Inducible Site-Specific Somatic Mutagenesis in Mouse Hepatocytes

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To perform ligand-dependent site-specific mutagenesis in the liver, we established transgenic mice expressing the tamoxifen-inducible Cre-ER^T recombinase (Feil *et al.*, 1996; Brocard et al., 1997) under the control of the human α 1-antitrypsin (α AT) promoter (Fig. 1) that is specifically active in hepatocytes (Jallat et al., 1990). Out of 20 transgenic founder animals, three lines expressing Cre-ER^T in the liver, but not in the other tested organs, were obtained (Fig. 2A and data not shown). The expression pattern of Cre-ER^T in the liver was analysed in the transgenic line exhibiting the highest Cre-ER^T expression level. In situ hybridization as well as immunohistochemistry analysis revealed that Cre-ER^T was expressed in approximately 40%-50% of the hepatocytes (Fig. 2B,C). Mosaic transgene expression using the α AT promoter was previously reported (Ponder, 1996; Veniant et al., 1996). To analyse the recombinase activity, the α AT-Cre-ER^T transgenic line was crossed with reporter mice containing a loxP-flanked (floxed) allele of a target gene (Feil et al., 1997; Mascrez et al., 1998), and offspring harboring both the Cre-ER^T and the floxed reporter allele were treated with tamoxifen. Excision occurred for 40%-50% of the reporter allele in the liver and was undetectable in all other organs analysed, as well as in oil-treated animals (Fig. 2D). Thus, recombination occurred in almost all hepatocytes expressing Cre-ER^T

Taken together, our results show that the α AT-Cre-ER^T transgenic mice allow conditional site-specific somatic mutagenesis in the liver. The mosaic Cre-ER^T expression in hepatocytes should be useful to produce genetic chimeras to study gene function during postnatal liver growth, regeneration, and carcinogenesis, and to perform genetic marking studies (Ponder, 1996).

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FIG. 1. Schematic diagram of the α AT-Cre-ER^T transgene. The blunt-ended 1.8 kb EcoR I-Sal I fragment, containing the human α 1-antitrypsin promoter and the first non-coding exon (grey box), the 5' sequences of the first intron (dotted line), and the c-myc splice acceptor site (hatched box), was isolated from pTG3925 plasmid (Jallat *et al.*, 1990) and cloned into the blunt-ended Sall site of pGS-Cre-ER^T (Indra *et al.*, 1999), resulting in α AT-Cre-ER^T. The black boxes represent rabbit β -globin exonic sequences. Transgenic mice were established with the 4.6 kb Not I fragment of $p\alpha$ AT-Cre-ER^T, as described (Feil *et al.*, 1996).

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NT aAT-Cre-ERT (A)- Cre-ER^T HPBT B WT αAT-Cre-ERT Cre (S) Cre (AS) C αAT-Cre-ERT WT oil-treated mouse tamoxifen-treated mouse D WT and floxed allele "excised

FIG. 2. Characterization of the α AT-Cre-ER^T transgenic mouse line. (A) RT-PCR analysis of Cre-ER^T expression was performed on RNA extracted from liver of WT mice and from kidney, spleen, and liver of α AT-Cre-ER^T transgenic mice, as described (Feil *et al.*, 1996). The PCR products corresponding to Cre-ER^T and HPRT mRNA are indicated. (B) In situ analysis of Cre-ER^T expression was performed as indicated on cryosections of liver from WT or aAT-Cre-ERT transgenic mice, hybridized with Cre antisense (AS) and sense (S) riboprobes, according to Dollé and Duboule (1989). Magnification, $160 \times$. (C) Cre-ER^T immunohistochemistry was performed with the biotinylated mouse monoclonal antibody 2Cre1A1 (unpublished) on cryosections of liver from tamoxifen-treated WT and aAT-Cre-ERT transgenic mice as indicated, according to Brocard et al. (1997). Staining of Cre-ER^T with 2Cre1A1 (red) and nuclear DNA with DAPI (blue) were superimposed (pink). Magnification, $400 \times$. (D) Cre-ER^Tmediated DNA excision was determined by Southern blot analysis, performed on genomic DNA isolated from organs of αAT -Cre-ER^T/ $RXR\alpha^{+/af2(l)}$ double-transgenic mice three days after the last tamoxifen or oil injection and digested with BamHI and hybridized with probe B, as described (Feil *et al.*, 1996; Mascrez *et al.*, 1998). C, control DNA from $RXR\alpha^{+/af2(II)}$ mice. The WT, floxed [af2(I)], and "excised" [af2(II)] RXRα alleles (Mascrez et al., 1998) are indicated by arrows.

allele

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