The paradoxical patent ductus arteriosus

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The ductus arteriosus (DA) is a vessel whose patency is required for fetal survival but is incompatible with postnatal life. Because of developmental insufficiency, the DA in preterm infants often fails to close in a condition known as patent DA (PDA). Although COX inhibitors can be used to close the DA by lowering circulating prostaglandin levels, their effectiveness is correlated with birth weight, and severely premature infants often require surgical repair. Paradoxically, targeted deletion of COX pathway components in mice results in PDA. In this issue of the JCI, Yokoyama et al. describe dual roles for prostaglandins in DA development and closure, offering new insights into the mechanism of negative effects of COX inhibitors that may influence the treatment of severely premature infants with PDA and lead to improvement of their outcomes (see the related article beginning on page 3026).

Nonstandard abbreviations used: DA, ductus arteriosus; HA, hyaluronic acid; HAS2, HA synthase 2; PDA, patent DA.

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The ductus arteriosus (DA) is a fetal vessel that bridges the pulmonary and systemic circulation during gestation, when oxygenation of fetal blood occurs in the placenta (Figure 1). In utero, circulating prostaglandins, formed by the action of COX enzymes, exert their vasodilatory effect on the DA to maintain its patency (1). Activation of the G protein–coupled receptor EP4 by PGE2, the primary prostaglandin regulating ductal tone, leads to DA smooth muscle relaxation (1). During late gestation, in preparation for birth, SMCs within the DA migrate toward the endothelial lining, forming mounds known as intimal cushions (2). A sharp decline in prostaglandin levels at birth results in constriction of the DA, bringing the intimal cushions into contact and occluding the DA lumen (1). Failure of DA closure, called patent DA (PDA), is the third most common congenital heart defect, and its incidence is increased in preterm infants (3). Fortunately, pharmacological inhibition of the COXs is an effective means of closing
The paradox of PDA

Given the known role of prostaglandins in maintaining DA patency, a logical corollary would be that mice with targeted deletions of genes that negatively affect prostaglandin synthesis or signaling would result in premature closure of the DA. Surprisingly, mice lacking the DA-enriched prostaglandin receptor EP4, or mice with combined deficiency of the COX enzymes, COX-1 and COX-2, were found to have PDA (6, 7). How is it that depleting a signaling pathway in mice has the same effect as activating that pathway in humans?

Yokoyama et al. shed light on this question in their study published in this issue of the JCI (8). They establish that PGE₂, acting via its receptor, EP4, has an additional prenatal role in the maturation of the fetal DA, preparing it for closure at birth by promoting intimal cushion formation (Figure 2). Using agonists and antagonists of the EP4 receptor, the authors show that EP4 controls DA smooth muscle migration to form intimal cushions while stimulating synthesis and secretion of hyaluronic acid (HA; also known as hyaluronan), an extracellular matrix component through which the SMCs traverse. With elegant use of siRNA-mediated knockdown of HA synthase 2 (HAS2), as well as HA depletion from EP4-activated cultures, the authors confirm that PGE-induced SMC migration occurs through EP4 stimulation of HA production.

These findings offer a likely explanation for why mice lacking prostaglandin signaling in the DA do not experience premature closure of the fetal vessel. EP4-null mice fail to form intimal cushions, so the vasodilatory effects of prostaglandins are likely not required for maintenance of DA patency before birth. Instead, patency persists postnatally in the absence of prostaglandin signaling because, without intimal cushion formation, occlusion of the DA lumen cannot occur.

While we believe that the study by Yokoyama et al. (8) is the first to uncover a role for prostaglandins in intimal cushion formation, the idea that DA remodeling is more important than vasoconstriction for DA closure is not new. The DA of mice lacking smooth muscle myosin, which is required for contraction of DA smooth muscle, also closes appropriately after birth (9), demonstrating that functional closure can occur without vessel wall contraction. Interestingly, comparison of the effects of genetic COX deficiency and administration of COX inhibitors at varying points of mouse gestation also revealed that prostaglandins have a novel role in the development of the DA, beyond their direct vasodilatory effect (10).

Figure 1

DA during fetal-neonatal circulatory transition. Left: During fetal development, the DA shunts blood away from the lungs and directly into the systemic circulation via the aorta. Right: At birth, as PGE levels fall, SMCs in the DA constrict, bringing the intimal cushions of the DA into contact and occluding the DA lumen, so that blood flows to the neonatal lungs. This remodeling process supports the change from a fetal circulation, where blood is oxygenated in the placenta, to a neonatal circulation, where gas exchange occurs in the lungs. Blue represents oxygen-poor blood; red represents oxygen-rich blood. RA, right atrium; LA, left atrium.
In addition to explaining the paradox of PDA in mice deficient in prostaglandin signaling, the elucidation by Yokoyama et al. (8) of a developmental role for prostaglandins during DA maturation may also add to our understanding of important clinical findings that have yet gone unexplained. In spite of their vasoconstrictive effect on the DA at birth, use of COX inhibitors by pregnant mothers increases the incidence of PDA in newborns (11, 12). If prostaglandin signaling is, indeed, required for intimal cushion formation in humans as in rodents, then infants who develop under conditions of insufficient PGE, as in the case of maternal use of COX inhibitors, may be born with an underdeveloped DA that remains patent in spite of PGE withdrawal. The current results also imply that, while DA smooth muscle development is not as advanced in premature infants as in their full-term counterparts, DA smooth muscle immaturity may not be the primary cause for the lack of DA constriction in response to COX inhibitors, as previously thought. Instead, a lack of intimal cushions, whose further development is blocked by the delivery of COX inhibitors, may preclude DA closure in preterm infants.

Implications for pharmacological intervention for PDA

While COX inhibitors offer an effective means to close the DA in near-term infants, for severely premature babies, COX inhibitors may exacerbate the problem by preventing the formation of intimal cushions required for closure. Yokoyama et al. (8) suggest that HAS activation, combined with prostaglandin blockade, might be an effective means to treat PDA in severely premature infants. Certainly their results suggest that, upon appropriate delivery of directed HAS stimulation, this may be a successful tactic in rodents, and future studies will determine whether a combined therapy is sufficient to promote DA maturation and closure in a timely manner.

It is important to note that, while the underlying molecular pathways controlling DA development and closure are strikingly similar in rodents and larger mammals, there are significant differences in their developmental physiology and the timing of critical closure events. For instance, while functional closure of the DA happens quite rapidly in rodents, occurring within the first few hours after birth (13), normal closure in newborn humans takes place over several days (14). More relevant to the early effects of prostaglandins, intimal cushion formation occurs much earlier and is far more pronounced in larger mammals than in small rodents (15). It has been hypothesized that, due to their increased luminal diameter, larger mammals require highly developed intimal cushions for contraction of the surrounding smooth muscle to effectively close the lumen (15). So while mice, rats, and even rabbits offer amenable models for genetic and initial pharmacological studies of DA development and closure, other large animals, such as lambs, pigs, or dogs, must be used to more directly model the physiology and pathophysiology of the human DA. Given the impact and potential clinical application of the study by Yokoyama et al., further studies of this kind will no doubt be undertaken in larger model organisms, hopefully offering answers that can be directly applicable to neonatal medicine and improve the outcomes of the multitude of low-birth-weight babies that struggle with this disease.

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Role for IKK2 in muscle: waste not, want not

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Activation of transcription factor NF-κB, the major regulator of the inflammatory response, depends on the inhibitor of NF-κB kinase (IKK) complex, which is composed of 2 catalytic subunits, IKK1 and IKK2 (also known as IKKα and IKKβ), and a regulatory subunit, IKKγ (also known as NEMO). In this issue of the JCI, Mourkioti et al. show that muscle-specific disruption in mice of the gene encoding IKK2 prevents NF-κB activation in response to denervation or toxin-induced injury (see the related article beginning on page 2945). Importantly, this genetic manipulation prevents muscle wasting, thereby providing strong evidence in support of a major pathogenic role for inflammation in a variety of muscular dystrophies characterized by progressive muscle fiber degeneration.

Transcription factor NF-κB is one of the major activators of the inflammatory response (1). In most cells, the majority of the NF-κB pool resides in the cytoplasm but undergoes rapid nuclear translocation upon activation of innate immune receptors or exposure to proinflammatory cytokines, such as TNF-α or IL-1. Nuclear translocation of NF-κB or its activation depends on degradation of specific inhibitory proteins called inhibitors of NF-κB (IkBs) via a process that requires IkB phosphorylation and polyubiquitination (2). Phosphorylation of IkBs is mediated by a specialized protein kinase complex, the IkB kinase (IKK) complex, which is composed of 2 catalytic subunits, IKK1 (also known as IKKα) and IKK2 (also known as IKKβ), and a regulatory subunit called NEMO or IKKγ (3). Gene disruption experiments have established that in most cell types, in response to most stimuli, IkB degradation and NF-κB activation are mainly dependent on IKK2 (4). Hence, targeted disruption of the IKK2 gene has been used to demonstrate its involvement in a variety of inflammatory disorders, such as multiorgan dysfunction (5), colitis-associated cancer (6), and even obesity-induced insulin resistance (7). Now, in this issue of the JCI, Mourkioti et al. (8) report on the addition of muscle degeneration and atrophy to the list of pathologies that are remarkably ameliorated upon targeted deletion of IKK2. This suggests that muscle degenerative diseases may have a common inflammatory component and thus may respond to antiinflammatory therapy. A role for inflammation in muscular dystrophies? Muscle degeneration and atrophy is a rather common pathology associated with a variety of diseases, including those that specifically target the muscle, for instance muscular dystrophies (9), or those that target other organs, such as cancer and immune disorders (10, 11). Even old age and immobilization can result in muscle wasting. Yet apart from cachexia, in which TNF-α plays an important role, and specific muscle inflammation/myositis, muscular dystrophies in particular and muscle atrophy in general have not been universally thought of as inflammatory diseases. Yet the new work of Mourkioti et al. (8) as well as previous work by Shoelson’s group that demonstrated severe muscle wasting following the constitutive activation of NF-κB in muscle cells (12) clearly demonstrate the involvement of the IKK/NF-κB signaling system in different types of muscle degeneration. These data therefore strongly suggest that muscular dystrophies and atrophies should also be considered inflammatory diseases and raise the prospects of novel therapies that target IKK2 or other steps in the NF-κB activation pathway.

Mourkioti et al. (8) used a mouse strain homozygous for a conditional Ikk2 allele (a so-called floxed allele) that can be deleted upon expression of Cre-recombinase to specifically inactivate IKK2 in


Nonstandard abbreviations used: IkB, inhibitor of NF-κB; IKK, IkB kinase; MufR1, muscle ring finger protein 1.

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