Cloning and nutritional regulation of polyunsaturated fatty acid desaturase and elongase of a marine teleost, the nibe croaker Nibea mitsukurii

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Abstract

We identified fatty acid desaturase (fads)-like and elongase (elovl)-like genes from nibe croaker to better understand the molecular basis of n-3 highly unsaturated fatty acid metabolism in marine fish. Phylogenetic analysis revealed that the fads-like and elovl-like genes were classified into the fads6 and elovl5 groups, respectively. We investigated the effects of various levels of docosahexaenoic acid (DHA)-enriched live feed, Artemia nauplii, on larval growth, survival, and fads-like and elovl-like gene expression. After a 15-day feed trial, total length, body weight, and survival were all significantly improved by the supplementation of Artemia

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with DHA. This result indicates that nibe croaker cannot endogenously produce enough DHA. Furthermore, the fads-like gene transcripts in larvae fed on oleic acid-enriched Artemia were significantly higher than those in larvae on 100% DHA-enriched Artemia. In contrast, no significant differences were observed in the transcript levels of the elovl-like gene. These data indicate that the fads6-like gene was controlled by negative feedback from the quantity of DHA stored in the larval body. These results have implications for the functionality of the fads-like gene in nibe croaker.

Keywords Artemia - Docosahexaenoic acid - n-3 Highly unsaturated fatty acid - Fatty acid desaturase - Fatty acid elongase - Nibea mitsukurii