

REVIEW

Intermedin (adrenomedullin-2): a novel counter-regulatory peptide in the cardiovascular and renal systems

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Intermedin (IMD) is a novel peptide related to calcitonin gene-related peptide (CGRP) and adrenomedullin (AM). Proteolytic processing of a larger precursor yields a series of biologically active C-terminal fragments, IMD_{1-53} , IMD_{1-47} and IMD_{8-47} . IMD shares a family of receptors with AM and CGRP composed of a calcitonin-receptor like receptor (CALCRL) associated with one of three receptor activity modifying proteins (RAMP). Compared to CGRP, IMD is less potent at $CGRP_1$ receptors but more potent at AM_1 receptors and AM_2 receptors; compared to AM, IMD is more potent at $CGRP_1$ receptors but less potent at AM_1 and AM_2 receptors. The cellular and tissue distribution of IMD overlaps in some aspects with that of CGRP and AM but is distinct from both. IMD is present in neonatal but absent or expressed sparsely, in adult heart and vasculature and present at low levels in plasma. The prominent localization of IMD in hypothalamus and pituitary and in kidney is consistent with a physiological role in the central and peripheral regulation of the circulation and water-electrolyte homeostasis. IMD is a potent systemic and pulmonary vasodilator, influences regional blood flow and augments cardiac contractility. IMD protects myocardium from the deleterious effects of oxidative stress associated with ischaemia-reperfusion injury and exerts an anti-growth effect directly on cardiomyocytes to oppose the influence of hypertrophic stimuli. The robust increase in expression of the peptide in hypertrophied and ischaemic myocardium indicates an important protective role for IMD as an endogenous counter-regulatory peptide in the heart.

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Abbreviations: AM, adrenomedullin; AMY, amylin; CGRP, calcitonin gene-related peptide; CL, calcitonin receptor-like receptor; CT, calcitonin; L-NAME, N^G -nitro-L-arginine-methyl-ester; RAMP, receptor activity-modifying protein; SHR, spontaneously hypertensive rat

Emergence of a peptide superfamily comprising calcitonin, calcitonin gene-related peptide, amylin and adrenomedullin and the discovery of intermedin

The regulatory influence of the peptide, calcitonin (CT), on calcium metabolism has long been recognized. In 1983, alternative tissue-specific processing of primary mRNA from the rat calcitonin gene (CT) was shown to generate two distinct single-chain polypeptides, namely calcitonin (rCT) comprising 32 amino acids and calcitonin gene-related peptide (r α CGRP) comprising 37 amino acids; CGRP is expressed predominantly in the nervous system and CT in parafollicular cells of the thyroid (Rosenfeld *et al.*, 1983).

Existence of a structurally similar peptide in humans was confirmed by isolation of h α CGRP from a medullary thyroid carcinoma (Morris *et al.*, 1984). A second CGRP/CT gene (β -gene), thought to have arisen by exon duplication, generates alternative β -forms of CGRP which differ from the α -forms by 1 and 3 amino acids in rats and humans, respectively. There is no evidence for expression of an alternative form of CT from this gene (Steenbergh *et al.*, 1985; Alevizaki *et al.*, 1986). Disparity in regional distribution of CGRP and its binding sites in the central nervous system (Kruger *et al.*, 1988) provided strong indication of the likely existence of additional member(s) of this peptide superfamily. The superfamily was enlarged with the subsequent discovery of amylin (AMY), a peptide identical in length to CGRP, which was extracted from amyloid deposits in pancreatic tissue of diabetics (Rink *et al.*, 1993), and adrenomedullin (AM), comprising 52 amino acids, extracted from human pheochromocytoma (Kitamura *et al.*, 1993). The superfamily expanded to varying degrees in different vertebrate lineages probably due to variation between these

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in regard to prevailing living conditions (Ogoshi *et al.*, 2003). Existence of multiple family members, capable of activating overlapping signalling mechanisms imparts enormous versatility and plasticity in responding to the environment. Members of this peptide superfamily play important roles in a variety of physiological processes including ionic balance, neurotransmission, glucose metabolism and cardiovascular and endocrine homeostasis.

Based on searches of genome sequences in humans and other mammals and non-vertebrates, Roh *et al.* (2004) identified a novel gene encoding a prepro peptide of 146–150 amino acids. Although its overall sequence possessed little similarity to other known proteins, the C-terminal region of this prepro-peptide encompassed a putative 47 amino-acid peptide sharing ~28% sequence identity with AM and <20% with CGRP and predicted to possess the common structural features characteristic of members of the superfamily. This novel peptide was named 'intermedin' (IMD) due to its abundant expression in the intermediate lobe of the anterior pituitary. In the same year, Takei *et al.* (2004b) examined mammalian complementary DNA libraries for potential mammalian homologues of five orthologues of AM (AM1–AM5) which they had identified in pufferfish. Using this approach, they also identified a 146–150 amino-acid mammalian prepro-peptide which yielded a mature peptide containing 47 amino acids; this displayed ~30% sequence homology to AM and >70% homology to the pufferfish AM2 orthologue; this novel mammalian peptide was named adrenomedullin-2 (AM2). Confirmation that the primary sequences of human prepro-AM2 and human prepro-IMD, each 148 amino acids, were identical and that the mature peptides elicited similar biological effects led to general acceptance that IMD and AM-2, discovered independently by the two research groups, were the same peptide.

The purpose of this article is to summarize progress made in regard to understanding the biological significance of IMD since the review published shortly after the discovery of this peptide (Chang *et al.*, 2004). The general distribution of IMD, the nature of its interaction with a common receptor signalling system shared with other members of this peptide superfamily, and emerging evidence of the unique physiological profile of IMD are addressed. Particular emphasis has been placed on appraising the effects of IMD in the heart and vasculature and in the kidney, and potential involvement in various cardiovascular pathologies. Detailed examination of the actions of AM and CGRP is beyond the scope of the current article; these have been reviewed recently elsewhere (for example, Brain and Grant, 2004; Ishimitsu *et al.*, 2006). However, where appropriate their actions have been compared and contrasted with those of IMD, to inform discussion of the physiological and pathophysiological relevance and unique contribution of the newest member of this peptide superfamily.

Synthesis and structure of intermedin

The 148 prepro-peptide encoded by a gene located on the distal arm of human chromosome 22q3 encompasses a

putative signal sequence of 20 amino acids at the N terminal. Pairs of basic amino acids are found at Arg100–Thr101 and Arg107–Val108 (Figure 1); proteolytic cleavage at these sites followed by cleavage of Tyr147–Gly148 and amidation of the C-terminal amino acid yields a long 47 amino-acid form of the peptide, IMD_{1–47} (prepro-IMD_{101–147}, IMDL) and a short 40 amino-acid peptide IMD_{8–47} (prepro-IMD_{108–147}, IMDS) (Roh *et al.*, 2004). Analysis of orthologous intermedins indicates that the position of the N-terminal dibasic cleavage site varies by several amino acids between species whereas the arginine residue seven amino acids downstream from this cleavage site in human IMD is conserved in all species, supporting an important physiological role for the shorter form (Chang *et al.*, 2004; Roh *et al.*, 2004). Identification of a further proteolytic cleavage site in rat prepro-IMD between Arg92–His93 indicates the existence of an additional larger peptide, prepro-IMD_{93–145} or rIMD_{1–53}, which also possesses biological activity (Yang *et al.*, 2005a; Ren *et al.*, 2006). It has been noted that similar pairs of basic amino acids at Lys93–Arg94 and Arg146–Arg147 demark mature AM_{1–52} within prepro-AM (Kitamura *et al.*, 1993) and that only the terminal 40 amino acids, AM_{13–52}, are absolutely required for biological activity (Chu *et al.*, 2000). Additional structurally distinct peptides are derived from the prepro-AM precursor and are also biologically active: an N-terminal peptide, prepro-AM_{22–41}, named pro-AM N-terminal peptide (Kitamura *et al.*, 1994), and a C-terminal peptide, prepro-AM_{153–185}, named adrenotensin (Gumusel *et al.*, 1996). A further peptide is demarked by prepro-AM_{45–92}; its physiological function, if any, remains unknown (Morgenthaler *et al.*, 2005). In contrast, the IMD gene is unlikely to encode additional active peptides since the putative prepro region is not conserved between species. IMD itself is highly conserved; mouse and rat IMD_{1–47} differ by one amino acid, while human IMD_{1–47} is 87% identical to rat and >60% to pufferfish (Roh *et al.*, 2004; Takei *et al.*, 2004b). Closer structural similarity of IMD to CGRP and AM than to AMY and CT indicates that CGRP, AM and IMD represent an evolutionary sub-branch of the superfamily (Chang *et al.*, 2004). IMD shares the structural features characteristic of the superfamily, namely an N-terminal intramolecular loop of six amino acids flanked by a disulphide bond, followed by a well-defined α -helical region leading into a predominantly disordered structure terminating in a C-terminal amide (Roh *et al.*, 2004). The length of the N-terminal sequence proximal to the intramolecular loop varies depending on the form of IMD, being greatest for IMD_{1–53}. It is highly probable that several biologically active forms of IMD are generated within the same species dependent on post-translational processing events occurring within the cell or after secretion.

Distribution of intermedin in mammalian tissues and plasma

Evidence obtained from transcriptional expression and/or immunohistochemical investigation in rodents and human, indicates that IMD is found at particularly high levels in kidney, within the gastrointestinal tract, especially the muscularis mucosa of stomach and in jejunum, in brain,

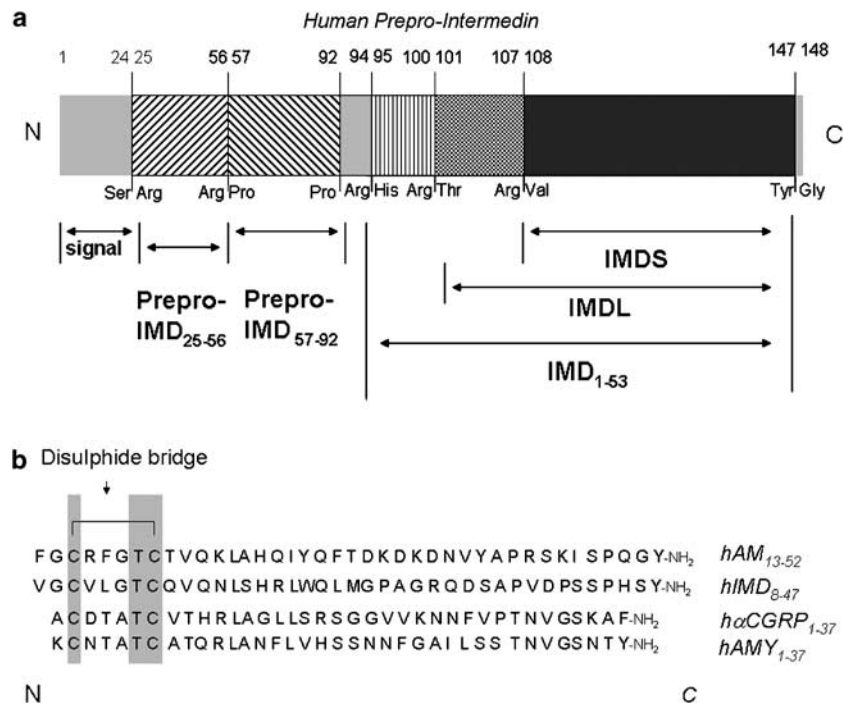


Figure 1 Schematic diagram (a) of human prepro-IMD showing the fragments derived from processing of the 148 amino-acid prepro-peptide precursor at the putative cleavage sites indicated; (b) comparison of the primary sequences of adrenomedullin_{13–52}, intermedin_{8–47} (IMDS), α CGRP_{1–37} and AMY_{1–37} in human; note the cysteine residues common to all of these peptides which contribute to the disulphide bridge giving rise to the characteristic intramolecular loop structure (adapted from Chang *et al.*, 2004; Roh *et al.*, 2004). AMY, amylin.

especially hypothalamus, where IMD is colocalized with arginine vasopressin, and intermediate and anterior lobes of pituitary (Roh *et al.*, 2004; Takei *et al.*, 2004b; Taylor *et al.*, 2005; Takahashi *et al.*, 2006), in skin (Kindt *et al.*, 2007) and in submaxillary gland, pancreas, lung, spleen, thymus and ovary, but not in testis or adrenal gland (Takei *et al.*, 2004b). Prepro-IMD is present, but much less abundantly so than prepro-AM in the heart of adult mouse (Takei *et al.*, 2004b) but is not detected in myocardium of adult rat (Pan *et al.*, 2005). In mouse, IMD is localized in heart to endothelial cells of coronary arteries and veins, and in kidney to endothelial cells of glomerular capillaries and the vasa recta running parallel to renal tubules (Takei *et al.*, 2004a). At variance with this observation, prepro-IMD is expressed in rat in neonatal cardiomyocytes (Pan *et al.*, 2005) and is detected, but present at substantially lower levels than prepro-AM, in adult cardiomyocytes (Zhao *et al.*, 2006) consistent with greater prominence of IMD during embryonic development of the myocardium. In human tissue obtained at autopsy from subjects without known cardiac or renal disease, Takahashi *et al.* (2006) reported that IMD is localized to myocardial cells but not endocardium, pericardial adipose tissue or vasculature of heart, and to renal tubular cells in both cortex and medulla, but not glomeruli or vasculature of kidney; the transcript is detected in kidney and cardiac left ventricle although expression levels vary considerably between individuals. It is unclear if such observations reflect genuine interspecies variation in the distribution of IMD, or arise as a consequence of disparity in the nature and duration of experimental protocols employed to detect the peptide or its transcript. Indeed, further

investigations by Morimoto *et al.* (2007) utilizing antiserum with higher sensitivity have indicated that IMD is localized, albeit less abundantly, in human pericardial adipocytes, vascular endothelial cells of pericardial veins and vascular smooth muscle of coronary arteries and renal arterioles in addition to cardiomyocytes and renal tubular cells. A general consensus is that the peptide is scarce in healthy adult myocardium. It is possible that the stress encountered leading up to death and the underlying pathology contributing to its cause influenced levels of IMD present in patients at autopsy.

IMD has been quantified in plasma of rat and human using primary antibodies which recognize a region common to all three bioactive fragments, IMD_{1–53}, IMD_{1–47} and IMD_{8–47}, but which do not crossreact with other members of this peptide superfamily. Plasma levels of IMD, like those of AM, are broadly similar or slightly greater in rat than human (Table 1), correlate positively with body mass index, particularly in females, but unlike those of AM are not increased with increasing age in adult human volunteers (Bell and Harbinson, unpublished observation). Plasma levels of IMD detected by radioimmunoassay are likely to be dependent on the sensitivity of the antibody used (Morimoto *et al.*, 2007). Reliable quantification of AM in plasma is also influenced by its biological activity, short half-life, existence of a binding protein in plasma which facilitates the presence of high concentrations of AM at receptor sites and influences degradation (Morgenthaler *et al.*, 2005). The extent to which such factors influence levels of IMD in plasma is not currently known. Neutral endopeptidase, a membrane-bound metalloproteinase, has

Table 1 Peptide levels in mammalian plasma

Peptide	Species	Plasma level pg ml^{-1} mean + s.e. (n)	Reference
IMD	Rat, adult SD	205.2 + 59.3 (5)	Taylor <i>et al.</i> (2006)
	Rat, adult SD	78.2 + 14.1 (10)	Bell <i>et al.</i> (2007)
	Rat, adult WKY	106.6 + 8.7 (8)	McDermott <i>et al.</i> (2007)
	Human, adult	7.2 + 0.6 (111)	Bell and Harbinson ^a (unpublished)
	Human, adult	126.0 + 9.1 (3)	Morimoto <i>et al.</i> (2007)
AM	Rat, adult SD	121.1 + 6.3 (10)	Bell <i>et al.</i> (2007)
	Rat, adult WKY	92.5 + 10.4 (8)	McDermott <i>et al.</i> (2007)
	Human, adult	27.1 + 1.5 (111)	Bell and Harbinson ^a (unpublished)
	Human, adult	37.0 + 1.5 (46)	Kato <i>et al.</i> (1999)
	Human, adult	39.7 + 8.3 (184)	Kato <i>et al.</i> (2002)
	Human, adult	13.7 + 6.1 (21)	Caliumi <i>et al.</i> (2004)
	Human, adult	8.7 + 2.1 (12)	Letizia <i>et al.</i> (2002)

AM, adrenomedullin; IMD, intermedin; WKY, Wistar–Kyoto rat.

^aMeasured in the same population of healthy individuals.

been implicated in degradation of AM (Rademaker *et al.*, 2002) and CGRP (Katayama *et al.*, 1991). It is likely that IMD is also a substrate for neutral endopeptidase although CGRP, AM and IMD might not represent the preferred substrates for this enzyme: more specific enzymatic pathways probably remain to be identified. At least part of the post-translational processing of the prepro peptides is likely to occur extracellularly since fragments of AM and IMD precursor peptides have been identified in plasma. Mid-region pro-AM_{45–92} has attracted interest; the biological function, if any, of this fragment is unclear, it is less readily metabolized and may provide a more reliable biomarker in human plasma than the biologically active AM (Morgenthaler *et al.*, 2005). Radioimmunoassay specifically recognizing fragments of the human IMD precursor, prepro-IMD_{25–56} and prepro-IMD_{57–92}, have also become available recently and may offer similar utility.

The cellular and tissue distribution of IMD overlaps in some aspects with that of CGRP and AM (reviewed in Bell and McDermott, 1996; Brain and Grant, 2004; Ishimitsu *et al.*, 2006) but there are significant differences (for example, Kindt *et al.*, 2007). The prominent localization of IMD in hypothalamus and pituitary is consistent with emerging evidence for involvement of IMD, like AM, in central regulation of thirst, fluid and electrolyte balance (Taylor *et al.*, 2005; Taylor and Samson, 2005), like CGRP and AM, in the endocrine stress response (Chang *et al.*, 2005; Taylor and Samson, 2005; Taylor *et al.*, 2006), and secretion of oxytocin which influences reproductive function and complex, poorly defined behavioural responses in both males and females (Hashimoto *et al.*, 2005; Taylor and Samson, 2005). The extensive distribution of the peptide throughout the digestive tract, notably in the mucosa of the stomach, supports involvement of IMD as a neurotransmitter or paracrine mediator in gastrointestinal regulation (Roh *et al.*, 2004). Among other putative roles assigned to IMD are regulation of keratinocyte growth and maturation in skin (Kindt *et al.*, 2007), regulation of fetal–placental growth (Chauhan *et al.*, 2006), suppression of food intake (Roh *et al.*, 2004) and release of cytokines from adipose tissue which modify insulin resistance and cardiovascular morbidity associated with obesity (Morimoto *et al.*, 2007). Detailed discussion of

these functions is not possible within the current review. Instead, emphasis has been placed on appraisal of the biological significance of IMD within the cardiovascular and renal systems.

Plasma levels of CGRP predominantly reflect overspill of this peptide from sensory nerves (Zaidi *et al.*, 1985) and plasma levels of AM reflect secretion from atria and vasculature (Jougasaki and Burnett, 2000). The origin of IMD in plasma is unclear; the pituitary is likely to make a significant contribution. The relative paucity of IMD in the vicinity of the vasculature indicates that IMD is primarily an endocrine peptide. The prominent distribution of IMD in the renal tubule indicates a likely physiological role in local regulation of water-electrolyte balance and circulatory blood volume, while the robust expression of the peptide in diseased myocardium in contrast to the sparse levels found in healthy myocardium supports an important influence on cardiac pathology.

Receptor pharmacology

The biological actions of this peptide superfamily are mediated by two closely related G-protein-coupled receptors, the calcitonin receptor (CT) and calcitonin receptor-like receptor (CL) (Poyner *et al.*, 2002). G_s-mediated activation of adenylate cyclase represents the major signalling pathway coupled with these receptors (Kuwasako *et al.*, 2004), although G_{q/11}-mediated coupling to phospholipase C- β , G_i and/or G_o-mediated regulation of potassium channels, activation of phosphatidylinositol 3-kinase and guanylate cyclase have been reported (reviewed in Poyner *et al.*, 2002). A membrane-associated receptor component protein is required for optimum activation of signal transduction (Prado *et al.*, 2001). In addition, receptor activity-modifying proteins (RAMPs) act as molecular chaperones for trafficking of CL and CT from endoplasmic reticulum and Golgi apparatus to the cell surface and exert a pivotal influence over the receptors' pharmacological characteristics (McLatchie *et al.*, 1998).

With the discovery of AM it became apparent that CGRP and AM shared common receptors with varying degrees of

affinity (Husmann *et al.*, 2003; Hay *et al.*, 2004). Coexpression of CL and RAMP1 constitutes a functional receptor with CGRP selectivity and the classical pharmacological characteristics of the CGRP₁ receptor defined by Dennis *et al.* (1990); coexpression of CL with RAMP2 or RAMP3 gives rise to two subtypes of AM-selective receptors, AM₁ and AM₂ (reviewed by Poyner *et al.*, 2002). While CGRP₈₋₃₇ displays some selectivity as an antagonist at CGRP₁ receptors, its ability to discriminate satisfactorily between CGRP₁ and AM receptors is limited (Hay *et al.*, 2003). Similarly, while AM₂₂₋₅₂ displays some preference for AM receptors, it is a relatively low-affinity antagonist and cannot distinguish adequately between AM₁ and AM₂ receptors; high concentrations of this fragment also antagonize the CGRP₁ receptor (Hay *et al.*, 2003). In contrast, 1-piperidinecarboxamide N-[2-[[5-amino-1-[4-(pyridinyl)-1-piperazinyl] carbonyl]pentyl]amino]-1-[(3,5-dibromo-4-hydroxyphenyl)-methyl]-2-oxoethyl]-4-(1,4-dihydro-2-oxo-3(2H)-quinazoliny)-[R-(R*,S*)], the first non-peptide antagonist targeting this receptor family, is highly selective for the CGRP₁ receptor and has greatly assisted elucidation of effects mediated by this receptor (Doods *et al.*, 2000). Non-peptide antagonists displaying selectivity for AM₁ and AM₂ receptors, respectively, are eagerly awaited. RAMPs 1, 2 and 3 also interact with CT to form a series of receptors, AMY₁, AMY₂ and AMY₃, which bind AMY with varying degrees of affinity and selectivity (Hay *et al.*, 2005). It has been proposed (Hay *et al.*, 2005) that the AMY₁ receptor might account for the pharmacological profile of the classical CGRP₂ receptor defined by Dennis *et al.* (1990), since there are no remaining orphan G-protein-coupled receptor in the human genome closely related to CT or CL.

In the absence of convincing evidence that established or orphan G-protein-coupled receptor proteins form functional IMD-specific receptors by interacting with novel binding partners, it is hypothesized on account of the close structural

similarity between IMD, AM and CGRP that IMD's biological effects are mediated via interaction with the various RAMP/CL complexes. Indeed, IMD interacts with RAMP/CL complexes in transfected 293T cells (Roh *et al.*, 2004). Also, IMD₁₋₄₇ and IMD₈₋₄₇ stimulate accumulation of cyclic AMP in rat L6 myoblasts and human neuroblastoma SK-N-MC cells, both of which express RAMPs and CL, while the fragment IMD₁₇₋₄₇ is devoid of activity but antagonizes the action of IMD (Chang *et al.*, 2004; Roh *et al.*, 2004). IMD interacts non-selectively with all three RAMP/CL complexes and possesses a pharmacological profile distinct from either AM or CGRP (Figure 2). Compared to CGRP, IMD has greater potency to stimulate cyclic AMP production in 293T cells transfected with RAMP3/CL (AM₂) but lower potency in cells transfected with RAMP1/CL (CGRP₁); compared to AM, IMD has lower potency in 293T cells transfected with either RAMP2/CL (AM₁) or RAMP3/CL (AM₂). The weak interaction of IMD at AMY₁ receptors mirrors that of CGRP (Hay *et al.*, 2005).

Distribution of receptors

CL is ubiquitously expressed, especially in areas of the central nervous system related to neuroendocrine and autonomic control, feeding and thirst, and in lung, adrenal gland, spleen, kidney and skin (Hagner *et al.*, 2002; Cottrell *et al.*, 2005). CL is also present within blood vessels, being extensively distributed throughout the systemic and pulmonary vasculature in endothelial cells and vascular smooth muscle cells, and in heart (Totsune *et al.*, 2000; Cueille *et al.*, 2002) including cardiomyocytes (Autelitano and Ridings, 2001; Zhao *et al.*, 2006). RAMP1 is widely expressed in brain and spinal cord, gastrointestinal tract, adrenal gland, fat tissue, thymus and spleen (Nagae *et al.*, 2000; Cottrell *et al.*, 2005). Within the vasculature, RAMP1 is prominent in

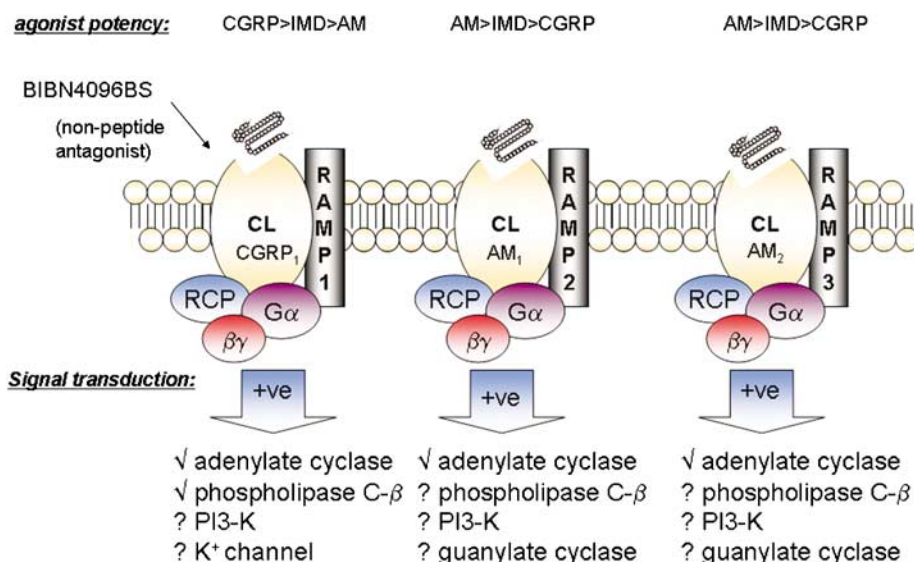


Figure 2 CGRP, AM and IMD share a common family of G-protein ($G\alpha\beta\gamma$)-coupled receptors formed by the association of the calcitonin receptor-like receptor (CL) with one of three receptor activity-modifying proteins (RAMPs). A receptor component protein (RCP) is also required for optimum activation of signal transduction. Pharmacological characteristics and putative signalling mechanisms associated with each of the three recognized receptor subtypes (CGRP₁, AM₁ and AM₂) are indicated (based on Poyner *et al.*, 2002; Roh *et al.*, 2004). CGRP, calcitonin gene-related peptide; AM, adrenomedullin; IMD, intermedin.

perivascular nerves and smooth muscle of smaller arteries and arterioles (Oliver *et al.*, 2002), and is also expressed in cardiomyocytes and non-myocytes (Autelitano and Ridings, 2001), inferring the presence of CGRP₁ receptors. RAMP2 is highly expressed in lung, spleen, fat tissue and kidney (Nagae *et al.*, 2000). Within the cardiovascular system, RAMP2 is found in macrovascular and microvascular endothelial cells, vascular smooth muscle (Kamitani *et al.*, 1999), cardiomyocytes, and less abundantly in cardiac non-myocytes (Autelitano and Ridings, 2001), inferring the presence of AM₁ receptors. RAMP3 is less extensively distributed: high levels are found in kidney and to a lesser extent in lung, spleen and thymus (Nagae *et al.*, 2000); RAMP3 is detected at low levels in adult myocardium and cardiomyocytes, being the least abundant RAMP (Zhao *et al.*, 2006) which infers that AM₂ receptors are sparse. However, RAMPs exhibit varying degrees of affinity for CL in different cell types in which they are coexpressed: the interaction of RAMP3 with CL predominates in rabbit aortic endothelial cells over that of RAMP1 or RAMP2 (Muff *et al.*, 1998). In addition, the distribution of RAMPs extends further than CL, indicating a much broader role for RAMPs in cellular function; additional functions of RAMPs, unrelated to their interaction with CL have emerged (Christopoulos *et al.*, 2003; Tfelt-Hansen *et al.*, 2006).

The sharing of a receptor family by CGRP, AM and IMD has confounded interpretation of radioligand binding and functional data pertaining to these naturally occurring ligands. However, a general finding is that the distribution of binding sites within the cardiovascular system mirrors that of the individual peptides themselves and of their receptor components. CGRP binding is particularly evident in intima and media of smaller blood vessels, cardiac valves and conduction tissue and more abundant in atria than ventricular myocardium (reviewed by Bell and McDermott, 1996). AM binding is prominent in vascular endothelium, especially microvasculature, but is also found in myocardium (reviewed by Ishimitsu *et al.*, 2006). There are few reports of binding studies using IMD as a radioligand. Within rat heart, IMD₁₋₅₃ recognizes a single population of binding sites on myocardial sarcolemmal membranes (Jia *et al.*,

2006). In view of the expression of all three RAMPs in myocardium, this finding could reflect inability of IMD to discriminate between CGRP₁ and AM receptors.

Haemodynamic effects

The initial observations of Takei *et al.* (2004b) indicated that intravenous injection of IMD₁₋₄₇ decreased arterial pressure more potently than AM in mice. Roh *et al.* (2004) reported that intraperitoneal injection of IMD₁₋₄₇ produced a hypotensive effect in rat; IMD₁₋₄₇ elicited a greater reduction in systolic blood pressure than an equivalent dose of IMD₈₋₄₇ but was less effective than AM. The response to IMD₁₋₄₇ was attenuated markedly by concurrent administration of CGRP₈₋₃₇ in normotensive and spontaneously hypertensive rat (SHR). A weak antagonistic effect of AM₂₂₋₅₂ on IMD response was demonstrated in SHR; the effect of this fragment was not examined in normotensive animals (Roh *et al.*, 2004). Intravenous administration of IMD₁₋₄₇ evokes a rapid and marked hypotensive effect in rat (Table 2): Taylor *et al.* (2005) found the response to 1 nmol kg⁻¹ IMD₁₋₄₇ to be of similar magnitude to that of CGRP but considerably less than that evoked by AM; Ren *et al.* (2006) reported that the hypotensive effect of IMD₁₋₄₇ was more prominent than that of an equivalent dose (3 nmol kg⁻¹) of either AM or IMD₁₋₅₃, while Fujisawa *et al.* (2006) noted that the response to IMD was similar in magnitude to an equivalent dose of AM but less sustained. Pan *et al.* (2005), employing exceptionally large doses of the peptides (~600 nmol kg⁻¹), found IMD₁₋₄₇ and IMD₈₋₄₇ to elicit responses of similar magnitude. Overall, these findings indicate that the longer form of IMD possesses greater biological activity than the truncated form *in vivo* and that the hypotensive effect of IMD is qualitatively and quantitatively different to that of AM. In contrast, the influence of IMD₁₋₄₇ on regional haemodynamics in rat is more similar to that of AM than CGRP but distinct from both (Fujisawa *et al.*, 2007). IMD increases the percent distribution of cardiac output particularly to kidney, consistent with the marked expression of the peptide in this organ, but also like AM to heart, lungs and spleen and like CGRP to liver,

Table 2 Infusion studies in rodents

Peptide	Dose		mABP maximum change (mm Hg)		
IMD ₁₋₄₇	1.0 nmol kg ⁻¹	iv	-7.9	Rat	Taylor <i>et al.</i> (2005)
AM	1.0 nmol kg ⁻¹	iv	-15.0	Rat	Taylor <i>et al.</i> (2005)
CGRP	1.0 nmol kg ⁻¹	iv	-5.0	Rat	Taylor <i>et al.</i> (2005)
AM	3.0 nmol kg ⁻¹	iv	-18.3	Rat	Ren <i>et al.</i> (2006)
IMD ₁₋₅₃	3.0 nmol kg ⁻¹	iv	-15.8	Rat	Ren <i>et al.</i> (2006)
IMD ₁₋₄₇	3.0 nmol kg ⁻¹	iv	-19.2	Rat	Ren <i>et al.</i> (2006)
IMD ₁₋₄₇	5.0 nmol kg ⁻¹	iv	-12.0	Rat	Fujisawa <i>et al.</i> (2006)
AM	5.0 nmol kg ⁻¹	iv	-10.0	Rat	Fujisawa <i>et al.</i> (2006)
AM	10 nmol kg ⁻¹	iv	-17.0	Mouse	Takei <i>et al.</i> (2004a)
IMD ₁₋₄₇	10 nmol kg ⁻¹	iv	-27.0	Mouse	Takei <i>et al.</i> (2004a)
IMD ₁₋₄₇	~600 nmol kg ⁻¹	iv	-39.0	Rat	Pan <i>et al.</i> (2005)
IMD ₈₋₄₇	~600 nmol kg ⁻¹	iv	-45.0	Rat	Pan <i>et al.</i> (2005)
IMD ₁₋₄₇	195 pmol	icv	+11.0	Rat	Taylor <i>et al.</i> (2005)
CGRP	195 pmol	icv	+9.0	Rat	Taylor <i>et al.</i> (2005)
AM	195 pmol	icv	No change	Rat	Taylor <i>et al.</i> (2005)

AM, adrenomedullin; CGRP, calcitonin gene-related peptide; icv, intracerebroventricular; IMD, intermedin; iv, intravenous; mABP, mean arterial blood pressure.

stomach and small intestine. In rat, the hypotensive effect of IMD, like that of AM, is enhanced in pregnancy (Chauhan *et al.*, 2007). In addition to an effect of IMD on resistance vessels, reduction of mean circulatory filling pressure in the absence of an effect on blood volume is evoked by IMD in rats subjected to ganglionic blockade, consistent with a direct action on capacitance vessels (Abdelrahman and Pang, 2006). Intravenous infusion of IMD into a large conscious mammal, namely the sheep, also results in marked reduction in mean arterial blood pressure; like that of AM, the effect of IMD is more pronounced on diastolic than systolic blood pressure (Charles *et al.*, 2006).

It is probable that the actions of exogenous IMD in vasculature are associated with activation of both CGRP and AM receptors dependent on route of administration and ease of access to these receptors. The relative abundance of these receptor populations, their localization on endothelium or smooth muscle and associated coupling mechanisms vary with regard to species and vessel and influence the characteristics of responses evoked in specific vascular beds. The action of CGRP is independent of endothelium in the majority of tissues investigated and associated with CGRP₁ receptor-stimulated accumulation of cyclic AMP and activation of potassium channels resulting in membrane hyperpolarization in vascular smooth muscle; less commonly CGRP, acting at CGRP₁ receptors or AM receptors located on endothelial cells stimulates nitric oxide release (reviewed in Bell and McDermott, 1996; Brain and Grant, 2004) (Figure 3). AM is also heterologous in respect to species and vessel, both CGRP₈₋₃₇- and AM₂₂₋₅₂-sensitive and endothelium-dependent and -independent effects have been reported (reviewed in Brain and Grant, 2004). IMD₁₋₄₇ and IMD₈₋₄₇ increase cyclic AMP levels and relax norepinephrine precontracted rat aorta *in vitro* (Pan *et al.*, 2005). Vasodilatation of this vessel in response to IMD₁₋₅₃ has been attributed to endothelium-dependent and -independent mechanisms (Yang *et al.*, 2006). Indeed IMD₁₋₅₃, like AM, stimulates constitutive, but not inducible nitric oxide synthase leading to enhanced nitric oxide production in aortic endothelial cells and also stimulates vascular L-arginine transport and expression of cationic amino-acid transporters, CAT-1 and CAT-2B (Yang *et al.*, 2006). Rat aorta is one of the few vessels in which endothelium-dependent nitric oxide release has been demonstrated in response to CGRP (Brain and Grant, 2004). However, the effect of IMD in aorta might also be accounted for by activation of AM-preferring receptors on vascular smooth muscle since AM-stimulated accumulation of cyclic AMP in aortic smooth muscle cells is attenuated by AM₂₂₋₅₂ (Eguchi *et al.*, 1994), although AM₂₂₋₅₂ lacks adequate specificity to enable CGRP₁ receptor involvement to be discounted. In rat mesenteric artery, the action of IMD is dependent on production of nitric oxide, accumulation of cyclic AMP and cyclic GMP and activation of potassium channels, again consistent with both endothelium-dependent and -independent signalling events (Chauhan *et al.*, 2007). Kandilci *et al.* (2006) reported that IMD₁₋₄₇ decreases pulmonary arterial pressure and decreased pulmonary vascular resistance in rat under conditions of elevated pulmonary vasoconstrictor tone. The response to IMD is attenuated by but less sensitive to CGRP₈₋₃₇ than that of

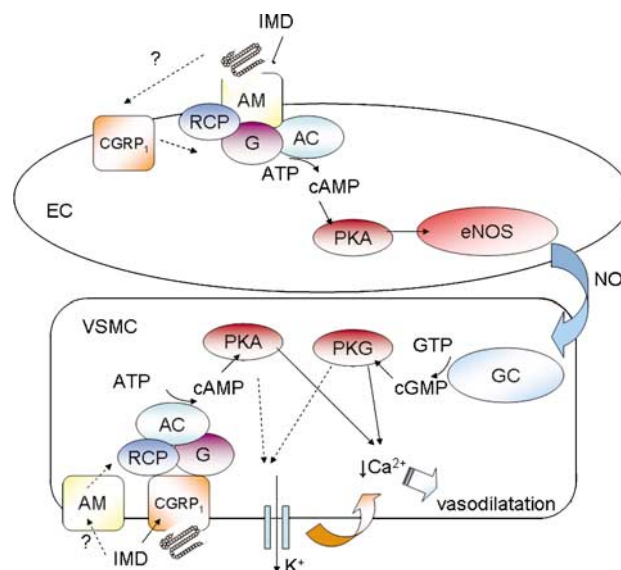


Figure 3 Schematic diagram outlining the proposed endothelium-dependent and -independent signalling pathways by which IMD might evoke vasodilatation. The relative contribution of such processes is likely to vary significantly with respect to vessel and species. Endothelial cell (EC); vascular smooth muscle cell (VSMC); adenylate cyclase (AC); nitric oxide synthase (NOS); protein kinase A (PKA); protein kinase G (PKG); guanylate cyclase (GC). The endothelium-dependent effect of IMD is attributed to cyclic AMP-dependent activation of constitutive endothelial NOS and synthesis of nitric oxide which then diffuses to VSMC to stimulate GC leading to accumulation of cyclic GMP and activation of PKG. Accumulation of cyclic AMP in EC is stimulated mainly by activation of endothelial AM receptors although a possible contribution of CGRP₁ receptors, at least in some vessels, cannot be discounted. Stimulation of CGRP₁ receptors, or alternatively in some vessels of AM receptors, on VSMC directly is associated with accumulation of cyclic AMP and activation of PKA. PKA and PKG trigger processes in VSMC which lead to reduced intracellular calcium ion concentration resulting in vasodilatation; such processes may involve activation of outward potassium currents leading to membrane hyperpolarization and reduced calcium entry. CGRP, calcitonin gene-related peptide; IMD, intermedin; NOS, nitric oxide synthase.

CGRP. Although IMD induces tachyphylaxis to CGRP, IMD does not induce self-tachyphylaxis, and CGRP does not induce tachyphylaxis to IMD. Pulmonary vasodilatation in response to IMD is dependent largely on production of nitric oxide and not cyclooxygenase products or activation of K_{ATP} channels (Kandilci *et al.*, 2006). This finding is consistent with the endothelial dependence of CGRP reported previously in rat pulmonary artery (Wisskirchen *et al.*, 1998). However, there is conflicting evidence regarding involvement of nitric oxide in the vasorelaxant responses to CGRP in this vessel (Heaton *et al.*, 1995; Wisskirchen *et al.*, 1998). The action of IMD cannot be entirely accounted for by stimulation of CGRP₁ receptors on endothelium coupled to production of nitric oxide. The action of AM in this vessel is not attenuated by CGRP₈₋₃₇ (Wisskirchen *et al.*, 1998) or N^G-nitro-L-arginine-methyl-ester (L-NAME) (Heaton *et al.*, 1995) consistent with the presence of additional receptors, coupled to alternative mechanisms, at which IMD might also act.

Hypotension evoked by IMD₁₋₄₇ in rat *in vivo* is not attenuated by L-N^G-monomethyl arginine (L-NMMA), prompting Taylor *et al.* (2005) to conclude that the systemic

vasodilator response to IMD, like that to CGRP, is much less dependent on nitric oxide production than that of AM. Taken together with the similar magnitude of the responses to IMD and CGRP, this observation is more consistent with an action of IMD at CGRP₁ receptors, located predominantly on smooth muscle, than at AM receptors, found mainly on endothelium. However, the lack of effect of L-NMMA could be interpreted as relative inhibition since vasoconstriction evoked by L-NMMA directly could mask an inhibitory effect by raising tone thereby favouring the action of a vasodilator mediator (Gardiner *et al.*, 1995). In all *in vitro* studies to date IMD has been applied to vessels in which endothelium-dependence of CGRP's action has been demonstrated, at least in part. Further studies are warranted to characterize the vasorelaxant effect of IMD in smaller resistance arteries and arterioles in which the vasorelaxant effect of CGRP is endothelium-independent and CGRP₁ receptors are found abundantly on smooth muscle (Oliver *et al.*, 2002).

The physiological as opposed to pharmacological influence of members of this family on haemodynamics is unclear. CGRP₈₋₃₇ does not affect basal blood pressure (Gardiner *et al.*, 1991) and results obtained from measurement of blood pressure in various strains of CT/ α CGRP gene knockout mice relative to wild-type mice have been conflicting (Lu *et al.*, 1999; Gangula *et al.*, 2000; Ohhashi *et al.*, 2001). CGRP₈₋₃₇ does influence regional blood flow (Gardiner *et al.*, 1990) which indicates that CGRP might regulate tissue blood flow under physiological conditions and provide a braking system for microvascular resistance to protect tissues from injury. Homozygous knockout of the AM gene in mice results in a lethal phenotype due to insufficient development of blood vessels (Caron and Smithies, 2001; Hay and Smith, 2001). Evidence obtained from viable heterozygous AM gene knockout mice indicates a role for endogenous AM in regulation of blood pressure (Shindo *et al.*, 2001). The physiological influence of IMD is unclear in the absence of suitable transgenic mice in which to investigate the unique contribution of this peptide.

Effects on the kidney

Takei *et al.* (2004b) initially reported that bolus injection of a hypotensive dose of IMD, but not AM, inhibited urine flow and reduced urine sodium concentration in mice. In conscious sheep infusion of a markedly hypotensive dose of IMD, like AM, also evoked a modest reduction in urine sodium concentration but did not influence urine volume (Charles *et al.*, 2006). In contrast, diuretic and natriuretic effects of AM have been reported previously in other species, including human (Nagaya *et al.*, 1999; Lainchbury *et al.*, 2000). It is difficult to dissociate effects exerted locally on the kidney from the consequences of the systemic hemodynamic effects of a vasodilator peptide. It is possible that the systemic hypotensive effect of IMD resulted in reduced renal perfusion thereby masking a direct diuretic and natriuretic effect of IMD. Furthermore, the strain of mouse used by Takei *et al.* (2004b) originated in a dry region of Central Asia and these mice urinate scarcely to conserve sodium and water; some natriuresis and diuresis was evoked

by IMD at higher rates of infusion in overhydrated mice. Direct infusion of IMD into rat renal artery at doses sufficient to profoundly lower systemic blood pressure also inhibits urine flow (Fujisawa *et al.*, 2004), while intravenous administration of less hypotensive doses of IMD, like AM, increases renal blood flow and the proportion of cardiac output distributed to the kidney indicative of a direct role for IMD in renal vasculature (Fujisawa *et al.*, 2006, 2007). Renal vasodilatation in response to AM has been attributed to a nitric oxide-dependent mechanism (Edwards *et al.*, 1996; Majid *et al.*, 1996). Activation of renal sympathetic nerve activity (RSNA) accompanies the hypotensive actions of IMD, AM and also of sodium nitroprusside; however, the effects of the peptides on RSNA are not completely suppressed while that of sodium nitroprusside is abolished by baroreceptor denervation (Fujisawa *et al.*, 2006). Similarly in sheep, enhancement of RSNA precedes the maximum reduction in blood pressure (Charles *et al.*, 2006). The baroreceptor reflex cannot therefore account for all of the augmentation of RSNA by these peptides and additional mechanisms such as direct enhancement of central sympathetic outflow are likely to contribute. The actions of IMD on renal blood flow persist when blood pressure, heart rate and RSNA return to normal (Fujisawa *et al.*, 2006); furthermore, subdepressor doses of the peptide also increase renal blood flow and decrease renal vascular resistance strongly supporting a direct effect on the renal vasculature (Fujisawa *et al.*, 2004). Lack of influence of IMD on glomerular filtration rate infers that IMD influences renal afferent and efferent arterioles equally; in contrast, intrarenal infusion of subdepressor doses of CGRP increases renal blood flow and glomerular filtration rate (Kurtz *et al.*, 1989; Edwards and Trizna, 1990). Increased urinary flow and urinary sodium excretion in response to IMD in the absence of an increase in glomerular filtration rate is indicative of a direct inhibitory influence of IMD on tubular re-absorption of sodium (Fujisawa *et al.*, 2004). The actions of AM are qualitatively similar but more persistent. At variance with these findings, AM stimulates sodium uptake in distal tubule luminal membranes *in vitro*, indicative of an antinatriuretic effect (Leclerc and Brunette, 2000). The direct action of IMD on tubular sodium handling, receptor involvement and associated signalling mechanism remains unresolved. Central administration of IMD, like AM, increases plasma levels of arginine vasopressin (Taylor and Samson, 2005) which exerts an action predominantly on distal tubule and collecting duct to conserve water, but not salt, by concentrating the urine and reducing urine volume. The influence of IMD on renal tubular water and electrolyte balance is likely to reflect a composite of the direct and indirect actions of the peptide.

Effects on the heart

IMD influences cardiac output as a consequence of direct and indirect action on the heart (Figure 4). Hypotension evoked by intraperitoneal or intravenous administration of IMD, like that of AM and CGRP, is accompanied by increased heart rate (Roh *et al.*, 2004; Takei *et al.*, 2004b; Charles *et al.*, 2006). Tachycardia evoked by intravenous administration of

IMD₁₋₄₇ in rat is blunted by ganglionic blockade consistent with autonomic nervous system involvement (Abdelrahman and Pang, 2006). In rat, tachycardia evoked by IMD is qualitatively similar but less sustained than that of AM (Roh *et al.*, 2004) but greater than that of sodium nitroprusside despite an equivalent decrease in mean arterial pressure. Tachycardia evoked by sodium nitroprusside is abolished by baroreceptor denervation while that of IMD is attenuated but not abolished (Fujisawa *et al.*, 2006). In sheep, augmentation of heart rate occurs before the maximum extent of reduction in blood pressure by IMD is achieved (Charles *et al.*, 2006). The action of IMD clearly cannot be accounted for solely by baroreceptor activation in response to hypotension and additional mechanism(s) are likely to contribute. Indeed, intracerebroventricular administration of IMD₁₋₄₇ (Taylor *et al.*, 2005) evokes a marked increase in blood pressure and heart rate in rat; in regard to potency and duration the actions of IMD₁₋₄₇ are more similar to those of CGRP than AM, are antagonized by CGRP₈₋₃₇ suggestive of CGRP₁ receptor involvement, and attenuated by phentolamine indicating that IMD's effect is attributable to sympathetic outflow from the brain (Taylor *et al.*, 2005). Intracerebroventricular administration of IMD₁₋₅₃ also increases blood pressure and heart rate; these actions are markedly attenuated by an anti-prepro-IMD antibody, or by either CGRP₈₋₃₇ or AM₂₂₋₅₂ (Ren *et al.*, 2006). However, the antibody did not influence blood pressure or heart rate in the absence of exogenous IMD₁₋₅₃ indicating that the endogenous peptide is unlikely to participate in central cardiovascular regulation under physiological conditions. IMD₈₋₄₇ also increases heart rate in isolated perfused hearts of rats consistent with a direct chronotropic action (Pan *et al.*, 2005). It is possible that the direct component of the positive chronotropic effect of IMD₈₋₄₇ is associated with activation

of CGRP₁ receptors since CGRP evokes a direct chronotropic response in spontaneously beating right atria and neonatal cardiomyocytes of rat (Fisher *et al.*, 1988; Wang and Fiscus, 1989).

Intravenous administration of IMD₈₋₄₇ or IMD₁₋₄₇ in rat at concentrations sufficiently large to evoke marked hypotension reduce left ventricular end-systolic pressure and maximum rates of pressure development and decline, indicative of a negative inotropic action (Pan *et al.*, 2005). IMD₈₋₄₇ evoked similar effects in isolated perfused hearts while IMD₁₋₄₇ was almost without effect; baseline parameters were greater in the IMD₈₋₄₇-treated group which may have accounted for the difference observed. Myocardial cyclic AMP content is elevated in response to intravenous administration *in vivo* or perfusion of isolated hearts with the peptides (Pan *et al.*, 2005); this observation is inconsistent with a negative inotropic action. In subsequent studies in the same laboratory, intraperitoneal injection of IMD₁₋₅₃ over four consecutive days in rat had no influence on cardiac function (Jia *et al.*, 2006) but paradoxically IMD₁₋₅₃ evoked a positive inotropic effect acutely in isolated perfused rat heart at variance with earlier findings (Yang *et al.*, 2005a). Fujisawa *et al.* (2007) also report that intravenous administration of IMD₁₋₄₇ enhances cardiac output in rats. Reports of augmented cardiac output in response to CGRP and AM are influenced by variations in regard to species studied, dose chosen, and therefore the extent of reduction in afterload and sympathetic activation. The latter influences are absent in isolated perfused hearts. CGRP elicits a positive inotropic action in isolated perfused hearts of guinea-pigs but not of rats or rabbits (reviewed in Bell and McDermott, 1996). AM evokes a positive inotropic effect with gradual onset in isolated perfused hearts of rats; this effect is not attenuated by CGRP₈₋₃₇, indicative of an AM receptor-dependent

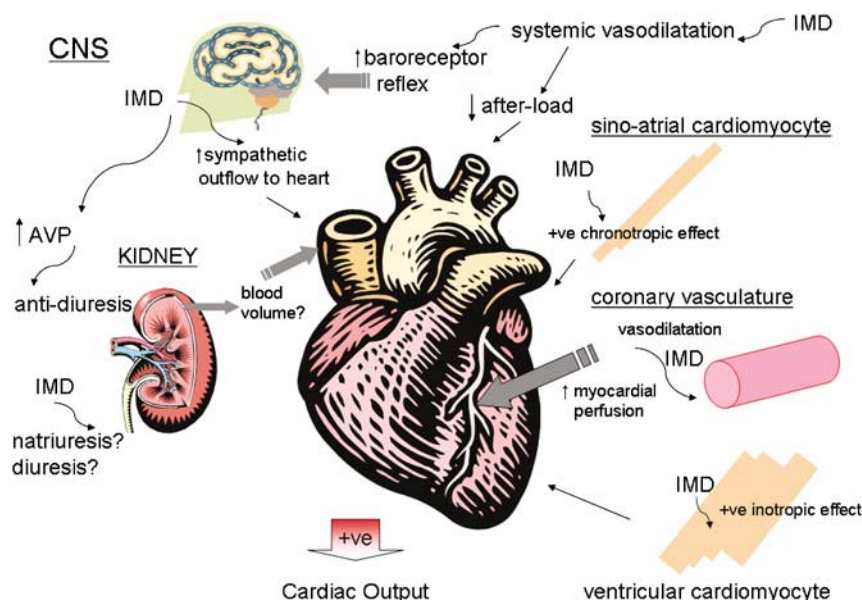


Figure 4 Schematic diagram depicting the combination of direct and indirect effects exerted upon the heart by IMD, which influence cardiac output. IMD exerts chronotropic and inotropic effects directly on myocardium. Coronary vasodilatation enhances myocardial perfusion, which augments contractility indirectly. Systemic vasodilatation reduces afterload and activates the baroreceptor mechanism to increase central sympathetic outflow, which also enhances cardiac contractility. The actions of IMD centrally and locally within the kidney influence blood volume which in turn influences venous return and cardiac output. IMD, intermedin.

response. Furthermore, the effect is independent of cyclic AMP and associated with initial discharge of calcium stores and subsequent protein kinase C-dependent phosphorylation of L-type calcium channels (Szokodi *et al.*, 1998). The positive inotropic response to IMD in rat might be attributed in part to enhanced myocardial perfusion since IMD₈₋₄₇ and IMD₁₋₅₃ dilate coronary arteries and enhance coronary flow in isolated perfused rat hearts (Pan *et al.*, 2005; Yang *et al.*, 2005a). The vasorelaxant effect of AM in human coronary artery is attenuated by AM₂₂₋₅₂ (Terata *et al.*, 2000). However, the effect of AM in rat is less potent than that of CGRP and sensitive to CGRP₈₋₃₇, suggesting CGRP₁ receptor involvement, at least in this species, in the response to IMD. Evidence for a direct action on myocardium requires use of isolated atria, papillary muscle or cardiomyocytes. IMD evokes a positive inotropic effect in adult rat (Bell, unpublished observation) and mouse ventricular cardiomyocytes (Dong *et al.*, 2006). In the latter, both PKA- and protein kinase C-dependent mechanisms are implicated; the inotropic effect is associated with intracellular calcium release and is qualitatively similar to those of CGRP and AM, but receptor involvement has not been explored. It is not known if IMD evokes similar responses directly in cardiomyocytes of other species including man. CGRP evokes a direct positive inotropic action in atria of rat, guinea-pig and human, human myocardial trabeculae and in ventricular cardiomyocytes of rat, but not rabbit or human (reviewed in Bell and McDermott, 1996; Saetrum Opgaard *et al.*, 2000). In adult rat ventricular cardiomyocytes, the action of CGRP is attenuated by CGRP₈₋₃₇ indicative of CGRP₁ receptor involvement, but independent of cyclic AMP (Bell and McDermott, 1994) and associated with calcium-induced calcium release (Huang *et al.*, 1999). AM elicits a positive inotropic effect on papillary muscle and atrial cardiomyocytes of rat (Ihara *et al.*, 2000), has no activity on human ventricular cardiomyocytes (Saetrum Opgaard *et al.*, 2000) and evokes a negative response in rabbit cardiomyocytes attributed to nitric oxide synthesis and activation of a cyclic GMP-dependent mechanism (Ikenouchi *et al.*, 1997). Ability to demonstrate a contractile response directly in myocardium is dependent on species, age, cardiac chamber and frequency and duration of electrical stimulation. Heterogeneity in regard to receptor involvement and signalling mechanism is likely to be influenced by selectivity of the antagonists and signal transduction inhibitors chosen and concentrations at which these are employed. Activation of CGRP₁ receptors coupled to phosphatidylinositol signalling and mobilization of intracellular calcium is likely to contribute to the inotropic effect of IMD in rodent ventricular cardiomyocytes; the involvement of additional receptors and/or signalling mechanisms is possible (Figure 5a).

Pathophysiological significance in the cardiovascular system

Expression of IMD is augmented in hypertrophied myocardium induced by chronic administration of isoprenaline in rat (Pan *et al.*, 2005). The transcript is also upregulated while

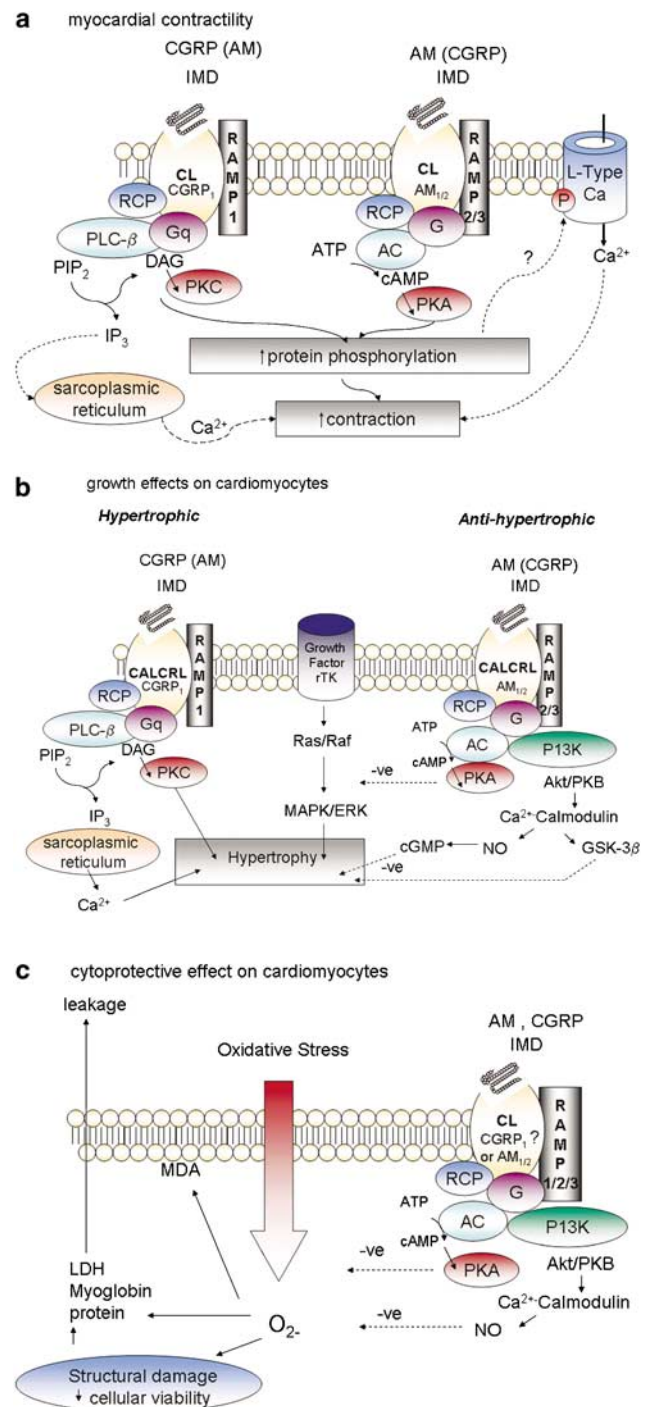


Figure 5 Schematic diagram outlining the proposed signalling pathways by which IMD might influence (a) contraction; (b) hypertrophic remodelling; (c) protection against the deleterious consequences of oxidative stress directly in ventricular cardiomyocytes. Adenylate cyclase (AC); protein kinase A (PKA); protein kinase C (PKC); diacylglycerol (DAG); mitogen-activated protein kinase (MAPK); receptor tyrosine kinase (rTK); phospholipase C-β (PLC-β); extracellular signal-regulated kinase (ERK); phosphoinositide 3-kinase (PI3K); receptor component protein (RCP). IMD, intermedin.

that of AM is unchanged (Figure 6) and neutral endopeptidase reduced (McDermott *et al.*, 2007) in hypertrophied left ventricular cardiomyocytes of SHR relative to that of

normotensive Wistar–Kyoto rat. IMD does not alter basal protein synthesis in neonatal (Pan *et al.*, 2005) or adult (Figure 7) rat cardiomyocytes but abolishes protein synthesis in response to a hypertrophic stimulus. These findings support an important counter-regulatory influence of IMD locally on hypertrophic growth and remodelling. It is probable that this action of IMD is mediated by AM-preferring receptors since AM attenuates cardiomyocyte hypertrophy (Bell *et al.*, 2006) while CGRP is a prohypertrophic stimulus in rat (Bell *et al.*, 1995) (Figure 5b). Increased expression of RAMP1, and to a lesser extent RAMP3 mRNA, is evident but RAMP2 mRNA is unchanged in hypertrophied cardiomyocytes of SHRs. Paradoxically, at protein level, all three RAMPs are less abundant in sarcolemmal membranes of SHR cardiomyocytes than those of Wistar–Kyoto rat (McDermott *et al.*, 2007); the consequences for IMD signalling and other functions regulated by RAMP proteins is unclear. Such changes could represent a primary defect in IMD receptor signalling or downregulation secondary to augmented levels of IMD.

Global ischaemia followed by reperfusion in rat heart *in vitro* is associated with bradycardia, inhibition of contractile function and significant myocardial injury. Calcium overload and excessive production of reactive oxygen species are central to ischaemia–reperfusion injury. IMD, administered during the reperfusion period ameliorates the reduction in cardiac contractility, demonstrated by reversal of bradycardia and of the decline in maximum change in left ventricular pressure and rates of pressure development and decay (Yang *et al.*, 2005b). IMD attenuates ischaemia–reperfusion injury, as evidenced by reduced leakage of lactate dehydrogenase, total protein and myoglobin from myocardium and less formation of malondialdehyde, an end product of lipid peroxidation by reactive oxygen species. The cardio-protective effects of IMD are similar to those of CGRP and AM (Kato *et al.*, 2003), which are mediated in part by cyclic AMP (Beltowski and Jamroz, 2004). IMD also stimulates accumulation of cyclic AMP in myocardium (Yang *et al.*, 2005a). The number of binding sites for IMD in myocardium is increased

during ischaemia–reperfusion, indicating transport of more receptors to the cell surface; receptor affinity is also increased. It is not clear which receptor subtype(s) are involved in the protective effects of IMD (Yang *et al.*, 2005a). Levels of the prepro-IMD precursor are reduced, indicative of enhanced processing to the active form. All three forms of the peptide, IMD_{1–53}, IMD_{1–47} and IMD_{8–47} are equally protective (Yang *et al.*, 2005a, b). At least part of the beneficial action of IMD could be attributed to improved coronary perfusion (Yang *et al.*, 2005a), but it is likely that a positive inotropic effect and cytoprotective effect is also exerted directly on myocardium (Figure 5c).

Subcutaneous injection of a high dose of isoprenaline over two consecutive days provides an alternative means of inducing myocardial ischaemic injury. This approach increases plasma lactate dehydrogenase activity and plasma and myocardial malondialdehyde content, lowers maximum rate of increase and decrease of left ventricular pressure development and raises left ventricular end diastolic pressure, indicative of severe heart failure and ischaemic injury. Myocardial structural disorder, subendocardial necrosis, fibroblast hyperplasia, capillary dilatation and leukocyte infiltration are evident. Concurrent administration of IMD_{1–53} improves myocardial contractile function, as evidenced by reversal of isoprenaline-evoked decline in functional parameters, attenuated myocardial lactate dehydrogenase leakage and malondialdehyde formation and isoprenaline-induced decline in myocardial cyclic AMP content (Jia *et al.*, 2006). IMD_{1–53} binding capacity is increased in myocardial sarcolemmal membranes and this is associated with augmented mRNA expression of CL and each of the RAMPs, especially RAMP3. This finding is indicative of enhanced synthesis of receptor components, in addition to an influence on transport of receptors to the cell surface, demonstrated in the acute setting *in vitro*. The absence of an effect of IMD_{1–53} on contractile function in the absence of isoprenaline-induced injury, together with marked upregulation of the IMD receptor system indicates that IMD plays a pivotal cardio-protective role in response to ischaemic insult.

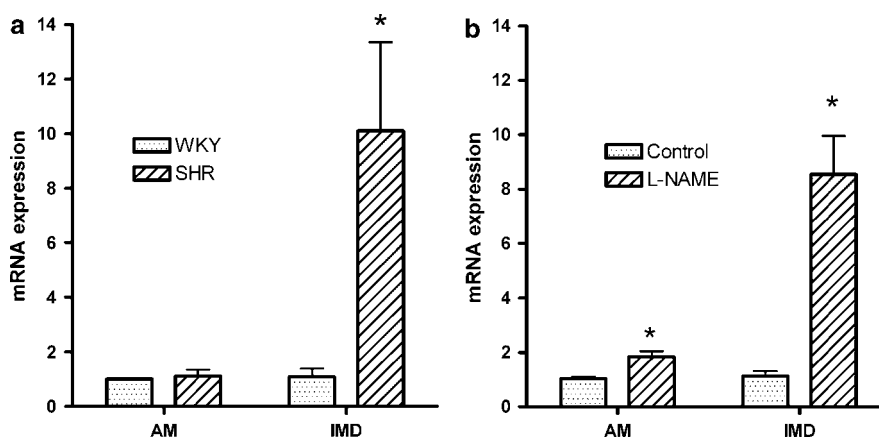


Figure 6 Expression of prepro-IMD and prepro-AM mRNAs in left ventricular cardiomyocytes of (a) SHR and WKY rat at 20 weeks of age (mean + s.e., $n=8$), data taken from McDermott *et al.*, 2007; (b) 8-week-old SD rat given L-NAME ($35 \text{ mg kg}^{-1} \text{ day}^{-1}$) in drinking water for 8 weeks and age-matched untreated rat for comparison, (mean + s.e., $n=10$), data taken from Bell *et al.* (2007). Data were expressed relative to glyceraldehyde 3-phosphate dehydrogenase mRNA levels. *Denotes significant difference between treatment groups, $*P<0.05$. AM, adrenomedullin; IMD, intermedin; SHR, spontaneously hypertensive rat; WKY, Wistar–Kyoto; L-NAME, N^G -nitro-L-arginine-methyl-ester.

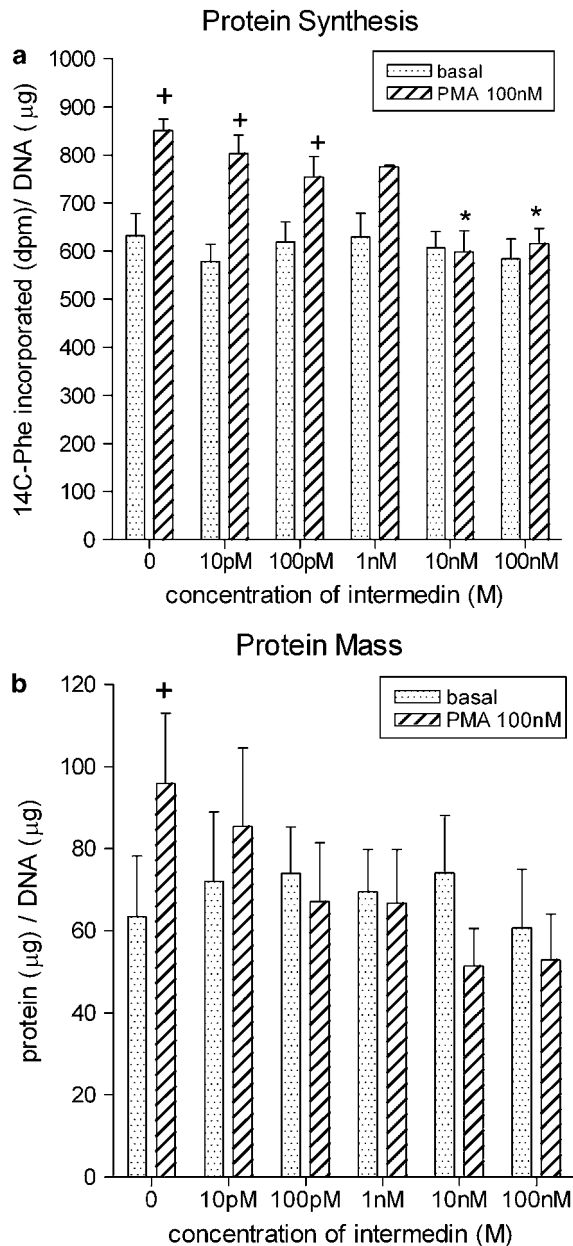


Figure 7 Effect of IMD (10 pM–100 nM) on (a) protein synthesis; (b) protein mass in ventricular cardiomyocytes (combined population obtained from left and right ventricles of 12-week-old normotensive SD control rat) and maintained in short-term (24 h) culture under basal conditions and in the presence of the hypertrophic stimulus, phorbol 12-myristate 13-acetate (PMA) (100 nM). Data are the mean values \pm s.e. of 4–6 heart cell preparations. *Denotes significant difference between PMA response in the absence and presence of IMD. ⁺Denotes significant difference between basal value and PMA-stimulated response ($P < 0.05$). Unpublished observation; methodology employed as described for AM in Bell *et al.* (2006). IMD, intermedin.

Chronic administration of the inhibitor of nitric oxide synthase, N^G -nitro-L-arginine methyl ester (L-NAME), to rats results in hypertension which depending on the dose given is accompanied by activation of the renin–angiotensin system, myocardial remodelling (cardiac hypertrophy and fibrosis), vascular remodelling (medial thickening and

perivascular fibrosis), cardiac ischaemia and necrosis, and mechanical dysfunction (reviewed in Zatz and Baylis, 1998; Bell *et al.*, 2006). In our own laboratory in which L-NAME ($35 \text{ mg kg}^{-1} \text{ day}^{-1}$) was administered to rats for 8 weeks, myocardial hypertrophy is evidenced by increased cardiomyocyte width and acquisition of a hypertrophic phenotype (Zhao *et al.*, 2006) while oxidative stress is shown by increased cardiomyocyte membrane protein oxidation and augmented expression of pro-oxidant enzymes (Bell *et al.*, 2007). Robust increase in prepro-IMD mRNA relative to prepro-AM in cardiomyocytes of rats treated chronically with L-NAME (Figure 6) indicates an important role for IMD in the cardiac pathology of nitric oxide deficiency. Furthermore, IMD, AM and their receptor components are differentially regulated: upregulation of AM, RAMP2 and RAMP3 expression in cardiomyocytes of rats subjected to chronic nitric oxide deficiency is prevented when blood pressure is normalized by the smooth muscle relaxant, hydralazine given concurrently with the diuretic, hydrochlorothiazide (Zhao *et al.*, 2006) while that of IMD and RAMP1 is not attenuated. Conversely, upregulation of cardiomyocyte IMD and RAMP1 is abolished when oxidative stress is normalized by vitamin C and Tempol; this antioxidant combination does not lower blood pressure and does not attenuate the augmented expression of AM, RAMP2 and RAMP3 in L-NAME-treated rats (Bell *et al.*, 2007). These studies indicate two important influences on AM and IMD signalling: cardiomyocyte expression of AM, RAMP2 and RAMP3 is influenced by pressure loading on the myocardium while that of IMD/RAMP1 is regulated by the extent of local oxidative stress in the absence of pressure loading. Neuronal release of CGRP from perivascular nerves is enhanced in response to myocardial ischaemia (Franco-Cereceda *et al.*, 1987). CGRP enhances blood flow to ischaemic myocardium, improves contractile indices, attenuates ischaemia–reperfusion arrhythmias and protects cardiomyocytes directly against the detrimental effects of oxidative stress (Ren *et al.*, 1993; Zhang *et al.*, 1994). Enhanced levels of non-neuronally derived IMD acting on CGRP₁ receptors, may, like neuronal CGRP, serve primarily an anti-ischaemic cardio-protective function in cardiomyocytes subjected to chronic nitric oxide deficiency. An additional attenuating influence for IMD on cardiomyocyte hypertrophy in this model, mediated by stimulation of AM₂ receptors cannot be discounted, particularly since RAMP3 expression is also enhanced.

Oxidative stress within the cerebral circulation is a common cardiovascular complication of diabetes and is also observed in response to restriction in cerebral blood flow due to stroke or transient ischaemic attack. Cerebral endothelial cells, a major component of blood–brain barrier, normally form tight junctions and have specialized transport systems to maintain strict ionic and metabolic homeostasis of brain parenchyma. Cerebral endothelial cells have a large energy requirement by virtue of the abundance and activity of active transport systems operational across the blood–brain barrier. Oxidative stress is associated with endothelial cell contraction, reduced blood–brain barrier integrity and leakage of serum components into the central nervous system resulting in oedema and neuronal damage. IMD, like

AM, preserves cerebral endothelial cell viability *in vitro* as evidenced by attenuation of hydrogen peroxide induced reduction in trans-endothelial electrical resistance and increased intercellular gap formation leading to increased permeability (Chen *et al.*, 2006). IMD (and AM) increases cyclic AMP and nitric oxide production, enhances peripheral localization of F-actin bands improving stability of tight junctions and preserves mitochondrial membrane potential and integrity. Prepro-AM is abundantly expressed in cerebral endothelium, indicative of an important regulatory role (Kis *et al.*, 2002). These effects of exogenously applied IMD are likely to be mediated by stimulation of endothelial AM receptors. Less is known currently about the presence and physiological influence of IMD in the cerebral vasculature *in vivo*, particularly whether expression of IMD is augmented locally in response to oxidative stress.

Future perspectives

Emerging evidence indicates that IMD is a potent vasodilator and enhances cardiac contractility and renal function directly and indirectly. Many of the effects demonstrated can be accounted for by the propensity of IMD to crossreact at receptors for CGRP and AM. It is unclear whether IMD reaches such receptors at appreciable levels to activate similar responses under physiological conditions. IMD is present constitutively at very low levels in adult myocardium and vasculature: plasma IMD could primarily reflect secretion from the pituitary, where the peptide is expressed abundantly. Species variation and lack of consistency in regard to use of IMD fragments of varying length has added to the current confusion in the literature. Elucidation of receptor involvement in the actions of IMD (and AM, CGRP) has also been hampered by lack of antagonists of sufficient potency and selectivity to distinguish between each receptor subtype. Non-peptide antagonists with selectivity for CGRP₁ receptors (Doods *et al.*, 2000) represent a major advance and identification of suitable antagonists displaying selectivity for AM₁ and AM₂ receptors is urgently required. The use of such pharmacological tools cannot however by itself provide information regarding which member of this peptide family might serve as the endogenous ligand at a particular receptor population in a given cell or tissue, or indeed the extent to which family members interact at the same receptor. Functional characterization of viable strains of mice in which the IMD gene has been deleted should provide clarity regarding the physiological role of IMD; the availability of such mice is eagerly awaited.

Potentially more interesting is the robust upregulation of the prepro-IMD transcript specifically in hypertrophied and ischaemic myocardium. Oxidative stress regulates expression of transcription factors such as nuclear factor- κ B, c-jun/c-fos heterodimer (AP-1) hypoxia-inducible factor HIF-1 α and endothelial PAS domain protein 1 (Valko *et al.*, 2007). The AM and CL genes possess binding motifs for one or more of these factors within their promoter regions (Nikitenko *et al.*, 2003; Ishimitsu *et al.*, 2006) and confirmation of the presence of similar motifs within the promoter region of the IMD gene could explain induction of this gene within

cardiomyocytes in response to oxidative stress. Reactive oxygen species also modify activity of various signalling kinases implicated in hypertrophic remodelling including protein kinase C, src tyrosine kinase and mitogen-activated protein kinase all of which influence gene expression. Taken together with the beneficial effects on contractile function and attenuation of cardiomyocyte hypertrophy and ischaemia-reperfusion injury, it is likely that IMD provides the myocardium with an endogenous mechanism to counter the detrimental consequences of oxidative stress and ischaemic insult. The importance of IMD in this regard should now be examined through investigation of the consequences of cardiac-restricted deletion of the IMD gene or localized application of small interfering RNA or IMD-specific antibodies in suitable experimental models of myocardial ischaemic insult. Confirmation of a crucial beneficial role of IMD should pave the way for translational research to the clinical arena. Firstly, the potential of measuring IMD as a cardiac biomarker could be addressed. Secondly, strategies could be devised to deliver IMD in the acute or chronic setting, such as infusion of the recombinant peptide or adenoviral vector-mediated delivery of the IMD gene to prevent or lessen the extent of myocardial injury in patients with ischaemic heart disease.

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Conflict of interest

The authors state no conflict of interest.

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