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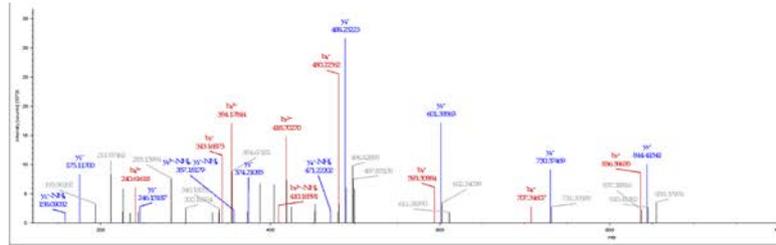
2 **Supplementary Figure 1** The full SDS-PAGE image of Figure 3a. The area within the rectangle was

3 adopted for Figure 3a.

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Spr0693

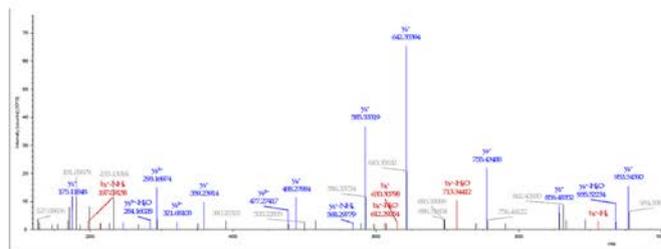
ADRHINELNQAR

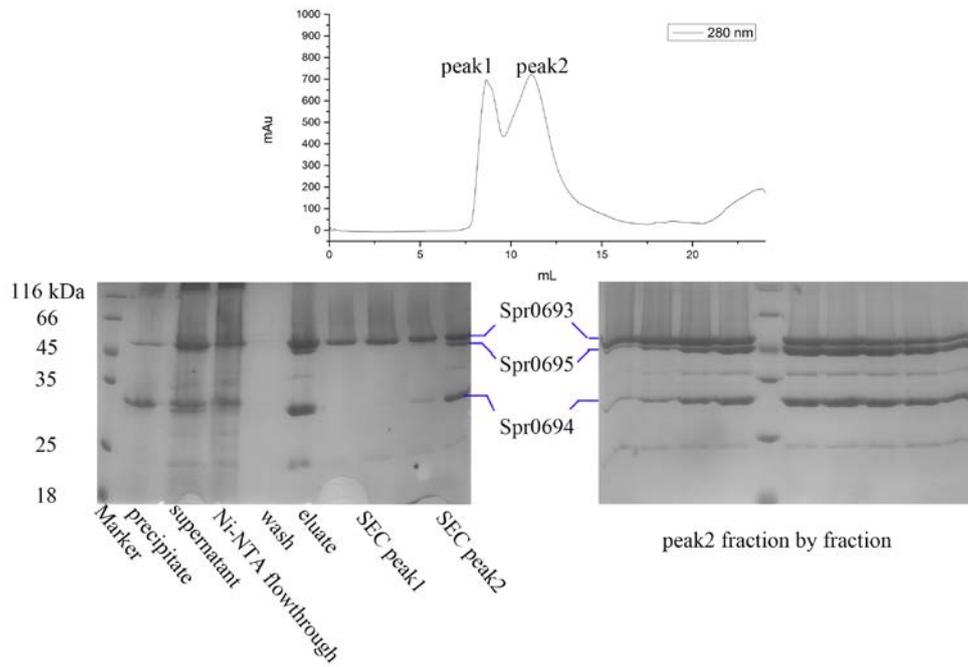


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Spr0695

VNDTSLPTLGPPEAR





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18 **Supplementary Figure 3** Size-exclusion chromatography profile with co-purification of  
 19 Spr0693-0694-0695.

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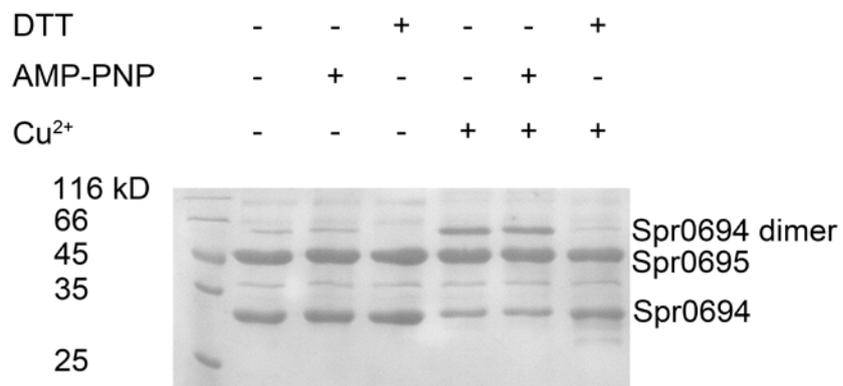
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40 **Supplementary Figure 4** The disulfide-crosslinking experiments of Spr0694<sub>G173C</sub>-0695 mutant at  
 41 different conditions. The reaction mix was separated by SDS-PAGE. Molecular masses are indicated  
 42 on the left and the corresponding proteins are indicated on the right.

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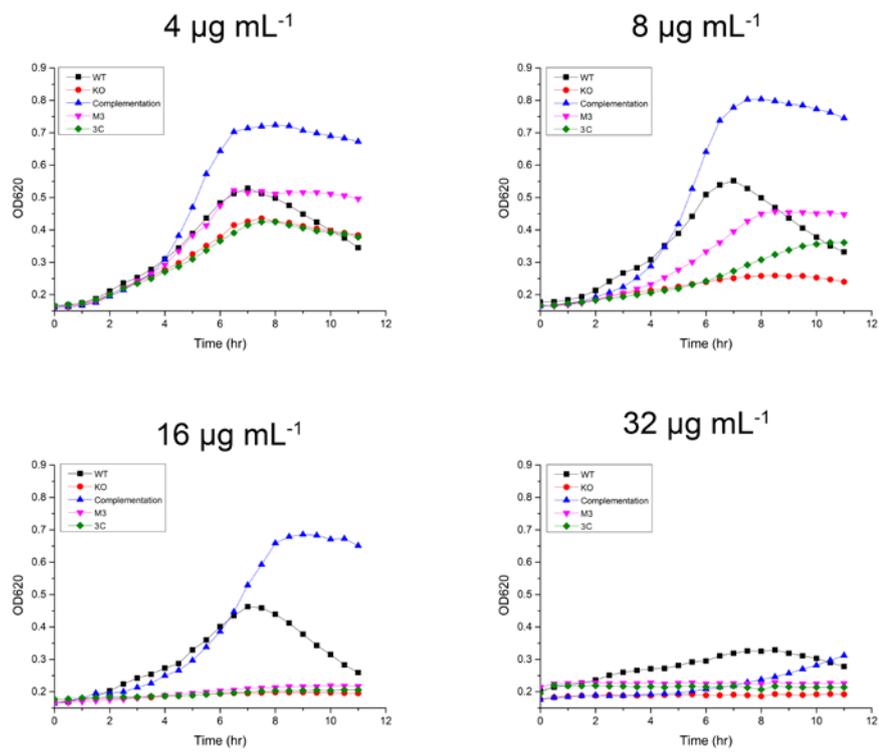
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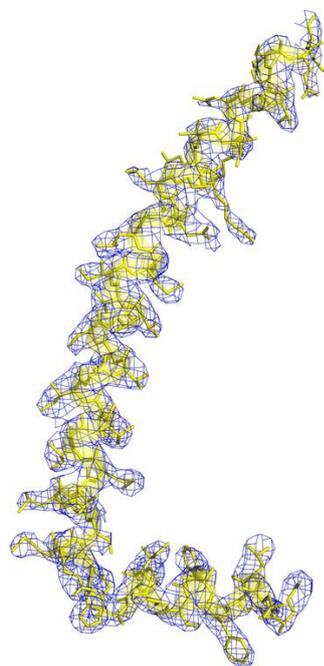
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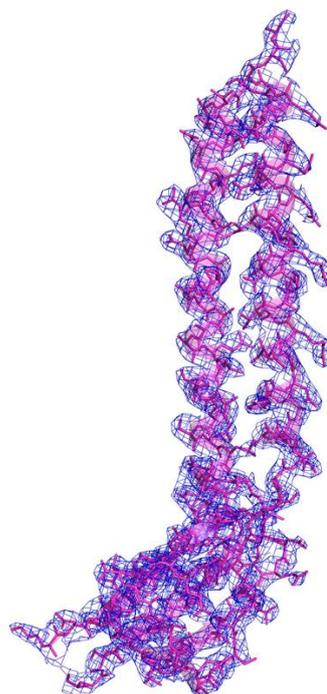
**Supplementary Figure 5** Growth curves of *S. pneumoniae* R6 and mutants in the presence of LL-37 at different concentrations.

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Spr0694-0695

b



Spr0693

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83 **Supplementary Figure 6** A portion of the *2Fo-Fc* electron density map of Spr0694-0695 and Spr0693,  
84 countered at 1.5  $\sigma$ . **(a)** Electron density map of TM1 of Spr0695. **(b)** Electron density map of the  
85  $\alpha$ -helical hairpin and lipoyl domains of Spr0693 monomer.

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| Strains, plasmids or primers            | Description  | Source       |
|---|--|--------------|
| <b>Strains</b>                          |  |              |
| WT                                      | <i>S. pneumoniae</i> R6; <i>spr0693-0694-0695</i> <sup>+</sup> ; Kan <sup>s</sup>  | <sup>1</sup> |
| KO                                      | <i>S. pneumoniae</i> R6 derivative; $\Delta$ <i>spr0693-0694-0695</i> ::Janus cassette; Kan <sup>r</sup>   | This study   |
| Complementation                         | KO strain transformed with pJWV25- <i>spr0693-0694-0695</i> ; Kan <sup>r</sup> , Tet <sup>r</sup>  | This study   |
| M3                                      | KO strain transformed with pJWV25-M3; Kan <sup>r</sup> , Tet <sup>r</sup>  | This study   |
| 3C                                      | KO strain transformed with pJWV25-3C; Kan <sup>r</sup> , Tet <sup>r</sup>  | This study   |
| <b>Plasmids</b>                         |  |              |
| pJWV25                                  | amp <sup>r</sup> , tet <sup>r</sup> , bgaA, P <sub>czeD</sub> ; integrate Zn <sup>2+</sup> -controlled target genes by double cross-over at the <i>S. pneumoniae</i> bgaA locus. | <sup>2</sup> |
| pJWV25- <i>spr0693-0695</i>             | amp <sup>r</sup> , tet <sup>r</sup> , bgaA, P <sub>czeD</sub> - <i>spr0693-0694-0695</i>   | This study   |
| pJWV25-M3                               | amp <sup>r</sup> , tet <sup>r</sup> , bgaA, P <sub>czeD</sub> - <i>spr0693-0694-0695</i> <sub>K207A-R208A-K210A</sub>  | This study   |
| pJWV25-3C                               | amp <sup>r</sup> , tet <sup>r</sup> , bgaA, P <sub>czeD</sub> - <i>spr0693-0694-0695</i> <sub>D49C-R208C/K52C-E205C/E374C-K207C</sub>  | This study   |
| <b>Primers for MICs</b>                 |  |              |
| <b>Nucleotide sequence (5' to 3')</b>   |  |              |
| Up-F                                    | GCGATATAACCAGCGCCTAGAATGGCAACTG  | This study   |
| Up-R                                    | CAAACCTCTGTGCTCGAGCACTCGCAGCACCGATTGCTGC<br>ATAC   | This study   |
| Down-F                                  | CAGCATTATCCTCTAGAGAAGTTGGAGTATCAATCCCAGT<br>TGCCC  | This study   |
| Down-R                                  | GCGTAGCCAAGAGCTGTGCGCATAGTTAAGAAG  | This study   |
| JC-F                                    | GGTGCTGCGAGTGCTCGAGCACAGAGTTTGTAGAAACGC<br>AAAAAGG   | This study   |
| JC-R                                    | GATTGATACTCCAACCTTCTCTAGAGGATAATGCTGAAAAC<br>TCCTTGAAG   | This study   |
| pJWV25-F                                | TGATAAGCTGAGCGCCGGTCGCTACCA  | This study   |
| pJWV25-R                                | CTTTTTCTTCATCTTGTCATCGTCATCCTTGTAGTCG  | This study   |
| pJWV25-Spr0693-F                        | GACCGGCGCTCAGCTTATCATTACATAACGAAGGGCTTC  | This study   |
| pJWV25-Spr0695-R                        | GACGATGACAAGATGAAGAAAAAGAATGGTAAAG   | This study   |
| Spr0695 <sub>K207A-R208A-K210A</sub> -F | GGAGGCTGCAGCGTCAGCAATATATGGGTTTGGTG  | This study   |
| Spr0695 <sub>K207A-R208A-K210A</sub> -R | CCCATATATTGCTGACGCTGCAGCCTCCGGACTAGTA  | This study   |
| Spr0695 <sub>D49C-K52C</sub> -F         | CAAGTCAATAAATGTATGACTTGCTCTCAGAAAAATATT<br>AGCGTC  | This study   |
| Spr0695 <sub>D49C-K52C</sub> -R         | GCTAATATTTTTCTGAGAGCAAGTCATACATTTATTGACT<br>TGACGAGATAGG   | This study   |
| Spr0695 <sub>E205C-K207C-R208C</sub> -F | GTTTATACTAGTCCGTGTGCTTGTGCTCAAAAATATATG<br>GGTTTGG   | This study   |



