

Table 1. Data collection and refinement statistics

<i>Data collection statistics:</i>							
<i>Dataset:</i>	<b>L28B2 (native)</b>	<b>L28C6 HgCl<sub>2</sub></b>	<b>L28D2 K<sub>2</sub>PtCl<sub>4</sub></b>	<b>L88A4 (NH<sub>4</sub>)<sub>2</sub>WS<sub>4</sub></b>	<b>L88C1 K<sub>2</sub>OsO<sub>4</sub></b>	<b>L88A2 EtOHg</b>	<b>L88B3_2 Ta<sub>6</sub>Br<sub>14</sub></b>
Temperature	RT (290K)						
Spacegroup	I4						
a=b=, Å	140.7	140.9	140.8	140.2	140.6	141.0	140.9
c=, Å	97.4	97.8	97.8	97.3	98.7	96.8	96.4
Resolution, Å (final shell)	48.8 – 3.5 (3.7 – 3.5)	48.8 – 3.4 (3.6 – 3.4)	40.5 – 3.5 (3.7 – 3.5)	42.3 – 2.5 (2.7 – 2.5)	40.5 – 3.0 (3.16 – 3.0)	28.7 – 3.4 (3.58 – 3.4)	48.2 – 2.8 (2.97 – 2.8)
Reflections unique (total)	12100 (45984)	13281 (49702)	12002 (35629)	30889 (114411)	18290 (36609)	12785 (30504)	22711 (85677)
Completeness (%) overall (final shell)	100.0 (100.0)	100.0 (100.0)	98.9 (99.8)	99.0 (93.7)	95.2 (97.6)	97.4 (98.8)	99.2 (95.3)
I/σ <sub>1</sub> overall (final shell)	2.8 (1.3)	4.1 (2.2)	3.3 (1.3)	7.9 (2.4)	2.9 (0.6)	4.5 (2.1)	2.6 (1.5)
<sup>a</sup> R <sub>merge</sub> overall	21 %	15.7 %	18.3 %	7.7 %	19.4 %	13.7 %	15.4 %
Number of heavy atom sites	-	4	10	4	3	12	1×6
<sup>b</sup> Phasing power (centric/acentric)	-	0.92/1.2	0.79/1.2	0.56/0.89	0.81/1.03	1.07/1.45	1.05/1.15
<sup>c</sup> FOM (MIR)	0.58	-	-	-	-	-	-
<i>Refinement statistics:</i>							
<i>Dataset:</i>	<b>L18B1 (native)</b>						
Temperature	90K	Number of atoms			6445		
Spacegroup	I4	Number of solvent molecules			468		
a=b=, Å	138.9	Number of bounded buffer molecule atoms			66		
c=, Å	94.1						
Resolution, Å (final shell)	28.0 – 1.9 (2.0 – 1.9)	Test set size			10% random		
Reflections unique (total)	70374 (268268)	<sup>d</sup> R <sub>cryst</sub> (R <sub>free</sub> )			0.187 (0.218)		
Completeness (%) overall (final shell)	99.9 (100.0)	RMS bonds/angles			0.017 Å/1.6°		
I/σ <sub>1</sub> overall (final shell)	10.4 (2.7)	Average B-factors (Å <sup>2</sup> )					
		main chain			25.2		
		side chains			26.6		
		solvent			35.5		
		cofactors			56.7		
<sup>a</sup> R <sub>merge</sub> overall	4.9 %						

<sup>a</sup>  $R_{merge} = \frac{\sum_{\mathbf{h}} \sum_{i=1}^{n_{\mathbf{h}}} | \langle I_{\mathbf{h}} \rangle - I_{hi} |}{\sum_{\mathbf{h}} \sum_{i=1}^{n_{\mathbf{h}}} | I_{hi} |}$ , where  $I_{hi}$  is an intensity value of  $i$ -th measurement of reflection  $\mathbf{h}$ ,  $\mathbf{h} = (h, k, l)$ , sum  $\sum_{\mathbf{h}}$  runs over all measured reflections, and  $\langle I_{\mathbf{h}} \rangle$  is an average measured intensity of the reflection  $\mathbf{h}$ . Number  $n_{\mathbf{h}}$

is a number of measurements of reflection  $h$ . Data were processed with MOSFLM [Leslie 2003] and further processed SCALA [Evans 1997] and TRUNCATE [French 1978] from CCP4 [CCP4 1994] package.

<sup>b</sup> Phasing power =  $\langle |F_h^{obs}| \rangle / r.m.s.d. \varepsilon$ , where  $\varepsilon$  is lack of closure.

<sup>c</sup> FOM – Figure of merit

<sup>d</sup>  $R_{crys, free} = \sum_h |F_h^{obs} - F_h^{calc}| / \sum_h |F_h^{obs}|$ , where  $F_h^{obs}$  and  $F_h^{calc}$  are observed and calculated structure factors, respectively.