

Combined effects of temperature and metal contamination on membrane fatty acid composition, desaturases and elongases in fathead minnow (*Pimephales promelas*)

Introduction

Long-chain polyunsaturated fatty acids (LC-PUFA), are essential for multiple physiological processes, including the maintenance of cell membrane structural integrity [1,2,3,4]. Biosynthesis of these fatty acids involves sequential desaturation and elongation of PUFA precursors. Two groups of enzymes are implicated in this process: desaturases (FADS) which incorporate double bonds into fatty acyl chains and elongases (ELOVL) which catalyze the condensation step in the elongation process [5]. In ectotherms, temperature influences the extent of unsaturation of biological membranes, cold-acclimated animals expressing a higher percentage of membrane phospholipid polyunsaturation compared to warm-acclimated conspecifics [6]. This process is known as homeoviscous adaptation (HVA) and it ensures membrane function and integrity for a range of acclimation temperatures, likely through modulations of desaturase and elongase gene transcription and activity. Some metals, such as Cd and Ni, can induce the production of reactive oxygen species (ROS), which may in turn lead to lipid peroxidation, PUFA being particularly vulnerable to ROS.

Study objectives

The aim of this study is to understand the combined effects of temperature and metal contamination (Cd and Ni) in fathead minnow (*Pimephales promelas*) muscle and brain on (i) the fatty acid composition of membrane phospholipids; (ii) the transcription level of desaturase and elongase genes; (iii) Differences in desaturase and elongase transcription between the two tissues

Materials and methods



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During warm acclimation (30°C) PUFA percentage decreased FA composition of brain phospholipids varied little with and SFA increased compared to 15°C and 25°C. temperature acclimation compared to muscle. Agrees with the HVA theory. Except for a decrease at 15°C under Ni exposure, PUFA

Cd exposure interfered with the normal warm acclimation percentage remained largely unaffected by metals. response of cell membrane composition at 30°C



Transcription level of genes encoding for desaturases (*fads2*: $\Delta 5/\Delta 6$ desaturase; *scd2*: Steroyl-CoA desaturase) and elongases (elovl6 and elovl5 encoding respectively for elongase6 and elongase5) varied between tissues: brain tissue showed a higher

temperatures. Temperature and metal combinations had different effects on the transcription level of desaturase and elongase genes,

In the muscle of Cd-exposed fish, desaturase and elongase transcription levels remained mostly unchanged in spite of a decrease in PUFA percentages. Inversely for Ni at 15°C, despite the increase of gene transcription levels (elovl6, scd2), PUFA In the brain, desaturase and elongase genes were mostly induced at low temperature and their level of transcription was lower at 25°C and 30°C. In contrast, PUFA percentage in warm-acclimated fish did not differ from the percentage in coldacclimated fish. Upregulation of gene expression at colder temperatures likely compensates for cold-induced reductions in enzyme activity, allowing maintenance of PUFA concentrations. This regulation is important as the functions of neural tissues

- in brain.
- Temperature and metal combinations had different effects on desaturase and elongase gene transcription levels.