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Stem cell-derived cell cultures and organoids for protozoan parasite propagation and studying host-parasite interaction

Christian Klotz, Toni Aebischer and Frank Seeber

FG 16 Parasitologie, Robert Koch Institut, 13353 Berlin, Germany

Address for correspondence: Frank Seeber, FG 16, Robert Koch Institut, Nordufer 20, 13353

Berlin, Germany. Tel: +49-30-18754-2856. Fax: +49-30-18754-2615. E-mail: see-

berf@rki.de

Summary

Possibilities to study the biology of human protozoan parasites and their interaction with the host remain severely limited, either because of non-existent or inappropriate animal models or because parasites cannot even be cultured in vitro due to strict human-host specificity or physiology. Here we discuss the prospects of using induced pluripotent stem cell (iPSC)-derived culture systems including organoids as a strategy to address many of these experimental bottlenecks. iPSCs already allow the generation of differentiated cell cultures for many human organs, and these cells and derivatives are amenable to reverse genetics in combination with advanced tools for genetic manipulation. We present examples of blood, neuron, liver, and intestine-dwelling protozoa, i.e. *Plasmodium falciparum*, *Toxoplasma gondii* and *Giardia duodenalis*, where iPSCs or organoids would allow addressing questions of cell and developmental biology, immunology, and pharmacology in unprecedented ways. Starting points and resources for iPSC experimentation are briefly discussed.

Keywords

Induced pluripotent stem cells; organoids; intestinal infection; *Giardia duodenalis*; *Toxoplasma gondii*; *Plasmodium falciparum*

Introduction

The ability to propagate pathogens in vitro in order to study their biology and interaction with their host is a prerequisite for understanding infectious diseases, and this is as important today as it was for Robert Koch when he first identified and cultivated *Bacillus anthracis* as the etiological agent causing anthrax (Koch, 1876). The study of protozoan parasites is no exception to this. Although a substantial body of literature exists describing in vitro methods for their propagation (e.g. Taylor and Baker, 1987; Visvesvara and Garcia, 2002) a large number of problems and restrictions remain.

Some parasites cannot be cultivated in vitro at all, e.g. *Cyclospora cayetanensis* (Ortega and Sanchez, 2010); for others such as *Cryptosporidium* sp. (Arrowood, 2002) or *Giardia duodenalis* (Benere et al., 2010), existing in vitro or in vivo systems are unsatisfactory, and axenic culture systems that would allow one to dissect biochemical dependencies for growth are not available for most medically important parasites, including all Apicomplexa. Host specificity is another important obstacle against the propagation of protozoan, in particular intracellular parasites, such as *Plasmodium* sp. or *Eimeria* sp., or for studying host species-restricted life cycle segments such as sexual reproduction of the apicomplexan parasites *Toxoplasma gondii* and *Neospora caninum*, which only occur in feline or canine hosts, respectively. The recent progress in stem cell biology, i.e. the in vitro generation of pluripotent stem cells and organoids from those, holds great promise as a solution for a number of the aforementioned problems in the study of specific, mainly human host-parasite combinations. In this review we aim to highlight some of the opportunities and remaining challenges of these developments for the study of protozoan parasites by focusing on select examples.

What are stem cells, and how can they be generated in vitro?

In 1998 the propagation of pluripotent cells from human blastocysts, cultured on mouse embryonic fibroblasts in serum-containing media, was reported (Thomson et al., 1998). Thus, a protocol became available for the derivation of human – in this case embryonic – stem cells, i.e. cells with the capacity for long-term self-renewal and the potential to differentiate into the components and derivatives of all three embryonic germ layers. While this was a technical breakthrough, it immediately provoked a necessary ethical discussion (Miller and Bloom, 1998) and subsequently led to legislation in different countries to regulate when and how the technique or working with human embryonic stem cells (hESCs) would be acceptable (e.g. in Germany see http://www.rki.de/DE/Content/Kommissionen/ZES/zes_node.html). Eight years later Takahashi and Yamanaka published a protocol that allowed for the reprogramming of differentiated murine fibroblasts by expression of just four transcription factors, Oct-4, Sox-2, Krüppel-like factor 4 (Klf-4), and c-myc (Takahashi and Yamanaka, 2006). Thereby they were able to re-establish the intracellular signaling circuitry that induces pluripotent stem cells (iPSCs). Since then several solutions for delivering such reprogramming factors into mouse and human somatic cells have been developed even to the level of commercial products, and these include alternatives such as using small molecules and miRNAs (for review see Hanna et al., 2010; Robinton and Daley, 2012; see also Supplementary Table 1 for selected iPSC-related resources). The iPSC technology not only poses less severe ethical issues, it also offers a real possibility to study the basis of genetic diseases, because iPSC lines can be established from affected patients (Sadelain et al., 2011). Additionally, protocols to re-derive differentiated cells of different types from iPSC are rapidly being developed. Needless to say, these

advances together with initiatives to establish iPSC banks (Rao et al., 2012) open opportunities to study host species- and host genotype-specific parasite interactions. Moreover, the possibility to genetically engineer human iPSCs with a high efficiency (Hockemeyer et al., 2011) in order to introduce at a defined site a transgene or disrupt the endogenous gene(s), thus creating functional mutants, will add even more impetus to this (see next section). In the context of protozoan parasitic infections, unlimited access to cells of the hematopoietic lineages, to neuronal cells, to liver and gastrointestinal cell types, or to organoids mimicking the respective organs is the promise that iPSCs of relevant host species hold (see below and Table 1). Methods for efficient transdifferentiation of readily available cells such as fibroblasts to more difficult-to-get cells complement the choices (Vierbuchen and Wernig, 2011).

Generation of human hematopoietic cells with designed genetic makeup – a novel resource for studying *Plasmodium*-erythrocyte interactions

iPSCs are an attractive platform for the propagation of human *Plasmodium* species due to the strict host cell specificity of the latter for human erythrocytes or hepatocytes (Schuster, 2002). Recent advances in the field of large-scale in vitro production of functional red blood cells (RBCs) from iPSCs (for review see Dravid and Crooks, 2011; Timmins and Nielsen, 2011) provide exciting opportunities for studying the respective host-parasite interactions. Through evolution, the burden of human malaria has resulted in the selection of a number of gene mutations in the human population that affect RBC function. These mutations lower morbidity and mortality due to malaria in the affected human beings (Bauduer, 2012; Gouagna et al., 2010) and have provided important clues to the intraerythrocytic biology of *P. falciparum*. It can be assumed that more host erythrocyte proteins are important in this context,

as suggested by recent studies (Bei et al., 2010; Crosnier et al., 2011). Based on protocols described previously (Giarratana et al., 2005), these investigators used CD34⁺ precursor stem cells purified from human bone marrow and knocked down glycophorin A and basigin, respectively, by RNAi before differentiating these cells to RBC and infecting them with *P. falciparum*. However, the potential of designing specific gene deletions, site-directed specific mutations and modifications or corrections with the aid of so-called zinc finger nucleases (ZFNs; Collin and Lako, 2011) or transcription activator-like effector nucleases (TALENs; Mussolino and Cathomen, 2012) in iPSCs (Hockemeyer et al., 2009; Hockemeyer et al., 2011) allows one to refine and extend the design of such experiments (see Fig. 1). The subsequent differentiation of such mutated cells into erythrocytes would greatly expand the repertoire of mutant erythrocytes amenable to functional testing. Moreover, access to human CD34⁺ precursor stem cells or already known mutant erythrocytes from patients is limited, and iPSC-derived erythrocytes from such individuals would facilitate research on such *P. falciparum*-infected cells.

A further application of this technology is to humanize mice with such cells (Arnold et al., 2011; Legrand et al., 2009) in order to study many aspects of host-parasite relation in vivo in an unprecedented way. The generation of mice with gene defects that prevent formation of T and B cells (*scid* or *rag* mutations) and natural killer cells (through IL-2 receptor γ chain deficiency) resulted in models that sustain meaningful numbers of human blood cells (Ito et al., 2002; Traggiai et al., 2004), and these models are continuously being improved (Brehm et al., 2012; Strowig and Flavell, 2012). The individual techniques have all been described (see Fig. 1 and Table 1), and it is only a matter of time until the first studies implementing it for *P. falciparum* are published.

Human neuronal cell lines and *Toxoplasma gondii*

One reason that *T. gondii* has emerged as a prominent model organisms for protozoan parasites is due to its ease of cultivation and its very broad host species and host cell range, including mutant cell lines that have been important to the study of many aspects of host-parasite interaction (Pfefferkorn, 1990). It could be argued that stem cell-derived systems offer little extra when studying *T. gondii* host-parasite interplay. However, the latent, chronic stage of *T. gondii*, the so-called bradyzoites, are found predominantly in brain, heart, and muscle tissue. These tissues are readily available, e.g. from experimentally infected mice, but equivalent human cells are difficult to obtain. iPSC-derived neuronal, cardiac, or muscle cells could, however, deliver sufficient quantities of well-defined samples that would enable comparative analyses relevant to human disease. For example, the recently reported direct reprogramming (transdifferentiation) of human fibroblasts into neurons (Ambasudhan et al., 2011; Pang et al., 2011; Pfisterer et al., 2011) opens exciting possibilities to study this particular host cell-parasite interaction in more detail. It would allow the evaluation of results obtained with infected rodent neurons that suggested an influence of *T. gondii* on neuronal dopamine metabolism (Prandovszky et al., 2011).

Human stem cell-derived liver cells for *Plasmodium* research on sporozoite invasion and development of exoerythrocytic forms (EEF)

The clinically silent malaria liver stage has become a major research area, because it represents one of the bottlenecks in the *Plasmodium* life cycle and is probably the most promising stage for vaccine development (Mueller et al., 2005; Nussenzweig et al., 1967). To date, the liver stage can only be studied in depth in animal models of malaria (there is no option for the human-specific *Plasmodium* species) and

hepatocyte cell lines. Adult, untransformed human liver cells are not routinely used due to the fact that they do not proliferate in vitro and are difficult to cryopreserve (Terry and Hughes, 2009). Still very little is known as to why mammalian *Plasmodium* sporozoites specifically target hepatocytes as their primary host cells for subsequent EEF development (Hafalla et al., 2011; Silvie et al., 2008). Recent advances in differentiating human ESCs or iPSCs into hepatocyte-like cells (Chen et al., 2012; Schwartz et al., 2005; Song et al., 2009; Touboul et al., 2010) or to transdifferentiate human fibroblasts into cells with hepatocyte-like properties (Sekiya and Suzuki, 2011) may for the first time offer opportunities to study aspects of this host cell specificity of human-specific liver stages of *P. falciparum* or *P. vivax* in more depth. Additionally, we lack knowledge about the full repertoire of antigens that are presented during the arrested EEF development. Genetically attenuated parasites showing this arrest can confer protection in murine models (Mueller et al., 2005). Stem cell-derived human hepatocytes could help identify or test presented (protective) antigens from respective mutants of human *Plasmodium* species in order to develop subunit vaccines against sporozoites and liver stages, as this is of high priority in combating malaria (Kappe et al., 2010).

Non-parenchymatic liver cells (NPLCs), i.e. stellate cells, liver sinusoidal endothelial cells (LSECs) and Kupffer cells, are highly specific cells with unique properties that contribute significantly to the liver tissue and are vitally important for liver function during homeostasis and infection (Protzer et al. 2012). While little is known about the role of NPLCs during the hepatic phase of human malaria, mouse data revealed that during the transmigration event, sporozoites are in intimate contact with LSECs and pass through Kupffer cells in order to enter the liver (Frevort et al., 2008; Klotz and Frevort, 2008). Obviously, the development of a method to derive NPLCs from stem cells would enable stud-

ies on the role of these cells (e.g. for antigen presentation during the hepatic stage of *Plasmodium*) to be intensified, but such a protocol is currently not available. Studies of *E. histolytica*-LSEC interactions (Faust et al., 2011) in order to better understand the process behind amoebiasis-induced liver abscesses could also benefit from stem cell-derived NPLCs.

Intestinal organoids as a new tool for enteric protozoa and for studying habitat-parasite interactions

Intestinal protozoan parasites are phylogenetically quite divergent but share the gut as a common habitat or entry point into the host. These organisms possess diverse biology for survival and replication in different niches; for example *G. duodenalis* or *Entamoeba* spp. dwell in the intestinal lumen, and *Cryptosporidium* spp. or *T. gondii* live within intestinal cells (Stark et al., 2009). Research on many of these organisms is hampered by a lack of appropriate in vitro culture systems. Stem cell-derived systems can offer new avenues to investigate the organ-specific interactions of these parasites.

In seminal papers published in 2009, two independent research groups reported different methods for the establishment of stem cell-based organoids from primary mouse intestinal tissue (Ootani et al., 2009; Sato et al., 2009). Ootani and colleagues established a more complex culture system from minced whole intestinal tissue embedded in a three-dimensional collagen structure with support of stromal cells (Ootani et al., 2009). Sato et al. succeeded in establishing organoid cultures from isolated crypts and from single Lgr5 marker-positive cells using a matrigel matrix and serum-free medium supplemented with Wnt antagonist R-Spondin1, TGF-beta signaling inhibitor Noggin and epidermal growth factor (Sato et al., 2009; Sato et al., 2011b). The latter system is particularly attractive, as no stromal cell support is

necessary and it can be implemented relatively easily in most laboratories. We established such organoids following the protocol of Sato et al. from primary mouse crypts isolated from various mouse strains (Fig. 2). The observed crypt-villus structures consist of all epithelial, stem cell-derived progenitor cells, i.e. enterocytes, paneth cells, goblet cells and enteroendocrine cells, but lack cells of mesenchymal or hematopoietic lineages (Sato et al., 2009). Organoids simulate the in vivo situation much better than for instance the intestinal cell lines currently used for studying host-parasite interactions in vitro. Additionally, they are amenable to live cell imaging techniques akin to intravital microscopy (Chieppa et al., 2006). Moreover, since organoids can be generated from different tissues of the gut, including the small intestine and colon (Sato et al., 2009; Sato et al., 2011a, b; Yui et al., 2012), site-specific interaction with a particular pathogen can be studied.

Recently, human intestinal organoids have been successfully established on the one hand from hESCs and hiPSCs (McCracken et al., 2011; Spence et al., 2011), and on the other hand starting from primary human tissue out of healthy and diseased patients (Sato et al., 2011a). Therefore in the future, we might be able to establish new in vitro culture systems for the sparsely studied human intestinal protozoan parasites that are still inadequately cultured or not culturable (see Table 1).

Giardia duodenalis

While in vitro studies on *G. duodenalis* are relatively easy to perform, it is much more difficult to investigate the events in the host gut that lead to disease and pathology, particularly in humans. Currently, researchers resort to animal models or studies of parasites with immortalized intestinal cell lines, both of which have their obvious and distinct limitations. The impact of the various proposed disease mechanisms on parasite pathogenicity (reviewed in Ankarklev et al., 2010; Cotton et al.,

2011), such as induction of apoptosis and loss of intestinal barrier function, and the role of putative parasite virulence factors in these processes is currently unknown. Intestinal organoid cultures derived from iPSCs or from human intestinal tissue would greatly advance our means to study *G. duodenalis* pathogenicity and could help define virulence factors relevant for human infection. Currently, intestinal organoids lack cells of the immune system but offer the possibility to study the direct interaction of the parasites with well-differentiated intestinal epithelial tissue as well as early events during infection. Furthermore, it seems possible to elucidate host factors affecting *G. duodenalis* infection outcome by using organoids derived either from patients with particular disease manifestations (e.g. from volunteers with symptomatic vs. asymptomatic infections) or from genetically modified iPSCs.

Another attractive application for intestinal organoids is to use them to improve the success rate of parasite adaptation to in vitro culture from clinical material. Although cell-free culture systems with trophozoites derived from biopsies or generated from cysts via in vitro excystation have been pioneered (Bingham and Meyer, 1979; Meyer, 1970), it is still not possible to derive cultures from all distinct *G. duodenalis* genotypes. This resulted in a large research bias towards easily culturable isolates (Smith et al., 1982). An organoid infection system enabling the cultivation of so far non-culturable isolates would represent a breakthrough and, for example, establish the basis for functional epidemiology of giardiasis.

Toxoplasma gondii

T. gondii's usual port of entry into its hosts is through the intestinal route via ingestion of tissue cysts or oocysts. While this initial infection process is documented to some degree in mice (Barragan and Hitziger, 2008; Barragan and Sibley, 2002) intestinal organoids could provide a relevant model for studying this aspect in the hu-

man system.

An exciting prospect of the use of intestinal organoids is the possibility to approach the in vitro reconstruction of the sexual reproductive cycle of *T. gondii*. The latter plays a central role in the creation of genotypes of different virulence, but at a scale that has previously not been anticipated (Rao et al., 2012; Wendte et al., 2011). Sexual differentiation only takes place in cats and other felids. Thus, the challenge is to establish feline intestinal organoids and find in vitro conditions that would allow the sexual cycle to commence (and finish). It would provide an opportunity to analyze this poorly understood process under defined conditions and follow the process in real time through live cell imaging. Furthermore, such a culture system may offer an initial platform for vaccine or drug testing aimed at blocking oocyst formation and thereby dissemination.

iPSCs as screening and evaluation platforms for new drugs

Although most of the excitement over using iPSCs for drug screening purposes is currently in the field of monogenic human diseases (Ebert and Svendsen, 2010; Grskovic et al., 2011), screening programs intended to identify novel anti-protozoan parasite drugs would also benefit from the availability of differentiated cells usually not easily accessible, for example neurons. Stem cell-derived, differentiated cells are thought to allow more sensitive pre-clinical toxicology testing (i.e. for cardiotoxicity, hepatotoxicity, genotoxicity, and reproductive toxicology) of drug candidates than conventional cell systems do (Davila et al., 2004). Also, exploring the basis of differential drug pharmacology, kinetics, and metabolism in different individuals (frequently reflected as apparent "resistance" against drug X) using iPSCs derived from such patients for a given parasitic disease could greatly help in tailoring treatment regimens with approved, cheap, and safe drugs, thereby possibly extending the half-life of some anti-

infectives in the clinics as in the case of malaria (Kerb et al., 2009; Paganotti et al., 2011).

Conclusions

Research exploiting iPSCs and their derived cells is still a very young field. Obviously, iPSC-derived cells and organ-like models have a number of caveats and disadvantages. Establishing protocols for one's own special purposes is not trivial. The requirement for expensive growth and differentiation factors currently make this approach very costly. Organ-like models still do not reproduce the situation *in vivo*; e.g. organoids often lack particular cell types such as immune cells or other components of the mimicked organ (e.g. the flora in gastrointestinal organoids), and this might be problematic. However, many step-by-step protocols and commercially

available reagents (including established human iPSCs) already exist, and methods improve at a remarkable speed. Prices for reagents depend on demand and can drop quickly. For particular research questions, lack of complexity (e.g. additional cell types) may also be advantageous, offering a basic cellular model where complexity levels can be added as desired. Thus, we believe that the promises iPSC-based approaches offer to the study of protozoan parasites and host-specific interactions are realistic, and the opportunities should be seized.

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Table 1: Potential applications of stem cell-derived primary tissues/cells for research on the host-pathogen interaction of protozoan parasites

Differentiated cells/tissue *	(Stem) Cell source and technical references *	Protozoan pathogen	Examples of possible application
Hemopoietic cells			
Erythrocytes	hiPSCs (Dravid and Crooks, 2011; Timmins and Nielsen, 2011)	<i>P. falciparum</i> and <i>P. vivax</i> (blood stage) <i>Babesia</i> spp.	Mutant erythrocytes for studying metabolite transport
Neuronal tissue			
Neurons	Fibroblasts (transdifferentiation) (Ambasudhan et al., 2011; Pang et al., 2011; Pfisterer et al., 2011) hESCs, hiPSCs (Lie et al., 2012)	<i>T. gondii</i>	Dissecting neuronal host-parasite interaction of the latent parasite stage (bradyzoites)
Liver			
Hepatocytes	hiPSCs, hESCs (Cai et al., 2007; Chen et al., 2012; Duan et al., 2007; Schwartz et al., 2005; Song et al., 2009; Touboul et al., 2010) Fibroblasts (transdifferentiation) (Sekiya and Suzuki, 2011)	<i>P. falciparum</i> and <i>P. vivax</i> <i>Entamoeba</i> spp.	Studying nutrient acquisition, replication, and antigen presentation in primary human (mutant) hepatocytes Revealing underlying pathogenicity factors of human liver/hepatocyte destruction
Hollow fiber / organoids	miPSCs and mESCs (Amimoto et al., 2011; Mizumoto et al., 2008)		
Intestine			
Intestinal organoids	hiPSCs, human intestinal crypts (McCracken et al., 2011; Sato et al., 2011a; Spence et al., 2011) Mouse Lrg5 ⁺ cells, mouse intestinal crypts (Ootani et al., 2009; Sato et al., 2011a, b, 2009)	<i>G. duodenalis</i> <i>Entamoeba</i> spp. <i>Balantidium</i> spp. <i>Cryptosporidium</i> spp. <i>Cyclospora</i> spp.	Improved in vitro culture of specific genotypes, studying virulence/ pathogenicity factors Studying pathogenicity of intestinal amoebiasis and balantidiosis Improved in vitro culture, drug and disinfectant screening, studying

biology of life cycle
stages, viability assays

T. gondii

Recapitulating host-
parasite interactions at the
primary infection site

* The different stem cell sources and differentiated cells can be combined with different protozoa, depending on the intended experiment.

Legends to illustrations

Figure 1

Driven by specific mixes of growth and differentiation factors, a variety of defined, differentiated cell types and even organoids can be derived from iPSCs (for generic iPSC protocols see Rajarajan et al., 2012). Direct re-programming, i.e. transdifferentiation of fibroblasts etc., to neurons by respective growth and differentiation factors is also possible. Engineered proteins containing zinc fingers or transcription activator-like effector domains fused to nucleases, called ZFN and TALEN, respectively, are used to create double-strand DNA breaks at predetermined sites in the genome (for technical resources see Sanjana et al., 2012). The breaks in turn are repaired by end joining and, in the presence of DNA constructs homologous to the chosen region, by homologous recombination-mediated repair. Thereby site-specific mutations/insertions can be introduced at the level of starting cells (e.g. fibroblasts) or iPSCs. Infection of iPSCs, of iPSC-derived cell types or organoids, or even of mice humanized by the engraftment of hematopoietic precursor cells etc., with the parasite of choice offer different avenues for pathogen propagation and host interaction studies.

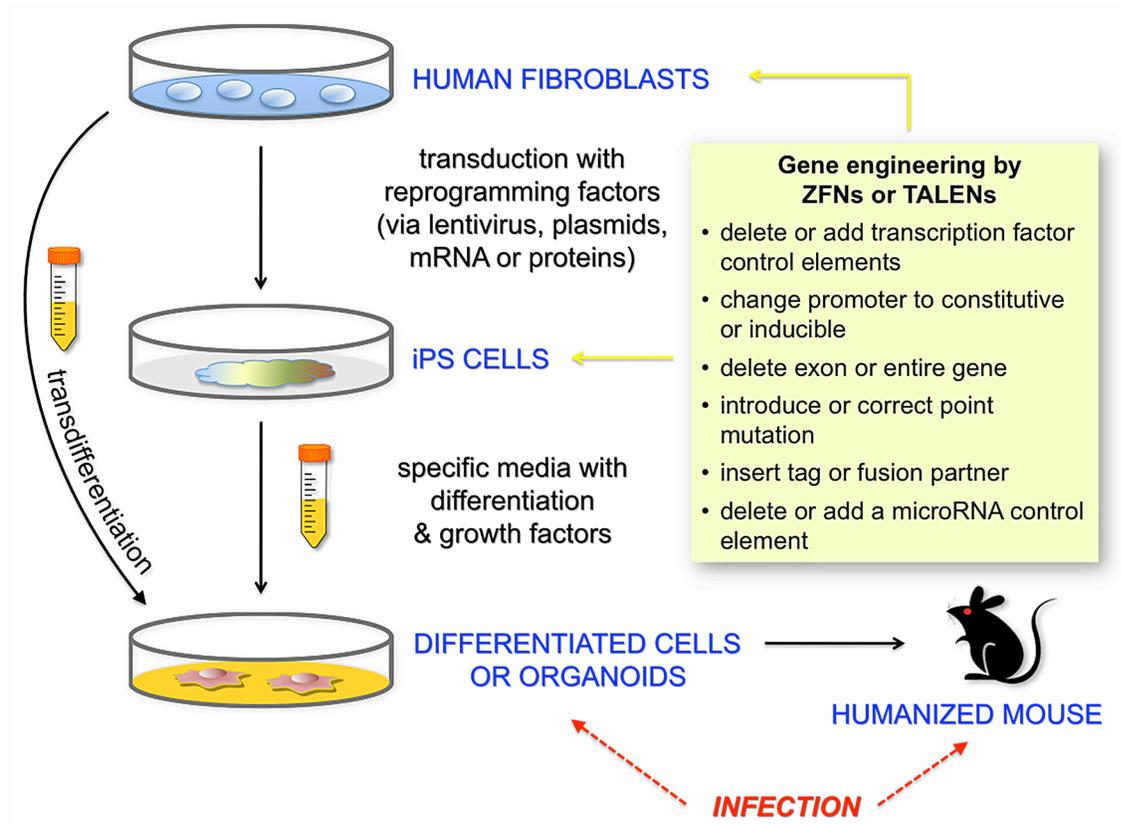
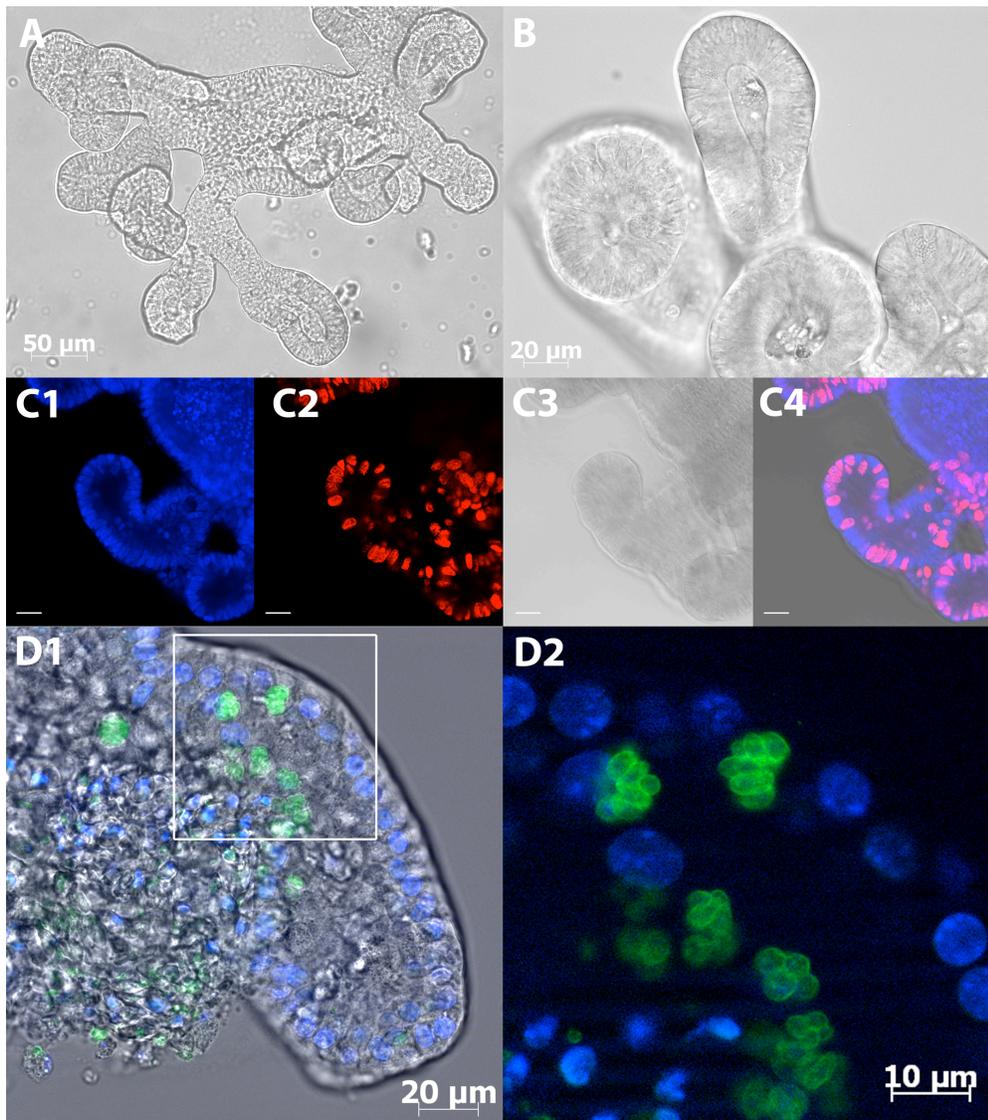


Figure 2

Intestinal organoid cultures to study host-parasite interaction. A, B: Long-term cultures of crypt-derived intestinal organoids. Note the enclosed crypt-villous-like structures with luminal shedding of jaded enterocytes. C1-C4: Confocal imaging of proliferating cells of organoid cultures visualized by 5-ethynyl-2'-deoxyuridine (Edu)-incorporation. C1, Dapi (blue.); C2, Edu (red); C3, Nomarski interference; C4, merge of C2 and C3. Scale bar = 20 μm . D1: Immunofluorescence image of an intestinal organoid infected with tachyzoites of the RH strain of *T. gondii* 48 h post infection. Dapi (blue); tachyzoite surface was stained with anti-SAG1 mab DG52 (green). The fluorescence image was merged with a Nomarski interference image. D2: Enlargement of the fluorescent image from D1 (white square). Note the rosette-like structures of intracellular parasites in the epithelial layer, typical of RH strain tachyzoite replication.



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Supplementary Table 1: Selection of internet addresses of iPSC and related resources

Mouse stem cell resources	URL
RIKEN	http://www2.brc.riken.jp/lab/cell/s5_list.cgi?key=e
Mutant Mouse Regional Resource Centers (MMRRC) Baygenomics	http://www.mmrrc.org/catalog/overview_BG.php
EMMA – The European Mouse Mutant Archive	http://www.emmanet.org/
Mount Sinai School of Medicine Soriano Lab	http://research.mssm.edu/soriano/lab/gene_trap.html
Sanger Gene Trap Consortium	http://www.sanger.ac.uk/resources/mouse/sigtr/
International Genetrap Consortium	http://www.genetrap.org/
Center for iPS Cell Research and Application	http://www.cira.kyoto-u.ac.jp/e/research/material_1.html
Depositories of Human stem cells & primary cells	
Boston Univ. Med. School iPS Bank Project	http://www.bumc.bu.edu/evanscenteribr/the-arcs/the-arcs/regenerative-medicine/
WiCELL Research Institute iPS Distribution	http://www.wicell.org/index.php?option=com_content&task=category&id=348&Itemid=262&sectionid=19
International Stem Cell Repository	http://www.umassmed.edu/iscr/index.aspx
The Human Genetic Cell Repository (NIGMS)	http://ccr.coriell.org/Sections/Collections/NIGMS/
LGC Standards/ATCC	http://www.lgcstandards-atcc.org/LGCAdvancedCatalogueSearch/ProductDescription/tabid/1068/Default.aspx#aPrope3e6d
HSCI iPS Cell Core Facility	http://www.hsci.harvard.edu/ipscore/node/3
European Human Embryonic Stem Cell Registry	http://www.hescreg.eu/
HPA Culture Collections	http://www.hpacultures.org.uk/products/celllines/primarycells/index.jsp
Health Science Research Resources Bank (Japan)	http://www.jhsf.or.jp/bank/Category-Index.html
System Biosciences (commercial)	http://www.systembio.com/stem-cell-research/ips-cell-lines
Sigma-Aldrich iPS Cell Design Studio (commercial)	http://www.sigmaaldrich.com/life-science/stem-cell-biology/ips-cell-design-studio.html
Plasmids for iPSC generation and modification at Addgene	
Plasmids for Stem Cell Research	http://www.addgene.org/stemcell/
Zinc Finger Consortium Reagents	http://www.addgene.org/zfc/
TALEN Kits	http://www.addgene.org/TALEN/