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Background

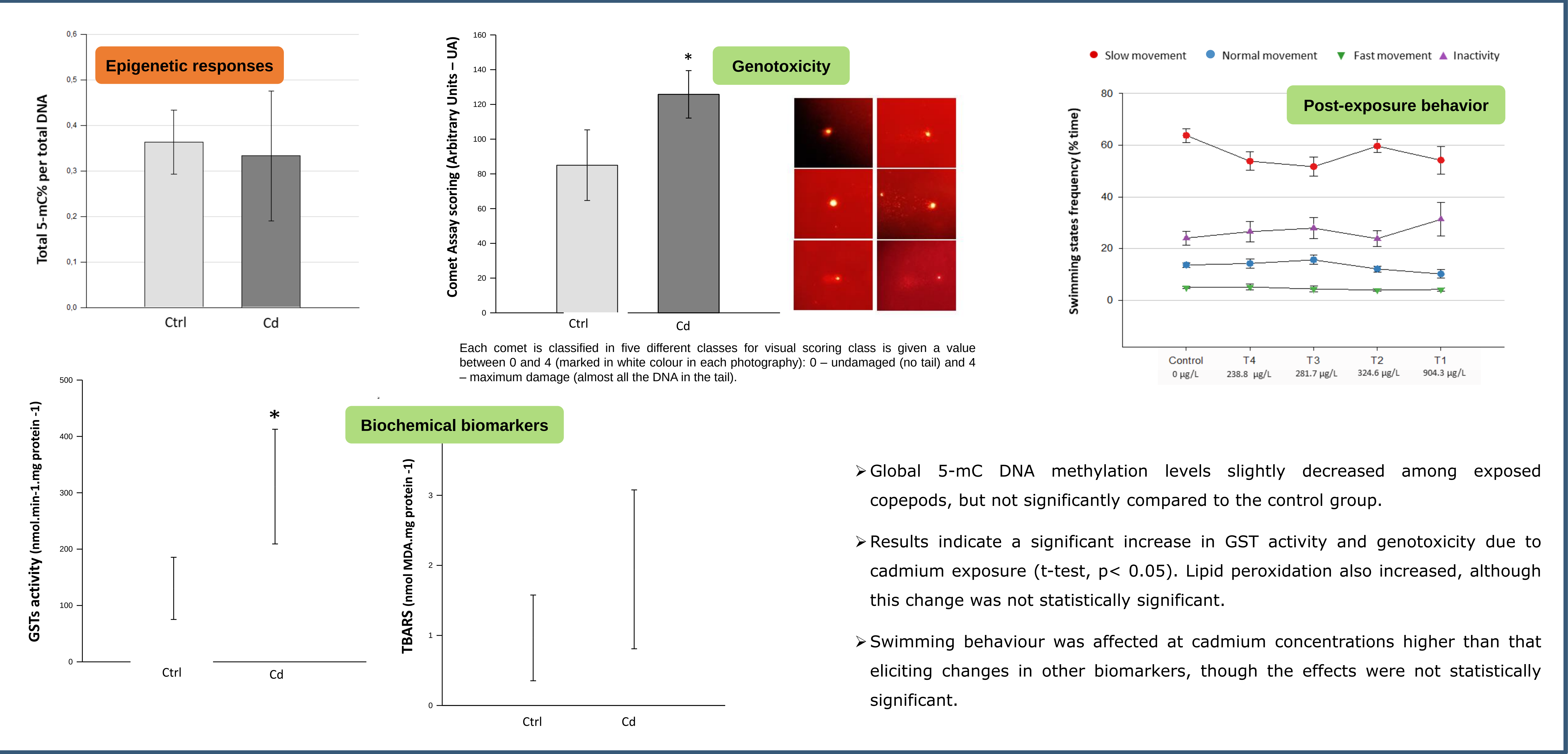
Increasing anthropogenic activities are impacting aquatic ecosystems, leading to contamination events that pose potential deleterious effects to organisms [1]. Cadmium is a trace metal with no essential biological role known and particularly harmful due to its high toxicity, persistence, and bioaccumulative properties[2].

Epigenetic and phenotypic responses that are responsive to Cd exposure can be integrated to understand the toxicity of the metal to copepods.

Aim

This study aims to understand the effects of short-term cadmium exposure on the copepod *Acartia tonsa*, a zooplanktonic marine species with important ecological role [3], integrating total **DNA methylation** changes with phenotypic changes concerning **DNA damage** and **oxidative stress** at the sub-individual level, as well as **behaviour** at the individual level.

Results



References

[1] Intergovernmental Panel on Climate Change (IPCC). (2023). *Climate change 2023: Synthesis report. Contribution of Working Groups I, II, and III to the Sixth Assessment Report of the Intergovernmental Panel on Climate Change* (H. Lee & J. Romero, Eds.). IPCC. <https://doi.org/10.59327/IPCC/AR6-9789291691647>

[2] Sharma, M., Kant, R., Sharma, A. K., & Sharma, A. K. (2024). Exploring the impact of heavy metals toxicity in the aquatic ecosystem. *International Journal of Energy and Water Resources*, Table 1. <https://doi.org/10.1007/s42108-024-00284->

[3] Turner, J. T. (2019). Trends in Copepod Studies. Distribution, Biology and Ecology. *Journal of Plankton Research*, 41(2), 203–205. <https://doi.org/10.1093/plankt/fbz001>

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Takeaway

- These findings highlight the sub-lethal effects of cadmium on *Acartia tonsa* and the convenience of multi-level biomarkers in assessing ecosystem health under human-induced stress.
- Despite limited knowledge on copepod epigenetics, evidence suggests epigenetic changes may aid rapid adaptation and phenotypic plasticity.
- This approach may clarify the links between epigenetic changes, gene expression, and phenotypic responses, helping to characterize the molecular basis of Adverse Outcome Pathways (AOPs) for cadmium-induced toxicity in *Acartia tonsa*.