

Phylogenetic Networks

part of “Graphen und Netzwerke in der Biologie”

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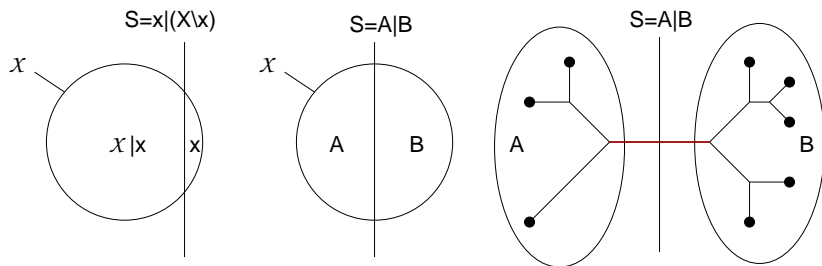
Computational EvoDevo
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- are unrooted phylogenetic networks
- contain at least as much information as a phylogenetic tree
- usually contain more information than a tree
- visualize information that is conflicting in tree representation
- monophyletic groups of sequences are more easily identified

What is a split?

A split $S = A|B$ is a bipartition of the sequences X into two non-empty, mutually exclusive subsets A and B . A and B are called *split parts*.

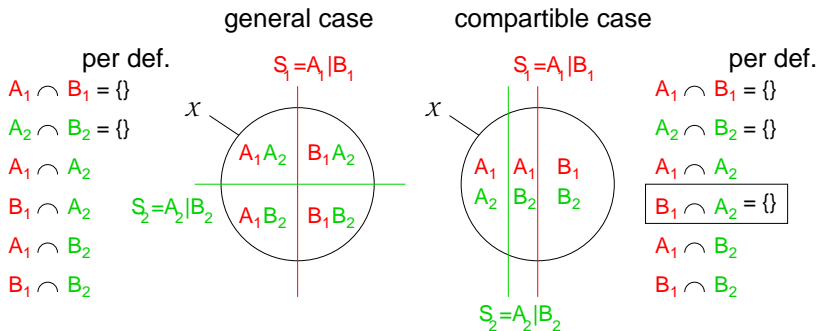


A *trivial split* $S = x|(X \setminus x)$ separates one sequence x from the other sequences $X \setminus x$.

Relation Between Splits and Trees

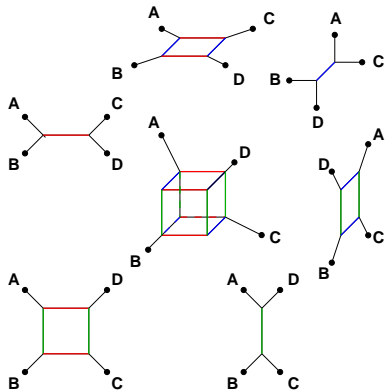
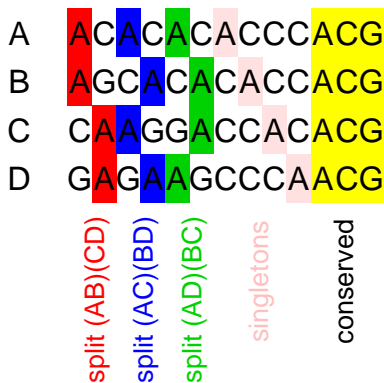
Given a set of splits S on X , can it be represented by a tree?

Compatibility: two splits $S_1 = A_1|B_1$ and $S_2 = A_2|B_2$ are *compatible* if one of the four possible intersections of the split parts is empty (or splits don't "cross" each other).



A set of splits is compatible if all pairs of splits are compatible.
A compatible set of splits can be represented as a tree.

The Information to be represented

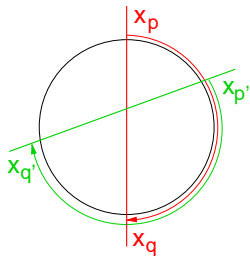


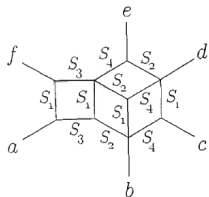
Circular splits

A set of splits S on X is called circular, if there exists a linear ordering (x_1, x_2, \dots, x_n) of the elements of X such that each split S_i has the form

$$S_i = \frac{x_p, x_{p+1}, \dots, x_q}{X - x_p, x_{p+1}, \dots, x_q} \quad (1)$$

for $1 < p \leq q \leq n$.

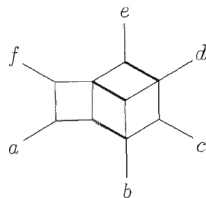




(a) Split network N

$$\begin{aligned}
 S_1 &= \{a, b, c\} \parallel \{d, e, f\} \\
 S_2 &= \{a, e, f\} \parallel \{b, c, d\} \\
 S_3 &= \{a, f\} \parallel \{b, c, d, e\} \\
 S_4 &= \{a, b, f\} \parallel \{c, d, e\}
 \end{aligned}$$

(b) Non-trivial splits



(c) Split $S_2 = \frac{\{a, e, f\}}{\{b, c, d\}}$

Figure 5.6 (a) A split network N representing all trivial splits on $\mathcal{X} = \{a, \dots, f\}$ and the four non-trivial splits listed in (b). All the edges representing a particular non-trivial split are labeled by that split. However, the labeling of edges by splits is usually omitted, as shown in (c). Edges representing the same split, such as the three edges shown in bold lines representing S_2 , are drawn parallel and with the same length.

Is the splits system a circular splits system? Why?

Circular Splits Systems Can be Represented by a Planar Network

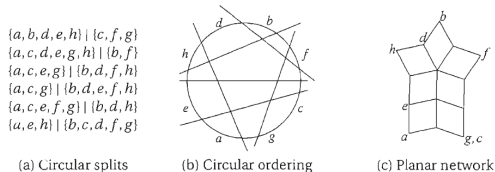


Figure 5.9 (a) A set of six circular splits S on $\mathcal{X} = \{a, b, \dots, h\}$. (b) An arrangement of the taxa around a circle such that every split $S = A | B \in S$ can be realized by a straight line through the circle that separates the two split parts A and B . A circular ordering is given by (a, g, c, f, b, d, h, e) . (c) An outer-labeled planar split network representing S .

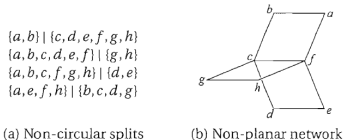


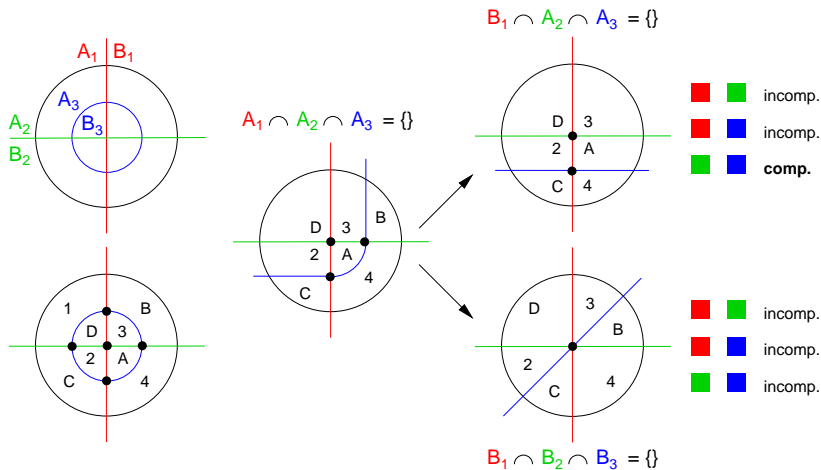
Figure 5.10 (a) A set of four non-circular splits S on $\mathcal{X} = \{a, b, \dots, h\}$. (b) A non-planar split network representing S .

Three distinct splits are weakly compatible if

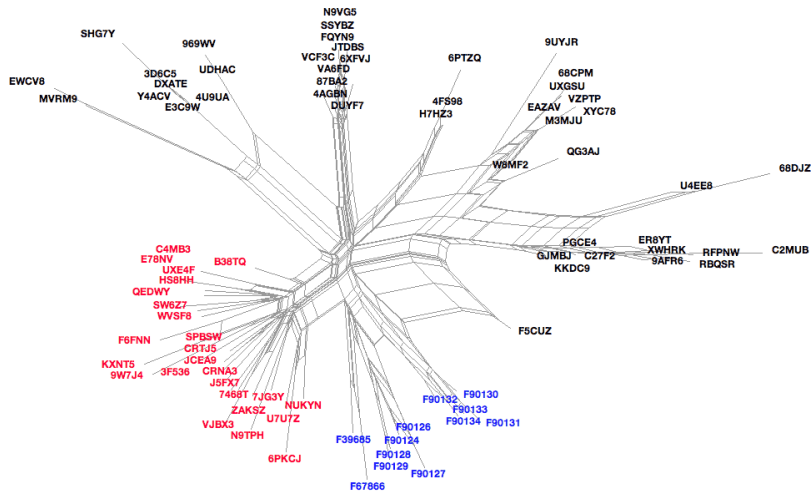
- at least one of the following intersections is empty:
 - $A_1 \cap A_2 \cap A_3$ (1)
 - $A_1 \cap B_2 \cap B_3$ (2)
 - $B_1 \cap A_2 \cap B_3$ (3)
 - $B_1 \cap B_2 \cap A_3$ (4)
- and symmetrically, at least one of the following intersections is empty:
 - $B_1 \cap B_2 \cap B_3$ (A)
 - $B_1 \cap A_2 \cap A_3$ (B)
 - $A_1 \cap B_2 \cap A_3$ (C)
 - $A_1 \cap A_2 \cap B_3$ (D)

A set of splits is weakly compatible if all triples are weakly compatible.

Weak compatibility



A Phylogenetic Network



Are the two group of sequences marked in red and blue supported by a split?

Phylogenetic Networks, Concepts, Algorithms and Applications
by Daniel H Huson, Regula Rupp and Celine Scornavacca.
Cambridge University Press, 2010.

<http://www.cambridge.org/9780521755962>

Software **SplitsTree** <http://www.splitstree.org/>