

Research Symposium

Pseudogenization of a resistance-nodulation-division (RND) efflux pump subunit underlies drug and cell envelope stress sensitivity in *Brucella ovis*

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Brucella species are intracellular bacterial pathogens that pose a significant threat to both animal and human health. To identify compounds that reduce *Brucella ovis* fitness in mammalian phagocytes, we conducted a high-throughput luminescence-based screen, which revealed dihydropyridine-class Ca^{2+} channel blockers nicardipine and cilnidipine as potential host-targeting anti-infectives. However, follow-up dose-response studies in pure culture revealed that these compounds directly inhibit *B. ovis* growth. To investigate possible dihydropyridine resistance mechanisms, we selected for *B. ovis* mutants tolerant to these drugs and identified single-base deletions in the *bepE* pseudogene. These mutations restored a functional open reading frame for BepE, a subunit of a resistance-nodulation-division (RND) efflux pump, increasing *B. ovis* resistance to dihydropyridine treatment. Given that *B. ovis* has undergone extensive pseudogenization and exhibits greater chemical

susceptibility than other *Brucella* species, we examined whether *bepE* influenced cell envelope integrity. *B. ovis* mutants with an intact *bepE* gene displayed enhanced resistance to membrane disruptors, including deoxycholate. To extend these findings, we investigated *bepE* function in *Brucella abortus*, a closely related zoonotic pathogen that encodes a fully intact BepE protein. Deleting *bepE* in *B. abortus* increased its susceptibility to deoxycholate and its sensitivity to cilnidipine during macrophage infection, indicating that *bepE* not only contributes to drug resistance in the intracellular niche but also supports *B. abortus* resistance to cell envelope stress. The results define *bepE* as a determinant of *Brucella* resistance to antimicrobial compounds, and demonstrate that its pseudogenization contribute to the heightened chemical sensitivity of *B. ovis* relative to other classical *Brucella* species.

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