# **Greedy Algorithms**

- Given a set of activities (s<sub>i</sub>, f<sub>i</sub>), we want to schedule the maximum number of non-overlapping activities.
- <u>GREEDY-ACTIVITY-SELECTOR</u> (s, f)
  - 1. n = length[s]
  - 2.  $S = \{a_1\}$
  - 3. i = 1
  - 4. for m = 2 to n do 5. if  $s_m$  is not before  $f_i$  then
  - **6**.  $S = S \cup \{a_m\}$
  - 7. i = m
  - 8. return S

### Example

- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14] -- <u>Sorted</u> by finish times
- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14]
- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14]
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- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14]
- [1,4], [3,5], [0,6], [5,7], [3,8], [5,9], [6,10], [8,11], [8,12], [2,13], [12,14]

## Why does it work?

#### • THEOREM

Let A be a set of activities and let  $a_1$  be the activity with the earliest finish time. Then activity  $a_1$  is in some maximum-sized subset of non-overlapping activities.

#### • PROOF

Let S' be a solution that does not contain  $a_1$ . Let  $a'_1$  be the activity with the earliest finish time in S'. Then replacing  $a'_1$  by  $a_1$  gives a solution S of the same size.

Why are we allowed to replace? Why is it of the same size?

### Then apply induction! How?

### Greedy Algorithms – Huffman Coding

#### • Huffman Coding Problem

Example: Release 29.1 of 15-Feb-2005 of <u>TrEMBL</u> Protein Database contains 1,614,107 sequence entries, comprising 505,947,503 amino acids. There are 20 possible amino acids. What is the minimum number of bits to store the compressed database?

~2.5 G bits or 300MB.

- How to improve this?
- <u>Information</u>: Frequencies are not the same.

Ala (A) 7.72	<mark>6ln</mark> (Q) 3.91	Leu (L) 9.56	<mark>Ser</mark> (S) 6.98
Arg (R) 5.24	<mark>6lu</mark> (E) 6.54	Lys (K) 5.96	Thr (T) 5.52
<mark>Asn</mark> (N) 4.28	<mark>Gly</mark> (G) 6.90	Met (M) 2.36	Trp (W) 1.18
Asp (D) 5.28	His (H) 2.26	Phe (F) 4.06	<b>Туг</b> (У) 3.13
Cys (C) 1.60	Ile (I) 5.88	Pro (P) 4.87	Val (V) 6.66

• Idea: Use shorter codes for more frequent amino acids and longer codes for less frequent ones.

### Huffman Coding

2 million characters in file.

A, C, G, T, N, Y, R, S, M

IDEA 1: Use ASCII Code Each need at least 8 bits, Total = 16 M bits = 2 MB	IDEA 3: Use Variable Length Codes A 22 11 T 22 10	How to Decode? Need Unique decoding! Easy for Ideas 1 & 2. What about Idea 32	
IDEA 2: Use 4-bit Codes Each need at least 4 bits, Total = 8 M bits = 1 MB	C 18 011 G 18 010 M 10 001	11010110111001000110000000110	
Percentage Frequencies	Y 5 00011   R 4 00010   S 4 00001   M 3 00000	11010110111001000110000000110	
<b>2 million characters in file</b> . Length = ?			

Expected length = ?

Sum up products of frequency times the code length, i.e.,

(.22x2 + .22x2 + .18x3 + .18x3 + .10x3 + .05x5 + .04x5 + .04x5 + .03x5 ) x 2 M bits =

3.24 M bits = .4 MB