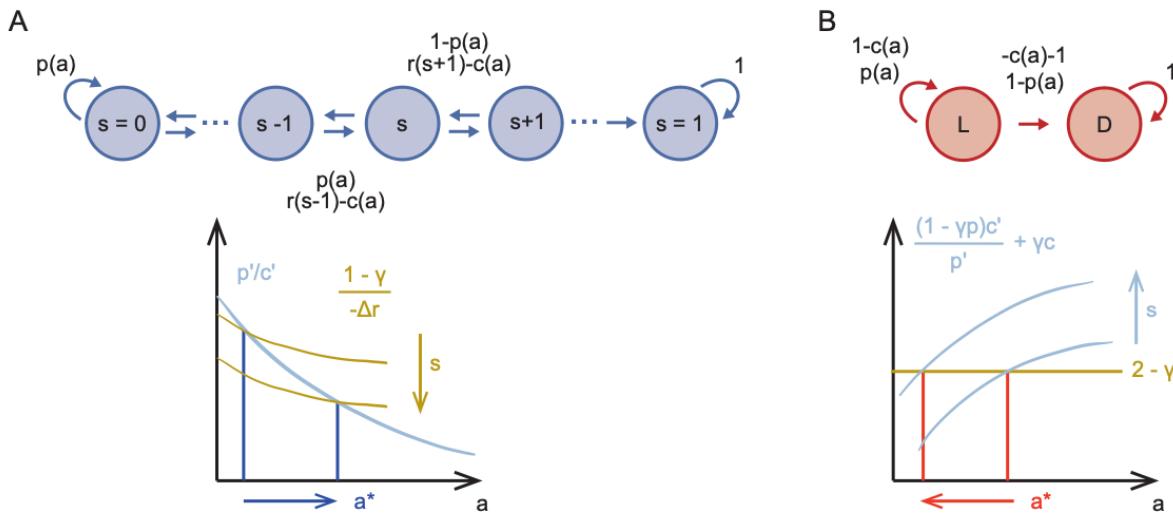


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**Figure 2—figure supplement 1. Sketch of Markov Decision Processes model and predictions for stinging.**

- 865 A) Directed graph representing the Markov Decision Process for predatory stinging (top) including states of  
866 starvation  $s$ , actions  $a$ , and transitions to adjacent states depending on the probability to catch prey  $p(a)$ .  
867 Graphical representation of the result that optimal predatory stinging increases with starvation (bottom).
- 868 B) Directed graph representing the Markov Decision Process for defensive stinging (top) including states of  
869 safety and danger  $L$  and  $D$ , actions  $a$ , and transitions between  $L$  and  $D$  depending on the probability to  
870 successfully stinging the predator  $p(a)$ . Graphical representation of the result that the optimal defensive  
871 stinging decreases with starvation (bottom).
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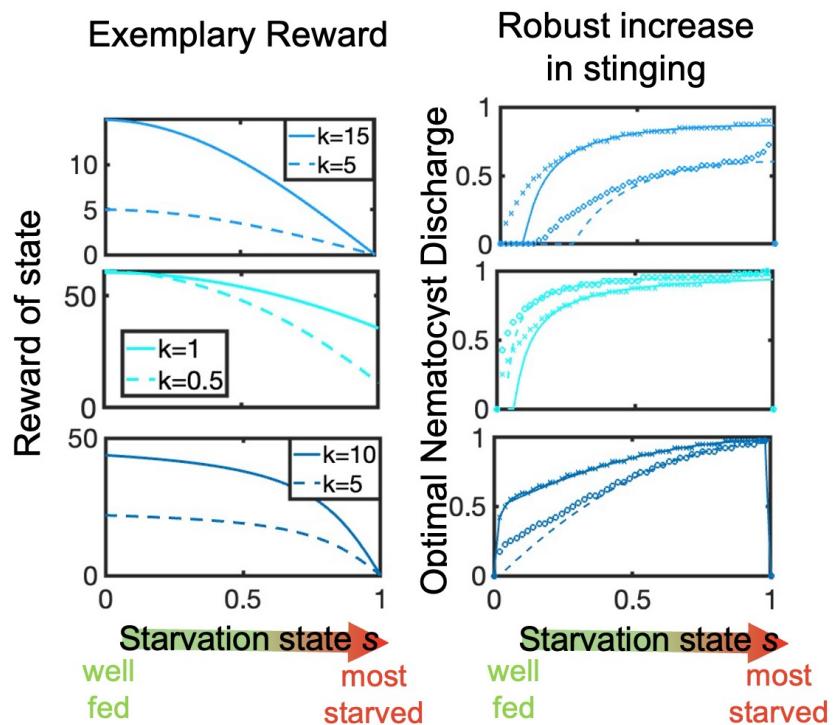
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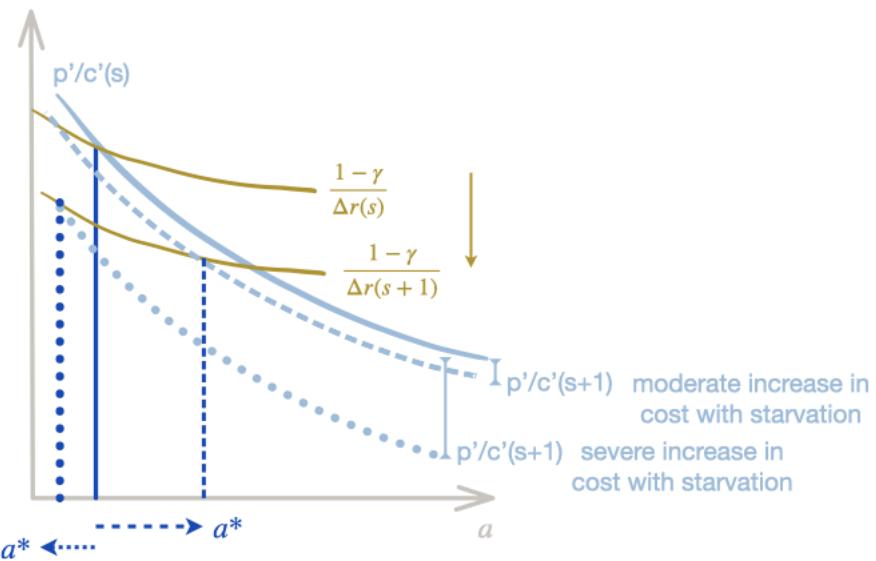
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881 **Figure 2—figure supplement 2. Optimal policy predicted by Bellman's theory for the MDP sketched in Figure**  
882 **2—figure supplement 1A.**

883 *Left:* three choices of concave reward functions  $r(s')$ :  $r(s) = k \cos(s\pi/2)$ , upper left;  $r = k(1 - 50s^2) + 60$ ,  
884 middle left;  $r = k \tan^{-1}(5(1 - s)/(\pi/10))$ , lower left. Solid and dashed lines correspond to two choices of the  
885 parameter  $k$  for each reward as in the legend. The cost of full discharge is constant  $c_0 = 1.5$  and the likelihood of  
886 successful discharge is  $p = p_M a(2 - a)$  with  $p_M = 0.6$ .

887 *Right:* the asymptotic solution for the optimal policy  $a^*(s)$  (solid and dashed lines matching the corresponding reward  
888 on the left) reproduces well the numerical solution obtained from solving Bellman's Equation (1) with the value  
889 iteration algorithm (crosses and circles correspond to the solid and dashed rewards on the left). Optimal nematocyst  
890 discharge increases with the starvation state, independently on the shape of the reward function.



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900 **Figure 2—figure supplement 3. Sketch of theoretical prediction for predatory stinging with increasing cost.**  
901 Similar to **Figure 2—figure supplement 1A bottom**, for the case where the cost per nematocyte varies with  
902 starvation  $c = c_0(s)a$ . Moderate increase in the cost per starvation (dashed light-blue line) do not affect the  
903 qualitative results as the green curve still intersects the light-blue curve for increasing values of  $a^*$  (marked by dashed  
904 dark-blue line). More dramatic increases of cost with starvation (light-blue dotted line) do lead to a decrease in  
905 predatory stinging with starvation as the intercept now moves backward with increasing  $s$  (marked by dark-blue  
906 dotted line).

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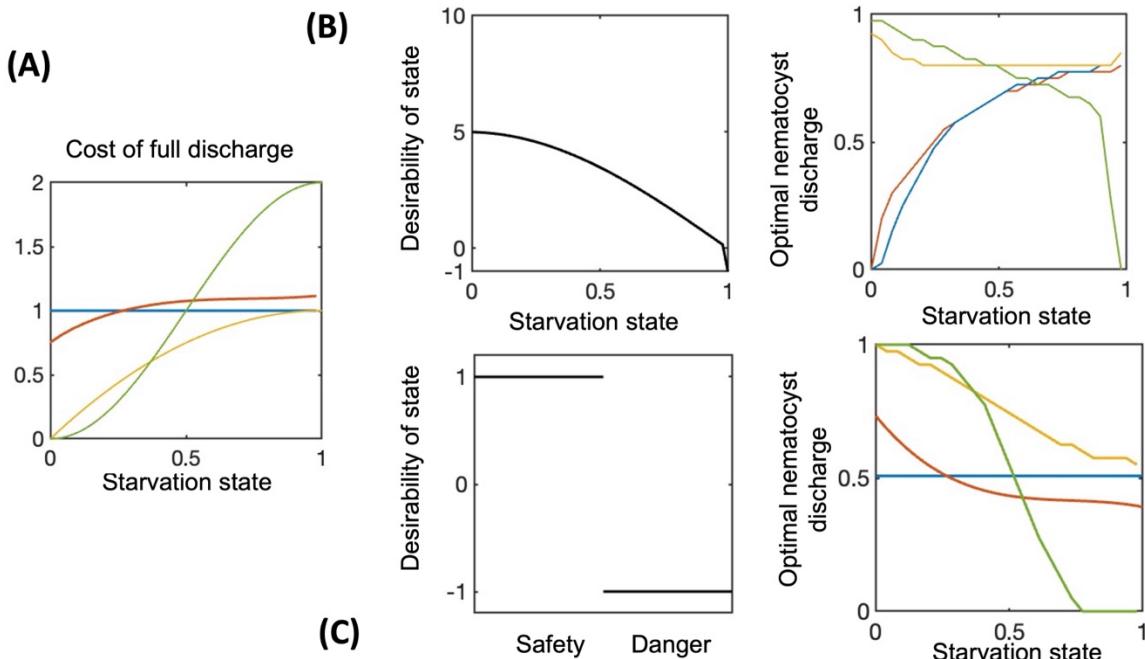
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**Figure 2—figure supplement 4. Effects of a moderately vs dramatically increasing cost with starvation.**

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For a constant cost of full discharge or moderately increasing cost with starvation, predatory stinging always increases, whereas defensive stinging decreases or stays constant (results discussed in main text, **Figure 2**, and reproduced here for comparison, red and blue curves in Panels A-C. For predation, we use desirability 2 from **Figure 2B**). When the cost function increases dramatically with starvation (panel A, yellow and green lines), defensive stinging keeps decreasing with starvation (panel C, right), but now also predatory stinging decreases with starvation (panel B, right, yellow and green lines). Results are obtained with numerical simulations.

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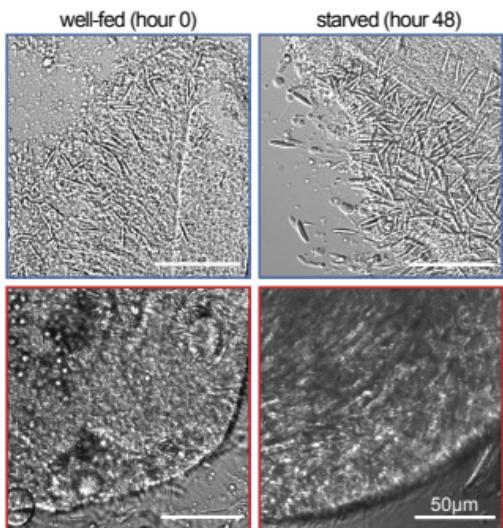
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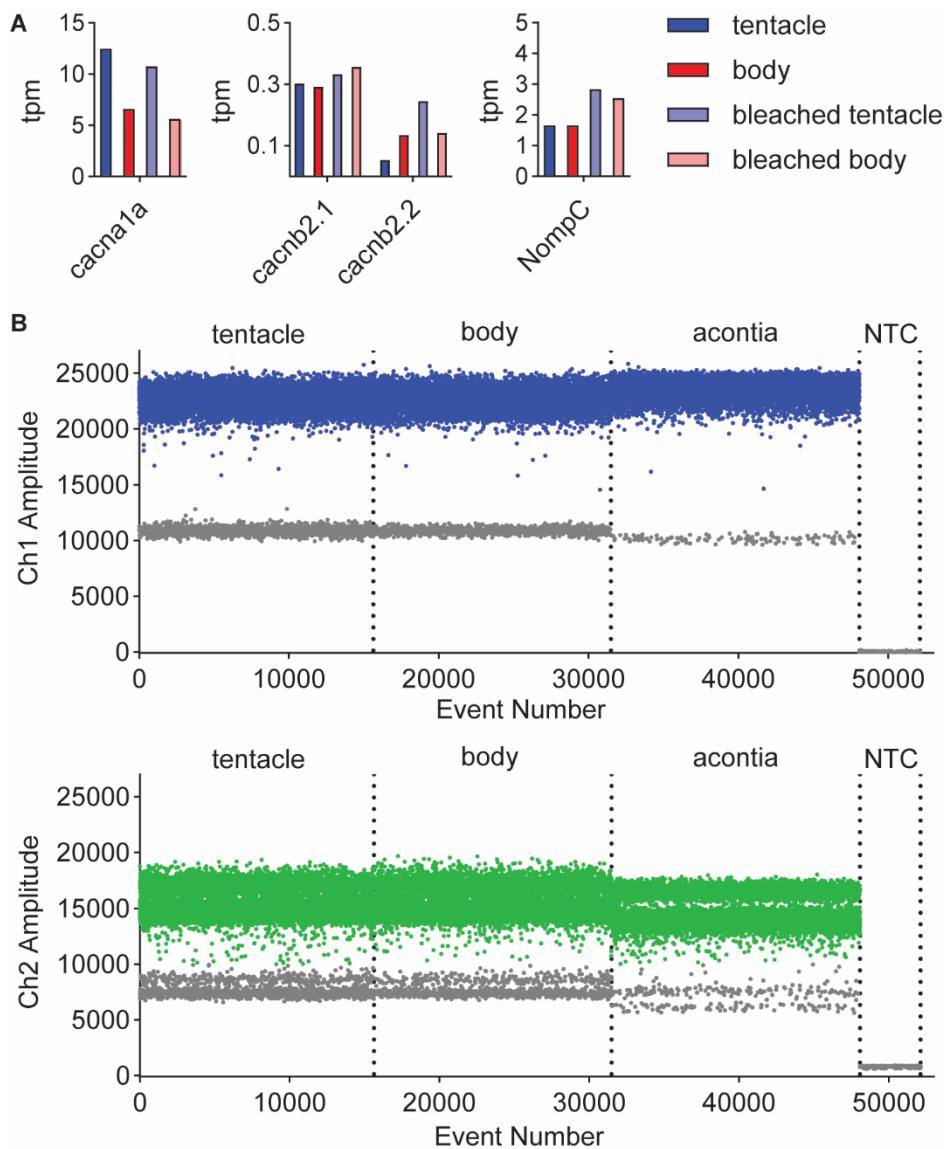
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929 **Figure 2—figure supplement 5. Modulation of *Nematostella* and *Exaiptasia* stinging is not due to changes in  
930 the abundance of nematocytes.**

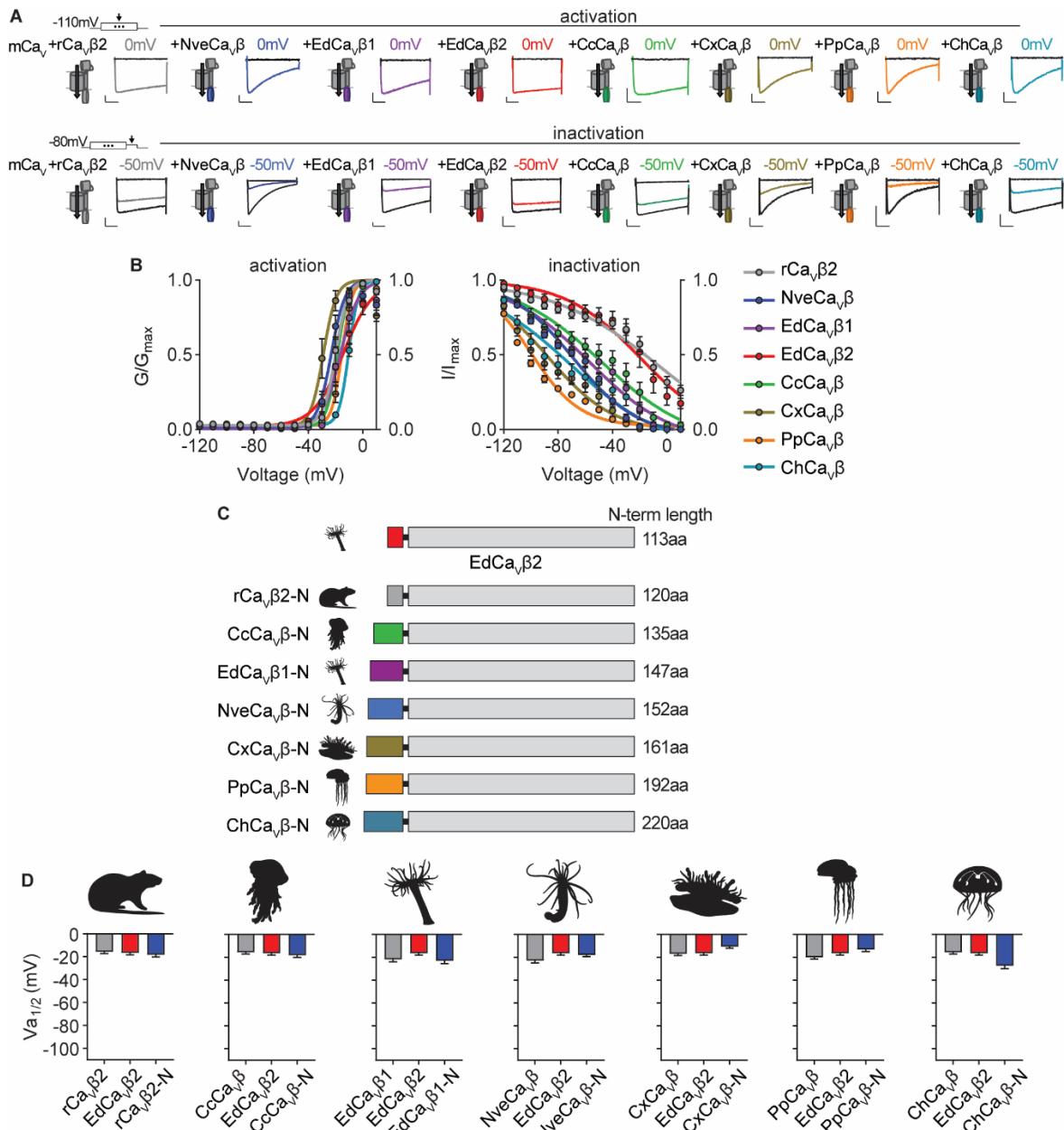
931 Nematocytes were highly abundant in tentacles from *Nematostella* (top) and *Exaiptasia* (bottom) before and after  
932 starvation. Representative of n = 3 animals. Scale bar = 50µm.  
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976 **Figure 4—figure supplement 1. Transcriptomic and molecular analyses of *Exaiptasia*  $\beta$  subunit isoforms.**

- 977 A) mRNA expression (transcripts per million, TPM) of voltage-gated calcium ( $\text{Ca}_V$ ) channel  $\alpha$  and  $\beta$  subunits  
978 in *Exaiptasia* tentacle (nematocyte abundant, blue), body (nematocyte non-abundant, red), bleached (minimal  
979 symbionts) tentacle (light blue), bleached body (light red) tissues. The  $\text{Ca}_V \alpha$  subunit was identified by  
980 homology to the sequence of the cnidarian  $\text{Ca}_V 2.1$  homolog found enriched in *Nematostella* nematocyte-rich  
981 tissues (Weir et al., 2020). NompC, the putative mechanoreceptor in *Nematostella* nematocytes (Schüler et  
982 al., 2015; Weir et al., 2020), was also detected in *Exaiptasia* tentacles.  
983 B) Representative plots of fluorescent amplitude across event number (droplet events) from amplification of  
984 unique regions of EdCav $\beta$ 1 (Ch1, Top) and EdCav $\beta$ 2 (Ch2, Bottom) sequences using droplet digital PCR  
985 (ddPCR, Bio-Rad Laboratories). Individual lanes correspond to tentacle RNA, body RNA, acontia RNA, and  
986 no template control (NTC). Blue and green points indicate positive PCR droplets after thresholding and gray  
987 points indicate negative droplets.  
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1015 **Figure 5—figure supplement 1. Voltage-dependent activation of Cav channels is conserved across cnidarian β  
1016 subunits.**

1017 A) Top: Voltage-gated currents from heterologously-expressed chimeric Cavs with the indicated β subunits  
1018 elicited by voltage pulses to -120mV (no current, black) and 0mV (colored). Abbreviations of species: Nve,  
1019 *Nematostella vectensis*; Ed, *Exaiptasia diaphana*; Cc, *Cyanea capillata* (jellyfish); Pp, *Physalia physalis*  
1020 (siphonophore); Ch, *Clytia hemisphaerica* (jellyfish); Cx, *Cassiopea xamachana* (jellyfish); r, *Rattus*  
1021 *norvegicus*. Bottom: Voltage-gated currents elicited by a maximally activating voltage pulse following 1 s  
1022 pre-pulses to -110 mV (max current, black), -50 mV (colored), or 20 mV (inactivated, no current, black).  
1023 Scalebars = 100pA, 50ms.

1024 B) Activation and inactivation curves for heterologously-expressed chimeric Cavs with different β subunits.  
1025 Activation: rCavβ2 V<sub>a1/2</sub> = -19.76 ± 1.16mV, n = 12; NveCavβ V<sub>a1/2</sub> = -23.07 ± 1.16mV, n = 5; EdCavβ1  
1026 V<sub>a1/2</sub> = -18.27 ± 1.08mV, n = 8; EdCavβ2 V<sub>a1/2</sub> = -14.22 ± 1.46mV, n = 5; CcCavβ V<sub>a1/2</sub> = -18.47 ± 1.59mV,

1027 n = 6; CxCav $\beta$  V<sub>a1/2</sub> = -28.89 ± 1.54mV, n = 15; PpCav $\beta$  V<sub>a1/2</sub> = -15.29 ± 1.23mV, n = 10; ChCav $\beta$  V<sub>a1/2</sub> = -  
1028 10.30 ± 1.04mV, n = 12. rCav $\beta$ 2 V<sub>i1/2</sub> = -2.98 ± 13.51mV, n = 12; NveCav $\beta$  V<sub>i1/2</sub> = -68.93 ± 1.53mV, n = 5;  
1029 EdCav $\beta$ 1 V<sub>i1/2</sub> = -56.76 ± 3.18mV, n = 8; EdCav $\beta$ 2 V<sub>i1/2</sub> = -18.84 ± 8.00mV, n = 5; CcCav $\beta$  subunit V<sub>i1/2</sub> = -  
1030 47.81 ± 5.57mV, n = 6; CxCav $\beta$  V<sub>i1/2</sub> = -87.75 ± 1.72mV, n = 15; PpCav $\beta$  V<sub>i1/2</sub> = -99.80 ± 0.92mV, n = 10;  
1031 ChCav $\beta$  V<sub>i1/2</sub> = -70.25 ± 4.67mV, n = 12 cells.

1032 C) Diagram of Cav  $\beta$  subunit domain swaps and the length of the N-terminus swapped in amino acids.

1033 D) Cnidarian Cav  $\beta$  N-termini do not greatly affect voltage-dependent activation of Cav channels containing  
1034 EdCav $\beta$ 2. Voltage-dependent activation (V<sub>a1/2</sub>) of heterologously-expressed Cavs with WT EdCav $\beta$ 2,  $\beta$   
1035 subunits from the indicated cnidarians, and chimeras with their N-termini on EdCav $\beta$ 2, p = 0.5830 for  
1036 average V<sub>i1/2</sub> values across mutant beta subunits, one-way ANOVA with Bartlett's test and post-hoc Tukey  
1037 test, n = 4-7 cells. Data represented as mean ± sem.

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**Figure 5—supplement table 1: Wild type and Chimeric Cav $\beta$  amino acid sequences.**

Protein name	Amino Acid Sequence
<i>Exaiptasia diaphana</i> Cav $\beta$ 1 (EdCav $\beta$ 1)	MAQDFALSNRDIELDSLEHDSTGSSTPSEIQRWHMYSDRSGRVVCKDSEPAYRASD TSSVDEDKETSRRELERRAWEALQAARSKPVAFAVRTNIAYEGLSEDDDSPVHGAA VSFNVKDFLHVKEKFNDWWIGRVVKEGCDIGFIPTPSKLKSLQQVGPATGGRPV RGSSKTVFHNDMVNQAQSPTNTSPSRHSSASVDAENGMEYNEEQHSPTSPSKT STLPRSASGNTVTSQSAPGQQGKSKKAFFKKQEQLPPYDVVPSMRPIVLVGPSLKG YEVTDMQMKAFLFDYMKHFQFSGRVLISRVTSDISLAKRSNLANPSKRNIERSNSKN SGLAEVQQEIERIFELSRGGLNLVVLDCDTVNHPTQLAKTS LAPLVVYVKISAPKVLQ RLIKTRGKTQSRALNVQLVAAEKLAQCSEDLYDLILDETQLQDACHHLGEFLEY WRATHPPNQPGSRPPNMQQSTPQYNVIEAGERPSVYL
<i>Exaiptasia diaphana</i> Cav $\beta$ 2 (EdCav $\beta$ 2)	MGNTDSVQSFTKDSEPAYRASDTSSVDEDKETSRRRELERRAWEALQAARSKPVAF AVRTNIAYEGLSEDDDSPVHGAAVSFNVKDFLHVKEKFNDWWIGRVVKEGCDIG FIPTPSKLKSLQQVGPATGGRPVRGSSKTVFHNDMVNQAQSPTNTSPSRHSSASV VDAENGMEYNEEQHSPTSPSKTSTLPRSASGNTVTSQSAPGQQGKSKKAFFKKQ EQLPPYDVVPSMRPIVLVGPSLKGYEVTDMQMKAFLFDYMKHFQFSGRVLISRVTSDI SLAKRSNLANPSKRNIERSNSKNGLAEVQQEIERIFELSRGGLNLVVLDCDTVNHPT QLAKTS LAPLVVYVKISAPKVLQRLIKTRGKTQSRALNVQLVAAEKLAQCSEDLY DLILDETQLQDACHHLGEFLEYWRATHPPNQPGSRPPNMQQSTPQYNVIEAGERPSVYL
<i>Cyanea Capillata</i> Cav $\beta$ (CcCav $\beta$ )	MWFGTKKSKDSERRKRQPIDVYREQALSVPAYIWGDDLSRKTSCTSSEYGEDD IEQIRVQALEQLAAARVKPVAFAMRANYGYNGAEDDDSPHIHMALSFEPKDFLHI KEKFNNDWLIGRVVREGCDIGFIPSPSKLESRLSGLAGRKMRQSSTSSNLHLQDAF SASSPSEDQRQNSFDDESLPPSSPVKSVNPVGIVGQPN SKTAKKGIFKKNDLSLPPYDV PSMRP VIFVGPSLKGYEVTDMQMKAFLDYLKHFQGRIVITRVTA DISTAKKSTIQ NLAKKPIIKERGATQASQE VNQ EIERIFELCRNLQLVVLDSYTVNYP A QVAKTSLAP IIVYIKISSPKV LTRLVKSRGKSQSKNLNVQLVA AVKLGQCSED MYDV VLDE TQLE DACEHLGEFLEAYWRAAHPSQS NFGAAGAPGSFTANGQPVV NVNSMDPFAQS PTRHLRTAQV
<i>Physalia physalis</i> Cav $\beta$ (PpCav $\beta$ )	MVTASYNVPLDNTSATHSFNYPHAFLLTHSSCSYHSNEGFIN SSTEVDIVDENDFKP LFEGNSNEPHCQKKVISFSSL DVVAPIWYFFEMGDEFDSRKTSCTSSEYGEEDVE ALRVQALEQLAAAASKPVAF AVR ANYGYNGSE DEDCPVNGMAV SFEAKDCLHIK VKFNN DWLIGRVVKEGH DIGFIP S ASR LDN IRQSGIS GKL KLRQ S STSS NM LEDQ SQPLSREQDN RSPSE ERG TS FDD DSP ASPL RNP SGSS LTANNNNNN NTAS NVN NSQ PKGKKGIFKKSENLPYDV VPSMRPII FVGPSLKGYEVTNMQMKAFLDYLKHFQG RIVITRVGADISLAKRSAFQHPGKQPVIQKKGNTQSGIVEVQ EIERIFELCRSMQLV VLD CESINHPSQVAKTSL API I AMIRI A SPKVL TR LIKSRGKSQT KHLNFQL VAAEKL NQCTEDMF D VILDENQLED ACEHLGDFLEAYWRS A VPPR RPYVNSDNRSYNNAG GQSIGNYNGGGQYNGTPQRHLRTAQV
<i>Cassiopea xamachana</i> Cav $\beta$ (CxCav $\beta$ )	MVQKSGMSRGPYPPSQEIPMEVFDPSPQGKYSKRKGRFKRSDGSTSSDTSNSFVR QGSAESYTSR PSDS DV SLEEDREALRKEAERQALA QLEKAKTKPVAFAVRTNVGY NPSPGDEV PVQGVAITFEPKDFLHIKEKYNN DWLIGRVVKEGCEVGFIPSPV KLDS LRL LQEQT LRQN RLSSSKSGDNSSS LGDV VTGTRR PT PPA SKQK QK STEH VPPY DV VPSMRPII LVGPSLKG YEVTDMQMKAFLDFLKHF DGRISIT RVTA DISL AKRSV LNNPSK HII ERSNTR SSLAEVQSEI ERI FELARTLQLVAL DADT INH PAQ LSKT SL API IVYIKITSPKVLQRLIKSRGKSQSKHLNVQIAASEKLAQC PPEMFDI ILDENQLEDAC EHLAEYLEA YWKATHPPSSTPPNPLL NRTMATA ALAASPAPVSNLQGPY LASGDQ PLDRATGEHASVHEY PGELGQPPG L YPSNHPP GRAGTL RALS RQDT FDADTPGSRN SAYTEPGDSCVDMETDPSEGPGPGDPAGGGT PPARQGSWEEE DYEEEMTDNRNR GRNKARYCAEGGGPVLGRNKNELEGWGQGVYIR

<i>Clytia hemisphaerica</i> Cavβ (ChCavβ)	MMHGSQTEPAISSMTSERHKNLSHGSRTSINSQRSTNKKVNSHVSFDESTAAPSS KKPGALSAAGGKKSVDNFSSVLQTVFALRWQKAAQKKKKPDDFQQMYMHS MSGALGSIIGDEFDGRKTSGTSSEYGDGEDLEALRILALEKLQAARTRPVAFAVRA NYGYNGSEDDDSPVHGMAVSFEKDDCLHIKDKFNKDWWIGRVVKEGHNIGFVP PDKLESIRQSGVSGKLKMRQSSTSSNMNLHDDPNQRSPLGEAGGNNSFDDETVN SPVRNVSTESTESNNNTNNNTNSLNAQKGKKGIFKKNEQLHPYYVIPSMRPIIFVGPSL KGYEVTDMMQKALFDYLKHRFSERIIFTRVNADISLAKRSNLNNQNRPFPKKS GQAGLAEVQEEVNRIFELCRSSQLVVLDCTINNPSPQVIKTSLAPIVAIKIASPKVLT RLIKSRGKNQVKHLNIQMAADKLSQCNEEMFDVVLDENQLEDACEHLGEFLEAY WRAAVPGAQEGLISQENGGFVNQGGPNGAGYNGVDQYGTPQRNLRTAQV
<i>Nematostella vectensis</i> cacnb2.1 (NVE β)	MEPEPGLSEQDIELDSLEQVSTASSFHSDIQRHYNDGREASRFIGADDNRDSDPAY RASDTSSIEEDRETSRRELERRAWDALQAARSKPVAFAVRTNLRYDGSEDDDSPVH GAAVSFEAKDFLVKEKFNDDWWIGRVVKEGCDIGFIPTPSKLQLQQIGGTASGR GMRNSKRDVFQFDMVNQAQSPTNTSPSRHSSTSVDAAENGVEYDDDQQSPTSPNK TLPRSASGITVSSQPGTATGTQGKPKGLFKQEQLPPYDVPSMRPIVLVGPSLK GYEVTDMMMQKALLDFMKHRFSGRVLIAVRTSDISLAKRTNMSNPGKQTIMERTKN KNTGLAEVQQEIERIFELARGLNLVVLDCTVNHTQLAKTSLAPMIVYIKIAAPKV LQRLIKTRGKSQLRNLSIQLVAAEKLAAQCSEDMYDLVLEETQLDDACEHLGEFLES YWRATHPPNQPGSRPPNVQPSNSTPQYNVIEGGERPSVYL
Rat cacnb2a (Rat β)	MQCCGLVHRRRVRSYGSADSYSRPSDSDVSLEEDREAVRREAERQAQAQLEK AKTKPVAFAVRTNVRYSAAQEDDVPPVPGMAISFEAKDFLVKEKFNNDWWIGRL VKEGCEIGFIPSPVKLEMNRLQHEQRAKQGKFYSSKSGGNSSSLGDIVPSSRKSTPP SSAIDIDATGLDAEENDIPANHRSPKPSANSVTSPHSKEKRMPFFKKTEHTPPYDV PSMRPVVLVGPSLKGYEVTDMMMQKALFDLKHRFEGRISITRVTAISLAKRSVLN NPSKHAIERSNTRSSLAEVQSEIERIFELARTLQLVVLADTINHPAQLSKTSAPII YVKISSPKVLQRLIKSRGKSQAKHLNVQMVAADKLAQCPPQESFDVILDENQLEDA CEHLADYLEAYWKATHPPSSNLPNPLLSRTLATSTLPLSPTLASNSQGSQGDQRTD RSAPRSASQAEEEPCLEPVKKSQHRSSTATHQNHRSGTGRGLSRQETFDSETQESRD SAYVEPKEDYSHEHVDRYVPHREHNHREESHSSNGHRHREPRHRTDMGRDQDH NECSKQRSRHKSKDRYCDKEGEVISKRRSEAGEWNRDVYIRQ
Rat β with NVE Hook	MQCCGLVHRRRVRSYGSADSYSRPSDSDVSLEEDREAVRREAERQAQAQLEK AKTKPVAFAVRTNVRYSAAQEDDVPPVPGMAISFEAKDFLVKEKFNNDWWIGRL VKEGCEIGFIPSPSKLQLQQIGGTASGRGMRNSKRDVFQFDMVNQAQSPTNTSPS RHSSSTSVDAENGVEYDDDQQSPTSPNKTLPRSASGTTVSSQPGTATGTQGKPKKG LFKKQEQLPPYDVPSMRPVVLVGPSLKGYEVTDMMMQKALFDLKHRFEGRISITR VTADISLAKRSVLNNPSKHAIERSNTRSSLAEVQSEIERIFELARTLQLVVLADTIN HPAQLSKTSAPIIVYVKISSPKVLQRLIKSRGKSQAKHLNVQMVAADKLAQCPPQ ESFDVILDENQLEACEHLADYLEAYWKATHPPSSNLPNPLLSRTLATSTLPLSPTL ASNSQGSQGDQRTDRSAPRSASQAEEEPCLEPVKKSQHRSSTATHQNHRSGTGRGL SRQETFDSETQESRDSA YVEPKEDYSHEHVDRYVPHREHNHREESHSSNGHRHREP RHRTRDMGRDQDHNECSKQRSRHKSKDRYCDKEGEVISKRRSEAGEWNRDVYIRQ Q
NVE β with Rat Hook	MEPEPGLSEQDIELDSLEQVSTASSFHSDIQRHYNDGREASRFIGADDNRDSDPAY RASDTSSIEEDRETSRRELERRAWDALQAARSKPVAFAVRTNLRYDGSEDDDSPV HGAAVSFEAKDFLVKEKFNDDWWIGRVVKEGCDIGFIPTPVKLENMRLQHEQR AKQGKFYSSKSGGNSSSLGDIVPSSRKSTPPSSAIDIDATGLDAEENDIPANHRSPK PSANSVTSPHSKEKRMPFFKKTEHTPPYDVPSMRPIVLVGPSLKGYEVTDMMQK ALLDFMKHRFSGRVLIAVRTSDISLAKRTNMSNPGKQTIMERTKNKNTGLAEVQQ EIERIFELARGLNLVVLDCTVNHTQLAKTSLAPMIVYIKIAAPKVLQRLIKTRGKS QSRNLSIQLVAAEKLAAQCSEDMYDLVLEETQLDDACEHLGEFLESYWRATHPPNQ PGSRPPNVQPSNSTPQYNVIEGGERPSVYL
Rat β with NVE GK	MQCCGLVHRRRVRSYGSADSYSRPSDSDVSLEEDREAVRREAERQAQAQLEK AKTKPVAFAVRTNVRYSAAQEDDVPPVPGMAISFEAKDFLVKEKFNNDWWIGRL

domain	VKEGCEIGFIPSPVKLENMRLQHEQRAKQGKFYSSKGNNSSSLGDIVPSSRKSTPPSSAIDIDATGLDAEENDIPANHRSPKPSANSVTPHSKEKRMPFFKTEHTPPYDVVPSPMRPIVLVGPSLKGYEVTDMMMQKALLDFMKHRFSGRVLIAVRTSDISLAKRTNMSNPGKQTIMERTKNKNTGLAEVQQEIERIFELARGLNLVLDCTVNHPTQLAKTSLAPMIVYIKIAAPKVLQRLIKTRGKSQRNLSQLVAAEKLAQCSEDMYDLVLEETQLDDACEHLGEFLESYWRATHPPNQPGSRPPNVQPSNTPQYNVIEGGERPSVYL
NVE $\beta$ with Rat GK domain	MEPEPGLSEQDIELDSLEQVSTASSFHSDIQRHYNDGREASRFIGADDNRDSDPAYRASDTSSIEEDRETSRRELERRAWDALQAARSKPVAFAVRTNLRYDGSEDDDSPVHGAAVSFEAKDFLVKEKFNDDWWIGRVVKEGCDIGFIPTPSKLKSLQQIGGTASGRGMRNSKRDVFQFDMVNQAQSPTNTSPSKTLPRASAGTTVSSQPGTATGTQGPKKKLFKKQEQLPPYDVVPSPMRPIVLVGPSLKGYEVTDMMMQKALLDFMKHRFSGRVLIAVRTSDISLAKRTNMSNPGKQTIMERTKNKNTGLAEVQQEIERIFELARGLNLVLDCTVNHPTQLAKTSLAPMIVYIKIAAPKVLQRLIKTRGKSQRNLSQLVAAEKLAQCSEDMYDLVLEETQLDDACEHLGEFLESYWRATHPPNQPGSRPPNVQPSNTPQYNVIEGGERPSVYL
Rat 5' on NVE $\beta$	MQCCGLVHRRRVRVSYGSADSYTSRPSDSDVSLEEDREAVRREAERQAQAQLEKAKTKPVAFAVRTNLRYDGSEDDDSPVHGAAVSFEAKDFLVKEKFNDDWWIGRVVKEGCDIGFIPTPSKLKSLQQIGGTASGRGMRNSKRDVFQFDMVNQAQSPTNTSPSKTLPRASAGTTVSSQPGTATGTQGPKKKLFKKQEQLPPYDVVPSPMRPIVLVGPSLKGYEVTDMMMQKALLDFMKHRFSGRVLIAVRTSDISLAKRTNMSNPGKQTIMERTKNKNTGLAEVQQEIERIFELARGLNLVLDCTVNHPTQLAKTSLAPMIVYIKIAAPKVLQRLIKTRGKSQRNLSQLVAAEKLAQCSEDMYDLVLEETQLDDACEHLGEFLESYWRATHPPNQPGSRPPNVQPSNTPQYNVIEGGERPSVYL
Rat 5' + SH3 on NVE $\beta$	MQCCGLVHRRRVRVSYGSADSYTSRPSDSDVSLEEDREAVRREAERQAQAQLEKAKTKPVAFAVRTNVRYSAAQEDDVPPVPGMAISFEAKDFLVKEKFNNDWWIGRVLVKEGCEIGFIPSPSKLKSLQQIGGTASGRGMRNSKRDVFQFDMVNQAQSPTNTSPSKTLPRASAGTTVSSQPGTATGTQGPKKKLFKKQEQLPPYDVVPSPMRPIVLVGPSLKGYEVTDMMMQKALLDFMKHRFSGRVLIAVRTSDISLAKRTNMSNPGKQTIMERTKNKNTGLAEVQQEIERIFELARGLNLVLDCTVNHPTQLAKTSLAPMIVYIKIAAPKVLQRLIKTRGKSQRNLSQLVAAEKLAQCSEDMYDLVLEETQLDDACEHLGEFLESYWRATHPPNQPGSRPPNVQPSNTPQYNVIEGGERPSVYL
NVE 5' + SH3 on Rat $\beta$	MEPEPGLSEQDIELDSLEQVSTASSFHSDIQRHYNDGREASRFIGADDNRDSDPAYRASDTSSIEEDRETSRRELERRAWDALQAARSKPVAFAVRTNVRYSAQEDDVPPVPGMAISFEAKDFLVKEKFNNDWWIGRVLVKEGCEIGFIPSPVKLENMRLQHEQRAKQGKFYSSKGNNSSSLGDIVPSSRKSTPPSSAIDIDATGLDAEENDIPANHRSPKPSANSVTPHSKEKRMPFFKTEHTPPYDVVPSPRVVLVGPSLKGYEVTDMMMQKALFDLKHRFEGRISITRVTAISLAKRSVNNPSKHAIERSNTRSSLAEVQQEIERIFELARTLQLVLDADTINHPAQLSKTSAPIIVYVKISSPKVLQRLIKSRGKSQAKHLNVQMVAADKLAQCPPQESFDVILDENQLEDACEHLADYLEAYWKATHPPSSNLPNPLLSRTLATSTLPLSPTLASNSQGSQGDQRTDRSAPRSASQAEEPCLEPVKKSQHRSSSATHQNHRSGTGRGLSRQETFDSETQESRDSAYPEPKEDYSHEHVDRYVPHREHNHREESSHSSNGHRHREPRHTRDMGRDQDHNECSKQRSRHKSQDRYCDKEVISKRRSEAGEWNRDVYIRQ
NVE 5' on Rat $\beta$	MEPEPGLSEQDIELDSLEQVSTASSFHSDIQRHYNDGREASRFIGADDNRDSDPAYRASDTSSIEEDRETSRRELERRAWDALQAARSKPVAFAVRTNLRYDGSEDDDSPVHGAAVSFEAKDFLVKEKFNDDWWIGRVVKEGCDIGFIPTPVKLENMRLQHEQRAKQGKFYSSKGNNSSSLGDIVPSSRKSTPPSSAIDIDATGLDAEENDIPANHRSPKPSA

	NSVTSPHSKEKRMPFFKKTEHTPPYDVVPSMRPVVLVGPLKGYESVTDMQQKALFDFLKHRF EGRISITRVTADISLAKRSVNNPSKHAIERSNTRSSLAEVQSEIERIFELARTLQLVV LDADTINHPAQLSKTSAPIIVYVKISSPKVLQRLIKSRGKSQAKHNVQMVAADK LAQCQQESFDVILDENQLEDACEHLADYLEAYWKATHPPSSNLNPNPLLSRTLATS TLPLSPTLASNSQGSQGDQRTDRSAPRSASQAEEEPCLEPVKKSQHRSQQATHQNH RSGTGRGLSRQETFDSETQESRDSAYVEPKEDYSHEHVDRYVPHREHNHREESHSS NGHRHREPRHRTRDMGRDQDHNECSKQRSRHKSKDRYCDKEGEVISKRRSEAGE WNRDVYIRQ
EdCav $\beta$ 2 with NVE $\beta$ NTerm	MEPEPGLSEQDIELDSLEQVSTASSFHSDIQRHYNDGREASRFIGADDFRNDRSDPAY RASDTSSIEEDRETSRRELERRAWDALQAARSKPVAFAVRTNLRYDGSEDDDSPV HGAAVSFEAKDFLHVKEKFNFDDWWIGRVVKEGCDIGFIPTPSKLKSLQQVGPATG GRPVRGSSKTVFHNDMVNQAQSPTNTSPSRHSSASVVAENGMEYNEEQHSPT SPTSKTSTLPRSASGNTVTSQSAPGQQGKSKKAFFKKQEQLPPYDVVPSMRPIVLV GPSLKGYESVTDMQQKALFDYMKHQFSGRVLISRTSDISLAKRSNLANPSKRNIIE RSNSKNGLAEVQQEIERIFELSRLGLNLVVLDCDTVNHPTQLAKTSLAPLVVYVKIS APKVLQRLIKTRGKTQSRALNVQLVAAEKLAQCSLEDYDLILDETQLQDACHHLG EFLESYWRATHPPNQPGSRPPNMQQSTPQYNVIEAGERPSVYL
EdCav $\beta$ 2 with CcCav $\beta$ NTerm	MWFGTJKSKDSERRKRQPIDVYREQALSVNPAYIWGDDLSRKTSCTSSEYGEDD IEQIRVQALEQLAAARVKPVAFAMRANYGYNGAEDEDDSPIHGMALSFEPKDFLHI KEKFNNDWLIGRVVREGCDIGFIPSPSKLKSLQQVGPATGGRPVRGSSKTVFHND MVNQAQSPTNTSPSRHSSASVVAENGMEYNEEQHSPTSPSKTSTLPRSASGNT VTSQSAPGQQGKSKKAFFKKQEQLPPYDVVPSMRPIVLVPSLKGYESVTDMQQK ALFDYMKHQFSGRVLISRTSDISLAKRSNLANPSKRNIIERSNSKNGLAEVQQEI ERIFELSRLGLNLVVLDCDTVNHPTQLAKTSLAPLVVYVKISAPKVLQRLIKTRGKT QSRALNVQLVAAEKLAQCSLEDYDLILDETQLQDACHHLGEFLESYWRATHPPNQ PGSRPPNMQQSTPQYNVIEAGERPSVYL
EdCav $\beta$ 2 with PpCav $\beta$ NTerm	MVTASYNVPLDNTSATHSFNYPHAFLLTHSSCSYHSNEGFINSTEVDIVDENDFKP LFEGNSNEPHCQKKVISFSSLNDNVVAPIWYFFEMGDEFDSRKTSCTSSEYGEEDV EALRVQALEQLAAASKPVAFAVRANYGYNGSEDEDCPVNGMAVSFEAKDCLHI KVKFNNNDWWIGRVVKEGHDIGFIPSPSKLKSLQQVGPATGGRPVRGSSKTVFHFN DMVNQAQSPTNTSPSRHSSASVVAENGMEYNEEQHSPTSPSKTSTLPRSASGNT VTSQSAPGQQGKSKKAFFKKQEQLPPYDVVPSMRPIVLVPSLKGYESVTDMQQK ALFDYMKHQFSGRVLISRTSDISLAKRSNLANPSKRNIIERSNSKNGLAEVQQEI ERIFELSRLGLNLVVLDCDTVNHPTQLAKTSLAPLVVYVKISAPKVLQRLIKTRGKT QSRALNVQLVAAEKLAQCSLEDYDLILDETQLQDACHHLGEFLESYWRATHPPNQ PGSRPPNMQQSTPQYNVIEAGERPSVYL
EdCav $\beta$ 2 with Rat $\beta$ NTerm	MQCCGLVHRRRVRSYGSADSYTSRPSDSDVSLEEDREAVRREAERQAQAQLEK AKTKPVAFAVRTNVRYSAAQEDEDVPVPGMAISFEAKDFLHVKEKFNNNDWWIGRL VKEGCEIGFIPSPSKLKSLQQVGPATGGRPVRGSSKTVFHNDMVNQAQSPTNTSP SRHSSASVVAENGMEYNEEQHSPTSPSKTSTLPRSASGNTVTSQSAPGQQGKSK KAFFKKQEQLPPYDVVPSMRPIVLVPSLKGYESVTDMQQKALFDYMKHQFSGR VLISRTSDISLAKRSNLANPSKRNIIERSNSKNGLAEVQQEIERIFELSRLGLNLVVL DCDTVNHPTQLAKTSLAPLVVYVKISAPKVLQRLIKTRGKTQSRALNVQLVAAEK LAQCSLEDYDLILDETQLQDACHHLGEFLESYWRATHPPNQPGSRPPNMQQSTPQ YNTVIEAGERPSVYL
EdCav $\beta$ 2 with CxCav $\beta$ NTerm	MVQKSGMSRGPYPPSQEIPMEVFDPSPQGKYSRKGRFKRSDGSTSDTTNSFVR QGSAESYTSRPSDSDVSLEEDREALRKEAERQALAQLEKAKTKPVAFAVRTNVGY NPSPGDEVVPQGVAITFEPKDFLHIKEKYNDWWIGRLVKEGCEVGFIPSPSKLKS LQQVGPATGGRPVRGSSKTVFHNDMVNQAQSPTNTSPSRHSSASVVAENGME YNEEQHSPTSPSKTSTLPRSASGNTVTSQSAPGQQGKSKKAFFKKQEQLPPYDV VPSMRPIVLVPSLKGYESVTDMQQKALFDYMKHQFSGRVLISRTSDISLAKRSNL ANPSKRNIIERSNSKNGLAEVQQEIERIFELSRLGLNLVVLDCDTVNHPTQLAKTS

	LAPLVVYVKISAPKVLQRLIKTRGKTQSRALNVQLVAAEKLAQCSEDLYDLILDETQLQDACHHLGEFLESYWRATHPPNQPGSRPPNMQQSTPQYNVIEAGERPSVYL
EdCav $\beta$ 2 with ChCav $\beta$ NTerm	MMHGSQTEPAISSMTSERNHKNLSHGSRTSINSQRSTNKVVNSHVSFDESTAAPSSKKPGALSAAGGKKSVDDNFSSVLQTVFALRWQKKAQKKKKPDDFQQMYMHSMSGALGSIIGDEFDGRKTSGTSSEYGDGEDLEALRILALEKLQAARTRPVAFAVRANYGYNGSEDDDSPVHGMAVSFEKDDCLHIKDKFNKDWWIGRVVKEGHNIGFVPSPSKLKSLQQVGPATGGRPVRGSSKTVFHFNMDMVNQAQSPTNTSPSRHSSASVVDASENGMEYNEEEQHSPTSPSKTSTLPRSASGNTVTQSAPGQQGKSKKAFFKKQEQLPPYDVVPSMRPIVLVGPSLKGYEVTDMMQKALFDYMKHQFSGRVLISRTSDISLAKRSNLANPSKRNIIERSNSKNSGLAEVQQEIERIFELSRLGNLVVLDCDTVNHPTQLAKTSLAPLVVYVKISAPKVLQRLIKTRGKTQSRALNVQLVAAEKLAQCSEDLYDILDETQLQDACHHLGEFLESYWRATHPPNQPGSRPPNMQQSTPQYNVIEAGERPSVYL
EdCav $\beta$ 2 with EdCav $\beta$ 1 NTerm	MAQDFALSNRDIELDSLEHVSTGSSTPSEIQRWHMYSDRSGRVVCKDSEPAYRASDTSSVDEDKETSRELLERAWEALQAARSKPVAFAVRTNIA YEGSEDDDSPVHGA AVSFNVKDFLHVKEKFNFDDWWIGRVVKEGCDIGIPTPSKLKSLQQVGPATGGRPVRGSSKTVFHFNMDMVNQAQSPTNTSPSRHSSASVVDASENGMEYNEEEQHSPTSPSKTSTLPRSASGNTVTQSAPGQQGKSKKAFFKKQEQLPPYDVVPSMRPIVLVGPSLKGYEVTDMMQKALFDYMKHQFSGRVLISRTSDISLAKRSNLANPSKRNIIERSNSKNSGLAEVQQEIERIFELSRLGNLVVLDCDTVNHPTQLAKTSLAPLVVYVKISAPKVLQRLIKTRGKTQSRALNVQLVAAEKLAQCSEDLYDILDETQLQDACHHLGEFLESYWRATHPPNQPGSRPPNMQQSTPQYNVIEAGERPSVYL

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