Human 5-HT4 and 5-HT7 Receptor Splice Variants: Are they Important?

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Abstract: G-protein-coupled receptors (GPCRs), which are encoded by >300 genes in the human genome, are by far the largest class of targets for modern drugs. These macromolecules display inherent adaptability of function, which is partly due to the production of different forms of the receptor protein. These are commonly called ‘isofoms’ or ‘splice variants’ denoting the molecular process of their production/assembly. Not all GPCRs are expressed as splice variants, but certain subclasses of 5-HT receptors are for example, the 5-HT4 and 5-HT7 receptors. There are at least 11 human 5-HT4 and three human 5-HT7 receptor splice variants. This review describes their discovery, nomenclature and structures. The discovery that particular splice variants are tissue specific (or prominent) has highlighted their potential as future drug targets. In particular, this review examines the functional relevance of different 5-HT4 and 5-HT7 receptor splice variants. Examples are given to illustrate that splice variants have differential modulatory influences on signalling processes. Differences in agonist potency and efficacies and also differences in desensitisation rates to 5-HT occur with both 5-HT4 and 5-HT7 receptor splice variants. The known and candidate signalling systems that allow for splice variant specific responses include GPCR interacting proteins (GIPs) and GPCR receptor kinases (GRKs) which are examined. Finally, the relevance of 5-HT receptor splice variants to clinical medicine and to the pharmaceutical industry is discussed.

Key Words: Serotonin receptors, GPCR receptor isoforms, GPCR receptor splice variants, GPCR interacting proteins, desensitisation, functional intestinal disorders, irritable bowel syndrome.

INTRODUCTION

G-protein-coupled receptors (GPCRs), are by far the largest class of targets for modern drugs. These macromolecules are encoded by >300 genes in the human genome. Once formed, they are transported and embedded in the cell-surface, where they take on their functions of detecting and responding to a diverse array of ligands. Numerous diseases and disorders have been linked to mutations and polymorphisms in GPCRs and in their natural states these receptors are the targets of an increasingly large number of therapeutic agents. It has been estimated that 50% of all modern drugs and almost one-quarter of the top 200 best-selling drugs in 2000 modulate GPCR activity (see [32] for review). Studies into GPCR splice variants or isoforms is a new research area that opens the possibilities to further refine safety margins of therapeutic drugs.

A general property of GPCRs is that they have inherent adaptability built into their function, which is partly due to the production of different forms of the receptor protein. Thus, different products can be generated from a single GPCR gene by the combination of alternative forms of particular exons. This process is referred to as ‘alternative splicing’ and translated products are called ‘splice variants’ or more commonly ‘isofoms’ (see Fig. 1). Over 70% of multi-exon genes expressed in humans are alternatively spliced to form various splice variants and the proteins involved in cellular communication are common amongst examples [41]. The discovery that particular splice variants are tissue specific (or prominent) has highlighted their potential as future drug targets. Therefore, just as the discovery of different receptor subgroups opened up vast opportunities to develop new drug treatments, the discovery of splice variants promises to further expand and refine these opportunities. Examples are to be found with prostaglandin EP3 receptors, which are subject to splice variance at the C-terminus and, to date, 10 splice variants have been identified across species, six of these being expressed in man. In addition, there is evidence for a splice variant form of EP1 receptor that lacks the highly conserved seventh transmembrane domain (TM). Amongst adrenoceptors, four splice variants of the α1A-AR have been reported as well as variants of the β2-AR. Of the serotonin (5-hydroxytryptamine, 5-HT) receptors, the 5-HT3 and 5-HT7 receptors in particular are noteworthy for the production of several splice variants [40].

Fig. (1). Methods of isoform or splice variant generation. RNA editing from adenosine deamination to form inosine which is read as guanosine and so codes for different amino acids. Alternatively if additional splice junctions are present (e.g. at * but also at the ends of exons depicted by boxes), exons can be incorporated or removed (drawn) resulting in different length mRNAs and so different proteins.

Several questions arise when considering the purpose of these GPCR splice variants. Are they expressed in specific tissues? Do they modulate signalling processes in different fashions? If so, could they act as potential therapeutic targets? Here we will discuss these questions in relation to serotonin receptor splice variants with an emphasis on the distribution of 5-HT1A and 5-HT7 receptor splice variants and their potential as therapeutic targets. We will emphasise the neuropharmacological aspects. However, 5-HT receptors are widely distributed, so it is important to note that substantial information has been acquired about their splice variants by studying peripheral tissues. Although 5-HT receptor splice variants are localised to nerve elements in many tissues, others are found in effector cells, such as smooth muscle. It is also important to note, that much of our knowledge about the function of splice variants comes from cell culture techniques. Although these can be criticised for being artificial and minimalist, they are the only practical way of studying individual splice variants, because most native cells express a mixture. In this review we first outline a brief commentary on the functions and classification system of 5-HT receptors. We will also refer to some excellent and extensive reviews on other aspects of the subject which are not covered in this paper. For instance, the structural and operational characteristics of 5-HT re-
5-HT receptors are described [38] and the distribution and function of 5-HT receptors has been reviewed [48,73]. There are also extensive reviews of the medicinal chemistry and pharmacology of 5-HT4 agonists and antagonists [46] and of 5-HT1 receptor ligands [33]. In addition, the clinical relevance of 5-HT receptors in pathophysiological conditions and their targeting with therapeutic drugs is covered [29].

5-HT

5-HT has diverse physiological effects and broadly these encompass regulation of the cardiovascular, digestive and central nervous systems. Mammalian physiologists had known since the 1800s that a vasoconstrictor substance was formed when blood was allowed to clot. In 1948, Rapport named the serum vasoconstrictor ‘serotonin’ and a year later he discovered that the active substance was 5-hydroxytryptamine. 5-HT is widely distributed in nature, occurring in both plants and animals. In mammals the largest amount of 5-HT is present along the length of the gastrointestinal tract (60-90%), mainly in enterochromaffin (EC) cells of the mucosal layer. The remainder occurs in the enteric nervous system with significant distribution in the brain and spinal cord, the heart and adrenocortical cells. 5-HT is synthesised from tryptophan in the liver, where the 5-HT content of platelets is acquired mainly by the gut. It is now recognised that disturbances in the levels of 5-HT and/or the densities of its receptors contributes to the pathogenesis of many clinical conditions. Some examples are the carcinoid syndrome and gastrointestinal motility disorders in the periphery, migraine, depression, anxiety, schizophrenia, obsessive compulsive disorders, eating disorders and the serotonin syndrome centrally.

5-HT RECEPTORS

The physiological and pathophysiological effects of 5-HT are mediated by at least 14 different receptor subtypes [2,37]. This relatively large number is attributed to the long evolutionary history of the 5-HT signalling system, which pre dates the separation of vertebrates and invertebrates, some 600 million years ago. Consequently, there has been abundant time for gene duplications, followed by mutations and sequence shifts to form the different genes encoding for the different subtypes. It has been speculated that the ancestral 5-HT receptors functioned to facilitate cell to cell communication and to promote growth and differentiation. This diversified to the level of complexity now apparent in the mammalian brain (for reviews, see [5,18,71]).

All mammalian 5-HT receptors are members of the GPCR superfamily of membrane-bound receptors (also called metabotropic receptors) with the exception of the 5-HT3 receptor which is an ion channel. These are classified into seven distinct classes, or families, according to their structure, pharmacological properties and preferred effector mechanisms (see the current IUPHAR appellation of 5-HT1, 5-HT2, 5-HT3, 5-HT4, 5-HT5, 5-HT6 and 5-HT7). Some of these classes include multiple receptors, which share similar structural and effector properties, but display very different operational profiles. The lower case character denotes that the class denotation as h5-HT4(a) [14]; several other splice variants have been cloned and characterised. Following its discovery in 1997 and naming as h5-HT4(a) [14]; several other splice variants have been cloned: h5-HT4(b) [15,72], two different h5-HT4(d) splice variants [13,15], h5-HT4(e) [15], h5-HT4(f) [13,24] (formerly called (e) [58] see [46]), h5-HT4(g) [20], h5-HT4(i) [13] the only example of an internal splice variant insert occurring in the 2nd extracellular loop, and h5-HT4(j) [74] (Fig. 2A). From the viewpoint of drug discovery, the tissue distribution of these human 5-HT4 receptor splice variants shows a degree of specificity (see [46] for review).

In addition, cellular studies show that desensitisation rates of 5-HT4 receptor splice variants depends upon the GPCR kinases (GRK) present [10] and the splice variant [64].

The h5-HT4(a) is dominant in most tissues and the h5-HT4(b) is also common, so opportunities for drug discovery need to take advantage of other tissue-specific splice variants. Table I summarises the current state of knowledge. Focussing on the atrial tissue of the heart, it can be seen that it is relatively well endowed with splice variants compared to the kidney and bladder. Also, it expresses the (n) splice variant, as does the brain and oesophagus, but this is absent from the kidney, bladder, stomach, ileum and colon. The current state of knowledge shows the human small intestine and colon also express h5-HT4(a) and that this is not found in other tissues [13,15,56]. Many of the other splice variants are more widely distributed, so it is important to map both their distribution and quantity as their actual levels of expression may be low.

Interestingly, 5-HT4 receptor splice variant expression has been shown to change dramatically in cancerous tissue. Normal adrenal tissue expressed both 5-HT4(a) and 5-HT4(b) and rarely 5-HT4 splice variants could potentially be used as a basis for new drug development. For instance, discovery of heart-selective drugs is achievable if the heart is found to express a 5-HT4 splice variant as the therapeutic target that is unique to the heart or more prevalent to the heart than other organs.

The 5-HT4 receptor present in the human atrium was the first to be cloned and characterised. Following its discovery in 1997 and naming as h5-HT4(a) [14]; several other splice variants have been cloned: h5-HT4(b) [15,72], two different h5-HT4(d) splice variants [13,15], h5-HT4(e) [15], h5-HT4(f) [13,24] (formerly called (e) [58] see [46]), h5-HT4(g) [20], h5-HT4(i) [13] the only example of an internal splice variant insert occurring in the 2nd extracellular loop, and h5-HT4(j) [74] (Fig. 2A). From the viewpoint of drug discovery, the tissue distribution of these human 5-HT4 receptor splice variants shows a degree of specificity (see [46] for review).
The C-terminal tails of 5-HT$_4$ receptor splice variants have been found to directly influence their functional properties and this is most dramatically seen in their transduction of agonist responses. A notable example is renzapride, which is nearly 20 times more potent at the h5-HT$_4$(d) than at the (g) splice variants in inducing cyclic AMP formation in COS cells. Another difference is that renzapride behaves as a full agonist at the h5-HT$_4$(d), but is a partial agonist at the (g) variant [59]. One interpretation for this phenomenon is that the C-terminus regions exert different torsion forces on the conserved transmembrane loops causing different stereic presentations of the active site to its ligands. Some indirect support for this explanation comes from our observation that the different splice variants exhibit over 10-fold variations in their affinities for ligands in binding studies (Coupar, Tochon-Danguy, Irving unpublished observations). Another explanation for the functional differences between 5-HT$_4$ splice variants is that they can link to different G proteins. Experiments using human embryonic kidney (HEK) cells have shown that the potencies of 5-methoxytryptamine at 5-HT$_4$(a) and 5-HT$_4$(d) splice variants are different and that this is correlated to coupling to only G$_q$ and to G$_i/o$ plus G$_q$ proteins, respectively [63]. More recent experiments using adenoviral expression of h5-HT$_4$(a) and h5-HT$_4$(d) splice variants in rodent cardiac myocytes that do not naturally express 5-HT$_4$ receptors demonstrated that the h5-HT$_4$(d) receptor was more efficiently coupled to adenylyl cyclase [23]. In addition, it was shown that pertussis toxin potentiated the stimulatory effect of 5-HT on L-type Ca$^{2+}$ current in rat myocytes expressing the 5-HT$_4$(d) splice variant but not the 5-HT$_4$(a) [23] providing further support for the suggestion that the (b) splice variant couples to both G$_q$ and G$_i/o$ proteins.

**5-HT$_7$ Receptor Splice Variants**

The 5-HT$_7$ receptor was first cloned from human tissues in 1993 [7]. Three human 5-HT$_7$ receptor splice variants were then discovered and named, h5-HT$_7$(a), h5-HT$_7$(b) and h5-HT$_7$(d) that differ in their C-terminal tails, but share an identical sequence up to Leu 358 for the h5-HT$_7$(d) fragment [43]. The known distribution of 5-HT$_7$ receptor (or more prevalent) is summarised in Table I.

**Fig. (2).** Serotonin receptor splice variants. (A) The 11 human 5-HT$_4$ receptor splice variants. Most of the splice variants occur in the C terminal tail following the splice site coding for the amino acid leucine 358 (shown as first amino acid in each splice variant). However, splice variant h5-HT$_4$(d) occurs in the second extracellular domain and has been reported to contain the “b” splice variant tail. (B) The 3 human 5-HT$_7$ receptor splice variants which occur in the C terminal tail following the splice site coding for the amino acid leucine 432 (shown as first amino acid in each splice variant). Potential phosphorylation sites (S/T) are in bold; and * indicates a PDZ ligand.
extensively, the 5-HT7 antagonist, SB-269970, induced a partial translocation of the h5-HT7(d) variant from cytoplasm to plasma membrane. Another noted difference was that the h5-HT7(d) variant was associated with a lesser efficacy at stimulating adenylyl cyclase. As a result, it was suggested that the C terminal tail of the h5-HT7(d) splice variant, which is the longest of the three human splice variants, may contain a motif that interacts with cellular transport systems to limit the amount destined for the plasma membrane [34].

The ability of a ligand to provoke a GPCR-mediated response is measured in terms of 'efficacy' and the ligand is referred to as an 'agonist' in classical pharmacological terms. It is now apparent that a balance occurs between the molecular mechanisms controlling receptor signalling, desensitisation and resensitisation or down regulation. Hence, the selectivity of agonists may be influenced by differences in the individual rates at which their splice variants desensitise and/or interact with numerous intracellular GPCR interacting proteins (GIPs).

**Differences in Intracellular Signalling Modulated by GIPs**

The most common GIPs interact with the C terminal tails of GPCRs; and PDZ domain containing proteins are the abundant members of this class. PDZ domains were first recognised as sequence repeats contained in three separate proteins: Post synaptic density (PSD) protein PSD-95, Discs large protein (the Drosophila homologue) and tight junction protein ZO-1. PDZ proteins are involved in scaffolding multi-protein complexes and have roles in protein trafficking (see [62] for a review). PDZ proteins bind to specific conserved consensus sequences that are found at the C-terminal end of proteins. These consensus ligands have been classified as class I PDZ ligand of S/TXΩ, class II of ΩXΩ and class III contains E/DXΩ where Ω represents a hydrophobic residue and X any amino acid [62]. Many GPCRs express a PDZ ligand at the extremity of their C terminus (usually the last 3-4 amino acids). Thus the C terminal tail of GPCRs can contain a PDZ ligand at its extremity while residues upstream of the last 3-4 are important in modulating the specificity of interactions with other proteins [17]. An example of how different splice variant tails can influence protein interactions is depicted in Fig. (3). This example is based on results of proteomic studies on mouse brain where two 5-HT1 receptor splice variants with different PDZ ligands in the extremity of their C terminal tails were used as a bait ligand to isolate the interacting proteins [42].

![Image of PDZ proteins interacting with mouse 5-HT1 receptor splice variants](image)

**Fig. (3).** Example of PDZ proteins interacting with mouse 5-HT1 receptor splice variants (a) and (c). The PDZ proteins, Veli, MPP3, NHERF and SNX27a interact with the extreme C-tail of 5-HT1R which contains the amino acids SCF. NHERF interacts with the scaffold protein ezrin and is involved in localising the 5-HT1R splice variant to specific parts of the membrane and SNX27a is involved at a later stage in assisting the receptor to be targeted to endosomes for desensitisation. The 5-HT1R receptor splice variant associates with nNOS, Sec23 and CIPP (another scaffolding protein). Arrows indicate protein interactions. For further details see text.

Several serotonin receptors contain PDZ ligands at the extremity of their C-terminal tails and have been the focus of investigation. GIPs containing PDZ domains specifically regulate receptors in the serotonin receptor family. The 5-HT2A receptor contains a PDZ ligand at its extreme C terminus and directly binds to PSD-95 which augments signal transduction and inhibits agonist-induced receptor internalisation [75]. Moreover, recent proteomic experiments have demonstrated that the PDZ ligand of the 5-HT2A receptor interacts with a different set of PDZ proteins to that of the 5-HT3C receptor [12]. It is noteworthy that PSD-95 interacts with 5-HT2A and 5-HT3C but not 5-HT4a receptors indicating that the different tails containing PDZ ligands interact with specific sets of proteins [12,42]. These specific sets of different PDZ proteins probably contribute to the different signal transduction properties of these receptors. Recently, another example of a GIP involved in serotonin receptor function has been reported where a small protein, p11, is involved in specifically transporting 5-HT1a receptors to the plasma membrane [69]. These findings provide support to the tenet that "specific sets of GIPs interact with different sets of receptor splice variants and that this is also tissue dependent".

There are five 5-HT1 and one 5-HT7 receptor splice variants with canonical PDZ C-terminal extremity ligand binding sites (Fig. 2). So far, there have been no reports on GIP interactions with any 5-HT1 or 5-HT7 receptor splice variants in human tissue. However, a recent study identified 13 GIPs (mainly PDZ domain proteins) that interact with either the mouse 5-HT4a, or 5-HT4c receptors [42]. Of these proteins, 10 interact specifically with the mouse 5-HT4a receptor splice variant. One of these is sorting nexin (SNX27) which is enriched in the brain and is involved in escorting the 5-HT4a receptor splice variant to early endosomes for desensitisation. Another protein, NHERF appears to be involved in directing the 5-HT4a receptor to the microvilli area where the two proteins are co-localised with another protein called ezrin [42] that interacts with the membrane phosphatidylinositol-(4,5)-bisphosphate [9]. The 5-HT4b splice variant does not contain a PDZ domain (Fig. 2A) and did not concentrate in the microvilli area. While the 5-HT4c receptor splice variant co-localised with CIPP which is a scaffolding protein and importantly this was not seen with the 5-HT4d receptor splice variant which does not contain a PDZ ligand [42]. These exciting results do indicate that the PDZ ligand domains of 5-HT1 or 5-HT7 receptor splice variants contribute to the receptor localisation and also suggest that rates of receptor desensitisation may vary with the splice variant expressed.

**Differences in Desensitisation Rate**

The diversity in C-terminus sequences probably also contributes to the well known pharmacological fact that 5-HT induces desensitisation (tachyphylaxis) at different rates and magnitudes in different tissues. This natural phenomenon functions to limit the biological response to endogenous substances, such as 5-HT by uncoupling the GPCR from its signal transduction pathway (for reviews, see [30,57]). Desensitisation generally begins with agonist-induced phosphorylation of the GPCR by GPCR receptor kinases (GRKs). The cytosolic proteins, arrestins then bind the phosphorylated GPCRs and prevent further coupling of that GPCR with G proteins and so reduce second messenger synthesis. The arrestin-GPCR complex is internalised by endocytosis where it can be recycled back to the membrane or degraded [30,57] (Fig. 4). GRKs play a critical role in GPCR desensitisation. There are three GRK subfamilies: rhodopsin kinase containing GRK1 and 7; β adrenergic receptor kinase (GRK2 and 3); and GRK4 group (GRK4, 5 and 6). Both GRK2 and GRK5 can phosphorylate many GPCRs including 5-HT1 receptor splice variants [10] and GRK2 in particular is crucial for embryonic cardiac development [57]. The role of GRKs and receptor desensitisation has been the focus of many studies but their effects on serotonin receptors are less well documented.
Desensitisation also has the potential to lessen the beneficial effects of agonists when used as therapeutic treatments. Indeed, we have established that desensitisation of native 5-HT4 receptors occurs in vivo in the rat. We found that 5 day infusions of 5-HT (75 µg/kg per hour) induced rightward shifts of the 5-HT4 receptor-mediated concentration-effect curves to 5-HT and the partial 5-HT4 agonist, SC 53116, in the oesophagus [55]. Therefore, knowledge of the extent and mechanisms of desensitisation in human tissue is crucial to the success of drug discovery programs aimed at developing 5-HT receptor agonists. In spite of this, only two studies have investigated the desensitisation properties of h5-HT4 receptor splice variants (human c and d variants [60]; mouse a, b, e, and f variants [10]) in cell culture. On the other hand, the h5-HT4 receptor splice variants all mediate heterologous desensitisation, which seems to be induced by both the agonist 5-HT and some antagonists [45]. In addition, all h5-HT4 receptor splice variants mediate heterologous desensitisation of endogenous Gs-coupled receptors in HEK 293 cells through unknown mechanisms that are independent of cAMP dependent protein kinase activation [3,45].

CLINICAL RELEVANCE

5-HT4 Receptors

More than 50 patents have been lodged since 2000 covering potential clinical applications for 5-HT4 agonists (ISI Web of Knowledge, Derwent Innovations Index, (http://portal.isiknowledge.com/portal.cgi)). The claims are mainly for treatments of the digestive system, where there is potential for using tegaserod for the treatment of IBS is outlined next.

Irritable Bowel Syndrome

It has been said that “A good set of bowels is worth more to a man than any quantity of brains” (Josh Billings, 1818–1885). Although this quotation was intended to be humorous there is some truth which is illustrated by the symptoms of the irritable bowel syndrome (IBS), which is a common functional bowel disorder.

The serotoninergic interneurons and motor neurons [16], as well as the muscularis mucosa of the rat oesophagus, where they mediate relaxation directly [11]. In the intestines, 5-HT4 receptor activation was originally shown to enhance acetylcholine released in isolated preparations of the guinea-pig ileum [25-27]. Patents also cover treatment of CNS disorders, such as Alzheimer's disease, anorexia, drug dependence, migraine and pain. As in the digestive tract, 5-HT4 receptor function in the brain is to enhance the release of acetylcholine, but they also modulate the release of dopamine, GABA and 5-HT [8], hence the interest in 5-HT4 agonists for treating cognitive disorders. Encouraging results have been obtained in rats, where the 5-HT4 agonist, RS 67333, improved associative memory and the 5-HT4 antagonist, RS 67532, decreased memory [52]. Their potential in the treatment of anorexia is suggested from experiments using obese mice which showed that the 5-HT4 agonist, mifepristone, improved food intake [4]. None of the patents claim the novel 5-HT4 receptor agonists are splice variant-selective, however, there are patents covering the DNA of splice variants themselves. One suggests gene therapy using the nucleic acid sequence of the 5-HT4 splice variant for gastrointestinal diseases, while another the 5-HT4 splice variant for treating cardiac and bladder disorders. References to Table 1 shows there is potential for refining drug selectivity because the target organs for these conditions contain different combinations of splice variants. To date the digestive tract has received the greatest attention as an organ for treating various disorders using 5-HT4 agonists. Currently, these drugs are used to treat several types of motility disorders including gastro-oesophageal reflux disease (GORD). The most commonly used drugs were cispalmitate and metoclopramide, but cisapride has been virtually withdrawn (available on limited access) due to its ability to induce rare, but potentially fatal cardiac arrhythmias. Metoclopramide is a relatively old 5-HT4 agonist with significant affinity at other receptor types. The most selective 5-HT4 agonist to date is tegaserod, which is used to treat IBS. Consequently, the rational for using tegaserod for the treatment of IBS is outlined next.
matched control subjects [49]. The human small intestine and colon express various h5-HT4 splice variants (see Table 1, [13,15,56]). It is possible that different 5-HT4 receptor splice variants are expressed in the various locations and so contribute to the regulation of bowel functions in different ways. However, there is a lack of quantitative data for 5-HT4 receptor splice variant expression relative to the colon to substantiate such speculations.

**5-HT7 Receptors**

5-HT7 receptors have also been identified in the circular muscle of the human colon, but as yet not in nerves [39,65]. All 5-HT7 receptor splice variants are expressed in the human small intestine and colon including a certain amount of the h5-HT7(d+5) fragment [43].

The patent literature also covers the potential use of novel full and partial 5-HT7 receptor agonists. The proposed applications are for the treatments of depression, anxiety and eating disorders, correction of circadian rhythm and migraine. As with the 5-HT4 ligands, none are suggested to be splice variant-selective. These applications are well-founded given the widespread distribution of the 5-HT7 receptor in the CNS and accumulating evidence for its functional effects. This includes the findings that some antidepressants, such as amitriptyline, have relatively high affinities at the 5-HT7 receptor (see [48,50,73] for reviews). The 5-HT7 receptor was originally discovered in the CNS where particularly high expression was found in the thalamus and hypothalamus. Functional studies of these areas led to the suggestion that 5-HT7 receptors localised in the suprachiasmatic nuclei of the hypothalamus are involved in controlling circadian rhythm [51]. Although subsequent studies have generated some uncertainty, it has been established that the selective 5-HT7 receptor antagonist, SB-269970 reduces paradoxical sleep in rats [35,48]. It has also been suggested that the 5-HT7 receptor is involved in migraine (see [70] for review). Several lines of evidence support this, such as the findings that the receptor is expressed in cranial blood vessels of experimental animals and that 5-HT and 5-HT7-preferring agonists cause them to dilate. The transcript of the 5-HT7 receptor has been detected in human internal carotid and meningeal arteries, but functional evidence for its potential role in migraine is lacking. Other lines of evidence point to the involvement of the 5-HT7 receptor in hyperalgesic pain and neurogenic inflammation. The fact that prophylactic anti migraine 5-HT receptor antagonists have relatively high affinity for the 5-HT7 receptor further implicates this receptor in the abnormal vascular and neurogenic alterations that account for migraine headache.

**CONCLUDING REMARKS**

The intricate differences in structural forms of 5-HT4 and/or 5-HT7 splice variants suggest that they perform separate functions. This would have occurred over a long evolutionary period resulting in the refinement to the ancestral serotonin signalling system. Specific functions are also implied by the varying degrees of tissue-specific distribution of most splice variants and that such functions are important to the organs in which they are expressed (Table 1) and the disease state of the tissue. Refinement of function is conferred by the C terminal tails of the splice variants which allow them to interact with different GIPs, GRKs and even different G proteins. To date, the established net outcome of splice variant-specific interactions within the cell determine factors such as the rate, duration and intensity of the response to 5-HT.

Although it has been shown that some splice variants have different sensitivities to ligands, it has only been demonstrated with a small number of agonists. It remains a considerable challenge to identify and develop splice variant-selective drugs (no antagonists have yet been identified). However, the pressure to do so may increase as a result of the recent voluntary restriction of tegaserod to Special Access use only. This was necessary as IBS patients have been shown to experience a higher chance of cardiovascular events, such as heart attack, stroke, or severe heart-related chest pain. This predictable problem could be avoided if it were possible to discover an agonist with reasonable selectivity for a splice variant(s) present in the intestines, but not in the heart (e.g. 5-HT4d or 5-HT7). Future drug discovery projects may also turn to the GIPs and GRKs in order to achieve tissue-specific effects.

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