

CodeMyPlant (2016-2018)

When high school students join scientists to barcode the flora of Geneva



Candice Yvon¹, Louis Nusbaumer², Sofia Wyler³, Romain Dewaele¹, Bruno Strasser¹, Yamama Naciri²

¹Bioscope - University of Geneva, Switzerland

²Conservatoire et Jardin botaniques de la Ville de Genève & Laboratoire de Systématique végétale et biodiversité – University of Geneva, Switzerland

³SwissBOL – Swiss Barcode of Life, Switzerland



DNA barcoding is a method that uses a short fragment of an organism's DNA - the "barcode" - to identify it as belonging to a particular species. It is dependent upon a DNA reference library, which is a database containing DNA sequences generated from specimens already identified. This molecular technique is especially useful to study samples that are morphologically unrecognizable (seedlings, roots, processed products,...). What is also exciting is that building up the DNA reference library gives us insight into biodiversity at its most subtle level. Comparing millions of DNA sequences substantially enhances our understanding of kinship within and between species, which may eventually help us improve biodiversity management.

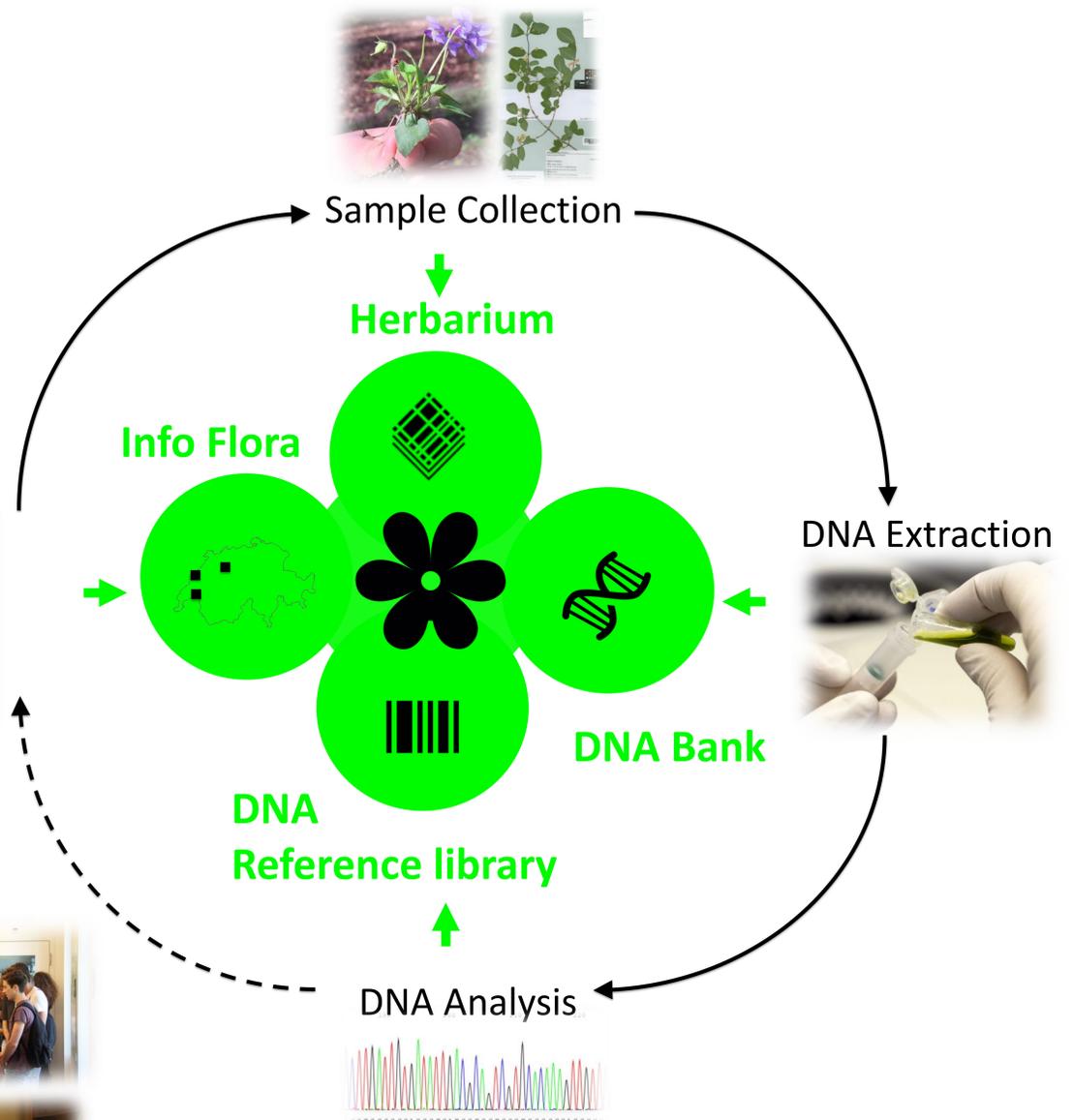
CodeMyPlant project mobilizes high school students and the general public alongside experts in drawing up a genetic inventory of Geneva's flora. The specimens collected are prepared and conserved in the Botanical garden's herbarium, while DNA samples are stored in a DNA bank. Floristic data are transmitted to the National data center *Info Flora* and genetic data are openly shared on the international DNA reference library "BOLD".

2016-2017
 
349 participants **245 DNA barcodes**

Plant Selection & Floristic data



Triggering curiosity and critical thinking among participants is another goal of the project, with a special focus on the concept of species and the Nature of Science. For that purpose, students are involved in several educational activities throughout the year and are asked to complete evaluation forms before and after hand.



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Candice Yvon, candice.yvon@unige.ch

Yamama Naciri, yamama.naciri@ville-ge.ch