

**Structural and dynamic insight into substrate binding and catalysis of Human Lipocalin  
Prostaglandin D Synthase  
Lim et al**

**Table S1:** Peak count for NMR amino acid assignment

<b>Total no. of residues</b>	176			
<b>Not observable residues</b>	8 (Pro) ; 6 (His)			
<b>Residues with two amide</b>	4 (Asn) ; 12 (Gln) ; 8 (Arg)			
<b>Observable residues</b>	210			
<b>Peak count for 2D <sup>15</sup>N-HSQC spectra</b>				
	Yellow fraction	Colourless fraction	Middle (colourless) fraction	Middle fraction added U46619
<b>Total peaks observed</b>	113 (53.8)	200 (95)	154 (73.3)	196 (93.6)
<b>Assignment</b>				
<b>Total residues cross peaks can be observed</b>	162			
<b>Total cross peaks observed in TROSY-HSQC</b>	145 (89.5)			
<b>Total assigned peaks</b>	121 (74.7)			

\*value in parentheses indicate percentage