



● MARCOS MÉNDEZ

### ABSTRACT

*Tree-thinking is a required skill for evolutionary literacy, but misconceptions are frequent in interpreting phylogenetic trees. Here, an easy, cheap way of building phylogenetic trees with drinking straws is suggested. It can be used to illustrate how to build phylogenetic trees from root to tips, to rotate nodes and to collapse branches.*

**Key Words:** *evolution; phylogenetic trees; misconceptions; hands-on learning.*

Tree-thinking, the ability to conceptualize evolution in terms of phylogenetic trees (Baum & Offner, 2008), is a skill needed to increase evolutionary literacy. However, misconceptions about how to read and interpret phylogenetic trees are common (Meir et al., 2007; Gregory, 2008; Staton, 2015; Kummer et al., 2016). These misconceptions are usually addressed by means of activities that involve building phylogenetic trees with different kinds of data (Campo et al., 2009; Rau, 2012; David, 2018; Punyasettiro & Yasri, 2021), and plotting phylogenetic trees by means of drawings (Bilardello & Valdés, 1998; Kozłowski, 2010; Dees & Momsen, 2016) or computers (Perry et al., 2008; Schneider et al., 2012; Zhang, 2012; Duffus, 2019). Here, a way of building phylogenetic trees by hand is suggested, using actual physical branches, a relatively little explored approach (Halverson, 2010). Physical phylogenetic trees facilitate the acquisition of topological skills—node rotation, branch regrafting—key to tree-thinking, that can be challenging when performed mentally or using drawings.

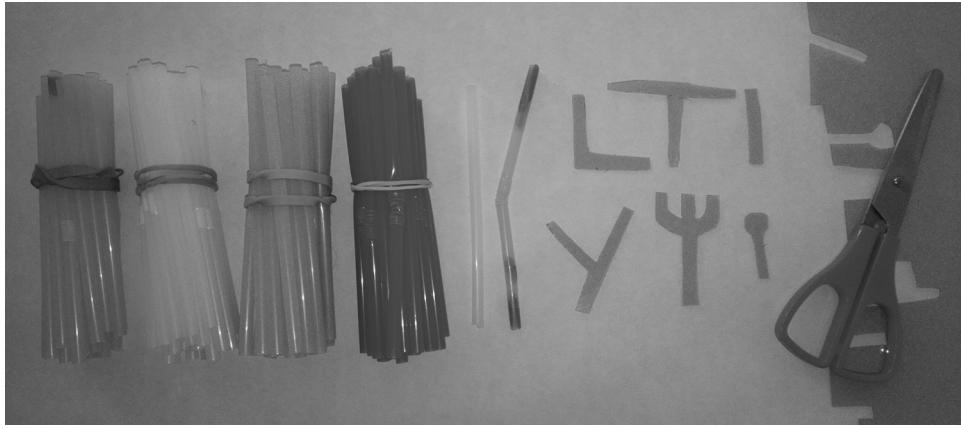
Physical phylogenetic trees can be built in an easy, cheap way using drinking straws. All that is needed are straws of different colors, plasticine, and a folder made of hard plastic (Figure 1). Either plastic or cardboard straws can be used, but the latter are more environmentally friendly. Tree branches are made with straws trimmed to a length of either 5 or 10 cm (Figure 1).

Plasticine is plugged at each end of the straws (Figure 1) to help with fitting the connectors. Several kinds of connectors can be cut from the plastic folder, using scissors or a cutter, with a width that fits the diameter of the straws (Figure 1 and Figure S1 in Supplementary Material provided with the online version of this article). Rectangular trees are assembled by joining the straws using T or L connectors (Figure 2). I connectors allow students to expand the length of a branch (Figure 2). This basic design can be modified at will. Possibilities are almost endless; only a few are listed here. First, pieces of paper with drawings of different organisms (e.g., caminalcules: Sokal, 1983; Gendron, 2000) can be glued at the tips of the tree using “lollypop” connectors (Figure 1). Second, straws of different colors can be used to flag different clades in the tree. Third, straws can be combined with pipe cleaners (Halverson, 2010) or coated with sandpaper of different grit, such that visually impaired students can touch the trees and figure out not only the shape but also the different clades that branch along the tree.

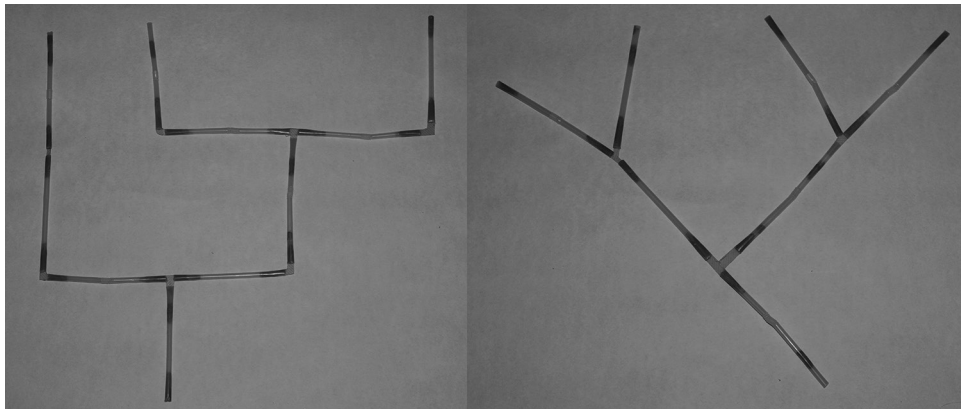
Building physical phylogenetic trees can be included in the teaching of tree-thinking in at least five ways. First, these trees can be easily assembled by the teacher or the students during lectures to illustrate different topologies of trees, such as symmetric and asymmetric branching. Second, building physical phylogenetic trees can be easily included in hands-on activities designed to allow students getting familiar with establishing phylogenetic relationships between organisms, such as caminalcules (Gendron, 2000). These hands-on activities can reinforce tree-thinking and tree-building skills (Schramm et al., 2019).

Third, phylogenetic trees built with drinking straws can help students in dispelling several usual misconceptions (Gregory, 2008) (Figure 3 and Further Misconceptions in Supplementary Material provided with the online version of this article). For example, several misconceptions about phylogenetic trees identified by

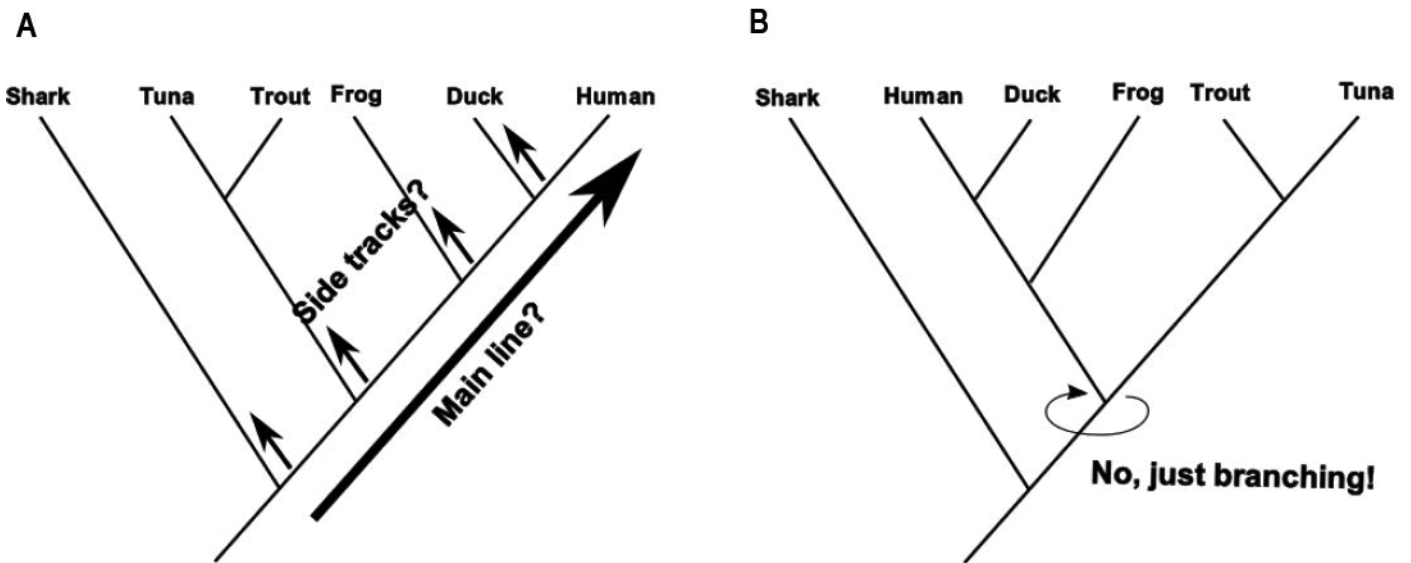
*Physical phylogenetic trees facilitate the acquisition of topological skills—node rotation, branch regrafting—key to tree-thinking, that can be challenging when performed mentally or using drawings.*



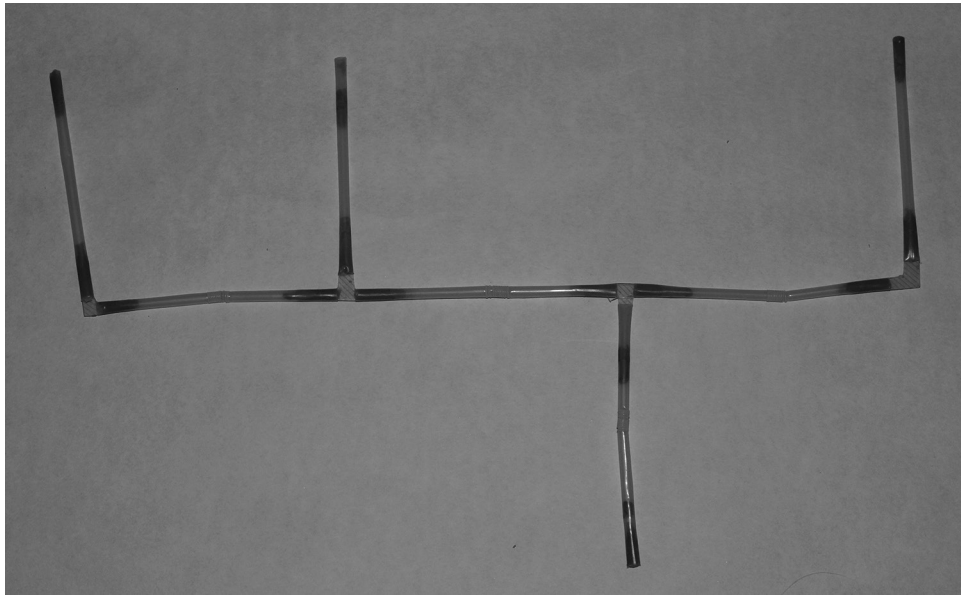
**Figure 1.** Materials needed to build phylogenetic trees with drinking straws. From left to right, drinking straws of different colors, trimmed to a length of 10 cm, straws with plugs of plasticine at the ends, one of them with a fold, and L, T, I, Y, psi, and lollipop connectors. Connectors are cut with scissors from a folder of hard plastic, visible at the right of the figure.



**Figure 2.** Left, a rectangular tree with three branches joined with T and L connectors; the leftmost branch is extended using an I connector. Right, a tree in diagonal version, using Y connectors.



**Figure 3.** A: a tree showing the apparent main line of evolution toward a target, humans, while the rest of the tree is interpreted as side tracks away from that target. B: a simple rotation shows that this main line is only a product of the arrangement of the branches; actually, the tree is showing branching only from a common ancestor and no evolution toward a target.



**Figure 4.** A tree with a polytomy, created with an inverted T connector.

Gregory (2008) can be dispelled by rotating branches around a node. Here, the misconception “main line and side tracks” is illustrated, as an on-class exercise. In asymmetrically branching trees, students tend to see one “main track,” showing the “progress toward a target,” while the rest of the branches are side tracks (Figure 3A). Rotation of nodes in a physical tree built with drinking straws graphically makes the point that such main track is only apparent, not real. The exercise has four steps. In step 1, show your students the phylogenetic tree on Figure 3A. In step 2, provide material to build trees with drinking straws and ask the students to build the tree, in which the misconception is apparent. In step 3, ask your students to rotate nodes of the tree so that they turn it into the tree in which the apparent track is not present. A potential solution is given in Figure 3B. In step 4, discuss how the apparent pattern in the left tree disappears without actually changing the tree, in other words, the phylogenetic relationships between species in the tree. This in-class exercise can be easily expanded to solve other misconceptions included in Gregory (2008) such as “reading across the tips” and “sibling vs. ancestor” (Further Misconceptions in Supplementary Material provided with the online version of this article).

Fourth, building physical trees helps realizing that trees are read from bottom to top, while drawing trees on a paper does not convey as easily the way in which a tree adds branches. Fifth, these trees support teaching how differences in topology involve branch regrafting, as well as how to collapse branches in order to create consensus trees with polytomies (Regrafting Branches in Supplementary Material provided with the online version of this article), both of which are more difficult with pipe cleaners. Trees with polytomies can be built simply using inverted T connectors (Figure 4), or by means of  $\Psi$  (psi) connectors (Figure 1). These suggestions can be easily developed by readers to fit their needs when teaching tree-thinking at primary, secondary, or college levels.

## References

Baum, D. A. & Offner, S. (2008). Phylogenies & tree-thinking. *American Biology Teacher*, 70, 222–229.

- Bilardello, N. & Valdés, L. (1998). Constructing phylogenies. *American Biology Teacher*, 60, 369–373.
- Campo, D., Alvarado, A., Machado-Schiaffino, G., Naji, L., Pelaez, R., Quiros, F., Rodríguez, O., Castillo, A.G.F., & Garcia-Vazquez, E. (2009). Inquiry-based learning of molecular phylogenetics II: The phylogeny of Camelidae. *Journal of Biological Education*, 43, 78–80.
- David, A. A. (2018). Using project-based learning to teach phylogenetic reconstruction for advanced undergraduate biology students: molluscan evolution as a case study. *American Biology Teacher*, 80, 278–284.
- Dees, J. & Momsen, J. L. (2016). Student construction of phylogenetic trees in an introductory biology course. *Evolution: Education & Outreach*, 9, ar3.
- Duffus, A. L. J. (2019). An emerging amphibian infection as a model for teaching phylogenetic reconstruction. *American Biology Teacher*, 81, 32–39.
- Gendron, R. P. (2000). The classification & evolution of caminalcules. *American Biology Teacher*, 62, 570–576.
- Gregory, T. R. (2008). Understanding evolutionary trees. *Evolution: Education & Outreach*, 1, 121–137.
- Halverson, K. L. (2010). Using pipe cleaners to bring the tree of life to life. *American Biology Teacher*, 72, 223–224.
- Kozlowski, C. (2010). Bioinformatics with pen and paper: building a phylogenetic tree. *Science in School*, 17, 29–33.
- Kummer, T. A., Whipple, C. J. & Jensen, J. L. (2016). Prevalence and persistence of misconceptions in tree thinking. *Journal of Microbiology & Biology Education*, 17, 389–398.
- Meir, E., Perry, J., Herron, J. C. & Kingsolver, J. (2007). College students’ misconceptions about evolutionary trees. *American Biology Teacher*, 69, 71–76.
- Perry, J., Meir, E., Herron, J. C., Maruca, S. & Stal, D. (2008). Evaluating two approaches to helping college students understand evolutionary trees through diagramming tasks. *CBE-Life Sciences Education*, 7, 193–201.
- Punyasettro, S. & Yasri, P. (2021). A game-based learning activity to promote conceptual understanding of chordates’ phylogeny and self-efficacy to learn evolutionary biology. *European Journal of Educational Research*, 10, 1937–1951.
- Rau, G. (2012). Make your own phylogenetic tree. *The Science Teacher*, 79, 44–49.
- Schneider, B., Strait, M., Muller, L., Efenbein, S., Shaer, O. & Shen, C. (2012). Phylo-Genie: engaging students in collaborative ‘tree-thinking’ through tabletop techniques. *CHI ’12: Proceedings of the SIGCHI Conference on Human Factors in Computing Systems*, 3071–3080.

Schramm, T., Schachtschneider, Y. & Schmiemann, P. (2019). Understanding the tree of life: and overview of tree-reading skill frameworks. *Evolution: Education & Outreach*, 12, ar11.

Sokal, R. R. (1983). A phylogenetic analysis of the caminalcules. I. The data base. *Systematic Zoology*, 32, 159–184.

Staton, J. L. (2015). Understanding phylogenies: Constructing and interpreting phylogenetic trees. *Journal of the South Carolina Academy of Science*, 13, 24–29.

Zhang, X. (2012). Teaching molecular phylogenetics through investigating a real-world phylogenetic problem. *Journal of Biological Education*, 46, 103–109.

MARCOS MÉNDEZ (marcos.mendez@urjc.es) is a full professor at the Area of Biodiversity and Conservation in the Evolutionary Ecology Research Group at the Universidad Rey Juan Carlos in Madrid, Spain.



# THANK YOU to our SUSTAINING MEMBERS!

## PROGRAMMATIC PARTNER

HHMI BioInteractive

## NON-PROFIT PARTNER

National Center for Science Education

## CORPORATE PARTNERS

3D Molecular Designs  
 10K Science  
 Bedford, Freeman & Worth High School Publishers  
 Bio-Rad Laboratories  
 Carolina Biological Supply Company  
 miniPCR

Sustaining Members share NABT's mission to promote biology and life science education. Learn more at [www.NABT.org](http://www.NABT.org).

