

COMMENTARY

Clinical considerations in study designs that use cotinine as a biomarker

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Received 1 November 2002, revised form accepted 11 April 2003

Subjects enrolled in studies are not always screened for routine habits such as smoking. Personal history is not always reliable and therefore an objective biomarker is necessary to screen for smokers. The objectives of this article were to review the metabolism of nicotine and other metabolic considerations associated with smoking; to review some of the routine methods used to assess exposure to nicotine-containing products; to revisit cotinine breakpoints utilized to distinguish smokers from non-smokers during screening for clinical trials; to assess the utility of screening questions regarding smoking practices; and to recommend standards for clinical pharmacology studies. The results indicated that cotinine levels serve as a useful biomarker of tobacco exposure; racial issues may be clinically relevant in determining smoking status; cessation of smoking should occur at least 14 days prior to the start of the study; adverse effects from nicotine withdrawal such as craving, hunger and weight gain may persist for more than 6 months; potential metabolic interactions via cytochrome P2A6 and P1A2 need to be considered when designing a study; and the use of a single calibrator as a breakpoint is acceptable if a categorical outcome such as 'smoker' versus 'non-smoker' is desired. Nicotine from food products is not expected to impact assay sensitivity or to be clinically relevant; a serum cotinine concentration of 10 ng ml⁻¹ be employed as a breakpoint for non-smokers versus smokers; other non-invasive alternatives are collection of urine, saliva, or hair (with suggested breakpoints of 200 ng ml⁻¹, 5 ng ml⁻¹ and 0.3 ng mg⁻¹ 1, respectively; screening questions be accompanied by testing for cotinine; and the inclusion of smokers in studies should be considered once the impact of smoking on the targeted population is understood.

Keywords: Cotinine, biomarker, assays, screening, questionnaire, clinical study designs.

Introduction

Clinical pharmacology studies typically exclude unsuitable subjects based on medical history, undesirable demographic characteristics, dietary issues, and/or use of drugs, including recreational drugs. However, subjects enrolled in a study are not always screened for routine habits such as smoking. It is well established that tobacco products may alter metabolism as well as response to drugs, as can the adverse events associated with withdrawal from tobacco (Hughes et al. 1984, Benowitz 1988, Miller 1989). Adverse effects from nicotine withdrawal such as craving, hunger and weight gain may persist for more than 6 months (Hughes et al. 1994).

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As demonstrated by Apseloff et al. (1994), self-reported personal history is not always reliable as a means of excluding smokers. Thus an objective marker is needed to screen for smokers.

Cotinine is a major metabolite of nicotine. Many studies have used this substance as a biomarker of tobacco smoking (Ehrlich et al. 1992, Apseloff et al. 1994, Benowitz 1996, Haufroid and Lison 1998). However, before this biomarker can be successfully employed for assessing exposure to tobacco or other sources of cotinine, it is important to establish a specific range of cotinine levels in the various types of biological samples (e.g. plasma, urine). The objectives of this article are: (i) to review the metabolism of nicotine and other metabolic considerations associated with smoking; (ii) to review some of the routine methods used to assess exposure to nicotine-containing products; (iii) to revisit cotinine breakpoints utilized to distinguish smokers from non-smokers during screening for clinical trials; (iv) to assess the utility of screening questions regarding smoking practices; and (v) to recommend standards for clinical pharmacology studies.

Metabolism of nicotine and cotinine

Nicotine is a major constituent of tobacco and undergoes extensive oxidative metabolism in the human liver. The major pathway of nicotine metabolism is Coxidation to cotinine. It has been estimated that an average of 70-80% of the nicotine absorbed by a smoker is metabolized to cotinine, and approximately 10-15% of the circulating cotinine is excreted in the urine. The remainder of the cotinine is further converted to other metabolites, including trans-3'-hydroxycotinine, cotinine glucuronide, cotinine-N-oxide, norcotinine and trans-3'-hydroxycotinine glucuronide. The other metabolites of nicotine, such as nicotine glucuronide, nicotine-1'-N-oxide, and nornicotine, are eliminated in the urine. About 10% of the nicotine is eliminated unchanged in urine (Benowitz 1996). The mean ± SD protein binding values of nicotine and cotinine in the plasma are 4.9 + 2.8% and $2.6\pm3.5\%$, respectively (Benowitz et al. 1982, 1983a). Thus protein binding does not meaningfully alter the pharmacokinetics of nicotine or cotinine. Cotinine is widely distributed throughout the body, with values 25–33% more than the volume of total body water ($\sim 0.60 \text{ l kg}^{-1}$). The mean plasma half-lives of nicotine, cotinine, and trans-3'-hydroxycotinine in smokers are about 2-3, 17 and 6 h, respectively (Benowitz 1996, Perez-Stable et al. 1998).

Other metabolites of nicotine such as trans-3'-hydroxycotinine have been investigated. Although the amount of trans-3'-hydroxycotinine excreted in urine is similar or higher compared with cotinine, the serum concentrations of trans-3'hydroxycotinine were much lower throughout an observation period of 60 h after infusion of nicotine or cotinine. When trans-3'-hydroxycotinine is infused and the half-life determination post-infusion calculated, the plasma half-life appeared shorter (6 h) than following post-infusion of cotinine (15-17 h) (Scherer et al. 1988, Curvall et al. 1990, Benowitz and Jacob 2001). There is a need for further research to clarify the relative elimination of trans-3'-hydroxycotinine following cotinine or nicotine administration.



The cotinine level in the blood of smokers is approximately 10 times that of nicotine (Benowitz et al. 1983b). Because of the longer half-life and higher plasma concentrations of cotinine, measurement of cotinine is less dependent on the exact timing for blood sampling. Therefore, it is a preferred surrogate measurement to estimate nicotine acquired from tobacco.

In two studies, the clearance and half-life of cotinine have been shown to be ethnically dependent (see Table 1). In the study comparing African Americans to Caucasians, the average clearance and non-renal clearance were about 18% and 23% lower in the African Americans, respectively, (Perez-Stable et al. 1998). The half-life appeared to be slightly longer in African Americans, but this difference was not statistically significant. In the study comparing the pharmacokinetics of cotinine in Latinos, Chinese Americans and Caucasians, a statistically significant difference was detected for clearance (total, renal and non-renal) and for half-life (Benowitz et al. 2002). Further statistical analysis revealed that both the renal and the non-renal clearance of Chinese Americans was statistically different from that of Caucasians: the average renal clearance was 40% slower and the average non-renal clearance was 30% faster in Chinese Americans. The half-life appears to be race dependent, but not enough data is available to show statistical differences among race comparisons.

There are individual variations in the quantitative relationship between cotinine level and nicotine intake. This variation is thought to be due to individual differences in the percentage of conversion of nicotine to cotinine (ranging from 55–92%) and the clearance rate for cotinine (ranging from 19–75 ml min⁻¹). Several studies have demonstrated the important role of cytochrome P450 2A6 (CYP2A6) in the C-oxidation of nicotine to generate cotinine (Nakajima et al. 1996, Murphy et al. 1999, Yamazaki et al. 1999). The degree of variation in the metabolism of nicotine in humans is possibly due to variable expression of CYP2A6 (Messina et al. 1997). Nakajima et al. (2000) recently showed that deficient cotinine formation in humans after cigarette smoking is associated with deletion of the CYP2A6 gene.

Table 1. Reported pharmacokinetics of cotinine.

Race	Clearance (ml min ⁻¹ kg ⁻¹)	Renal clearance (ml min ⁻¹ kg ⁻¹)	Non-renal clearance (ml min ⁻¹ kg ⁻¹)	Vss (l kg ⁻¹)	Half-life (h)	
Caucasian ^a African American ^a p value ^c	$\begin{array}{c} 0.68 \pm 0.24 \\ 0.56 \pm 0.17 \\ 0.009 \end{array}$	0.08 ± 0.04 0.10 ± 0.04 0.10	$\begin{array}{c} 0.61 \pm 0.25 \\ 0.47 \pm 0.18 \\ 0.009 \end{array}$	0.80 ± 0.19 0.75 ± 0.11 0.16	$15.8 \pm 4.9 \\ 17.7 \pm 4.3 \\ 0.07$	
Latino ^b Chinese American ^b Caucasian ^b p value ^c	$\begin{array}{c} 0.76 \pm 0.32 \\ 0.60 \pm 0.23 \\ 0.76 \pm 0.35 \\ 0.001 \end{array}$	0.10 ± 0.04 0.14 ± 0.06 0.09 ± 0.05 0.006*	0.66 ± 0.33 0.46 ± 0.22 0.67 ± 0.34 < 0.001*	0.79 ± 0.21 0.83 ± 0.19 0.93 ± 0.24 0.12	$14.6 \pm 5.2 \\ 18.3 \pm 5.4 \\ 16.1 \pm 5.2 \\ < 0.001$	

Values are mean \pm SD.



^aPerez-Stable et al. 1998.

^bBenowitz et al. 2002.

 $^{^{}c}p < 0.05$ indicates statistical significance. *The value for Chinese Americans is statistically different from that for Caucasians

Despite interindividual and racial variation, it has been demonstrated that the apparent shape of the disposition profile of nicotine and cotinine are similar in smokers and non-smokers. The half-life and volume of distribution in smokers and non-smokers was not significantly different. However, it is unresolved as to whether the clearance of nicotine in non-smokers is faster or slower compared with smokers (Benowitz and Jacob 1993). After cessation of smoking, blood and urine cotinine concentrations declined to non-smoking levels within 3-4 days (Jarvis et al. 1988, Benowitz and Jacob 1993). Therefore, given the population or interindividual variation, a normal range for cotinine should be established to reflect the level of nicotine exposure for individuals before they are enrolled into clinical trials.

Nicotine, smoking and drug interactions

Table 2 provides examples of drugs that are substrates, inhibitors and inducers of CYP2A6 (Pritchard and Wolf 2000). Drugs that are competitive substrates or potent inhibitors of CYP2A6 could result in elevated nicotine concentrations and diminished cotinine concentrations in subjects using tobacco products. Drugs that induce CYP2A6 could result in lower nicotine concentrations and may increase or decrease cotinine levels, since cotinine is both generated by and metabolized by CYP2A6. While nicotine itself is metabolized by CYP2A6, the inhalation of smoke induces cytochrome P450 1A2 (CYP1A2) (Schein 1995, Pelkonen et al. 1998, Zevin and Benowitz 1999). CYP1A2 is responsible for the metabolism of numerous drugs (see Table 3) and the activation of some procarcinogens (Ahn et al. 2000, Lode 2001, Physicians' Desk Reference 2001, Yew 2002). It has been shown that cigarette smoking can affect the pharmacokinetic and pharmacodynamic properties of many drugs (Miller 1989, 1990, Zevin and Benowitz 1999). The pharmacokinetic interactions between cigarette smoking and certain concomitant medications or xenobiotic exposure may be mediated by the induction of CYP1A2. The polycyclic aromatic hydrocarbons (PAHs) rather than the nicotine in the tobacco smoke are believed to be responsible for this induction. In humans, CYP1A2 is detected only in the liver, where it is regulated by at least two mechanisms: one controlling the constitutive expression levels and the other regulating its inducibility (Butler et al. 1992, Landi et al. 1999). It has been shown

Table 2. Examples of drugs that are substrates, inducers and/or inhibitors of CYP2A6.

Substrates	Inducers	Inhibitors		
Coumarin	Phenobarbitone	Coumarin		
Dicoumarol	Dexamethasone	Nicotine		
Warfarin	Rifampicin	Methoxsalen		
Nicotine	_	Tranylcypromine		
Cotinine		Pilocarpine		
Halothane		Menadione		
Losigamone		Miconazole		
Valproic acid		Clotrimazole		
Methoxyflurane		Ketoconazole		
Sevoflurane		Psoralen		
Methyl tert-butyl ether		Metyrapone		



Table 3. Examples of drugs that are substrates, inducers and/or inhibitors of CYP1A2.

Carvedilol Rifampin Cimetidine Cilostazol Ritonavir Fluorquinolones Cinnarizine Ciprofloxacin Clomipramine Enoxacin Clozapine Gatifloxacin Clozapine Levofloxacin Clunarizine Lomefloxacin Cluvoxamine Lomefloxacin Cluvoxamine Moxifloxacin Cluvoxamine Moxifloxacin Cluvoxamine Moxifloxacin Cluvoxamine Moxifloxacin Cluvoxamine Moxifloxacin Cluvoxamine Moxifloxacin Cluvoxamine Interferon Cluvoxamine Interferon Cluvoxamine Interferon Cluvoxamine Fluvoxamine Clinarizine Methoxsalen Cluvoxamine Interferon Clivoxamine Clivo	Substrates	Inducers	Inhibitors
Ellostazol Ritonavir Fluorquinolones Cinnarizine Ciprofloxacin Clozapine Enoxacin Clozapine Gatifloxacin Clozapine Levofloxacin Cluvoxamine Lomefloxacin Cluvoxamine Moxifloxacin Cluvoxamine Interferon Cluvoxamine Interferon Cluvoxamine Interferon Cluvoxamine Interferon Cluvoxamine Interferon Clozacin Clozacin Clozacin Clozacin Clozacin Clovacin C	Amitriptyline	Omeprazole	Amiodarone
Ciprofloxacin Clomipramine Clozapine Clozacin Cl	Carvedilol	Rifampin	Cimetidine
Clomipramine Enoxacin Clozapine Gatifloxacin Clozapine Levofloxacin Cluvoxamine Lomefloxacin Cluvoxamine Moxifloxacin Moxignocaine (injected) Morfloxacin Sparfloxacin Mexiletine Fluvoxamine Mianserin Interferon Mirtazapine Methoxsalen Marpoxen Methimazole (thiamazole) Destradiol, oestrone Moclobemide Delanzapine Propofol Tropafenone Ticlopidine Troparenolol Cluzole Copinirole	Cilostazol	Ritonavir	Fluorquinolones
Clozapine Gatifloxacin Cyclobenzaprine Levofloxacin Clunarizine Lomefloxacin Cluvoxamine Moxifloxacin Cluvoxamine Moxifloxacin Cluvoxamine Moxifloxacin Cluvoxamine Moxifloxacin Moxifloxacin Moxifloxacin Moxifloxacin Moxifloxacin Morfloxacin Morfloxacin Morfloxacin Morfloxacin Cluvoxamine Cluvo	Cinnarizine		Ciprofloxacin
Acyclobenzaprine Ilunarizine Ilunarizine Iluvoxamine I	Clomipramine		Enoxacin
Illunarizine Illuvoxamine Illuvoxamine Illuvoxamine Illuvoxamine Illuvoxamine Illuvoxamine Illuvoxamine Illuvoxamine Illuvoxamine Illuroxacin Morfloxacin Morfloxacin Morfloxacin Morfloxacin Morfloxacin Sparfloxacin Illuroxamine Illuroxamin	Clozapine		Gatifloxacin
Tuvoxamine Idaloperidol Moxifloxacin Morfloxacin Sparfloxacin Sparfloxacin Fluvoxamine Minereron Minereron Minereron Mexiletine Methoxsalen Methoxsalen Methimazole (thiamazole) Moclobemide Propofol Propafenone Propofol Triclopidine Propofol Triclopidine	Cyclobenzaprine		Levofloxacin
Italoperidol Norfloxacin mipramine Ofloxacin Sparfloxacin Mexiletine Fluvoxamine Mintazapine Methoxsalen Methoxsalen Methomazole (thiamazole) Destradiol, oestrone Meclobemide Planzapine Propofol Tropafenone Ticlopidine Tropranolol Miluzole Lopinirole Lopivacaine Tacirine Theophylline Thiabendazole Gerapamil Gileuton	Flunarizine		Lomefloxacin
mipramine Ofloxacin signocaine (injected) Sparfloxacin Aexiletine Fluvoxamine Aianserin Interferon Airtzapine Methoxsalen Aproxen Methoxsalen Abertradiol, oestrone Moclobemide Postradiol, oestrone Propofol Aropafenone Ticlopidine Aropranolol Aiduzole Atopivacaine Acarrine Cheophylline Chiabendazole Erapamil Gileuton	Fluvoxamine		Moxifloxacin
Agnocaine (injected) Agrocaine Interferon Agrocaine Methoxsalen Methoxsalen Methoxsalen Moclobemide Propofol Tropafenone Tropafenone Tropafenone Tropafenone Tropidine Tropidin	Haloperidol		
Aexiletine Aianserin Airtazapine Aproxen Aprox	Imipramine		Ofloxacin
Mianserin Mirtazapine Mirtazapine Methoxsalen Methoxsalen Methimazole (thiamazole) Moclobemide Moclobemide Moclobemide Propofol Ticlopidine Tropafenone Tropafenone Tropafenole Moclobemide Tropafenone Tropafenon	Lignocaine (injected)		Sparfloxacin
Airtazapine Methoxsalen Maproxen Methimazole (thiamazole) Destradiol, oestrone Moclobemide Dlanzapine Propofol Tropafenone Ticlopidine Tropranolol Giluzole Ropinirole Ropivacaine Tacrine Theophylline Thiabendazole Gerapamil Gileuton	Mexiletine		Fluvoxamine
Japroxen Methimazole (thiamazole) Destradiol, oestrone Moclobemide Dlanzapine Propofol Tropafenone Ticlopidine Tropranolol Giluzole Lopinirole Lopivacaine Tacrine Theophylline Thiabendazole Gerapamil Gileuton Ticlopidine	Mianserin		Interferon
Destradiol, oestrone Ticlopidine Ticlop	Mirtazapine		Methoxsalen
Dianzapine Propofol Tropafenone Ticlopidine Troparanolol Liluzole Lopinirole Lopinirole Lopivacaine Cacrine Cheophylline Chiabendazole Lerapamil Lileuton	Naproxen		Methimazole (thiamazole)
ropafenone Ticlopidine ropranolol kiluzole kopinirole kopivacaine Gacrine Theophylline Thiabendazole ferapamil kileuton	Oestradiol, oestrone		Moclobemide
ropranolol ciluzole copinirole co	Olanzapine		Propofol
kiluzole kopinirole kopivacaine facrine Theophylline Thiabendazole ferapamil kileuton	Propafenone		Ticlopidine
Appinirole Appivacaine Factine Theophylline Thiabendazole Ferapamil Fileuton	Propranolol		
Ropivacaine Pacrine Pheophylline Phiabendazole Perapamil Poileuton	Riluzole		
acrine Theophylline Thiabendazole Terapamil Gileuton	Ropinirole		
'heophylline 'hiabendazole 'erapamil iileuton	Ropivacaine		
'hiabendazole 'erapamil Gileuton	Tacrine		
erapamil Gileuton	Theophylline		
illeuton	Thiabendazole		
	Verapamil		
olmitriptan	Zileuton		
	Zolmitriptan		

that cigarette smoking upregulates CYP1A2 expression at the transcriptional level and is probably tissue-specific (Quattrochi et al. 1994). It may take several days to recover from CYP1A2 induction at the mRNA level, but it may take a much longer time at the protein level. In a study on hepatic cytochrome P450-dependent drug metabolism in rabbits, it was shown that the drug-metabolizing capacity and total P450 enzyme content returned to normal 7 days after the first injection of a stimulating agent (Saitoh et al. 1999). In a study assessing the metabolism of theophylline after cessation of smoking for 16 days, the results suggested that clearance of theophylline returned to the level of non-smokers as early as after 7 days (Benowitz et al. 1989). In studying the interaction between oltipraz and CYP1A2 in humans, using caffeine as an in vivo probe, Sofowora et al. (2001) recently reported that the reduction in CYP1A2 activity was reversible: after discontinuing oltipraz, the activity of CYP1A2 returned to 57% and 100% of base activity after 2 and 14 days, respectively. These factors should be taken into consideration when designing studies and interpreting safety, pharmacodynamic and pharmacokinetic data.

Cotinine as a biomarker for tobacco smoking

Chemicals in tobacco, such as carbon monoxide, thiocyanate, cyanide, anabasine and anatabine, can be detected and measured in the blood (SNRT



Subcommittee on Biochemical Verification 2002). Testing for these components of tobacco can require specialized equipment and may be time consuming, and it is expensive to obtain rapid results. In addition, the levels of these chemicals are either non-specific or insensitive (Benowitz 1996, SNRT Subcommittee on Biochemical Verification 2002). Plasma nicotine levels correlate well with smoking exposure, but blood samples must be collected during or immediately after exposure because of the short half-life; 30 min after 6.5 h of smoking, the plasma nicotine concentrations were $0.012-0.044 \text{ mg l}^{-1}$ in smokers compared with $0-0.006 \text{ mg l}^{-1}$ in nonsmokers (Baselt 1995).

Cotinine is highly specific and sensitive for tobacco use in the absence of nicotine replacement therapy (SNRT Subcommittee on Biochemical Verification 2002). Cotinine concentrations have been shown to correlate well with nicotine exposure, and plasma cotinine levels of 2-10 ng ml⁻¹ was considered to indicate environmental tobacco smoke exposure (Eskenazi et al. 1995). Another study determined that steady-state cotinine levels were linearly and directly related to the daily available nicotine (r = 0.919), with an increase in correlation coefficient directly related to the increase in tar and nicotine yield (Rosa et al. 1992). However, other research has indicated that nicotine intake per cigarette smoked, as estimated from salivary cotinine levels, does not correspond with expected machine-smoked vields of nicotine. Machine-smoked nicotine yields are poor predictors of nicotine intake in smokers. Cotinine levels in blood, saliva, semen and urine were correlated quantitatively during passive and active smoking (Curvall et al. 1990, Vine et al. 1993). Cotinine levels in semen and blood were similar in magnitude, but levels were an order of magnitude higher in urine. For non-smokers, smokers of 1-19 cigarettes and smokers of > 20 cigarettes, the median cotinine concentrations were 0.8, 220 and 602 ng ml⁻¹, respectively, in semen, 0.8, 137 and 467 ng ml⁻¹, respectively, in blood, and 17, 3516 and 7179 ng ml⁻¹, respectively, in urine. Cotinine levels increased significantly with the increase in the number of cigarettes smoked per day (Vine et al. 1993). Since cotinine concentrations are substantial in these three body fluids, any of them can be used to assess smoking status. The advantage of urine collection is that it is non-invasive.

Besides tobacco exposure, some foods, including aubergine, potato, tomato, cauliflower, bell pepper and black tea, contain small amounts of nicotine (Castro and Monji 1986, Davis et al. 1991). For the population of countries for which consumption data were available, the mean daily dietary nicotine intake was approximately 1.4 µg per day (2.25 µg per day at the 95th percentile) (Siegmund et al. 1999). Davis et al. (1991) estimated the average daily intake of nicotine from food sources is about 9 µg, and that the maximum daily consumption of nicotine from these foods would be about 100 µg. They also reported average and maximal daily urinary cotinine levels from food consumption of 0.6 and 6.2 ng ml⁻¹, respectively. Most of these calculations are based on the ingestion of black tea; however, the necessary volume to achieve these amounts would be about 4 l per day. This would result in diuresis, dilute urine and concentrations below 6.2 ng ml⁻¹ (Benowitz 1996). Therefore, for most people, nicotine intake from food sources is insignificant compared with cigarette smoking, and is not expected to impact assay sensitivity or be clinically relevant. Dietary restrictions of food sources



of nicotine are not considered necessary in designing a study. In summary, cotinine is a specific and sensitive biomarker for cigarette smoking.

Determination of cotinine levels

Plasma/serum

A typical plasma cotinine level for a smoker is 300 ng ml⁻¹, which corresponds to a daily intake of 24 mg of nicotine. On average, a cigarette smoker absorbs about 1 mg of nicotine per cigarette smoked (Benowitz 1996). Multiple methodologies (enzyme immunoassay, high performance liquid chromatography, radioimmunoassay, Immulite, gas chromatography, liquid chromatography-mass spectrometry) have been employed in the determination of plasma and serum cotinine levels. Examples of some of these methodologies used for screening the smoking status of subjects are discussed briefly below; detailed consideration of very sensitive and specific research tools such as gas chromatography and liquid chromatographymass spectrometry are beyond the scope of this review.

A competitive-inhibition enzyme immunoassay (EIA) technique was used to analyse serum samples from 30 current smokers, 34 former smokers and 15 nonsmokers collected over a 1 year period (Gonzalez et al. 1996). Since the mean cotinine level did not change in an individual over time, it was concluded that it could be used reliably to assess the level of smoking. In non-smokers the maximum serum cotinine level was 19.7 ng ml⁻¹, with a mean of 11.2 ng ml⁻¹, and in current smokers the minimum serum cotinine level was 47.8 ng ml⁻¹, with a mean of 564.9 ng ml⁻¹. These results were consistent with radioimmunoassay results (Vine et al. 1993).

Lawson et al. (1998) determined plasma cotinine concentrations using high performance liquid chromatography (HPLC) and found cotinine levels of 179, 266 and 304 ng ml⁻¹, respectively, in light (10-15 cigarettes per day), moderate (16-30 cigarettes per day) and heavy (>30 cigarettes per day) smokers. Pirkle et al. (1996) summarized the distribution of serum cotinine levels based on a survey in the US population aged 4 years and older. The highest cotinine levels were 0.1, 0.8 and 600 ng ml⁻¹ in those without reported environmental tobacco smoking exposure, those with reported environmental exposure, and tobacco smokers, respectively.

In our own research, Immulite assays from Diagnostic Products Corporation (Los Angeles, California, USA) were used for the measurement of cotinine levels in serum. Immulite assays are chemiluminescent immunoassays performed on an automated analyser that measure cotinine quantitatively. The detection limit for the method is 2 ng ml⁻¹ with 20 µl of serum. The lower and upper limits of quantification are 10 and 500 ng ml⁻¹, respectively. Samples were assayed in duplicate during the validation period of 20 days, two runs per day, for a total of 40 runs and 80 replicates. The precision was 6.3% within run and 16% overall. The antibody used was highly specific to cotinine, with non-detectable cross-reactivity to other naturally occurring compounds that might be present in patient samples. Cross-reactivity to 3-hydroxycotinine was 28%. Values of \leq 20 ng ml⁻¹ were used to distinguish non-smokers from smokers. This assay was used to screen 30 healthy



Table 4. Screening questions for tobacco use.

- Are you exposed to tobacco products by handling them (e.g. farming tobacco, manufacturing cigars, etc)?
- Do you use tobacco products of any form (e.g. cigarettes, pipes, cigars, chewing tobacco, etc.)?
- - If so, in what form: \Box cigarettes \Box pipe \Box cigars
- Approximately how may cigarettes, cigars, or pipes do you smoke per day?
- Are you an ex-smoker? If so, about when [month/year] did you quit? Have you used tobacco products within the last 6 months?
- Do you use any products to help quit or prevent smoking (e.g. gum, patch, etc.)?
- Do you live with a smoker?
- Do you frequent establishments where smoking is allowed?
- Are there any circumstances where you are routinely (at least once a day) exposed to 'second hand' smoke?

subjects who claimed not to smoke. These subjects were asked several screening questions (Table 4) prior to serum cotinine testing. All the enrolled subjects had serum cotinine levels of < 10 ng ml⁻¹ (the lower limit of quantification). Thus, appropriate screening questions are effective, and the serum cotinine concentrations in non-smokers should be $< 10 \text{ ng ml}^{-1}$.

Urine

The correlation between urinary cotinine and daily tobacco smoke is approximately 75% or greater (Haufroid and Lison 1998). Urinary cotinine reaches peak levels at 6 h after stopping environmental tobacco smoke exposure, remains at an apparent plateau for about half a day, and then decreases with a mean half-life of 19 h (Willers et al. 1995). For urinary cotinine analysis, urine samples are stable for up to 30 h at room temperature or for up to 8 days at -25° C (Lequang et al. 1994). Another study showed that cotinine measurements made based on the urine samples stored at -20° C for 10 years still allowed a clear separation of smokers and non-smokers, with 92% sensitivity and 100% specificity (Riboli et al. 1995).

Urine cotinine levels can be used to follow occupational exposure in tobacco pickers. The urinary cotinine level averaged 35 ng g⁻¹ creatinine in unexposed persons and 890 ng g⁻¹ creatinine in 16 persons who were heavily exposed at work (Gehlbach et al. 1975). In another study, researchers used a urinary cotinine/ creatinine ratio (CCR) ≥ 30 ng mg⁻¹ to identify children exposed to passive smoking at home (Ehrlich et al. 1992).

Table 5 summarizes the results of three different assays routinely used to screen subjects for urine cotinine. The first method is a double antibody radioimmunoassay (RIA) kit from DPC (Diagnostic Products Corporation). In a liquid-phase RIA, ¹²⁵I-labelled cotinine competes for a fixed time with cotinine (and other nicotine metabolites) in the urine sample for antibody sites. After incubation, separation of bound sites from free is achieved using the phosphoethylene glycol (PEG)-accelerated double antibody method. The antibody-bound fraction is then precipitated and counted. The samples were analysed in duplicate and the concentrations were read from a calibration curve. The detection limit of the method was 5 ng ml⁻¹ using 50 μl of urine. The lower and upper limits of quantification were 100 and 15 000 mg ml⁻¹, respectively. Correlation coefficients



Table 5. Screen failures attributed to smoking identified by post-screening questions in various studies.

Study .	Breakpoint used Assay type $(ng ml^{-1})$		Total no. screened	Screen failures identified by screening questions		Subjects passing screening questions but positive for cotinine		Cotinine in non-smokers (ng ml ⁻¹)	
		used		No.	%	No.	%	Mean \pm SD	Range
European study 1	RIA	≥ 300	40	8	20	4	10	89 ± 51	16-201
European study 2	RIA	≥ 300	14	0	0	0	0	$< \overline{100}^{\rm b}$	
European study 3	RIA	≥ 300	31	0	0	0	0	115 ± 46	99 - 297
European study 4	RIA	≥ 300	37	3	8.1	0	0	104 ± 17	99 - 173
European study 5	RIA	≥ 300	51	0	0	0	0	99 ± 1	99 - 106
European study 6	Immulite	≥ 300	57	2	3.5	2	3.5	101 ± 10	100 - 163
European study 7	Immulite	$< 20^{a}$	57	2	3.5	0	0	$< 1\overline{0}^{ m b}$	
US study 1	EIA	≥ 300	12	0	0	0	0	ND	
US study 2	RIA	_ ≥ 300	850	58	6.8	_	_	ND	

RIA, radioimmunoassay; EIA, enzyme immunoassay; ND, not determinable.

^aCut-off point in serum.

^bAll values less than the value indicated.

of calibration curves recorded during the validation period were above 0.9930. According to the manufacturer's information, the anti-serum is highly specific for nicotine metabolites, with no cross-reactivity detected with other drugs.

The second method is a chemiluminescent immunoassay called Immulite (Diagnostic Products Corporation) that can also be used for the measurement of cotinine level in serum as described above.

The third method is a cotinine EIA developed by Diagnostic Reagents, Inc. This cotinine assay is a liquid, ready-to-use homogeneous EIA based on the competition of a cotinine-labelled enzyme glucose-6-phosphate dehydrogenase (G6PDH) and the free cotinine in the sample for a fixed amount of cotininespecific antibody binding sites. In the absence of cotinine, the cotinine-labelled G6PDH is bound by the cotinine-specific antibody and the enzyme activity is inhibited. The enzyme G6PDH activity is determined spectrophotometrically at 340 nm by measuring its ability to convert nicotinamide adenine dinucleotide (NAD) to reduced NAD (NADH). Acceptable control ranges are established with the selected controls supplied. The negative, 100 ng ml⁻¹, 250 ng ml⁻¹, 500 ng ml⁻¹, 1000 ng ml⁻¹ and 2000 ng ml⁻¹ cotinine calibrators should be used for calibration of the assay. Controls should be used at least once a day to validate the assay performance. The cross-reactivity in the assay was 50% for 3'-hydroxycotinine and 0.2% for theophylline, but no cross-reactivity was reported for other drugs.

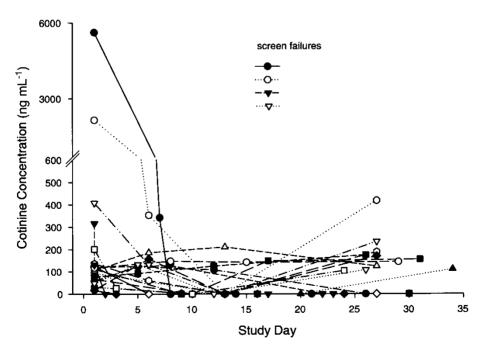
Proper advertising and pre-screening by phone can reduce the potential for screen failures when recruiting non-smokers. European studies 1-6 were performed sequentially. As can be seen in Table 5, the greatest number of screen failures, as well as subjects passing screening questions but determined to be positive for cotinine, occurred in European study 1. After retraining staff on the importance of smoking habits to our studies, the percentage of failures during screening and subsequent testing of cotinine decreased considerably. In a large US trial investigating a drug metabolized by CYP1A2 (US study 2), proper screening of smokers was important. Even then, as shown in Table 5, there was screen failure of about 7% due to no phone screening prior to subjects arriving at the clinic. This reinforces the necessity for proper phone screening and confirmation of smoking status by testing for cotinine. This table re-emphasizes the usefulness of proper screening questions.

The advantage of using calibrators to define a standard curve is that actual concentrations of cotinine can be obtained. Using a single calibrator for a breakpoint will only provide 'yes' or 'no' answers with regard to cotinine; if the clinical study already has a predefined breakpoint, 'yes' and 'no' answers are quicker to obtain.

The cotinine EIA kit insert published in March 1997 (Diagnostic Reagents, Inc.) states, 'The urine cotinine concentration for non-smokers is normally below 500 ng ml⁻¹, as reported by Jarvis' (Jarvis et al. 1987). Urine cotinine levels in non-smokers have been reported to be $< 100 \text{ ng ml}^{-1}$, with the majority of subjects having levels less than 80 ng ml⁻¹ in urine (Biber et al. 1987, Kolonen and Puhakainen 1991, Eremin et al. 1992). Riboli et al. (1995) suggested 150 ng mg⁻¹ creatinine as a cut-off point for tobacco exposure, with 1.5% of alleged non-



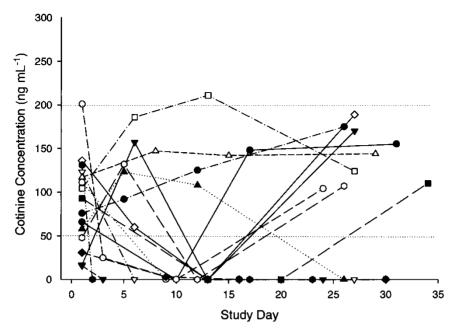
smokers being light smokers in their study involved 1369 women in 10 countries. To be sure there is no enrolment of smokers in phase I studies, Apseloff et al. (1994) suggested a threshold value of 50 ng ml⁻¹. However, this suggestion may eliminate a substantial number of subjects who have no smoking history and who are not exposed to environmental tobacco smoke, as shown by our own research in Figures 1 and 2. Fifteen out of 19 subjects who were not screen failures had values > 50 ng ml⁻¹, and four subjects were screen failures in terms of having urinary cotinine levels of $> 300 \text{ ng ml}^{-1}$ but were allowed to continue in the study. Upon revisiting the screening questions, it was determined that these screen failure subjects either lived with a smoker or travelled on trains where smoking was allowed. In three of the four screen failures the cotinine levels plateaued off to levels < 299 ng ml⁻¹ within 8 days of confinement. The other subject left the facility and returned again after exposure to environmental tobacco smoke. Only two subjects consistently had cotinine concentrations of <50 ng ml⁻¹ throughout the confinement. Excluding the screen failures, only three subjects had cotinine concentrations on day 1 that were > 50 ng ml⁻¹; these fell to < 50 ng ml⁻¹ and stabilized below this breakpoint. Table 5 shows the mean + SD and range of urine cotinine concentrations in non-smokers from five of the studies. Of the 216 subjects from these five studies, only two (0.93%) had cotinine concentrations > 200 ng ml $^{-1}$. Based on these results, a breakpoint separating smokers from non-



···O·· Subject was allowed to leave clinic following day 14 and determined to be exposed to second-hand smoke upon return, as evident from cotinine level.

Figure 1. Urine cotinine concentrations in healthy subjects maintained in a smoke-free study unit for up to 35 days in a supervised, contained, controlled environment.





Urine cotinine concentrations in healthy subjects (with a urinary cotinine < 300 ng ml⁻¹) maintained in a smoke-free study unit for up to 35 days in a supervised, contained, controlled environment.

smokers of 200 ng ml⁻¹ in urine is recommended when utilizing kits such as the cotinine EIA kit for routine screening of potential study volunteers.

Saliva

Levels of cotinine in saliva have been a somewhat useful measure of recent cigarette consumption. Similar terminal half-lives were seen in saliva and plasma, as the clearance and distribution values in saliva were directly proportional to the corresponding values in plasma. Thus, salivary concentrations would give the same information about cotinine disposition in the body as plasma concentrations (Curvall et al. 1990). Saliva cotinine concentrations are about 10-40% higher than in plasma, and can vary depending on stimulation of saliva production (SNRT Subcommittee on Biochemical Verification 2002). Greater accuracy was obtained by correcting for age and a cubic non-linear component (Swan et al. 1993). Passive smokers usually have saliva cotinine concentrations < 5 ng ml⁻¹, but heavy passive exposure can result in saliva cotinine levels ≥ 10 ng ml⁻¹. Levels between 10 and 100 ng ml⁻¹ may result from infrequent active smoking or regular active smoking with a low nicotine intake. Levels greater than 100 ng ml⁻¹ are probably the result of regular active smoking (Etzel 1990). Parazzini et al. (1996) suggested a breakpoint of 10 ng ml⁻¹ to distinguish non-smokers (≤ 10 ng ml⁻¹) from smokers (>10 ng ml⁻¹). The breakpoint used will depend on whether infrequent active smoking or regular smoking with a low nicotine intake as well as the use of saliva stimulants would affect the experimental design under investigation.



Hair

While plasma, serum, urine and saliva cotinine levels can be used to detect recent exposure to tobacco smoking, cotinine in hair may reveal information on long-term cigarette smoking. It may indicate tobacco smoking history in subjects who might deliberately abstain for several days before a sample was taken. Hair analysis is a sensitive marker of chemical exposure over a prolonged period of time because the substance is incorporated into the growing hair shaft. Cotinine within the hair shaft could not be removed by hexane washing and thereby must be extracted for analysis by a strong solvent (Haley and Hoffmann 1985).

Cotinine levels in the hair have been shown to reflect systemic exposure to tobacco (Uematsu 1993, Knight et al. 1998). Eliopoulos et al. (1996) reported a significant correlation between the number of cigarette smoked and measurement of cotinine levels in hair (r = 0.57, p = 0.0008). Cotinine concentrations in hair were also correlated to those in plasma (r = 0.42, p = 0.02). The average hair cotinine levels were 1.18 and 1.95 ng mg⁻¹ hair in smoking women and men, respectively, while the plasma cotinine concentrations were 297 and 330 ng ml $^{-1}$, respectively. A RIA method has a lower limit of sensitivity of 0.1 ng mg⁻¹ hair for cotinine, based on a 2 mg sample of hair (Klein et al. 1994). It has been shown that the average hair cotinine concentrations are 6.3 and 0.3 ng mg⁻¹ hair for smoking and non-smoking mothers, respectively (Eliopoulos et al. 1994, Klein and Koren 1999). This was very close to the results of Haley and Hoffmann (1985), who reported mean cotinine concentrations of 0.5 and 0.12 ng mg⁻¹ hair for smokers and non-smokers, respectively (Haley and Hoffmann 1985). Using a gas chromatographic method, Kintz (1992) reported that cotinine concentrations as low as 0.01 ng mg⁻¹ hair could be detected and quantified. In this study the average hair cotinine concentrations were 2.54 and 0.07 ng mg⁻¹ hair for smoking and non-smoking mothers, respectively. In summary, measurement of hair cotinine is an economic, non-invasive way to detect tobacco smoking. A cotinine concentration of 0.3 ng mg⁻¹ hair may be used as a cut-off to distinguish nonsmokers from smokers.

It has been reported that the hair colour may affect nicotine uptake by the hair. White or fair hair has lower nicotine levels than black hair. In order to minimize inter-individual variability, it is advisable to include hair colour as a covariate in the analysis of hair cotinine (Al-Delaimy 2002).

Other considerations in determining nicotine exposure

Geographical location may be important in studies where smoking status is a concern. In the US there is a heightened awareness of exposure to 'second hand' smoke. In fact, many public facilities do not allow smoking or have restricted areas. In Europe, there currently appears to be a higher prevalence of 'second hand' smoke, even though there is a move towards facilities for non-smokers. However, analysing data from studies involving patients enrolled in US and European trials, the prevalence of smoking is about the same (41% in the US versus 42% in Europe). Specific trial information is proprietary, but these transatlantic studies had the same design. The prevalence of smoking may be higher than in the general



population since the disease population was prone to lifestyle compliance issues. The prevalence of tobacco smoking in many countries around the world, such as Canada, Australia, New Zealand, the UK, Africa, Latin America, many parts of Asia, Norway, Chile, Ecuador, Venezuela, Cuba and France, need to be taken into consideration when providing medical treatment.

Conclusions

Cotinine levels may serve as a useful biomarker of tobacco exposure. Cotinine clearance and elimination half-life are race dependent and these factors must be considered in defining breakpoints.

Potential metabolic interactions via CYP2A6 and CYP1A2 need to be considered when designing a study. From a metabolic standpoint, cessation of smoking should occur at least 14 days prior to the start of the study.

There are well-established assays and kits available for detecting cotinine. The choice of the assay depends on the sensitivity and specificity needed for the research being conducted. As a screening tool, the use of calibrators to define a breakpoint is acceptable. Nicotine from food products is not expected to impact assay sensitivity or be clinically relevant. Thus, dietary restrictions related to nicotine are not considered necessary when determining exposure to environmental tobacco smoke or smoking status.

Generally, blood cotinine measurements are chosen only when venipuncture is being performed for another purpose such as part of clinical chemistry testing. Since urine collection is easier and non-invasive, urinary cotinine is often measured. The investigators' comfort level in defining breakpoints to differentiate smokers from non-smokers depends on assay sensitivity and specificity, and the potential clinical impact on the study. The choice of assay must be balanced with costs and the specificity and sensitivity necessary to assess clinical outcomes.

It is our recommendation that a serum cotinine concentration of 10 ng ml⁻¹ and a urinary cotinine concentration of 200 ng ml⁻¹ be employed as breakpoints to aid in the confirmation of smoking status during study screening periods. Other non-invasive alternatives are salivary cotinine levels (smoker > 10 ng ml⁻¹) and hair cotinine levels (smoker ≥ 0.3 ng mg⁻¹ hair). Since concentrations in urine are substantially greater and thus require a less sensitive assay, urine testing is the best alternative. It is highly recommended that screening questions be accompanied by testing for cotinine where smoking or nicotine-containing products may affect the study outcome. Also, geographical location may affect the ability to effectively recruit non-smokers if required for the study.

In the early phases of drug development, a full understanding of metabolism in humans nor the expected adverse event profile of the drug are not known. Hence, it is critical for most early phase clinical trials to exclude the tobacco-smoking population in order to avoid capturing 'noisy data' that will interfere with the interpretation of pharmacokinetic and drug safety results. The inclusion of smokers in studies should be considered once the impact of smoking on the targeted population is understood.



Acknowledgements

The authors are grateful to Yi Yuan Hu for her contributions to the historic references through literature searches, and to Celine Lese and Jerry Brisson for their support in preparing this manuscript.

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