Supplementary Table S1

Crystallographic statistics

Data collection	
Space group	P6 ₅
Unit cell parameters	<i>a</i> = <i>b</i> =98.2 Å, <i>c</i> =80.3 Å
Wavelength (Å)	0.9700
Resolution range (Å)	50 – 1.6
Redundancy	6.7
Unique reflections	386876
Completeness (%)	99.3 (97.6)
$I/\sigma(I)$	32.3 (2.8)
R_{sym}^{-1}	0.054 (0.424)
Refinement	
Resolution range (Å)	32.1 – 1.60 (1.70-1.60)
No. of reflections	57522
<i>R</i> -factor/Free <i>R</i> -factor ²	0.187 / 0.206
RMS deviations	
Bond lengths (Å)	0.005
Bond angles (deg.)	1.2
Average B-value (Å ²)	25.6

All numbers in parentheses refer to the highest resolution shell statistics.

 $^{^{1}}R_{\text{sym}}=\Sigma |I_{\text{avg}}-Ii| / \Sigma Ii$, where Ii is the observed intensity and I_{avg} is the average intensity.

 $^{^2}$ Free *R*-factor is calculated for 10 % of randomly selected reflections excluded from refinement.