

A Small Rna that Regulates Pyrimidine and Methionine Metabolism is Necessary for Establishing *Coxiella Burnetii*'s Intracellular Niche During Early Stages of Infection

Shaun Wachter *, Department of Biological Sciences, The University of Montana, Missoula

Matteo Bonazzi, CNRS, Universite Montpellier, Montpellier

Kyle Shifflett, Department of Biological Sciences, The University of Montana, Missoula

Abraham S. Moses, Department of Biology and Center for Life in Extreme Environments, Portland State University, Portland

Rahul Raghavan, Department of Biology and Center for Life in Extreme Environments, Portland State University, Portland

Michael F. Minnick, Department of Biological Sciences, The University of Montana, Missoula

*Indicates Presenter

Coxiella burnetii is an obligate intracellular gammaproteobacterium and zoonotic agent of Q fever in humans. We previously identified 15 small RNAs (sRNAs) in *C. burnetii* with differential expression in the large and small cell developmental forms grown axenically and in infected host cells. Here, we describe the function of one of these sRNAs, termed *Coxiella burnetii* small RNA 12 (CbsR12). CbsR12 is highly expressed in both large- and small-cell variants in vitro, and is the dominant non-tRNA/rRNA/tmRNA transcript in both morphotypes during mammalian tissue culture infection. Through a combination of in vitro and in vivo assays, we have identified several targets of CbsR12. Of these, we have confirmed that CbsR12 binds to and upregulates translation of *carA* transcripts coding for carbamoyl phosphate synthetase A; an enzyme that catalyzes the first step of pyrimidine biosynthesis. In addition, CbsR12 binds and downregulates translation of *metK* transcripts coding for S-adenosyl methionine (SAM) synthase, an essential component of the methionine cycle. Furthermore, we have established that CbsR12 is necessary for full expansion of *Coxiella*-containing vacuoles (CCVs) and is linked to growth rate in a dose-dependent manner in the early phase of infection of Vero and THP-1 cell lines. This is the first characterization of a trans-acting sRNA of *C. burnetii* and the first description of a bacterial sRNA that regulates *carA* and *metK* expression. This study also illustrates the utility of transposon insertion mutants in elucidating *Coxiella*'s sRNAs and the importance of sRNA regulation in establishment of the intracellular CCV niche.