



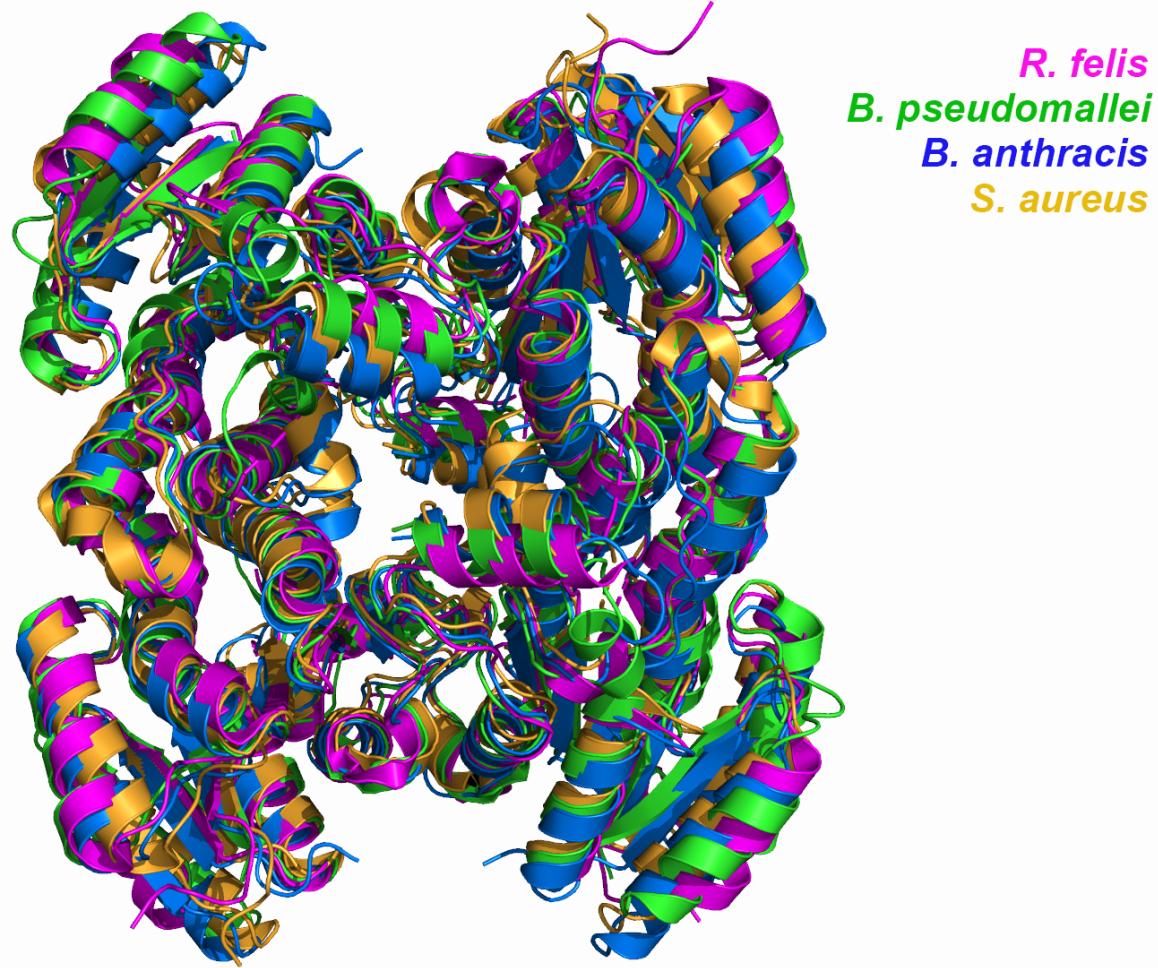
STRUCTURAL BIOLOGY  
COMMUNICATIONS

Volume 77 (2021)

Supporting information for article:

Crystal structure of acetoacetyl-CoA reductase from *Rickettsia felis*

Justas V. Rodarte, Jan Abendroth, Thomas E. Edwards, Donald D. Lorimer,  
Bart L. Staker, Sunny Zhang, Peter J. Myler and Krystle J. McLaughlin



**Supplementary Figure S1.** Structural Alignment of acetoacetyl-CoA reductases from *Rickettsia felis* (PDB entry 4kms) and *Burkholderia pseudomallei* (PDB entry 3ezl), and FabGs from *Staphylococcus aureus* (PDB entry 3osu) and *Bacillus anthracis* (PDB entry 2uvd). Tetramers were generated using crystallographic symmetry and aligned in PyMOL.