

Development of teleost intermuscular bones undergoing intramembranous ossification based on histological-transcriptomic-proteomic data

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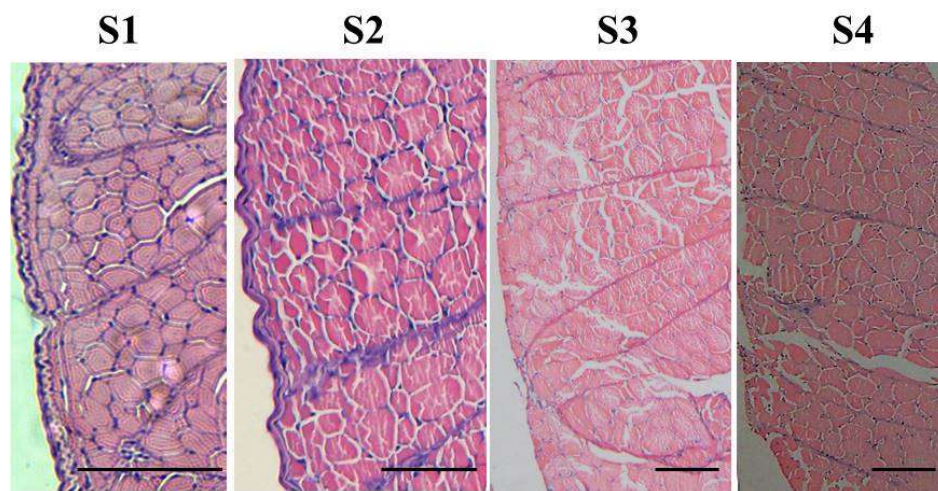


Figure S1 The histology structure with HE staining of muscle tissues at four stages. Scales bars = 50 μ m.

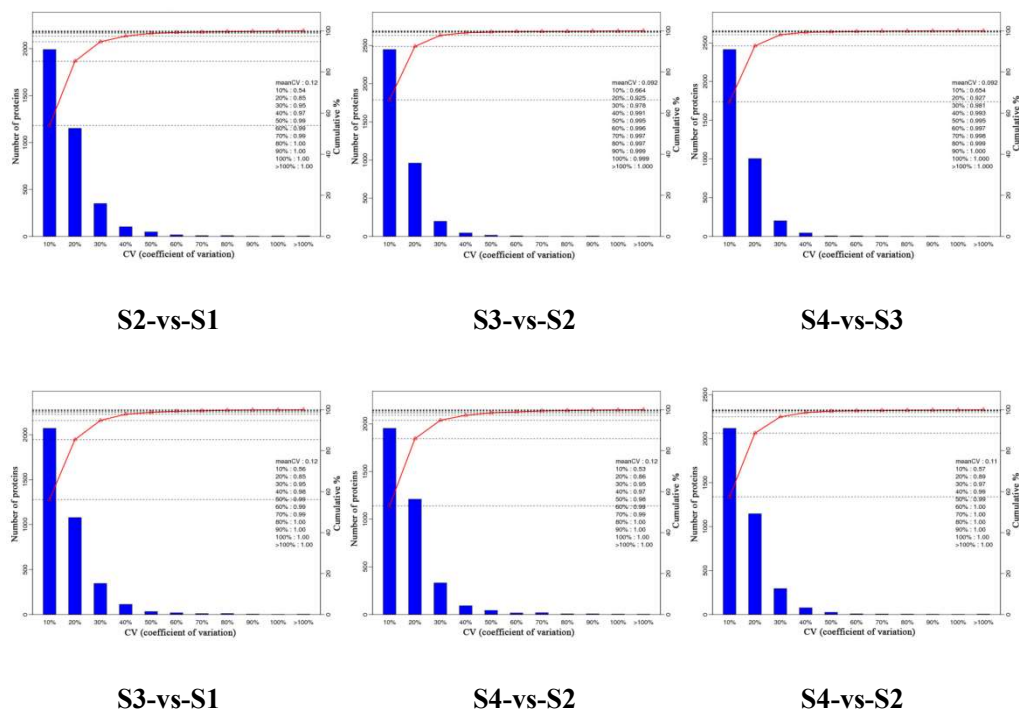
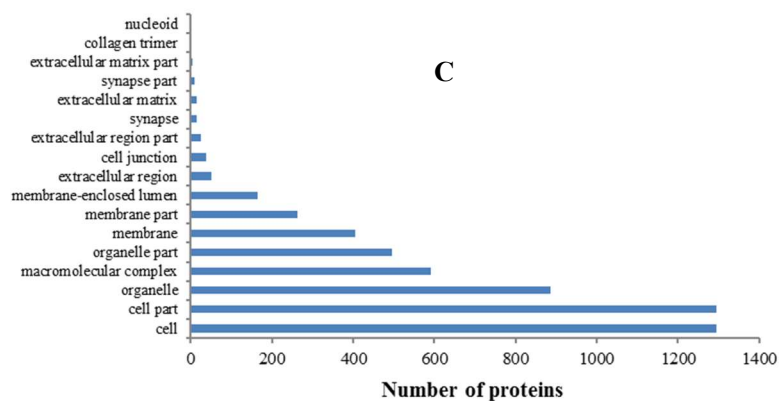
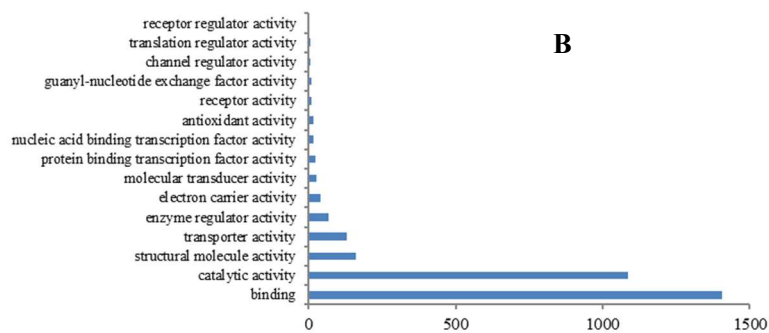
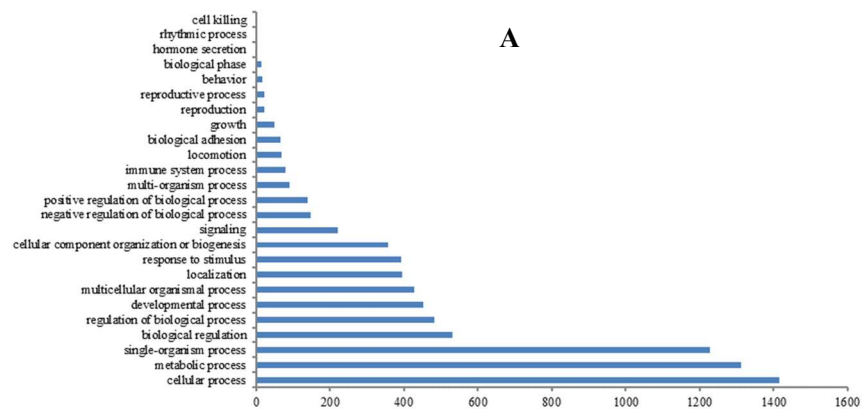


Figure S2 Repeatability analysis of data obtained from iTRAQ in different groups.



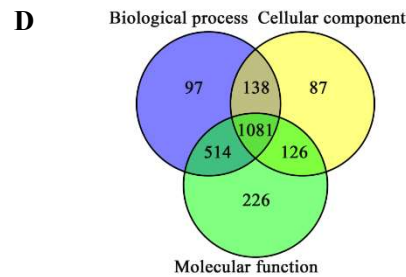


Figure S3 Proteins were functionally annotated for (A) biological process, (B) molecular function, as well as (C) cellular component and Venn diagram of proteins annotated for three processes (D).

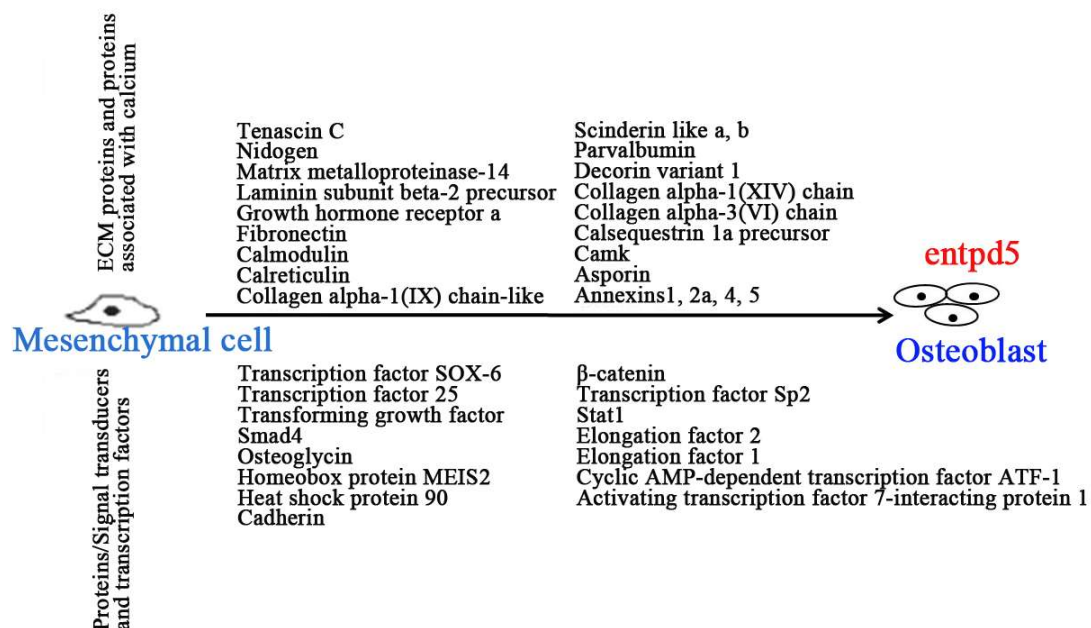


Figure S4 Schematic representation of the key proteins in osteoblastogenesis during IB ossification process.

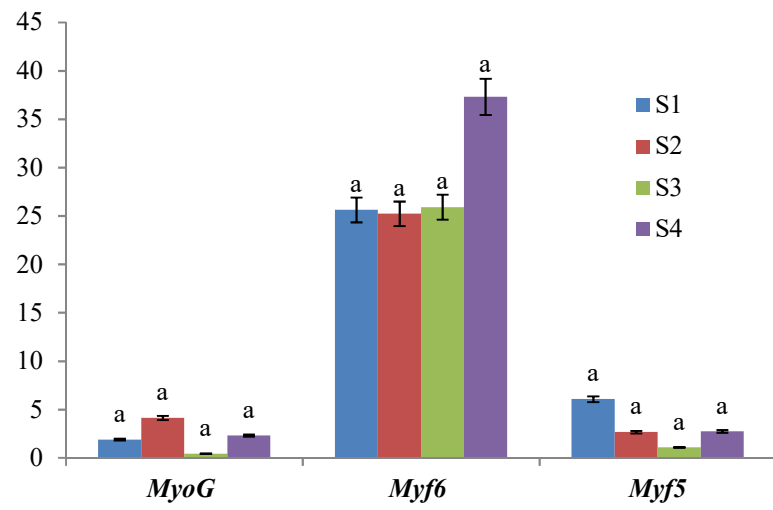


Figure S5 Expression of muscle-related genes at different stages based on RNA-seq data.

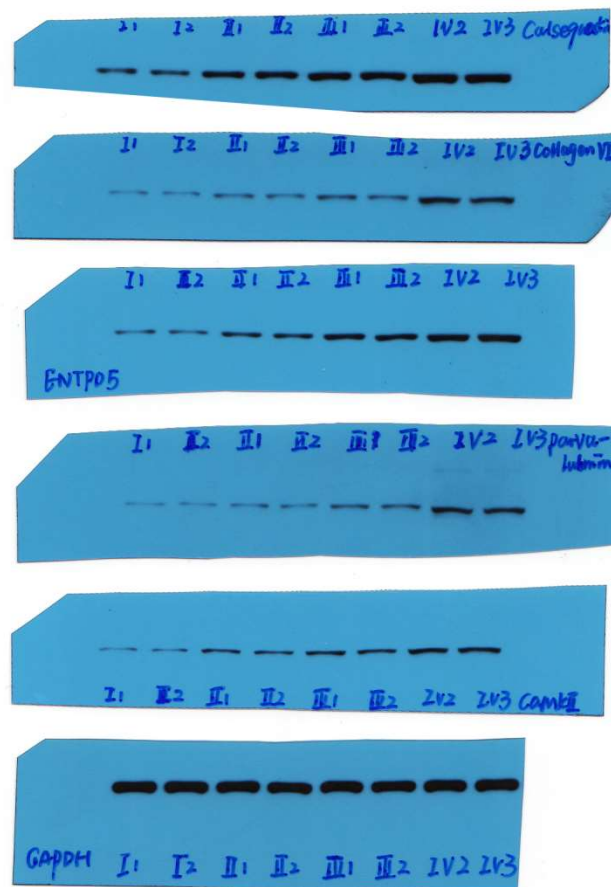


Figure S6 The raw data photo of western blot for five proteins; GAPDH is used as a loading control.

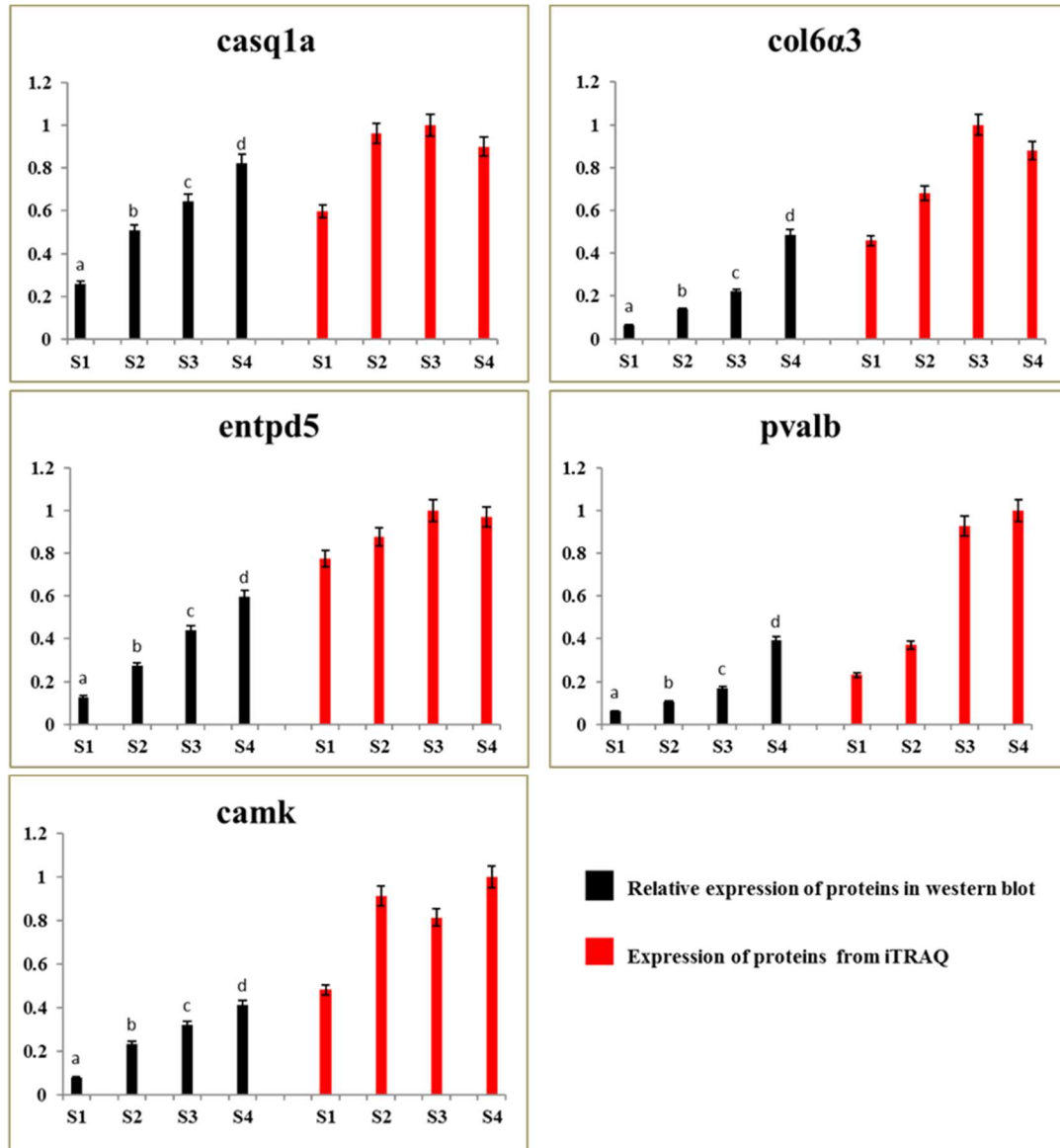


Figure S7 Comparison of expression levels for the 5 detected bone-related proteins based on iTRAQ and western blot results. Different letters above pillars for each stage indicated statistical significance at the level of $p < 0.05$.

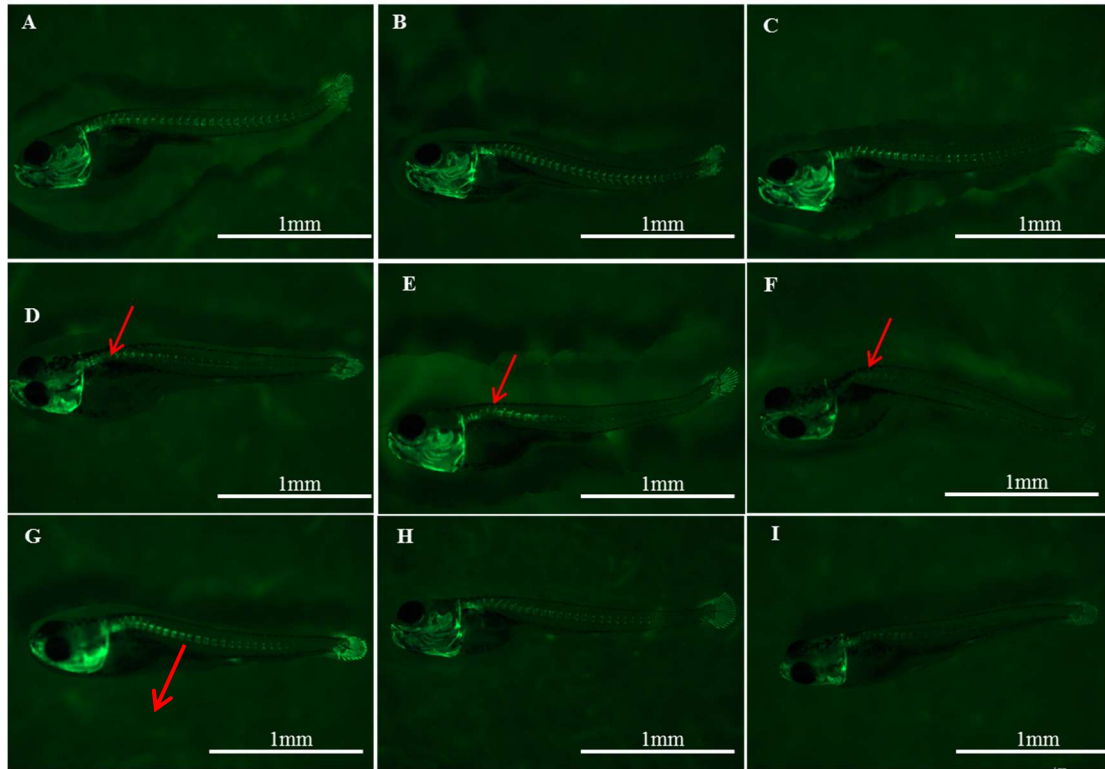


Figure S8 Effects of IWP-L6 (D–F) and K02288 (G–I) on Tg (*sp7/eGFP*) expression of vertebrate in zebrafish larvae. Control larvae (A, B, C) were treated with DMSO. Red arrows represent that part of vertebrate had a curved phenomenon.