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Early development of histiocytic sarcomas in p53 knockout mice treated with N-bis(2-hydroxypropyl)nitrosamine

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Abstract. p53 knockout mice have been utilized for the functional analysis of p53 in carcinogenic processes and for the evaluation of the carcinogenic potential of chemicals. In this study, we established that p53 knockout mice have an elevated susceptibility to the induction of histiocytic sarcoma (HS) by N-bis(2-hydroxy-propyl)nitrosamine (BHP). p53 heterozygous (+/-) and wild-type (++) mice were treated with 20 or 200 ppm BHP in their drinking water for 15 weeks or with 20 ppm BHP for 40 weeks. An additional group of p53 nullizygous (-/-) mice were treated with 20 ppm BHP for 15 weeks. In a 15-week experiment, hepatic HSs were unexpectedly observed in BHP-treated p53 (-/-) mice (30.8%) but not in p53 (+/-) and (++) mice and untreated (-/-) mice, indicating that a complete loss of p53 dramatically accelerates the genesis of HS. In a 40-week experiment, HSs were significantly increased in female p53 (+/-) mice (37.5%) as compared with female (++) mice (5.0%). Additionally, PCR-SSCP and sequencing analysis revealed a high frequency of p53 gene mutations in HSs, demonstrating the involvement of p53 gene mutations in HS development. Our data add to the understanding of the carcinogenic susceptibility of p53 knockout mice, and may help to elucidate the pathogenesis of HS development.

Introduction

The tumor suppressor p53 plays a critical role in preventing the accumulation of the multiple genetic changes involved in tumor development (1). p53 knockout mice, first established

by Donehower *et al* (2), have been utilized for the functional analysis of p53 in carcinogenic processes and, additionally, for the evaluation of the carcinogenic potential of chemicals (3).

Histiocytic sarcoma (HS) is classified as a non-lymphoid hematopoietic neoplasm (4) and is thought, based on its immunohistochemical and ultrastructural features, to be derived from cells of mononuclear phagocyte lineage (5,6). It is a rare, spontaneous tumor in mice; at 24 months of age, incidences of it in common strains are between 0 and 5.5% (4,7). However, more frequent development has been reported in genetically engineered mice (8-12). It is noteworthy that this is also the case in animals with p53-related alterations. For example, it was reported that HS is the most common spontaneous tumor in p21-deficient mice, accounting for approximately half of all the lesions which develop (8). Loss of BAX alters the tumor spectrum in ARF-deficient mice and results in the emergence of HS (10). Using double-mutant mice, it was also demonstrated that coincident loss of ARF, a regulator of the p53 pathway, with INK4A and PTEN induced the development of HS (12). However, the relevance of p53 itself to HS development has remained unclear. Spontaneous development of HS has not been reported in p53 knockout mice, either in nullizygotes [p53 (-/-)] or heterozygotes [p53 (+/-)], though they show markedly enhanced genesis of tumors of various types (2,13-18). The mechanisms underlying HS development are poorly understood, so it would appear that studies using genetically engineered mice are warranted.

N-bis(2-hydroxypropyl)nitrosamine (BHP) is a multi-organ genotoxic carcinogen in rodents, whose oral administration induces both lung and hepatic vascular tumors in mice (19). We previously reported on the carcinogenicity of BHP in p53 knockout mice, with a particular focus on differences in susceptibility with reference to lung and hepatic vascular tumors (20). BHP also rapidly induced HS in the livers of p53 nullizygous (-/-) and heterozygous (+/-) mice, though it was not expected that histiocytic cells would be the targets of this carcinogen. We therefore further investigated, in this study, HS development in BHP-treated p53 knockout mice, with a focus on the molecular mechanisms linked to p53 gene alteration.

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Key words: p53 knockout mouse, histiocytic sarcoma, N-bis(2-hydroxypropyl)nitrosamine, liver, p53 mutation

Table I. PCR primers for genotyping and for SSCP analysis of mouse p53.

Target		Primer sequence	Product length (bp)	
Genotype	Wild-type	10681	5'-GTGTTTCATTAGTTCCCCACCTTGAC-3'	320
		10480	5'-ATGGGAGGCTGCCAGTCCTAACCC-3'	
	Mutant	10588	5'-GTGGGAGGGACAAAAGTTTCGAGGCC-3'	150
		10930	5'-TTTACGGAGCCCTGGCGCTCGATGT-3'	
PCR-SSCP	Exon 5	Sense	5'-TCTCTTCCAGTACTCTCCTC-3'	214
		Antisense	5'-AGGCGGTGTTGAGGGCTTA-3'	
	Exon 6	Sense	5'-GGCTTCTGACTTATTCTTGC-3'	181
		Antisense	5'-CAACTGTCTCTAAGACGCAC-3'	
	Exon 7	Sense	5'-TCACCTGGATCCTGTGTCTT-3'	170
		Antisense	5'-CAGGCTAACCTAACCTACCA-3'	
	Exon 8	Sense	5'-ACTGCCTTGTGCTGGTCCCTT-3'	279
		Antisense	5'-GGAGAGGCGCTTGTGCAGGT-3'	

Materials and methods

Animals. p53 knockout mice produced by Donehower *et al* (2) were backcrossed to C57BL/6J (N4) and maintained at the Animal Facility of Aichi Cancer Center Research Institute. Five-week-old males and females were used in the present experiment. They were housed at a maximum of 5 per plastic cage on wood chips in an air-conditioned room with a 12-h light-dark cycle, and given a basal diet (Oriental NMF, Oriental Yeast, Tokyo, Japan) and drinking water *ad libitum*. Genotyping of each mouse was performed by PCR using 2 sets of primers (Table I), basically as previously described (21).

Experimental design. The experimental design is shown in Fig. 1. BHP (Nacalai Tesque, Kyoto, Japan) was dissolved in distilled water and freshly prepared two times per week. In experiment I, male and female p53 (+/+) and (+/-) mice were given drinking water *ad libitum* containing 20 or 200 ppm BHP in light-shielded bottles for 15 weeks. An additional group of p53 (-/-) male mice were treated with only 20 ppm for 15 weeks. In experiment II, long-term observation was performed using male and female p53 (+/+) and (+/-) mice that were given drinking water containing 20 ppm BHP for 40 weeks. As controls, matching numbers of mice of each genotype were given distilled water for the same period of time. Necropsies were performed on all animals that died or were sacrificed when they became moribund. Data for these were not included in experiment I, but, in experiment II, the mice were considered as effective animals when prematurely sacrificed as moribund after experimental week 36. At the end of the experiment, all surviving mice were sacrificed and autopsied.

Histopathological analysis. Tissues were fixed in 4% paraformaldehyde in 0.1 M phosphate buffer and were routinely

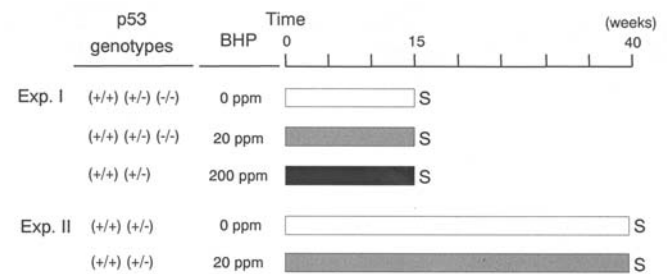


Figure 1. Experimental design. In experiment I, male and female p53 (+/+), (+/-) and male (-/-) mice were used. In experiment II, animals were (+/+) and (+/-) mice of both sexes. BHP at 20 (gray bar) or 200 (closed bar) ppm and water (open bar) as a control were given. Exp., experiment; BHP, N-bis(2-hydroxypropyl)nitrosamine; S, sacrifice.

processed to sections stained with hematoxylin and eosin (H&E). For the differential diagnosis of HS and lymphoma, the following immunohistochemical markers were used: rabbit anti-human CD3 polyclonal antibody (Dako Japan, Kyoto, Japan) for T-lymphocyte; rat anti-mouse CD45R/B220 monoclonal antibody (Pharmingen, San Diego, CA) for B-lymphocyte; and rat anti-mouse F4/80 monoclonal antibody (Serotec, Oxford, UK) for monocyte/macrophage detection. A standard avidin-biotin technique (Vectastain ABC kit, Vector Laboratories, Burlingame, CA) was employed with diaminobenzidine as the chromogen. Sections were counterstained with Meyer's hematoxylin to aid orientation.

Analysis of p53 gene alterations. Twelve hepatic HSs from BHP-treated p53 knockout mice, 2 from p53 (+/+) and 10 from (+/-) mice, were subjected to analysis of p53 gene alterations. In order to detect the loss of heterozygosity (LOH) of the p53 gene in the lesions of p53 (+/-) mice, PCR was

Table II. Incidence of histiocytic sarcoma.

Experimental period	Sex	BHP (ppm)	p53 genotypes	No. of animals			No. of mice with tumors (%)		
				Initial	Survival	Effective	Liver	Spleen	Lung
Exp. I (15 weeks)	Male	0	+/+	20	20	20	0	0	0
			+/-	20	19 ^e	19	0	0	0
			-/-	20	17 ^f	17	0	0	0
		20	+/+	32	32	32	0	0	0
			+/-	32	32	32	0	0	0
			-/-	41	26 ^g	26	8 (30.8) ^a	3 (11.5)	2 (7.7)
	Female	0	+/+	20	20	20	0	0	0
			+/-	20	20	20	0	0	0
			-/-	20	20	20	0	0	0
		20	+/+	32	32	32	0	0	0
			+/-	32	32	32	0	0	0
			-/-	29	15 ⁱ	15	0	0	0
Exp. II (40 weeks)	Male	0	+/+	20	20	20	0	0	0
			+/-	20	19 ^j	19	0	0	0
		20	+/+	20	19 ^k	19	0	0	0
	Female	0	+/-	20	14 ^l	17 ^c	2 (11.8)	0	0
			+/+	20	19 ^m	19	0	0	0
		20	+/+	20	20	20	0	0	0
		20	+/-	20	15 ⁿ	16 ^d	1 (5.0)	0	0
			+/+	20	15 ⁿ	16 ^d	6 (37.5) ^b	0	0

^aSignificantly different from male 20 ppm BHP-treated p53 (+/+) and (+/-) mice and control p53 (-/-) mice using the Fisher's exact test, $P < 0.001$, < 0.001 and < 0.05 ; ^bSignificantly different from 20-ppm BHP-treated female p53 (+/+) and control female p53 (+/-) mice, $^dP < 0.05$ and $^eP < 0.005$; ^cData from three mice sacrificed at experimental week 38 and 39 were included; ^dData from one mouse sacrificed at experimental week 37 was included; ^eCause of death was lymphoma; ^fThe deaths included one from fibrosarcoma, one from malignant fibrous histiocytoma and one from hydrocephalus; ^gThe 15 deaths included eight from lymphomas, three from sarcomas, one from malignant fibrous histiocytoma and one of unknown cases; ^hThe 12 deaths included ten from hepatic hemangiosarcomas and two of unknown causes; ⁱThe causes of all of fourteen deaths were hepatic hemangiosarcomas; ^{j,k}Causes of deaths were not determined; ^lThree moribund mice were sacrificed at experimental weeks 38 and 39. Cause of deaths of the other three mice was lymphoma development; ^mThe cause of the death was a carcinoma; ⁿOne moribund mouse was sacrificed at week 37. The deaths of the other four mice included hepatic histiocytic sarcoma, lymphoma, hemangio sarcoma and one unknown cause.

performed to specifically amplify the wild or mutant allele with PCR primers for the genotyping of mice (Table I). PCR-single strand conformation polymorphism (SSCP) analysis was conducted basically as previously described (21). The four pairs of primers applied for mouse p53 exons 5-8 are listed in Table I. The primers for exon 5 were designed to specifically amplify the wild-type allele. Abnormal bands, separated by electrophoresis, were excised from the gels and subjected to sequencing analysis utilizing an ABI PRISM 3100 with a BigDye Terminator v3.1 Cycle Sequencing Ready Kit (Applied Biosystems, Forester City, CA). For tumor samples with p53 mutations at exons 7 and 8 in p53 (+/-) mice, long PCR was performed to selectively amplify the wild-type allele

of the p53 gene using a Takara LA-PCR kit (Takara Bio, Shiga, Japan) as previously described (22). Due to the localization of mutations, a PCR sense primer for exon 5 was used in combination with antisense primers for exons 7 or 8 (Table I).

Statistical analysis. Data for incidences of tumors were analyzed using Fisher's exact probability test. The survival of each genotype of mice was analyzed with the log-rank test.

Results

Mortality of mice of each genotype. In Experiment I, 3 untreated and 15 BHP-treated p53 (-/-) mice died before week

15. The causes of their deaths were mainly the development of lymphomas and sarcomas. In the 20 ppm BHP-treated case, p53 (-/-) mice demonstrated a significantly higher mortality rate than their p53 (+/+) and (+/-) counterparts ($P < 0.005$). In the 200 ppm BHP-treated group composed of p53 (+/+) and (+/-) mice, survival rates of both male and female p53 (+/-) mice were remarkably decreased at weeks 12 and later due to hemorrhaging into the abdominal cavity caused by the rupturing of hepatic vascular tumors. On the other hand, all the p53 (+/+) mice survived.

In Experiment II, the mortality of 20 ppm BHP-treated p53 (+/-) mice was significantly higher than in both male and female control p53 (+/-) and BHP-treated p53 (+/+) mice ($P < 0.05$).

Incidence of HSs. The observed incidences of HSs are summarized in Table II. In the short-term experiment, at 15 weeks HSs were only observed in 20 ppm BHP-treated p53 (-/-) mice, the incidence of this being significantly higher than the lack of it in p53 (+/+) and (+/-) mice of the same dose group ($P < 0.001$) and untreated (-/-) mice ($P < 0.05$) (Table II, note a).

In the long-term experiment, the development of HS in BHP-treated p53 (+/-) female mice was significantly increased as compared with BHP-treated (+/+) mice and untreated (+/-) mice ($P < 0.05$ and < 0.005) (Table II, note b). Similarly, in male mice, the incidence was higher in BHP-treated p53 (+/-) mice than in treated (+/+) mice, but without statistical significance. Additionally, in both p53 (+/+) and (+/-) mice, the incidences of HS were higher in female mice than in their male counterparts, but statistical significance was again not attained.

Pathological analysis. Macroscopically, HSs sometimes appeared as yellow-white nodules raised above the surface of the liver. However, some cases were detected by histological observation without any gross findings. In two p53 (-/-) mice with systemic invasion of HSs, markedly enlarged livers and spleens were grossly observed.

Microscopically, HSs were composed of large cells with pale eosinophilic cytoplasm and pleomorphic or folded nuclei. Multinucleated giant cells were sometimes present in the neoplastic lesions (Fig. 2A) and, as previously reported, the extramedullary hematopoiesis of erythrocytic series cells was common (23). Vascular invasion of tumor cells was common in the liver, but metastasis to other organs was relatively infrequent. In two p53 (-/-) mice, HS cells were found in multiple organs, including the spleen, lungs and blood vessels. It was possible to classify hepatic HSs into diffuse and nodular types with reference to their proliferation patterns. In the former, HS cells diffusely infiltrated hepatic sinusoids, forming lesions of various sizes in all liver lobes (Fig. 2A). With the latter pattern, HS cells were gathered in multicentric groups (Fig. 2B). The diffuse pattern was observed in 4 of p53 (-/-) mice bearing HS in the short-term experiment and 2 female (+/-) mice in the long-term experiment. All other lesions were of the nodular pattern. HSs need to be differentiated from lymphomas (24,25). We therefore paid special attention to the differential diagnosis as lymphoma is the most frequent spontaneous tumor in p53 knockout mice (2,13,14,17). Immunohistochemical analyses, using the murine macrophage surface marker F4/80, were used to determine tumor origin. Clusters of positively stained tumor cells were found (Fig. 2C),

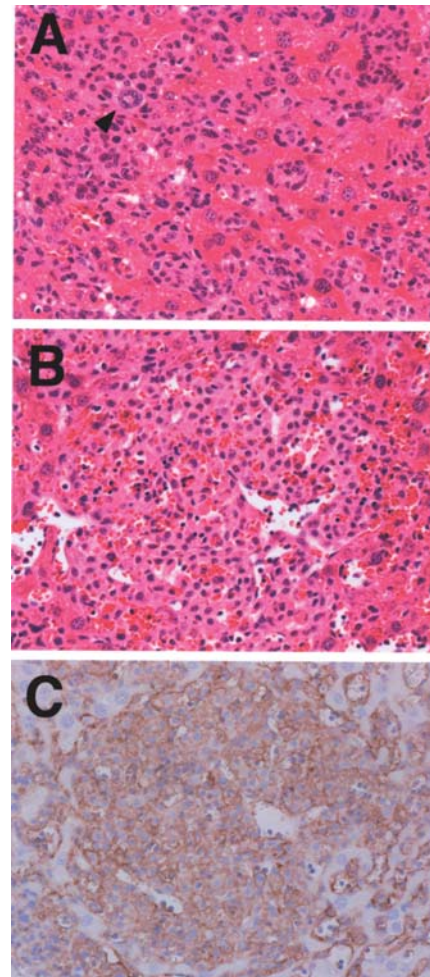


Figure 2. Histopathology of histiocytic sarcomas in p53 knockout mice. (A) Histiocytic sarcoma diffusely infiltrating in sinusoids with multinucleated giant cell formation (arrowheads). H&E staining. (B and C) Serial sections of a histiocytic sarcoma growing as a nodule. (B) H&E staining. (C) Immunohistochemistry for F4/80. Original magnification, x200.

although F4/80 yielded various degrees of tumor cell staining within and between tumors. Additionally, HS cells did not react to either of the B- and T-lymphocyte markers, CD45R and CD3, respectively.

Alterations of the p53 gene in HSs. The 12 HSs which developed in BHP-treated mice, 2 from p53 (+/+) and 10 from (+/-) animals, were subjected to p53 gene analysis. p53 gene mutations were identified in 8 of the 12 (66.7%) lesions. Several mutations were concurrently found in 6 of the HSs and a total of 21 mutations, concentrated in exons 5 and 8, were detected (Table III). Three (in tumors HS01, HS04 and HS10; all of them with other mutations) were silent and 2 (tumor IDs: HS04 and HS11) were of nonsense type. All the others were missense mutations. Most gave rise to transitions ($19/21 = 90.5\%$), with G:C→A:T transitions accounting for 61.9% (13/21) of all the mutations in HSs. In HS04, a multicentric nodular lesion, 2 and 3 mutations were found in exons 5 and 8, respectively, indicating a heterogeneous cell population. To determine the alleles in which the mutations occurred, LA-PCR was performed to specifically amplify the wild-type allele of the p53 gene. Only one HS (tumor ID: HS01) from p53 (+/-) mice was successfully amplified for

Table III. p53 gene alterations identified in histiocytic sarcomas.

No.	Tumor ID	Sex	p53 genotype	p53 LOH	p53 gene mutation						
					Exon	Mutated allele	Codon	Nucleotide change	Amino acid change	Event	
1	HS 03	F	(+/+)	ND	(-)						
2	HS 04	F	(+/+)	ND	5		153	CGT→CAT	Arg→His	Transition	
							155	CGC→CGT	Silent	Transition	
					8		283	GAA→TAA	Glu→Stop	Transversion	
							297	CCC→TCC	Pro→Ser	Transition	
							301	GCA→ACA	Ala→Thr	Transition	
3	HS 01	M	(+/-)	(-)	5	Wild	179	TGC→TGT	Silent	Transition	
					7	Mutant	251	ATC→ACC	Ile→Thr	Transition	
					8	Unknown	269	GTT→GCT	Val→Ala	Transition	
4	HS 02	M	(+/-)	(-)	(-)						
5	HS 05	F	(+/-)	(-)	(-)						
6	HS 06	F	(+/-)	(-)	8	Unknown	278	GAC→GGC	Asp→Gly	Transition	
7	HS 07	F	(+/-)	(-)	5	Wild	152	AGC→GGC	Ser→Gly	Transition	
							172	CGC→CAC	Arg→His	Transition	
					8	Unknown	278	GAC→GGC	Asp→Gly	Transition	
							293	TGC→CGC	Cys→Arg	Transition	
8	HS 08	F	(+/-)	(-)	(-)						
9	HS 09	F	(+/-)	(-)	5	Wild	155	CGC→TGC	Arg→Cys	Transition	
					8	Unknown	292	CTT→TTT	Leu→Phe	Transition	
							301	GCA→ACA	Ala→Thr	Transition	
10	HS 10	F	(+/-)	(-)	5	Wild	153	CGT→CAT	Arg→His	Transition	
					8	Unknown	296	CTG→CTA	Silent	Transition	
11	HS 11	F	(+/-)	(-)	5	Wild	155	CGC→TGC	Arg→Cys	Transition	
					8	Unknown	283	GAA→TAA	Glu→Stop	Transversion	
12	HS 12	F	(+/-)	(-)	5	Wild	149	CCA→TCA	Pro→Ser	Transition	

exon 7, and no mutation was identified in the wild-type allele. The other microdissected samples could not be subjected to LA-PCR analysis due to poor preservation of genomic DNA in the paraffin-embedded samples. Primers for the PCR-SSCP analysis were used to amplify the wild-type allele of exon 5, and it was revealed that 5 of 7 HSs had missense mutations in exon 5 (tumor IDs: HS07, 09, 10, 11 and 12) of the p53 wild-type allele in p53 (+/-) mice. This indicated the anticipated complete loss of functional p53 protein.

Discussion

This study clearly demonstrated that p53 knockout mice are highly susceptible to BHP-induction of HS. Thus, in the short-term experiment, lesions were found in only BHP-treated p53 (-/-) mice, indicating that the complete loss of p53 dramatically accelerated the genesis of HS. The high susceptibility of p53 (-/-) mice was consistent with data for BHP-induced lung and hepatic vascular tumors (20), as well as stomach (21), esophageal (22), tongue (26), colon (27), and urinary bladder (28) carcinogenesis due to other carcinogens.

With long-term treatment of BHP, the significantly greater yield of HSs in p53 (+/-) mice as compared to p53 (+/+) mice again pointed to elevated susceptibility, especially in females. Furthermore, PCR-SSCP and sequencing analysis revealed a high frequency of p53 gene mutations, with the expected complete loss of functional protein in the BHP-induced HS. This is in agreement with earlier reports that the accelerated induction of neoplasms by chemical carcinogens in p53 (+/-) mice is accompanied by the loss of functional p53 through mutational inactivation (20,22,29,30) or the loss of heterozygosity (31). In accordance with Knudson's two-hit hypothesis, the data point to a necessity for a 'second hit' to inactivate the residual normal allele.

To the best of our knowledge, this is the first report to directly demonstrate the relevance of p53 to murine HS development. The tumor suppressor p53 prevents the propagation of genetically damaged cells by the transcriptional regulation of downstream genes (1). As well, increased development of HS has been described in animals genetically engineered for genes downstream of p53, such as p21-deficient and BAX/ARF double-deficient mice (8,10). Furthermore, Carrasco *et al*

demonstrated that p53 expression was frequently repressed, correlating with ARF promoter methylation in human HS (12). The fact that a deficiency of p53 accelerated the development of HSs is in line with our expectations.

BHP is a multi-organ genotoxic carcinogen which induces various types of tumors in rodents. Through oral administration, it induces lung and hepatic vascular tumors in mice (19). In addition to these tumors (20), though HS induction has not previously been reported in any rodents, including other genetically engineered mice (32,33), BHP unexpectedly induced HSs in p53 knockout mice. Target organs may become altered in genetically engineered mice, reflecting organ-specific sensitivity as indicated for rasH2 mice (34). Thus, p53 knockout mice might not be expected to share the same target organs as wild-type mice because the susceptibility of p53 (+/-) mice to chemical carcinogens is organ-dependent (20,35,36). Ozaki *et al* reported that, when comparing susceptibilities of rasH2 and p53 (+/-) mice under the same experimental conditions, different types of tumors - lung tumors in rasH2 mice and hepatic hemangiomas in p53 (+/-) mice - were predominantly induced by urethane treatment (37). When genetically engineered mice are used in carcinogenic studies, we must take this characteristic into consideration. Besides the deficiency of p53, the genetic background of p53 knockout mice might partly affect the development of HS. It appears that the spontaneous development of HS is more frequent in the C57BL/6J strain than in other common strains. In all cases, the development of HS is infrequent until 18 months of age, but rapidly increases in mice older than 24 months, specifically in C57BL/6J mice (4,5,7,24).

In conclusion, this study clearly shows an elevated susceptibility of nullizygous and heterozygous p53 knockout mice to BHP-induction of HS, with the direct involvement of p53 gene mutations. Our data add to the understanding of the carcinogenic susceptibility of p53 knockout animals and aid in the elucidation of the mechanisms involved in HS development.

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