

## SUPPLEMENTARY MATERIAL

**Supplementary Table 1. Correlations between the different *in vitro* markers within the short-term experiment (1B).**

	p16INK4a	TAF	ROS	SAβ-gal
<b>Non-stressed</b>				
p16INK4a	n/a	-	-	-
TAF	<b>0.253 (0.011)</b>	n/a	-	-
ROS	<b>0.321 (0.001)</b>	-0.048 (0.636)	n/a	-
SAβ-gal	0.151 (0.134)	0.054 (0.598)	<b>0.232 (0.021)</b>	n/a
<b>Stressed</b>				
p16INK4a	n/a	-	-	-
TAF	<b>0.227 (0.024)</b>	n/a	-	-
ROS	<b>0.299 (0.003)</b>	0.038 (0.706)	n/a	-
SAβ-gal	0.162 (0.109)	<b>0.215 (0.032)</b>	0.115 (0.255)	n/a

N=100. Values are depicted as Pearson's correlation coefficient (P-value), partial correlation with adjustment for batch. n/a: not applicable. P16INK4a: % of p16 positive cells; Telomere-associated foci (TAF): % of nuclei with  $\geq 1$  53BP1 foci coinciding with telomeric DNA per nucleus; ROS: mean fluorescence intensity peak; SAβ-gal: median fluorescence intensity peak. All *in vitro* variables are the mean of duplicate experiments.

**Supplementary Table 2. Correlations between the different *in vitro* markers within the long-term experiment (2B).**

	ROS	SAβ-gal	Telomere shortening	Telomere length
<b>Non-stressed</b>				
ROS	n/a	-	-	-
SAβ-gal	<b>0.341 (0.042)</b>	n/a	-	-
Telomere shortening	<b>-0.466 (0.004)</b>	-0.011 (0.949)	n/a	n/a
Telomere length	-0.100 (0.550)	0.280 (0.093)	n/a	n/a
<b>Stressed</b>				
ROS	n/a	-	-	-
SAβ-gal	<b>0.436 (0.008)</b>	n/a	-	-
Telomere shortening	-0.220 (0.198)	-0.030 (0.862)	n/a	n/a
Telomere length	0.156 (0.358)	0.276 (0.104)	n/a	n/a

N=40. Values are depicted as Pearson's correlation coefficient (P-value), partial correlation with adjustment for batch. n/a: not applicable. ROS: mean fluorescence intensity peak; SAβ-gal: median fluorescence intensity peak; Telomere shortening: percentage of shortening per population doubling; Telomere length: the percentage compared to the reference cell line