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Genetic variation on the NFKB1 genes associates with the outcomes of HCV infection among Chinese Han population

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## **ACCEPTED MANUSCRIPT**

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Chinese Han population Hao-zhi Fan<sup>1,2¶</sup>, Peng Huang<sup>1,2¶</sup>, Jian-guo Shao<sup>3</sup>, Ting Tian<sup>1,2</sup>, Jun Li<sup>4</sup>, Feng Zang <sup>1,2</sup>, Mei Liu <sup>1,2</sup>, Hong Xue <sup>3</sup>, Jing-jing Wu <sup>1,2</sup>, Rong-bin Yu<sup>1,2</sup>, Ming Yue<sup>4,\*</sup>

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