

**Table S1. Ion Channels Used in the Phylogenetic Analysis, Related to Figure 4**

Organism	Phylum	Channel	Alternative names	GenBank accession
<i>Aiptasia pallida</i> (sea anemone)	Cnidaria	ApNa <sub>v</sub> 1		AAB96953
<i>Anopheles gambiae</i> (African malaria mosquito)	Arthropoda	AgCa <sub>v</sub> 1	-	ABQ88369
<i>Aplysia californica</i> (sea hare)	Mollusca	AcNa <sub>v</sub> 1	SCAP1	AAC47457
<i>Blatella germanica</i> (German cockroach)	Arthropoda	BgNa <sub>v</sub> 1	Para	AAC47483
		BgNa <sub>v</sub> 2	BSC1	AAK01090
<i>Branchiostoma floridae</i> (lancelet)	Cephalochordata	BfNa <sub>v</sub> 1	-	HQ877456
		BfNa <sub>v</sub> 2	-	HQ877455
<i>Cancer borealis</i> (Jonah crab)	Arthropoda	CbNa <sub>v</sub> 1	-	ABL10360
<i>Capitella teleta</i> (polychete worm)	Annelida	CtNa <sub>v</sub> 1	-	None, JGI protein ID 210954
		CtNa <sub>v</sub> 2	-	None, JGI protein ID 134859
<i>Clytia hemisphaerica</i> (hydrozoan jellyfish)	Cnidaria	ChNa <sub>v</sub> 2.1	-	JQ066821
		Ch Na <sub>v</sub> 2.5	-	JQ066822
<i>Cyanea capillata</i> (lion mane's jellyfish)	Cnidaria	CcNa <sub>v</sub> 2.5	-	AAA75572
		CcCa <sub>v</sub> 1	-	AAC63050
<i>Danio rerio</i> (zebrafish)	Chordata	DrNa <sub>v</sub> 1.6	-	AAG18440
<i>Drosophila melanogaster</i> (fruit fly)	Arthropoda	DmNa <sub>v</sub> 1	Para	AAB59195
		DmNa <sub>v</sub> 2	DSC1, 60E	Q9W0Y8
		DmCa <sub>v</sub> 1	-	AAA81883
<i>Halocynthia roretzi</i> (ascidian)	Chordata	HrNa <sub>v</sub> 1	TuNa1	BAA04133
		HrNa <sub>v</sub> 2	TuNa2	BAA95896
		HrCa <sub>v</sub> 1	-	BAA34927
<i>Homo sapiens</i> (human)	Chordata	hNa <sub>v</sub> 1.1	SCN1A, Brain	P35498

			type I channel	
		hNa <sub>v</sub> 1.2	SCN2A, Brain type II channel	Q99250
		hNa <sub>v</sub> 1.3	SCN3A, Brain type III channel	Q9NY46
		hNa <sub>v</sub> 1.4	SCN4A, Skeletal muscle type IV channel	AAO83647
		hNa <sub>v</sub> 1.5	SCN5A, Cardiac muscle type V	Q14524
		hNa <sub>v</sub> 1.6	SCN8A	Q9UQD0
		hNa <sub>v</sub> 1.7	SCN9A, Peripheral sodium channel 1, PN1	Q15858
		hNa <sub>v</sub> 1.8	SCN10A	Q9Y5Y9
		hCa <sub>v</sub> 1.1	CAC1S, L-type calcium channel	Q13698
<i>Hydra magnipapillata</i>	Cnidaria	HmNa <sub>v</sub> 2.1	-	JQ066819
		HmNa <sub>v</sub> 2.5	-	JQ066820
<i>Loligo bleekeri</i> (squid)	Mollusca	Lb Na <sub>v</sub> 2	-	BAA03398
<i>Loligo opalescens</i> (squid)	Mollusca	LbNa <sub>v</sub> 2	-	AAA16202
<i>Mnemiopsis leidy</i> (comb jelly)	Ctenophora	MI Na <sub>v</sub> 2a	-	AEF59085
		MI Na <sub>v</sub> 2b	-	AEF59086
<i>Monosiga brevicollis</i>	Opisthokonta	MbNa <sub>v</sub> 2		HQ877454
<i>Mus musculus</i> (mouse)	Chordata	Nax	NaG, SCN6A, SCN7A	AAA66192
<i>Nematostella vectensis</i>	Cnidaria	NvNa <sub>v</sub> 2.1	-	HQ877457

(starlet sea anemone)		NvNa <sub>v</sub> 2.2	-	HQ877458
		NvNa <sub>v</sub> 2.3	-	HQ877459
		NvNa <sub>v</sub> 2.4	-	HQ877460
		NvNa <sub>v</sub> 2.5	-	HQ877461
		NvCa <sub>v</sub> 1	-	XP_001639054
<i>Polyorchis penicillatus</i> (hydrozoan jellyfish)	Cnidaria	PpNa <sub>v</sub> 2.5	-	AAC38974
<i>Rattus norvegicus</i> (rat)	Chordata	rNa <sub>v</sub> 1.4	SCN4A, Skeletal muscle type IV channel	P15390
		rCa <sub>v</sub> 1.2	CAC1S, L- type calcium channel	P22002
<i>Strongylocentrotus purpuratus</i> (purple sea urchin)	Echinodermata	SpuNa <sub>v</sub> 2	-	XP_001189610
<i>Stylophora pistillata</i> (stony coral)	Cnidaria	SpiCa <sub>v</sub> 1	-	AAD11470
<i>Thalassiosira pseudonana</i> (diatom)	Bacillariophyta	TpCa <sub>v</sub>	-	XP_002289136
<i>Thecamonas trahens</i>	Apusozoa	TtNa <sub>v</sub> 2	TtrNa <sub>v</sub>	None, Broad Institute protein model
<i>Trichoplax adhaerens</i>	Placozoa	TaNa <sub>v</sub> 2a	-	HQ877452
<i>Trichoplax adhaerens</i>	Placozoa	TaNa <sub>v</sub> 2b	-	HQ877453
<i>Varroa destructor</i>	Arthropoda	VdNa <sub>v</sub> 1	-	AAP13992

**Table S1.** Ion channels used in the phylogenetic analysis. Most channels are derived from cloned transcripts that were deposited in Genbank. Channel transcripts cloned in the present study appear on gray background. Channel obtained from public genome sequencing projects supported by EST data appear in red. The channel protein models of the annelid *Capitella teleta* are from the genome sequencing project performed by the Joint Genome Institute (JGI) available at <http://genome.jgi.doe.gov/Capca1/Capca1.home.html>. The

channel protein model of *Thecamonas trahens* is from the Origins of Multicellularity project by the Broad Institute available at [http://broadinstitute.org/annotation/genome/multicellularity\\_project](http://broadinstitute.org/annotation/genome/multicellularity_project).