

1 **Supplementary Table 2. Data collection and refinement statistics**

| <i>A. aeolicus</i> LpxC/LPC-069 | |
|-------------------------------------|-------------------------------|
| Data collection | |
| Wavelength | 1.0000 Å |
| Space group | P 61 |
| Cell dimensions | |
| <i>a</i> , <i>b</i> , <i>c</i> (Å) | 65.99, 65.99, 132.42 |
| α , β , γ (°) | 90.0, 90.0, 120.0 |
| Resolution (Å) | 50.0 – 1.62 (1.66 – 1.62) |
| R_{meas} | 9.7 (76.2) |
| $I / \sigma I$ | 20.13 (4.95) |
| Completeness (%) | 100 (100) |
| Redundancy | 12.4 (12.4) |
| Refinement | |
| Resolution (Å) | 29.53 – 1.62 (1.66 – 1.62) |
| Unique reflections | 41379 (2956) |
| Redundancy | 12.4 (12.4) |
| $R_{\text{work}} / R_{\text{free}}$ | 0.139/0.154 |
| No. atoms | 2869 |
| protein | 2323 |
| ligand/ion | 98 |
| water | 448 |
| Average <i>B</i> -factors | 18.29 |
| protein | 15.36 |
| ligand/ion | 19.49 |
| water | 33.19 |
| R.m.s. deviations | |
| bond lengths (Å) | 0.006 |
| bond angles (°) | 0.90 |
| Ramachandran | |
| Favored (%) | 96.9 |
| allowed (%) | 3.1 |
| outliers (%) | 0.0 |

2 *Values in parentheses correspond to the highest-resolution shell.

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