

Target enrichment for plant/animal systematics - methodological workshop

9.-17.6.2025

Department of Botany, Charles University, Prague
Roswitha Schmickl, Tomáš Fér, Vojta Zeisek, Soňa Pířová

Monday	Tuesday	Wednesday	Thursday	Friday
9.6.	10.6.	11.6.	12.6.	13.6.
<p><i>Morning</i></p> <p>General introduction</p> <p>Theory - library preparation (Rosi, Tomáš)</p> <p>Theory - target enrichment principle (Rosi) & discussion</p> <p><i>Afternoon</i></p> <p>Lab work demonstration - DNA conc. (Nanodrop, Qubit), Covaris sonication, gel... (Soňa)</p> <p>Independent/group work - preparing presentation of Ufimov et al. (2021) paper – custom vs. universal probes</p>	<p><i>Morning</i></p> <p>Lab work demonstration - library preparation, size selection, gel, barcoding... (Soňa)</p> <p>Paper presentation & discussion (custom vs. universal probes)</p> <p><i>Afternoon</i></p> <p>Lab work demonstration (contin.)</p> <p style="text-align: center;">&</p> <p>Independent work on paper of choice about target enrichment in plant/animal systematics</p>	<p><i>Morning</i></p> <p>Theory - approaches for data analysis (Rosi)</p> <p>Theory - target enrichment data structure (Vojta)</p> <p>Computer work - data cleaning, gene alignments with HybPiper (Vojta)</p> <p><i>Afternoon</i></p> <p>Theory - gene trees vs species trees (Tomáš)</p>	<p><i>Morning</i></p> <p>Theory - gene trees vs species trees (continue) (Tomáš)</p> <p><i>Afternoon</i></p> <p>Computer work - gene tree, species tree building (Vojta)</p>	<p><i>Morning</i></p> <p>Computer work - HybPhyloMaker - initial steps (cleaning, mapping, alignment, filtering) (Tomáš)</p> <p><i>Afternoon</i></p> <p>Computer work - HybPhyloMaker - species tree methods, discordance, networks (Tomáš)</p> <p>Theory&discussion - discordance, networks, hybridization (Rosi, Tomáš)</p>

Monday	Tuesday
16.6.	17.6.
<p><i>Morning</i></p> <p>Student presentations - papers of their choice (5mins + 10mins discussion)</p> <p><i>Afternoon</i></p> <p>Group work/discussion - reading Joyce et al. (2025), discussion of tools&approaches</p>	<p><i>Morning</i></p> <p>Plastome 'assembly' - HybPhyloMaker, FastPlast (Tomáš)</p> <p><i>Afternoon</i></p> <p>Wrap-up, varia (Rosi, Tomáš, Vojta)</p> <p>Hands-on session with own data etc. (Rosi, Tomáš, Vojta)</p>