

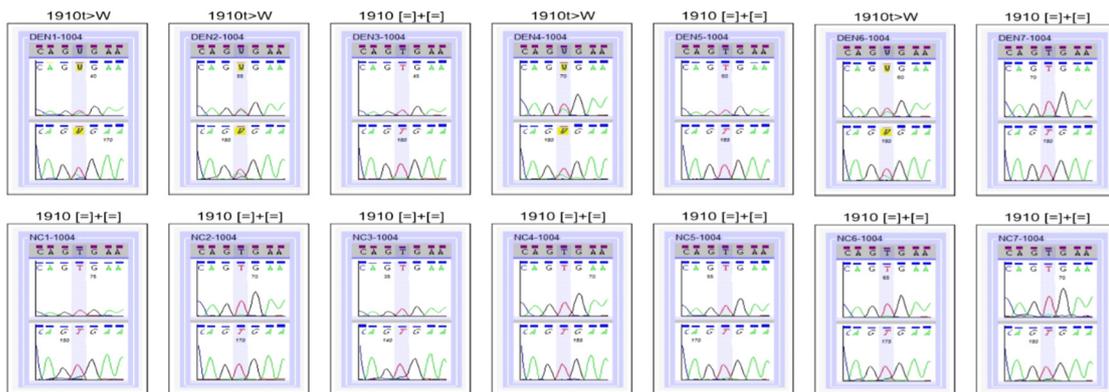
**Supplementary Table S1. Detailed information of all liver specimens from vehicle- and DEN-treated mice.**

Specimen ID	Genotype	Base Position	Genotype Result	User Edit	QV	Base Position Coverage	Amplicon ID	Layer ID	ROI ID	ROI Position	Genotype Comments
DEN1-1004	1910t>W	1910	-	no	65	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
DEN2-1004	1910t>W	1910	-	no	65	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
DEN3-1004	[=]+[=]	1910	-	no	65	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
DEN4-1004	1910t>W	1910	-	no	52	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
DEN5-1004	[=]+[=]	1910	-	no	65	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
DEN6-1004	1910t>W	1910	-	no	65	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
DEN7-1004	[=]+[=]	1910	-	no	65	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
NC1-1004	[=]+[=]	1910	-	no	49	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
NC2-1004	[=]+[=]	1910	-	no	65	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
NC3-1004	[=]+[=]	1910	-	no	47	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
NC4-1004	[=]+[=]	1910	-	no	65	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
NC5-1004	[=]+[=]	1910	-	no	65	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
NC6-1004	[=]+[=]	1910	-	no	46	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	
NC7-1004	[=]+[=]	1910	-	no	40	2X	Mus_Braf_V637 E	NP_647455.3	NP_647455.3_	2077 region_1	

**A**

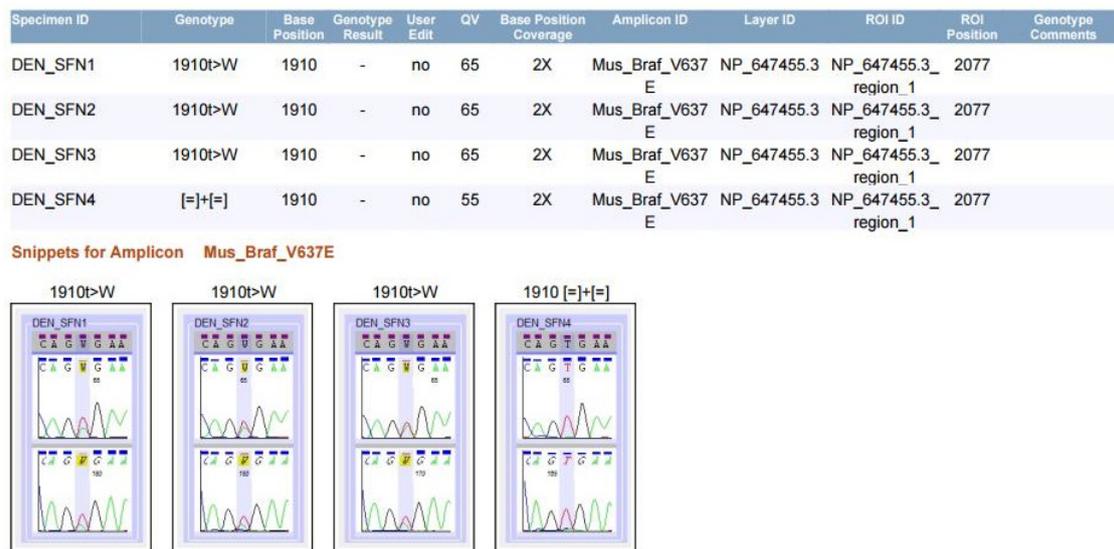
Primer name	Sequence	PCR size (bp)	Tm.
Mus_Braf_V637E_2F_M13F	<b>GTA<sup>+</sup>AAACGAGCGGCCAGT<sup>+</sup>TTCTCTTACTTACTGCAC</b>	290	60
Mus_Braf_V637E_R	<b>GCAATTATGCCTGGCTTACA</b>		

**B**

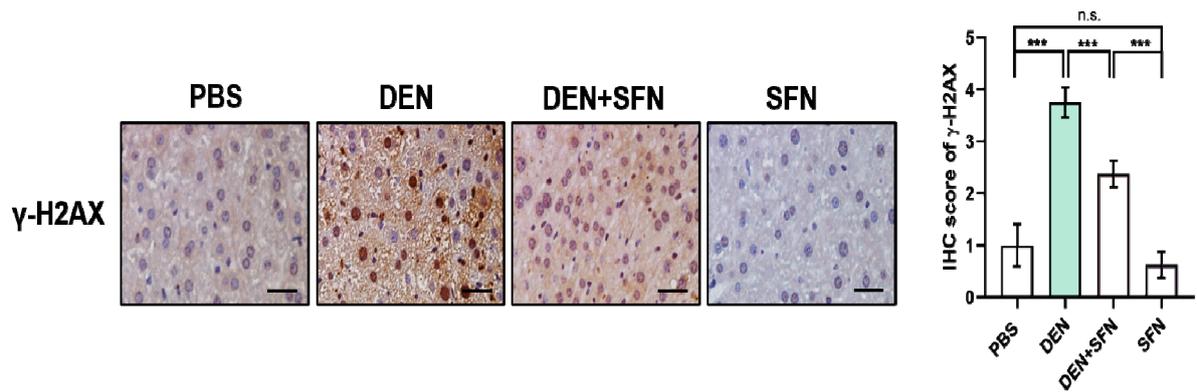


**Supplementary Figure S1. *B-Raf*<sup>V637E</sup> mutation detected in DEN-induced liver tumors.** Total RNA was extracted from vehicle- or DEN-treated liver tissue with the DNeasy® Blood & Tissue Kit, and converted to cDNA using reverse transcriptase

following the standard procedure. The cDNA region bearing murine *B-Raf* gene was amplified using PCR with HiPi Tag polymerase. The PCR products were purified by the QIAquick Purification Kit. **(A)** Designed primers for sequencing are listed, GTAAAACGACGGCCAGT sequence represents M13F primer. **(B)** Sequences of the genetic variants in murine *B-Raf*<sup>V637E</sup> were detected by capillary electrophoresis.



**Supplementary Figure S2. *B-Raf*<sup>V637E</sup> mutation detected in SFN-treated DEN-induced liver tumors.** Sequencing was performed and cleaned up with the BigDye® Terminator v3.1 Cycle Sequencing Kit. Sequences were detected and analyzed by ABI PRISM 3730XL Analyzer. Sequences of the genetic variants in murine *B-Raf*<sup>V637E</sup> were detected by capillary electrophoresis. Three out of 4 cases of *B-Raf*<sup>V637E</sup> mutation (labelled by yellow color) were detected by capillary electrophoresis in SFN plus DEN-induced liver tumors.



**Supplementary Figure S3. Immunohistochemical analysis of  $\gamma$ -H2AX in DEN-induced murine hepatocarcinogenesis with and without SFN administration.** The paraffin sections of liver tissues were subjected to immunohistochemical staining with an antibody against  $\gamma$ -H2AX. The IHC score was analyzed by the image processing program Image ,J and results are shown as the mean  $\pm$  SD of 4 samples for each group. \*\*\* $p < 0.001$ ; n.s.: non-significant. Representative images of stained sections are displayed. Scale bar, 100  $\mu$ m.