



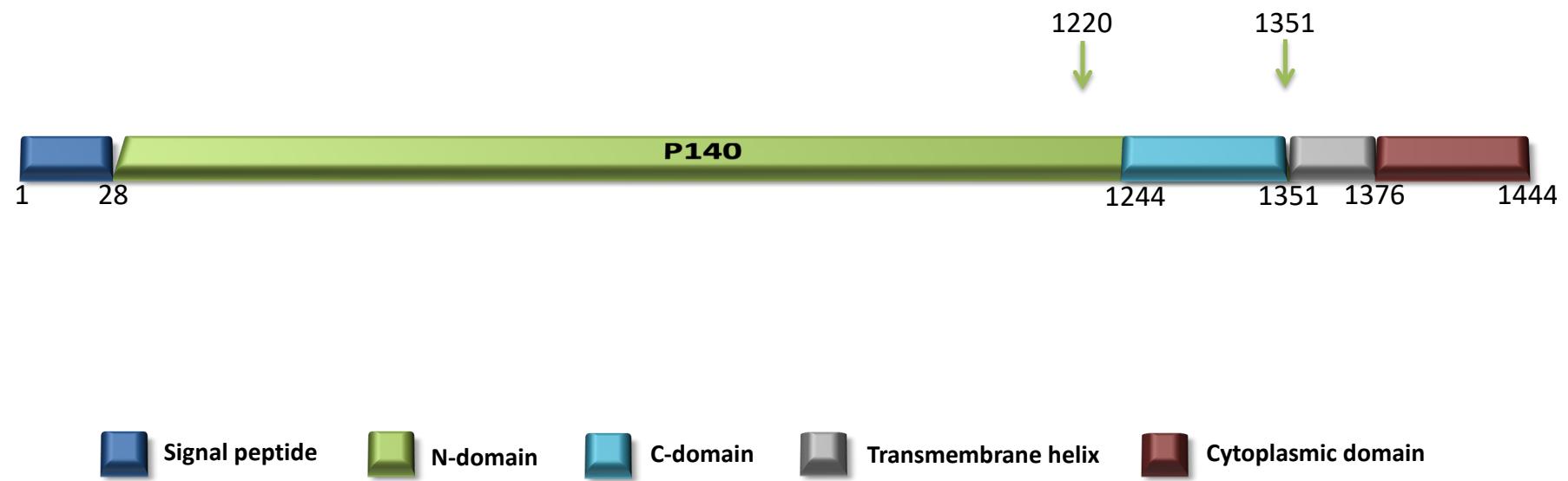
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Supporting information for article:

Alternative conformation of the C-domain of the P140 protein from *Mycoplasma genitalium*

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Supplementary Figure S1. P140 regions and C-Domain construct.

Schematic representation of the P140 regions predicted with the Psipred server¹: The signal peptide (blue), the extracellular region (orange), the transmembrane helix (gray) and the cytoplasmic region (brown). Green arrows indicate the C-domain construct used for crystallographic and biological studies.

| Primers | Sequence |
|---------|-------------------------------------|
| P140cF | AAGTTCTGTTCAGGGCCCGAAAATGGATGGTAAAC |
| P140cR | ATGGTCTAGAAAGCTTAATCTGCCACTGATTAAAC |

Supplementary Table S1.

Primers used in this study

References for the Supplementary information

1. McGuffin, L.J., Bryson, K. & Jones, D.T. The PSIPRED protein structure prediction server. *Bioinformatics* 16, 404-5 (2000).