

## Biostatistics 615/815 Lecture 9: Dynamic Programming

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February 3rd, 2011

## Recap: Hash Tables

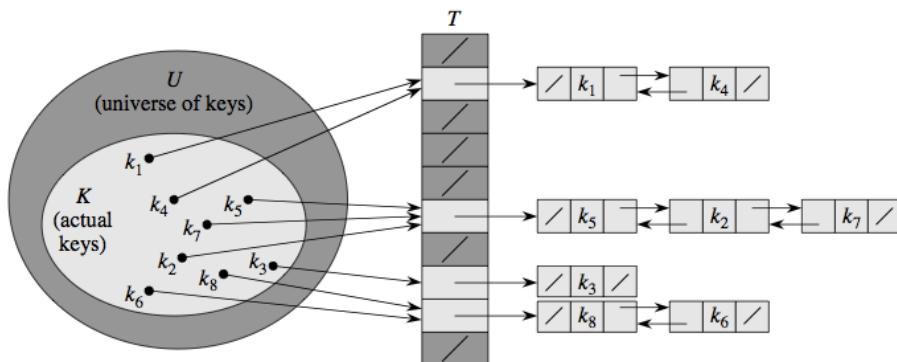
### Key features

- $\Theta(1)$  complexity for INSERT, SEARCH, and REMOVE
- Requires large memory space than the actual content for maintaining good performance
- But uses much smaller memory than direct-address tables

### Key components

- Hash function
  - $h(x.key)$  mapping key onto smaller 'addressable' space  $H$
  - Total required memory is the possible number of hash values
  - Good hash function minimize the possibility of key collisions
- Collision-resolution strategy, when  $h(k_1) = h(k_2)$ .

## Recap: Illustration of CHAINEDHASH



## Recap : Open hash

### Probing strategies

- Linear probing
- Quadratic probing
- Double hashing

### Double Hashing

- $h(k, i) = (h_1(k) + ih_2(k)) \bmod m$
- The probe sequence depends in two ways upon  $k$ .
- For example,  $h_1(k) = k \bmod m$ ,  $h_2(k) = 1 + (k \bmod m')$
- Avoid clustering problem
- Performance close to ideal scheme of uniform hashing.

## Today

## Dynamic Programming

- Fibonacci numbers
- Manhattan tourist problems
- Edit distance problem

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## A divide-and-conquer algorithms for Fibonacci numbers

## Fibonacci numbers

$$F_n = \begin{cases} F_{n-1} + F_{n-2} & n > 1 \\ 1 & n = 1 \\ 0 & n = 0 \end{cases}$$

## A recursive implementation of fibonacci numbers

```
int fibonacci(int n) {
    if ( n < 2 ) return n;
    else return fibonacci(n-1)+fibonacci(n-2);
}
```

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## Recap: Divide and conquer algorithms

## Good examples of divide and conquer algorithms

- TOWEROFHANOI
- MERGESORT
- QUICKSORT
- BINARYSEARCHTREE algorithms

These algorithms divide a problem into smaller and disjoint subproblems until they become trivial.

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## Performance of recursive FIBONACCI

## Computational time

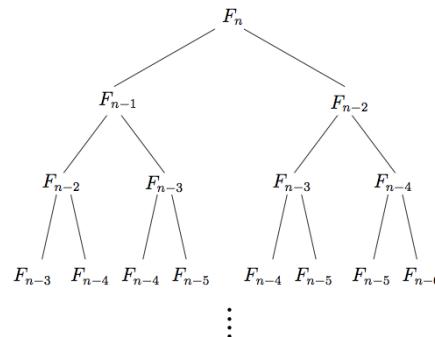
- 4.4 seconds for calculating  $F_{40}$
- 49 seconds for calculating  $F_{45}$
- $\infty$  seconds for calculating  $F_{100}$ !

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## What is happening in the recursive FIBONACCI



## A non-redundant FIBONACCI

```

int fibonacci(int n) {
    int* fibs = new int[n+1];
    fibs[0] = 0;
    fibs[1] = 1;
    for(int i=2; i <= n; ++i) {
        fibs[i] = fibs[i-1]+fibs[i-2];
    }
    int ret = fibs[n];
    delete [] fibs;
    return ret;
}
  
```

## Time complexity of redundant FIBONACCI

$$\begin{aligned}
 T(n) &= T(n-1) + T(n-2) \\
 T(1) &= 1 \\
 T(0) &= 1 \\
 T(n) &= F_{n+1}
 \end{aligned}$$

The time complexity is exponential

## Key idea in non-redundant FIBONACCI

- Each  $F_n$  will be reused to calculate  $F_{n+1}$  and  $F_{n+2}$
- Store  $F_n$  into an array so that we don't have to recalculate it

## A recursive, but non-redundant FIBONACCI

```
int fibonacci(int* fibs, int n) {
    if ( fibs[n] > 0 ) {
        return fibs[n]; // reuse stored solution if available
    }
    else if ( n < 2 ) {
        return n; // terminal condition
    }
    fibs[n] = fibonacci(n-1) + fibonacci(n-2); // store the solution once computed
    return fibs[n];
}
```

## Steps of dynamic programming

- Characterize the structure of an (optimal) solution
- Recursively define the value of an (optimal) solution
- Compute the value of an (optimal) solution, typically in a bottom-up fashion
- Construct an optimal solution from computed information.

## Dynamic programming

### Key components of dynamic programming

- Problems that can be divided into subproblems
- Overlapping subproblems - subproblems share subsubproblems
- Solves each subsubproblem just once and then saves its answer

### Why dynamic programming?

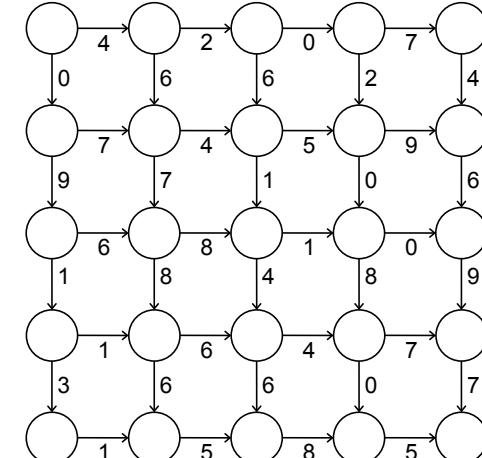
According to wikipedia... "The word 'dynamic' was chosen because it sounded impressive, not because how the method works"

### Examples of dynamic programming

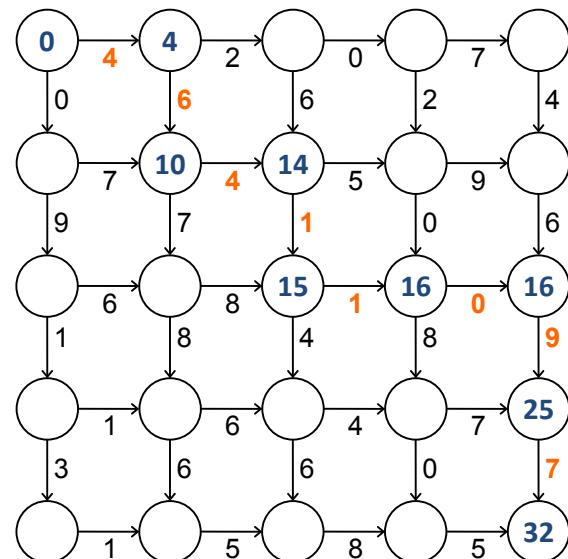
- Shortest path finding algorithms
- DNA sequence alignment
- Hidden markov models

## The Manhattan tourist problem

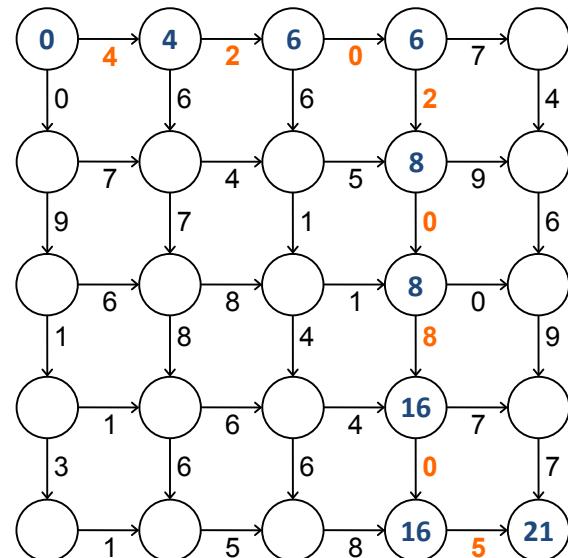
Find the cost-optimal path from left-top corner to right-bottom corner



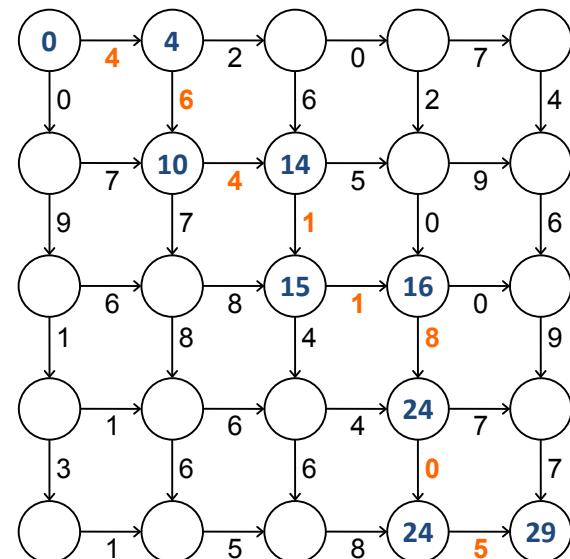
## One possible (but not optimal) solution



## And here comes an optimal solution



## A slightly better, but still not an optimal solution



## A brute-force algorithm

## Algorithm BRUTEFORCEMTP

- ① Enumerate all the possible paths
- ② Calculate the cost of each possible path
- ③ Pick the path that produces a minimum cost

## Time complexity

- Number of possible paths are  $\binom{n_r+n_c}{n_r}$
- Super-exponential growth when  $n_r$  and  $n_c$  are similar.



## Implementing Manhattan tourist algorithm

```
template<class T>
class Matrix { // Matrix data type to store the costs
    T* data; // internal data as one-dimensional array
    int nr, nc; // # rows and # cols
    Matrix(const Matrix<T>& m) {}; // prevent copy
public:
    Matrix(int nrows, int ncols) : nr(nrows), nc(ncols) {
        data = new T[nrows*ncols](); // initialize matrix
    }
    ~Matrix() {
        if ( data != NULL ) delete [] data;
    }
    // accessor function : possible to use to read/write elements
    // value1 = M.at(i,j);
    // M.at(i,j) = value2;
    T& at(int r, int c) { return data[r*nc+c]; }
    void print(); // print the content of the matrix (omitted)
};
```

## Calculating optimal cost

```
// hw, vw : horizontal and vertical input weights
// cost : stored optimal cost from (0,0) to (r,c)
// move : stored optimal decision to reach (r,c)
// r,c : the position of interest
int optimalCost(Matrix<int>& hw, Matrix<int>& vw,
    Matrix<int>& cost, Matrix<int>& move, int r, int c) {
    // if cost is stored already, skip the cost evaluation
    if ( cost.at(r,c) == 0 ) {
        if ( ( r == 0 ) && ( c == 0 ) ) cost.at(r,c) = 0; // terminal condition
        else if ( r == 0 ) { // only horizontal move is possible
            move.at(r,c) = 0; // 0 means horitontal move to (r,c)
            cost.at(r,c) = optimalCost(hw,vw,cost,move,r,c-1) + hw.at(r,c-1);
        }
        else if ( c == 0 ) { // only vertical move is possible
            move.at(r,c) = 1; // 1 means vertical move to (r,c)
            cost.at(r,c) = optimalCost(hw,vw,cost,move,r-1,c) + vw.at(r-1,c);
        }
    }
```

## Manhattan tourist problem : main()

```
int main(int argc, char** argv) {
    int nrows = 5, ncols = 5;
    Matrix<int> hw(nrows,ncols-1), vw(nrows-1,ncols); // weight matrices
    hw.at(0,0) = 4; hw.at(0,1) = 2; ... // initialize horizontal weights
    vw.at(0,0) = 0; vw.at(0,1) = 6; ... // initialize vertical weights

    // optimal costs and decisions for backtracking
    Matrix<int> cost(nrows,ncols), move(nrows,ncols);
    // calculate the optimal cost, recording the backtracking info
    int optCost = optimalCost(hw,vw,cost,move,nrows-1,ncols-1);
    std::cout << "Optimal cost is " << optCost << std::endl;
    // backtrack the stored decision to reconstruct an optimal path
    trackOptimalPath(hw,vw,cost,move,nrows-1,ncols-1);
    return 0;
}
```

## Calculating optimal cost (cont'd)

```
else { // evaluate the cumulative cost of horizontal and vertical move
    int hcost = optimalCost(hw,vw,cost,move,r,c-1) + hw.at(r,c-1);
    int vcost = optimalCost(hw,vw,cost,move,r-1,c) + vw.at(r-1,c);
    if ( hcost > vcost ) { // when vertical move is optimal
        move.at(r,c) = 1; // store the decision
        cost.at(r,c) = vcost; // and store the optimal cost
    }
    else { // when horizontal move is optimal
        move.at(r,c) = 0;
        cost.at(r,c) = hcost;
    }
}
return cost.at(r,c); // return the optimal cost
}
```

## Dynamic programming : A smart recursion

- Dynamic programming is recursion without repetition
  - ① Formulate the problem recursively
  - ② Build solutions to your recurrence from the bottom up
- Dynamic programming is not about filling in tables; it's about smart recursion (Jeff Erickson)

## More examples of edit distance

F	O	O	D							
M	O	N	E	Y						
A	L	G	O	R	I	T	H	M		
A	L		T	R	U	I	S	T	I	C

- Similar representation to DNA sequence alignment
- Does the above alignment provides an optimal edit distance?

## Minimum edit distance problem

### Edit distance

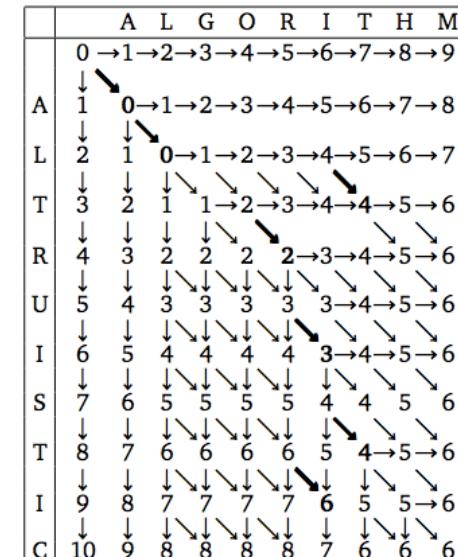
Minimum number of letter insertions, deletions, substitutions required to transform one word into another

### An example

FOOD → MOD → MOND → MONED → MONEY

Edit distance is 4 in the example above

## A dynamic programming solution



## Recursively formulating the problem

- Input strings are  $x[1, \dots, m]$  and  $y[1, \dots, n]$ .
- Let  $x_i = x[1, \dots, i]$  and  $y_j = y[1, \dots, j]$  be substrings of  $x$  and  $y$ .
- Edit distance  $d(x, y)$  can be recursively defined as follows

$$d(x_i, y_j) = \begin{cases} i & j = 0 \\ j & i = 0 \\ \min \left\{ \begin{array}{l} d(x_{i-1}, y_j) + 1 \\ d(x_i, y_{j-1}) + 1 \\ d(x_{i-1}, y_{j-1}) + I(x[i] \neq y[j]) \end{array} \right\} & \text{otherwise} \end{cases}$$

- Similar to the Manhattan tourist problem, but with 3-way choice.
- Time complexity is  $\Theta(mn)$ .

## Summary

### Today

- Dynamic programming is a smart recursion avoiding redundancy
- Divide a problem into subproblems that can be shared
- Examples of dynamic programming
  - Fibonacci numbers
  - Manhattan tourist problem
  - Edit distance problem

### Next lecture

- Algorithms in graphs
  - Using boost library
  - Dijkstra's algorithm (CLRS Chapter 24)
- Introduction to hidden Markov model