Accepted Manuscript

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PII: S0378-1119(17)30787-4

DOI: doi:10.1016/j.gene.2017.09.054

Reference: GENE 42203

To appear in: Gene

Received date: 29 March 2017 Revised date: 24 September 2017 Accepted date: 25 September 2017

Please cite this article as: Siyu He, Lulu Zhu, Fengchun Liu, Quan Liu, Yanjing Shao, Mengqing Hua, Han Ding, Wei Shao, Yinan Du, Xin Hou, Cuiping Ren, Miao Liu, Jijia Shen, Functions of the Vasa gene in Schistosoma japonicum as assessed by RNA interference. The address for the corresponding author was captured as affiliation for all authors. Please check if appropriate. Gene(2017), doi:10.1016/j.gene.2017.09.054

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ACCEPTED MANUSCRIPT

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Abstract

Vasa, an enzyme belonging to the helicase family, contributes to the regulation of reproductive system development in many species. Thus, we hypothesized that the *Vasa3* gene may function in the reproductive system of the parasite *Schistosoma japonicum* (*S. japonicum*), which is a major causative agent of schistosomiasis. It is a severe disease globally affecting humans and animals. To test this hypothesis, we firstly conducted whole mount *in situ* hybridization analyses and found that the *S. japonicum Vasa3* (*SjVasa3*) gene was expressed mainly in the reproductive organs. We then explored the reproductive functions of *Vasa3* in *S. japonicum* using RNA interference (RNAi) techniques. Coupled schistosomes collected from mice 28 days post infection (dpi) were transfected three times with *SjVasa3*-specific small interfering RNA (siRNA) and cultured *in vitro* for up to 10 days. As measured by quantitative PCR (qPCR) and Western blot analysis, levels of

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