

Accepted Manuscript

Simultaneous RNA-seq based transcriptional profiling of intracellular *Brucella abortus* and *B. abortus*-infected murine macrophages

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PII: S0882-4010(17)31142-7

DOI: [10.1016/j.micpath.2017.10.029](https://doi.org/10.1016/j.micpath.2017.10.029)

Reference: YMPAT 2538

To appear in: *Microbial Pathogenesis*

Received Date: 9 September 2017

Revised Date: 15 October 2017

Accepted Date: 16 October 2017

Please cite this article as: Hop HT, Arayan LT, Reyes AWB, Huy TXN, Min W, Lee HJ, Son JS, Kim S, Simultaneous RNA-seq based transcriptional profiling of intracellular *Brucella abortus* and *B. abortus*-infected murine macrophages, *Microbial Pathogenesis* (2017), doi: 10.1016/j.micpath.2017.10.029.

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Title: Simultaneous RNA-seq based transcriptional profiling of intracellular *Brucella abortus* and *B. abortus*-infected murine macrophages

Running title: RNA-seq analysis of *B. abortus* and *B. abortus*-infected macrophages

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