Escape Mutations, Ganciclovir Resistance, and Teratoma Formation in Human iPSCs Expressing an HSVtk Suicide Gene

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Human pluripotent stem cells (hPSCs) hold great promise for cell therapy. However, a major concern is the risk of tumor formation by residual undifferentiated cells contaminating the hPSC-derived cell product. Suicide genes could safeguard against such adverse events by enabling elimination of cells gone astray, but the efficacy of this approach has not yet been thoroughly tested. Here, we engineered a lentivirally encoded herpes simplex virus thymidine kinase (HSVtk) with expression restricted to undifferentiated hPSCs through regulation by the let7 family of miRNAs. We show that induced pluripotent stem cells (iPSCs) expressing a let7-regulated HSVtk transgene are selectively killed by ganciclovir (GCV), whereas differentiated cells are fully protected. However, in contrast to previous studies, we find that in vivo GCV administration results in longer latency but does not prevent teratoma formation by iPSCs expressing either a constitutive or a let7-regulated HSVtk, without evidence of silencing of the HSVtk. Clonal analyses of iPSCs expressing HSVtk revealed frequent emergence of GCV resistance which, at least in some cases, could be attributed to preexisting inactivating mutations in the HSVtk coding sequence, selected for upon GCV treatment. Our findings have important consequences for the future use of suicide genes in hPSC-based cell therapies.

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Introduction

Induced pluripotent stem cells (iPSCs) have opened exciting new prospects for regenerative medicine. However, the same properties of unlimited self-renewal and pluripotency which are unique to human pluripotent stem cells (hPSCs) and central to their therapeutic promise are also intrinsically linked to their potential for extensive growth and tumorigenicity.1 Several preclinical studies in rodent and nonhuman primate models have established the risk of tumorigenicity associated with administration of hPSC-derived cell grafts.2–7 Tumor formation may follow dedifferentiation and transformation of terminally differentiated cells, aberrant proliferation of incompletely differentiated progenitors, or teratoma development by residual pluripotent cells that may persist after differentiation and contaminate the graft.8–10 Although the latter are benign tumors, their growth can have catastrophic consequences in particular anatomic sites such as joints, brain, spinal cord, or the eye.11–13 The immune-privileged status of undifferentiated, as opposed to differentiated cells—at least in part due to low major histocompatibility complex antigen expression by the former—may further increase this risk, due to evasion of immune surveillance even in a nonautologous cell therapy setting.14

The use of suicide genes has been proposed as a promising strategy to safeguard against teratoma formation of hPSC-derived cell products. Herpes simplex virus thymidine kinase (HSVtk) is the most common suicide gene with a long track record in the clinic.15 It encodes an enzyme that converts a prodrug, ganciclovir (GCV), into a cytotoxic metabolite. Several recent studies have suggested that HSVtk can prevent and/or ablate teratomas derived from human or murine PSCs.16–19 Ideally a suicide approach should be selective to pluripotent undifferentiated cells, while sparing their differentiated progeny. Attempts to engineer such an expression pattern in suicide genes have included the use of pluripotent cell-specific promoter/enhancers, such as the Oct4 (knockin at the endogenous Oct4 locus),20 NANO2 (knockin at the endogenous NANOlocus19 or on a lentiviral vector18), or the synthetic promoter/enhancer element Early Transposon promoter and Oct4 and Sox2 enhancers, EOE-C(3+)

We and others have previously shown that regulation by endogenous miRNAs is a very effective and robust strategy to endow transgenes with tissue- and developmental stage-specific expression.23–25 Tagging of a transgene with sequences complementary to a mature miRNA subjects the former to posttranscriptional regulation by the specific miRNA, so that its expression is suppressed only in the presence of the miRNA but not in its absence. Because expression of most miRNAs is transcriptionally regulated in a cell type- and differentiation stage-specific manner, this mechanism of transgene regulation has found several applications in genetic engineering.24

The let7 family of miRNAs are ubiquitously expressed in differentiated tissues, but absent from pluripotent cells. Here,
in order to sequester expression of HSVtk strictly in undifferentiated hPSCs and render the latter selectively sensitive to GCV-induced cell death, we engineered HSVtk to be regulated by the let7 family of miRNAs. While we observed no teratoma formation for up to 5 weeks in mice that received HSVtk-expressing iPSCs with administration of GCV, no protection from teratomas was observed in the longer term, likely due to inactivating mutations of the HSVtk transgene. Our study has fundamental implications for the use of HSVtk as a suicide gene in hPSC-based cell and gene therapy. It also highlights the need for longer observation periods in preclinical testing for teratoma formation.

Results
Let7 regulation suppresses expression of HSVtk in differentiated but not pluripotent cells
We set to engineer a lentivirally encoded HSVtk to be specifically expressed in undifferentiated hPSCs, rendering them sensitive to cell death by GCV, but not in their differentiated progeny, which would thus be resistant to GCV using miRNA regulation (Figure 1a). We selected the let7 family of miRNAs on the basis of its well-documented widespread and high-level expression in practically all differentiated cells, but not in undifferentiated pluripotent stem cells. We constructed bidirectional lentiviral vectors expressing HSVtk linked by a P2A peptide to GFP driven by the EF1α promoter or the pluripotent-specific EOS-C(3+) promoter and constitutively expressed hPGK-driven mCherry (vectors pRG-tk and pRG-EOS-tk, respectively) (Figure 1b). EF1α-driven HSVtk was tagged to four tandem miRNA recognition elements (MRE) complementary to mature miRNAs let7a, let7d, let7f, and let7g (vector pRG-tk-let7) or to four MREs complementary to mature miR-142. miR-142 is a miRNA predominantly expressed in hematopoietic cells, but not in pluripotent stem cells or fibroblasts. The four vectors were transduced into an iPSC line (thai-ips thal-5.10-Cre8) derived from a beta-thalassemia patient, as well as in bone marrow fibroblasts derived from the same individual (thali-fibs). As expected, both iPSCs and fibroblasts coexpressed mCherry with GFP-HSVtk expressed by the nonregulated (pRG-tk) and miR-142-regulated (pRG-tk-miR142) vectors (Figure 1c). In contrast, let7 regulation resulted in suppression of GFP-HSVtk expression in fibroblasts but not in undifferentiated iPSCs (Figure 1c and Supplementary Figure S1). GFP-HSVtk expression levels relative to mCherry were practically identical in iPSCs with or without let7 regulation, showing that let7 regulation has no detectable effect in HSVtk expression in pluripotent cells (Supplementary Figure S2a). EOS-driven HSVtk expression was also undetectable in fibroblasts but maintained in iPSCs at low levels (Figure 1c). This vector was thus not further used.

Let7-regulated HSVtk enables selective killing of pluripotent but not differentiated cells
To next test if let7-regulated HSVtk could operate as a “death switch,” iPSC colonies homogeneously expressing GFP-HSVtk and GFP-HSVtk-let7 were individually picked under a microscope and expanded (Supplementary Figure S2b). iPSCs expressing GFP-HSVtk and GFP-HSVtk-let7 showed unchanged morphology (Figure 2a) and expression of pluripotency markers (Figure 2b) and could form trilineage teratomas upon injection into immunodeficient mice (Figure 2c).

Once clonal iPSC cultures homogeneously expressing GFP-HSVtk were established, GCV sensitivity was assessed. iPSCs expressing either HSVtk or HSVtk-let7 were completely eliminated over 14 days of GCV treatment (Figure 3a). No live cells could be detected after treatment of 1–2 million iPSCs with GCV at 0.5 mg/ml or 1 mg/ml. In contrast, fibroblasts expressing let7-regulated, but not constitutive, HSVtk were completely protected over a range of GCV concentrations up to 1 mg/ml (Figure 3b). No cell death was observed by assaying cell counts even at the highest GCV dose (Figure 3c,d). In contrast, fibroblasts expressing constitutive or miR-142-regulated HSVtk were sensitive even at very low GCV concentrations (Figure 3c). Assessed further by a more sensitive flow cytometry-based assay, fibroblasts expressing HSVtk-let7 (Figure 3d, mCherry+) remained resistant to GCV, similarly to untransduced cells (mCherry-), as the % of mCherry- cells remained stable over a range of GCV doses and similar to that of untreated cells (Figure 3d). These results collectively demonstrate that let7-regulated HSVtk is robustly expressed in undifferentiated iPSCs, rendering them sensitive to killing by GCV, while it is posttranscriptionally suppressed in differentiated cells, preserving their resistance to GCV.

HSVtk delays but does not abolish the emergence of teratomas
To next test if HSVtk with or without let7 regulation can prevent teratoma formation, we injected the iPSC clones transduced with pRG-tk, pRG-tk-let7, or a control GFP vector subcutaneously into NSG mice. One week later, administration of GCV or phosphate-buffered saline through a minipump was initiated and maintained for 2 weeks. Tumor size was monitored over 8 weeks (Figure 4a). By week 5 after cell injection, over half of the mice that received iPSCs not expressing HSVtk or were not treated with GCV developed palpable teratomas (Figure 4b). In contrast, none of the mice that received iPSCs harboring HSVtk or HSVtk-let7 and GCV developed teratomas. However, by week 6 after injection, tumors appeared in several GCV-treated mice from both groups injected with iPSCs harboring pRG-tk or pRG-tk-let7. These had initially overall smaller size than the tumors in untreated mice but continued growing over the next 2 weeks until they reached a size comparable to the tumors in the untreated mice and in the mice that received iPSCs expressing GFP only (Figure 4b). By week 8, teratomas had developed in 8 out of 10 injection sites of iPSCs expressing HSVtk and in 9 out of 10 injection sites of iPSCs expressing HSVtk-let7 with GCV treatment. All (10 out of 10) sites of injection of iPSCs not expressing HSVtk or not treated with GCV showed tumor growth.

These results show that HSVtk (with or without let7 regulation) does not ultimately prevent teratoma formation. Teratomas derived from HSVtk-expressing iPSCs developed with a longer latency than teratomas derived from iPSCs not expressing HSVtk in the presence of GCV. These kinetics are consistent with initial sensitivity to GCV and emergence of resistance over the course of treatment.
GCV resistance is mediated by HSVtk mutation

Resistance to GCV-mediated killing in cells expressing HSVtk has previously been reported to occur through various genetic and epigenetic mechanisms.30–32 To first test if GCV resistance was due to HSVtk silencing, teratomas grown in mice injected with iPSCs transduced with pRG-tk with or without GCV treatment were collected and their cellular components were analyzed by flow cytometry for GFP-HSVtk expression. All four teratomas tested showed robust expression of GFP-HSVtk in the majority of the cells (Figure 5). This result precludes transgene silencing as the cause of GCV resistance.

To further investigate the mechanism of GCV resistance, we derived single-cell iPSC clones expressing either pRG-tk or pRG-tk-let7. Eleven clones were expanded for ~3 weeks and their sensitivity to GCV was tested. The clones showed variable sensitivity to GCV with few but detectable resistant colonies remaining in 6 of the 11 clones after exposure to GCV (1 mg/ml) for 14 days (Figure 6a). Sequencing of the HSVtk in two of the GCV-resistant clones revealed nonsense mutations in the HSVtk coding region (Figure 6b). These results show that HSVtk mutation is a frequent mechanism of resistance of human iPSCs to GCV.
Discussion

Given the very high stakes for the stem cell field, translation of stem cell therapies needs to proceed with extreme caution to avoid the occurrence of serious adverse events in early-stage trials. Suicide genes, among other strategies, will be valuable tools to increase safety in stem cell therapies. There is thus an urgent need to evaluate the efficacy of such an approach.

Here, we showed that miRNA regulation by let7 can be used to engineer a robust genetic “switch” in a suicide gene, so that the latter is “on” in undifferentiated hPSCs but gets turned “off” once the cells are differentiated to the desired cell type. To the best of our knowledge, this constitutes the most effective pluripotent cell-specific regulation of a vector-encoded transgene reported to date and can find many applications given the general scarcity of tools for engineering pluripotent-specific gene expression. Alternative approaches involve either the much more cumbersome targeting of a suicide gene in an endogenous pluripotent cell-specific locus, such as OCT4 or NANOG. This approach requires multiple engineering steps and it may interfere with expression of the endogenous gene and the pluripotent state and the properties of the target cell. On the other hand, incorporation of promoter/enhancer elements from genes such as NANOG and OCT4 in vectors has not so far resulted in robust transgene expression and/or regulation. Synthetic composite promoters/enhancers, like EOS-C(3+), may be more promising. In our study, expression of HSVtk by the EOS-C(3+) promoter was relatively low (Figure 1c) but might have been sufficient for effective suicide gene therapy. Since our miRNA-based strategy is robust and very simple to implement, as it only requires the incorporation of short sequences in the vector, it can have broader application in transgene regulation for basic stem cell research, in addition to suicide gene therapy. While expressing exogenous MREs can potentially exert some inhibitory effects on endogenous miRNAs, this would require very high levels of MRE expression that are unlikely to be reached with our strategy.

Previous studies proposed the prevention of teratomas by an HSVtk suicide gene.
Differences in various experimental parameters, such as the onset and the duration of GCV treatment following cell injection, the vector, the dose of GCV, and the HSVtk variant, can be noted between studies. A major difference that can account for the difference in the observed outcome is that, in contrast to the previous studies, we used a larger cohort of mice and a longer observation period. Indeed, if we had evaluated our mice at week 5 with no further follow-up, our conclusion would also have been that HSVtk completely abolishes teratoma formation (Figure 4b). This illustrates that short-term studies can miss heterochronic teratoma formation between test and control groups as a result of resistance acquired later in the course of an experiment and highlights the need for long observation periods in future teratoma formation studies.

Our findings document acquisition of resistance to GCV by iPSCs expressing HSVtk. The injected cells were clonal and confirmed to be sensitive to GCV before injection. Furthermore, the kinetics of teratoma growth were consistent with initial sensitivity and subsequent emergence of resistance to GCV. GCV resistance has been observed in various murine and human cell lines, including human iPSCs30–32 and relapse after GCV...
suicide treatment has been reported even in clonal tumors.\textsuperscript{30} Epigenetic silencing as well as a variety of genetic events including point mutations, small frameshifting insertions/deletions, deletions of the entire gene, loss of part or of an entire chromosome, and postinsertional recombination of retroviral encoded HSVtk have all been documented as mechanisms mediating GCV resistance.\textsuperscript{30–32} Of all the different mechanisms, we were able to document inactivation of the HSVtk gene by point mutations. Since our cells were transduced at high multiplicity of infection and likely harbored more than one HSVtk copies and we did not test for large-scale rearrangements or chromosomal deletions, we cannot exclude the possibility that additional mechanisms of resistance—genetic and epigenetic—might also have contributed to escape and teratoma formation in our study.

Our clonal analyses suggest that escape mutants arise frequently (Figure 6a). In a translational setting, many factors can influence the frequency of mutational HSVtk inactivation, such as the vector system used and the degree of expansion of the iPSCs before and after differentiation and before and after administration to the patient. Both mutations in our clones were G-to-A substitutions (Figure 6b). Wild-type HIV-1 shows a high rate of G-to-A substitutions, due to the error-prone retroviral reverse transcriptase and/or RNA editing by mammalian cytidine deaminases of the APOBEC family.\textsuperscript{33} Therefore, the genetic stability and mutation rate of the viral vector may need to be considered in suicide gene transfer into hPSCs. Since malignant transformation often results in genetic and epigenetic instability, the emergence of GCV-resistant clones in the event of malignancy arising...
from iPSC derivatives may be even more likely. The specific type of tumor and the driver genetic alterations that underlie transformation and tumor development can also influence the genomic stability and frequency of GCV resistance.

Acquired GCV resistance and teratoma growth through escape mutational events is akin to drug resistance in malignant tumors. The cancer field has extensively documented and studied the emergence of tumor resistance to cytotoxic agents through genetic events that are subject to selection and clonal evolution in a Darwinian process.

Due to their vast proliferation capacity, processes of clonal evolution are also at play in hPSCs and have fundamental implications for stem cell therapies. A mutation that confers resistance of cancer cells to chemotherapy is usually a passenger mutation before treatment, arising by chance as a result of background mutation rate. Similarly, in our study, HSVtk mutations were found to preexist in rare cells of the original population and were selected for upon treatment with GCV. Thus, in a translational setting, HSVtk mutations can be acquired by rare hPSCs at any stage of cell expansion prior to as well as following administration of the final cell product to the patient. The cells that harbor resistance mutations can lurk in the cell graft and remain inconsequential until treatment with the prodrug is initiated. Once treatment is applied, these cells immediately have a selective advantage and can readily give rise to tumor growth. In view of this, more emphasis should be placed on methods for detection by deep sequencing and purging of cells harboring potential resistance mutations to suicide gene therapy at an early stage, as well as after subsequent expansion and prior to infusion.

Suicide genes other than HSVtk have been developed and tested in human and nonhuman primate PSCs. These include the yeast cytosine deaminase, an enzyme that converts the nontoxic prodrug 5-fluorocytosine into a potent antimetabolite, 5-fluorouracil, and the inducible Caspase-9, a proapoptotic molecule that can be activated by a dimerizer drug (specific chemical inducer of dimerization). Whereas experience with these suicide genes is much more limited, resistance has been observed in studies utilizing inducible Caspase-9 in rhesus macaque hematopoietic stem and progenitor cells and human iPSCs, with teratoma regrowth after chemical inducer of dimerization treatment also observed in the latter case. Although the mechanisms of resistance to chemical inducer of dimerization treatment have not yet been thoroughly investigated, they are likely to involve the same array of genetic and epigenetic mechanisms that inactivate HSVtk and hPSCs harboring inducible Caspase-9 or other suicide genes will most likely also be subject to the same clonal dynamics. In view of this, safe stem cell therapies may require combinations of suicide genes or, more likely, the combinatorial use of more than one redundant
approaches for purging and/or suicide gene therapy of a cell graft. Alternative strategies that can be combined with suicide genes are methods for purging of hPSCs in vitro prior to transplantation. These include antibodies against hPSC-specific surface antigens—either cytotoxic or to be used for fluorescent-activated cell sorting- or magnetic-activated cell sorting-based depletion. Another possibility is the development of small molecule inhibitors driven by the identification of hPSC-specific vulnerabilities arising from dependence on particular signaling or metabolic pathways.

In conclusion, suicide genes can enhance the safety of stem cell therapies. For this promise to materialize, a better understanding of both the potential and the limitations of suicide gene therapy and the possibilities of combining it with other approaches will need to be the focus of future preclinical studies.

**Materials and methods**

**Cell culture.** The thal-iPSC line thal-5-10-Cre8 and the bone marrow fibroblasts (thal-fibs) that was derived from have previously been described. Human iPSCs were cultured on a feeder layer of mitomycin C-treated mouse embryonic fibroblasts (GlobalStem, Gaithersburg, MD) or in feeder-free conditions and passaged with dispase or accutase as previously described.

**Lentiviral vector construction and production.** pRG-tk was derived from vector NG by replacing NTP with mCherry and GFP with a GFP-P2A-HSVtk cassette in which the P2A peptide is preceded by a Gly-Ser-Gly linker generated by overlapping PCR. The wild-type full-length HSVtk was used. MREs were inserted in AvrII/XmaI sites immediately downstream of HSVtk. The let7 MRE consists of four sequences complementary to each of the four mature miRNAs let7g, let7a, let7d, and let7f (in this order) with sequence 5′ to 3′ (relative to transcription of HSVtk, which is encoded in reverse orientation in the pRG-tk lentiviral vector):

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AACTGTACAAACTACTACCTCAtaAACTATACAACCTACTACCTCAtaAACTATGCAACCTACTACCTCTtaAACTATAC
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The miR-142 MRE consists of four sequences complementary to miR-142-3p and miR-142-p5 with sequence 5′ to 3′ (relative to transcription of HSVtk, which is encoded in reverse orientation in the pRG-tk lentiviral vector):
AAGTAGGCTTTTACTTTTGTaTCCATAAAGTAGAAAAAC ACTACAtaAGTAGGCCTTTACTTTTGTaTCCATAAAGT AGGAAACACTACA

The Eos-C(3+) promoter enhancer was obtained from plasmid PL-SIN-EOS-C(3+)-EiP plasmid.22 Lentiviral vector packaging and transduction of iPSCs and fibroblasts were performed as described previously.43 The pRG-tk, pRG-tk-le17, and pRG-tk-miR142 vectors yielded comparable titers.

Flow cytometry. Undifferentiated iPSCs were dissociated with accutase and stained with Alexa Fluor 647-conjugated anti-Tra-1–81 or anti-Tra-1–60 or anti-SSEA3 or anti-SSEA4 (BD Biosciences, Franklin Lakes, NJ). Data were acquired in a LSRII cytometer (BD Biosciences) and analyzed with the FlowJo software (Tree Star, Ashland, OR). iPSCs transduced with control vectors expressing mCherry only and GFP only were used for compensation setup.

Teratoma formation assays. For teratoma formation assays, iPSCs were suspended in hES medium containing 10 µmol/l of the Rho-associated kinase (Rok) inhibitor Y-27632 (Tocris Bristol, UK). Approximately 2 million cells were injected subcutaneously in the flank of female 8-week-old NOD-SCID IL2Rgnull (NSG) mice (Jackson Laboratory, Bar Harbor, ME) together with Matrigel (BD Bioscience). One week after cell implantation, the mice were randomized to receive either phosphate-buffered saline or ganciclovir (at 5 mg/kg per day) administered via an osmotic minipump (Alzet #1007D) implanted subcutaneously. Minipumps were replaced every 7 days, for a total of 2 weeks of treatment. Mice were observed daily throughout the treatment period for signs of morbidity/mortality. Tumors were measured twice weekly using calipers and volume was calculated using the formula: length × width² × 0.52. Body weight was measured twice weekly using calipers, and volume was calculated of treatment. Mice were observed daily throughout the treatment.

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