Clinical applications of patient-specific induced pluripotent stem cells in cardiovascular medicine

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ABSTRACT
The emergence of induced pluripotent stem cell (iPSC) technology has had a great impact on the field of medicine ever since the groundbreaking discovery in 2006 that overexpression of four specific transcription factors was able to turn back the developmental clock of somatic cells into an embryonic-like state. The resulting iPSCs carry the developmental potential of human embryonic stem cells (hESC) without the embryo and have been heralded as a powerful tool to study development and disease. This technology has made it possible for the first time for researchers to transform end-differentiated cells from a particular individual into another cell type that remains specific to that individual, paving the way for novel methods of in vitro disease modelling and therapeutic applications. This paper reviews some of the key areas in cardiovascular medicine in which iPSC technology has been applied and discusses the future directions and ongoing challenges ahead in this exciting field.

INTRODUCTION
The emergence of induced pluripotent stem cell (iPSC) technology has had a great impact on the field of medicine ever since groundbreaking work by Yamanaka and colleagues demonstrated that the overexpression of four genes (or ‘Yamanaka factors’: Oct4, Sox2, Klf4, c-Myc) was able to turn back the developmental clock of somatic cells into an embryonic-like state.1 The resulting iPSCs carry the developmental potential of human embryonic stem cells (hESC) without the embryo and have been heralded as a powerful tool to study development and disease. The importance of Yamanaka’s pioneering work is that, for the first time, researchers have a tool to transform end-differentiated cells from a particular individual into another cell type that remains specific to the individual while, at the same time, circumventing many of the ethical dilemmas that have hampered traditional embryonic stem cell research to date. The past few years have witnessed an exponential increase in the number of publications on iPSCs, with a number of genetic cardiovascular diseases being successfully modelled in vitro using this technology.2,3 Researchers in this fast-moving field are also moving towards other applications of iPSCs, including regenerative medicine and drug testing and discovery.4,5 There is therefore enormous potential for this new technology, both in a greater understanding of disease mechanisms and improved treatments. Moreover, the fact that the iPSCs are produced from a patient’s own tissue means that the results of in vitro testing are directly relevant to that individual, allowing the new era of ‘personalised medicine’ to become more of a reality. At this 5-year mark after the initial iPSC discovery we revisit what has been achieved to date in relation to cardiovascular medicine, examine the future directions and critically appraise the ongoing issues and challenges ahead that need to be overcome prior to widespread clinical application.

GENERATION OF PATIENT-SPECIFIC IPSCS AND CARDIOMYOCYTES
Several methods have been employed to reprogramme somatic cells into iPSCs, which can broadly be divided into integrating or non-integrating methods.3 Integrating methods involve viral delivery (using retro- or lentviruses) and integration of the Yamanaka factors into the host genome while non-viral non-integrating methods involve the transient expression of reprogramming factors (eg, episomal plasmid vectors, minicircle vectors, RNA and protein delivery) without genomic integration. In addition, some laboratories have reported success using viral non-integrating methods. Methods that involve genomic integration carry several disadvantages, including random integration into the host genome resulting in the expression of endogenous genes and the development of a tumorigenic phenotype. This may not be a major issue if the iPSCs are used for disease modelling and in vitro testing, but presents a major barrier to use of these cells as a therapeutic option, as will be discussed later. Non-integrating methods are therefore a more suitable option for iPSC-based therapeutic applications. In our laboratory we have successfully reprogrammed patients’ dermal fibroblasts into iPSCs using both integrating and non-integrating methods. A schematic diagram of the stages involved for the generation of patient-specific iPSCs and cardiomyocytes from dermal fibroblasts using the viral integrating method is shown in figure 1.

iPSC colonies can be differentiated into functional cardiomyocytes using a variety of methods, which are very similar to those traditionally employed to produce cardiomyocytes from hESCs since both iPSCs and hESCs share very similar characteristics and differentiation potential. Currently, the most common method of generating cardiomyocytes from iPSCs is the embryoid body (EB) differentiation system which coaxes the iPSCs to differentiate into the cardiac lineage.3 Light microscopic images of iPSC-derived cardiomyocytes (iPSC-CMs) produced from patient dermal fibroblast samples are shown in figure 2. Although the iPSC-CMs show many of the functional...
characteristics of normal human cardiomyocytes (discussed below), some other aspects of these cells, including their morphology, remain relatively immature and 'fetal-like', as can be seen in figure 2.

FUNCTIONAL CHARACTERISATION OF IPSC-DERIVED CARDIOMYOCYTES

Before iPSC-CMs can be considered for use as disease models or in regenerative medicine, the cells produced need to be confirmed to exhibit the essential functional characteristics of human cardiomyocytes. The cellular phenotype of iPSC-CMs can be characterised using several methods including electrophysiological and molecular biological techniques. Whole cell patch clamping can be performed on individual cardiomyocytes after cells are dissociated from the EBs, allowing for measurement of specific ionic currents and action potential (AP) profiles. Three types of AP morphologies can usually be recorded from iPSC-CMs: ‘atrial-like’, ‘ventricular-like’ and ‘nodal-like’ morphologies (figure 3A). In our experience, the majority of cardiomyocytes produced using viral-based methods of reprogramming (approximately 65%) exhibit a ventricular-like AP profile. The iPSC-CMs appear to respond to known drugs affecting the AP in the expected way—for example, the AP is prolonged in the presence of the delayed rectifier potassium current (IKr) blocker E-4031 (figure 3C). Electrophysiological properties of the iPSC-CMs can also be evaluated at the multicellular level using microelectrode array mapping techniques; areas within EBs observed to be contracting are dissected and placed on microelectrode array plates where field potentials and morphologies across the cells of interest can be recorded. Traditional methods of assessing intracellular Ca²⁺ handling, such as use of Ca²⁺-sensitive fluorescent indicators and confocal microscopy, can be applied to iPSC-CMs to confirm the presence of intact and functional Ca²⁺ handling apparatus (figure 3D). Molecular biological techniques to confirm functionality of the iPSC-CMs include immunocytochemistry, phosphoproteomic assays and quantitative reverse transcriptase PCR (qRT-PCR) to identify molecular targets and expression levels of key proteins found in normal functional cardiomyocytes.

Figure 2  Light microscopic images of typical dermal fibroblasts, human induced pluripotent stem cells (iPSCs) and iPSC-derived cardiomyocytes (iPSC-CMs). (A) Parental human dermal fibroblasts (FB) in culture (magnification ×40). (B) Colony of human iPSCs derived from dermal fibroblasts via retroviral-based reprogramming using Yamanaka factors (magnification ×40). (C) Small cluster of contracting cardiomyocytes (iPSC-CMs) dissociated from the contracting embryoid bodies derived from iPSCs (magnification ×200).

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phenotype ‘in the petri dish’ and thus establish an in vitro model of the disease. A number of pioneering proof-of-concept studies have been published over the past year in patients with inherited arrhythmogenic diseases, most notably different subtypes of long QT syndrome, which confirm that patient-specific iPSC-CMs can indeed recapitulate the clinical phenotype in vitro and thus represent a potential and exciting new method for disease modelling. These models will allow researchers to better understand disease mechanisms as well as to investigate the effects of different compounds on the cellular electrophysiology of the cells, allowing for novel drug discovery or treatments. Table 1 summarises the studies in humans to date for which iPSC technology has been used to generate in vitro models of a number of cardiovascular diseases.

Modelling the long QT syndrome

Moretti et al were the first to show that iPSC-CMs can recapitulate the disease phenotype in humans. They identified an autosomal dominant inheritance of a 596G→A missense mutation in the KCNQ1 gene within a family which has previously been shown to be associated with long QT syndrome type 1 (LQTS1). Electrophysiological parameters in iPSC-CMs generated from two patients with LQTS1 were compared with those generated from unrelated healthy control subjects using whole-cell patch clamping. AP duration was found to be significantly prolonged and the repolarisation velocity reduced in LQTS1 iPSC-CMs compared with controls. Furthermore, single cell electrophysiological analysis showed a reduction specifically in the I_{Ks} current in ventricular LQTS1 iPSC-CMs compared with controls with no significant differences in the other currents measured.

Similarly, two other groups reported their findings on the use of iPSC-CM disease models for LQTS2. Itzhaki et al obtained dermal fibroblasts from a patient with LQTS2 harbouring a known KCNH2 gene mutation (affecting I_{Kr}) and showed that AP duration was prolonged and repolarisation velocity reduced in LQTS2 iPSC-CMs compared with controls. They measured and verified that I_{Kr} was significantly reduced in LQTS2 iPSC-CMs. The investigators further tested drugs that could either relieve or aggravate the clinical phenotype of LQTS2 and found AP duration to be prolonged with E-4031, which was associated with new early after depolarisations (EADs) or an increase in the number or complexity of EADs. The potential therapeutic effects of nifedipine and the K_{ATP} channel opener pinacidil (which augments the outward potassium current) was tested in this in vitro model; the drugs were demonstrated to shorten the AP duration and abolish EADs. Matsa et al also successfully generated iPSC-CMs from a patient with LQTS2 with a known KCNH2 mutation and her mother, a carrier for the same mutation. The investigators further tested drugs that could either relieve or aggravate the clinical phenotype of LQTS2 and found AP duration to be prolonged with E-4031, which was associated with new early after depolarisations (EADs) or an increase in the number or complexity of EADs. The potential therapeutic effects of nifedipine and the K_{ATP} channel opener pinacidil (which augments the outward potassium current) was tested in this in vitro model; the drugs were demonstrated to shorten the AP duration and abolish EADs. Matsa et al also successfully generated iPSC-CMs from a patient with LQTS2 with a known KCNH2 mutation and her mother, a carrier for the same mutation. The investigators further tested drugs that could either relieve or aggravate the clinical phenotype of LQTS2 and found AP duration to be prolonged with E-4031, which was associated with new early after depolarisations (EADs) or an increase in the number or complexity of EADs. The potential therapeutic effects of nifedipine and the K_{ATP} channel opener pinacidil (which augments the outward potassium current) was tested in this in vitro model; the drugs were demonstrated to shorten the AP duration and abolish EADs. Matsa et al also successfully generated iPSC-CMs from a patient with LQTS2 with a known KCNH2 mutation and her mother, a carrier for the same mutation. The investigators further tested drugs that could either relieve or aggravate the clinical phenotype of LQTS2 and found AP duration to be prolonged with E-4031, which was associated with new early after depolarisations (EADs) or an increase in the number or complexity of EADs. The potential therapeutic effects of nifedipine and the K_{ATP} channel opener pinacidil (which augments the outward potassium current) was tested in this in vitro model; the drugs were demonstrated to shorten the AP duration and abolish EADs. Matsa et al also successfully generated iPSC-CMs from a patient with LQTS2 with a known KCNH2 mutation and her mother, a carrier for the same mutation. The investigators further tested drugs that could either relieve or aggravate the clinical phenotype of LQTS2 and found AP duration to be prolonged with E-4031, which was associated with new early after depolarisations (EADs) or an increase in the number or complexity of EADs. The potential therapeutic effects of nifedipine and the K_{ATP} channel opener pinacidil (which augments the outward potassium current) was tested in this in vitro model; the drugs were demonstrated to shorten the AP duration and abolisi...
Table 1

| Disease modelled      | Methods used to evaluate cellular phenotype                                                                 | Number of patients in study | Controls used                                                                 | Reference |
|-----------------------|---------------------------------------------------------------------------------------------------------------|-----------------------------|-------------------------------------------------------------------------------|-----------|
| LQTS type 1 (father and son from affected family) | Whole cell patch clamp; immunohistochemistry                                                                | 1                           | 1 healthy individual                                                          | 9         |
| LQTS type 2 (mother and daughter) | Whole cell patch clamp; microelectrode recordings; K+ channel blockers and openers                           | 2                            | 2 healthy individuals                                                          | 10        |
| LQTS type 2 (unrelated) | Whole cell patch clamp; control cell lines and genetically unrelated hESC-derived fibroblasts               | 2 (unrelated)                | 2 healthy individuals                                                          | 12        |
| Ca 2+ (unrelated)     | Microscopic morphometry; immunocytochemistry; antibody array and western blot analyses                        | 2 (unrelated)                | 2 healthy individuals                                                          | 13        |

In vitro abnormality of patient-derived iPSC-CMs

#### Disease

**LQTS type 1**
-**Reprogramming method:** Retroviral integration
-**Gene mutation:** R190Q mutation in KCNQ1
-**Controlled phenotype:** APD prolongation, decrease in IKs

**LQTS type 2**
-**Reprogramming method:** Lentiviral integration
-**Gene mutation:** A614V mutation in the KCNH2 gene
-**Controlled phenotype:** APD prolongation, increased after depolarisations

**Ca 2+**
-**Reprogramming method:** Retroviral integration
-**Gene mutation:** G1681A mutation in the CACNA1C gene
-**Controlled phenotype:** APD prolongation, increased after depolarisations

**Ca 2+ (unrelated)**
-**Reprogramming method:** Retroviral integration
-**Gene mutation:** G406R mutation in the PTEN gene
-**Controlled phenotype:** APD prolongation, increased after depolarisations

**LEOPARD syndrome**
-**Reprogramming method:** Retroviral integration
-**Gene mutation:** R144W mutation in the TGFBR1 gene
-**Controlled phenotype:** Increased arrhythmogenicity in ARVC, myocardial disarray in HCM

APD, action potential duration; hESC, human embryonic stem cell; iPSC-CM, induced pluripotent stem cell-derived cardiomyocytes; LQTS, long QT syndrome.

In theory, other cardiovascular diseases with a hereditary component may be modelled using iPSC technology providing the iPSC-CMs display the key phenotypic features associated with the disease of interest. However, the process becomes more complex and difficult with increased heterogeneity of the disease (both phenotypically and genetically) or with greater involvement of external factors during the natural history of the disease. LEOPARD syndrome, an autosomal dominant developmental disorder in which the PTPN11 gene encoding a SHP2 phosphatase is mutated, and the associated hypertrophic cardiomyopathy (HCM) phenotype of the disease has been modelled successfully using iPSC technology. The investigators found that iPSC-CMs generated from two patients with LEOPARD syndrome had a larger median surface area, a greater degree of sarcomeric assembly and a greater degree of NFAT nuclear localisation than iPSC-CMs from control patients without disease. Protein molecular targets that could be potentially implicated in the LEOPARD syndrome phenotype were identified through proteomic assays, and certain proteins were found to be more abundant or had increased phosphorylation in cardiomyocytes derived from patients with LEOPARD syndrome compared with control subjects. Our group has successfully generated iPSC-CMs from a patient with arrhythmogenic right ventricular cardiomyopathy (ARVC) harbouring a PKP2 mutation and are in the process of modelling the disease in vitro. However, significant hurdles still exist in modelling the more complex cardiovascular diseases using iPSC technology. These include: (1) difficulty in ensuring a purified cardiomyocyte population from iPSCs through standard cardiomyocyte differentiation protocols; (2) the complexities of reproducing a heterogeneous disease phenotype (e.g., fibrofatty change and increased arrhythmogenicity in ARVC, myofibril disarray in HCM) which may involve other systemic factors in vitro using only cardiomyocytes; and (3) limitations of modelling essentially adult-onset diseases (for certain types of cardiomyopathies) using iPSC-CMs with a predominantly fetal-like phenotype. It is likely that further in vitro manipulation of iPSC-CMs will be required to model the more complex diseases successfully. This may involve measures such as over-expression of apolipoprotein-A1 cDNA or enhancement of the syndactyly and autism. They analysed time-lapse images of spontaneously contracting EBs and found that Timothy syndrome EBs contracted at a slower rate and more irregularly than control EBs. They also found that the L-type channel current in iPSC-CMs derived from the patients with Timothy syndrome had significantly reduced voltage-dependent inactivation compared with control cells. Only ventricular-like iPSC-CMs from patients with Timothy syndrome had APs that were significantly prolonged compared with controls, while atrial and nodal Timothy syndrome iPSC-CMs did not. This is in contrast to the study by Moretti et al on LQTS1 iPSC-CMs in which both ventricular and atrial-like iPSC-CMs displayed the disease phenotype. In addition, Yazawa et al observed that arrhythmias and delayed depolarisations were already evident in spontaneously beating Timothy syndrome iPSC-CMs while arrhythmias were only elicited under condition of adrenergic stress with the application of isoproterenol in LQTS1. Given that Cav1.2 plays a role in Ca2+-induced Ca2+ release from the sarcoplasmic reticulum, the investigators demonstrated that Cav1.2 inactivation in Timothy syndrome could affect the timing and amplitude of ventricular Ca2+ release, leading to larger and more prolonged Ca2+ elevations.

**Modelling other inherited cardiovascular diseases**

In theory, other cardiovascular diseases with a hereditary component may be modelled using iPSC technology providing the iPSC-CMs display the key phenotypic features associated with the disease of interest. However, the process becomes more complex and difficult with increased heterogeneity of the disease (both phenotypically and genetically) or with greater involvement of external factors during the natural history of the disease. LEOPARD syndrome, an autosomal dominant developmental disorder in which the PTPN11 gene encoding a SHP2 phosphatase is mutated, and the associated hypertrophic cardiomyopathy (HCM) phenotype of the disease has been modelled successfully using iPSC technology. The investigators found that iPSC-CMs generated from two patients with LEOPARD syndrome had a larger median surface area, a greater degree of sarcomeric assembly and a greater degree of NFAT nuclear localisation than iPSC-CMs from control patients without disease. Protein molecular targets that could be potentially implicated in the LEOPARD syndrome phenotype were identified through proteomic assays, and certain proteins were found to be more abundant or had increased phosphorylation in cardiomyocytes derived from patients with LEOPARD syndrome compared with control subjects. Our group has successfully generated iPSC-CMs from a patient with arrhythmogenic right ventricular cardiomyopathy (ARVC) harbouring a PKP2 mutation and are in the process of modelling the disease in vitro. However, significant hurdles still exist in modelling the more complex cardiovascular diseases using iPSC technology. These include: (1) difficulty in ensuring a purified cardiomyocyte population from iPSCs through standard cardiomyocyte differentiation protocols; (2) the complexities of reproducing a heterogeneous disease phenotype (e.g., fibrofatty change and increased arrhythmogenicity in ARVC, myofibril disarray in HCM) which may involve other systemic factors in vitro using only cardiomyocytes; and (3) limitations of modelling essentially adult-onset diseases (for certain types of cardiomyopathies) using iPSC-CMs with a predominantly fetal-like phenotype. It is likely that further in vitro manipulation of iPSC-CMs will be required to model the more complex diseases successfully. This may involve measures such as over-expression of apolipoprotein-A1 cDNA or enhancement of the syndactyly and autism. They analysed time-lapse images of spontaneously contracting EBs and found that Timothy syndrome EBs contracted at a slower rate and more irregularly than control EBs. They also found that the L-type channel current in iPSC-CMs derived from the patients with Timothy syndrome had significantly reduced voltage-dependent inactivation compared with control cells. Only ventricular-like iPSC-CMs from patients with Timothy syndrome had APs that were significantly prolonged compared with controls, while atrial and nodal Timothy syndrome iPSC-CMs did not. This is in contrast to the study by Moretti et al on LQTS1 iPSC-CMs in which both ventricular and atrial-like iPSC-CMs displayed the disease phenotype. In addition, Yazawa et al observed that arrhythmias and delayed depolarisations were already evident in spontaneously beating Timothy syndrome iPSC-CMs while arrhythmias were only elicited under condition of adrenergic stress with the application of isoproterenol in LQTS1. Given that Cav1.2 plays a role in Ca2+-induced Ca2+ release from the sarcoplasmic reticulum, the investigators demonstrated that Cav1.2 inactivation in Timothy syndrome could affect the timing and amplitude of ventricular Ca2+ release, leading to larger and more prolonged Ca2+ elevations.
Utilities until such methods are optimised and further research is performed in the area of maturation of iPSC-CMs, modelling of the more complex cardiovascular diseases including the cardiomyopathies will remain a major challenge.

APPLICATIONS IN DRUG TESTING AND DISCOVERY

In vitro drug screening
Part of the process of drug development and testing of new products for clinical use includes the demonstration that the product does not have any significant cardiac toxicities or proarrhythmic actions. Early and accurate detection of potential cardiac toxicities could save the pharmaceutical industry millions of dollars, avoiding additional funds being wasted to develop the drug further. There is an increasing demand for good human models of healthy as well as diseased hearts, given the limitations of current animal models that are currently used in drug testing and fundamental differences in cardiac electrophysiology between animals and humans. Recent studies have demonstrated that hESC-CMs may represent a novel platform for electrophysiological drug screening as the cells respond to a number of cardiac and non-cardiac drugs in the predicted way.\(^{19 20}\)

Applications in personalised medicine
As demonstrated from the human cardiac disease models generated so far, several drugs have been tested on patient-specific iPSC-CMs and found either to ameliorate or alleviate the disease phenotype.\(^ {9-11}\) The iPSC-CM disease model allows the concept of ‘personalised medicine’ to be put forward as these cells will in theory provide information on the effects of a particular drug on the individual from which the cells were obtained. Thus, using this technology, clinicians and scientists may be able to ascertain the cardiac response or side effects of an individual to a new drug in vitro before the drug is actually tested out in the individual. Further work needs to be done before this application can be realised, including evidence that the iPSC-CMs generated faithfully reproduce the clinical phenotype and response to drugs in vitro (in both normal and diseased individuals) and improved methods of iPSC-CM generation, since current methods are too laborious and time-consuming for routine clinical application. However, this remains an exciting and novel application of iPSC technology which is not possible with other currently available methods.

Another promising future application of iPSC technology is the concept of targeted gene modification of patient-specific iPSCs using homologous recombination techniques and zinc finger nucleases (ZFNs) with the exciting prospect of correcting disease-causing mutations.\(^ {21}\) This strategy has several potential advantages over conventional gene therapy, including avoidance of the need for immunosuppression and elimination of the risk of insertional mutagenesis caused by therapeutic vectors. The technique has recently been successfully applied in iPSCs to correct the genetic mutations responsible for A1-antitrypsin deficiency,\(^ {22}\) and sickle cell anaemia,\(^ {23}\) although it is still currently in its infancy and thus remains challenging and inefficient.

Applications in regenerative medicine

Myocardial repair
To date, a wide variety of different stem cell types have been investigated for their ability to repair the infarcted heart in both animal studies and human clinical trials.\(^ {24}\) The emergence of iPSC technology adds an additional cell source that may have a potential clinical application in this area, although research in regenerative medicine using these cells is still at an early stage.\(^ {25}\) The major advantage of iPSCs over other cell types for cell transplantation is the fact that these cells are patient-specific, thereby circumventing the important issue of tissue rejection associated with transplantation procedures. However, enthusiasm for use of iPSC technology in regenerative medicine has been hampered by the increasingly recognised problems of the unpredictable nature and behaviour of transplanted iPSCs and the potential for these cells to cause harm. Additional issues such as the potential teratogenicity of undifferentiated cells and proarrhythmia of iPSC-CMs need to be addressed before clinical grade iPSCs can be used in regenerative medicine.

To date, several disease models such as sickle cell anaemia,\(^ {26}\) Parkinson’s disease,\(^ {27}\) and haemophilia A\(^ {28}\) have been successfully treated using iPSC-based therapy. In relation to cardiovascular medicine, Nelson et al showed that intramyocardial delivery of iPSCs into the infarcted hearts of mice following ligation of the left anterior descending artery yielded progeny that properly engrafted without disrupting cytoarchitecture.\(^ {29}\) Furthermore, the investigators found that iPSC treatment restored contractile performance, ventricular wall thickness and electrical stability, while also achieving in situ regeneration of cardiac, smooth muscle and endothelial tissue. In another proof-of-concept study, Mauritz et al showed that injection of iPSC-derived fetal liver kinase-1 (Flk-1) positive progenitor cells into the ischaemic myocardium of left anterior descending artery-ligated mice improved cardiac function following myocardial infarction and that the Flk-1 positive cells differentiated into cardiovascular lineages in vitro and in vivo.\(^ {30}\)

Biological pacemakers
Another potential and exciting application of iPSC technology in cardiovascular medicine is in the development of cell-based therapies for biological pacing. The advent of electronic pacemakers has transformed the treatment and lives of many patients suffering from symptomatic bradyarrhythmias over the past few decades. However, a number of important limitations still exist with permanent pacemakers, including the need for a surgical procedure and its associated risks, the ongoing risk of infection and the need for frequent generator changes and possible lead changes over the lifetime of the individual (this becomes more of an issue the younger the patient is at first implant). The concept of a ‘biological pacemaker’ which can integrate with the host cardiac tissue, generate the electrical impulses required by the individual (whose intrinsic pacemaker function is defective/inadequate) and physiologically adapt to the cardiac requirements of that individual is certainly an attractive one. One of the first major reports on the use of stem cells for biological pacing involved the implantation of cardiomyocytes derived from hESCs into the ventricles of pigs with complete heart block.\(^ {31}\) The investigators performed histological examination and detailed 3D-electroanatomical mapping to show that the hESC-CMs integrated with the host cardiac tissue and were able to pace the swine hearts at a higher rate than the back-up rate set by the electronic pacemaker. Similar findings were obtained by another group using genetically engineered hESCs transplanted into guinea pig hearts.\(^ {32}\) This approach, however, requires immunosuppression if applied to humans since the hESCs are not obtained from the host. The use of patient-specific iPSC-CMs may overcome this problem, although other potential issues regarding tumorigenicity,
Table 2  Summary of current and future applications of iPSC technology in relation to cardiovascular medicine and ongoing challenges ahead

| Cardiovascular application | Advantage of using iPSC-technology | Ongoing challenges |
|----------------------------|-----------------------------------|-------------------|
| Disease modelling          | iPSC-CMs carry same genetic mutation allowing rarer genetic diseases to be modelled | Relatively immature phenotype |
| Drug testing and discovery | Unlimited source of human cardiomyocytes | Relatively immature phenotype |
| Regenerative medicine      | No need for immunosuppression | Need to improve efficiency of iPSC-CM production and upscaling of cells |

Summary of current and future applications of iPSC technology in relation to cardiovascular medicine and ongoing challenges ahead

Table 2 summarises the current and future applications of iPSC technology in relation to cardiovascular medicine and the ongoing challenges ahead. In the immediate future the most promising application of iPSCs in cardiovascular medicine is in the areas of disease modelling and in vitro drug testing. However, an important issue for researchers to address is in designing strategies that will allow iPSC-CMs to become more mature in vitro and thus more applicable to human disease modelling. Other challenges, particularly in relation to the potential use of iPSCs in regenerative medicine, include the issues of teratogenicity and potential proarrhythmias. In addition, current methods of iPSC reprogramming and differentiation into functional cardiomyocytes are still relatively slow, labour intensive and inefficient. In our laboratory, for example, it still takes approximately 2–4 months to transform somatic undifferentiated iPSCs to form teratomas.

CONCLUSIONS

Differentiation of human iPSCs in vitro into functional cardiomyocytes is now becoming a process which can be effectively manipulated and directed. New and improved protocols for cardiomyocyte differentiation of iPSCs are rapidly accumulating in the literature.45 44 Generation of other cell types related to the heart and cardiovascular system using iPSC technology is still in its infancy, although preclinical proof-of-concept studies have shown some potential for the use of such cells in the treatment of peripheral vascular disease45 and valvular heart diseases.46 Table 2 summarises the current and future applications of iPSC technology in relation to cardiovascular medicine and the ongoing challenges ahead. In the immediate future the most promising application of iPSCs in cardiovascular medicine is in the areas of disease modelling and in vitro drug testing. However, an important issue for researchers to address is in designing strategies that will allow iPSC-CMs to become more mature in vitro and thus more applicable to human disease modelling. Other challenges, particularly in relation to the potential use of iPSCs in regenerative medicine, include the issues of teratogenicity and potential proarrhythmias. In addition, current methods of iPSC reprogramming and differentiation into functional cardiomyocytes are still relatively slow, labour intensive and inefficient. In our laboratory, for example, it still takes approximately 2–4 months to transform somatic undifferentiated iPSCs to form teratomas.

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