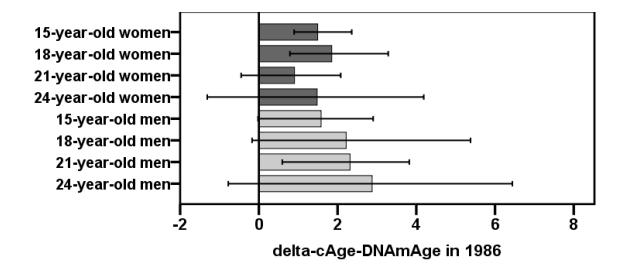
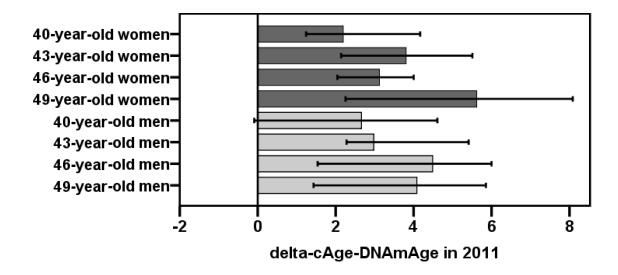
ADDITIONAL MATERIAL (2)

Supplementary Table 1. The summary of DNAmAge values among samples collected in YFS in 1986 and 2011. The values were categorized by cAge and gender. The gender-wise differences in DNAmAge values within the cAge groups did not meet statistical significance (Mann Whitney U-test, P>0.05).

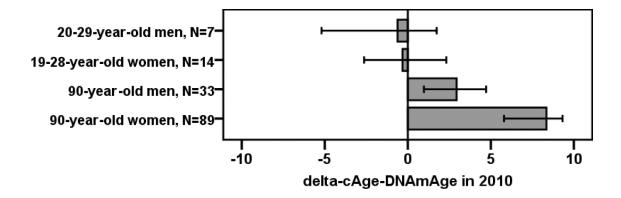
	YFS in 1986				YFS in 2011			
cAge	15	18	21	24	40	43	46	49
	49	44	55	35	49	44	55	35
N (N, women)	(29)	(30)	(31)	(21)	(29)	(30)	(31)	(21)
	13.42	15.78	18.68	21.12	37.34	40.02	41.51	44.92
median (MAD) of DNAmAge values, men	(2.35)	(1.70)	(2.57)	(5.14)	(3.82)	(1.21)	(3.62)	(2.71)
_	13.50	16.15	20.09	22.52	37.81	39.19	42.88	43.38
median (MAD) of DNAmAge values, women	(1.21)	(2.84)	(2.40)	(4.28)	(2.93)	(3.44)	(2.07)	(4.83)



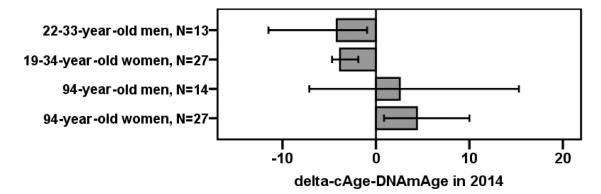
Supplementary figure 1. The differences between cAge and DNAmAge (Δ -cAge-DNAmAge) in the Young Finns study among samples collected in 1986 are shown as bars. Sample origin was whole blood leukocytes. Each bar illustrate gender-wise median of Δ -cAge-DNAmAge within a cAge group. Error bars correspond to 95% confidence intervals.



Supplementary figure 2. The differences between cAge and DNAmAge (Δ -cAge-DNAmAge) in the Young Finns study among samples collected in 2011 are shown as bars. Sample origin was whole blood leukocytes. Each bar illustrate gender-wise median of Δ -cAge-DNAmAge within a cAge group. Error bars correspond to 95% confidence intervals.



Supplementary figure 3. The differences between cAge and DNAmAge (Δ -cAge-DNAmAge) in the Vitality 90+ study among samples collected in 2010 are shown as bars. Sample origin was peripheral blood mononuclear cells. Each bar illustrate gender-wise median of Δ -cAge-DNAmAge within a cAge group. Error bars correspond to 95% confidence intervals.



Supplementary figure 4. The differences between cAge and DNAmAge (Δ -cAge-DNAmAge) in the Vitality 90+ study among samples collected in 2014 are shown as bars. Each bar illustrate gender-wise median of Δ -cAge-DNAmAge within a cAge group. Sample origin was peripheral blood mononuclear cells. Error bars correspond to 95% confidence intervals.