Running title: Structure of Mil, an archaeal Imp4-like protein. Authors: Ng *et al.* 

## Supplementary information

**Fig 1.** Stereo-view of the electron density map  $(2DF_o-mF_c)$  in the region around the disulphide bridge. The map is shown at  $1\sigma$  level. Secondary structural elements are colored as in Fig 2. A dashed line represents the hydrogen bond between Glu101 and Gly99. The electron density shows that the two cysteines present in the molecule form a disulphide bridge. Since these cysteines are not conserved, not even within the archaeal group, this disulphide bridge seems to be a unique feature in the structure of Mil. Cys96 is in the middle of the loop connecting the  $\beta$ F strand to the  $\alpha$ 3 helix, while Cys104 sits on the latter helix. Thus, the disulphide bridge is likely to stabilize the conformation of the loop containing Cys104. Further stabilization could be attained by a hydrogen bond between the side chain of Glu101 and the backbone amide of Gly99.

