

SplitsTree

SplitsTree is a program for inferring [phylogenetic trees](#) and [networks](#). It handles various sources for phylogenetic information such as sequence alignments, distance matrices, and [tree sets](#). SplitsTree offers various methods for building phylogenetic trees such as split decomposition, [NeighborNet](#), consensus bootstrap trees.

References

- <http://www.splitstree.org/> Accessed 28 October 2015.
- Huson, Daniel H., and David Bryant. 2006. "Application of Phylogenetic Networks in Evolutionary Studies." *Molecular Biology and Evolution* 23 (2): 254–267.

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