## Accepted Manuscript

Estimation of Rickettsia rickettsii copy number in the blood of patients 1 with Rocky Mountain spotted fever suggests cyclic diurnal trends in bacteremia

Kato Cecilia, Chung Ida, Paddock Christopher

PII: S1198-743X(15)01086-1

DOI: 10.1016/j.cmi.2015.12.019

Reference: CMI 477

To appear in: Clinical Microbiology and Infection

Received Date: 14 December 2015

Accepted Date: 18 December 2015

Please cite this article as: Cecilia K, Ida C, Christopher P, Estimation of Rickettsia rickettsii copy number in the blood of patients 1 with Rocky Mountain spotted fever suggests cyclic diurnal trends in bacteremia, *Clinical Microbiology and Infection* (2016), doi: 10.1016/j.cmi.2015.12.019.

This is a PDF file of an unedited manuscript that has been accepted for publication. As a service to our customers we are providing this early version of the manuscript. The manuscript will undergo copyediting, typesetting, and review of the resulting proof before it is published in its final form. Please note that during the production process errors may be discovered which could affect the content, and all legal disclaimers that apply to the journal pertain.



## ACCEPTED MANUSCRIPT

1 Estimation of *Rickettsia rickettsii* copy number in the blood of patients with Rocky

## 2 Mountain spotted fever suggests cyclic diurnal trends in bacteremia

3 Sir,

4 Infection with Rickettsia rickettsii, an obligate intracellular gram-negative bacterium, causes Rocky Mountain spotted fever (RMSF), one of the most lethal of all known infectious diseases. 5 Because R. rickettsii infects predominantly fixed endothelial cells of small and medium-sized 6 blood vessels, there are few quantitative assessments of bacteremia for RMSF and limited data 7 suggest that low numbers of R. rickettsii circulate in the peripheral blood of ill patients [1]. This 8 observation was described by early investigators who recognized that R. rickettsii bacteria could 9 scarcely be found in stained peripheral blood smears of patients, even those with severe disease 10 [2]. Herein we use TaqMan real-time PCR to evaluate levels of rickettsial DNA in a series of 11 clinical samples obtained from patients with RMSF. Whole blood (n=23), serum (n=6), and 12 plasma (n=1) specimens were collected from 13 US patients with RMSF from 2010-2015 and 13 submitted to the Rickettsial Zoonoses Branch Reference Diagnostic Laboratory at CDC 14 (Supplemental Table). Individual specimens were accompanied by clinical information that 15 varied in completeness. Each sample was positive for PanR8 *Rickettsia* spp., and RRi6 *R*. 16 *rickettsii*-specific targets [3], with controls performing as expected (Supplemental Figure). 17 RMSF specimens were quantified with PanR8 in duplicate (undiluted or 1:10), using ten-fold 18

dilutions of *R. rickettsii* DNA from 0.1 fg to 10.0 pg (1 fg = 0.8282 copies). Samples from

patients with fatal (n=20) and nonfatal (n=10) outcomes ranged from 1.41 x  $10^3 \pm 2.78 \times 10^2$  to

21  $2.05 \times 10^6 \pm 1.89 \times 10^3$  copies /mL (median = 1.63 x 10<sup>5</sup>), and 8.40 x 10<sup>1</sup> ± 4.19 x 10<sup>1</sup> to 3.95 x

22  $10^5 \pm 3.33 \times 10^4$  copies /mL (median = 2.82 x 10<sup>3</sup>), respectively (Supplemental Table). Copy

numbers of *R. rickettsii* DNA were significantly greater in patients with fatal outcomes (p < 0.05,

Download English Version:

## https://daneshyari.com/en/article/6128904

Download Persian Version:

https://daneshyari.com/article/6128904

Daneshyari.com