

Macromolecular adducts and related biomarkers in biomonitoring and epidemiology of complex exposures

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Summary. In order to evaluate the potential of biological markers in epidemiology and risk assessment of complex exposures, we review recent studies of macromolecular adducts and oncogene activation in human populations. Results are discussed in terms of the strengths and weaknesses of various study designs in order to identify the most promising approaches for future research.

Introduction

Environmental carcinogenesis (broadly defined to include both lifestyle and non-lifestyle factors) represents an important area for cancer prevention. In particular, mixtures and multiple exposures pose a particular challenge to researchers since humans are rarely exposed to single agents.

Biological markers have a strong potential to contribute to cancer prevention in a number of ways: by permitting early identification of potential risks (hazards) to humans and by providing comparative molecular dosimetry and response data in both humans and experimental animals, facilitating quantitative risk extrapolation between species. Biological markers can also be incorporated into analytic epidemiological studies in such a way as to increase their ability to identify a causal relationship between exposure to specific agents and increased risk of cancer. Very importantly, biomarkers should significantly expand our knowledge of the mechanisms by which individual carcinogens or mixtures exert their effects. Such information would permit more effective intervention and prevention strategies.

Despite this broad potential, the field is at an early stage of development in which many of the newer laboratory methods are still being validated in the laboratory and in small-scale pilot studies. Although the slow pace may cause epidemiologists and risk-assessors to become impatient, this groundwork is essential to the future use of biological markers and molecular epidemiology. At the very least, recent research has demonstrated that a number of methods are adequately sensitive and reproducible for human studies of environmental carcinogenesis and that the

Table 1. Studies of macromolecular adducts in humans

| Compound analysed | Exposure source | Biological sample | Population | Reference |
|--|--|--|---|--|
| <i>Chemical-specific adducts</i> | | | | |
| <i>N</i> -3-(2-Hydroxyethyl)-histidine; <i>N</i> -(2-hydroxyethyl)valine | Ethylene oxide | RBC | Workers, smokers, unexposed | Calleman <i>et al.</i> , 1978; Van Sittert <i>et al.</i> , 1985; Farmer <i>et al.</i> , 1986; Törnqvist <i>et al.</i> , 1986; Latrino <i>et al.</i> , unpub. |
| Alkylated Hb | Propylene oxide | RBC | Workers | Osterman-Golkar <i>et al.</i> , 1984 |
| 4-Aminobiphenyl-Hb | Cigarette smoke | RBC | Smokers, non-smokers | Bryant <i>et al.</i> , 1987 |
| Styrene-Hb | Styrene | RBC | Workers | Brenner <i>et al.</i> , unpub. |
| AFB ₁ -guanine | Diet | Urine | Chinese and Kenyans residing in a high exposure and high and low risk area respectively | Groopman <i>et al.</i> , 1985; Autrup <i>et al.</i> , 1983 |
| 3-Methyladenine | Methylating agents | Urine | Unexposed | Shuker & Farmer, 1988 |
| PAH-DNA | PAH in cigarette smoke, in workplace | WBC, lung tissue, placenta | Lung cancer patients, smokers, workers | Perera <i>et al.</i> , 1989b; Shamsuddin <i>et al.</i> , 1985; Vähäkangas <i>et al.</i> , 1985; Harris <i>et al.</i> , 1985; Haugen <i>et al.</i> , 1986; Everson <i>et al.</i> , 1986, 1988; Perera <i>et al.</i> , 1987, 1988a; Garner <i>et al.</i> , 1988; Weston <i>et al.</i> , 1988; Brenner <i>et al.</i> , unpub. |
| Antibodies to PAH-DNA | PAH in workplace | Serum | Workers | Haugen <i>et al.</i> , 1986 |
| <i>O</i> ⁶ -Methyldeoxy-guanosine | Nitrosamines in diet | Oesophageal and stomach mucosa | Chinese and European cancer patients | Umbenhauer <i>et al.</i> , 1985; Wild <i>et al.</i> , 1986 |
| Cisplatinum-DNA, Cisplatinum-protein | Cisplatin | WBC, Hb, plasma proteins | Chemotherapy patients | Reed <i>et al.</i> , 1986; Fichtinger-Schepman <i>et al.</i> , 1987; Mustonen <i>et al.</i> , 1988; Den Engelse <i>et al.</i> , 1988; Perera <i>et al.</i> , 1989a |
| 8-Methoxypsoralen-DNA | 8-Methoxy-psoralen | Skin | Psoriasis patients | Santella <i>et al.</i> , 1988 |
| Styrene oxide-DNA | Styrene | WBC | Workers | Liu <i>et al.</i> , 1988 |
| <i>Non-chemical-specific adducts</i> | | | | |
| Multiple carcinogens by ³² P-postlabelling | Betel and tobacco chewing, smoking, industrial exposures, wood smoke | Placenta, lung tissue, oral mucosa, WBC, bone marrow, colonic mucosa | Smokers, workers, volunteers | Everson <i>et al.</i> , 1986; Dunn & Stich, 1986; Chacko & Gupta, 1987; Phillips <i>et al.</i> , 1986, 1988a, b; Reddy & Randerath, 1987 |

Source: Perera (1989)

Abbreviations: PAH, polycyclic aromatic hydrocarbon; RBC, red blood cells; WBC, white blood cells; Hb, haemoglobin, AFB, aflatoxin B.

Table 2. Biological markers in smokers and non-smokers

| Marker | Smokers | Non-smokers |
|-----------------------------|--------------------------------|------------------------------|
| 4-Aminobiphenyl-Hb (pg/g) | 154.5 [11.3*] <i>n</i> = 19 | 32.2 [2.9] <i>n</i> = 18 |
| SCE (average per metaphase) | 10.8 [0.603†] <i>n</i> = 11 | 8.1 [0.47] <i>n</i> = 10 |
| Cotinine (ng/ml) | 419.2 [47.4*] <i>n</i> = 10 | 0.3 [0.032] <i>n</i> = 10 |

Values are means [S.E.]

**p* = 0.0001

†*p* = 0.002

Source: Perera *et al.* (1987)

overall approach is feasible. The experience gained to date also allows some recommendations as to the optimal design of future molecular epidemiological studies.

This paper reviews recent developments in the use of macromolecular adducts and complementary markers in biomonitoring and molecular epidemiology, highlighting those which are most pertinent to the evaluation of carcinogenic risks of complex mixtures. Three types of exposures will be evaluated: lifestyle-related (cigarette smoking); occupational; and clinical. The common denominator for the first two is exposure to multiple polycyclic aromatic hydrocarbons (PAHs) and aromatic amines. Because study design dictates the interpretation of results, examples of cross-sectional, longitudinal serial sampling, retrospective and nested case-control studies will be described, with a brief discussion of the advantages and disadvantages of each approach.

As shown in Table 1, considerable research has been directed towards validating macromolecular adducts as markers of the biologically effective dose of carcinogens. Carcinogen-DNA adducts resulting from covalent interaction of electrophilic carcinogens with DNA are considered to be a necessary although not sufficient event in the initiation of chemical carcinogenesis; in certain instances carcinogen-protein

Table 3. Correlation between 4-aminobiphenyl-Hb and ethylene oxide-Hb adducts in smokers and non-smokers

| | Smokers (13) | Non-smokers (7) |
|-------------------------------|-------------------------|----------------------|
| Ethylene oxide-Hb (nmol/mol) | 238 [29.36] (75-393) | 18 [3.04] (10-40) |
| 4-Aminobiphenyl-Hb (pmol/mol) | 54 [3.88] (30-68) | 11 [1.39] (4-7) |

Values are means [S.E.], with range in parentheses

r = 0.83 smokers and non-smokers, *p* < 0.005

r = 0.50 smokers only, *p* < 0.005

Source: Latriano *et al.* (unpublished)

adducts can satisfactorily serve as a surrogate for DNA binding (Miller & Miller, 1981; Weinstein *et al.*, 1984; Wogan & Gorelick, 1985; Harris, 1985; Farmer *et al.*, 1986; Yuspa & Poirier, 1989).

Cross-sectional studies

Most of the research on macromolecular adducts has been cross-sectional in nature, aimed at elucidating the relationship between carcinogen-DNA or carcinogen-protein adducts and estimated exposure to carcinogens. In most cases, multiple exposures or complex mixtures (for example, cigarette smoke, diet, industrial pollution and combination chemotherapy) have been involved.

Cigarette smoke is a classical complex mixture which has been the subject of much molecular research. For example, in a study of 22 smoking and 24 non-smoking volunteers, a battery of markers was evaluated in repeat peripheral blood samples drawn several days apart (Perera *et al.*, 1987) (see Table 2). Environmental histories were obtained by questionnaire. 4-Aminobiphenyl (4-ABP)-haemoglobin (Hb) adducts were found to be more specific to cigarette smoke than DNA adducts of polycyclic aromatic hydrocarbons (PAHs). Unlike PAH-DNA adducts, for which there was a high and variable background, 4-ABP-Hb levels were significantly different ($p = 0.0001$) between smokers and non-smokers. A repeat blood sample (two days later) gave comparable results for 4-ABP-Hb: 139.0 (S.E. 7.9) pg/g for the smokers versus 36.2 (S.E. 5.02) for the non-smokers. Sister chromatid exchanges were also significantly elevated in the smokers and were positively correlated with 4-ABP-Hb. This finding suggests that aromatic amines contribute significantly to the integrated genotoxic effect of cigarette smoke.

Recently samples from a subset of the original group were analysed for ethylene oxide-haemoglobin (EtO-Hb) adducts by the procedure of Törnqvist *et al.* (1986). The levels of these (hydroxyethylvaline) adducts were significantly higher in smokers ($n = 13$) than in non-smokers ($n = 7$) (Latrino, *et al.* unpublished). Ethylene oxide and 4-aminobiphenyl adduct levels were highly correlated ($r = 0.83$) in the smokers and non-smokers combined and in the smokers as a group ($r = 0.50$, $p < 0.005$) (see Table 3). The measured levels of ethylene oxide-haemoglobin adducts were approximately 10-fold higher than those seen by Törnqvist *et al.* (1986) in smokers and non-smokers, possibly as a result of the different standards used (here hydroxyethylvaline instead of ethylene oxide-modified haemoglobin).

This study design has the advantage of simplicity. However, had the differences between the two groups (smokers and non-smokers) been smaller, they might have been missed because of swamping by interindividual variability. For example, a 30% coefficient of variation was seen in levels of 4-ABP-Hb among the smokers, all of whom smoked between one and two packs per day (mean, 1.4 ± 0.4 packs per day).

In contrast to the smoker/non-smoker study, a cross-sectional study of PAH-DNA adducts in long-term employees in an iron foundry in Finland showed a clear dose-response relationship between estimated source-specific exposure to PAHs and levels of adducts ($p = 0.0001$) after adjusting for cigarette smoking and time since vacation (Perera *et al.*, 1988a). Workers were classified as having high exposure

Table 4. PAH-DNA adducts and serum oncogene protein expression in foundry workers and unexposed controls

| | | | | | | Serum oncogene proteins ^a | | | | | |
|---|---------|-----------------------------------|---|--------|--------|--------------------------------------|--------|------------|---------------|--------|--------|
| Ambient exposure ($\mu\text{g BP}/\text{m}^3$) | Subject | Daily cigarette consumption | PAH-DNA adduct levels ^a (fmol/ μg) | | | <i>fes</i> | | <i>ras</i> | | | |
| | | | Post-vacation | Work 1 | Work 2 | Post-vacation | Work 1 | Work 2 | Post-vacation | Work 1 | Work 2 |
| >0.2 (high exposure) | 1 | 0 | 0.11 | 0.8 | NA | — | — | | — | — | |
| | 2 | 20 | 0.13 | 2.0 | NA | + | + | | — | — | |
| | 3 | 0 | NA | 2.8 | NA | — | — | | — | — | |
| 0.05–0.2 (medium exposure) | 4 | 15 | 0.13 | 0.36 | NA | — | — | | — | — | |
| | 5 | 20–25 | NA | 0.32 | 0.42 | — | — | — | — | + | — |
| | 6 | 15 | NA | 0.8 | 0.5 | — | — | — | — | — | — |
| | 7 | 0 | 0.48 | 1.28 | NA | + | + | | — | — | |
| | 8 | 20 | ND | 0.4 | NA | — | — | | — | — | |
| 0.05 (low exposure) | 9 | 20 | ND | 0.38 | | | | | | | |
| | 10 | | ND | 1.52 | | | | | | | |
| | 11 | | ND | 0.32 | | | | | | | |
| | 12 | | 0.08 | 1.20 | | | | | | | |
| 0 (unexposed controls) | 9 | 15 | | 0.08 | | | — | | | — | |
| | 10 | 10 | | ND | | | — | | | — | |
| | 11 | 0 | | 0.14 | | | — | | | — | |
| | 12 | 0 | | ND | | | — | | | — | |
| | 13 | 0 | | ND | | | — | | | — | |
| | 14 | 0 | | 0.3 | | | — | | | — | |
| | 15 | 0 | | ND | | | — | | | — | |
| | 16 | 10 | | ND | | | — | | | — | |
| 17 | 15 | | 0.1 | | | — | | | — | | |
| 18 | 0 | | 0.1 | | | — | | | — | | |

NA, not available; ND, non-detectable

^a Post-vacation samples were drawn immediately after a one-month vacation; work 1 samples were drawn six weeks after return to work; work 2 samples were drawn more than two months after return to work.

Source: Brandt-Rauf et al. (1989a)

to benzo[a]pyrene (BP) as a representative PAH (8 h time-weighted average greater than $0.2 \mu\text{g}/\text{m}^3$), medium exposure ($0.05\text{--}0.2 \mu\text{g}/\text{m}^3$) or low exposure (less than $0.05 \mu\text{g}/\text{m}^3$). However, here also significant interindividual variation was seen in adduct levels in the exposed workers. For example, in 13 foundry workers with comparable medium exposure ($0.05\text{--}0.2 \mu\text{g}/\text{m}^3$ BP) there was a 20-fold range of PAH-DNA adducts ($0.10\text{--}2 \text{ fmol}/\mu\text{g}$). Among 18 workers with low exposure ($<0.05 \mu\text{g}/\text{m}^3$ BP) there was a 29-fold range ($0.030\text{--}0.86 \text{ fmol}/\mu\text{g}$).

Recently, we have investigated whether levels of PAH-DNA adducts (as a marker of biologically effective dose) correlated with activation of oncogenes (as a marker of biological effect), since various PAHs have been shown experimentally to activate the *ras* oncogene (Balmain & Pragnell, 1983; Marshall *et al.*, 1984). For this purpose, we assayed sera from foundry workers and controls, using a modified western blotting technique, for oncogene protein products (*ras*, *fes*, *myc*, *myb*, *sis*, *B-EGF*, *int-1*, *myb*, *src* and *mos*) as previously described (Brandt-Rauf *et al.*, 1989b). The method was able to reveal whether the oncogene was overexpressed, but not whether it had undergone a point mutation. A five-fold increase in protein expression was considered a positive result. As shown in Table 4, one individual in the medium exposure group showed elevated levels of the *ras* oncogene product in one of three serum samples tested. Repeat serum samples from two other individuals in the medium- and high-exposure groups had significantly elevated levels of the *fes* oncogene protein product. Samples from the 10 unexposed controls were uniformly negative. Both *fes* and *ras* serum proteins have previously been detected in lung cancer patients and cigarette smokers (see Brandt-Rauf *et al.*, 1989a, for review). Since foundry workers are also at elevated risk of lung cancer (IARC, 1984), these results suggest that oncogene activation may prove to be a useful early marker for respiratory cancer related to PAH exposure. Obviously, this clue must be followed up by further studies. Two of the three individuals with elevated serum levels of oncogene protein products were smokers of approximately one pack of cigarettes per day. However, it is likely that the major contribution of PAHs to their body burden was from workplace exposure, since daily intake of BP resulting from exposure to $0.2 \mu\text{g}/\text{m}^3$ in the foundry is comparable to BP intake from smoking 5–7 packs per day.

Longitudinal studies

As with all cross-sectional studies, which provide a 'snapshot' at a particular point in time, this research cannot establish temporal or causal relationships between exposure, adduct formation and oncogene activation. For this reason, longitudinal studies in individuals whose exposure changes significantly over time are preferable in terms of establishing the relationship between exposure and biological markers such as macromolecular adducts. Few such studies have yet been conducted, but several 'natural experiments' have occurred such as studies of individuals in smoking cessation programmes, workers sampled before beginning employment in a coke-oven plant (Vähäkangas *et al.*, this volume), workers sampled after an interruption

Table 5. Levels of biological markers in subjects in a smoking cessation programme

| | Stoppers | | | Non-stoppers | | |
|----------------------------|----------------------|---------------------|---------------------|----------------------|----------------------|----------------------|
| | Sample 1 | Sample 2 | Sample 3 | Sample 1 | Sample 2 | Sample 3 |
| PAH-DNA (fmol/ μ g) | 0.30 [0.095] (4) | — | 0.19 [0.009] (4) | 0.42 [0.07] (8) | — | 0.21 [0.04] (8) |
| SCE (ave. met.) | 10.94 [0.71] (8) | 10.82 [0.35] (8) | 9.65 [0.67†] (8) | 12.57 [0.77] (10) | 12.81 [0.55] (10) | 12.23 [1.15] (10) |
| 4-ABP-Hb (pg/g) | 117.3 [30.85] (4) | 74.8 [15.9] (4) | 28.6 [7.73] (2) | 140.8 [15.71] (6) | 116.3 [26.78] (3) | — |
| Cotinine (ng/ml) | 194 [77.55] (4) | 1.75 [0.65] (4) | — | 255.5 [31.6] (4) | 163 [158.44] (2) | — |

Sample 1 was taken at enrolment into the programme, sample 2 after 21 days and sample 3 at 3–6 months. Values are means [S.E.] with number of samples in parentheses.

† $p = 0.03$, Wilcoxon test

Source: Brenner *et al.* (unpublished)

in exposure (Haugen *et al.*, 1986) and chemotherapy patients from whom pretreatment and serial post-treatment samples have been obtained (Reed *et al.*, 1986; Fichtinger-Schepman *et al.*, 1987; Mustonen *et al.*, 1988; Perera *et al.*, 1989a).

Recently, we carried out a pilot study of smokers in a smoking cessation programme, obtaining repeat blood samples from 21 subjects upon enrolment into the programme, and then 21 days and 3–6 months after enrolment. Roughly half of the individuals quit smoking entirely within three weeks ('stoppers') and the other half either reduced smoking or continued to smoke at the same level ('non-stoppers'). As shown in Table 5, SCEs were significantly decreased in the stoppers (but not the non-stoppers) 3–6 months after quitting, while levels of 4-ABP-Hb and cotinine fell sharply within 21 days. However, PAH-DNA adducts were not significantly affected by smoking cessation, possibly because of their lack of specificity for cigarette smoke constituents and/or the short follow-up.

To build upon these results, we are beginning a more definitive study of similar subjects who will be followed for two years after quitting to evaluate persistence of DNA and haemoglobin adducts and oncogene protein products. We shall also evaluate several potential genetic/metabolic markers of susceptibility, such as aryl-hydrocarbon hydroxylase (AHH) activity and glutathione-S-transferase activity. This approach will allow detailed evaluation of the persistence of biological markers in human peripheral blood cells.

Another natural experiment has been provided by the annual vacation taken by Finnish foundry workers during the month of July. As shown in Table 4, white blood cell DNA from 12 individuals was assayed for PAH-DNA adducts after the workers returned from their four-week vacation. Nine of these subjects were sampled six weeks after their return to work. Four individuals were plant workers with low

exposure to PAH. The levels of adducts were significantly higher (0.92 ± 0.61 fmol/ μ g; $p = 0.004$, Wilcoxon test) in the post-work sample compared to the post-vacation sample (0.12 ± 0.14 fmol/ μ g), showing a clear biological effect of exposure.

Another longitudinal, serial sampling study has evaluated biological markers in patients treated with cisplatin-based chemotherapy: cisplatin (*cis*DDP) given in combination with Velban, mitomycin C, VP-16, adriamycin and/or 5-fluorouracil. Peripheral blood samples were collected just before and after the first, second and last cycles of chemotherapy and were assayed as shown in Table 6. In subjects with a baseline and at least two post-treatment samples, levels of SCE, plasma protein binding and haemoglobin-binding measured by atomic absorption spectrometry were significantly increased following treatment ($p < 0.01$). These results were consistent with prior studies of DNA and protein adducts and SCE in cisplatin-treated patients (Reed *et al.*, 1986; R. Mustonen *et al.*, unpublished; Fichtinger-Schepman *et al.*, 1987). We have recently begun a follow-up study which will assess a full battery of markers (*cis*DDP-DNA and *cis*DDP-haemoglobin adducts, SCE and micronuclei, chromosomal aberrations, gene mutation (HPRT and GPA) and oncogene activation in peripheral blood cells and/or semen). This study will allow the effect of *cis*DDP to be separated from that of other genotoxicants administered, since adduct measurements are specific to *cis*DDP but the other markers will give an integrated measure of total genetic toxicity of chemotherapy. Correlation between *cis*DDP adducts and the non-specific markers will be particularly sought.

To summarize, the advantage of longitudinal serial sampling studies such as those described above is that the problem of interindividual variation is mitigated by having each individual serve as his or her own control. In addition, in the foundry workers study, all individuals were sampled on the same days so that possible seasonal variation in binding was controlled for. However, such research opportunities are rare, and although they address the temporal relationship between exposure and biomarkers, they leave unanswered the question of whether biomarkers are related to cancer risk.

Case-control studies

The first step in exploring the role of various biomarkers as risk factors for cancer has been taken in several modified case-control studies. These have a practical advantage over prospective studies in terms of feasibility and can provide circumstantial evidence of an association between biomarkers and cancer risk. However, as with cross-sectional studies, they cannot establish the causal sequence between adduct formation, gene mutation, oncogene activation and cancer. Moreover, the latency from exposure to cancer induction is usually years to decades and biological markers formed during exposure are likely to have been lost or diluted out by the time cancer is diagnosed. Unless exposure has been continuous and unchanged and metabolic processes have not altered over time, biological measurements made now are not relevant to present risk of cancer.

We have recently evaluated 81 cases with primary carcinoma of the lung and 67 controls for levels of PAH-DNA adducts and SCEs in order to determine whether,

Table 6. Levels of biomarkers in cisplatin-treated patients by time of sample collection

| | Baseline ^a | Post Cycle 1 ^b | Pre Cycle 2 ^a | Post Cycle 2 ^c | Post Final Cycle ^c |
|---|-----------------------|---------------------------|--------------------------|---------------------------|-------------------------------|
| Cumulative <i>cis</i> DPP dose (mg/m ²) | 0 | 99.23 ± 13.24 | 105.45 ± 28.41 | 197.00 ± 35.82 | 351.9 ± 60 |
| SCE | 10.80 ± 3.01 (12) | 17.39 ± 6.52* (12) | 17.45 ± 5.03* (12) | — | — |
| Plasma protein binding | 0.47 ± 1.07 (10) | 27.50 ± 9.34* (10) | 2.37 ± 3.34 (2) | 33.37 ± 11.07* (10) | 33.49 ± 9.77* (10) |
| Haemoglobin binding | 0.01 ± 0.04 (11) | 2.08 ± 1.04* (11) | | 2.97 ± 1.25* (11) | 2.89 ± 1.04 (4) |

* $p < 0.01$ (Wilcoxon and *t*-tests)^d

Values are means ± SD (standard deviation). Only subjects with measurements of baseline and two post-treatment samples are included; numbers of sample are given in parentheses.

^a Samples drawn within 24 h before treatment

^b SCE sample drawn 2 weeks after treatment; protein sample drawn 12–24 h post-treatment

^c SCE and/or protein sample drawn 12–24 h post-treatment

^d Analysis of difference between baseline and respective post-treatment values; both tests were two-tailed

Source: Perera et al. (1989a)

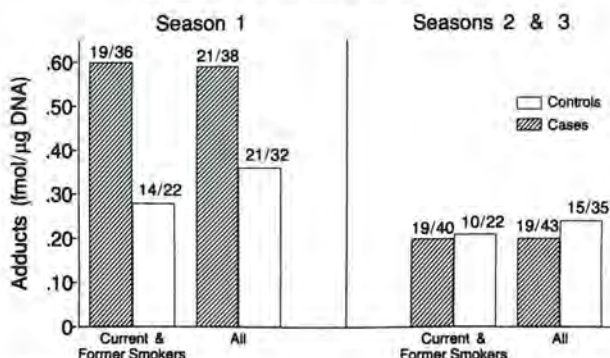


Figure 1. Polycyclic aromatic hydrocarbon-DNA adducts in white blood cells of lung cancer cases and controls (current smokers and former smokers; all subjects)

Values plotted are the means for positive samples; numbers above the bars represent the ratio of positive samples to all samples assayed. In Season 1 samples, the mean adduct level in lung cancer patients (former and current smokers combined) exceeded that in former and current smoker controls ($p = 0.02$, Wilcoxon test).

when exposure to PAHs and other mutagens or carcinogens was comparable, cases had higher levels of biomarkers than controls (Perera *et al.*, 1989b). This would suggest that the ability to efficiently activate and bind carcinogens may have been a risk factor in their disease. SCEs in lymphocytes did not differ between cases and controls. As in the smokers/non-smokers study, PAH-DNA adducts were not specific to cigarette smoking but apparently reflected the numerous background sources of PAHs such as diet and ambient air (Table 7). Among current smokers, white blood cell DNA from lung cancer cases had significantly higher levels of PAH-DNA adducts ($p = 0.01$, Wilcoxon test). As shown in Figure 1, this case-control difference was contributed by blood samples collected during the months July-October. This seasonal effect is consistent with the observation of seasonal variation in AHH activity in human lymphocytes (Paigen *et al.*, 1981). Genetically regulated metabolism of PAHs by the AHH microsomal enzyme system has been linked (in some but not all studies reported) to lung cancer risk (Kouri *et al.*, 1982; Kärki *et al.*, 1987).

As in the foundry workers study, we used western blotting techniques to screen sera of lung cancer cases and controls for peptide sequences of nine different proteins including *ras*, *fes* and *myc* (Brandt-Rauf *et al.*, 1989a). All 18 cases were positive for at least one oncogene protein, in contrast to two of the 20 healthy controls (Table 7). In 15 of these cases, all but one of which were non-squamous cell cancers, *ras* gene protein products were identified. Eleven of the 18 cases had activated *fes* compared to none of the controls. All 18 lung cancer cases were current or former cigarette smokers with impressive smoking histories (up to 105 pack years). These results are consistent with prior studies in that PAHs are known to activate the *ras* gene and a high prevalence of activated *ras* has been seen in non-small cell carcinomas and in

Table 7. DNA adducts and serum oncogene proteins in lung cancer patients

| Patient | Age | Sex | Smoking history (pack-years) | Tumour type | Lymphocyte adducts (fmol/μg) | Oncogene protein | | | | | | | | |
|-----------------|-----|-----|---------------------------------|----------------|---------------------------------|-------------------------|------------|------------|------------|------------|------------|------------|------------|-----|
| | | | | | | <i>ras</i> | <i>fes</i> | <i>myb</i> | <i>int</i> | <i>sis</i> | <i>myc</i> | <i>mos</i> | <i>src</i> | EGF |
| 1 | 75 | F | S(NA) | A | — | ++ | — | + | + | — | — | — | — | — |
| 2 | 71 | F | S(90) | A(S) | — | ++ | ++ | — | — | — | — | — | — | — |
| 3 | 67 | F | S(31) | A | — | ++ | — | — | — | — | — | — | — | — |
| 4 | 42 | M | S(20) | L | — | ++ | — | — | — | — | — | — | — | — |
| 5 | 67 | F | S(40) | A | — | ++ | — | — | — | — | — | — | — | — |
| 6 | 82 | M | S(55) | A | +(0.45) | ++ | — | — | — | — | — | — | — | — |
| 7 | 49 | M | S(63) | S | — | ++ | — | — | — | — | — | — | — | — |
| 8 | 65 | M | S(40) | L | +(0.16) | ++ | ++ | + | — | + | — | — | — | — |
| 9 | 68 | F | S(62) | A | — | ++ | ++ | — | — | — | — | — | — | — |
| 10 | 62 | M | S(61) | S | — | — | ++ | — | — | — | — | — | — | — |
| 11 | 69 | F | S(40) | A | — | — | ++ | — | — | — | — | — | — | — |
| 12 | 73 | F | S(84) | S | — | — | — | — | + | — | — | — | — | — |
| 13 | 67 | F | S(105) | A | — | ++ | ++ | — | — | — | — | — | — | — |
| 14 | 67 | F | S(13) | L | — | ++ | ++ | — | — | — | — | — | — | — |
| 15 | 60 | F | S(20) | L | — | ++ | — | — | — | — | — | — | — | — |
| 16 | 63 | M | S(84) | A | — | ++ | ++ | — | — | — | — | — | — | — |
| 17 | 69 | M | S(50) | A | +(0.13) | ++ | ++ | — | — | — | — | — | — | — |
| 18 | 62 | M | S(30) | A | — | ++ | ++ | — | — | — | — | — | — | — |
| <i>Controls</i> | | | | | | | | | | | | | | |
| 19 | 58 | M | NS | | | — | | | | | | | | |
| 20 | 25 | M | NS | | | — | | | | | | | | |
| 21 | 30 | M | ES | | | — | | | | | | | | |
| 22 | 31 | M | CS | | | + [All others negative] | | | | | | | | |
| 23 | 52 | M | ES | | | + | | | | | | | | |
| 24 | 39 | M | CS | | | — | | | | | | | | |
| 25 | 29 | M | NS | | | — | | | | | | | | |
| 26 | 34 | M | CS | | | — | | | | | | | | |
| 27 | 26 | M | NS | | | — | | | | | | | | |
| 28 | 35 | M | CS | | | — | | | | | | | | |
| 29 | 35 | M | NS | | | — | | | | | | | | |
| 30 | 52 | M | CS | | | — | | | | | | | | |
| 31 | 36 | M | NS | | | — | | | | | | | | |
| 32 | 56 | M | NS | | | — | | | | | | | | |
| 33 | 56 | M | NS | | | — | | | | | | | | |
| 34 | 32 | M | NS | | | — | | | | | | | | |
| 35 | 48 | M | CS | | | — | | | | | | | | |
| 36 | 44 | M | NS | | | — | | | | | | | | |

S, smoker; NS, non-smoker; CS, current smoker; ES, ex-smoker
 ++, strongly positive; +, positive; —, negative

A, adeno; AS, adenosquamous; L, large cell; S, squamous
 Source: Brandt-Rauf *et al.* (1989a)

tumours from cigarette smokers. High levels of *fos* expression have also been detected in human lung tumours (reviewed in Brandt-Rauf *et al.*, 1989a).

PAH-DNA adducts and SCEs were measured in white blood cell DNA from these 18 patients. Three of these individuals had detectable levels of adducts, and all three were positive for the *ras* oncoprotein, but 11 of the *ras*-positive cases did not have detectable levels of adducts. Thus, there was no apparent correlation between adducts and oncogene activation, possibly because of the small number of subjects studied.

Nested case-control studies

As noted above, case-control studies have the serious limitation that if exposure or pharmacokinetic processes have altered between the time of cancer induction and diagnosis, current levels of macromolecular adduct levels may bear little relationship to cancer risk. Thus, the nested case-control is the optimal design for evaluating the relationship between biological markers and cancer risk, provided, of course, that the marker is stable in samples collected at the outset of a prospective study. In this approach, the cohort is followed and when cancer cases appear among the cohort, their stored biological samples are analysed and compared with those from one or more matched controls. This allows calculation of the relative risk, i.e., the measure of association between the incidence rates of cancer and the biomarker(s) being evaluated. Examples include a recent study of serum retinol and retinol binding protein and lung cancer (Friedman *et al.*, 1986) and an ongoing study of hepatitis B chronic carriers involving retrospective measurement of aflatoxin B₁ adducts on albumin and other markers once incident hepatocellular carcinoma cases are identified (Bosch & Muñoz, 1988).

Chemotherapy patients provide a valuable model for determining the predictive value of biomarkers to risk of cancer, since they may be at risk of second malignancy resulting from genotoxic chemotherapy. A multi-centre nested case-control study storing baseline and serial samples from cancer patients treated with cisplatinum-based chemotherapy would be both a practical and effective approach to marker validation.

Conclusions

Returning to the assessment of complex mixtures and multiple exposures, in light of the above discussion, our recommendations are as follows:

- (1) Chemical-specific dosimeters (DNA and protein adducts) should be used in conjunction with non-specific markers such as SCE, micronuclei, gene mutation and oncogene activation which give an integrated estimate of the genotoxic/procarcinogenic effect of exposure.

- (2) In order to separate the effect of specific constituents of the mixture, a step-wise evaluation can be carried out, as illustrated by the reverse pyramid in Figure 2. First, external exposure is characterized as completely as possible through ambient or personal monitoring and questionnaires. This provides an estimate of the

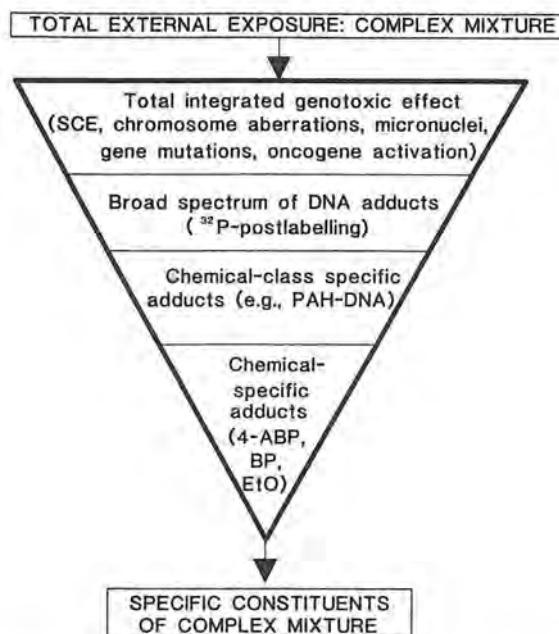


Figure 2. Assessment of the potential carcinogenic risk of complex mixtures

level and pattern of exposure both to the mixture and to its individual components. The next step is to analyse the relationship between integrated and chemical-specific exposure variables on the one hand and (a) total genotoxic/procarcinogenic effect, (b) the broad spectrum of DNA adducts by the postlabelling assay, (c) class-specific adducts (e.g., PAHs by immunoassay) and (d) individual chemical-specific adducts (e.g., 4-ABP-Hb or BP-DNA). Correlations between biological markers are also examined. The main question of interest is what proportion of the total genotoxic/procarcinogenic effect of exposure to complex mixtures is attributable to specific constituents in the mixture. Also, is there apparent interaction between individual constituents or do the effects appear to be additive? The answers to these questions may allow identification of the major pathogenic agents present in a chemical mixture.

(3) The most informative designs for identifying the relationship between exposure to complex mixtures and markers of biological dose and effect are serial sampling studies, which provide measurements at baseline and at several subsequent times after exposure has either ceased (e.g., smokers giving up the habit) or begun (e.g., newly appointed workers).

(4) To evaluate the role of complex mixtures in cancer risk, the nested case-control study is the optimal design. Suitable exposures for study include cigarette smoking, occupational exposures to fossil-fuel combustion products and combination chemotherapy.

In conclusion, the dramatic progress in the molecular biology of cancer during

the last five years has provided the impetus and the methods for a new, more powerful approach to cancer epidemiology and risk assessment. Provided the necessary groundwork is carried out, biological markers should become a valuable tool in cancer prevention.

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