

Table 1 - Data collection and refinement statistics

	Anisotropic cutoff	Spherical cutoff
<b>Data Collection</b>		
Beamline	APS 24-IDE	
Wavelength (Å)	0.97915	
Number of crystals	2	
Space group	P2 <sub>1</sub> 2 <sub>1</sub> 2 <sub>1</sub>	
Cell dimensions		
a,b,c (Å)	72.45, 154.51, 202.85	
α=β=γ (°)	90	
Resolution (Å)	122.91-2.774	
shell where  I σ ~ 2 (Å)	3.141-2.774 ( I σ = 2.0)	3.755-3.571 ( I σ = 1.885)
R <sub>merge</sub>	0.213 (1.589)	0.452 (34.263)
R <sub>meas</sub>	0.221 (1.655)	0.469 (35.536)
R <sub>pim</sub>	0.059 (0.457)	0.124 (9.357)
CC <sub>1/2</sub>	(0.602)	(0.165)
I  / σI	9.4 (2.0)	4.3 (0.1)
Completeness (%)	93.2 (80.2)	44.6 (7.3)
Multiplicity	14.1 (13.0)	14.5 (14.3)
<b>Refinement</b>		
No. of reflections used†	24830	
R <sub>work</sub> (%)	24.8	
R <sub>free</sub> (%)	28.6	
No. of total atoms	10250	
No. of K <sup>+</sup> ions	6	
Average B factor (Å <sup>2</sup> ), all atoms	82.526	
Clashscore	7.3	
Molprobity score	1.77	
Ramachandran plot		
favored (%)	95.70	
allowed (%)	3.99	
disallowed (%)	0.31	
R.m.s. deviations		
Bond lengths (Å)	0.006	
Bond angles (°)	0.923	

\*Values in parentheses are for the highest resolution shell.

†5% of these reflections were used to calculate Rfree