ an extra root element (a))
2 data Digit a = One a (RBTTree a) | Two a (RBTTree a) a (RBTTree a)
3
4 incr :: a -> RBTTree a -> [Digit a] -> [Digit a]
5 incr x t [] = [One x t]
6 incr x t ((One y u) : ds) = Two x t y u : ds
7 incr x t ((Two y u z v) : ds) = One x t : incr y (Node Black u z v) ds

```

We can convert a list of digits back to a red black tree by:

```
1  — fold left to combine the digits together into a tree
2  fromList :: [a] -> RBTREE a
3  fromList xs = foldl link Empty (foldr add xs)
4
5  link :: RBTREE a -> Digit a -> RBTREE a
6  link l (One x t) = Node Black l x t
7  link l (Two x t y u) = Node Black (Node Red l x t) y u
```

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Randomized Algorithms

An algorithm that uses random values to produce a result.

Algorithm Type	Running time	Correct Result
Monte Carlo	Predicatable	Unpredictably
Las Vegas	Unpredictable	Predictably

Random Generation

Functions are deterministic (always map same inputs to same outputs), this is known as **Leibniz's law** or the **Law of indiscernibles**:

$$x = y \Rightarrow fx = fy$$

We can exhibit pseudo random behaviour using an input that varies explicitly (e.g Random numbers through seeds) implicitly (e.g Microphone or camera noise)

Inside IO Monad

We can use basic random through the IO monad like this:

```

1 import Control.Monad.Random (getRandom)
2
3 main :: IO ()
4 main = do
5     x <- getRandom :: IO Int
6     print (42 + x)
```

However using the **IO monad** is too specific, we may want to use random numbers in other contexts.

StdGen

In haskell we can use **Stdgen**.

```

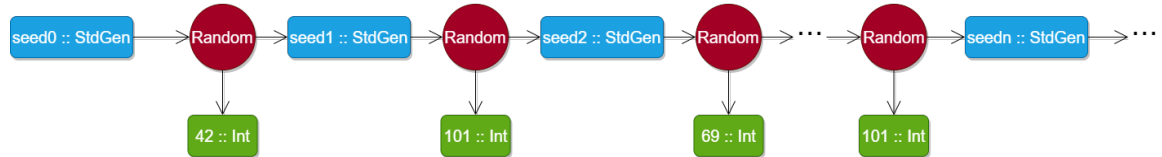
1 import System.Random (StdGen)
2
3 — Create a source of randomness from an integer seed
4 mkStdGen :: Int -> StdGen
5
6 — Generate a random interger , and a new source of randomness
7 random :: StdGen -> (Int , StdGen)
8
9 — Generate an infinite list of random numbers using an initial seed
10 — (source of random).
11 randoms :: StdGen -> [Int]
12 randoms seed = x:randoms seed' where (x, seed') = random seed
13
14 — In order to generate random value for any type, a typeclass is used
15 class Random a where
```

```

16 random  :: StdGen -> (a, StdGen)
17 randoms :: StdGen -> [a]
18
19 — Random over a (R)ange
20 randomR :: (a, a) -> StdGen -> (a, StdGen)
21 randomRs :: (a, a) -> StdGen -> [a]

```

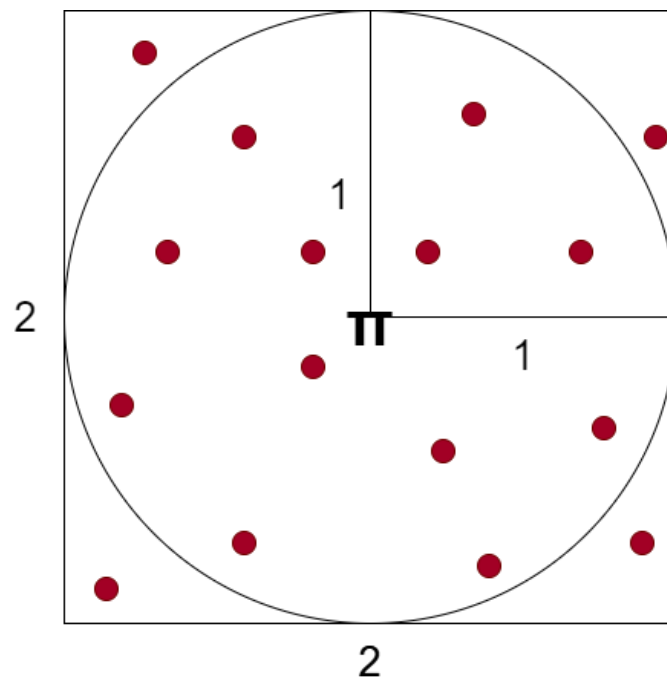
By passing the newly generated **StdGen** we can generate new values based on the original seed.



With Random Monad

Rather than passing **StdGen** seeds through the program, we can use the **MonadRandom** monad which internally uses this value.

Randomized π



(Monte Carlo Algorithm - known number of samples, known running time per sample) To estimate π , find the proportion of randomly selected spots that are within the circle.

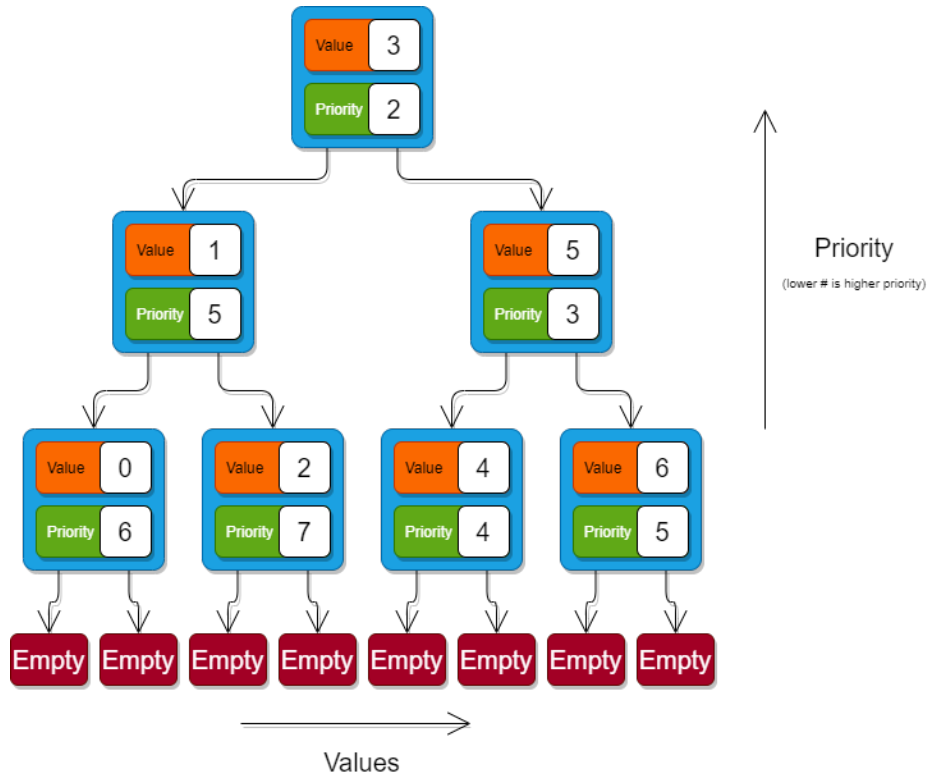
Area of square	$2 \times 2 = 4$
Area of circle	$\pi \times 1^2 = \pi$
Probability in circle	$\frac{\pi}{4}$

Once we have the proportion, we can multiply by 4 to get an estimate of π .

```
1 import Control.Monad.Random (getRandomR, randomRs, MonadRandom)
2 import System.Random (mkStdGen, StdGen)
3
4 — Here we can use one quarter of the circle, hence if the distance from the
5 — bottom left (0,0) to the point is within 1 then it is in the circle.
6 inside :: Double -> Double -> Bool
7 inside x y = 1 >= x * x + y * y
8
9 — Take 1000 samples and return 4 * the proportion.
10 montePi :: MonadRandom m => m Double
11 montePi = loop samples 0
12   where
13     samples = 10000
14     loop 0 m = return (4 * fromIntegral m / fromIntegral samples)
15     loop n m = do
16       x <- getRandomR (0,1)
17       y <- getRandomR (0,1)
18       loop (n+1) (if inside x y then m + 1 else m)
19
20
21
22
23 — Using a stream of random numbers (RandomRs)
24
25 — Get pairs of random numbers from the stream
26 pairs :: [a] -> [(a,a)]
27 pairs (x:y:ls) = (x,y):pairs ls
28
29 — From the pairs of random numbers, get the proportion of points inside the
30 — circle and use to get pi.
31 montePi' :: Double
32 montePi' = 4 * hits src / fromIntegral samples
33   where
34     samples = 10000
35     hits    = fromIntegral .
36               length .
37               filter (uncurry inside) .
38               take samples .
39               pairs
40     src     = randomRs (0, 1) (mkStdGen 42) :: [Double]
```

Treaps

Simultaneously a **Tree** and a **Heap**. Stores values in order, while promoting higher priority nodes to the top of the tree.



```

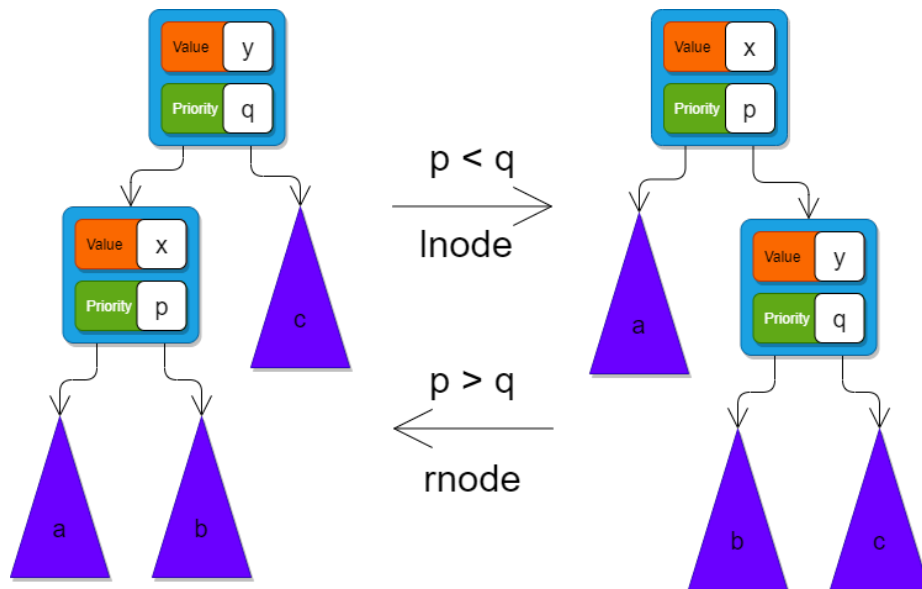
1  — Node contains child treaps, as well as value (a) and the priority (Int)
2  data Treap a = Empty | Node (Treap a) a Int (Treap a)
3
4  — Normal tree search using values
5  member :: Ord a => a -> Treap a -> Bool
6  member x (Node l y r)
7  | x == y    = True
8  | x < y     = member x l
9  | otherwise = member x r
10 member _ Empty = False
11
12 — Priority based insert
13 pininsert :: Ord a => a -> Int -> Treap a -> Treap a
14 pininsert x p Empty = Node Empty x p Empty
15 pininsert x p t@(Node l y q r)
16 | x == y    = t
17 | x < y     = lnode (pininsert x p l) y q r
18 | otherwise = rnode l y q (pininsert x p r)
19
20 — rotate right (check left node)
21 lnode :: Treap a -> a -> Int -> Treap a -> Treap a
22 lnode Empty y q r = Node Empty y q r
23 lnode l@(Node a x p b) y q c
24 | q > p    = Node a x p (Node b y q c)
25 | otherwise = Node l y q c
26
27 — rotate left (check right node)
28 rnode :: Treap a -> a -> Int -> Treap a -> Treap a
29 rnode l y q Empty = Node l y q Empty

```

```

30 rnode a x p r@(Node b y q c)
31 | q < p      = Node (Node a x p b) y q c
32 | otherwise = Node a x p r
33
34 — delete node by recursively searching, then delete and merge subtrees
35 delete :: Ord a => a -> Treap a -> Treap a
36 delete x Empty = Empty
37 delete x (Node a y q b)
38 | x == y      = merge a b
39 | x < y       = Node (delete x a) y q b
40 | otherwise   = Node a y q (delete x b)
41
42 merge :: Treap a -> Treap a -> Treap a
43 merge Empty r = r
44 merge l Empty = l
45 merge l@(Node a x p b) r@(Node c y q d)
46 | p < q      = Node a x p (merge b r)
47 | otherwise  = Node (merge l c) y q d

```



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Randomized Treaps

By using a random value for priority when inserting values into the treap, we can ensure a high likelihood of balancing, without complex balancing being required.

We can use this to create a randomized quicksort.

```

1 import System.Random (StdGen, mkStdGen, random)
2 — node random :: StdGen -> (Int, StdGen)
3
4 data RTreap a = RTreap StdGen (Treap a)
5
6 insert :: Ord a => a -> RTreap a -> RTreap a
7 insert x (RTreap seed t) = RTreap seed' (pinsert x p t)
8   where (p, seed') = random seed
9
10 — note 42 is used for
11 empty :: RTreap a
12 empty = RTreap (mkStdGen 42) Empty
13
14 — Build up tree, requires O(n log n)
15 fromList :: Ord a => [a] -> RTreap a
16 fromList xs = foldr insert empty xs
17
18 — Linear time conversion (use treap tolist)
19 toList :: RTreap a -> [a]
20 toList (RTreap _ t) = tolist t
21
22 — Randomized Quicksort O(n log n)
23 — Effectively the random priorities are the partitions, first pivot is the
24 — highest priority.
25 rquicksort :: Ord a => [a] -> [a]
26 rquicksort = toList . fromList

```

Randomized Binary Trees

We can balance a binary tree without using a treap, by inserting at the root (and rotating the tree to ensure it is ordered) with a certain probability.

```

1 import System.Random (StdGen, mkStdGen, randomR)
2
3 data BTree a = Empty | Node (BTree a) a (BTree a)
4
5 insert :: Ord a => a -> BTree a -> BTree a
6 insert x Empty = Node Empty x Empty
7 insert x t@(Node l y r)
8   | x == y = t
9   | x < y = Node (insert x l) y r
10  | otherwise = Node l y (insert x r)
11

```

```

12 — basic lefty/right rotations
13 rotr :: BTree a -> a -> BTree a -> BTree a
14 rotr (Node a x b) y c = Node a x (Node b y c)
15 rotr _ _ = error "(rotr): left was empty"
16
17 rotl :: BTree a -> a -> BTree a -> BTree a
18 rotl a x (Node b y c) = Node (Node a x b) y c
19 rotl _ _ = error "(rtol): right was empty"
20
21
22 — Insert to the root of the tree (maintaining order)
23 insertRoot :: Ord a => a -> BTree a -> BTree a
24 insertRoot x Empty = Node Empty x Empty
25 insertRoot x t@(Node l y r)
26   | x == y = t
27   | x < y = rotr (insertRoot x l) y r
28   | otherwise = rotl l y (insertRoot x r)
29
30 — Randomized binary tree
31 data RBTREE a = RBTREE StdGen Int (BTree a)
32
33 empty :: RBTREE a
34 empty = RBTREE (mkStdGen 42) 0 Empty
35
36 — chance of 1 / n+1 of inserting at root.
37 insert' :: Ord a => a -> RBTREE a -> RBTREE a
38 insert' x (RBTREE seed n t) = RBTREE seed' (n+1) (f x t)
39   where
40     f = case p of
41         0 -> insertRoot
42         _ -> insert
43     (p, seed') = randomR (0,n) seed

```

For every insert we have chance $\frac{1}{n+1}$ of inserting at the root of the tree. Then this occurs, the contents are rotated to ensure the tree's ordering is maintained.

This means that there is a very high probability of balance being maintained, however correct results are only returned when distinct elements are inserted at most once.

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Mutable Algorithms

We can use **STRef s a** to hold a mutable reference to **a** that can be created, read and modified.

```

1  — State Transformer (ST) takes a state s and return value a.
2  — "Give me any program with any state s, and if it returns an a, so will I".
3  runST :: (forall s . ST s a) -> a
4
5  — Take a value a, produces a program with some state s, that returns a
6  — reference with that state and the value stored.
7  newSTRef :: a -> ST s (STRef s a)
8
9  — Takes in a reference in some state s, performs a computation to get a
10 readSTRef :: STRef s a -> ST s a
11
12 — Take in a reference, and a new value a, performing a computation to update
13 — the referenced value (update does not return anything itself, hence []).
14 writeSTRef :: STRef s a -> a -> ST s ()
15
16 — Takes a value, returns a computation that executes in any state s to return
17 — an a.
18 return :: a -> ST s a
19
```

We can use this to create a mutable version of fibonacci.

```

1  import Data.STRef (newSTRef, readSTRef, writeSTRef)
2  import Control.Monad.ST (runST)
3
4  — immutable looping fibonacci
5  fib :: Int -> Integer
6  fib n = loop n 0 1
7  where
8      loop :: Int -> Integer -> Integer -> Integer
9      loop 0 x y = x
10     loop n x y = loop (n-1) y (x+y)
11
12 — mutable looping fibonacci
13 fib0 :: Int -> Integer
14 fib0 n = runST $ do
15     rx <- newSTRef 0
16     ry <- newSTRef 1
17     let loop 0 = do readSTRef rx
18         loop n = do {
19             x <- readSTRef rx;
20             y <- readSTRef ry;
21             writeSTRef rx y;
22             writeSTRef ry (x + y);
23             loop (n - 1); }
24     loop n

```


Mutable Datastructures

Array

```
1 import Control.Monad.ST
2 import Data.Array.ST
3
4 — Use indexable i, range, default to get a mutable array in some state
5 newArray :: Ix i => (i,i) -> a -> ST s (STArray s i a)
6
7 — Use indexable i, a mutable array and return the value a with state s
8 readArray :: Ix i -> STArray s i a -> i -> ST s a
9
10 — Use indexable i, a mutable array, and the value to place, return a
11 — computation that updates the array (new state) but returns no value.
12 — note:
13 writeArray :: Ix i => STArray s i a -> i -> a -> ST s ()
```

Each operation is assumed to take constant time.

For example, an algorithm to find the smallest natural number not in a list.

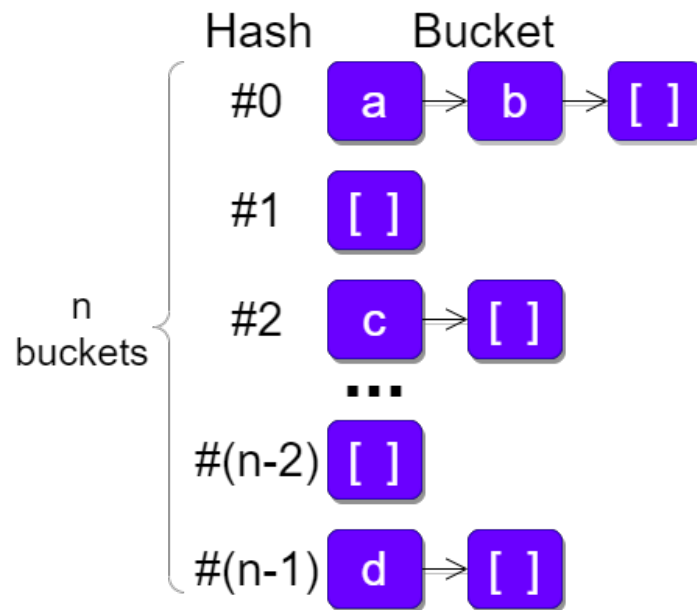
```
1 import Data.Array.MArray (MArray(newArray))
2 import Data.Array.ST
3 import Control.Monad.ST
4 import Data.List ((\\))
5
6 — immutable
7 minfree :: [Int] -> Int
8 minfree xs = head ( [0..] \\ xs )
9
10 — effectively the same as minfree
11 minfree' :: [Int] -> Int
12 minfree' xs = head . filter(not . ('elem' xs)) $ [0..]
13
14
15 — Builds up an array of which are present,
16 minfreeMut :: [Int] -> Int
17 minfreeMut = length . takeWhile id . checklist
18
19 {-
20 Build an array, at each index True/False for if the index is in xs
21 We only need to use an array of size (length xs) as we do not care about
22 natural numbers larger than this (if they are in the list, then a smaller
23 natural number was missed).
24
25 xs [0,1,2,3,6,7,8]
26 ys [T,T,T,T,F,F,T] ... (don't care about 7 or 8)
27
28 -}
29 checklist :: [Int] -> [Bool]
30 checklist xs = runST $ do {
31   ys <- newArray (0,l-1) False :: ST s (STArray s Int Bool);
32   sequence [writeArray ys x True | x <- xs, x < l];
33   getElems ys;
34 }
35 where
36   l = length xs
```

Hash

```
1 class Hashable a where
2   hash :: a -> Int
```

A hash generates an integer from some data. Typically range restricted (e.g hashmap can hold a finite number of entries), and the hash function should be designed to reduce collisions (two distinct data having the same hash).

Below is an example of a bucket based hash map, using linked list buckets.



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Mutable Nub

This can be useful for the **nub** function (removes duplicates from a list) by using a bucket based hashmap of items from the list to determine duplication.

```

1 import Control.Monad (when)
2 import Data.Array.ST
3   ( getElems, newListArray, readArray, writeArray, STArray )
4 import Control.Monad.ST ( ST, runST )
5
6
7 — immutable version:
8 nub :: Eq a => [a] -> [a]
9 nub = reverse . foldl nubHelper []
10   where
11     nubHelper :: Eq a => [a] -> a -> [a]
12     nubHelper ns c
13       | c `elem` ns = ns
14       | otherwise  = c:ns
15
16 — mutable version, create a hash table of characters to track which have
17 — already been seen.
18 nubMut :: (Hashable a, Eq a) => [a] -> [a]
19 nubMut xs = concat $ runST $ do
20   axss <- newListArray (0, n - 1) (replicate 256 [ ]) :: ST s (STArray s Int [a])
21   sequence [do {
22     let hx = hash x `mod` (n - 1)
23     ys <- readArray axss hx
24     unless (x `elem` ys) $ do writeArray axss hx (x : ys)}
25   | x <- xs]
26   getElems axss
27   where
28     — number of buckets in the hash table
29     n = 256

```

Quicksort

We can also implement quicksort, which can be done in place in an array (saved memory and time as accesses are constant time).

By using mutable data structures we can swap elements when reordering by reading and writing from the array.

```

1 import Data.Array.ST ( readArray, writeArray, STArray, getElems )
2 import Control.Monad.ST ( ST )
3
4 — swap elements over (classic store in temp & swap over other before writeback)
5 swap :: STArray s Int a -> Int -> Int -> ST s ()
6 swap axs i j = do
7   temp <- readArray axs i

```

```

8   readArray axs j >>= writeArray axs i
9   writeArray axs j temp
10
11  qsort :: Ord a => [a] -> [a]
12  qsort xs = runST $ do
13    axs <- newListArray (0,n) xs
14    aqsort axs 0 n
15    getElems axs
16    where
17      n = length xs - 1
18
19  {-
20  Partition around a pivot (k) (all smaller to left, larger to right
21  (unsorted)), then recur on these partitions
22  Index:      0      (k-1) k (k+1)      n
23  Contents: [ ..<= x .. ][x][ .. >x .. ]
24  -}
25  aqsort :: Ord a => STArray s Int a -> Int -> Int -> ST s ()
26  aqsort axs i j
27  | i >= j = return ()
28  | otherwise = do
29    k <- apartition axs i j
30    aqsort axs i (k - 1)
31    aqsort axs (k + 1) j
32
33  apartition :: Ord a => STArray s Int a -> Int -> Int -> ST s Int
34  apartition axs p q = do
35    x <- readArray axs p
36    let loop i j
37      | i > j = do
38        swap axs p j
39        return j
40      | otherwise = do
41        u <- readArray axs i
42        if u < x
43          then do loop (i + 1) j
44          else do
45            swap axs i j
46            loop i (j - 1)
47    loop (p+1) q

```