## **Accepted Manuscript**

Identification of microRNAs with heat stress responsive and immune properties in *Marsupenaeus japonicus* based on next-generation sequencing and bioinformatics analysis: Essential regulators in the heat stress-host interactions

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PII: \$1050-4648(18)30299-7

DOI: 10.1016/j.fsi.2018.05.030

Reference: YFSIM 5311

To appear in: Fish and Shellfish Immunology

Received Date: 31 March 2018

Revised Date: 9 May 2018

Accepted Date: 16 May 2018

Please cite this article as: Zheng J, Cao J, Mao Y, Su Y, Wang J, Identification of microRNAs with heat stress responsive and immune properties in *Marsupenaeus japonicus* based on next-generation sequencing and bioinformatics analysis: Essential regulators in the heat stress-host interactions, *Fish and Shellfish Immunology* (2018), doi: 10.1016/j.fsi.2018.05.030.

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## ACCEPTED MANUSCRIPT

1	Identification of microRNAs with heat stress responsive and immune properties
2	in Marsupenaeus japonicus based on next-generation sequencing and
3	bioinformatics analysis: essential regulators in the heat stress-host interactions
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17	Abstract
18	Summer mortality syndrome is one of the most serious issue for Marsupenaeus
19	japonicus aquaculture in China. Since it causes massive economic loss and threatens
20	sustainability of M. japonicus aquaculture industry, thus, there is an urgent desire to
21	reveal the heat stress-host interactions mechanisms that lead to mass mortalities of $M$ .
22	japonicus in hot summer months. MicroRNAs (miRNAs) are small noncoding RNAs
23	that involved in regulation of diverse biological processes, including stress and
24	immune response, and might serve as potential regulators in the heat stress-host
25	interactions. In the present study, miRNAs with heat stress responsive and immune
26	properties were identified and characterized in M. japonicus by small RNA
27	sequencing and bioinformatics analysis. In total, 79 host miRNAs were identified,
28	among which 15 miRNAs were differentially expressed in response to heat stress.
29	Target genes prediction and function annotation revealed that a variety of host cellular
30	processes, such as signal transduction, transcription, anti-stress response, ribosomal

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