

Table S1. Primers for mouse genotyping

| <i>Genotyping for Keap1^{A/A} background mouse</i> | |
|---|--------------------------------|
| Primers | Sequence (5' - 3') |
| 5-cko 4int1 | GCACATCCTTCATCTCTCCGCACTGGGGAG |
| 3-Kp1-4Ex | CCTCCGTGTCAACATTGGCGCGACTAG |
| R260-EGFP | GACTTGAAGAAGTCGTGCTGCTTCATGTG |
| <i>Genotyping for Keap1^{B/B} background mouse</i> | |
| Keap1-BF-F | CGAGGAAGCGTTTGCTTTAC |
| BF-R1 | AGCCCCCTGCTGCATAGATAC |
| NeoI-3R | GAGTCACCGTAAGCCTGGTC |
| Keap1-4F | GAGTCCACAGTGTGTGGCC |
| <i>Genotyping for Rosa^{NIC/NIC} background mouse</i> | |
| IMR0883 | AAAGTCGCTCTGAGTTGTTAT |
| IMR8038 | TAAGCCTGCCCAGAAGACTC |
| IMR8039 | GAAAGACCGCGAAGAGTTTG |
| PGK-3740 | GATGTGGAATGTGTGCGAGGCCAGAGGC |
| NICD-5477 | GATTGTCGTCCATCAGAGCACCATCTGAGG |
| <i>Genotyping for Cre background mouse</i> | |
| Cre1 | ACGTTCAACGGCATCAACGT |
| Cre2 | CTGCATTACCGGTCGATGCA |

Table S2. PCR programs for mouse genotyping

| <i>Keap1^{A/A} background mouse</i> | | | |
|--|---------|--------|--------------------------------|
| Step # | Temp °C | Time | Note |
| 1 | 95 | 1 min | - |
| 2 | 95 | 30 sec | - |
| 3 | 68.5 | 30 sec | - |
| 4 | 72 | 30 sec | repeat steps 2-4 for 35 cycles |
| 5 | 72 | 1 min | - |
| 6 | 4 | - | hold |
| Product; Flox A : ~350 bp, Disrupted : ~550 bp, Wt : ~250 bp | | | |
| <i>Keap1^{B/B} background mouse</i> | | | |
| Regular genotyping Primer: Keap1 BF-F and BF-R1 | | | |
| Step # | Temp °C | Time | Note |
| 1 | 98 | 3 min | - |

| | | | |
|--|----------------|-------------|--------------------------------|
| 2 | 95 | 30 sec | - |
| 3 | 68 | 30 sec | - |
| 4 | 72 | 30 sec | repeat steps 2-4 for 35 cycles |
| 5 | 4 | - | hold |
| Product; Flox B : 445 bp, Wt : 267 bp | | | |
| Excision confirmation Primer: Keap1-4F, BF-F and NeoI-3R | | | |
| Step # | Temp °C | Time | Note |
| 1 | 95 | 1 min | - |
| 2 | 95 | 30 sec | - |
| 3 | 68.5 | 30 sec | - |
| 4 | 72 | 30 sec | repeat steps 2-4 for 35 cycles |
| 5 | 72 | 1 min | - |
| 6 | 4 | - | hold |
| Product; Undisrupted : 383 bp, Disrupted : 288 bp, Wt : ~205 bp | | | |
| <i>Rosa^{NIC/NIC}</i> mouse (https://www.jax.org/strain/008159) accessed 8/21/2023 | | | |
| Regular genotyping Primers: IMR0883, IMR8038 and IMR8039 | | | |
| Step # | Temp °C | Time | Note |
| 1 | 94 | 3 min | - |
| 2 | 94 | 30 sec | - |
| 3 | 54 | 1 min | - |
| 4 | 72 | 1 min | repeat steps 2-4 for 35 cycles |
| 5 | 72 | 2 min | - |
| 6 | 10 | - | hold |
| Product; Tg : 320 bp, Wt : 235 bp | | | |
| <i>Rosa^{NIC}</i> active allele by Cre Primers; IMR0883, NICD-5477 and PGK-3740 | | | |
| Step # | Temp °C | Time | Note |
| 1 | 95 | 1 min | - |
| 2 | 95 | 30 sec | - |
| 3 | 65.4 | 30 sec | - |
| 4 | 72 | 45 sec | repeat steps 2-4 for 35 cycles |
| 5 | 72 | 2 min | - |
| 6 | 4 | - | hold |
| Product; Cre-Active : 650 bp, Cre-Inactive : 550 bp | | | |
| <i>Cre transgenic allele</i> | | | |
| Step # | Temp °C | Time | Note |
| 1 | 94 | 1 min | - |
| 2 | 94 | 30 sec | - |

| | | | |
|---------------------------------|----|--------|--------------------------------|
| 3 | 60 | 30 sec | - |
| 4 | 72 | 30 sec | repeat steps 2-4 for 35 cycles |
| 5 | 4 | - | hold |
| Product; <i>Cre-Tg</i> : 355 bp | | | |

Table S3. Primers for transgene confirmation

| Primers | Sequence (5' - 3') |
|----------------|------------------------|
| CAGGs | CTCTAGAGCCTCTGCTAACC |
| PB-6679080a1-R | AGGCGTCCTTCCTTATATGCTA |
| pCAG-F | GCAACGTGCTGGTTATTGTG |
| PB-6754832a1-R | GTTGAAACTGAGCGAAAAAGGC |

Table S4. PCR program for HTI-transgene confirmation

Primer for DA-*Nrf2*: pCAG-F and PB-6754832a1-R

Primer for *Nqo1*: CAGGs and PB-6679080a1-R

| Step # | Temp °C | Time | Note |
|--------|---------|--------|--------------------------------|
| 1 | 94 | 1 min | - |
| 2 | 95 | 30 sec | - |
| 3 | 65.7 | 30 sec | - |
| 4 | 72 | 20 sec | repeat steps 2-4 for 35 cycles |
| 5 | 4 | - | hold |

Product; DA-*Nrf2* : 294 bp, *Nqo1*: 401bp

Table S5. Primers used in mutagenesis

| Primer | Sequence (5' - 3') |
|-------------------|--|
| 5-Nrf2-XN-ATG | CTAGTCTAGACATATGATGGACTTGGAGTTGCCACCGCC |
| 5-Nrf2 DLG-A SfcI | AGGACTACAGTCCCAGCAGGACATGGATTTGATTGACATCGCATGGAGGGCAGCAAT AGCAGCTGCAGTAAGTCGAGAAGTGTTTGACTTTAGTCAG |
| 5-KA-Nrf2 BglII | ATAGATCTTGGAGTAAGTCGAGAAGTGTTTGACTTTAGTCAGCGACAGGCAGACTATG AGCTGGAAGCACAGGCAGCACTCGAAGCGGAAAGACAAGAGC |
| 5'-PstI-KA-Nhe2 | GAGCAACTGCAGGCGGAACAGGAGGCGG |
| 3'-PstI-KA-Nhe2 | TTCCGCCTGCAGTTGCTCTTGTCTTTCC |
| 3-BamHI | CCTGGGAGTAGCTGGCGGATCCACTG |
| 3-KA Nrf2 EcoRI | AGGAATTCTCCTGTTTCTTCATCCAGTTGAAACTGAGCGAAAAAGGCCGCCTCCTGTTC CGCCTGGAGTTGCTCTTGTCTTTCCGCTTCGAGTGCTGCCT |

Table S6. Antibodies used in experiments

| Target Protein | Provider | Dilution |
|--|-----------------------------|-----------------|
| Nrf2 | Invitrogen PA5-27882 | 2,000 |
| LaminB1 | Proteintech 12987-1-AP | 5,000 |
| Nqo1 | Abcam ab2346 | IB, IHC: x 500 |
| Gclc | Proteintech 12601-1-AP | 2,000 |
| GstA1-5 | Invitrogen PA5-79335 | 2,000 |
| Acc1 | Proteintech 21923-1-AP | 1,000 |
| Fasn | Proteintech 10624-2-AP | 1,000 |
| Keap1 | Original | 3,000 |
| Luciferase | Novus Biological NB100-1677 | 1,000 |
| Rabbit anti-Goat IgG (H+L)-HRP | Invitrogen 31402 | 10,000 |
| Goat Anti-Rabbit IgG (H + L)-HRP | BIO RAD 1706515 | 3,000 |
| Horse anti-Goat IgG (H+L), Biotinylated | Vector Laboratories BA-9500 | x 200 |

Supplemental Figure S1. *Keap1* and flox mutant gene structure and primer positions for genotyping. (A) *Kp1^{A/A}* flox allele. (B) *Kp1^{B/B}* flox allele. Grey and blue boxes indicate non-coding and coding region of each exon of *Keap1* gene. The triangles show position of flanking lox sequences in mutant mice.

Supplemental Figure S2. Representative results of confirmative genotyping. The positive controls were from tail DNA isolated from each heterozygote of *Kp1^{A/+}*, *Kp1^{B/+}*, *Rosa^{NIC/+}* and *Rosa^{NIC/+}::AdiCre* mice for *Keap1 A flox* (top), *Keap1 B flox* (second), *Rosa* (third), and *Adipoq Cre* (bottom) genotyping, respectively.

Supplemental Figure S3. Confirmation of transgene by HTI. The primer set positions for *Nqo1* (CAGGs, PB-6679080a-1-R) and *DA-NRF2* expression vectors (pCAG-F, PB-6754832a1-R) are depicted in (A). The representative PCR results are shown in (B). White, steel grey and mercury boxes show the representative result of transgene detection from *pCAG Mock*, *pCAG Nqo1* and *pCAG DA-Nrf2* HTI mice (N=4), respectively. 1μL of template DNA was utilized as control which includes 10ng of each plasmid DNA mixed with wild-type tail genomic DNA treated as per usual genotyping.

Supplemental Figure S4. Immunohistochemical analysis of NQO1 expression in the HTI-liver of *Rosa^{NIC/NIC}::AdiCre* mice. 5-week old male *Rosa^{NIC/NIC}::AdiCre* mice were hydrodynamically injected with *pCAG Mock* (A) or *pCAG Nqo1* (B) through the tail vein. Five weeks following HTI and feeding with HFD, livers were isolated from the mice and its sections were prepared and analyzed with anti-NQO1 antibody immunohistochemically. NQO1 derived from *pCAG Nqo1* HTI liver was stained strongly by the precipitated DAB reaction product. Scale bar: 100 μm.

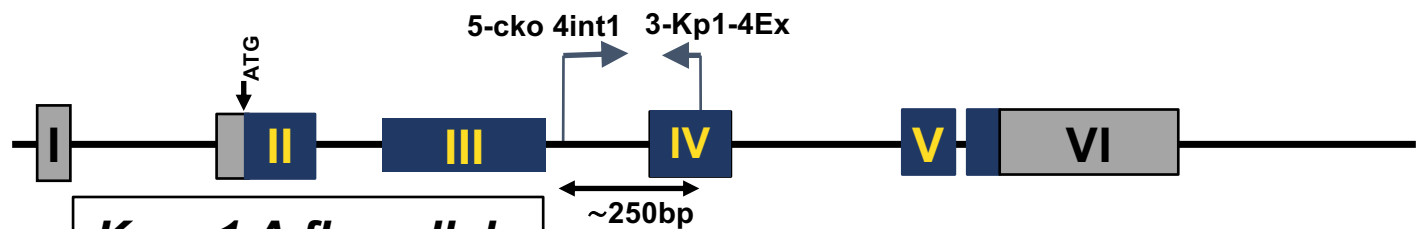
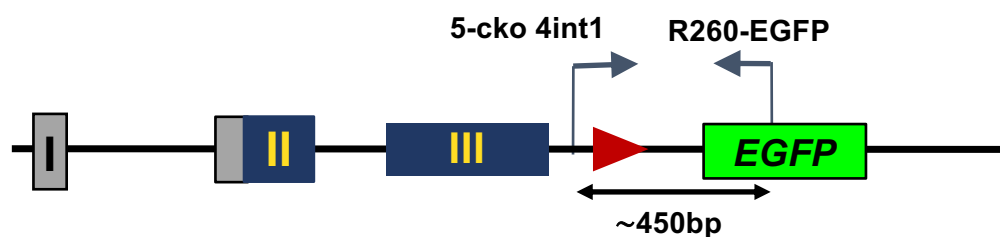
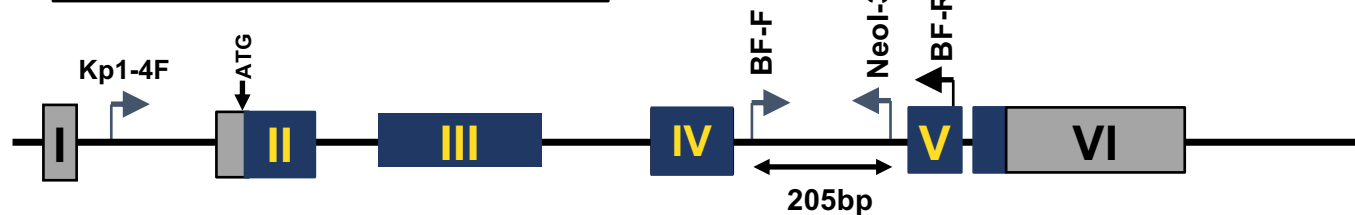
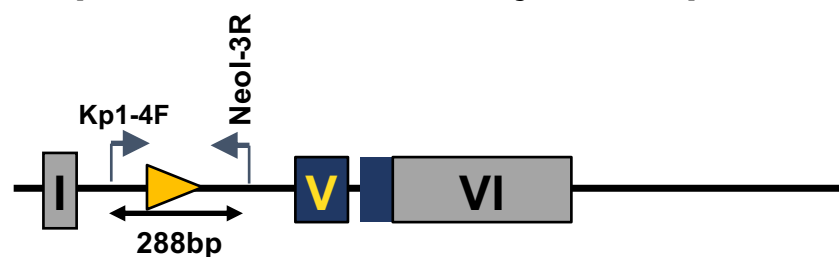
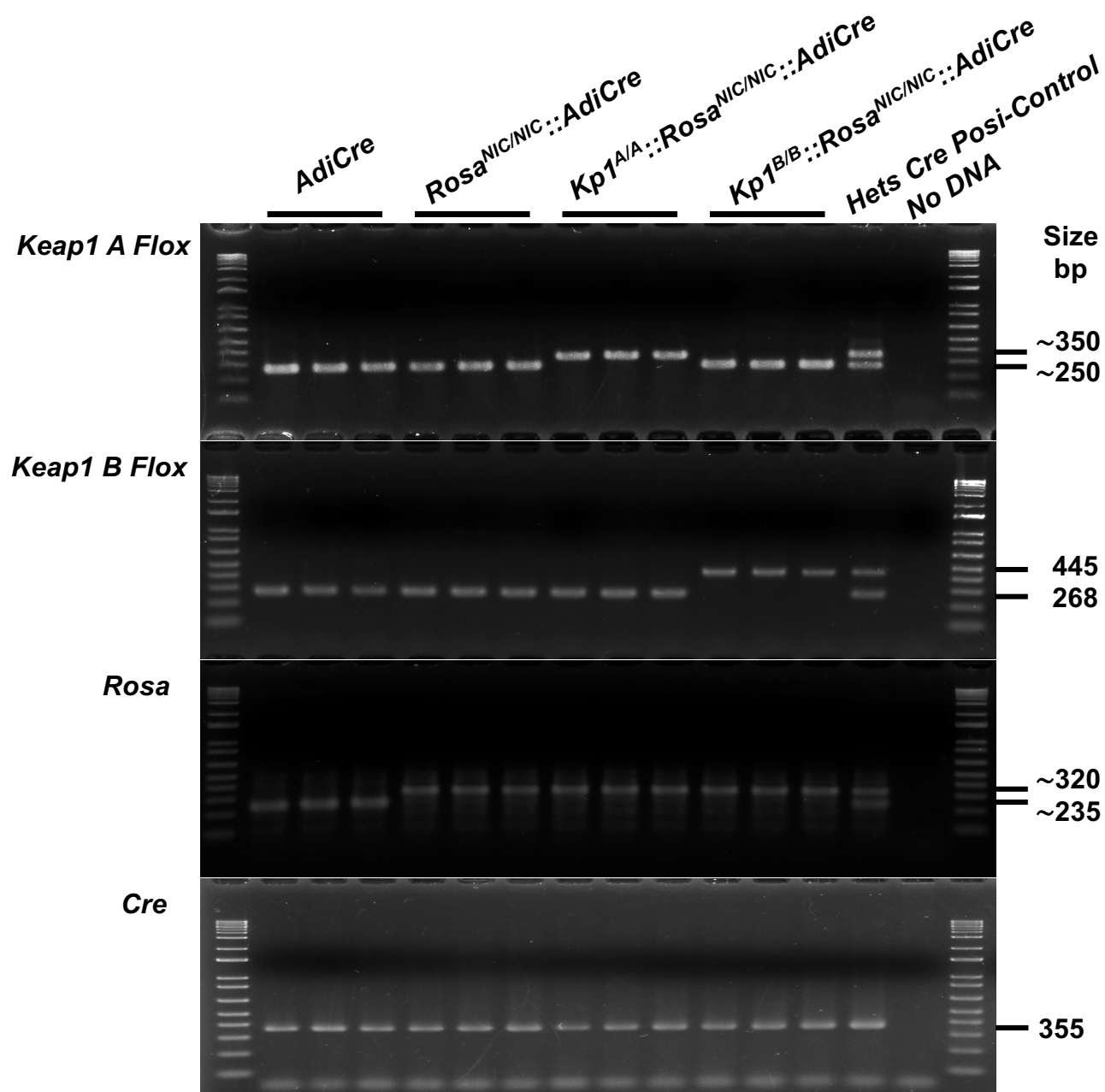
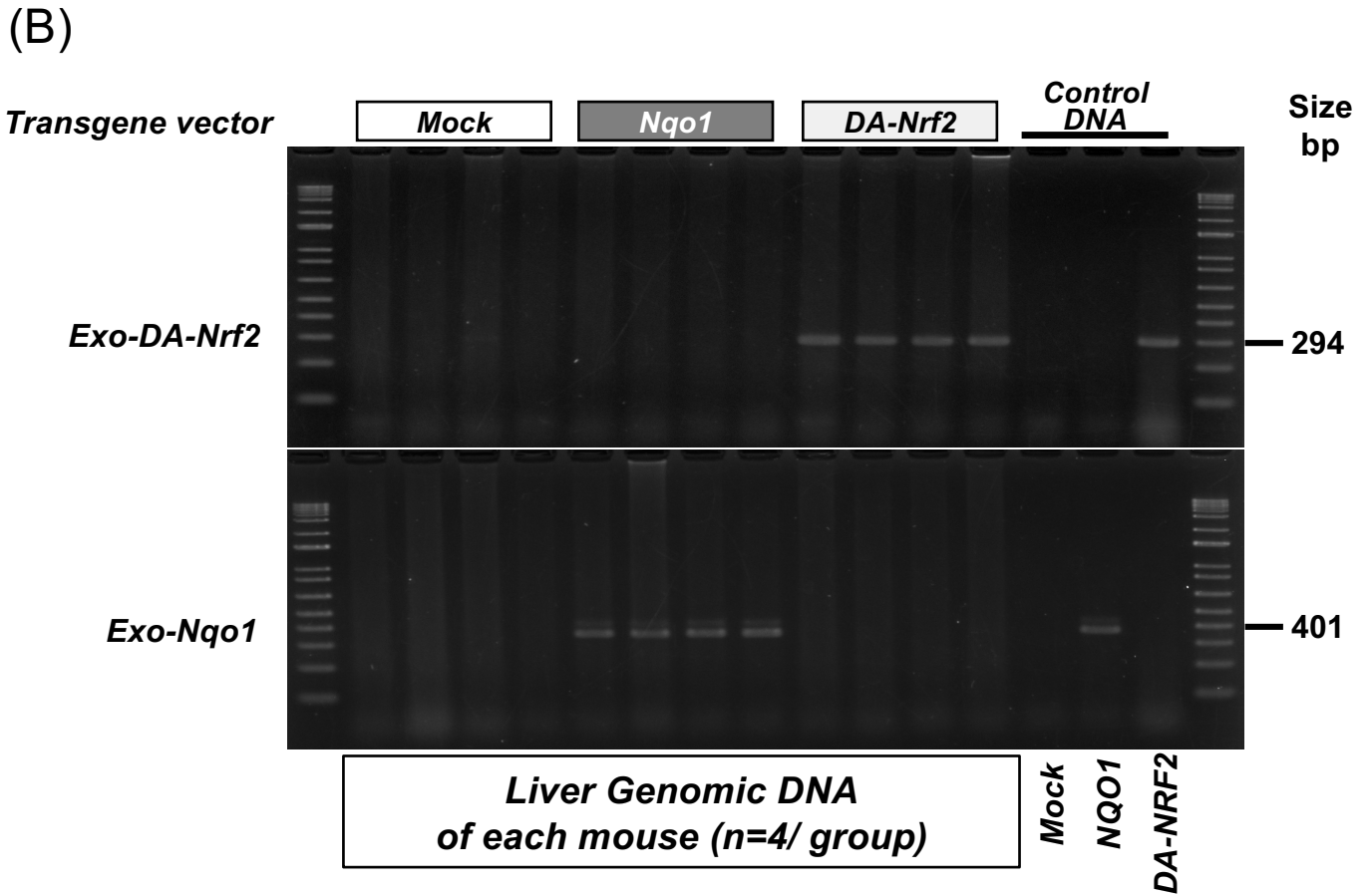
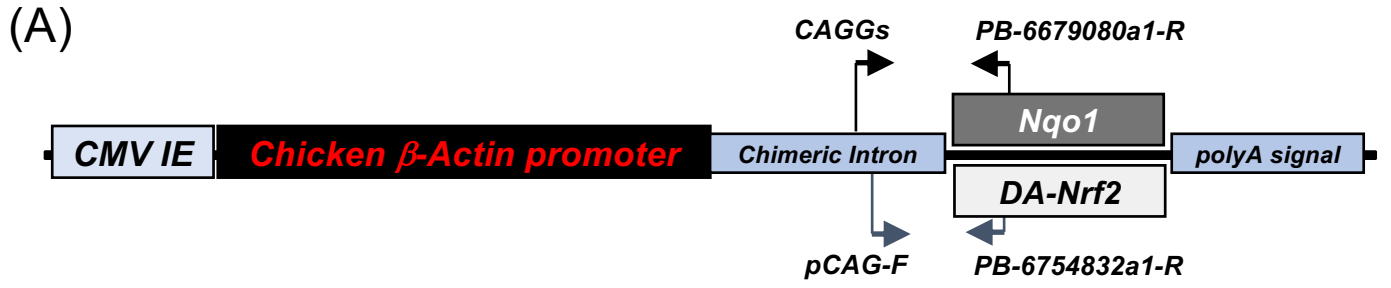
(A) *Keap1* Wild type allele***Keap1 A* excision allele by Cre Expression****(B) *Keap1* Wild type allele*****Keap1 B* excision allele by Cre Expression**

Figure S2

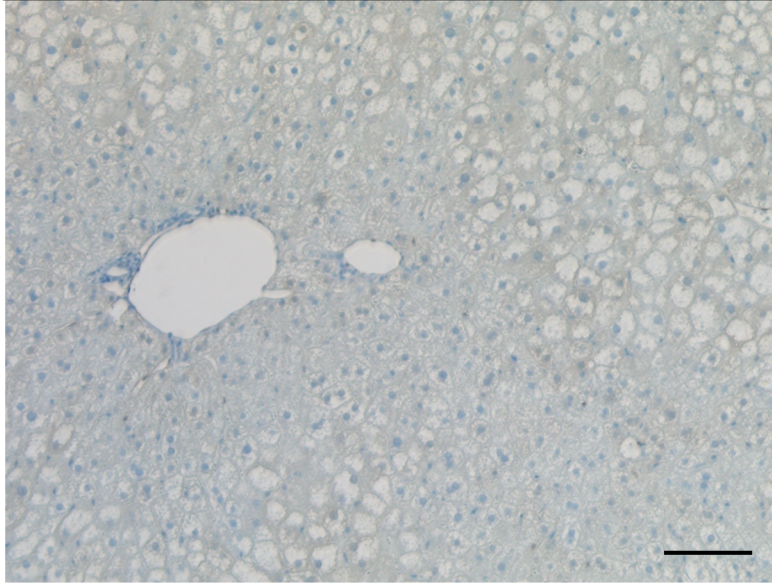


**Liver Genomic DNA
of each mouse (n=3/ group)**



(A)

pCAG Mock



(B)

pCAG Nqo1

