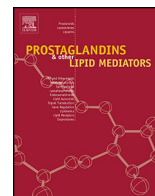




Prostaglandins and Other Lipid Mediators



Review

Why do a wide variety of animals retain multiple isoforms of cyclooxygenase?



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ABSTRACT

Cyclooxygenase (COX) has been cloned from the phyla Cnidaria, Mollusca, Arthropoda, and Chordata of the animal kingdom. Many organisms have multiple COX isoforms that have arisen from gene duplication. It is not well understood why there are multiple COX isoforms in the same organism, or when duplication of the COX gene occurred. Here, we summarize the current knowledge of the evolutionary history of COX in the animal kingdom and discuss the reasons why the multiple COX system has been retained so widely. The phylogenetic analysis suggests that all COX genes in animals may descend from a common ancestor and that the duplication of an ancestral COX gene might occur within each lineage after the divergence of the animal. In most instances, the expressions of multiple COX isoforms are separately regulated and these isoforms play different and important pathophysiological roles in each organism. This may be the reason why multiple COX isoforms are widely retained.

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Contents

1. Introduction	15
2. Characteristics of COX and its products	15
3. The evolutionary history of COX in the animal kingdom	16
3.1. Placental mammals	16
3.2. Marsupials and egg-laying mammals	16
3.3. Birds, reptiles, amphibians, and lobe-finned fishes	17
3.4. Ray-finned fishes	17
3.5. Cartilaginous fishes and jawless vertebrates	18
3.6. Protochordates	18
3.7. Mollusks and arthropods	19
3.8. Cnidarians	19
4. Why are multiple COX genes widely retained in the animal kingdom?	19
5. Conclusion	20
References	20

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1. Introduction

Oxygenated fatty acids are collectively named oxylipins. Many oxylipins have biological activities. One of the most pathophysiologically important groups of oxylipins in animals is the eicosanoids, which include prostanoids, leukotrienes, lipoxins, and hydroxyeicosatetraenoic acids [1]. Cyclooxygenase (COX), which is also known as prostaglandin (PG) H synthase or PG endoperoxide synthase (E.C. 1.14.99.1), catalyzes the initial step of prostanoid biosynthesis from polyunsaturated fatty acids. COX belongs to the heme-dependent myeloperoxidase superfamily and is a distant homolog of the pathogen-induced oxygenases and linoleate diol synthase of plants and fungi, respectively [2]. COX or COX-like genes have been found in animals, fungi, plants, unicellular organisms, and even in microbes [3,4].

All eutherians (also called placental mammals) that have been cloned so far have two isoforms of COX: COX-1 and -2. Moreover, multiple COX isoforms have been found in the many organisms, including invertebrates and vertebrates, which are known to express COX. It is not well understood when the evolutionary duplication of the COX gene occurred and why there are multiple isoforms of COX in a single organism. Here, we summarize the current knowledge of the evolutionary history of COX in the animal kingdom and discuss the reasons why the multiple COX system has been so widely retained (Table 1).

2. Characteristics of COX and its products

Prostanoids are produced when polyunsaturated fatty acids are released from the cellular membrane by phospholipase A₂ and then metabolized by COX and specific terminal prostanoid synthases. In mammals, arachidonic acid (5,8,11,14-eicosatetraenoic acid) is the major substrate for COX. This fatty acid is abundant in cell membranes, where it is esterified into glycerophospholipids. Upon stimulation of a target cell by relevant stimuli, phospholipase A₂ is activated and arachidonic acid is released from the phospholipids. COX has some unique characteristics. First, COX is a monotopic membrane-bound protein [5]; hence, it can only integrate into a single leaflet of the lipid bilayer. COX is largely located on the luminal side of the endoplasmic reticulum membrane and the nuclear envelope [6]. After post-translational processing in the endoplasmic reticulum, the mature protein has an apparent molecular mass of 67–72 kDa and exists as a homodimer [7]. Additionally, COX is bifunctional: it catalyzes both a *bis*-oxygenase reaction (habitually referred to as the cyclooxygenase reaction), in which arachidonic acid is converted to PGG₂, and a peroxidase reaction, in which PGG₂ undergoes a 2-electron reduction to PGH₂ [8]. The *bis*-oxygenase and peroxidase sites are physically and functionally distinct within the COX protein [8]. Pharmacologists have long been interested in COX because it is the primary target of non-steroidal anti-inflammatory drugs. These drugs, including aspirin and indomethacin, selectively inhibit the *bis*-oxygenase reaction. With the exception of aspirin, which covalently modifies the *bis*-oxygenase site by acetylating Ser-530 (in sheep COX-1) [9,10], these drugs compete directly with the substrate, arachidonic acid, for binding to the *bis*-oxygenase site and inhibit its activity, but they have little effect on peroxidase activity [11].

The product of the COX-catalyzed reaction, PGH₂, is the intermediate from which a variety of primary prostanoids, including PGD₂, PGE₂, PGF_{2α}, PGI₂, and thromboxane (TX) A₂, are produced by terminal prostanoid synthases through reactions of isomerization, oxidation, or reduction. Terminal prostanoid synthases are a heterogeneous family of enzymes that are only expressed in specific cell types. In mammals, single types of PGI synthase [12] and TXA synthase [13], two types of PGD synthase [14],

Table 1

The protein IDs of COX isoforms mentioned in the text.^a

Placental mammals

Human (*Homo sapiens*)

COX-1, NP.000953.2; COX-2, NP.000954.1

Sheep (*Ovis aries*)

COX-1, NP.001009476.1; COX-2, NP.001009432.1

Marsupials and egg-laying mammals

Tasmanian devil (*Sarcophilus harrisii*)

COX-1, ENSSHAP00000001746; COX-2, ENSSHAP00000019049

Tammar wallaby (*Macropus eugenii*)

COX-1, ENSMEUP00000006100; COX-2, ENSMEUP00000000284

The opossum (*Monodelphis domestica*)

COX-1a, XP.001370514.1; COX-1b1, XP.001370542.2; COX-1b2, XP.001370595.1;

COX-2-like, XP.001375945.1

Platypus (*Ornithorhynchus anatinus*)

COX-1-like, XP.001512281.2; COX-2-like, XP.001516281.2

Birds, reptiles, amphibians, and lobe-finned fishes

Turkey (*Meleagris gallopavo*)

COX-1-like, XP.003211401.1; COX-2-like, XP.003208549.1

Chicken or red jungle fowle (*Gallus gallus*)

COX-1, XP.425326.4; COX-2, NP.001161190.1

Zebra finch (*Taeniopygia guttata*)

COX-1, ENSTGUP00000007069; COX-2, ENSTGUP00000003584

Green Anole (*Anolis carolinensis*)

COX-1-like, XP.003230294.1; COX-2-like, XP.003223472.1

Chinese softshell turtle (*Pelodiscus sinensis*)

COX-1, ENSPSIP00000005203; COX-2, ENSPSIP00000011001

Western clawed frog (*Xenopus (Silurana) tropicalis*)

COX-1, NP.001258184.1; COX-2, NP.001025697.1

African clawed frog (*Xenopus laevis*)

COX-1, NP.001091389.1; COX-2, NP.001086946.1

Coelacanth (*Latimeria chalumnae*)

COX-1, ENSLACP00000001481; COX-2, ENSLACP000000021975

Ray-finned fishes

Platyfish (*Xiphophorus maculatus*)

COX-1-1, ENSXMAP000000003268; COX-1-2, ENSXMAP000000005187;

COX-2b, ENSXMAP000000002436

Green spotted puffer (*Tetraodon nigroviridis*)

COX-1a, GSTENP00029010001; COX-1b, GSTENP00012293001; COX-2b,

GSTENP00033631001

Pufferfish (*Takifugu rubripes*)

COX-1a, ENSTRUP000000042693; COX-1b, ENSTRUP000000046266; COX-2b,

ENSTRUP00000010041

Atlantic croaker (*Micropogonias undulatus*)

COX-1, BAF52621.1; COX-2, BAF52620.1

Nile tilapia (*Oreochromis niloticus*)

COX-1-like, XP.003451513.1; COX-1-like, XP.003454212.1; COX-2-like, XP.003445100.2

Longhorn sculpin (*Myoxocephalus octodecemspinosus*)

COX-1a, ACO34913.1; COX-1b, ACH73272.1; COX-2b, ACH73267.1

Three-spined stickleback (*Gasterosteus aculeatus*)

COX-1a, ENSGACP00000018397; COX-1b, ENSGACP000000021267; COX-2b, ENSGACP000000009851

Japanese medaka (*Oryzias latipes*)

COX-1a, ENSORLP00000000625; COX-1b, ENSORLP000000006426; COX-2b, ENSORLP00000011863

Mummichog or Euryhaline Killifish (*Fundulus heteroclitus*)

COX-1a, ACH73266.1; COX-1b, ACH73265.1; COX-2, AAS21313.2

Atlantic cod (*Gadus morhua*)

COX-1, ENSGMOP00000015572; COX-2b, ENSGMOP00000011635

The brook trout (*Salvelinus fontinalis*)

COX-1, AAF14529.1; COX-2, AAD45896.1

Rainbow trout (*Oncorhynchus mykiss*)

COX-1b, NP.001117833.1; COX-2, NP.001117820.1; COX-2b, NP.001118139.1

Zebrafish (*Danio rerio*)

COX-1b, NP.705942.1; COX-2a, NP.705943.1; COX-2b, NP.001020675.1

Cartilaginous fishes and jawless vertebrates

Spiny dogfish shark (*Squalus acanthias*)

sCOX, AAL37727.1

Epaullette shark (*Hemiscyllium ocellatum*)

Partial COX-2, ADD59908.1

Sea lamprey (*Petromyzon marinus*)

ICOX, ACH73269.1; COX-1, ENSPMAP00000001004 (gene ID, ENSPMAG00000000916); COX-2a, ENSPMAP000000003793 (gene ID, ENSPMAG000000003485)

Atrantac hagfish (*Myxine glutinosa*)

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