

Table S1 List of the α -CA sub-families with the species and accession number of the amino acid sequences used in this study.

α -CA Sub-Family	Species Name	Accession #
CA 10	<i>Homo sapiens</i> _CA10	AAH47456.1
	<i>Mus musculus</i> _CA10	AHH17606.1
	<i>Macaca mulatta</i> _CA10	AF64410.1
	<i>Equus caballus</i> _CA10	XP_001503286.2
	<i>Bos taurus</i> _CA10	AAI26508.1
	<i>Gallus gallus</i> _CA10	XP_415644.1
	<i>Taenioptygia guttata</i> _CA10	XP_002198310.1
	<i>Anolis carolinensis</i> _CA10	XP_003216966.1
	<i>Rattus norvegicus</i> _CA10	XP_002727818.1
	<i>Xenopus tropicalis</i> _CA10	XP_002932546.1
	<i>Danio rerio</i> _CA10	NP_001032198.1
	<i>Oreochromis niloticus</i> _CA10	XP_003438532.1
CA11	<i>Macaca mulatta</i> _CA11	AF64868.1
	<i>Bos taurus</i> _CA11	AAI23828.1
	<i>Homo sapiens</i> _CA11	AAH02662.1
	<i>Mus musculus</i> _CA11	AAH19392.1
	<i>Rattus norvegicus</i> _CA11	AAI78173.1
	<i>Loxodonta africana</i> _CA11	XP_003406877.1
	<i>Anolis carolinensis</i> _CA11	XP_003225067.1
	<i>Strongylocentrotus purpuratus</i> _CA11	XP_784796.2
	<i>Branchiostoma floridae</i> _CA11	XP_002611575
	<i>Drosophila melanogaster</i> _CA11a	NP_572581.1
	<i>Tribolium castaneum</i> _CA11a	XP_972474.2
	<i>Nasonia vitripennis</i> _CA11a	XP_003424234.1
	<i>Drosophila melanogaster</i> _CA11b	NP_572407.3
	<i>Aedes aegypti</i> _CA11	XP_0016588761.1
	<i>Tribolium castaneum</i> _CA11b	XP_974173.2
	<i>Daphnia pulex</i> _CA4	EFX88377.1
	<i>Daphnia galeata</i> _CA4	unpublished
	<i>Daphnia pulex</i> _CA3	EFX72738.1
	<i>Daphnia galeata</i> _CA3	unpublished
	<i>Loa loa</i> _CA11	XP_003137734.1
	<i>Brugia malayi</i> _CA11	XP_001900835.1
	<i>Caenorhabditis elegans</i> _CA11a	NP_495567.3
	<i>Trichinella spiralis</i> _CA11	XP_003376534.1
	<i>Caenorhabditis elegans</i> _CA11b	NP_498083.1
CA 8	<i>Homo sapiens</i> _CA8	EAW86826.1
	<i>Macaca mulatta</i> _CA8	EHH28528.1
	<i>Mus musculus</i> _CA8	AAH10773.1
	<i>Rattus norvegicus</i> _CA8	AAH87586.1
	<i>Bos taurus</i> _CA8	NP_001077159.1
	<i>Equus caballus</i> _CA8	XP_001496523.1
	<i>Gallus gallus</i> _CA8	XP_419221.3
	<i>Meleagris gallopavo</i> _CA8	XP_003205121.1
	<i>Taenioptygia guttata</i> _CA8	XP_002197204>1
	<i>Anolis carolinensis</i> _CA8	XP_003219630.1
	<i>Xenopus tropicalis</i> _CA8	NP_001011213.1
	<i>Danio rerio</i> _CA8	AAH92740.1
	<i>Oncorhynchus mykiss</i> _CA8	NP_001118116.1
CA2	<i>Strongylocentrotus purpuratus</i> _CA8	XP_795365.3
	<i>Homo sapiens</i> _CA2	NP_000058.1
	<i>Macaca mulatta</i> _CA2	NP_001182346.1
	<i>Equus caballus</i> _CA2	XP_001488540.1
	<i>Bos taurus</i> _CA2	NP_348667.1
	<i>Mus musculus</i> _CA2	EDL05130.1
	<i>Rattus norvegicus</i> _CA2	NP_062164.1
	<i>Gallus gallus</i> _CA2	NP_990648.1
	<i>Taenioptygia guttata</i> _CA2	XP_002199833.1
	<i>Anolis carolinensis</i> _CA2	XP_003219585.1
	<i>Xenopus tropicalis</i> _CA2	AAH89661.1
	<i>Xenopus laevis</i> _CA2	NP_001080080.1

Piscean CA1/2	Oncorhynchus mykiss_CA1	NP_001117692.1
	Dicentrarchus labrax_CA1	CBN82139.1
	Oncorhynchus mykiss_CA2	NP_001117693.1
	Ictalurus punctatus_CA2	NP_001187560.1
	Danio rerio_CA2	NP_954685.1
CA 1	Homo sapiens_CA1	NP_001729.1
	Macaca mulatta_CA1	EHH28617.1
	Equus caballus_CA1	XP_001488605
	Mus musculus_CA1	AAA37354.1
	Rattus norvegicus_CA1	AAI58889.1
	Bos taurus_CA1	AAI16127.1
	Xenopus tropicalis_CA1	XP_002939198
CA 13	Homo sapiens_CA13	AAH52602.1
	Macaca mulatta_CA13	XP_001095487.1
	Equus caballus_CA13	XP_001489984.2
	Mus musculus_CA13	AAK16672.1
	Rattus norvegicus_CA13	NP_001128465.1
	Bos taurus_CA13	DAA22563.1
	Gallus gallus_CA13	XP_003640859.1
	Meleagris gallopavo_CA13	XP_003205200.1
	Python molurus_CA13	AEA49961.1
	Anolis carolinensis_CA13	XP_003219587.1
	Xenopus tropicalis_CA13	NP_001072448.1
	Xenopus laevis_CA13	NP_001086981.1
	Petromyzon marinus_CA2	
	Homo sapiens_CA7	NP_005173.1
	Macaca mulatta_CA7	XP_001085299.2
CA 7	Mus musculus_CA7	NP_444300.1
	Rattus norvegicus_CA7	NM_001106165.1
	Loxodonta africana_CA7	XM_003417135.1
	Bos taurus_CA7	NP_001179451.1
	Gallus gallus_CA7	XP_41415.2
	Meleagris gallopavo_CA7	XP_003209867.1
	Taeniopygia guttata_CA7	XP_002190292.1
	Xenopus tropicalis_CA7	NP_001015903.1
	Danio rerio_CA7	NM_200813.1
	Ictalurus punctatus_CA7	NP_001187680.1
	Oreochromis niloticus_CA7	XM_003445759.1
	Homo sapiens_CA5a	NP_001730.1
	Macaca mulatta_CA5a	EHH31930.1
	Equus caballus_CA5a	XP_001500362.2
CA 5	Bos taurus_CA5a	DAA20167.1
	Mus musculus_CA5a	AAH30174.1
	Rattus norvegicus_CA5a	AAH88147.1
	Homo sapiens_CA5b	NP_00915.1
	Macaca mulatta_CA5b	EHH30558.1
	Equus caballus_CA5b	XP_001490399.1
	Bos taurus_CA5b	NP_001074377.2
	Mus musculus_CA5b	CAM26357.1
	Rattus norvegicus_CA5b	AAH81872.1
	Gallus gallus_CA5b	XP_414195.3
	Meleagris gallopavo_CA5b	XP_003209935.1
	Taeniopygia guttata_CA5b	XP_002192230.2
	Xenopus tropicalis_CA5b	CAJ82585.1
	Xenopus laevis_CA5b	NP_001084659.1
	Danio rerio_CA5b	NP_001104671.1
CA 3	Homo sapiens_CA3	3UYN
	Macaca mulatta_CA3	EHH28618.1
	Mus musculus_CA3	AAH11129.1
	Rattus norvegicus_CA3	AAH61980.1
	Equus caballus_CA3	NP_001157426.1
	Bos taurus_CA3	DAA22526.1

	Gallus gallus_CA3	NP_001264340.1
	Meleagris gallopavo_CA3	XP_003205196.1
	Taeniopygia guttata_CA3	XP_002199827.1
	Anolis carolinensis_CA3	XP_003219614.1
Invertebrate Cytosolic CA	Branchiostoma floridae_ccA2	XP_00259304.1
	Branchiostoma floridae_ccA3	XP_002605865.1
	Ixodes scapularis_ccA	XP_002416504.1
	Anthopleura elegantissima_ccA	AAD32675.1
	Strongylocentrotus purpuratus_ccA	XP_782997.2
	Drosophila melanogaster_ccA	AAF53332.1
	Anopheles gambiae_ccA	ABF66618.1
	Danaus plexippus_ccA	EHJ77985.1
	Tribolium castaneum_ccA	XP_974322.1
	Daphnia pulicaria_CA1	unpublished
	Daphnia pulex_CA1	EFX86103.1
	Daphnia galeata_CA1	unpublished
	Daphnia pulicaria_CA2	unpublished
	Daphnia pulex_CA2	EFX86394.1
	Daphnia galeata_CA2	unpublished
	Panaeus monodon_ccA	ABV65904.1
	Litopenaeus vannamei_ccA	ADM16544.2
	Callinectes sapidus_ccA	ABN51213.1
	Portunus trituberculatus_ccA	AFV46144.1
	Carcinus maenus_ccA	ABX71208.1
	Caligus rogercresseyi_ccA	ACO11717.1
	Caligus clemensi_ccA	ACO15216.1
	Lepeophtheirus salmonis_ccA	ACO12707.1
CA 4	Homo sapiens_CA4	NP_000708.1
	Macaca mulatta_CA4	XP_001107970.2
	Bos taurus_CA4	AAB09466.1
	Equus caballus_CA4	XP_001501182.1
	Mus musculus_CA4	AAH12704.1
	Rattus norvegicus_CA4	AAH97329.1
	Gallus gallus_CA4	XP_415893.1
	Python molurus_CA4	AEA49062.1
	Danio rerio_CA4	AAI09406
	Oreochromis niloticus_CA4	XP_003456174.1
	Xenopus tropicalis_CA4	XP_002983882.1
	Xenopus laevis_CA4	NP_001079788.1
	mus musculus_CA15	NP_085035.1
	Rattus norvegicus_CA15	NP_001099371.1
CA15	Equus caballus_CA15	XP_005613908.1
	Gallus gallus_CA15	XP_415218
	Meleagris gallopavo_CA15	XP_003211059.1
	Taeniopygia guttata_CA15	XP_002195938.2
	Anolis carolinensis_CA15	XP_003225153.1
	Xenopus tropicalis_CA15	XP_002931952.2
	Homo sapiens_CA6	NP_001206.2
CA 6	Macaca mulatta_CA6	XP_001099188.2
	Bos taurus_CA6	DAA21262.1
	Equus caballus_CA6	NP_001137421.2
	Mus musculus_CA6	NP_033932.2
	Rattus norvegicus_CA6	NP_001128313.1
	Gallus gallus_CA6	XP_425745.3
	Meleagris gallopavo_CA6	XP_003212306.1
	Taeniopygia guttata_CA6	XP_002187446.1
	Xenopus laevis_CA6	NP_001123372.1
	Xenopus tropicalis_CA6	NP_001085550.1
	Danio rerio_CA6	XP_002666525.1
	Homo sapiens_CA9	NP_001207.2
	Macaca mulatta_CA9	XP_001088481.1
	Equus caballus_CA9	XP_001504561.3

CA9	Bos taurus_CA9	XP_002689744.1
	Mus musculus_CA9	AAI20545.11
	Rattus norvegicus_CA9	EDL98746.1
	Gallus gallus_CA9	XP_001233320.2
	Danio rerio_CA9	XP_694982.2
CA12	Homo sapiens_CA12	NP_001209.1
	Macaca mulatta_CA12	EHH27385.1
	Mus musculus_CA12	AAH35941.1
	Rattus norvegicus_CA12	NP_001074225.1
	Bos taurus_CA12	XP_870878.3
	Gallus gallus_CA12	XP_413756.2
	Meleagris gallopavo_CA12	XP_003209429.1
	Xenopus tropicalis_CA12	CAJ81489.1
	Xenopus laevis_CA12	NP_001091227.1
	Homo sapiens_CA14	BAA85002.1
CA14	Macaca mulatta_CA14	NP_001244832.1
	Equus caballus_CA14	XP_001489157.1
	Bos taurus_CA14	NP_001179134.1
	Mus musculus_CA14	NP_035927.1
	Rattus norvegicus_CA14	NP_001103125.1
	Gallus gallus_CA14	XP_003642713.1
	Meleagris gallopavo_CA14	XP_00312828.1
	Xenopus tropicalis_CA14	NP_001103521.1
	Danio rerio_CA14	NP_001032782.1
	Oreochromis niloticus_CA14	XP_005810348.1
Daphnia CA 5/6/7	Daphnia pulex_CA6B	EFX88008.1
	Daphnia galeata_CA6B	unpublished
	Daphnia pulex_CA6E	EFX88011.1
	Daphnia galeata_CA6E	unpublished
	Daphnia pulex_CA6F	EFX88012.1
	Daphnia galeata_CA6F	unpublished
	Daphnia pulex_CA6C	EFX88009.1
	Daphnia galeata_CA6C	unpublished
	Daphnia pulex_CA6D	EFX88010.1
	Daphnia pulex_CA6G	EFX88013.1
	Daphnia pulicaria_CA5	unpublished
	Daphnia pulex_CA5	EFX81683.1
	Daphnia galeata_CA5	unpublished
	Daphnia pulex_CA6A	EFX88007.1
	Daphnia galeata_CA6A	unpublished
	Daphnia pulex_CA7D	EFX88103.1
	Daphnia galeata_CA7D	unpublished
	Daphnia pulex_CA7A	EFX88100.1
	Daphnia galeata_CA7A	unpublished
	Daphnia pulex_CA7B	EFX880101.1
	Daphnia galeata_CA7B	unpublished
	Daphnia pulex_CA7I	EFX88114.1
	Daphnia pulex_CA7M	EFX88119.1
	Daphnia galeata_CA7I	unpublished
	Daphnia galeata_CA7G	unpublished
	Daphnia pulex_CA7H	EFX88113.1
	Daphnia pulex_CA7L	EFX88118.1
	Daphnia pulex_CA7K	EFX88116.1
	Daphnia pulex_CA7O	EFX88121.1
	Daphnia pulex_CA7P	EFX88123.1
	Daphnia galeata_CA7P	unpublished
	Daphnia pulex_CA7J	EFX88115.1
	Daphnia pulex_CA7N	EFX88120.1
	Daphnia pulex_CA7Q	EFX77528.1
	Daphnia galeata_CA7H	unpublished
	Daphnia galeata_CA7J	unpublished
	Daphnia pulex_CA7C	EFX88102.1
	Daphnia pulex_CA7E	EFX88104.1

	<i>Daphnia galeata</i> _CA7C	unpublished
	<i>Daphnia pulex</i> _CA6H	EFX88180.1
Invertebrate GPI-anchored CA	<i>Callinectes sapidus</i> _gCA	ABN51214.1
	<i>Portunus trituberculatus</i> _gCA	AFV46145.1
	<i>Carcinus maenius</i> _gCA	ABX71209.1
	<i>Branchiostoma floridae</i> _gCA	XP_002601262.1
	<i>Strongylocentrotus purpuratus</i> _gCA	XP_796525.1
	<i>Anopheles gambiae</i> _gCA	AC528257.1
	<i>Danaus plexippus</i> _gCA	EHJ70026.1
	<i>Patella vulgata</i> _gCA	CCJ509593.1
	<i>Neisseria gonorrhoeae</i> _CA	YP_207719.1
	<i>Kingella denitrificans</i> _CA	WP_003782874.1
Bacteria CA	<i>Pastuerella multocida</i> _CA	WP_005752370.1
	<i>Streptococcus mutans</i> _CA	WP_002278508.1
	<i>Thiorhodococcus drewsii</i> _CA	WP_007041651.1
	<i>Aeromonas caviae</i> _CA	WP_010675444.1
	<i>Yersinia enterocolotica</i> _CA	CCQ39919.1
	<i>Serratia</i> sp._CA	YP_006027343.1
	<i>Enterobacter cloacae</i> _CA	EUL61977.1
	<i>Klebsiella oxytoca</i> _CA	EH590795.1
	<i>Pantoea vagans</i> _CA	WP_13358087.1
	<i>Thermovibrio ammonificans</i> _CA	YP_004152175.1

Table S2 Listing of conserved amino acid residues for each group of α -CAs along with *Daphnia* α -CAs. The gray highlighted residues represent conserved sites throughout the phylogeny.

Dark green represents sites conserved in all extracellular α -CAs, medium green the GPI-anchored CAs, light green the transmembrane CAs, pink the secretory CAs. The dark blue represents sites that are conserved in all intracellular α -CAs, medium blue the cytosolic CAs and red the CA-RPs. The bacterial group shows conserved sites across α -CAs. AS = active site, ZB = zinc-binding site, SB = substrate-binding site, GPI = purported GPI-anchoring site, SuB = disulfide bond site.

		Residue	
GPI-anchored CAs	Extracellular CAs		
Invert GPI CAs		W 180	
Daphnia CA5		W 182	
Daphnia CA6s		Y 183	
Daphnia CA7s		Y 184	
Vert GPI CAs		W 185	
Secretory CAs (CA6s)		W 191	
Transmembrane CAs		W 193	
CA-RP	Intracellular CAs		
CA-RP 8		P F W G L N W	
CA-RP 11		P F W G L N F P D	
Daphnia CA3		Y G V E W G L N W	
Daphnia CA4		P A F W G L I N P E W S	
CA-RP 10		P S F W G L I N P Q W M	
Cytosolic CAs		V P S F W G L V N S A W N	
Invert cCAs		P W	
Daphnia CA1		A E H Y E	
Daphnia CA2		C G P H S Y T A K H F L H	
Vert cCAs		P W	
Bacterial CAs		W Y G	
GPI-anchored CAs	Extracellular CAs	305	
Invert GPI CAs		306	
Daphnia CA5		308	
Daphnia CA6s		309	
Daphnia CA7s		310	
Vert GPI CAs		311	
Secretory CAs (CA6s)		312	
Transmembrane CAs		N N G H T	
CA-RP	Intracellular CAs		
CA-RP 8		N T G K V N G G P P	
CA-RP 11		N D G H I Q L K S K V L G P L P	
Daphnia CA3		G L N T G F I T V D N S S K P S I N L T G G P P	
Daphnia CA4		V A G Y L S N M Q G L I F T V D N S S K P S I N L T G G P P	
CA-RP 10		R K V S G T M Y N T G R H V S L R L D K E H L S L A V N V S A G P M T Y S H R L E E I R L H F G S E D	
Cytosolic CAs		N G W L G P L Y L Q F H H W G	
Invert cCAs		P K H T V D V T N T G Y C W K A H V H G E E S S L E G G P L K D K Y L Q H C H W G V N N V G	
Daphnia CA1		P K E T E L L N N G S V F D D S T L S G G P L S H Y R L Q F H H W G	
Daphnia CA2			
Vert cCAs		N N G H T G L Q H F H P	
Bacterial CAs			
GPI-anchored CAs	Extracellular CAs	371	
Invert GPI CAs		372	
Daphnia CA5		373	
Daphnia CA6s		374	
Daphnia CA7s		375	
Vert GPI CAs		376	
Secretory CAs (CA6s)		377	
Transmembrane CAs		378	
CA-RP	Intracellular CAs		
CA-RP 8		G N G T D C V N D G H I Q L K S K V L G P L P	
CA-RP 11		G L N T G F I T V D N S S K P S I N L T G G P P	
Daphnia CA3		V A G Y L S N M Q G L I F T V D N S S K P S I N L T G G P P	
Daphnia CA4		V S G V L N T G Q S L V F R P E T G P K H S L A V N V S A G P M T Y S H R L E E I R L H F G S E D	
CA-RP 10		R K V S G T M Y N T G R H V S L R L D K E H L S L A V N V S A G P M T Y S H R L E E I R L H F G S E D	
Cytosolic CAs		N G W L G P L Y L Q F H H W G	
Invert cCAs		P K H T V D V T N T G Y C W K A H V H G E E S S L E G G P L K D K Y L Q H C H W G V N N V G	
Daphnia CA1		P K E T E L L N N G S V F D D S T L S G G P L S H Y R L Q F H H W G	
Daphnia CA2			
Vert cCAs		N N G H T G L Q H F H P	
Bacterial CAs			
GPI-anchored CAs	Extracellular CAs	407	
Invert GPI CAs		408	
Daphnia CA5		409	
Daphnia CA6s		410	
Daphnia CA7s		402	
Vert GPI CAs		403	
Secretory CAs (CA6s)		404	
Transmembrane CAs		405	
CA-RP	Intracellular CAs		
CA-RP 8		S E H T D G M E L H V H N K Y G S L G A 407	
CA-RP 11		S E H T I N N P A M E L H V H Y F V H F 408	
Daphnia CA3		S E H T I S K A A E L H V H N K Y G S L G A 409	
Daphnia CA4		S E H T I Y A E H I V H Y N K Y G S L G A 410	
CA-RP 10		S E H T I N G M E H V H N K Y G S L G A 411	
Cytosolic CAs		S E H T D G R E H V H Y N S K Y S E A D P 412	
Invert cCAs		S E H T G A E H V H Y N K Y S A D P 413	
Daphnia CA1		S E H T G A E H V H Y N K Y S A D P 414	
Daphnia CA2		S E H T G A E H V H Y N K Y S A D P 415	
Vert cCAs		S E H T D G I E L Q R D N P A 416	
Bacterial CAs		S E H T D G I E L Q R D N P A 417	
GPI-anchored CAs	Extracellular CAs	418	
Invert GPI CAs		E S 418	
Daphnia CA5		D 420	
Daphnia CA6s		D 421	
Daphnia CA7s		D 422	
Vert GPI CAs		D 433	
Secretory CAs (CA6s)		D N P A 434	
Transmembrane CAs		D N P A 435	
CA-RP	Intracellular CAs		
CA-RP 8		S E H T V N F K A F P M E L H L I H W N S T L S I D E A G K H G I I I A L F V Q I G K E H G L K A	
CA-RP 11		S E H T V N F K A F P M E L H L I H W N S T L S I D E A G K H G I I I A L F V Q I G K E H G L K A	
Daphnia CA3		S E H T M I A G R A F P A E L Q V F G F N T Q L Y S T Y E A L E S P H G I V A L S V L I Q I G D A S L Q T T E F R L	
Daphnia CA4		S E H T I Q G A F P A E I Q F Y G Y N A E L Y A N A S E A R H K S Q G L V A A V M Q I G E T P N E L R S	
CA-RP 10		S E H T L N G Q A F S G E V Q L I H Y N H E L Y T N V T E A A K S P N G L V V V S I F K V S S S N P F L N R M L	
Cytosolic CAs		S E H T V D G Y E L H L V H W N K Y S F E A G L A V L G V F E G K E H E E K	
Invert cCAs		S E H T V D G K A Y A E L H L V H W N T K Y S F E A G L A V L G V F V E G K E H E E K	
Daphnia CA1		S E H T V N G Q Y G A E I H F V H W N T K Y G S F E A L K H C G G L A V L G V F V E G K E H E E K	
Daphnia CA2		S E H T V D G Y A E L H L V H W N T K Y G S F E A L K Y G D G L A V L G V F L K V G K E N P E L N K Q	
Vert cCAs		S E H T V D G P E H V H L A V L G V F E G K E H E E K	
Bacterial CAs		S E N G P E H V H L A V L G V F E G K E H E E K	

		Residue	
GPI-anchored CAs			
Invert GPI CAs	Extracellular CAs	Residue	
Daphnia CA5		441	
Daphnia CA6s		442	
Daphnia CA7s		443	
Vert GPI CAs		444	
Secretory CAs (CA6s)		445	
Transmembrane CAs		446	
CA-RP	Intracellular CAs	447	
CA-RP 8		448	
CA-RP 11		449	
Daphnia CA3		450	
Daphnia CA4		451	
CA-RP 10		452	
Cytosolic CAs		453	
Invert cCAs		454	
Daphnia CA1		455	
Daphnia CA2		456	
Vert cCAs		457	
Bacterial CAs		458	
GPI-anchored CAs			
Invert GPI CAs	Extracellular CAs	Residue	
Daphnia CA5		508	
Daphnia CA6s		510	
Daphnia CA7s		511	
Vert GPI CAs		512	
Secretory CAs (CA6s)		513	
Transmembrane CAs		514	
CA-RP	Intracellular CAs	515	
CA-RP 8		516	
CA-RP 11		517	
Daphnia CA3		518	
Daphnia CA4		519	
CA-RP 10		520	
Cytosolic CAs		521	
Invert cCAs		522	
Daphnia CA1		523	
Daphnia CA2		524	
Vert cCAs		525	
Bacterial CAs		526	
GPI-anchored CAs			
Invert GPI CAs	Extracellular CAs	Residue	
Daphnia CA5		W	V
Daphnia CA6s		T	F
Daphnia CA7s		T	F
Vert GPI CAs		T	F
Secretory CAs (CA6s)		V	V
Transmembrane CAs		V	S
CA-RP	Intracellular CAs	P	I
CA-RP 8		T	P
CA-RP 11		T	P
Daphnia CA3		A	I
Daphnia CA4		A	I
CA-RP 10		I	T
Cytosolic CAs		M	N
Invert cCAs		K	P
Daphnia CA1		P	V
Daphnia CA2		V	E
Vert cCAs		V	E
Bacterial CAs		K	F
GPI-anchored CAs			
Invert GPI CAs	Extracellular CAs	Residue	
Daphnia CA5		L	R
Daphnia CA6s		L	V
Daphnia CA7s		L	R
Vert GPI CAs		H	V
Secretory CAs (CA6s)		R	V
Transmembrane CAs		V	E
CA-RP	Intracellular CAs	R	V
CA-RP 8		V	E
CA-RP 11		V	F
Daphnia CA3		R	V
Daphnia CA4		R	V
CA-RP 10		C	I
Cytosolic CAs		R	V
Invert cCAs		E	R
Daphnia CA1		R	C
Daphnia CA2		V	R
Vert cCAs		V	A
Bacterial CAs		F	S

Table S3 Ancestral states of amino acids at informative residues. Ancestral states were inferred using a Maximum Likelihood model in MEGA 5.0 and the most probable states at a given residue were selected. CA groups are based on the most recent common ancestor as determined by the phylogenetic analysis, for example the hypothetical ancestral Animalia α -CA state is the predicted sequence of the most recent common ancestor of the extracellular α -CAs and intracellular α -CAs, while the extracellular α -CA hypothetical ancestor is the most recent common ancestor to the invertebrate and vertebrate clades of Extracellular α -CAs (Figure 1). The residues are based on the alignment performed for phylogeny construction. Codes on the residues are as follows: SuB = disulfide bond site, AS = active site, ZB = zinc-binding site, and SB = substrate-binding site. Shaded amino acids reflect an amino acid change from the most common recent ancestor and the shaded amino acids in a box are amino acid change that resulted in convergent evolution. The no. of amino acid changes reflects the number of changes from the most common recent ancestor for that group.

Residue	214	232	233	314	315	AS	318	319	351	363	AS	354	355	ZB	356	381	385	ZB	387	419	434	445	448	491	492	499	SB	501	505	512	573	578	No. of Amino Acid Changes
CA Group																																	
Bacterial	C	P	I	N	H	I	Q	L	Q	F	H	F	P	H	V	G	N	L	W	Y	R	T	P	G	K	P	N						
Ancestral Animalia CA	C	D	I	N	H	I	Q	L	Q	F	H	F	P	H	V	G	N	L	I	Y	R	T	P	G	K	P	N	2					
Extracellular	C	D	I	N	H	V	D	L	Q	F	H	F	P	H	V	S	N	Y	I	Y	R	T	P	V	K	P	N	5					
Invertebrate	C	D	I	N	H	V	T	A	Q	L	H	F	P	H	V	S	N	F	I	Y	R	T	T	V	E	P	N	6					
Vertebrate	C	D	S	N	H	V	D	A	Q	F	H	F	P	H	V	S	N	Y	I	Y	R	T	P	V	E	P	N	3					
GPI	C	D	S	N	H	V	D	A	Q	F	H	F	P	H	V	S	N	Y	I	Y	R	T	P	V	E	P	N	0					
CA 4	C	D	S	N	H	V	D	A	Q	F	H	F	P	H	V	S	N	Y	I	Y	R	T	P	T	E	P	N	1					
CA 15	C	D	S	D	H	L	S	A	Q	F	H	F	P	H	V	S	N	Y	V	Y	R	T	D	V	E	P	K	6					
Secretory																																	
CA6	C	D	P	N	H	I	S	A	Q	M	H	L	V	H	V	G	E	Y	I	Y	R	T	P	S	D	A	N	11					
Transmembrane	C	D	P	N	H	L	S	A	Q	L	H	L	P	H	V	G	N	Y	L	Y	R	T	P	S	N	A	N	10					
CA9	C	S	P	N	H	L	T	A	Q	L	H	L	P	H	V	G	N	Y	L	F	R	T	P	S	N	A	N	3					
CA12	C	D	S	N	H	M	S	A	Q	L	H	L	A	H	V	G	N	Y	I	Y	R	T	P	S	R	Q	N	6					
CA14	C	D	P	N	H	L	S	A	Q	L	H	L	P	H	V	G	N	Y	L	Y	R	T	P	S	N	A	N	0					
Intracellular	C	D	P	T	H	S	F	L	Q	F	H	F	P	H	V	G	N	L	T	W	T	T	P	S	K	P	N	8					
CA-RP	C	D	P	T	H	T	V	L	E	I	R	F	P	H	I	G	N	L	T	W	T	T	P	S	K	P	N	6					
CA8	N	D	P	I	H	T	V	L	E	V	R	F	P	H	I	G	H	L	T	W	T	T	P	G	R	P	S	7					
CA10	C	D	P	R	H	S	D	L	E	I	R	L	S	Q	I	G	N	N	N	I	T	I	P	T	N	P	N	12					
CA11	C	D	P	R	S	S	D	L	E	I	R	L	P	Q	I	G	N	L	T	M	T	T	A	T	N	P	H	11					
Cytolsolic	A	D	P	T	N	S	F	L	Q	F	H	F	P	H	V	G	H	L	T	W	T	T	P	S	K	P	K	4					
Invertebrate	A	D	P	T	T	S	F	L	Q	F	H	A	P	H	V	G	H	L	T	W	T	T	P	S	K	P	K	2					
Vertebrate	A	D	P	S	N	S	F	L	Q	F	H	F	P	H	V	G	H	L	T	W	T	T	P	S	K	P	K	1					
CA5	G	D	P	W	N	S	F	L	Q	F	H	F	P	H	V	G	H	L	V	W	T	T	P	S	K	P	K	3					
CA7	E	D	P	S	H	S	V	L	Q	F	H	F	P	H	V	G	H	L	T	W	T	T	P	S	K	P	K	3					
Fish CA1/2	N	D	A	L	H	S	F	L	Q	F	H	F	P	H	V	G	N	L	L	W	T	T	P	S	K	P	K	5					
CA1	N	D	P	V	H	S	F	L	Q	F	H	F	A	H	V	G	N	L	L	W	T	T	H	P	C	K	P	4					
CA2	N	D	P	V	H	S	F	L	Q	F	H	F	A	H	V	G	K	L	L	W	T	T	P	C	K	P	K	4					
CA3	K	D	P	L	N	T	C	L	Q	F	H	F	A	H	V	G	K	M	L	W	T	T	P	C	K	P	K	7					
CA13	N	D	P	L	H	S	F	L	Q	F	H	F	A	H	V	G	N	L	T	W	T	T	V	P	S	K	P	K	3				

Figure S1. Phylogeny of α -CAs inferred from a maximum-likelihood analysis performed with RaxML version 8.0 with 1000 iterations. Bootstrap values are indicated at the nodes. Species are collapsed within a larger taxonomical grouping.



