

# Leveraging the ion channel platform for lead generation in atrial fibrillation

Competition in the pharmaceutical industry requires a reduction in research and development project cycle times and, accordingly, an increase in productivity and efficiency. Pharmaceutical companies also are faced with the challenge of translating massive amounts of new genomic data into innovative drugs. More than a thousand possible new drug targets have emerged from sequencing the human genome, but currently available drugs only target approximately 500 different proteins<sup>1</sup>. In order to effectively handle data overload from genomic technologies, progress in finding chemical drug leads is needed. A good strategy is a chemical biology-driven, knowledge-based approach, whereby chemical information and expertise within target families (eg ion channels, kinases, etc) are acquired and applied. Thereby, a match of chemical and biological information within target families is achieved. **Figure 1** illustrates the concept of matching chemical and biological spaces. This article highlights methods towards such an ion channel target family approach to lead generation in atrial fibrillation.

**R**esearchers believe the human genome encodes more than 300 different ion channels, which are protein pores that control the flow of charged particles through a cell. Yet only 5% of drugs marketed today target ion channels. However, drugs targeting ion channels and transporters generate more than \$24 billion in worldwide sales per year. According to the FDA, the number of

new approved drugs targeting ion channels is equal to or even higher than that for drugs targeting proteases, polymerases and reverse transcriptases.

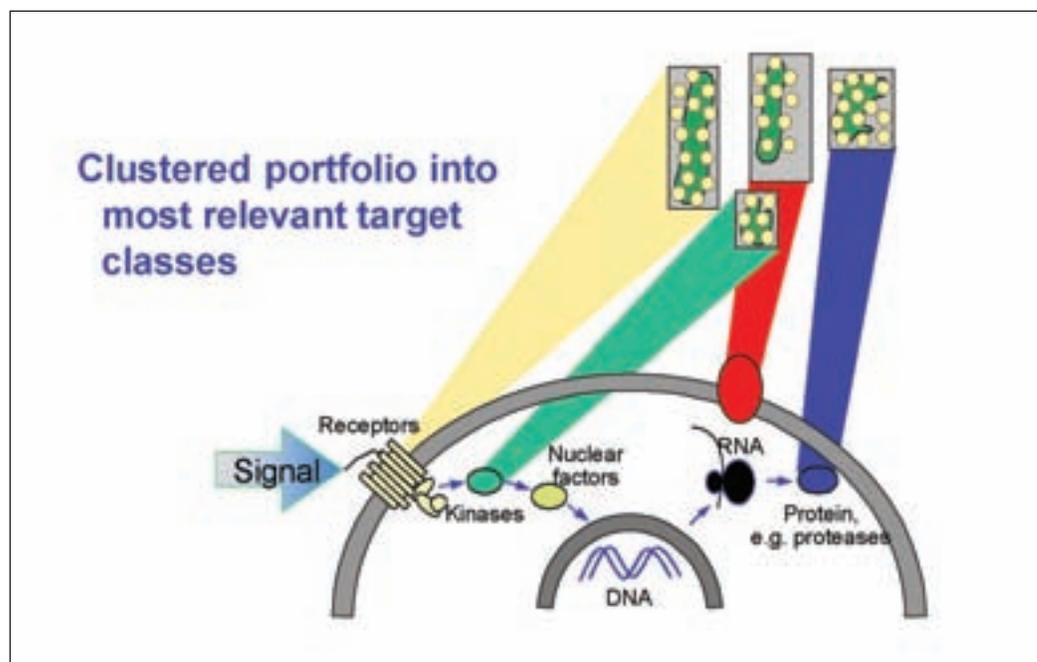
Lack of interest is not what has hindered the development of drugs targeting ion channels since many 'channelopathies' such as cystic fibrosis, diabetes, heart disease, deafness, blindness and epilepsy can all result from malfunctions in ion channels.

**By Professor Andreas Busch**

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**Figure 1**

Using the chemical biology approach to match chemical and biological spaces based on target classes



Ion channels, which are ubiquitous, often tissue specific and sensitive to pharmaceuticals, also make excellent targets because the proteins protrude out in the membrane exposed to the extracellular space. **Figure 2** gives an example of the heterogeneous ion channel targets involved in many disease states.

In the post-genomic era, progress in transcription profiling revealing the tissue specific distributions of ion channels and a greater understanding of these proteins means that ion channels will play an increasingly important role as therapeutic drug targets in the future. These advances, combined with improved cell-based assays, will make ion channels a favourable class of selective, tissue specific, drug targets.

Despite their remarkable physiological value, ion channels have remained a relatively unexploited therapeutic target class, especially in comparison to target areas such as G-protein coupled receptors or kinases. Major challenges have been the lack of high throughput screening assays to test several hundred thousand compounds against a particular ion channel and the availability of targeted libraries of candidate ion channel modulators<sup>2</sup>. Therefore, a biased ion channel library is of high interest for lead finding in the field.

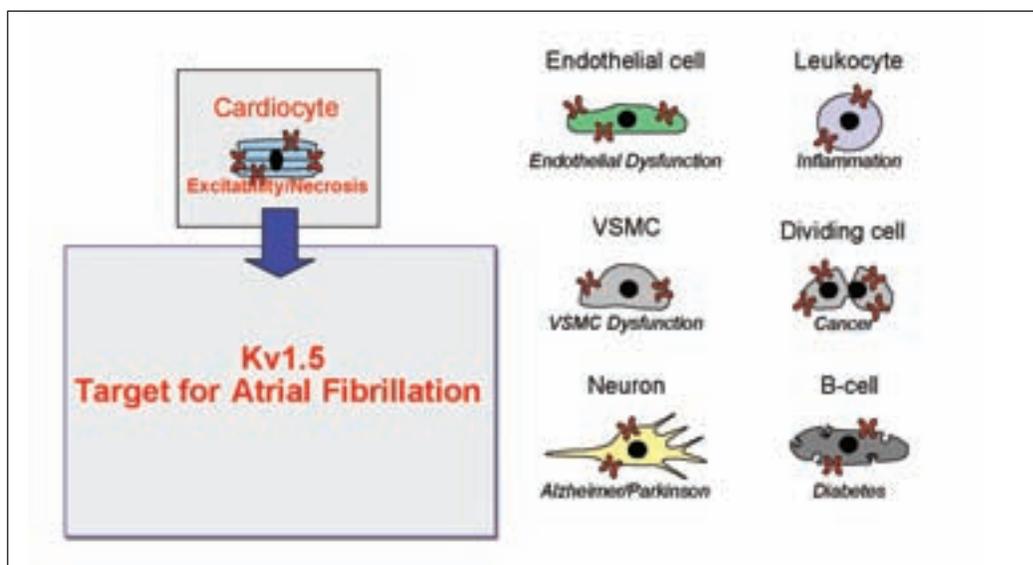
### **The Aventis ion channel library**

Molecular informatics is involved in several areas of the value chain of drug discovery, mainly in lead finding and lead optimisation through

appropriate techniques, such as high throughput screening data analysis, database mining and pharmacophore modelling<sup>3</sup>. The ion channel target family-related approach at Aventis consists of biased libraries, which were built by matching chemical and biological space. Therefore, the intersection of biological structures and functionalities with chemical structures and properties is derived to perform a knowledge-driven biased library design. This allows an extraction of common structural features for target families out of an essentially infinite chemical space. Thereby, chemical libraries could be built and enriched by preferred features of biologically active compounds.

The Aventis ion channel library was composed using public and proprietary ion channel lead compounds. Similarity searches and virtual screening based on pharmacophores and homology models yielded representatives of this library. The addition of several combinatorial libraries of ion channel-privileged scaffolds also supplemented this biased library.

As stated previously, potassium channels play critical roles in a wide variety of physiological processes, including the regulation of heart rate, muscle contraction, neurotransmitter release, neuronal excitability, insulin secretion, epithelial electrolyte transport, cell volume regulation, immune function and cell proliferation. As such, potassium channels have been recognised as potential therapeutic drug targets for many years. Today, more



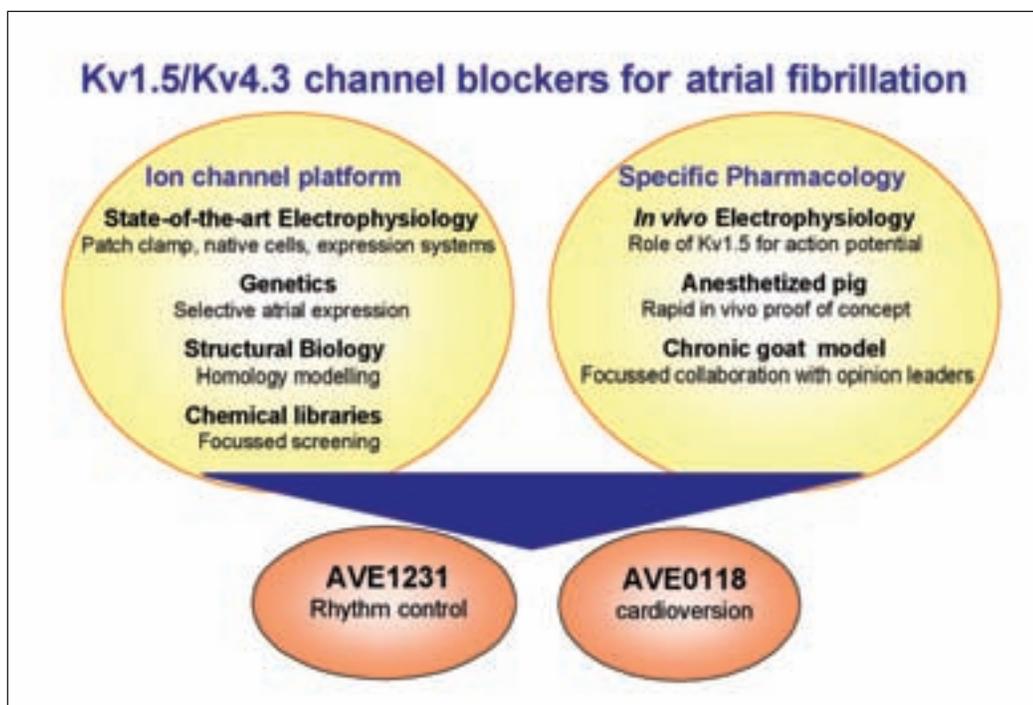
**Figure 2**  
The heterogeneous family of ion channel targets are involved in many disease states

than 80 potassium channel and potassium channel-related genes have been identified. In addition, the molecular compositions and the functions of many important channels are being elucidated. Currently, there are active pharmaceutical research and development programmes involving ion channels in a number of areas, including asthma, inflammation, arrhythmia and CNS disorders. This field represents a major growth area with the emergence of, for example, the Kv1.5 channel as a target for atrial fibrillation.

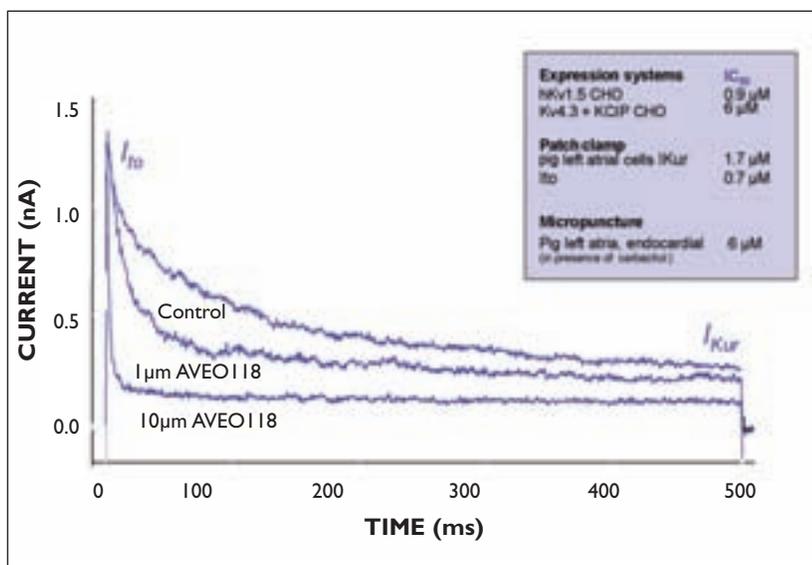
**Blocking the Kv1.5/Kv4.3 channels for the treatment of atrial fibrillation**

Atrial arrhythmias are a common cardiovascular problem. Despite several anti-arrhythmic drugs being available commercially, a medical need for safer and more efficient anti-arrhythmic treatment remains. Atrial selective compounds that terminate or prevent atrial arrhythmias while not prolonging the QTc-time would be of high value and potentially commercially successful.

In this context, Aventis has leveraged our ion



**Figure 3**  
Translating target class expertise into potentially specific and safe compounds for atrial fibrillation



**Figure 4**  
Whole-cell patch-clamp recording of pig atrial myocyte treated with control or AVE0118

channel platform to generate and investigate leads for a promising ion channel target. Figure 3 illustrates the steps Aventis has taken in translating target class expertise into potentially specific and safe compounds for atrial fibrillation. The atrial voltage-gated potassium channels Kv1.5 and Kv4.3 are regarded as promising targets, which should not be associated with ventricular side-effects unlike classical anti-arrhythmics. Our ion channel platform has aided the development of the drug candidates AVE0118 and AVE1231, two Kv1.5 and Kv4.3 blockers demonstrating efficacy in several atrial fibrillation animal models with no pro-arrhythmic effects<sup>4</sup>.

The chemistry, selectivity and structure-activity data disclosed in the literature demonstrate the anti-arrhythmic efficacy of Kv1.5/Kv4.3 blockers *in vivo*. Several studies in pig, dog or goat models have confirmed their proposed atrial selective anti-arrhythmic effect *in vivo*. AVE0118 prolongs atrial refractoriness without any effect on the QT-interval. Most of the more intensively characterised Kv1.5 blockers have turned out not to be selective but also block other ion channels. Based on the currently available data it seems that additional inhibition of Kv4.3 and KACH is beneficial for the desired anti-arrhythmic effect without hampering the ventricular safety of a Kv1.5 blocker. Figure 4 shows the effects of AVE0118 on whole-cell currents in pig atrial myocytes. The concentrations of AVE0118 are indicated in the figure. In patch clamp recordings, it can be observed that increasing concentrations of AVE0118 decreased the transient K currents ( $I_{to}$ ) and the sustained K current ( $I_{Kur}$ ) related to the Kv4.3 and Kv1.5 channels, respectively.

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### Conclusions

At Aventis, knowledge-based lead finding is being achieved by matching chemical and biological space through the collection of target family-related ligands and the identification of privileged structural motifs to build target family biased libraries for focused screening to find better hits faster. Ion channels occur as large families of related genes with cell-specific expression patterns. Ion channels have been an untapped resource because of the difficulty in studying and screening these targets. However, this may change as advances have been made over the past decade in automating relatively high-throughput screening technologies for ion channels.

Progress in transcription profiling, which is revealing the tissue-specific distributions of ion channels and a greater understanding of these proteins, means that ion channels will play an increasingly important role as therapeutic targets. These advances, combined with improved cell-based assays, will make ion channels a favourable class of selective, tissue-specific targets.

### Acknowledgement

The editorial assistance of Keith Lantz was invaluable for the preparation of this article. **DDW**

### Publisher's note

This article was written prior to the Sanofi-Aventis merger and therefore the views expressed by the author may not reflect the future strategy of the new organisation.

*Professor Andreas Busch is presently Global Head of the Therapeutic Department Cardiovascular Diseases at Sanofi-Aventis. An internationally recognised scientist in the field of pharmacology and physiology, with more than 100 publications in peer-reviewed international journals, he is Expert Pharmacologist of the German Society for Pharmacology and Toxicology and Extraordinary Professor of Pharmacology at the University of Frankfurt. Along with these academic distinctions, Professor Busch has also gained great consideration from his peers from industry, with important achievements in strategy development and implementation, restructuring and productivity increase. As Head of Cardiovascular Research, he holds the responsibility for the overall portfolio from target identification to the identification of developmental candidates.*