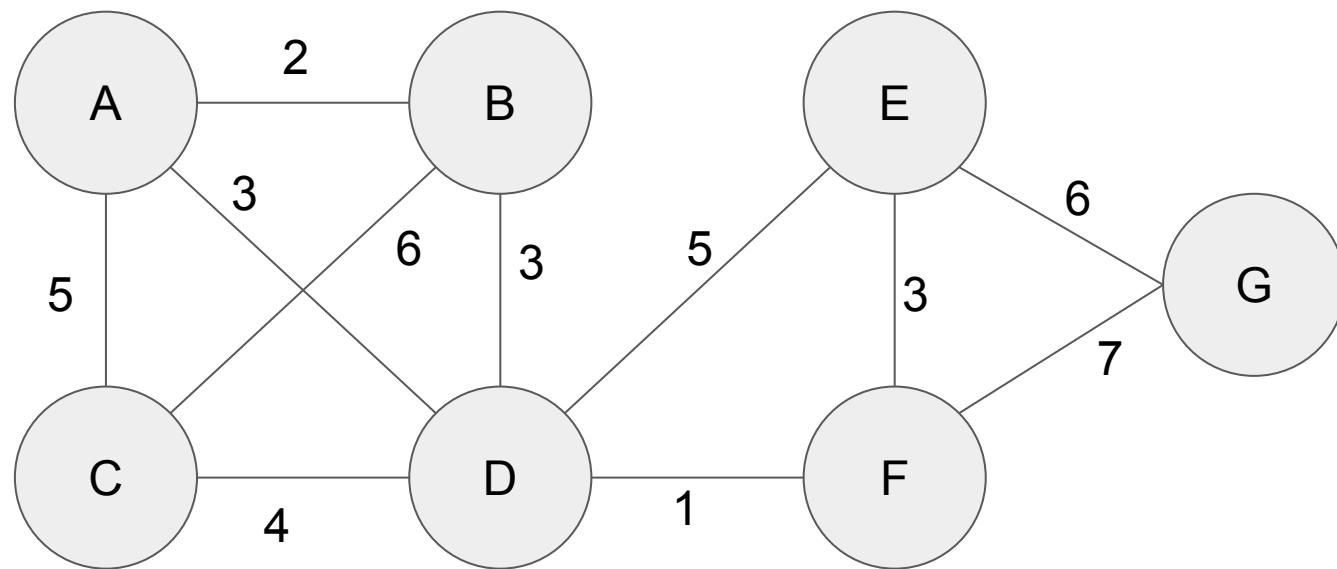


Greedy algorithms

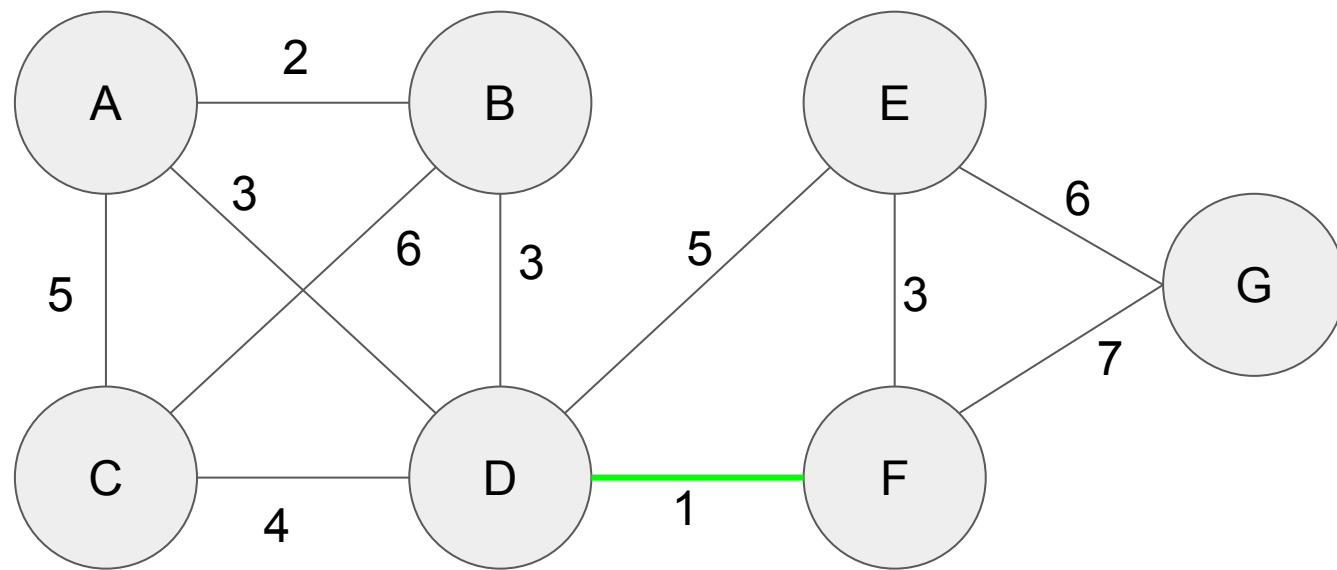
CMPSC 465 - Yana Safonova

MST and clustering

Kruskal's algorithm aka single-linkage clustering

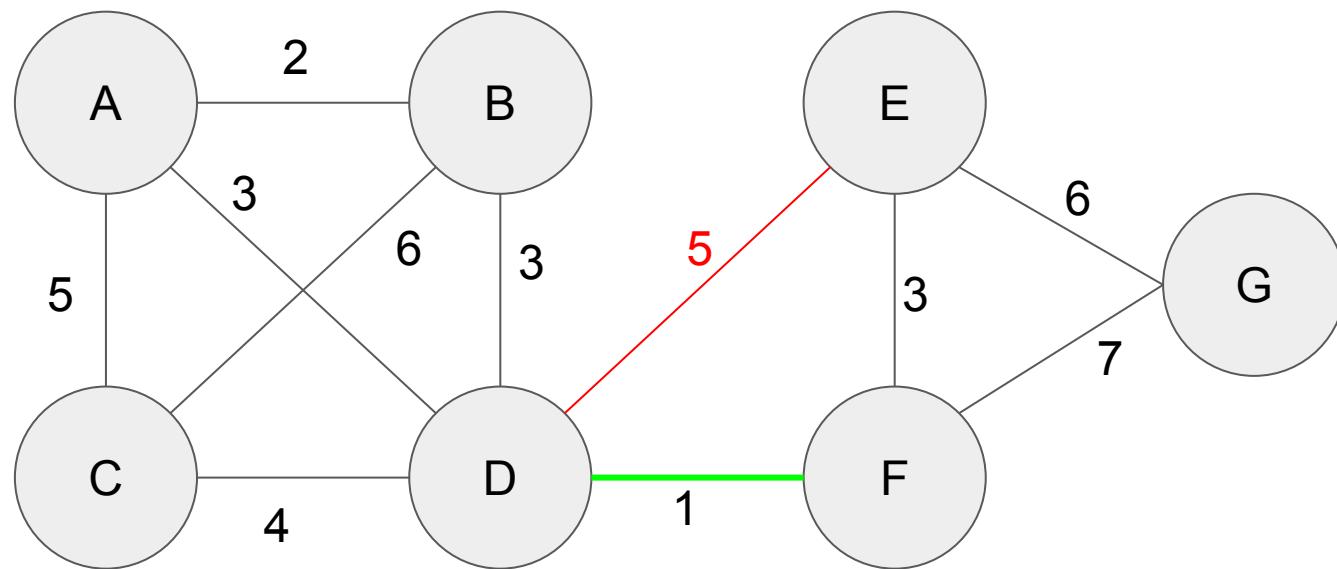


Kruskal's algorithm - weight 1



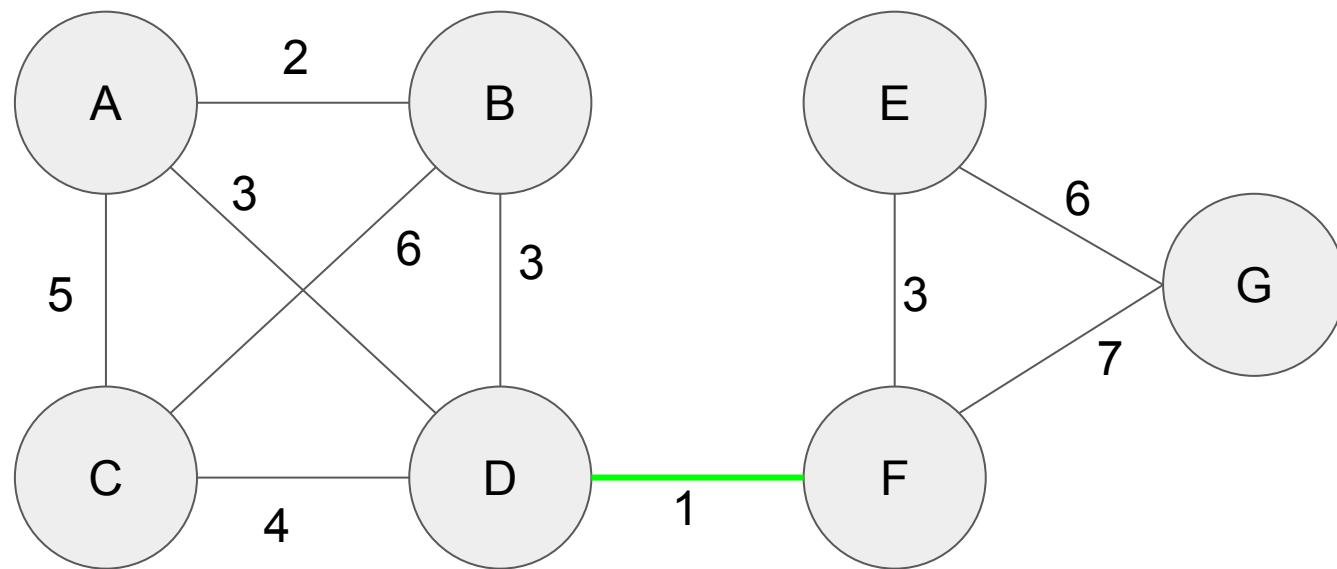
D F

Kruskal's algorithm - weight 1



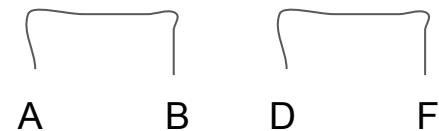
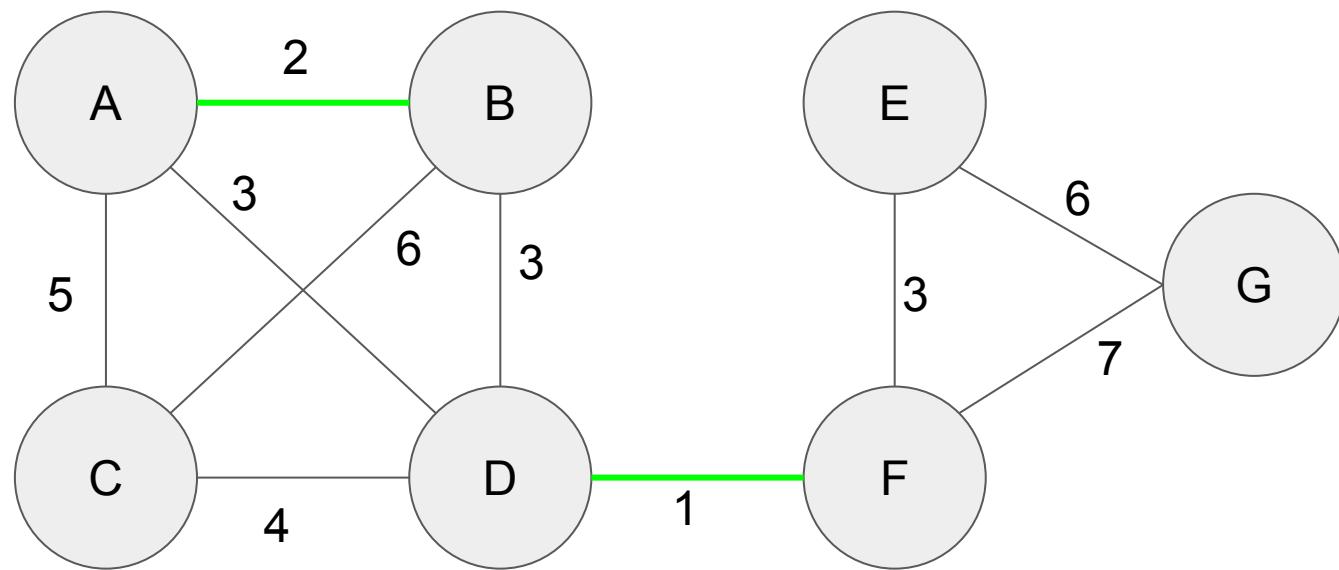
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Kruskal's algorithm - weight 1

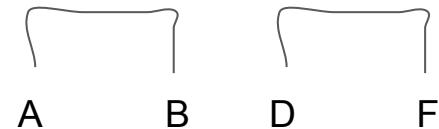
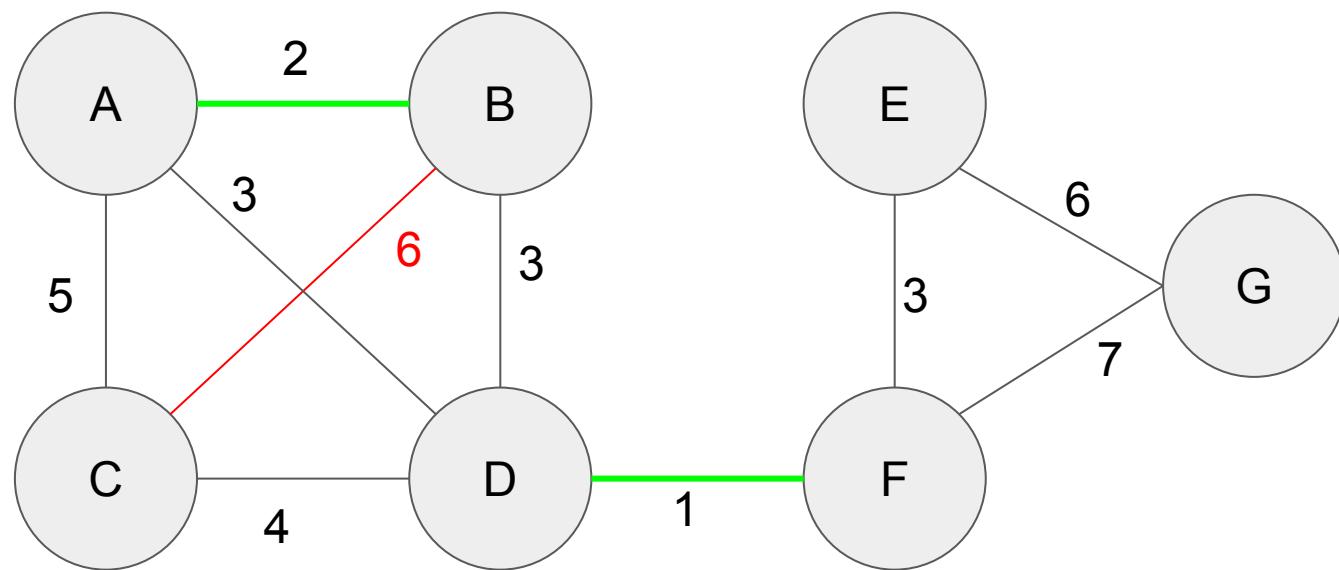


D F

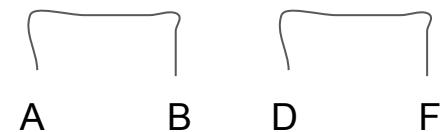
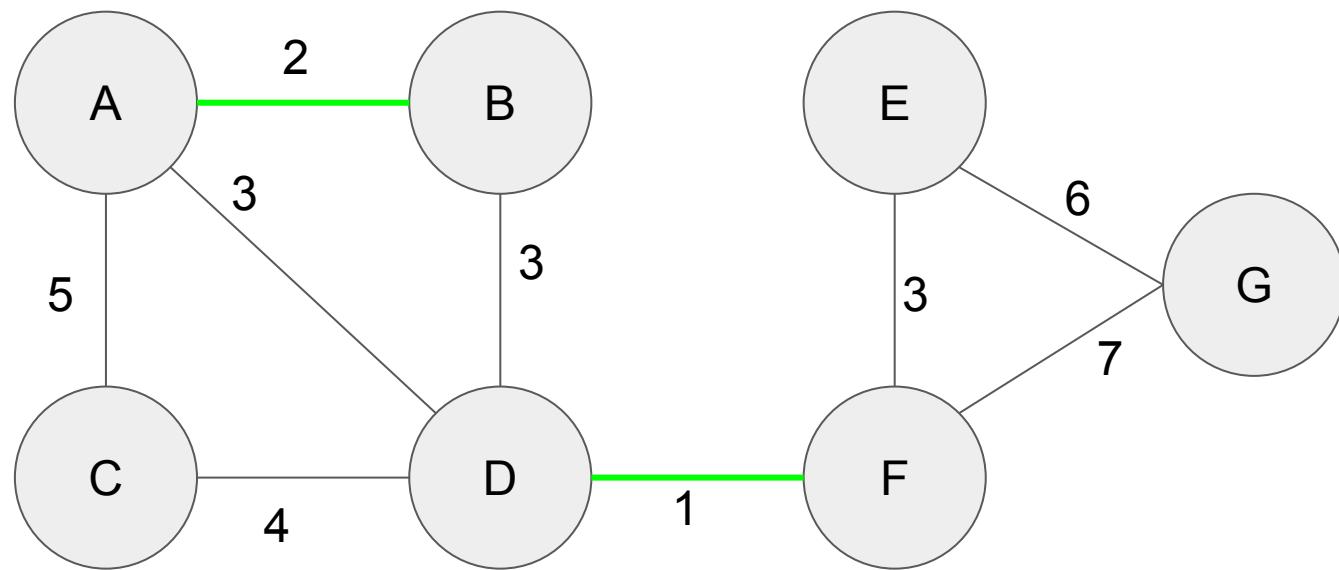
Kruskal's algorithm - weight 2



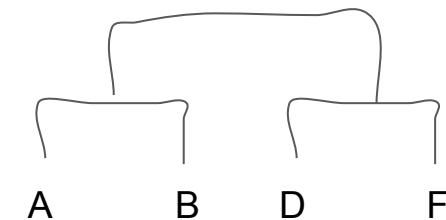
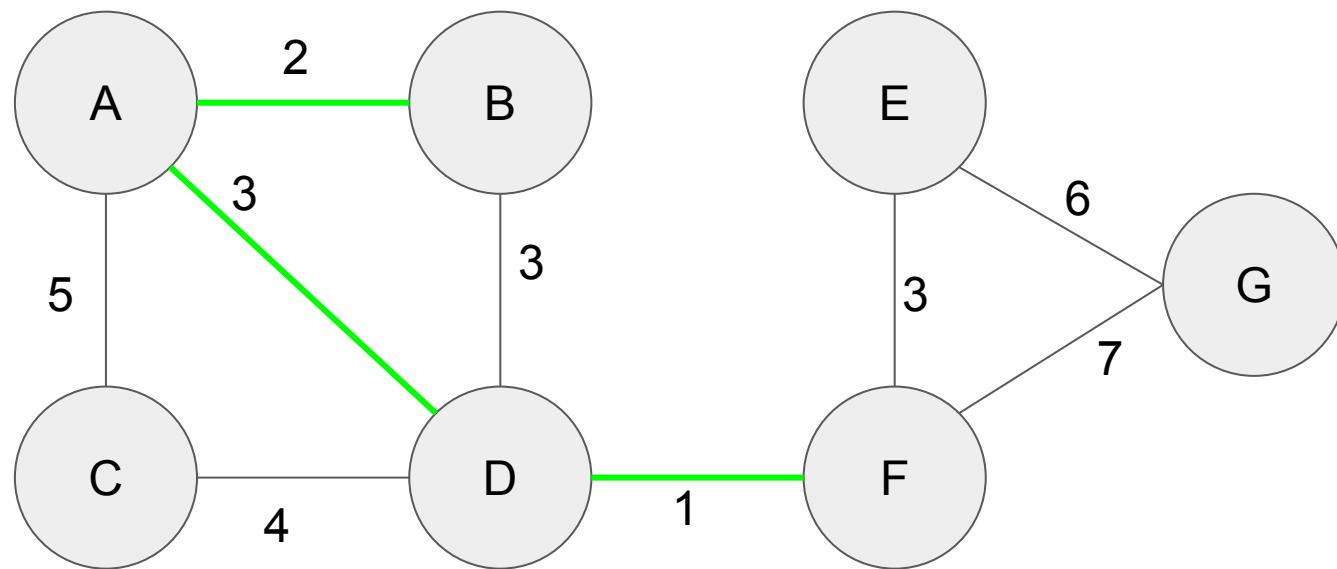
Kruskal's algorithm - weight 2



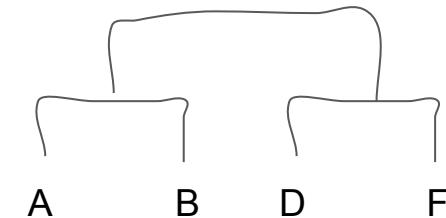
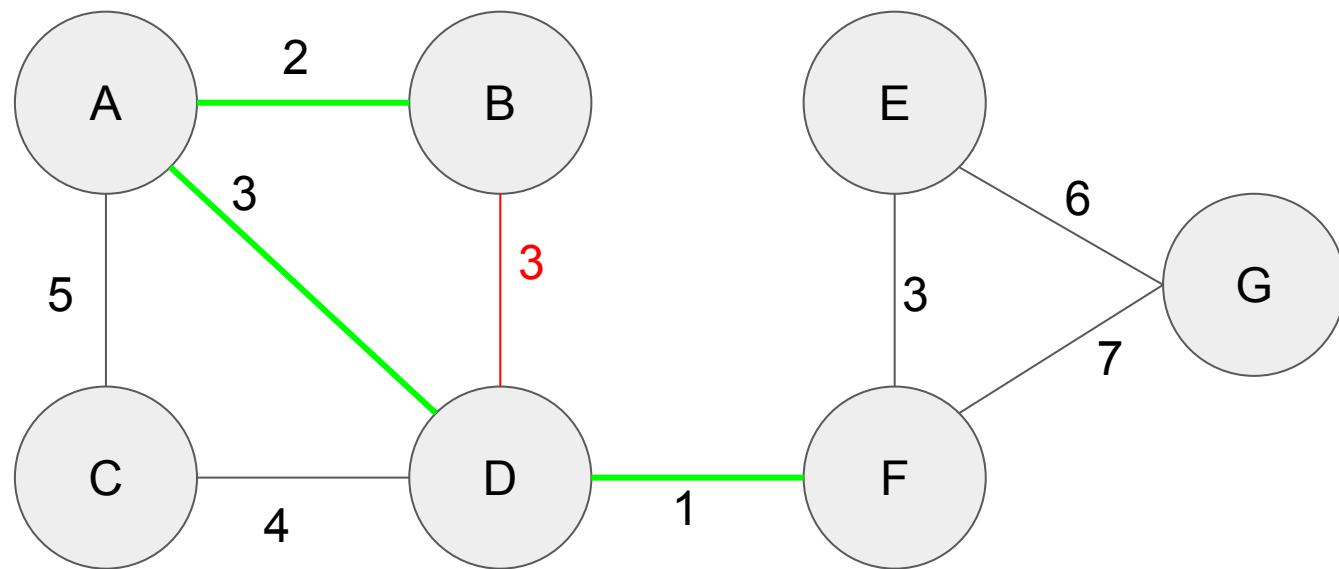
Kruskal's algorithm - weight 2



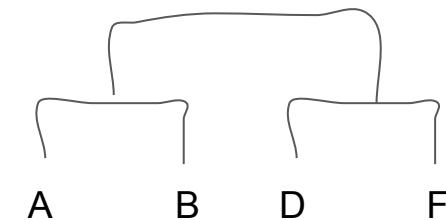
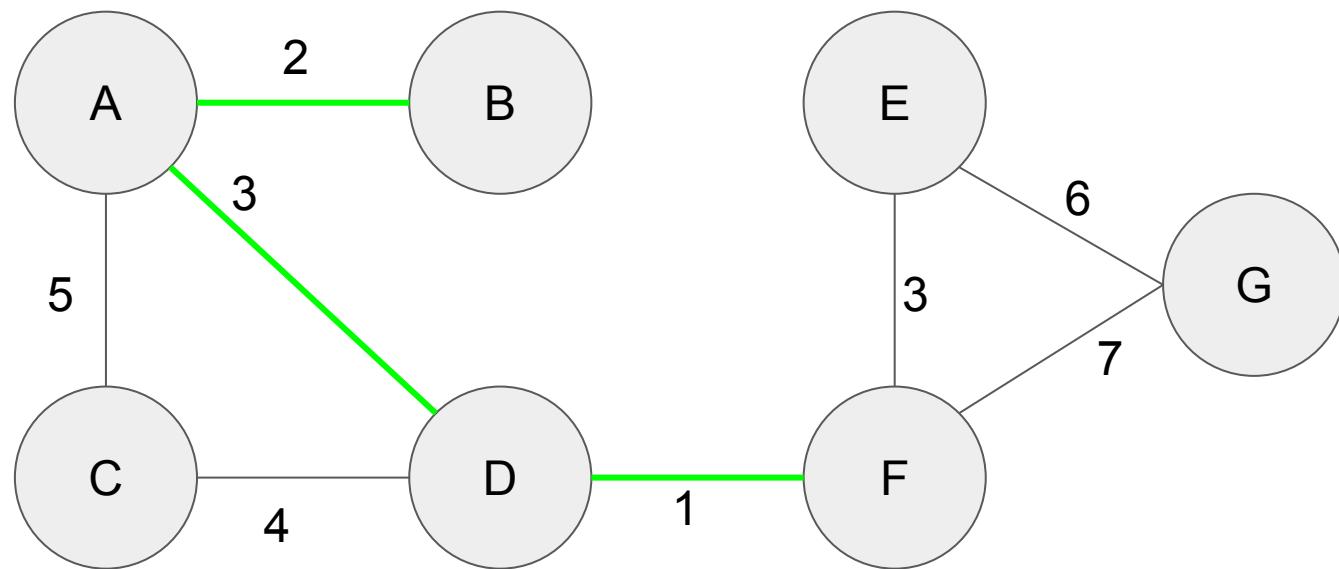
Kruskal's algorithm - weight 3



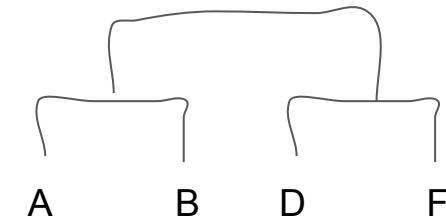
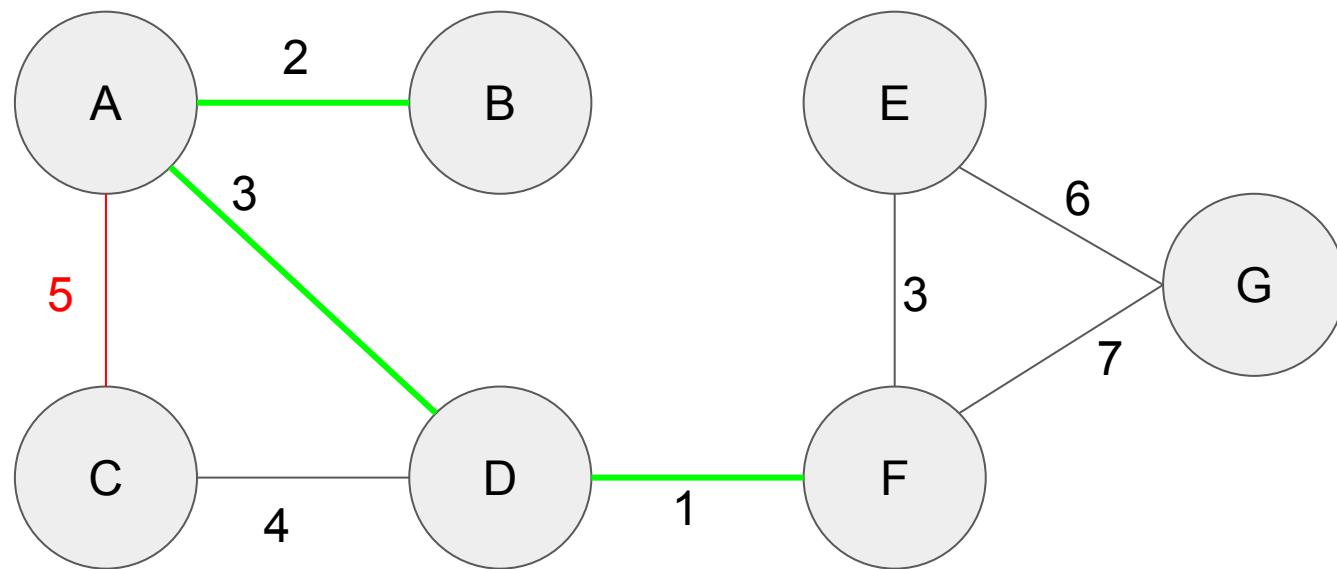
Kruskal's algorithm - weight 3



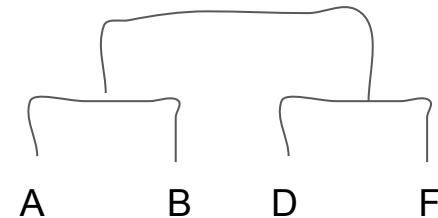
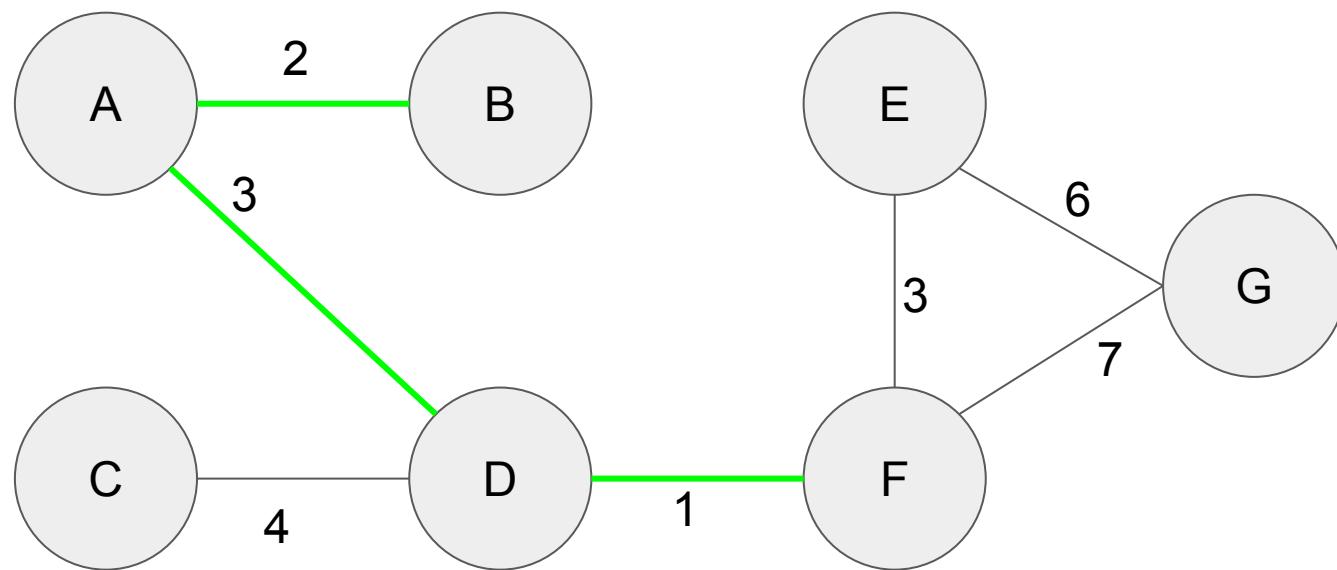
Kruskal's algorithm - weight 3



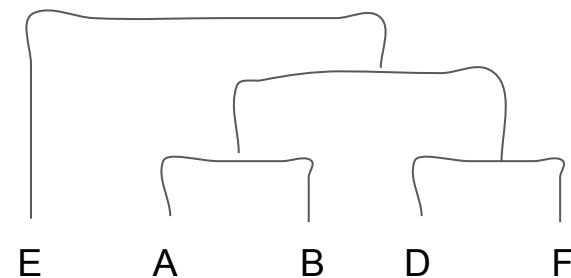
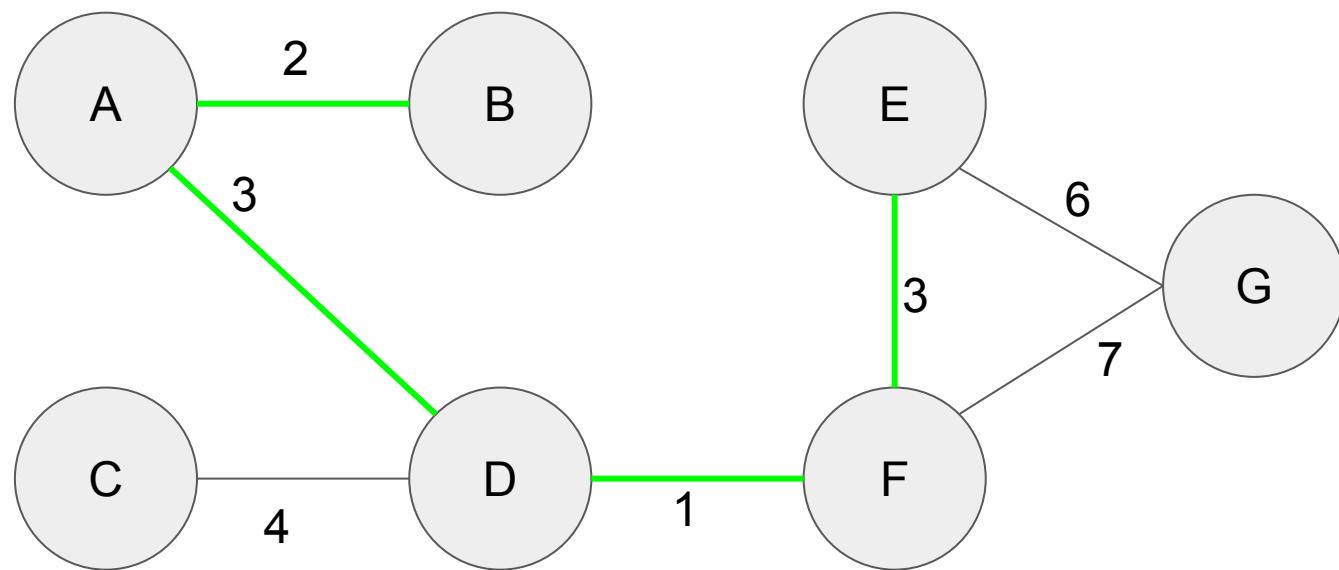
Kruskal's algorithm - weight 3



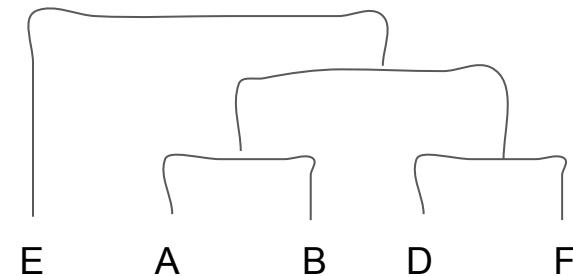
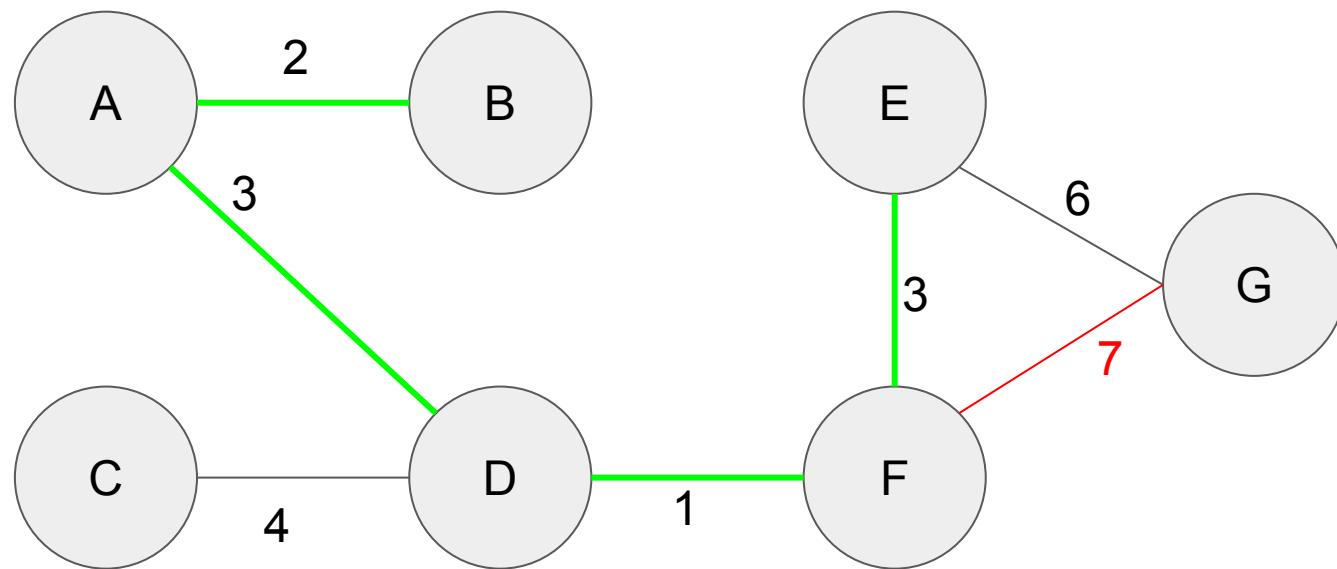
Kruskal's algorithm - weight 3



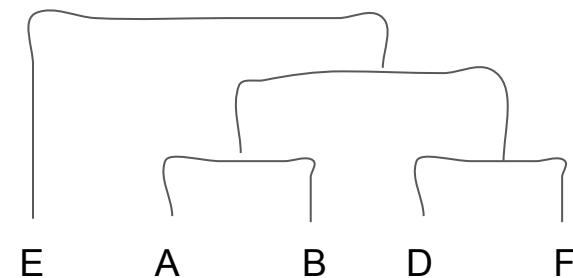
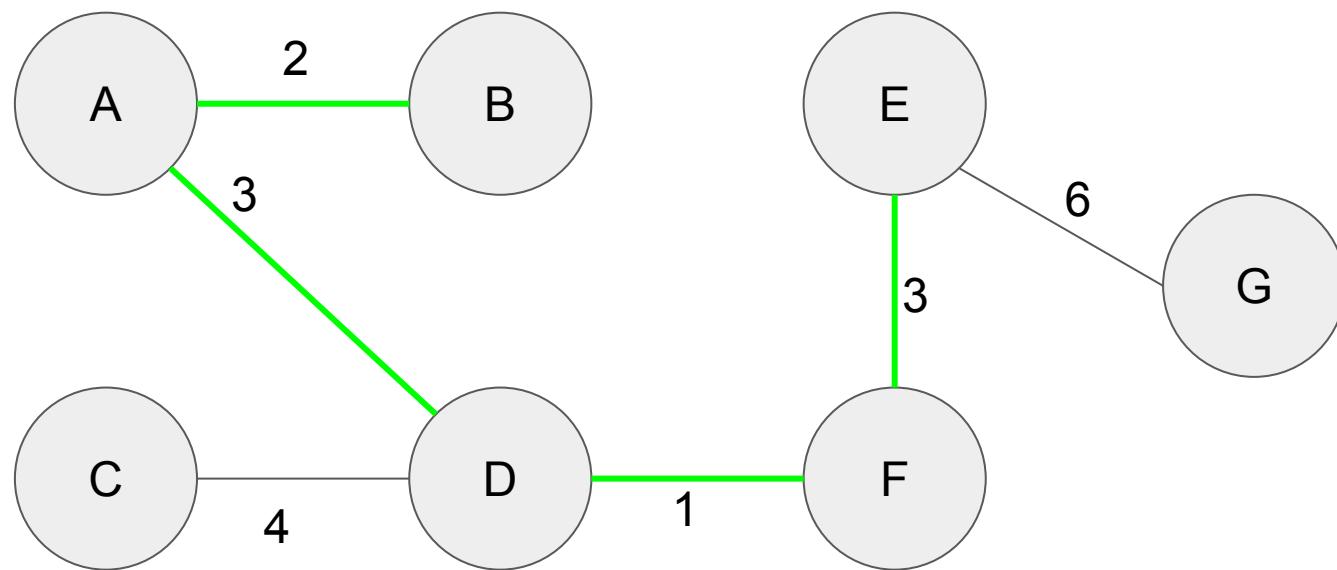
Kruskal's algorithm - weight 3



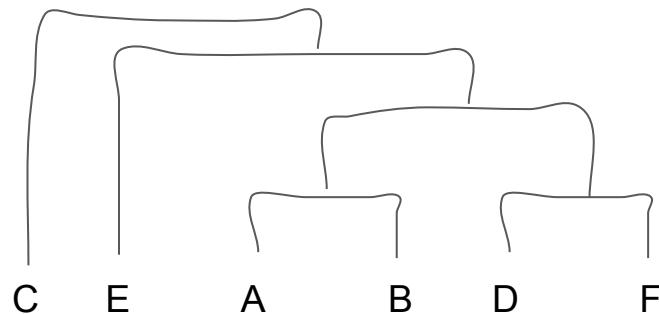
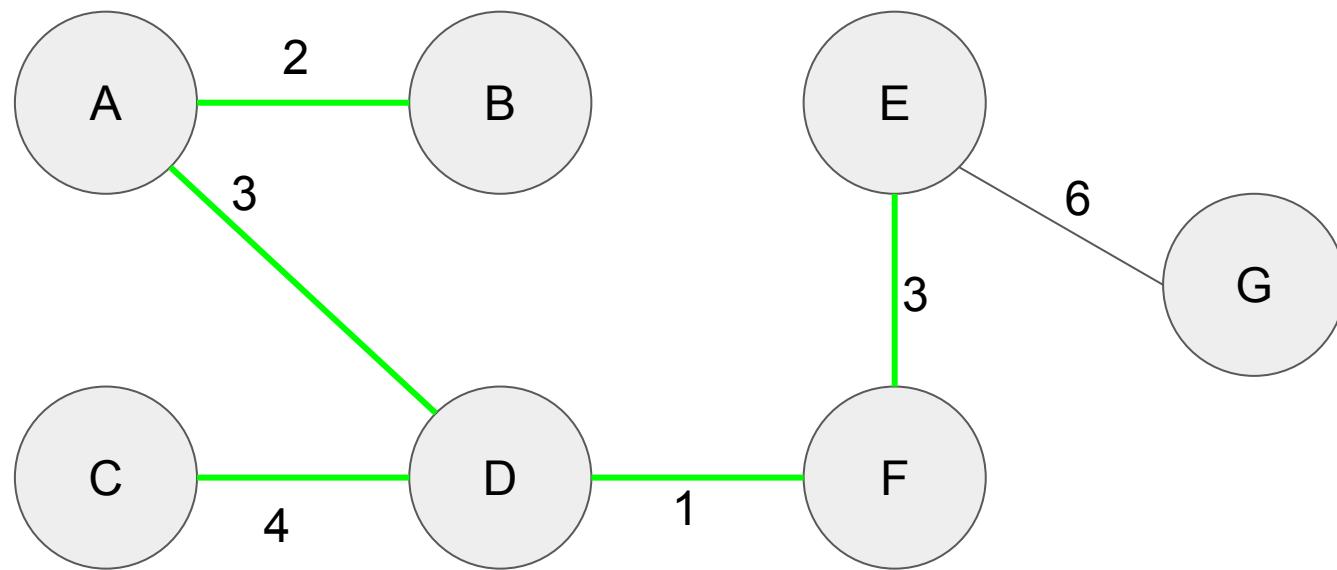
Kruskal's algorithm - weight 3



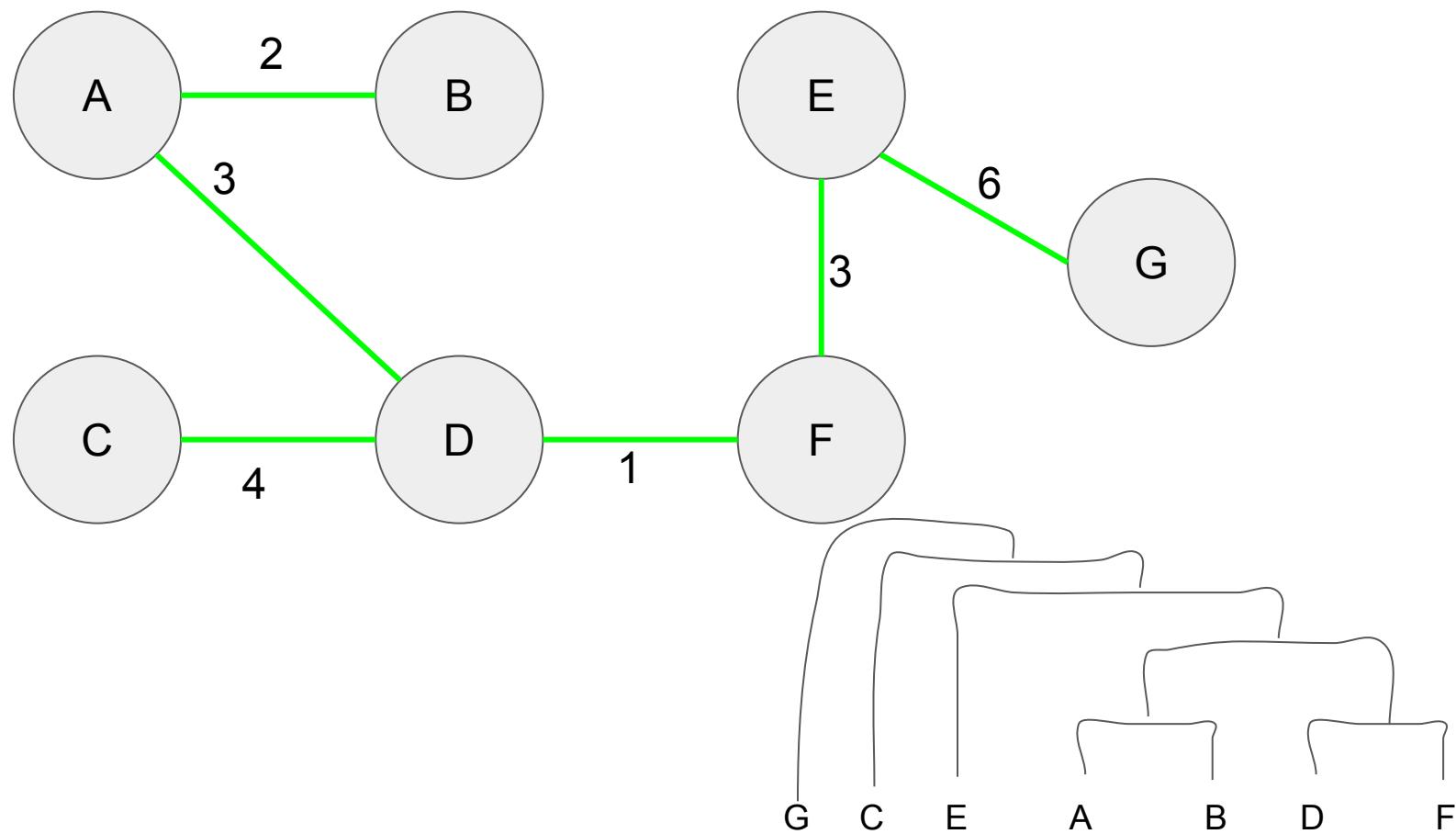
Kruskal's algorithm - weight 3



Kruskal's algorithm - weight 4



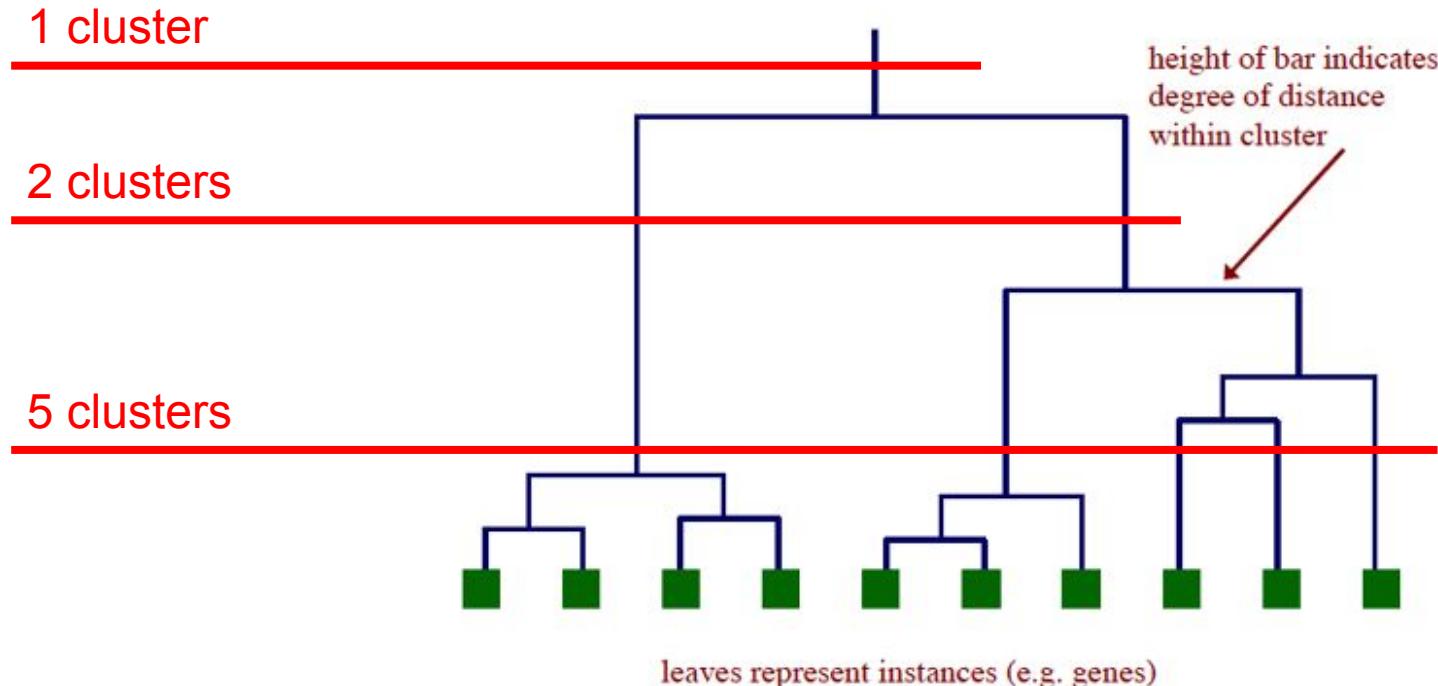
Kruskal's algorithm - weight 6



Dendrogram

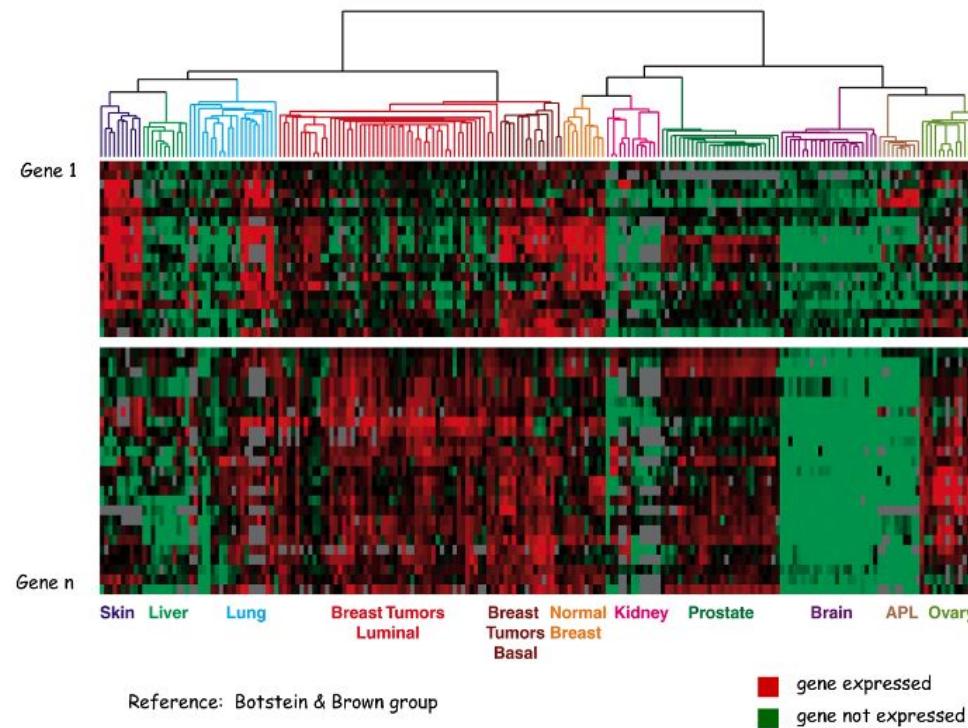
Dendrogram. Scientific visualization of hypothetical sequence of evolutionary events.

- Leaves = genes.
- Internal nodes = hypothetical ancestors.



Applications in data analysis

Tumors in similar tissues cluster together.



Applications: taxonomy



Morphological features

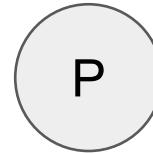
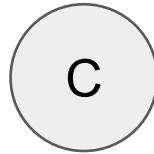
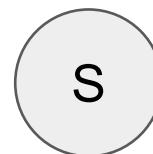
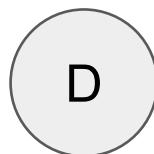
Dolphin (D): aquatic, eats fish

Seal (S): semi-aquatic, eats fish

Cow (C): terrestrial, eats grass

Panda (P): terrestrial, eats grass

Morphological MST:



Applications: taxonomy

Morphological features

Dolphin (D): aquatic, eats fish

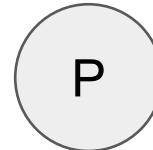
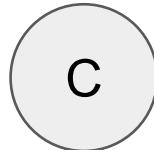
Seal (S): semi-aquatic, eats fish

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Panda (P): terrestrial, eats grass



Morphological MST:



Applications: taxonomy

Morphological features

Dolphin (D): aquatic, eats fish

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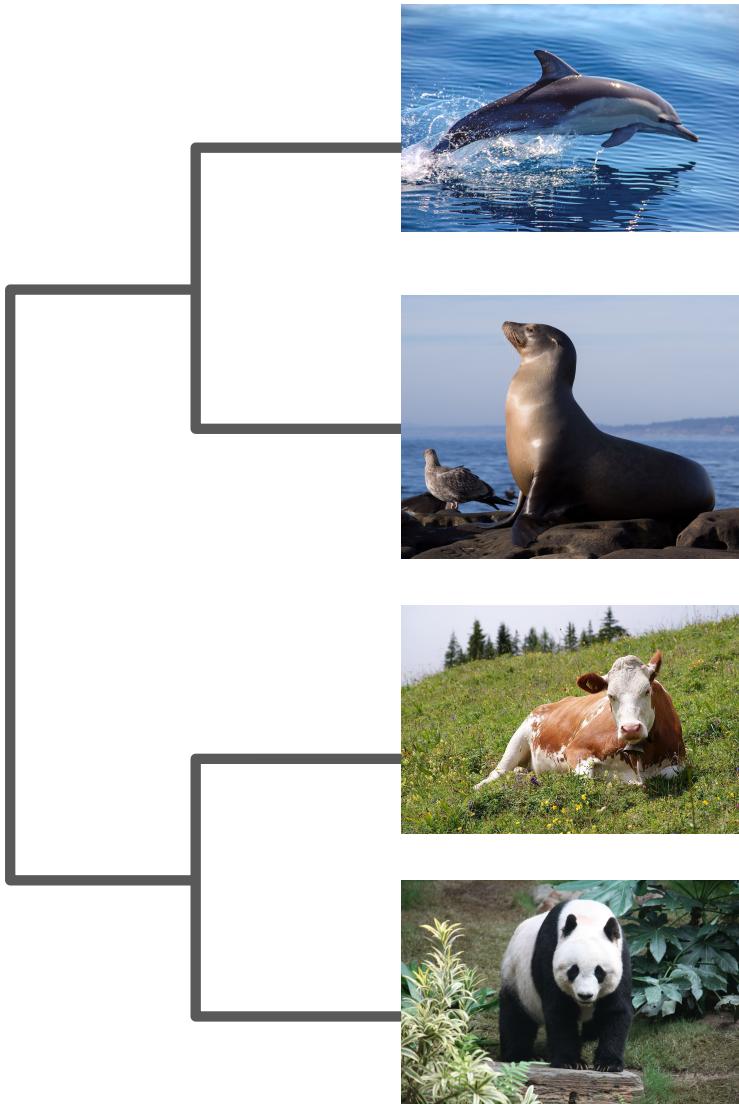
Panda (P): terrestrial, eats grass



Morphological MST:



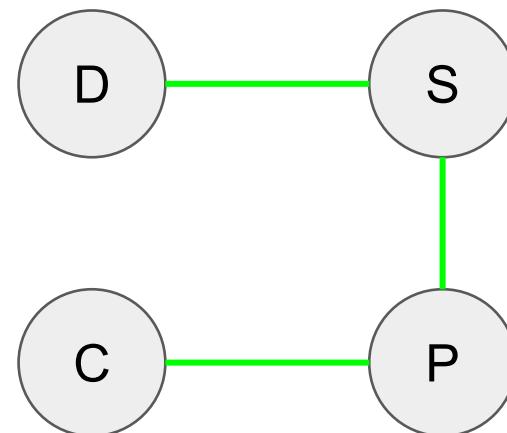
Applications: taxonomy



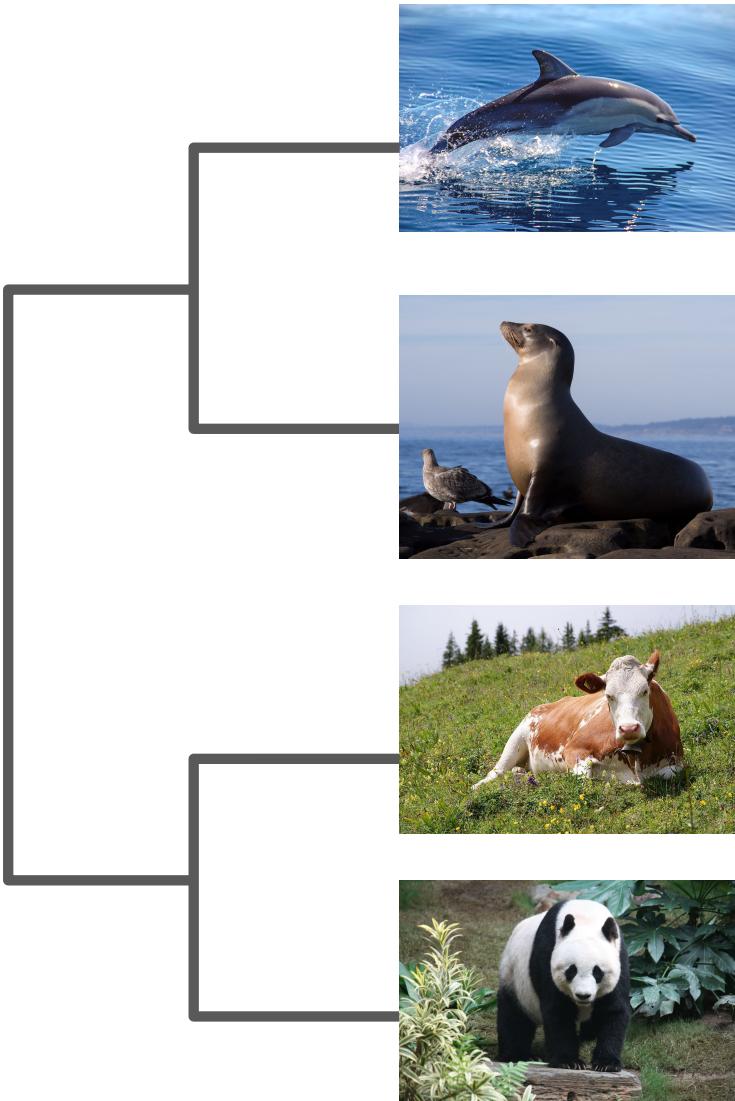
Morphological features

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Morphological MST:



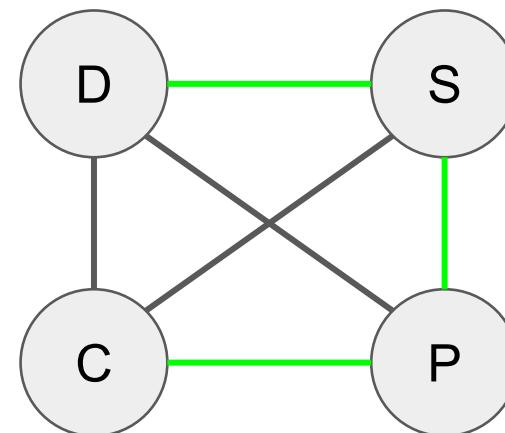
Applications: taxonomy



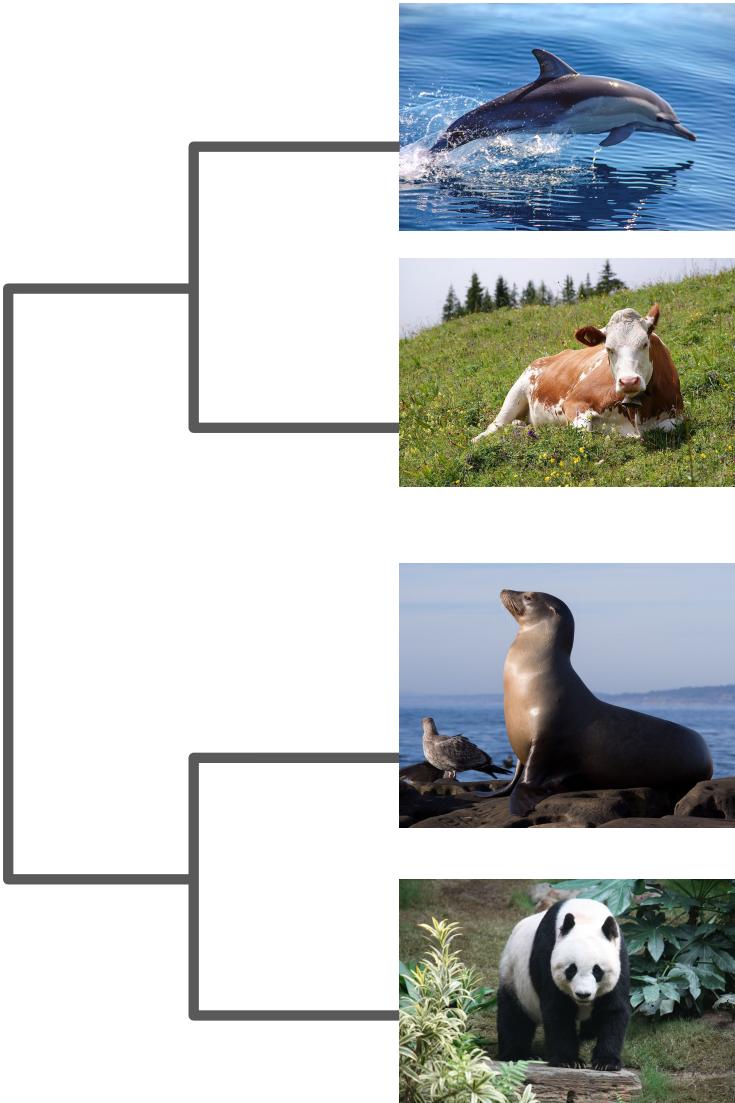
Morphological features

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Morphological MST:



Applications: taxonomy



Morphological features

Dolphin (D): aquatic, eats fish

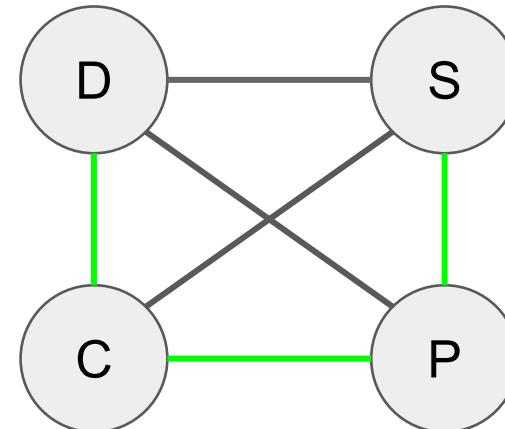
Seal (S): semi-aquatic, eats fish

Cow (C): terrestrial, eats grass

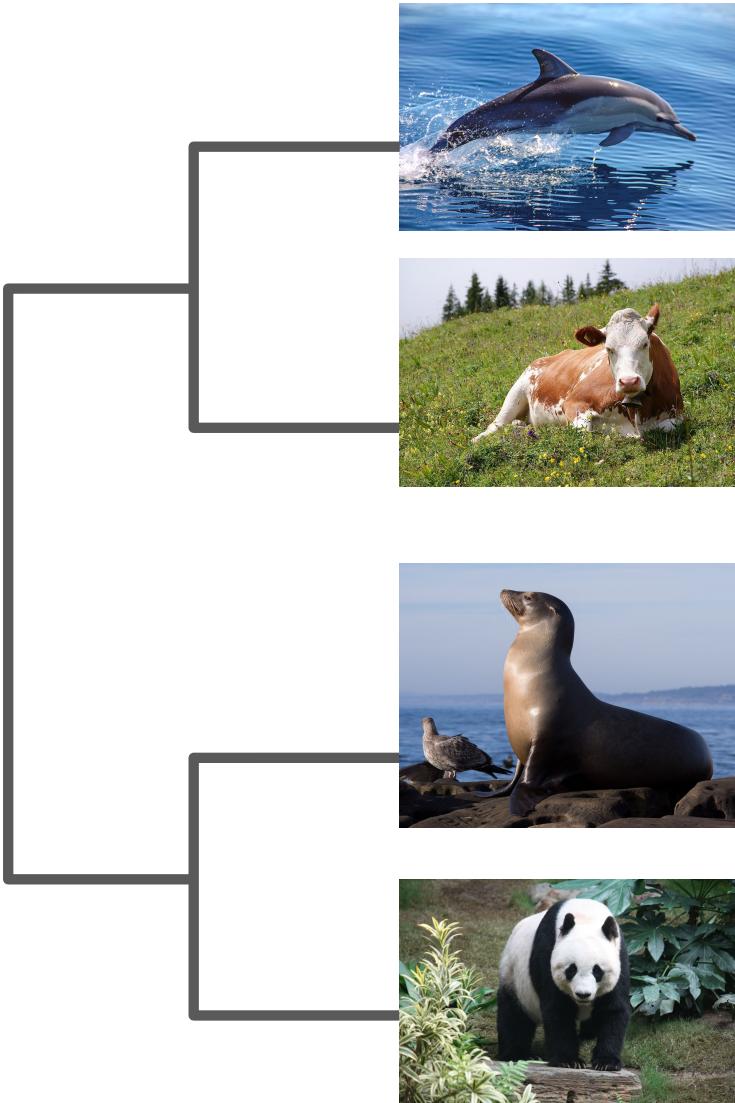
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Genetic MST

(the actual evolutionary path):

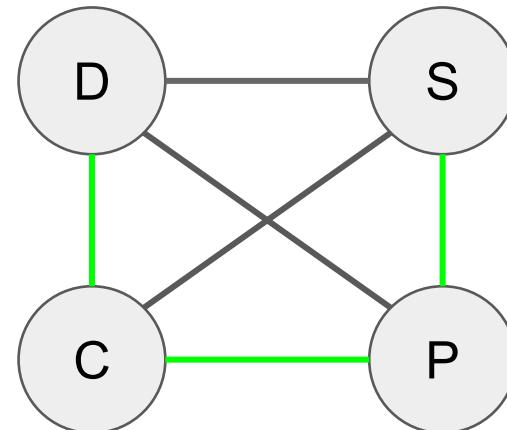


Applications: taxonomy



Computing genetic distances
will be discussed in the
dynamic programming section

Genetic MST
(the actual evolutionary path):



Huffman coding

Huffman encoding

An encoding scheme used in, e.g., MP3 encoding

Data: a string S of symbols over an alphabet Γ

Goal: find a binary encoding e of Γ resulting in minimum encoded length of S

Denote the encoded string by S_e

$a \quad 0110001$

Example: ASCII encoding $b \quad 01100010$

$\vdots \quad \vdots$

Different encodings

Consider $\Gamma = \{a, b, c\}$

Stats on S : a appears 45 times, b 16 times, and c twice

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- Fixed-length encoding

$$a \rightarrow 00$$

$$e_1 : b \rightarrow 01 \quad |S_{e_1}| = 45 \times 2 + 16 \times 2 + 2 \times 2 = 126$$

$$c \rightarrow 10$$

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- Variable-length encoding

$$a \rightarrow 0$$

$$e_2 : b \rightarrow 10 \quad |S_{e_2}| = 45 \times 1 + 16 \times 2 + 2 \times 2 = 81$$

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$$c \rightarrow 11$$

$$a \rightarrow 0$$

- Be careful! $e_2 : b \rightarrow 1$ Decoding will lead to ambiguity

$$c \rightarrow 01$$

Prefix-free encoding

$$a \rightarrow 0$$

Consider the bad encoding e_2 : $b \rightarrow 1$ How to decode 010110?

$$c \rightarrow 01$$

ababba?, ccba?, abcba?, or ... ?

Prefix-free encoding

$$a \rightarrow 0$$

Consider the bad encoding e_2 : $b \rightarrow 1$ How to decode 010110?

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ababba?, ccba?, abcba?, or ... ?

To avoid ambiguity, we need the encoding to be **prefix-free**

Definition

An encoding is **prefix-free** if no codeword is a prefix of any other codewords

Tree representation of a prefix-free encoding

Definition

A **full binary tree** is a binary tree where each node is either a leaf or it has two children

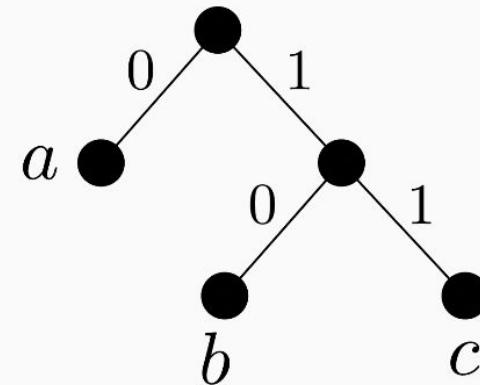
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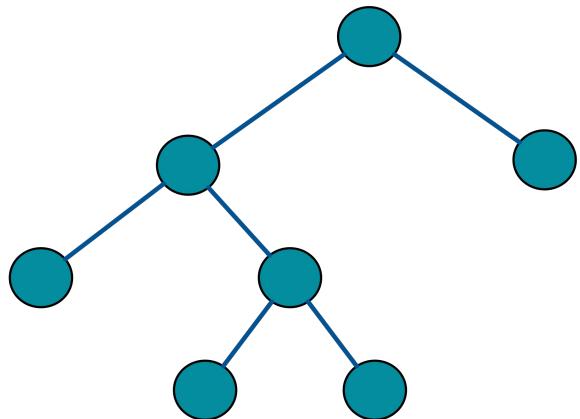
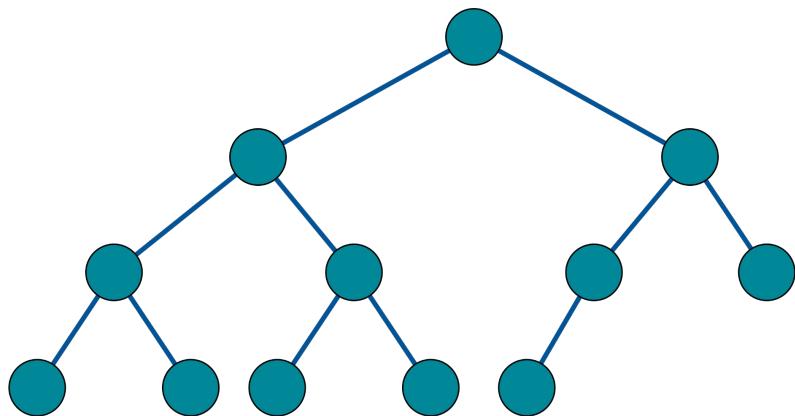
A **full binary tree** is a binary tree where each node is either a leaf or it has two children

We use a full binary tree to represent a prefix-free encoding

- leaves are corresponding to symbols in Γ
- label edge to the left child with 0
- label edge to the right child with 1



Is it a full binary tree?



Tree representation of a prefix-free encoding

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We use a full binary tree to represent a prefix-free encoding

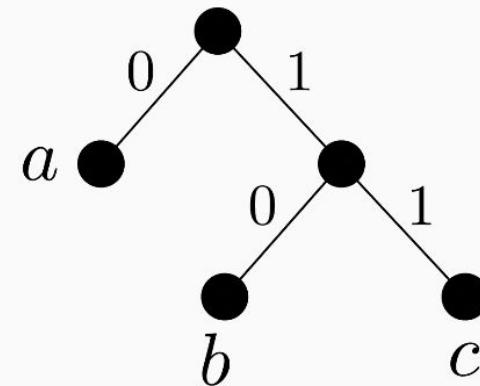
- leaves are corresponding to symbols in Γ
- label edge to the left child with 0
- label edge to the right child with 1

To obtain the encoding, read edge labels from root to a symbol

$a \rightarrow 0, b \rightarrow 10, c \rightarrow 11,$

Depth of a leaf \equiv length of its codeword

It guarantees to be prefix-free



Tree representation of a prefix-free encoding

Let e be an encoding represented by a tree

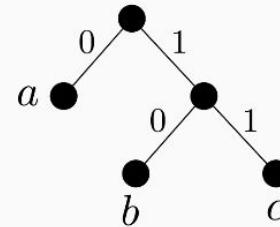
For string S , let f_v be the symbol count in S for each $v \in \Gamma$

Tree representation of a prefix-free encoding

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For string S , let f_v be the symbol count in S for each $v \in \Gamma$

$$|S_e| = \sum_{v \in \Gamma} f_v \cdot \text{depth}(v)$$



$$|S_e| = 45 \times 1 + 16 \times 2 + 2 \times 2 = 81$$

Consider $\Gamma = \{a, b, c\}$

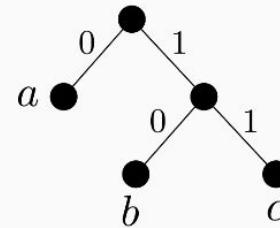
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A useful re-write: label internal nodes with counts of descendants

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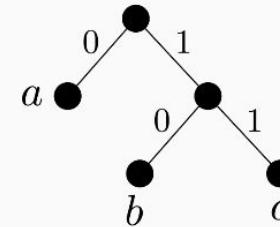
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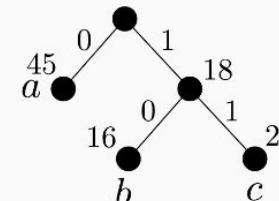
$$|S_e| = 45 \times 1 + 16 \times 2 + 2 \times 2 = 81$$

A useful re-write: label internal nodes with counts of descendants

For all non-root node v , define

$\text{cost}(v) := \text{sum of leaf node counts descending from } v$

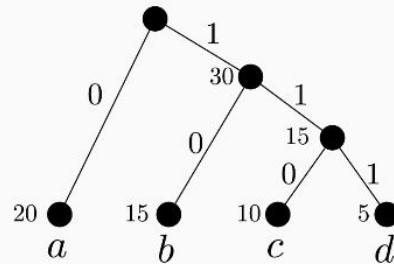
$$|S_e| = \sum_{v \in T - \{\text{root}\}} \text{cost}(v)$$



$$|S_e| = 45 + 16 + 2 + 18 = 81$$

Constructing the prefix-free encoding tree: examples

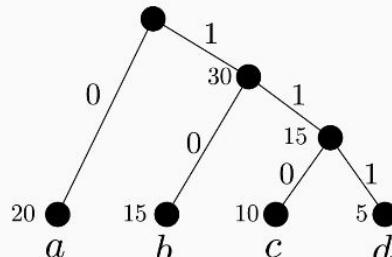
- $a : 20, b : 15, c : 10, d : 5$



$a \rightarrow 0$
 $b \rightarrow 10$
 $c \rightarrow 110$
 $d \rightarrow 111$

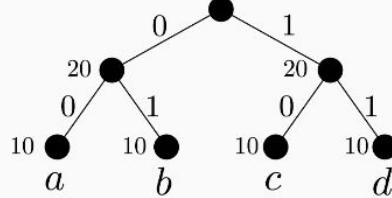
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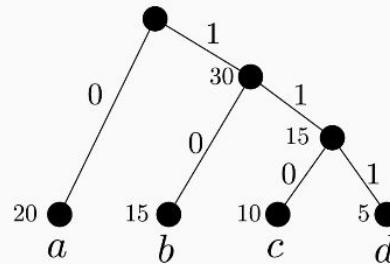
- $a : 10, b : 10, c : 10, d : 10$



$a \rightarrow 00$
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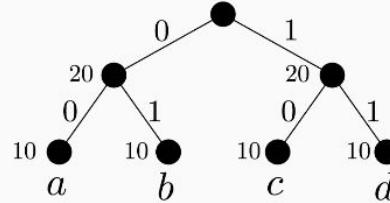
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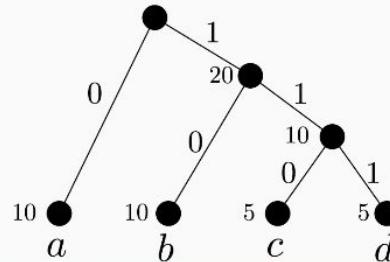
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- $a : 10, b : 10, c : 5, d : 5$



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What are the total costs of these encodings?