# **Best-First Search Minimizing Space or Time**

#### **RBFS**

Take more space than IDA\*
Take less time than IDA\*

♦ Similar to A\* algorithm developed for heuristic search

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  - » Both are recursive in the same sense

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Difference between A\* and RBFS

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- Difference between A\* and RBFS
  - » A\* keeps in memory all of the already generated nodes
  - » RBFS only keeps the current search path and the sibling nodes along the path

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- » What is the space complexity?
  - > Linear with depth of the search
    - Same as IDA\*

# **RBFS** memory

» When RBFS suspends searching a subtree, what does it remember?

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#### RBFS memory – 2

» When RBFS suspends searching a subtree, what does it remember?

> An updated f-value of the root of the subtree

# **Updated f-values**

» How does RBFS update the f-values?

# **Updated f-values – 2**

- » How does RBFS update the f-values?
  - > Backing up the f-values in the same way as A\* does

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#### f-value notation

- ♦ Static f-value
  - >> f(N)
    - > Value returned by the evaluation function
    - > Always the same

#### f-value notation – 2

- Static f-value
  - >> f(N)
    - > Value returned by the evaluation function
    - > Always the same
- Backed-up value
  - >> **F(N)** 
    - > Changes during the search
      - Depends upon descendants of N

RBFS backs up f-values in the same way as A\*

» How is F(N) defined?

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RBFS backs up f-values in the same way as A\*

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> If N has never been expanded?

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- RBFS backs up f-values in the same way as A\*
  - » How is F(N) defined?
    - > If N has never been expanded?
      - F(N) = f(N)
    - > If N has been expanded?
      - $F(N) = \min (F(S_j))$
      - Where  $S_j$  are the subtrees of N

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- » How is the bound determined?

# **RBFS** subtree exploration – 4

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- » How is the bound determined?
  - > From the F-values of the siblings along the current search path

- » How does RBFS explore subtrees?
  - > As in A\*, within a given f-bound
- » How is the bound determined?
  - > From the F-values of the siblings along the current search path
  - > The smallest F-value
    - The closest competitor

Suppose N is currently the best node

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> N is expanded

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  - > N's children are expanded

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    - > **F**(**N**) > **Bound**
  - » Then what happens?

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    - > **F(N)** > **Bound**
  - » Then what happens?
    - > Nodes below N are forgotten

### **Subtree exploration – 12**

- Suppose N is currently the best node
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  - » Until when?
    - > **F(N)** > **Bound**
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    - > N's F-value is updated

### **Subtree exploration – 13**

- Suppose N is currently the best node
  - > N is expanded
  - > N's children are expanded
  - » Until when?
    - > F(N) > Bound
  - » Then what happens?
    - > Nodes below N are forgotten
    - > N's F-value is updated
    - > RBFS selects which node to expand next

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- Suppose a child N<sub>k</sub> of N is generated again

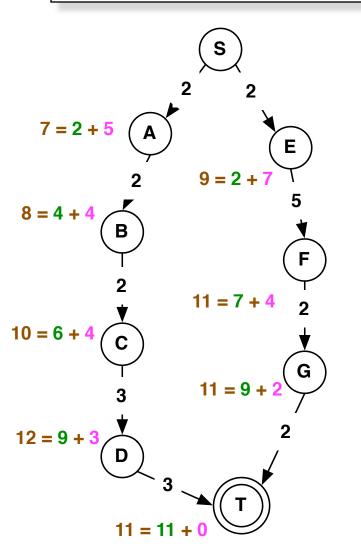
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- Suppose a child N<sub>k</sub> of N is generated again
  - » Compute f(N<sub>k</sub>)

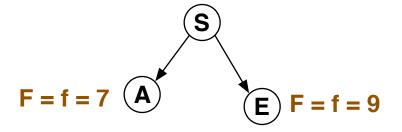
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    - > N<sub>k</sub>'s F-value can be inherited from N
      - $-N_k$  was generated earlier
      - $F(N_k)$  was ≥ F(N), otherwise F(N) would be smaller

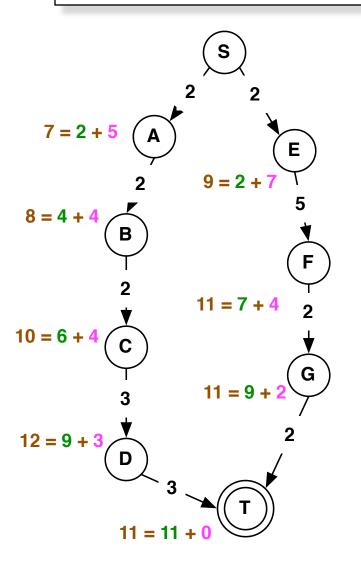


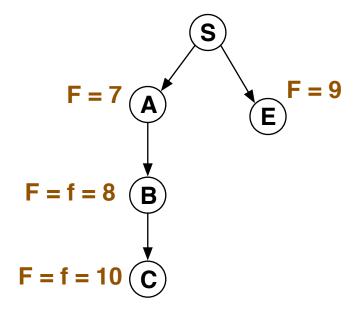


S is expanded

A is found to be the best child

f(n) in mocha = g(n) in clover + h(n) in magenta



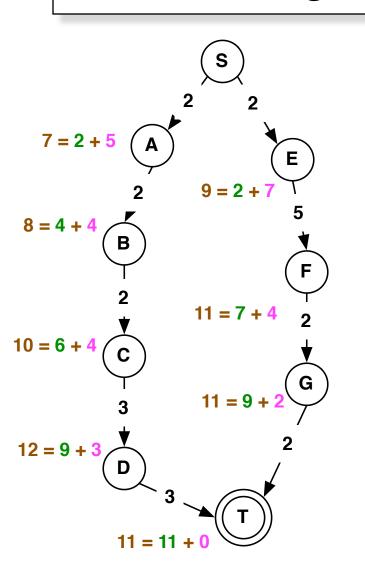


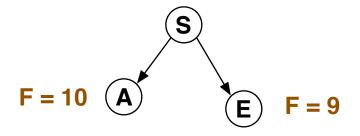
A is expanded with bound 9

C has F-value 10

Stop expansion, backup F value

f(n) in mocha = g(n) in clover + h(n) in magenta



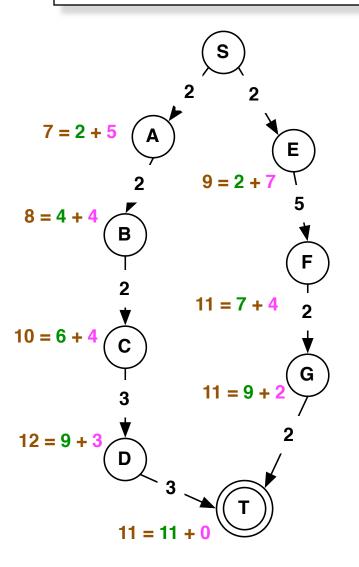


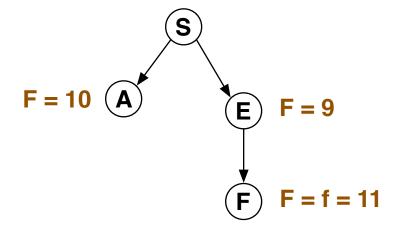
Forget expansion from A

A has backed up F value 10

E is best to expand next

f(n) in mocha = g(n) in clover + h(n) in magenta



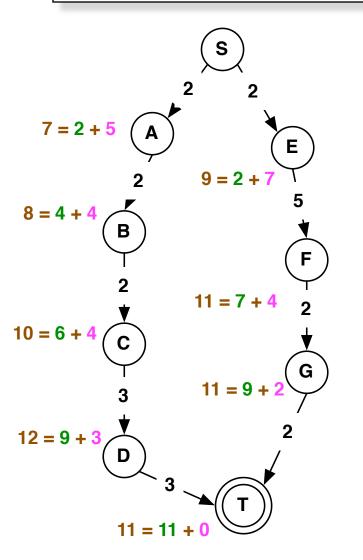


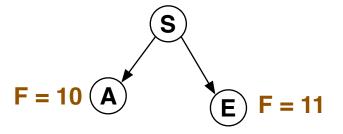
E is expanded with bound 10

F has F-value 11

Stop expansion, backup F value

f(n) in mocha = g(n) in clover + h(n) in magenta



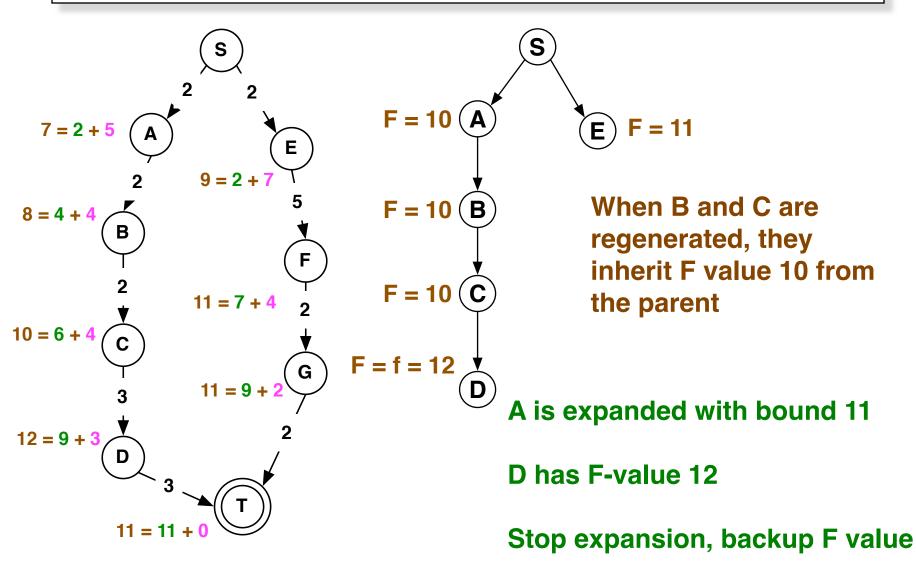


Forget expansion from E

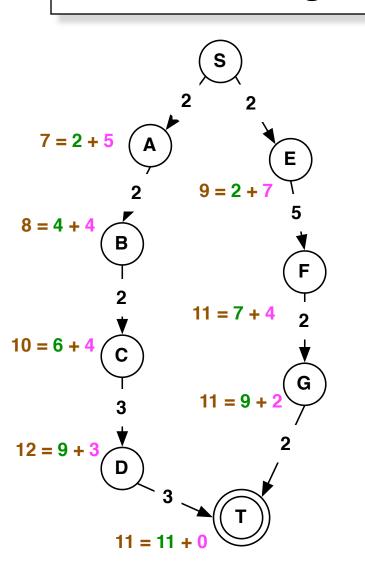
E has backed up F value 11

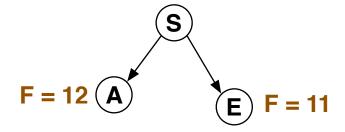
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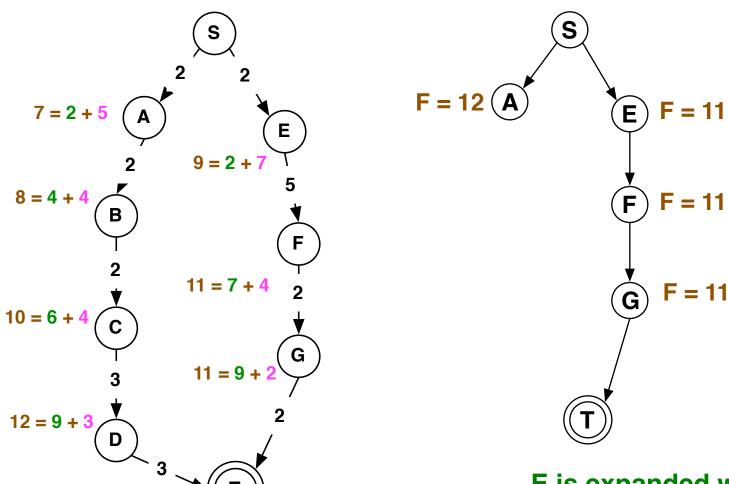


Forget expansion from A

A has backed up F value 12

E is best to expand next

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11 = 11 +

E is expanded with bound 12

Reach goal, search ends

### **Algorithm**

```
function NewF (N, F(N), Bound)
  if F(N) > Bound then NewF := F(N)
  else if goal(N) then exit search with success
  else if N has no children then NewF := infinity - dead end
  else for each child N<sub>k</sub> of N do
        if f(N) < F(N) then F(N_k) := max(F(N), f(N_k))
                      else F(N_k) := f(N_k)
        sort children N<sub>k</sub> in increasing order of F-value
        while F(N_1) \leq Bound and F(N_1) < infinity do
            Bound1 := min (Bound, F-value of sibling N_1)
            F(N_1) := NewF(N_1, F(N_1), Bound1)
            reorder nodes N_1, N_2, \dots according to new F(N_1)
           end
        end
        NewF := F(N_1)
```

### **Summary RBFS properties**

- Space complexity
  - » Linear in depth of search
- Cost of minimizing space
  - » Regenerate previously generated nodes
    - > Overhead substantially less than IDA\*

- Expands nodes in best-order
  - » Like A\*, even with non-monotonic f-function
  - » Unlike IDA\*, which requires a monotonic f-function