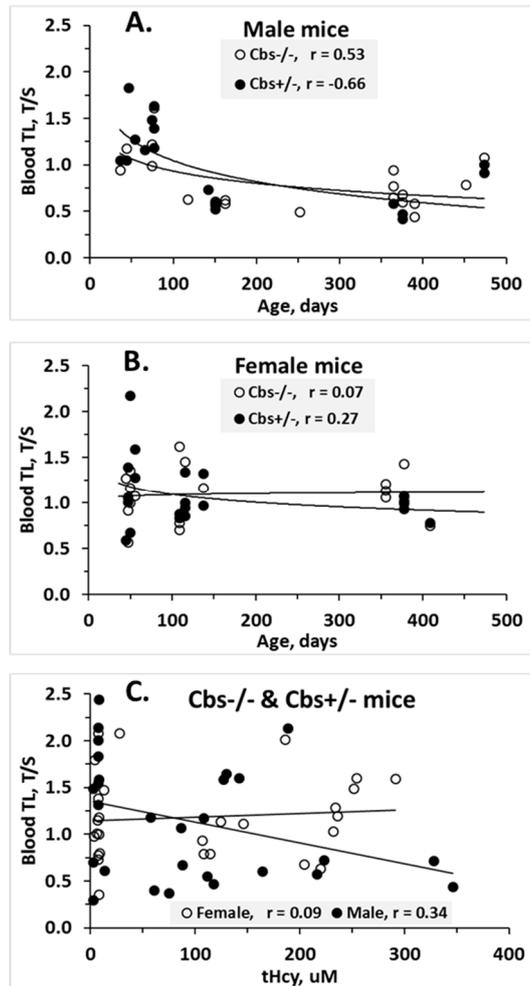
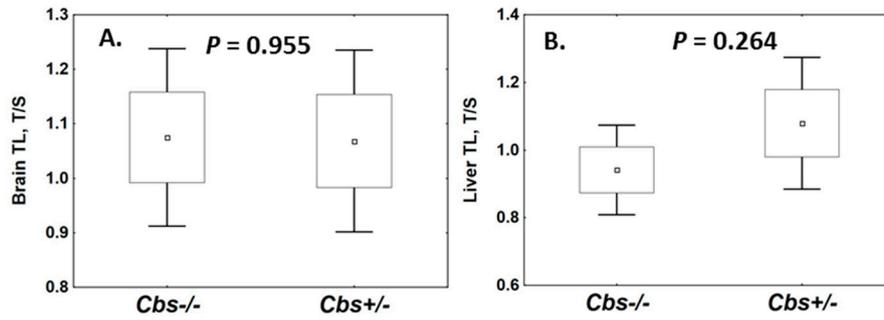


## Supplementary Material

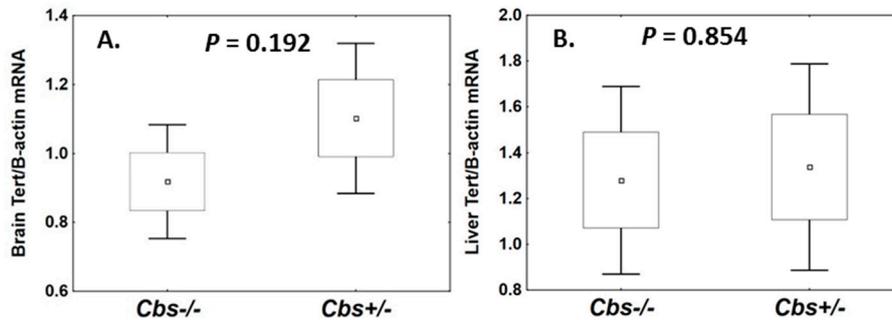
Supplementary Figure S1.  
Supplementary Figure S2.  
Supplementary Figure S3.  
Supplementary Figure S4.  
Supplementary Table S1.  
Supplementary Table S2.  
Supplementary Table S3.  
Supplementary Table S4.  
Supplementary Table S5.  
Supplementary Table S6.  
Supplementary Table S7.  
Supplementary Table S8.



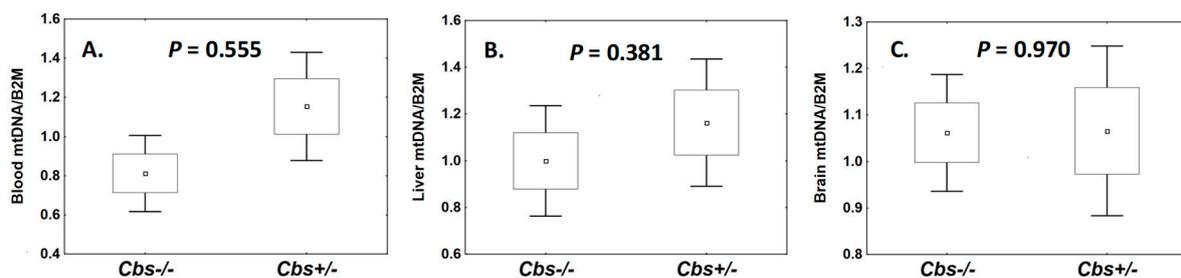
**Supplementary Figure S1.** Relationships between blood TL, age, and tHcy in *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice. Blood TL vs. age in (A) female and (B) male mice. (C) Blood TL vs. tHcy.



**Supplementary Figure S2.** Telomere length (TL) in brains (A) and livers (B) of *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice. TL was quantified by qPCR in *Cbs*<sup>-/-</sup> mice ( $n = 21$ ) and sex- and age-matched control *Cbs*<sup>+/-</sup> siblings ( $n = 22$ ) as described in the Materials and Methods. Box and whiskers represent the mean  $\pm$  SEM and the mean  $\pm$  95% CI, respectively.



**Supplementary Figure S3.** *Tert* mRNA levels in brains (A) and livers (B) of *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice. *Tert* mRNA was quantified by qPCR in *Cbs*<sup>-/-</sup> mice ( $n = 21$ ) and sex- and age-matched control *Cbs*<sup>+/-</sup> siblings ( $n = 22$ ) as described in the Materials and Methods. Box and whiskers represent the mean  $\pm$  SEM and the mean  $\pm$  95% CI, respectively.



**Supplementary Figure S4.** mtDNA levels in blood (A), livers (B), and brains (C) of *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice. mtDNA was quantified by qPCR in *Cbs*<sup>-/-</sup> mice (blood,  $n = 40$ ; liver and brain,  $n = 21$ ) and sex- and age-matched control *Cbs*<sup>+/-</sup> siblings (blood,  $n = 40$ ; liver and brain,  $n = 22$ ) as described in the Materials and Methods. Box and whiskers represent the mean  $\pm$  SEM and the mean  $\pm$  95% CI, respectively.

Supplementary Table S1. Aging/senescence-related mRNAs in the liver of <i>Cbs<sup>-/-</sup></i> and <i>Cbs<sup>+/-</sup></i> mice stratified by sex and age.														
Genotype (n)	Sex (n)	Age*	<i>Pai-1/β-actin</i> mRNA			<i>p21/β-actin</i> mRNA			<i>Mcp1/β-actin</i> mRNA			<i>Il-6/β-actin</i> mRNA		
			Mean ± SD (n)	FC <sup>#</sup> ; <i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>	Mean ± SD (n)	FC <sup>#</sup> ; <i>P</i> <sub>genotype</sub>	FC <sup>#</sup> ; <i>P</i> <sub>sex</sub>	Mean ± SD (n)	FC <sup>#</sup> ; <i>P</i> <sub>genotype</sub>	FC <sup>#</sup> ; <i>P</i> <sub>sex</sub>	Mean ± SD (n)	FC <sup>#</sup> ; <i>P</i> <sub>genotype</sub>	FC <sup>#</sup> ; <i>P</i> <sub>sex</sub>
<i>Cbs<sup>-/-</sup></i> (18–20)	♀	Young	10.7 ± 6.4 (4)	10.7; <b>0.025</b>	13.4; 0.060	5.3 ± 3.2 (4)	7.6; <b>0.016</b>	0.87; 0.272	4.2 ± 2.2 (4)	11.0; <b>0.014</b>	2.0; 0.202	0.78 ± 1.11 (3)	4.6; 0.311	0.55; 0.500
		Old	3.1 ± 1.5 (6)	4.4; <b>0.006</b>	1.47; 0.318	6.3 ± 4.8 (6)	4.2; 0.059	1.75; 0.124	4.3 ± 1.8 (6)	1.4; 0.174	0.34; <b>0.004</b>	2.87 ± 1.50 (6)	0.8; 0.525	0.71; 0.252
	FC <sup>#</sup> ; <i>P</i> <sub>age</sub>		0.29; <b>0.021</b>			1.2; 0.589			1.0; 0.906			1.2; 0.072		
	♂	Young	1.3 ± 1.3 (3)	0.8; 0.571		10.4 ± 7.5 (2)	8.7; <b>0.005</b>		2.1 ± 1.0 (3)	3.0; <b>0.005</b>		1.43 ± 1.05 (3)	8.4; <b>0.063</b>	
		Old	4.5 ± 2.9 (5)	3.0; 0.106		3.1 ± 2.8 (6)	2.4; 0.123		12.7 ± 5.3 (6)	3.2; <b>0.034</b>		4.06 ± 1.98 (7)	1.0; 0.923	
	FC <sup>#</sup> ; <i>P</i> <sub>age</sub>		3.5; 0.132			0.30; 0.065			6.0; <b>0.013</b>			3.0; 0.066		
<i>Cbs<sup>+/-</sup></i> (20–23)	♀	Young	1.0 ± 0.9 (4)		0.59; 0.230	0.70 ± 0.37 (5)		0.58;0 .124	0.38 ± 0.12 (4)		0.49; 0.307	0.17 ± 0.10 (4)		0.29; <b>0.010</b>
		Old	0.7 ± 0.2 (5)		0.45; 0.216	1.50 ± 0.62 (4)		1.15;0 .676	3.12 ± 0.39 (5)		0.69; 0.715	3.42 ± 1.29 (5)		0.88; 0.796
	FC <sup>#</sup> ; <i>P</i> <sub>age</sub>		0.7; 0.384			2.1; <b>0.047</b>			8.2; <b>0.000</b>			20.1; <b>0.002</b>		
	♂	Young	1.7 ± 0.8 (7)			1.2 ± 0.5 (7)			0.78 ± 0.62 (7)			0.59 ± 0.24 (7)		
		Old	1.5 ± 1.4 (4)			1.3 ± 0.9 (7)			1.66 ± 1.58 (4)			3.90 ± 3.72 (6)		
	FC <sup>#</sup> ; <i>P</i> <sub>age</sub>		0.9; 0.784			1.1; 0.749			2.1; 0.217			6.6; <b>0.037</b>		

\* Young = 63- to 66-day-old; Old = 354- to 408-day-old. # FC, fold change.

Supplementary Table S2. Aging/senescence-related mRNAs in the brain of *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice stratified by sex and age.

Genotype (n)	Sex (n)	Age*	<i>Pai-1</i> /β-actin mRNA			<i>p21</i> /β-actin mRNA			<i>Mcp1</i> /β-actin mRNA			<i>Il-6</i> /β-actin mRNA			<i>Kl</i> /β-actin mRNA		
			Mean ± SD (n)	FC <sup>#</sup> ; <i>P</i> <sub>gen</sub>	<i>P</i> <sub>sex</sub>	Mean ± SD (n)	FC <sup>#</sup> ; <i>P</i> <sub>gen</sub>	FC <sup>#</sup> ; <i>P</i> <sub>sex</sub>	Mean ± SD (n)	FC <sup>#</sup> ; <i>P</i> <sub>gen</sub>	<i>P</i> <sub>sex</sub>	Mean ± SD (n)	FC <sup>#</sup> ; <i>P</i> <sub>gen</sub>	FC <sup>#</sup> ; <i>P</i> <sub>sex</sub>	Mean ± SD (n)	FC <sup>#</sup> ; <i>P</i> <sub>gen</sub>	FC <sup>#</sup> ; <i>P</i> <sub>sex</sub>
<i>Cbs</i> <sup>-/-</sup> (17-21)	♀ (8-11)	Young	1.38 ± 0.14 (3)	1.3; 0.404	0.167	1.87 ± 0.63 (4)	<b>1.9</b> ; <b>0.035</b>	0.206	0.66 ± 0.12 (3)	0.7; 0.259	0.266	1.54 ± 1.24 (4)	2.0; 0.286	1.9; 0.383	2.00 ± 0.34 (4)	1.4; 0.192	<b>2.5</b> ; <b>0.010</b>
		Old	0.97 ± 0.33 (5)	0.9; 0.361	0.623	1.83 ± 0.77 (7)	<b>2.9</b> ; <b>0.004</b>	0.360	1.03 ± 0.21 (7)	<b>1.4</b> ; <b>0.020</b>	0.622	1.34 ± 0.29 (7)	0.83; 0.417	<b>1.6</b> ; <b>0.032</b>	1.17 ± 0.56 (7)	1.4; 0.996	0.168
		<i>P</i> <sub>age</sub>	0.7; 0.093		1.0; 0.936		1.6; <b>0.02</b>		0.9; 0.678		<b>0.6</b> ; <b>0.022</b>						
	♂ (8-10)	Young	0.76 ± 0.60 (2)	0.5; 0.366		1.09 ± 0.80 (3)	0.750		0.89 ± 0.29 (3)	0.8; 0.421		0.81 ± 0.45 (3)	0.71; 0.520		0.80 ± 0.46 (3)	0.53; 0.138	
		Old	0.71 ± 0.54 (7)	0.7; 0.411		0.64 ± 0.46 (7)	0.818		1.16 ± 0.68 (5)	0.8; 0.773		0.83 ± 0.47 (7)	0.83; 0.495		0.80 ± 0.41 (7)	0.99; 0.465	
		<i>P</i> <sub>age</sub>	0.911		0.286		0.547		0.949		0.996						
<i>Cbs</i> <sup>+/-</sup> (19-22)	♀ (8-9)	Young	1.03 ± 0.63 (4)		0.527	0.96 ± 0.23 (4)		0.254	0.95 ± 0.37 (4)		0.599	0.77 ± 0.46 (4)		0.270	1.44 ± 0.69 (4)		0.878
		Old	1.05 ± 0.29 (5)		0.989	1.43 ± 0.66 (5)		<b>2.1</b> ; <b>0.036</b>	0.72 ± 0.08 (4)		0.052	1.61 ± 0.77 (5)		0.121	0.83 ± 0.28 (4)		0.947
		<i>P</i> <sub>age</sub>	0.951		0.231		0.273		0.098		0.156						
	♂ (10-13)	Young	1.42 ± 1.04 (6)			1.41 ± 0.69 (7)			1.07 ± 0.27 (6)			1.14 ± 0.53 (7)			1.50 ± 0.65 (7)		
		Old	1.05 ± 0.55 (4)			0.69 ± 0.28 (6)			1.41 ± 0.54 (6)			1.00 ± 0.37 (6)			0.81 ± 0.54 (6)		
		<i>P</i> <sub>age</sub>	0.7; 0.535		<b>0.5</b> ; <b>0.038</b>		1.3; 0.238		0.9; 0.596		0.5; 0.064						

\* Young = 63- to 66-day-old; Old = 354- to 408-day-old. # FC, fold change.

**Supplementary Table S3. Blood TL, plasma tHcy levels in *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice stratified by sex and age.**

Genotype ( <i>n</i> )	Sex ( <i>n</i> )	Age* ( <i>n</i> )	tHcy, $\mu$ M			TL, T/S		
			Mean $\pm$ SD	<i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>	Mean $\pm$ SD	<i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>
<i>Cbs</i> <sup>-/-</sup> (40)	♀ (20)	Young (12)	133 $\pm$ 64	<0.000	0.990	1.08 $\pm$ 0.31	0.820	0.507
		Old (8)	238 $\pm$ 33	<0.000	<b>0.044</b>	1.10 $\pm$ 0.19	0.374	<0.000
		<i>P</i> <sub>age</sub>	<b>0.002</b>			0.855		
	♂ (20)	Young (6)	133 $\pm$ 42	<0.000		1.19 $\pm$ 0.27	0.335	
		Old (14)	156 $\pm$ 96	<0.000		0.68 $\pm$ 0.17	0.615	
		<i>P</i> <sub>age</sub>	0.531			<0.000		
<i>Cbs</i> <sup>+/-</sup> (40)	♀ (20)	Young (14)	6.4 $\pm$ 1.8		0.180	1.12 $\pm$ 0.41		0.165
		Old (6)	8.7 $\pm$ 2.8		0.669	1.01 $\pm$ 0.18		<b>0.002</b>
		<i>P</i> <sub>age</sub>	0.096			0.564		
	♂ (20)	Young (10)	7.3 $\pm$ 0.2			1.34 $\pm$ 0.27		
		Old (10)	5.1 $\pm$ 5.4			0.64 $\pm$ 0.19		
		<i>P</i> <sub>age</sub>	0.272			<0.000		

\* Young = 36- to 115-day-old; Old = 117- to 473-day-old; ♀ *Cbs*<sup>-/-</sup>, ♂ *Cbs*<sup>-/-</sup>, ♀ *Cbs*<sup>+/-</sup> and ♂ *Cbs*<sup>+/-</sup>.

**Supplementary Table S4. Brain TL and *Tert* mRNA expression in *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice stratified by sex and age.**

Genotype ( <i>n</i> )	Sex ( <i>n</i> )	Age*	Rel. brain TL, T/S			Rel. brain <i>Tert</i> /β-actin mRNA		
			Mean $\pm$ SD	<i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>	Mean $\pm$ SD	<i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>
<i>Cbs</i> <sup>-/-</sup> (19)	♀ (10)	young	0.73 $\pm$ 0.09	0.527	0.159	1.29 $\pm$ 0.33	0.916	<b>0.017</b>
		old	0.95 $\pm$ 0.22	0.948	<b>0.010</b>	0.87 $\pm$ 0.35	0.275	0.352
		<i>P</i> <sub>age</sub>	0.095			0.154		
	♂ (9)	young	0.94 $\pm$ 0.22	0.504		0.55 $\pm$ 0.17	<b>0.035</b>	
		old	1.53 $\pm$ 0.34	0.155		0.78 $\pm$ 0.23	0.423	
		<i>P</i> <sub>age</sub>	0.063			0.184		
<i>Cbs</i> <sup>+/-</sup> (24)	♀ (10)	young	1.00 $\pm$ 0.78		0.746	1.27 $\pm$ 0.32		0.978
		old	0.96 $\pm$ 0.32		0.273	1.28 $\pm$ 0.61		0.067
		<i>P</i> <sub>age</sub>	0.916			0.977		
	♂ (14)	young	1.11 $\pm$ 0.24			1.26 $\pm$ 0.39		
		old	1.17 $\pm$ 0.26			0.66 $\pm$ 0.21		
		<i>P</i> <sub>age</sub>	0.702			<b>0.022</b>		

\* Young = 63- to 66-day-old; Old = 354- to 408-day-old.

**Supplementary Table S5. Liver TL and *Tert* mRNA expression in *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice stratified by sex and age.**

Genotype ( <i>n</i> )	Sex ( <i>n</i> )	Age*	Rel. liver TL, T/S			Rel. liver <i>Tert</i> /β-actin mRNA		
			Mean ± SD	<i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>	Mean ± SD	<i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>
<i>Cbs</i> <sup>-/-</sup> (19)	♀ (10)	young	1.08 ± 0.36	0.909	0.353	0.24 ± 0.10	0.264	<b>0.020</b>
		old	1.11 ± 0.30	0.186	<b>0.014</b>	2.22 ± 0.81	0.512	<b>0.029</b>
		<i>P</i> <sub>age</sub>	0.892			<b>0.001</b>		
	♂ (9)	young	0.81 ± 0.35	0.415		0.43 ± 0.03	0.859	
		old	0.75 ± 0.19	0.735		1.43 ± 0.47	0.428	
		<i>P</i> <sub>age</sub>	0.737			<b>0.008</b>		
<i>Cbs</i> <sup>+/-</sup> (24)	♀ (10)	young	1.12 ± 0.47		0.681	0.90 ± 0.79		0.318
		old	1.50 ± 0.58		<b>0.018</b>	2.89 ± 1.07		<b>0.012</b>
		<i>P</i> <sub>age</sub>	0.325			<b>0.010</b>		
	♂ (14)	young	1.01 ± 0.40			0.56 ± 0.14		
		old	0.78 ± 0.23			1.43 ± 0.39		
		<i>P</i> <sub>age</sub>	0.228			<b>0.000</b>		

\* Young = 63- to 66-day-old; Old = 354- to 408-day-old.

Supplementary Table S6. mtDNA levels in the blood, brain, and liver of *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice stratified by sex and age.

Genotype ( <i>n</i> )	Sex ( <i>n</i> )	Age*	Rel. blood mtDNA/B2M DNA			Rel. brain mtDNA/B2M DNA			Rel. liver mtDNA/B2M DNA		
			Mean ± SD ( <i>n</i> )	FC; <i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>	Mean ± SD ( <i>n</i> )	FC; <i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>	Mean ± SD ( <i>n</i> )	FC; <i>P</i> <sub>genotype</sub>	<i>P</i> <sub>sex</sub>
<i>Cbs</i> <sup>-/-</sup> (21–39)	♀ (11–19)	Young	1.39 ± 0.78 (8)	0.9; 0.861	0.322	0.73 ± 0.08 (4)	0.9; 0.269	<b>0.045</b>	0.44 ± 0.29 (4)	0.7; 0.321	0.250
		Old	0.68 ± 0.13 (11)	0.8; 0.139	0.613	0.98 ± 0.23 (7)	1.0; 0.916	<b>0.002</b>	1.12 ± 0.49 (7)	0.6; 0.059	0.525
		<i>P</i> <sub>age</sub>	<b>0.004</b>			0.062			<b>0.036</b>		
	♂ (10–20)	Young	1.51 ± 1.09 (6)	0.7; 0.118		0.95 ± 0.14 (3)	1.1; 0.514		0.75 ± 0.32 (3)	1.1; 0.897	
		Old	0.76 ± 0.40 (14)	0.9; 0.641		1.38 ± 0.14 (7)	0.94; 0.922		1.31 ± 0.58 (7)	0.8; 0.426	
		<i>P</i> <sub>age</sub>	<b>0.046</b>			<b>0.002</b>			0.160		
<i>Cbs</i> <sup>+/-</sup> (22–38)	♀ (9–20)	Young	1.46 ± 0.72 (14)		<b>0.041</b>	0.84 ± 0.17 (4)		0.561	0.68 ± 0.25 (4)		0.858
		Old	0.84 ± 0.75 (6)		0.927	1.00 ± 0.38 (5)		0.264	1.75 ± 0.53 (5)		0.620
		<i>P</i> <sub>age</sub>	0.103			0.463			<b>0.004</b>		
	♂ (13–18)	Young	2.15 ± 0.96 (10)			0.89 ± 0.13 (7)			0.68 ± 0.22 (7)		
		Old	0.87 ± 0.62 (8)			1.40 ± 0.60 (7)			1.58 ± 0.69 (6)		
		<i>P</i> <sub>age</sub>	<b>0.003</b>			<b>0.048</b>			<b>0.004</b>		

\* Young = 44 to 77-day-old; Old = 108–390-day-old.

\* Young = 63 to 66-day-old; Old = 354 to 408-day-old.

**Supplementary Table S7. Determinants of *Tert* mRNA in the brain and liver of *Cbs*<sup>-/-</sup> and *Cbs*<sup>+/-</sup> mice: Multiple regression analysis.**

Variable	<i>Tert</i> mRNA			
	Brain		Liver	
	$\beta$	<i>p</i>	$\beta$	<i>p</i>
Female ( <i>n</i> = 19)				
Age*	-0.18	0.438	<b>0.81</b>	<b>0.000</b>
<i>Cbs</i> genotype	0.23	0.343	0.08	0.531
TL	0.08	0.765	<b>0.34</b>	<b>0.015</b>
	F = 0.83, <i>p</i> = 0.451, R <sup>2</sup> = 0.09		F = 25.6, <i>p</i> = 0.000, R <sup>2</sup> = 0.76	
Male ( <i>n</i> = 22)				
Age*	-0.31	0.207	<b>0.84</b>	<b>0.000</b>
<i>Cbs</i> genotype	0.41	0.100	0.02	0.908
TL	0.15	0.716	0.13	0.380
	F = 1.9, <i>p</i> = 0.187, R <sup>2</sup> = 0.21		F = 18.1, <i>p</i> = 0.003, R <sup>2</sup> = 0.61	
All ( <i>n</i> = 41)				
Sex	<b>-0.43</b>	<b>0.006</b>	-0.13	0.194
Age*	-0.22	0.152	<b>0.72</b>	<b>0.000</b>
<i>Cbs</i> genotype	0.24	0.115	0.07	0.484
TL	0.05	0.796	<b>0.37</b>	<b>0.000</b>
	F = 4.3, <i>p</i> = 0.011, R <sup>2</sup> = 0.28		F = 28.9, <i>p</i> = 0.000, R <sup>2</sup> = 0.69	
*Mouse age varied from 63 to 408 days.				

**Supplementary Table S8. List of primers used for mouse mRNA quantification by RT-qPCR.**

Gene name	Forward 5'→3'	Reverse 5'→3'
<i>Tert</i>	GTGAACAGCCTCCAGACAG	TTCCTAACACGCTGGTCAAA
<i>Pai-1</i>	ACGCCTGGTGCTGGTGAATGC	ACGGTGCTGCCATCAGACTTGTG
<i>Mcp1</i>	GGGATCATCTTGCTGGTGAA	AGGTCCTGTCATGCTTCTG
<i>p21</i>	GGCAGACCAGCCTGACAGAT	TTCAGGGTTTTCTTTCAGAGAAG
<i>Il-6</i>	TGGTACTCCAGAAGACCAGAGG	AACGATGATGCACTTGCAGA
<i>Kl</i>	TGTATGTGACAGCCAATGGAATCG	GAATACGCAAAGTAGCCACAAAGG
<i>p16</i>	GAACTCTTTCGGTCGTACCC	CGAATCTGCACCGTAGTTGA
<i>Il-1<math>\beta</math></i>	CAACCAACAAGTGATATTCTCCATG	GATCCACACTCTCCAGCTGCA
$\beta$ -actin	TGTTACCAACTGGGACGACA	GGGGTGTGAAGGTCTCAAA
<i>Gapdh</i>	CGTCCCGTAGACAAAATGGT	TTGATGGCAACAATCTCCAC