

## Supplementary Material

# Characterization of the Complete Mitochondrial Genome of Sea Duck *Mergus serrator* and Comparison with other Anseriformes Species



Peng Chen<sup>1</sup>, Jiaqi Li<sup>1</sup>, Hongbo Li<sup>2</sup>, Qin Lu<sup>3</sup>, Wei Liu<sup>1,4,5\*</sup> and Jianliang Zhang<sup>1\*</sup>

<sup>1</sup>Nanjing Institute of Environmental Sciences, Ministry of Ecology and Environment, Nanjing 210042, China

<sup>2</sup>National Marine Environmental Monitoring Center, Ministry of Ecology and Environment, Dalian 116023, China.

<sup>3</sup>Nantong Museum, Nantong 226001, China

<sup>4</sup>College of Marine Life Sciences and Frontiers Science Center for Deep Multispheres and Earth System, Ocean University of China, Qingdao 266003, China

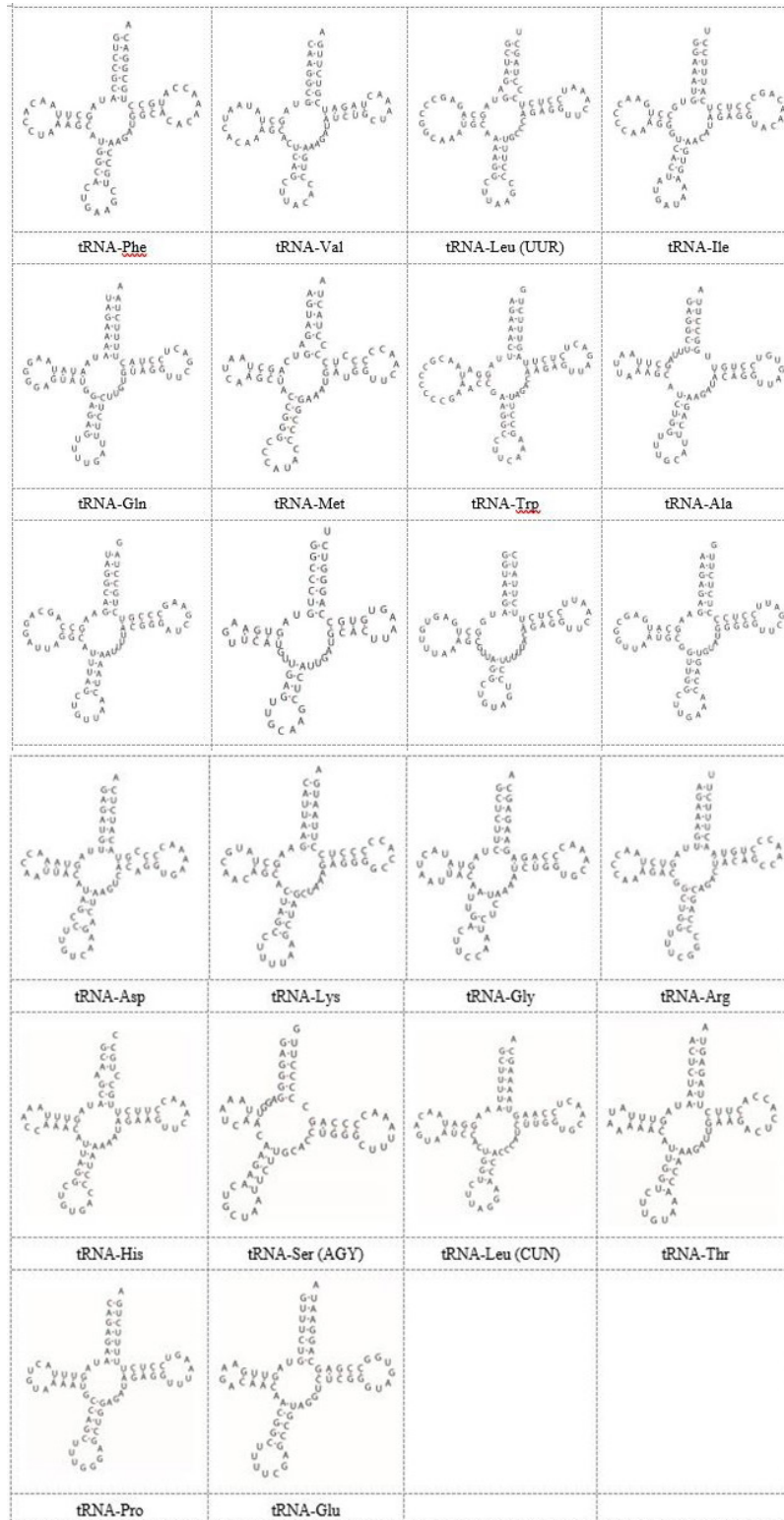
<sup>5</sup>Institute of Evolution and Marine Biodiversity, Ocean University of China, Qingdao 266003, China.

\* Corresponding author: lwecology@126.com  
0030-9923/2023/0005-2365 \$ 9.00/0



Copyright 2023 by the authors. Licensee Zoological Society of Pakistan.

This article is an open access  article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (<https://creativecommons.org/licenses/by/4.0/>).

Supplementary Fig. 1. General structure of 22 transfer RNAs of *M serrator*.