Algorithms and Data Structures 2 Recap Lectures 17-18

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Topics we covered so far

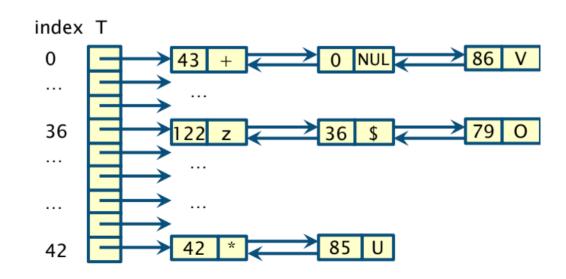
- •Map ADT
- **Direct-address tables**
- Hash table data structure
- Hash functions
- Collision resolution by chaining
- Collision resolution by open addressing
 - Linear probing
 - Quadratic probing
 - Double hashing
- Perfect hashing

Question 1

 Briefly describe collision resolution by chaining and open addressing in the hash table data structure

- In collision resolution by chaining, all elements that hash to the same slot are stored in a linked list (called chain)
 - Doubly linked list to support fast deletion
 - Usually, no check is performed to prevent the insertion of elements with duplicate keys
 - Some implementations store the first element of each list directly in T
 - Operations are based on the standard operations for doubly linked lists
- Example

```
CHAINED-HASH-INSERT(T,x)
insert x at the head of list T[h(x.key)]
```



- In collision resolution by open addressing, alternative cells are probed until an empty cell is found
- To determine which slots to probe, the hash function is extended to include the probe number (starting from 0) as a second input: h: U x {0,1, ..., m-1} → {0,1,..., m-1}
 - Linear probing
 - Quadratic probing
 - Double hashing
- With open addressing, we require that every hash-table position is eventually considered as a slot for a new key as the table fills up

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- Linear probing, uses a hash function of the form $h(k,i) = (h'(k) + i) \mod m$ where h' is an ordinary hash function
- Linear probing is easy to implement, but it suffers from primary clustering
 - Long runs of occupied slots build up, increasing the average search time

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- Quadratic probing uses a hash function of the form $h(k,i) = (h'(k) + c_1i + c_2i^2)$ mod m where h' is an ordinary hash function and c_1, c_2 constants
- If two keys have the same initial probe position, then their probe sequences are the same
 - This leads to a milder form of clustering, called secondary clustering
 - In general, better than linear probing

 There is no guarantee of finding an empty cell once the table gets more than half full

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• Double hashing uses a hash function of the form $h(k,i) = (h_1(k) + ih_2(k))$ mod m where h_1 and h_2 are ordinary hash functions

- The value h₂(k) must be relatively prime to the hash-table size m for the entire hash table to be searched
 - 1. Let m be a power of 2 and design h₂ to always produce an odd number
 - 2. Let m be prime and design h₂ to always return a positive integer less than m
- One of the best methods available for open addressing

Question 2

- Draw a representation of a 7-celled hash table T and its contents after the hash function $h(x) = (3x+1) \mod 7$ has been used to insert the elements in the sequence 100,19,3,5,4,7 into T, assuming collisions are handled using
 - a) the chaining method
 - b) linear probing

- $h(x) = (3x+1) \mod 7$
- insert the elements in the sequence 100,19,3,5,4,7 into H
- assuming collisions are handled using chaining

Hashes

h(100) = 0

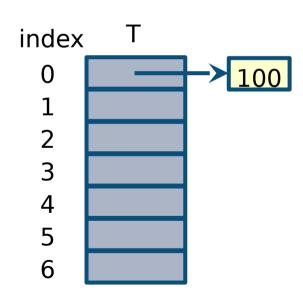
h(19) = 2

h(3) = 3

h(5) = 2

h(4) = 6

h(7) = 1



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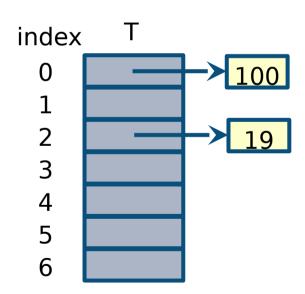
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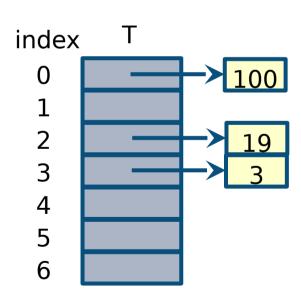
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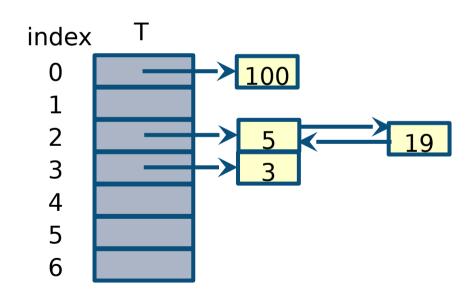
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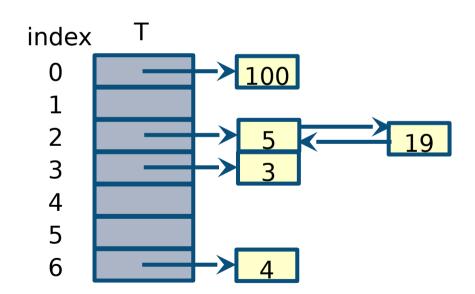
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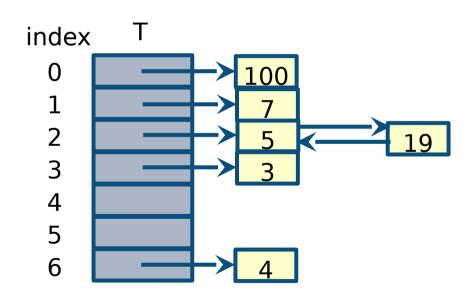
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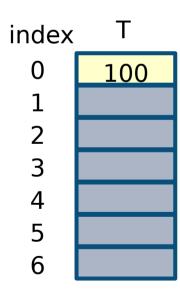
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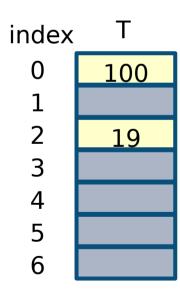
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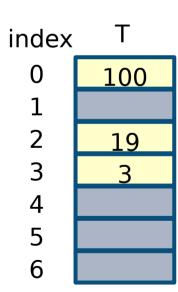
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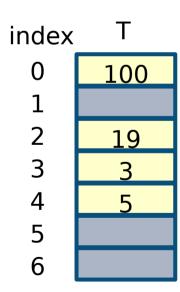
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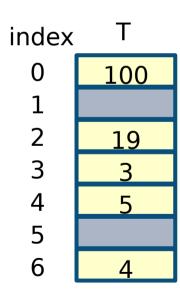
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