

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

CLUSTAL 2.0.12 multiple sequence alignment

```
3EKJ_A|PDBID|CHAIN|SEQUENCE      MRGSHHHHHHGMSAMTGGQQMGRDLYDDDDKDLATMVDSSRRKWNKTGHA 50
3EVP_A|PDBID|CHAIN|SEQUENCE      -----
3EKH_A|PDBID|CHAIN|SEQUENCE      MRGSHHHHHHGMSAMTGGQQMGRDLYDDDDKDLATMVDSSRRKWNKTGHA 50
3EVR_A|PDBID|CHAIN|SEQUENCE      -----DSSRRKWNKTGHA 13
3EK7_A|PDBID|CHAIN|SEQUENCE      MRGSHHHHHHGMSAMTGGQQMGRDLYDDDDKDLATMVDSSRRKWNKTGHA 50
3EVL_A|PDBID|CHAIN|SEQUENCE      MRGSHHHHHHGMSAMTGGQQMGRDLYDDDDKDLATMVDSSRRKWNKTGHA 50
3EVV_A|PDBID|CHAIN|SEQUENCE      MRGSHHHHHHGMSAMTGGQQMGRDLYDDDDKDLATMVDSSRRKWNKTGHA 50
3EK4_A|PDBID|CHAIN|SEQUENCE      MRGSHHHHHHGMSAMTGGQQMGRDLYDDDDKDLATMVDSSRRKWNKTGHA 50
3EK8_A|PDBID|CHAIN|SEQUENCE      MRGSHHHHHHGMSAMTGGQQMGRDLYDDDDKDLATMVDSSRRKWNKTGHA 50
2G3D_A|PDBID|CHAIN|SEQUENCE      -----
2G2S_A|PDBID|CHAIN|SEQUENCE      -----
1QYQ_A|PDBID|CHAIN|SEQUENCE      -----
2G6E_A|PDBID|CHAIN|SEQUENCE      -----
1YHH_A|PDBID|CHAIN|SEQUENCE      -----
1YHG_B|PDBID|CHAIN|SEQUENCE      -----
2G5Z_A|PDBID|CHAIN|SEQUENCE      -----
2G16_A|PDBID|CHAIN|SEQUENCE      -----
2HG0_A|PDBID|CHAIN|SEQUENCE      -----
2HGY_A|PDBID|CHAIN|SEQUENCE      -----
1YJF_A|PDBID|CHAIN|SEQUENCE      -----
1YHI_A|PDBID|CHAIN|SEQUENCE      -----
1YJ2_A|PDBID|CHAIN|SEQUENCE      -----
1QYF_A|PDBID|CHAIN|SEQUENCE      -----
1QXT_A|PDBID|CHAIN|SEQUENCE      -----
2A1M_A|PDBID|CHAIN|SEQUENCE      -----
2A1K_A|PDBID|CHAIN|SEQUENCE      -----
2A1J_A|PDBID|CHAIN|SEQUENCE      -----
2A1L_A|PDBID|CHAIN|SEQUENCE      -----
2HFC_A|PDBID|CHAIN|SEQUENCE      -----
2HCG_A|PDBID|CHAIN|SEQUENCE      -----
2Q20_A|PDBID|CHAIN|SEQUENCE      -----
2FZU_A|PDBID|CHAIN|SEQUENCE      -----
1KYR_A|PDBID|CHAIN|SEQUENCE      -----
1KYP_A|PDBID|CHAIN|SEQUENCE      -----
1KYS_A|PDBID|CHAIN|SEQUENCE      -----
2FWQ_A|PDBID|CHAIN|SEQUENCE      -----
1B9C_D|PDBID|CHAIN|SEQUENCE      -----
1MYW_A|PDBID|CHAIN|SEQUENCE      -----
3E08_A|PDBID|CHAIN|SEQUENCE      -----
1QY0_A|PDBID|CHAIN|SEQUENCE      -----
2Q11_A|PDBID|CHAIN|SEQUENCE      -----
1QY3_A|PDBID|CHAIN|SEQUENCE      -----
2QT2_A|PDBID|CHAIN|SEQUENCE      -----
2QRF_A|PDBID|CHAIN|SEQUENCE      -----
2HRS_A|PDBID|CHAIN|SEQUENCE      -----
2B3Q_D|PDBID|CHAIN|SEQUENCE      -----
2B3P_A|PDBID|CHAIN|SEQUENCE      -----
1KPS_A|PDBID|CHAIN|SEQUENCE      -----
3G11_A|PDBID|CHAIN|SEQUENCE      -----
3GJ2_A|PDBID|CHAIN|SEQUENCE      -----
3G9A_A|PDBID|CHAIN|SEQUENCE      -----
3K1K_B|PDBID|CHAIN|SEQUENCE      -----
1J00_A|PDBID|CHAIN|SEQUENCE      -----
1J11_A|PDBID|CHAIN|SEQUENCE      -----
2A8B_B|PDBID|CHAIN|SEQUENCE      -----
2A8A_A|PDBID|CHAIN|SEQUENCE      -----
3CD1_A|PDBID|CHAIN|SEQUENCE      -----
3CD9_A|PDBID|CHAIN|SEQUENCE      -----
3CB9_A|PDBID|CHAIN|SEQUENCE      -----
3CBE_A|PDBID|CHAIN|SEQUENCE      -----
20KW_F|PDBID|CHAIN|SEQUENCE      -----
20KY_B|PDBID|CHAIN|SEQUENCE      -----
2HQZ_A|PDBID|CHAIN|SEQUENCE      -----
2HJQ_A|PDBID|CHAIN|SEQUENCE      -----
202B_A|PDBID|CHAIN|SEQUENCE      -----
2H9W_A|PDBID|CHAIN|SEQUENCE      -----
2024_A|PDBID|CHAIN|SEQUENCE      -----
2029_A|PDBID|CHAIN|SEQUENCE      -----
2H6V_A|PDBID|CHAIN|SEQUENCE      -----
1RM0_A|PDBID|CHAIN|SEQUENCE      -----
1RM1_A|PDBID|CHAIN|SEQUENCE      -----
1RMP_A|PDBID|CHAIN|SEQUENCE      -----
1RRX_A|PDBID|CHAIN|SEQUENCE      -----
2Q6P_A|PDBID|CHAIN|SEQUENCE      -----
1Z1Q_A|PDBID|CHAIN|SEQUENCE      -----
156Z_A|PDBID|CHAIN|SEQUENCE      -----
1Z1P_A|PDBID|CHAIN|SEQUENCE      -----
1F06_A|PDBID|CHAIN|SEQUENCE      -----
1F09_A|PDBID|CHAIN|SEQUENCE      -----
2YFF_A|PDBID|CHAIN|SEQUENCE      -----
3D08_A|PDBID|CHAIN|SEQUENCE      -----
3D01_A|PDBID|CHAIN|SEQUENCE      -----
3D09_A|PDBID|CHAIN|SEQUENCE      -----
3D0A_A|PDBID|CHAIN|SEQUENCE      -----
3D0C_A|PDBID|CHAIN|SEQUENCE      -----
3D0D_A|PDBID|CHAIN|SEQUENCE      -----
3D0E_A|PDBID|CHAIN|SEQUENCE      -----
3D0F_A|PDBID|CHAIN|SEQUENCE      -----
3D0H_A|PDBID|CHAIN|SEQUENCE      -----
3D0I_A|PDBID|CHAIN|SEQUENCE      -----
3D0J_A|PDBID|CHAIN|SEQUENCE      -----
3D0K_A|PDBID|CHAIN|SEQUENCE      -----
3D0L_A|PDBID|CHAIN|SEQUENCE      -----
3D0M_A|PDBID|CHAIN|SEQUENCE      -----
3D0N_A|PDBID|CHAIN|SEQUENCE      -----
3D0O_A|PDBID|CHAIN|SEQUENCE      -----
3D0Q_A|PDBID|CHAIN|SEQUENCE      -----
3D0R_A|PDBID|CHAIN|SEQUENCE      -----
3D0S_A|PDBID|CHAIN|SEQUENCE      -----
3D06_A|PDBID|CHAIN|SEQUENCE      -----
3D07_A|PDBID|CHAIN|SEQUENCE      -----
3DPW_A|PDBID|CHAIN|SEQUENCE      -----
```

3DPX_A | PDBID | CHAIN | SEQUENCE -----
3DPZ_A | PDBID | CHAIN | SEQUENCE -----
3DQ2_A | PDBID | CHAIN | SEQUENCE -----
3DQ3_A | PDBID | CHAIN | SEQUENCE -----
1HUJ_A | PDBID | CHAIN | SEQUENCE -----
1YFP_A | PDBID | CHAIN | SEQUENCE -----
2JAD_A | PDBID | CHAIN | SEQUENCE -----
1H6R_A | PDBID | CHAIN | SEQUENCE -----
1C4F_A | PDBID | CHAIN | SEQUENCE -----
1EMB_A | PDBID | CHAIN | SEQUENCE -----
1Q4A_A | PDBID | CHAIN | SEQUENCE -----
1EMA_A | PDBID | CHAIN | SEQUENCE -----
1Q4B_A | PDBID | CHAIN | SEQUENCE -----
1EMG_A | PDBID | CHAIN | SEQUENCE -----
2DUF_A | PDBID | CHAIN | SEQUENCE -----
2DUJ_A | PDBID | CHAIN | SEQUENCE -----
2DUI_A | PDBID | CHAIN | SEQUENCE -----
2DUG_A | PDBID | CHAIN | SEQUENCE -----
2DUH_A | PDBID | CHAIN | SEQUENCE -----
1Q4E_A | PDBID | CHAIN | SEQUENCE -----
1Q73_A | PDBID | CHAIN | SEQUENCE -----
1Q4D_A | PDBID | CHAIN | SEQUENCE -----
1Q4C_A | PDBID | CHAIN | SEQUENCE -----
1JBZ_A | PDBID | CHAIN | SEQUENCE -----
1JBY_A | PDBID | CHAIN | SEQUENCE -----
1W75_D | PDBID | CHAIN | SEQUENCE -----
1W7U_D | PDBID | CHAIN | SEQUENCE -----
1W7T_D | PDBID | CHAIN | SEQUENCE -----
1HCJ_A | PDBID | CHAIN | SEQUENCE -----
2QLE_A | PDBID | CHAIN | SEQUENCE -----
1BFF_A | PDBID | CHAIN | SEQUENCE -----
2EMO_A | PDBID | CHAIN | SEQUENCE -----
1EMF_A | PDBID | CHAIN | SEQUENCE -----
2EMD_A | PDBID | CHAIN | SEQUENCE -----
2EMN_A | PDBID | CHAIN | SEQUENCE -----
1EMM_A | PDBID | CHAIN | SEQUENCE -----
1EMC_A | PDBID | CHAIN | SEQUENCE -----
1EML_A | PDBID | CHAIN | SEQUENCE -----
1EME_A | PDBID | CHAIN | SEQUENCE -----
1EMK_A | PDBID | CHAIN | SEQUENCE -----
2WUR_A | PDBID | CHAIN | SEQUENCE -----
1GFL_A | PDBID | CHAIN | SEQUENCE -----
1RM9_A | PDBID | CHAIN | SEQUENCE -----
2WSN_A | PDBID | CHAIN | SEQUENCE -----
2Q57_A | PDBID | CHAIN | SEQUENCE -----
2WSO_A | PDBID | CHAIN | SEQUENCE -----
1OXF_A | PDBID | CHAIN | SEQUENCE -----
1CV7_A | PDBID | CHAIN | SEQUENCE -----
3GEX_A | PDBID | CHAIN | SEQUENCE -----
1OXD_A | PDBID | CHAIN | SEQUENCE -----
1OXE_A | PDBID | CHAIN | SEQUENCE -----
3LVC_A | PDBID | CHAIN | SEQUENCE -----
3LVD_A | PDBID | CHAIN | SEQUENCE -----
3LVA_A | PDBID | CHAIN | SEQUENCE -----
2HPW_A | PDBID | CHAIN | SEQUENCE -----
2WIQ_A | PDBID | CHAIN | SEQUENCE -----
3A8S_A | PDBID | CHAIN | SEQUENCE -----
2WIS_A | PDBID | CHAIN | SEQUENCE -----
3GB3_A | PDBID | CHAIN | SEQUENCE -----
3GL4_A | PDBID | CHAIN | SEQUENCE -----
2A46_A | PDBID | CHAIN | SEQUENCE -----
2A48_A | PDBID | CHAIN | SEQUENCE -----
2A47_A | PDBID | CHAIN | SEQUENCE -----
2ZMW_A | PDBID | CHAIN | SEQUENCE -----
2ZMJ_A | PDBID | CHAIN | SEQUENCE -----
2ZOG_A | PDBID | CHAIN | SEQUENCE -----
2ZOT_A | PDBID | CHAIN | SEQUENCE -----
3BX9_B | PDBID | CHAIN | SEQUENCE -----
3BXA_A | PDBID | CHAIN | SEQUENCE -----
3BXC_A | PDBID | CHAIN | SEQUENCE -----
3BXC_A | PDBID | CHAIN | SEQUENCE -----
3H10_A | PDBID | CHAIN | SEQUENCE -----
3H1R_A | PDBID | CHAIN | SEQUENCE -----
3M22_A | PDBID | CHAIN | SEQUENCE -----
3M24_A | PDBID | CHAIN | SEQUENCE -----
3IP2_A | PDBID | CHAIN | SEQUENCE -----
3EST_A | PDBID | CHAIN | SEQUENCE -----
3E5V_A | PDBID | CHAIN | SEQUENCE -----
1UIS_B | PDBID | CHAIN | SEQUENCE -----
3E5W_A | PDBID | CHAIN | SEQUENCE -----
2A50_A | PDBID | CHAIN | SEQUENCE -----
2A52_A | PDBID | CHAIN | SEQUENCE -----
1XMZ_A | PDBID | CHAIN | SEQUENCE -----
2A54_C | PDBID | CHAIN | SEQUENCE -----
2A53_A | PDBID | CHAIN | SEQUENCE -----
2A56_A | PDBID | CHAIN | SEQUENCE -----
1XQM_A | PDBID | CHAIN | SEQUENCE -----
3CFE_L | PDBID | CHAIN | SEQUENCE -----
3CFH_S | PDBID | CHAIN | SEQUENCE -----
3CFA_L | PDBID | CHAIN | SEQUENCE -----
1YZW_A | PDBID | CHAIN | SEQUENCE -----
2IB5_A | PDBID | CHAIN | SEQUENCE -----
2IB6_A | PDBID | CHAIN | SEQUENCE -----
2C9T_A | PDBID | CHAIN | SEQUENCE -----
1G7K_A | PDBID | CHAIN | SEQUENCE -----
1ZGO_A | PDBID | CHAIN | SEQUENCE -----
1GGX_A | PDBID | CHAIN | SEQUENCE -----
1ZGQ_H | PDBID | CHAIN | SEQUENCE -----
1ZGP_A | PDBID | CHAIN | SEQUENCE -----
2H8Q_A | PDBID | CHAIN | SEQUENCE -----
2VAE_A | PDBID | CHAIN | SEQUENCE -----
2V4E_A | PDBID | CHAIN | SEQUENCE -----
2VAD_A | PDBID | CHAIN | SEQUENCE -----
2H5P_A | PDBID | CHAIN | SEQUENCE -----
2H5R_A | PDBID | CHAIN | SEQUENCE -----

2HRS_A | PDBID | CHAIN | SEQUENCE -----
2B3Q_D | PDBID | CHAIN | SEQUENCE -----
2B3P_A | PDBID | CHAIN | SEQUENCE -----
1KPS_A | PDBID | CHAIN | SEQUENCE -----
3GJ1_A | PDBID | CHAIN | SEQUENCE -----
3GJ2_A | PDBID | CHAIN | SEQUENCE -----
3G9A_A | PDBID | CHAIN | SEQUENCE -----
3K1K_B | PDBID | CHAIN | SEQUENCE -----
1JCO_A | PDBID | CHAIN | SEQUENCE -----
1JCL_A | PDBID | CHAIN | SEQUENCE -----
2A8B_B | PDBID | CHAIN | SEQUENCE -----
2AHA_A | PDBID | CHAIN | SEQUENCE -----
3CD1_A | PDBID | CHAIN | SEQUENCE -----
3CD9_A | PDBID | CHAIN | SEQUENCE -----
3CB9_A | PDBID | CHAIN | SEQUENCE -----
3CBE_A | PDBID | CHAIN | SEQUENCE -----
2OKW_F | PDBID | CHAIN | SEQUENCE -----
2OKY_B | PDBID | CHAIN | SEQUENCE -----
2HQZ_A | PDBID | CHAIN | SEQUENCE -----
2HJO_A | PDBID | CHAIN | SEQUENCE -----
2Q2B_A | PDBID | CHAIN | SEQUENCE -----
2H9W_A | PDBID | CHAIN | SEQUENCE -----
2O24_A | PDBID | CHAIN | SEQUENCE -----
2O29_A | PDBID | CHAIN | SEQUENCE -----
2H6V_A | PDBID | CHAIN | SEQUENCE -----
1RMQ_A | PDBID | CHAIN | SEQUENCE -----
1RMM_A | PDBID | CHAIN | SEQUENCE -----
1RMP_A | PDBID | CHAIN | SEQUENCE -----
1RRX_A | PDBID | CHAIN | SEQUENCE -----
2Q6P_A | PDBID | CHAIN | SEQUENCE -----
1Z1Q_A | PDBID | CHAIN | SEQUENCE -----
156Z_A | PDBID | CHAIN | SEQUENCE -----
1Z1P_A | PDBID | CHAIN | SEQUENCE -----
1F0B_A | PDBID | CHAIN | SEQUENCE -----
1F09_A | PDBID | CHAIN | SEQUENCE -----
2YFP_A | PDBID | CHAIN | SEQUENCE -----
3DQ8_A | PDBID | CHAIN | SEQUENCE -----
3DQ1_A | PDBID | CHAIN | SEQUENCE -----
3DQ9_A | PDBID | CHAIN | SEQUENCE -----
3DQA_A | PDBID | CHAIN | SEQUENCE -----
3DQC_A | PDBID | CHAIN | SEQUENCE -----
3DQD_A | PDBID | CHAIN | SEQUENCE -----
3DQE_A | PDBID | CHAIN | SEQUENCE -----
3DQF_A | PDBID | CHAIN | SEQUENCE -----
3DQH_A | PDBID | CHAIN | SEQUENCE -----
3DQI_A | PDBID | CHAIN | SEQUENCE -----
3DQJ_A | PDBID | CHAIN | SEQUENCE -----
3DQK_A | PDBID | CHAIN | SEQUENCE -----
3DQL_A | PDBID | CHAIN | SEQUENCE -----
3DQM_A | PDBID | CHAIN | SEQUENCE -----
3DQN_A | PDBID | CHAIN | SEQUENCE -----
3DQO_A | PDBID | CHAIN | SEQUENCE -----
3DQU_A | PDBID | CHAIN | SEQUENCE -----
3DQA_A | PDBID | CHAIN | SEQUENCE -----
3DQ5_A | PDBID | CHAIN | SEQUENCE -----
3DQ6_A | PDBID | CHAIN | SEQUENCE -----
3DQ7_A | PDBID | CHAIN | SEQUENCE -----
3DPW_A | PDBID | CHAIN | SEQUENCE -----
3DPX_A | PDBID | CHAIN | SEQUENCE -----
3DPZ_A | PDBID | CHAIN | SEQUENCE -----
3DQ2_A | PDBID | CHAIN | SEQUENCE -----
3DQ3_A | PDBID | CHAIN | SEQUENCE -----
1HUY_A | PDBID | CHAIN | SEQUENCE -----
1YFP_A | PDBID | CHAIN | SEQUENCE -----
2JAD_A | PDBID | CHAIN | SEQUENCE -----
1H6R_A | PDBID | CHAIN | SEQUENCE -----
1C4F_A | PDBID | CHAIN | SEQUENCE -----
1EMB_A | PDBID | CHAIN | SEQUENCE -----
1Q4A_A | PDBID | CHAIN | SEQUENCE -----
1EMA_A | PDBID | CHAIN | SEQUENCE -----
1Q4B_A | PDBID | CHAIN | SEQUENCE -----
1EMG_A | PDBID | CHAIN | SEQUENCE -----
2DUF_A | PDBID | CHAIN | SEQUENCE -----
2DUJ_A | PDBID | CHAIN | SEQUENCE -----
2DUT_A | PDBID | CHAIN | SEQUENCE -----
2DUG_A | PDBID | CHAIN | SEQUENCE -----
2DUH_A | PDBID | CHAIN | SEQUENCE -----
1Q4E_A | PDBID | CHAIN | SEQUENCE -----
1Q73_A | PDBID | CHAIN | SEQUENCE -----
1Q4D_A | PDBID | CHAIN | SEQUENCE -----
1Q4C_A | PDBID | CHAIN | SEQUENCE -----
1J8Z_A | PDBID | CHAIN | SEQUENCE -----
1J8Y_A | PDBID | CHAIN | SEQUENCE -----
1W75_D | PDBID | CHAIN | SEQUENCE -----
1W7U_D | PDBID | CHAIN | SEQUENCE -----
1W7T_D | PDBID | CHAIN | SEQUENCE -----
1HCJ_A | PDBID | CHAIN | SEQUENCE -----
2QLE_A | PDBID | CHAIN | SEQUENCE -----
1BFP_A | PDBID | CHAIN | SEQUENCE -----
2EMO_A | PDBID | CHAIN | SEQUENCE -----
1EMF_A | PDBID | CHAIN | SEQUENCE -----
2EMD_A | PDBID | CHAIN | SEQUENCE -----
2EMN_A | PDBID | CHAIN | SEQUENCE -----
1EMM_A | PDBID | CHAIN | SEQUENCE -----
1EMC_A | PDBID | CHAIN | SEQUENCE -----
1EML_A | PDBID | CHAIN | SEQUENCE -----
1EME_A | PDBID | CHAIN | SEQUENCE -----
1EMK_A | PDBID | CHAIN | SEQUENCE -----
2WUR_A | PDBID | CHAIN | SEQUENCE -----
1GFL_A | PDBID | CHAIN | SEQUENCE -----
1RM9_A | PDBID | CHAIN | SEQUENCE -----
2WSN_A | PDBID | CHAIN | SEQUENCE -----
2Q57_A | PDBID | CHAIN | SEQUENCE -----
2WSO_A | PDBID | CHAIN | SEQUENCE -----
1OXF_A | PDBID | CHAIN | SEQUENCE -----

1CV7_A | PDBID | CHAIN | SEQUENCE -----
3GEX_A | PDBID | CHAIN | SEQUENCE -----
10XD_A | PDBID | CHAIN | SEQUENCE -----
10XE_A | PDBID | CHAIN | SEQUENCE -----
3LVC_A | PDBID | CHAIN | SEQUENCE -----
3LVD_A | PDBID | CHAIN | SEQUENCE -----
3LVA_A | PDBID | CHAIN | SEQUENCE -----
2HPW_A | PDBID | CHAIN | SEQUENCE -----
2WIQ_A | PDBID | CHAIN | SEQUENCE -----
3A8S_A | PDBID | CHAIN | SEQUENCE -----
2WIS_A | PDBID | CHAIN | SEQUENCE -----
3GB3_A | PDBID | CHAIN | SEQUENCE -----
3GL4_A | PDBID | CHAIN | SEQUENCE -----
2A46_A | PDBID | CHAIN | SEQUENCE -----
2A48_A | PDBID | CHAIN | SEQUENCE -----
2A47_A | PDBID | CHAIN | SEQUENCE -----
2ZM0_A | PDBID | CHAIN | SEQUENCE -----
2ZML_A | PDBID | CHAIN | SEQUENCE -----
2Z06_A | PDBID | CHAIN | SEQUENCE -----
2Z07_A | PDBID | CHAIN | SEQUENCE -----
3BX9_B | PDBID | CHAIN | SEQUENCE -----
3BXA_A | PDBID | CHAIN | SEQUENCE -----
3BXB_A | PDBID | CHAIN | SEQUENCE -----
3BXC_A | PDBID | CHAIN | SEQUENCE -----
3H10_A | PDBID | CHAIN | SEQUENCE -----
3H1R_A | PDBID | CHAIN | SEQUENCE -----
3M22_A | PDBID | CHAIN | SEQUENCE -----
3M24_A | PDBID | CHAIN | SEQUENCE -----
3IP2_A | PDBID | CHAIN | SEQUENCE -----
3EST_A | PDBID | CHAIN | SEQUENCE -----
3E5V_A | PDBID | CHAIN | SEQUENCE -----
1UIS_B | PDBID | CHAIN | SEQUENCE -----
3E5W_A | PDBID | CHAIN | SEQUENCE -----
2A50_A | PDBID | CHAIN | SEQUENCE -----
2A52_A | PDBID | CHAIN | SEQUENCE -----
1XM2_A | PDBID | CHAIN | SEQUENCE -----
2A54_C | PDBID | CHAIN | SEQUENCE -----
2A53_A | PDBID | CHAIN | SEQUENCE -----
2A56_A | PDBID | CHAIN | SEQUENCE -----
1XQM_A | PDBID | CHAIN | SEQUENCE -----
3CF_F | PDBID | CHAIN | SEQUENCE -----
3CFH_S | PDBID | CHAIN | SEQUENCE -----
3CFA_L | PDBID | CHAIN | SEQUENCE -----
1YZW_A | PDBID | CHAIN | SEQUENCE -----
2IB5_A | PDBID | CHAIN | SEQUENCE -----
2IB6_A | PDBID | CHAIN | SEQUENCE -----
2C9L_A | PDBID | CHAIN | SEQUENCE -----
1G7K_A | PDBID | CHAIN | SEQUENCE -----
1ZGO_A | PDBID | CHAIN | SEQUENCE -----
1GG_X | PDBID | CHAIN | SEQUENCE -----
1ZGQ_H | PDBID | CHAIN | SEQUENCE -----
1ZGP_A | PDBID | CHAIN | SEQUENCE -----
2H8Q_A | PDBID | CHAIN | SEQUENCE -----
2VAE_A | PDBID | CHAIN | SEQUENCE -----
2V4E_A | PDBID | CHAIN | SEQUENCE -----
2VAD_A | PDBID | CHAIN | SEQUENCE -----
2H5P_A | PDBID | CHAIN | SEQUENCE -----
2H5R_A | PDBID | CHAIN | SEQUENCE -----
2H50_A | PDBID | CHAIN | SEQUENCE -----
2QLG_A | PDBID | CHAIN | SEQUENCE -----
2QLL_A | PDBID | CHAIN | SEQUENCE -----
2QLH_A | PDBID | CHAIN | SEQUENCE -----
2H5Q_A | PDBID | CHAIN | SEQUENCE -----
3KCS_A | PDBID | CHAIN | SEQUENCE -----
3KCT_A | PDBID | CHAIN | SEQUENCE -----
3LF3_A | PDBID | CHAIN | SEQUENCE -----
3LF4_A | PDBID | CHAIN | SEQUENCE -----
2P4M_A | PDBID | CHAIN | SEQUENCE -----
2ARL_A | PDBID | CHAIN | SEQUENCE -----
1MOV_A | PDBID | CHAIN | SEQUENCE -----
1M0J_A | PDBID | CHAIN | SEQUENCE -----
2WHS_A | PDBID | CHAIN | SEQUENCE -----
2WHT_A | PDBID | CHAIN | SEQUENCE -----
2WHU_A | PDBID | CHAIN | SEQUENCE -----
3IR8_A | PDBID | CHAIN | SEQUENCE -----
3CGL_A | PDBID | CHAIN | SEQUENCE -----
2Z6Z_A | PDBID | CHAIN | SEQUENCE -----
2T0V_A | PDBID | CHAIN | SEQUENCE -----
2POX_A | PDBID | CHAIN | SEQUENCE -----
2IE2_A | PDBID | CHAIN | SEQUENCE -----
2Z10_A | PDBID | CHAIN | SEQUENCE -----
2Z6Y_A | PDBID | CHAIN | SEQUENCE -----
2GX2_A | PDBID | CHAIN | SEQUENCE -----
2GX0_B | PDBID | CHAIN | SEQUENCE -----
2Z6X_A | PDBID | CHAIN | SEQUENCE -----
3LS3_A | PDBID | CHAIN | SEQUENCE -----
3LSA_A | PDBID | CHAIN | SEQUENCE -----
2DDC_A | PDBID | CHAIN | SEQUENCE -----
2DD0_A | PDBID | CHAIN | SEQUENCE -----
1XSS_A | PDBID | CHAIN | SEQUENCE -----
2VVI_A | PDBID | CHAIN | SEQUENCE -----
2VVI_A | PDBID | CHAIN | SEQUENCE -----
2VVJ_A | PDBID | CHAIN | SEQUENCE -----
1ZUX_A | PDBID | CHAIN | SEQUENCE -----
2BTJ_A | PDBID | CHAIN | SEQUENCE -----
2GW3_A | PDBID | CHAIN | SEQUENCE -----
2GW4_A | PDBID | CHAIN | SEQUENCE -----
2OTE_A | PDBID | CHAIN | SEQUENCE -----
20TB_A | PDBID | CHAIN | SEQUENCE -----
2HQK_A | PDBID | CHAIN | SEQUENCE -----
2VZX_A | PDBID | CHAIN | SEQUENCE -----
3ADF_A | PDBID | CHAIN | SEQUENCE -----
2ICR_A | PDBID | CHAIN | SEQUENCE -----
2FL1_A | PDBID | CHAIN | SEQUENCE -----
2OGR_A | PDBID | CHAIN | SEQUENCE -----

1XAE_A | PDBID | CHAIN | SEQUENCE -----
1XA9_A | PDBID | CHAIN | SEQUENCE -----
2PX5_A | PDBID | CHAIN | SEQUENCE -----
2PXW_A | PDBID | CHAIN | SEQUENCE -----
2OJK_A | PDBID | CHAIN | SEQUENCE -----
2RH7_A | PDBID | CHAIN | SEQUENCE -----
2C9J_H | PDBID | CHAIN | SEQUENCE -----
2G6X_A | PDBID | CHAIN | SEQUENCE -----
2G6Y_A | PDBID | CHAIN | SEQUENCE -----
2G3O_A | PDBID | CHAIN | SEQUENCE -----
2DD7_A | PDBID | CHAIN | SEQUENCE -----
2DD9_D | PDBID | CHAIN | SEQUENCE -----

3EKJ_A | PDBID | CHAIN | SEQUENCE IGDGPVLLPDNHYLSTQSKLSKDPNEKRDHMVLLFVFTAAGITLGMDELY 150
3EVP_A | PDBID | CHAIN | SEQUENCE IGDGPVLLPDNHYLSTQSKLSKDPNEKRDHMVLLFVFTAAGITLGMDELY 93
3EKH_A | PDBID | CHAIN | SEQUENCE IGDGPVLLPDNHYLSTQSKLSKDPNEKRDHMVLLFVFTAAGITLGMDELY 150
3EVR_A | PDBID | CHAIN | SEQUENCE IGDGPVLLPDNHYLSTQSKLSKDPNEKRDHMVLLFVFTAAGITLGMDELY 113
3EK7_A | PDBID | CHAIN | SEQUENCE IGDGPVLLPDNHYLSTQSKLSKDPNEKRDHMVLLFVFTAAGITLGMDELY 150
3EVL_A | PDBID | CHAIN | SEQUENCE IGDGPVLLPDNHYLSTQSKLSKDPNEKRDHMVLLFVFTAAGITLGMDELY 150
3EVV_A | PDBID | CHAIN | SEQUENCE IGDGPVLLPDNHYLSTQSKLSKDPNEKRDHMVLLFVFTAAGITLGMDELY 150
3EK4_A | PDBID | CHAIN | SEQUENCE IGDGPVLLPDNHYLSTQSKLSKDPNEKRDHMVLLFVFTAAGITLGMDELY 150
3EK8_A | PDBID | CHAIN | SEQUENCE IGDGPVLLPDNHYLSTQSKLSKDPNEKRDHMVLLFVFTAAGITLGMDELY 150
2G3D_A | PDBID | CHAIN | SEQUENCE -----
2G2S_A | PDBID | CHAIN | SEQUENCE -----
1QYQ_A | PDBID | CHAIN | SEQUENCE -----
2G6E_A | PDBID | CHAIN | SEQUENCE -----
1YHL_A | PDBID | CHAIN | SEQUENCE -----
1YHG_B | PDBID | CHAIN | SEQUENCE -----
2G5Z_A | PDBID | CHAIN | SEQUENCE -----
2G16_A | PDBID | CHAIN | SEQUENCE -----
2HG0_A | PDBID | CHAIN | SEQUENCE -----
2HG_Y | PDBID | CHAIN | SEQUENCE -----
1YJF_A | PDBID | CHAIN | SEQUENCE -----
1YHL_A | PDBID | CHAIN | SEQUENCE -----
1YJZ_A | PDBID | CHAIN | SEQUENCE -----
1QYF_A | PDBID | CHAIN | SEQUENCE -----
1QXT_A | PDBID | CHAIN | SEQUENCE -----
2A1M_A | PDBID | CHAIN | SEQUENCE -----
2A1K_A | PDBID | CHAIN | SEQUENCE -----
2A1J_A | PDBID | CHAIN | SEQUENCE -----
2A1L_A | PDBID | CHAIN | SEQUENCE -----
2HFC_A | PDBID | CHAIN | SEQUENCE -----
2HCG_A | PDBID | CHAIN | SEQUENCE -----
2Q20_A | PDBID | CHAIN | SEQUENCE -----
2FZU_A | PDBID | CHAIN | SEQUENCE -----
1KYR_A | PDBID | CHAIN | SEQUENCE -----
1KYP_A | PDBID | CHAIN | SEQUENCE -----
1KYS_A | PDBID | CHAIN | SEQUENCE -----
2FWQ_A | PDBID | CHAIN | SEQUENCE -----
1B9C_D | PDBID | CHAIN | SEQUENCE -----
1MYW_A | PDBID | CHAIN | SEQUENCE -----
3ED8_A | PDBID | CHAIN | SEQUENCE -----
1QYQ_A | PDBID | CHAIN | SEQUENCE -----
2QU1_A | PDBID | CHAIN | SEQUENCE -----
1QY3_A | PDBID | CHAIN | SEQUENCE -----
2QT2_A | PDBID | CHAIN | SEQUENCE -----
2QRF_A | PDBID | CHAIN | SEQUENCE -----
2HRS_A | PDBID | CHAIN | SEQUENCE -----
2B3Q_D | PDBID | CHAIN | SEQUENCE -----
2B3P_A | PDBID | CHAIN | SEQUENCE -----
1KPS_A | PDBID | CHAIN | SEQUENCE -----TG 2
3G1L_A | PDBID | CHAIN | SEQUENCE -----
3G1Z_A | PDBID | CHAIN | SEQUENCE -----
3G9A_A | PDBID | CHAIN | SEQUENCE -----
3K1K_B | PDBID | CHAIN | SEQUENCE -----
1JCO_A | PDBID | CHAIN | SEQUENCE -----
1J1L_A | PDBID | CHAIN | SEQUENCE -----
2A88_B | PDBID | CHAIN | SEQUENCE -----
2AHA_A | PDBID | CHAIN | SEQUENCE -----
3CD1_A | PDBID | CHAIN | SEQUENCE -----MRG 3
3CD9_A | PDBID | CHAIN | SEQUENCE -----MRG 3
3CB9_A | PDBID | CHAIN | SEQUENCE -----MRG 3
3CBE_A | PDBID | CHAIN | SEQUENCE -----MRG 3
2OKW_F | PDBID | CHAIN | SEQUENCE -----
2OKY_B | PDBID | CHAIN | SEQUENCE -----
2HOZ_A | PDBID | CHAIN | SEQUENCE -----
2HJO_A | PDBID | CHAIN | SEQUENCE -----
2O2B_A | PDBID | CHAIN | SEQUENCE -----
2H9W_A | PDBID | CHAIN | SEQUENCE -----
2O24_A | PDBID | CHAIN | SEQUENCE -----
2O29_A | PDBID | CHAIN | SEQUENCE -----
2H6V_A | PDBID | CHAIN | SEQUENCE -----
1RMO_A | PDBID | CHAIN | SEQUENCE -----
1RML_A | PDBID | CHAIN | SEQUENCE -----
1RMP_A | PDBID | CHAIN | SEQUENCE -----
1RRX_A | PDBID | CHAIN | SEQUENCE -----
2O6P_A | PDBID | CHAIN | SEQUENCE -----
1Z1Q_A | PDBID | CHAIN | SEQUENCE -----
1S6Z_A | PDBID | CHAIN | SEQUENCE -----
1Z1P_A | PDBID | CHAIN | SEQUENCE -----
1F0B_A | PDBID | CHAIN | SEQUENCE -----
1F09_A | PDBID | CHAIN | SEQUENCE -----
2YFP_A | PDBID | CHAIN | SEQUENCE -----
3DQ8_A | PDBID | CHAIN | SEQUENCE -----
3DQ1_A | PDBID | CHAIN | SEQUENCE -----
3DQ9_A | PDBID | CHAIN | SEQUENCE -----
3DQA_A | PDBID | CHAIN | SEQUENCE -----
3DQC_A | PDBID | CHAIN | SEQUENCE -----
3DQD_A | PDBID | CHAIN | SEQUENCE -----
3DQE_A | PDBID | CHAIN | SEQUENCE -----
3DQF_A | PDBID | CHAIN | SEQUENCE -----
3DQH_A | PDBID | CHAIN | SEQUENCE -----
3DQI_A | PDBID | CHAIN | SEQUENCE -----
3DQJ_A | PDBID | CHAIN | SEQUENCE -----

3DQK_A | PDBID | CHAIN | SEQUENCE -----
3DQL_A | PDBID | CHAIN | SEQUENCE -----
3DQM_A | PDBID | CHAIN | SEQUENCE -----
3DQN_A | PDBID | CHAIN | SEQUENCE -----
3DQO_A | PDBID | CHAIN | SEQUENCE -----
3DQU_A | PDBID | CHAIN | SEQUENCE -----
3DQ4_A | PDBID | CHAIN | SEQUENCE -----
3DQ5_A | PDBID | CHAIN | SEQUENCE -----
3DQ6_A | PDBID | CHAIN | SEQUENCE -----
3DQ7_A | PDBID | CHAIN | SEQUENCE -----
3DPW_A | PDBID | CHAIN | SEQUENCE -----
3DPX_A | PDBID | CHAIN | SEQUENCE -----
3DPZ_A | PDBID | CHAIN | SEQUENCE -----
3DQ2_A | PDBID | CHAIN | SEQUENCE -----
3DQ3_A | PDBID | CHAIN | SEQUENCE -----
1HUY_A | PDBID | CHAIN | SEQUENCE -----
1YFP_A | PDBID | CHAIN | SEQUENCE -----
2JAD_A | PDBID | CHAIN | SEQUENCE -----
1H6R_A | PDBID | CHAIN | SEQUENCE -----
1C4F_A | PDBID | CHAIN | SEQUENCE -----
1EM8_A | PDBID | CHAIN | SEQUENCE -----
1Q4A_A | PDBID | CHAIN | SEQUENCE -----
1EMA_A | PDBID | CHAIN | SEQUENCE -----
1Q4B_A | PDBID | CHAIN | SEQUENCE -----
1EMG_A | PDBID | CHAIN | SEQUENCE -----
2DUF_A | PDBID | CHAIN | SEQUENCE -----
2DUJ_A | PDBID | CHAIN | SEQUENCE -----
2DUT_A | PDBID | CHAIN | SEQUENCE -----
2DUG_A | PDBID | CHAIN | SEQUENCE -----
2DUH_A | PDBID | CHAIN | SEQUENCE -----
1Q4E_A | PDBID | CHAIN | SEQUENCE -----
1Q73_A | PDBID | CHAIN | SEQUENCE -----
1Q4D_A | PDBID | CHAIN | SEQUENCE -----
1Q4C_A | PDBID | CHAIN | SEQUENCE -----
1J8Z_A | PDBID | CHAIN | SEQUENCE -----
1J8Y_A | PDBID | CHAIN | SEQUENCE -----
1W75_D | PDBID | CHAIN | SEQUENCE -----
1W7U_D | PDBID | CHAIN | SEQUENCE -----
1W7T_D | PDBID | CHAIN | SEQUENCE -----
1HCJ_A | PDBID | CHAIN | SEQUENCE -----
2QLE_A | PDBID | CHAIN | SEQUENCE -----
1BFP_A | PDBID | CHAIN | SEQUENCE -----
2EMO_A | PDBID | CHAIN | SEQUENCE -----
1EMF_A | PDBID | CHAIN | SEQUENCE -----
2EMD_A | PDBID | CHAIN | SEQUENCE -----
2EMN_A | PDBID | CHAIN | SEQUENCE -----
1EMM_A | PDBID | CHAIN | SEQUENCE -----
1EMC_A | PDBID | CHAIN | SEQUENCE -----
1EML_A | PDBID | CHAIN | SEQUENCE -----
1EME_A | PDBID | CHAIN | SEQUENCE -----
1EMK_A | PDBID | CHAIN | SEQUENCE -----
2WUR_A | PDBID | CHAIN | SEQUENCE -----
1GFL_A | PDBID | CHAIN | SEQUENCE -----
1RM9_A | PDBID | CHAIN | SEQUENCE -----
2WSN_A | PDBID | CHAIN | SEQUENCE -----
2Q57_A | PDBID | CHAIN | SEQUENCE -----
2WSO_A | PDBID | CHAIN | SEQUENCE -----
10XF_A | PDBID | CHAIN | SEQUENCE -----
1CV7_A | PDBID | CHAIN | SEQUENCE -----
3GEX_A | PDBID | CHAIN | SEQUENCE -----
10XD_A | PDBID | CHAIN | SEQUENCE -----
10XE_A | PDBID | CHAIN | SEQUENCE -----
3LVC_A | PDBID | CHAIN | SEQUENCE -----
3LVD_A | PDBID | CHAIN | SEQUENCE -----
3LVA_A | PDBID | CHAIN | SEQUENCE -----
2HPW_A | PDBID | CHAIN | SEQUENCE -----
2WIQ_A | PDBID | CHAIN | SEQUENCE -----
3A85_A | PDBID | CHAIN | SEQUENCE -----
2WIS_A | PDBID | CHAIN | SEQUENCE -----
3GB3_A | PDBID | CHAIN | SEQUENCE -----
3GL4_A | PDBID | CHAIN | SEQUENCE -----
2A46_A | PDBID | CHAIN | SEQUENCE -----
2A48_A | PDBID | CHAIN | SEQUENCE -----
2A47_A | PDBID | CHAIN | SEQUENCE -----
2ZMW_A | PDBID | CHAIN | SEQUENCE -----
2ZMU_A | PDBID | CHAIN | SEQUENCE -----
2Z06_A | PDBID | CHAIN | SEQUENCE -----
2Z07_A | PDBID | CHAIN | SEQUENCE -----
3BX9_B | PDBID | CHAIN | SEQUENCE -----
3BXA_A | PDBID | CHAIN | SEQUENCE -----
3BXB_A | PDBID | CHAIN | SEQUENCE -----
3BXC_A | PDBID | CHAIN | SEQUENCE -----
3H10_A | PDBID | CHAIN | SEQUENCE -----
3H1R_A | PDBID | CHAIN | SEQUENCE -----
3M22_A | PDBID | CHAIN | SEQUENCE -----
3M24_A | PDBID | CHAIN | SEQUENCE -----
3IP2_A | PDBID | CHAIN | SEQUENCE -----
3E5T_A | PDBID | CHAIN | SEQUENCE -----
3E5V_A | PDBID | CHAIN | SEQUENCE -----
1UI5_B | PDBID | CHAIN | SEQUENCE -----
3E5W_A | PDBID | CHAIN | SEQUENCE -----
2A50_A | PDBID | CHAIN | SEQUENCE -----
2A52_A | PDBID | CHAIN | SEQUENCE -----
1XMZ_A | PDBID | CHAIN | SEQUENCE -----
2A54_C | PDBID | CHAIN | SEQUENCE -----
2A53_A | PDBID | CHAIN | SEQUENCE -----
2A56_A | PDBID | CHAIN | SEQUENCE -----
1XQM_A | PDBID | CHAIN | SEQUENCE -----
3CFE_L | PDBID | CHAIN | SEQUENCE -----
3CFH_S | PDBID | CHAIN | SEQUENCE -----
3CFA_L | PDBID | CHAIN | SEQUENCE -----
1YZW_A | PDBID | CHAIN | SEQUENCE -----
2IB5_A | PDBID | CHAIN | SEQUENCE -----
2IB6_A | PDBID | CHAIN | SEQUENCE -----
2C9L_A | PDBID | CHAIN | SEQUENCE -----

-----MRGSHHHHHH 10

-----MGSSHHHHHSSGL 14

-----MGSSHHHHHSSGL 14

-----MRGSHHHHHHG 11

-----MRGSHHHHHHG 11

-----MRGSHHHHHHG 11

-----MRGSHHHHHHGMSMTGGQ 19

-----M 1

-----M 1

-----M 1

-----M 1

-----M 1

-----M 2

-----M 2

-----M 2

-----M 2

1EMC_A1|PDBID|CHAIN|SEQUENCE -----MASKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 43
1EML_A1|PDBID|CHAIN|SEQUENCE -----MASKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 43
1EME_A1|PDBID|CHAIN|SEQUENCE -----MASKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 43
1EMK_A1|PDBID|CHAIN|SEQUENCE -----MASKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 43
2WJR_A1|PDBID|CHAIN|SEQUENCE -----MSKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 42
1GFL_A1|PDBID|CHAIN|SEQUENCE -----ASKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 42
1RM9_A1|PDBID|CHAIN|SEQUENCE -----VSKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 42
2WSN_A1|PDBID|CHAIN|SEQUENCE -----MVSKEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 43
2Q57_A1|PDBID|CHAIN|SEQUENCE GLALPVATMSKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 60
2WS0_A1|PDBID|CHAIN|SEQUENCE -----MVSKEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 43
10XF_A1|PDBID|CHAIN|SEQUENCE -----SKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 41
1CV7_A1|PDBID|CHAIN|SEQUENCE -----MSKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 42
3GEX_A1|PDBID|CHAIN|SEQUENCE --LYFQGMVSKGEELFGGLVPIILVELDGDVNGHKFVSVSSEGEDATYGL 48
10XD_A1|PDBID|CHAIN|SEQUENCE -----MSKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 42
10XE_A1|PDBID|CHAIN|SEQUENCE -----MSKGEELFTGVVPIILVELDGDVNGHKFVSVSSEGEDATYGL 42
3LVC_A1|PDBID|CHAIN|SEQUENCE -----MSKGAELFTGLVPIILVELDGDVNGHKFVSVSSEGEDATYGL 42
3LVD_A1|PDBID|CHAIN|SEQUENCE -----MSKGAELFTGLVPIILVELDGDVNGHKFVSVSSEGEDATYGL 42
3LVA_A1|PDBID|CHAIN|SEQUENCE -----MSKGAELFTGLVPIILVELDGDVNGHKFVSVSSEGEDATYGL 42
2HPW_A1|PDBID|CHAIN|SEQUENCE ----MTALTEGAKLFEKEIPYITIELGDEVGMKFIKKEGTDATTGI 45
2WIQ_A1|PDBID|CHAIN|SEQUENCE VPRGSHMGSEGGPALFQSDMTFKIFIDGDEVNGQKFTIVADGSKKPFPHGF 64
348S_A1|PDBID|CHAIN|SEQUENCE ----HMGEPPALFQSDMTFKIFIDGDEVNGQKFTIVADGSKKPFPHGF 45
2WIS_A1|PDBID|CHAIN|SEQUENCE VPRGSHMGSEGGPALFQSDMTFKIFIDGDEVNGQKFTIVADGSKKPFPHGF 64
3GB3_A1|PDBID|CHAIN|SEQUENCE -----MEGGPALFQSDMTFKIFIDGDEVNGQKFTIVADGSKKPFPHGF 42
3GL4_A1|PDBID|CHAIN|SEQUENCE -----MEGGPALFQSDMTFKIFIDGDEVNGQKFTIVADGSKKPFPHGF 42
244E_A1|PDBID|CHAIN|SEQUENCE -----SALSNKFIGDDMKTYHMDGCVNGHYFTVKGEKPKPYEGT 53
2448_A1|PDBID|CHAIN|SEQUENCE -----SALSNKFIGDDMKTYHMDGCVNGHYFTVKGEKPKPYEGT 53
2447_A1|PDBID|CHAIN|SEQUENCE -----SALSNKFIGDDMKTYHMDGCVNGHYFTVKGEKPKPYEGT 53
22M_A1|PDBID|CHAIN|SEQUENCE -----MVSIVKPEKMKRYMDGCVNGHEFTIEGEGTRPYEGH 39
22ML_A1|PDBID|CHAIN|SEQUENCE -----MVSIVKPEKMKRYMDGCVNGHEFTIEGEGTRPYEGH 39
220E_A1|PDBID|CHAIN|SEQUENCE QMGRDLYDDDDKMSVIKPEKMKRYMDGCVNGHEFTIEGEGTRPYEGH 69
2207_A1|PDBID|CHAIN|SEQUENCE -----DPTMVSIVKPEKMKRYMDGCVNGHEFTIEGEGTRPYEGH 42
3BX9_B1|PDBID|CHAIN|SEQUENCE RGSHHHHHGSMSLELITENHMKLYMEGTVNNHFKCTSEGEKPKPYEGT 51
3BXA_A1|PDBID|CHAIN|SEQUENCE RGSHHHHHGSMSLELITENHMKLYMEGTVNNHFKCTSEGEKPKPYEGT 51
3BXB_A1|PDBID|CHAIN|SEQUENCE RGSHHHHHGSMSLELITENHMKLYMEGTVNNHFKCTSEGEKPKPYEGT 51
3BXC_A1|PDBID|CHAIN|SEQUENCE RGSHHHHHGSMSLELITENHMKLYMEGTVNNHFKCTSEGEKPKPYEGT 51
3H10_A1|PDBID|CHAIN|SEQUENCE -----MSELITENHMKLYMEGTVNNHFKCTSEGEKPKPYEGT 39
3H1R_A1|PDBID|CHAIN|SEQUENCE -----MSELITENHMKLYMEGTVNNHFKCTSEGEKPKPYEGT 39
3M2Z_A1|PDBID|CHAIN|SEQUENCE -----MVSKEELIKENHMKLYMEGTVNNHFKCTSEGEKPKPYEGT 43
3M24_A1|PDBID|CHAIN|SEQUENCE -----MSEELIKENHMKLYMEGTVNNHFKCTSEGEKPKPYEGT 40
3IPZ_A1|PDBID|CHAIN|SEQUENCE -----GVSKEELIKEDNHMKLYMEGTVNNHFKCTSEGEKPKPYEGT 43
3E5T_A1|PDBID|CHAIN|SEQUENCE GSHHHHHHGHMNSLKENMRMVMYMEGSVNGYQFKCTGEGDGNPYMG 52
3E5V_A1|PDBID|CHAIN|SEQUENCE GSHHHHHHGHMNSLKENMRMVMYMEGSVNGYQFKCTGEGDGNPYMG 52
1UIS_B1|PDBID|CHAIN|SEQUENCE -----IHMNSLKENMRMVMYMEGSVNGYQFKCTGEGDGNPYMG 41
3E5W_A1|PDBID|CHAIN|SEQUENCE GSHHHHHHGHMNSLKENMRMVMYMEGSVNGYQFKCTGEGDGNPYMG 52
2A50_A1|PDBID|CHAIN|SEQUENCE MRGSHHHHHHGSASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKPKPYEGT 50
2A52_A1|PDBID|CHAIN|SEQUENCE MRGSHHHHHHGSASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKPKPYEGT 50
1XMZ_A1|PDBID|CHAIN|SEQUENCE MRGSHHHHHHGSASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKPKPYEGT 50
2A54_C1|PDBID|CHAIN|SEQUENCE MRGSHHHHHHGSASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKPKPYEGT 50
2A53_A1|PDBID|CHAIN|SEQUENCE MRGSHHHHHHGSASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKPKPYEGT 50
2A56_A1|PDBID|CHAIN|SEQUENCE MRGSHHHHHHGSASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKPKPYEGT 50
1XQL_A1|PDBID|CHAIN|SEQUENCE -----MASLLETMPFKTTIEGTVNGHYFKCTGKGEKPKPYEGT 39
3CF_L1|PDBID|CHAIN|SEQUENCE -----MASLLETMPFRMTEGTVNGHYFKCTGKGEKPKPYEGT 39
3CFH_S1|PDBID|CHAIN|SEQUENCE -----MASLLETMPFRMTEGTVNGHYFKCTGKGEKPKPYEGT 39
3CFA_L1|PDBID|CHAIN|SEQUENCE -----MASLLETMPFRMTEGTVNGHYFKCTGKGEKPKPYEGT 39
1YW_A1|PDBID|CHAIN|SEQUENCE -----MAGLLKESMRKMYMEGTVNGHYFKCEGEGDGNPYMG 39
2IB5_A1|PDBID|CHAIN|SEQUENCE -----GSHMASKISDNVRIKLYMEGTVNNHFMCEAEGEKPKPYEGT 42
2IB6_A1|PDBID|CHAIN|SEQUENCE -----GSHMASKISDNVRIKLYMEGTVNNHFMCEAEGEKPKPYEGT 42
2C91_A1|PDBID|CHAIN|SEQUENCE -----MYSIKETMRVQLSMEGSVNGHYFKCTGKGEKPKPYEGT 39
SHHHHHHGRSSKNVKEFMRFKVRMEGTVNGHEFIEGEGGRPYEGH 53
167K_A1|PDBID|CHAIN|SEQUENCE -----MRSSKNVKEFMRFKVRMEGTVNGHEFIEGEGGRPYEGH 42
12G0_A1|PDBID|CHAIN|SEQUENCE -----MRSSKNVKEFMRFKVRMEGTVNGHEFIEGEGGRPYEGH 42
16G_A1|PDBID|CHAIN|SEQUENCE -----MRSSKNVKEFMRFKVRMEGTVNGHEFIEGEGGRPYEGH 42
12G0_H1|PDBID|CHAIN|SEQUENCE -----MRSSKNVKEFMRFKVRMEGTVNGHEFIEGEGGRPYEGH 42
12GP_A1|PDBID|CHAIN|SEQUENCE -----MRSSKNVKEFMRFKVRMEGTVNGHEFIEGEGGRPYEGH 42
2H8Q_A1|PDBID|CHAIN|SEQUENCE -----VIKEFMRFKVRMEGTVNGHEFIEGEGGRPYEGH 42
2VAE_A1|PDBID|CHAIN|SEQUENCE -----MASSEVDIKEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 42
2V4E_A1|PDBID|CHAIN|SEQUENCE -----NVIKPFMRKVRMEGSVNGHEFIEGEGGRPYEGT 37
2VAD_A1|PDBID|CHAIN|SEQUENCE -----MDNTEVDIKEFMRFKVRMEGSVNGHYFIEGEGGRPYEGT 42
2H5P_A1|PDBID|CHAIN|SEQUENCE ---MVSKEENMAIIEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 47
2H5R_A1|PDBID|CHAIN|SEQUENCE ---MVSKEENMAIIEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 47
2H5Q_A1|PDBID|CHAIN|SEQUENCE ---MVSKEENMAIIEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 47
2QLG_A1|PDBID|CHAIN|SEQUENCE ---MVSKEE---VIKEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 43
2QLT_A1|PDBID|CHAIN|SEQUENCE ---MVSKEE---VIKEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 43
2QLH_A1|PDBID|CHAIN|SEQUENCE ---MVSKEE---VIKEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 43
2H5Q_A1|PDBID|CHAIN|SEQUENCE ---MVSKEEDMAIIEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 47
3KCS_A1|PDBID|CHAIN|SEQUENCE ---MVSKEEDMAIIEFMRFKVRMEGSVNGHYFIEGEGGRPYEGT 47
3KCT_A1|PDBID|CHAIN|SEQUENCE ---MVSKEEDMAIIEFMRFKVRMEGSVNGHYFIEGEGGRPYEGT 47
3LF3_A1|PDBID|CHAIN|SEQUENCE -----EEDMAIIEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 42
3LF4_A1|PDBID|CHAIN|SEQUENCE -----EEDMAIIEFMRFKVRMEGSVNGHEFIEGEGGRPYEGT 42
2P4M_A1|PDBID|CHAIN|SEQUENCE -----MSVIATQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 38
2ARL_A1|PDBID|CHAIN|SEQUENCE -----GSVIATQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 38
1MOV_A1|PDBID|CHAIN|SEQUENCE -----GCVIATQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 38
1M0U_A1|PDBID|CHAIN|SEQUENCE -----GCVIATQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 38
2WH5_A1|PDBID|CHAIN|SEQUENCE HSSGLVPRGSHMVSIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 59
2WHT_A1|PDBID|CHAIN|SEQUENCE HSSGLVPRGSHMVSIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 59
2WHU_A1|PDBID|CHAIN|SEQUENCE HSSGLVPRGSHMVSIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 59
3TR8_A1|PDBID|CHAIN|SEQUENCE --SS-----VSVIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 40
3CGL_A1|PDBID|CHAIN|SEQUENCE SHHHHHHGSCKSVIKEMLIDLHLEGTNGHYFIEKGGKQPPNEG 53
226Z_A1|PDBID|CHAIN|SEQUENCE GRDLYDDDDKDPMSVIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 71
210V_A1|PDBID|CHAIN|SEQUENCE GRDLYDDDDKDPMSVIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 71
2POX_A1|PDBID|CHAIN|SEQUENCE GRDLYDDDDKDPMSVIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 71
2IEZ_A1|PDBID|CHAIN|SEQUENCE -----MVSIVKDPDKIKLRMEGAVNGHPPFAIEGVLGKPFPEGK 38
2210_A1|PDBID|CHAIN|SEQUENCE -----SHMSVIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 41
226Y_A1|PDBID|CHAIN|SEQUENCE -----SHMSVIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 41
2GXZ_A1|PDBID|CHAIN|SEQUENCE -----SMVSIVKDPDKIKLRMEGAVNGHPPFAIEGVLGKPFPEGK 57
2GX0_B1|PDBID|CHAIN|SEQUENCE -----SMVSIVKDPDKIKLRMEGAVNGHPPFAIEGVLGKPFPEGK 57
226X_A1|PDBID|CHAIN|SEQUENCE GRDLYDDDDKDPMSVIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 71
3L53_A1|PDBID|CHAIN|SEQUENCE -----TDPMSVIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 51
3L5A_A1|PDBID|CHAIN|SEQUENCE -----TDPMSVIAKQMTYKYVMSGTVNGHYFEVEGDGKPKPYEG 51
2DDC_A1|PDBID|CHAIN|SEQUENCE -----VSVITSEMKEIVRMEGAVNGHFKVITTKGSGQPPFEG 38
2DDD_A1|PDBID|CHAIN|SEQUENCE -----VSVITSEMKEIVRMEGAVNGHFKVITTKGSGQPPFEG 38
1X55_A1|PDBID|CHAIN|SEQUENCE -----VSVITSEMKEIVRMEGAVNGHFKVITTKGSGQPPFEG 38
2VVL_A1|PDBID|CHAIN|SEQUENCE -----HMSAIKDPDKIKLRMEGAVNGHFFVTDGDTGKPFPEGK 40
2VHH_A1|PDBID|CHAIN|SEQUENCE -----HMSAIKDPDKIKLRMEGAVNGHFFVTDGDTGKPFPEGK 40
2VJ3_A1|PDBID|CHAIN|SEQUENCE -----HMSAIKDPDKIKLRMEGAVNGHFFVTDGDTGKPFPEGK 40
12UX_A1|PDBID|CHAIN|SEQUENCE -----MSAIKDPDKIKLRMEGAVNGHFFVTDGDTGKPFPEGK 40

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

```
2BTJ_A | PDBID | CHAIN | SEQUENCE -----GGAIKPDMKINLRMEGNVNGHFVVDGDKTGPFEKG 38
2GW3_A | PDBID | CHAIN | SEQUENCE -----APMSLIKPEMKIKLLMEGNVNGHFVIEGDGKHPFEKG 40
2GW4_A | PDBID | CHAIN | SEQUENCE -----APMSLIKPEMKIKLLMEGNVNGHFVIEGDGKHPFEKG 40
2OTE_A | PDBID | CHAIN | SEQUENCE -----GVIKPDMKIKLMEGNVNGHFVIEGEGEKPPYDGTN 37
2OTB_A | PDBID | CHAIN | SEQUENCE -----GVIKPDMKIKLMEGNVNGHFVIEGEGEKPPYDGTN 37
2HQK_A | PDBID | CHAIN | SEQUENCE -----GVIKPDMKIKLMEGNVNGHFVIEGEGEKPPYDGTN 37
2VZK_A | PDBID | CHAIN | SEQUENCE -----MNLKEDMRVKVHMEGNVNGHFVIEGEGEKPPYDGTN 38
3ADF_A | PDBID | CHAIN | SEQUENCE -----GAHMSVIKPEMKIKLMRGTVNHNHFVIEGEGKNPYEGTQ 41
21CR_A | PDBID | CHAIN | SEQUENCE MRGSHHHHHHGSAGLTDMMHFRMEGCVDGHKFVIEGNGNPNPKGK 50
2FL1_A | PDBID | CHAIN | SEQUENCE -----SAHGLTDDMTMFRMEGCVDGHKFVIEGNGNPNPKGK 39
20GR_A | PDBID | CHAIN | SEQUENCE -----MAHSHKGLKEEMTKYHMEGCVNGHKFVITGEGIYPPFKGK 42
1XAE_A | PDBID | CHAIN | SEQUENCE -----MAHSHKGLKEEMTKYHMEGCVNGHKFVITGEGIYPPFKGK 42
1XA9_A | PDBID | CHAIN | SEQUENCE -----MAHSHKGLKEEMTKYHMEGCVNGHKFVITGEGIYPPFKGK 42
2PX5_A | PDBID | CHAIN | SEQUENCE -----SKHGLTKEMTKYHMEGCVNGHKFVITGEGIYPPFKGK 39
2PXW_A | PDBID | CHAIN | SEQUENCE -----SKHGLTKEMTKYHMEGCVNGHKFVITGEGIYPPFKGK 39
20JK_A | PDBID | CHAIN | SEQUENCE -----MAQSKHGLTKEMTKYHMEGCVNGHKFVITGEGIYPPFKGK 42
2RH7_A | PDBID | CHAIN | SEQUENCE -----MDLAKLGLKEEMTKYHMEGCVNGHKFVITGEGIYPPFKGK 42
2C9J_H | PDBID | CHAIN | SEQUENCE -----MSQLDNNLSVSYMKGNNVHFEYDGGDDSPSGQF 38
2G6X_A | PDBID | CHAIN | SEQUENCE -----LPAMEIECRITGTLNGVEFELVGGEGTPEQGRM 34
2G6Y_A | PDBID | CHAIN | SEQUENCE -----LPAMEIECRITGTLNGVEFELVGGEGTPEQGRM 34
2G3O_A | PDBID | CHAIN | SEQUENCE -----MPAMKIECRITGTLNGVEFELVGGEGTPEQGRM 34
2DD7_A | PDBID | CHAIN | SEQUENCE -----TTFKIESRIHGNLNGEKFELVGGVYG--EEGR 31
2DD9_D | PDBID | CHAIN | SEQUENCE -----TTFKIESRIHGNLNGEKFELVGGVYG--EEGR 31
. . . * . . . *
3EKJ_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 247
3EVR_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 190
3EKH_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 247
3EVR_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 210
3EK7_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 247
3EVL_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 247
3EVD_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 247
3EK4_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 247
3EK8_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 247
2G3O_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2G2S_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
1QYQ_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYG--VQCFSRYPDHMKHQDFFSAMP 88
2G6E_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 90
1YH1_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 90
1YH2_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 90
2G5Z_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 90
2G16_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 89
2HGD_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 90
2HG4_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 90
1YJ3_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 90
1YH1_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 90
1YJ2_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGAVQCFSRYPDHMKHQDFFSAMP 90
1QYF_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
1QXT_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2AIM_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2AIK_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2AIJ_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2AIL_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2HFC_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2HCG_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2QZ0_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2FZU_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
1KYR_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
1KYP_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
1KY5_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
2FQW_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
189C_D | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
1MYW_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGGLQCFARYPDHMKHQDFFSAMP 90
3ED8_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGGLQCFARYPDHMKHQDFFSAMP 89
1QY0_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGGLQCFARYPDHMKHQDFFSAMP 89
2QUL_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 92
1QY3_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
2QZT_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
2QRF_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
2HRS_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
2B3Q_D | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
2B3P_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
1KPS_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 99
3G11_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
3G12_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 90
3G9A_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
3K1K_B | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
1J08_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
1J1C_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
2A88_B | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
2A8A_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
3CD1_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 100
3CD9_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 100
3CB9_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 100
3CBE_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 100
20KW_F | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
20KY_B | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
2HQZ_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
2HJ0_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
202B_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 95
2H9W_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 92
2024_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 95
2029_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 95
2H6V_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 95
1RMO_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
1RMM_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 88
1RMP_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 88
1RRX_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 88
2Q6P_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
121Q_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
156Z_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
121P_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGVQCFSRYPDHMKHQDFFSAMP 89
1F0B_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGGLQCFARYPDHMKHQDFFSAMP 89
1F09_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGGLQCFARYPDHMKHQDFFSAMP 89
2YFP_A | PDBID | CHAIN | SEQUENCE TLKFICT---TGKLPVWPPTLVTTLYGGLQCFARYPDHMKHQDFFSAMP 89
```


Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

2GX0_B | PDBID | CHAIN | SEQUENCE EG-YSWERSMNYEDGGICNATNDITL--DG-DCYIYEIRFDGVNFPANGP 149
2Z6X_A | PDBID | CHAIN | SEQUENCE EG-YSWERSMNYEDGGICATNDITL--DG-DCFYIYEIRFDGVNFPANGP 163
3LSA_A | PDBID | CHAIN | SEQUENCE EG-YSWERSMIYEDGGICNATNDITL--DG-DCYICEIRFDGVNFPANGP 143
3LSA_A | PDBID | CHAIN | SEQUENCE EG-YSWERSMIYEDGGICNATNDITL--DG-DCYICEIRFDGVNFPANGP 143
2DDC_A | PDBID | CHAIN | SEQUENCE EG-YSWERSMNYEDGGICLATNNTMKKDGSCNFVEIRFDGVNFPANGP 133
2DDD_A | PDBID | CHAIN | SEQUENCE EG-YSWERSMNYEDGGICLATNNTMKKDGSCNFVEIRFDGVNFPANGP 133
1X5S_A | PDBID | CHAIN | SEQUENCE EG-YSWERSMNYEDGGICLATNNTMKKDGSCNFVEIRFDGVNFPANGP 133
2VVL_A | PDBID | CHAIN | SEQUENCE KG-YSWERSLTFEDGGICIAIARNITM--EG-DTFYKVRFHGVNFPANGP 132
2VHL_A | PDBID | CHAIN | SEQUENCE KG-YSWERSLTFEDGGICIAIARNITM--EG-DTFYKVRFHGVNFPANGP 132
2VJ3_A | PDBID | CHAIN | SEQUENCE KG-YSWERSLTFEDGGICIAIARNITM--EG-DTFYKVRFHGVNFPANGP 132
12UX_A | PDBID | CHAIN | SEQUENCE KG-YSWERSLTFEDGGICIAIARNITM--EG-DTFYKVRFHGVNFPANGP 130
2BTJ_A | PDBID | CHAIN | SEQUENCE KG-YSWERSLTFEDGGICIAIARNITM--EG-DTFYKVRFHGVNFPANGP 130
2GW3_A | PDBID | CHAIN | SEQUENCE KG-FSWERSLTFEDGGICIAIARNITL--KG-DTFYKVRFHGVNFPANGP 132
2GWA_A | PDBID | CHAIN | SEQUENCE KG-FSWERSLTFEDGGICIAIARNITL--KG-DTFYKVRFHGVNFPANGP 132
2OTE_A | PDBID | CHAIN | SEQUENCE EG-YSWERTMTFEDKGIKVKKSDISM--EE-DSFYIYIHLKGFNFPANGP 129
2OTB_A | PDBID | CHAIN | SEQUENCE EG-YSWERTMTFEDKGIKVKKSDISM--EE-DSFYIYIHLKGFNFPANGP 129
2HQK_A | PDBID | CHAIN | SEQUENCE EG-YSWERTMTFEDKGIKVKKSDISM--EE-DSFYIYIHLKGFNFPANGP 129
2VZX_A | PDBID | CHAIN | SEQUENCE EG-YSWERTMTFEDKGIKVKKSDISM--EE-DSFYIYIHLKGFNFPANGP 130
3ADF_A | PDBID | CHAIN | SEQUENCE EG-YHWERSMTYEDGGICTATSNTSM--RG-DCFFYDIFDGVNFPANGP 133
21CR_A | PDBID | CHAIN | SEQUENCE AG-YTWRSRFRFEDGAVCICSADITVN-VRENCIYHSTFYGVNFPADGP 144
2FL1_A | PDBID | CHAIN | SEQUENCE AG-YTWRSRFRFEDGAVCICSADITVN-VRENCIYHSTFYGVNFPADGP 133
20GR_A | PDBID | CHAIN | SEQUENCE AG-YTWRSRFLFEDGAVCICNVDITVS-VKNCIYHKSIFGNMFPADGP 136
1XAE_A | PDBID | CHAIN | SEQUENCE AG-YTWRSRFLFEDGAVCICNVDITVS-VKNCIYHKSIFGNMFPADGP 136
1XAG_A | PDBID | CHAIN | SEQUENCE AG-YTWRSRFLFEDGAVCICNVDITVS-VKNCIYHKSIFGNMFPADGP 136
2PKX_A | PDBID | CHAIN | SEQUENCE AG-YTWRSRFLFEDGAVCICNVDITVS-VKNCIYHKSIFGNMFPADGP 133
2PKW_A | PDBID | CHAIN | SEQUENCE AG-YTWRSRFLFEDGAVCICNVDITVS-VKNCIYHKSIFGNMFPADGP 133
20JK_A | PDBID | CHAIN | SEQUENCE AG-YTWRSRFLFEDGAVCICNVDITVS-VKNCIYHKSIFGNMFPADGP 136
2RH7_A | PDBID | CHAIN | SEQUENCE EG-FTYERNIRYQDGTAVKVSISLE--DGKFTVNVDFKAKDLRMRGP 134
2C9J_H | PDBID | CHAIN | SEQUENCE EA-FQNNRIEFEDGGIVNMSDITYK--DKVLHGDVWALGVNFPANGP 130
266X_A | PDBID | CHAIN | SEQUENCE NGGYTNTRIEKYEDGGVLHVSFSYRY--EAGRVIGDFKVMGTGFPEPDSV 126
266Y_A | PDBID | CHAIN | SEQUENCE NGGYTNTRIEKYEDGGVLHVSFSYRY--EAGRVIGDFKVMGTGFPEPDSV 126
2630_A | PDBID | CHAIN | SEQUENCE NGGYTNTRIEKYEDGGVLHVSFSYRY--EAGRVIGDFKVMGTGFPEPDSV 126
2D07_A | PDBID | CHAIN | SEQUENCE NGGYTNTRIEKYEDGGILEVNFRTY--EFNKIIGDVEICIGHGFPSPQ 123
2D09_D | PDBID | CHAIN | SEQUENCE NGGYTNTRIEKYEDGGILEVNFRTY--EFNKIIGDVEICIGHGFPSPQ 123

3EKJ_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNTRDQLTEEQIAEFKEAFS-----LFDKQDGD 328
3EVP_A | PDBID | CHAIN | SEQUENCE ILGHKLEYN-----LFDKQDGD 245
3EKH_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNTRDQLTEEQIAEFKEAFS-----LFDKQDGD 328
3EVR_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNTRDQLTEEQIAEFKEAFS-----LFDKQDGD 291
3EK7_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNTRDQLTEEQIAEFKEAFS-----LFDKQDGD 328
3EVL_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNTRDQLTEEQIAEFKEAFS-----LFDKQDGD 328
3EVV_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNTRDQLTEEQIAEFKEAFS-----LFDKQDGD 328
3EK4_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNTRDQLTEEQIAEFKEAFS-----LFDKQDGD 328
3EK8_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNTRDQLTEEQIAEFKEAFS-----LFDKQDGD 328
2G3D_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2G2S_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
1QYQ_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 159
2G6E_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
1YHH_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
1YHG_B | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2G5Z_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2G16_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
2HG0_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2HGY_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
1YJF_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
1YHL_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
1YJ2_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSG-NVYITA-----DKQKNG 161
1QFY_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
1QXT_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2AWM_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2AWK_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2AWJ_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2AWL_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2HFC_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2HCG_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2QZ0_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
2FZU_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
1KYR_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNFNSG-NVYITA-----DKQKNG 161
1KYP_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNFNSG-NVYITA-----DKQKNG 161
1KYS_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNFNSG-NVYITA-----DKQKNG 161
2FWQ_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNFNSG-NVYITA-----DKQKNG 161
1B9C_D | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
1MYW_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
3E08_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNFNSH-NVYITATRGTAARPACKIPNDLKQKVMHNDKQKNG 184
1QY0_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
2QU1_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 163
1QY3_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
2QT2_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
2QRF_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
2HRS_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
2B3Q_D | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
2B3P_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNFNSH-NVYITA-----DKQKNG 160
1KPS_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 170
3G1L_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
3GJ2_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 161
3G9A_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
3K1K_B | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 160
1JCB_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 160
1JCL_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 160
2A8H_B | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 160
2A8A_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 160
3CD1_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 172
3CD9_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 172
3CB9_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 172
3CBE_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYITA-----DKQKNG 172
20KW_F | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 160
20KY_B | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 160
2HQZ_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 160
2HJO_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 160
202B_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 166
2H9W_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 163
2024_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 166
2029_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 166
2H6V_A | PDBID | CHAIN | SEQUENCE ILGHKLEYNNYNSH-NVYIMA-----DKQKNG 166

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

1RM0_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1RM1_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 159
1RMP_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 159
1RRX_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 159
2Q6P_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
121Q_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
156Z_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
121P_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1F0B_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSQ-NVYIMA-----DKQKNG 160
1F09_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSQ-NVYIMA-----DKQKNG 160
2YFP_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSG-NVYIMA-----DKQKNG 160
3D08_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D01_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D09_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0A_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0C_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0D_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0E_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0F_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0H_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0I_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0J_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0K_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0L_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0M_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0N_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0O_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0Q_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0R_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0S_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0T_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0U_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0V_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0W_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0X_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3D0Z_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3DQ2_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
3DQ3_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 165
1HUY_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 163
1YFP_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 158
2JAD_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-CVYIVA-----DKQKNG 160
1H6R_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-CVYIVA-----DKQKNG 160
1C4F_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1EMB_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1Q4A_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1EMA_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1Q4B_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1EMG_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
2DUF_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSD-NVYIMA-----DKQKNG 160
2DUJ_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSD-NVYIMA-----DKQKNG 160
2DU1_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSD-NVYIMA-----DKQKNG 160
2DUG_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSN-NVYIMA-----DKQKNG 160
2DUH_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSN-NVYIMA-----DKQKNG 160
1Q4E_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1Q73_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1Q4D_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1Q4C_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1J8Z_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSG-NVYIMA-----DKQKNG 160
1J8Y_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSG-NVYIMA-----DKQKNG 160
1W75_D PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1W7U_D PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1W7T_D PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1HCJ_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
2QLE_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1BFP_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
2EMO_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 161
1EMF_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 161
2EMD_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 161
2EMN_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 161
1EMM_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 161
1EMC_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 161
1EML_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 161
1EME_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 161
1EMK_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 161
2WUR_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1GFL_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYIMA-----DKQKNG 160
1RM9_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYITA-----DKQKNG 160
2WSN_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYITA-----DKQKNG 161
2Q57_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSD-NVYITA-----DKQKNG 178
2WSO_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSD-NVYITA-----DKQKNG 161
10XF_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYITA-----DKQKNG 159
1CV7_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYITA-----DKQKNG 160
3GEX_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYITA-----DKQKNG 166
10XD_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYITA-----DKQKNG 160
10XE_A PDBID CHAIN SEQUENCE	ILGHKLEYNNSH-NVYITA-----DKQKNG 160
3LVC_A PDBID CHAIN SEQUENCE	ILGNKMEYNNAH-NVYITM-----DKAKNG 160
3LVD_A PDBID CHAIN SEQUENCE	ILGNKMEYNNAH-NVYITM-----DKAKNG 160
3LVA_A PDBID CHAIN SEQUENCE	ILGNKMEYNNAH-NVYITM-----DKAKNG 160
2HPW_A PDBID CHAIN SEQUENCE	ILGMRVLYHSPPH-AVYILP-----DRKNGG 161
2WIQ_A PDBID CHAIN SEQUENCE	IMRDQLVDIILPNE THMFPHG-----PNA 178
3A85_A PDBID CHAIN SEQUENCE	IMRDQLVDIILPNE THMFPHG-----PNA 159
2WIS_A PDBID CHAIN SEQUENCE	IMRDQLVDIILPNE THMFPHG-----PNA 178
3G83_A PDBID CHAIN SEQUENCE	IMRDQLVDIILPNE THMFPHG-----PNA 156
3GL4_A PDBID CHAIN SEQUENCE	IMRDQLVDIILPNE THMFPHG-----PNA 156
2A46_A PDBID CHAIN SEQUENCE	VMAKKTGWDPSFEKMTVC-----DGI 169
2A48_A PDBID CHAIN SEQUENCE	VMAKKTGWDPSFEKMTVC-----DGI 169
2A47_A PDBID CHAIN SEQUENCE	VMAKKTGWDPSFEKMTVC-----DGI 169
2ZMW_A PDBID CHAIN SEQUENCE	IMQNSVDWEPSTEKITAS-----DGV 155
2ZMU_A PDBID CHAIN SEQUENCE	IMQNSVDWEPSTEKITAS-----DGV 155
2Z06_A PDBID CHAIN SEQUENCE	IAQNKTLGWEPSTEKMTVR-----DGS 182
2Z07_A PDBID CHAIN SEQUENCE	IMQNKTLGWEPSTEKMTVR-----DGS 155
3B8X_B PDBID CHAIN SEQUENCE	VMQKKTLGWEASTEMLYPA-----DGG 165
3B8X_A PDBID CHAIN SEQUENCE	VMQKKTLGWEASTEMLYPA-----DGG 165
3B8B_A PDBID CHAIN SEQUENCE	VMQKKTLGWEASTEMLYPA-----DGG 165
3B8C_A PDBID CHAIN SEQUENCE	VMQKKTLGWEASTEMLYPA-----DGG 165
3H10_A PDBID CHAIN SEQUENCE	VMQKKTLGWEASTEMLYPA-----DGG 153

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

3H1R_A PDBID CHAIN SEQUENCE	VMQKKTLGWEAHEMLYPA-----DGG 153
3M22_A PDBID CHAIN SEQUENCE	VMQKKTLGWEANTEMLYPA-----DGG 157
3M24_A PDBID CHAIN SEQUENCE	VMQKKTLGWEAFETLYPA-----DGG 154
3IP2_A PDBID CHAIN SEQUENCE	VMQKKTLGWEASTEMLYPA-----DGG 157
3E5T_A PDBID CHAIN SEQUENCE	VMQKKTGWEPNTEMLYPA-----DGG 166
3E5V_A PDBID CHAIN SEQUENCE	VMQKKTGWEPSTEMLYPA-----DGG 166
1UI5_B PDBID CHAIN SEQUENCE	VMQKKTGWEPNTEMLYPA-----DGG 155
3E5W_A PDBID CHAIN SEQUENCE	VMQKKTGWEPSTEMLYPA-----DGG 166
2A50_A PDBID CHAIN SEQUENCE	VMQNKAGRWEPAEIVYEV-----DGV 164
2A52_A PDBID CHAIN SEQUENCE	VMQNKAGRWEPAEIVYEV-----DGV 164
1XMZ_A PDBID CHAIN SEQUENCE	VMQNKAGRWEPAEIVYEV-----DGV 164
2A54_C PDBID CHAIN SEQUENCE	VMQNKAGRWEPAEIVYEV-----DGV 164
2A53_A PDBID CHAIN SEQUENCE	VMQNKAGRWEPAEIVYEV-----DGV 164
2A56_A PDBID CHAIN SEQUENCE	VMQNKAGRWEPAEIVYEV-----DGV 164
1XQM_A PDBID CHAIN SEQUENCE	VMQNKVGRWEPAEIVYEV-----DGV 153
3CFH_L PDBID CHAIN SEQUENCE	VMQNKAGWEPGCEILYEV-----DGV 153
3CFH_S PDBID CHAIN SEQUENCE	VMQNKAGWEPGCEILYEV-----DGV 153
3CFA_L PDBID CHAIN SEQUENCE	VMQNKAGWEPGCEILYEV-----DGV 153
1Y2W_A PDBID CHAIN SEQUENCE	VMQNKSGWEPSTEIVYEV-----NGV 151
2I85_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPCEMRYTR-----DGV 156
2I86_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPCEMRYTR-----DGV 156
2C9T_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPSTEIVYEV-----DGG 153
1G7K_A PDBID CHAIN SEQUENCE	VMQKKTGWAEATERLYPR-----DGV 167
1ZG0_A PDBID CHAIN SEQUENCE	VMQKKTGWAEATERLYPR-----DGV 156
1G6X_A PDBID CHAIN SEQUENCE	VMQKKTGWAEATERLYPR-----DGV 156
1ZG0_H PDBID CHAIN SEQUENCE	VMQKKTGWAEATERLYPR-----DGV 156
1ZGP_A PDBID CHAIN SEQUENCE	VMQKKTGWAEATERLYPR-----DGV 156
2H80_A PDBID CHAIN SEQUENCE	VMQKKTGWAEATERLYPR-----DGV 150
2V4E_A PDBID CHAIN SEQUENCE	VMQKKTGWEPSTERLYPR-----DGV 156
2V4F_A PDBID CHAIN SEQUENCE	VMQKKTGWEPSTERLYPR-----DGV 151
2V4D_A PDBID CHAIN SEQUENCE	VMQKKTGWEPSTEKLYPQ-----DGV 156
2H5P_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 161
2H5R_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 161
2H50_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 161
2QLG_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 157
2QLL_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 157
2QLH_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 157
2H5Q_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 161
3KCS_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 161
3KCT_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 161
3LF3_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 156
3LF4_A PDBID CHAIN SEQUENCE	VMQKKTGWAEASERMYPE-----DGA 156
2P4M_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPSESLFAR-----GGM 152
2ARL_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPSESLFAR-----GGM 152
1M0V_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPSESLFAR-----GGM 152
1M0U_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPSESLFAR-----GGM 152
2WH5_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPSTERLFAR-----DGM 173
2WHT_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPSTERLFAR-----DGM 173
2WHU_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPSTERLFAR-----DGM 173
3IR8_A PDBID CHAIN SEQUENCE	VMQKKTQGWEPSTERLFAR-----DGM 154
3CGL_A PDBID CHAIN SEQUENCE	VVQKKTGWEPSTERLYPR-----DGV 167
2Z6Z_A PDBID CHAIN SEQUENCE	VMQKRTVKWEPSTEKLYVR-----DGV 185
2I0V_A PDBID CHAIN SEQUENCE	VMQKRTVKWEPSTEKLYVR-----DGV 185
2POX_A PDBID CHAIN SEQUENCE	VMQKRTVKWEPSTEKLYVR-----DGV 185
2IEZ_A PDBID CHAIN SEQUENCE	VMQKRTVKWEPSTEKLYVR-----DGV 152
2Z10_A PDBID CHAIN SEQUENCE	VMQKRTVKWEPSTEKLYVR-----DGV 155
2Z6Y_A PDBID CHAIN SEQUENCE	VMQKRTVKWEPSTEKLYVR-----DGV 155
2GZ2_A PDBID CHAIN SEQUENCE	VMQKRTVKWEPSTEKLYVR-----DGV 171
2GX0_B PDBID CHAIN SEQUENCE	VMQKRTVKWEPSTEKLYVR-----DGV 171
2Z6X_A PDBID CHAIN SEQUENCE	VMQKRTVKWEPSTEKLYVR-----DGV 185
3LS3_A PDBID CHAIN SEQUENCE	VMQKRTVKWELSTEKLYVR-----DGV 165
3LSA_A PDBID CHAIN SEQUENCE	VMQKRTVKWELSTEKLYVR-----DGV 165
2DDC_A PDBID CHAIN SEQUENCE	VMQRKTVKWESSTEKMYVR-----DGV 155
2DD0_A PDBID CHAIN SEQUENCE	VMQRKTVKWESSTEKMYVR-----DGV 155
1X5S_A PDBID CHAIN SEQUENCE	VMQRKTVKWESSTEKMYVR-----DGV 155
2VVI_A PDBID CHAIN SEQUENCE	VMQKKTWKWEPSTEKMYVR-----DGV 154
2VHL_A PDBID CHAIN SEQUENCE	VMQKKTWKWEPSTEKMYVR-----DGV 154
2VJ2_A PDBID CHAIN SEQUENCE	VMQKKTWKWEPSTEKMYVR-----DGV 154
1ZUX_A PDBID CHAIN SEQUENCE	VMQKKTWKWEPSTEKMYVR-----DGV 152
2BTJ_A PDBID CHAIN SEQUENCE	VMQKKTWKWEPSTEKMYVR-----DGV 152
2GW3_A PDBID CHAIN SEQUENCE	VMQKKTWKWAEATEKMYLR-----DGV 154
2GW4_A PDBID CHAIN SEQUENCE	VMQKKTWKWAEATEKMYLR-----DGV 154
2OTE_A PDBID CHAIN SEQUENCE	VMQKKTWKWEPSTEILYVR-----DGV 151
2OT8_A PDBID CHAIN SEQUENCE	VMQKKTWKWEPSTEILYVR-----DGV 151
2HQK_A PDBID CHAIN SEQUENCE	VMQKKTGWDASTERMYVR-----DGV 151
2VZX_A PDBID CHAIN SEQUENCE	VMQKKTWKWEPSTEKLHVR-----DGL 152
3ADF_A PDBID CHAIN SEQUENCE	VMQKKTWKWEPSTEKMYVE-----DGV 155
2ICR_A PDBID CHAIN SEQUENCE	VVKKMTTNWEPSECKIIPVPS-----GKI 168
2FL1_A PDBID CHAIN SEQUENCE	VVKKMTTNWEPSECKIIPVPS-----GKI 157
20GR_A PDBID CHAIN SEQUENCE	VVKKMTTNWEPSECKIIPVPS-----GKI 160
1XAE_A PDBID CHAIN SEQUENCE	VVKKMTTNWEPSECKIIPVPS-----GKI 160
1XAG_A PDBID CHAIN SEQUENCE	VVKKMTTNWEPSECKIIPVPS-----GKI 160
2PXS_A PDBID CHAIN SEQUENCE	VVKKMTTNWEPSECKIIPVPS-----GKI 157
2PXW_A PDBID CHAIN SEQUENCE	VVKKMTTNWEPSECKIIPVPS-----GKI 157
20JK_A PDBID CHAIN SEQUENCE	VVKKMTTNWEPSECKIIPVPS-----GKI 160
2RH7_A PDBID CHAIN SEQUENCE	VMQDQIVGMQPSYSEMYTN-----VTS 156
2C9J_H PDBID CHAIN SEQUENCE	VMKNEIVMEPAEETLAK-----NGV 152
2G6X_A PDBID CHAIN SEQUENCE	IFTDKIIRSNATVEHLHPMG-----DND 149
2G6Y_A PDBID CHAIN SEQUENCE	IFTDKIIRSNATVEHLHPMG-----DND 149
2G30_A PDBID CHAIN SEQUENCE	IFTDKIIRSNATVEHLHPMG-----DND 149
2DD7_A PDBID CHAIN SEQUENCE	IFKDTIVKSCPTVDLMLPMS-----GNI 146
2DD9_D PDBID CHAIN SEQUENCE	IFKDTIVKSCPTVDLMLPMS-----GNI 146
:	:
3EKJ_A PDBID CHAIN SEQUENCE	GITTKQLGTVMRSLGQNPTEAELQDMIN---EVDADGNGTIDFPFLTMM 375
3EVP_A PDBID CHAIN SEQUENCE	-----
3EKH_A PDBID CHAIN SEQUENCE	TITTKELGTVMRSLGQNPTEAELQDMIN---EVDADGNGTIDFPFLTMM 375
3EVR_A PDBID CHAIN SEQUENCE	TITTKELGTVMRSLGQNPTEAELQDMIN---EVDADGNGTIDFPFLTMM 338
3EK7_A PDBID CHAIN SEQUENCE	TITTKELGTVMRSLGQNPTEAELQDMIN---EVDADGNGTIDFPFLTMM 375
3EVL_A PDBID CHAIN SEQUENCE	TITTKELGTVMRSLGQNPTEAELQDMIN---EVDADGNGTIDFPFLTMM 375
3EVL_A PDBID CHAIN SEQUENCE	TITTKELGTVMRSLGQNPTEAELQDMIN---EVDADGNGTIDFPFLTMM 375
3EVA_A PDBID CHAIN SEQUENCE	TITTKELGTVMRSLGQNPTEAELQDMIN---EVDADGNGTIDFPFLTMM 375
3EK4_A PDBID CHAIN SEQUENCE	TITTKELGTVMRSLGQNPTEAELQDMIN---EVDADGNGTIDFPFLTMM 375
3EK8_A PDBID CHAIN SEQUENCE	TITTKELGTVMRSLGQNPTEAELQDMIN---EVDADGNGTIDFPFLTMM 375
2G3D_A PDBID CHAIN SEQUENCE	IKANFKIRHNIE-----DGSVQLADHYQ---QNTPIGDGPVLLPDNHYLS 203
2G2S_A PDBID CHAIN SEQUENCE	IKANFKIRHNIE-----DGSVQLADHYQ---QNTPIGDGPVLLPDNHYLS 203

Supplementary Material (ESI) for Molecular BioSystems
 This journal is (c) The Royal Society of Chemistry, 2011

2WHT_A1 PDBID CHAIN SEQUENCE	LIGNDYMALKLE-----GGHYLCEPKS--TYKAKP---VRMPGRHEID 213
2WHU_A1 PDBID CHAIN SEQUENCE	LIGNDYMALKLE-----GGHYLCEPKS--TYKAKP---VRMPGRHEID 213
31R8_A1 PDBID CHAIN SEQUENCE	LIGNDYMALKLE-----GGHYLCEPKS--TYKAKP---VRMPGRHEID 194
3CGL_A1 PDBID CHAIN SEQUENCE	LIGDIHHALTV-----GGHYACDIKT--VYRAKAA--LKMPGYHYVD 208
2Z6Z_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYHFVD 225
21OV_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYHFVD 225
2POX_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYHFVD 225
21E2_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYHFVD 192
221Q_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYHFVD 195
2Z6Y_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYHFVD 195
2G2Z_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYHFVD 211
2G00_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYHFVD 211
2Z6X_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYHFVD 225
3L53_A1 PDBID CHAIN SEQUENCE	LKSDGNYSALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYH5VD 205
3LSA_A1 PDBID CHAIN SEQUENCE	LKSDGNYSALSLE-----GGHYRCDFKT--TYKAK-KV--VQLPDYH5VD 205
20DC_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALLQ-----GGHYRCDFRT--TYKAK-KV--VQLPDYHFVD 195
20DD_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALLQ-----GGHYRCDFRT--TYKAK-KV--VQLPDYHFVD 195
1X55_A1 PDBID CHAIN SEQUENCE	LKGDVNMMALLQ-----GGHYRCDFRT--TYKAK-KV--VQLPDYHFVD 195
2VVT_A1 PDBID CHAIN SEQUENCE	LTGDITMALLLE-----GNAHYRCDSRT--TYKAKEKQ--VKLPYGHYVD 195
2VHV_A1 PDBID CHAIN SEQUENCE	LTGDITMALLLE-----GNAHYRCDSRT--TYKAKEKQ--VKLPYGHYVD 195
2VJ3_A1 PDBID CHAIN SEQUENCE	LTGDITMALLLE-----GNAHYRCDSRT--TYKAKEKQ--VKLPYGHYVD 195
12UX_A1 PDBID CHAIN SEQUENCE	LTGDITMALLLE-----GNAHYRCDFRT--TYKAKEKQ--VKLPYGHYVD 193
28TJ_A1 PDBID CHAIN SEQUENCE	LTGDITMALLLE-----GNAHYRCDFRT--TYKAKEKQ--VKLPYGHYVD 193
2G03_A1 PDBID CHAIN SEQUENCE	LTGDITMALLLE-----GNAHYRCDFRT--TYKSRQEQ--VKLPYGHYVD 195
2G04_A1 PDBID CHAIN SEQUENCE	LTGDITMALLLE-----GNAHYRCDFRT--TYKSRQEQ--VKLPYGHYVD 195
20TE_A1 PDBID CHAIN SEQUENCE	LVGDIKHKLLE-----GGHHRVDFKT--IYRAK-KA--VKLPDYHFVD 191
20TR_A1 PDBID CHAIN SEQUENCE	LVGDIKHKLLE-----GGHHRVDFKT--IYRAK-KA--VKLPDYHFVD 191
2H0K_A1 PDBID CHAIN SEQUENCE	LKGDVXKLLLE-----GGHHRVDFKT--IYRAK-KA--VKLPDYHFVD 191
2VZX_A1 PDBID CHAIN SEQUENCE	LVGINMALLE-----GGHYRCDFKT--TYKAK-KV--VQLPDHAFVD 192
3ADF_A1 PDBID CHAIN SEQUENCE	LKGDVWMLLLE-----GGHYRCDFKT--TYKAK-KE--VRLPDAHKVD 195
21CR_A1 PDBID CHAIN SEQUENCE	LKGDVSMYLLK-----DGGRYRCQFDT--IYKAKTEP--KEMPDWHFIQ 209
2FL1_A1 PDBID CHAIN SEQUENCE	LKGDVSMYLLK-----DGGRYRCQFDT--IYKAKTEP--KEMPDWHFIQ 198
20GR_A1 PDBID CHAIN SEQUENCE	LKGDVSMYLLK-----DGGRYRCQFDT--IYKAKSVP--SKMPFHFIFQ 201
1XA6_A1 PDBID CHAIN SEQUENCE	LKGDVSMYLLK-----DGGRYRCQFDT--IYKAKSVP--SKMPFHFIFQ 201
1XA9_A1 PDBID CHAIN SEQUENCE	LKGDVSMYLLK-----DGGRYRCQFDT--IYKAKSVP--SKMPFHFIFQ 201
2PXS_A1 PDBID CHAIN SEQUENCE	LKGDVSMYLLK-----DGGRLRCQFDT--IYKAKSVP--RKMPDWHFIQ 198
2PXW_A1 PDBID CHAIN SEQUENCE	LKGDVSMYLLK-----DGGRLRCQFDT--IYKAKSVP--RKMPDWHFIQ 198
20JK_A1 PDBID CHAIN SEQUENCE	LKGDVSMYLLK-----DGGRLRCQFDT--IYKAKSVP--RKMPDWHFIQ 201
2RH7_A1 PDBID CHAIN SEQUENCE	VIGECIIAFLKQ-----TGKHFYHMR--VYKSQKP--VTMLPYHFIQ 197
2C9J_H PDBID CHAIN SEQUENCE	LVGFCPKAYLLK-----DGSYYGHMT--IYRYSKQG--QLPFGFIK 193
2G6X_A1 PDBID CHAIN SEQUENCE	LVGSDVTRFSLR-----DGGYYSVVDSSHMFKSAIHPISLQNGGPMFAF 194
2G6Y_A1 PDBID CHAIN SEQUENCE	LVGSDVTRFSLR-----DGGYYSVVDSSHMFKSAIHPISLQNGGPMFAF 194
2G3Q_A1 PDBID CHAIN SEQUENCE	LVGSDVTRFSLR-----DGGYYSVVDSSHMFKSAIHPISLQNGGPMFAF 194
2D07_A1 PDBID CHAIN SEQUENCE	IASSYARAFQKL-----DGSFYAEVKNINDFKNIPHEFSGSKG-PMFTH 190
2D09_D PDBID CHAIN SEQUENCE	IASSYARAFQKL-----DGSFYAEVKNINDFKNIPHEFSGSKG-PMFTH 190
3EKJ_A1 PDBID CHAIN SEQUENCE	ARKMKDSEEEIREAFRVFKDGNQYISAAQLRHVMNLGKLTDEEVD 425
3EVP_A1 PDBID CHAIN SEQUENCE	ARKMKDSEEEIREAFRVFKDGNQYISAAQLRHVMNLGKLTDEEVD 425
3EKH_A1 PDBID CHAIN SEQUENCE	ARKMKDSEEEIREAFRVFKDGNQYISAAQLRHVMNLGKLTDEEVD 425
3EVR_A1 PDBID CHAIN SEQUENCE	ARKMKDSEEEIREAFRVFKDGNQYISAAQLRHVMNLGKLTDEEVD 388
3EK7_A1 PDBID CHAIN SEQUENCE	ARKMKDSEEEIREAFRVFKDGNQYISAAQLRHVMNLGKLTDEEVD 425
3EVL_A1 PDBID CHAIN SEQUENCE	ARKMKDSEEEIREAFRVFKDGNQYISAAQLRHVMNLGKLTDEEVD 425
3EVV_A1 PDBID CHAIN SEQUENCE	ARKMKDSEEEIREAFRVFKDGNQYISAAQLRHVMNLGKLTDEEVD 425
3EK4_A1 PDBID CHAIN SEQUENCE	ARKMKDSEEEIREAFRVFKDGNQYISAAQLRHVMNLGKLTDEEVD 425
3EK8_A1 PDBID CHAIN SEQUENCE	ARKMKDSEEEIREAFRVFKDGNQYISAAQLRHVMNLGKLTDEEVD 425
2G3D_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
2G2S_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1QYQ_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 237
2G6E_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1YHI_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1YHG_B PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
2G5Z_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
2G1G_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 238
2HG0_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
2HCY_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1YJF_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1YHI_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1YJ2_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1QYF_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGI----- 230
1QXT_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGI----- 230
2AWM_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGI----- 230
2AWK_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGI----- 230
2AWJ_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGI----- 230
2AWL_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGI----- 230
2HFC_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
2HCG_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
2QZ0_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGI----- 230
2FZU_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1KYR_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1KYP_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1KYS_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
2FWQ_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
1B9C_D PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 238
1MW_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 239
3E0B_A1 PDBID CHAIN SEQUENCE	YQTVLSDPNE-KRDHMLLE-----FVTAAGITLGDMLYK----- 262
1QYQ_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 238
2QUL_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 241
1QY3_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGIT----- 229
2QI2_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 238
2QRF_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGIT----- 230
2HRS_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FHTAAGITHGMDLYK----- 238
2B3Q_D PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYKSGHHHHH 246
2B3P_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYKSGHHHHH 246
1KP5_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGIVPRGTGLYK----- 248
3G11_A1 PDBID CHAIN SEQUENCE	HQSALSKDPNE-KRDHMLLE-----FVTAAGIT----- 231
3GJ2_A1 PDBID CHAIN SEQUENCE	HQSALSKDPNE-KRDHMLLE-----FVTAAGIT----- 231
3G9A_A1 PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 238
3K1K_B PDBID CHAIN SEQUENCE	TQSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 238
1JC0_A1 PDBID CHAIN SEQUENCE	TCSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 238
1J1C_A1 PDBID CHAIN SEQUENCE	TCSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 238
2AH8_B PDBID CHAIN SEQUENCE	TCSALSKDPNE-KRDHMLLE-----RVTAAGITHGMDLYK----- 238
2AHA_A1 PDBID CHAIN SEQUENCE	TCSALSKDPNE-KRDHMLLE-----RVTAAGITHGMDLYK----- 238
3CD1_A1 PDBID CHAIN SEQUENCE	TCSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 250
3CD9_A1 PDBID CHAIN SEQUENCE	TCSALSKDPNE-KRDHMLLE-----FVTAAGITHGMDLYK----- 250

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

2A48_A1	PDBID CHAIN SEQUENCE	HRIARTDLKGGN-SVQLTEH-----AVAHITSVVPF-----	240
2A47_A1	PDBID CHAIN SEQUENCE	TRIARTDLKGGN-SVQLTEH-----AVAHITSVVPF-----	240
22M0_A1	PDBID CHAIN SEQUENCE	HRLVLR---KTEGN-ITELVED-----AVAHSELEHHHHH-----	226
22M1_A1	PDBID CHAIN SEQUENCE	HRLVLR---KTEGN-ITELVED-----AVAHSELEHHHHH-----	226
2206_A1	PDBID CHAIN SEQUENCE	HR1SS---VWGT-KIQMED-----ATAHLYPFGVPCQ-----	254
2207_A1	PDBID CHAIN SEQUENCE	HR1SS---VWGT-KIQMED-----ATAHLYPFGS-----	224
38X9_B1	PDBID CHAIN SEQUENCE	RRLERIKEADKET-YVEQHEV-----AVARYCD-LPSKLGHLN-----	245
38X9_A1	PDBID CHAIN SEQUENCE	RRLERIKEADKET-YVEQHEV-----AVARYCD-LPSKLGHLN-----	245
38X8_A1	PDBID CHAIN SEQUENCE	RRLERIKEADKET-YVEQHEV-----AVARYCD-LPSKLGHLN-----	245
38X7_A1	PDBID CHAIN SEQUENCE	RRLERIKEADKET-YVEQHEV-----AVARYCD-LPSKLGHLN-----	245
3H10_A1	PDBID CHAIN SEQUENCE	YRLERIKEADKET-YVEQHEV-----AVARYCD-LPSKLGHLN-----	233
3H1R_A1	PDBID CHAIN SEQUENCE	YRLERIKEADKET-YVEQHEV-----AVARYCD-LPSKLGHLN-----	233
3M22_A1	PDBID CHAIN SEQUENCE	HLRERIKEADKET-YVEQHEV-----AVARYCD-LPSKLGHLN-----	237
3M24_A1	PDBID CHAIN SEQUENCE	YRLERIKEANNET-YVEQHEV-----AVARYCD-LPSKLGHLN-----	234
3IP2_A1	PDBID CHAIN SEQUENCE	RRLERIKEADNET-YVEQHEV-----AVARYCD-LPSKLGHLNGMDELY	243
3E5T_A1	PDBID CHAIN SEQUENCE	HLRERLEESDKEM-FVQHEH-----AVAKFCD-LPSKLGRL-----	244
3E5V_A1	PDBID CHAIN SEQUENCE	HLRERLEESDKEM-FVQHEH-----AVAKFCD-LPSKLGRL-----	244
1U1S_B1	PDBID CHAIN SEQUENCE	HLRERLEESDKEM-FVQHEH-----AVAKFCD-LPSKLGRLGRL-----	237
3E5W_A1	PDBID CHAIN SEQUENCE	HLRERLEESDKEM-FVQHEH-----AVAKFCD-LPSKLGRL-----	244
2A50_A1	PDBID CHAIN SEQUENCE	HR1E1MEEVEKGG-CYKQYEA-----AVGRYCDAAAPSKLGHN-----	243
2A52_A1	PDBID CHAIN SEQUENCE	HR1E1MEEVEKGG-CYKQYEA-----AVGRYCDAAAPSKLGHN-----	243
1XM2_A1	PDBID CHAIN SEQUENCE	HR1E1MEEVEKGG-CYKQYEA-----AVGRYCDAAAPSKLGHN-----	243
2A54_C1	PDBID CHAIN SEQUENCE	HR1E1MEEVEKGG-CYKQYEA-----AVGRYCDAAAPSKLGHN-----	243
2A53_A1	PDBID CHAIN SEQUENCE	HR1E1MEEVEKGG-CYKQYEA-----AVGRYCDAAAPSKLGHN-----	243
2A56_A1	PDBID CHAIN SEQUENCE	HR1E1MEEVEKGG-CYKQYEA-----AVGRYCDAAAPSKLGHN-----	243
1XQ0_A1	PDBID CHAIN SEQUENCE	HR1E1MEEVEKGG-CYKQYEA-----AVGRYCDAAAPSKLGHN-----	232
3CFE_L1	PDBID CHAIN SEQUENCE	HR1EVK-EVQGG-HYEQYEA-----AVARYCDAAAPSKLGH-----	231
3CFH_S1	PDBID CHAIN SEQUENCE	HR1EVK-EVQGG-HYEQYEA-----AVARYCDAAAPSKLGH-----	231
3CFA_L1	PDBID CHAIN SEQUENCE	HR1EVK-EVQGG-HYEQYEA-----AVARYCDAAAPSKLGH-----	231
1Z7W_A1	PDBID CHAIN SEQUENCE	IRLQML-RKKKDE-YFELYEA-----SVARYSD-LPEKAN-----	225
2185_A1	PDBID CHAIN SEQUENCE	HRPE1VKVSENGT-LFEQHE-----SVARYCQTCPKSLGH-----	235
2186_A1	PDBID CHAIN SEQUENCE	HRPE1VKVSENGT-LFEQHE-----SVARYCQTCPKSLGH-----	235
2C91_A1	PDBID CHAIN SEQUENCE	LRMEKLNISDDWK-TVEQHE-----VWASYSQ-VPSKLGHN-----	228
167K_A1	PDBID CHAIN SEQUENCE	SKLDITSHNEDYT-IVEQYER-----TEGRHHLFL-----	226
1ZG0_A1	PDBID CHAIN SEQUENCE	SKLDITSHNEDYT-IVEQYER-----TEGRHHLFL-----	225
16X0_A1	PDBID CHAIN SEQUENCE	SKLDITSHNEDYT-IVEQYER-----TEGRHHLFL-----	225
1ZG0_H1	PDBID CHAIN SEQUENCE	SKLDITSHNEDYT-IVEQYER-----TEGRHHLFL-----	225
1ZG0_P1	PDBID CHAIN SEQUENCE	SKLDITSHNEDYT-IVEQYER-----TEGRHHLFL-----	225
2H8Q_A1	PDBID CHAIN SEQUENCE	SKLDITSHNEDYT-IVEQYER-----TEGRHHLFL-----	219
2VAE_A1	PDBID CHAIN SEQUENCE	SKLDITSHNEDYT-IVEQYER-----AEGRHHLFL-----	225
2V4E_A1	PDBID CHAIN SEQUENCE	SKLDITSHNEDYT-IVEQYER-----TEARHHLFL-----	220
2VAD_A1	PDBID CHAIN SEQUENCE	SKLDITSHNEDYT-IVEQYER-----AEGRHSGSQ-----	225
2H5P_A1	PDBID CHAIN SEQUENCE	IKLDITSHNEDYT-IVELYER-----AEGRHSTGMDELYK-----	236
2H5R_A1	PDBID CHAIN SEQUENCE	IKLDITSHNEDYT-IVELYER-----AEGRHSTGMDELYK-----	236
2H50_A1	PDBID CHAIN SEQUENCE	IKLDITSHNEDYT-IVEQYER-----AEGRHSTGMDELYK-----	236
2QLG_A1	PDBID CHAIN SEQUENCE	IKLDITSHNEDYT-IVEQYER-----AEGRHSTGA-----	226
2QLL_A1	PDBID CHAIN SEQUENCE	IKLDITSHNEDYT-IVEQYER-----AEGRHSTGA-----	226
2QLH_A1	PDBID CHAIN SEQUENCE	IKLDITSHNEDYT-IVEQYER-----AEGRHSTGA-----	226
2H5Q_A1	PDBID CHAIN SEQUENCE	IKLDITSHNEDYT-IVEQYER-----AEGRHSTGMDELYK-----	236
3KCS_A1	PDBID CHAIN SEQUENCE	RKLDITSHNEDYT-IVEQYER-----AEGRHSTGMDELYK-----	236
3KCT_A1	PDBID CHAIN SEQUENCE	RKLDITSHNEDYT-IVEQYER-----AEGRHSTGMDELYK-----	236
3LF4_A1	PDBID CHAIN SEQUENCE	IKLDITSHNEDYT-IVEQYER-----SEGRHSTGMDELYK-----	231
3LF3_A1	PDBID CHAIN SEQUENCE	IKLDITSHNEDYT-IVEQYER-----AEGRHSTGMDELYK-----	231
2P4M_A1	PDBID CHAIN SEQUENCE	RKLDVTHNKKDYT-SVEQCEI-----SIARKPVVA-----	221
2ARL_A1	PDBID CHAIN SEQUENCE	RKLDVTHNKKDYT-SVEQCEI-----SIARKPVVA-----	221
1MOV_A1	PDBID CHAIN SEQUENCE	RKLDVTHNKKDYT-SVEQCEI-----SIARKPVVA-----	221
1M0U_A1	PDBID CHAIN SEQUENCE	RKLDVTHNKKDYT-SVEQCEI-----SIARKPVVA-----	221
2WH5_A1	PDBID CHAIN SEQUENCE	RKLDVTHNKKDYT-SVEQCEI-----SIARKPVVA-----	221
2WHT_A1	PDBID CHAIN SEQUENCE	RKLDVTHNKKDYT-SVEQCEI-----SIARKPVVA-----	221
2WHU_A1	PDBID CHAIN SEQUENCE	RKLDVTHNKKDYT-SVEQCEI-----SIARKPVVA-----	221
31R8_A1	PDBID CHAIN SEQUENCE	RKLDVTHNKKDYT-SVEQCEI-----SIARKPVVA-----	221
3CGL_A1	PDBID CHAIN SEQUENCE	TKLVWVNNKDFM-KVEEHEI-----AVARHPFVPEPKDK-----	243
2262_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	257
210V_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	257
2P0X_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	257
21E2_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	224
2210_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	227
226Y_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	227
2GX2_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	243
2GX0_B1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	243
226X_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	257
31S3_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	237
31SA_A1	PDBID CHAIN SEQUENCE	HHTEIKSHDKDY-SVNLHEH-----AEAS-ELPRAK-----	237
2DDC_A1	PDBID CHAIN SEQUENCE	HLMETTSHDKDY-KVKLYEH-----AKAS-GLPRLAK-----	227
2DD0_A1	PDBID CHAIN SEQUENCE	HLMETTSHDKDY-KVKLYEH-----AKAS-GLPRLAK-----	227
1X5S_A1	PDBID CHAIN SEQUENCE	HR1ETTSHDKDY-KVKLYEH-----AKAS-GLPRLAK-----	227
2VVT_A1	PDBID CHAIN SEQUENCE	HCTEILSHDKDY-KVKLYEH-----AVAS-GLPDNARR-----	228
2VWH_A1	PDBID CHAIN SEQUENCE	HCTEILSHDKDY-KVKLYEH-----AVAS-GLPDNARR-----	228
2VJ3_A1	PDBID CHAIN SEQUENCE	HCTEILSHDKDY-KVKLYEH-----AVAS-GLPDNARR-----	228
1ZUX_A1	PDBID CHAIN SEQUENCE	HCTEILSHDKDY-KVKLYEH-----AVAS-GLPDNARR-----	226
2BT3_A1	PDBID CHAIN SEQUENCE	HCTEILSHDKDY-KVKLYEH-----AVAS-GLPD-----	222
2GW3_A1	PDBID CHAIN SEQUENCE	HCTEILSHDKDY-KVKLYEH-----AVAS-GLPDNARR-----	227
2G14_A1	PDBID CHAIN SEQUENCE	HCTEILSHDKDY-KVKLYEH-----AVAS-GLPDNARR-----	227
20TE_A1	PDBID CHAIN SEQUENCE	HR1EILNHDKDY-KVTVYES-----AVARY-----	216
20TB_A1	PDBID CHAIN SEQUENCE	HR1EILNHDKDY-KVTVYES-----AVARY-----	216
2HQK_A1	PDBID CHAIN SEQUENCE	HR1EILNHDKDY-KVTVYES-----AVARNSTDGM-----	221
2VZK_A1	PDBID CHAIN SEQUENCE	HR1EILNHDKDY-KVKLYEH-----AVARYSPLPSQVHHHHHH-----	231
3ADF_A1	PDBID CHAIN SEQUENCE	HR1EILNHDKDY-KVKLYEH-----AVARYSMLPSQAK-----	228
21CR_A1	PDBID CHAIN SEQUENCE	HKLNRDRSDAKNKKWQLTEH-----ATASRSALP-----	239
2FL1_A1	PDBID CHAIN SEQUENCE	HKLNRDRSDAKNKKWQLTEH-----ATASRSALP-----	228
20GR_A1	PDBID CHAIN SEQUENCE	HKLNRDRSDAKNKKWQLTEH-----ATAFPSALA-----	231
1XAE_A1	PDBID CHAIN SEQUENCE	HKLNRDRSDAKNKKWQLTEH-----ATAFPSALA-----	231
1XA9_A1	PDBID CHAIN SEQUENCE	HKLNRDRSDAKNKKWQLTEH-----ATAFPSALA-----	231
2PX5_A1	PDBID CHAIN SEQUENCE	HKLNRDRSDAKNKKWHLTEH-----ATASGSALP-----	228
2PXW_A1	PDBID CHAIN SEQUENCE	HKLNRDRSDAKNKKWHLTEH-----ATASGSALP-----	228
20JK_A1	PDBID CHAIN SEQUENCE	HKLNRDRSDAKNKKWHLTEH-----ATASGSALP-----	231
2RH7_A1	PDBID CHAIN SEQUENCE	HLR1VKTNDTASG-YVQHE-----ATAAHTIKKIEGSLPDVHHHHHH-----	241
2C93_H1	PDBID CHAIN SEQUENCE	HLR1VKTNDTASG-YVQHE-----ATAAHTIKKIEGSLPDVHHHHHH-----	225
2G6X_A1	PDBID CHAIN SEQUENCE	RRVEEDHSNTELG-IVEYQH-----AFKTPD-----	219
2G6Y_A1	PDBID CHAIN SEQUENCE	RRVEEDHSNTELG-IVEYQH-----AFKTPD-----	219
2G30_A1	PDBID CHAIN SEQUENCE	RRVEELHSNTELG-IVEYQH-----AFKTPDIAFA-----	222
2DD7_A1	PDBID CHAIN SEQUENCE	RRVEETHKENLA-MVEYQ-----VFNSAPRDM-----	218
2DD9_D1	PDBID CHAIN SEQUENCE	RRVEETHKENLA-MVEYQ-----VFNSAPRDM-----	218

3EKJ_A | PDBID | CHAIN | SEQUENCE EMIREAGIDGGQVNYEEFVQMMTAK 451

3EVP_A | PDBID | CHAIN | SEQUENCE EMIREADIDGGQVNYEEFVQMMTAK 451
3EKH_A | PDBID | CHAIN | SEQUENCE EMIREADIDGGQVNYEEFVQMMTAK- 413
3EVR_A | PDBID | CHAIN | SEQUENCE EMIREADIDGGQVNYEEFVQMMTAK 451
3EK7_A | PDBID | CHAIN | SEQUENCE EMIREADIDGGQVNYEEFVQMMTAK 451
3EVU_A | PDBID | CHAIN | SEQUENCE EMIREADIDGGQVNYEEFVQMMTAK 451
3EUV_A | PDBID | CHAIN | SEQUENCE EMIREADIDGGQVNYEEFVQMMTAK 451
3EK4_A | PDBID | CHAIN | SEQUENCE EMIREADIDGGQVNYEEFVQMMTAK 451
3EK8_A | PDBID | CHAIN | SEQUENCE EMIREADIDGGQVNYEEFVQMMTAK 451

2G3D_A | PDBID | CHAIN | SEQUENCE -----
2G2S_A | PDBID | CHAIN | SEQUENCE -----
1QYQ_A | PDBID | CHAIN | SEQUENCE -----
2G6E_A | PDBID | CHAIN | SEQUENCE -----
1YHH_A | PDBID | CHAIN | SEQUENCE -----
1YHG_B | PDBID | CHAIN | SEQUENCE -----
2G5Z_A | PDBID | CHAIN | SEQUENCE -----
2G16_A | PDBID | CHAIN | SEQUENCE -----
2HGD_A | PDBID | CHAIN | SEQUENCE -----
2HGY_A | PDBID | CHAIN | SEQUENCE -----
1YJF_A | PDBID | CHAIN | SEQUENCE -----
1YHI_A | PDBID | CHAIN | SEQUENCE -----
1YJZ_A | PDBID | CHAIN | SEQUENCE -----
1QYF_A | PDBID | CHAIN | SEQUENCE -----
1QXT_A | PDBID | CHAIN | SEQUENCE -----
2AWM_A | PDBID | CHAIN | SEQUENCE -----
2AWK_A | PDBID | CHAIN | SEQUENCE -----
2AWJ_A | PDBID | CHAIN | SEQUENCE -----
2AWL_A | PDBID | CHAIN | SEQUENCE -----
2HFC_A | PDBID | CHAIN | SEQUENCE -----
2HCG_A | PDBID | CHAIN | SEQUENCE -----
2QZB_A | PDBID | CHAIN | SEQUENCE -----
2FZU_A | PDBID | CHAIN | SEQUENCE -----
1K1R_A | PDBID | CHAIN | SEQUENCE -----
1K1P_A | PDBID | CHAIN | SEQUENCE -----
1K1S_A | PDBID | CHAIN | SEQUENCE -----
2FWQ_A | PDBID | CHAIN | SEQUENCE -----
1B9C_D | PDBID | CHAIN | SEQUENCE -----
1MW_A | PDBID | CHAIN | SEQUENCE -----
3ED8_A | PDBID | CHAIN | SEQUENCE -----
1QY0_A | PDBID | CHAIN | SEQUENCE -----
2QU1_A | PDBID | CHAIN | SEQUENCE -----
1QY3_A | PDBID | CHAIN | SEQUENCE -----
2QTZ_A | PDBID | CHAIN | SEQUENCE -----
2QRF_A | PDBID | CHAIN | SEQUENCE -----
2HRS_A | PDBID | CHAIN | SEQUENCE -----
2B3Q_D | PDBID | CHAIN | SEQUENCE -----
2B3P_A | PDBID | CHAIN | SEQUENCE -----
1KPS_A | PDBID | CHAIN | SEQUENCE -----
3G1L_A | PDBID | CHAIN | SEQUENCE -----
3GJZ_A | PDBID | CHAIN | SEQUENCE -----
3G9A_A | PDBID | CHAIN | SEQUENCE -----
3K1K_B | PDBID | CHAIN | SEQUENCE -----
1JCO_A | PDBID | CHAIN | SEQUENCE -----
1JCL_A | PDBID | CHAIN | SEQUENCE -----
2A8B_B | PDBID | CHAIN | SEQUENCE -----
2AHA_A | PDBID | CHAIN | SEQUENCE -----
3CD1_A | PDBID | CHAIN | SEQUENCE -----
3CD9_A | PDBID | CHAIN | SEQUENCE -----
3CB9_A | PDBID | CHAIN | SEQUENCE -----
3CBE_A | PDBID | CHAIN | SEQUENCE -----
2OKW_F | PDBID | CHAIN | SEQUENCE -----
2OKY_B | PDBID | CHAIN | SEQUENCE -----
2HOZ_A | PDBID | CHAIN | SEQUENCE -----
2HJO_A | PDBID | CHAIN | SEQUENCE -----
2O2B_A | PDBID | CHAIN | SEQUENCE -----
2H9W_A | PDBID | CHAIN | SEQUENCE -----
2O24_A | PDBID | CHAIN | SEQUENCE -----
2O29_A | PDBID | CHAIN | SEQUENCE -----
2H6V_A | PDBID | CHAIN | SEQUENCE -----
1RMO_A | PDBID | CHAIN | SEQUENCE -----
1RMM_A | PDBID | CHAIN | SEQUENCE -----
1RMP_A | PDBID | CHAIN | SEQUENCE -----
1RRX_A | PDBID | CHAIN | SEQUENCE -----
2Q6P_A | PDBID | CHAIN | SEQUENCE -----
1Z1Q_A | PDBID | CHAIN | SEQUENCE -----
1S6Z_A | PDBID | CHAIN | SEQUENCE -----
1Z1P_A | PDBID | CHAIN | SEQUENCE -----
1F0B_A | PDBID | CHAIN | SEQUENCE -----
1F09_A | PDBID | CHAIN | SEQUENCE -----
2YFP_A | PDBID | CHAIN | SEQUENCE -----
3D08_A | PDBID | CHAIN | SEQUENCE -----
3DQ1_A | PDBID | CHAIN | SEQUENCE -----
3DQ9_A | PDBID | CHAIN | SEQUENCE -----
3DQA_A | PDBID | CHAIN | SEQUENCE -----
3DQC_A | PDBID | CHAIN | SEQUENCE -----
3DQD_A | PDBID | CHAIN | SEQUENCE -----
3DQE_A | PDBID | CHAIN | SEQUENCE -----
3DQF_A | PDBID | CHAIN | SEQUENCE -----
3DQH_A | PDBID | CHAIN | SEQUENCE -----
3DQI_A | PDBID | CHAIN | SEQUENCE -----
3DQJ_A | PDBID | CHAIN | SEQUENCE -----
3DQK_A | PDBID | CHAIN | SEQUENCE -----
3DQL_A | PDBID | CHAIN | SEQUENCE -----
3DQM_A | PDBID | CHAIN | SEQUENCE -----
3DQN_A | PDBID | CHAIN | SEQUENCE -----
3DQO_A | PDBID | CHAIN | SEQUENCE -----
3DQU_A | PDBID | CHAIN | SEQUENCE -----
3DQ4_A | PDBID | CHAIN | SEQUENCE -----
3DQ5_A | PDBID | CHAIN | SEQUENCE -----
3DQ6_A | PDBID | CHAIN | SEQUENCE -----
3DQ7_A | PDBID | CHAIN | SEQUENCE -----
3DPW_A | PDBID | CHAIN | SEQUENCE -----
3DPX_A | PDBID | CHAIN | SEQUENCE -----
3DPZ_A | PDBID | CHAIN | SEQUENCE -----
3DQ2_A | PDBID | CHAIN | SEQUENCE -----

3DQ3_A | PDBID | CHAIN | SEQUENCE -----
1HU7_A | PDBID | CHAIN | SEQUENCE -----
1YFP_A | PDBID | CHAIN | SEQUENCE -----
2JAD_A | PDBID | CHAIN | SEQUENCE -----
1H6R_A | PDBID | CHAIN | SEQUENCE -----
1C4F_A | PDBID | CHAIN | SEQUENCE -----
1EMB_A | PDBID | CHAIN | SEQUENCE -----
1Q4A_A | PDBID | CHAIN | SEQUENCE -----
1EMA_A | PDBID | CHAIN | SEQUENCE -----
1Q4B_A | PDBID | CHAIN | SEQUENCE -----
1EMG_A | PDBID | CHAIN | SEQUENCE -----
2DUF_A | PDBID | CHAIN | SEQUENCE -----
2DUJ_A | PDBID | CHAIN | SEQUENCE -----
2DUJ_A | PDBID | CHAIN | SEQUENCE -----
2DUG_A | PDBID | CHAIN | SEQUENCE -----
2DUH_A | PDBID | CHAIN | SEQUENCE -----
1Q4E_A | PDBID | CHAIN | SEQUENCE -----
1Q73_A | PDBID | CHAIN | SEQUENCE -----
1Q4D_A | PDBID | CHAIN | SEQUENCE -----
1Q4C_A | PDBID | CHAIN | SEQUENCE -----
1J8Z_A | PDBID | CHAIN | SEQUENCE -----
1JBY_A | PDBID | CHAIN | SEQUENCE -----
1W75_D | PDBID | CHAIN | SEQUENCE -----
1W7U_D | PDBID | CHAIN | SEQUENCE -----
1W7T_D | PDBID | CHAIN | SEQUENCE -----
1HCJ_A | PDBID | CHAIN | SEQUENCE -----
2QLE_A | PDBID | CHAIN | SEQUENCE -----
1BFP_A | PDBID | CHAIN | SEQUENCE -----
2EMQ_A | PDBID | CHAIN | SEQUENCE -----
1EMF_A | PDBID | CHAIN | SEQUENCE -----
2EMD_A | PDBID | CHAIN | SEQUENCE -----
2EMN_A | PDBID | CHAIN | SEQUENCE -----
1EMM_A | PDBID | CHAIN | SEQUENCE -----
1EMC_A | PDBID | CHAIN | SEQUENCE -----
1EML_A | PDBID | CHAIN | SEQUENCE -----
1EME_A | PDBID | CHAIN | SEQUENCE -----
1EMK_A | PDBID | CHAIN | SEQUENCE -----
2WUR_A | PDBID | CHAIN | SEQUENCE -----
1GFL_A | PDBID | CHAIN | SEQUENCE -----
1RM9_A | PDBID | CHAIN | SEQUENCE -----
2WSN_A | PDBID | CHAIN | SEQUENCE -----
2Q57_A | PDBID | CHAIN | SEQUENCE -----
2WSO_A | PDBID | CHAIN | SEQUENCE -----
1OXF_A | PDBID | CHAIN | SEQUENCE -----
1CV7_A | PDBID | CHAIN | SEQUENCE -----
3GEX_A | PDBID | CHAIN | SEQUENCE -----
1OXD_A | PDBID | CHAIN | SEQUENCE -----
1OXE_A | PDBID | CHAIN | SEQUENCE -----
3LVC_A | PDBID | CHAIN | SEQUENCE -----
3LVD_A | PDBID | CHAIN | SEQUENCE -----
3LVA_A | PDBID | CHAIN | SEQUENCE -----
2HPW_A | PDBID | CHAIN | SEQUENCE -----
2W1Q_A | PDBID | CHAIN | SEQUENCE -----
3A8S_A | PDBID | CHAIN | SEQUENCE -----
2W1S_A | PDBID | CHAIN | SEQUENCE -----
3GB3_A | PDBID | CHAIN | SEQUENCE -----
3GL4_A | PDBID | CHAIN | SEQUENCE -----
2A46_A | PDBID | CHAIN | SEQUENCE -----
2A48_A | PDBID | CHAIN | SEQUENCE -----
2A47_A | PDBID | CHAIN | SEQUENCE -----
2ZMW_A | PDBID | CHAIN | SEQUENCE -----
2ZML_A | PDBID | CHAIN | SEQUENCE -----
2Z06_A | PDBID | CHAIN | SEQUENCE -----
2Z07_A | PDBID | CHAIN | SEQUENCE -----
3BX9_B | PDBID | CHAIN | SEQUENCE -----
3BXA_A | PDBID | CHAIN | SEQUENCE -----
3BXB_A | PDBID | CHAIN | SEQUENCE -----
3BXC_A | PDBID | CHAIN | SEQUENCE -----
3H1O_A | PDBID | CHAIN | SEQUENCE -----
3H1R_A | PDBID | CHAIN | SEQUENCE -----
3M22_A | PDBID | CHAIN | SEQUENCE -----
3M24_A | PDBID | CHAIN | SEQUENCE -----
3IP2_A | PDBID | CHAIN | SEQUENCE -----
3EST_A | PDBID | CHAIN | SEQUENCE -----
3ESV_A | PDBID | CHAIN | SEQUENCE -----
1UIS_B | PDBID | CHAIN | SEQUENCE -----
3E5W_A | PDBID | CHAIN | SEQUENCE -----
2A50_A | PDBID | CHAIN | SEQUENCE -----
2A52_A | PDBID | CHAIN | SEQUENCE -----
1XMZ_A | PDBID | CHAIN | SEQUENCE -----
2A54_C | PDBID | CHAIN | SEQUENCE -----
2A53_A | PDBID | CHAIN | SEQUENCE -----
2A56_A | PDBID | CHAIN | SEQUENCE -----
1XQM_A | PDBID | CHAIN | SEQUENCE -----
3CFE_L | PDBID | CHAIN | SEQUENCE -----
3CFH_S | PDBID | CHAIN | SEQUENCE -----
3CFA_L | PDBID | CHAIN | SEQUENCE -----
1YZW_A | PDBID | CHAIN | SEQUENCE -----
2IB5_A | PDBID | CHAIN | SEQUENCE -----
2IB6_A | PDBID | CHAIN | SEQUENCE -----
2C9T_A | PDBID | CHAIN | SEQUENCE -----
1G7K_A | PDBID | CHAIN | SEQUENCE -----
1ZG0_A | PDBID | CHAIN | SEQUENCE -----
1GGX_A | PDBID | CHAIN | SEQUENCE -----
1ZGQ_H | PDBID | CHAIN | SEQUENCE -----
1ZGP_A | PDBID | CHAIN | SEQUENCE -----
2HRQ_A | PDBID | CHAIN | SEQUENCE -----
2VAE_A | PDBID | CHAIN | SEQUENCE -----
2V4E_A | PDBID | CHAIN | SEQUENCE -----
2VAD_A | PDBID | CHAIN | SEQUENCE -----
2H5P_A | PDBID | CHAIN | SEQUENCE -----
2H5R_A | PDBID | CHAIN | SEQUENCE -----
2H5O_A | PDBID | CHAIN | SEQUENCE -----
2QLG_A | PDBID | CHAIN | SEQUENCE -----
2QLL_A | PDBID | CHAIN | SEQUENCE -----

2QLH_A | PDBID | CHAIN | SEQUENCE -----
2H5Q_A | PDBID | CHAIN | SEQUENCE -----
3KCS_A | PDBID | CHAIN | SEQUENCE -----
3KCT_A | PDBID | CHAIN | SEQUENCE -----
3LF3_A | PDBID | CHAIN | SEQUENCE -----
3LF4_A | PDBID | CHAIN | SEQUENCE -----
2P4M_A | PDBID | CHAIN | SEQUENCE -----
2ARL_A | PDBID | CHAIN | SEQUENCE -----
1MOV_A | PDBID | CHAIN | SEQUENCE -----
1MOL_A | PDBID | CHAIN | SEQUENCE -----
2WHS_A | PDBID | CHAIN | SEQUENCE -----
2WHT_A | PDBID | CHAIN | SEQUENCE -----
2WHU_A | PDBID | CHAIN | SEQUENCE -----
3IR8_A | PDBID | CHAIN | SEQUENCE -----
3CGL_A | PDBID | CHAIN | SEQUENCE -----
2Z6Z_A | PDBID | CHAIN | SEQUENCE -----
2TOV_A | PDBID | CHAIN | SEQUENCE -----
2POX_A | PDBID | CHAIN | SEQUENCE -----
2IE2_A | PDBID | CHAIN | SEQUENCE -----
2Z1O_A | PDBID | CHAIN | SEQUENCE -----
2Z6Y_A | PDBID | CHAIN | SEQUENCE -----
2GXZ_A | PDBID | CHAIN | SEQUENCE -----
2GX8_B | PDBID | CHAIN | SEQUENCE -----
2Z6X_A | PDBID | CHAIN | SEQUENCE -----
3LS3_A | PDBID | CHAIN | SEQUENCE -----
3LSA_A | PDBID | CHAIN | SEQUENCE -----
2DDC_A | PDBID | CHAIN | SEQUENCE -----
2DDD_A | PDBID | CHAIN | SEQUENCE -----
1XSS_A | PDBID | CHAIN | SEQUENCE -----
2VVT_A | PDBID | CHAIN | SEQUENCE -----
2VVH_A | PDBID | CHAIN | SEQUENCE -----
2VVJ_A | PDBID | CHAIN | SEQUENCE -----
1ZUK_A | PDBID | CHAIN | SEQUENCE -----
2BTJ_A | PDBID | CHAIN | SEQUENCE -----
2GW3_A | PDBID | CHAIN | SEQUENCE -----
2GW4_A | PDBID | CHAIN | SEQUENCE -----
2OTE_A | PDBID | CHAIN | SEQUENCE -----
2OTB_A | PDBID | CHAIN | SEQUENCE -----
2HQK_A | PDBID | CHAIN | SEQUENCE -----
2VZX_A | PDBID | CHAIN | SEQUENCE -----
3ADF_A | PDBID | CHAIN | SEQUENCE -----
2ICR_A | PDBID | CHAIN | SEQUENCE -----
2FL1_A | PDBID | CHAIN | SEQUENCE -----
2OGR_A | PDBID | CHAIN | SEQUENCE -----
1XAE_A | PDBID | CHAIN | SEQUENCE -----
1XA9_A | PDBID | CHAIN | SEQUENCE -----
2PXS_A | PDBID | CHAIN | SEQUENCE -----
2PXW_A | PDBID | CHAIN | SEQUENCE -----
2OJK_A | PDBID | CHAIN | SEQUENCE -----
2RH7_A | PDBID | CHAIN | SEQUENCE -----
2C9J_H | PDBID | CHAIN | SEQUENCE -----
2G6X_A | PDBID | CHAIN | SEQUENCE -----
2G6Y_A | PDBID | CHAIN | SEQUENCE -----
2G3O_A | PDBID | CHAIN | SEQUENCE -----
2DD7_A | PDBID | CHAIN | SEQUENCE -----
2DD9_D | PDBID | CHAIN | SEQUENCE -----

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

CLUSTAL 2.0.12 multiple sequence alignment

```
Q17106/5-229      --EELFTGVVPIILVELDGDVNGQKFSVRGEGEDATYGLTLKFICTTG----KLVPWPW 54
Q17105/5-229      --EELFTGVVPIILVELDGDVNGQKFSVSGEGEDATYGLTLNFICTTG----KLVPWPW 54
P42212/5-229      --EELFTGVVPIILVELDGDVNGHKFSVSGEGEDATYGLTLKFICTTG----KLVPWPW 54
Q6YG20/5-229      --AELFTGVVPIILVELDGDVNGHKFSVSGEGEDATYGLTLKFICTTG----KLVPWPW 54
Q8WT5/5-229       --EELFTGVVPIILVELDGDVHGHKFSVRGEGEDADYGLKLEIKFICTTG----KLVPWPW 54
Q8WT4/5-229       --EELFTGVVPIILVELDGDVHGHKFSVRGEGEDADYGLKLEIKFICTTG----KLVPWPW 54
Q8WT9/5-229       --EELFTGIVPVLIELDGDVHGHKFSVRGEGEDADYGLKLEIKFICTTG----KLVPWPW 54
Q8WT0/5-229       --EELFTGIVPVLIELDGDVHGHKFSVRGEGEDADYGLKLEIKFICTTG----KLVPWPW 54
Q8WT8/5-229       --EELFTGIVPVLIELDGDVHGHKFSVRGEGEDADYGLKLEIKFICTTG----KLVPWPW 54
Q8WT6/5-229       --EELFTGIVPVLIELDGDVHGHKFSVRGEGEDADYGLKLEIKFICTTG----KLVPWPW 54
Q8WP95/5-229      --EELFTGIVPVLIELDGDVHGHKFSVRGEGEDADYGLKLEIKFICTTG----KLVPWPW 54
Q8WT7/5-229       --EELFTGIVPVLIELDGDVHGHKFSVRGEGEDADYGLKLEIKFICTTG----KLVPWPW 54
Q6R57/5-229       --ALLFHGKIPYVVEGMEGNDVGHFTFIRGKYGDASVGVKVDQFICTTG----DVVPWWS 54
Q6R56/39-258      --TETFQKLPYKLELDGDVGGQTFKVIKGEVGDATGVIEGKYVCTEG----EVPISWV 54
Q6R55/2-222       --AEYFEKLPYKVELDGDVGGQTFKVIKGEVGDASTGRVEGKYVCTEG----EVPISWV 54
Q6R54/5-225       --PALFQSDMTFKIFIDGVVNDQKFTIADGSSKFPHGDFNVHVCETG----KLPMWPK 54
Q2TCH5/5-225      --PALFQSDMTFKIFIDGVEVNGQKFTIVADGSSKFPHGDFNVHVCETG----KLPMWPK 54
Q53UG7/1-221      --MSVITSEMKLRLMEGAVNGHKFVITGKGSQPFEGIQN--MDLTVIE--GGPLPFAFD 55
Q53UG8/1-221      --MSVITSEMKIQLRMEGAVNGHKFVITGKGSQPFEGIQN--VDLTVIE--GGPLPFAFD 55
Q8MU48/1-218      --MSMIKPEMKIKMRMDGAVNGHKFVITGEGSGEPFEGKQT--MNLTVID--GGPLPFAFD 55
A8CLW1/1-218      --MNVIKPDMKIKLRMEGAVNGHKFAIEGEGNGQPFEGKQT--MNLKVKE--GGPLPFAFD 55
B6CT24/1-218      --MSVIKPDMKIKLRMEGAVNGHKFAIEGEGKQPFEGKQT--MNLKVKE--GGPLPFAFD 55
Q5TL66/1-218      --MSVIKPDMKIKLRMEGAVNGHKFAIEGVGLGKPFEGKQS--MDLKVKE--GGPLPFAFD 55
A8CLQ1/1-223      --MSVINPDMKIKLYMEGAVNGHKFIEKGEKNGKPFEGKQT--MDLAVVD--GGPLPFAFD 55
Q2LI65/1-219      --MSAIKPDMLIKLRMEGAVNGHTFEIEGEGKGPFGKQK--MDLKVVD--GGPLPFAFD 55
A8CLW4/1-222      --MNVIKPDMKIKLRMEGAVNGHKFVIEGKGDGKPYEGTQT--MDLEVIE--GGPLPFAFD 55
B6CT25/4-224      --EDVIKRDMLIKLRMEGAVNGHKFVIEGNGDQKPYEGTQT--IDLEVIE--GAPLPFAFD 55
A3F207/1-221      --MSVKSVMNVKLRLELGAVNGHFPVIEGKGNHGFEGTQD--IKLTVKE--GGPLPFAFD 55
Q962P9/1-221      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGTQS--MDLTVKE--GAPLPFAFD 55
Q7Z0W8/1-221      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGTQS--MDLTVKE--GAPLPFAFD 55
Q66ND5/1-221      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGKQT--MDLTVIE--GAPLPFAFD 55
Q66ND4/1-221      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGKQT--MDLTVIE--GAPLPFAFD 55
Q66ND2/1-221      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGKQT--MDLTVIE--GAPLPFAFD 55
A7UAL7/1-221      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGKQT--MDLTVIE--GAPLPFAFD 55
Q7Z0W6/1-221      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGKQT--MDLTVIE--GAPLPFAFD 55
A7UAL9/1-221      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGKQT--MDLTVIE--GAPLPFAFD 55
Q8T5F2/8-228      --KGVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGTQT--MDLTVIE--GAPLPFAFD 55
Q7Z0W7/8-228      --KGVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGTQT--MDLTVIE--GAPLPFAFD 55
Q8MU47/8-228      --KGVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGTQT--MDLTVIE--GAPLPFAFD 55
Q95V07/1-221      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGTQS--MDLTVIE--GAPLPFAFD 55
A7UAL3/1-221      --MSVIKTDMLIKLRMEGAVNGHKFVIEGEGKPFEGKQT--MDLTVIE--GAPLPFAFD 55
A7UAL6/1-218      --MSVIKTDMLIKLRMEGAVNGHKFVIEGEGKPFEGKQT--MDLTVIE--GAPLPFAFD 55
A7UAL8/1-218      --MSVIKPDMKIKLRMEGAVNGHKFVIEGDKGKPFEGKQT--MDLTVIE--GAPLPFAFD 55
A8CL52/1-219      --MSLINPEMKIKLLEMEGAVNGHFPVIEGDKGHPFEGKQS--MDLVKE--GAPLPFAFD 55
Q8I638/1-219      --MSLIKPEMKIKLLEMEGAVNGHFPVIEGDKGHPFEGKQS--MDLVKE--GAPLPFAFD 55
A3F208/1-219      --MSVIKEMKIKLRMEGAVNGHAFVIEGNGNHPFEGKQS--MDLEVKK--GAPLPFAFD 55
A8CLR7/1-219      --MSVIKPDMKIKLRMEGAVNGHFPVIEGDKGRPFEGKQS--MDLDVKE--GGPLPFAFD 55
Q5S6Z9/1-219      --MSAIKPDMLIKLRMEGAVNGHFPVIEGDKGKPFEGKQS--MDLEVKE--GGPLPFAFD 55
Q66PW1/5-223      --KTVVQFMKIKMSLEGTVNGHYFKIVGEGDTPFEGKQT--LHLKVKE--GAPLPFAFD 55
Q8T5F1/1-219      --MSVKSVMKIKLRMEGAVNGHNFVIVGEGEKPYEGTQS--MDLTVKE--GAPLPFAFD 55
Q7Z0W9/1-219      --MSVKSVMKIKLRMEGAVNGHNFVIVGEGEKPYEGTQS--MDLTVKE--GAPLPFAFD 55
Q6USK3/1-219      --MSVKSVMKIKRRMEGAVNGHNFVIVGEGEKPYEGTQS--MDLTVKE--GAPLPFAFD 55
A7UAL4/1-219      --MSVIKPDMKIKLRMEGAVNGHFPVIEGEGKPFEGTQS--MDLTVKE--GAPLPFAFD 55
A7UAM0/1-219      --MSVKSVMKIKLRMEGAVNGHNFVIVGEGEKPYEGTQS--MDLTVKE--GGPLPFAFD 55
Q7Z0W5/1-218      --MSVKSVMKIKLHMDGIVNGHKFMITGEGEKPFEGTHT--IILKVKE--GGPLPFAFD 55
Q95UA7/1-218      --MSVKSVMKIKLRMDGIVNGHKFMITGEGEKPFEGTHT--IILKVKE--GGPLPFAFD 55
A7UAL1/1-218      --MSVKSVMKIKLRMDGIVNGHKFMITGEGEKPFEGTHT--IILKVKE--GGPLPFAFD 55
A8CLP7/1-218      --MSVIKTDMLIKLRQMGAVNGHFPVITGEGEKPYEGKHT--INLTVQD--GGPLPFAFD 55
A7UAL2/1-218      --MSVKSVMKIKLRMDGIVNGHKFMITGEGEKPFEGTHT--IILKVKE--GGPLPFAFD 55
A7UAL5/1-218      --MSVIKPDMKIKLRMQGVVNGHKFVIEGEGKPFEGTQT--INLTVKE--GAPLPFAFD 55
Q963F5/1-218      --MSVIKPDMKIKLRMQGVVNGHKFVIEGEGKPFEGTQT--INLTVKE--GAPLPFAFD 55
Q7Z0W4/1-218      --MSVIKPDMKIKLRMEGAVNGHNFVIEGEGKPFEGTQT--INLTVKE--GGPLPFAFD 55
Q66ND3/1-218      --MSVIKPDMKIKLRMEGAVNGHNFVIEGEGKPFEGTQT--INLTVKE--GGPLPFAFD 55
Q2LI64/1-35       --MSVMKSEMKIKLRMEGAVNGHTFEIEGEGKGPFE----- 35
Q60I25/1-218      --MSVIKPEMKIKLCMRGTVNGHNFVIEGEGKPNPYEGTQI--LDLNVTE--GAPLPFAFD 55
Q60I24/1-218      --MSVIKPEMKIKLCMRGTVNGHNFVIEGEGKPNPYEGTQI--LDLNVTE--GAPLPFAFD 55
A8CLT2/1-218      --MNVIKPDMKIKLCMRGTINGHNFVIEGEGKPNPYEGTQI--LDLNVTE--GAPLPFAFD 55
B5BR3/1-218       --MSVIKQEMKIKLHMDGAVNGHAFVIEGDKGKPYDGTQT--LNLVKE--GAPLPFSYD 55
B6CT26/1-219      --MSVIKQEMKIKLHMDGAVNGHAFVIEGDKGKPYDGTQT--LNLVKE--GAPLPFSYD 55
Q9U6Y3/43-260     --MGVIKPDMKIKLRMEGAVNGHAFVIEGEGKPYDGTHT--LNLVKE--GAPLPFSYD 55
Q8T6U0/1-218      --MNLIKEDMRVVKHMEGAVNGHAFVIEGEGKPYDGTQT--LNLVKE--GAPLPFSYD 55
Q8T5F0/7-229      --KNVVKDFMKITLRMDGAVNGKFAVNGTGDGPNYGGIQS--LKLTVDG--NKPLPFAFD 55
Q8T5F3/6-223      --MKVKEHMKIKLRMGVTVNGKFAVNGTGDGYPYQKQI--LKLIVEG--SEPLPFAFD 54
Q66PW0/1-221      --MNALQEMKIKLTMVGVVNGQSFKIDGKGGKPYEGSQE--LTLKVVE--GGPLFSYD 55
Q8MU46/1-221      --MNALQEMKIKLTMVGVVNGQSFKIDGKGGKPYEGSQE--LTLKVVE--GGPLFSYD 55
Q8T5E9/1-221      --MSALKEEMKIKLTMVGVVNGQSFQIDGEGKGGKPYEGSQE--LTLVVE--GGPLFSYD 55
Q8T5E8/1-221      --MSALKEEMKIXLTLVGVVNGHFPKIDGKGGKPYEGSQE--LTLVVE--GGPLFSYD 55
Q8T6T8/1-221      --MSALKEEMKINLTMVGVVNGLPFKIRGDKGGKPYEGSQE--LTLVVK--GGPLFSYD 55
Q66PU5/1-220      --MSVIVKEMTKLHMEGTVNGHAFVIEGEGKGDYVNGVQS--MNLVKG--GAPLPFSFD 55
A8CL57/1-220      --MSVIVKEMTKLHMEGTVNGHAFVIEGEGKGDYVNGVQS--MNLVKG--GAPLPFSFD 55
```

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

Q8MMA2/34-254 --TNGIAEEMKTRVHLEGTVNGHSFTIKGEGRGPYKGEQF-MSLEVVN--GAPLPFSFD 55
Q8MMA1/12-234 --VNGIAEEMKTDLIMEGIVNGHSFTIEGKGTGYXKGDQF-MKLEVVK--GAPLPFSFD 55
Q66ND6/5-222 --KSVIKEEMLINLHLEGTFNHGYFEIKGKGGKPNNEGTNT-VTLEVTK--GGPLPFGWH 55
Q9U67/5-223 --KSVIKEEMLIDLHLEGTFNHGYFEIKGKGGKPNNEGTNT-VTLEVTK--GGPLPFGWH 55
Q55367/6-223 --EEVIKEFMRFKVMEGVSNGHEFEIEGEGEGRPYEGTQT-ARLKVTK--GGPLPFAWD 55
Q55368/6-223 --EEVIKEFMRFKVMEGVSNGHEFEIEGEGEGRPYEGTQT-AKLKVTK--GGPLPFAWD 55
Q2LI66/5-222 --KNVIKEFMRFKVMEGVSNGHEFEIEGEGEGRPYEGHNT-VKLKVTK--GGPLPFAWD 55
Q66ND8/5-222 --KNVIKEFMRFKVMEGVSNGHEFEIEGEGEGRPYEGHNT-VKLKVTK--GGPLPFAWD 55
Q66ND7/5-222 --KNVIKEFMRFKVMEGVSNGHEFEIEGEGEGRPYEGHNT-VKLKVTK--GGPLPFAWD 55
Q9U6Y8/5-222 --KNVIKEFMRFKVMEGVSNGHEFEIEGEGEGRPYEGHNT-VKLKVTK--GGPLPFAWD 55
Q6K85/5-222 --KNVIKEFMRFKVMEGVSNGHEFEIEGEGEGRPYEGHNT-VKLKVTK--GGPLPFAWD 55
Q2WG74/7-224 --ENVITFMRFKVMEGVSNGHEFEIEGEGEGRPYEGHNT-VKLKVTK--GGPLPFAWD 55
Q9GTJ7/5-224 --KNVIKEFMRFKVMEGVSNGHEFEIKGEGEGRPYEGHCS-VKLMVTK--GGPLPFAFD 55
A8CLM7/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VRLTVTK--GGPLPFAWD 55
A8CLM1/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
Q1JU63/2-219 --VSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
Q1JV71/2-219 --VSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
Q1JV72/2-219 --VSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
Q1JV70/2-219 --VSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
B5T1L1/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VRLAVTK--GGPLPFAWD 55
A8CLV6/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
Q66P0/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
Q66P8/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
Q9SP04/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
A8CLL3/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
Q66P1/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
Q66P9/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VRLTVTK--GGPLPFAWD 55
P83690/1-218 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
A8CLN8/6-226 --QSGVDVMTLHMDGIVNGHDFIEGKGGKPNPYEGVQT-MKLVTK--GAPLPFSID 55
B6CTZ7/1-220 --MSVIKQVMKTLHLEGTVNGHDFIEGKGGKPNPYEGLQH-MKMTVTK--GAPLPFSVH 55
Q9U6Y5/5-228 --KHGLTKEMTKYRMEGCVGDGHKVFITGEGIGYPPFKGQA-INLCVVE--GGPLPFAED 55
Q9U6Y4/5-228 --KHGLTKEMTKYRMEGCVGDGHKVFITGEGIGYPPFKGQA-INLCVVE--GGPLPFSED 55
Q8T4U4/5-228 --KHGLTDDMTMFRMEGCVGDGHKVFIEGNGNPNPYEGVQT-INLCVVE--GGPLPFSED 55
B2ZAG2/5-226 --KHGLTDDMTMFRMEGCVGDGHKVFITGEGIGYPPFKGQA-INLCVVE--GGPLPFSED 55
Q69ZV6/5-228 --KHGLTKDMTKYRMEGCVGDGHKVFITGEGIGYPPFKGQA-INLCVVE--GGPLPFSED 55

B5LIN1/1-161 -----
B5T1L5/5-228 --KHGLTKDMTKYRMEGCVGDGHKVFITGEGIGYPPFKGQA-INLCVVE--GGPLPFSED 55
Q66PV2/5-228 --KHGLTKDMTKYRMEGCVGDGHKVFITGEGIGYPPFKGQA-INLCVVE--GGPLPFSED 55
Q86LC2/5-228 --KHGLDFDMTKYRMEGCVGDGHKVFITGEGIGYPPFKGQA-INLCVVE--GGPLPFSED 55
B2ZAG1/5-228 --KQGLTDNMTKYRMEGCVGDGHKVFITGEGIGYPPFKGQA-INLCVVE--GGPLPFSED 55
A8CLN4/5-231 --KNGLTKNMTKYRMEGCVGDGHKVFITGEGIGYPPFKGQA-INLCVVE--GGPLPFSED 56
A8CLV1/5-230 --KNGVDRMMLKFKHMEGVSNGHEFTIEGEGTGPYEGTQS-IQLRVEK--GGPLPFSVD 55
A8CLU7/1-223 ---MALPKQMKLYHMEGVSNGHEFTIEGEGGPEYEGTHT-IKLQVVE--GSPLPFSPD 54
A8CLP2/5-228 --KQSLPDSMKLIYHMDGIVNGHSFVIEGEGGKPNPYEGTHT-IKLQVVE--GSPLPFSAD 55
B6CTZ2/5-226 --KNVLPNMMLTYHMEGVSNGHEFTIEGEGTGNPKGKHT-ITLQVVK--GGPLPFSVD 55
Q2VTM9/5-219 --KQVLPDRVDMRYHMDGIVNGHDFIEGEGTGPYEGKKT-LKLRTVK--GGPLPFAFD 55
Q1ALD3/5-219 --KQVLPDRVDMRYHMDGIVNGHSFTIEGEGTGPYEGKKT-LKLRTVK--GGPLPFAFD 55
Q1ALD5/5-219 --KQVLPDRVDMRYHMDGIVNGHSFTIEGEGTGPYEGKKT-LKLRTVK--GGPLPFAFD 55
Q1ALD4/5-219 --KQVLPDRVDMRYHMDGIVNGHSFTIEGEGTGPYEGKKT-LKLRTVK--GGPLPFAFD 55
Q09HR5/5-219 --KQVLPDRVDMRYHMDGIVNGHSFTIEGEGTGPYEGKKT-LKLRTVK--GGPLPFAFD 55
Q2VTM8/5-219 --KQVLPDRVDMRYHMDGIVNGHSFTIEGEGTGPYEGKKT-LKLRTVK--GGPLPFAFD 55
A8CLU1/1-218 ---MALSKQVTKYHMDGIVNGHDFIEGEGTGPYEGKKT-VTLVTK--GAPLPFSFD 54
Q9U6Y6/5-224 --NFIQDDMKMTYHMDGIVNGHDFIEGEGTGPYEGKKT-STFKVTMANGKPLPFSFD 57
A8CLT7/5-222 --KQCIANEMTMFHMDCVNGHYFTIEGEGSGRPYEGKQM-SKFKVTK--GGPLPFSFD 55
Q66ND1/5-220 --KQVILKDMMLKFKHMEGVSNGHEFTIEGEGTGPYEGKQM-STFKVTK--GGPLPFSFD 55
Q66PV5/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
B5T1L2/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
Q66PV3/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
B5T1L0/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
Q66PV4/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
B5LIN0/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
B5T1L3/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
B5T1L4/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
Q66PV7/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
A8CLY3/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
Q66PV8/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
Q66PV9/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
A8CLX9/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
B6CTZ3/5-226 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
Q6I7B1/6-227 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
Q66PV6/5-226 --KHGITQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-AELVITKPAKPLPFSFD 57
Q86LV8/1-213 ---MAVPTQVKMYSMDGNFNGQSFVVEGEGTGNPYEGHQ-SLKLTV---KGEPLPFAFD 53
Q86LV7/1-213 ---MAVPTQVKMYSMDGNFNGQSFVVEGEGTGNPYEGHQ-SLKLTV---KGEPLPFAFD 53
A8CLR1/5-220 --KQGIQVQEMTKYHMEGVSNGHEFTIEGEGTGPYEGKQM-SELVITKPAKPLPFSFD 57
Q6R8F5/5-223 --KQVIGKEMNMTYHMDGIVNGHDFIEGEGTGPYEGHQ-SLKLRT---KGEPLPFAFD 55
Q6R8F4/1-219 ---MVLKQNMNTYHMDGIVNGHDFIEGEGTGPYEGHQ-SLKLRT---KGEPLPFAFD 54
Q6R8F3/5-220 --KQVIGKEMNMTYHMDGIVNGHDFIEGEGTGPYEGHQ-SLKLRT---KGEPLPFAFD 55
Q6I7B2/2-218 --VSVIKPEMKRYHMDGIVNGHDFIEGEGTGPYEGHQ-MTLRVTMAKGGPMPFAFD 57
A7VMR3/2-218 --VSVIKPEMKRYHMDGIVNGHDFIEGEGTGPYEGHQ-MTLRVTMAKGGPMPFAFD 57
Q6I7B4/1-210 -----MKMYFMDGVSNGHEFTIEGEGTGPYEGHQ-MTLRVTMAKGGPMPFAFD 50
Q6I7B3/1-217 --MSVIAKQMTYKYVMSGTVNGHYFEVQDGGKGPYEGEQT-VKLTVTK--GGPLPFAWD 55
A8CLQ6/5-221 --KQAIKQDMKINYFMDGVSNGHEFTIEGEGTGPYEGHQ-MTLRVTMAKGGPMPFAFD 57
A8CLW9/4-223 --SKQVLNDVMTYFMDGVSNGHDFIEGEGTGPYEGHQ-FNLRVT--KGAPLPFAVD 55

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

Q95W86/2-220 --AGLLKESMRKIYMEGTVNGYHFCKEAGEGDNPFEGTQN-MRIRVTE--GAPLPFAFD 55
Q95W11/2-220 --AGLLKESMRKIYMEGTVNGYHFCKEAGEGDNPFEGTQN-MRIRVTE--GAPLPFAFD 55
Q8MU45/2-220 --AGLLKESMRKIYMEGTVNGYHFCKEAGEGDNPFEGTQN-MRIRVTE--GAPLPFAFD 55
Q95W85/2-220 --AGLLKESMRKIYMEGTVNGYHFCKEAGEGDNPFEGTQS-MRIRVTE--GAPLPFAFD 55
ASYTR7/2-222 --ASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKNPFEGTQE-MKIEVIE--GGPLPFAFH 55
Q9GZ28/2-222 --ASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKNPFEGTQE-MKIEVIE--GGPLPFAFH 55
ASYTR8/2-222 --ASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKNPFEGTQE-MKIEVIE--GGPLPFAFH 55
Q9GP15/2-222 --ASFLKKTMPFKTTIEGTVNGHYFKCTGKGEKNPFEGTQE-MKIEVIE--GGPLPFAFH 55
Q2VFP3/2-221 --ASLVKDDMCIKMTMEGTVNGHHFKCVGEGEGKPFEGTQV-EKIRITE--GGPLPFAFD 55
A0AQQ7/2-222 --ASKISDNVRKILYMEGTVNNHHFMCFAEGEGKPYEGTQM-ENIKVTK--GGPLPFSFD 55
A0AQQ8/2-222 --ASKISDNVRKILYMEGTVNNHHFMCFAEGEGKPYEGTQM-ENIKVTK--GGPLPFSFD 55
Q8T5F8/2-222 --NSLIKENMRMVMVMEGSVNGYQFKCTGEGDGNPYMGQT-MRIKVVE--GGPLPFAFD 55
Q86LV4/2-220 --YSYIKETMRKSVYMEGNVNNHAFKCTAEGEGKPYKGSQK-LTITVTE--GGPLPFAFD 55
Q5QIU5/2-220 --YSYIKETMRKSVYMEGNVNNHAFKCTAEGEGKPYKGSQK-LTITVTE--GGPLPFAFD 55
Q8T6T9/2-220 --CSYIKETMQSKVYMEGNVNNHAFKCTAEGEGKPYKGSQK-LTITVTE--GGPLPFAFD 55
Q8T5E7/2-220 --YPIWIKETMRKSVYMEGNVNNHAFKCTAEGEGKPYKGSQK-LTITVTE--GGPLPFAFD 55
ASYTR6/2-219 --YPIWIKETMRVQLSMEGSVNYHAFKCTGKGEKPYEGTQS-LNITITE--GGPLPFAFD 55
Q9GP16/2-219 --YPIWIKETMRVQLSMEGSVNYHAFKCTGKGEKPYEGTQS-LNITITE--GGPLPFAFD 55
Q5ZQ06/2-221 --DPLVNGNLKIKMFMNGSVNKHFKCEAEGEGNIHSGTHI-MKITVTE--GGPLPFAFD 55
B9UPG6/1-219 --MSQFPEYLPVTVHMRGNVNNLEFEYDGGGGDPRAGFT-MNMQLRG--RKPLPFSYD 55
Q5ZQ05/1-219 --MSQDNNLSVSVYMKGNVNNHFEFEYDGGGGDPSGQFS-LKTKLRG--GKPLPFSYD 55
Q7Z168/1-218 ---MNLKSNVSVSVYMKGNVNNHFEFEYDGGGGDPTGKYS-MKMTLRG--QNCLPFSYD 54
Q9BLZ0/8-225 --NTGLKEIMSASAKAVEGIVNNHVFMEGFGKGNVLFQGNL-MQIRVTK--GGPLPFAFD 55
Q9BLY9/8-225 --NTCLQEVMSYKVNLEGI VNNHVTFMEGCKGNILFQGNL-VQIRVTK--GAPLPFAFD 55
Q96319/5-223 --KLGKLEVMPTKINLEGLVGDHAFSMEGVGEGNILEGTQE-VKISVTK--GAPLPFAFD 55
A8CLX6/5-222 --KQALADTMKMTWLMEGSVNGHAFTEIEGTGKPYEGKQS-GTFRVTK--GGPLPFAFD 55
A7S3R2/57-279 --YSLKSDAMKLFHFKMEGSVNGHCFEIQGVGEGKAFDGEHW-SKLCVVK--GKHLPFSYD 55
A7RQ05/1-174 ---MKLHFLEGSVNGHCFEIQGEVGEKPFEGEQW-AKHCVVK--GKHLPFLSD 48
A7S1H5/1-224 --MSLSDAMKLLHILEGSVNGHCFEIHGEVGEKAFEGEQW-SKFTVVK--GGPLPFSYD 55
A7S1H3/1-82 -----
B1PNC2/5-218 -----ATHELHIFGSFNGVDFDMVGHGTGNPNDGYEE---LNLKSTKGALQFSPW 47
B1PNC0/4-217 -----PATHELHIFGSFNGVDFDMVGRGTGNPNDGYEE---LNLKSTKGALQFSPW 48
B1PNC1/5-218 -----ATHELHIFGSFNGVDFDMVGRGTGNPNDGYEE---LNLKSTKGALQFSPW 47
B1PNC4/5-218 -----ATHELHIFGSFNGVDFDMVGRGTGNPNDGYEED---LHLKSTKGALQFSPW 47
C3YR99/48-261 -----LPTTHELHIFGSFNGVDFDMVGRGTGNPNDGYEE---LNLKSTKGALQFSPW 49
B1PNC3/4-217 -----PATHELHIFGSINSLEFDLVGRGTGNPKEGYEE---LHLKSTKSALQFSPW 48
B1PNB8/5-217 -----ATHELHIFGSINSLEFDLVGRGTGNPKEGYEE---LHLKSTKSALQFSPW 47
B1PNB9/4-216 -----PATHELHIFGSINSLEFDLVGRGTGNPREGYEE---LHLKSTKSALQFSPW 48
C3YRA1/51-267 -----LPKTHELHIFGSFNGVDFDMVGEGTGNPNEGSEE---LKLKSTNGPLKFSY 49
C3YRA2/11-223 -----PKTHELHIFGSFNGVDFDMVGRGTGNPNEGSEE---LNAKFTKGPLKFSY 48
B1PND7/4-187 -----PATHELHIFGKINGHEYDMRGKGTGNPNDGYED---LDLKSTKDDLQFSPW 48
B1PND6/5-189 -----KTHELHIFGKINGHEYDMRGKGTGNPNDGYED---LDLKS-KDDLQFSPW 46
B1PND8/4-184 -----PATHELHIFGKINGHEYDMRGKGTGNPNDGYED---LDLKSTKDDLQFSPW 48
B1PND5/5-186 -----ATHELHIFDINGHEYDMRGKGTGNPNDGYED---LDLKSTKDDLQFSPW 47
C3YRA3/4-216 -----PTTHEVHYGSINGVEFDLVGSGKGNPKDGESE---IQVKSTKGPLKFSY 48
B1PND3/4-217 -----PATHDLHISGSINGHEFDLEGSKGNAKEYQE---LHLKSNKGDLSFSPW 48
B1PND2/5-217 -----ATHDLHISGSINGHEFDLEGSKGNAKEYQE---LHLKSNKGDLSFSPW 47
B1PNC9/4-217 -----PATHDLHISGSINGHEFDLEGSKGNAKEYQE---LHLKSNKGDLSFSPW 48
B1PNC6/5-217 -----ATHDLHISGSINGHEFDLEGSKGNAKEYQE---LHLKSNRGLSFSPW 47
B1PNC7/4-220 -----PATHDLHISGSINGHEFDLEGSKGNAKEYQE---LHLKSNKGDLSFSPW 48
B1PND0/5-220 -----ATHDLHISGSINGHEFDLEGSKGNAKEYQE---LHLKSNKGDLSFSPW 47
B1PNC5/5-220 -----ATHDLHISGSINGHEFDLEGSKGNAKEYQE---LHLKSNKGDLSFSPW 47
B1PND1/5-217 -----KTHDLHISGSVNGHEFDLEGSKGNAKEYQE---LHLKSNRGLSFSPW 47
B1PND4/5-217 -----KTHDLHISGSVNGHEFDLEGSKGDAKEYQE---LHLKSNRGLSFSPW 47
B1PNC8/4-69 -----PATHDLHISGSINGHEFDLEGSKGNAKEYQE---LHLKSNRGLSFSPW 48
C3YRA0/8-104 TSAVTEPLPTTHELHIFGSFNGVDFDLVGRGEGNPKDGSQK---LHLKSTKGALQFSPW 56
C3YKQ3/5-218 -----ATHDLHISGSINGHEFDLVGGGKGNPAGSLV---TTAKSTKGALQFSPY 47
C3YK7/5-217 -----ATHEIHLHGSVNGHEFDLVGGGKGNPAGSLV---TEVKSTMGPLKFSY 47
C3YKQ2/53-191 -----
A8QZ9/4-215 -----PTTHDLHIFGSVNGAEFDLVGGGKGNPNDGTLE---TSVKSTRGALPCSPL 48
C3YK9/5-215 -----TAHDLHIFGSVNGAEFDLVGGGKGNPNDGTLE---TSVKSTRGALPCSPL 47
C3YKQ1/4-215 -----PTTHDLHIFGSVNGAEFDLVGGGKGNPNDGTLE---TSVKSTRGALPCSPL 48
C3ZLQ0/369-585 ---TMSVPTNLDLHIYGSINGMEFDMVGGGSGNPNDSLS---VNVKSTKGALRVSP 52
C3YK10/4-218 -----KSVPTNLDLHIYGSINGMEFDMVGGGSGNPKDGSLSA---VNVKSTKGALRVSP 51
C3YKQ0/5-214 -----TAHCHMFGSINGHEFDLVGGGKGNPNDGTLE---TKVRSTKGALPFSV 47
Q6W10/1-215 -----MPDMKLECHISGTMNGEEFELVAGDGNTEGRMT---NKMKSIGPLSFSY 50
Q6W09/1-217 -----MPDMKLECHISGTMNGEEFELVAGDGNTEGRMT---NKMKSIGPLSFSY 50
Q6W11/1-215 -----MPVMKIECRISGTMNGEEFELVAGDGNTEGRMT---NKMKSIGPLSFSY 50
Q6W12/1-213 -----MPAMKIECRITGLNGVFEFELVGGGEGTPEQGRMT---NKMKSIGPLSFSY 50
Q6W13/1-214 -----MPAMKIECRISGTLNGVFEFELVGGGEGTPEQGRMT---NKMKSIGPLSFSY 50
Q6W08/1-214 -----MAAMKIECRITGLNGVFEFELVGGGEGTPEQGRMT---NKMKSIGPLSFSY 50
Q2MHN7/1-215 -----MTTFKIESRIHGNLNGEKFELVGGG---EEGRLE---IEMKTKDKPLAFSP 48

Q17106/5-229 TLVTTFSYGVQCFSRYPDHMKQHDFLKSAMPEG---YVQERTIFYK-DDGNYKTRAEV- 108
Q17105/5-229 TLVTTFSYGVQCFSRYPDHMKQHDFLKSAMPEG---YVQERTIFYK-DDGNYKTRAEV- 108
P42212/5-229 TLVTTFSYGVQCFSRYPDHMKQHDFLKSAMPEG---YVQERTIFFK-DDGNYKTRAEV- 108
Q6YGZ0/5-229 TLVTTFSYGVQCFSRYPDHMKQHDFLKSAMPEG---YIQERTIFFK-DDGNYKTRAEV- 108
Q8WTC5/5-229 TLVTTLYGILCFARYPEHMKMNDFFKSAMPEG---YIQERTIFFQ-DDGKYKTRGEV- 108
Q8WTC4/5-229 TLVTTLYGILCFARYPEHMKMNDFFKSAMPEG---YIQERTIFFQ-DDGKYKTRGEV- 108
Q8WTC9/5-229 TLVTTLYGIQCFARYPEHMKMNDFFKSAMPEG---YIQERTIFFQ-DDGKYKTRGEV- 108
Q8WTD0/5-229 TLVTTLYGIQCFARYPEHMKMNDFFKSAMPEG---YIQERTIFFQ-DDGKYKTRGEV- 108
Q8WTC8/5-229 TLVTTLYGIQCFARYPEHMKMNDFFKSAMPEG---YIQERTIFFQ-DDGKYKTRGEV- 108

Q8WTC6/5-229 TLVTTLSYGIQCFARYPEHMKMNDFFKSAMPEG----YIQERTIFFQ--DDGKYKTRGEV- 108
Q8WP95/5-229 TLVTTFSYGIQCFARYPEHMKMNDFFKSAMPEG----YIQERTIFFQ--DDGKYKTRGEV- 108
Q8WTC7/5-229 TLVTTLSYGIQCFARYPEHMKMNDFFKSAMPEG----YIQERTIFFQ--DDGKYKTRGEV- 108
Q6RYS7/5-229 TLVTTLYTGAQCFARYPELK--DFYKSCMPEG----YVQERTITFE--GDGVFKTRAEV- 106
Q6RYS6/39-258 SLITLSYGAQCFARYPNEIN--DFKSTFSPG----YHQERKITYE--NDGVLTAALKI- 106
Q6RYS5/2-222 SLITLSYGGKCFVRYPNVIK--DFKSTFPTG----YHQERKITYE--DDGVLTAALKV- 106
Q6RYS4/5-225 PICHLIQYGEPPFAKYPNGIS--HFAQCFPEG----LIDRTVRFE--NDGTMTSHHTY- 106
Q2TCH5/5-225 PICHLIQYGEPPFAKYPDGIS--HFAQCFPEG----LIDRTVRFE--NDGTMTSHHTY- 106
Q53UG7/1-221 ILTTVFYDGNRVFVKYPEEIV--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q53UG8/1-221 ILTTAFHYGNRVFVYEPEDIV--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q8MU48/1-218 ILTTAFYDGNRVFVKYPEDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A8CLW1/1-218 ILTTIFNYGNRVFVKYPPDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
B6CT24/1-218 ILTTIFNYGNRVFVKYPPDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q5TLG6/1-218 ILTTVFYDGNRVFVKYPENIV--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A8CLQ1/1-223 ILTTSFNYGNRVFVKYPTDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q2LI65/1-219 ILTTVFHYGNRVFVKYPGDIA--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A8CLW4/1-222 ILTTVFYDGNRVFVKYPTDIA--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
B6CT25/4-224 ILTTVFYDGNRVFVKYPEIV--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A3F207/1-221 ILTTVFHYGNRVFVKYPPDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q962P9/1-221 ILTTVFYDGNRVFVKYQDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q7Z0W8/1-221 ILTTVFYDGNRVFVKYQDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q66ND5/1-221 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q66ND4/1-221 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q66ND2/1-221 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A7UAL7/1-221 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q7Z0W6/1-221 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A7UAL9/1-221 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q8T5F2/8-228 ILTTVFYDGNRVFVKYPEDIA--DYFKQSFP---EGYFWERSMSYE--DGGICLATNMI- 107
Q7Z0W7/8-228 ILTTVFYDGNRVFVKYPEDIA--DYFKQSFP---EGYFWERSMSYE--DGGICLATNMI- 107
Q8MU47/8-228 ILTTVFYDGNRVFVKYPEDIA--DYFKQSFP---EGYFWERSMSYE--DGGICLATNMI- 107
Q95V10/1-221 ILTTVFYDGNRVFVKYQDIP--DYFKQSFP---EGYFWERSMSYE--DGGICLATNMI- 107
A7UAL3/1-221 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A7UAL6/1-218 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A7UAL8/1-218 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A8CL52/1-219 ILTTAFHYGNRVFVKYPDHII--DYFKQSFP---NGFSWERSLMFE--DGGICLATNMI- 107
Q8I6J8/1-219 ILTTAFHYGNRVFVKYPDHII--DYFKQSFP---KGSWERSLMFE--DGGICLATNMI- 107
A3F208/1-219 ILTTAFHYGNRVFVKYGNIP--DYFKQSFP---KGSWERSLMFE--DGGICLATNMI- 107
A8CLR7/1-219 ILTTAFHYGNRVFVKYPDHII--DYFKQSFP---GGYSWERSLMFE--DGGICLATNMI- 107
Q55629/1-219 ILTTAFHYGNRVFVKYPDHII--DYFKQSFP---KGSWERSLMFE--DGGICLATNMI- 107
Q66PW1/5-223 ILTTAFHYGNRVFVKYPENIP--DYFKQSFP---KGSWERSLMFE--DGGICLATNMI- 107
Q8T5F1/1-219 IMTTVFHYGNRVFVKYKHIP--DYFKQSFP---EYSWERSMSYE--DGGICLATNMI- 107
Q7Z0W9/1-219 IMTTVFHYGNRVFVKYKHIP--DYFKQSFP---EYSWERSMSYE--DGGICLATNMI- 107
Q6UKS3/1-219 IMTTVFHYGNRVFVKYKHIP--DYFKQSFP---EYSWERSMSYE--DGGICLATNMI- 107
A7UAL4/1-219 IMTTAFHYGNRVFVKYKHIP--DYFKQSFP---EYSWERSMSYE--DGGICLATNMI- 107
A7UAM0/1-219 ILTTVFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q7Z0W5/1-218 ILTTAFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q95UA7/1-218 ILTTAFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A7UAL1/1-218 ILTTAFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A8CLP7/1-218 ILTTAFYDGNRVFVKYPKDIP--DYFKQSFP---AGYSWERSMSYE--DGGICLATNMI- 107
A7UAL2/1-218 ILTTAFYDGNRVFVKYPKDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
A7UAL5/1-218 ILTSAFYDGNRVFVKYPPDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q963F5/1-218 ILTSAFYDGNRVFVKYPPDIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q7Z0W4/1-218 ILTAAFYDGNRVFVKYPRDIA--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q66ND3/1-218 ILTAAFYDGNRVFVKYPRDIA--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q2LI64/1-35 -----
Q60I25/1-218 ILTTVFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q60I24/1-218 ILTTVFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
A8CLT2/1-218 ILTTVFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
B5BR3/1-218 ILTNAFYDGNRVFVKYKADIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
B6CT26/1-219 ILTNAFYDGNRVFVKYKADIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q9U6Y3/43-260 ILSNAFYDGNRVFVKYKADIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q8T6U0/1-218 ILTTALHYGNRVFVKYKADIP--DYFKQSFP---EGYSWERSMSYE--DGGICLATNMI- 107
Q8T5F0/7-229 ILSAIFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 109
Q8T5F3/6-223 ILSAIFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 109
Q66PW0/1-221 ILTTIFQYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 110
Q8MU46/1-221 ILTTIFQYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 110
Q8T5E9/1-221 ILTTIFQYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 110
Q8T5E8/1-221 ILTTIVHYGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 110
Q8T6T8/1-221 ILTTMFQYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 110
Q66PUS/1-220 LLTPAFMYGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
A8CL57/1-220 LLTPAFMYGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q8MA2/34-254 LLTPAFMYGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q8MA1/12-234 LLTPAFMYGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q66ND6/5-222 ILCPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q9U6Y7/5-223 ILCPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q553G7/6-223 ILSPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q553G8/6-223 ILSPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q2LI66/5-222 ILSPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q66ND8/5-222 ILSPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q66ND7/5-222 ILSPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q9U6Y8/5-222 ILSPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q6K6F85/5-222 ILSPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107
Q2WG74/7-224 ILSPOFYDGNRVFVKYKADIP--DYFKQSFP---EGYHWSERSMSYE--DGGICLATNMI- 107

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

Q9GTJ7/5-223 ILSPQFYQGSKYVVKHPADIP--DYKLSFP---EGFKWERVMNFE-DGGVVTVSQDS- 107
A8CLM7/1-218 ILSPLSQYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
A8CLM1/1-218 ILSPLSQYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
Q1JU63/2-219 ILSPLMICYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
Q1JV71/2-219 ILSPLFQYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
Q1JV72/2-219 ILSPLFQYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
Q1JV70/2-219 ILSPLQYGSIPFTKYPEDIP--DYFKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
B5T1L1/1-218 ILSPQCYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
A8CLV6/1-218 ILSPQCYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
Q66PV0/1-218 ILSPQCYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
Q66PU8/1-218 ILSPQCYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
Q9SP04/1-218 ILSPQCYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
A8CLL3/1-218 ILSPQAQYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
Q66PV1/1-218 ILSPQFYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DDAVCTVSND- 107
Q66PU9/1-218 ILSPQCYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
P83690/1-218 ILSPQCYGSIPFTKYPEDIP--DYVKQSFPP---EGYTWERIMNFE-DGAVCTVSND- 107
A8CLN8/6-226 ILLPQHMYSKPFKYPESIP--DYIKLSFP---EGITWERSMTFE-DGAVCTVSND- 107
B6CTZ7/1-220 ILTPSHMYSKPFKYPADIP--DYHKQSFPP---EGMSWERSMIFE-DGGVCTASNHSS 108
Q9U6Y5/5-228 ILSAAFYGNRVFTEYPQDIV--DYFKNSCPAG---YTWDRSFLFE-DGAVCICNADIT 108
Q9U6Y4/5-228 ILSAGFKYGRIFTEYPQDIV--DYFKNSCPAG---YTWGRSFLFE-DGAVCICNV DIT 108
Q8T4U4/5-228 ILSAAFYGNRLFTEYPEGIV--DYFKNSCPAG---YTWHRSRFRFE-DGAVCICSADIT 108
B2ZAG2/5-226 ILSAAFYGNRLFTEYPOGIV--DYFKNSCPAG---YTWQRSFLFE-DGAVCIASADIS 108
Q692V6/5-228 ILSAAFYGNRVFTEYPOGMV--DFFKNSCPAG---YTWHRSL LFE-DGAVCTTSADIT 108
B5LIN1/1-161 -----GNRVFTEYPOGMV--DFFKNSCPAG---YTWHRSL LFE-DGAVCTTSADIT 45
B5T1L5/5-228 ILSAAFYGNRVFTEYPOGMV--DFFKNSCPAG---YTWHRSL LFE-DGAVCTTSADIT 108
Q66PV2/5-228 ILSAAFYGNRVFTEYPOGMV--DFFKNSCPAG---YTWHRSL LFE-DGAVCTTSADIT 108
Q86LC2/5-228 ILSVTFYGNRIFTEYPSIV--DYFKNSCPAG---HTWQRSL LFE-DGAVCIASADIS 108
B2ZAG1/5-228 ILSAVFTYGNRIFADYQPID--DYFKNSCPAG---YTWTRSFLFE-DGAVAIASADIR 108
A8CLN4/5-231 ILSAVFYGNRVFTKYPQDLV--DYFKNSCPAG---YTWQRSFLFE-DGAVCTASADIT 109
A8CLV1/5-230 ILSAVFLYGNRVFTKYPQDLV--DYFKNSCPAG---YTWQRSFLFE-DGAVCTASADIT 108
A8CLU7/1-223 ILSVTFYGNRCFTKYPNPVIV--DYFKNSCSGGG---YTFGRSFLYE-DGAVCTASGDIT 108
A8CLP2/5-228 ILSVTFYGNRCFTKYPNPVIV--DYFKNSCSGGG---YKFGRSFLYE-DGAVCTASGDIT 108
B6CTZ2/5-226 ILSVTFYGNRCFTKYPNPTV--DYFKNSCPPG---YTFERSFLYE-DGAVCTASGDIT 109
Q2VTM9/5-219 ILSVTFYGNRCFCEYPEDMP--DYFKQSLPEG---HSWERTLMFE-DGGCGTASAHIS 108
Q1ALD3/5-219 ILSATFTYGNRCFCEYPEDMP--DYFKQSLPEG---YSWERTMMYE-DGACGTASAHIS 108
Q1ALD5/5-219 ILSATFTYGNRCFCEYPEDMP--DYFKQSLPEG---YSWERTMMYE-DGGCGTSAHIR 108
Q1ALD4/5-219 ILSATFTYGNRCFCEYPEDMP--DYFKQSLPEG---YSWERTMMYE-DGACGTASAHIS 108
Q09HR5/5-219 ILSATFTYGNRCFCEYPEDMP--DYFKQSLPEG---YSWERTMMYE-DGGCGTASAHIS 108
Q2VTM8/5-219 ILTATFYGNRCFCEYPEDMP--DYVKQSFPEG---YSFERTMMFE-DGACCTTSVHLS 108
A8CLU1/1-218 ILSAVFLYGNRAFTDYPKIV--DYFKQSFPEG---YSFERTLFE-DGGYCTASADIS 107
Q9U6Y6/5-224 ILSVTFYGNRCFTAYPTSMV--DYFKQAFPDPG---MSYERTFTYE-DGGVATASWEIS 110
A8CLT7/5-222 ILSAFYGNRCFTAYPAGMH--DYFKQAFPPEG---MSYERTFTFE-DGGVATASGDIS 108
Q66ND1/5-220 ILSAFYGNRCFTKYPADMP--DYFKQAFPAG---MSFERTFTFE-DGGVATASGHIC 108
Q66PV5/5-226 ILSVTFYGNRCFTKYPEGMP--DYFKQAFPDPG---MSFERSFLYE-DGGVATASWNIR 110
B5T1L2/5-226 ILSVTFYGNRCFTKYPEGMP--DYFKQAFPDPG---MSYERSFLYE-DGGVATASWNIR 110
Q66PV3/5-226 ILSVTFYGNRCFTKYPEGMP--DYFKQAFPDPG---MSYERSFLYE-DGGVATASWNIR 110
B5T1L0/5-226 ILSVTFYGNRCFTKYPEGMP--DYFKQAFPDPG---MSYERSFLYE-DGGVATASWNIR 110
Q66PV4/5-226 ILSVTFYGNRCFTKYPEGMP--DYFKQAFPDPG---MSYERSFLYE-DGGVATAGWNIG 110
B5LIN0/5-226 ILSVTFYGNRCFTKYPEGMP--DYFKQAFPDPG---MSYERSFLYE-DGGVATASWNIR 110
B5T1L3/5-226 ILSVTFYGNRCFTKYPEGMP--DYFKQAFPDPG---MSYERSFLYE-DGGVATASWNIR 110
B5T1L4/5-226 ILSVTFYGNRCFTKYPADMP--DYFKQAFPDPG---MSYERSFLYE-DGAVATASWKIR 110
Q66PV7/5-226 ILSVTFYGNRCFTKYPADMP--DYFKQAFPDPG---MSYERSFLYE-DGAVATASWKIR 110
A8CLY3/5-226 ILSVTFYGNRCFTKYPADMP--DYFKQAFPDPG---MSYERSFLYE-DGAVATASWNIR 110
Q66PV8/5-226 ILSVTFYGNRCFTKYPADMP--DYFKQAFPDPG---MSYERSFLYE-DGAVATASWNIR 110
Q66PV9/5-226 ILSVTFYGNRCFTKYPADMP--DYFKQAFPDPG---MSYERSFLYE-DGAVATASWNIR 110
A8CLX9/5-226 ILSVTFYGNRCFTKYPADMP--DYFKQAFPDPG---MSYERSFLYE-DGAVATASWNIR 110
B6CTZ3/5-226 ILSVTFYGNRCFTKYPADIP--DYFKQAFPDPG---MSYERSFLYE-DGAVATASWNIR 110
Q6I7B1/6-227 ILSVTFYGNRCFTKYPADMP--DYFKQAFPDPG---MSYERSFLYE-DGGVATASWSIR 110
Q66PV6/5-226 ILSVTFYGNRCFTKYPADLP--DYFKQAYPGG---MSYERSFYVQ-DGGIATASWNVQ 110
Q86LV8/1-213 ILSATFTYGNRVFTKYPEGKT--DYFKQAFPDPG---LWERTMTFE-DGGICTVAEIS 106
Q86LV7/1-213 ILSATFTYGNRVFTKYPEGKT--DYFKQAFPDPG---LWERTMTFE-DGGICTVAEIS 106
A8CLR1/5-220 ILSAAFYGNRCFTKYPDEIR--DYFKQAFPDPG---LWERSMAFE-DGASAATAEIS 110
Q6R8F5/5-223 ILTAAFCYGNRCFVNYPAEIA--DYFKQAFPPEG---LWERSMTFE-DGGFAAVSADIS 108
Q6R8F4/1-219 ILTAAFCYGNRCFVNYPAEIA--DYFKQAFPPEG---LWERSMTFE-DGGFAAVSADIS 107
Q6R8F3/5-220 ILSAAFYGNRCFTKYPKEIP--DMFKQAFPDPG---MSWERTMTFE-DGGVASVSADIS 108
Q6I7B2/2-218 LVSHVFCYGNRFTKYPEEIP--DYFKQAFPPEG---LWERSLQFE-DGGSASVSADIS 110
A7VMR3/2-218 LVSHVFCYGNRFTKYPEEIP--DYFKQAFPPEG---LWERSLQFE-DGGSASVSADIS 110
Q6I7B4/1-210 LVSHVFCYGNRFTKYPEEIP--DYFKQAFPPEG---LWERSLQFE-DGGFAAVSADIS 103
Q6I7B3/1-217 LVSHVFCYGNRFTKYPEEIP--DYFKQAFPPEG---LWERSLQFE-DGGFAAVSADIS 110
A8CLQ6/5-221 ILSHTFCYGNRFTKYPEEIP--DYFKQAFPPEG---LWERSLQFE-DGGFAAVNANIS 110
A8CLW9/4-223 ILSAAFYGNRCFTKYPKEIP--DFFKQSLPEG---MSWERTMTFE-DGGIVAI SAHIR 108
Q9SW86/2-220 ILSPPCAYGSKTFIKHTSGIP--DYFKQSFPEG---FTWERTT IYE-DGGVLT AHQDT- 107
Q9SW11/2-220 ILSPPCAYGSKTFIKHTSGIP--DYFKQSFPEG---FTWERTT IYE-DGGVLT AHQDT- 107
Q8MU45/2-220 ILSPPCAYGSKTFIKHTSGIP--DYFKRSFPEG---FTWERTT IYE-DGRLVTAHQDT- 107
Q9SW85/2-222 ILAPCCYGRSFTVHHTAEIP--DFFKQSFPEG---FTWERTT IYE-DGGILT AHQDT- 107
ASYTR7/2-222 ILSSTCMYGSKTFIKYVSGIP--DYFKQSFPEG---FTWERTT IYE-DGGFLT AHQDT- 107
Q9GZ28/2-222 ILSSTCMYGSKTFIKYVSGIP--DYFKQSFPEG---FTWERTT IYE-DGGFLT AHQDT- 107
ASYTR8/2-222 ILSSTCMYGSKTFIKYVSGIP--DYFKQSFPEG---FTWERTT IYE-DGGFLT AHQDT- 107
Q9GP15/2-222 ILSSTCMYGSKTFIKYVSGIP--DYFKQSFPEG---FTWERTT IYE-DGGFLT AHQDT- 107
Q2VFP3/2-221 ILAPCCYGSKTFIKHVSIGIP--DYFKRSFPEG---FTWERTQ IFE-DGGYLT IAHQDT- 107
A0AQQ7/2-222 ILSPPCAYGSAITKYTSIGIP--DYFKQSFPEG---FTWERTT IYE-DGAYLT TQOET- 107
A0AQQ8/2-222 ILTPNCQLGSAITKYTSIGIP--DYFKQSFPEG---FTWERTT IYE-DGAYLT TQOET- 107
Q81SF8/2-222 ILATSFMYGSKTFIKHTKGIP--DFFKQSFPEG---FTWERTVRYE-DGGVFTVMQDT- 107

Q86LV4/2-220 ILSHAFQYGNKVFTKYPDDIP--DFFKQSLSGG----FTWKRVSNYE-DGGVLTVDQKT- 107
Q5QIU5/2-220 ILSHAFRYGNKVFTKYPDDIP--DFFKQSLSGG----FTWKRVSNYE-DGGVLTVTQET- 107
Q8T6T9/2-220 ILSHAFRYGNKVFAKYPKDPH--DFFKQSLPEG----FTWERSVNYE-DGGVLTVKQET- 107
Q8T5E7/2-220 ILSHAFQYGNKVFTDYPDDIP--DFFKQSLSDG----FTWRRVSXYX-XGGVLTVTQDT- 107
ASYTR6/2-219 ILSHAFQYGIKVFAYKYPKEIP--DFFKQSLPGG----FSWERSVSTYE-DGGVLSATQET- 107
Q9GPI6/2-219 ILSHAFQYGIKVFAYKYPKEIP--DFFKQSLPGG----FSWERSVSTYE-DGGVLSATQET- 107
Q5ZQ06/2-221 ILSRTFCYGHKMLDNSGNIP--DFFKQSLKDG----YSWERETTYE-DGGVLNSKQKH- 107
B9UPG6/1-219 IITTFGQYGVRAFTKYPDSIA--DYFKGSFPEA----FQWNRRIEFE-DGGVINMSSDI- 107
Q5ZQ05/1-219 IITMFGQYGFRAFTKYPEGIA--DYFKGSFPEA----FQWNRRIEFE-DGGVINMSSDI- 107
Q7Z168/1-218 IITTAFAQYGFVFTKYPEGIV--DYFKDSLPA--FQWNRRIEFE-DGGVLNMSDDI- 106
Q9BLZ0/8-225 IVSIAFYGNRTFTKYPDDIA--DYVQSFPAQ----FFYERNLRFE-DGAIIVDIRSDI- 107
Q9BL9/8-225 IVSPAIFYGNRTFTKYPNDIS--DYFIQSFPAQ----FMYERTLRYE-DGGLVEIRSDI- 107
Q96319/5-223 IVSVAFSYGNRAYTGYPEEIS--DYFLQSFPEG----FTYERNIRYQ-DGGTAIVKSDI- 107
A8CLX6/5-222 IVAPTLKYGFKCFMKYPADIP--DYFLAFPEG----LTYDRKIAFE-DGGCATAVEM- 107
A7S3R2/57-279 ILMPSMSYGTQKFAKYPAGMT--DFFKA AVENG--GLSWERTMTFE-DGGYCTIVNTSE 109
A7RGQ5/1-174 IIMPNI7----FAKYPDGMT--GFFKA AVVNG--GLSWERTMAFE-DRGYCTAVNTSE 97
A7S1H5/1-224 LIAPCLKYKSGKPFVKYPDDMT--DFFKA AVENG--GLSWERTMSLEQDGGFCSVVNTSK 110
A7S1H3/1-82 ----MFYGSKAFAYKYPDGMT--DFFKA AVENG--GLSWERTKTFE-DGGYCT--HST 46
B1PNC2/5-218 ILVPIQIGYGFHQYLPFPDG-M--SPFQAAMKDG--GYQVHRTMQFE-DGASLTSNYRYT 101
B1PNC0/4-217 ILVPIQIGYGFHQYLPFPDG-M--SPFQAAMKDG--GYQVHRTMQFE-DGASLTSNYRYT 102
B1PNC1/5-218 ILVPIQIGYGFHQYLPFPDG-M--SPFQAAMKDG--GYQVHRTMQFE-DGASLTSNYRYT 101
B1PNC4/5-218 ILIPHIQYGFHQYLPFPDG-M--SPFQAAMQDG--GYQVHRTMQFE-DGASLTANFRYT 101
C3YR99/48-261 ILVPIQIGYGFHQYLPFPDG-M--SPFQAAMQDG--GYQVHRTMQFE-DGASLTAHFRYT 103
B1PNC3/4-217 ILVPIQIGYGFHQYLPFPDG-M--SPFQAAMQDG--GYQVHRTMQFE-DGATLTGIYRYT 103
B1PNC8/5-217 ILVPIQIGYGFHQYLPFPDG-M--SPFQAAMQDG--GYQVHRTMQFE-DGATLTGIYRYT 102
B1PNC9/4-216 ILVPIQIGYGFHQYLPFPDG-M--SPFQAAMQDG--GYQVHRTMQFE-DGATLTGIYRYT 103
C3YRA1/51-267 ILVPHLGYAFNQYLPFPDG-M--SPFQAAMQDG--GYQVHRTMQFE-DGAFVTAHLRYT 103
C3YRA2/11-223 ILVPHLGYAYYQYLPFPDG-M--SPFQAAMQDG--GYQVHRTMQFE-DGASVTAHRYT 102
B1PND7/4-187 ILVQNIQYGFNQYLPFPDG-M--SPFQAAMQDG--GYEVHREMEFE-DGATVTGIYRYT 103
B1PND6/5-189 ILVQNIQYGFNQYLPFPDG-M--SPFQAAMQDG--GYEVHREMEFE-DGATLTGIYRYT 101
B1PND8/4-184 ILVQNIQYGFNQYLPFPDG-M--SPFQAAMQDG--GYEVHREMEFE-DGATLTGIYRYT 103
B1PND5/5-186 ILVQNIQYGFNQYLPFPDG-M--SPFQAAMQDG--GYEVHREMEFE-DGATVTGIYRYT 102
C3YRA3/4-216 IVPVNIQYGFHQYLPFPDG-M--SPFQAAMQDG--GYVHRTMQFE-DGASLTGNRYT 102
B1PND3/4-217 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG--GYVMHRAMRFE-DGAMLHSDHRYT 103
B1PND2/5-217 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG--GYVMHRAMRFE-DGAMLHSDHRYT 102
B1PNC9/4-217 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG--GYVMHRAMRFE-DGAMLHSDHRYT 103
B1PNC6/5-217 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG--GYVMHRAMRFE-DGAMLHSDHRYT 102
B1PNC7/4-220 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG--GYVMHRAMRFE-DGAMLHSDHRYT 103
B1PND0/5-220 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG--GYVMHRAMRFE-DGAMLHSDHRYT 102
B1PNC5/5-220 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG--GYVMHRAMRFE-DGAMLHSDHRYT 102
B1PND1/5-217 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG--GYVMHRAMRFE-DGAMLHSDHRYT 102
B1PND4/5-217 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG--GYVMHRAMRFE-DGAMLHSDHRYT 102
B1PNC8/4-69 ILVPIQIGYGFHQYLPFPDG-M--SPYQAAMHDG----- 66
C3YRA0/8-104 LMVPHIGYGFHQYLPFPDGEM--SPYQAAMHDG--G-----AQ 91
C3YKQ3/5-218 LMIPLHIGYGFHQYLPFPDG-P--SPFQTSMLG--GYAVYRVDFE-DGGKLTTEFKYS 101
C3YK7/5-217 LMIPLHIGYGFHQYLPFPDG-P--SPFQTSMLG--GYAVYRVDFE-DGGKLTTEFKYS 101
C3YKQ2/53-191 -IGPNLGY-----QVHRVFKFE-DGGVLNCFRYT 28
A8QZ9/4-215 LIGPNLGYGFHQYLPFPDG-A--SPFQTAITDG--GYQVHRVFKFE-DGGVLNCFRYT 101
C3YK9/5-215 LIGPNLGYGFHQYLPFPDG-A--SPFQTAITDG--GYQVHRVFKFE-DGGVLNCFRYT 100
C3YKQ1/4-215 LIGPNLGYGFHQYLPFPDG-A--SPFQTAITDG--GYQVHRVFKFE-DGGVLNCFRYT 101
C3ZLQ0/369-585 LVGPHLGYGHYQYLPFPDG-P--SPFQA AVNNG--GYQMHRSFNFE-DGAVLTATYNY 105
C3YK10/4-218 LVGPHLGYGHYQYLPFPDG-A--SPFQA AVNNG--GYQMHRSFNFE-DGAVLTATYNY 104
C3YKQ0/5-214 ILAPNLGYGHYQYLPFPDG-T--SPYQAAITNG--VYQKHTFKFE-DGGVMTINFRT 100
Q6WV10/1-215 LLSHILGYGYYHFATFPAGYE--NIYLHAMKNG--GYSNVRTERYE-DGGIISITFNRY 104
Q6WV09/1-217 LLSHILGYGYYHFATFPAGYE--NIYLHAMKNG--GYSNVRTERYE-DGGIISITFNRY 104
Q6WV11/1-215 LLSHIMZYGYYHFATFPAGYE--NVYLHAAKNG--GYNTNTRTERYE-DGGIISVNFTRY 104
Q6WV12/1-213 LLSHVMGYGFYHFGTYPGSGYE--NPLHAAINNG--GYNTNTRIEKYE-DGGVLHVSFSYR 104
Q6WV13/1-214 LLSHVMGYGFYHFGTYPGSGYE--NPLHAAINNG--GYNTNTRIEKYE-DGGVLHVSFSYR 104
Q6WV08/1-214 LLSHVMGYGFYHFGTYPGSGYE--NPLHAAINNG--GYNTNTRIEKYE-DGGVLHVSFSYR 104
Q2MHN7/1-215 LLSHCMGYGFYHFAFPKGTG--NIYLHAAINNG--GYNTNTRIEKYE-DGGIILEVNFRT 102

Q17106/5-229 -KFEGDT----LVNRIELKIDFKEDGNILGHKMEYNNSHNVYIMGDKPKNGIKVNFK 162
Q17105/5-229 -KFEGDT----LVNRIELKIDFKEDGNILGHKMEYNNSHNVYIMGDKPKNGIKVNFK 162
P42212/5-229 -KFEGDT----LVNRIELKIDFKEDGNILGHKLENNYNSHNVYIMADKQKNGIKVNFK 162
Q6YQZ0/5-229 -KFEGDT----LVNRIELTGTDFKEDGNILGNKMEYNNYNAHNVYIMTDKAKNGIKVNFK 162
Q8WTC5/5-229 -KFEGDT----LVNRIELKGMDFKEDGNILGHKLENNYNSHNVYIMPDKANGLKVNFK 162
Q8WTC4/5-229 -KFEGDT----LVNRIELKGMDFKEDGNILGHKLENNYNSHNVYIMPDKANGLKVNFK 162
Q8WTC9/5-229 -KFEGDT----LVNRIELKGMDFKEDGNILGHKLENNYNSHNVYIMPDKANGLKVNFK 162
Q8WTD0/5-229 -KFEGDT----LVNRIELKGMDFKEDGNILGHKLENNYNSHNVYIMPDKANGLKVNFK 162
Q8WTC8/5-229 -KFEGDT----LVNRIELKGMDFKEDGNILGHKLENNYNSHNVYIMPDKANGLKVNFK 162
Q8WTC6/5-229 -KFEGDT----LVNRIELKGMDFKEDGNILGHKLENNYNSHNVYIMPDKANGLKVNFK 162
Q8WP95/5-229 -KFEGDT----LVNRIELKGMDFKEDGNILGHKLENNYNSHNVYIMPDKANGLKVNFK 162
Q8WTC7/5-229 -KFEGDT----LVNRIELKGMDFKEDGNILGHKLENNYNSHNVYIMPDKANGLKVNFK 162
Q6RYS7/5-229 -TFENG-----VYNRVKNLQKQKDGHLVGNLEFNFTPHCLYIWDQANHGLKSAFK 160
Q6RYS6/39-258 -TMESGA-----IVNRINVKGTGFDKDGHCQKNESSP-PSITYVYVPE--GEGIRIYR 157
Q6RYS5/2-222 -TLESQA-----IYNRISVKGVGFKKDGNCCKRKHSSP-PQVSYVYVY--GEGIRIYR 157
Q6RYS4/5-225 -ELDGTG-----VISRITVNCDFGQPDGPIKMDQL--VDILPTETHMFPHGSAVRQLCYI 159
Q2TCH5/5-225 -ELDGTG-----VVSRTVNCDFGQPDGPIKMDQL--VDILPNETHMFPHGSAVRQLAFI 159
Q53UG7/1-221 -TMKKDG--SNCFYVEIRFDGVNFPANGPVMQRKTKVKEP--STEKMY--VRDGLVKGDN 161
Q53UG8/1-221 -TMKKDG--SNCFYVEIRFDGVNFPANGPVMQRKTKVKEP--STEKMY--VRDGLVKGDN 161
Q8MU48/1-218 -KMGD-----CFSYVEIRFDGVNFPANGPVMQRKTKVKEP--CTXEMY--VRDGLVKGGLN 158
A8CLW1/1-218 -TLEGDC-----FVYKIRFDGVNFPANGPVMQRKTKVKEP--STEKLY--VRDGLVKGDN 158

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

B6CTZ4/1-218 -TLDGDC----FVYKIRFDGVNFPKSPVLQKKTQKWE-STEKLY--VRDGLLKGVDN 158
Q5TL66/1-218 -TLDGDC----YIYEIRFDGVNFPANGPVMQKRTVKWE-STEKLY--VRDGLVKGVDN 158
A8CLQ1/1-223 -TLLKDTDDSNFYFYKIRFDGVNFAANGPVMQKKTAKWE-STEKMY--VRDGLVKGVEV 163
Q2LI65/1-219 -TLVGDVFSRS----EIRFEGVNFANGPVMQKKTQKWE-STEKMY--VRDGLVKGVDN 159
A8CLW4/1-222 -TLSKDI--ANCFDYNIRFDGVNFPKSPVLQKKTQKWE-STEMMY--VRDGLVKGVDN 161
B6CTZ5/4-224 -TLSKVI--DNCFHYNIRFDGVNFPKSPVLQKKTQKWE-STEMMY--VRDGLVKGVDN 161
A3F207/1-221 -KLLKDE--ENCFHKNIRFDGVNFPANSPVMLKTTQKWE-STEKMY--ARDGVVKGVDN 161
Q962P9/1-221 -TLMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEKMY--VRDGLVKGVDN 161
Q7Z0W8/1-221 -TLMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEKMY--VRDGLVKGVDN 161
Q66ND5/1-221 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEKMY--VRDGLVKGVDN 161
Q66ND4/1-221 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEKMY--VRDGLVKGVDN 161
Q66ND2/1-221 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEKMY--VRDGLVKGVDN 161
A7UAL7/1-221 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEKMY--VRDGLVKGVDN 161
Q7Z0W6/1-221 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEKMY--VRDGLVKGVDN 161
A7UAL9/1-221 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEMMY--VRDGLVKGVDN 161
Q875F2/8-228 -TMEGV--DDCFAYKIRFDGVNFPANGPVMQKKTQKWE-STEMMY--ARDGLVKGVDN 161
Q7Z0W7/8-228 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEMMY--VRDGLVKGVDN 161
Q8MU47/8-228 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEKMY--XRDGLVKGVDN 161
Q95VT0/1-221 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEMMY--ARDGLVKGVDN 161
A7UAL3/1-221 -TMMKGV--DDCFVYKIRFDGVNFPANGPVMQKKTQKWE-STEMMY--VRDGLVKGVDN 161
A7UAL6/1-218 -KLEG--DCFFYEIRFYGVNFPSSGPMQKKTQKWE-STEMMY--VRDGLVKGVDN 158
A7UAL8/1-218 -KLEG--DCFFYEIRFYGVNFPSSGPMQKKTQKWE-STEMMY--VRDGLVKGVDN 158
A8CL52/1-219 -TLEGDT----FFNKVRFYGVNFPNGPVMQKKTQKWE-STEKMY--LRDGLVTDIT 158
Q816J8/1-219 -TLEGDT----FFNKVRFYGVNFPNGPVMQKKTQKWE-STEKMY--LRDGLVTDIT 158
A3F208/1-219 -TMEGDT----FVNKVRFYGVNFPNGPVMQKKTQKWE-STEKMY--VRDGLVTDIT 158
A8CLR7/1-219 -KMGVDT----FYNTVRFYGVNFPNGPVMQKKTQKWE-STEKMY--VRDGLVTDIT 158
Q556Z9/1-219 -TMEGDT----FVNKVRFHGVNFPANGPVMQKKTQKWE-STEKMY--VRDGLVTDIT 158
Q66PW1/5-223 -KMGVDT----FHNEVQFYGVNFPANGPVMQKKTQKWE-STEKMY--VRDGLVTDIT 158
Q875F1/1-219 -TMEGDC----FFNKVRFYGVNFPNGPVMQKKTQKWE-STEKMY--VRDGLVTDIT 158
Q7Z0W9/1-219 -TMEGDC----FFNKVRFYGVNFPNGPVMQKKTQKWE-STEKMY--VRDGLVTDIT 158
Q6USK3/1-219 -TMEGDC----FFNKVRFYGVNFPNGPVMQKKTQKWE-STEKMY--VRDGLVTDIT 158
A7UAL4/1-219 -TMEGDC----FFNKVRFYGVNFPNGPVMQKKTQKWE-STEKMY--VRDGLVTDIT 158
A7UAM0/1-219 -KMEGDC----FIYKIRFDGVNFPKSPVMQKKTQKWE-STEKMY--VRDGLVKGDI 158
Q7Z0W5/1-218 -KLEGDC----FFYEIRFYGVNFPSSGPMQKKTQKWE-STEMMY--VRDGLVKGDI 158
Q95UA7/1-218 -KLEGDC----FFYEIRFYGVNFPSSGPMQKKTQKWE-STEMMY--VRDGLVKGDI 158
A7UAL1/1-218 -KLEGDC----FFYEIRFYGVNFPSSGPMQKKTQKWE-STEMMY--VRDGLVKGDI 158
A8CLP7/1-218 -KIEGDC----FTYDIFRFGVNFPPGPMQKKTQKWE-STEMMY--VRDGLVGEVE 158
A7UAL2/1-218 -KLEGDC----FFYEIRFYGVNFPSSGPMQKKTQKWE-STEMMY--VRDGLVKGDI 158
A7UAL5/1-218 -KMEGDC----FIYEIQFHGVNFPNGPVMQKKTQKWE-STEKMY--VRDGLVKGDI 158
Q963F5/1-218 -KMEGDC----FIYEIQFHGVNFPNGPVMQKKTQKWE-STEKMY--VRDGLVKGDI 158
Q7Z0W4/1-218 -RMEGDC----FIYEIRYDGVNFPSSGPMQKKTQKWE-STEKMY--VRDGLVKGDI 158
Q66ND3/1-218 -RMEGDC----FIYEIRYDGVNFPSSGPMQKKTQKWE-STEKMY--VRDGLVKGDI 158
Q2LI64/1-35 -----
Q60I25/1-218 -SMRGDC----FFYDIFRFGVNFPPGPMQKKTQKWE-STEKMY--VRDGLVKGDI 158
Q60I24/1-218 -SMRGDC----FFYDIFRFGVNFPPGPMQKKTQKWE-STEKMY--VRDGLVKGDI 158
A8CLT2/1-218 -SMRGDC----FFYDIFRFGVNFPPGPMQKKTQKWE-STEKMY--VRDGLVKGDI 158
B5BR3/1-218 -SMEGDC----FIYKIRFDGVNFPNGPVMQKKTQKWE-STEMMY--VRDGLMVDN 158
B6CTZ6/1-219 -SMEGDC----FIYKIRFDGVNFPNGPVMQKKTQKWE-STEMMY--ARDGLMVDN 158
Q9U6Y3/43-260 -SMEEDS----FIYEIRFDGMNFPNGPVMQKKTQKWE-STEMMY--VRDGLVGDII 158
Q876U0/1-218 -SLEGDC----FFQNVRFYGVNFPNGPVMQKKTQKWE-STEKLY--VRDGLVGNIN 158
Q875F0/7-229 -KMGVDE----FQYNIRFDGVNFPEDGPMQKKTQKWE-STEMMY--VQGGVKGVEV 160
Q875F3/6-223 -TMVGGC----FQYDIFRFGVNFPPGPMQKKTQKWE-STEMMY--MQNGVKGVEV 159
Q66PW0/1-221 -SVVGDG----FNYEIHFMGANFPNGPVMQKKTQKWE-STEKMY--ERDGLRGDVP 161
Q8MU46/1-221 -SVVGDG----FNYEIHFMGANFPNGPVMQKKTQKWE-STEKMY--ERDGLRGDVP 161
Q875E9/1-221 -SVVGDG----FYYIRFNGENFPNGPVMQKKTQKWE-STEMMY--ERDGLRGDIA 161
Q875E8/1-221 -RVVDGT----FNYDIHFMGADPLNFPVMQKKTQKWE-STEMMY--QCDGLRGDIA 161
Q876T8/1-221 -SVVGDG----FNYDIHFMGANPLDGPVMQKKTQKWE-STEMMY--ERDGLRGDIA 161
Q66PU5/1-220 -RLDQKEM---CFIYDVRFHGVNFPANGPVMQKKTQKWE-STEKMY--ARDGLVKGDI 160
A8CL57/1-220 -RLDQKEM---CFIYDVRFHGVNFPANGPVMQKKTQKWE-STEMMY--ARDGLVKGDI 160
Q8MAA2/34-254 -RLEEER---RFVNVRFHGVNFPNGPVMQKKTQKWE-STEMMY--PRDGLFEGVD 160
Q8MAA1/12-234 -RLEEER---RFVNVRFHGVNFPNGPVMQKKTQKWE-STEMMY--PRNGLLEGYDE 163
Q66ND6/5-222 -SLTGNC----FNYDIKFTGLNFPNGPVMQKKTQKWE-STERLY--PRDGLVIGDII 158
Q9U6Y7/5-223 -SLTGNC----FNYDIKFTGLNFPNGPVMQKKTQKWE-STERLY--PRDGLVIGDII 158
Q553G7/6-223 -SLQDGC----FIYKVKRGTNFPSDGPMQKKTQKWE-SSERMY--PEDGALKGEMK 158
Q553G8/6-223 -SLQDGC----FIYKVKRGTNFPSDGPMQKKTQKWE-SSERMY--PEDGALKGEMK 158
Q2LI66/5-222 -SLQDGC----FIYKVKRGTNFPSDGPMQKKTQKWE-STERLY--PRDGLVIGDII 158
Q66ND8/5-222 -SLQDGC----FIYKVKRGTNFPSDGPMQKKTQKWE-STERLY--PRDGLVIGDII 158
Q66ND7/5-222 -SLQDGC----FIYKVKRGTNFPSDGPMQKKTQKWE-STERLY--PRDGLVIGDII 158
Q9U6Y8/5-222 -SLQDGC----FIYKVKRGTNFPSDGPMQKKTQKWE-STERLY--PRDGLVIGDII 158
Q6K85/5-222 -SLQDGC----FIYKVKRGTNFPSDGPMQKKTQKWE-STERLY--PRDGLVIGDII 158
Q2W674/7-224 -SLQDGC----FIYKVKRGTNFPSDGPMQKKTQKWE-STERLY--PRDGLVIGDII 158
Q9GT37/5-223 -SLQDGC----FIYKVKRGTNFPSDGPMQKKTQKWE-STERLY--PRDGLVIGDII 158
A8CLM7/1-218 -SIQGNC----FIYNVKISGVNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
A8CLM1/1-218 -SIQGNC----FIYNVKISGVNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
Q1JU63/2-219 -SIQGNC----FIYNVKISGVNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
Q1JV71/2-219 -SIQGNC----FIYNVKISGVNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
Q1JV72/2-219 -SIQGNC----FIYNVKISGVNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
Q1JV70/2-219 -SIQGNC----FIYNVKISGVNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
B5T1L1/1-218 -SIQGNC----FIYHVKFSGLNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
A8CLV6/1-218 -SIQGNC----FIYHVKFSGLNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
Q66PW0/1-218 -SIQGNC----FIYHVKFSGLNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
Q66PU8/1-218 -SIQGNC----FIYHVKFSGLNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158
Q95P04/1-218 -SIQGNC----FIYHVKFSGLNFPNGPVMQKKTQKWE-STERLY--ARDGMLGNIN 158

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

A8CLL3/1-218 -SIQGN-----FIYNVKSGLNFPSPGPMQKKTQGWEP-NTERLL--ARDGMLIGNNF 158
Q66P1/1-218 -SIQGN-----FIYHVKFSGVNFPNPGPMQKKTQGWEP-NTERLF--ARDGMLIGNNF 158
Q66P9/1-218 -SIQGN-----FIYHVKFSGLNFPNPGPMQKKTQGWEP-NTERLF--ARDGMLIGNNF 158
P83690/1-218 -SIQGN-----FTYHVKFSGLNFPNPGPMQKKTQGWEP-HSERLF--ARGGMLIGNNF 158
A8CLN8/6-226 -RLDGD5-----FIYEVRF LGVNFPRDGPVMQKKTTRGWD-STERLY--ECGGWQRGDVH 158
B6CT27/1-220 INLQENC-----FIYDVKFHGVNLPDGPVMQKTIAGWEP-SVETLY--VRDGMKSDTA 160
Q9U6Y5/5-228 VSVEENC-----MYHESKFYGVNFPADGPVMKMTDNWEP-SCEKIIPVPKQIGILKGDVS 162
Q9U6Y4/5-228 VSVKENC-----IYHKSIFNGMNFADGPVMKMTTNWEA-SCEKIMPVPKQIGILKGDVS 162
Q8T4U4/5-228 VNVRENC-----IYHSTFYGVNFPADGPVMKMTTNWEP-SCEKIIPINSQKILKGDVS 162
B2ZAG2/5-226 V--KENC-----IHHSTFYGVNFPDGPVMKKTVTYNWEP-SCEKIIPISQKILKGDVS 160
Q692V6/5-228 VSVEENC-----FYHNSKFHGVNFPADGPVMKMTTNWEP-SCEKIIPVPRQIGILKGDIA 162
B5LIN1/1-161 VSVEENY-----FYHNSKFHGVNFPADGPVMKMTTNWEP-SCEKIIPVPRQIGILKGDIA 99
B5T1L5/5-228 VSVEENC-----FYHNSKFHGVNFPADGPVMKMTTNWEP-SCEKIIPVPRQIGILKGDIA 162
Q66PV2/5-228 VSVEENC-----FYHNSKFHGVNFPADGPVMKMTTNWEP-SCEKIIPVPRQIGILKGDIA 162
Q86LC2/5-228 LSVVEERC-----FYHESKFGVNFADGPVMKMTTNWEP-SCEKIIPVPRQIGILKGDVS 162
B2ZAG1/5-228 LSVQEK-----FHHVSRFYGVNFPADGPVMKMTTNWEP-SCEKIIPVPSQIGILKGDAS 162
A8CLN4/5-231 VSVEENC-----FYHESKFHGVNFPADGPVMKMTTNWEP-SCEKIIPVPRQIGILKGDVT 163
A8CLV1/5-230 VSVEENC-----FYHESKFHGVNFPADGPVMKMTTNWEP-SCEKIIPVPRQIGILKGDVT 162
A8CLU7/1-223 LSSDKS-----FEHKSFLGVNFPADGPVMKKTETTNWEP-SCEKMT--PNGMTLIGDVT 160
A8CLP2/5-228 LSADKKS-----FEHKSFLGVNFPADGPVMKKTETTNWEP-SCEKMT--PNGMTLIGDVT 161
B6CT22/5-226 LSDDKAS-----FHHKSFFGVNFPDGPVMKKTETTNWEP-SCEKMT--PSGKTLKGDVI 160
Q2VTM9/5-219 LD--KNC-----FVHKSFTFGVNFANGPMQKKTTLNWE-SSELIT--AGDGILKGDVT 158
Q1ALD3/5-219 LD--KDC-----FIHNSFTFGVNFANGPMQKKTAMNWE-SSELIT--PCDGILKGDVT 158
Q1ALD5/5-219 LE--KNC-----FVHQSFTFGVNFANGPMQKKTALNWE-SSELIT--PCDGILKGDVT 158
Q1ALD4/5-219 LD--KNG-----FVHNSFTFGVNFANGPMKKTGVNWE-SSEKIT--ACDGILKGDVT 158
Q09HR5/5-219 LN--KTG-----FVHKSFTFGVNFANGPMQKKTTLNWE-SSEKIT--ACDGILKGDVT 158
Q2VTM8/5-219 LT--KNC-----FVHNSFTFGVNFANGPMQKKTTLNWE-SSEKIT--PFEGNLKGDVT 158
A8CLU1/1-218 LDSASNC-----FIHKSFFGVNFPDGPVMKKTETTNWEP-SSEKIT--VRDGLKGDVT 159
Q9U6Y6/5-224 LKG--NC-----FEHKSFTFGVNFADGPVMKKTGWDP-SFEKMT--VCDGILKGDVT 160
A8CLT7/5-224 LKG--NC-----FVHKSFMHGVNFPADGPVMKKTGWDP-SFEKMT--VCNGILKGDVT 158
Q66ND1/5-220 LEG--NW-----FKHSMFHGVNFPANGPMQKKTIGWEP-SFEKMT--VSNILKGDVT 158
Q66PV5/5-226 LER--DC-----FIHKSIFHGVNFPADGPVMKKTIGWEP-AFEKMT--VSKDVLKGDVT 160
B5T1L2/5-226 LER--DC-----FIHKSIFHGVNFPADGPVMKKTIGWEP-AFEKMT--VSKDVLKGDVT 160
Q66PV3/5-226 LER--DC-----FIHKSIFHGVNFPADGPVMKKTIGWEP-AFEKMT--VSKDVLKGDVT 160
B5T1L0/5-226 LER--DC-----FIHKSIFHGVNFPADGPVMKKTIGWEP-AFEKMT--VSKDVLKGDVT 160
Q66PV4/5-226 LER--DC-----FIHKSIFHGVNFPADGPVMKKTIGWEP-AFEKMT--VSKDVLKGDVT 160
B5LIN0/5-226 LER--DC-----FVHKSIFHGVNFPADGPVMKKTIGWEP-AFEKMT--VSKDVLKGDVT 160
B5T1L3/5-226 LEG--DC-----FIHKSIFHGVNFPADGPVMKKTIGWEP-SFEKMT--VSKDVLKGDVT 160
B5T1L4/5-226 LEG--NC-----FIHNSIFHGVNFPADGPVMKKTIGWEP-SFEKMT--VSKEVLRGDVT 160
Q66P7/5-226 LEG--NC-----FIHNSIFHGVNFPADGPVMKKTIGWEP-SFEKMT--VSKEVLRGDVT 160
A8CLY3/5-226 LEG--NC-----FIHKSIFHGVNFPADGPVMKKTIGWEP-SFEKMT--VSKEVLRGDVT 160
Q66PV8/5-226 LEG--NC-----FIHKSIFHGVNFPADGPVMKKTIGWEP-SFEKMT--VSKEVLRGDVT 160
Q66PV9/5-226 LEG--NC-----FIHNSIFHGVNFPADGPVMKKTIGWEP-SFEKMT--VSKEVLRGDVT 160
A8CLX9/5-226 LEG--NC-----FIHNSIFHGVNFPADGPVMKKTIGWEP-SFEKMT--VSKEVLRGDVT 160
B6CT23/5-226 LEG--NC-----FIHNSIFHGVNFPADGPVMKKTIGWEP-SFEKMT--VSKEVLRGDVT 160
Q6I7B1/6-227 LEG--NC-----FIHNSIFHGVNFPADGPVMKKTIGWEP-SFEKMT--VAKEVLRGDVT 160
Q66PV6/5-226 LEG--NC-----FIHKSITLGVNFPADGPVMKKTIGWEP-AFEKMT--GFNEVLRGDVT 160
Q86LV8/1-213 L--TGSV-----FEHKSIFVGVNFPANGPMQKKTIGWEP-STEKMA--ANDGSVQGYDT 156
Q86LV7/1-213 L--TGNV-----FEHKSIFVGVNFPANGPMQKKTIGWEP-STEKMA--ANDGSVQGYDT 156
A8CLR1/5-220 L--EGDC-----FEHECFVGVNFPANGPMQKKTIGWEP-STEKMT--AHGKVVQGNVP 160
Q6R8F5/5-223 L--NGNC-----FVHQSIFVGVNFPADGPVMQKKTIGWEP-STEKMI--VRDGLKGDVT 158
Q6R8F4/1-219 LK--GNW-----FVHQSIFVGVNFPADGPVMQKKTIGWEP-STEKMI--VRDGLKGDVT 158
Q6R8F3/5-220 IE--KDG-----FKHRSKFVGVNFPNPGPMQKKTTRGWD-SSEKIT--VRDGLKGDVT 159
Q6I7B2/2-218 L--RGNT-----FYHKSFTGVNFPADGPVMQKKTIGWEP-STEKIT--ASDGLKGDVT 160
A7VMR3/2-218 L--RGNT-----FYHKSFTGVNFPADGPVMQKKTIGWEP-STEKIT--ASDGLKGDVT 160
Q6I7B4/1-210 L--RGNC-----FEHKSIFVGVNFPADGPVMQKKTIGWEP-STEKIT--TYDGLKGDVT 153
Q6I7B3/1-217 L--RGNC-----FEHKSIFVGVNFPADGPVMQKKTIGWEP-STEKIT--TCDGLKGDVT 160
A8CLQ6/5-221 L--KGDC-----FEHNSKFVGVNFPADGPVMQKKTIGWEP-STEKIT--VSDGLKGDVT 160
A8CLW9/4-223 L--IGNR-----FEHKSIFVGVNFPADGPVMQKKTIGWEP-SSEKIT--PRDGLKGYVP 158
Q9S86/2-220 -SLEGN-----LTYKVKVLTGNFPADGPVMKKTIGWEP-CTEIVY--QDNGVLRGRNV 158
Q9S11/2-220 -SLEGN-----LNYKVKVLTGNFPADGPVMKKTIGWEP-CTEIVY--QDNGVLRGRNV 158
Q8MU45/2-220 -SLEGN-----PIYKVKVLTGNFPAASPVMKKTIGWEP-STEIVY--QDNGVLRGRNV 158
Q9S85/2-220 -SLEGN-----LTYKVKVLTGNFPADGPVMKKTIGWEP-CTEVVY--PENGVLKGRNV 158
ASYTR7/2-222 -SLDGD-----LVYKVKILGNFPADGPVMQKKTIGWEP-ATEIVY--EVDGLKGRQSL 158
Q9G28/2-222 -SLDGD-----LVYKVKILGNFPADGPVMQKKTIGWEP-ATEIVY--EVDGLKGRQSL 158
ASYTR8/2-222 -SLDGD-----LVYKVKILGNFPADGPVMQKKTIGWEP-ATEIVY--EVDGLKGRQSL 158
Q9GP15/2-222 -SLDGD-----LVYKVKILGNFPADGPVMQKKTIGWEP-ATEIVY--EVDGLKGRQSL 158
Q2VFP3/2-221 -SLQGNN-----FIFKVVNIGANFPANGPMQKKTIGWEP-CVEMLY--PRDGLKGRQSL 158
A0AQ7/2-222 -KLDGNC-----LVYNIKILGCFNFPNPGPMQKKTIGWEP-CCEMRY--TRDGLKGRQSL 158
A0AQ8/2-222 -KLDGNC-----LVYNIKILGCFNFPNPGPMQKKTIGWEP-CCEMRY--TRDGLKGRQSL 158
Q8T5F8/2-222 -SLEDGC-----LVYHAKVGTGNFSPNGAVMQKKTIGWEP-NTEMLY--PADGGLRGYSQ 158
Q86LV4/2-220 -SLEDGC-----IICNIKVHGTNFPADGPVMQKKTIGWEP-STETVI--PRDGLKGRQSL 158
Q5QIU5/2-220 -SLKGD-----IICNIKVHGTNFPADGPVMQKKTIGWEP-STETVI--PRDGLKGRQSL 158
Q8T6T9/2-220 -SLEDGC-----IICKIKAHGTNFPADGPVMQKKTIGWEP-STETVI--PRDGLKGRQSL 158
Q8T5E7/2-220 -SLKGD-----IICNIKVHGTNFPADGPVMQKKTIGWEP-STETVI--PQDGGIVAARS 159
ASYTR6/2-219 -SLQGD-----IICKVKVLTGNFPANGPMQKKTIGWEP-STETVI--PRDGLKGRQSL 158
Q9GP16/2-219 -SLQGD-----IICKVKVLTGNFPANGPMQKKTIGWEP-STETVI--PRDGLKGRQSL 158
Q5ZQ06/2-221 -TLKGD-----ITTYSEVTGT--FPPNGPMKKTIGWEP-SCENVI--PRDGLKGRQSL 157
B9UPG6/1-219 -TFKDN-----VYGDVWALGVNFPNPGPMKKTIGWEP-AEETLI--PQNGVLKGRQSL 158
Q5ZQ05/1-219 -TYKDK-----LHGDVWALGVNFPNPGPMKKTIGWEP-AEETLI--AKNGVLKGRQSL 158
Q7Z168/1-218 -TYKDN-----LHGDVWAVGVNFPNPGPMKKTIGWEP-TEETFT--PKNGVLKGRQSL 157
Q9BLZ0/8-225 -SLEDD-----FHYKVEYRGNFSPNGPMQKKTIGWEP-SFEVYV--MNSGVLKGRQSL 158
Q9BLY9/8-225 -NLIED-----FVYRVEYKGSNFPDGPVMQKKTIGWEP-SFEAMY--MNGVLKGRQSL 158

Q963I9/5-223 -SLEDGK----FIVNVDFKAKDLRRMGPMVQQDIVGMQP--SYESMY--TNVTSVIGECI 158
A8CLX6/5-222 -SLKGN-----LVHKTNFQGNFIDGPMVQKRTLWEP--TSEKMT--PCDGIIGDITI 158
A753R2/57-279 LKDG-----LHYHTNFHGINLKPDPVPMQKRTMGWLP--SVETNI--PRRDTLLGDIN 159
A7RGQ5/1-174 LKNR-----LHYHTTFHGINLNPDKPLMQRKRTMGWLP--SVETNI--PRGDTLVGDIN 147
A751H5/1-224 LDKDG-----LHYHMTFQGINLDPNGPMVKMTMGWLP--SVETNI--PRGTLVGDIN 160
A751H3/1-82 HQNGS-----LHYHVTFHGINPKTDGPMVQKITMGWSP--SVE----- 82
B1PNC2/5-218 YEGSHIKG-----EFQVNGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 152
B1PNC0/4-217 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 153
B1PNC1/5-218 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 152
B1PNC4/5-218 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 152
B1PNC4/5-218 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 152
C3YR99/48-261 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 154
B1PNC3/4-217 YEGTHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 154
B1PNC8/5-217 YEGTHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 154
B1PNC9/4-216 YEGTHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 154
C3YRA1/51-267 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 154
C3YRA2/11-223 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 153
B1PND7/4-187 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 154
B1PND6/5-189 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 152
B1PND8/4-184 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 154
B1PND5/5-186 YEGSHIKG-----EFQVIGTGFPADGPVMTNSLTTADWCVTMPLY--PNDKTIISTFD 153
C3YRA3/4-216 YDGGHIK-----EFHVGSGFLDGGPVMTKSLTAVDWSVATMLF--PNDTIVSTID 153
B1PND3/4-217 YNGNHIK-----EFRLTSGGFPADGPVMTNSLTTADWCVDKLLY--PDNTIIGKFD 154
B1PND2/5-217 YNGNHIK-----EFRLTSGGFPADGPVMTNSLTTADWCVDKLLY--PDNTIIGKFD 153
B1PNC9/4-217 YKGNHIK-----EFRLTSGGFPADGPVMTNSLTTADWCVDKLLY--PDNTIIGKFD 154
B1PNC6/5-217 YKGNHIK-----EFRLTSGGFPADGPVMTNSLTTADWCVDKLLY--PDNTIIGKFD 153
B1PNC7/4-220 YKGNHIK-----EFRLTSGGFPADGPVMTNSLTTADWCVDKLLY--PDNTIIGKFD 154
B1PND0/5-220 YKGNHIK-----EFRLTSGGFPADGPVMTNSLTTADWCVDKLLY--PDNTIIGKFD 153
B1PNC5/5-220 YKGNHIK-----EFRLTSGGFPADGPVMTNSLTTADWCVDKLLY--PDNTIIGKFD 153
B1PND1/5-217 YNGNHIK-----EFRLTSGGFPADGPVMTNSLTTADWCVDKLLY--PDNTIIGKFD 153
B1PND4/5-217 YNGNHIK-----EFRLTSGGFPADGPVMTNSLTTADWCVDKLLY--PDNTIIGKFD 153
B1PNC8/4-69 ----- 97
C3YRA0/8-104 YDGNHE----- 97
C3YKQ3/5-218 YEGSHIKA-----DMKLMGSGFPDDGPVMTSQQIVDQDGCVSCKTY--LNNNTIVDSFD 152
C3YK7/5-217 YEGTHIKS-----DFKLMGSGFPDDGPVMTSQQIVDQDGCVSCKTY--LNDNTIVDSFD 152
C3YKQ2/53-191 YEGGKIKG-----EFQLIGSGFPAGGPVMSGGLTTLDRSVAKLQC--SDDCTITGTNN 79
A8QV29/4-215 YEGGKIKG-----EFQLIGSGFPAGGPVMSGGLTTLDRSVAKLQC--SDDCTITGTNN 152
C3YK9/5-215 YEGGKIKG-----EFQLIGSGFPAGGPVMSGGLTTLDRSVAKLQC--SDDCTITGTNN 151
C3YKQ1/4-215 YEGGKIKG-----EFQLIGSGFPAGGPVMSGGLTTLDRSVAKLQC--SDDCTITGTNN 152
C3ZL00/369-585 YSGGKIQ-----EFHLVSGFCPPDGPVMTNLTGLDRSVAKLQC--VSDDKLAEFVD 156
C3YK10/4-218 YSGGKIQ-----EFHLVSGFCPPDGPVMTNLTGLDRSVAKLQC--VSDDKLAEFVD 155
C3YKQ0/5-214 YSGNKIKG-----EFHVGSGFPDDGPVMTNSLQQHDHNVRLMV--LGDKTIIGSDNM 151
Q6W10/1-215 YEGNKIG-----DFKVVGTGFPPTNSLIFTDKIISNPTCEMFY--KADNVLVNAVY 155
Q6W09/1-217 YEGNKIG-----DFKVVGTGFPPTNSLIFTDKIISNPTCEMFY--KADNVLVNAVY 155
Q6W11/1-215 YEGNKIG-----DFKVVGTGFPPTNSLIFTDKIISNPTCEMFY--KADNVLVNAVY 155
Q6W12/1-213 YEGNKIG-----DFKVVGTGFPPTNSLIFTDKIISNPTCEMFY--KADNVLVNAVY 155
Q6W13/1-214 YEGNKIG-----DFKVVGTGFPPTNSLIFTDKIISNPTCEMFY--KADNVLVNAVY 155
Q6W08/1-214 YEGNKIG-----DFKVVGTGFPPTNSLIFTDKIISNPTCEMFY--KADNVLVNAVY 155
Q2MH7/1-215 YEFNKIIG-----DVEICIGHFSPQSPFKDITVKSCTPVDLMLP--MSGNIIASSYA 153

Q17106/5-229 IRHNKIDG---SVQLADHYQQNTPIGD--GPVLLPDN--HYLSTQSALSQDP--HGKRDHML 216
Q17105/5-229 IRHNKIDG---SVQLADHYQQNTPIGD--GPVLLPDN--HYLSTQSALSQDP--NEKRDHML 216
P42212/5-229 IRHNIEEG---SVQLADHYQQNTPIGD--GPVLLPDN--HYLSTQSALSQDP--NEKRDHML 216
Q6Y20/5-229 IRHNIEEG---SVQLADHYQQNTPIGD--GPVLLPDN--HYLSTQSALSQDP--NEKRDHMY 216
Q8WT5/5-229 IRHNIEEG---GVQLADHYQTNVPLGD--GPVLIPIIN--HYLSYQTAISKDR--NETRDHMF 216
Q8WT4/5-229 IRHNIEEG---GVQLADHYQTNVPLGD--GPVLIPIIN--HYLSYQTAISKDR--NETRDHMF 216
Q8WT9/5-229 IRHNIEEG---GVQLADHYQTNVPLGD--GPVLIPIIN--HYLSYQTAISKDR--NETRDHMF 216
Q8WT0/5-229 IRHNIEEG---GVQLADHYQTNVPLGD--GPVLIPIIN--HYLSYQTAISKDR--NETRDHMF 216
Q8WT8/5-229 IRHNIEEG---GVQLADHYQTNVPLGD--GPVLIPIIN--HYLSYQTAISKDR--NETRDHMF 216
Q8WT6/5-229 IRHNIEEG---GVQLADHYQTNVPLGD--GPVLIPIIN--HYLSYQTAISKDR--NETRDHMF 216
Q8WP95/5-229 IRHNIEEG---GVQLADHYQTNVPLGD--GPVLIPIIN--HYLSYQTAISKDR--NETRDHMF 216
Q8WT7/5-229 IRHNIEEG---GVQLADHYQTNVPLGD--GPVLIPIIN--HYLSYQTAISKDR--NETRDHMF 216
Q6RY57/5-229 IMHEITGSKE--DFIVADHTQMNTPIGG--GPVHVPEY--HHITYHTVLSKDV--TDHRDNMSL 216
Q6RY56/39-258 NIYPTKDG---HYVADTQVNRPIRAQGTSAIPTY--HHIKSKVDLSTDP--EENKDHIII 212
Q6RY55/2-222 NIYPTKDG---GYVADTRQVNRPIKAEGKSAIPKY--HYIKSKIDLSTDP--NERKDHIII 212
Q6RY54/5-225 GFTTADGG---LMMSHFDSKLTFNRSRAIKIPGP--HFVTIITKQMDT--SDKRDHVCQ 212
Q2TCH5/5-225 GFTTADGG---LMMGHFDSKMTFNRSRAIEIPGP--HFVTIITKQMDT--SDKRDHVCQ 212
Q53UG7/1-221 MALLLQGG---GHYRCDFRTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 212
Q53UG8/1-221 MALLLQGG---GHYRCDFRTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 212
Q8MU48/1-218 MALLLEGG---GHFRCDLKTYYKAK--KVV--QMPDY--HFVNHRIEIVSHD--EDYNNVKL 209
A8CLW1/1-218 MALLLEGG---GHFRCDFKTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 209
B6CTZ4/1-218 MALLLEGG---GHYRCDFKTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 209
Q5TL66/1-218 MALLLEGG---GHYRCDFKTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 209
A8CLQ1/1-223 MALLLEGG---GHYRCDFKTTYKAK--KVV--RPSY--HFVDHRIEIVSHD--KDYNNVKL 214
Q2LI65/1-219 MALLLEGG---GHYRCDFKTTYKAK--KVV--QLPEY--HFVDHRIEIVSHD--RDYNNVKL 210
A8CLW4/1-222 MALLLEGG---AGHYRCDFKTTYKAK--KAV--KLPDY--HFVDHRIEIVSHD--KDYNNVKL 213
B6CTZ5/4-224 MALLLEGG---GHYRCDFKTTYKAK--KVV--ELPDY--HFVDHRIEIVSHD--KDYNNVKL 212
A3F207/1-221 MALLLQGG---GHYRCDFKTTYKAK--KVV--PLPDY--HFVDHRIEIVSHD--KDYNNVKL 212
Q962P9/1-221 MALLLEGG---GHYRCDFKTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 212
Q7Z0W8/1-221 MALLLEGG---GHYRCDFKTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 212
Q66ND5/1-221 MALLLEGG---GHYRCDFKTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 212
Q66ND4/1-221 MALLLEGG---GHYRCDFKTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 212
Q66ND2/1-221 MALLLEGG---GHYRCDFKTTYKAK--KVV--QLPDY--HFVDHRIEIVSHD--KDYNNVKL 212

Supplementary Material (ESI) for Molecular BioSystems
This journal is (c) The Royal Society of Chemistry, 2011

A7UAL7/1-221 MALLLEGG---GHYRCDFKTTYKAK-KVV--QLPDY-HFVDHRIEIVSHD--KDYNKVKL 212
Q7Z0W6/1-221 MALLLEGG---GHYRCDFKTTYKAK-KVV--QLPDY-HFVDHRIEIVSHD--KDYNKVKL 212
A7UAL9/1-221 MALLLEGG---GHYRCDFKTTYKAK-KVV--QLPDY-HFVDHRIEIVSHD--KDYNKVKL 212
Q8T5F2/8-228 MALLLEGG---GHYRCDFKTTYKAK-KVV--RLPDY-HFVDHRIEIVSHD--KDYNKVKL 212
Q7Z0W7/8-228 MALLLEGG---GHYRCDFKTTYKAK-KVV--RLPDY-HFVDHRIEIVSHD--KDYNKVKL 212
Q8MU47/8-228 MALLLEGG---GHYRXDXKTTYRAK-KVV--QLPDY-HFVDHRIEIVSHD--KDYNKVKL 212
Q95VT0/1-221 MALLLEGG---GHYRCDFKTTYKAK-KVV--RLPDY-HFVDHRIEIVSHD--KDYNKVKL 212
A7UAL3/1-221 MALLLEGG---GHYRCDFKTTYKAK-KVV--RLPDY-HFVDHRIEIVSHD--KDYNKVKL 212
A7UAL6/1-218 MALLLEGG---GHYRCDFKTTYKAK-KFV--QLPDY-HFVDHRIEIVSHD--KDYNKVKL 209
A7UAL8/1-218 MALLLEGG---GHYRCDFKTTYKAK-KVV--QLPDY-HFVDHRIEIVSHD--KDYNKVKL 209
A8CL52/1-219 MALLLEGG---VHYRCDFRTTYKSRQEGV--KLPY-HFVDHCISIVSHD--KDYTKVKL 210
Q8T6J8/1-219 MALLLEGG---VHYRCDFRTTYKSRQEGV--KLPY-HFVDHCISILRHD--KDYNEVKL 210
A3F208/1-219 MALLLEGG---VHYRCDFRTTYKTQEDV--KLPY-HFVDHCISILRHD--KDYTKVKL 210
A8CLR7/1-219 MALLLEGG---VHYRCDFRTTYKAKEKGV--KLPY-HFVDHCIEILSHD--KDYNKVKL 210
Q55629/1-219 MALLLEGG---VHYRCDFRTTYKAKEKGV--KLPY-HFVDHCIEILSHD--KDYNKVKL 210
Q66PW1/5-223 MALLLEGG---VHYRCDFRTTYKAKEKGP--KFPY-HLVDHCIEITSHD--KDYNVVEL 210
Q8T5F1/1-219 MALLLEGG---GHYRCDFRTTYRAKKGV--KLPY-HFVDHCIEILRHD--KEYTEVKL 210
Q7Z0W9/1-219 MALLLEGG---GHYRCDFRTTYRAKKGV--KLPY-HFVDHCIEILRHD--KEYTEVKL 210
Q6USK3/1-219 MALLLEGG---GHYRCDFRTTYRAKKGV--KLPY-HFVDHCIEILRHD--KEYTEVKL 210
A7UAL4/1-219 MALLLEGG---VHYRCDFRTTYKAKEKGV--ELPGY-HFVDHCIEILSHD--KDYNKVKL 210
A7UAM0/1-219 MALLLEGG---GHYRCDFRTTYKAKEKGV--ELPGY-HFVDHCIEILRHD--KDYTEVKL 210
Q7Z0W5/1-218 RTLLLEGG---KHRCNFRSTYRAK-KGV--VLPEY-HFVDHCIEILSHD--KDYNTVEV 209
Q95UA7/1-218 RTLLLEGG---KHRCNFRSTYRAK-KGV--VLPEY-HFVDHCIEILSHD--KDYNTVEV 209
A7UAL1/1-218 RTLLLEGG---KHRCNFRSTYRAK-KGV--VLPEY-HFVDHCIEILSHD--KDYNTVEV 209
A8CLP7/1-218 RTLLLEGG---KHRCNFRSTYRAK-KEV--VLPEY-HFVDHCIEILSHD--KDYNVVVEV 209
A7UAL2/1-218 MALLLEGG---GHYRCDFKTTYKAK-KVV--QLPDY-HFVDHRIEIVSHD--KDYNKVKL 209
A7UAL5/1-218 MALLLEGG---GHYRCDFKTTYKAK-KVV--RLPDY-HFVDHRIEIVSHD--KDYNKVKL 209
Q963F5/1-218 MALLLEGG---GHYRCDFRSTYKAK-KRV--QLPDY-HFVDHRIEILSHD--NDYNTVKL 209
Q7Z0W4/1-218 MALLLEGG---GHYRCDFRSTYKAK-KRV--QLPDY-HFVDHRIEILSHD--NDYNTVKL 209
Q66ND3/1-218 MALLLEGG---GHYRCDFRSTYKAK-KRV--QLPDY-HFVDHRIEILSHD--NDYNTVKL 209
Q2LI64/1-35 -----
Q60I25/1-218 MALLLEGG---GHYRCDFKTTYKAK-KVDV--RLPDY-HFVDHRIEILKHD--KDYNKVKL 209
Q60I24/1-218 MALLLEGG---GHYRCDFKTTYKAK-KVDV--RLPDY-HFVDHRIEILKHD--KDYNKVKL 209
A8CLT2/1-218 MALLLEGG---GHYRCDFKTTYKAK-KVDV--RLPDY-HFVDHRIEILKHD--KDYNKVEL 209
B5BR3C/1-218 MALLLEGG---GHYRCDFKTSYKAK-KVV--QLPDY-HFVDHRIEILSHD--RDYSKVKL 209
B6CT26/1-219 MALLLEGG---GHYRCDFKTSYKAK-KVV--QLPDY-HFVDHRIEILSHD--RDYSKVKL 210
Q9U6Y3/43-260 HSLLEGG---GHYRCDFKTSYKAK-KVV--QLPDY-HFVDHRIEILSHD--KDYNKVTL 209
Q8T6U0/1-218 MALLLEGG---GHYRCDFKTTYKAK-KVDV--QLPDY-HFVDHRIEILSHD--SDYNKVKL 209
Q8T5F0/7-229 MALLLEGG---SHYRCDFKTTYKAKNPVPTALPXV-HYVDHCIEITEEN--RDY--VKL 212
Q8T5F3/6-223 MALLLEGG---SHYRCDFKTTYKAKNPVPTALPXV-HYVDHCIEITEEN--RDY--VKL 209
Q66PW0/1-221 MSLLLEGG---DHYRCDFKTIYKPN-KKV--KLPY-HFVDHCIEIKSQE--NDYNMVAL 212
Q8MU46/1-221 MSLLLEGG---DHYRCDFKTIYKPN-KKV--KLPY-HFVDHCIEIKSQE--NDYNMVAL 212
Q8T5E9/1-221 MSLLLEGG---GHYRCDFKTIYTPK-KV--NMPY-HFVDHCIEIKSQE--NDYNMVAL 212
Q8T5E8/1-221 MSLLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--DDYNVVEL 212
Q8T6T8/1-221 MSLLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--DDYNVVEL 212
Q66PUS/1-220 MTLRVEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--KDYNTVAL 211
A8CL57/1-220 MTLRVEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--KDYNTVAL 211
Q8MAA2/34-254 MTLRVEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 212
Q8MAA1/12-234 KTRLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 214
Q66ND6/5-222 HALTVAGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--KEFMKVEE 210
Q9U6Y7/5-223 HALTVAGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--KEFMKVEE 210
Q553G7/6-223 MRLKLDG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 209
Q553G8/6-223 MRLKLDG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 209
Q2LI66/5-222 KALKLDG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--KDYTNVEL 209
Q66ND8/5-222 KALKLDG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--KDYTNVEL 209
Q66ND7/5-222 KALKLDG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--KDYTNVEL 209
Q9U6Y8/5-222 KALKLDG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 209
Q6KF85/5-222 KALKLDG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 209
Q2W74/7-224 KALKLDG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 209
Q9GTJ7/5-223 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
A8CLM7/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
A8CLM1/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q1JU63/2-219 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q1JV71/2-219 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q1JV72/2-219 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q1JV70/2-219 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
B5T1L1/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
A8CLV6/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q66PV0/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q66PV8/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q95P04/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
A8CLL3/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q66PV1/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q66PV9/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
P83690/1-218 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
A8CLN8/6-226 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
B6CT27/1-220 MALLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 210
Q9U6Y5/5-228 MYLLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 215
Q9U6Y4/5-228 MYLLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 215
Q8T4U4/5-228 MYLLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 215
B2ZAG2/5-226 MYLLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 213
Q692V6/5-228 MYLLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 215
B5LIN1/1-161 MYLLLEGG---GHYRCDFKTIYKPN-KV--NMPY-HFVDHCIEIKSQE--EDYTNVEL 152

B5T1L5/5-228 MYL L L K D G --- G R Y R C Q F D T T Y K A K S D P K -- E M P E W - H F I Q H K L T R E D R S - D A K N Q K W Q L 215
Q66PV2/5-228 MYL L L K D G --- G R Y R C Q F D T T Y K A K S D P K -- E M P E W - H F I Q H K L T R E D R S - D A K N Q K W Q L 215
Q86L2/5-228 S Y L L L K D G --- G R Y H C Q I D S V Y K A K S E P K -- K M P D R - H F I Q H K L T R E D R S - D A T N Q K W Q L 215
B22A61/5-228 M Y L L L K D G --- G R Y R C Q F D S V Y K A K S K P K -- V M P D W - H F I Q H K L T R E D R S - D A K N Q K W Q L 215
A8CLN4/5-231 M F L L L K D G --- G R Y R C Q F D T V Y K A K S D P K T I M P D W - H F I Q H K L N R E D R S - D A K H Q K W R L 218
A8CLV1/5-230 M F L L L K D G --- G R Y R C Q F D T V Y K A K S D P K T I M P D W - H F I Q H K L N R E D R S - D A K H Q K W R L 217
A8CLU7/1-223 E F L L K D G --- K R Y K C Q F H T F H D A K E K S R N M P M P D F - H F V Q H E I E R K D L P - G P M - Q T W Q L 214
A8CLP2/5-228 G F L L K D G --- K R Y K C Q F H T F H D A K D K S K M P M P D F - H F V Q H K I E R K D L P - G S M - Q T W R L 215
B6CT22/5-226 E F L L L E G G --- G R Y K C Q F H T V Y R A K T E P K -- R M P E F - H F V Q H K L T R T D V S - D P L K Q Q W Q L 213
Q2VTM9/5-219 M F L M L E G G --- H R L K C Q F T T S Y K - A K K A V -- K M P P N - H I E H R L V R K E V A D A --- V Q I 206
Q1ALD3/5-219 M F L L L E G G --- H R H K C Q F T T S Y K - A H K A V -- K I P P N - H I E H R L V R K E V G D A --- V Q I 206
Q1ALD5/5-219 M F L M L E G G --- H R Q K C Q F T T S Y K - A S K A V -- K M P P N - H I E H V V W K G E D S D G --- F Q I 206
Q1ALD4/5-219 M F L V L E G G --- H R L K C L F Q T T Y K - A D K V V -- K M P P N - H I E H R L V R S E D G D A --- V Q I 206
Q09HR5/5-219 M F L L L E G G --- Y Q Q K C Q F Q T T Y R - T D K T V -- K M P P N - H I E H R L V R S E D G D A --- V Q I 206
Q2VTM8/5-219 M F L K L E G G --- Q Q H R C Q F Q T T Y K - A H K A V -- K M P P N - H I E H R L V R S Q G D A --- V Q L 206
A8CLU1/1-218 M F L S L T D G --- G N H R C Q F S T L Y K - A K K A V -- K L P T E S H Y V E H R L V R T D L P N G K --- V Q L 209
Q9U6Y6/5-224 A F L M L Q G G --- G N Y R C Q F H T S Y K - T K K P V -- T M P P N - H V V E H R I A R T D L D K G -- G N S V Q L 211
A8CLT7/5-222 M F L M L E G G --- K N Y K C Q F H T S Y K - T K K P V -- T L P S N - H V V E H R I V R T N L D K A -- G N H V Q L 209
Q66ND1/5-220 M F L L K G G --- G Y H S C Q F H T S Y K - T K E P V -- T L P Q N - H V V E H R I T R T D I E D --- K K V L L 207
Q66PV5/5-226 E F L M L E G G --- G Y H S C Q F H S T Y K - P E K P V -- T L P P N - H V V E H I V R T D L G Q T A K G F T V K L 213
B5T1L2/5-226 E F L M L E G G --- G Y H S C Q F H S T Y K - P E K P V -- T L P P N - H V V E H I V R T D L G Q R A K G F T V K L 213
Q66PV3/5-226 E F L M L E G G --- G Y H S C Q F H S T Y K - P E K P A -- A L P P N - H V V E H I V R T D L G Q S A K G F T V K L 213
B5T1L0/5-226 E F L M L E G G --- G Y H S C Q F H S T Y K - P E K P A -- A L P P N - H V V E H I V R T D L G Q S A K G F T V K L 213
Q66PV4/5-226 G F L M L E G G --- G Y H S C Q F H S T Y K - P E K P A -- A L P P N - H V V E H I V R T D L G Q S A K G F T V K L 213
B5L1N0/5-226 E F L M L E G G --- G Y H S C Q F H S T Y K - P E K P G -- T L P P N - H V V E H I V R T D L G Q S A K G F A V K L 213
B5T1L3/5-226 M F L M L E G G --- G Y H R C Q F H S T Y K - P E K P G -- T L P P N - H V V E H I V R T D L G Q S A K G F T V K L 213
B5T1L4/5-226 M F L M L E G G --- G S H R C Q F H S T Y K - T E K P V -- T L P P N - H V V E H Q I V R T D L G Q S A K G F T V K L 213
Q66PV7/5-226 M F L M L E G G --- G S H R C Q F H S T Y K - T E K P V -- T L P P N - H V V E H Q I V R T D L G Q S A K G F T V K L 213
A8CLY3/5-226 M F L M L E G G --- G S H R C Q F H S T Y K - T E K P V -- T L P P N - H V V E H Q I V R T D L G Q T A K G F T V K L 213
Q66PV8/5-226 M F L M L E G G --- G S H R C Q F H S T Y K - T E K P V -- T L P P N - H V V E H Q I V R T D L G Q S A K G F T V K L 213
Q66PV9/5-226 M F L M L E G G --- G Y H R C Q F H S T Y K - T E K P V -- E L P P N - H V V E H Q I V R T D L G Q S A K G F T V K L 213
A8CLX9/5-226 M F L M L E G G --- G Y H R C Q F H S T Y K - T E K P V -- E L P P N - H V V E H Q I V R T D L G Q S A K G F T V K L 213
B6CT23/5-226 M F L M L E G G --- G Y H R C Q F H S T Y K - T E K P V -- T L P P N - H V V E H Q I V R T D L G Q S A K G F T V K L 213
Q6I7B1/6-227 Q F L L L E G G --- G Y Q R C R F H S T Y K - T E K P V -- A M P P S - H V V E H Q I V R T D L G Q T A K G F K V K L 213
Q66PV6/5-226 E F L M L E G G --- G Y H S C Q F H S T Y K - P E K P V -- E L P P N - H V I E H I V R T D L G K T A K G F M V K L 213
Q86LV8/1-213 M F L K L E G G --- G R H K C Y F V T N H K - A K R A V -- K V P D N - H F V W H R L V R N --- G D G N T V E L 204
Q86LV7/1-213 M F L K L E G G --- G R H K C Y F V T N H K - A K R A V -- K V P D N - H F V W H R L V R N --- G D G N T V E L 204
A8CLR1/5-220 M F L K L E G G --- G R H R C D F R T T Y K - A K K D V -- K M P N S - H F I T H C L V R K --- G D G N N T E L 208
Q6R8F5/5-223 M Y L K L E G G --- G N Y R C E Y R T I Y K - A K K D V -- K M P K S - H F V E H T L V R N N I K K D A K G N T I E L 211
Q6R8F4/1-219 M Y L K L E G G --- G N Y R C D Y R T I Y K - A K K D V -- K M P K S - H F V E H T L V R N N I K K D A K G N T I E L 211
Q6R8F3/5-220 M F L K L E G G --- G N Y R C D F E T T Y K - A K K A V -- K M L D S - H F I E H R L V T T --- I R G N N V E L 207
Q6I7B2/2-218 M Y L K L E G G --- G N H K C Q F K T T Y K A A K K I L -- K M P G S - H Y I S H R L V R K --- T E G N I T E L 209
A7VMR3/2-218 M Y L K L E G G --- G N H K C Q F K T T Y K A A K E I L -- E M P G D - H Y I G H R L V R K --- T E G N I T E L 209
Q6I7B4/1-210 M F L K L E G G --- G N H K C Q F K T T Y K A A K K V L -- K M P Q S - H F I G H R L V R K --- T K G N I T E L 202
Q6I7B3/1-217 M F L K L A G G --- G N H K C Q F K T T Y K A A K K I L -- K M P Q S - H F I G H R L V R K --- T E G N I T E L 209
A8CLQ6/5-221 M F L K L V G G --- G N H K C Q F T T T Y K A A K K V L -- D M P Q S - H F I F H R L V R K --- T E G N I T K 209
A8CLM9/4-223 S F L V L Q G G --- G N Y R C D Y T S Y R - A M K P V -- E M P G G - H F I Q H R I V R R D I K K D S N G N T W Q I 211
Q95W86/2-220 M A L K V S G -- R - P P L I C H L H S T Y R S K K A C - A L T M P G F - H F A D L R I Q M P K - K -- K K D E Y F E L 210
Q95W11/2-220 M A L K V S G -- R - P P L I C H L H S T Y R S K K A C - A L T M P G F - H F A D L R I Q M P K - K -- K K D E Y F E L 210
Q8M145/2-220 M A L K V S G -- R - P P L I C H L H S T Y R S K K A C - A L T M P G F - H F A D L R I Q M P K - K -- K K D E Y F E L 210
Q95W85/2-220 M A L K V G D -- R - R - L I C H L Y S Y R S K K A V R A L T M P G F - H F T D I R L Q M P R - K -- K K D E Y F E L 210
A5YTR7/2-222 M A L K C P G -- G - R H L T C H L H T T Y R S K K P A S A L K M P G F - H F E D H R I E I M E E V -- E K G C Y K Q 212
Q9GZ28/2-222 M A L K C P G -- G - R H L T C H L H T T Y R S K K P A S A L K M P G F - H F E D H R I E I M E E V -- E K G C Y K Q 212
A5YTR8/2-222 M A L K C P G -- G - R H L T C H L H T T Y R S K K P A S A L K M P G F - H F E D H R I E I M E E V -- E K G C Y K Q 212
Q9GP15/2-222 M A L K C P G -- G - R H L T C H L H S T Y R S K K P A S A L K M P G F - H F G D H R I E I M E E V -- E K G C Y K Q 212
Q2VFP3/2-221 M A L K C T D -- G - N H L T S H L R T T Y R S R K P S N A V N M P E F - H F G D H R I E I L K - A -- E Q G K F Y E Q 211
A0AQQ7/2-222 M A L K C A D -- G - N H L T C H L R T T Y R S K K A A K A L Q M P P F - H F S D H R P E I V K V S -- E N G T L F E Q 212
A0AQQ8/2-222 M A L K C A D -- G - N H L T C H L R T T Y R S K K A A K A L Q M P P F - H F S D H R P E I V K V S -- E N G T L F E Q 212
Q81SF8/2-222 M A L N V D G -- G - G Y L S C S F E T T Y R S K K T V E N F K M P G F - H F V D H R L E R L E E S -- D K E M F V V Q 212
Q86LV4/2-220 P A L K L R N N - K - G H L L C V M E T T Y K P N K --- K V N L P K L - H F H H L R M E K D S I S -- D D E K T I Q 210
Q5QIU5/2-220 P A L K L L D N - K - G H L L C V M E T T Y K P N K --- K V N L P K L - H F H H L R M E K D S I S -- D D E K T I Q 210
Q8T6T9/2-220 P A L K L L G N - K - G H L L C V M E T T Y K S K K --- K V N L P K P - H F H H L R M E K D S V S -- D D E K T I E Q 210
Q8T5E7/2-220 P A L R L R D -- K - G H L I C H M E T T Y K P N K --- E V K L P E L - H F H H L R M E K L S V S -- D D G K T I Q 210
A5YTR6/2-219 P A L M L A D -- G - G H L S C F M E T T Y K S K K --- E V K L P E L - H F H H L R M E K L N I S -- D D W K T V E Q 209
Q9GPI6/2-219 P A L M L A D -- G - G H L S C F M E T T Y K S K K --- E V K L P E L - H F H H L R M E K L N I S -- D D W K T V E Q 209
Q5ZQ06/2-221 I A L K L D D D H G - G H L T G N F K A T Y K S N N P -- N I N L P K H - H F V E N R V E M L E G R -- D G G N Y V K L 211
B9UP66/1-219 K A Y L L K D G --- S Y Y Y G M T T F Y R S K K S G Q -- A L P G F - H F I Q H R L V K T K V E -- P G F K M V E Q 210
Q5ZQ05/1-219 K A Y L L K D G --- S Y Y Y G M T T F Y R S K K S G Q -- P L P G F - H F I K H R L V K T K V E -- P G F K M V E Q 210
Q7Z168/1-218 K A Y L L K D G --- S Y Y Y G M T T F Y R S K K S G Q -- A P P G Y - H F V K H R L V K T N V G -- H G F K T V E Q 209
Q9BL20/8-225 L V Y K L E S G --- N Y S C H M K T F Y R S K G G V K -- E F P E Y - H F I H R L E K T Y V E -- E G S - F V E Q 209
Q9BLY9/8-225 L V Y K L N S G --- K Y S C H M K T L M K S G V V K -- E F P S Y - H F I Q H R L E K T Y V E -- D G G - F V E Q 209
Q963I9/5-223 I A F K L Q T G --- K H F T Y H M R T V Y K S K P V E -- T M P L Y - H F I Q H R L V K T N V D -- T A S G Y V V Q 210
A8CLX6/5-222 M Y L M V E G G --- K T L K C R Y E N N Y R A N K P V L -- M P P S - H F V D L R L T R T N L D -- K E G L A F K L 209
A7S3R2/57-279 M L L K V N D G --- S F L R V Q F E T V Y R F M K P V P A G F K M P P H F M A Y R L T R V D N D -- E H C D T V I Q 214
A7RG05/1-174 M L L K V N D G --- S F L R V K F E T V Y R Y V N ----- L Y Q K ----- 174
A7S1H5/1-224 M L L K V N D G --- S F L R V K F V T V Y R F M K P V P A G F K M P P H F I A F K L T R V E N D -- A D C N V V L L 215
A7S1H3/1-82 -----
B1PNC2/5-218 W T Y T T G S G K --- R Y Q S T V R T N Y T F A K P M A A N I L Q N Q - P M F V F R K T E L K H S --- K T E L N F 204
B1PNC0/4-217 W T Y T T G S G K --- R Y Q S T V R T N Y T F A K P M A A N I L K N Q - P M F V F R K T E L K H S --- K T E L N F 205
B1PNC1/5-218 W T Y N T A S G K --- R Y Q S T V R T N Y T F N K P M A A N I L K N Q - P M F V F R K T E L K H S --- K T E L N F 204
B1PNC4/5-218 W T Y T T G S G K --- R Y Q S T V R T N Y T F A K P M A A N I L K N Q - P M F V F R K T E L K H S --- K T E L N F 204
C3YR99/48-261 W T Y T T T E G K --- R Y Q S T V R T N Y T F A K P M A A N I L Q K Q - P M F V F R K T E L Q H S --- K T E L T F 206
B1PNC3/4-217 W T Y T T T S G K --- R Y Q S N V R S N F T F A K P I A A N I L Q K Q - P M F V F R K T E L K H S --- K T E L N F 206

B1PNB8/5-217 WYTTTSGK---RYQSNVRSNFTFAKPIAANILQKQ-PMFVFRKTELKHS----KTELN 205
 B1PNB9/4-216 WYTTTSGK---RYHSNVRSNFTFAKPIAANILQKQ-PMFVFRKTELKHS----KTELN 206
 C3YRA1/51-267 KTYTTTSGK---RYQCTFRENNTFAKPMADILQKQ-PMFIFHKTELQHSN---NAELTF 207
 C3YRA2/11-223 CSYTTTAGK---RYQSKMRENNTFAKPMADILQKQ-PMFVFRKSELQHS----KTELTF 205
 B1PND7/4-187 QTYTTTSGK---GYKSTVRTNNTFAKPIAADMM----- 184
 B1PND6/5-189 QTYTTASGK---GYKSTVRTNNTFAKPMADMMQNO----- 185
 B1PND8/4-184 QTYTTTSGK---GYQSTVRTNNTFAKPIAA----- 181
 B1PND5/5-186 QTYTTASGK---RYQGTVRTNNTFAKPIAANI----- 182
 C3YRA3/4-216 WTCPTTSGK---RYHATVRTNYTFAKPIAASILQKQ-PMFVFRKTEVKAS----DAEINL 205
 B1PND3/4-217 WYTTTSGK---RYQSDVQTNVTFKPIASADILKKQ-PMFVFRKVELKHS----KTELN 206
 B1PND2/5-217 WYTTTSGK---RYQSDVQTNVTFKPIASADILKKQ-PMFVFRKVELKHS----KTELN 205
 B1PNC9/4-217 WYTTTSGK---RYQSDVQTNVTFKPIAADILKKQ-PMFVFRKVELKHS----KTELN 206
 B1PNC6/5-217 WYTTTSGK---RYQSDVQTNVTFKPMADILKKQ-PMFVFRKVELKHT----KTELN 205
 B1PNC7/4-220 WYTTTSGK---RYQSDVQTNVTFKPIAADILKKQ-PMFVFRKVELKHT----KTELN 206
 B1PND0/5-220 WYTTTSGK---RYQSDVQTNVTFKPIAADILKKQ-PMFVFRKVELKHT----KTELN 205
 B1PNC5/5-220 WYTTTSGK---RYQSDVQTNVTFKPIAADILKKQ-PMFVFRMELKHT----KTELN 205
 B1PND1/5-217 WYTTTSGK---RYQSDVQTNVTFKPIASADILKKQ-PMFVFRKVELKHS----KTELN 205
 B1PND4/5-217 WYTTTSGK---RYQSDVQTNVTFKPIASADILKKQ-PMFVFRKVELKHS----KTELN 205
 B1PNC8/4-69 -----
 C3YRA0/8-104 -----
 C3YKQ3/5-218 WSYNLQNGK---RYRARSVSHYIFDKPFSADLKKQ-PVFVYRKHVVKAS----KTEVTL 204
 C3YKQ7/5-217 WSYNLQNGK---RYRARTVSNYIFGKPLAADVMKKQ-PVFVYRKYVKST---QTEITL 204
 C3YKQ2/53-191 WSFCTTDGK---RHQADVQTNVTFKPLPAGLKEKM-PIFLGHQIEVKAS----KTEITL 131
 A8QK29/4-215 WSFCTTDGK---RHQADVQTNVTFKPLPAGLKEKM-PIFLGHQIEVKAS----KTEITL 204
 C3YKQ9/5-215 WSFCTTDGK---RHQADVQTNVTFKPLPAGLKEKM-PIFLGHQIEVKAS----KTEITL 203
 C3YKQ1/4-215 WSFCTTDGK---RHQADVQTNVTFKPLPAGLKEKM-PIFLGHQIEVQAS----KTEINL 204
 C3ZLQ0/369-585 WTYRTSSGG---RYRATVQTNVTFKPIAAGLKNM-PMFVFRQLEVTGS----KTEISL 208
 C3YK10/4-218 WSYRTSSGG---RYRATVQTNVTFKPIAAGLKNM-PMFVFRQLEVTGS----KTEIGL 207
 C3YKQ0/5-214 WFTFTEKGGK---KRYKAEVMTNATFAQNLQPLKNVM-PLFVFRKVDIDCS----KTEVTL 205
 Q6WV10/1-215 RTYLLKDDG---YSAQVNNHMHFKSAIHTTMLQNGGSMFTYRVEEHT---QNEVAI 208
 Q6WV09/1-217 RTYLLKDDG---YSAQVNNHMHFKSAIHPTMLQNGGSMFTHRVEENHT---KTNVAI 208
 Q6WV11/1-215 RTWMLRDGG---YSAQVNNHMHFKTAMHPTMLQNGGSMFTYRVEELHS---QSDVGI 208
 Q6WV12/1-213 RTFSLRDGG---YYSFVVDSHMHFKSAIHPSILQNGGPMFAFRRVEELHS---NTELGI 208
 Q6WV13/1-214 RTFSLRDGG---YYSFVVDSHMHFKSAIHPSILQNGGPMFAFRRVEELHS---NTELGI 208
 Q6WV08/1-214 RTLSLRDGG---YYSQVTSIHIFKNAIHPSILHNGGSMFTYRVEELHT---QTDLGI 208
 Q2MHN7/1-215 RAFQLKDDG---FYTAEVKNIDFKNPIHESFSKSG-PMFTHRVEEHT---KENLAM 205

Q17106/5-229 LEFVTSAGI--- 225
 Q17105/5-229 LEFVTAARI--- 225
 P42212/5-229 LEFVTAAGI--- 225
 Q6YGZ0/5-229 FEFVTAAGI--- 225
 Q8WTC5/5-229 LEFFSACGH--- 225
 Q8WTC4/5-229 LEFFSACGH--- 225
 Q8WTC9/5-229 LEFFSACGH--- 225
 Q8WTD0/5-229 LEFFSACGH--- 225
 Q8WTC8/5-229 LEFFSACGH--- 225
 Q8WTC6/5-229 LEFFSACGH--- 225
 Q8WP95/5-229 LEFFSACGH--- 225
 Q8WTC7/5-229 LEFFSACGH--- 225
 Q6RYS7/5-229 VETVRAVDC--- 225
 Q6RYS6/39-258 KETNCAFDF--- 220
 Q6RYS5/2-222 KEVNVASGI--- 221
 Q6RYS4/5-225 REVTYAHSV--- 221
 Q2TCH5/5-225 REVAYAHSV--- 221
 Q53UG7/1-221 YEHAKAHSG--- 221
 Q53UG8/1-221 YEHAKAHSG--- 221
 Q8MU48/1-218 SEHAEAHSG--- 218
 A8CLW1/1-218 CEHAEAHSG--- 218
 B6CTZ4/1-218 CEHAEAHSG--- 218
 Q5TLG6/1-218 HEHAEAHSE--- 218
 A8CLQ1/1-223 HEHAEAHSG--- 223
 Q2LI65/1-219 REHAEAHFG--- 219
 A8CLW4/1-222 REHAEAHSG--- 222
 B6CTZ5/4-224 HEHAKARYG--- 221
 A3F207/1-221 FEDAVAHSG--- 221
 Q962P9/1-221 YEHAEAHSG--- 221
 Q7Z0N8/1-221 YEHAEAHSG--- 221
 Q66ND5/1-221 YEHAEAHFG--- 221
 Q66ND4/1-221 YEHAEAHFG--- 221
 Q66ND2/1-221 YEHAEAHSG--- 221
 A7UAL7/1-221 YEHAEAHSG--- 221
 Q7Z0W6/1-221 YEHAEAHSG--- 221
 A7UAL9/1-221 YEHAEAHSG--- 221
 Q8T5F2/8-228 HEHAEARHG--- 221
 Q7Z0W7/8-228 HEHAEARHG--- 221
 Q8MU47/8-228 YEHAEAHSG--- 221
 Q95VT0/1-221 YEHAEAHSG--- 221
 A7UAL3/1-221 HEHAEAHSG--- 221
 A7UAL6/1-218 YEHAEAHSG--- 218
 A7UAL8/1-218 YEHAEAHSG--- 218
 A8CL52/1-219 YEHAHAHLG--- 219
 Q8I6J8/1-219 YEHAHAHSG--- 219

A3F208/1-219	YEHAVAHSG---	219
A8CLR7/1-219	YEHAVAHSG---	219
Q5S6Z9/1-219	YEHAVAHSG---	219
Q66PW1/5-223	YEHAVAHSG---	219
Q8T5F1/1-219	YEHAEAHSG---	219
Q7Z0W9/1-219	YEHAEAHSG---	219
Q6USK3/1-219	YEHAEAHSG---	219
A7UAL4/1-219	YEYEAHSG---	219
A7UAM0/1-219	CEHAEAHSG---	219
Q7Z0W5/1-218	YENAVARPS---	218
Q95UA7/1-218	YENAVARPS---	218
A7UAL1/1-218	YENAVARPS---	218
A8CLP7/1-218	YENAVARQQ---	218
A7UAL2/1-218	YEHAEAHSG---	218
A7UAL5/1-218	YEHAEAHSG---	218
Q963F5/1-218	SEDAEARYS---	218
Q7Z0W4/1-218	SENAEARYS---	218
Q66ND3/1-218	SENAEARYS---	218
Q2LI64/1-35	-----	
Q60I25/1-218	YENAVARYS---	218
Q60I24/1-218	YENAVARYS---	218
A8CLT2/1-218	YENAVARYS---	218
B5BRC3/1-218	YENAVARYS---	218
B6CTZ6/1-219	YENAVARYS---	219
Q9U6Y3/43-260	YENAVARYS---	218
Q8T6U0/1-218	YEHGVARYS---	218
Q8T5F0/7-229	QEYAKARSLH-	223
Q8T5F3/6-223	REHAKARSS---	218
Q66PW0/1-221	FEDAVAHYS---	221
Q8MU46/1-221	FEDAVAHYS---	221
Q8T5E9/1-221	SEDAVAHNS---	221
Q8T5E8/1-221	YEGAVAHYS---	221
Q8T6T8/1-221	TEVAEARYS---	221
Q66PU5/1-220	YEAARVARS---	220
A8CL57/1-220	YEAARVARS---	220
Q8MMA2/34-254	HDVSWARYS---	221
Q8MMA1/12-234	REVALARYS---	223
Q66ND6/5-222	HEIIVARXH---	218
Q9U6Y7/5-223	HEIIVARHH---	219
Q5S3G7/6-223	YERAEGRHS---	218
Q5S3G8/6-223	YERAEGRHS---	218
Q2LI66/5-222	YERTEGRHH---	218
Q66ND8/5-222	YERTEGRHH---	218
Q66ND7/5-222	YERTEGRHH---	218
Q9U6Y8/5-222	YERTEGRHH---	218
Q6KF85/5-222	YERTEGRHH---	218
Q2WG74/7-224	YERTEGRHH---	218
Q9GTJ7/5-223	YEKTQGRHH---	219
A8CLM7/1-218	CEISTARHS---	218
A8CLM1/1-218	CEISTARHS---	218
Q1JU63/2-219	CETAIARHS---	218
Q1JV71/2-219	CETAIARHS---	218
Q1JV72/2-219	CETAIARHS---	218
Q1JV70/2-219	CETAIARHS---	218
B5T1L1/1-218	CEISTARHP---	218
A8CLV6/1-218	CEISTARHS---	218
Q66PV0/1-218	CEISTARHP---	218
Q66PU8/1-218	CEISTARHP---	218
Q95P04/1-218	CEISTARHP---	218
A8CLL3/1-218	CEISTARHP---	218
Q66PV1/1-218	CEISTARHP---	218
Q66PU9/1-218	REISTARHP---	218
P83690/1-218	CEISTARHP---	218
A8CLN8/6-226	FEQREIAHLS	221
B6CTZ7/1-220	QEAAEGHFS---	220
Q9U6Y5/5-228	TEHAIASGS---	224
Q9U6Y4/5-228	TEHAIAFPS---	224
Q8T4U4/5-228	IEHAIASRS---	224
B2ZAG2/5-226	VEHAVASRS---	222
Q692V6/5-228	VEHAVASRS---	224
B5LIN1/1-161	VEHAVASRS---	161
B5T1L5/5-228	VEHAVASRS---	224
Q66PV2/5-228	VEHAVASRS---	224
Q86LC2/5-228	KERAVASLS---	224
B2ZAG1/5-228	VEHASASRS---	224
A8CLN4/5-231	VENAIAYRS---	227
A8CLV1/5-230	VENAIAYRS---	226
A8CLU7/1-223	TEHAAACKN---	223
A8CLP2/5-228	TEHAAACKT---	224
B6CTZ2/5-226	TEDAAACES---	222
Q2VTM9/5-219	QEHAVAKHF---	215
Q1ALD3/5-219	QEHAVAKHF---	215
Q1ALD5/5-219	KEHAVAKHF---	215

Q1ALD4/5-219	QEHAVAKYF---	215
Q09HR5/5-219	QEHAVAKYF---	215
Q2VTM8/5-219	KEHAVAKCF---	215
A8CLU1/1-218	EEHAAARLN---	218
Q9U6Y6/5-224	TEHAVAHIT---	220
A8CLT7/5-222	DEHAVAHVN---	218
Q66ND1/5-220	EETAVAHVN---	216
Q66PV5/5-226	EEHAAAHHV---	222
B5T1L2/5-226	EEHAAAHHV---	222
Q66PV3/5-226	EEHAAAHHV---	222
B5T1L0/5-226	EEHAAAHHV---	222
Q66PV4/5-226	EEHAAAHHV---	222
B5LIN0/5-226	EEHAAAHHV---	222
B5T1L3/5-226	EEHAAAHHV---	222
B5T1L4/5-226	EAHAAAHHV---	222
Q66PV7/5-226	EAHAAAHHV---	222
A8CLY3/5-226	EEHAAAHSV---	222
Q66PV8/5-226	EAHAAAHHV---	222
Q66PV9/5-226	EAHAAAHHV---	222
A8CLX9/5-226	EAHAAAHHV---	222
B6CTZ3/5-226	EALAAAHHV---	222
Q6I7B1/6-227	EEHAEHHV---	222
Q66PV6/5-226	VQHAAAHHV---	222
Q86LV8/1-213	EETAEARYD---	213
Q86LV7/1-213	EETAEARYD---	213
A8CLR1/5-220	IEDAEEARN---	216
Q6R8F5/5-223	IEDASARY---	219
Q6R8F4/1-219	IEDASARY---	219
Q6R8F3/5-220	QEDAEARNS---	216
Q6I7B2/2-218	VEDAVAHS---	217
A7VMR3/2-218	VEDAVAHS---	217
Q6I7B4/1-210	VEDAVAHC---	210
Q6I7B3/1-217	VEDAVAHC---	217
A8CLQ6/5-221	VEDVEAHN---	217
A8CLW9/4-223	QEDAFAHNS---	220
Q95W86/2-220	YEASVARYS---	219
Q95W11/2-220	YEASVARYS---	219
Q8MU45/2-220	YEASVARYS---	219
Q95W85/2-220	YEASVARYS---	219
ASVTR7/2-222	YEAAVGRYC---	221
Q9GZ28/2-222	YEAAVGRYC---	221
ASVTR8/2-222	YEAAVGRYC---	221
Q9GP15/2-222	YEAAVARYC---	221
Q2VFP3/2-221	YESAVARYC---	220
A0AQQ7/2-222	HESVARYC---	221
A0AQQ8/2-222	HESVARYC---	221
Q8ISF8/2-222	HEHAVAKFC---	221
Q86LV4/2-220	HEDVRASYF---	219
Q5QIU5/2-220	HEHVVASYF---	219
Q8T6T9/2-220	HENVRASYF---	219
Q8T5E7/2-220	HEYVVASYS---	219
ASVTR6/2-219	HESVVASYS---	218
Q9GP16/2-219	HESVVASYS---	218
Q5ZQ06/2-221	HERNKARYS---	220
B9UPG6/1-219	SEYATAFVS---	219
Q5ZQ05/1-219	AEYATAHVC---	219
Q7Z168/1-218	TEYATAHVS---	218
Q9BLZ0/8-225	HETAIAQLT---	218
Q9BLY9/8-225	HETAIAQMT---	218
Q963I9/5-223	HETAIAAHS---	219
A8CLX6/5-222	EYAVARVL---	218
A7S3R2/57-279	HEWSEAFSC---	223
A7RGQ5/1-174	-----	
A7S1H5/1-224	HEWGKAFSC---	224
A7S1H3/1-82	-----	
B1PNC2/5-218	KEWQKAFADV---	214
B1PNC0/4-217	KEWQKAFD---	214
B1PNC1/5-218	KEWQMAFADV---	214
B1PNC4/5-218	KEWQKAFADV---	214
C3YR99/48-261	KEWQKAFD---	214
B1PNC3/4-217	KEWQTAFS---	214
B1PNB8/5-217	KEWQTAFS---	213
B1PNB9/4-216	KEWQTAF---	213
C3YRA1/51-267	KEKQTAFSDM---	217
C3YRA2/11-223	KEWQKAFD---	213
B1PND7/4-187	-----	
B1PND6/5-189	-----	
B1PND8/4-184	-----	
B1PND5/5-186	-----	
C3YRA3/4-216	KEWQKAFH---	213
B1PND3/4-217	KQWQKAFQ---	214
B1PND2/5-217	KQWQKAFQ---	213
B1PNC9/4-217	KQWQKAFQ---	214

B1PNC6/5-217	KQWQKAFQ----	213
B1PNC7/4-220	KQWQKAFQDIA-	217
B1PND0/5-220	KQWQKAFQDIA-	216
B1PNC5/5-220	KQWQKAFQDIA-	216
B1PND1/5-217	KQWQKAFQ----	213
B1PND4/5-217	KQWQKAFQ----	213
B1PNC8/4-69	-----	
C3YRA0/8-104	-----	
C3YKQ3/5-218	DEREKAFYEL--	214
C3YKP7/5-217	DEREKAFYE---	213
C3YKQ2/53-191	SEKVKAFI----	139
A8QVZ9/4-215	SEKVKAFI----	212
C3YKP9/5-215	CEKVKAFI----	211
C3YKQ1/4-215	SEKVKAFI----	212
C3ZLQ0/369-585	QEQQKAFST---	217
C3YKI0/4-218	QEQQKAFS----	215
C3YKQ0/5-214	IEREK-----	210
Q6WV10/1-215	VEYQNVF-----	215
Q6WV09/1-217	VEYQNVFKT---	217
Q6WV11/1-215	VEYQHVF-----	215
Q6WV12/1-213	VEYQH-----	213
Q6WV13/1-214	VEYQHA-----	214
Q6WV08/1-214	VEYQHV-----	214
Q2MHN7/1-215	VEYQQVFNSA--	215