	1
104% shair D MOGHEDI adh	- Camp I
1Q4K_chain_D_MQSpTPL.pdb,	
<pre>1UMW_chain_E_PMQSpTPL.pdb,</pre>	P
20JX_chain_E_LLCSTPNGL.pdb,	L_C <mark>ST</mark> PN
<pre>3BZI_chain_E_LLCSpTPNG.pdb,</pre>	LLLLLLLL
<pre>3CL5_chain_B_PPHSpT.pdb,</pre>	PPPPPP
<pre>3FVH_chain_B_LHSpTA.pdb,</pre>	X
<pre>3HIK_chain_B_PLHSpT.pdb,</pre>	
3P2Z_chain_B_PLHSpTA.pdb,	
3P34_chain_B_MQSpTPL.pdb,	X
3P35_chain_E_MQSpSPL.pdb,	X
<pre>3P36_chain_B_DPPLHSpTA.pdb,</pre>	XDL.H <mark>ST</mark> A
<pre>3P37_chain_D_FDPPLHSpTA.pdb,</pre>	L.H.STAL.H.STA
<pre>3Q1L_chain_E_FMPPPMSpSM.pdb,</pre>	XFMPPP
4E67_chain_B_PLHSpTA.pdb,	
4E9C_chain_B_LDPPLHSpTA.pdb,	
4HAB_chain_D_PLHSpTM.pdb,	
4HY2_chain_D_LHSpTM.pdb,	L.H.S.T.MXL.H.S.T.MX
4JB7_chain_C_LDDLVAESPRKEFARINMDG	MGHHHHHLDDLV.AESPRK
4LKM chain D PLHSpTM.pdb,	
5J19_chain_C_ESCFTNAAFSSpTPK.pdb,	. ESCFINAAAF.SSTPK
5NFU_chain_C_HSpTA.pdb,	XH <mark>ST</mark> A
6GY2 chain C ATPPTLSSpTVLI.pdb,	
	The state of the s

1Q4K_chain_D_MQSpTPL.pdb,	
<pre>1UMW_chain_E_PMQSpTPL.pdb,</pre>	
20JX_chain_E_LLCSTPNGL.pdb,	GL
3BZI_chain_E_LLCSpTPNG.pdb,	
<pre>3CL5_chain_B_PPHSpT.pdb,</pre>	
3FVH_chain_B_LHSpTA.pdb,	
3HIK_chain_B_PLHSpT.pdb,	
<pre>3P2Z_chain_B_PLHSpTA.pdb,</pre>	X
3P34_chain_B_MQSpTPL.pdb,	
<pre>3P35_chain_E_MQSpSPL.pdb,</pre>	X
3P36_chain_B_DPPLHSpTA.pdb,	X
<pre>3P37_chain_D_FDPPLHSpTA.pdb,</pre>	X
<pre>3Q1L_chain_E_FMPPPMSpSM.pdb,</pre>	
4E67_chain_B_PLHSpTA.pdb,	X.
4E9C_chain_B_LDPPLHSpTA.pdb,	. X
4HAB_chain_D_PLHSpTM.pdb,	XX
4HY2_chain_D_LHSpTM.pdb,	
4JB7_chain_C_LDDLVAESPRKEFARINMDG	EFARINMDGIAVPDEREFDIEADMRPHELEQESDTFGAG
4LKM_chain_D_PLHSpTM.pdb,	X
5J19_chain_C_ESCFTNAAFSSpTPK.pdb,	KK
5NFU_chain_C_HSpTA.pdb,	
<pre>6GY2_chain_C_ATPPTLSSpTVLI.pdb,</pre>	