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### Comparative Analysis of Enzyme and Pathway Engineering Strategies for 5FC-Mediated Suicide Gene Therapy Applications

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#### Abstract

Bacterial and yeast encoded cytosine deaminases (bCD and yCD, respectively) are widely investigated suicide enzymes used in combination with the prodrug 5-fluorocytosine (5FC) to achieve localized cytotoxicity. Yet characteristics such as poor turnover rates of 5FC (bCD) and enzyme thermolability (yCD) preclude their full therapeutic potential. We previously applied regio-specific random mutagenesis and computational design to create novel bCD and yCD variants with altered substrate preference (bCD1525) or increased thermostability (yCDdouble, yCD<sub>triple</sub>) to aid in overcoming these limitations. Others have utilized pathway engineering in which the microbial enzyme uracil phosphoribosyltransferase (UPRT) is fused with its respective CD, creating bCD/bUPRT or yCD/yUPRT. In this study, we evaluated whether the overlay of CD mutants onto their respective CD/UPRT fusion construct would further enhance 5FC activation, cancer cell prodrug sensitivity and bystander activity in vitro and in vivo. We show that all mutant fusion enzymes allowed for significant reductions in  $IC_{50}$  values relative to their mutant CD counterparts. However, in vivo the CD mutants displayed enhanced tumor growth inhibition capacity relative to the mutant fusions, with bCD<sub>1525</sub> displaying the greatest tumor growth inhibition and bystander activity. In summary, mutant bCD1525 appears to be the most effective of all bacterial or yeast CD or CD/UPRT enzymes examined and as such is likely to be the best choice to significantly improve the clinical outcome of CD/5FC suicide gene therapy applications.

#### Keywords

cytosine deaminase; uracil phosphoribosyltransferase; suicide gene therapy; bystander effect; 5-fluorocytosine

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#### Introduction

The cytosine deaminase (CD; EC 3.5.4.1)/5-fluorocytosine (5FC) system is one of the most well-characterized enzyme/prodrug systems in suicide gene therapy (SGT). Found primarily in microbes such as *Escherichia coli* and *Saccharomyces cerevisiae*, CD is a pyrimidine salvage pathway enzyme responsible for catalyzing the hydrolytic deamination of cytosine to form uracil and ammonia[1]. Similarly, CD (*codA* in *E. coli*, *FCY1* in *S. cerevisiae*) deaminates the prodrug 5FC into the potent antimetabolite and radiosensitizing agent, 5-fluorouracil (5FU). 5FU is further converted to the irreversible thymidylate synthase inhibitor, 5FdUMP, leading to depleted dTTP pools, DNA synthesis inhibition, and subsequent apoptosis[2-4]. Its cytotoxic activity is also extended by passive diffusion from CD-expressing to neighboring untransfected cells resulting in a phenomenon known as the bystander effect. This phenomenon is an essential part of SGT and serves to compensate, at least in part, for inefficient gene transfer seen with current delivery systems[1].

The CD/5FC system was originally developed using bacterial CD (bCD)[5]. Although promising, the therapeutic application of this system has been limited by the poor turnover rate ( $k_{cat}$ ) of 5FC by bCD. Alternatively, yeast CD (yCD) was later evaluated and shown to display superior kinetic properties toward 5FC (22-fold lower K<sub>m</sub> for 5FC) and increase the efficacy of the CD/5FC system approach[6, 7]. However, the thermolabile nature of yCD at 37°C precludes its full therapeutic potential. To overcome this limitation, computational design studies aimed at repacking the hydrophobic core of yCD identified two enzyme mutants (yCD<sub>double</sub> and yCD<sub>triple</sub>) with increased half-lives and melting temperatures that significantly increased the sensitivity of cancer cells to 5FC treatment as compared to wildtype yCD[8, 9]. Similarly, to alleviate the enzymatic limitations seen with bCD, regiospecific random mutagenesis studies targeting two regions flanking the bCD active site identified a superior bCD mutant, bCD<sub>1525</sub>, with improved kinetic properties to wild-type bCD. With a 19-fold shift in substrate preference for 5FC, bCD<sub>1525</sub> significantly reduced IC<sub>50</sub> levels, improved the bystander effect, and prevented tumor growth at 5FC levels where tumors expressing wild-type bCD were unresponsive[10].

In microbes, 5FU is directly converted to the downstream metabolite 5FUMP by uracil phosphoribosyltransferase (UPRT; EC 2.4.2.9). UPRT (*upp* in *E. coli, FUR1* in *S. cerevisiae*) is a member of the pyrimidine salvage pathway and is responsible for catalyzing the synthesis of UMP or 5FUMP from uracil or 5FU, respectively[11, 12]. Overexpression of UPRT in multiple cancer cell lines resulted in a marked increase in 5FU sensitization. However, it has been shown that systemic treatment of 5FU can lead to gastrointestinal and hematological toxicity[13, 14]. In addition, some human tumor cell lines display resistance to 5FU as a result of possible defects in downstream cellular metabolism or overexpression of 5FU catabolic enzymes[15]. To circumvent 5FU systemic toxicity and bypass 5FU resistance, co-expression of CD with UPRT and expression of bCD or yCD fused to UPRT (bCD/bUPRT and yCD/yUPRT) have both been tested and shown to enhance 5FC-mediated cell killing[16-20].

In the present study, we evaluated whether the overlay of previously created yeast and bacterial CD mutants onto their corresponding CD/UPRT fusion construct would result in a

cooperative increase in 5FC metabolism and cancer cell prodrug sensitivity *in vitro* and *in vivo*. Our data suggest that although previous studies have shown wild-type CD/UPRT fusions enhance prodrug sensitivity *in vitro* and *in vivo* compared to their CD counterparts, UPRT activity in fusion with our previously identified CD mutants negatively influences bystander activity *in vivo* and diminishes the therapeutic value of the CD/5FC system. Taken together, this comparative study reveals that the optimized bacterial mutant bCD<sub>1525</sub> alone displays ideal cell and tumor ablative characteristics for suicide gene therapy applications.

#### Materials and Methods

#### Materials

Oligonucleotides used to mutate and/or sequence bCD, yCD, bUPRT and yUPRT were obtained from Integrated DNA Technologies (Coralville, IA). Restriction endonucleases and T4 DNA ligase were purchased from New England Biolabs (Beverly, MA). The Wizard PCR prep kit (Promega, Madison, WI), HiSpeed Plasmid Midi Kit (Qiagen, Valencia, CA) and StrataPrep EF Plasmid Midikit (Stratagene, La Jolla, CA) were all used for DNA purification. Alamar Blue was purchased from Serotec Limited (Oxford, UK). All cell culture reagents were purchased from Gibco (Carlsbad, CA), and all other reagents were purchased from Sigma-Aldrich (St. Louis, MO) unless otherwise noted.

#### **Bacterial strains**

Escherichia coli strain XL1-Blue [F': :Tn10 proA<sup>+</sup>B<sup>+</sup> lacI<sup>q</sup> D(lacZ) M15/recA1 endA1 gyrA96 (NaI<sup>r</sup>) thi hsdR17 ( $r_k m_k^+$ ) supE44 relA1 lac] and NM522 [F<sup>+</sup> lacI<sup>q</sup> (lacZ)-M15 proA<sup>+</sup>B<sup>+</sup>/supE thi (lac-proAB) (hsdMS-mcrB)5( $r_k m_k McrBC^-$ )] were used for cloning and mutagenesis procedures. E. coli strains BM604 [HfrH thi galE (att $\lambda$ -bio) deoA103 deoC argA lysA cytR upp udp pyrF30] and GIA39 [thr dadB3 fhuA21 codA1 lacY1 tsk-95 glnV44(AS)  $\lambda^-$  pyrF101 his-108 argG6 ilvA634 thi-1 deoC1 glt-15] were first lysogenized with  $\lambda$ DE3 according to manufacturer protocol (Novagen, Madison, WI) and then used for UPRT and CD genetic complementation, respectively. The BM604 strain, a strain deficient in UPRT activity and orotidine 5-phosphate decarboxylase, was a kind gift from F. Bringel[21, 22], and the GIA39 strain, a strain deficient in CD and orotidine 5-phosphate decarboxylase, was obtained from the E. coli Genetic Stock Center (CGSC # 5594).

### Construction of bCD/bUPRT and yCD/yUPRT fusion mutant constructs and genetic complementation

The bCD/bUPRT fusion gene was first removed from pORF-*codA*: :*upp* (Invivogen, San Diego, CA), which carries functional bCD and bUPRT, using *NcoI/Hin*DIII and subcloned into *NcoI/Hin*DIII digested pETHT[23]. The yCD/yUPRT fusion gene was isolated as an *Xho*I(blunt-ended)/*Not*I fragment from pCI-neo*FCU1* (Transgene, France) and subcloned into *Eco*RI(blunt-ended)/*Not*I digested pETHT. Both bCD/bUPRT and yCD/yUPRT were then subjected to mutagenesis using the QuikChange site-directed mutagenesis kit from Stratagene (La Jolla, CA) to create the respective mutant fusion counterparts. Two pairs of oligonucleotides (MB455 5'-GGATTGATCTGCAAATCGCCGCTTTCCCTCAGG-3'/ MB456 5'-CCTGAGGGAAAGCGGCGATTTGCAGATCAATCC-3' and MB457 5'-CTTTGGTCACGATGATGTCTGCGGTCCGTGGTAT-3'/MB458 5'-

ATACCACGGACCGCAGACATCATCGTGACCAAAG-3') were used to introduce the bCD<sub>1525</sub> (V152A/F316C/D317G) sequence. Oligonucleotide pairs (MB462 5'-GCCTATGAGGAGGCGCTCTTAGGTTACAAAGAGGG-3'/MB463 5'-CCCTCTTTGTAACCTAAGAGCGCCTCCTCATAGGC-3' and MB466 5'-GACGATGAGAGGTGTAAAAAGCTCATGAAACAATTTATCG-3'/MB467 5'-CGATAAATTGTTTCATGAGCTTTTTACACCTCTCATCGTC-3') were used to introduce the yCD<sub>double</sub> (A23L/I140L) sequence, while yCD<sub>triple</sub> (A23L/V108I/I140L) used (MB464 5'-GGTATTCCACGCTGTGTTATCGGT-3'/MB465 5'-CGTTCTCACCGATAACACAGCGTGGAATACC-3') in addition to those used for the yCD<sub>double</sub> sequence. DNA sequencing analysis confirmed the presence of correct mutations in the mutant fusion constructs. The resulting plasmids were designated pETHT:bCD<sub>1525</sub>/ bUPRT, pETHT:yCD<sub>double</sub>/yUPRT and pETHT:yCD<sub>triple</sub>/yUPRT, respectively. The construction of pETHT:bCD, pETHT:bCD<sub>1525</sub>, pETHT:yCD, pETHT:yCD<sub>double</sub> and pETHT:yCD<sub>triple</sub> were as previously described[9,10]. For complementation studies, E. coli strains GIA39(DE3) or BM604(DE3) harboring either pETHT, wild-type or mutant CD, or wild-type or mutant CD/UPRT constructs were grown at 37°C on either CD[24,25] or UPRT[21,22] complementation plates.

#### Construction of mammalian expression vectors

The bCD/bUPRT fusion gene was subcloned from pETHT:bCD/bUPRT into *Eco*RV/*Xho*I digested pcDNA6/myc-HisB (Invitrogen, Carlsbad, CA) as an *Nco*I(blunt-ended)/*Xho*I fragment, while the yCD/yUPRT fusion gene was subcloned from pETHT:yCD/yUPRT into *Eco*RV/*Not*I digested pcDNA6/myc-HisB as an *Nco*I(blunt ended)/*Not*I fragment creating pcDNA:bCD/bUPRT and pcDNA:yCD/yUPRT, respectively. Following restriction enzyme verification, DNA sequencing analysis confirmed the presence of both fusion genes. Both pcDNA:bCD/bUPRT and pcDNA:yCD/yUPRT were then subjected to mutagenesis using the Stratagene QuikChange site-directed mutagenesis kit to create their mutant fusion counterparts as described above. Resulting plasmids were designated pcDNA:bCD<sub>1525</sub>/ bUPRT, pcDNA:yCD<sub>double</sub>/yUPRT and pcDNA:yCD<sub>triple</sub>/yUPRT. DNA sequencing analysis confirmed the presence of poth sequencing analysis confirmed the mutant fusion constructs. The construction of pcDNA:bCD, pcDNA:yCD, and their mutant constructs was described previously by Fuchita *et al.* (2009) and Stolworthy *et al.* (2008), respectively.

#### Cell line

Rat C6 glioma cells were purchased from ATCC (Manasass, VA) and grown in Dulbecco's Modified Essential Medium containing 5% fetal bovine serum, 1mM sodium pyruvate, 10mM HEPES, 100µM nonessential amino acids, 100U/mL penicillin G, 10µg/mL streptomycin sulfate, 292µg/mL L-glutamine, 100µM sodium citrate and 0.0014% NaCl. Cells were maintained in a humidified incubator at 37°C in 5% CO<sub>2</sub>.

#### In vitro cytotoxicity assay

Stable transfectants were created by the addition of 1µg purified endotoxin-free DNA (Stratagene, La Jolla, CA) (pcDNA (empty vector), pcDNA:bCD, pcDNA:bCD<sub>1525</sub>, pcDNA:bCD/bUPRT, pcDNA:bCD<sub>1525</sub>/bUPRT, pcDNA:yCD, pcDNA:yCD<sub>double</sub>,

pcDNA:yCD<sub>triple</sub>, pcDNA:yCD/yUPRT, pcDNA:yCD<sub>double</sub>/yUPRT, or pcDNA:yCD<sub>triple</sub>/ yUPRT) to  $1\times10^5$  rat C6 glioma cells by lipofection using FuGENE 6 transfection reagent (Roche Diagnostic, Penzberg, Germany) at a 3:1 ratio as described by the manufacturer. Stable transfectants were selected by culture in media supplemented with 4µg/mL blasticidin.

Immunoblot analysis using rabbit polyclonal antiserum followed by goat anti-rabbit-AP conjugated secondary antibody (Bio-Rad, Hercules, CA) was performed to determine protein expression levels, as previously described[9]. Antisera to purified bCD, bUPRT or yCD was generated by Harlan Laboratories (Indianapolis, IN), while antiserum for yUPRT was generated by Invitrogen (Carlsbad, CA). An AP Conjugate Substrate Kit (Bio-Rad) was used for colorimetric detection. Stable transfectants confirmed by immunoblot were used for *in vitro* cytotoxicity assays. Pools of transfectants were plated at an initial density of 500 cells/well in 96-well microtiter plates. Following cell adherence overnight, 5FC (0-10mM) was added in sets of eight wells for each concentration tested. After a 7 day incubation period, cell viability was determined by the addition of the redox indicator dye Alamar Blue and fluorometric readings at 530/590nm using a Biotek Synergy HT microplate reader several hours later as described by the manufacturer. Data were plotted with the standard error of the mean and at least three replicates were performed.

#### In vitro bystander experiment

Rat C6 glioma cells stably transfected with pcDNA were mixed at different ratios with stable transfectants harboring pcDNA:bCD, pcDNA:bCD<sub>1525</sub>, pcDNA:bCD/bUPRT, pcDNA:bCD<sub>1525</sub>/bUPRT, pcDNA:yCD, pcDNA:yCD<sub>double</sub>, pcDNA:yCD<sub>triple</sub>, pcDNA:yCD/yUPRT, pcDNA:yCD<sub>double</sub>/yUPRT, or pcDNA:yCD<sub>triple</sub>/yUPRT. Mixed cells were then plated at an initial density of 500 cells/well in 96-well microtiter plates. Following cell adherence overnight, 4 or 10mM 5FC was added to the plated cells. Cell survival was determined as described above.

#### Xenograft tumor model

Pools of rat C6 glioma cells stably transfected with pcDNA, pcDNA:bCD<sub>1525</sub>, pcDNA:bCD/bUPRT, pcDNA:bCD<sub>1525</sub>/bUPRT, pcDNA:yCD<sub>double</sub>, pcDNA:yCD<sub>triple</sub>, pcDNA:yCD/yUPRT, or pcDNA:yCD<sub>double</sub>/yUPRT ( $0.5 \times 10^6$  cells in 200µl PBS at pH 7.2, n=5 for each group) were injected subcutaneously into the left and right flanks of five to six week old female, athymic NCr-*nu/nu* mice (National Cancer Institute, Frederick, MD). For the bystander xenograft tumor model experiments, pools of C6 glioma cells stably transfected with pcDNA:bCD<sub>1525</sub>, pcDNA:bCD<sub>1525</sub>/bUPRT, pcDNA:yCD<sub>double</sub>, pcDNA:yCD<sub>double</sub>/yUPRT, pcDNA:yCD<sub>triple</sub>, or pcDNA:yCD<sub>double</sub>, pcDNA:yCD<sub>double</sub>/yUPRT, pcDNA:yCD<sub>triple</sub>, or pcDNA:yCD<sub>triple</sub>/UPRT were mixed with pools of C6 glioma cells stably transfected with pcDNA at a ratio of 80:20 (vector : suicide gene) then subcutaneously injected into the left and right flanks of nude mice (n=5). When tumors reached 3-4mm in diameter (day 0), PBS or 5FC at 375mg kg<sup>-1</sup> was administered via intraperitoneal injection twice a day for up to 21 days. Tumor volume was monitored by caliper measurement (length, width, and height) every other day, starting at day 0, until mice were euthanized. Tumor volume was calculated using the formula:  $4/3\pi$  ((width × length × height)/2) and analyzed for statistical significance using the Student's T-test. All mice with

heavy tumor burden were sacrificed in compliance with IACUC recommended practices and guidelines.

#### Results

#### Genetic complementation

To determine whether bCD or yCD mutations overlaid onto their respective CD/UPRT constructs maintained dual functionality, previously established *E. coli* genetic complementation assays evaluating CD or UPRT activity were performed using GIA39 (DE3) or BM604 (DE3) *E. coli*, respectively (see Materials and Methods). In these assays, pETHT, pETHT:bCD, pETHT:bCD<sub>1525</sub>, pETHT:bCD/bUPRT, pETHT:bCD<sub>1525</sub>/bUPRT, pETHT:yCD, pETHT:yCD<sub>double</sub>, pETHT:yCD<sub>triple</sub>, pETHT:yCD/yUPRT, pETHT:yCD<sub>double</sub>/yUPRT and pETHT:yCD<sub>triple</sub>/yUPRT were used to transform *E. coli* and bacteria were then plated onto appropriate selective media supplemented with carbenicillin. For CD genetic complementation, all CD and CD/UPRT constructs were able to complement the UPRT deficiency of BM604 (DE3). Cells harboring pETHT were unable to complement the deficiency of either *E. coli* strain and all cultures grew on non-selective media (data not shown). These results indicate all wild-type and mutant fusion constructs maintain dual functionality, which is in accordance to previous studies [16, 17].

#### In vitro prodrug sensitivity assays

To examine the ability of mutant fusion enzymes to sensitize cancer cells to 5FC, wild-type or mutant CD or CD/UPRT constructs were first cloned into the expression vector pcDNA and the resulting plasmids were used to transfect rat C6 glioma cells. Pools of stable transfectants were then exposed to a wide concentration range of 5FC (0-10mM) and cell survival determined after 7 days (Figure 1). Consistent with a previous report by Fuchita et al. (2009)[10], bCD<sub>1525</sub>-transfected cells conferred increased sensitivity to 5FC compared to wild-type bCD-transfected cells. This sensitivity is a direct reflection of the increased 5FU levels produced in the bCD<sub>1525</sub>-transfected cells[10]. While wild-type bCD/bUPRTtransfected cells failed to lower IC<sub>50</sub> levels in comparison to bCD<sub>1525</sub>, bCD<sub>1525</sub>/bUPRTtransfected cells displayed a 4.5-fold decrease (<0.1mM) in IC<sub>50</sub> value (Figure 1A). As previously reported by Stolworthy et al. (2008)[9], both yCD<sub>double</sub> and yCD<sub>triple</sub> had lower IC50 values than wild-type yCD-transfected C6 cells. Wild-type yCD/yUPRT displayed a similar IC<sub>50</sub> value to yCD<sub>triple</sub> (1.8mM) and a 2.6-fold lower IC<sub>50</sub> than yCD<sub>double</sub> (4.75mM). The yCD/yUPRT fusion mutants further lowered IC<sub>50</sub> values 3.3-fold (yCD<sub>double</sub>/yUPRT) and 6-fold (yCD<sub>triple</sub>/yUPRT) compared to wild-type yCD/yUPRT. Interestingly, of the two yCD/yUPRT fusion mutants, yCD<sub>double</sub>/yUPRT showed the greatest reduction in IC<sub>50</sub> (8.5-fold) in comparison to its yCD mutant counterpart, yCD<sub>double</sub>. Both yCD<sub>double</sub>/yUPRT and yCD<sub>triple</sub>/yUPRT exhibited similar IC<sub>50</sub> values of 0.55 and 0.3mM, respectively (Figure 1B). When wild-type or mutant CD/UPRT-transfected cells were analyzed together, bCD1525/bUPRT-transfected cells displayed the greatest degree of 5FC sensitivity (Figure 1C).

#### *In vitro* bystander activity analyses

One key feature of the CD/5FC system that poses a potential advantage over other SGT systems, such as the HSVTK/GCV system, is the potent bystander effect that results from the passive diffusion of 5FU from transfected to untransfected cancer cells. This potent bystander effect can help overcome poor transduction efficiencies seen with current delivery systems and may increase the likelihood of complete tumor eradication [26, 27]. In the case of the CD/UPRT fusion constructs, UPRT activity may induce bystander characteristics that more heavily rely on alternative mechanisms of action, such as the transfer of active metabolites via gap junctions, such as that observed with the HSVTK/GCV system. To investigate the ability of mutant CD/UPRT fusion constructs to elicit a potent bystander effect, cocultures of vector and wild-type or mutant CD or CD/UPRT-transfected cells were mixed at different ratios and subjected to a set concentration of 5FC for 7 days (see Materials and Methods). Whereas a cell population containing 15% bCD<sub>1525</sub>-transfected cells is sufficient to induce 50% cell killing, bCD1525/bUPRT-transfected cells failed to display significant bystander cell killing activity at the 5FC concentration used (Figure 2A). Alternatively, cell populations containing yCD<sub>double</sub>/yUPRT or yCD<sub>triple</sub>/yUPRTtransfected cells elicited a robust bystander effect as indicated by 50% tumor cell killing when only 10-15% of the cell population expressed these mutant fusion enzymes. Indeed, when all wild-type or mutant CD/UPRT constructs were compared together, the yeast fusion mutants demonstrated the most significant bystander activity (Figure 2B). Both wild-type bCD/bUPRT and yCD/yUPRT showed slight to modest cell killing effects (Figure 2B), while yCD<sub>double</sub> and yCD<sub>triple</sub> failed to display significant bystander activity at the 5FC concentration used (data not shown). Clonogenic assays[28] examining bystander cell populations containing 100% suicide gene-transfected cells revealed these populations are unable to form colonies (data not shown). Thus detection of some survival for the yeast fusion mutants (Figure 2B) most likely reflects an artifact of the Alamar Blue dye assays rather than cell viability.

#### Xenograft tumor model

Constructs that conferred the best combined cell killing and bystander activity were selected for further characterization in an *in vivo* xenograft tumor model. The yCD<sub>double</sub>/yUPRT construct was chosen for further *in vivo* characterization over yCD<sub>triple</sub>/yUPRT as both displayed similar *in vitro* prodrug sensitivity and bystander activity, yet yCD<sub>double</sub>/yUPRT showed the greatest improvement over its respective yCD mutant counterpart. Figure 3 illustrates tumor growth in athymic NCr-*nu/nu* mice treated with 5FC (375mg kg<sup>-1</sup>, twice a day) or phosphate buffered saline (PBS) over time. In Figure 3A, mice bearing bCD<sub>1525</sub> or bCD<sub>1525</sub>/bUPRT-transfected tumors showed indistinguishable growth rates to those bearing vector-transfected tumors when treated with PBS. However, when mice bearing bCD<sub>1525</sub> or bCD<sub>1525</sub>/bUPRT-transfected tumors were treated with 5FC a significant growth restriction emerged starting day 4 (*P* 0.05) in comparison to those bearing vector-transfected tumors. Mice bearing bCD<sub>1525</sub>-transfected tumors and treated with 5FC displayed the greatest restriction in tumor growth (day 16 mean tumor volume: 27.16mm<sup>3</sup>) (Figure 3B; Table 1). Figure 3C shows all PBS treated groups to have statistically insignificant differences in growth rates. In contrast, tumor cell growth rates in mice bearing yCD<sub>double</sub> or yCD<sub>double</sub>/

yUPRT-transfected tumors were significantly reduced starting day 8 relative to mice bearing vector-transfected tumors when treated with 5FC. Following the same trend as the bCD<sub>1525</sub> versus bCD<sub>1525</sub>/bUPRT *in vivo* results, the single gene construct yCD<sub>double</sub> displayed the greatest ability to restrict tumor growth (day 16 mean tumor volume: 269.31mm<sup>3</sup>) (Figure 3D; Table 2). Neither wild-type bCD/bUPRT nor yCD/yUPRT-transfected tumors displayed significant sensitivity to 5FC (data not shown). Based on tumor growth inhibition rates, bCD<sub>1525</sub>-transfected tumors appear to be the most sensitive to 5FC treatment of all tested constructs. All PBS treated mice and 5FC treated mice bearing vector-transfected tumors were sacrificed at day 16 or 18 due to heavy tumor burden.

#### Bystander xenograft tumor model

As the xenograft tumor model results did not directly substantiate our *in vitro* sensitivity assay results, we next sought to determine whether the tumor microenvironment may alter or influence bystander activity in vivo. To this end, an 80:20 mixture of vector transfected cells to cells harboring previously tested mutant CD or mutant CD/UPRT fusion gene were injected into athymic NCr-nu/nu mice. Figure 4 illustrates tumor growth in mice treated with 5FC (375mg kg<sup>-1</sup>, twice a day) or PBS over time. Results show  $bCD_{1525}$  to display the greatest bystander activity and response to 5FC treatment as indicated by substantial tumor growth inhibition (day 16 mean tumor volume: 179.57mm<sup>3</sup>), while bCD<sub>1525</sub>/bUPRT also exhibited a significant bystander killing effect (day 16 mean tumor volume: 432.51 mm<sup>3</sup>). Conversely, yCD<sub>double</sub>/yUPRT did not evoke sufficient bystander activity able to suppress tumor growth and yCD<sub>double</sub> displayed minimal bystander activity (significance starting day 14 (P 0.05) in comparison to PBS treated vector: yCD<sub>double</sub>-expressing tumors) (Figure 4B, Table 3). Similar results were seen with yCD<sub>triple</sub> and yCD<sub>triple</sub>/yUPRT (data not shown). All tumors from PBS treated mice groups grew at approximately the same rate ( $P \, 0.05$ ) (Figure 4A). The variability in tumor volumes at day 16 may be attributed to necrotic centers or the amorphous nature of the large tumors that prevented consistent measurements.

#### Discussion

Although multiple preclinical studies have demonstrated the feasibility of suicide gene therapy for cancer treatment, it is evident that further improvements are necessary to achieve clinical success. To this end, we and other research teams have utilized enzyme and pathway engineering strategies to enhance the catalytic properties or increase the stability of multiple suicide enzymes. These engineered enzymes have been shown to lower necessary prodrug doses used during therapy, enhance cancer cell prodrug sensitivity and improve bystander activity (reviewed in Ardiani *et al.*, manuscript in preparation). Our primary goal for this study was to evaluate whether the combination of enzyme and pathway engineering strategies using the CD/5FC approach would further augment prodrug activation and cooperatively increase cancer cell prodrug sensitivity. This comparative analysis study was also used as a means to identify the most suitable bacterial or yeast wild-type or mutant CD or CD/UPRT enzyme for clinical application.

Earlier studies directly comparing wild-type bacterial and yeast CD revealed that, despite its thermolability, the use of yCD enhanced HT29 human colon cancer cell radiosensitization,

the bystander effect, and the overall therapeutic value of the CD/5FC treatment strategy over bCD[6, 7]. Later studies comparing wild-type bCD, yCD, and the yCD/yUPRT fusion in a rat hepatoma model confirmed these findings and also provided evidence that the yCD/ yUPRT fusion further enhanced cancer cell 5FC sensitivity in vitro and in vivo over both bCD and yCD alone[29]. Prior to these studies, the yCD/yUPRT fusion was characterized in multiple cancer cell lines and shown to significantly improve bystander activity in vitro relative to wild-type yCD[17]. Similarly, glioblastoma and colon cancer cell lines expressing the bCD/bUPRT fusion displayed enhanced 5FC sensitivity and bystander activity in *vitro*[30,31] and significant tumor growth inhibition *in vivo*[30] over wild-type bCD. Our results recapitulate the superiority of the CD/UPRT fusions over their wild-type CD counterparts and also reveal that wild-type yCD/yUPRT-transfected rat C6 glioma cells are slightly more sensitive to 5FC than wild-type bCD/bUPRT-transfected cells and maintain greater bystander activity. The combination of elevated yCD activity[17], more efficient kinetic parameters toward 5FC[6], and fusion to yUPRT[17,29] may account for these enhanced in vitro results. In the context of wild-type CD or CD/UPRT enzymes, all combined results indicate the yCD/yUPRT fusion to be the most effective prodrug activation system.

An advantage of SGT over conventional treatments such as systemic chemotherapy is its ability to selectively convert prodrugs at targeted tumor sites resulting in local, high antimetabolite concentrations. However, use of the CD/5FC system to date has been limited to only a handful of conducted clinical trials[32-34]. These trials have been hindered in part by inefficient gene delivery and poor prodrug activation at the 5FC doses used. Previously, we reported that yCD<sub>double</sub> and yCD<sub>triple</sub>-transfected rat C6 glioma cells and several bCD<sub>1525</sub>-transfected cell lines were rendered more sensitive to 5FC than wild-type CDtransfected cells at low 5FC concentrations[9, 10]. Our current data are consistent with these reports and also show that bCD1525 displayed the lowest relative IC50 value of all CD mutants tested. Furthermore, bCD<sub>1525</sub> was the only CD mutant able to display significant bystander activity in vitro at the 5FC dose tested. Wild-type yCD, although previously revealed to significantly increase the therapeutic effect of the CD/5FC system relative to wild-type bCD, showed minimal relative therapeutic value in vitro as compared to the bacterial mutant bCD<sub>1525</sub>. When tested in vivo, our xenograft and bystander tumor model studies reiterated the superior antitumor and bystander activity of bCD<sub>1525</sub> relative to the yCD mutant, yCD<sub>double</sub>, at a low 5FC concentration. Similar results for yCD<sub>double</sub> were seen with yCD<sub>triple</sub> in the bystander xenograft tumor model study (data not shown). As a single gene, the bacterial mutant  $bCD_{1525}$  is therefore the most effective CD mutant tested and its use in cancer gene therapy may allow for reduced, less debilitating 5FC treatment doses for tumor ablation. Based on previous work with these variants[9, 10] and the reports by many other investigators regarding the ability to sensitize many different cancer cell lines to 5FC using CD or CD/UPRT fusions[3-7, 13, 15-20, 26,27, 29-37,38] the results shown here will likely be relevant to a number of cancer types.

As current delivery systems pose a major limitation for successful cancer gene therapy, a robust bystander effect is crucial for its successful clinical application[1]. By introducing UPRT activity to the CD/5FC system, it is feasible that more efficient conversion of 5FU to

its active metabolites may diminish available, diffusible 5FU pools thereby reducing the bystander effect and in turn its therapeutic value. However, as described above, for wild-type CD/UPRT constructs the presence of UPRT appears to enhance bystander activity *in vitro*. This phenomenon is not likewise observed when the CD mutants are present in the CD/UPRT fusion constructs.

Although it displayed the greatest reduction in IC<sub>50</sub> value of all constructs tested, bCD<sub>1525</sub>/ bUPRT lacked significant bystander activity *in vitro*. Prior HPLC studies measuring 5FU levels released into culture medium revealed SW480 colon cancer cells expressing wild-type bCD or bCD/bUPRT produce similar 5FU levels; yet, the bCD/bUPRT fusion displayed superior bystander activity to bCD alone[30]. While the concerted action of bCD and bUPRT may account for the ability of bCD<sub>1525</sub>/bUPRT to confer increased cancer cell 5FC sensitivity *in vitro*, it may also account for the decreased bystander activity seen *in vitro* and *in vivo* for bCD<sub>1525</sub>/bUPRT relative to bCD<sub>1525</sub> alone. In this case, the transport of 5FUMP and other downstream metabolites to neighboring cells may be restricted by the inability of charged molecules to diffuse across membranes, thus diminishing the bystander activity. Our *in vitro* and *in vivo* bystander results are consistent with this hypothesis.

Like bCD1525/bUPRT, both yCDdouble/yUPRT and yCDtriple/yUPRT displayed significant reductions in IC50 values compared to their CD mutant counterparts. However, unlike bCD<sub>1525</sub>/bUPRT, the yeast fusion mutants exhibited increased bystander activity in vitro (data not shown). This increased bystander activity may be directly related to elevated CD activity and extracellular 5FU levels. For example, previous studies have shown that cells co-expressing yCD and yUPRT enzymes fail to exhibit enhanced bystander cell killing, or increased 5FU levels, relative to those expressing yCD alone [17]. Enhanced yCD half-life and melting temperature may also be contributing factors. Indeed, the fusion of yCD with other proteins has been shown to prolong yCD half-life and increase its thermostability[35, 36]. For instance, fusion of wild-type yCD to the hyaluronan binding domain of tumor necrosis factor alpha-stimulated gene-6 (TSG-6) prolonged the enzymatic activity of yCD at physiological temperatures and increased its apparent melting temperature[35]. This may explain the similar IC<sub>50</sub> values seen between wild-type yCD/yUPRT and the thermostabilized yCD<sub>triple</sub>. Our in vitro results also point to the overlay of yCD mutants onto yCD/yUPRT as a potential means to further elevate yCD melting temperature or half-life activity as the mutant fusions displayed lower IC50 values relative to the yCD mutants alone. This was more prevalent for yCD<sub>double</sub>/yUPRT presumably due to the already increased half-life of yCD<sub>triple</sub> (117 hrs at 50°C) compared to yCD<sub>double</sub> (21 hrs at 50°C)[8]. Although superior in cell killing and bystander activity in vitro, decreased amounts of diffusible 5FU due to UPRT activity may account for the decreased therapeutic value of yCD<sub>double</sub>/yUPRT compared to yCD<sub>double</sub> alone in vivo. We are currently investigating 5FC metabolite levels produced from mutant CD and CD/UPRT fusion transfected cells using HPLC analysis to ascertain the metabolic flux in these various transfected cells lines as a means to correlate antimetabolite levels with cell killing potency.

In conclusion, this comparative analysis study demonstrates that while enzyme and pathway engineering approaches separately enhance 5FC-mediated cancer cell death for the CD/5FC system, their concomitant use may negatively influence the system's therapeutic value. With

its enhanced catalytic properties and striking antitumor activity at low prodrug doses, the bacterial mutant  $bCD_{1525}$  has promising clinical application and is a prime candidate for use in combinatorial strategies. Most notably,  $bCD_{1525}$  may selectively enhance adjuvant radiation therapy as its product, 5FU, is a potent radiosensitizer[2, 37]. Likewise antineoplastic agents[19], immunomodulatory molecules[38,39], and oncolytic virus delivery vehicles[40] have all enhanced the therapeutic outcome of the CD/5FC treatment strategy. Thus, a multimodal approach using  $bCD_{1525}$  may move suicide gene therapy from a promising, alternative anticancer strategy to a potent treatment modality for complete tumor ablation.

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#### Figure 1.

Sensitivity of wild-type or mutant CD or CD/UPRT-transfected rat C6 glioma cells to 5FC. Pools of stable transfectants containing pcDNA, (**A**) wild-type or mutant bCD or bCD/ bUPRT, (**B**) wild-type or mutant yCD or yCD/yUPRT, or (**C**) wild-type or mutant bCD/ bUPRT or yCD/yUPRT were evaluated for 5FC sensitivity. After 7 days of 5FC treatment, cell survival was measured by Alamar Blue treatment according to the manufacturer's protocol with untreated (control) samples (0mM 5FC) set to 100% cell survival. Each data

point (mean  $\pm$  SEM; n = 3; performed with 24 replicates) is expressed as a percentage of the value for control samples.

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#### Figure 2.

*In vitro* bystander analyses of rat C6 glioma transfectants treated with 5FC. Pools of C6 cells containing pcDNA were mixed with cells harboring (**A**) wild-type or mutant bCD or bCD/ bUPRT or (**B**) wild-type or mutant bCD/bUPRT or yCD/yUPRT at different ratios and subjected to 4mM (**A**) or 10mM (**B**) 5FC for 7 days. Cell survival was measured by Alamar Blue treatment and each data point (mean  $\pm$  SEM; n = 3; performed with 24 replicates) is expressed as a percentage of the value for control wells containing 100% pcDNA-transfected cells.

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#### Figure 3.

Xenograft tumor growth course of bCD or yCD mutant or mutant fusion enzymes. Pools of stably transfected rat C6 glioma cells expressing pcDNA (×), pcDNA:bCD<sub>1525</sub> ( $\Box$ ), pcDNA:bCD<sub>1525</sub>/bUPRT ( $\blacksquare$ ), pcDNA:yCD<sub>double</sub> ( $\triangle$ ), or pcDNA:yCD<sub>double</sub>/yUPRT ( $\blacktriangle$ ) were used to seed tumors in nude mice (n = 5 for each group). When tumor size reached 3-4mm (Day 0), PBS (**A**,**C**) or 375mg kg<sup>-1</sup> 5FC (**B**, **D**) was administered intraperitoneally twice a day for up to 21 days. Asterisks (\*) denote statistical significance (P 0.05) in tumor sizes between mice bearing mutant CD or CD/UPRT-transfected cells to those bearing pcDNA-transfected cells, while (†) denotes statistical significance (P 0.05) in tumor sizes between mice bearing mutant CD-transfected cells to those bearing mutant CD/UPRT-transfected cells (**B**,**D**).

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#### Figure 4.

Bystander xenograft tumor model of bCD or yCD mutant and mutant fusion enzymes. Pools of rat C6 glioma cells transfected with pcDNA were mixed with cells stably transfected with pcDNA:bCD<sub>1525</sub> ( $\Box$ ), pcDNA:bCD<sub>1525</sub>/bUPRT ( $\blacksquare$ ), pcDNA:yCD<sub>double</sub>( $\triangle$ ), or pcDNA:yCD<sub>double</sub>/yUPRT ( $\blacktriangle$ ) and then used to seed tumors in nude mice (n = 5 for each group). When tumor size reached 3-4mm (Day 0), PBS ( $\blacktriangle$ ) or 375mg kg<sup>-1</sup> 5FC ( $\blacksquare$ ) was administered intraperitoneally twice a day for up to 21 days. Asterisks (\*) denote statistical significance (P 0.05) in tumor sizes between mice bearing 20% bCD<sub>1525</sub>-transfected cells to those bearing 20% yCD<sub>double</sub> or yCD<sub>double</sub>/yUPRT-transfected cells, while (†) denotes statistical significance (P 0.05) in tumor sizes between mice bearing 20% bCD<sub>1525</sub>-transfected cells to those bearing 20% bCD<sub>1525</sub>-transfected cells ( $\blacksquare$ ).

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## Table 1

In vivo growth of rat C6 tumor cells transfected with pcDNA, pcDNA:bCD<sub>1525</sub> or pcDNA:bCD<sub>1525</sub>/bUPRT in response to PBS or 5FC treatment.

	Mea	n tumor volume (m	.m <sup>3</sup> )
	Day 0	Da	y 16
	5FC (375mg kg <sup>-1</sup> )	PBS	5FC (375mg kg <sup>-1</sup> )
pcDNA	14.98 (1.26) <sup>a</sup>	1303.79 (123.55)	754.86 (131.38) <sup>**</sup>
bCD <sub>1525</sub>	18.37 (4.09)	1282.14 (234.76)	27.16 (4.63)
bCD <sub>1525</sub> /bUPRT	24.17 (4.53)	1119.38 (200.54)	87.42 (20.44) <sup>**</sup>
<sup>a</sup> Standard error mea	ın.		

\*\* P 0.01 (bCD1525 vs. bCD1525/bUPRT tumor volume) or 0.001 (bCD1525 vs. pcDNA tumor volume) (n = 5).

5FC, 5-fluorocytosine; bCD, bacterial cytosine deaminase; bUPRT, bacterial uracil phosphoribosyltransferase; PBS, phosphate buffered saline.

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# Table 2

In vivo growth of rat C6 tumor cells transfected with pcDNA, pcDNA; yCD<sub>double</sub> or pcDNA; yCD<sub>double</sub>/yUPRT in response to PBS or 5FC treatment.

	Mea	n tumor volume (r	am <sup>3</sup> )
	Day 0	Ds	iy 16
	5FC (375mg kg <sup>-1</sup> )	PBS	5FC (375mg kg <sup>-1</sup> )
pcDNA	$8.82\ (0.91)^{d}$	890.51 (103.22)	864.23 (142.17) <sup>**</sup>
yCD <sub>double</sub>	11.75 (1.4)	735.92 (83.99)	269.31 (43.47)
yCD <sub>double</sub> /yUPRT	10.55 (1.07)	728.28 (166.02)	529.37 (67.67)**
a Standard error mean			

\*\* P = 0.05 (yCDdouble vs. yCDdouble/yUPRT tumor volume) or 0.005 (yCDdouble vs. pcDNA tumor volume) (n = 5).

5FC, 5-fluorocytosine; yCD, yeast cytosine deaminase; yUPRT, yeast uracil phosphoribosyltransferase; PBS, phosphate buffered saline.

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# Table 3

Bystander in vivo growth of rat C6 tumors containing an 80:20 ratio of vector transfected (pcDNA) to transfected (yCD<sub>double</sub>, yCD<sub>double</sub>/yUPRT, bCD<sub>1525</sub>, or bCD<sub>1525</sub>/bUPRT) cells in response to PBS or 5FC treatment.

	Mea	n tumor volume (n	nm <sup>3</sup> )
	Day 0	D	ay 16
	5FC (375mg kg <sup>-1</sup> )	PBS	<b>5</b> FC (375mg kg <sup>-1</sup> )
yCD <sub>double</sub>	$10.42 (0.96)^{d}$	1010.8 (128.84)	$718.88 \left(82.2\right)^{**}$
yCD <sub>double</sub> /yUPRT	6.78 (0.76)	705.98 (170.6)	$810.62\ (142.01)^{**}$
bCD <sub>1525</sub>	8.88 (1.19)	1066.5 (196.17)	179.57 (28.9)
bCD <sub>1525</sub> /bUPRT	8.23 (0.65)	638.0 (70.33)	432.51 (53.68) <sup>**</sup>

<sup>a</sup>Standard error mean.

\*\* 0.005 (bCD1525 vs. bCD152S/bUPRT tumor volume) or 0.0005 (bCD1525 vs. yCDdouble and yCDdouble/yUPRT tumor volumes) (n = 5).

5FC, 5-fluorocytosine; bCD or yCD, bacterial or yeast cytosine deaminase; bUPRT or yUPRT, bacterial or yeast uracil phosphoribosyltransferase; PBS, phosphate buffered saline.