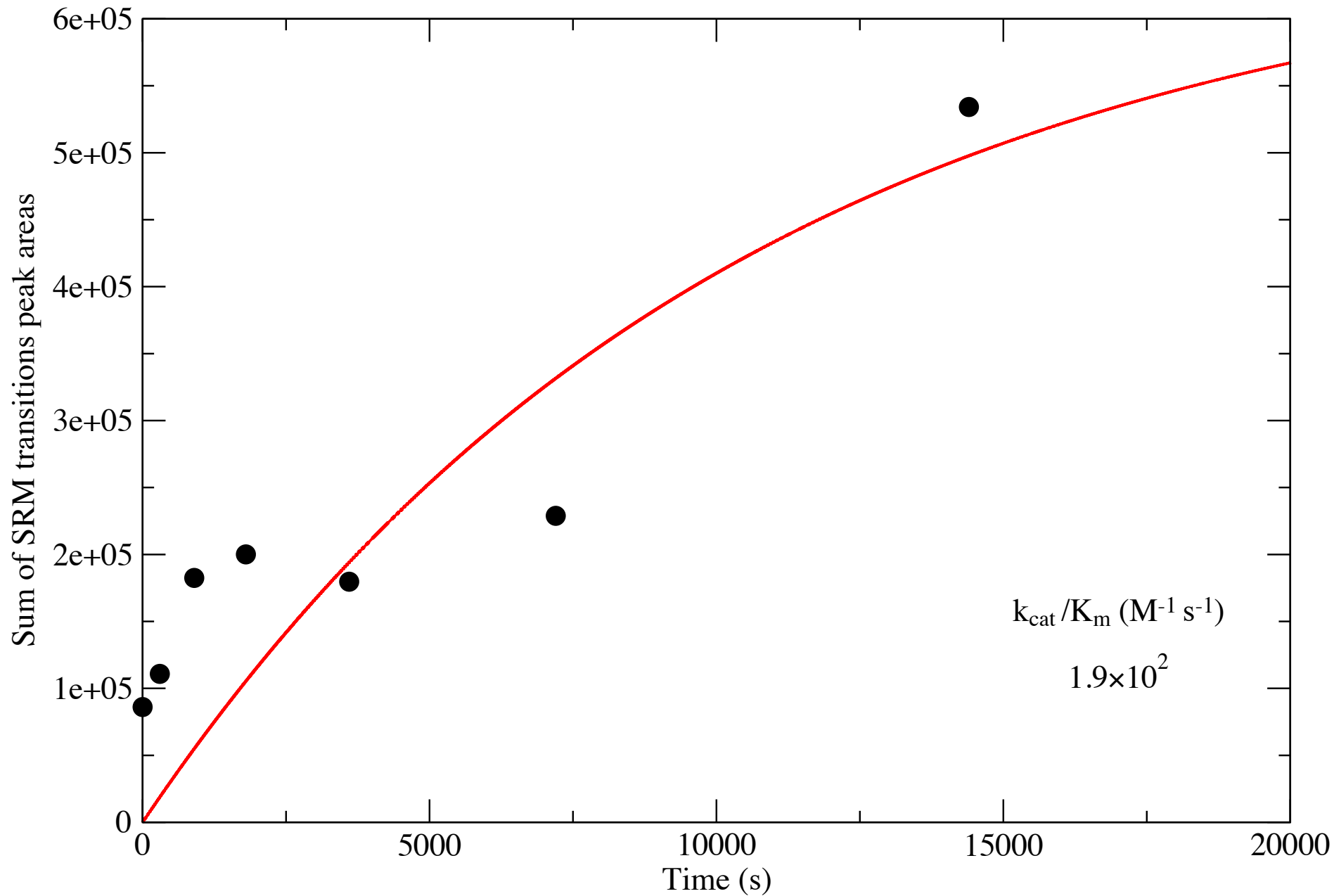
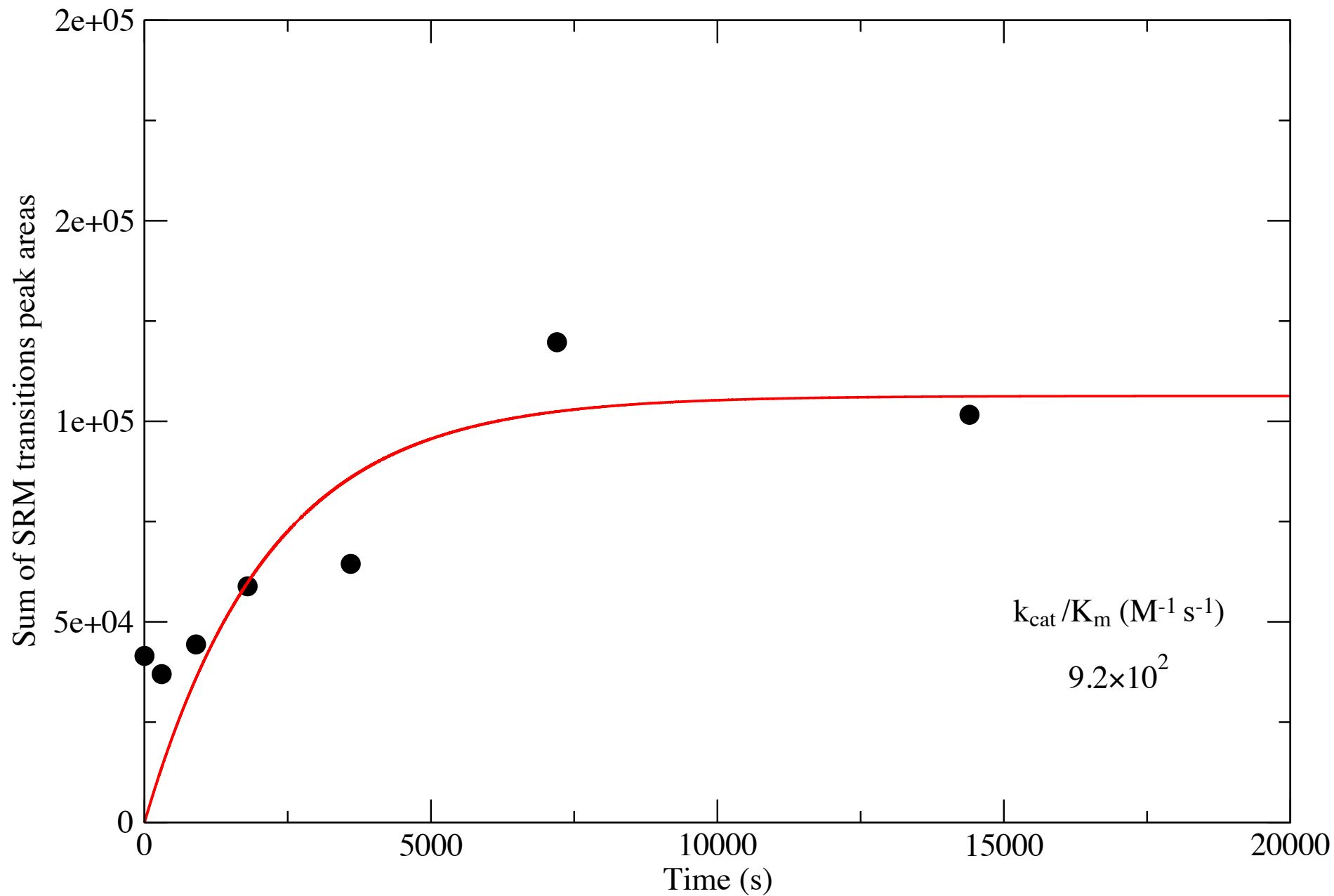


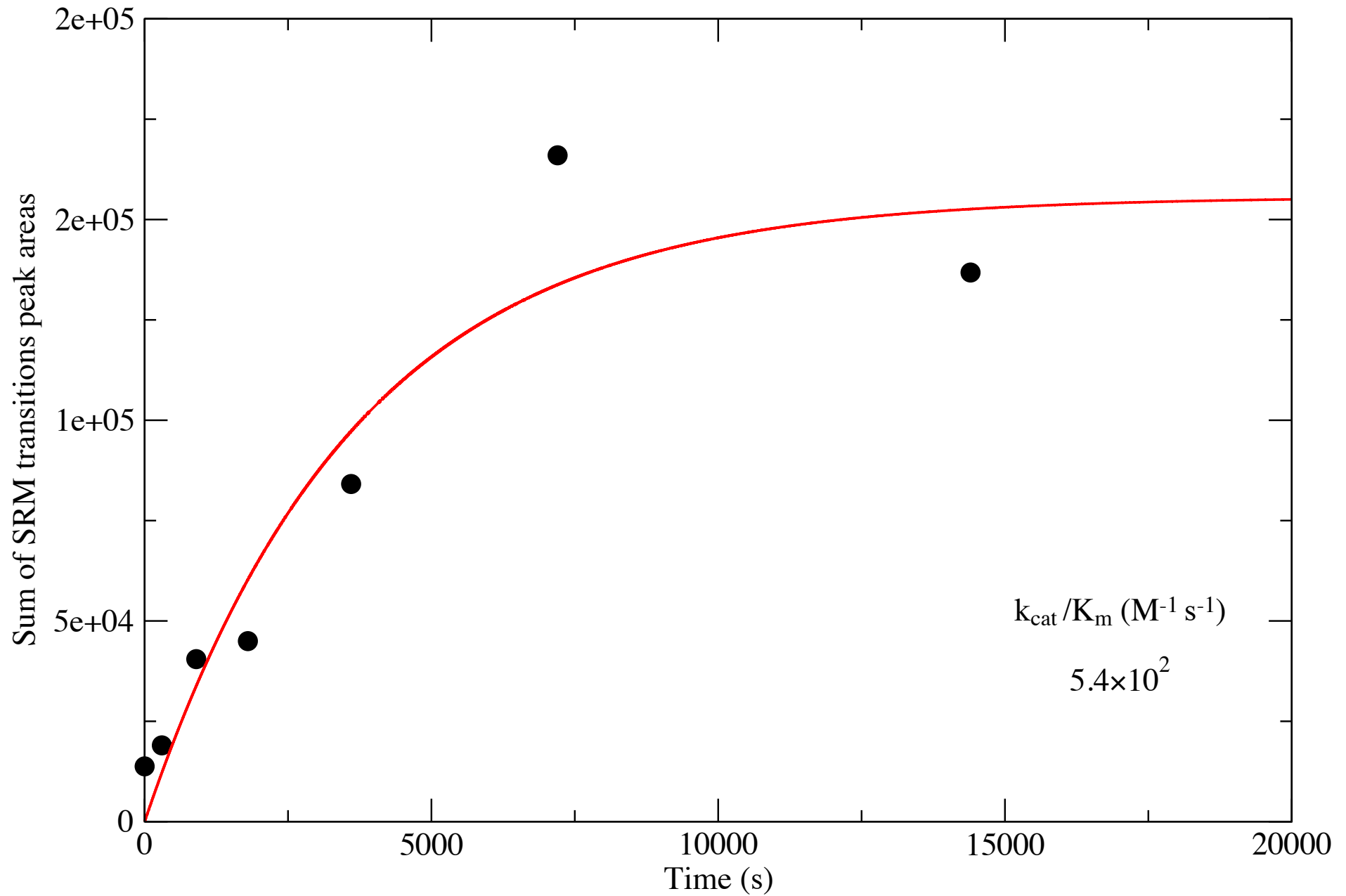
ACLY_HUMAN (SVPSPR)



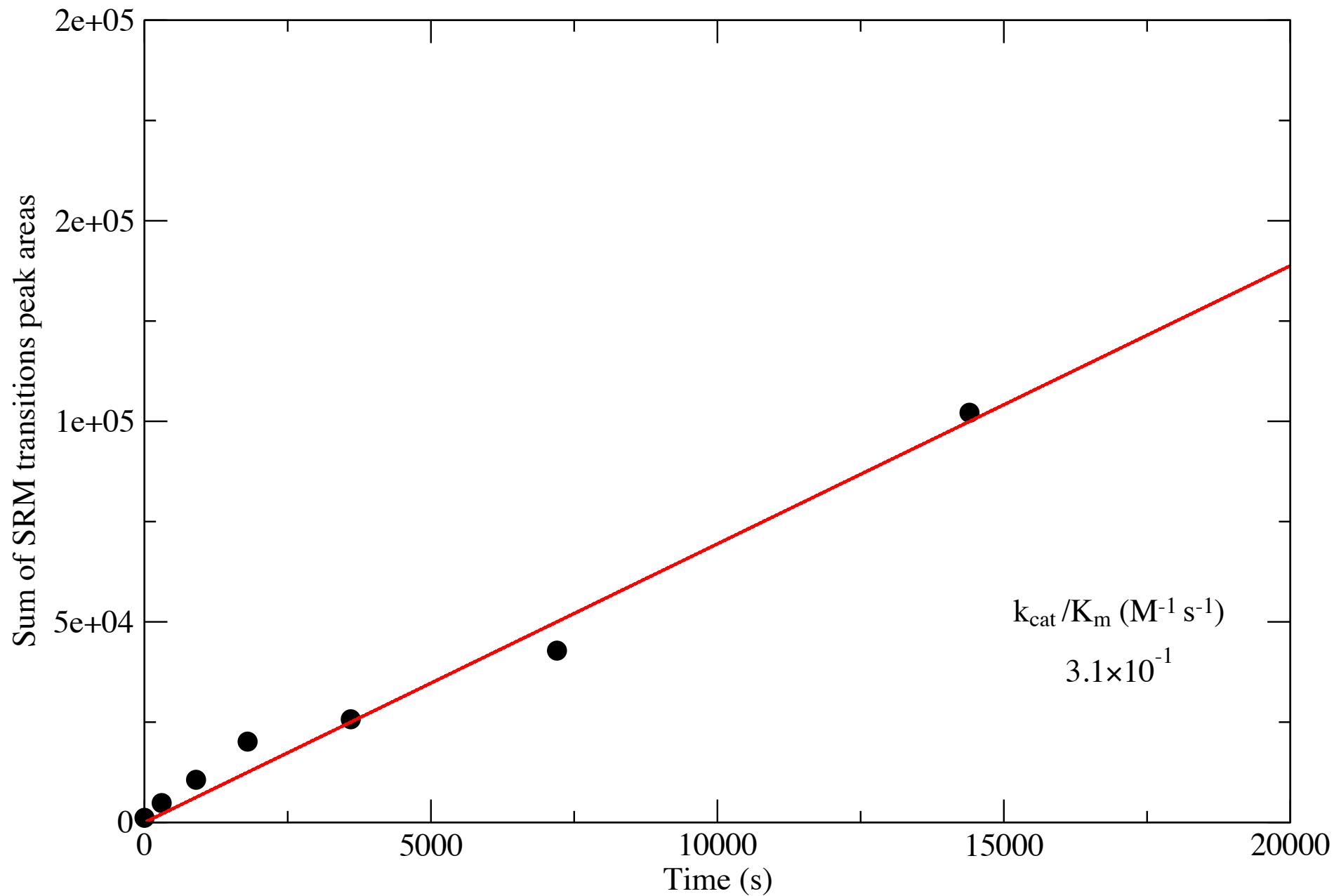
ACTB_HUMAN (GQVITIGNER)



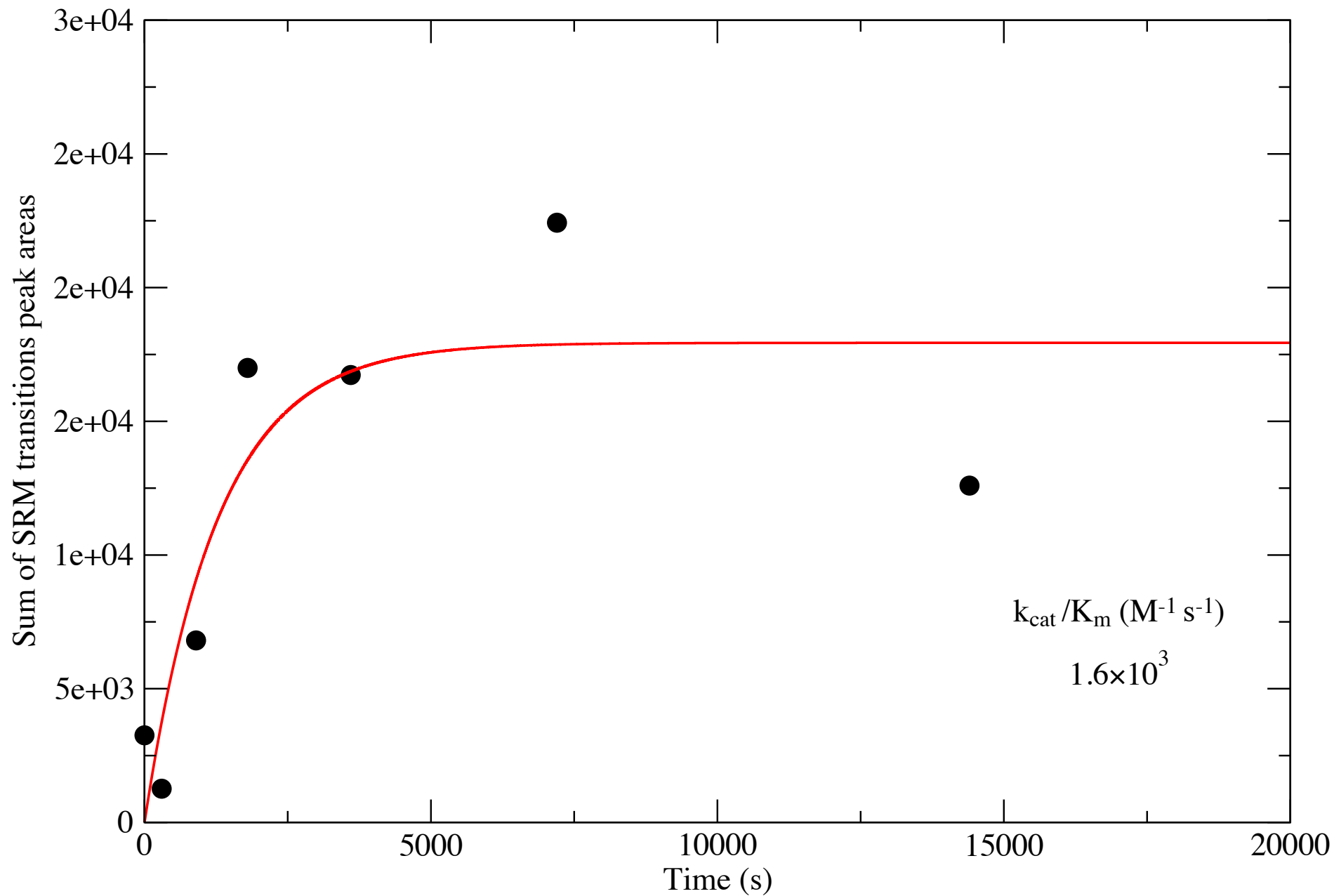
ACTB_HUMAN (GVTHTVPIYEGYALPHAILR)



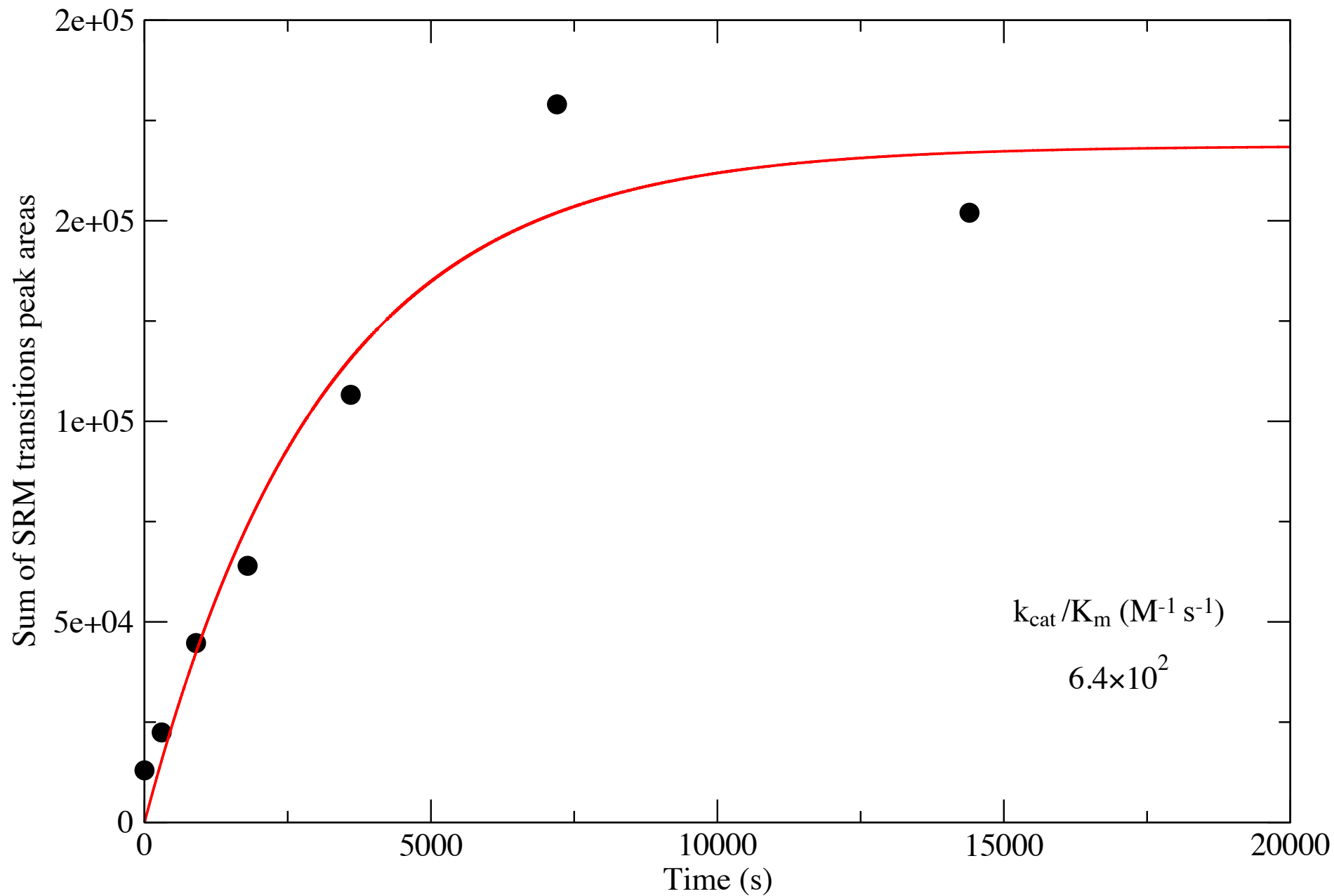
AKNA_HUMAN (GVAATPGKAEATR)



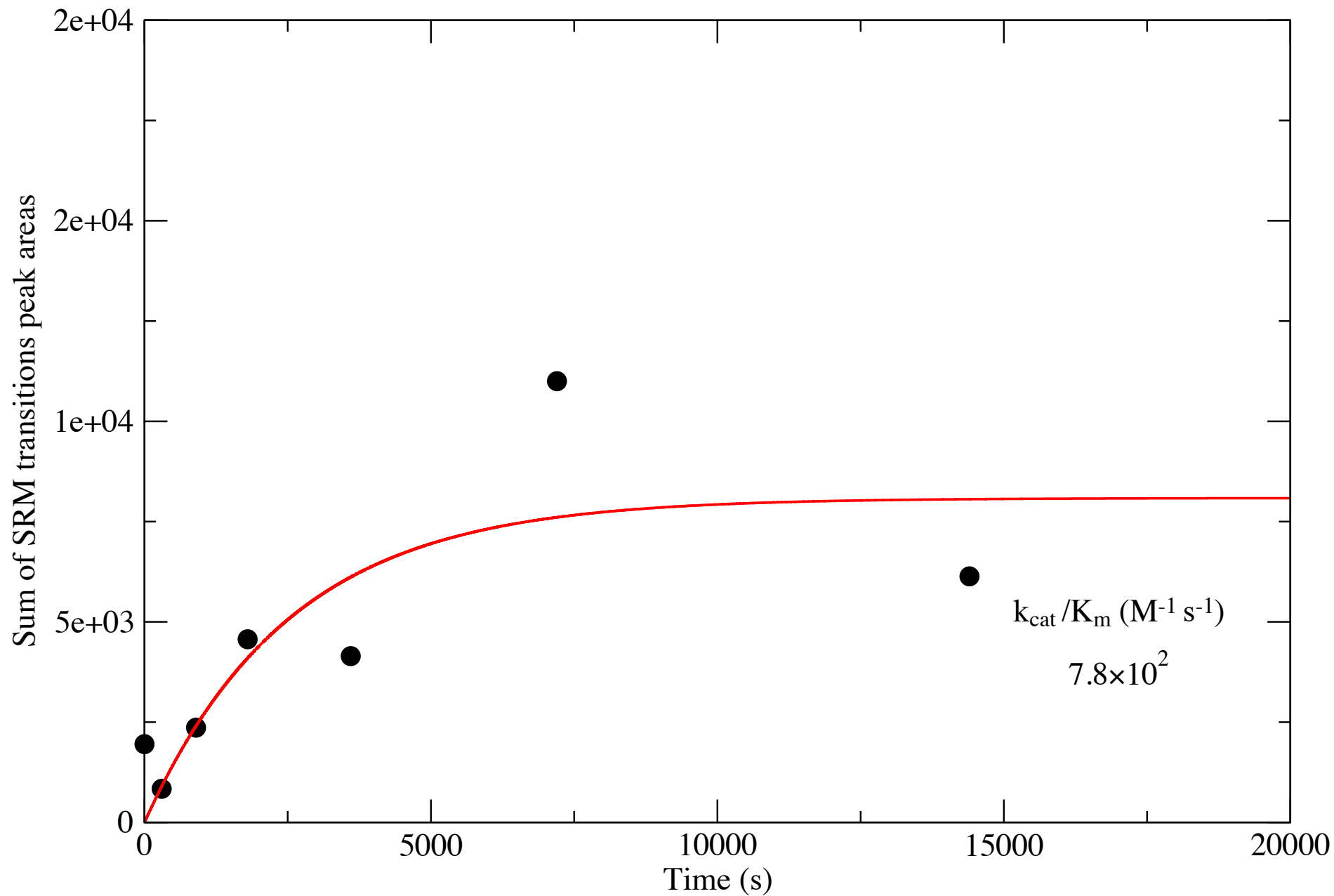
BAG6_HUMAN (GASAETEPWAAA VPPEWVPPIQQDIQSQRK)



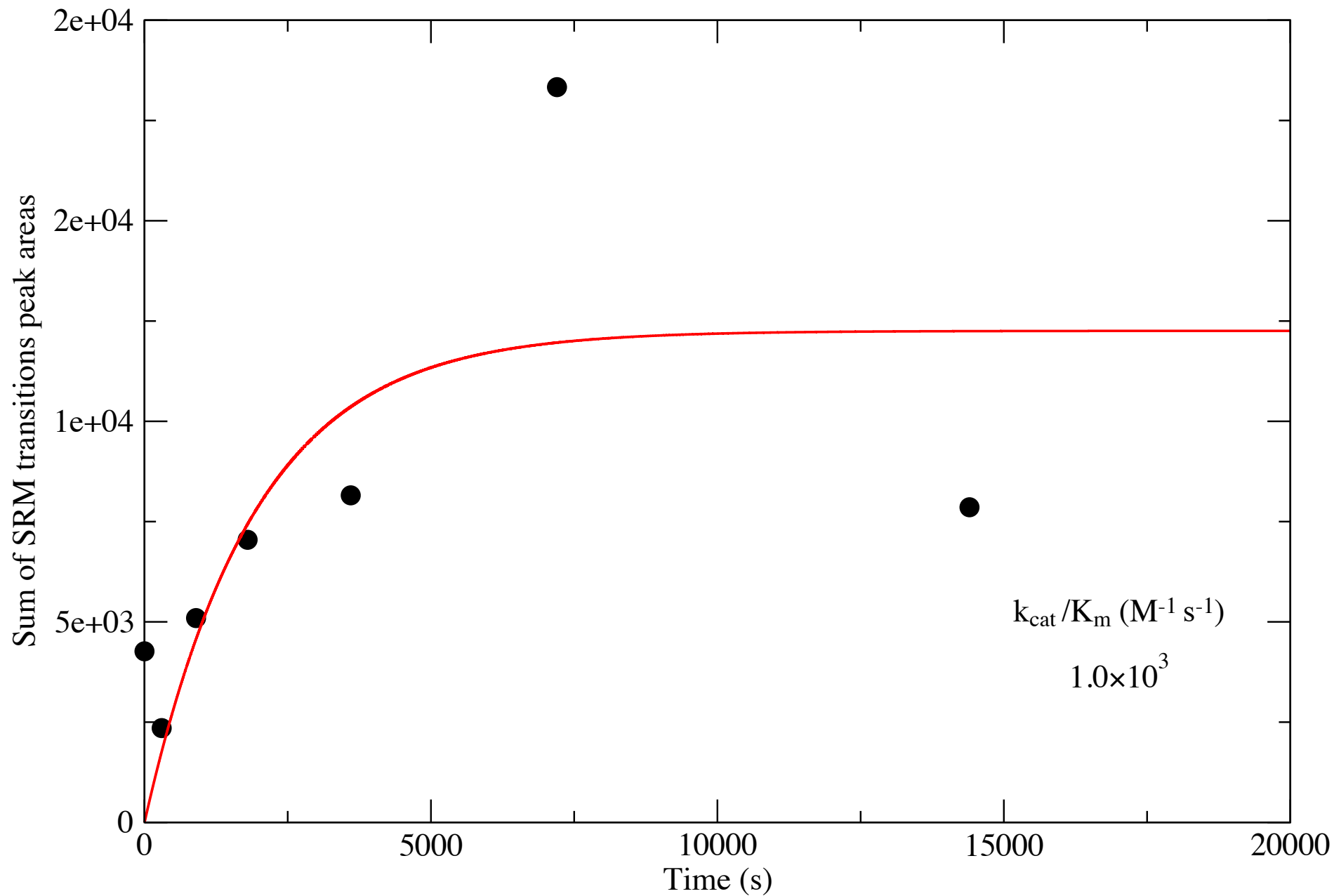
CC50A_HUMAN (GGPPCAPGGTAK)



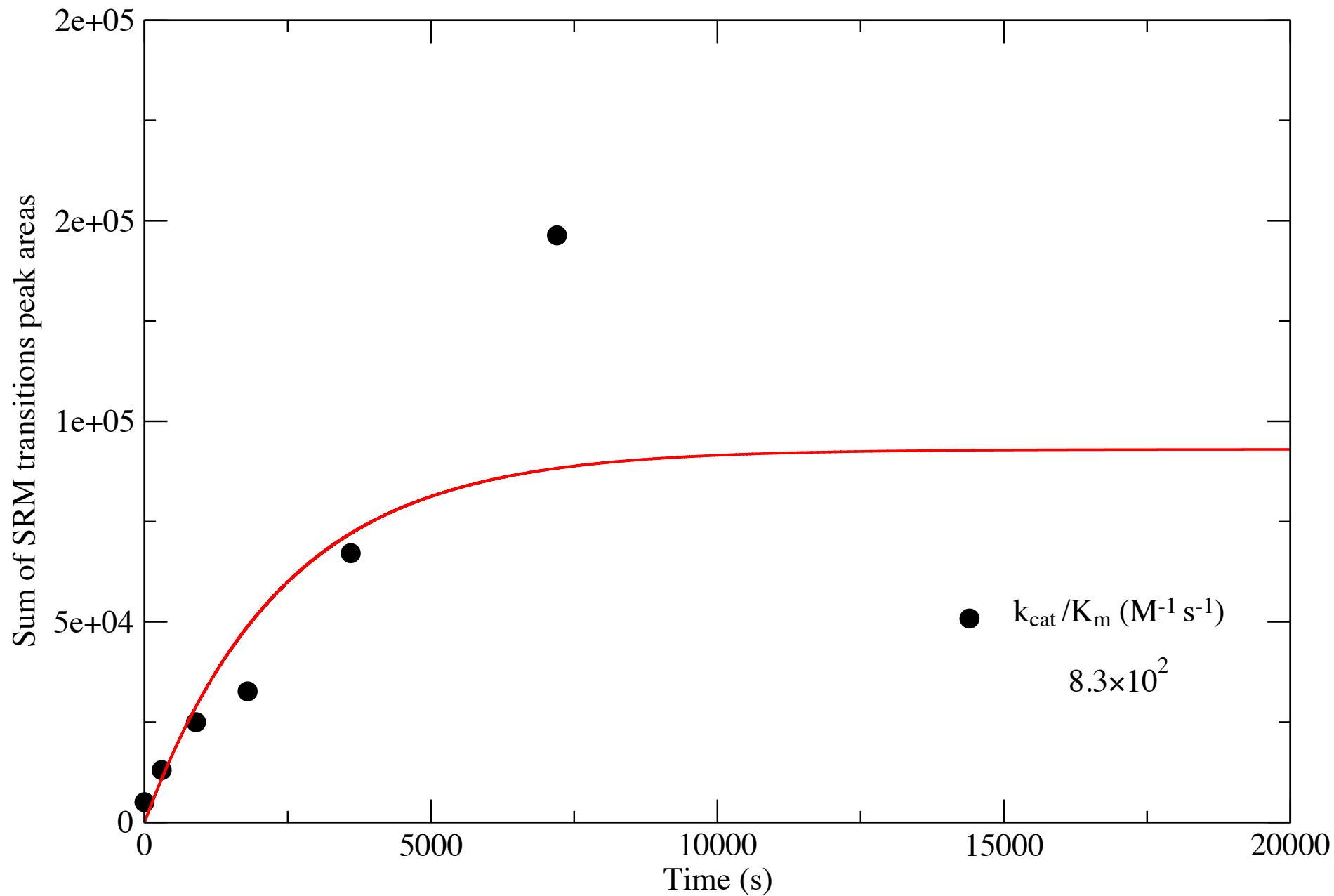
CLAP1_HUMAN (GGAASPATEGRGGSEVEGGR)



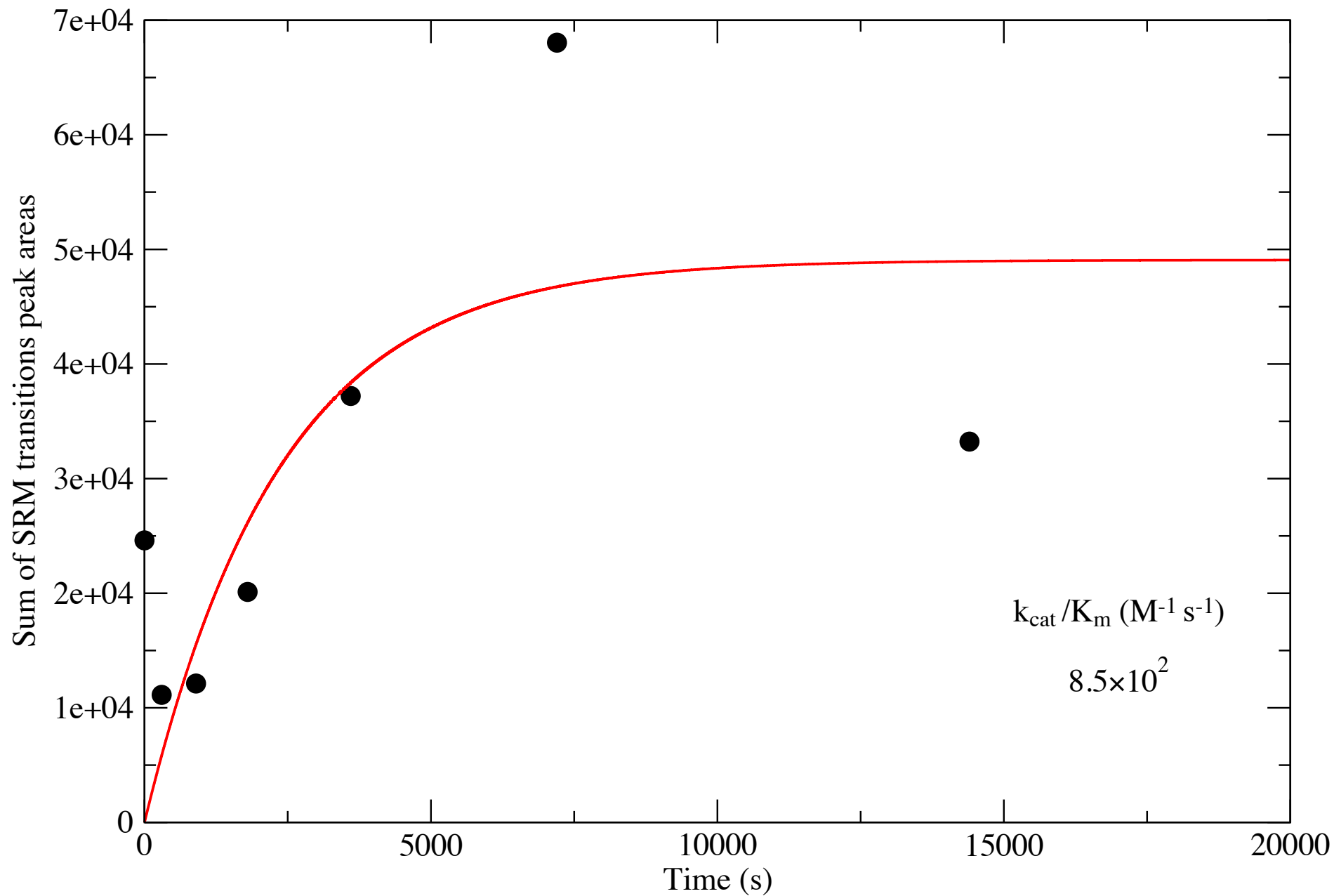
CR025_HUMAN (GVADSTVISSMPCLLMELRR)



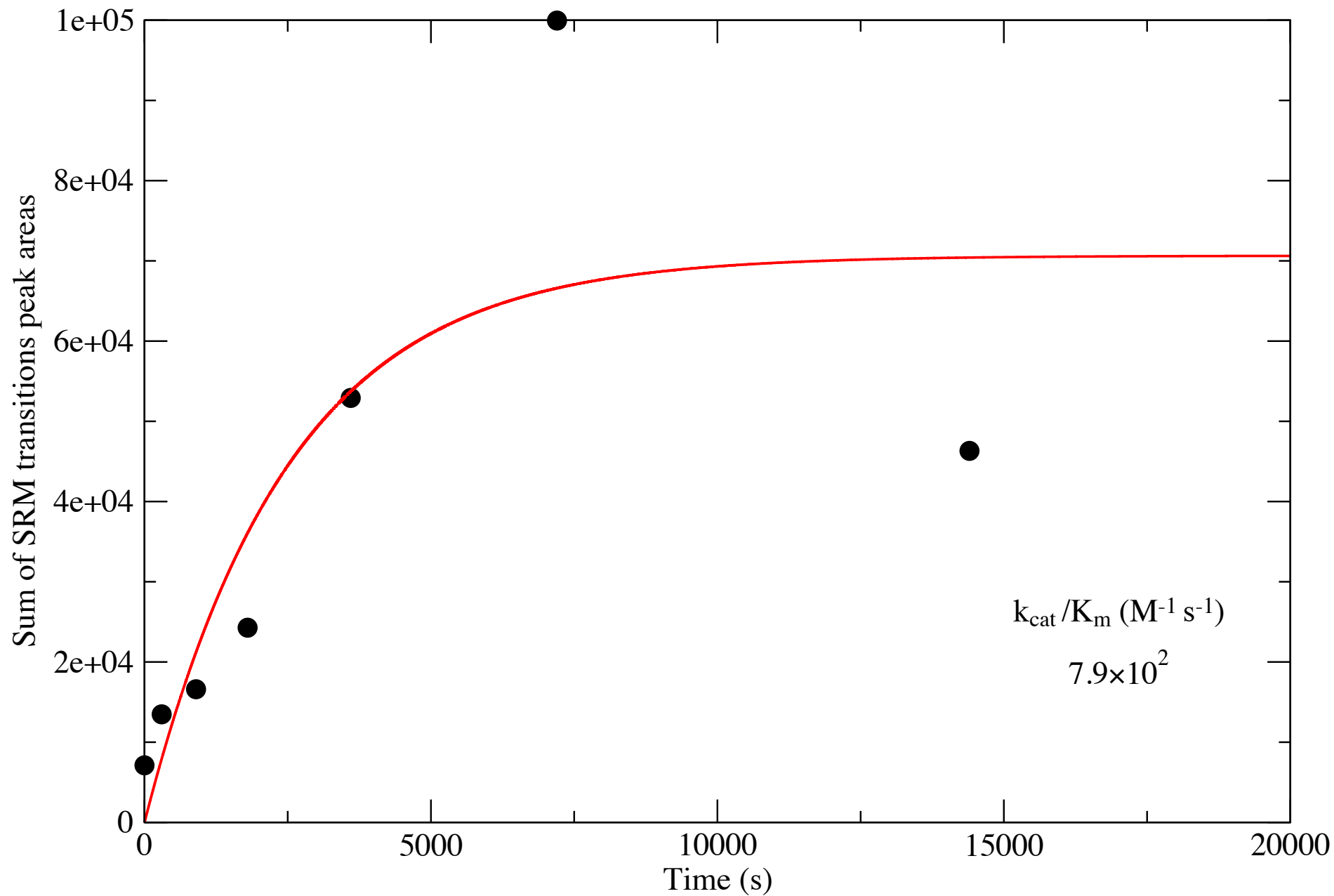
CTU2_HUMAN (SATAFGAQTSSR)



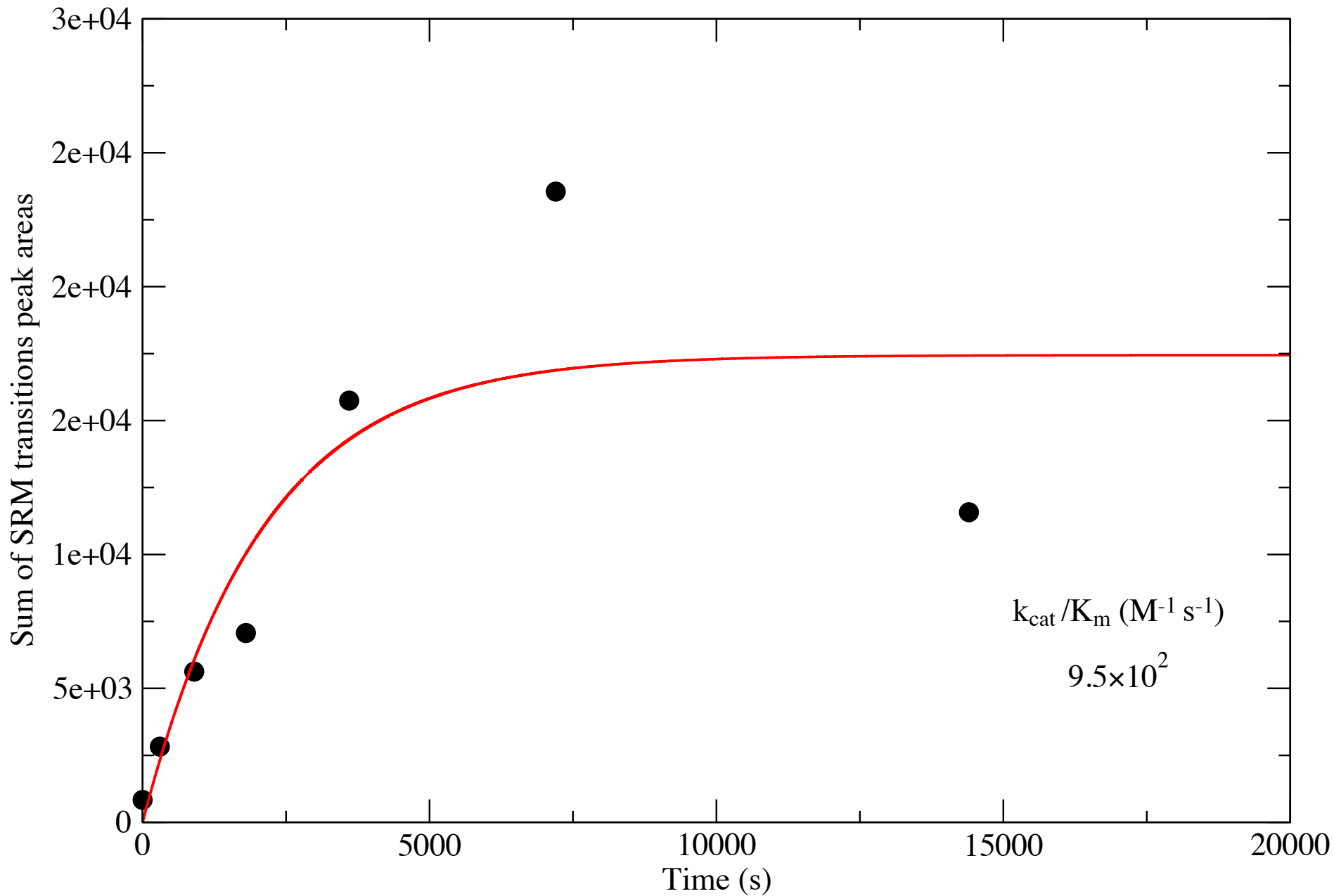
DGCR8_HUMAN (SMGADPGPPDEKDPPLGAEAAPGALGQVK)



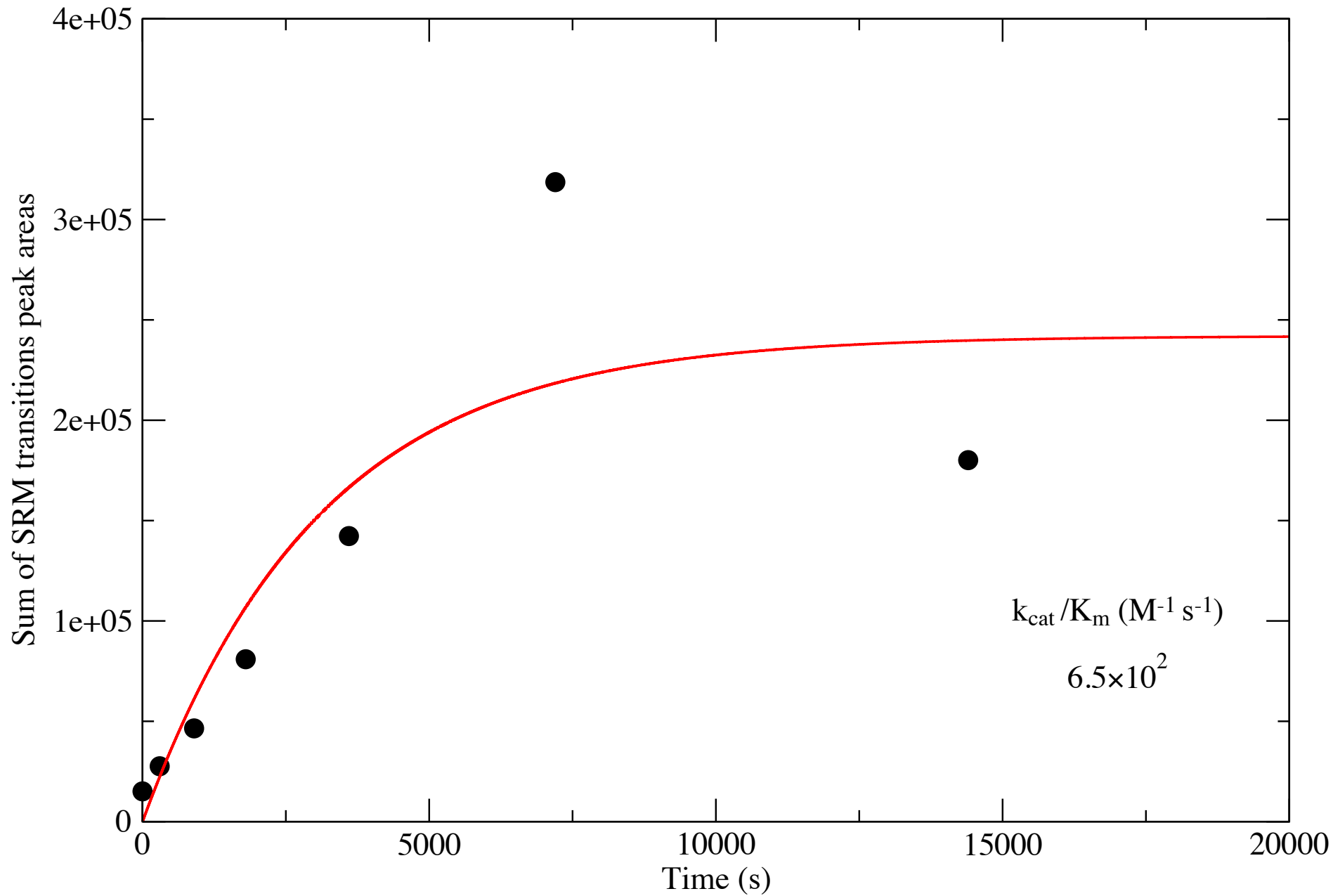
DHX30_HUMAN (FLSMTQQDSHAPLRDSR)



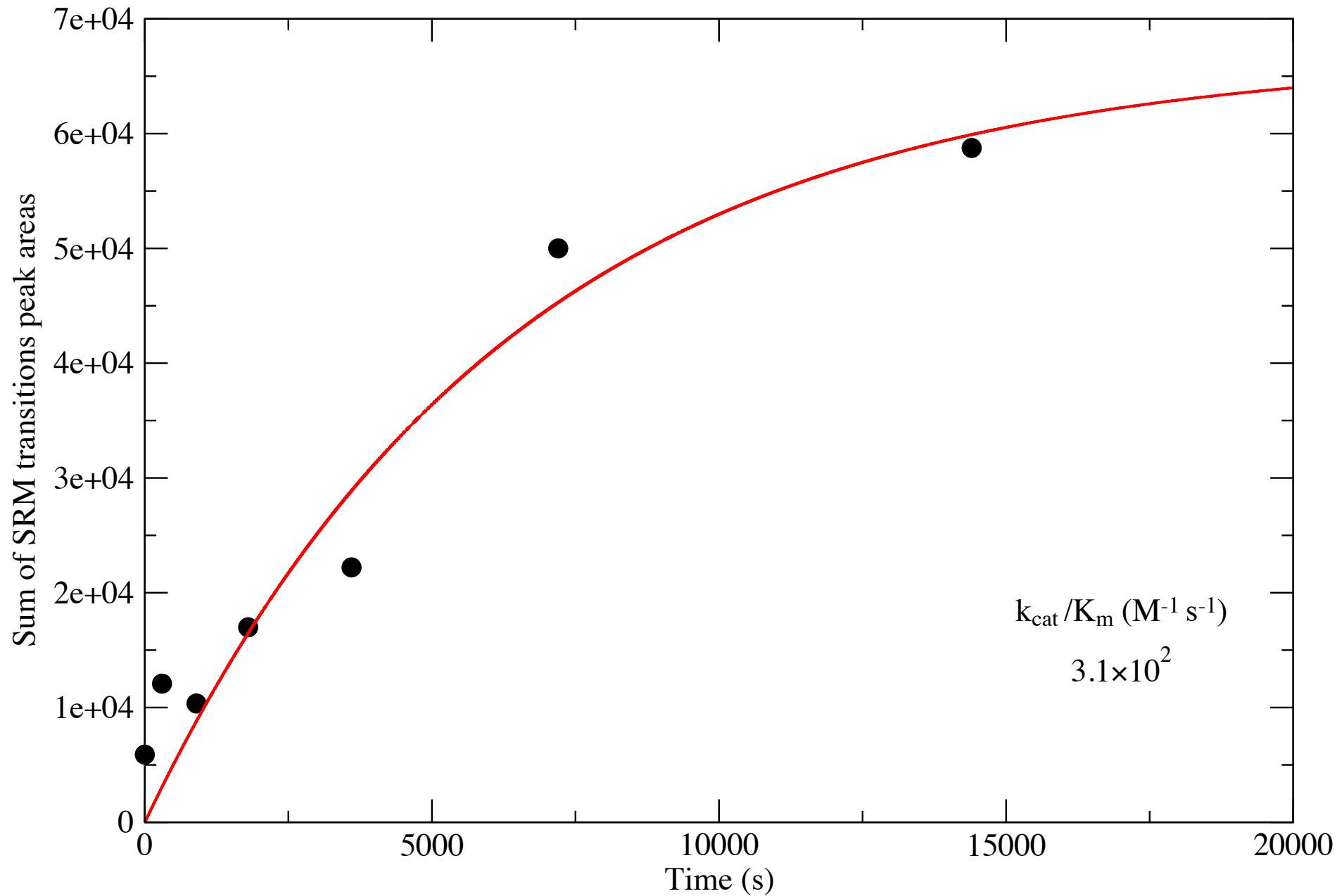
DSRAD_HUMAN (GHSQGAPNSDPSLEPEDR)



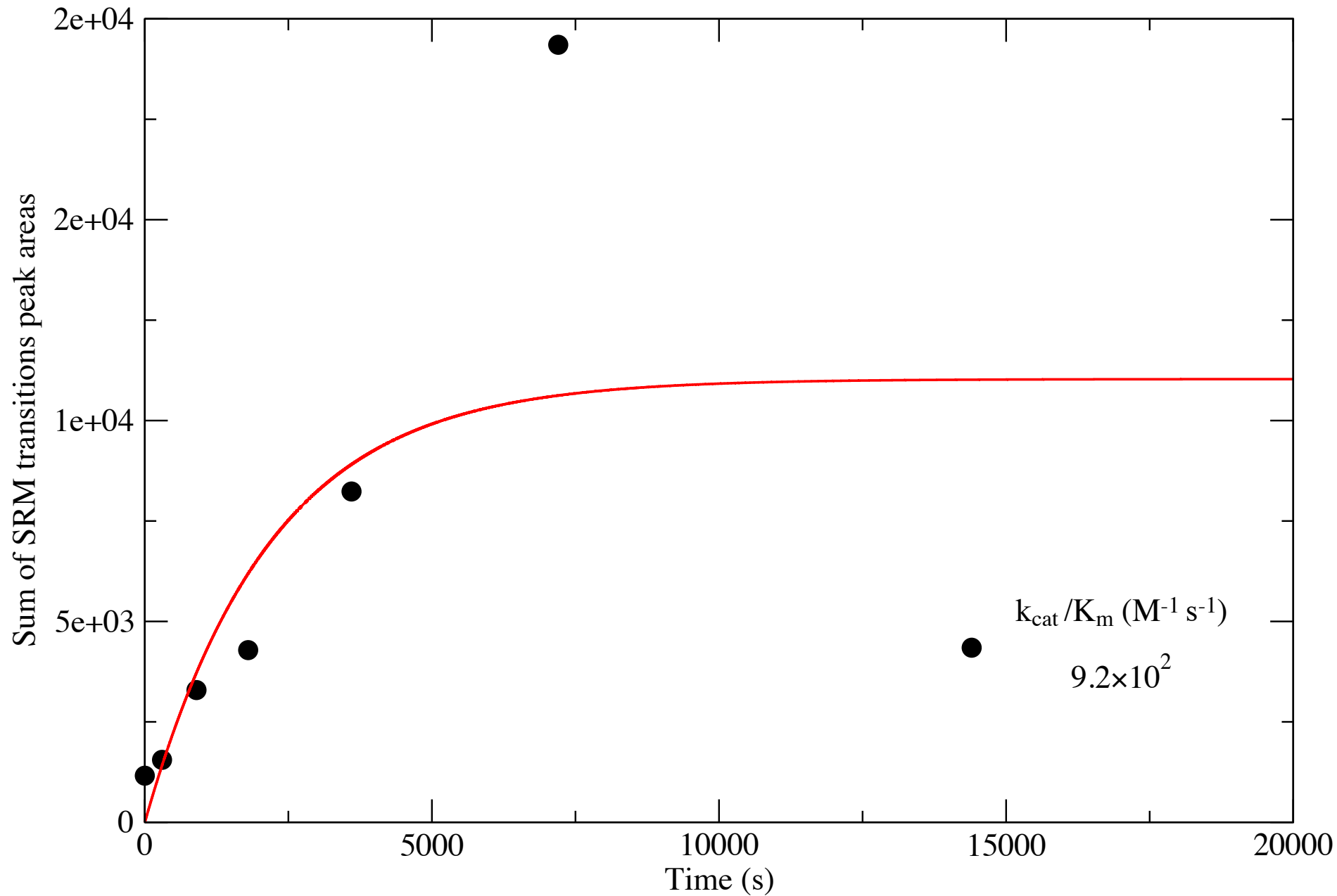
ENPL_HUMAN (GLNASQIR)



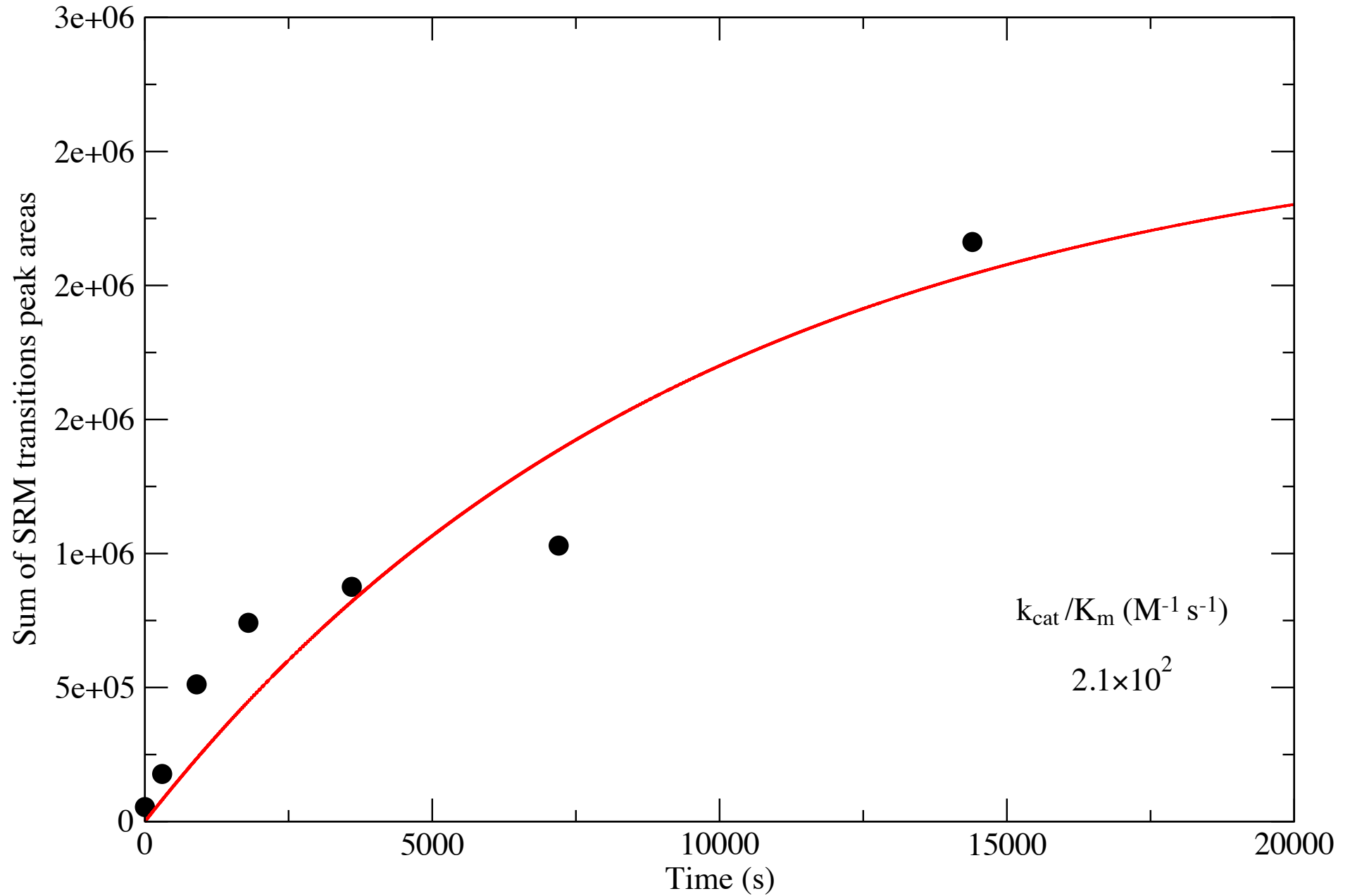
ENPL_HUMAN (GTVEEDLGK)



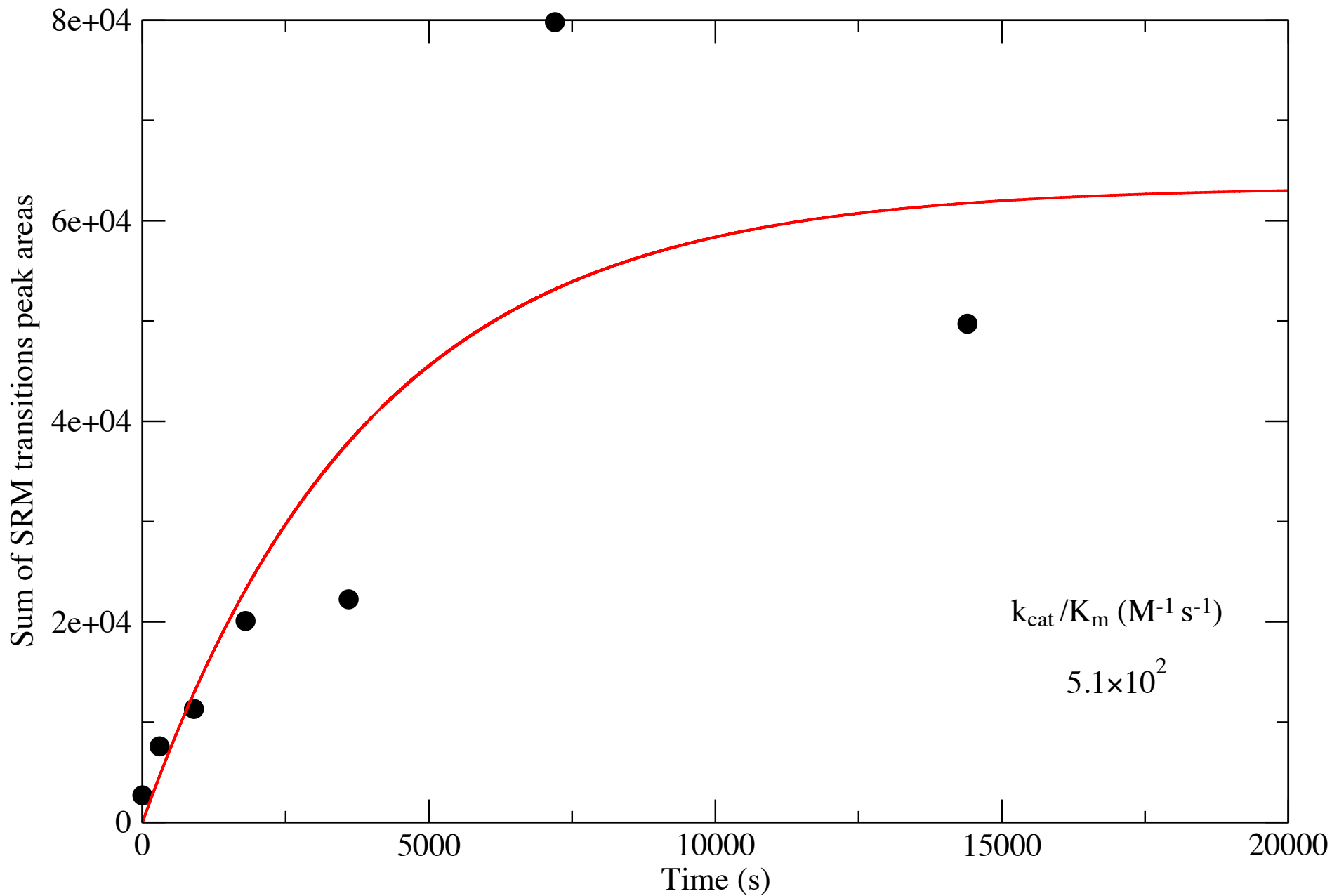
ERO1A_HUMAN (GIKSASYKYSEEANNLIEECEQAER)



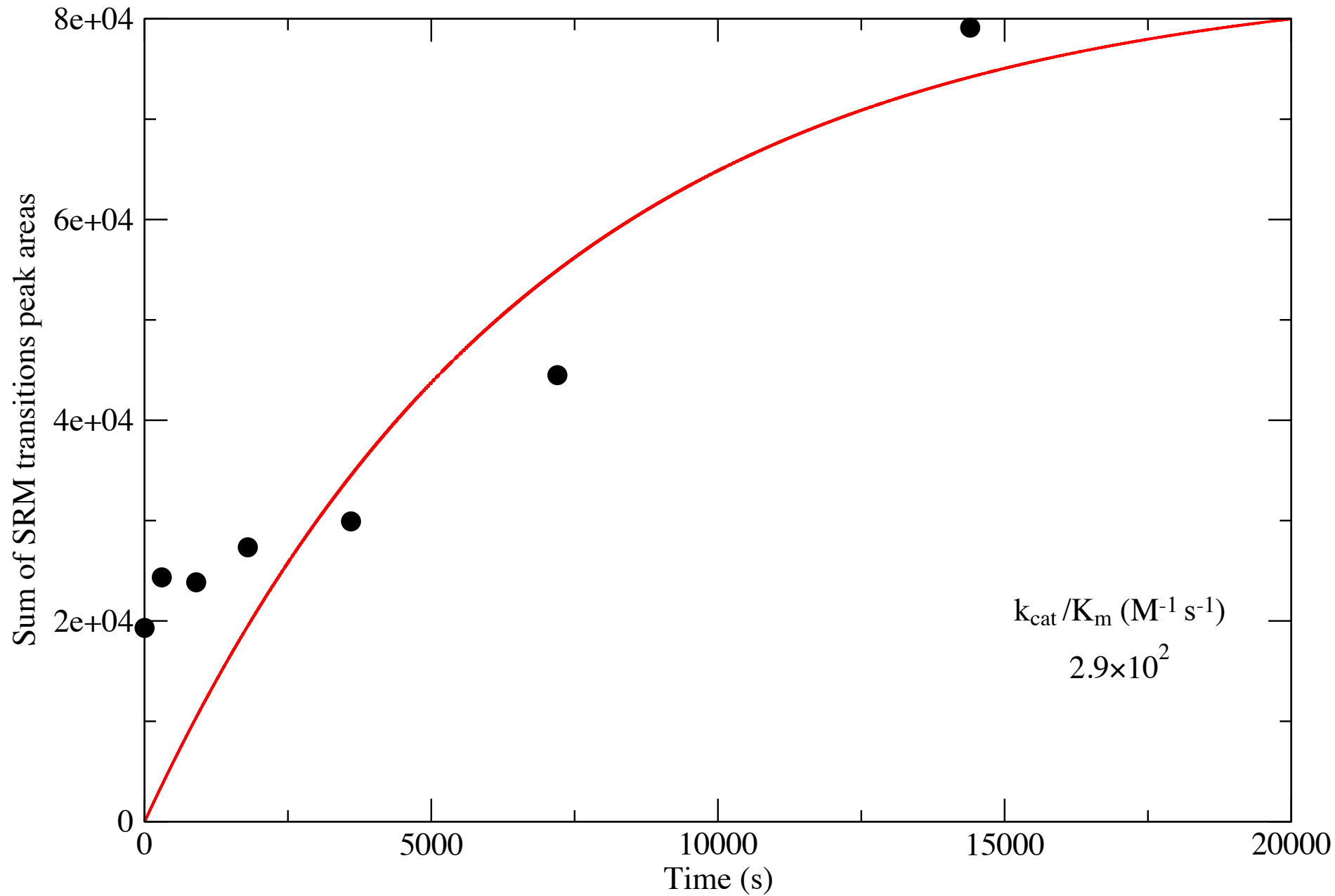
F117B_HUMAN (GHRAPPPLVQR)



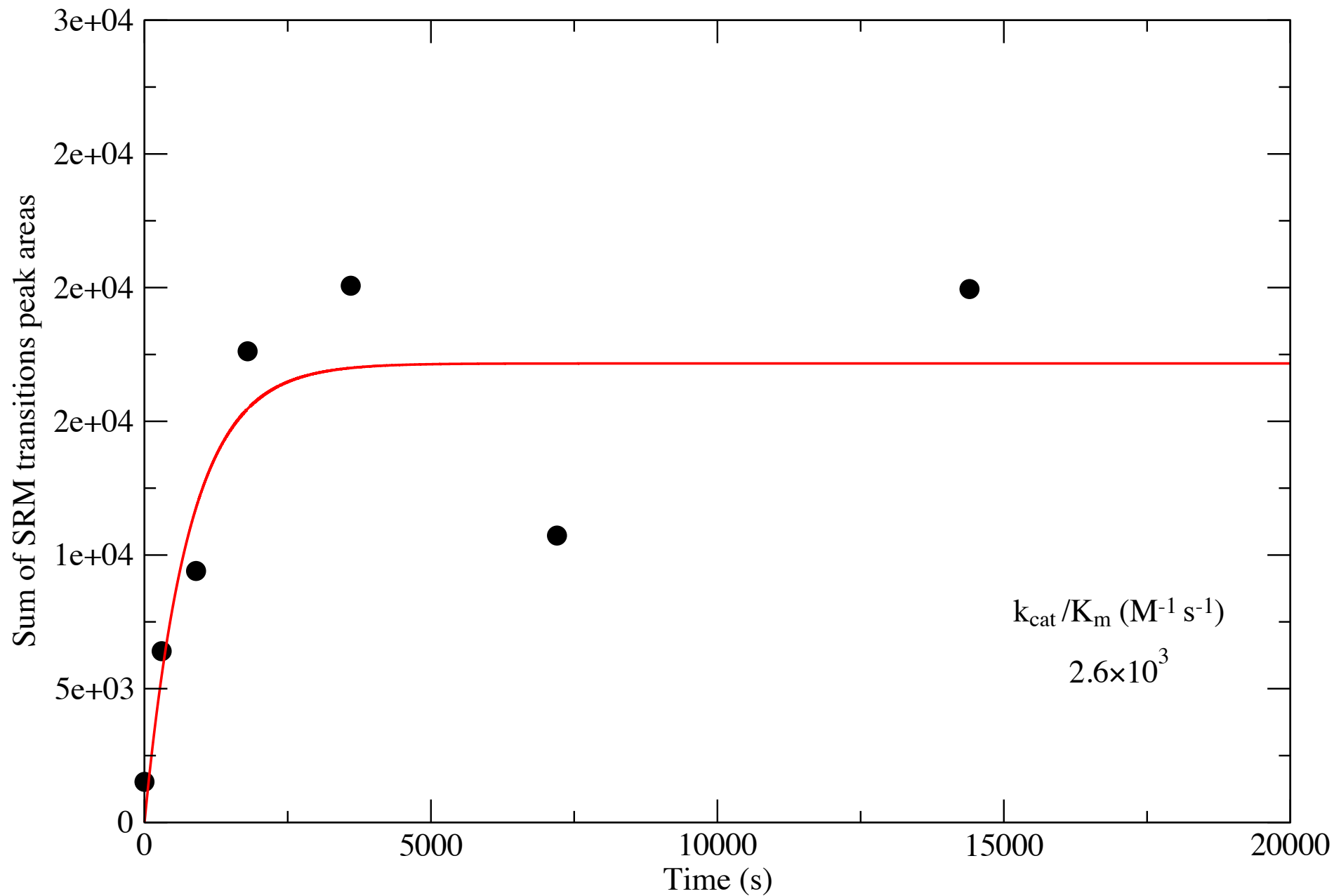
FYB_HUMAN (GFPAPPK)



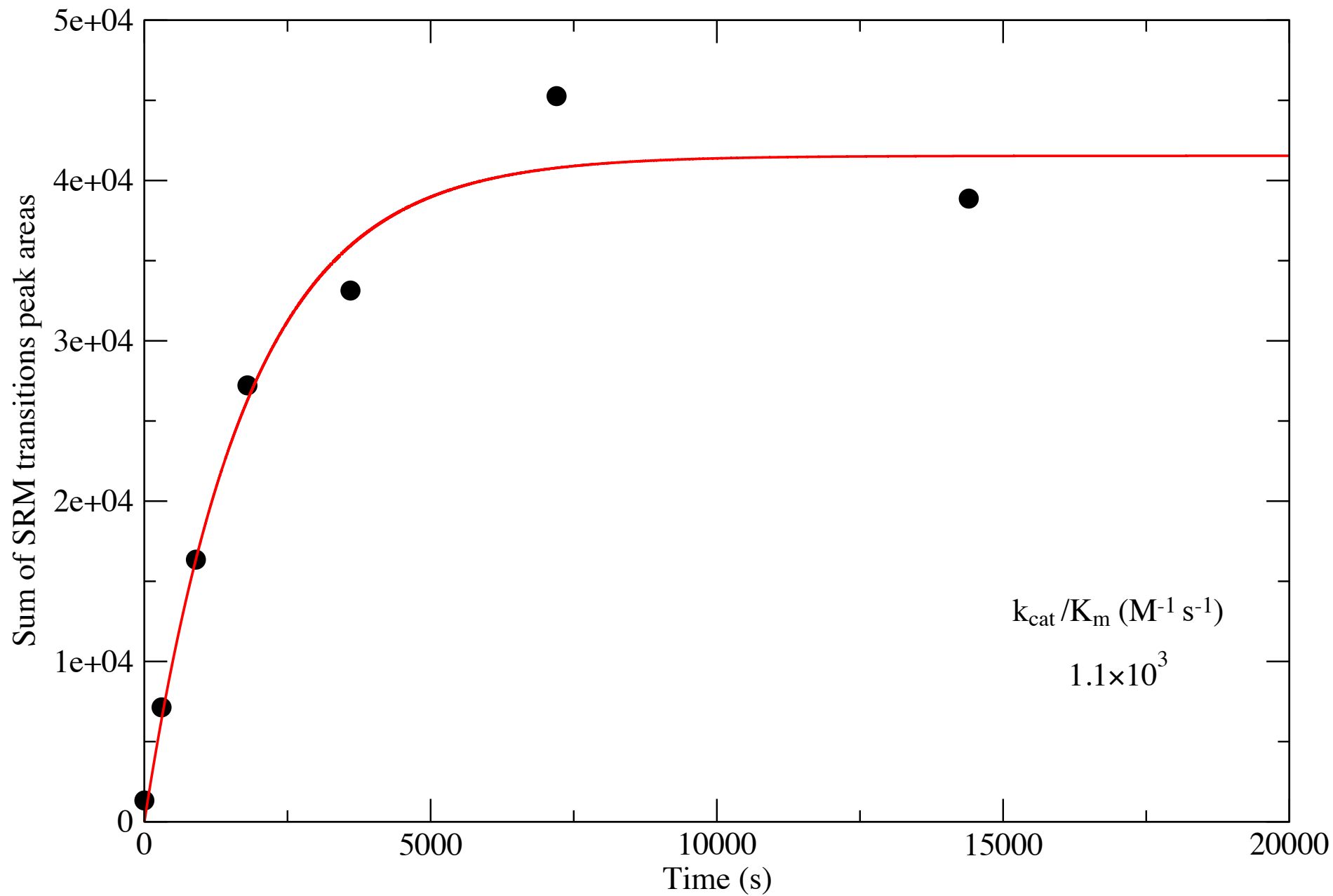
GDIR2_HUMAN (SKLNYKPPPQK)



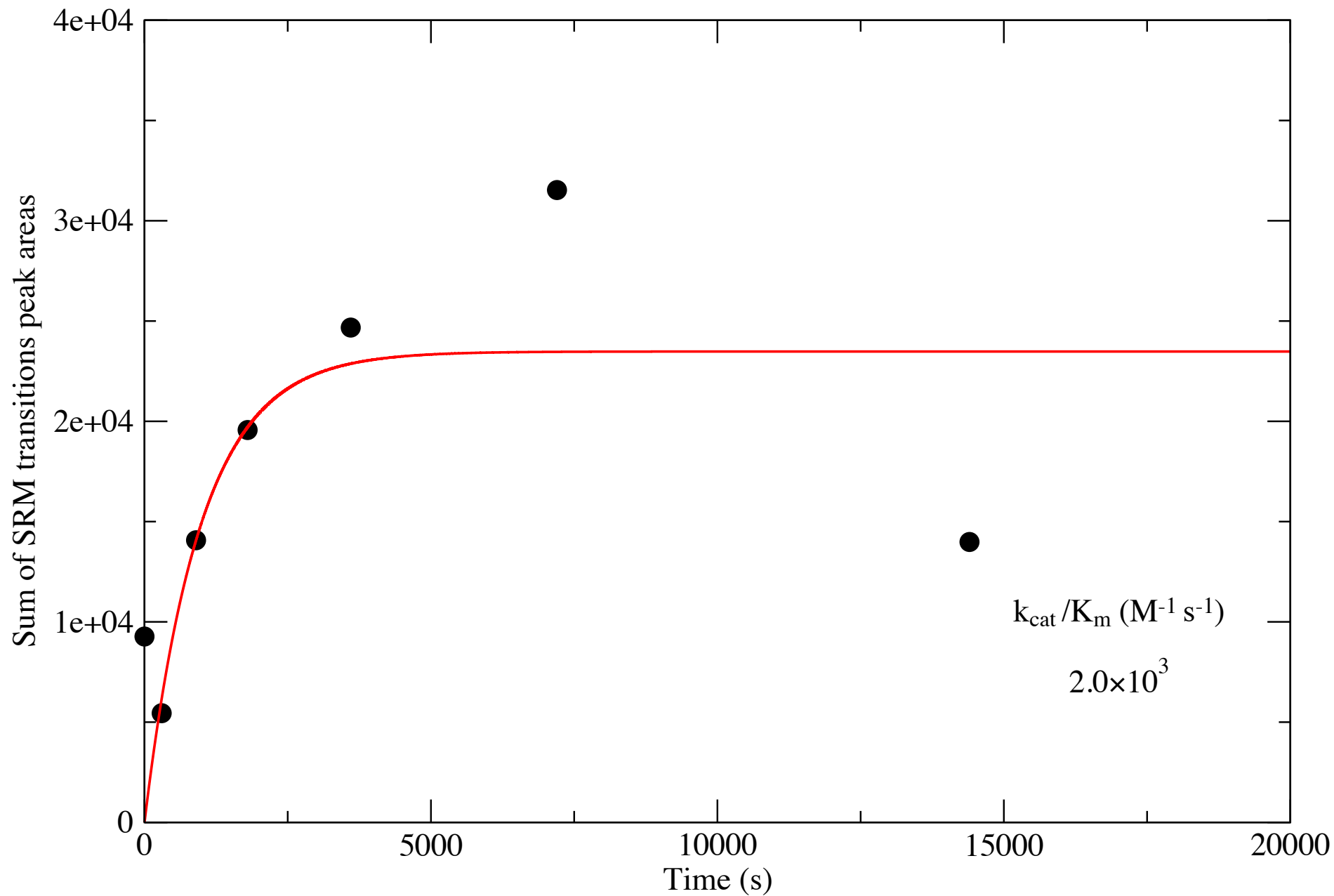
GMIP_HUMAN (GGGEVSSQGPEDSLLGTQSR)



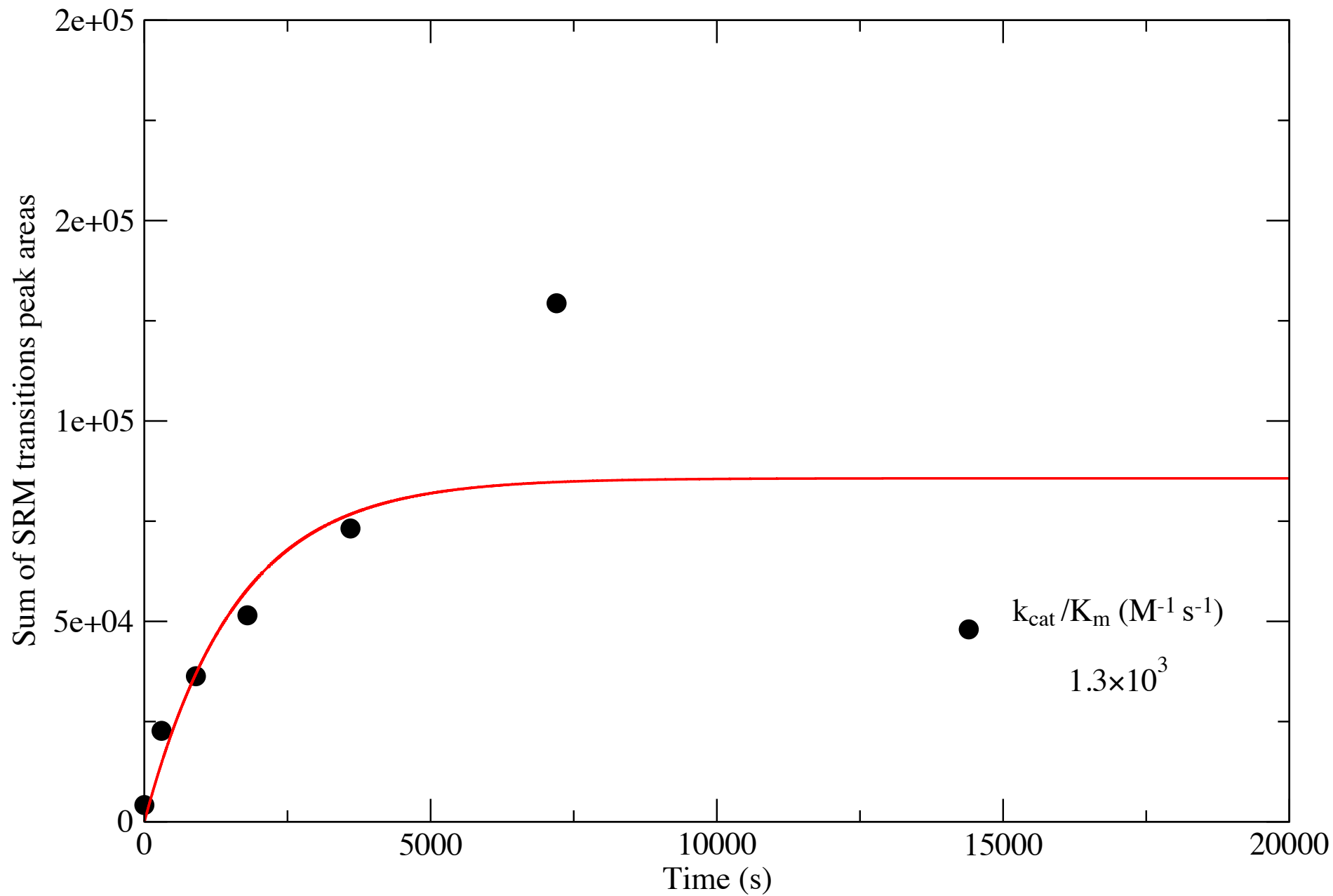
GNL1_HUMAN (GESVTHHIR)



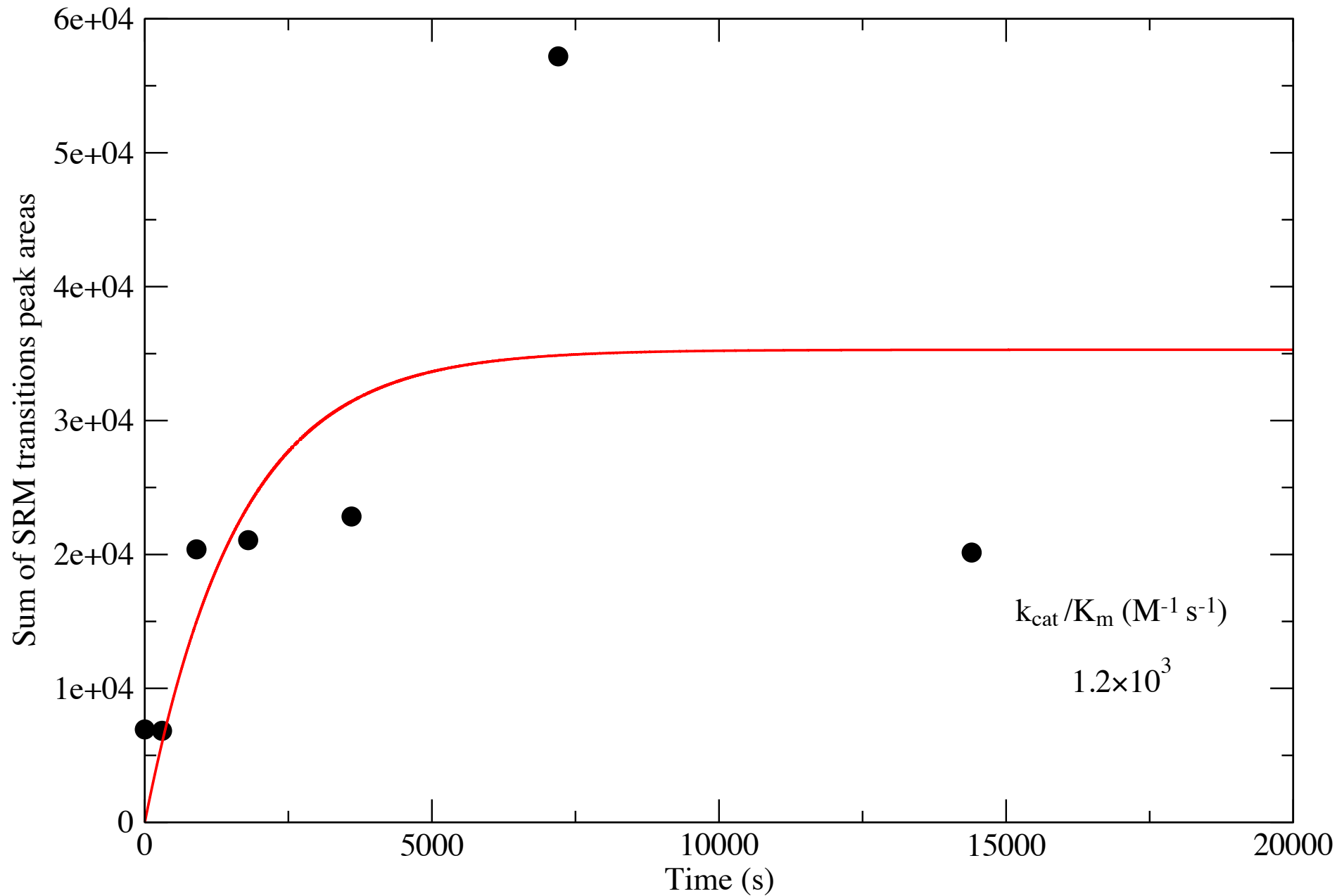
GRAP2_HUMAN (GHC GTGLGSEMNAALMHR)



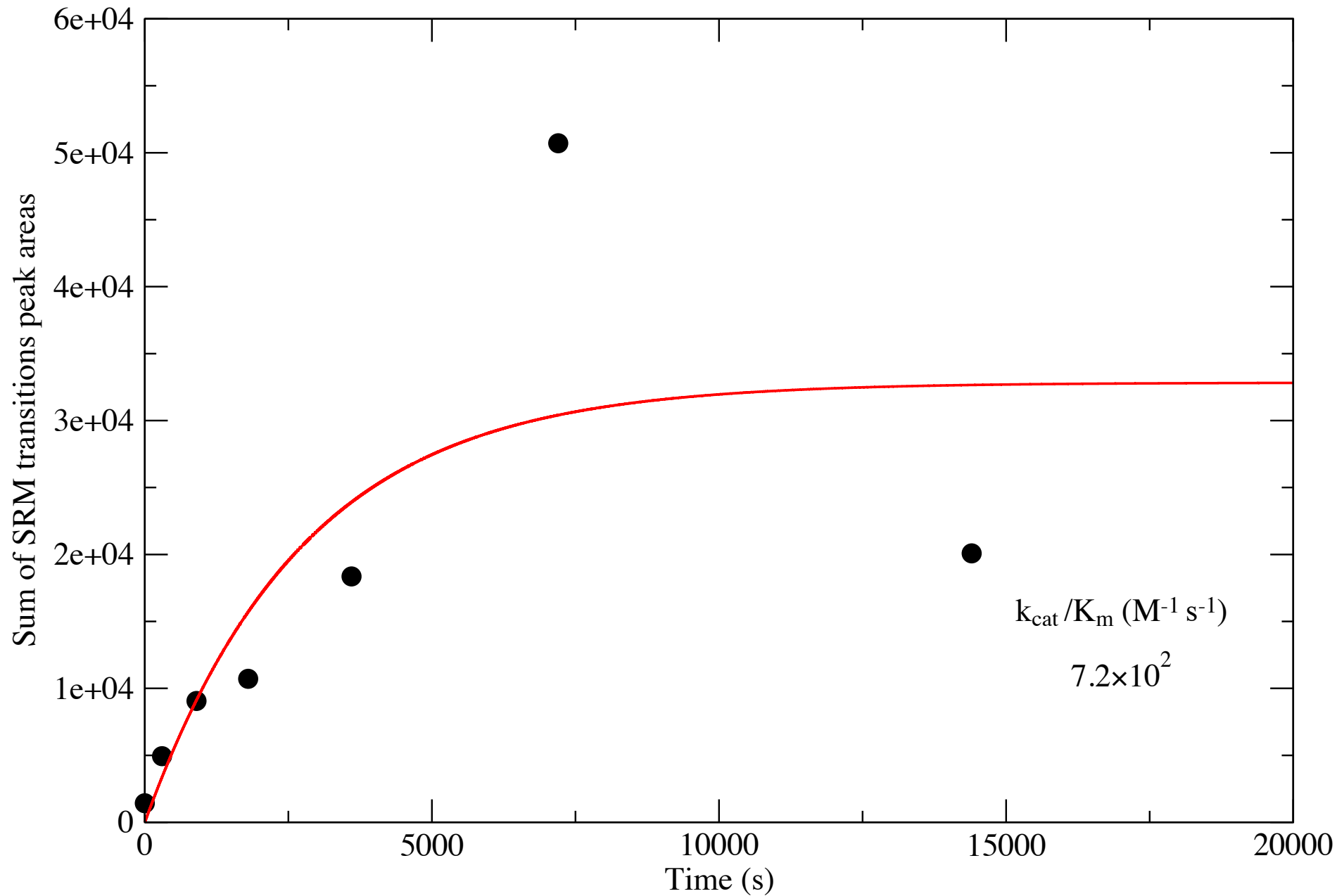
HDAC6_HUMAN (SMLMQGSR)



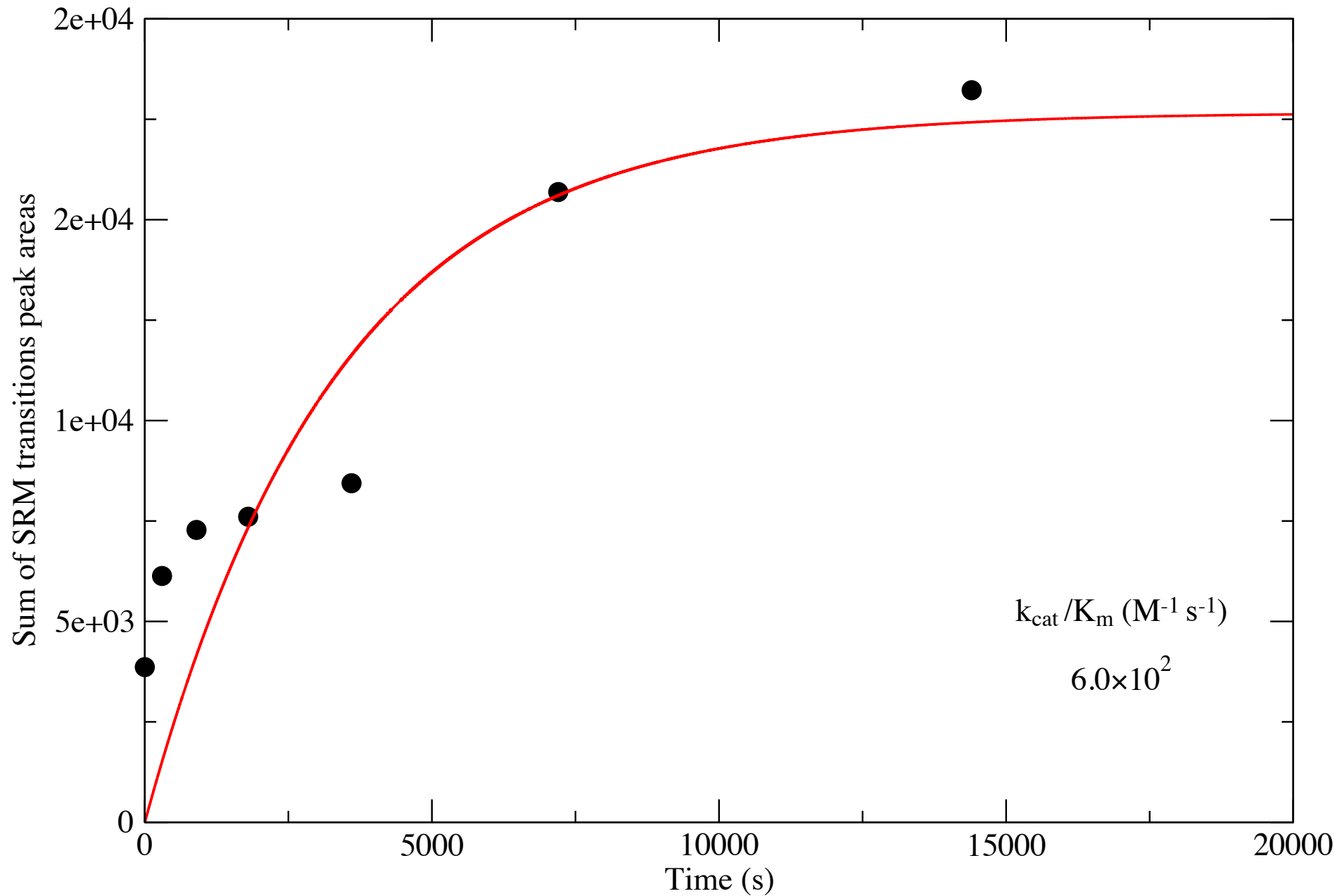
HJURP_HUMAN (GSVQAAA WGPELPSHR)



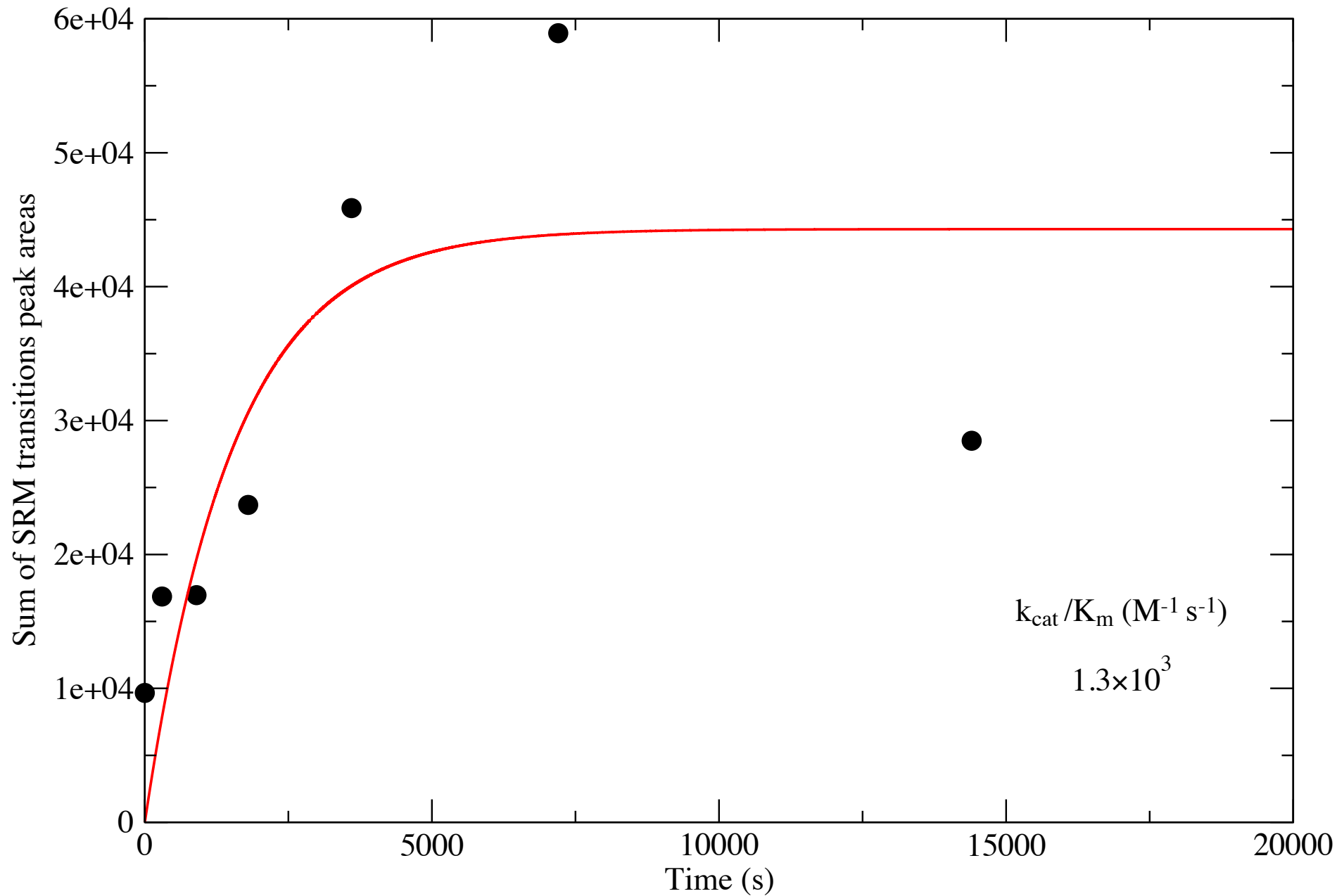
HUWE1_HUMAN (GGSGNSTIIVSR)



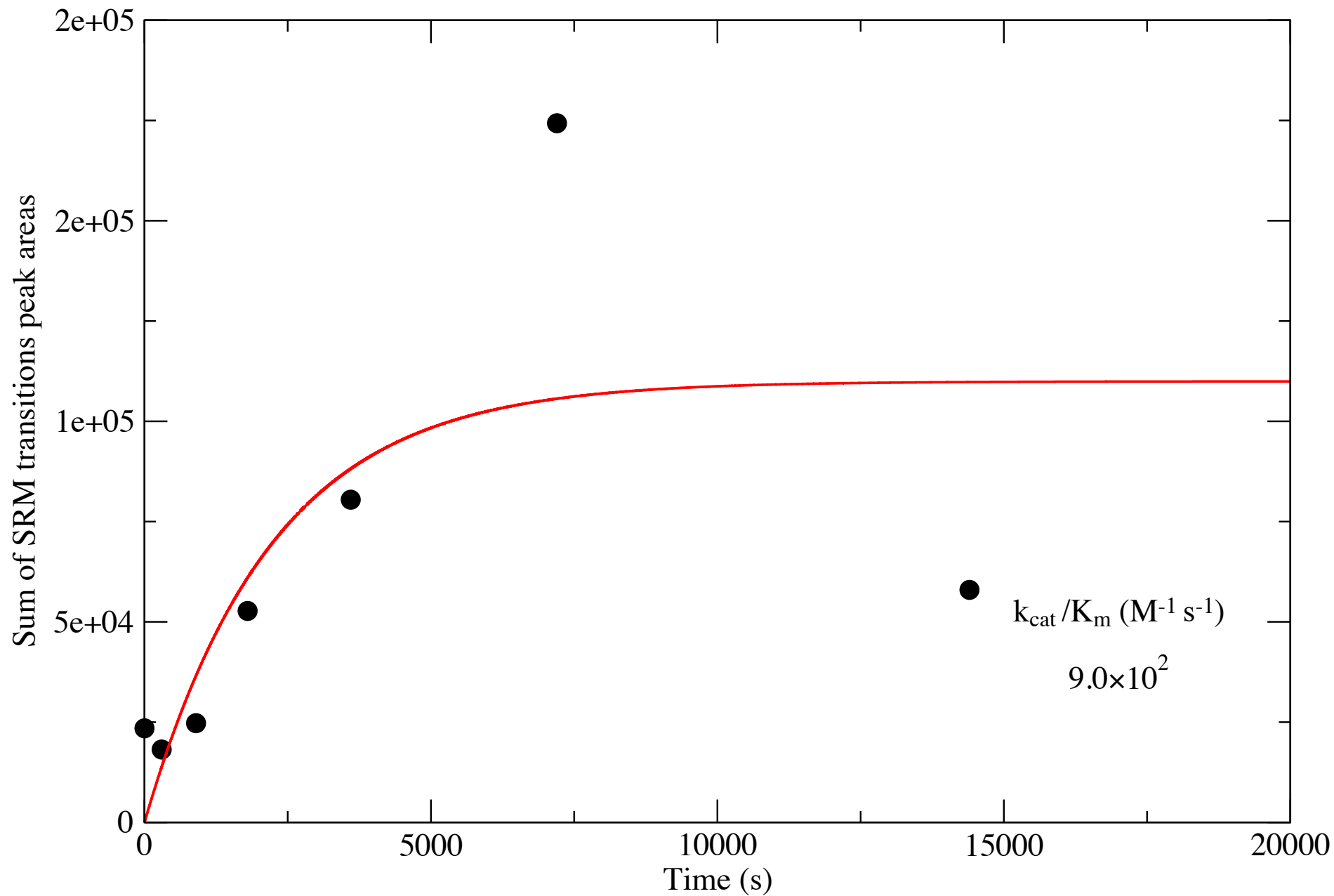
IF4H_HUMAN (GALLGDR)



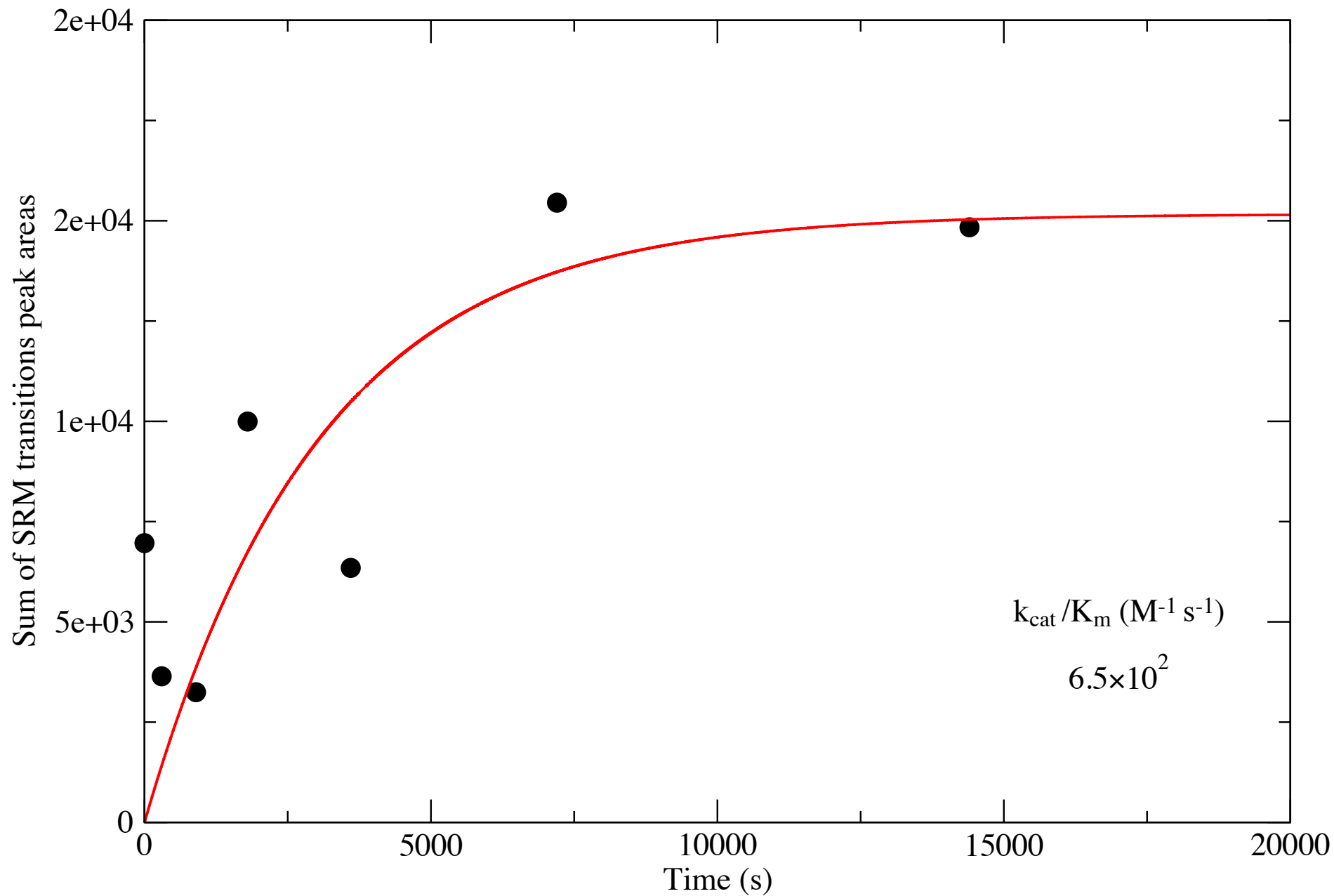
KMT2D_HUMAN (ALYVACQGQPK)



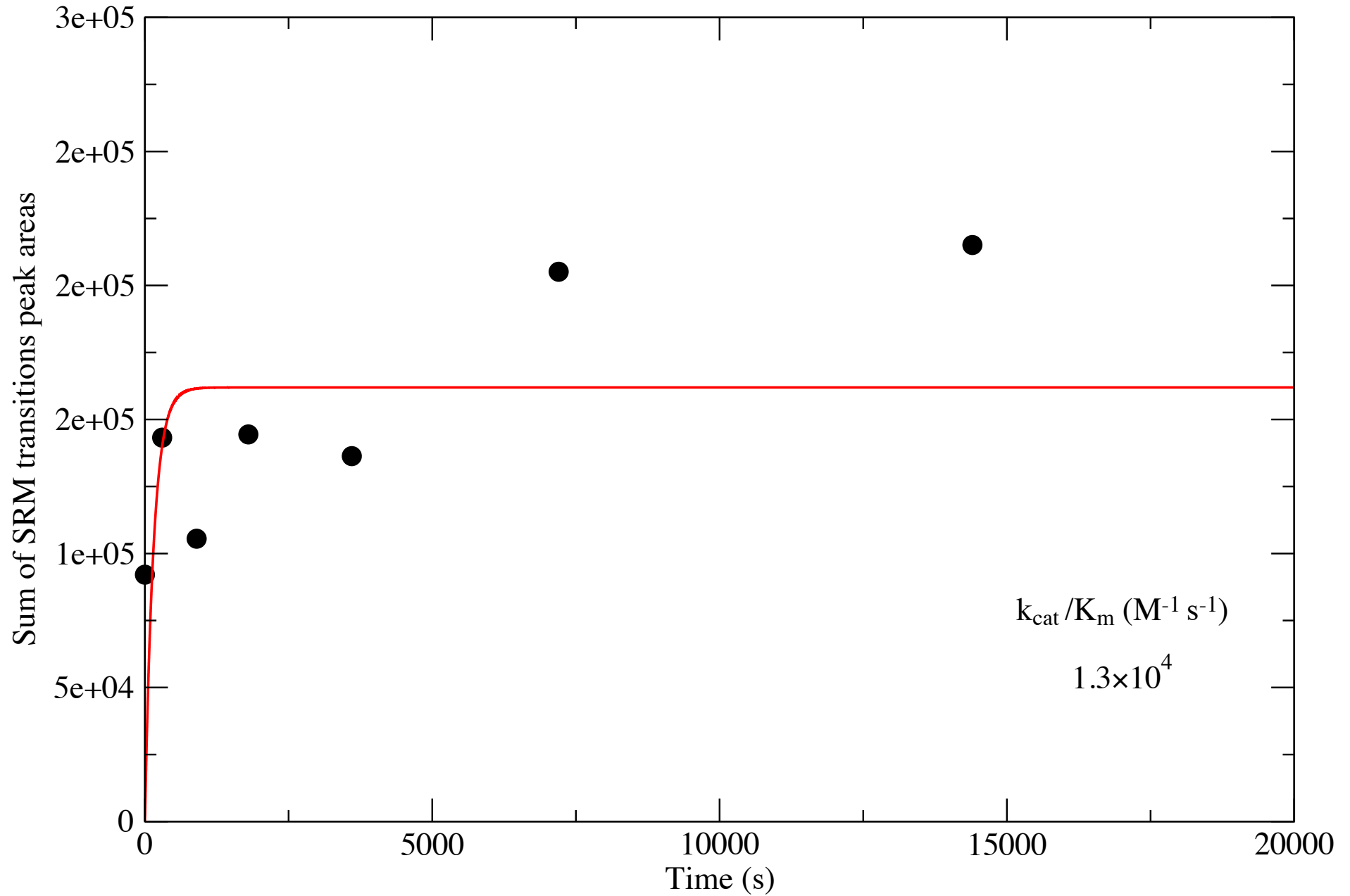
MCM4_HUMAN (GAAAEDIVASEQSLGQK)



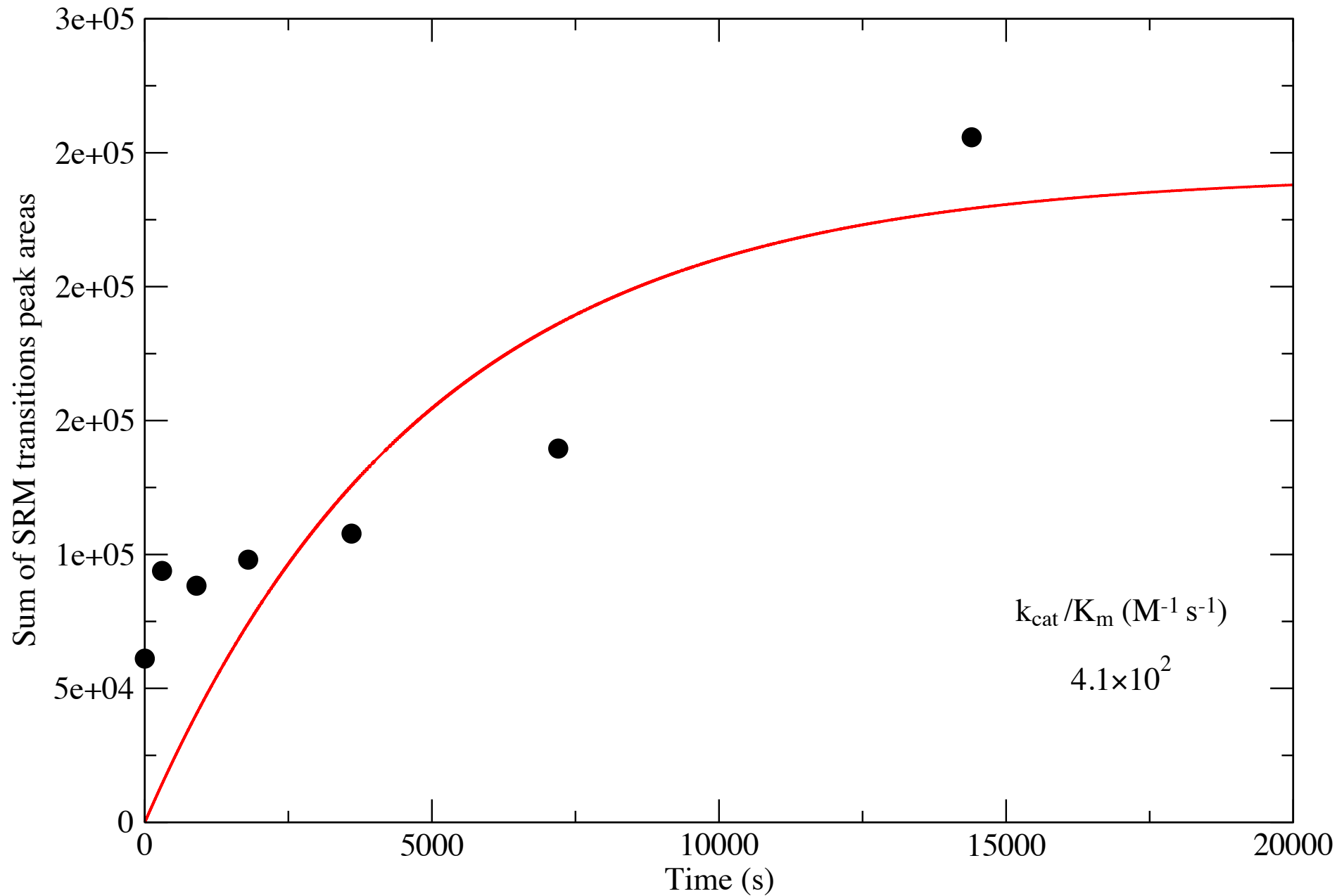
NMT1_HUMAN (SAQDQPVKMNSLPAER)



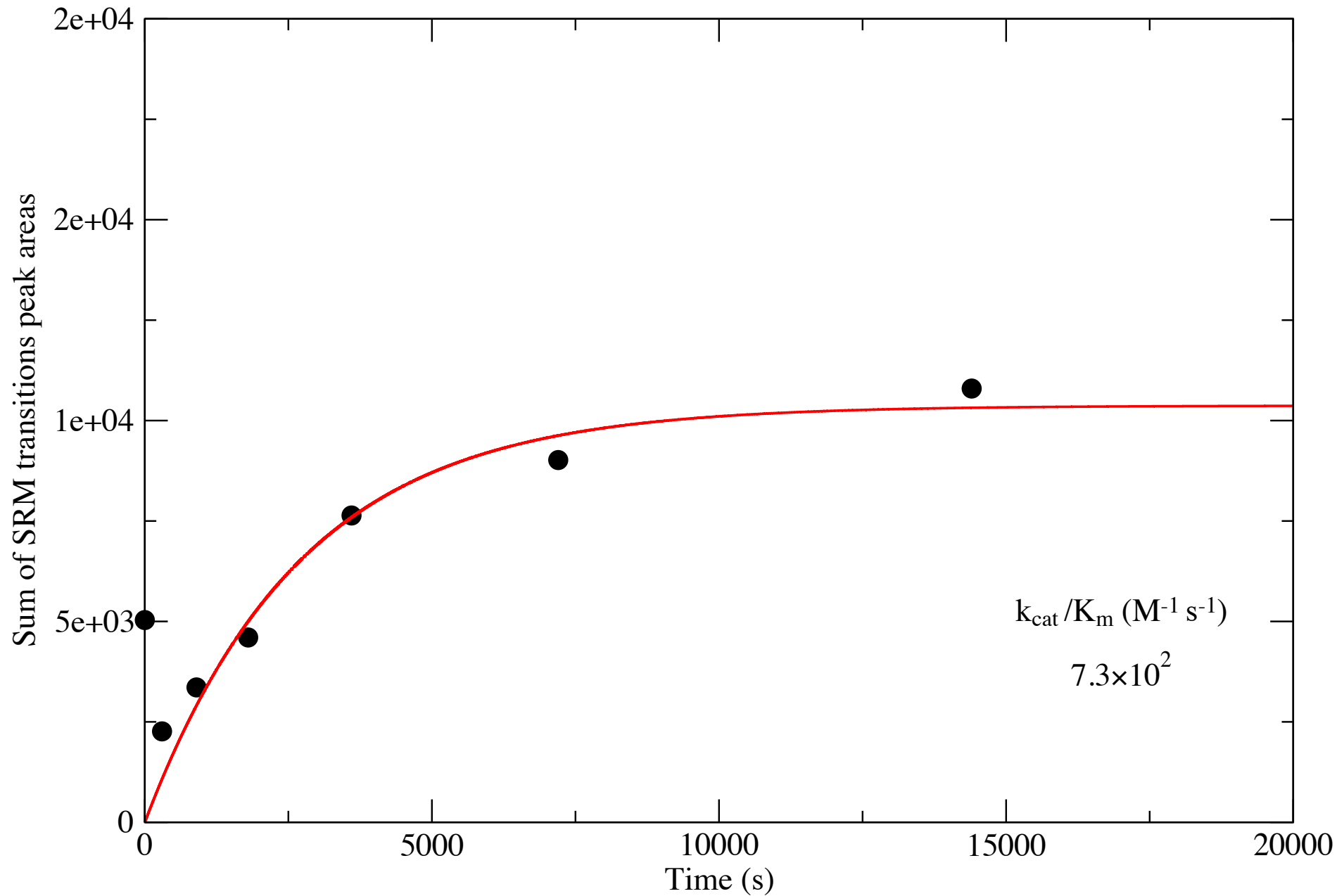
NP1L4_HUMAN (GVPSDSVEAAK)



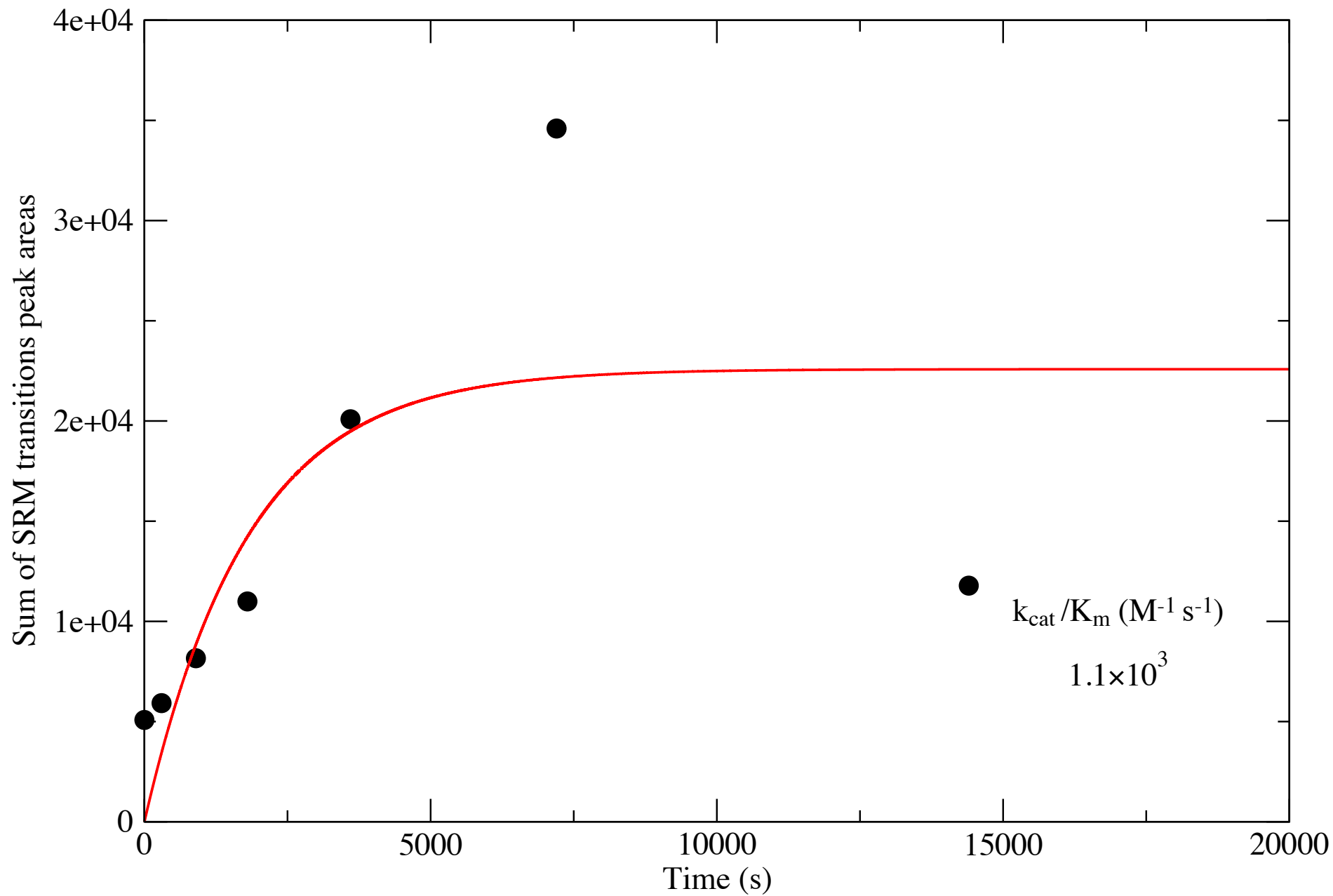
PARP1_HUMAN (GFSELR)



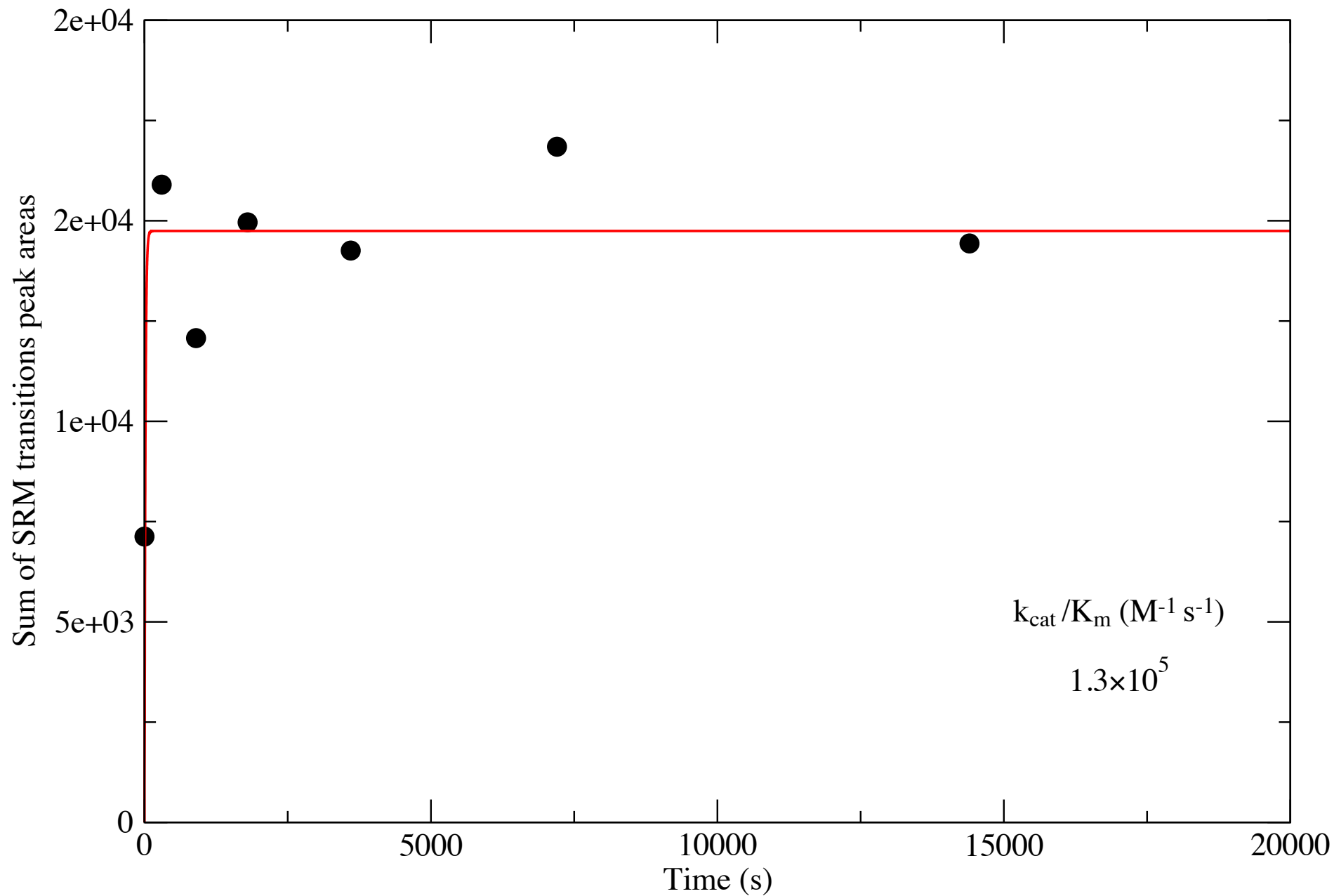
PRC2B_HUMAN (SVLPVPPIEFGVSPK)



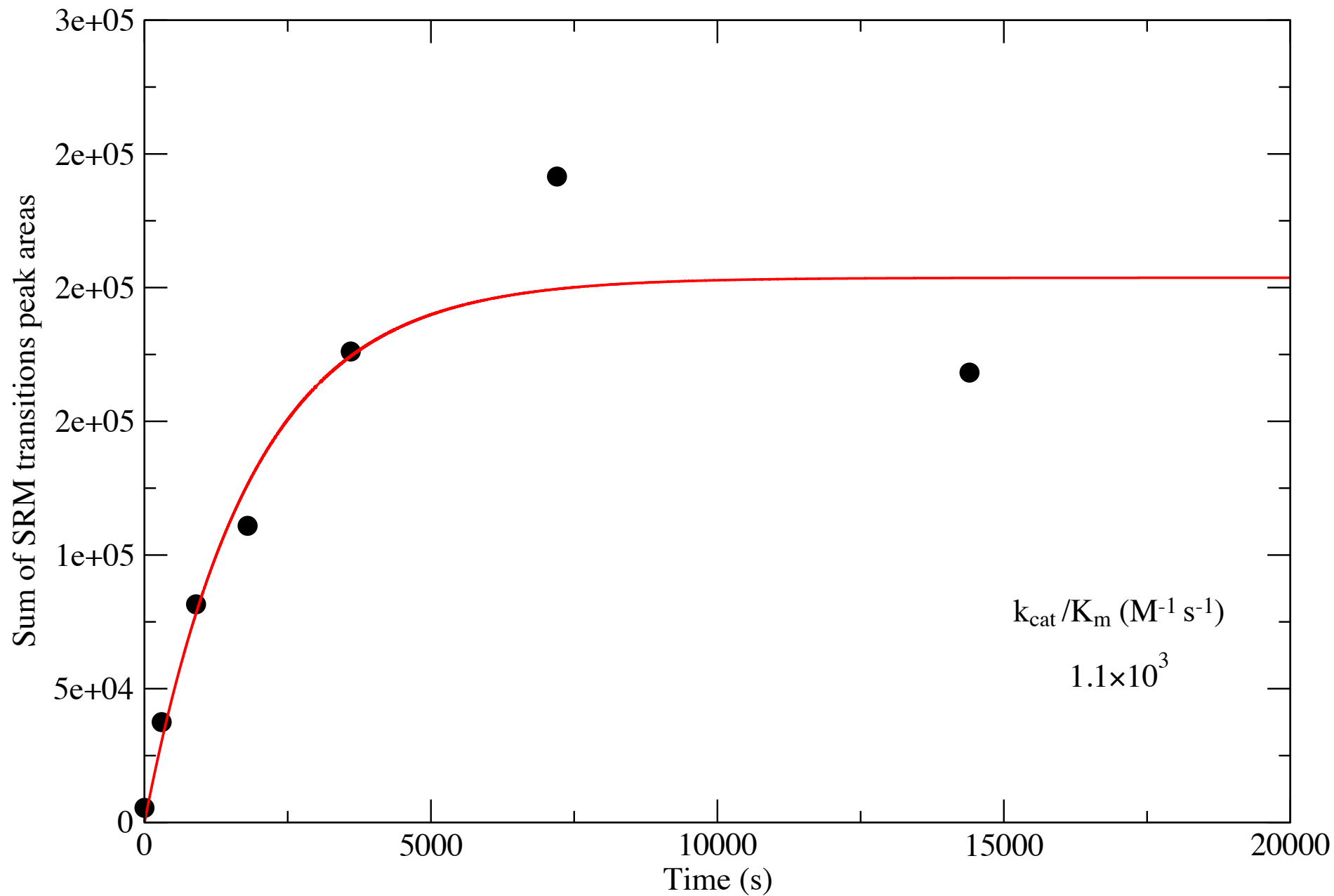
PREX1_HUMAN (SAFPLLSLGPR)



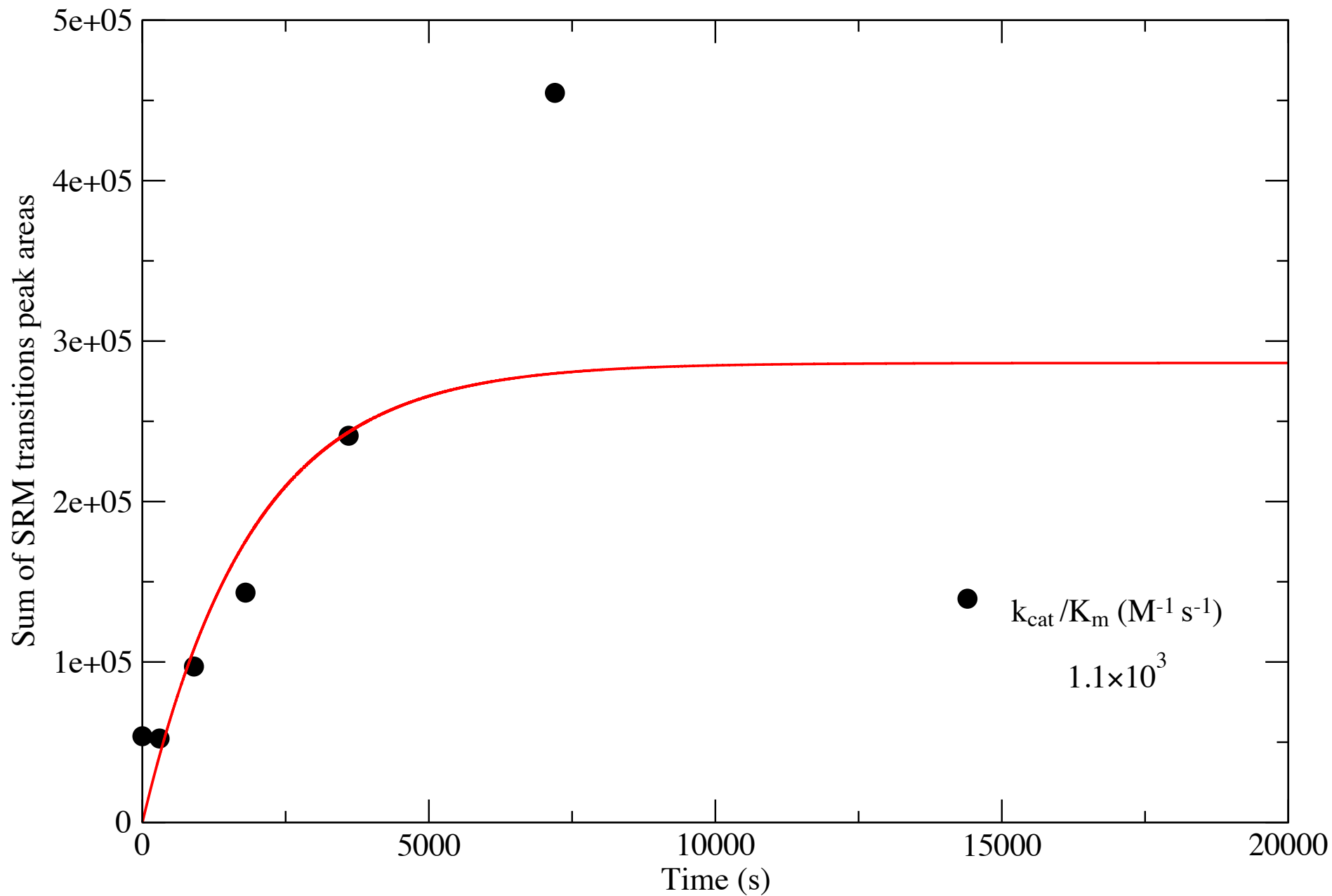
PTMA_HUMAN (AAVDTSSSEITTK)



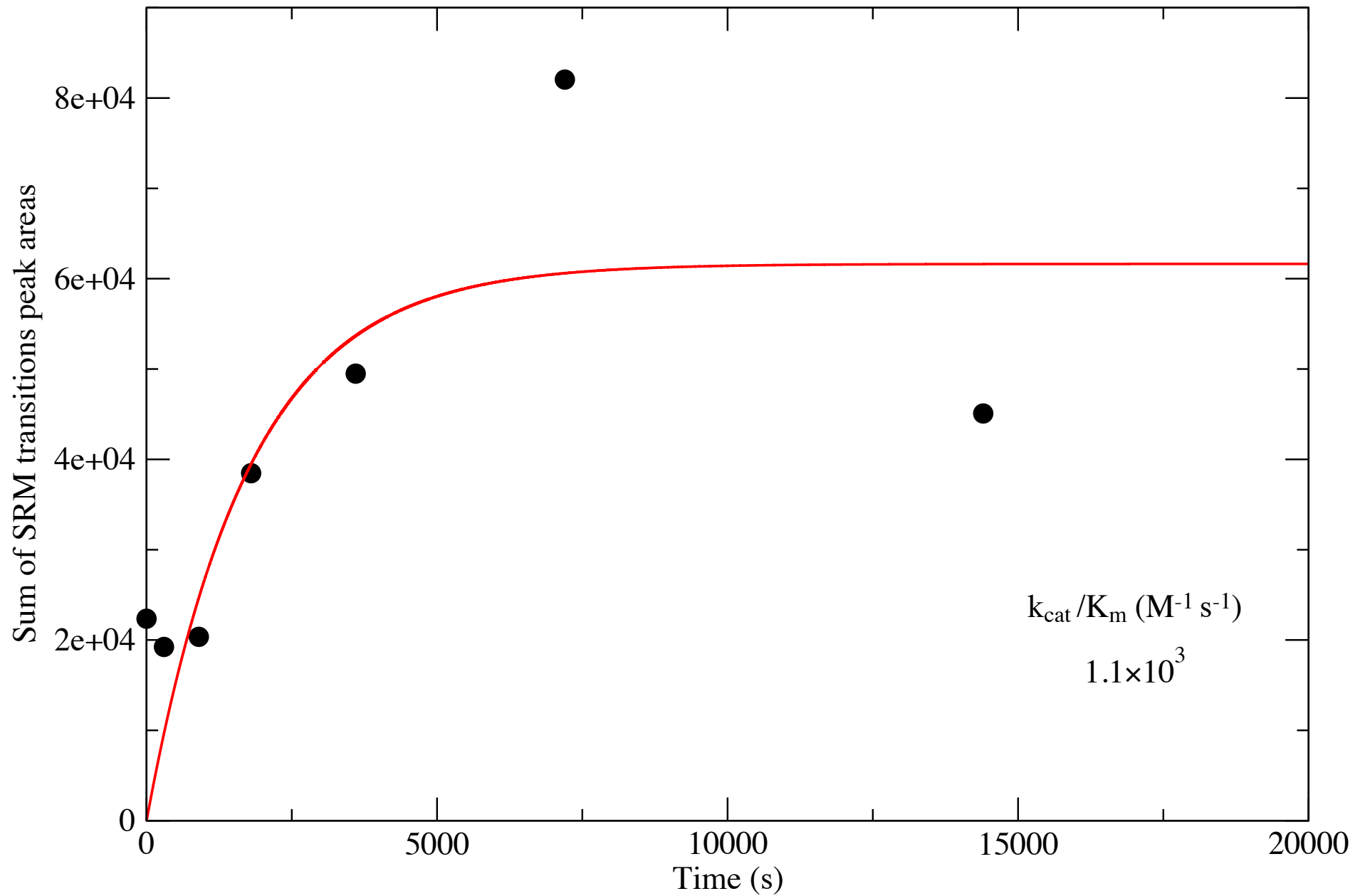
RGPA1_HUMAN (GLSLQFK)



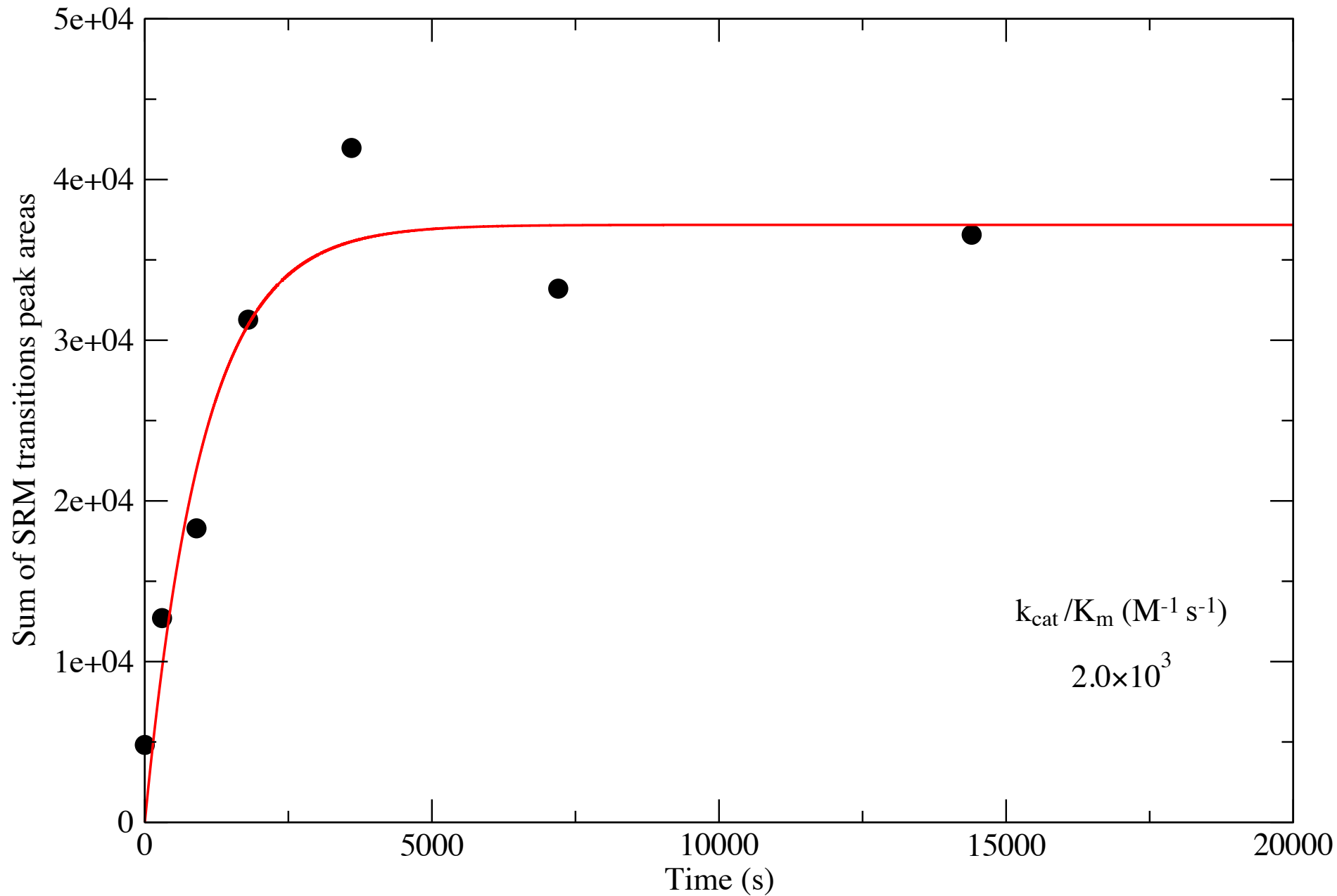
RSRC1_HUMAN (SFVQQTFR)



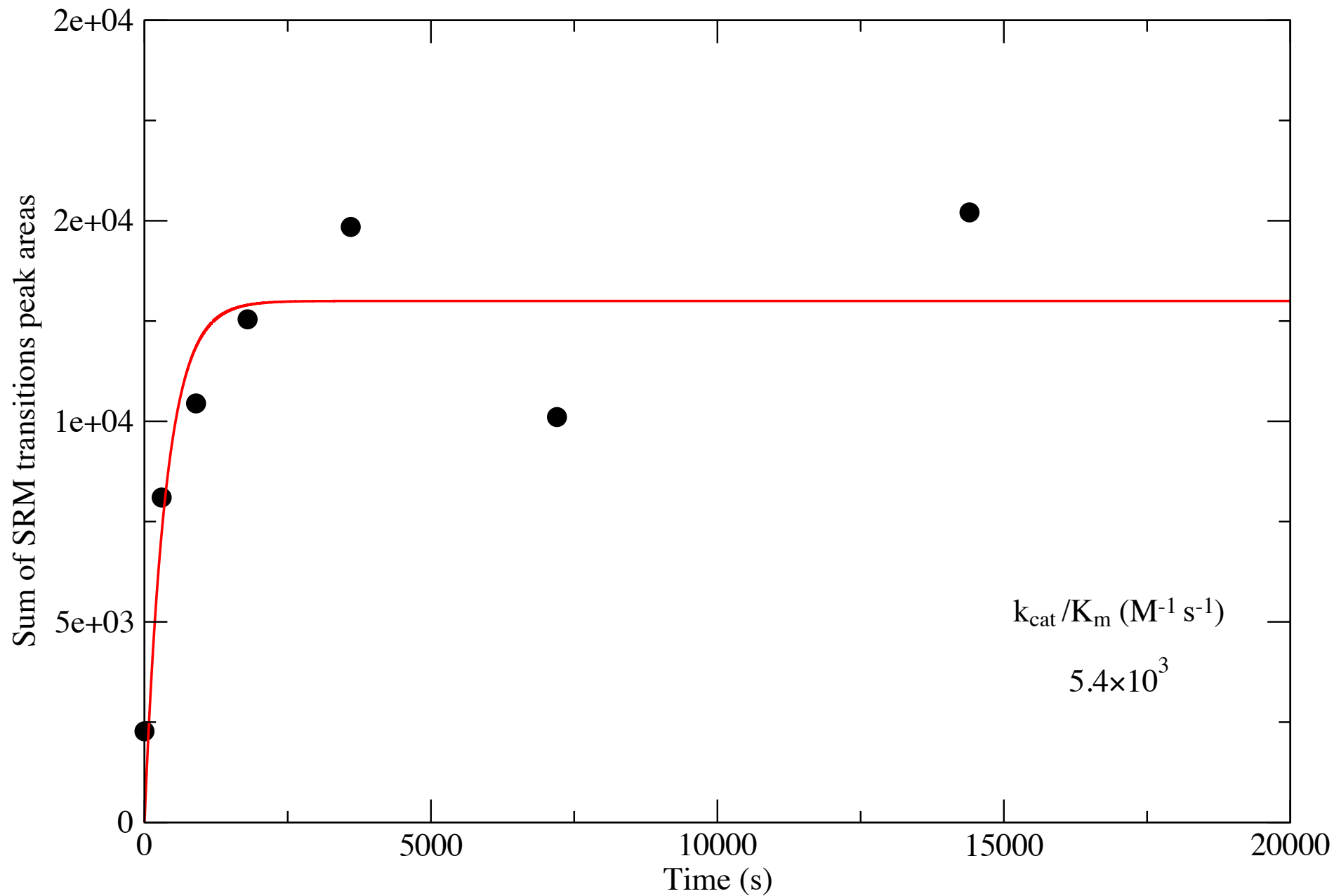
SAFB2_HUMAN (GLLDSFCDSK)



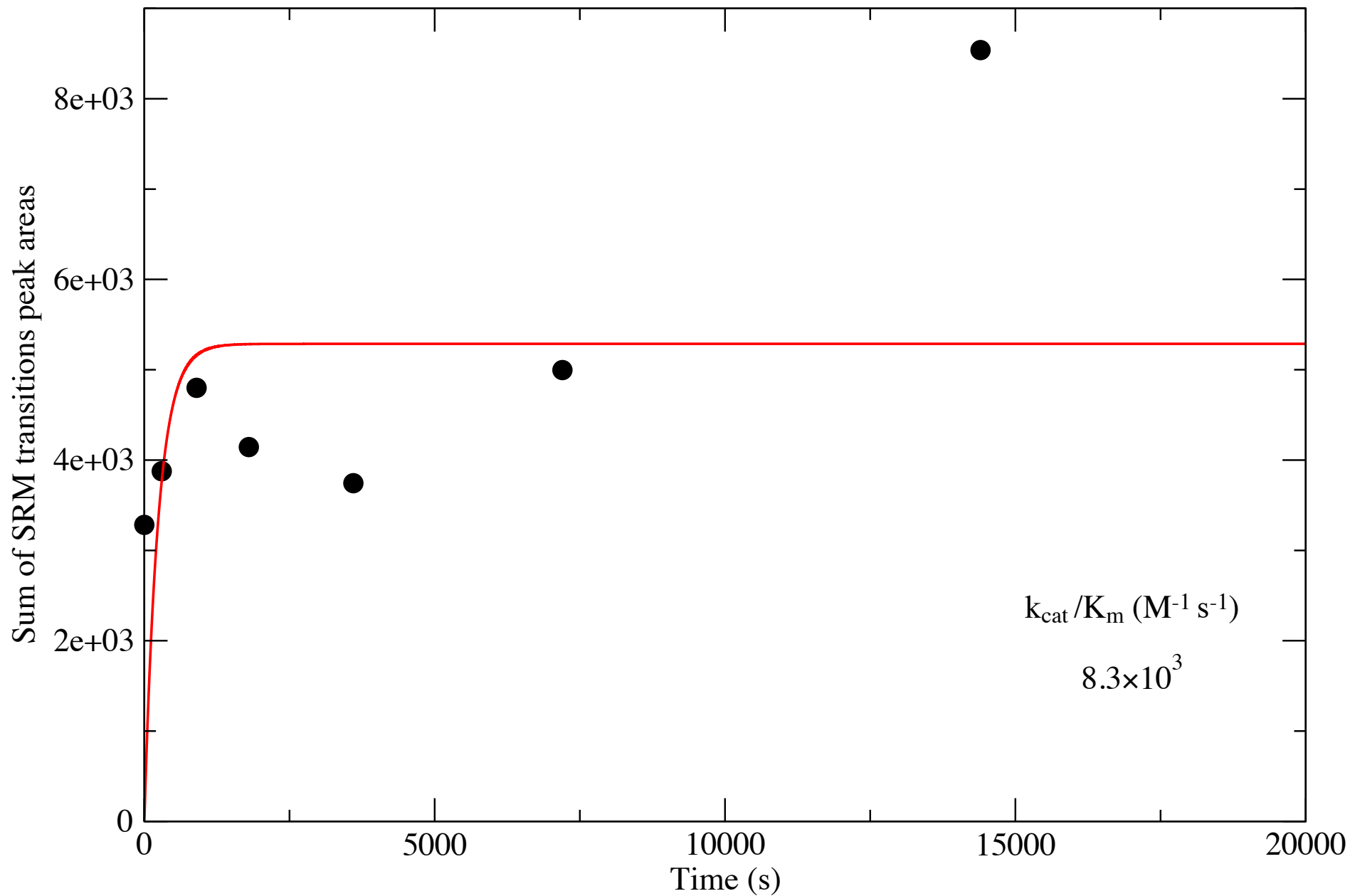
SAFB2_HUMAN (GWGGYGSDKR)



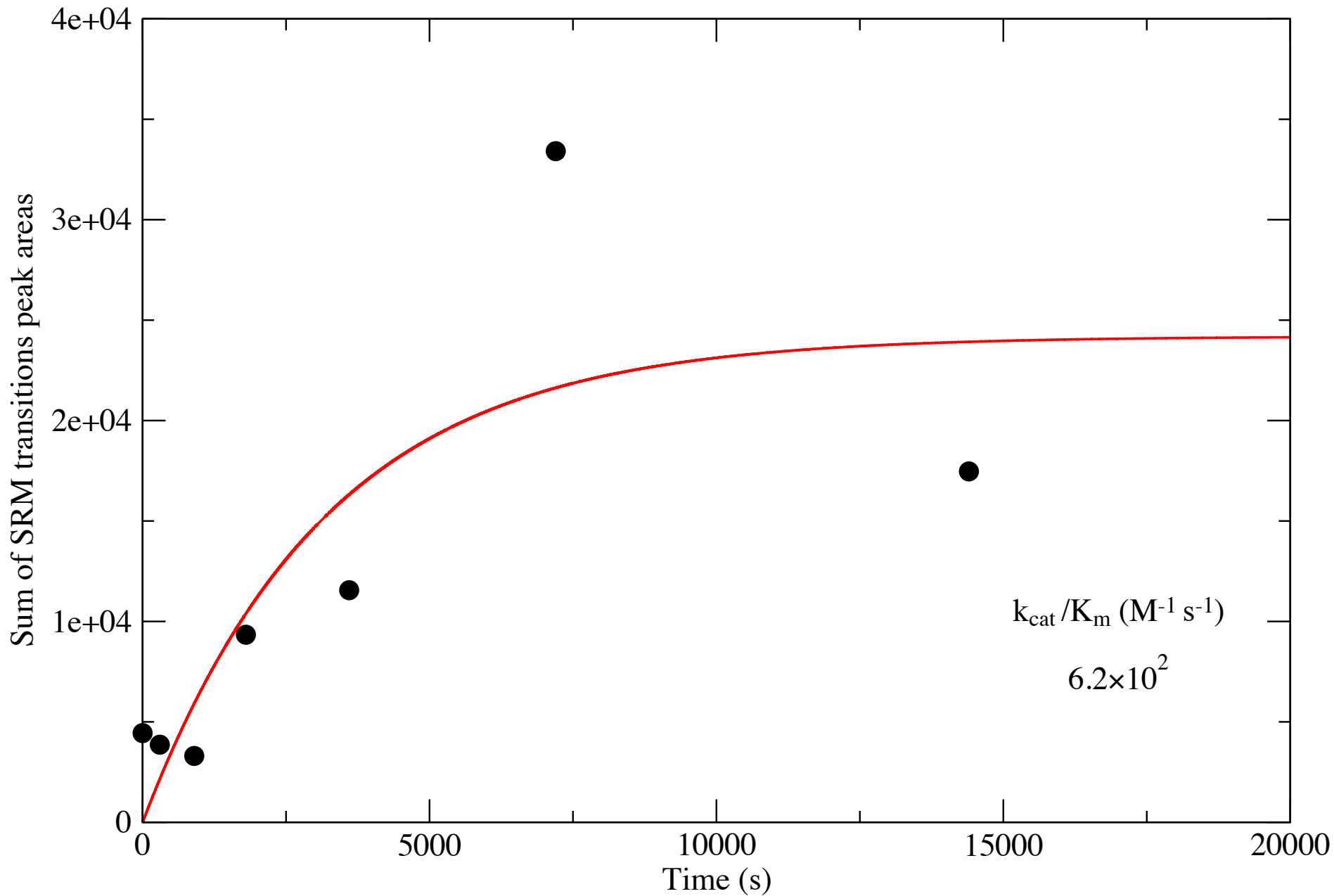
SC16A_HUMAN (SGPTQPPLSLSPAPETK)



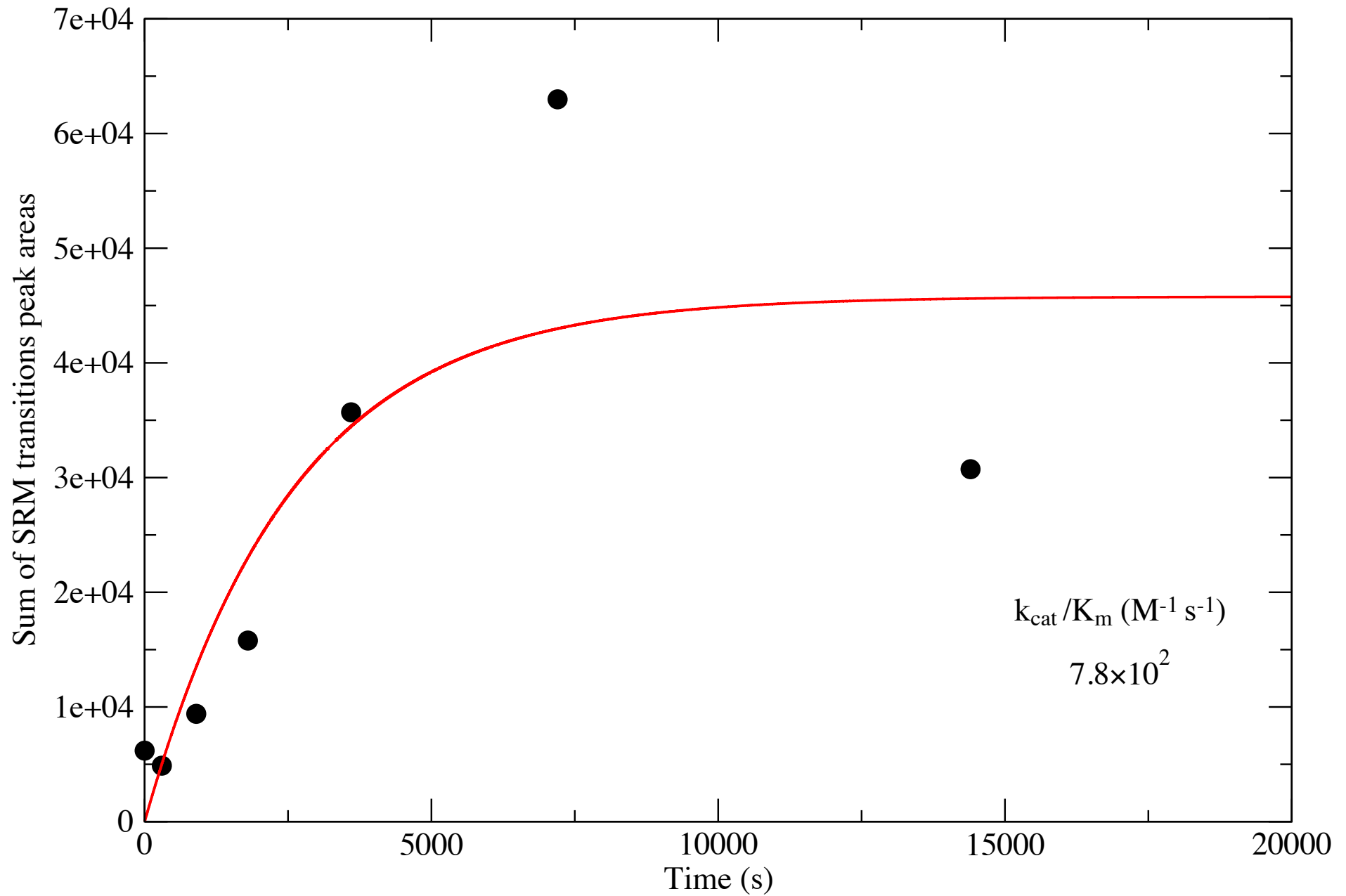
SC16A_HUMAN (SVSSSYSSR)



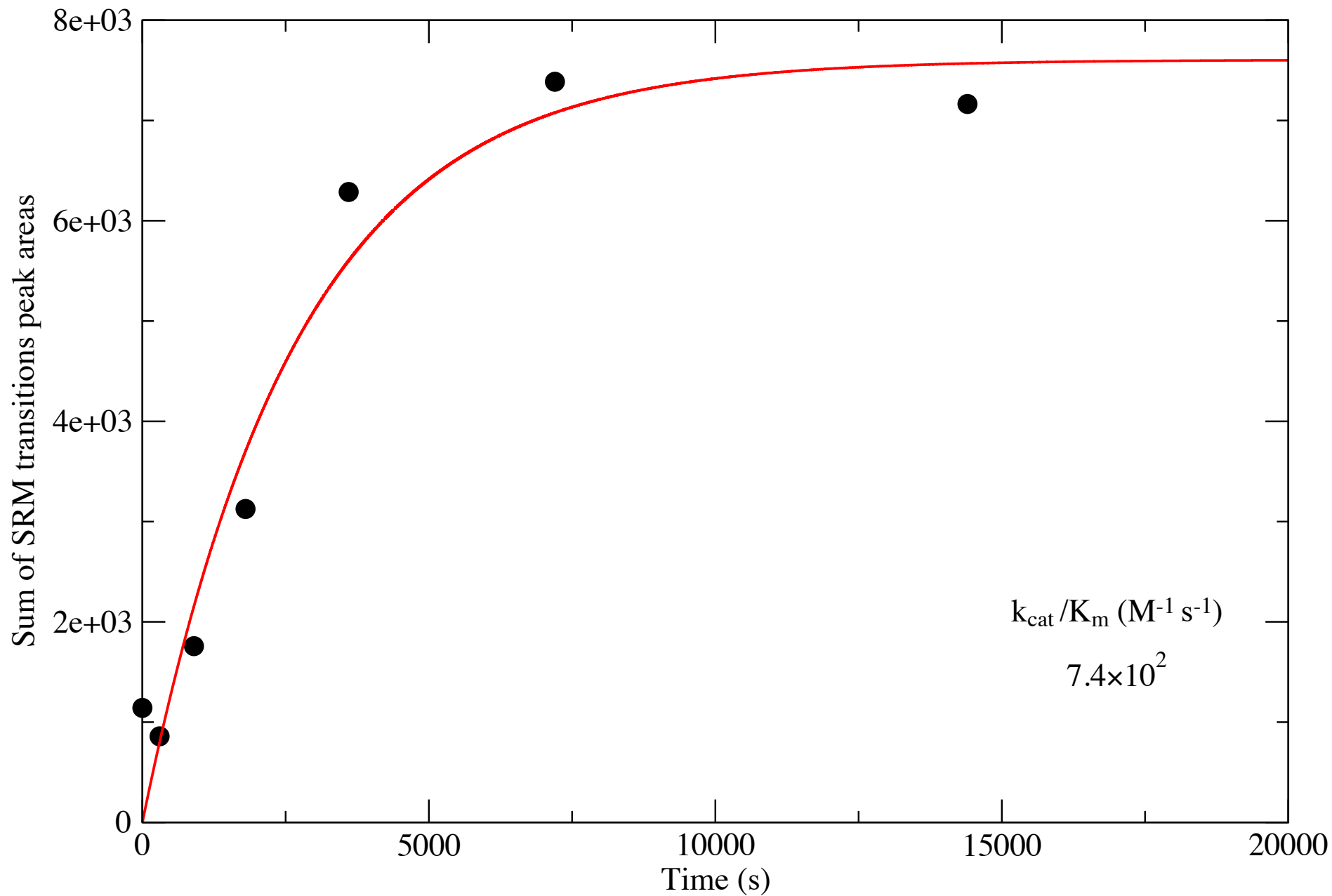
SR140_HUMAN (GVPIKSLDDDDLDGVPLDATEDSK)



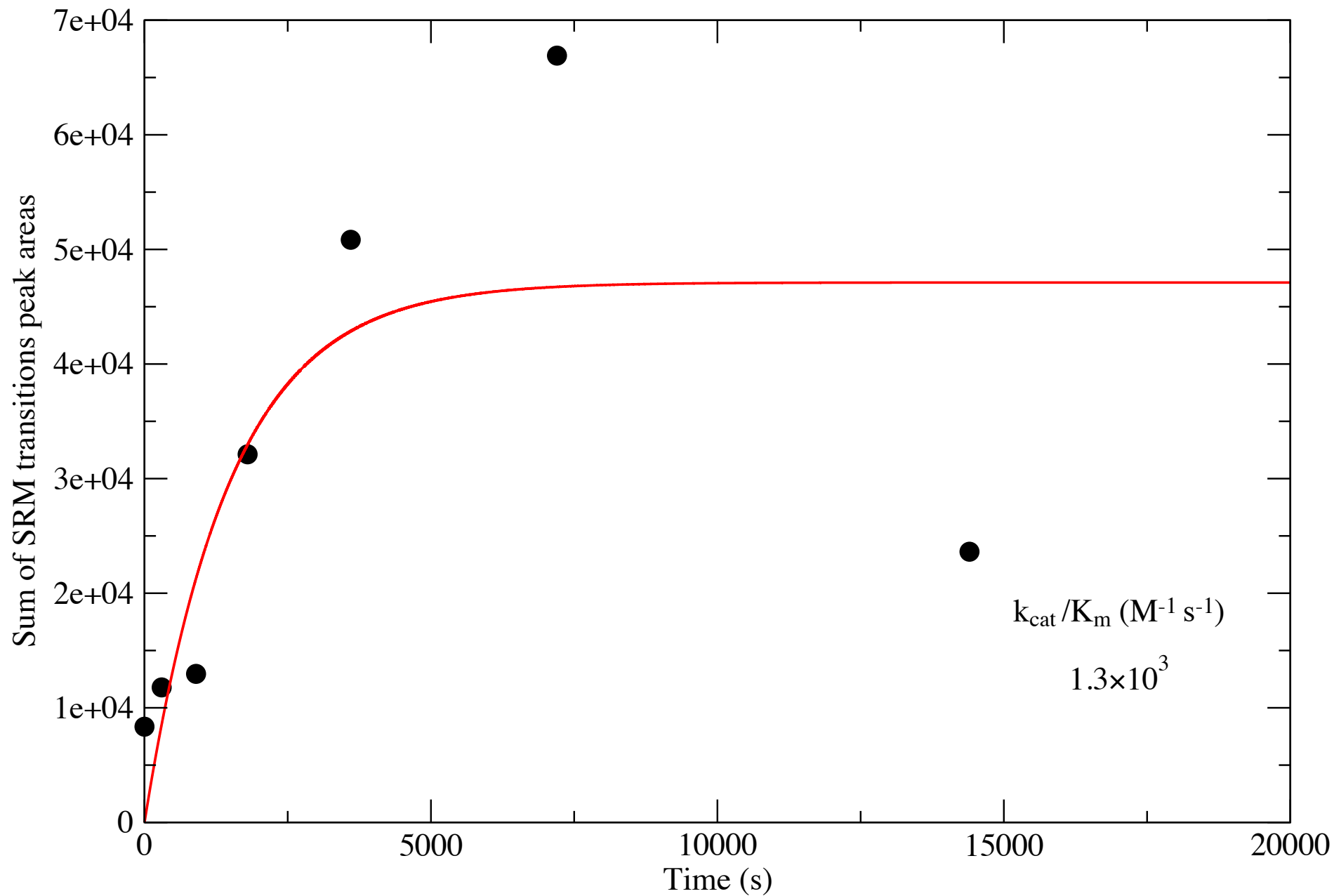
TCOF_HUMAN (GEVPTAVAPAQEK)



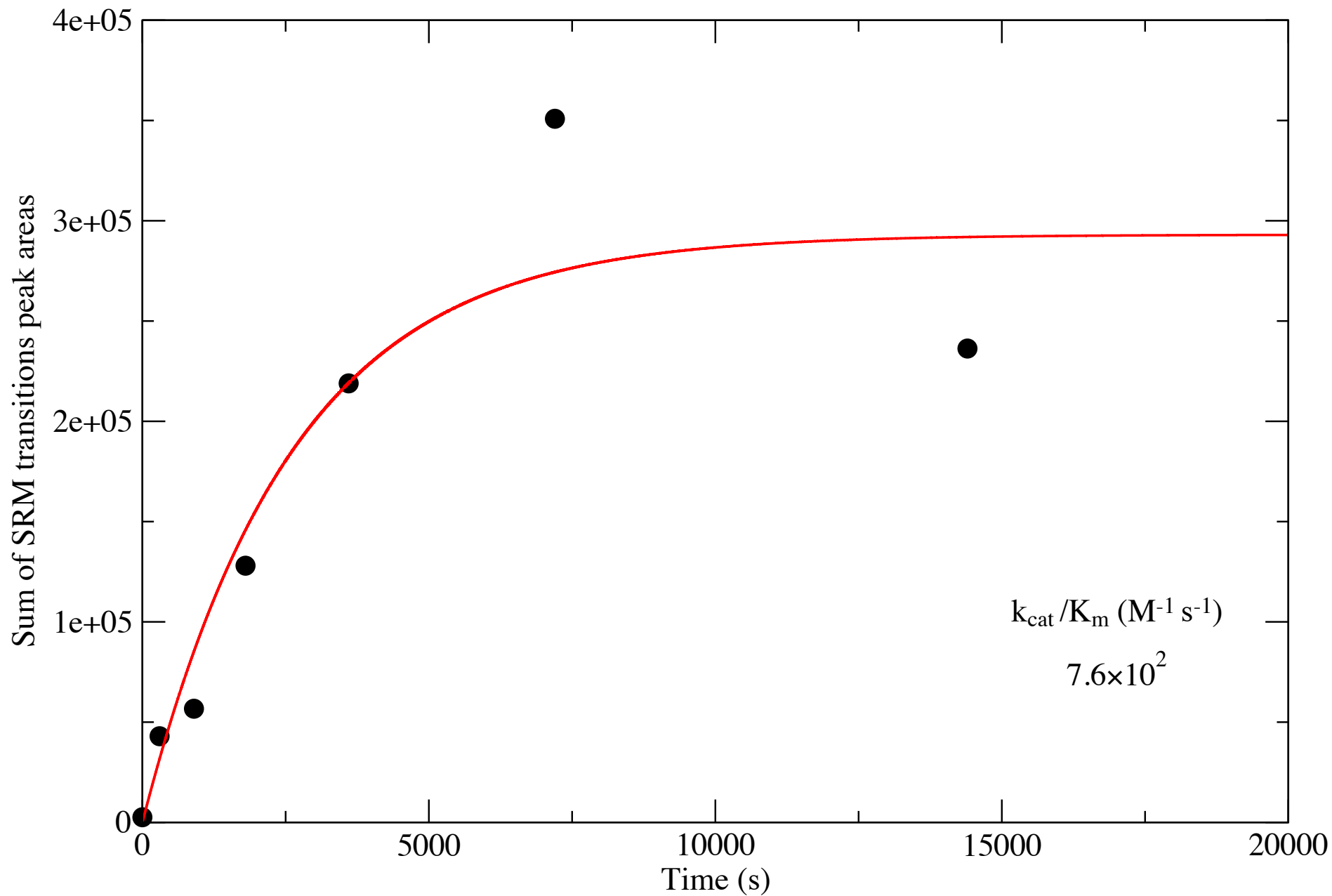
TIF1B_HUMAN (GADSTGVVAK)



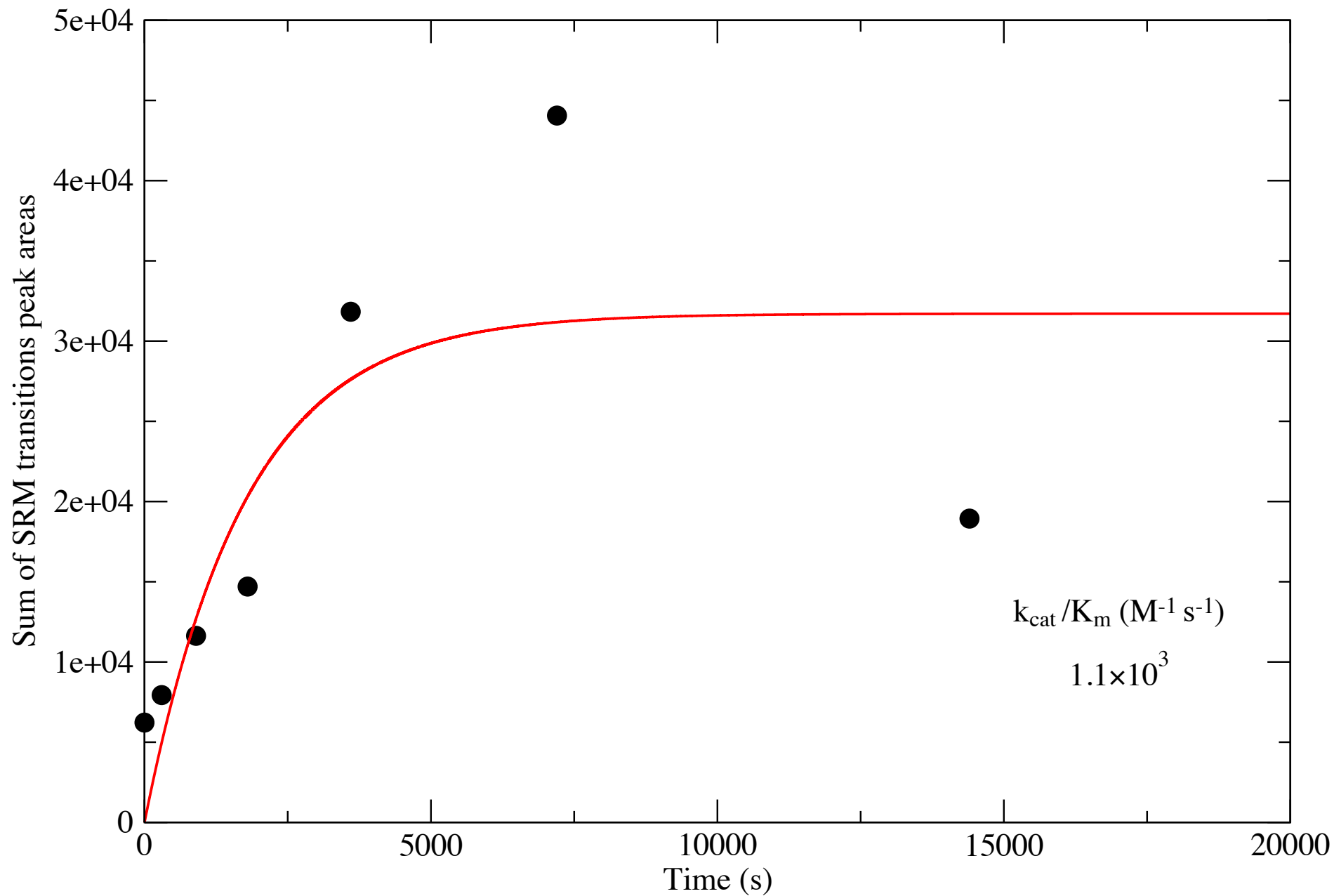
TREF1_HUMAN (GLGLPVGSK)



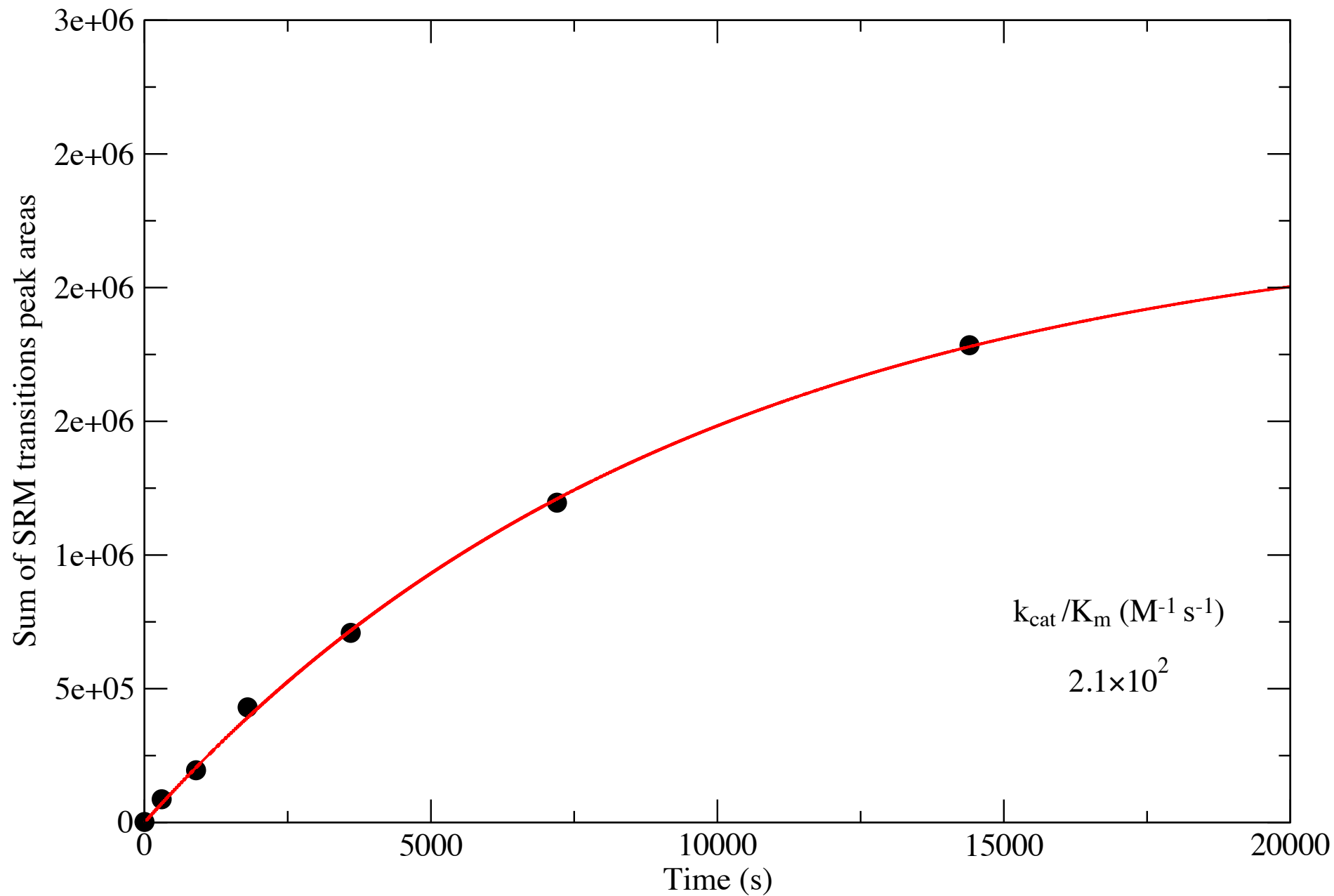
TYB4_HUMAN (MAEIEKFDK)



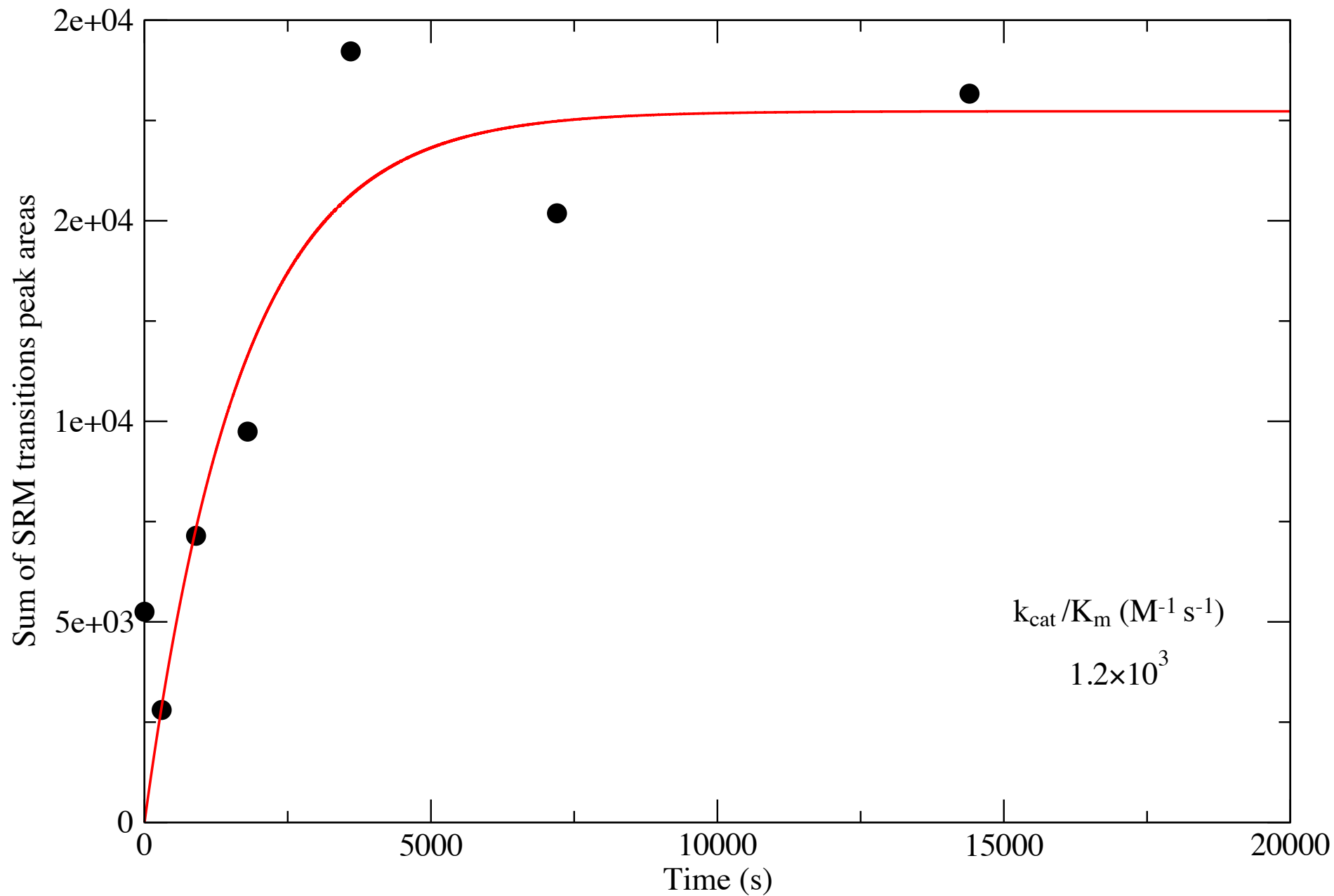
UBN1_HUMAN (SFIDNSEA YDELVPASLTK)



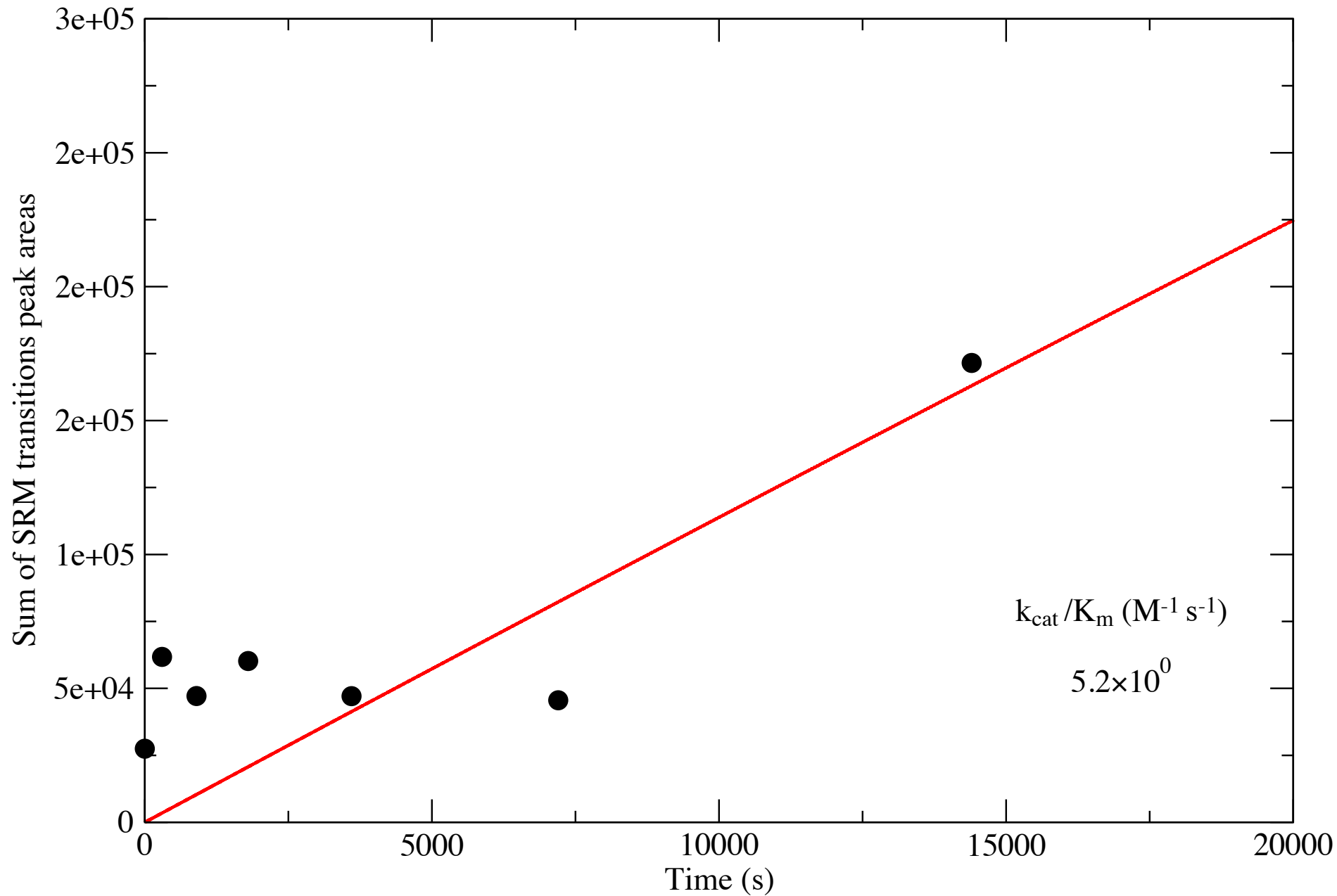
UBP10_HUMAN (GVSGGLGQR)



VCIP1_HUMAN (GCVADALGAAAFATR)



YBOX3_HUMAN (GVPVEGSR)



ZAP70_HUMAN (GYTPEPAR)

