5. Arbres phylogénétiques

- L'arbre des espèces
- L'arbre, objet abstrait
- Remplir un tableau de distances
- L'algorithme UPGMA
- Quand les différences sont trompeuses
- La diversité des algorithmes informatiques
- Les applications en microbiologie

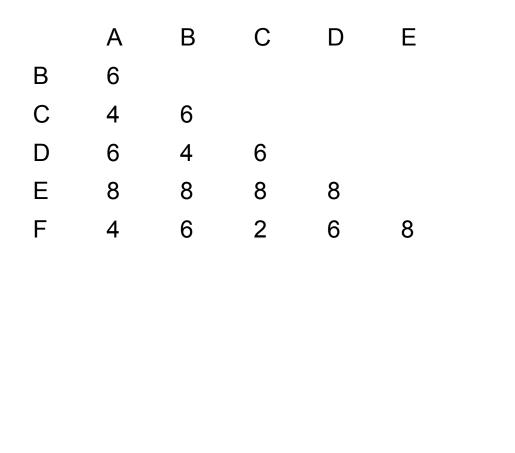


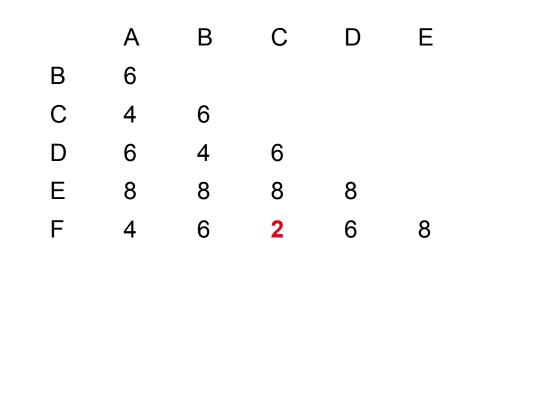
UPGMA

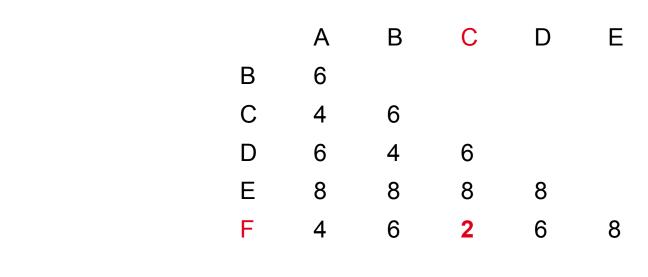
Unweighted Pair Group Method with Arithmetic Mean

Un nom plutôt compliqué pour une méthode plutôt simple

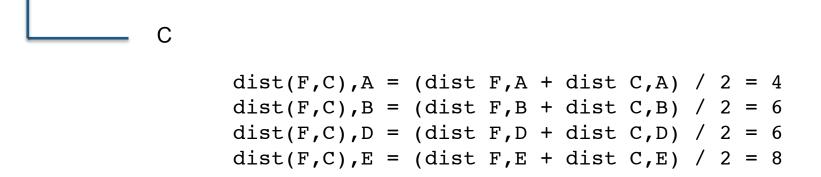
Le point de départ, un tableau de distances











Ε

8 8 8 8



Ε

8 8 8 8

















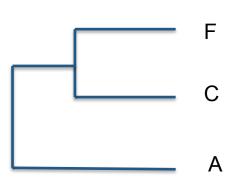


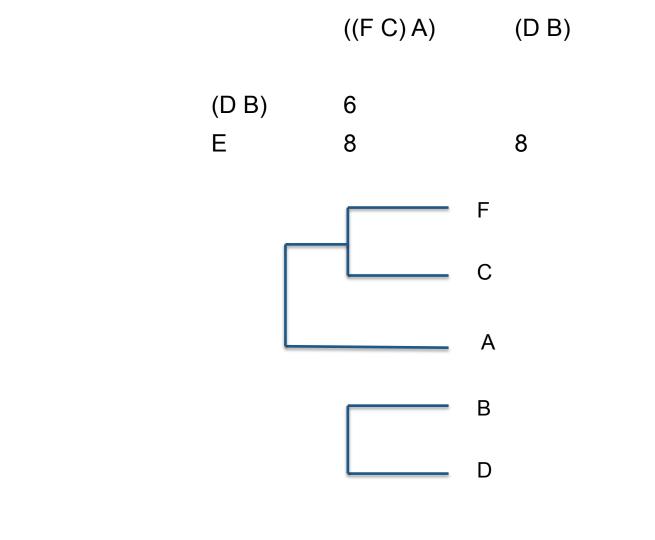
D 6 6 4

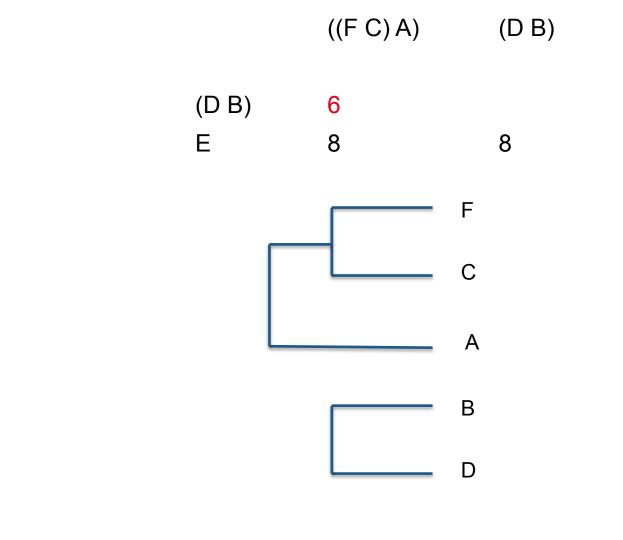
E 8 8 8 8

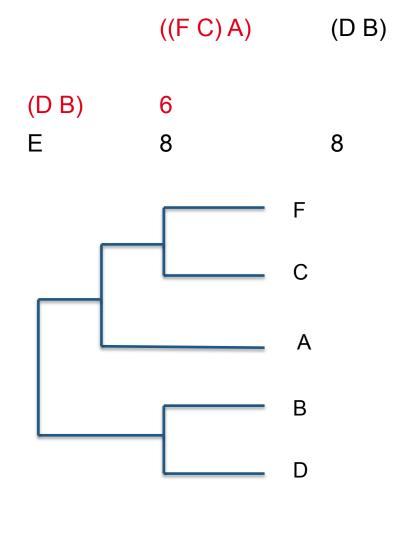


E 8 8 8 8

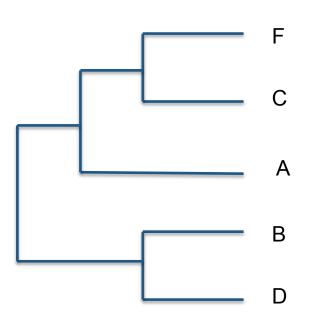








```
dist (((F C) A)(D B)) E
= (dist ((F C) A) E + dist (D B) E) / 2
= 8
```



((F C) A) (D B)

