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Interaction of benzo[a]pyrene diol epoxide isomers with human serum albumin: Site specific characterisation of adducts and associated kinetics

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Carcinogenicity of benzo[a]pyrene {B[a]P, a polycyclic aromatic hydrocarbon (PAH)} involves DNA-modification by B[a]P diol epoxide (BPDE) metabolites. Adducts to serum albumin (SA) are not repaired, unlike DNA adducts, and therefore considered advantageous in assessment of *in vivo* dose of BPDEs. In the present work, kinetic experiments were performed in relation to the dose (i.e. concentration over time) of different BPDE isomers, where human SA (hSA) was incubated with respective BPDEs under physiological conditions. A liquid chromatography (LC) tandem mass spectrometry methodology was employed for characterising respective BPDE-adducts at histidine and lysine. This strategy allowed to structurally distinguish between the adducts from racemic *anti*- and *syn*-BPDE and between (+)- and (–)-*anti*-BPDE, which has not been attained earlier. The adduct levels quantified by LC-UV and the estimated rate of disappearance of BPDEs in presence of hSA gave an insight into the reactivity of the diol epoxides towards the N-sites on SA. The structure specific method and dosimetry described in this work could be used for accurate estimation of *in vivo* dose of the BPDEs following exposure to B[a]P, primarily in dose response studies of genotoxicity, e.g. in mice, to aid in quantitative risk assessment of PAHs.

Polycyclic aromatic hydrocarbons (PAHs), several of which are carcinogens, are formed during incomplete combustion of organic matters and are known to be ubiquitous in the environment. Exposure of the general population to PAHs occurs from food and other sources such as cigarette smoke, vehicle exhausts and domestic wood burning^{1–3}. Food can be contaminated by PAHs from the environment or during preparation, e.g. when smoked or grilled^{4,5}.

Benzo[a]pyrene (B[a]P), classified as a human carcinogen (IARC 2012)², has widely been used as an indicator of carcinogenic PAHs, and is considered one of the most prevalent and carcinogenic of PAHs³. Metabolism of B[a]P gives rise to the electrophilic reactive B[a]P diol epoxides (BPDEs, Fig. 1)^{6,7}: (+)- and (–)-*anti*-7β, 8α-dihydroxy-9α,10α-epoxy-7,8,9,10-tetrahydrobenzo[a]pyrene (*anti*-BPDE), and (+)- and (–)-*syn*-7β, 8α-dihydroxy-9β,10β-epoxy-7,8,9,10-tetrahydrobenzo[a]pyrene (*syn*-BPDE). The diol epoxides are the causative factors responsible for DNA damage from B[a]P-exposure, and amongst them the most potent mutagenic and carcinogenic metabolite has been identified as (+)-*anti*-BPDE⁸. The BPDE stereoisomers covalently bind with varying degree of reactivity to DNA, e.g. to exocyclic amino groups on deoxyguanosine, resulting in formation of DNA adducts⁷.

Adducts formed through covalent binding of low molecular weight electrophilic compounds to nucleophilic sites on blood proteins are often used as a surrogate biomarker for the corresponding DNA adducts⁹. Analysis of adducts to haemoglobin (Hb) and serum albumin (SA) have, compared to DNA adducts, advantages of high abundance of the proteins. Unlike DNA, damage to the protein is not repaired and hence protein adducts have an inherent advantage as a measure for human exposure, e.g. in epidemiologic studies^{9,10}. Hb adducts have been

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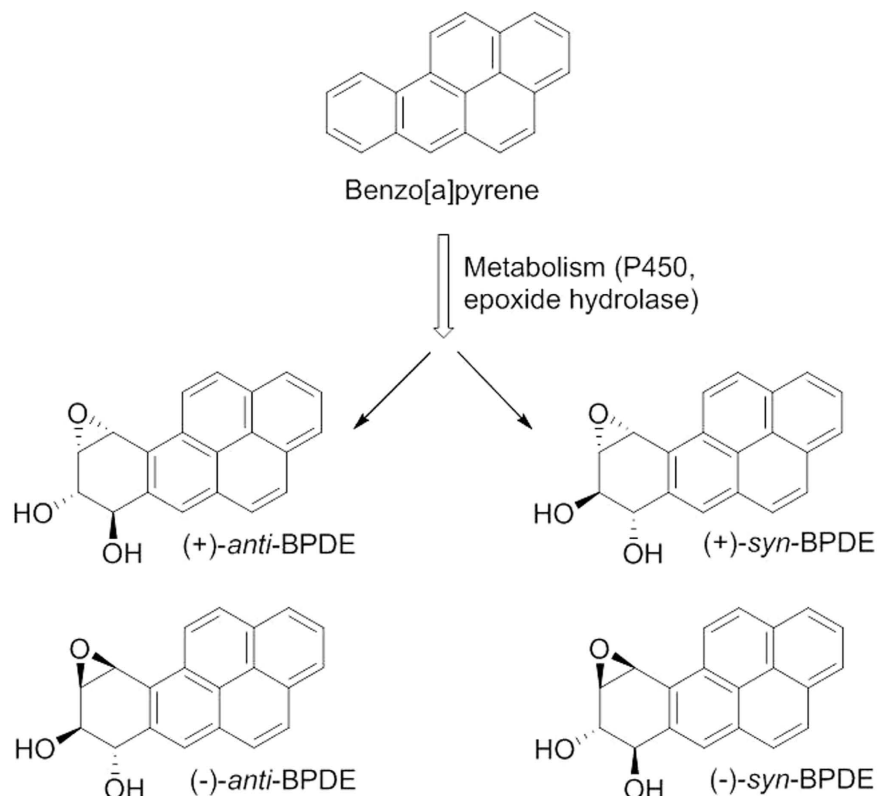


Figure 1. Benzo[a]pyrene and its four diol epoxide metabolites, the enantiomers (+)-*anti*-, (-)-*anti*-, (+)-*syn*- and (-)-*syn*-BPDE.

used for measurement of dose of the precursor electrophile in blood, i.e. referred as the *in vivo* dose or “the area under the concentration-time curve” (AUC), which can be used as a basis in cancer risk assessment^{11–13}. However, with regard to adducts from diol epoxides of PAHs there is no sufficiently established method for protein adduct measurements that can be used for accurate measurement of the *in vivo* dose.

Albumin is the most abundant protein in serum. Albumin, possessing large hydrophobic pockets, is involved in the transport of various endogenous substrates as well as lipophilic xenobiotics^{14,15}. SA adducts of BPDEs have been studied with different methods (reviewed in Boysen and Hecht, 2003¹⁶ and Kafferlein *et al.*¹⁷). One approach to measure the BPDE adducts to SA or Hb is to detach the adduct by hydrolysis to form corresponding tetrols, which are then detected by gas chromatography (GC) or liquid chromatography (LC) in combination with fluorescence or mass spectrometry (MS). Carboxylic ester adducts in albumin with aspartate and glutamate, considered favourable for binding the (+)-*anti*-BPDE¹⁸, could be released as tetrols following mild acidic hydrolysis *in vitro*. This approach has shown to be sensitive enough for application to studies of PAH-exposed humans, for instance occupationally exposed workers¹⁷. Disadvantages are though that the measured hydrolysed products have no tag from the protein proving that they correspond to released adducts and the analytes also lack specificity with regard to the binding site of BPDE in the protein. Moreover, the ester adducts are prone to undergo hydrolysis also *in vivo*. These circumstances make it difficult to calculate dose *in vivo* from measured tetrols and they are therefore not suitable as biomarkers for *in vivo* dose assessment. Further, currently available immunoassays to measure BPDE-adducts from SA provides a measure for general PAH exposure rather than specifically for B[a]P, due to cross-reactivity of the antibody with other PAHs¹⁷.

Adducts to N-sites of amino acids, such as histidine and lysine, are more stable compared to the ester adducts. Indeed, His¹⁴⁶ and Lys¹⁹⁵ have been assigned as the sites on SA for binding of BPDEs¹⁹. In our earlier work, methods to measure histidine adducts from (±)-*anti*-BPDE in human SA (hSA) were studied. One approach tested was to cleave the albumin by hydrazinolysis and detect the histidine hydrazide adduct by LC tandem mass spectrometry (MS/MS)²⁰, though the approach was considered difficult to use for routine procedure due to the hazardous properties of hydrazine. Recently, we developed a method for LC-MS/MS measurement of histidine and lysine adducts from (±)-*anti*-BPDE in hSA based on enzymatic digestion²¹. Even though structural information was obtained from this study, separation of the (+)- and (-)-enantiomer adducts of *anti*-BPDE was not attained that could be useful to assess the relative reactivity of each enantiomer and to confirm their stereospecific conformation. Furthermore, the adduct formation from the *syn*-form is not well-characterised, and, thus there is a lack of knowledge about the extent of binding and reactivity of the *syn*-BPDEs towards the nucleophilic sites of SA. Consequently, so far it has not been possible to use these methods in their present form to measure *in vivo* dose of the individual BPDE isomers, e.g. in mice.

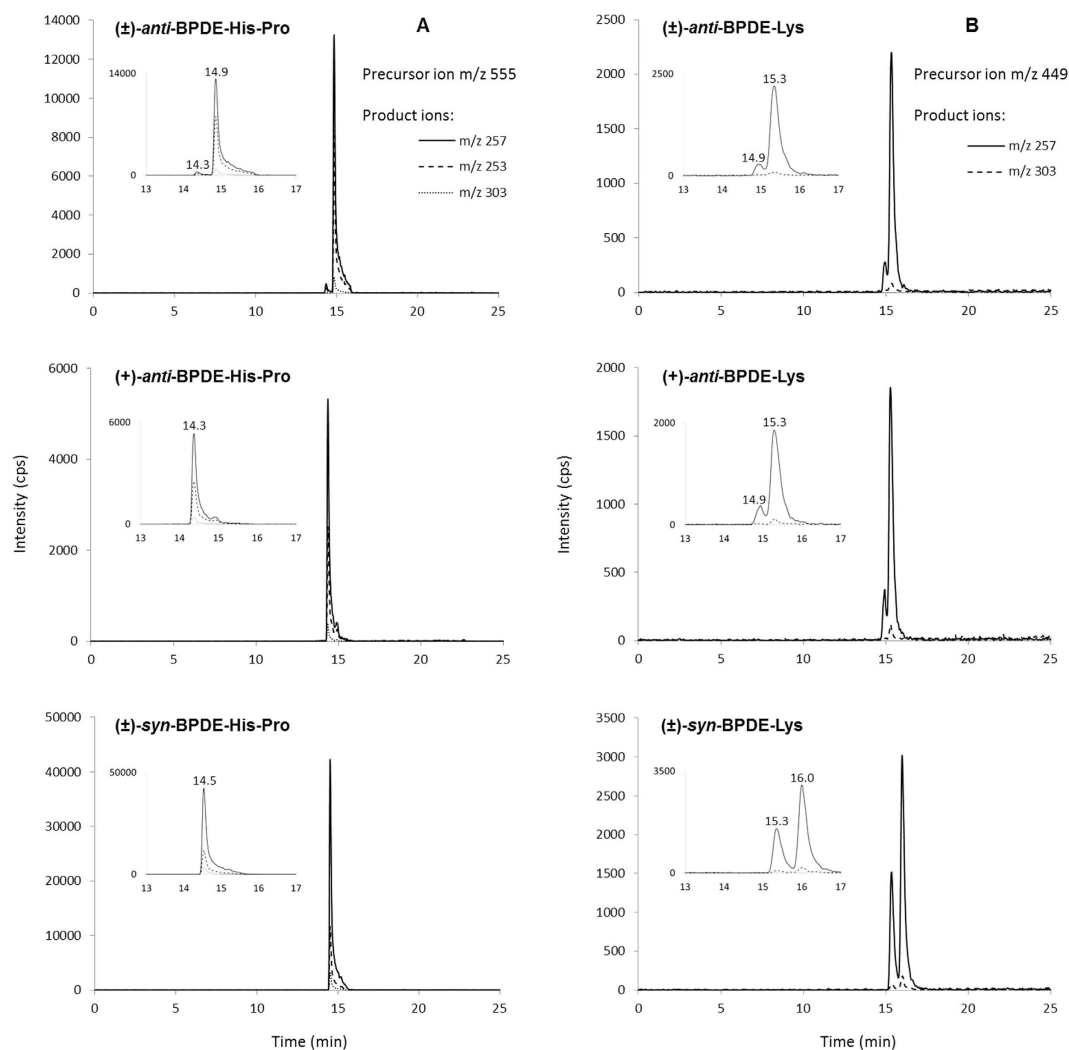


Figure 2. LC-MRM chromatograms showing peaks of BPDE-His-Pro (A) and BPDE-Lys (B) formed from hSA alkylated with (±)-*anti*-, (+)-*anti*- and (±)-*syn*-BPDE. Region between 13–17 min is inserted in each chromatogram.

In the present study we characterise the formation of histidine- and lysine-adducts from (±)-*anti*-, (+)-*anti*- and (±)-*syn*-BPDE in hSA by a structure specific LC-MS/MS methodology. Formed adduct levels and disappearance of respective BPDEs over time were measured under physiological conditions to obtain the rate of adduct formation and relationship between adduct level in hSA and dose (i.e. concentration over time) for the different BPDE isomers. This relationship could be applied in future *in vivo* studies to estimate the AUC of BPDEs from measured adduct levels of the diol epoxides arising from B[a]P exposure. The *in vivo* doses of electrophilic metabolites are potentially useful as a quantitative tool to facilitate species-extrapolation of genotoxic effects and cancer risk from animal studies to human exposure.

Results

Characterisation of serum albumin adducts to BPDEs. Human serum albumin was incubated *in vitro* under physiological conditions with (±)-*anti*-, (+)-*anti*- and (±)-*syn*-BPDE; albumin to BPDE molar ratio of 9:1. Following the incubation period (*t*) for 24 h, the protein was precipitated and enzymatically digested by pronase, and subsequently the adducts were enriched by solid phase extraction (SPE) and characterised by LC-MS/MS. A recently developed MS/MS method for analysis of the albumin adducts [with a limit of detection of about 1 fmol adduct/mg SA for (±)-*anti*-BPDE²¹] was adopted to measure the generated analytes, BPDE-His-Pro, BPDE-His-Pro-Tyr and BPDE-Lys. However, the tripeptide was not detected in any of the samples, probably due to a near complete digestion of the protein by pronase^{22,23}.

The LC-MS/MS (MRM) chromatograms in Fig. 2 show BPDE-His-Pro and BPDE-Lys from the respective incubation studies with (±)-*anti*-BPDE, (+)-*anti*-BPDE and (±)-*syn*-BPDE. LC conditions were adjusted to separate analytes from the *anti*- and *syn*-isomers, and between the enantiomers of each. This resulted in developing two setup of LC conditions, which are described in Methods. Setup 1 was applied to separate histidine analytes formed from the *anti*-enantiomers, (+) and (−); with BPDE-His-Pro from (+)-*anti*-BPDE (at 14.3 min) eluting nearly half a minute before that from (−)-*anti*-BPDE (at 14.9 min), and the His-Pro analyte from (±)-*syn*-BPDE

BPDE isomers	BPDE-His-Pro, Precursor ion m/z 555 [M+H] ⁺			BPDE-Lys, Precursor ion m/z 449 [M+H] ⁺		
	Retention time (min) ^b	Product ion (m/z)	Intensity	Retention time (min) ^c	Product ion (m/z)	Intensity
(±)-anti-	14.9/14.3	257	1.71 × 10 ⁵ au ^d	15.3/14.9	257	4.54 × 10 ⁴ au
		253	1.02 × 10 ⁵ au		253	n.d.
		303	1.13 × 10 ⁴ au		303	2.31 × 10 ³ au
(+) -anti-	14.3	257	14% ^e	15.3/14.9	257	43%
		253	18%		253	n.d.
		303	17%		303	41%
(−)-anti- ^a	14.9	257	86%	15.3/14.9	257	57%
		253	82%		253	n.d.
		303	83%		303	59%
(±)-syn-	14.5 (broad)	257	6.06 × 10 ⁴ au	15.3/16.0	257	7.94 × 10 ⁴ au
		253	2.91 × 10 ⁴ au		253	n.d.
		303	3.85 × 10 ³ au		303	4.39 × 10 ³ au

Table 1. Summary from LC-MS/MS analysis of BPDE-His-Pro and BPDE-Lys, formed from *in vitro* alkylation of hSA with (±)-anti-, (+)-anti- and (±)-syn-BPDE. ^a(−)-anti-BPDE values were estimated based on the racemic (±)-anti- and enantiomer (+)-anti-BPDE. ^bLC system setup 1. ^cLC system setup 2. ^darea, measured as arbitrary units by the Analyst software used for quantification. ^e% of (±)-anti-BPDE; n.d., not determined.

eluting between the analytes from (+) and (−)-anti-BPDE (Fig. 2 and Table 1). With regard to the BPDE-Lys analytes (setup 2), a complete separation was not obtained between those from the respective enantiomers, requiring further optimisation of the LC conditions if this would be desired.

The MS-method using ESI⁺ was effective in determining site of adduct formation, in this case to histidine or lysine, and peptide length (i.e. mono- and dipeptide) following digestion of the protein (Fig. 2 and Table 1). Table 1 summarizes the characterisation of the measured BPDE-adducts on histidine and lysine in SA, with retention times, the precursor and product ions used and relative intensities. The fragmentation pattern was in agreement with that described for the albumin adducts from (±)-anti-BPDE²¹, with m/z 257 as the predominant fragment from BPDE-His-Pro and BPDE-Lys (cf. Westberg *et al.*²¹ for proposed structures of the fragments).

Measurement of the adduct levels. Histidine and lysine adduct levels formed in hSA (0.6 mM) from (±)-anti-BPDE, (+)-anti-BPDE and (±)-syn-BPDE [initial concentration (C_0) of the different BPDEs were the same, 0.066 mM; t 24 h], following the digestion and isolation, were measured by LC-UV at 345 nm. Fig. 3 shows the chromatograms of respective adducts from the different BPDE isomers. The LC-UV system was coupled on-line to the mass spectrometer and peaks of BPDE-His-Pro and BPDE-Lys were confirmed based on their respective MS/MS spectra as described above. Additionally, there were minor unidentified peaks, e.g. peak 4 in Fig. 3. The respective adduct levels of BPDE-His-Pro ($A_{His-Pro}$) and BPDE-Lys (A_{Lys}), given in Table 2, were calculated from the area of the corresponding peaks in Fig. 3 and using equation (1) as described in Methods.

Adduct levels, $A_{His-Pro}$ and A_{Lys} , from the corresponding racemic (±)-anti- and (±)-syn-BPDE were similar, in the range of 1.3–1.9 pmol adduct/mg SA (Table 2). The (−)-anti-enantiomer of BPDE formed nearly 9 times more histidine adducts than the most potent mutagenic enantiomer, (+)-anti-BPDE, similar to that indicated earlier^{18,24}. Conversely, it was found that the selectivity for adduct formation was reversed for the lysine site, with (+)-anti-enantiomer forming ca. 3 times more adducts than the (−)-anti-enantiomer as seen from Table 2.

Kinetics of BPDEs in presence of albumin. BPDEs in aqueous mixture of SA, besides their reactions with the nucleophilic sites of the albumin, can also hydrolyse to tetrols, triols and ketones^{25,26}. Kinetics of the studied diol epoxides (±)-anti-, (+)-anti- and (±)-syn-BPDE were investigated in the present study by determining their respective apparent disappearance rate under the incubation conditions used for measurement of the adduct levels. For this purpose, aliquots were taken at certain time intervals from the incubation mixture, the BPDE in the aliquots was derivatised by β-mercaptoethanol (BME), and subsequently analysed by LC-UV at 345 nm. Representative LC-UV chromatograms from incubation at 20 min are shown in Fig. 4. Extraction efficiency of recovering BPDE-BME from the hSA mixture was estimated >90%.

The disappearance rate of the BPDEs in hSA was measured by following the thiol derivative of BPDE, i.e. BPDE-BME (peak a in Fig. 4). Pseudo-first order rate constants for disappearance (λ) of the diol epoxides were determined by plotting peak area of BPDE-BME at a given time (Area BPDE-BME_t) in relation to the area at time 0 min (Area BPDE-BME₀) versus the corresponding incubation time as shown in Fig. 5; and the obtained values of apparent λ from the respective slopes are given in Table 2. Thereby the corresponding second-order rate constants calculated from concentration of the albumin (6×10^{-4} M) for disappearance of (±)-anti-, (+)-anti- and (±)-syn-BPDE are 0.028, 0.39 and 0.31, in M⁻¹·s⁻¹, respectively. Earlier, hydrolysis rate of (±)-anti- and (±)-syn-BPDE in albumin-free water at 25 °C have been reported as 4.3×10^{-4} s⁻¹ and 1.5×10^{-2} s⁻¹, respectively^{26,27}. Altogether, these results show that the BPDEs are stabilized in presence of albumin, probably by formation of a physical association complex (BPDE-SA) as suggested earlier^{26,28,29}. Islam *et al.* estimated the apparent equilibrium constant for association of (±)-anti-BPDE to hSA as 2–3 times greater than that of (±)-syn-BPDE²⁶.

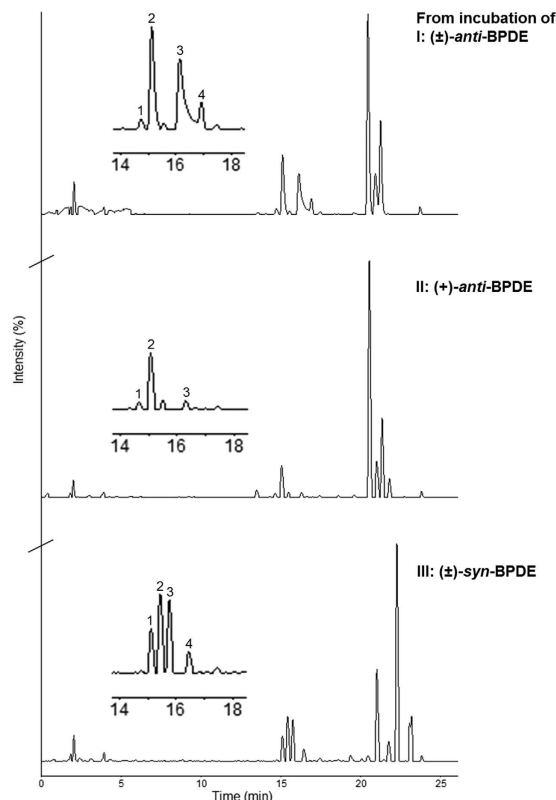


Figure 3. LC-UV chromatograms from incubation of (±)-*anti*-BPDE (I), (+)-*anti*-BPDE (II) and (±)-*syn*-BPDE (III) with hSA, at 37 °C, pH 7.5 for 24 h followed by enzymatic digestion of the albumin and isolation of the adducts; insert showing the region between 14–18 min. Where, peaks 1 and 2 correspond to BPDE-Lys, peak 3 to BPDE-His-Pro, and some unidentified peaks, e.g. peak 4, with array of peaks around 20–25 min from hydrolysed BPDEs e.g. tetrols. (Peaks 1, 2 and 3 were confirmed by MS/MS characterisation.)

BPDE isomers	$A_{His-Pro}$ (pmol adduct/mg SA)	A_{Lys} (pmol adduct/mg SA)	BPDE λ ($s^{-1} \times 10^{-5}$) [n , r^2] ^b
(±)- <i>anti</i> -	1.4	1.4	1.7 ± 0.1^c [9, 0.96]
(+)- <i>anti</i> -	0.28	2.2	24 ± 2 [6, 0.97]
(-)- <i>anti</i> - ^a	2.4	0.64	n.d.
(±)- <i>syn</i> -	1.3	1.9	19 ± 5 [6, 0.78]

Table 2. Histidine- and lysine-adduct levels ($A_{His-Pro}$ and A_{Lys} , respectively) from BPDEs, and disappearance rate (λ) of the epoxides. ^a(-)-*anti*-BPDE values were estimated based on the racemic (±)-*anti*- and enantiomer (+)-*anti*-BPDE. ^bNumber of time points n and correlation coefficient as r^2 are given in parenthesis (cf. Fig. 5). ^cSD calculated using LINEST; n.d., not determined.

The λ value of (+)-*anti*-BPDE was ca. 14 times higher than (±)-*anti*-BPDE (Table 2), implying that under the incubation conditions the disappearance of the (-)-enantiomer was slower than that of the (+)-form. Further, the disappearance rates in combination with the adduct levels presented in Table 2 give an insight into the reactivity of the BPDEs towards the nucleophilic sites of the albumin. For instance, after 24 h reaction at same initial concentrations, the sum of the adduct levels to histidine and lysine ($A_{His-Pro} + A_{Lys}$) from (±)-*anti*-BPDE and (±)-*syn*-BPDE were similar (in the range 2.8–3.2 pmol adduct/mg SA), despite that (±)-*syn*-BPDE eliminates ca. 11 times faster than the *anti*-form. This indicates that *syn*-BPDE could be considered to have a higher reactivity than the *anti*-form towards the N-sites of SA.

Dosimetry of BPDEs. From the *in vitro* incubation performed with hSA, where C_0 of BPDE was 0.066 mM, t 24 h and λ from Table 2, the dose D in mM·h of the BPDE isomers calculated using equation (3) (cf. Törnqvist *et al.*⁹) are given in Table 3. The D of BPDEs gives a measure of the concentration of the respective diol epoxides over time; referred to as AUC in *in vivo* situation^{9,13}. Using the measured levels of the adducts to histidine and lysine ($A_{His-Pro}$ and A_{Lys} , Table 2) and D , the second-order rate constants for the reaction of BPDE isomers with the nucleophilic sites, histidine and lysine ($k_{His-Pro}$ and k_{Lys} , respectively), in SA were calculated [equation (4)] and are included in Table 3.

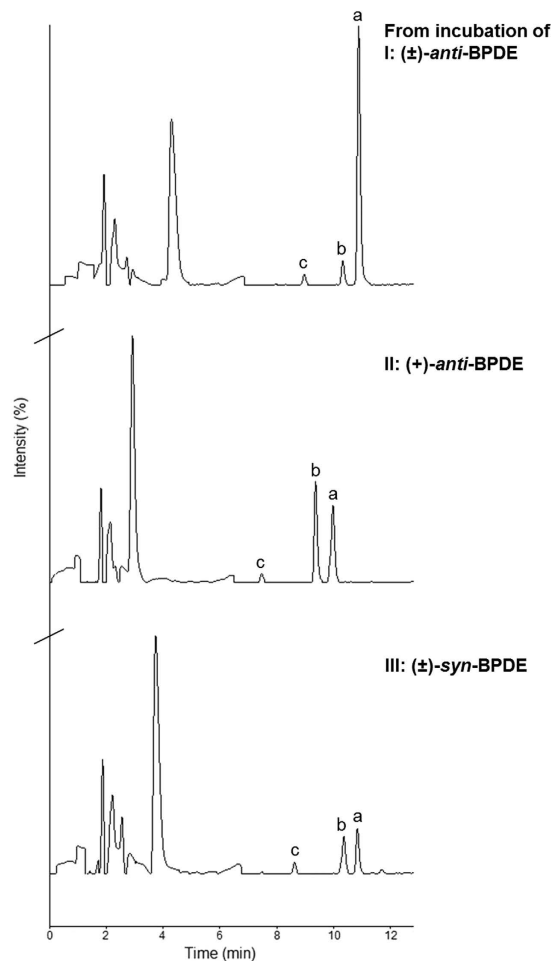


Figure 4. LC-UV chromatograms from incubation of (\pm)-*anti*-BPDE (I), (+)-*anti*-BPDE (II) and (\pm)-*syn*-BPDE (III) with hSA, at 37 °C, pH 7.5 for 20 min and subsequent derivatisation by BME. Where, peak **a** corresponds to BPDE-BME derivative, peaks **b** and **c** to hydrolysed products of BPDEs, as tetrols, and array of peaks before 5 min probably from polar components of the albumin. (Peak **a** in chromatograms I, II and III was confirmed by comparison with standard BPDE-BME formed under similar conditions. In II there was shift in retention time towards left since it was performed at a different occasion.)

It is seen from Table 3 that in general the measured reactivity of the albumin's N-sites is about a magnitude higher towards (\pm)-*syn*-BPDE than towards the (\pm)-*anti*-form under the applied conditions. The $k_{His-Pro}$ and k_{Lys} values in the table show that lysine is more prone to react with (+)-*anti*-BPDE in comparison to histidine site of the protein. The selectivity between the two N-sites is nullified for the racemic form.

Discussion

Measurement of adducts to blood proteins offers possibility to detect and quantify exposure of the precursor compound and to estimate the *in vivo* dose of the carcinogenic metabolites^{9,30}. A prerequisite for this is to have accurate structure-specific analytical methods, which could be obtained by MS. It is preferred if the analyte includes a tag from the protein, as this would confirm that the analyte is originating from a covalent bound adduct. The characterisation of adduct formation to hSA from BPDE isomers by LC-MS/MS in the present investigation is an important advancement towards this approach for PAHs.

The carrier protein hSA has been associated with binding, stabilizing and transporting of bulky epoxides^{15,26,28}. Most of the earlier work on studies of BPDE-induced changes of SA have been on the racemic *anti*-isomers (as in references^{18,20–22,31}), with only a few studies on the *syn*-forms (mainly by Islam and co-workers^{25,26}). With regard to specific MS methods for adduct characterisation, histidine and lysine adducts from racemic *anti*-BPDE have been resolved earlier by our group^{21,24}, but we found no studies on characterisation of adducts from the *syn*-form. Histidine adduct of (+)-*anti*-BPDE from mouse SA was also measured²⁴, but not with hSA and the two enantiomers of the *anti*-form were not separable in that study.

In the present work, a state-of-the-art MS methodology was used to characterise modification induced by BPDE isomers on SA at histidine and lysine. SA adduct levels formed during physiological conditions in hSA from the various BPDEs at the same initial concentrations were quantified (Table 2). The varying selectivity observed of the protein adduct formation could be explained by a preceding noncovalent binding,

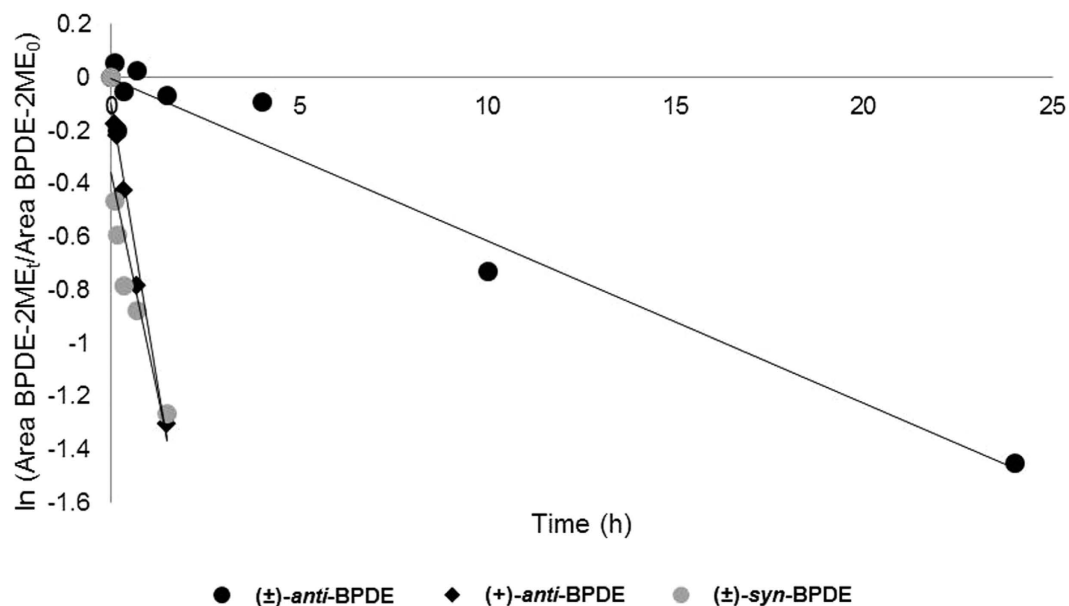


Figure 5. Plot showing disappearance of BPDEs in hSA, at 37°C, pH 7.5. Values of n and r^2 given in Table 2.

BPDE isomers	D (mM-h)	$k_{His-Pro}$ ($L \cdot g^{-1} \cdot h^{-1}$)	k_{Lys} ($L \cdot g^{-1} \cdot h^{-1}$)
(±)-anti-	0.83	1.7×10^{-6}	1.7×10^{-6}
(+)-anti-	0.078	3.6×10^{-6}	2.8×10^{-5}
(-)-anti- ^a	1.6	n.d.	n.d.
(±)-syn-	0.099	1.3×10^{-5}	1.9×10^{-5}

Table 3. Dose (D) of BPDEs in the hSA mixture and rate constants for reactions with histidine ($k_{His-Pro}$) and lysine (k_{Lys}). ^a(-)-anti-BPDE values were estimated based on the racemic (±)-anti- and enantiomer (+)-anti-BPDE; n.d., not determined.

which would control rotational and translational movement of the BPDE^{22,29}. Other studies on different PAHs epoxy-metabolites have also shown selective alkylation of the albumin sites, e.g. binding by different diol epoxides isomers of fluoranthene, chrysene, benz[a]anthracene and benzo[c]phenanthrene to histidine or lysine^{22,32}. From kinetic experiments it was indicated that in presence of SA, the BPDE isomers are protected from hydrolysis by varying degrees depending upon their conformation (cf. Table 2), probably by entering the hydrophobic pockets of the albumin^{26,28}. Further, the concentration over incubation time, i.e. dose D , of BPDEs in the hSA mixture was estimated and used to calculate the rate of formation of the adducts (Table 3). From the same initial concentration of anti-BPDE enantiomers, the dose D of the most potent mutagenic metabolite of B[a]P, (+)-anti-BPDE, is expected to be about an order of magnitude lower than the (-)-enantiomer (cf. λ in Table 2 and D in Table 3). The results overall give a mechanistic insight into the relative reactivity of the diol epoxides towards N-sites of the protein.

The rate of adduct formation to histidine of SA from the different isomers can be expressed as 1.7–13 pmol/g SA per μ M-h to simplify comparisons (from Table 3). This is much lower than the rate of adduct formation to DNA; earlier studies of adduct levels in human blood after *in vitro* reaction with (±)-anti-BPDE showed about 10^3 times higher adduct levels to deoxyguanosine in DNA than to histidine in SA²⁴. Though, the reactivity of BPDE towards the nucleophilic sites in SA is of the same magnitude as that of low molecular weight genotoxic epoxides towards N-terminal valine in Hb³³. Such adducts to Hb are established for monitoring of *in vivo* dose, or AUC, of e.g. glycidamide, genotoxic metabolite of acrylamide from dietary exposure in humans³³.

The present study gives results that have a potential to be applied in an approach that uses albumin adduct levels to estimate *in vivo* doses of metabolites of B[a]P as an indicator compound for PAHs. Primarily, the *in vivo* dosimetry will be used in our future work to estimate genotoxicity per dose unit of diol epoxide metabolites of B[a]P *in vivo*, for comparison with other carcinogens as well as with genotoxic potency *in vitro*^{11,12}. It might not be possible to measure the protein-adduct specifically from the (+)-anti-BPDE enantiomer in humans, e.g. from exposure via food, with the current available methods. In such case an inter-species comparison of the doses (AUC) of the carcinogenic metabolites in humans might be obtained from *in vivo* doses in exposed animals and metabolism studies through a parallelogram approach (cf. Motwani and Törnqvist, 2014³⁴). The derived *in vivo* doses based on the albumin adducts can then further be applied in different procedures to aid in quantitative cancer risk estimation of PAHs.

Methods

Chemicals and other materials used. β -Mercaptoethanol (BME, $\geq 99\%$) was obtained from Sigma-Aldrich (Steinheim, Germany). (\pm)-*anti*-, (\pm)-*syn*- and (+)-*anti*-Benzo[a]pyrene-7,8-diol-9,10-epoxide (BPDE) were purchased from the National Cancer Institute (NCI), Chemical Carcinogen References Standard Repository (Kansas City, KS, USA). Purity of (\pm)-*anti*-, (\pm)-*syn*- and (+)-*anti*-BPDE was ca. 95%, 95% and 80%, respectively, determined by derivatisation with BME (cf. Michaud *et al.*³⁵). Stock solutions of the three BPDEs (each 10 $\mu\text{g}/\mu\text{L}$) were prepared in ice-cold tetrahydrofuran (500 μL). Lyophilised human serum albumin (hSA) and pronase from *Streptomyces griseus* were obtained from Sigma-Aldrich (Steinheim, Germany). All solvents and chemicals used were of HPLC and analytical grade, respectively. Sep-Pak Plus C₁₈ solid phase extraction (SPE) cartridges (360 mg) were purchased from Waters (Milford, MA, USA). Eppendorf Protein LoBind tubes (Hamburg, Germany) were used for albumin samples.

Caution. BPDEs are known carcinogenic compounds. They were handled carefully in a well-ventilated fume-hood and destroyed immediately after use by 1 M aqueous H₂SO₄.

Incubation of hSA with (\pm)-*anti*-BPDE, (+)-*anti*-BPDE and (\pm)-*syn*-BPDE. BPDE solution in tetrahydrofuran was added to hSA (80 mg) in ice-cold water (2 mL) to give a final solution of 0.5 μg BPDE per mg hSA, and the pH was adjusted to 7.5 with NaOH (0.1 M, ca. 10 μL) as previously described²¹. The mixture was incubated at 37 °C for 24 h in a water-bath. Incubation with each of the three BPDE isomers was performed to characterise and measure the SA adducts, and to estimate the apparent disappearance rate constants as described below.

Kinetic studies of BPDE in hSA. Aliquots (200 μL) were taken out from the incubation at certain time intervals and mixed with BME (2 M in 0.4 M aq. NaOH, 500 μL) for 30 min at room temperature to form the BME derivative of BPDE (BPDE-BME). The mixture was cooled on ice-bath and centrifuged (14000 rpm, 4 °C, 20 min). The supernatant was extracted with ethyl acetate (400 $\mu\text{L} \times 3$), washed with water (200 $\mu\text{L} \times 2$), and the organic phase was dried under N₂. Methanol-water (1:1 v/v, 400 μL) was added to the samples and subsequently the respective BPDE-BME derivatives from (\pm)-*anti*-BPDE, (+)-*anti*-BPDE and (\pm)-*syn*-BPDE were analysed by LC-UV. Using the area from the LC-UV measurements, $\ln(\text{Area BPDE-BME}_t/\text{Area BPDE-BME}_0)$ versus the corresponding incubation time was plotted to determine the disappearance rate constant λ for the BPDE isomers. Values of n and r^2 for the individual plots are given in Table 2. Standard deviation (SD) of the slope, i.e. λ , was calculated using the Excel array function LINEST. The experiment using (\pm)-*anti*-BPDE with 20 min incubation time was repeated thrice, all giving similar peak area of BPDE-BME by LC-UV as shown in Fig. 4.

Extraction efficiency to recover BPDE-BME. (\pm)-*anti*-BPDE was used in a test to estimate the extraction efficiency (in %) of BPDE-BME following centrifugation and extraction as described above in kinetic studies. In three parallel experiments, free BPDE (4 μg in 200 μL), i.e. without albumin, was prepared from the stock solution in ice-cold water. This was added to BME (2 M in 0.4 M aq. NaOH, 500 μL) and the reaction was continued for 30 min. The aqueous samples were analysed by LC-UV and the area of BPDE-BME peak was compared with those from the kinetic study of (\pm)-*anti*-BPDE in hSA at 0 min.

Standard solution of BPDE derivatised with BME. Standard BPDE-BME derivative was prepared by a similar method as described earlier³⁵. An alkaline solution of BME (2 M, 10 mL) was prepared using 0.4 M aq. NaOH. Under ice-cold conditions water was added to a stock solution of respective BPDE in tetrahydrofuran (10 $\mu\text{g}/\mu\text{L}$) to give a concentration of 20 $\mu\text{g}/\text{mL}$. Two hundred microliter from this solution of BPDE was mixed with 500 μL of the alkaline BME solution for 30 min at room temperature. The mixture was extracted with ethyl acetate (400 $\mu\text{L} \times 3$), washed with water (200 $\mu\text{L} \times 2$), and the organic phase dried under N₂ to give BPDE-BME, which was diluted by methanol-water (1:1 v/v, 400 μL) and analysed by LC-UV.

Isolation of BPDE-adducts of hSA. Alkylation of hSA (80 mg) in 2 mL water with respective BPDEs was investigated following the incubation experiments described above at 37 °C, pH 7.5 for 24 h, with a concentration of 0.5 μg BPDE per mg hSA. The resulting adducts were isolated according to an earlier described method²¹. Briefly, following incubation after 24 h, the mixture was adjusted to pH 4 with 1 M HCl and a saturated solution of ammonium sulphate (2 mL) was added to precipitate the BPDE-alkylated hSA, at ca. 4 °C for 30 min. The precipitate was isolated, washed with methanol (10 mL), ethyl acetate (10 mL $\times 2$) and pentane (5 mL) and dried at room temperature. Subsequently, for enzymatic hydrolysis BPDE-alkylated hSA (10 mg) was mixed with pronase (0.9 mg) in aq. ammonium bicarbonate (50 mM, 450 μL), and the mixture was incubated at 37 °C for ca. 20 h. The obtained BPDE-alkylated peptide analytes were isolated by SPE and concentrated to 150 μL in water, and analysed by LC-UV and LC-MS/MS.

LC-UV and LC-MS/MS. A tandem LC-UV-MS/MS system that consisted of a Shimadzu Prominence LC 20 system and UV detector (Shimadzu Corp., Kyoto, Japan) interfaced to an API 3200 Q-trap instrument with a TurboIonSpray interface, from AB Sciex (Concord, ON, Canada) was used. BPDE-alkylated peptide analytes were analysed by two different LC setup conditions; setup 1 was favoured for obtaining separation between the His-Pro analytes from (\pm)-*anti*-, (+)-*anti*- and (\pm)-*syn*-BPDE, and setup 2 was used for separation between the respective analytes of BPDE-His-Pro and BPDE-Lys. In setup 1 the LC was coupled directly to the MS/MS via a column, while in setup 2 the UV detector, set at 345 nm, was interfaced prior to the MS/MS. In both setups, a Kinetex C18 2.6 μm , 150 \times 2.1 mm column (Phenomenex) was used and the mobile phase consisted of A, 0.1% (v/v) formic acid in methanol/water (5:95, v/v), and B, 0.1% (v/v) formic acid in methanol/water (95:5, v/v). For setup 1 gradient used was 5% B for 2 min, increased to 100% B in 24 min, at 100% B for 2 min, down to 5% B in 1 min and at 5% B for 10 min, with flow rate 0.2 mL/min, and injection volume of 5 μL . For setup 2, 0% B for 2 min,

increased to 100% B in 13 min, at 100% B for 2 min, down to 0% B in 1 min and at 0% B for 10 min, with flow rate 0.1 mL/min, and injection volume of 2 μ L. The mass spectrometer was operated with electrospray ionisation in positive ion mode (ESI⁺) using multiple reaction monitoring (MRM) mode, with parameters similar to that described earlier²¹. The MRM transitions used for BPDE-His-Pro were 555 > 253, 257 and 303, for BPDE-His-Pro-Tyr were 718 > 416, 257 and 303, and for BPDE-Lys were 449 > 257 and 303. Acquisition and processing of data from the mass spectrometer were performed using the Analyst software version 1.5 from AB Sciex.

The BPDE-BME derivative was analysed by LC-UV. An Ascentis Express F5 2.7 μ m, 150 \times 2.1 mm column (Supelco Analytical) from the LC system was coupled to the UV detector set at 345 nm. Eluent A was 0.1% (v/v) formic acid in acetonitrile/water (5:95, v/v), and B was 0.1% (v/v) formic acid in acetonitrile/water (95:5, v/v). The gradient was 10% B for 3 min, increased to 70% B in 8 min, then to 100% B in 1 min, at 100% B for 4 min, down to 10% B in 1 min and at 10% B for 5 min, with flow rate 0.2 mL/min, and injection volume of 2 μ L.

Estimation of BPDE-adduct levels. Adduct levels of BPDE-His-Pro and BPDE-Lys formed from incubation of (\pm)-*anti*-, (+)-*anti*- and (\pm)-*syn*-BPDE with hSA followed by enzymatic digestion and isolation, as described above, were estimated using the respective areas from LC-UV and incorporating them in equation (1) (cf. Westberg *et al.*²¹).

$$C_{inj} = a \times f/Q \times \varepsilon \times l \times V_{inj} \quad (1)$$

where, C_{inj} , concentration in the injected samples (in M); a , peak area (microvolt second); f , flow rate (L/sec); Q , output voltage/absorption unit (microvolt per AU); ε , extinction coefficient ($M^{-1}cm^{-1}$); l , cuvette length (cm); and V_{inj} , injection volume (L). Deoxyguanosine (dG) with ε 15400 $M^{-1}cm^{-1}$ at 259 nm was used as a reference to obtain Q , which is an instrument specific value calculated by analysing a compound with known extinction at a given concentration. The ε of BPDE-dG, 29000 $M^{-1}cm^{-1}$, at 345 nm was used as an approximate for the histidine and lysine adducts of BPDE. Influence on ε from the dG, His-Pro and Lys on the BPDE-adduct were considered to be negligible²¹.

Calculation for dosimetry. Dose (D) of a reactive compound can be defined as the concentration (C) over time (t)^{9,13}. D quantitatively takes into account the initial concentration (C_0), disappearance rate (λ) and t according to equation (2). The further derivatised equation (3) was used for calculation of D . Subsequently, D and the measured adduct levels (A) were used for determining the rate for formation of SA adducts (k_{SA}) as per equation (4).

$$D = \int C_0 \times e^{-\lambda \times t} dt \quad (2)$$

$$D = [C_0 \times (1 - e^{-\lambda \times t})/\lambda] \quad (3)$$

$$k_{SA} = A/D \quad (4)$$

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Author Contributions

H.V.M., E.W. and M.T. designed the experiments. H.V.M. performed the experiments and analysed the data. H.V.M. wrote the manuscript, and M.T. contributed with comments.

Additional Information

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