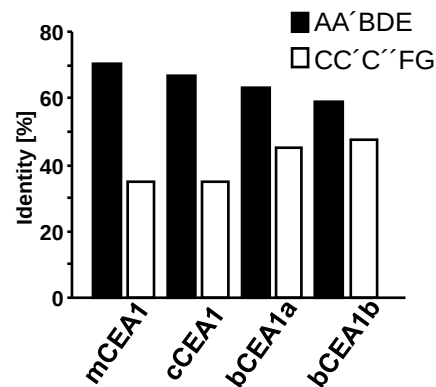


A

	<u>A</u>	<u>A'</u>	<u>B</u>	<u>C</u>	<u>C'</u>
hCEA1	QLTTESMPFNVAEGKEVLLL	VHNL	PQQLFGYSWYKGERVDG	NRQIVGYAIG	
mCEA1	EV...AV.PQ...DNN.....		LA.GAFA...NTTAIDKE	.ARFVPN	
cCEA1	.V.V...V.P.A...DA..R.L...		GDTASLT.F...T.LP	THK.LL.VID	
bCEA1a	..I.TV.PLA...SD...A..VT		KNPL..A..R....NSQL	.AS.---	
bCEA1b	...I.TV.PLA...SD...AQ.V		TKSPL.....R....NTQL	.AS.RVD	

	<u>C''</u>	<u>D</u>	<u>E</u>	<u>F</u>	<u>G</u>
hCEA1	TQQATPGPANSGRETIYP	NASLLIQ	NVTQ	NDTG	FYTLQVIKSDLVNEEATGQFHVY
mCEA1	SNMNFT.Q.Y....I..S.G...		F.MI.MK.M.V...DMTDEN	YRRTQ..VR...H	
cCEA1	.KIT.....Y.....G...F..		I.L...S.I..I.NQKFEALVRG	QLQVF	
bCEA1a	RVAT.K...H....L...GT.....		K...S...L.T.D..QT.RQ..HL	.IL	
bCEA1b	.NAT.K...H....L...GT....S...		K...S.M.L.T.D..QT.RP..HL..L		

B**C**