

Table 1. Statistics on the surfaces of apo structures within the canonical set of proteins

PDB ID	% Surf (SC res)	% Surf (LB res)	SC-LB overlap	# SC sites	# LB sites	# Overlapping sites	% LB sites identified
3pfk	51.0	20.4	0.255 (0.155)	19	3	3	100.0
4ake	45.4	17.8	0.274 (0.154)	29	2	2	100.0
1cd5	58.9	10.0	0.153 (0.096)	24	2	1	50.0
1j3h	6.6	8.0	0.25 (0.041)	2	1	1	100.0
1bks	34.3	9.7	0.079 (0.079)	24	4	1	25.0
1e5x	20.7	9.3	0.139 (0.077)	17	3	2	66.7
1efk	5.5	8.6	0.03 (0.036)	10	10	0	0.0
1nr7	14.9	17.5	0.187 (0.102)	45	24	6	25.0
1xtt	29.8	19.6	0.295 (0.154)	31	5	5	100.0
2hnp	73.9	13.3	0.16 (0.134)	25	2	2	100.0
3d7s	26.7	13.7	0.054 (0.064)	26	9	0	0.0
3ju5	1.6	3.9	0 (0.013)	1	2	0	0.0
mean	30.8	12.7	0.156 (0.092)	21.083	5.583	1.917	55.6