

Lysozymes constitute an ancient family of proteins that are important components of the innate immune system of animals. They hydrolyze the 1,4-P-linkages between N-acetyl-d-glucosamine and N-acetylmuramic acid in the peptidoglycan of bacterial cell walls, and are part of a nonimmunological ancestral bactericidal system in vertebrates. Hydrolysis of peptidoglycan compromises cell wall integrity, causing cell lysis and bacterial death. Most bacteria engaged in commensal or pathogenic interactions with an animal host have evolved various strategies to evade lysozyme action. These include shielding by the outer membrane, in the case of most gram-negative bacteria, peptidoglycan modification, or the more recently emerged production of lysozyme inhibitors. In the case of *Brucella*, after invasion of the host, *Brucella* is phagocytosed by neutrophils (PMNs), macrophages, and dendritic cells. *Brucella* spp. resists a wide range of bactericidal cationic peptides and lysozyme, and is able to prevent lysosome fusion - and to locate in autophagosome-like compartments. Moreover, *B. abortus* cells elicit little respiratory burst and only reduced levels of degranulation in PMNs, thus contributing to a limited exposure of *Brucella* to lysozyme. We have identified a gene that encodes for a putative lysozyme inhibitor of the MhC family in *B. abortus* 2308. The gene was overexpressed and purified as a His-tagged protein, and the protein activity was determined in vitro against suspensions of *Micrococcus lysodeikticus*, confirming that it is a real lysozyme inhibitor. The role of this inhibitor in resistance of *Brucella* to lysozyme has been studied in a knock-out mutant, both in vitro as in vivo, using purified lysozyme, macrophages and human neutrophils.