

Additional File 1. Bilaterians searched for *Dact* genes.

Abbreviations of species names are indicated on the left; others: o, order; so, superorder. *Dact* EST sequences are represented by a tick; for genomic sequences, the genomic localization is indicated, following the nomenclature of the 2012 edition of the Ensemble (<http://www.ensembl.org/index.html>), NCBI (<http://www.ncbi.nlm.nih.gov/>), elephant shark genome project (<http://esharkgenome.imcb.a-star.edu.sg/>) and Japanese lamprey genome project (<http://jlampreygenome.imcb.a-star.edu.sg/>) databases. For the Anole lizard, *Xenopus tropicalis*, the coelacanth, and the sea lamprey, in addition the nomenclature used in the 2008 edition of Ensemble is shown.

Deuterostomes									
	abbreviation	species	common name	Dact1	Dact2	Dact3 or dact3a	dact3b	Dact4	comments
classification									
Phylum: Chordata									
Subphylum: Vertebrata									
Infraphylum: Gnathostomata									
Superclass: Osteichthyes									
Class: Sarcopterygii									
Subclass: Tetrapodomorpha									
Amniota; Synapsids -	Hs	<i>Homo sapiens</i>	human	chr14✓	chr6✓	chr19✓		(locus chr11)	<i>Dact4</i> locus is identifiable but lacks a <i>Dact</i> gene
Infraclass: Mammals;	Mm	<i>Mus musculus</i>	mouse	chr12✓	chr17✓	chr7✓		(locus chr19)	
Placentalia	Bt	<i>Bos Taurus</i>	cattle	chr10	chr9	chr18		(locus chr29)	
	Cf	<i>Canis familiaris</i>	dog	chr8	chr1	chr1		(locus chr18)	
	La	<i>Loxodonta africana</i>	African elephant	s2755		s4446			
Infraclass: Mammals;	Md	<i>Monodelphis domestica</i>	opossum	chr1	chr2	chr4		(locus chr5)	
Marsupials	Oa	<i>Ornithorhynchus anatinus</i>	platypus	c214, 26117; c164635 (ex4)	c20694, c22096	c69853 (ex1), c289003 (ex4)			
Infraclass: Mammals;									
Monotremes									
Amniota; Diapsids;	Gg	<i>Gallus gallus</i>	chicken	chr5✓	chr3✓				<i>Dact3</i> and 4 loci not conserved
Archosauromorpha;	Mg	<i>Meleagris gallopavo</i>	turkey	chr5	chr2				
Infraclass: Aves	Tg	<i>Taeniopygia guttata</i>	zebrafinch	chr5	chr3				
	Ap	<i>Anas platyrhynchos</i>	duck	s626	s530				
	Mu	<i>Melopsittacus undulatus</i>	budgerigar	JH556483	JH556595				
Amniota; Diapsids;	Ac	<i>Anolis carolinensis</i>	Anole lizard	GL343459 (formerly s5, s29) and chr2	chr1 (formerly s549)	GL343568 (formerly s1268)		GL343928 (formerly s71, s1206)	
Infraclass: Lepidosauromorpha	Pym	<i>Python molurus</i>	Burmese python	c26513843 c26939700	c24659227 c26563497 c26659464 c26922077	c26582260		c26563039 c26110406 c25444438 c24942884	
Amniota; Anapsids;	Cpb	<i>Chrysemys picta bellii</i>	Western painted turtle	JH584658	JH584758	JH584472		JH584543	
Infraclass: Testudines (turtles and tortoises)	Ps	<i>Pelodiscus sinensis</i>	Chinese soft shield turtle	JH212506	JH208515	JH209944		JH206113	
Infraclass: Amphibia;	Xt	<i>Xenopus tropicalis</i>	Western clawed frog	GL172704 (formerly s68) ✓	(locus GL172638)	GL172667 (formerly s31) ✓		(locus GL173034, formerly s398)	Remnants of <i>Dact2</i> and 4 loci identifiable
Anura	XI	<i>Xenopus laevis</i>	African clawed frog	1-a, 1-b ✓		3-a, 3-b ✓			
	Rc	<i>Rana catesbeiana</i>		✓					
Subclass: Dipnoi (lungfishes)	Pa	<i>Protopterus aethiopicus</i>	African or marbled lungfish		✓				
Subclass: Coelacanthimorpha	Lc	<i>Latimeria chalumnae</i>	Coelacanth	JH126564 (formerly c002619-17)	JH129292 (formerly c230613-11)	JH128225 (formerly c190000,13,18,22)		JH126593 (formerly c015702)	
Class: Actinopterygii									
Subclass: Neopterygii									
Infraclass: Holostei (bowfins, gars)	Lo	<i>Lepisosteus oculatus</i>	Spotted gar	LG7	LG16	LG2		LG28	Intronless <i>dact4r</i> on LG14
Infraclass: Teleostei									
o: Cypriniformes	Dr	<i>Danio rerio</i>	zebrafish	chr17✓	chr13✓	chr18✓	chr10✓	chr14✓	Intronless <i>dact4r</i> on chr24 ✓
	Pp	<i>Pimephales promelas</i>	Fathead minnow	✓		✓		✓	
o: Siluriformes	Ip	<i>Ictalurus punctatus</i>	Channel catfish	✓					

so: Protacanthopterygii									
o: Salmoniformes	Ssa	<i>Salmo salar</i>	Altantic salmon	✓					
	Ot	<i>Oncorhynchus tshawytscha</i>	Chinook salmon					✓	
	Omy	<i>Oncorhynchus mykiss</i>	Rainbow trout	✓			✓		
o: Osmeriformes	Omo	<i>Osmerus mordax</i>	Rainbow smelt		✓				
so: Paracanthopterygii									
o: Gadiformes=Anacanthini	Gmo	<i>Gadus morhua</i>	Atlantic cod	s909/ c572298, c143687,	s609/ c360826	s3150	s4368	s3610; partial sequences on c76676	
so: Acanthopterygii									
o: Tetraodontiformes	Tr	<i>Takifugu rubripes</i>	Fugu	(locus on s136)	s53	s165	s455	c187	dact1 locus present, but no dact1 gene
	Tn	<i>Tetraodon nigroviridis</i>	Green spotted pufferfish	(locus on s7089)	chr17	chrUn/s14702	chr7	chr1	
o: Gasterosteiformes	Ga	<i>Gasterosteus aculeatus</i>	three-spined stickleback	grXV✓	grVI	grIV✓	(locus on grVII)	grIV✓	Sequence gap at the position of the dact3b gene
o: Beloniformes	Ol	<i>Oryzias latipes</i>	Medaka	chr22✓	chr15✓	chr13	chr14✓	chr10✓	
o: Perciformes	Pf	<i>Perca flavescens</i>	Yellow perch					✓	
	Di	<i>Dicentrarchus labrax</i>	European seabass					✓	
	On	<i>Oreochromis niloticus</i>	Nile Tilapia	GL831368	GL831145	GL831147	GL831515✓	GL831165	
	Hb	<i>Haplochromis burtoni</i>	African cichlid				✓		
	Sau	<i>Sparus aurata</i>	Gilthead seabream					✓	
o: Cyprinodontiformes	Xm	<i>Xiphophorus maculatus</i>	Southern platyfish	JH557083	JH556663	JH556880		AGAJ01047724 & JH556834	
Superclass Chondrichthyes									
Subclass: Elasmobranchii;									
Subdivision Selachii; so: Squalomorphi	Sac	<i>Squalus acanthias</i>	Spiny dogfish shark	✓					
Subdivision / so: Batoidea	Tc	<i>Torpedo californica</i>	Pacific electric ray		✓			✓	
	Le	<i>Leucoraja erinacea</i>	Little skate http://www.marinegenomics.org/node/27111					✓	
Subclass: Holocephali;	Cm	<i>Callorhinus milii</i>	Elephant shark	AAVX 014209261.1 01221604.1 01236760.1 01179725.1 01276763.1	AAVX 01057442.1 01184388.1	AAVX 01159308.1		AAVX 01316550.1	
Infraphylum: Gnathostomata									
Superclass: Cyclostomata									
Class: Hyperoartia/Petromyzontida	Pm	<i>Petromyzon marinus</i>	Sea lamprey	dactA (exons 3+4, GL476511/c36439); dactB (exons 3+4, c37220 and c20195); dactD (exon 4, c54804)				At least 3 genes	
	Lj	<i>Lenthenteron japonicum</i>	Japanese lamprey	dactA (KE993709); dactB (KE993739); dactC (exons 2-4, KE993726; similar exon 4 on KE999188/KE99520); dactD (exon 4, KE994909); orphan exon 1: APJL01152884/APJL01160608				At least 4 genes	
Phylum: Chordata									
Subphylum: Tunicata/Urochordata									
Class: Appendicularia	Oik	<i>Oikopleura dioica</i>							No significant hit
Class: Ascidiacea	Cii	<i>Ciona intestinalis</i>	Vase tunicate						
	Cis	<i>Ciona savignyi</i>							
Phylum: Chordata	Bfl	<i>Branchiostoma floridae</i>	Florida lancelet				s65 ✓		
Subphylum: Cephalochordata									
Class Leptocardii									
Phylum: Hemichordata	Sak	<i>Saccoglossus kowalevskii</i>	Acorn worm						No significant hit
Class: Enteropneusta									
Phylum: Echinodermata	Spu	<i>Strongylocentrotus purpuratus</i>	California purple sea urchin						No significant hit
Class: Echinoidea									

Protostomes - lophotrochozoa									
Phylum: Mollusca	Apc	<i>Aplysia californica</i>	California sea slug or California sea hare						No significant hit
Protostomes - ecdysozoa									
Phylum: Arthropoda									
Class: Insecta	Drm	<i>Drosophila melanogaster</i>	Fruit fly						No significant hit
	Trc	<i>Tribolium castaneum</i>	Red flour beetle						
	Bom	<i>Bombyx mori</i>	domesticated silkworm						

Phylum: Nematoda	Cel	<i>C.elegans</i>						No significant hit
	Cbr	<i>C.briggsae</i>						
	Loa	<i>Loa loa</i>	African eye worm					