Backbone and sidechain resonance assignments and secondary structure of Scc4 from *Chlamydia trachomatis*

Biomolecular NMR Assignments

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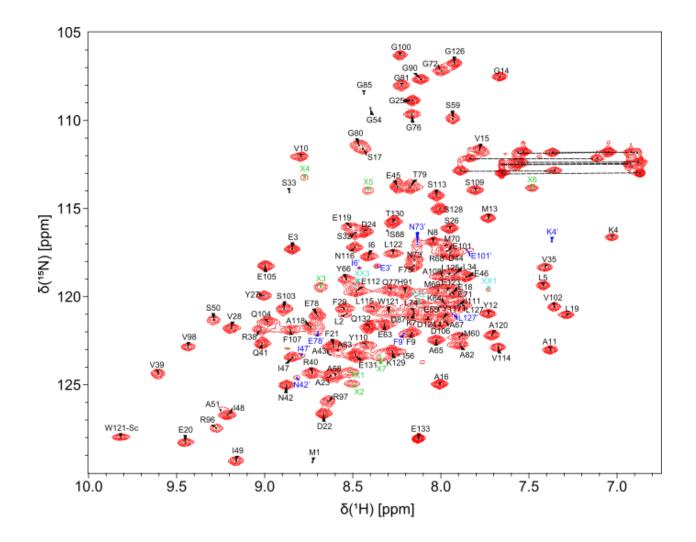
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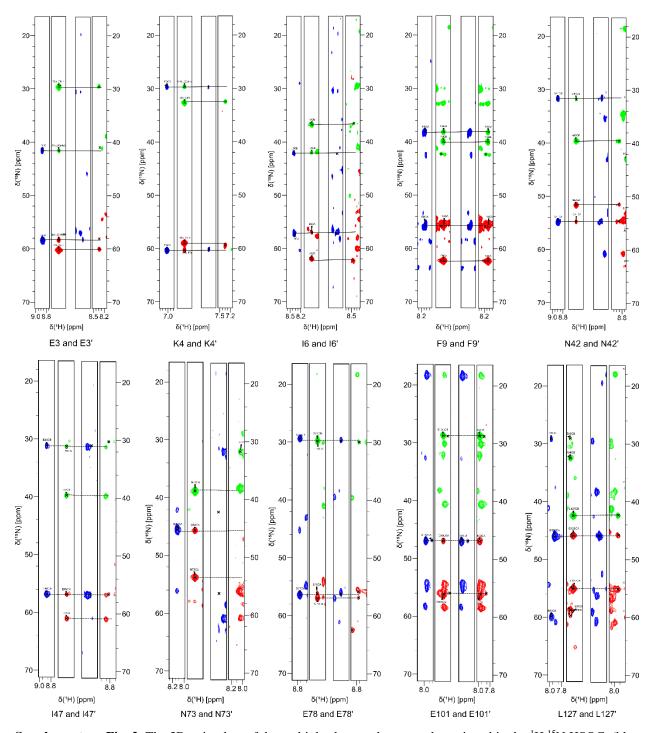
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Electronic Supplementary Material – Online Resource 1

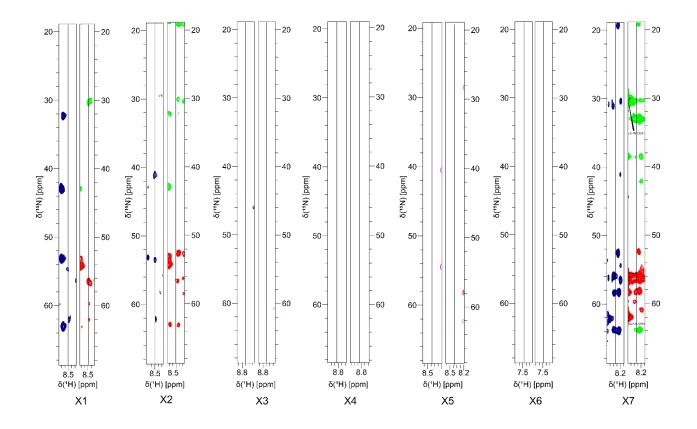
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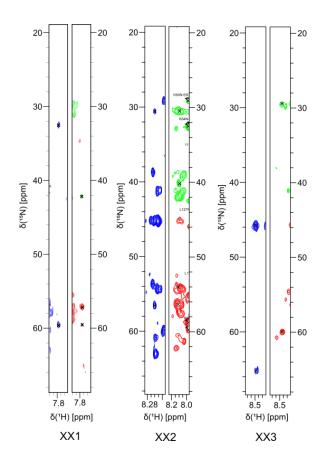
Supplementary Fig. 1 The ¹H,¹⁵N HSQC spectrum of Scc4 with assigned residues in black and the residues with multiple slow-exchange peaks in blue. The green labels (X1, X2, etc.) correspond to peaks with missing C^{α} and C^{β} resonances in the 3D NMR experiments. The cyan peaks (XX1, XX2, etc.) are unassigned peaks with ambiguous C^{α} and C^{β} resonances.



Supplementary Fig. 2 The 3D strip plots of the multiple slow-exchange peaks assigned in the ¹H,¹⁵N HSQC. (blue labels in Supplementary Fig. 1). In each pair of strip plots, the first strip is the ¹⁵N plane from the 3D CBCA(CO)NH experiment and the second strip is the corresponding ¹⁵N plane from the 3D HNCACB experiment. The horizontal lines indicate resonances used to assign the peaks. Assignment labels within the strip plots were not intended to be readable and can be disregarded. The contour levels in each plot are identical.



Supplementary Fig. 3 The 3D strip plots of the unassigned peaks in the ¹H,¹⁵N HSQC due to missing C^{α} and C^{β} resonances (green labels in Supplementary Fig. 1). In each pair of strip plots, the first strip is the ¹⁵N plane from the 3D CBCA(CO)NH experiment and the second strip is the corresponding ¹⁵N plane from the 3D HNCACB experiment. Assignment labels within the strip plots were not intended to be readable and can be disregarded. The contour levels in each plot are identical.



Supplementary Fig. 4 The 3D strip plots of the unassigned peaks in the ¹H,¹⁵N HSQC due to ambiguous C^{α} and C^{β} resonances (cyan labels in Supplementary Fig. 1). In each pair of strip plots, the first strip is the ¹⁵N plane from the 3D CBCA(CO)NH experiment and the second strip is the corresponding ¹⁵N plane from the 3D HNCACB experiment. Assignment labels within the strip plots were not intended to be readable and can be disregarded. The contour levels in each plot are identical.