Short Communication

Activation of the Antiviral Prodrug Oseltamivir Is Impaired by Two Newly Identified Carboxylesterase 1 Variants

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ABSTRACT:
Oseltamivir phosphate is an ethyl ester prodrug widely used in the treatment and prevention of both Influenzavirus A and B infections. The conversion of oseltamivir to its active metabolite oseltamivir carboxylate is dependent on ester hydrolysis mediated by carboxylesterase 1 (CES1). We recently identified two functional CES1 variants p.Gly143Glu and p.Asp260fs in a research subject who displayed significant impairment in his ability to metabolize the selective CES1 substrate, methylphenidate. In vitro functional studies demonstrated that the presence of either of the two mutations can result in severe reductions in the catalytic efficiency of CES1 toward methylphenidate, which is required for hydrolysis and pharmacological deactivation. The aim of the present study was to investigate the function of these mutations on activating (hydrolyzing) oseltamivir to oseltamivir carboxylate using the cell lines expressing wild type (WT) and each mutant CES1. In vitro incubation studies demonstrated that the S9 fractions prepared from the cells transfected with WT CES1 and human liver tissues rapidly convert oseltamivir to oseltamivir carboxylate. However, the catalytic activity of the mutant hydrolyases was dramatically hindered. The V_max value of p.Gly143Glu was approximately 25% of that of WT enzyme, whereas the catalytic activity of p.Asp260fs was negligible. These results suggest that the therapeutic efficacy of oseltamivir could be compromised in treated patients expressing either functional CES1 mutation. Furthermore, the potential for increased adverse effects or toxicity as a result of exposure to high concentrations of the nonhydrolyzed prodrug should be considered.

Oseltamivir phosphate (Tamiflu; Roche, Nutley, NJ) is widely used in the treatment and prophylaxis of both Influenzavirus A and B infections. In addition, oseltamivir may be effective in preventing or treating avian influenza or so-called “bird flu.” Oseltamivir is an ester prodrug and, in general, it is readily converted to its active form oseltamivir carboxylate mediated by hepatic carboxylesterase 1 (CES1) (Fig. 1) (Shi et al., 2006). The active metabolite exerts its antiviral effects via the selective inhibition of neuraminidase.

Carboxylesterases are members of the αβ hydrolase -fold family and expressed in many tissues, especially in the liver, small intestine, and lung (Satoh and Hosokawa, 2006; Ross and Crow, 2007). The major human carboxylesterases include CES1 (UniprotKB/Swiss-Prot P23141) and carboxylesterase 2 (CES2) (UniprotKB/Swiss-Prot O00748). CES1 and CES2 are largely distinguished from one another by their substrate specificity and tissue distribution (Imai et al., 2006; Satoh and Hosokawa, 2006). CES1 more readily catalyzes substrates with a relatively large acyl group and small alcohol group such as methylphenidate, temocapril, and oseltamivir (Sun et al., 2004; Imai et al., 2005; Shi et al., 2006). In contrast, CES2 preferentially hydrolyzes compounds bearing a small acyl moiety and bulky alcohol group, which includes agents such as cocaine and irinotecan. CES1 predominates in the human liver, whereas CES2 is the major carboxylesterase expressed in the intestine (Imai et al., 2006). Hepatic CES1 is the major esterase governing the metabolism of numerous and structurally diverse therapeutic agents formulated as carboxylic acid esters, carbamates, thiosteres, and amide compounds including those prodrugs formulated as esters. In addition, a number of endogenous substrates are recognized.

In a recent study, we identified two CES1 mutations, p.Gly143Glu and p.Asp260fs (Zhu et al., 2008), in a subject who displayed profound alteration of the pharmacokinetics of racemic (dl)-methylphenidate (Ritalin; Novartis Pharmaceuticals, Summit, NJ), a selective CES1 substrate, during a single-dose pharmacokinetic study (Patrick et al., 2007). The minor allele frequency of p.Gly143Glu was estimated to be 3.7, 4.3, 2.0, and 0% in white, black, Hispanic, and Asian populations, respectively, by a genotyping study that contains a total of 925 subjects with varied racial and ethnic backgrounds. It was concluded that the p.Asp260fs variant was extremely rare because none of the 925 screened subjects carried this mutation. The functional consequences of both mutations were investigated using cell lines stably expressing each individual mutant. The in vitro incubation study demonstrated that the catalytic function of both p.Gly143Glu and p.Asp260fs is impaired to such a significant degree that CES1-mediated methylphenidate hydrolysis was essentially nil using these two CES1 mutants, whereas wild-type (WT) CES1 readily cleaved the ester (Zhu et al., 2008).

Even though the two newly discovered CES1 mutations were determined to be dysfunctional enzymes in terms of hydrolyzing methylphenidate to its inactive metabolite ritalinic acid, the influence of these CES1 variants on prodrug activation has not been examined to date. Oseltamivir (a drug in wide therapeutic use) has recently been shown to be a selective

ABBREVIATIONS: CES1, carboxylesterase 1; CES2, carboxylesterase 2; WT, wild type; PNPA, p-nitrophenyl acetate; PNP, p-nitrophenol; HPLC, high-performance liquid chromatography.
The oseltamivir phosphate incubation study demonstrated that the S9 preparations of both WT CES1-transfected cells and human liver tissues efficiently convert oseltamivir to its active antiviral component, oseltamivir carboxylate, suggesting that oseltamivir serves as an excellent substrate of CES1 (Fig. 3). The $V_{\text{max}}$ and $K_m$ values were determined to be 145 ± 5 nmol/min/mg protein and 1.38 ± 0.13 mM, respectively, under our experimental conditions (Fig. 4). The CES1 variants p.Gly143Glu and p.Asp260fs displayed poor catalytic activity toward oseltamivir hydrolysis (Figs. 3 and 4). The $V_{\text{max}}$ value of p.Gly143Glu was found to be 37 ± 1 nmol/min/mg protein, which is approximately 25% of that of WT CES1. The $K_m$ value of p.Gly143Glu was estimated to be 2.15 ± 0.18 mM. In addition, p.Asp260fs failed to produce any detectable hydrolysis of oseltamivir as measured by the formation of oseltamivir carboxylate (Fig. 3). The human liver S9 fractions prepared from a healthy donor specimen produced similar catalytic activity toward both PNPA and oseltamivir phosphate, which was in excellent agreement with that of our WT CES1-transfected cells (Fig. 3).

Discussion

CES1 is the predominant hydrolase in the liver and plays an important role in the biotransformation of drugs and prodrugs that contain ester bonds. CES1 genetic variants and their potential for having therapeutic implications have been increasingly reported recently. Our previous study identified two nonsynonymous coding region variants, p.Gly143Glu and p.Asp260fs. In vitro functional studies have shown that the catalytic function mediating the typically efficient and rapid hydrolysis of methylphenidate was clearly disrupted in both the p.Gly143Glu variant and the p.Asp260fs mutation. The potential for clinically significant outcomes in the presence of these two mutations was investigated in the original subject found to carry both CES1 variants. That subject displayed an extremely abnormal pharmacokinetic profile after the administration of methylphenidate, displaying vastly higher overall blood concentrations of methylphenidate and an unprecedented distortion in the disposition of the respective isomers of the drug (Patrick et al., 2007; Zhu et al., 2008).
In addition, the subject experienced significantly higher cardiovascular vital signs relative to other 19 study subjects serving as a pharmacodynamic correlate to the pharmacokinetic observations (Zhu et al., 2008).

Because the activation of many ester prodrugs depends to a great degree upon functional CES1 enzyme to produce the therapeutic moiety, dysfunctional CES1 variants could hinder prodrug activation and lead to the alteration of therapeutic effects and accumulation of the parent prodrug with continued dosing. Such an outcome could lead to therapeutic failure and, depending on the compound administered, unanticipated adverse effects or toxicities. As a prodrug, oseltamivir does not exhibit activity toward the influenza virus unless it is converted to its active metabolite oseltamivir carboxylate by CES1 (Fig. 1). In the present study, the catalytic activity of p.Gly143Glu and p.Asp260fs toward oseltamivir hydrolysis (i.e., activation) was investigated using transfected cell lines stably expressing WT and individual mutant CES1 enzyme. The data indicated that the enzymatic activity of p.Gly143Glu is substantially decreased with a $V_{\text{max}}$ value approximately one fourth that of WT CES1, whereas p.Asp260fs failed to show any measurable hydrolytic activity toward oseltamivir. Acknowledging the limitations of in vitro methodologies, this fundamental alteration in the catalytic activity of CES1 strongly suggests that the activation of oseltamivir would be compromised in patients who express such CES1 variants. In addition to these two mutations, several other natural nonconservative CES1 variants were recently determined to also have functional significance (Shi et al., 2006; Tang et al., 2006). Furthermore, beyond the coding area mutations, a number of functional variants have been reported in the transcriptional regulation region of CES1 gene (Geshi et al., 2005; Hosokawa et al., 2008; Yoshimura et al., 2008). Among those, a single nucleotide
polymorphism. −816AC of the CES1A2 gene was found to be associated with an improved therapeutic response to an angiotensin-converting enzyme inhibitor imidapril, which is a produg and selectively activated by CES1 (Geshi et al., 2005).

It was noted that the observed $V_{\text{max}}$ value of WT CES1 is consistent with that reported by Shi et al. (2006), whereas the $K_m$ value is seven times higher. We suspect this difference is more likely than not the result of different experimental conditions used in these two independent studies. For example, the reaction buffer used in the present study was phosphate-buffered saline containing 10 mM HEPES (pH 7.4), whereas a Tris buffer was used in the study by Shi et al. (2006). Indeed, a recent in vitro study addressing this very issue indicates that different enzymatic activity of CES could be observed when different assay buffers were used (Williams et al., 2008). Finally, the S9 fractions used in the present study were prepared from a stable CES1 cell line rather than a transient expression assay.

Functional CES1 is not only critical for the conversion of oseltamivir to its active metabolite to achieve a favorable therapeutic response, but it is also related to the toxicity during oseltamivir therapy. Converging evidence suggests that CES1 function in juvenile animals remain at a significantly lower level than that of adult animals (Kadner et al., 1992; Morgan et al., 1994; Moser et al., 1998; Karanth and Pope, 2000; Padilla et al., 2004; Anand et al., 2006). Animal studies demonstrated that juvenile rats did not hydrolyze oseltamivir efficiently, and they are more susceptible than adults to oseltamivir toxicity (http://www.fda.gov/cder/foi/nda2000/21–246_TamiFlu_Phrarm.pdf). The present study suggests that, in addition to age, genetic variation is potentially an important factor influencing the enzymatic function of CES1 and could play a role in the therapeutic outcome and toxicity of pharmacotherapy with oseltamivir as well as other known CES1 substrates. Our previously published data with the psychostimulant methylphenidate indicate that the effects of p.Gly143Glu mutation on the pharmacological disposition and potential toxicities of oseltamivir.

In summary, two newly identified CES1 mutations p.Gly143Glu and p.Asp260fs were determined to be dysfunctional enzymes with respect to the activation of the prodrug oseltamivir. Impaired enzymatic function could have significant implications with regard to both the therapeutic efficacy and tolerability of oseltamivir. It should be noted that the extremely low prevalence of p.Asp260fs mutation relegates its clinical significance to being very minor even though it results in a nonfunctional enzyme. However, p.Gly143Glu is a common variant in all populations assessed thus far, with the exception of Asians. A clinical study, particularly one assessing patients who have been genotyped and found to be heterozygously expressing p.Gly143Glu, is warranted to elucidate the influence of the p.Gly143Glu mutation on the pharmacological disposition and potential toxicities of oseltamivir.

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References


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