

Accepted Manuscript

Transcriptomics analysis reveals candidate genes and pathways for susceptibility or resistance to Singapore grouper iridovirus in orange-spotted grouper (*Epinephelus coioides*)

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PII: S0145-305X(18)30341-0

DOI: [10.1016/j.dci.2018.09.003](https://doi.org/10.1016/j.dci.2018.09.003)

Reference: DCI 3250

To appear in: *Developmental and Comparative Immunology*

Received Date: 1 July 2018

Revised Date: 4 September 2018

Accepted Date: 4 September 2018

Please cite this article as: Yang, M., Wang, Q., Wang, S., Wang, Y., Zeng, Q., Qin, Q., Transcriptomics analysis reveals candidate genes and pathways for susceptibility or resistance to Singapore grouper iridovirus in orange-spotted grouper (*Epinephelus coioides*), *Developmental and Comparative Immunology* (2018), doi: 10.1016/j.dci.2018.09.003.

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