The aryl hydrocarbon receptor nuclear translocator (Arnt) is required for tumor initiation by benzo[a]pyrene

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Benzo[a]pyrene (B[a]P) is a ligand for the aryl hydrocarbon receptor (Ahr). After binding ligand, Ahr dimerizes with the aryl hydrocarbon receptor nuclear translocator (Arnt) protein, and the dimer upregulates the transcription of Cyp1a1, Cyp1b1 and other enzymes involved in the metabolic activation of B[a]P. Arnt null mice die in utero. Mice in which Arnt deletion occurs constitutively in the epidermis die perinatally. In the current study, mice were developed in which the Arnt gene could be deleted specifically in adult skin epidermis. This deletion had no overt pathological effect. Homozygosity for a null reduced nicotinamide adenine dinucleotide (phosphate): quinone oxidoreductase allele contained a K14-Cre-ERT2 allele which was excised by this procedure, and these genotypes are homozygotes by this procedure, and these genotypes are not distinguishable from heterozygotes. Nqo1, reduced nicotinamide adenine dinucleotide (phosphate): quinone oxidoreductase, TPA, 12-O-tetradecanoyl phorbol-13-acetate (B[a]P) is used as the initiator and 12-O-tetradecanoyl phorbol-13-acetate (TPA) is used as the promoter (1). Tumor initiation by B[a]P is presumed to require its metabolism to mutagenic derivatives (2). The principal such derivative is B[a]P 7,8-diol-9,10-epoxide, which is generated by the activities of certain cytochrome P450s (principally CYP1A1 and CYP1B1) and epoxide hydrolase. The levels of CYP1A1 and CYP1B1 are markedly upregulated by B[a]P in a process mediated by the aryl hydrocarbon receptor (Ahr). After binding ligands such as B[a]P, the Ahr translocates into the nucleus and dimerizes with the aryl hydrocarbon receptor nuclear translocator (Arnt) protein. The Ahr/Arnt dimer then binds to xenobiotic-responsive elements upstream of the Cyp1a1 and Cyp1b1 proximal promoter region of these and other genes and upregulates their rates of transcription (3).

Although it has been demonstrated that Ahr is required for complete carcinogenesis by B[a]P (4), it has not been determined whether Ahr is required in the initiation or promotion stages of this process or both. It is thus formally possible that Ahr is not required for tumor initiation by B[a]P. Furthermore, even if Ahr is required for tumor initiation by B[a]P, it is conceivable that Arnt is not required since some pathological effects of Ahr ligands appear to be mediated by Ahr independently of Arnt and of mutagenic events (5–8). This study addresses the hypothesis that Arnt is required for skin tumor initiation by B[a]P, by utilizing mice in which Arnt has been deleted specifically in the epidermis. Arnt (and Arnt2) can also dimerize with hypoxia-inducible factor (HIF)-1α or HIF-2α to form HIF, which represents the master regulator of the hypoxic response. In normal cells under normoxic conditions, HIF-1α and HIF-2α are stabilized and upregulated many genes involved in glycolysis, angiogenesis and cell survival. Portions of solid tumors are generally in a hypoxic state, resulting in activation of HIF-1α and/or HIF-2α (9). Furthermore, HIF-1α and HIF-2α are also upregulated in many cancer cells even under normoxic conditions, due to the effects of activated proto-oncogenes or the inactivation of tumor suppressor genes (10). A large number of studies have been performed to investigate the role of HIF in tumor growth. With few exceptions, these studies have indicated that HIF activity enhances tumor growth (11). However, these studies have utilized tumor xenograft models, which have the inherent limitations that the tumor cells so tested are already fully malignant and that the tumor cells are exposed to an artificial cellular environment. In contrast, our experiments provides us with the potential opportunity to investigate the effect of ablation of Arnt (and thus of HIF) on the growth kinetics of endogenously arising tumors in the skin. This issue represents a further objective of our studies.

Materials and methods

Breeding and genotyping of mice

The original Arnt2 allele contained a neo cassette (12). This was excised as described previously (13). The mice were originally of a mixed C57BL/6, 129/Sv and FVB/N genetic background. The mice were crossed to homozygous K14-Cre-ER12 mice (14), which were of a mixed C57BL/6 and FVB/N genetic background. Progeny from this cross were then backcrossed at least 10 successive times to the C57BL/6 strain before experiments were performed. These mice are therefore in a 100% C57BL/6 background. Genotyping of the Arnt and Arnt2 alleles was performed by polymerase chain reaction as described previously (15). The K14-Cre-ER12 transgene was genotyped by polymerase chain reaction primers directed at the Cre gene as described previously (15). K14-Cre-ER12 mice were crossed with reduced nicotinamide adenine dinucleotide (phosphate): quinone oxidoreductase (Nqo1−/−) mice that were of the C57BL/6 genetic

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background, and the presence of the wild-type and Nqo1−/− alleles was assessed by polymerase chain reaction (16).

Carcinogen treatment

Mice of 11 weeks of age were injected intraperitoneally with 100 μg tamoxifen (in 100 μl sunflower oil) for five consecutive days and again for three consecutive days every month. Carcinogen treatments started 4 weeks after the first tamoxifen dose. Mice were treated on an area of shaved dorsal skin with 200 μg of B[a]P dissolved in 200 μl acetone, with one dose of 5 μmol N-methyl-N'-nitro-N-nitrosoguanidine (MNNG) or with the vehicle, acetone. Twice weekly applications of 10 μg TPA started 1 week after initiator treatment and were continued for 25 weeks. Mice were monitored for tumors weekly for up to 40 weeks or were euthanized when tumors exceeded 1.5 cm in diameter, if they ulcerated or if the mice showed signs of distress or discomfort. Tumors >2 mm in diameter were counted. See Table I for the different types of carcionogen treatment. DNA from tumors was used for the DNeasy Blood and Tissue kit (Qiagen, Valencia, CA) following the manufacturer’s instructions.

Analysis of tumors

Diagnosis was performed by a board certified pathologist on tumor specimens fixed with formalin and stained with hematoxylin and eosin. The additional diagnosis was performed with the indicated antibody.

Isolation of epidermis and dermis from normal skin

Mouse skin was treated with 0.25% trypsin at 4°C for five consecutive days, and the epidermis and dermis were pelleted away from the epidermis with forceps. Epidermal sheets were stirred in Eagle’s minimal essential medium containing 8% fetal calf serum. The resulting epidermal cell suspension was then filtered through Nytex gauze to remove epidermal cell suspension to remove the stratum corneum and remaining hair, and the epidermal cells were pelleted by centrifugation. DNA was extracted from the dermis and epidermal cell pellets using the DNeasy Blood and Tissue kit (Qiagen, Valencia, CA) following the manufacturer’s instructions.

Results

Deletion of Arnt eliminates the development of tumors in a model of tumor initiation by B[a]P

Mice that are homozygous for an Arnt null allele die in utero (17,18). ArntF/F, K14-Cre mice, in which deletion of Arnt occurs in the skin epidermis, die within 1 day of birth due to loss of skin barrier function (15). In order to delete Arnt in the skin epidermis of adult mice, we constructed mice of the following genotype-ArntF/F: K14-Cre-ER[T2]+ (either homozygous or heterozygous for the Cre transgene). Cre-ER[T2] expression in the above construct is driven by the keratin 14 promoter, which is selective for the basal layer of stratified squamous epithelia. Cre-ER[T2] encodes a fusion protein between the Cre recombinase and a mutated ligand-binding domain of the human estrogen receptor α. The recombinase activity of this construct is activated by the anti-estrogen tamoxifen but not by endogenous estrogens (14). These mice were originally in a mixed C57BL/6:129/Sv: FVB/N genetic background, but we backcrossed them at least 10 times to the C57B1/6 strain, so that the experimental animals were in a pure C57BL/6 genetic background. The C57BL/6 mouse strain is particularly resistant to tumor formation in the B[a]P plus TPA protocol (19). In order to increase the number of tumors, we generated mice of the above genotype that were also homozygous for a Nqo1 null allele since Nqo1−/− mice exhibit much greater sensitivity to B[a]P plus TPA skin tumorigenesis than wild-type C57BL/6 mice (16,19).

We utilized a protocol involving a single topical application of B[a]P to mouse skin, followed by twice weekly applications of TPA. In this protocol, B[a]P acts as tumor initiator and TPA acts as tumor promoter. We compared tumor formation in ArntF/F: K14-Cre-ER[T2]+: Nqo1−/− mice that had previously been injected intra-peritoneally with tamoxifen or had been left untreated. Analysis of the epidermis and dermis isolated directly from the skin of these mice showed that Arnt deletion occurred rapidly and efficiently in the epidermis but not the dermis of these mice after tamoxifen treatment (Figure 1), consistent with the original observations on the K14-Cre-ER[T2] transgene by Li et al. (14). Tamoxifen treatment had no overt detrimental effect on the skin of Cre-positive mice not treated with B[a]P (group I5). Four weeks after the initial tamoxifen treatment, mice were treated once topically with 200 μg B[a]P and then twice weekly with TPA (groups I1 and I2 in Table I). As controls, we also examined mice untreated with B[a]P (groups I5 and I6) or mice treated with the alkylating agent MNNG (groups I3 and I4), which acts as a direct-acting tumor initiator, but whose activity is unlikely to be affected by the activity of Ahr, Arnt or their target genes (all mice received TPA). From 14 to 24 mice were treated per group (approximately the same number of males and females in each group). Fifty-eight per cent of ArntF/F: K14-Cre-ER[T2]+: Nqo1−/− mice untreated with tamoxifen but treated with B[a]P developed tumors, and there were an average of 1.8 tumors per mouse. No tumors were obtained in equivalent mice that were treated with tamoxifen to induce deletion in the ArntF allele (Figure 2A and B). MNNG + TPA treatment generated tumors in approximately the same proportion of mice (50 and 43%) at approximately the same frequency (1.1 and 0.70) in tamoxifen-untreated and in tamoxifen-treated mice, respectively. No tumors were obtained in mice treated with TPA without prior treatment with B[a]P or MNNG, irrespective of whether or not they had been treated with tamoxifen (Figure 2A and B). These results demonstrate that Arnt activity is required for tumor initiation by B[a]P in this experimental model.

All eight tumors induced by B[a]P plus TPA that were analyzed were squamous cell carcinomas. Thus, all the tumors were epithelial in origin, which is consistent with the topical application of B[a]P; its

<table>
<thead>
<tr>
<th>Group</th>
<th>Tamoxifen</th>
<th>B[a]P or MNNG</th>
<th>TPA</th>
<th>Number of mice</th>
<th>Expected Arnt phenotype</th>
</tr>
</thead>
<tbody>
<tr>
<td>I1</td>
<td>+</td>
<td>B[a]P</td>
<td>+</td>
<td>24</td>
<td>–</td>
</tr>
<tr>
<td>I2</td>
<td>+</td>
<td>B[a]P</td>
<td>+</td>
<td>19</td>
<td>+</td>
</tr>
<tr>
<td>I3</td>
<td>+</td>
<td>MNNG</td>
<td>+</td>
<td>14</td>
<td>–</td>
</tr>
<tr>
<td>I4</td>
<td>+</td>
<td>MNNG</td>
<td>+</td>
<td>14</td>
<td>+</td>
</tr>
<tr>
<td>I5</td>
<td>+</td>
<td>Acetone</td>
<td>+</td>
<td>14</td>
<td>–</td>
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<tr>
<td>I6</td>
<td>+</td>
<td>Acetone</td>
<td>+</td>
<td>15</td>
<td>+</td>
</tr>
</tbody>
</table>

All mice were of the genotype K14-Cre-ER[T2]+: ArntF/F: Nqo1−/−.

Fig. 1. Rapid deletion of Arnt in epidermis but not dermis of ArntF/F: K14-Cre-ER[T2]+ mice treated with tamoxifen. Mice were injected with tamoxifen for five consecutive days, and skin excised 3 days later. The floxed Arnt allele gives a polymerase chain reaction (PCR) product of 307 bp, whereas the deleted Arnt allele gives a PCR product of 328 bp. Both ArntF and ArntFb alleles were detected in whole skin of tamoxifen-treated Cre+ mice (lane 1). No deletion occurred in the epidermis (epi) from Cre− mice treated with tamoxifen (lane 2) or in the epidermis from Cre+ mice not treated with tamoxifen (lane 3) or in the dermis (derm) of Cre− mice treated with tamoxifen (lane 4). Complete deletion occurred in the epidermis of tamoxifen-treated Cre+ mice (lanes 5, 6 and 7, corresponding to three different mice). (The initial lane contains DNA molecular weight markers.)
greater rate of metabolism in epidermis than dermis (20,21) and its limited permeation to the dermis (22). In the case of MNNG plus TPA, 64% of the tumors were squamous cell carcinomas and 14% were papillomas and thus of epidermal origin. The remainder of the tumors were neurofibrosarcomas (Table II).

Although Arnt2 has been reported not to collaborate with Ahr in activation of gene transcription and was previously not detected in mouse skin by gross analysis (23–25), we nevertheless thought it prudent to test for Arnt2 protein expression in skin and in the tumors. Immunohistochemical analysis failed to detect the Arnt2 protein in normal skin epidermal keratinocytes or in representative tumors, whereas the protein was readily detectable in mouse kidney (data not shown).

**Role of Arnt in kinetics of tumor growth**

Since MNNG is presumed to act independently of Arnt, the kinetics of growth of the tumors induced by MNNG plus TPA provided a potential opportunity to investigate the effect of ablating Arnt on the growth characteristics of endogenously arising tumors. Mice treated with tamoxifen developed tumors from the MNNG plus TPA regimen (I3, Figure 2A and B) more quickly than those not treated with tamoxifen (I4). This suggested that loss of Arnt accelerated tumor growth in this system. In order to investigate this possibility further, we examined representative tumor specimens for the presence of the ArntF and ArntD alleles. Tumors from mice treated with tamoxifen (I3) exhibited only the ArntD allele. Unexpectedly, however, mice not treated with tamoxifen [whether exposed to B[a]P or MNNG] exhibited the ArntD allele as well as the ArntF allele (groups I2 and I4, Figure 3).

The proportion of the different alleles in these last tumors was not related to the sex of the animals, indicating that the degree of deletion was not related to the levels of circulating estrogens in the animals. Deletion of Arnt in non-tamoxifen-treated mice also did not occur in mice not treated with TPA. These observations are consistent with the notion that TPA activates Cre-ERT2 either directly or indirectly and caused partial deletion of Arnt. Since deletion of Arnt presumably occurred in the non-tamoxifen-treated, MNNG plus TPA-treated mice during the development of the tumors, the interpretation of the tumor growth kinetic data with regard to the role of Arnt in the growth of these tumors is compromised and conclusions regarding whether Arnt accelerates or retards tumor growth in this model cannot be made. It should be noted, however, that the single B[a]P treatment occurred before TPA treatment, and therefore, the conclusion that Arnt is required for tumor initiation by B[a]P is not compromised.

**Discussion**

In our study, we used mice in which Arnt could be deleted specifically in the adult epidermis. Furthermore, we introduced homozygosity for an Nqo1 null allele into these mice to enhance their sensitivity to the
tumor initiation–promotion protocol. B[α]P is metabolized by Cyp1a1 and other cytochrome P450s. Nqo1 null mice exhibit an increased sensitivity to B[α]P-induced skin cancer, probably via an altered redox state along with lack of induction of p53 and decreased apoptosis (26). The ArntF/F: K14-Cre-ERT2: Nqo1 null mice were completely resistant to induction at the site of application (28). Our results are consistent with this generalization. There is a considerable effort underway to develop HIF-1α and/or HIF-1β inhibitors for the treatment of cancer (reviewed in ref. 29). Since inhibition of Arnt would not only negate the hypoxic response to tumors but, as our results suggest, would also inhibit the genotoxic effects of the environmentally ubiquitous polycyclic aromatic hydrocarbons, Arnt may represent a better target for therapy than HIF-1α.

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Conflict of Interest Statement: None declared.

References

Fig. 3. Genotyping of tumor specimens. Tumors were excised and their DNA isolated. Genotyping of the Arntβ and Arntγ alleles was performed by polymerase chain reaction (PCR) as described in ‘Materials and Methods’. The corresponding PCR products in the ethidium bromide stained gels are indicated.

A

B

100 bp

100 bp

Arntβ

Arntγ

12 mice

13 mice

14 mice

I2 mice

1960

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