

# No correlation between side-chain oxidation and S-mephenytoin 4'-hydroxylase activity

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KEY WORDS propranolol; mephenytoin; cytochrome P-450; hydroxylases; pharmacokinetics

AIM: To determine any relation between the side-chain oxidative capacity for propranolol and S-mephenytoin 4-hydroxylase (cytochrome P-450 CYP2C19) activity in human. METHODS: S-mephenytoin oxidative metabolite 4'-hydroxymephenytoin (4'-OH-M) S- and R-mephenytoin and naphthoxyacetic acid (NLA) excretion in plasma were measured after 14 healthy extensive smokers. Data were given as geometric mean  $\pm$  SD. 100 mg and 80 mg propranolol were administered. S/R-mephenytoin was determined by chiral capillary HPLC with nitrogen-phosphorus detection. 4'-OH-M in urine by reversed-phase HPLC with ultraviolet detection and plasma propranolol or NLA by HPLC with fluorescence detection. RESULTS: No correlation was found between the plasma metabolic clearance ( $\alpha_m$ ) of propranolol and NLA and S/R-mephenytoin ( $r_s = -0.08$ ;  $P = 0.869$ ) nor between the  $\alpha_m$  of propranolol and NLA and S/R-mephenytoin ( $r_s = 0.11$ ;  $P = 0.714$ ). CONCLUSIONS: CYP2C19 is not a principal P-450 isozyme for the side-chain oxidation of propranolol.

Propranolol undergoes extensive first-pass metabolism by human liver microsomes.

<sup>1</sup>Project supported by the National Natural Science Foundation of China (No. 9200154 and 9330230) and by the Chinese Academy of Sciences (No. 92-568).  
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Received 1996-09-13 Accepted 1997-04-27

P450 (CYP) proteins via 3 main pathways: aromatic naphthalene hydroxylation at 1- and 5-positions, side-chain oxidation (N-desisopropylation of N-deethylpropylamine) and glucuronidation of aromatic hydroxylation is catalyzed predominantly by human debrisoquine 4-hydroxylase (CYP2D6) and the N-desisopropylation of propranolol to naphthoxyacetic acid (NLA) is dependent upon human CYP2C19. However, the side chain N-desisopropylation is mediated principally by recombinant and/or cloned human CYP2C19 and human liver microsomal CYP2C19 but not by human liver microsomal CYP2C19. This is in agreement with a lack of inhibition of propranolol N-desisopropylation by mephenytoin in human liver microsomes; propranolol N-desisopropylation was correlated with CYP2C19 content ( $r = 0.40$ ;  $P = 0.01$ ). Hydroxylation ( $r = 0.16$ ) catalyzed largely by CYP2C19 and S-mephenytoin 4-hydroxylation ( $r = 0.60$ ;  $P = 0.0005$ ) implying that CYP2C19 may be the major P450 isozyme in propranolol N-desisopropylation. Thus, a major role played by CYP2C19 in the involvement of propranolol N-desisopropylation was proposed on the basis of above evidence and the fact that the plasma metabolite clearance ( $\alpha_m$ ) of propranolol was much higher in smokers than in non-smokers. The induction of CYP2C19 by smoking and a controversy concerning the relative contribution of CYP2C19 to the side-chain oxidation of propranolol.

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**SUBJECTS AND METHODS**

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8 h urine (m) beion on of 4' 8 h urinary on of 4' I-OH M) bei 90 %-1 d S-mephenytoin d (19 D %). The study prot o. w approved by e Ethics t O {HZJ TZ Irlivmity and writtm idoomed nsent was each subject.

pmtMcmc19acMty d ch subj was determined by taking 8 single oral do 0l 10@ raem1c mephenytoin (Mesantoin<sup>o</sup> ndoz) after an overnight fast. Urine w Uoated during the xstdose period of 0 8 h and m limot stored at-30 .4'-OH-M levds m mne w e by an unproved LC method(11) and amounts of S- and R-meph ytoin in U!in@ by cl al capillrury gas ; m th a sli t nlodification<sup>13'</sup>). The CYP2C19 ac vity was pressed S/R ratio or 19 D %.

At least 2 wk after the mephenytoin test each subject received a single oral dose of 80 mg of racemic propranolol h chloride (Inderal tablet 10 mg each Beijing 2nd pharmacMid Co)Using a si protocol what o mephenytoin t t urine was J. pected for 24 h. Urin Y NLAwasmoured by HPLC<sup>14)</sup> - t mination of NLA in reolicate urine samples w s done within 1 d. Estimate a the fraction d to d excreted λin the urine ( fro) w mean value of t e 2 p:allet samples of each subject. Venous blcx (10 mL) was heparinized before and after oral propranolol PI ma propranolol w quantitated by HPLC

using 4-methylpropranolol stand d.

Data analysis Individual elimination rate constant (Ke) was timated from the slope of the terminal phase (3 -24 h) of the log plasma ncmntation-timeα rrv e. The oral clærance of propranolol (Cl 0) w: calculated with tne equations: Cl<sub>o</sub> = d /AUC Cl<sub>m</sub> = j x Cl<sub>o</sub>, in which f<sub>m</sub> is the fraction of the total dose excreted as the nletabolite NLA in urine. CO elation between Cl<sub>m</sub> of propranolol to NLA and CYP2C19 activity (S/R ratio 19 D %) was analyzed by nonparametic Spearman's r lk rrelation with e use of the SAS syst (SAS 6.10 version SAS Institute C NC USA).

**RESULTS AND DISCUSSION**

In this study relationship between S/R ratio and Cl<sub>m</sub> (r<sub>s</sub> = 0.0484; P = 0.8695) or between Cl<sub>m</sub> and S/R ratio (r<sub>s</sub> = -0.1077; P = 0.7140) had no sismificant Irrelations in the EM subjects with a lazprange of cymc19acd vity(Fig1). Although

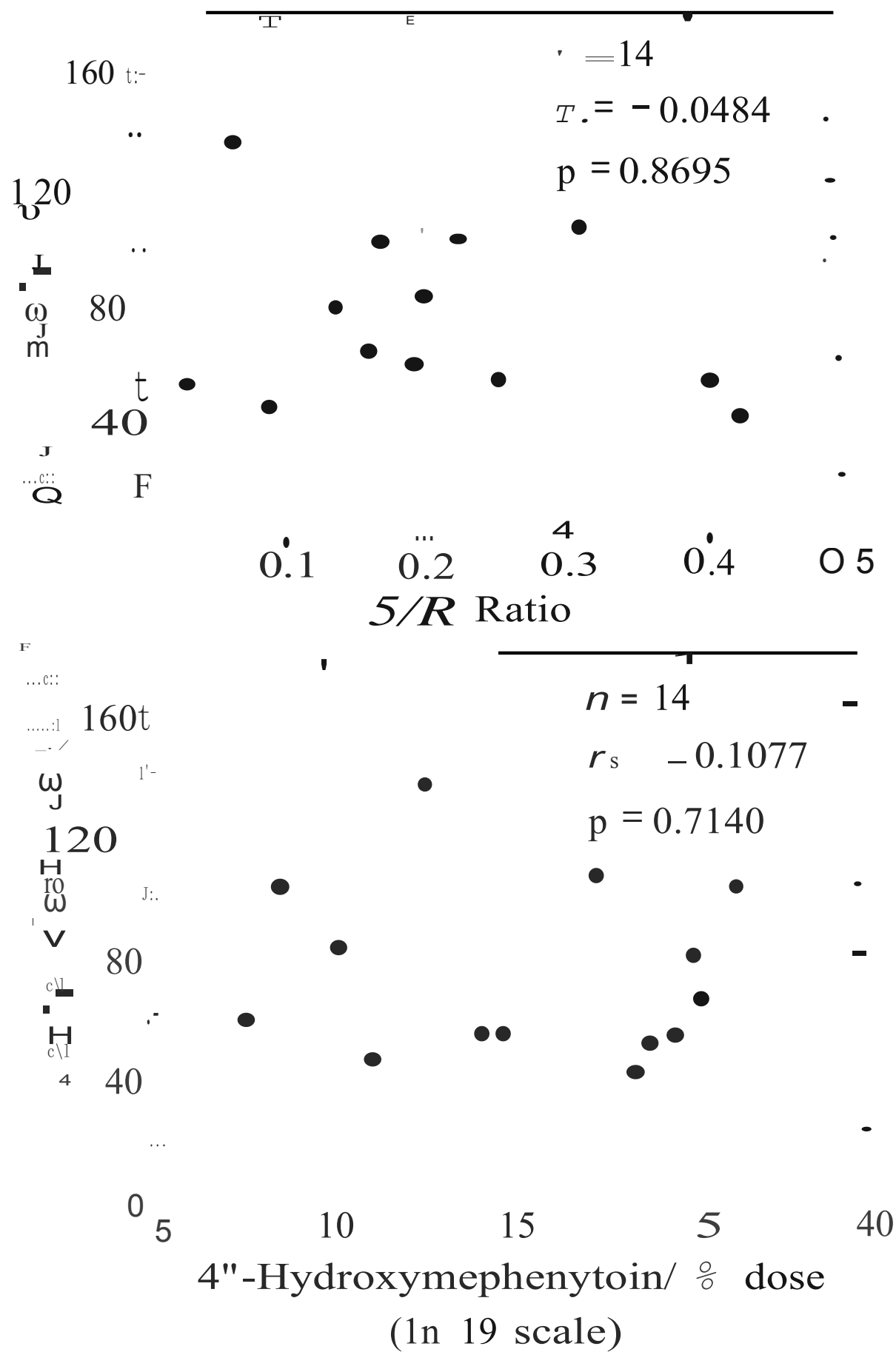


Fig 1. Correla nbe een Cl<sub>m</sub> of propranolol to NLA and S/R ratio of mephenytoin (up' ) and 4'-OH-M % d f S-mephenytoin (lower) excre d in the ineO-8h ilstdoe.

Ward et aZ(2) found that the Cl<sub>m</sub> of propranolol to NLA a further oxidative metabolite of N- desisopropylpropranolol correlated highly with the S/R ratio of mephenytoin in the postdose 0 - 8 h urine samples (r = 0.80') and was somewhat lower in the p r metabolizers (PM) of S-mephenytoin than in the EM subjects they also suggested that other CYF iso mes may be capable of catalyzing the same m. etaoolic reaction. Thus an under-

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