

**Longread-only approach to the organellar genome
assembly of a rare endemic non-model species
Crepis callicephalo Juz. (Asteraceae)**

Emirsaliiev A. ^{1,2*}, Tsyupka V. ^{1,2}, Grebennikova O. ^{1,2}, Bulavin I. ^{1,2}, Krivenko O. ^{1,2}, Ivanova N. ², Nikiforov A. ², Mitrofanova I. ^{1,2,3}

¹ Kurchatov Genomic Centre – NBG-NSC, Russian Federation, Yalta

² FSFIS "Nikita Botanical Gardens – National Scientific Center of RAS", Russian Federation, Yalta

³ FSBIS Main Botanical Gardens named after N.V. Tsitsin, Russian Federation, Moscow

Crepis callicephalala Juz.

C. callicephalala Juz. is a rare endemic species of Crimean Flora.

- perspective source of biologically active compounds
- wild relative of such important cultivated plants as *Lactuca* and *Cichorium*
- notable niche specialisation
- the genus has a great success story as an object in cytoembriology, but still is unclear in modern omics approaches
- endangered conservation status



Plastid genome assembly tools used

de novo tools:

- Canu, flye, pomoxis (*de novo* mode) and NECAT

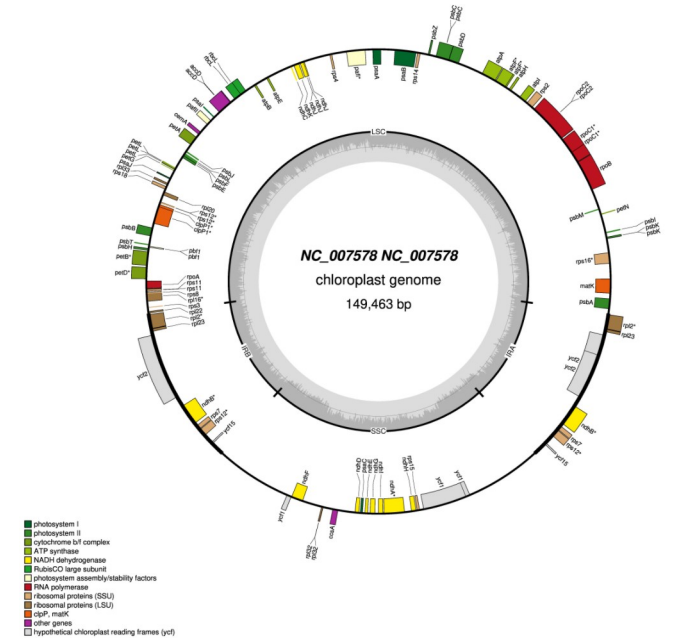
reference-guided tools:

- pomoxis (reference-guided mode)

None of widely used *de novo* assembly tools worked well

Reference was constructed from three plastomes – the one belonged to *L. sativa*, and the others were produced by pomoxis-medaka pipeline both in reference-guided and *de novo* modes. Mapped reads were used for assembly with pomoxis-medaka on *L. sativa* reference.

Resulting plastome of *C. callicephalo* had length 149,463 bp and contained 35 tRNA, 120 protein-coding genes and 13 putative pseudogenes.



Acknowledgments: The research was done within the framework of the grant assignment No. 075-15-2019-1670 and conducted on the base of the Unique Scientific Installation "Scientific Center of Plant Biotechnology, Genomics and Conservation" of FSFIS "NBG-NSC" and facilities of Shared Access Center Bioinformatics of FRC Institute of Cytology of Siberian Branch of the RAS.