

Supplementary Fig. 1 (Continued)

B

zCasp8	-----MDPQ-I FHEIDENLITSGDVDQLKFLCLDFIPKRRILESVTDAKDLIL	45
hCASP8	-----MDFSRNLYDIGEQLDSEDLASKFLSLDYIPQRKQEPIKDALMLFQ	46
mCasp8	-----MDFQSCLYAAEELGSEDLAALKFLCLDYIPHKKQETIEDAOKLFL	46
xCasp8	MSDLLSSTVDAMNLADMNKLIIFEISEDLDKTETIAMIFLCEKRVTAQEKENIKDAKTLFL	60
zCasp8	RIDEQGLLED--ELLFPPELLIAIGRIDLLEIILKKK-KEEVERNLLRCDSRKGVSAQRKM	102
hCASP8	RLOEKRMLEESNLNSFLKELLFRINRLDLLITYLNTRKEEMERELQTPGR--AQISAYRVM	104
mCasp8	RLREKGMLLEEGLNSFLKELLFHISRWDLLVNFLDCNREEMVRELDPDN--AQISPYRVM	104
xCasp8	CLKKKDILCYNDLSFLKELLYRIGRNDLLRGKLGVRTEEIKRTIEWS---PQISPYRIL	116
zCasp8	LKISEDMIEENFRAAKFLLD--LPRAKLGRSTSFLDALIEMEKQQRILGPDLNDELYRIL	160
hCASP8	LYQISEEVSRSELRSFKFLLOEEISKCKLDDDMNLIDIFIEMEKRVILGEGKLDILKRCV	164
mCasp8	LFKLTSEEVSEELRSFKFLLNNEIPKCKLEDDLSLLEIFVEMEKRTMLAENNLETLKSIC	164
xCasp8	LYDISQGLSKKEVEDILKYLLDLSTAKT---ENASILEIFILEEKVGKLEPDDLOKLIKHD	173
zCasp8	EKCD-KQLAVMIEFRNQSHRDQEQQGRPLPEEVF-----LNNPVSETMEERR	207
hCASP8	AQIN-KSLLKIIINDYEEFSKERSSSLEGSQSPDEFS-----NCEELCGVMTISDS	211
mCasp8	DQVN-KSLLGKIEDYERSSTERMSLEGREELPPSV-----LDEMSLKMAELCDS	213
xCasp8	ETIGCKNLSRNIEDYERISEADNHRPENLPDLFEKISVQEEQVNCTAQEPFNGEQRTQPQ	233
zCasp8	-RNSSAGAITTDATPPLNPNEYIILTQRPGLGYCLIINNYNFIKS-----TNLLKRTGT	260
hCASP8	PREQ-----DSES-QTLDKVYQMKSQPRGYCLIINNHNFPAKAREKVPKLHSIRDRNGT	263
mCasp8	PREQ-----DSES-RTSDKVYQMKNKPRGYCLIINNHDFSKAREDITQLRKMKDRKGT	265
xCasp8	ETESDYCQP--QQHS-HMNLEYHLEKNPHGWCVVINNYDFKEARSQDCKYT---DREGT	287
zCasp8	DMDKDRLAKLFSRMHFQIEVRNDLEAWAIKDEIKQFANRNHASMGAFVCCILSHGEKGTV	320
hCASP8	HIDAGALTITFEELHFEIKPHDDCTVEQIYEILKTYOLMDHSNMDCFICCIILSHGDKGII	323
mCasp8	DDKKEALSKTFKELHFEIVSYDDCTANEIHEILEGYQSADHKNKDCFICCIILSHGDKGVV	325
xCasp8	AKDAEETIRIFNARGYITEEHDLTAANIQKTLEMYSKKDHAEKDSFVCFILSHGGTV	347
zCasp8	I GTDGKEVEIREVTLPFAG--CRTLASKPKLFFIQACQGDENOAGVWTSDGREDAPEEDE	378
hCASP8	YGTDGQEPPPIYELTSQFTGLKCPSSLAGKPKVFFIQACQGDNYQKGIPVETDSE---EQP	379
mCasp8	YGTDGKEASIYDLTSYFTGSKCPSSLGKPKIFFIQACQGSNFQKGVPDEAGFE---QQNH	382
xCasp8	CGCDGEEVEIKRLTKYFNGQHCRSLINKPKIFFIQACQGKESHPK--VDMMDMD---TS	400
zCasp8	KYEEDAGIIVLRKIPLEADFIIGMATVEEYL-SYRHTKKGSILIQELCKKMEELCPKKEDM	438
hCASP8	YLEMDLSSPOTRYIPDEADFLLGMMATVNNCVSYRNPAGEGTWYIQSLCQSLRERCPRGDDI	439
mCasp8	TLEVDSSS-HKNYIPDEADFLLGMMATVKNCVSYRDPVNGTWYIQSLCQSLRERCPQGDDI	441
xCasp8	FYEFDANG---SHLPTEADFIATAFATVEDYTSIRHRENGSIYIQLCKALTTYN--QDL	455
zCasp8	LSILTKVNFEVSKRILKGY---KQMPEPRYTLTKKLVLPI	476
hCASP8	LSILTEVNYEVSNKDDK--KNGKQMPQPTFTLRKKLVFP	479
mCasp8	LSILTGVNVDVSNKDDR--RNKGKQMPQPTFTLRKKLFFPP-	480
xCasp8	IDILTSVNSDVAJMLFRLWRKNVTQMPSFKSELKKLTLPPPNA	500