Accepted Manuscript

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PII: S0016-6480(17)30782-7

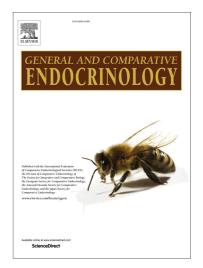
DOI: https://doi.org/10.1016/j.ygcen.2018.07.013

Reference: YGCEN 12992

To appear in: General and Comparative Endocrinology

Received Date: 17 November 2017

Revised Date: 8 June 2018 Accepted Date: 24 July 2018



Please cite this article as: Benjamin, H., Joanne, B., Pierre, G., Divergent evolution and clade-specific duplications of the Insulin-like Receptor in malacostracan crustaceans, *General and Comparative Endocrinology* (2018), doi: https://doi.org/10.1016/j.ygcen.2018.07.013

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ACCEPTED MANUSCRIPT

Divergent evolution and clade-specific duplications of the Insulin-like

Receptor in malacostracan crustaceans

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Abstract

The Insulin-like Receptors (IRs) are an important protein family, represented by three members in vertebrates, two of which are well-known for their implication in metabolism (Insulin Receptor) and growth (IGF Receptor). In contrast, little is known about these receptors in invertebrates, in which a single gene generally exists except for a part of insects and other occasional species-specific duplications. In this study, we used publicly available sequences as well as de novo assembled transcriptomes to investigate the IR evolution in malacostracan crustaceans, animals in which the Insulin/IGF pathway is known to be implicated in sexual development through the androgenic gland hormone. We described the evolutionary divergence of malacostracan IRs compared to all the other metazoan sequences, including other pancrustaceans. We also demonstrated two well conserved duplications of IRs:

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