

## Time Complexities

We define the cost for evaluating algorithms based on their input size. We analyse performance on this as it varies. For insert :: int -> [int] -> [int] we work out this cost by writing a recurrence relation. We perform worst case analysis - for insert this would be an insert right at the end:

```
Tinsert(n) = 1 + Tinsert(n-1) = 1 + 1 + ... 1 + Tinsert(n-n) = n + Tinsert(0) = n+1.
```

Thus insertion is O(n). We do all our calculations under strict evaluation.

```
isort :: [int] -> [int]
isort [] = []
isort (x::xs) = insert x (isort xs)
```

We have: Tisort(0) = 1, Tisort(n) = 1 + Tisort(n-1) + Tisort(n-1).

So Tisort(n) = 1 + Tisort(n-1) + Tisort(n-1)

= 1 + n + (1 + Tisort(n-2) + Tisort(n-2))

= 2n+1 + Tisort(n-2)

Isort(0) = 1, (1) = 3, (2) = 6, (3) = 1 + 3 + 6 = 10.

This pattern unfolds to: n+1 + n + n-1 + ... 2 + 1, and is thus the sum n+1 numbers. This is (n+1)(n+2)/2, and thus O(n²).

## 2) Evaluation

In our evaluation models, like lazy evaluation, minimum = head isort is actually O(n). This is hard to deal with thus I'll only do strict evaluation.

### 2.1) Normal Forms

There are 3 types of normal forms when reducing a function (a function is f(x) = y <-> f = λx y - **Lambda Calculus form**)

Normal Form: Cannot be evaluated any further - no further reductions.

Weak Head Normal Form: Just a constructor, we know what type it is from the constructor, but it is reducible. e.g. [3+2], λx.3+4, Just(8+9)

### 2.2) Expressions

The key to giving an algorithm an overall cost is to give a series of rules that will assign a cost to a function by allocating a cost to its constituent parts. **Expressions** can be broken down into terms of the following grammar:

```
e ::= x (variables)
    | k (constants)
    | f e₁..en (applications)
    | if e then e1 else e2 (conditionals)
```

A function in this language is assumed to be defined by the form f x₁ ... xₙ = e. **Infix operations** are written as f x₁ ... xₙ instead of (+) x₁ y. **Primitive constants** such as True, False, 0, 1, 2, are available, as are stand operations on them such as <, <=, (+), and (\*). **List constants and operations** are also primitive, such as [], (:), (c), null, head and tail. There are **two main types of evaluation order** which changes runtime as mentioned before: 1) **Applicative order**: Evaluating arguments to a function before evaluating the function itself. 2) **Normal order**: evaluating the function before evaluating its arguments. Applicative order doesn't always terminate, but if it does then these both always reduce to some normal form. In a strict setting we use applicative order, in a lazy setting we use normal order.

### 2.3) Strict Term Analysis

Given a function f of n arguments, T(f) x₁ ... xₙ - the number of steps it takes to evaluate f x₁ ... xₙ can be worked out as follows: 1) For primitive f, T(f) x₁ ... xₙ = 0, e.g. T(head) xs = 0, T((++) x y) = 0. 2) For any other function f: f x₁ ... xₙ = e, T(f) x₁ ... xₙ = 1 + T(e) (We are given that the arguments are already evaluated). Evaluating a function is 1 step, and then we need to carry out the function body e). Now, T(e) can be defined in terms of expressions, by induction on e:

**Primitives and Variables:** T(x) = 0, T(k) = 0

**Application:** T(f e₁ ... eₙ) = T(f) e₁ ... eₙ + T(e₁) + ... + T(eₙ).

**Conditional:** T(if e then e₁ else e₂) = T(e) + if e then T(e₁) else T(e₂)

Examples:

T(3+4) = T((+) 3 4) + T(3) + T(4) = 0 + 0 + 0 = 0

T(length xs) is: if null xs then 0 else T(++) 1 (length (tail xs)) + T(1) + T(length (tail xs)) (app rule)

= 1 + T(null xs) + if null xs then 0 else T(length (tail xs)) + T(++) 1 (length (tail xs)) (prim rule)

We applied **primitive/var rule** throughout multiple times, for terms like T(n).

So T(length) xs = 1 + if null xs then 0 else T(length) (tail xs), generating a recurrence relation.

**Composition Rule:** The cost of f(g(x)) is T(f(g(x))) = T(f) (g) + T(g) x

### 3) Asymptotic Functions

High to low cost: O(m²), O(k²), O(n!), O(n²), O(n log n), O(n), O(1)

(When dealing with asymptotics we restrict ourselves to a family of mathematical functions called **L-Functions**).

**3.1.1) L-Functions** - an L-Function is a **real, positive, monotonic** (only moves in 1 direction on y axis), one-valued function (each point has a unique value in the range) on a real variable **defined for all values greater than some definite value** by a finite combination of algebraic symbols, exponentials, and logarithms, operating on real constants and the variable. **Theorem:** Any L-function f is ultimately continuous, of constant sign, monotonic, and as n -> ∞, the value f(n) tends to one of 0, ∞, or some other definite limit. Can categorise L-Functions with Du Bois Raymond.

## 3.2) Du-Bois Raymond Notation

The first important concept when looking at complexity is the notion of the rate of increase of a function relative to another: a function can be seen to grow quickly or slowly with respect to some other function. As is standard, the rate of increase of two functions can be understood as the ratio between them.

Now suppose f and g are L-functions, and consider the ratio of the L-function f/g(n) = f(n)/g(n) as n tends to infinity. This gives rise to a family of operations, <, <=, >, and > that can be used to compare functions:

f < g ⇔ lim<sub>n→∞</sub> f(n)/g(n) < 0

f ≤ g ⇔ lim<sub>n→∞</sub> f(n)/g(n) ≤ 0

f > g ⇔ 0 < lim<sub>n→∞</sub> f(n)/g(n) < ∞

f ≥ g ⇔ lim<sub>n→∞</sub> f(n)/g(n) ≥ 0

f > g ⇔ lim<sub>n→∞</sub> f(n)/g(n) > 0 (3.5)

These operations have good calculational properties:

We have **trichotomy**: f is less than g, f is comparable to g, f is more than g.

**Converse:** f < g ⇔ g > f.

**Transitivity:** f < g && g < h ⇒ f < h (same with less and equal).

### 3.3) Bachman-Landau (Big-O) Notation

f = O(g(n)) ⇔ f < g

f = O(g(n)) ⇔ f ≤ g

f = O(g(n)) ⇔ f = g

f = Ω(g(n)) ⇔ f ≥ g (3.12)

f = ω(g(n)) ⇔ f > g (3.13)

These sets can also be defined directly:

O(g(n)) = { f | ∃ v0 > 0. ∃ n0 > 0. f(n) < v0 g(n) }

O(g(n)) = { f | ∃ v0 > 0. ∃ n0 > 0. v0 n0 < f(n) < v0 g(n) }

Ω(g(n)) = O(g(n)) ∩ Ω(g(n)) (3.16)

Ω(g(n)) = { f | ∃ v0 > 0. ∃ n0 > 0. v0 n0 < f(n) < v0 g(n) } (3.17)

ω(g(n)) = { f | ∃ v0 > 0. ∃ n0 > 0. v0 n0 > f(n) > v0 g(n) } (3.18) (3.19)

### 4) Lists

**Lists** in Haskell are defined like so: data [a] = [] | (:) a [a]

The data keyword indicates that a new datatype is being introduced.

The type itself is called [a], which can be constructed by means of the constructors, [] for empty lists, and (:) for adding an element to a list, introduced to the right of the equality symbol. The types of these constructors is: [] :: [a] :: (:) :: a -> [a] -> [a]

The ++ operation is O(n). The ++ operation is O(n).

(++) :: [a] -> [a] -> [a]

[ ] ++ ys = ys

(x : xs) ++ ys = x : (xs ++ ys)

The whole list must be traversed, so given length xs = n, then T(++)(n) ∈ O(n). This pattern of recursion on the structure of a list itself crops up frequently, and is captured by the **foldr** function:

foldr :: (a -> b -> b) -> b -> [a] -> b

foldr f k [ ] = k

foldr f k (x : xs) = f x (foldr f k xs)

foldr applies functions to a list like so f x₁ f x₂ (... f xₙ k)), effectively applying the two arg function to functions in the order we get the result. When foldr is used with a binary operator (x), the following holds: foldr (:) [x₁, x₂, ..., xₙ] [] = x₁ ∘ (x₂ ∘ (... (xₙ ∘ e)))

Furthermore, when (:) is associative and e is a neutral element, this is simply: foldr (:) [x₁, x₂, ..., xₙ] [] = x₁ ∘ x₂ ∘ ... ∘ xₙ

One way to interpret this is that applying foldr is a destructor of lists, dual to the operations (:) and [] which are constructors. This can be seen by considering the effect of foldr (:) [], it is equal to the identity function.

We can define some list operations in terms of foldr: concat [xs, xs₁, xs₂, ...] = xs ++ xs₁ ++ xs₂ ... = foldr (++) [] xs, thus concat = foldr (++) []. It is O(nm), n = num lists, m = length of longest list.

**foldl** is foldr but left recursive:

foldl :: (b -> a -> b) -> b -> [a] -> b

foldl f k [ ] = k

foldl f k (x : xs) = foldl f (f k x) xs

Some functions are **faster/slower** using left recursion rather than right recursion despite producing the same results. (concat using foldr is much slower, O(n²m) as we have our resultant list on the left hand side each time, rather than what we're adding. And we see the definition of concat means that we'd thus traverse much more space)

**Just because functions produce the same result doesn't mean they're the same speed.**

**Monoid:** a set X that is equipped with an associative binary operation (·): X × X -> X and a neutral (identity) element e: X.

### 5) Abstract Lists

Abstract interfaces for lists can be created that can be instantiated to different concrete implementations with varying complexity characteristics. Our interface looks like this: *class List* where

*fromList :: [a] -> list a*

*normalize :: list a -> list a*

*single :: a -> list a*

*snoc :: list a -> a -> list a*

*tail :: list a -> list a*

*last :: list a -> a*

*isSingle :: list a -> Bool*

*isSingle :: list a -> list a -> list a*

*(!!) :: list a -> Int -> a*

*fromList* takes us to our abstract implementation, *toList* from it, *normalize* applies *fromList* *toList*. (*toList* *fromList* = id, and should be maintained as such) *empty* is a function constructing the empty list, *single* :: a -> list a

*single* x = *fromList* [x]

*snoc* :: list a -> a -> list a

*snoc* xs x = *fromList* (*toList* xs ++ [x])

Here are some concrete implementations of lists:

**4.1) Default Lists** - the standard list implementation. *toList* and *fromList* are simply the identity function. We can define all functions by referring to their **Prelude** versions or redefining them in place.

**4.2) Tree Lists** - A binary tree with values at its leaves can be considered to be a list, where an in-order traversal of the list from left to right corresponds to the order of the list elements. This representation is good as **appending two trees together is achieved by simply placing them under a parent fork**.

**4.3) Difference Lists** - We get constant time cons, snoc and ++. Other operations become more expensive though. For a difference list we replace (++) with function composition. Thus appending lists together always ends up in a right-associated list which occurs because of the definition of composition.

(·) :: (b -> c) -> (a -> b) -> (a -> c)

(g · f) x = g (f x)

If f = (xs++), g = (ys++), and h = (zs++), then their composition: ((zs++) · (ys++) · (xs++)) [] = zs ++ (ys ++ (xs ++ []))

Thus we get right associative appends which is desirable as it is O(1).

This is how we implement them:

*newtype DList a = DList ([a] -> [a])*

Note carefully that the append function (++) used here has type:

(++) :: List list a -> list a -> list a This has a constraint that states that list must be a member of the List class, which allows it to work on different types, as annotated in the calculation. We get:

*instance List DList* where

*toList :: DList a -> [a]*

*toList (DList fxs) = fxs [ ]*

*fromList :: [a] -> DList a*

*fromList xs = DList (xs++)*

*(++) :: DList a -> DList a -> DList a*

*DList fxs ++ DList fys = DList (fxs · fys)*

*fxs* represents funcons on lists, *fys* = (xs++).

**6) Divide and Conquer** - fundamental algorithmic strategy. Consists of three parts: 1) Divide a problem into subproblems 2) Solve subproblems into subsolutions 3) Combine subsolutions into a solution

An example is merge sort:

*msort :: [Int] -> [Int]*

*msort [] = [ ]*

*msort (x : xs) = [x] ++ msort xs*

*msort xs = merge (msort us) (msort vs)*

where

*(us, vs) = splitAt (n `div` 2) xs*

*n = length xs*

*merge :: [Int] -> [Int] -> [Int]*

*merge [ ] ys = ys*

*merge xs [ ] = xs*

*merge (x : xs) (y : ys)*

*[x <= y :: merge xs (y : ys)]*

*[otherwise :: y :: merge (x : xs) ys]*

*msort* splits the list in halves, and then we merge back up starting with the smallest lists.

We see that merge does the bulk of the work in its recursive case, swapping the value of x and y. This repeated splitting gives us T<sub>msort</sub>(0) = 1, T<sub>msort</sub>(n) = 1 + T<sub>merge</sub>(n) = T<sub>merge</sub>(n) + T<sub>merge</sub>(n/2) + T<sub>merge</sub>(n/2) + 2 × T<sub>msort</sub>(n/2), and solving it gives us O(n log n). The key is that to merge two lists of size n (we only merge lists of similar size), it takes at most n/2 comparisons. And we build the list up merging as we go so we are comparing using small lists. Quicksort is another divide and conquer algorithm. The time complexity is usually O(n log n), but with bad pivot choice is O(n²) and thus the worst case (so some time complexity) is O(n²).

Here we choose pivot = first element always, split into a list of items smaller than it and one bigger. We then sort these two lists which are split, and combine results.

*qsort :: [Int] -> [Int]*

*qsort [ ] = [ ]*

*qsort [x] = [x]*

*qsort (x : xs) = qsort us ++ [x] ++ qsort vs*

where (us, vs) = partition (<x) xs

*partition :: (a -> Bool) -> [a] -> ([a], [a])*

*partition p xs = (filter p xs, filter (¬ p) xs)*

### 7) Dynamic Programming

We trade storage for speed using **memoization**. The speedup comes from caching subsolutions with memorization and later looking them up rather than recomputing these subsolutions. Our strategy is:

1. Write an inefficient recursive algorithm that solves the problem.

2. Improve efficiency by storing intermediate shared results. (using *tabulate* and *memo*)

We should use an array to store results in Haskell as arrays are O(1) compared to a list's O(n). We make arrays from lists using this function

*array :: Ix i => (i, i) -> [a] -> Array i a*

*array* can be used to look things up in constant time.

*fib* :: Int -> Integer

*fib 0 = 1*

*fib 1 = 1*

*fib n = fib (n - 1) + fib (n - 2)*

Here's an example, computing the nth Fibonacci number in O(n), using bottom up DP.

**memo** must be in the same scope as *fib*. We see *memo* mirrors our old fib recursive.

*fib' :: Int -> Integer*

*fib' n = table ! n*

where

*table :: Array Int Integer*

*table = tabulate (0, n) memo*

*memo 0 = 0*

*memo 1 = 1*

*memo n = table ! (n - 1) + table ! (n - 2)*

*tabulate :: Ix i => (i, i) -> (i -> a) -> Array i a*

*tabulate (u, v) f = array (u, v) [(i, f i) | i ← range (u, v)]*

Here's the standard *tabulate* function which produces our table by applying our function f to all values between x and y. (0, n) and n in this case). The magic comes from tabulating with our *memo* function. All we have to do now is customise our function supplied to *tabulate*, to get the values we need (it should follow the pattern of the old recursive version), and get the correct array index as the final answer.

**7.1) Edit Distance Problem** - this is a complex DP problem which requires a 2d array, and thus indexing. In this problem we need to find the number of insertions, deletions and updates it takes to turn one string into another. The problem is simplified by considering only deletions and updates - insertion of a char into a string is the same as deleting the char from the other. One way to visualise it is moving to the left = deletion of first char in first string, right = deletion of first from right, middle = deletion of both first chars if they match.

We capture this with this recursive algorithm:

*dist :: String -> String -> Int*

*dist xs [ ] = length xs*

*dist [ ] ys = length ys*

*dist xs@(x::xs') ys@(y::ys') = minimum [dist xs ys + 1, dist xs' yys + 1,*

*dist xs ys + if x == y then 0 else 1]*

Minimum lets us consider the cheapest cost of three choices at each node. This is very inefficient - O(3<sup>m+n</sup>). We can fix this with DP, but we need to make our problem amenable to DP as strings can't be used to index into an array. We do this by measuring our progress

across xs and ys with indices i and j rather than cutting the head of the string away:

*dist' :: String -> String -> Int -> Int -> Int*

*dist' xs ys i 0 = i*

*dist' xs ys 0 j = j*

*dist' xs ys i j = minimum [dist' xs ys i (j-1) + 1, dist' xs ys (i-1) j + 1,*

*dist' xs ys (i-1) (j-1) + if x == y then 0 else 1]*

where

*m = length xs*

*n = length ys*

*x = xs !! (m - i)*

*y = ys !! (n - j)*

*tabularizing* this is quite easy now. Just take our template, replace memo with the definition of *dist'*, pass in a 2d array index into *tabulate* (doesn't need to be redefined) We do have to use *fromList* to make arrays rather than lists - to remove the !!.

*dist'' :: String -> String -> Int*

*dist'' xs ys = table ! (m, n)*

where

*table = tabulate ((0, 0), (m, n)) (\uncurry memo)*

*memo :: Int -> Int -> Int*

*memo i 0 = i*

*memo 0 j = j*

*memo i j = minimum [table ! (i, j - 1) + 1, table ! (i - 1, j) + 1,*

*table ! (i - 1, j - 1) + if x == y then 0 else 1]*

where *x = xs !! (m - i)*

*y = ys !! (n - j)*

*m = length xs*

*n = length ys*

*axis, axis :: Array Int Char*

*axis = fromList xs*

*axis = fromList ys*

## Standard Access Lists

The standard list representation models itself on Peano Numbers. Random Access Lists model the structure on

**Binary Numbers** instead, which has its own benefits.

newType RAList a = RAList [Tree a]

The RAList has the same complexity as a tree. e.g. !!:

(!!) :: RAList a → Int → a

(!!) :: RAList (t :: ts) !! k

| k < m == !! k

| otherwise = RAList ts !! (k - m)

where m = length t

Given xs :: RAList a, the cost of performing xs !! k is O(log k) in the worst case. The interesting operation is the

function:

cons :: a → RAList a → RAList a

cons xs x = RAList (consTrees (Leaf x) xs)

where

consTrees :: Tree a → RAList a → [Tree a]

consTrees t (RAList []) = [t]

consTrees t (RAList (Tip::ts)) = t : ts

consTrees t (RAList (t'::ts)) = Tip :

consTrees (fork t t') (RAList ts)

Notice that this follows the structure of the Inc :: Binary →

Binary function, benefiting from similar amortized

complexity.

## 10) Searching

To search for an item in a structure, we must have some

notion of **equality**:

class Eq a where

(=) :: a → a → Bool

Valid implementations of eq need to have an equality

operator that behaves well - we need **reflexivity**,

**transitivity** and **antisymmetry**. The language

doesn't enforce this, so we must implement it validly.

Here's an example implementation of Eq:

data Fruit = Apple | Orange

instance Eq Fruit where

Apple == Apple = True

Orange == Orange = True

\_ == \_ = False

The simplest way to search a list is to just query the entire

thing until we find the elem:

runmunge x [] = False

runmunge x (y : ys) = x == y || runmunge x ys

We can improve on this by using a data structure with

more order

## 10.1) Ordered Lists:

The assumption that the elements can be ordered is recorded with an

Ord constraint, which

expects the (<) relation to be defined. The Ord class itself

relies on the existence of Eq, so that the order (<) can be

compatible with equality (=).

class Eq a → Ord a where

(<) :: a → a → Bool

We require that the < relation is a partial order

- transitive, reflexive, antisymmetric

(x < y ∧ y < x). We could implement a poset:

class Poset x where

toPoset :: Ord a => [a] → poset a

fromPoset :: poset a → [a]

empty :: poset a

insert :: Ord a => a → poset a → poset a

delete :: Ord a => a → poset a → poset a

member :: Ord a => a → poset a → Bool

union :: Ord a => poset a → poset a → poset a

inter :: Ord a => poset a → poset a → poset a

We write member like so, but it has the same

time time complexity

member :: Ord a => [a] → Bool

member x [] = False

member x (y : ys) = x == y || (x < y && member x ys)

**10.2) Search Trees** - Quicksort works by

taking a pivot that partitions data into

two parts. The structure of this recursion

can be captured in this tree:

data Tree a = Nil | Node (Tree a) a (Tree a)

Constructing this tree is like the splitting step of quicksort:

instance Poset Tree where

toPoset :: Ord a => [a] → Tree a

toPoset [] = Nil

toPoset (x : xs) = Node (toPoset (x : xs)) x (toPoset xs)

(where vs, us) = partition (< x) xs

This allows faster access to elements when balanced.

member :: Ord a => Tree a → Bool

member x Nil = False

member x (Node (hNode p x q) y r)

| x == y = True

| x < y = member x l

| otherwise = member x r

If the tree is balanced, then its depth will force T<sub>member</sub>(n)

O(log(n)), where n is the number of elements in the tree.

The worst case is still linear though.

## 10.2.1) Binary Search (AVL) Trees - balanced trees

By keeping track of height:

type Height = Int

data HTree a = HTip | HNode Height (HTree a) a (HTree a)

To make sure trees are constructed in the proper way where

Height is preserved, a smart constructor is used:

hNode :: HTree a → a → HTree a → HTree a

hNode l x r = HNode h l x r

where h = (height l ⊕ height r) + 1

height :: HTree a → Int

height HTip = 0

height (HNode h l x r) = h

Insert is the difficult case, as we must maintain balancedness of

our tree. We use the balance and balancer smart constructors,

to maintain the invariant:

1) **The difference in height between siblings is at most 1:**

instance Poset HTree where

insert :: Ord a => a → HTree a → HTree a

insert x HTip = hNode HTip x HTip

insert x t@(HNode l y r) =

| x == y = t

| x < y = balance (insert x l) y r

| otherwise = balancer l y (insert x r)

There are multiple cases to consider for balance and balancer:

1) The height of it and r differ by at most 1 already. Focusing on

the balance case - inserting into the left tree, we only need

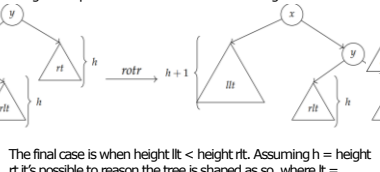
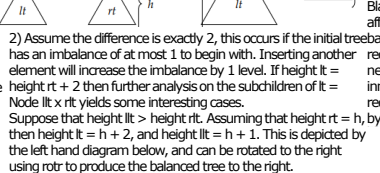
compare height l - height r:

balance :: HTree a → a → HTree a → HTree a

balance l y rt

| height l - height rt ≤ 1 = hNode l y rt

To fall into this case tree must be one of the two balanced trees



2) Assume the difference is exactly 2, this occurs if the initial tree

has an imbalance of at most 1 to begin with. Inserting another

element will increase the imbalance by 1 level. If height l =

height rt + 2 then further analysis on the subchildren of l =

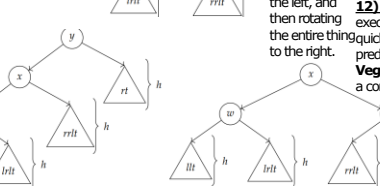
Node l l x r ylt shows some interesting cases.

Suppose that height l > height rt. Assuming that height rt = h,

then height l = h + 2, and height l l = h + 1. This is depicted by

the left hand diagram below, and can be rotated to the right

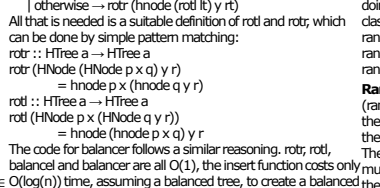
using rotr to produce the balanced tree to the right.



The final case is when height l < height rt. Assuming h = height

rt it's possible to reason the tree is shaped as so, where l =

Node l l w r l and r l = Node l r l x r r t:



These cases can be encoded by:

| otherwise = case l of hNode l x r l

| height l > height rt = rotr (hNode l y rt)

| otherwise = rotr (hNode (rotr l) y rt)

## 11) Red Black Trees - Another way of creating balanced trees,

they don't store height but rather the colour of a node - red/black.

data Colour = R | B

data RBTree a = E | N Colour (RBTree a) a (RBTree a)

We have **two invariants**:

1) **Every red node has a black parent node**

2) **Every path from the root node to a leaf must have the**

**same number of black nodes**

These invariants enforce that the tree is imbalanced by at most a

factor of 2 in one of the branches. Grants fast searching.

The insert x function inserts a

new red leaf at the bottom of

the tree that contains x. We

recursively call the go function

on the appropriate subtree

until an empty node is found.

Every node along the path to

that leaf is balanced by

applying the balance function.

To ensure that the parent

node is not red, the blacken

function is applied to the final

result.

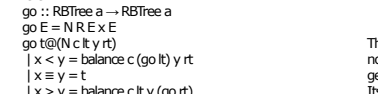


Figure 13.1: Valid Red-Black trees

Figure 13.2: Invalid Red-Black trees

instance Poset RBTree where

insert :: Ord a => a → RBTree a → RBTree a

insert :: Ord a => a → RBTree a → RBTree a

insert x t@(N c l x r) =

where

go :: RBTree a → RBTree a

go E = N R x E

go t@(N c l y r) =

| x < y = balance c (go l) y r

| x == y = t

| x > y = balance c l y (go r)

blacken :: RBTree a → RBTree a

blacken (N R l x r) = N B l x r

blacken t = t

Blacken is only ever applied to the result of go t, so it will only ever

affect the root node, enforcing variant 1 for that node. The

treebalance function balances the tree by ensuring no red nodes with

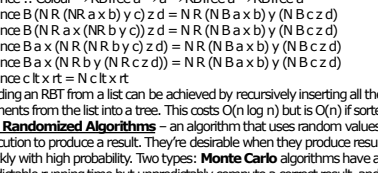
red children. Assuming that the tree is valid to start with, the only

new red node will have been inserted at one of the leaves. The

innermost application of balance will be able to fix a potential red

conflict, but may itself create a new red node that needs fixing

by the next call to balance. 4 cases for balance:



We do this with pattern matching all 4 cases:

balance :: Colour → RBTree a → a → RBTree a → RBTree a

balance B (N R (N R a x b) y) c z d = N R (N B a x b) y (N B c z d)

balance B (N R a x (N R b y c) z) d = N R (N B a x b) y (N B c z d)

balance B a x (N R (N R b y c) z) d = N R (N B a x b) y (N B c z d)

balance B a x (N R b y (N R c z d)) = N R (N B a x b) y (N B c z d)

balance c k x r t = N c k x r

Building an RBT from a list can be achieved by recursively inserting all the

elements from the list into a tree. This costs O(n log n) but is O(n) if sorted list

**12) Randomized Algorithms** - an algorithm that uses random values in its

execution to produce a result. They're desirable when they produce results

quickly with high probability. Two types: **Monte Carlo** algorithms have a

predictable running time but unpredictably compute a correct result, and **Las**

**Vegas** algorithms have an unpredictable running time, but predictably compute

a correct result. Leibniz's law (identity of indiscernibles): x = y ⇔ f x = f y.

holds true for any function. Thus the idea for random value

generation is to start with a seed value, from which a random

value and a new seed can be extracted. Seed values have

type StdGen and can be created with mkStdGen function:

mkStdGen :: Int → StdGen

mkStdGen :: Int → StdGen Once created this can be passed into

the random function.

random :: StdGen → (Int, StdGen)

We can easily produce a list of randoms by just calling random seed, and then

doing x :> randoms seed. We can do other types by implementing this interface:

class Random a where

randomR :: StdGen → (a, StdGen)

randomR :: StdGen → (a, StdGen)

randomRs :: (a, a) → StdGen → [a]

**Randomized Pi** - to compute pi, we can generate a random coordinate pair

(range 0 to 1), and see if the Pythagorean distance from 0 is less than 1. If it is

then we have a hit. We compute the num hits / sample size \* 4, to get pi. (as

the ratio of circlesize/squaresize = pi/4).

The important point to note here is that the variables seed, 'seed', and 'seed'

must be carefully scheduled to happen sequentially. Inside checks if we're inside

the circle:

inside :: (Double, Double) → Bool

inside (x, y) = x \* x + y \* y < 1

montePi :: Double

montePi = loop (mkStdGen 42) samples 0

where

loop :: StdGen → Int → Int → Double

loop seed 0 m = 4 \* fromIntegral m / fromIntegral samples

loop seed n m =

let (x, seed') = randomR (0, 1) seed

(y, seed'') = randomR (0, 1) seed'

m' = if inside (x, y) then m + 1 else m

n' = n - 1

in loop seed'' n' m'

samples :: Int

samples = 10000

**12.2) Sequencing Random Generators:** Threading seeds around can be

error prone, so instead we can handle seed gen automatically by using a

context m. The key change in the following code is the use of the do

keyword, which indicates that the following block of code is to be executed

sequentially, one line at a time:

montePi' :: MonadRandom m => m Double

montePi' = loop 100000000 0

where

loop :: MonadRandom m => Int → Int → m Double

loop 0 m = return (4 \* fromIntegral m / fromIntegral samples)

loop n m = do

x <- getRandomR (0, 1)

y <- getRandomR (0, 1)

let m' = if inside (x, y) then m + 1 else m

n' = n - 1

loop n' m

The base case has a return, and the assignment of x and y is through special

notation that indicates they're of a sequential operation getRandomR (0, 1):

getRandomR :: MonadRandom m => (Int, Int) → m Int

Its pure counterpart is randomR. Since we're wrapping a type into a higher

level monad type, all random functions have a monad version:

class Monad m => MonadRandom m where

getRandom :: Random a => m a

getRandoms :: Random a => m [a]

getRandomRs :: Random a => (a, a) → m [a]

(remember a monad consists of a type constructor that takes a type as

argument and returns a context: for the monad computations which can

involve side effects or state, a unit function - return or pure which takes a

type and wraps it in the monad lifting it to the monadic context, and a bind

function that takes a monad and transforms its inner value and returns a new

monad - allowing for sequencing of computations.) A different approach is to

use **12.3) Random Streams** - In this version, all of the random values are

generated before being transformed into an appropriate sample. In this

version, we use the randomRs function and all rand vals are generated

before transformation