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Pharmacokinetics of the potent redox modulating manganese porphyrin, MnTE-2-PyP⁵⁺ in plasma and major organs of B6C3F1 mice

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Abstract

Mn(III)tetrakis(*N*-ethylpyridinium-2-yl)porphyrin, MnTE-2-PyP⁵⁺, a potent catalytic superoxide and peroxynitrite scavenger, has been beneficial in several oxidative stress-related disease thus far examined. Pharmacokinetic studies are essential for the better assessment of the therapeutic potential of MnTE-2-PyP⁵⁺ and similar compounds, as well as for the modulation of their bioavailability and toxicity. Despite high hydrophilicity, this drug entered mitochondria after single 10 mg/kg intraperitoneal injection at levels high enough (5.1 μM; 2.95 ng/mg protein) to protect it against superoxide/peroxynitrite damage. Utilizing the same analytical approach, which involves the reduction of MnTE-2-PyP⁵⁺, followed by the exchange of Mn²⁺ with Zn²⁺, and HPLC/fluorescence detection of ZnTE-2-PyP⁴⁺, we measured levels of MnTE-2-PyP⁵⁺ in mouse plasma, liver, kidney, lung, heart, spleen, and brain over a period of 7 days after a single intraperitoneal injection of 10 mg/kg. Two B6C3F1 female mice per time point were used. The pharmacokinetic profile in plasma and organs was complex; thus a non-compartmental approach was utilized to calculate the area under the curve (AUC), c_{max}, t_{max}, and drug elimination half-time (t_{1/2}). In terms of levels of MnTE-2-PyP⁵⁺ found, the organs can be classified into 3 distinct groups: (1) high levels: kidney, liver, and spleen; (2) moderate levels: lung and heart; and (3) low levels: brain. The maximal levels in plasma, kidney, spleen, lung, and heart are reached within first 45 minutes whereas in case of liver a prolonged absorption phase was observed with the maximal concentration reached at 8 hours. Moreover, accumulation of the drug in brain continues beyond time of the experiment (7 days) and is likely driven by the presence of negatively charged phospholipids. For tissues other than brain, a slow elimination phase (single exponential decay, t_{1/2} = 60 to 135 hours) is observed. The calculated pharmacokinetic parameters will be used to design optimal dosing regimens in future preclinical studies utilizing this and similar compounds.

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Keywords

MnTE-2-PyP⁵⁺; AEOL 10113; SOD mimic; pharmacokinetics in mouse plasma; pharmacokinetics in mouse organs; pharmacokinetics in mouse tissues

Introduction

For more than a decade we have been investigating metalloporphyrin-based antioxidants, studying their *in vitro* efficacy in eliminating reactive species and *in vivo* efficacy in oxidative stress-related models of diseases [1–8]. From our studies on structure-activity relationships between the redox potential of metalloporphyrins and their ability to dismutate superoxide, O₂^{•-}, MnTE-2-PyP⁵⁺ emerged as a standard potent catalytic antioxidant now used a wide range of cellular and animal studies. The compound has a high efficacy to *dismute* (*disproportionate*) O₂^{•-} [1,2] (log k_{cat} = 7.76 as compared to log k_{cat} = 8.84 - 9.3 for SOD enzymes [9–14]). Moreover, its efficacy to dismutate O₂^{•-} parallels its efficacy to *reduce* peroxynitrite, ONOO⁻, (log k_{red} = 7.53) [15,16]. MnTE-2-PyP⁵⁺ is also very effective in scavenging carbonate radical, CO₃^{•-} [16]. The reduction of peroxynitrite or CO₃^{•-} is catalytic in nature *in vivo*, as it is coupled to cellular reductants; the overall reaction is more efficient with ascorbate, and urate than with potential targets such as thiols and amino acids [15–17]. The ability to eliminate O₂^{•-}, ONOO⁻, and CO₃^{•-} exceeds that of other compounds presently in preclinical and clinical studies such as Mn salen derivatives [18,19], Mn cyclic polyamines [20], nitrones [21,22], nitroxides [23], MitoQ [24] and analogous compounds [25]. MnTE-2-PyP⁵⁺ ameliorates oxidative stress-related diseases, presumably through elimination of reactive species, thus providing the protection of most biologically relevant targets against *primary oxidative stress*. Moreover, the elimination of reactive species prevents activation of transcription factors AP-1 [26,27], HIF-1α [28–31], and NF-κB [32,33] which in turn inhibits the expression of cytokines and/or enzymes that would otherwise perpetuate *secondary oxidative stress*. The catalytic power of MnTE-2-PyP⁵⁺ is enhanced by the presence of electron-withdrawing positively charged *ortho* alkylpyridyl groups close to the metal center. These proximal positive charges contribute to both *thermodynamic* (favorable redox potential) [1,6] and *kinetic* (electrostatic attraction of the negatively charged substrate) [6,34] facilitation of O₂^{•-} dismutation, and ONOO⁻ and CO₃^{•-} reduction. Several other metalloporphyrins [35–38] and corroles [39] have been developed that exploit the effects of cationic *ortho* quaternary nitrogens.

The beneficial effects of Mn alkylpyridylporphyrins have been observed in central nervous system injuries (stroke [40–42], spinal cord injury [43], ALS [44], Alzheimer's disease [45]), radiation injury [46–49], cancer [26,29], diabetes [50,51], sickle cell disease [52], ischemia/reperfusion conditions [53–55], arthritis [56] and others. However, very little is known about the bioavailability, site of action and pharmacokinetics and toxicology of the compound. It has been assumed in past that excessive hydrophilicity would limit the accumulation of MnTE-2-PyP⁵⁺ within the cell and its organelles, suggesting the *extracellular* localization of porphyrin as a likely site of action. However, the high efficacy observed *in vivo* strongly implicated not only *cytosol*, but also *mitochondrion* [26,27] and *nucleus* [32,33] as possible sites of action. In order to test the validity of such conclusions we have developed a sensitive HPLC/fluorescence method for determination of the positively charged Mn(III) *ortho N*-alkylpyridylporphyrins *in vivo*, and used it in a mouse mitochondrial study [57]. Herein, utilizing similar analytical approach, we undertook a pharmacokinetic study to detect levels of porphyrin in mouse plasma and different organs over the period of 7 days after single 10 mg/kg intraperitoneal injection of MnTE-2-PyP⁵⁺. This is the first detailed pharmacokinetic study on such an important class of Mn-porphyrin-based antioxidant therapeutics.

Experimental

General

Sodium L-ascorbate, mannitol, albumin from bovine serum (BSA), and ZnCl₂ were from Sigma and Zn(CH₃COO)₂ × 2 H₂O from J. T. Baker. Acetonitrile and trifluoroacetic acid (TFA) were from Fisher Scientific and triethylamine from Pierce. Methanol (anhydrous, absolute) was from Mallinckrodt. Glacial acetic acid was from EM Science. Phosphate-buffered saline (50 mM sodium phosphate, 0.9% NaCl, pH 7.4) (PBS) was from Gibco.

Porphyrins

MnTE-2-PyP⁵⁺ ($\lambda_{\max} = 454$ nm, $\log \epsilon = 5.14$) [1,2], and ZnTE-2-PyP⁴⁺ ($\lambda_{\max} = 425.5$ nm, $\log \epsilon = 5.46$) [57–59] were prepared as described previously. ZnTnBu-2-PyP⁴⁺ was used as an internal standard and was prepared and characterized as described [57].

Uv/vis spectroscopy

Uv/vis was done on a Shimadzu 2501 PC UV/Vis spectrophotometer.

Mice

The University of Kentucky Medical Center Research Animal Facility has a continuously accredited program from AAALAC International. All experiments using animals were performed according to the approved protocol for humane care and use of animals. Two B6C3F1/J female mice were used for each time point. (This strain is commonly used as a background strain for transgenic MnSOD mice used in studying oxidative stress injuries.) The mice weighed from 20 to 25 g and the volumes of porphyrin saline solutions ranged accordingly from 200 to 250 μ L. The mice were euthanized with high-dose pentobarbital. Before harvesting tissue, the animals were perfused with saline to avoid artifacts related to the retention of blood in the organs and tissue. All mice perfused had their blood drawn from the left ventricle. The perfusion was by gravity flow of saline via a blunt needle placed into the aorta through left ventricle and the saline flow through right atrium was continued until the liver cleared out.

Protein levels

The protein levels were determined in plasma and tissue homogenates by colorimetric Bradford assay as specified by manufacturer (Bio-Rad, Richmond CA). Protein levels were used to normalize tissue levels in homogenates.

MnTE-2-PyP⁵⁺ plasma/tissue extraction

Organs were cryo-pulverized in a Bessman tissue pulverizer (BioSpec Products, Bartlesville, OK) under liquid nitrogen and then homogenized in rotary homogenizer (PTFE pestle and glass tube) with 2 volumes of deionized water. 100 μ L of either plasma or tissue homogenate was transferred into a 2 mL polypropylene screw-cap vial and 200 μ L of deionized water and 300 μ L of 1% acetic acid in methanol were added. The content was mixed in a multi-vortexer for 60 sec and the homogenate was centrifuged 5 min at 16,000 g. 400 μ L of the supernatant was pipetted into a 5 mL polypropylene tube (10 × 50 mm) and the solvent was completely removed in a Savant Speed-Vac evaporator at 40 °C within 1 h. The dry residue was dissolved in 100 μ L of deionized water containing 20 mM triethylamine, the pH of which was adjusted with diluted TFA to pH 6.2, followed by 60 sec vortexing and centrifugation for 2 min at 2000 g. 80 μ L of the supernatant was transferred into a 2 mL polypropylene screw-cap tube, and 30 μ L of 1.0 M either Zn acetate or Zn chloride in water and 10 μ L of freshly prepared 0.11 M sodium ascorbate in water were added. The solution was left for one hour at either 50 °C or 70 °C. (If working at 70 °C the more acidic salt Zn chloride ought to be used due to a lower degree

of hydrolysis than that of Zn acetate). 20 μL of 2.7 M TFA and 20 μL of 800 nM ZnTnBu-2-PyP⁴⁺ were added, followed by vortexing for 60 sec and centrifugation for 5 min at 16,000 g. 110 μL of the supernatant was transferred to the HPLC autosampler polypropylene vial equipped with silicone/PTFE septum.

HPLC

Equipment: Waters 2695 HPLC system (pump, autosampler, column oven) and Waters, model 2475 fluorescence detector set on Gain 100, $\lambda_{\text{exc}} = 425 \text{ nm}$ and $\lambda_{\text{abs}} = 656 \text{ nm}$ [60–62]. Column: YMC-Pack, ODS-AM, C18 column (3 μm particle size, 120 \AA pore size, 150 \times 4.6 mm) at 45 $^{\circ}\text{C}$. Solvent A: 95% aqueous (deionized water, 20 mM triethylamine, pH 2.7 adjusted with concentrated TFA) and 5% acetonitrile. Solvent B: acetonitrile. Elution gradient at 1.5 mL/min: 0-15-20 min, 100% A - 80% A - 100% A, followed by 5 min column conditioning. Autosampler temperature: 4 $^{\circ}\text{C}$. Injection volume: 80 μL .

The *ortho* isomers of MnTE-2-PyP⁵⁺, ZnTE-2-PyP⁴⁺ and their parent metal-free ligand, H₂TE-2-PyP⁴⁺ have atropoisomers, which we previously characterized by HPLC/uv/vis/NMR and X-ray methods [59]. We observed that the abundance ratio of the four isomers of ZnTE-2-PyP⁴⁺ and their retention times on HPLC were the same in mouse heart mitochondrial homogenates [57], in processed mouse organ homogenates or in aqueous solution [58]. This indicated that the interactions of any of the four atropoisomers with cellular components [57, 58] are not strong enough to overcome the rotational barrier of the alkylpyridyl “arms” which would cause a change in the abundance ratio of atropoisomers. The experiments also implied that the Mn porphyrin does not undergo oxidative degradation or any other modification *in vivo*. Using any of the atropoisomers for calculation of the levels of MnTE-2-PyP⁵⁺ in mitochondria gave the same result. ZnTnBu-2-PyP⁴⁺ was used as an internal standard, and its atropoisomers were also present in the chromatogram at the expected abundance ratio [57, 58]. The assignment of the individual atropoisomers, $\alpha\alpha\alpha\alpha$, $\alpha\alpha\alpha\beta$, $\alpha\alpha\beta\beta$ and $\alpha\beta\alpha\beta$, of both Zn porphyrins was done based on our previous HPLC/¹H NMR/X-ray study [57,58]. In this study, the most abundant atropoisomer, $\alpha\alpha\alpha\beta$, was used for the quantification of both ZnTE-2-PyP⁵⁺ and ZnTnBu-2-PyP⁴⁺ (internal standard).

Calibration curves

A set of serially diluted standard samples of MnTE-2-PyP⁵⁺ were prepared for calibration. The range varied from 1.5 ng to 100 μg per mL of drug-free plasma or per g of drug-free wet tissue, and was dependent upon the type of the organ studied. MnTE-2-PyP⁵⁺ was added into the tissue homogenate or plasma, followed by 15-min incubation at room temperature. The rest of the assay procedure was identical to the procedure performed with organs and plasma of drug-treated animals. Response was calculated as the ratio between standard peak area/internal standard peak area.

Recovery of MnTE-2-PyP⁵⁺, i.e. the overall efficacy of extraction from plasma and organs plus Mn-to-Zn exchange, was determined in triplicates in the following way. A known amount of MnTE-2-PyP⁵⁺ was added to plasma or organ homogenate, and the procedure of extraction/metal exchange was followed. The response from this experiment was compared to the response of the sample where corresponding amount of ZnTE-2-PyP⁴⁺ (equal to 100% yield of MnTE-2-PyP⁵⁺) was diluted into mobile phase. The obtained recovery values are 83% in plasma, 89% in liver, 88% in kidney, 83% in spleen, 70% in lung, 85% in heart, 97% in brain.

Results and Discussion

MnTE-2-PyP⁵⁺ is very effective in eliminating reactive species (RS), particularly O₂⁻ and ONOO⁻, with rate constants that are among the highest when compared to other synthetic

antioxidants [1–8,15–17]. This allows the compound not only to decrease the *primary insult* of RS to biological molecules, but also to inhibit activation of transcription factors which in turn leads to suppression of expression of those cytokines and enzymes that perpetuate *secondary oxidative stress* [26–33]. Thus the MnTE-2-PyP⁵⁺ has a potential to be an efficacious therapeutic in all diseases that have oxidative stress as a common factor [40–56]. Due to its high hydrophilicity and positive charge, crossing the blood brain barrier and the mitochondrial envelope has always been a concern. Thus, we were surprised to find 5.1 μM (2.95 ng/mg protein) MnTE-2-PyP⁵⁺ in mouse heart mitochondria 4 (or 7) hours after a single 10 mg/kg intraperitoneal injection; as a note, 10 mg/kg is the most commonly used (as single or multiple) therapeutic dose of this porphyrin [57]. A previous study by Ferrer-Sueta *et al* [17] on submitochondrial particles indicate that at ≥3 μM MnTE-2-PyP⁵⁺ nearly fully eliminates ONOO⁻ mediated damage to the components of mitochondrial respiration machinery. The dose of 10 mg/kg was picked up based upon the rodent studies on the diabetes [50], cancer [28,29,63], and radioprotection [48,64] that have reported beneficial effects (delayed onset of diabetes, tumor growth delay, radioprotection) when MnTE-2-PyP⁵⁺ was administered at doses ranging from 6 to 15 mg/kg. The TD₅₀ was found to be 91.5 mg when MnTE-2-PyP⁵⁺ was administered subcutaneously.^a

We have reported that Mn porphyrins can mimic cyt P450, yet to a lower extent than Fe analogous porphyrins [65]. It was also shown that Mn porphyrin can redox cycle with cyt P450 reductase which action was claimed to have possible beneficial and/or adverse effects [66]. Thus far our studies did not indicate any metabolism of MnTE-2-PyP⁵⁺ and similar cationic porphyrins by enzymatic systems *in vivo*.

Herein, we studied the pharmacokinetics (PK) of MnTE-2-PyP⁵⁺ in mouse plasma and several important organs. Animal perfusion to remove blood from organs, and measurement of porphyrin levels were performed as in mitochondrial study [57] but with some necessary modifications primarily in sample preparation procedures. The HPLC/fluorescence method is based on the reduction of MnTE-2-PyP⁵⁺ by ascorbic acid, followed by *in situ* Mn²⁺ to Zn²⁺ exchange and fluorescence detection of ZnTE-2-PyP⁴⁺. Assay conditions are given in Scheme I. ZnTnBu-2-PyP⁴⁺ was used as an internal standard. Calibration curves, made with drug-free plasma and corresponding organ homogenates, were linear in the ranges needed for quantification. Lower limits of quantification (LLQ) in ng/mL (plasma) or ng/g wet tissue (organs), with corresponding sample size in brackets, were as follows: plasma 4.64, (100 μL), liver 11.9 (33 mg), spleen 72.5 (33 mg), kidney 215 (33 mg), lung 35.8 (33 mg), heart 35.8 (33 mg), and brain 3.6 (100 mg). Accuracy (relative error) acceptance criteria at each nominal concentration were <15% and <20% at LLQ [67]. The concentration in day 7 plasma samples was barely detectable and outside the standard curve range (below LLQ). Nonetheless, in the course of manuscript preparation, a new LC/MS/MS method for MnTE-2-PyP⁵⁺ in plasma has been developed in our lab with LLQ = 0.5 ng/mL using 250 μL sample. Thus, the pooled day 7 plasma sample (2 animals) was measured by LC/MS/MS and the method will be described in a subsequent publication.

Pharmacokinetic profiles are shown in Figure 1A–G along with corresponding calibration curves. The levels of porphyrin are given in μg/g wet tissue or μg/mL plasma. For each time-point the two concentrations measured, corresponding to two animals, are presented as open circles whereas the average of these values is given by the solid bold line. Average concentrations at each time point were used in the pharmacokinetic calculations. Semi-log plots are chosen to illustrate the level of complexity in the PK behavior. Namely, a single-exponential rise/decay in concentration would indicate drug absorption/distribution from/into one or more “compartments” of very similar blood perfusion and/or drug affinity properties. Such single-

^aPanni, M, The TD₅₀ of MnTE-2-PyP⁵⁺ given subcutaneously.

exponential process would appear linear in a semi-log plot. Thus, as it can be seen in Figure 1A (far left), plasma pharmacokinetics over 7 days appears to be very complex—several linear regions (compartments) can be recognized. This is in agreement with the pharmacokinetics seen in different organs (Figures 1B–G). For this reason, and because of limited number of time points, no attempt was made to fit a multi-compartmental model into the data. Instead, the non-compartmental approach (WinNonlin, Pharsight Inc.) was used: numerical integration of the PK profile is performed to calculate the area under the curve (AUC), and the terminal slope is used to calculate drug half-time ($t_{1/2}$) for each organ and plasma; c_{\max} is the highest concentration found along the PK profile, and t_{\max} is the corresponding time (Table 1). In terms of levels of MnTE-2-PyP⁵⁺ found, the organs can be classified into 3 distinct groups: (1) high levels: kidney, liver, and spleen; (2) moderate levels: lung and heart; and (3) low levels: brain. The plots clearly reveal differences in the drug penetration (diffusion or active transport), and drug-tissue affinity (binding or retention) among organs. During the first 15 minutes, the drug is rapidly absorbed from the intraperitoneal cavity into the plasma. The levels in kidney, spleen, lung, and heart also increase rapidly, presumably due to a passive diffusion - the drug levels in kidney reach especially high values, obviously due to both high penetration and a high affinity. At the same time, in liver, the drug shows a slow penetration ($t_{\max} = 8$ hrs) but reaching very high levels due to high affinity. The drug also exhibits a very slow penetration into brain. Due to different affinity (weak versus strong) for the different tissues, an intensive distribution/re-distribution of the drug in the body is going on for 2 days after which a slow drug elimination from all organs (except brain) takes place at a similar rate. The MnTE-2-PyP⁵⁺ levels in brain, however, start *increasing* after 2 days and continue the trend past 7 days meaning that the drug binds very strongly to the brain tissue; it is noteworthy that brain continues to accumulate MnTE-2-PyP⁵⁺ despite very low and ever decreasing concentration of the drug in circulating blood (plasma) (Figure 1G). This accumulation of the drug in the brain at late times is maintained by the high concentration of MnTE-2-PyP⁵⁺ in other tissues (liver, kidney, and spleen) from where it is supplied over days and weeks via re-distribution.

We have already found that positively charged isomeric Mn(III) methylpyridylporphyrins (MnTM-2(3,4)-PyP⁵⁺) associate with the nucleic acids of *E. coli* [1]. Its cationic nature further drives the accumulation of MnTE-2-PyP⁵⁺ in mitochondria [57]. Finally, a study in collaboration with Tse and Piganelli,^b showed that MnTE-2-PyP⁵⁺ accumulates 3-fold more in the nucleus than in the cytosol of macrophages, presumably driven by the negatively charged phosphate groups of the nucleic acids. Similarly, the accumulation of MnTE-2-PyP⁵⁺ in brain is likely driven by the abundance of negatively charged phospholipids in this organ, and may explain the beneficial effects of MnTE-2-PyP⁵⁺ in ALS [44], stroke models [41,42] and pain management [68]. Further it is well-documented that overexpression of MnSOD can suppress cancer phenotypes and the metastatic potential of cancer cells; all suggesting a high therapeutic potential of MnTE-2-PyP⁵⁺ for brain radioprotection and brain tumor treatment.

The ability of MnTE-2-PyP⁵⁺ to cross blood brain barrier has been indicated in an independent study where nearly full reversal of morphine tolerance was achieved by MnTE-2-PyP⁵⁺ therapy; moreover, identical effects were observed whether MnTE-2-PyP⁵⁺ was administered intraperitoneally or intrathecally [63,68]. Further, when *para* methyl analogue, MnTM-4-PyP⁵⁺ was used in a rabbit model of cerebral palsy its presence in fetal brain was observed by MRI.^c More thorough study is in progress to evaluate whether brain levels achieved by the intraperitoneal or subcutaneous routes are close to the therapeutic levels in different animal models of central nervous system injuries.

^bSpasojević, I.; Batinić-Haberle, I.; Tse, H.; Piganelli, J. Accumulation of MnTE-2-PyP⁵⁺ in nucleus, unpublished

^cSidhartha, T. MnTM-4-PyP⁵⁺ in cerebral palsy, unpublished.

The levels of MnTE-2-PyP⁵⁺ $\geq 3 \mu\text{M}$ in submitochondrial particles shown to be protective by Ferrer-Sueta et al [17], coincide with levels at which the aerobic growth SOD-deficient *E. coli* achieves $\geq 60\%$ of the growth of the wild type [1,69]. Based on Vinnakota and Bassingthwaite data on the density of myocardial tissue to be 1.053 g/mL [70], and assuming that it is valid for other tissues as well, the 1 $\mu\text{g/g}$ wet tissue is $\sim 1 \mu\text{M}$ MnTE-2-PyP⁵⁺ ($M_r = 965.13$). Consequently, after single intraperitoneal injection, therapeutic doses of MnTE-2-PyP⁵⁺ might have been achieved in liver, kidney and spleen. This is in agreement with Saba et al data [54] where kidney ischemia/reperfusion injury was significantly diminished by a single intravenous injection of hexyl analogue, MnTnHex-2-PyP⁵⁺. More studies are needed to determine the magnitude and the frequency of administrations needed to achieve therapeutic doses in other organs.

Conclusions

We report here the first detailed pharmacokinetic study on an important class of catalytic Mn-porphyrin-based antioxidant therapeutics. The findings presented are relevant for the proper dosing of MnTE-2-PyP⁵⁺ in future *in vivo* studies. The PK analysis shows a very efficient absorption of the drug from intraperitoneal space into blood stream (within 15 min) and a very slow elimination of the drug from the body ($t_{1/2} = 55\text{--}135$ hrs). After single dose, near-therapeutic levels were achieved in liver, kidneys, and spleen. The observed high accumulation and slow release from most tissues suggests no need for frequent dosing. An initial higher loading dose may be followed by a small periodic “maintenance” dose. However, in attempts to deliver a higher dose to organs like lung, heart and brain, one must bear in mind that very high and possibly toxic levels may be reached in liver, kidneys, and spleen.

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Abbreviations

SOD, superoxide dismutase
MnP, Mn^{III}TM-2(3,4)-PyP⁵⁺, Mn(III) *meso*-tetrakis(*N*-methylpyridinium-2(3,4)-yl) porphyrin; 2, 3, and 4 being *ortho*, *meta* and *para* isomer, respectively
Mn^{III}TE-2-PyP⁵⁺, Mn(III) *meso*-tetrakis(*N*-ethylpyridinium-2-yl)porphyrin (AEOL-10113)
Zn^{II}TE-2-PyP⁴⁺, Mn(III) *meso*-tetrakis(*N*-ethylpyridinium-2-yl)porphyrin
MnTnHex-2-PyP⁵⁺, Zn *meso*-tetrakis(*N*-n-hexylpyridinium-2-yl)porphyrin
Zn^{II}TnBu-2-PyP⁴⁺, Zn *meso*-tetrakis(*N*-n-butylpyridinium-2-yl)porphyrin
H₂Talkyl-2-PyP⁴⁺, *meso*-tetrakis(*N*-alkylpyridinium-2-yl)porphyrin *meso* is indicating substitution in 5,10,15,20 positions on porphyrin core
HIF-1, hypoxia inducible factor-1, NF- κ B, nuclear factor κ B
AP-1, activator protein-1
BSA, bovine serum albumin
TFA, trifluoroacetic acid
PBS, phosphate-buffered saline

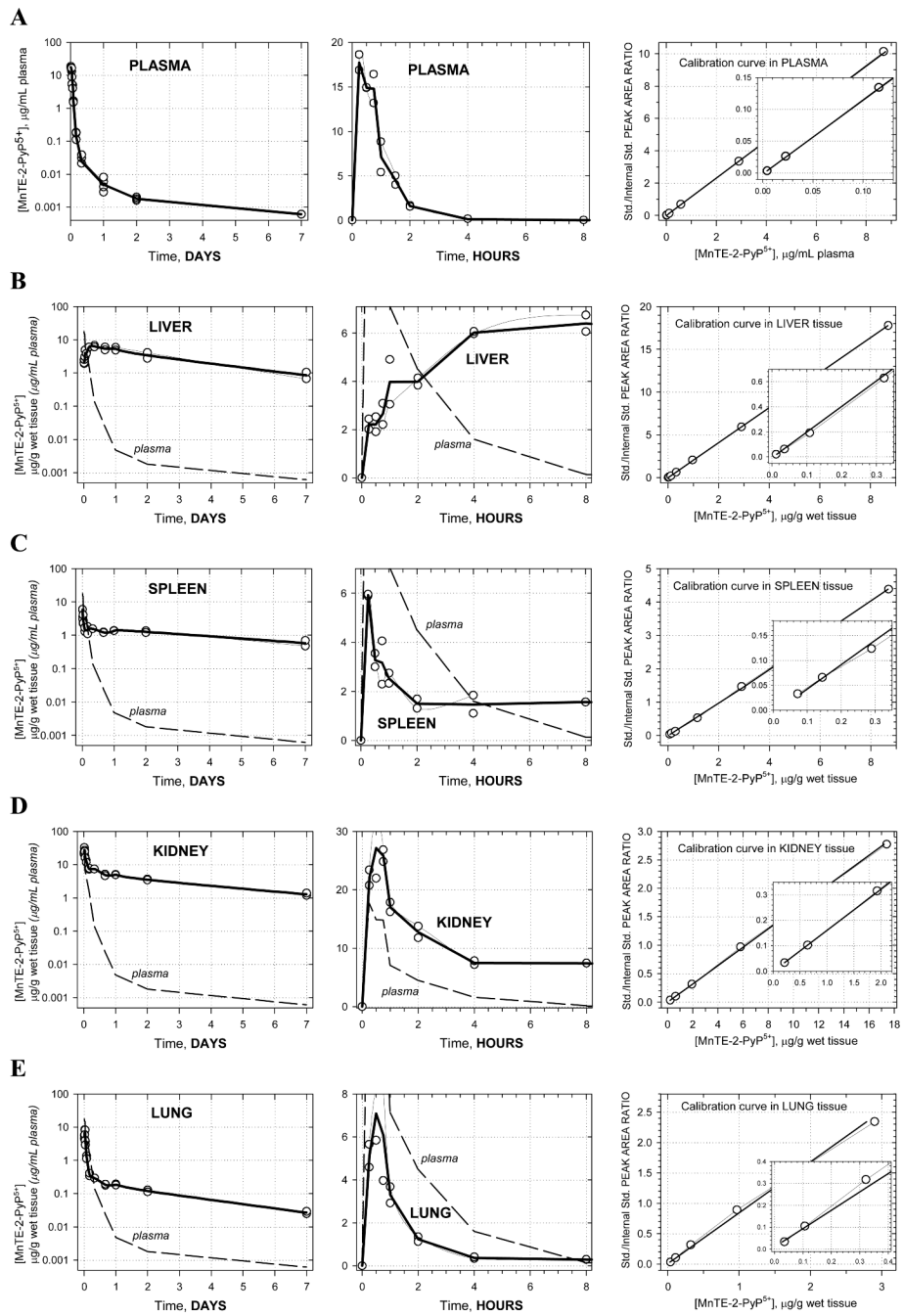
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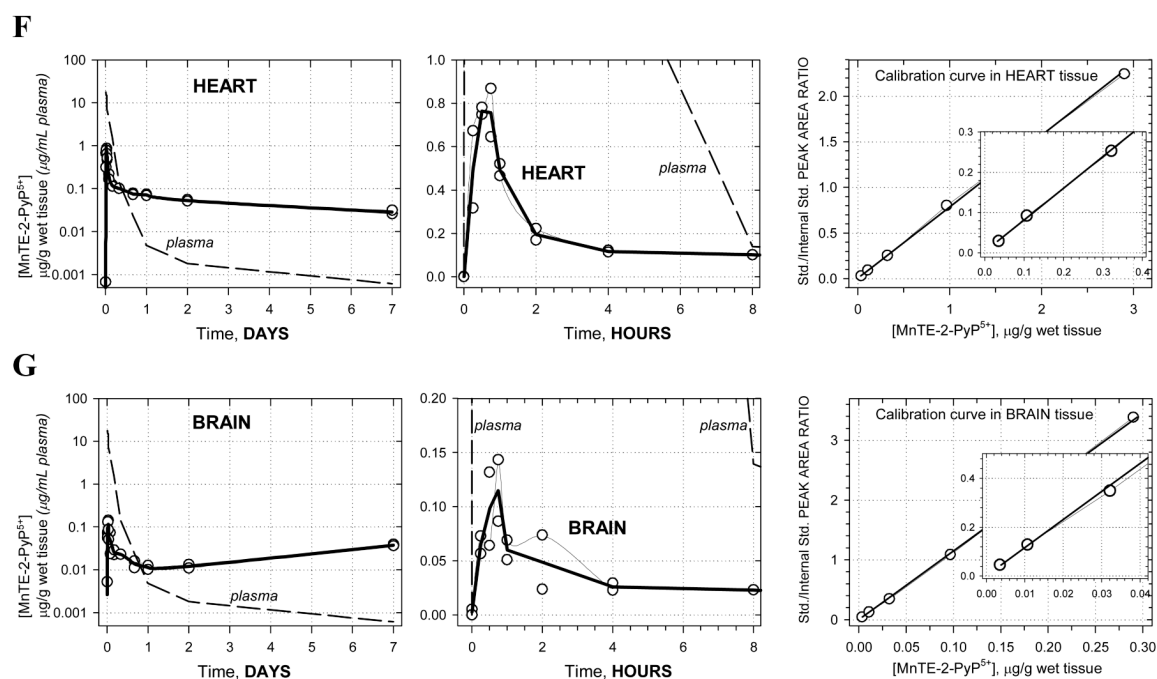
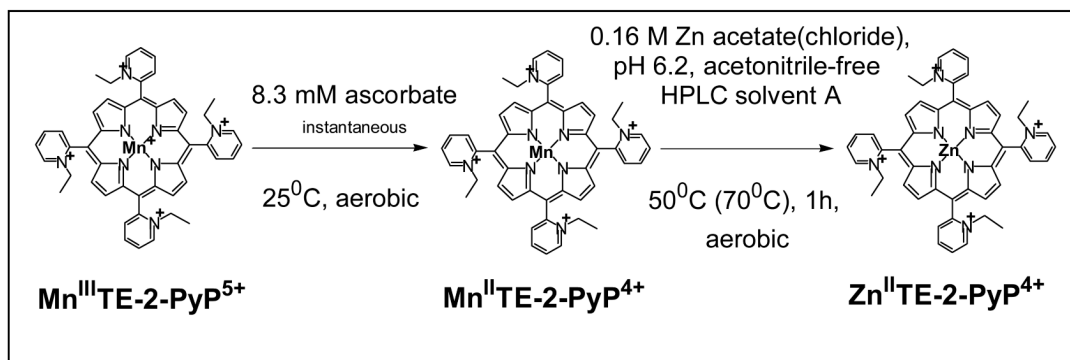


Figure 1. Pharmacokinetics of MnTE-2-PyP⁵⁺ in plasma (A), liver (B), spleen (C), kidney (D), lung (E), heart (F) and brain (G). Levels of porphyrin are given in µg per mL of plasma or per g of wet tissue. Drug was given at a single intraperitoneal dose of 10 mg/kg to female B6C3F1 mice. Animals were anesthetized, perfused and organs excised. 2 mice were used per data point. Calibration curves were done in plasma and related organs. The known amounts of MnTE-2-PyP⁵⁺ were added into the plasma and organ homogenates; the rest of the procedure was identical to the one used for drug-treated mice.

**Scheme I.**

The representation of the method used to determine levels of MnTE-2-PyP⁵⁺ in plasma and organs. After the proteins were precipitated from plasma and tissue homogenates, the Zn salt was added along with sodium ascorbate into the supernatant to exchange Mn(II) by Zn(II). While the reduction of the Mn(III)P occurred instantaneously, one hour of incubation at 50 or 70 °C is needed to yield a complete Mn by Zn exchange for the HPLC/fluorescence measurement of ZnTE-2-PyP⁴⁺.

Pharmacokinetic parameters of MnTE-2-PyP⁵⁺ in plasma, liver, kidney, spleen, lung, heart and brain: t_{max} (time point where maximum concentration, C_{max} was achieved), C_{max} (maximum observed concentration), $t_{1/2}$ (half life of MnTE-2-PyP⁵⁺ elimination obtained from terminal slope between days 2 and 7), AUC_{last} (area under the curve up to the last measurement, i.e. 7 days) and AUC_{inf} (projected total area under the curve at infinite time point) that indicates the exposure of the organ to the drug.

Table 1

organ	t_{max} , h	C_{max} , µg/g	$t_{1/2}$, h	AUC_{last} , µg · h/mL(g)	AUC_{inf} , µg · h/mL(g)
plasma	0.25	17.74	76.9	18.7	18.8
liver	8	6.40	60.1	459.1	533.7
spleen	0.25	5.92	107.0	174.5	265.2
kidney	0.5	27.17	82.0	543.4	695.8
lung	0.75	0.12	55.5	24.4	26.5
heart	0.5	0.76	135.0	9.2	14.8
brain	0.75	0.11 (0.04)*	-	3.6	-

Clearance (= dose/AUC) of the drug from body was calculated from plasma data to be 0.53 mg/kg/h.

* value at 7-day time point ($C_{max}[L-AST]$)

** no terminal phase achieved