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Analyses of the molecular mechanisms associated with salinity adaption of *Trachidermus fasciatus* through combined iTRAQ-based proteomics and RNA sequencing-based transcriptomics

Qian Ma, Xinfu Liu, Wenrong Feng, Shufang Liu, Zhimeng Zhuang

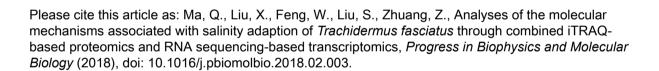
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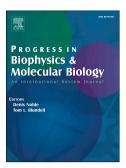
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ACCEPTED MANUSCRIPT

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4	Qian Ma ^{1,2} , Xinfu Liu ¹ , Wenrong Feng ¹ , Shufang Liu ^{1,2} , Zhimeng Zhuang ^{1,3*}
5 6 7 8 9	 Yellow Sea Fisheries Research Institute, Chinese Academy of Fishery Sciences, Qingdao 266071, China Function Laboratory for Marine Fisheries Science and Food Production Processes, Qingdao National Laboratory for Marine Science and Technology, Qingdao 266200, China Function Laboratory for Marine Biology and Biotechnology, Qingdao National Laboratory for Marine Science and Technology, Qingdao 266200, China
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	*Correspondence to: Zhimeng Zhuang, Yellow Sea Fisheries Research Institute, 106 Nanjing Road,

Qingdao 266071, P.R. China. Email: zhuangzm@ysfri.ac.cn; Fax: 0086-532-85811514; Tel: 0086-532-85836344

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