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## Cellular and molecular immune markers of aging and frailty

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# Appendices



## APPENDIX A

# Supplement to Chapter 2

### A.1 Supplementary methods

### Construction and validation of the frailty index

We constructed a frailty index to quantify health status based on previous studies (Collerton et al., 2012; K. Rockwood et al., 2011; Schoufour et al., 2017; Searle et al., 2008). Multiple health deficits were evaluated to determine their applicability in a frailty index. Deficits that could be evaluated on the basis of the latest available data from the Doetinchem Cohort Study (DCS) (round 5 (2008-2012) or the first four years of the ongoing round 6, (2013-2016)) were included if they had been used previously in an existing frailty index, if the prevalence of the deficit was at least one percent in the DCS, and if there is a known association with cognitive, physical, or psychological functioning. The final selection contained 36 deficits, including several parameters of physical functioning (e.g. handgrip strength), psychological functioning (e.g. the question ‘feeling depressed last week’), cognitive functioning (e.g. cognitive memory based on the Verbal Learning Task), and the presence of a number of chronic conditions (e.g. renal dysfunction). For a full description of all deficits, see Table A.1. Health deficits were either dichotomized or trichotomized, with 0= total absence of the deficit, 0.5= partial presence of the deficit, and 1= full presence of the deficit. For instance, the deficit ‘limitations in walking 100 meters’ was obtained from a self-reported questionnaire and was divided in no limitation (0), some limitation (0.5), and severe limitation (1). The frailty index is calculated by dividing the number deficits in an individual by the total number of deficits investigated (n=36).

We analyzed the frailty index distribution per sex (expected to be right-skewed and higher in women (Mitnitski et al., 2001; Searle et al., 2008)) and associations between frailty and all-cause mortality to further validate it. From all Doetinchem cohort participants who gave written consent, we obtained participants’ vital status information until Jan 1, 2018 from municipal personal records. Within each of the chosen 2.5-year age intervals, the frailty index of the deceased was compared with that of the survivors. To investigate whether frailty was

associated with all-cause mortality independently of age, the asymptotic Wilcoxon-Mann-Whitney test was used to estimate the overall association, combining the data from all age categories.

Within the Doetinchem Cohort Study (DCS), women had a higher frailty score than men ( $p < 0.001$ ) and for both men and women the distribution of the frailty index was right-skewed (Figure A.1a), which is consistent with previously constructed frailty indexes (Collerton et al., 2012; Mitnitski et al., 2001; Schoufour et al., 2017). Furthermore, frailty was strongly positively associated with age (Figure A.1b and A.1c) but inversely associated with educational level and smoking (all  $p < 0.001$ ). In the interval from January 2012 to January 2018, 223 out of 4311 participants died. The frailty index of the deceased individuals was consistently higher than the frailty index of the survivors in all but the lowest ( $< 45$  years) age interval (Figure A.1c). Overall, our results showed that a higher frailty index is strongly associated with all-cause mortality, also independent of age ( $p < 0.001$ ).

## A.2 Supplementary figures and tables

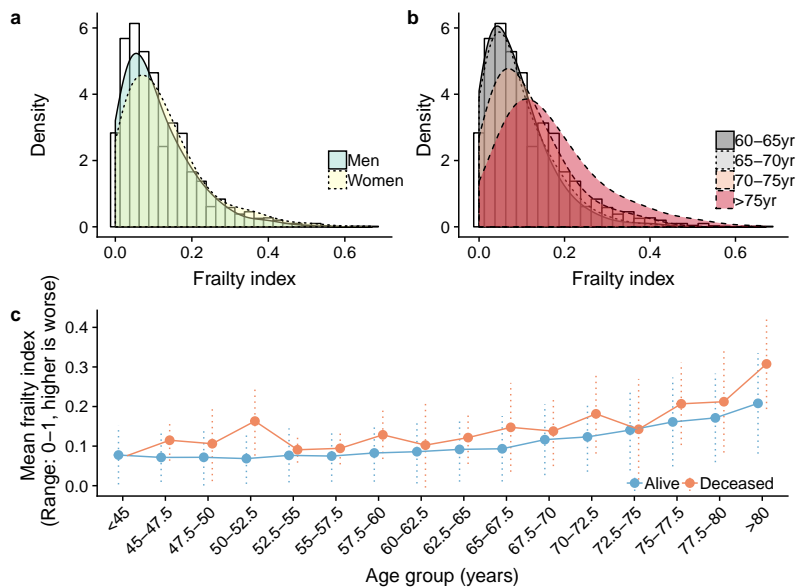


Figure A.1: Frailty index characteristics within the Doetinchem cohort. (a) Frailty index distribution per sex. (b) Frailty index density distribution per (5-year) age group. (c) Association of mortality and frailty index within the Doetinchem cohort. Colored dots show the mean frailty index grouped by (2.5-year) age category and vital status. Vital status is defined as either surviving the time period from January 2012 to January 2018 (Alive, n=4311), or dying within that time frame (Deceased, n=223).

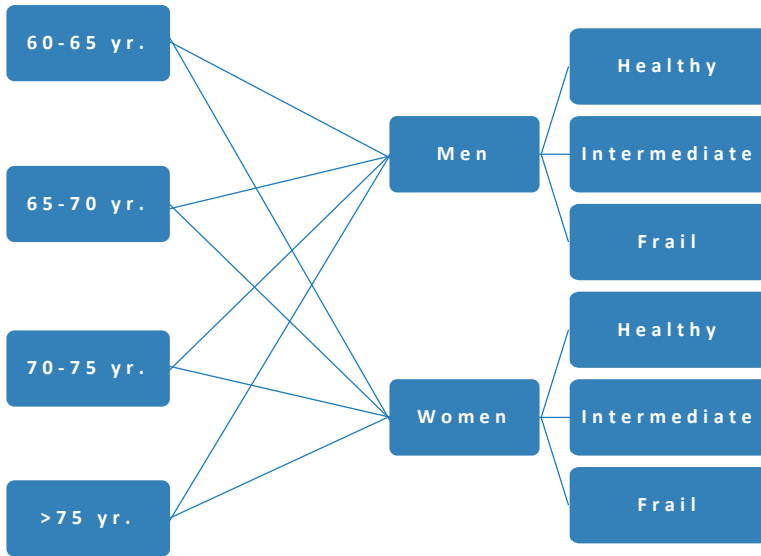


Figure A.2: Stratification scheme for sampling of individuals for DCS subcohort selection. Frail: 15% highest frailty index per age category and sex in the DCS. Healthy: 15% lowest frailty index per age category and sex in the DCS. Intermediate: participants not belonging to frail or healthy category.

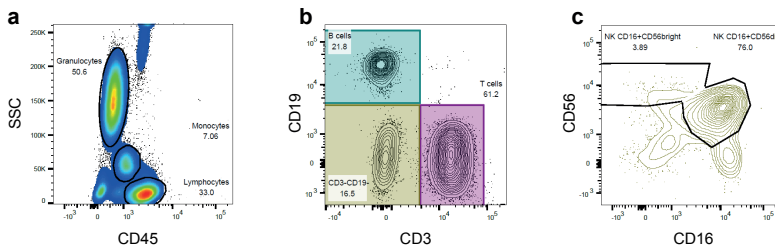


Figure A.3: Flow cytometry gating strategy to determine (a) lymphocytes, granulocytes, monocytes, (b) T cells, B cells, and (c) NK cells. SSC = Side scatter.



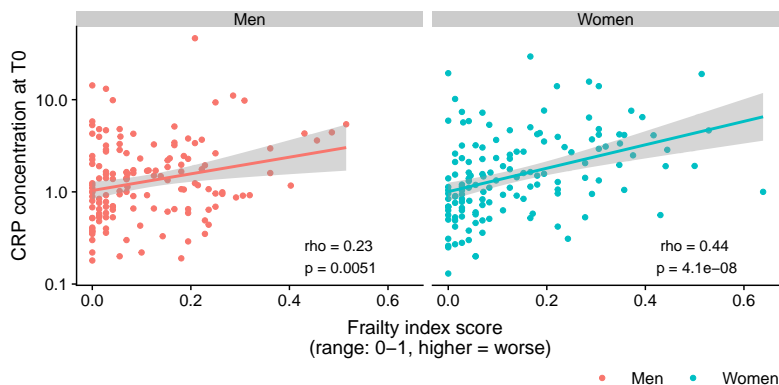


Figure A.4: The association between CRP concentrations and the frailty index score. The CRP measurements at the last time point (T0, 2008-2013) were plotted against the frailty index score for men and women separately. rho = spearman's rank correlation coefficient.

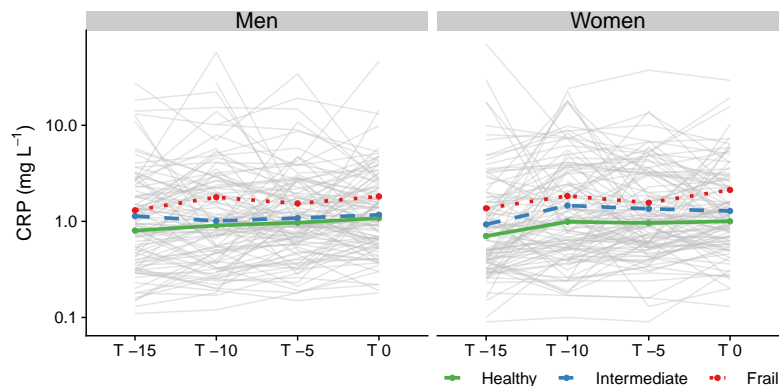


Figure A.5: CRP trajectories over a time period of approximately 15 years per frailty group, after excluding people with cardiovascular disease (CVD) and people with chronic joint inflammation. The x-axis displays the time in years according to the reference time point (T0). This T0 is the moment of the most recent blood withdrawal for CRP concentration measurement. For 58% of the participants (n=106), the frailty score was assessed at T0. For the others (n=183), it was assessed more recently (5 years after T0). The colored lines show the geometric mean values per measurement time point, stratified by sex and frailty group. Significant differences between frailty and CRP levels are presented separately in Table 2.

Table A.1: Frailty index components

No	Frailty index component	Description	Value: 0	Value: 0.5	Value: 1
1	RR	High (systolic) blood pressure	$RR < 160$		$RR \geq 160$
2	Cardiologic Disease	One or more of the following conditions: myocardial infarction, bypass-surgery, balloon dilatation, cardiac catheterization, pacemaker implantation, large blood vessel surgery, hospitalization due to cardiac failure	No		One or more prevalent
3	Diabetes	Prevalence of Diabetes	No		Yes
4	Hearing	Inability to maintain a conversation in a group of 3 or more people due to hearing impairment (with hearing aid if needed)	Yes or with some effort		No or with great effort
5	Malignancy	(History of) any form of malignancy	No		Yes
6	Joint Inflammation	Chronic joint inflammation in the past year	No		Yes
7	Osteoporosis	Osteoporosis diagnosed by a medical doctor, in the past year,	No		Yes
8	Lower Back Pain	Severe lower back complaints in the past year (including lumbar herniated nucleus pulposus)	No		Yes
9	CVA	(History of) stroke	No		Yes
10	Migraine	Migraine prevalence in the past year	No		Yes
11	Neurologic Disease	One or more of the following neurological diseases, diagnosed by a medical doctor: m. Parkinson, Multiple Sclerosis, epilepsy	No		Yes

Table A.1: Frailty index components (*continued*)

No	Frailty index component	Description	Value: 0	Value: 0.5	Value: 1
12	Asthma	Asthma diagnosed by a doctor and one or more asthma attacks in the past year	No		Yes
13	Spirometry Ratio	Poor lung function quantified by spirometry measurements: first second of forced expiration divided by the forced vital capacity (FEV1/FVC)	$FEV1/FVC > 0.70$		$FEV1/FVC \leq 0.70$
14	Digestive Tract	Severe bowel disorders in the past year, diagnosed by a medical doctor	No		Yes
15	Vertigo	Vertigo with falling the past 12 months	No vertigo	Some vertigo	Severe vertigo
16	BMI	Over- or underweight	$18.5 < BMI \leq 30$		$BMI \leq 18.5$ or $BMI > 30$
17	Pain	Limited in daily activities due to pain	No limitation	Some limitation	Severe limitation
18	Incontinence	Unintentional urine incontinence past 12 months	No		Yes
19	Ankle Brachial Index	Poor ankle brachial index (ABI, the ratio of the systolic blood pressure in the ankle to the blood pressure in the arm)	$ABI > 0.9$		$ABI \leq 0.9$
20	Health Perception	(Subjective) perception of poor health	No or some impairment		Severe impairment

Table A.1: Frailty index components (*continued*)

No	Frailty index component	Description	Value: 0	Value: 0.5	Value: 1
21	Eyesight	Bad eyesight perception, facial recognition within 4 meters (with glasses/ contact lenses if needed)	No		Yes
22	Renal Function	Poor renal function (estimated Glomular Filtration Rate (eGFR), calculated with plasma creatinine concentrations) (Inker et al., 2012)	$eGFR \geq 60$		$eGFR < 60$
23	Cognitive Speed	Poor cognitive speed. Z-scores corrected for measurements per person and for education level. Scores derived from the Stroop Color-Word Test and the Letter-Digit Substitution Test, as described previously (Nooyens et al., 2011)	Not belonging to 10% participants with lowest z-score within the Doetinchem cohort		Belonging to 10% participants with lowest z-score within the Doetinchem cohort
24	Cognitive Memory	Poor cognitive memory. Z-scores corrected for measurements per person and for education level. Scores derived from the Verbal Learning Test, as described previously (Nooyens et al., 2011)	Not belonging to 10% participants with lowest z-score within the Doetinchem cohort		Belonging to 10% participants with lowest z-score within the Doetinchem cohort

Table A.1: Frailty index components (*continued*)

No	Frailty index component	Description	Value: 0	Value: 0.5	Value: 1
25	Cognitive Flexibility	Poor cognitive flexibility. Z-scores corrected for measurements per person and education level. Scores derived from the Stroop Color-Word Test, as described previously (Nooyens et al., 2011)	Not belonging to 10% participants with lowest z-score within the Doetinchem cohort		Belonging to 10% participants with lowest z-score within the Doetinchem cohort
26	Physical Inactive	Not meeting the Dutch healthy exercise norm.(Kemper, 2000). In addition: belonging to the 25th lowest percentile of walking activity and the 10th percentile lowest low/medium/high intensive activities in the Doetinchem cohort	No		Yes
27	ADL	Limited in washing and dressing due to health	No limitation	Some limitation	Severe limitation
28	Household	Limited in daily activities (cooking, cleaning) due to health	No limitatlion	Some limitation	Severe limitation
29	Walking	Limited in in walking 100 meters	No limitatlion	Some limitation	Severe limitation

Table A.1: Frailty index components (*continued*)

No	Frailty index component	Description	Value: 0	Value: 0.5	Value: 1
30	Lifting	Limited in lifting or carrying groceries due to health	No limitation	Some limitation	Severe limitation
31	Walking Stairs	Limited in climbing stairs	No limitation	Some limitation	Severe limitation
32	Grip Strength	Poor grip strength (cutoff points as described previously) (Fried et al., 2001)			
		<i>Men, BMI</i> $\leq 24$	$> 29$		$\leq 29$
		<i>Men, 24 &lt; BMI</i> $\leq 26$	$> 30$		$\leq 30$
		<i>Men, 26 &lt; BMI</i> $\leq 28$	$> 30$		$\leq 30$
		<i>Men, BMI</i> $> 28$	$> 32$		$\leq 32$
		<i>Women, BMI</i> $\leq 24$	$> 17$		$\leq 17$
		<i>Women, 23 &lt; BMI</i> $\leq 26$	$> 17.3$		$\leq 17.3$
		<i>Women, 26 &lt; BMI</i> $\leq 29$	$> 18$		$\leq 18$
		<i>Women, BMI</i> $> 29$	$> 21$		$\leq 21$
33	Depressed	Feeling depressed the past week	No limitation	Some limitation	Severe limitation
34	Happiness	Feeling unhappy the past 4 weeks	No limitation	Some limitation	Severe limitation

Table A.1: Frailty index components (*continued*)

No	Frailty index component	Description	Value: 0	Value: 0.5	Value: 1
35	Mental Effort	Feeling as if every activity costs effort during the past week	No limitation	Some limitation	Severe limitation
36	Getting Going	Not being able to get going the past week	No limitation	Some limitation	Severe limitation

## A.2. Supplementary figures and tables

Table A.2: Linear mixed effects regression model of CRP concentration ( $\text{mg L}^{-1}$ ) for both sexes, after excluding people with CVD or joint inflammation

Predictors	log(CRP) for men			log(CRP) for women		
	Estimates	CI	p	Estimates	CI	p
(Intercept)	-0.27	-1.51 – 0.97	0.675	-0.63	-1.87 – 0.61	0.324
Frailty index	2.14	0.31 – 3.98	0.024	2.28	0.71 – 3.85	0.005
Age at baseline	0.00	-0.02 – 0.03	0.851	0.01	-0.02 – 0.03	0.531
Years after baseline age	0.01	-0.00 – 0.03	0.106	0.02	0.01 – 0.03	0.009
Observations	413			420		
Marginal R2 / Conditional R2	0.035/0.644			0.062/0.570		

*Note:*

CRP concentrations were log transformed before analysis. Four CRP concentration measurements per person were used, with approximately 5 years between measurements. Frailty category: the 'healthy' category was taken as the reference. Baseline age: The age at which the first CRP measurement took place. Time: time difference in years with the first baseline CRP measurement. CI= 95% confidence interval. P-values <0.05 are displayed in bold and were considered statistically significant.

Table A.3: CRP concentrations ( $\text{mg L}^{-1}$ ) dependent on age and leukocyte numbers

Predictors	Men (n=145)			Women (n=144)		
	Estimates	CI	p	Estimates	CI	p
(Intercept)	-0.63	-1.61 – 0.35	0.21	-0.58	-1.79 – 0.64	0.354
Granulocytes	0.28	0.14 – 0.41	<0.001	0.40	0.21 – 0.59	<0.001
Monocytes	0.00	-0.12 – 0.12	0.954	-0.13	-0.40 – 0.14	0.33
T cells	0.10	-0.06 – 0.27	0.229	0.01	-0.16 – 0.18	0.888
B cells	-0.09	-0.29 – 0.12	0.415	0.05	-0.09 – 0.19	0.484
NK cells	0.03	-0.11 – 0.17	0.659	-0.06	-0.21 – 0.09	0.405
Age at baseline	0.01	-0.00 – 0.03	0.141	0.01	-0.01 – 0.04	0.219
Time (years after baseline age)	0.01	-0.00 – 0.02	0.09	0.02	0.01 – 0.03	0.006
Observations	533			546		
Marginal R2 / Conditional R2	0.096 / 0.607			0.105 / 0.603		

*Note:*

CRP concentrations were log transformed before analysis. Four CRP concentration measurements per person were used, with approximately 5 years between measurements. Cell numbers were transformed to z-scores for easier comparison. Age at baseline: age at first CRP measurement. P-values <0.05 are displayed in bold and were considered statistically significant.





## APPENDIX B

# Supplement to Chapter 3

### B.1 Supplementary figures and tables

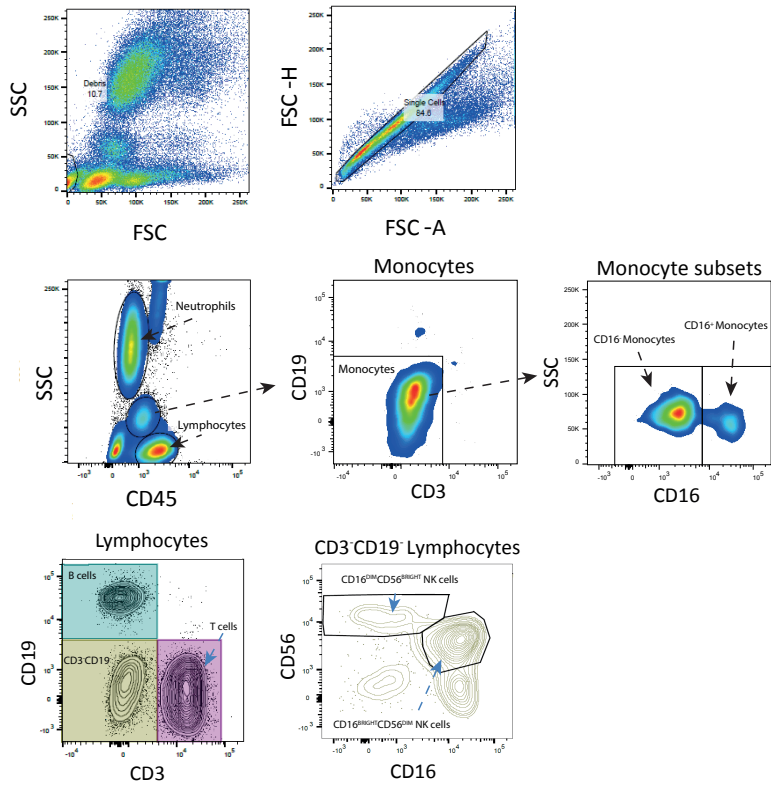


Figure B.1: General gating strategy showing gating of single cells by FSC-A versus FSC-H, leukocyte subsets by SSC and CD45, monocyte subpopulations by CD16, T and B-cells by CD3 and CD19, and NK cells by CD16 and CD56.

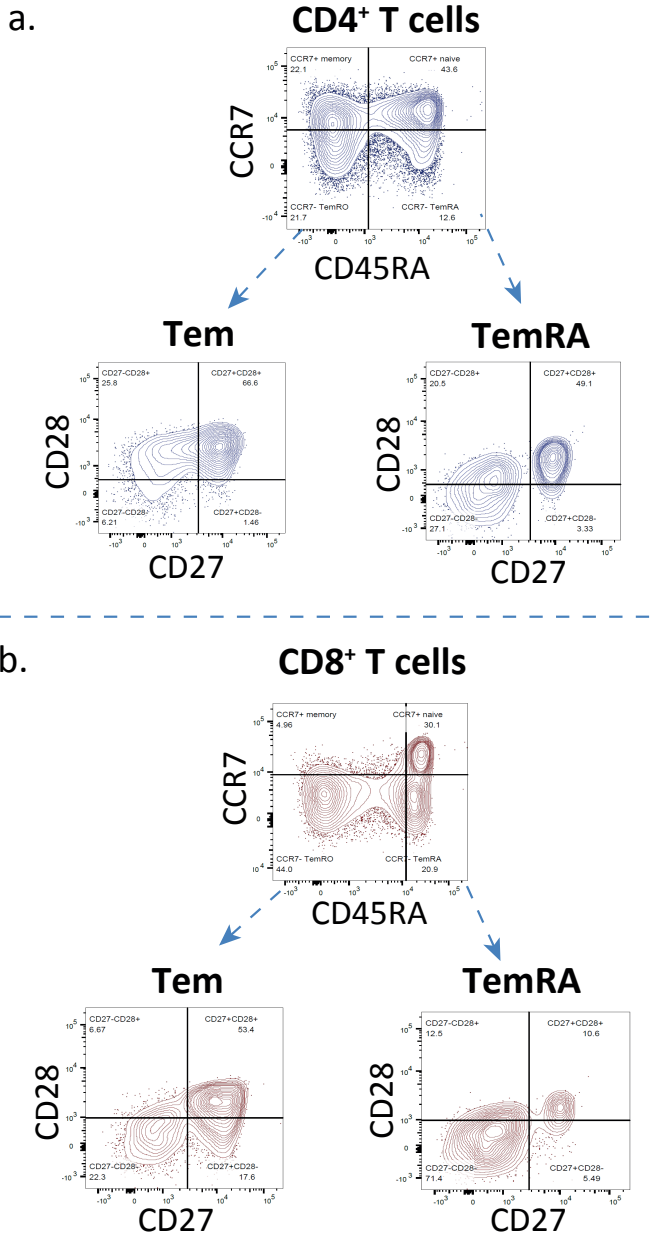


Figure B.2: Gating strategy for (a) memory CD4<sup>+</sup> T cells and (b) memory CD8<sup>+</sup> T cells. Tem: effector memory (CD4<sup>+</sup> or CD8<sup>+</sup>) T cells. TemRA: effector memory T cells re-expressing CD45RA.

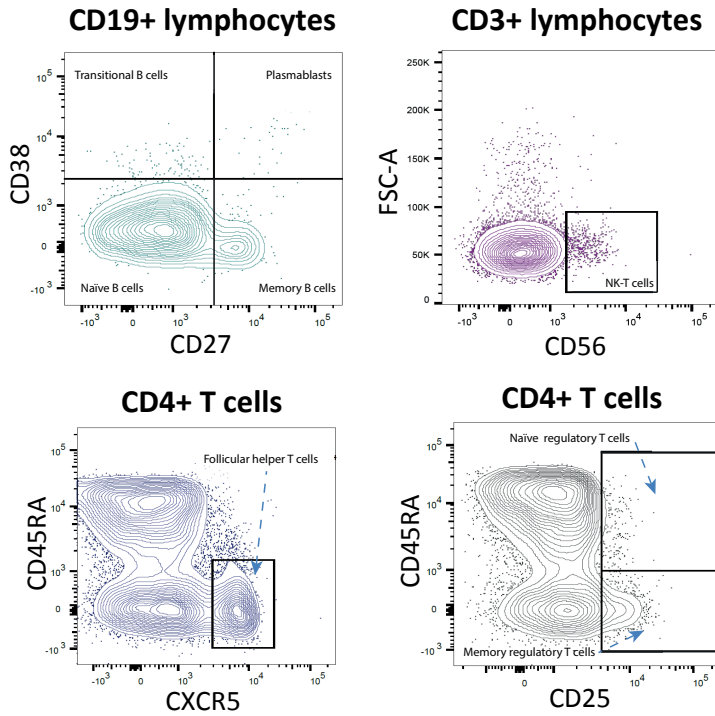


Figure B.3: Gating strategy for (a) B cell subsets and (b) follicular helper T cells and regulatory T cells.

## B.1. Supplementary figures and tables

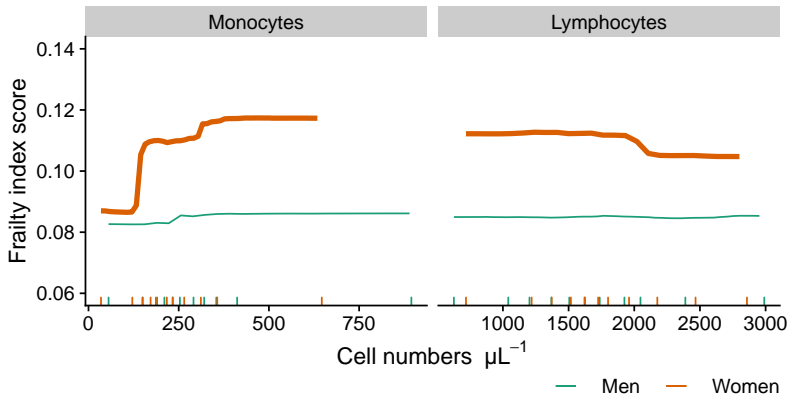


Figure B.4: Partial dependence plots showing how frailty ‘depends’ on total monocyte numbers and total lymphocyte numbers in men ( $n=140$ ) and women ( $n=137$ ). Participants with missing frailty index score data ( $n=12$ ) were excluded from analysis. The short vertical segments on the horizontal axis represent the deciles of the cell numbers in the data. Range of the figures is restricted to the part containing most of the data.

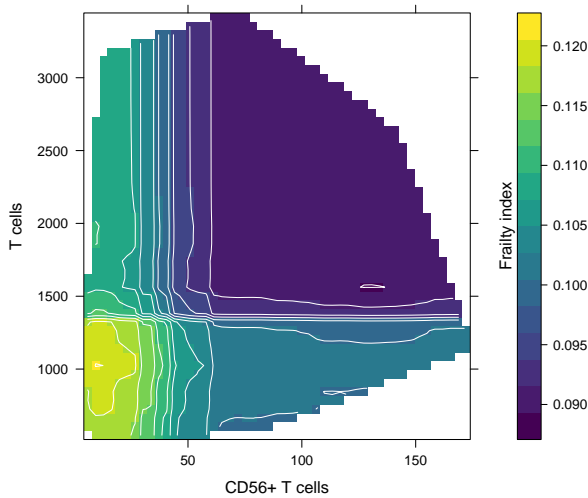


Figure B.5: Partial dependence plot with the possible joint relationship of total T cell numbers and CD56<sup>+</sup> T cell numbers on frailty. To avoid extrapolation, the plot is restricted to the area of which data is available.

Table B.1: Definition of cell phenotypes by expression of cell surface markers

Cell phenotype	Definition
<b>T cells</b>	
T cells	CD45 <sup>+</sup> CD3 <sup>+</sup>
T Follicular helper	CD3 <sup>+</sup> CD4 <sup>+</sup> CD45RA <sup>-</sup> CXCR5 <sup>+</sup>
CD56 <sup>+</sup> T cells	CD45 <sup>+</sup> CD3 <sup>+</sup> CD56 <sup>+</sup>
CD4/CD8 ratio	CD4 T cells/CD8 T cells
CD4 T cells	CD45 <sup>+</sup> CD3 <sup>+</sup> CD4 <sup>+</sup>
CD4 Naive	CD3 <sup>+</sup> CD4 <sup>+</sup> CD45RA <sup>+</sup> CCR7 <sup>+</sup>
CD4 CM	CD3 <sup>+</sup> CD4 <sup>+</sup> CD45RA <sup>-</sup> CCR7 <sup>+</sup>
CD4 TemRA	CD3 <sup>+</sup> CD4 <sup>+</sup> CD45RA <sup>+</sup> CCR7 <sup>-</sup>
CD4 TemRA Early	CD3 <sup>+</sup> CD4 <sup>+</sup> CD45RA <sup>+</sup> CCR7 <sup>-</sup> CD27 <sup>+</sup> CD28 <sup>+</sup>
CD4 TemRA Late	CD3 <sup>+</sup> CD4 <sup>+</sup> CD45RA <sup>+</sup> CCR7 <sup>-</sup> CD27 <sup>-</sup> CD28 <sup>-</sup>
CD4 Tem	CD3 <sup>+</sup> CD4 <sup>+</sup> CD45RA <sup>-</sup> CCR7 <sup>-</sup>
CD4 Tem Early	CD3 <sup>+</sup> CD4 <sup>+</sup> CD45RA <sup>-</sup> CCR7 <sup>-</sup> CD27 <sup>+</sup> CD28 <sup>+</sup>
CD4 Tem Late	CD3 <sup>+</sup> CD4 <sup>+</sup> CD45RA <sup>-</sup> CCR7 <sup>-</sup> CD27 <sup>-</sup> CD28 <sup>-</sup>
Regulatory T cells	CD3 <sup>+</sup> CD4 <sup>+</sup> CD25 <sup>BRIGHT</sup>
Treg naive	CD3 <sup>+</sup> CD4 <sup>+</sup> CD25 <sup>BRIGHT</sup> CD45RA <sup>+</sup>
Treg memory	CD3 <sup>+</sup> CD4 <sup>+</sup> CD25 <sup>BRIGHT</sup> CD45RA <sup>-</sup>
CD8 T cells	CD45 <sup>+</sup> CD3 <sup>+</sup> CD8 <sup>+</sup>
CD8 Naive	CD3 <sup>+</sup> CD8 <sup>+</sup> CD45RA <sup>+</sup> CCR7 <sup>+</sup>
CD8 CM	CD3 <sup>+</sup> CD8 <sup>+</sup> CD45RA <sup>-</sup> CCR7 <sup>+</sup>
CD8 TemRA	CD3 <sup>+</sup> CD8 <sup>+</sup> CD45RA <sup>+</sup> CCR7 <sup>-</sup>
CD8 TemRA Early	CD3 <sup>+</sup> CD8 <sup>+</sup> CD45RA <sup>+</sup> CCR7 <sup>-</sup> CD27 <sup>+</sup> CD28 <sup>+</sup>
CD8 TemRA Late	CD3 <sup>+</sup> CD8 <sup>+</sup> CD45RA <sup>+</sup> CCR7 <sup>-</sup> CD27 <sup>-</sup> CD28 <sup>-</sup>
CD8 Tem	CD3 <sup>+</sup> CD8 <sup>+</sup> CD45RA <sup>-</sup> CCR7 <sup>-</sup>
CD8 Tem Early	CD3 <sup>+</sup> CD8 <sup>+</sup> CD45RA <sup>-</sup> CCR7 <sup>-</sup> CD27 <sup>+</sup> CD28 <sup>+</sup>
CD8 Tem Late	CD3 <sup>+</sup> CD8 <sup>+</sup> CD45RA <sup>-</sup> CCR7 <sup>-</sup> CD27 <sup>-</sup> CD28 <sup>-</sup>
<b>Neutrophils</b>	
Neutrophils	SSC <sup>BRIGHT</sup> CD45 <sup>DIM</sup>
Neutrophils, CD16 expr.	SSC <sup>BRIGHT</sup> CD45 <sup>DIM</sup> ;CD16 <sup>expression</sup>
<b>Monocytes</b>	
Monocytes	SSC <sup>DIM</sup> CD45 <sup>+</sup> CD3 <sup>-</sup> CD19 <sup>-</sup>
CD16 <sup>-</sup> monocytes	SSC <sup>DIM</sup> CD45 <sup>+</sup> CD16 <sup>-</sup>
CD16 <sup>-</sup> mon. CD38 expr.	SSC <sup>DIM</sup> CD45 <sup>+</sup> CD16 <sup>-</sup> ;CD38 <sup>expression</sup>
CD16 <sup>-</sup> mon. HLADR expr.	SSC <sup>DIM</sup> CD45 <sup>+</sup> CD16 <sup>-</sup> ;HLADR <sup>expression</sup>
CD16 <sup>+</sup> monocytes	SSC <sup>DIM</sup> CD45 <sup>+</sup> CD16 <sup>+</sup>
CD16 <sup>+</sup> mon. CD38 expr.	SSC <sup>DIM</sup> CD45 <sup>+</sup> CD16 <sup>+</sup> ;CD38 <sup>expression</sup>
CD16 <sup>+</sup> mon. HLADR expr.	SSC <sup>DIM</sup> CD45 <sup>+</sup> CD16 <sup>+</sup> ;HLADR <sup>expression</sup>
<b>B cells</b>	
B cells	CD45 <sup>+</sup> CD19 <sup>+</sup>
Plasmablasts	CD45 <sup>+</sup> CD19 <sup>+</sup> CD38 <sup>BRIGHT</sup> CD27 <sup>BRIGHT</sup>
Transitional B cells	CD45 <sup>+</sup> CD19 <sup>+</sup> CD38 <sup>BRIGHT</sup> CD27 <sup>-</sup>

Table B.1: Definition of cell phenotypes by expression of cell surface markers (*continued*)

Cell phenotype	Definition
Naive B cells	$CD45^+ CD19^+ CD38^{DIM} CD27^-$
Memory B cells	$CD45^+ CD19^+ CD38^{DIM} CD27^+$
<b>NK cells</b>	
NK cells	Either NK $CD56^{BRIGHT}$ or NK $CD56^{DIM}$
NK $CD56^{BRIGHT}$	$CD45^+ CD3^- CD19^- CD16^{DIM} CD56^{BRIGHT}$
NK $CD56^{DIM}$	$CD45^+ CD3^- CD19^- CD16^{BRIGHT} CD56^{DIM}$



Table B.2: Leukocyte numbers per sex and CMV serostatus

	Men			Women		
	CMV- (n=69)	CMV+ (n=76)	P value <sup>1</sup>	CMV- (n=53)	CMV+ (n=91)	P value <sup>1</sup>
<b>T cells</b>						
T cells	1002.6 (428.9)	1202.5 (463.4)	<0.001*	1144.7 (436.2)	1272.4 (474.9)	0.026*
T Follicular helper	49.5 (28.8)	61.8 (34.1)	0.04*	67.5 (37.6)	66.7 (40.9)	0.374
CD56 <sup>+</sup> T cells	21.8 (27.8)	60.9 (108.5)	<0.001*	16 (16.7)	48.5 (61.9)	<0.001*
CD4/CD8 ratio	3.6 (2.7)	2.5 (1.8)	0.002*	4.2 (3.2)	2.8 (1.7)	<0.001*
CD4 T cells	669.7 (370.9)	766.5 (391.6)	0.041*	897.3 (371.9)	892.2 (363.3)	0.847
CD4 Naive	316.3 (283.4)	243.1 (215.2)	0.292	343.7 (262)	374.6 (295.9)	0.924
CD4 CM	175.6 (90.3)	217.1 (105)	0.01*	234.1 (118)	230.9 (149.4)	0.179
CD4 TemRA	27.8 (28.8)	46.7 (45.7)	<0.001*	33.9 (50.7)	52.6 (52.4)	0.003*
CD4 TemRA Early	22.4 (26.5)	19.7 (21.2)	0.666	29.7 (44.1)	28.2 (32.2)	0.839
CD4 TemRA Late	0.1 (0.2)	5.4 (24.2)	<0.001*	0.1 (0.2)	5.9 (13.9)	<0.001*
CD4 Tem	110.6 (70.1)	153.8 (108.3)	<0.001*	155.8 (91.6)	179.9 (112.8)	0.177
CD4 Tem Early	77.7 (51.5)	86.9 (54.3)	0.166	114.7 (59.8)	106 (73.2)	0.362
CD4 Tem Late	1.1 (1.5)	15 (25.5)	<0.001*	1.1 (1.6)	8.9 (14.2)	<0.001*
Regulatory T cells	34 (18.8)	34.7 (23.8)	0.556	42.6 (22.8)	42.1 (26.5)	0.377
Treg naive	6.7 (4.1)	7.2 (3.2)	0.607	8.3 (6.4)	9.4 (5.7)	0.479
Treg memory	9.3 (5.8)	10.2 (5.2)	0.343	10.5 (5.6)	9.1 (6.4)	0.049
CD8 T cells	188.9 (198.7)	337.2 (246.1)	<0.001*	203.5 (124.1)	333.2 (159.1)	<0.001*
CD8 Naive	20 (31.5)	18 (26.4)	0.532	35.4 (41.9)	34.1 (33.8)	0.948
CD8 CM	13.7 (11.7)	17.3 (15.6)	0.01*	17.3 (14)	16.8 (15.3)	0.76
CD8 TemRA	32.3 (36.9)	83.6 (109.9)	<0.001*	27.8 (26.9)	75 (74)	<0.001*
CD8 TemRA Early	6.3 (7.8)	7.4 (8.4)	0.73	6.7 (7.5)	9.1 (9.3)	0.129
CD8 TemRA Late	10.5 (16.1)	54.1 (66.2)	<0.001*	8.6 (13)	40.4 (49.1)	<0.001*
CD8 Tem	120.1 (116.9)	227.9 (135.7)	<0.001*	117.4 (62.9)	212.4 (111.9)	<0.001*
CD8 Tem Early	64.3 (63.6)	73 (68.1)	0.409	74.7 (53.3)	88.6 (59.3)	0.538

CD8 Tem Late	11.4 (14.5)	59 (72.8)	<0.001*	6.6 (7.8)	48.9 (55.5)	<0.001*
<b>Neutrophils</b>						
Neutrophils	2811.5 (2204.9)	2646.3 (1309.4)	0.197	2707.9 (1478.6)	2851.3 (1573.1)	0.733
Neutrophils, CD16 expr. <sup>2</sup>	3671 (5564)	3971 (5900.2)	0.393	3950 (5273)	4343 (5299)	0.86
<b>Monocytes</b>						
Monocytes	276.5 (136.6)	246.5 (126.3)	0.057*	233.4 (144.2)	218.4 (125.2)	0.365
CD16 <sup>-</sup> monocytes	273.2 (134.9)	246 (122.1)	0.064*	229.3 (143)	211.8 (125.8)	0.351
CD16 <sup>-</sup> mon. CD38 expr. <sup>2</sup>	1200.5 (338.2)	1273 (341.5)	0.039*	1168 (193)	1216 (285)	0.401
CD16 <sup>-</sup> mon. HLADR expr. <sup>2</sup>	9106 (6090.2)	8736 (4603)	0.78	8834 (3806)	8171 (6208)	0.907
CD16 <sup>+</sup> monocytes	4 (4.7)	4 (3.9)	0.397	4 (3.8)	3.4 (4.2)	0.628
CD16 <sup>+</sup> mon. CD38 expr. <sup>2</sup>	445.5 (192.5)	499 (199.5)	0.157	412 (139)	444 (218)	0.138
CD16 <sup>+</sup> mon. HLADR expr. <sup>2</sup>	56851 (27956)	52916 (41323.5)	0.587	42036 (41637)	39669 (39328)	0.935
<b>B cells</b>						
B cells	124 (86.8)	150.4 (104.2)	0.427	167.7 (115.2)	173.9 (118.8)	0.904
Plasmablasts	1.5 (1.5)	1.4 (1.4)	0.677	1.5 (1.5)	1.4 (1.2)	0.644
Transitional B cells	3.7 (3.7)	3.8 (5.6)	0.643	4.6 (4.8)	4.2 (4.4)	0.806
Naive B cells	92.1 (72.8)	111 (75.6)	0.487	124.8 (88.9)	120.4 (78.1)	0.826
Memory B cells	24 (24.4)	28.4 (30)	0.73	32.8 (33.4)	37.1 (32.2)	0.667
<b>NK cells</b>						
NK cells	274.9 (217)	217.4 (191.2)	0.109	236.1 (117)	212.3 (138.8)	0.39
NK CD56 <sup>BRIGHT</sup>	10.7 (7.2)	10.4 (7.7)	0.872	11.6 (8.9)	9.9 (6.6)	0.199
NK CD56 <sup>DIM</sup>	251 (219.4)	203.1 (183.8)	0.116	220.8 (109.8)	204.6 (139.5)	0.46

*Note:*

All values are median (interquartile range) cell numbers per  $\mu\text{L}$ , unless otherwise stated

<sup>1</sup> P values of Kruskal-Wallis rank sum test between CMV serostatus per sex, adjusted for age

<sup>2</sup> Values in Median fluorescence intensity

\* Selected outcomes when the False Discovery Rate is set to a maximum of 15%

Table B.3: Spearman associations between immune cell sub-populations and frailty in men (n=140)

Immune cell subset	$\rho$	P value	FDR*
<b>Neutrophils</b>	<b>0.25</b>	<b>0.002</b>	<b>0.077</b>
Transitional B cells	-0.19	0.015	0.324
Monocytes	0.17	0.027	0.389
CD16 <sup>-</sup> monocytes	0.16	0.036	0.383
CD4 Naive	0.17	0.040	0.341
CD4 TemRA Late	0.13	0.078	0.562
CD4 TemRA	0.17	0.090	0.553
Regulatory T cells	0.12	0.104	0.561
CD16 <sup>+</sup> monocytes	0.16	0.133	0.633
CD16 <sup>-</sup> mon. HLADR expr.	-0.15	0.133	0.571
NK CD56 <sup>DIM</sup>	-0.08	0.172	0.672
CD4 TemRA Early	0.14	0.191	0.686
CD4 T cells	0.09	0.207	0.683
NK cells	-0.06	0.208	0.638
CD4/CD8 ratio	0.10	0.219	0.628
Memory B cells	-0.07	0.319	0.857
T cells	0.09	0.321	0.812
CD8 Tem Early	-0.07	0.354	0.845
Treg naive	0.08	0.387	0.877
Neutrophils, CD16 expr.	-0.10	0.399	0.858
T Follicular helper	0.07	0.439	0.898
CD16 <sup>+</sup> mon. CD38 expr.	-0.05	0.440	0.860
Lymphocytes	0.04	0.574	1.073
NK CD56 <sup>BRIGHT</sup>	-0.03	0.582	1.042
CD8 Naive	0.05	0.590	1.014
CD8 CM	-0.05	0.650	1.074
CD8 T cells	0.03	0.658	1.048
B cells	-0.06	0.667	1.025
CD8 Tem Late	0.09	0.689	1.022
CD56 <sup>+</sup> T cells	0.03	0.694	0.995
CD8 Tem	0.02	0.697	0.967
CD16 <sup>-</sup> mon. CD38 expr.	-0.01	0.705	0.948
CD16 <sup>+</sup> mon. HLADR expr.	0.07	0.728	0.949
CD4 Tem Late	0.03	0.735	0.930

Table B.3: Spearman associations between immune cell sub-populati (*continued*)

Immune cell subset	$\rho$	P value	FDR*
CD4 CM	-0.05	0.757	0.931
CD4 Tem Early	-0.03	0.789	0.943
CD8 TemRA Late	0.02	0.796	0.925
Plasmablasts	0.04	0.810	0.916
Naive B cells	-0.05	0.839	0.925
CD8 TemRA	0.02	0.851	0.915
CD8 TemRA Early	-0.01	0.931	0.976
Treg memory	0.03	0.967	0.990
CD4 Tem	0.00	0.994	0.994

*Note:*

The (Spearman) associations are ordered by p value, with the lowest p values shown at the top. Associations that were selected with a FDR lower than 15% are shown in bold.

\* False Discovery Rate (estimated)

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Table B.4: Spearman associations between immune cell sub-populations and frailty in women (n=137)

Immune cell subset	$\rho$	P value	FDR*
<b>Neutrophils</b>	<b>0.40</b>	<b>&lt;0.001</b>	<b>0.000</b>
<b>CD16<sup>-</sup> monocytes</b>	<b>0.24</b>	<b>0.003</b>	<b>0.066</b>
<b>Monocytes</b>	<b>0.23</b>	<b>0.004</b>	<b>0.054</b>
<b>CD56<sup>+</sup> T cells</b>	<b>-0.20</b>	<b>0.01</b>	<b>0.109</b>
<b>CD4 TemRA Late</b>	<b>-0.13</b>	<b>0.015</b>	<b>0.128</b>
CD16 <sup>-</sup> mon. CD38 expr.	-0.20	0.022	0.157
CD4 Tem Early	0.20	0.032	0.194
CD16 <sup>+</sup> mon. CD38 expr.	-0.18	0.037	0.200
CD16 <sup>-</sup> mon. HLADR expr.	-0.12	0.063	0.302
CD4 Tem	0.15	0.073	0.313
NK CD56 <sup>BRIGHT</sup>	-0.15	0.08	0.311
CD8 TemRA	-0.09	0.091	0.325
Transitional B cells	-0.16	0.096	0.318
CD8 TemRA Late	-0.08	0.099	0.303
NK CD56 <sup>DIM</sup>	-0.15	0.133	0.382
NK cells	-0.13	0.149	0.401
CD8 CM	0.14	0.165	0.418
CD4 Tem Late	-0.04	0.235	0.562
CD4/CD8 ratio	0.11	0.25	0.567
CD4 TemRA Early	0.09	0.274	0.589
CD8 T cells	-0.10	0.322	0.658
CD4 CM	0.07	0.34	0.665
CD4 Naive	-0.11	0.344	0.643
T Follicular helper	0.09	0.347	0.621
CD8 Naive	-0.13	0.359	0.618
Treg memory	0.07	0.374	0.618
CD8 TemRA Early	-0.02	0.422	0.672
Memory B cells	-0.09	0.428	0.658
Plasmablasts	-0.07	0.461	0.683
T cells	-0.05	0.49	0.702
CD8 Tem Early	0.12	0.519	0.720
Lymphocytes	-0.02	0.526	0.707
CD16 <sup>+</sup> monocytes	0.07	0.551	0.718
B cells	-0.06	0.579	0.732

Table B.4: Spearman associations between immune cell subpopulations (*continued*)

Immune cell subset	$\rho$	P value	FDR*
Neutrophils, CD16 expr.	-0.01	<b>0.73</b>	<b>0.897</b>
Treg naive	-0.04	<b>0.779</b>	<b>0.930</b>
CD8 Tem	0.02	<b>0.814</b>	<b>0.946</b>
CD4 TemRA	0.00	<b>0.913</b>	<b>1.033</b>
CD16 <sup>+</sup> mon. HLADR expr.	0.01	<b>0.946</b>	<b>1.044</b>
CD8 Tem Late	0.01	<b>0.953</b>	<b>1.025</b>
Regulatory T cells	0.01	<b>0.966</b>	<b>1.013</b>
CD4 T cells	0.01	<b>0.967</b>	<b>0.990</b>
Naive B cells	0.00	<b>0.984</b>	<b>0.984</b>

*Note:*

The (Spearman) associations are ordered by p value, with the lowest p values shown at the top. Associations that were selected with a FDR lower than 15% are shown in bold.

\* False Discovery Rate (estimated)

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Table B.5: Repeated sensitivity analysis in men when monocytes were restricted to be HLADR<sup>+</sup>

Immune cell subset	$\rho$	P value	FDR*
<b>Neutrophils</b>	<b>0.25</b>	<b>0.002</b>	<b>0.061</b>
Transitional B cells	-0.19	0.014	0.276
Monocytes (HLADR <sup>+</sup> )	0.17	0.027	0.349
CD4 Naive	0.17	0.040	0.386
CD16 <sup>-</sup> monocytes (HLADR <sup>+</sup> )	0.14	0.048	0.378
CD4 TemRA Late	0.13	0.078	0.508
CD4 TemRA	0.17	0.091	0.508
Regulatory T cells	0.12	0.104	0.508
CD16 <sup>+</sup> monocytes (HLADR <sup>+</sup> )	0.17	0.108	0.470
NK CD56 <sup>DIM</sup>	-0.08	0.172	0.671
CD4 TemRA Early	0.14	0.191	0.677
CD4 T cells	0.09	0.205	0.666
NK cells	-0.06	0.209	0.626
CD4/CD8 ratio	0.10	0.222	0.617
T cells	0.09	0.319	0.828
Memory B cells	-0.07	0.320	0.780
CD8 Tem Early	-0.07	0.354	0.812
Treg naive	0.08	0.390	0.844
Neutrophils, CD16 expr.	-0.10	0.401	0.822
T Follicular helper	0.07	0.441	0.860
Lymphocytes	0.04	0.574	1.066
NK CD56 <sup>BRIGHT</sup>	-0.03	0.580	1.029
CD8 Naive	0.05	0.589	0.999
CD8 CM	-0.05	0.651	1.058
CD8 T cells	0.03	0.657	1.026
B cells	-0.06	0.669	1.003
CD8 Tem Late	0.09	0.688	0.994
CD56 <sup>+</sup> T cells	0.03	0.695	0.968
CD8 Tem	0.02	0.697	0.938
CD4 Tem Late	0.03	0.734	0.954
CD4 CM	-0.05	0.756	0.951
CD4 Tem Early	-0.03	0.789	0.962
CD8 TemRA Late	0.02	0.794	0.938
Plasmablasts	0.04	0.807	0.926

Table B.5: Repeated sensitivity analysis in men when monocytes were restricted to be HLADR<sup>+</sup> (*continued*)

Immune cell subset	$\rho$	P value	FDR*
Naive B cells	-0.05	0.838	0.934
CD8 TemRA	0.02	0.853	0.924
CD8 TemRA Early	-0.01	0.929	0.980
Treg memory	0.03	0.966	0.992
CD4 Tem	0.00	0.994	0.994

*Note:*

The (Spearman) associations are ordered by p value, with the lowest p values shown at the top. Associations that were selected with a FDR lower than 15% are shown in bold

\* False Discovery Rate (estimated)

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Table B.6: Repeated sensitivity analysis in women when monocytes were restricted to be HLADR<sup>+</sup>

Immune cell subset	$\rho$	P value	FDR*
<b>Neutrophils</b>	<b>0.40</b>	<b>&lt;0.001</b>	<b>0.000</b>
<b>Monocytes (HLADR<sup>+</sup>)</b>	<b>0.23</b>	<b>0.004</b>	<b>0.074</b>
<b>CD56<sup>+</sup> T cells</b>	<b>-0.20</b>	<b>0.01</b>	<b>0.124</b>
<b>CD16<sup>-</sup> monocytes (HLADR<sup>+</sup>)</b>	<b>0.18</b>	<b>0.014</b>	<b>0.139</b>
<b>CD4 TemRA Late</b>	<b>-0.13</b>	<b>0.015</b>	<b>0.120</b>
CD4 Tem Early	0.20	0.032	0.208
CD4 Tem	0.15	0.071	0.395
NK CD56 <sup>BRIGHT</sup>	-0.15	0.078	0.380
CD8 TemRA	-0.09	0.09	0.390
Transitional B cells	-0.16	0.096	0.375
CD8 TemRA Late	-0.08	0.099	0.350
NK CD56 <sup>DIM</sup>	-0.15	0.133	0.433
NK cells	-0.13	0.149	0.446
CD8 CM	0.14	0.167	0.466
CD4 Tem Late	-0.04	0.235	0.610
CD4/CD8 ratio	0.11	0.251	0.612
CD4 TemRA Early	0.09	0.274	0.628
CD8 T cells	-0.10	0.324	0.701
CD4 CM	0.07	0.34	0.697
T Follicular helper	0.09	0.345	0.674
CD4 Naive	-0.11	0.346	0.642
CD8 Naive	-0.13	0.359	0.637
Treg memory	0.07	0.375	0.636
CD8 TemRA Early	-0.02	0.423	0.688
Memory B cells	-0.09	0.428	0.668
CD16 <sup>+</sup> monocytes (HLADR <sup>+</sup> )	0.09	0.441	0.661
Plasmablasts	-0.07	0.458	0.662
T cells	-0.05	0.49	0.683
CD8 Tem Early	0.12	0.521	0.701
Lymphocytes	-0.02	0.528	0.686
B cells	-0.06	0.578	0.727
Neutrophils, CD16 expr.	-0.01	0.729	0.888
Treg naive	-0.04	0.78	0.921
CD8 Tem	0.02	0.814	0.934

Table B.6: Repeated sensitivity analysis in women when monocytes were restricted to be HLADR<sup>+</sup> (*continued*)

Immune cell subset	$\rho$	P value	FDR*
CD4 TemRA	0.00	0.914	1.018
CD8 Tem Late	0.01	0.953	1.032
Regulatory T cells	0.01	0.965	1.017
CD4 T cells	0.01	0.967	0.992
Naive B cells	0.00	0.985	0.985

*Note:*  
The (Spearman) associations are ordered by p value, with the lowest p values shown at the top. Associations that were selected with a FDR lower than 15% are shown in bold  
\* False Discovery Rate (estimated)

Table B.7: Additional analysis in men (n=140) to test associations with frailty when percentages of subpopulations are used instead of absolute numbers

Immune cell subset	$\rho$	P value	FDR*
Transitional B cells, (%)	-0.22	0.011	0.373
CD4 Naive, (%)	0.22	0.018	0.302
CD4 CM, (%)	-0.24	0.023	0.260
T cells, (%)	0.16	0.047	0.395
NK cells, (%)	-0.11	0.091	0.621
CD4 TemRA Late, (%)	0.11	0.169	0.958
B cells, (%)	-0.14	0.181	0.880
NK CD56 <sup>BRIGHT</sup> , (%)	0.10	0.185	0.787
CD4 Tem, (%)	-0.10	0.186	0.702
CD4 Tem Early, (%)	-0.11	0.204	0.695
CD8 T cells, (%)	-0.11	0.209	0.644
CD4 T cells, (%)	0.12	0.213	0.603
Treg memory, (%)	-0.09	0.215	0.562
NK CD56 <sup>DIM</sup> , (%)	-0.08	0.247	0.600
CD4 TemRA, (%)	0.10	0.260	0.589
CD8 Naive, (%)	0.08	0.342	0.727
Regulatory T cells, (%)	0.05	0.425	0.849
CD8 CM, (%)	-0.03	0.464	0.877
CD8 Tem Early, (%)	-0.07	0.469	0.840
CD8 Tem Late, (%)	0.11	0.474	0.806
CD4 TemRA Early, (%)	0.06	0.573	0.927
Plasmablasts, (%)	0.04	0.582	0.900
Memory B cells, (%)	-0.01	0.670	0.990
CD8 TemRA, (%)	0.03	0.676	0.958
Naive B cells, (%)	0.01	0.705	0.958
CD8 TemRA Late, (%)	0.00	0.719	0.940
CD8 TemRA Early, (%)	0.00	0.769	0.969
T Follicular helper, (%)	-0.03	0.793	0.963
CD16 <sup>-</sup> monocytes, (%)	0.01	0.838	0.983
CD4 Tem Late, (%)	-0.05	0.848	0.961
CD8 Tem, (%)	0.08	0.865	0.948
CD16 <sup>+</sup> monocytes, (%)	-0.01	0.890	0.945
Treg naive, (%)	-0.02	0.934	0.962

Table B.7: Additional analysis in men (n=140) to test associations with frailty when percentages of subpopulations are used instead of absolute numbers (*continued*)

Immune cell subset	$\rho$	P value	FDR*
CD56 <sup>+</sup> T cells, (%)	-0.03	0.992	0.992

*Note:*

The (Spearman) associations are ordered by p value, with the lowest p values shown at the top. Associations that were selected with a FDR lower than 15% are shown in bold

\* False Discovery Rate (estimated)

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Table B.8: Additional analysis in women (n=137) to test associations with frailty when percentages of subpopulations are used instead of absolute numbers

Immune cell subset	$\rho$	P value	FDR*
CD56 <sup>+</sup> T cells, (%)	-0.18	0.009	0.304
CD4 TemRA Late, (%)	-0.17	0.009	0.156
CD8 Tem, (%)	0.20	0.021	0.236
Transitional B cells, (%)	-0.21	0.022	0.188
CD4 Tem Early, (%)	0.17	0.029	0.195
CD8 Tem Early, (%)	0.18	0.056	0.319
CD8 CM, (%)	0.14	0.062	0.301
CD8 TemRA Late, (%)	-0.09	0.065	0.275
CD4 Tem, (%)	0.16	0.066	0.251
CD8 TemRA, (%)	-0.09	0.075	0.256
Naive B cells, (%)	0.13	0.086	0.264
NK cells, (%)	-0.17	0.093	0.264
CD4 Naive, (%)	-0.12	0.133	0.347
CD4 CM, (%)	0.14	0.138	0.336
CD4 T cells, (%)	0.10	0.205	0.465
Treg memory, (%)	0.07	0.229	0.486
CD4 Tem Late, (%)	-0.05	0.232	0.465
CD4 TemRA Early, (%)	0.10	0.258	0.487
T cells, (%)	0.16	0.293	0.524
CD8 T cells, (%)	-0.09	0.294	0.499
Memory B cells, (%)	-0.08	0.334	0.540
T Follicular helper, (%)	0.10	0.350	0.540
CD16 <sup>+</sup> monocytes, (%)	-0.02	0.458	0.677
Treg naive, (%)	-0.05	0.460	0.651
CD8 Tem Late, (%)	0.09	0.480	0.653
CD16 <sup>-</sup> monocytes, (%)	0.07	0.547	0.716
B cells, (%)	-0.06	0.572	0.720
CD8 Naive, (%)	-0.10	0.662	0.803
Plasmablasts, (%)	-0.04	0.694	0.814
Regulatory T cells, (%)	-0.04	0.726	0.823
CD8 TemRA Early, (%)	-0.04	0.728	0.798
NK CD56 <sup>DIM</sup> , (%)	0.02	0.825	0.877
NK CD56 <sup>BRIGHT</sup> , (%)	-0.02	0.825	0.850

Table B.8: Additional analysis in women (n=137) to test associations with frailty when percentages of subpopulations are used instead of absolute numbers (*continued*)

Immune cell subset	$\rho$	P value	FDR*
CD4 TemRA, (%)	0.00	0.893	0.893

*Note:*

The (Spearman) associations are ordered by p value, with the lowest p values shown at the top. Associations that were selected with a FDR lower than 15% are shown in bold

\* False Discovery Rate (estimated)

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## APPENDIX C

# Supplement to Chapter 4

### C.1 Supplementary methods

The initial response of the random sample of the population was 62%. Two-third\* of those who agreed to be measured again were approached for the second measurement 6 years later (T1 in our study). Everybody was invited for the subsequent measurements every 5 years with exclusion of those who died, moved too far away, emigrated and those who actively withdrew from the study. Respondents can skip one or more rounds and then participate again. Per round, 2% - 3% of the participants died or moved. From T2 onward, in every round 23% of those invited did not participate with almost 50% of them mentioning that they do not want to participate again or not this time (reasons given: no time, not interested, already often medically examined) and from 50% we got no response

#### Description HAI components

##### Systolic blood pressure

SBP was measured twice while the participant was in an upright sitting position. The two measurements were averaged. The categorization for the HAI was as follows: 2 = below 126



mmHg; 1 = 126-142 mmHg; 0 = above or equal to 143 mmHg. This is a similar to classification applied in previous studies (5, 6, 8). Subjects who reported a physician diagnosis of hypertension, or who were taking medication for hypertension, were classified in the most unhealthy category (score = 0).

### **Random blood glucose**

Random blood glucose (RBG) was determined in a peripheral blood plasma sample. Diabetes is diagnosed when the random glucose concentration is  $11.1 \text{ mmol L}^{-1}$  or higher. Hence, all values equal or above  $11.1 \text{ mmol L}^{-1}$  were assigned a score of 0. The International Diabetes Federation recommends additional screening for individuals with values between  $5.6 - 11.1 \text{ mmol L}^{-1}$  (17). Moreover, Bowen, Xuan (18) showed that a  $\text{RBG} \geq 5.6 \text{ mmol L}^{-1}$  was more strongly associated with undiagnosed diabetes than any single risk factor and remained strongly associated with undiagnosed diabetes after adjustment for traditional diabetes risk factors. Hence, values between  $5.6 - 11.1 \text{ mmol L}^{-1}$  were assigned a score of 1. Consequently, all values below  $5.6 \text{ mmol L}^{-1}$  received a score of 2. Participants who reported a physician diagnosis of diabetes or who were using medication for diabetes were coded in the highest RBG group (=0).

### **Creatinine**

Several biochemical markers were measured in all the available samples of the follow up rounds. We used plasma creatinine levels, which are associated with renal function (19). The cut-off points for creatinine were sex-specific and replicated from previous studies (6, 8). The cut-off points for women were the following: 2 = below  $70.7 \text{ mmol L}^{-1}$ ; 1 =  $70.7 - 88.4 \text{ mmol L}^{-1}$ ; 0 = above or equal to  $88.4 \text{ mmol L}^{-1}$ . For men the cut-off points were the following: 2 = below  $97.2 \text{ mmol L}^{-1}$ ; 1 =  $97.2 - 114.9 \text{ mmol L}^{-1}$ ; 0 = above or equal to  $114.9 \text{ mmol L}^{-1}$ .

**Forced vital capacity (FVC)**

Pulmonary function measurements were performed by trained paramedics using a heated pneumotachometer. Participant's expiratory pulmonary volume was measured in a sitting position while wearing a nose clip. At least three technically acceptable attempts for measuring FVC had to be achieved, of which two had to be reproducible according to ERS criteria (20). For comparability to other studies, the FVC was used as an indicator for lung function. Cut-off points were sex-specific, with the following values for FVC in men: 2 = above or equal to 3.84 L; 1 = 3.19-3.84 L; and 0 = below 3.19 L. For FVC in women, scores were the following: 2 = above or equal to 2.61 L; 1 = 2.14-2.61 L; 0 = below 2.14 L.

**Cognitive function**

Cognition was only measured in the study population aged 45 years and older. Cognitive function was measured by four neuropsychological tests: the 15 Words Verbal Learning Test (immediate and delayed recall), the Stroop Color-Word-Test, the Word Fluency Test and the Letter Digit Substitution Test. Nooyens, Bueno-de-Mesquita (21) have described these cognitive tests in more detail. From the separate test scores a summary score of global cognitive functioning was calculated. To capture the decline rate over time, the global cognitive functioning score was transformed into a z-score which was derived from the values of all the rounds together. Participants scoring below the 10th percentile on the global cognitive functioning z-score in T4 were considered cognitively frail, and assigned a score of 0 assigned for cognition. This cut-off point is consistent with the definition for cognitive frailty used in the DCS previously (22, 23). The z-score that corresponded to the 10th percentile in T4 (z-score = -1.05) was applied as the cut-off point for cognitive frailty in the other rounds of the DCS as well. Participants with a z-score between -1.05 – 0 were assigned a score of '1,' while participants with a z-score above 0

received a score of ‘2.’ As cognitive tests were only performed among participants aged 45 years or older, we assumed that participants younger than 45 years were cognitively healthy (24), and therefore they were assigned a score of ‘2.’

### **Dichotomization lifestyle factors regression analysis**

Lifestyle was defined by the following variables: sleep, physical activity, body mass index (weight and height were assessed by a health care worker and were used to calculate BMI (weight (kg) [height (m)]<sup>-2</sup>), smoking status and alcohol consumption. As these behaviors may vary over time, we accounted for this by creating dummy variables that reflected whether someone was engaged in the behavior every measurement round or not. This resulted for smoking status in the following dummy variable: non-smoker in every round versus the rest. For sleeping we took a cut-off point of 7 hours or more sleep duration per 24 hours period. Sufficient physical activity was defined as adherence to the Dutch physical activity guidelines, which recommends 30 minutes of moderate to vigorous physical activity per day on at least 5 days per week (11). BMI was dichotomized as being obese or not (BMI  $\geq$  30.0). We classified someone as an alcohol consumer when the respondent indicated to consume alcohol sometimes or regularly.

## **C.2 Supplementary figures and tables**

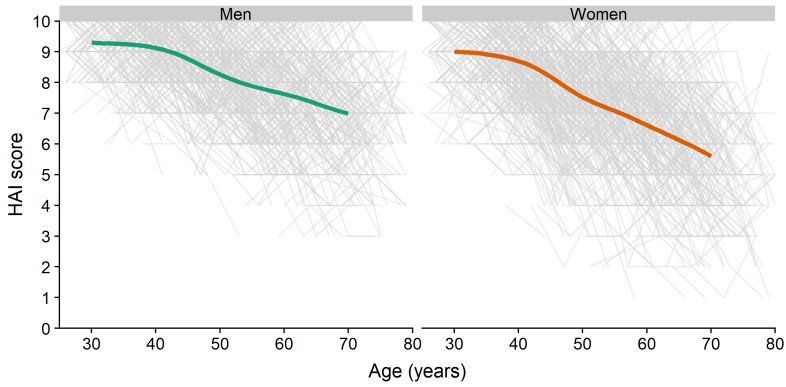


Figure C.1: Development over time of the average HAI scores for men and women

Table C.1: Response rates in the DCS of the included rounds

Period	Invited	Participated	Response	DCS round	This study
1987 – 1991	20154	12404	0.62	1	-
1993 – 1997	7768	6117	0.79	2	T1
1998 – 2002	6581	4918	0.75	3	T2
2003 – 2007	5783	4520	0.78	4	T3
2008 – 2012	5136	4018	0.78	5	T4

Table C.2: Scores at T1 for respondents who had data on all four measurements for the indicator

	Women		Men		Total
	Mean (SD)	Range	Mean (SD)	Range	N
Systolic blood pressure (mm Hg)	119.9 (15.9)	79-200	127.1 (14.6)	93-191	3439
Random glucose ( $mmol * L^{-1}$ )	5.1 (1.0)	1-17.2	5.4 (1.3)	1-18.3	3167
Creatinine	73.5 (12.6)	37-125	72.8 (12.5)	41-129	3064
FVC (L)	4.0 (0.6)	1.9-7.4	5.4 (1.0)	2.7-27.2	2303
Global cognitive function - T1	0.1 (0.7)	-2.2 - 1.9	0.1 (0.7)	-2.2 - 1.9	403
Global cognitive function - T2	0.0 (0.7)	-2.5 - 1.9	0.0 (0.7)	-2.6 - 2.2	1,390

*Note:*

Cognition was only measured in a subsample at T1

Table C.3: Scores at T4 for respondents who had data on all four measurements for the indicator

	Women		Men		Total
	Mean (SD)	Range	Mean (SD)	Range	N
Systolic blood pressure (mm Hg)	128.5 (18.0)	84-215	134.8 (17.5)	85-240	3439
Random glucose ( $mmol * L^{-1}$ )	5.2 (1.4)	2.1-24.2	5.4 (1.4)	2.4-17.2	3167
Creatinine	76.4 (14.4)	35-148	75.8 (15.0)	42-157	3064
FVC (L)	3.6 (0.6)	1.6-6.7	5.0 (0.9)	2.1-8.5	2303
Global cognitive function	-0.04 (0.8)	-4.3	-0.05 (0.7)	-4.2	1390

*Note:*

Cognition was only measured in a subsample at T1

Table C.4: Scores at T1 on the healthy aging indicators of the imputed sample

	Women		Men		Total
	Mean (SD)	Range	Mean (SD)	Range	N
Systolic blood pressure (mm Hg)	120.7 (16.2)	79-200	127.4 (14.8)	92-191	4461
Random glucose ( $mmol * L^{-1}$ )	5.2 (1.2)	1-19.6	5.4 (1.3)	1-18.3	4393
Creatinine	73.4 (12.9)	37-146	73.0 (12.6)	41-131	4330
FVC (L)	3.9 (0.6)	1.8-7.4	5.4 (1.0)	2.4-27.2	4002
Global cognitive function- T1	0.1 (0.7)	-2.2 -1.9	0.1 (0.7)	-2.2 - 1.9	403
Global cognitive function - T2	0.0 (0.7)	-2.9 - 1.9	0.0 (0.7)	-2.6 - 2.2	3,403

*Note:*

Cognition was only measured in a subsample at T1

## C.2. Supplementary figures and tables

Table C.5: Scores at T4 on the healthy aging indicators of the imputed sample

	Women		Men		Total
	Mean (SD)	Range	Mean (SD)	Range	N
Systolic blood pressure (mm Hg)	128.4 (18.0)	84-215	134.6 (17.4)	85-240	3903
Random glucose ( $mmol * L^{-1}$ )	5.2 (1.3)	2.1-24.2	5.4 (1.6)	1.5-18	3993
Creatinine	76.8 (17.3)	35-389	75.9 (16.1)	42-298	3823
FVC (L)	3.6 (0.7)	1.1-6.7	4.9 (0.9)	2.1-8.5	3518
Global cognitive function	-0.02 (0.8)	-2.5-2.7	-0.02 (0.7)	-2.5-2	3115

Table C.6: Score categories at T1 on the healthy aging indicators for complete and imputed sample

	Complete sample		Imputed sample	
	Women, N(%)	Men, N(%)	Women, N(%)	Men, N(%)
Systolic blood pressure (mm Hg)	1,818 (47.2)	1,624 (52.8)	2,342 (52.5)	2,119 (47.5)
0 (hypertension)	11.6	16.9	12.2	16.2
1 (high – normal)	20.2	33.6	21.4	34.9
2 (optimal)	68.2	49.5	66.4	48.9
Random glucose ( $mmol * L^{-1}$ )	1,656 (52.2)	1,515 (47.8)	2,301 (52.4)	2,092 (47.6)
0 (diabetes)	1.1	1.3	1.4	1.3
1 (pre stage diabetes)	20.6	30.1	20	32.4
2 (no diabetes)	78.3	68.6	78.6	66.3
Creatinine	1,600 (52.2)	1,464 (47.8)	2,314 (53.4)	2,016 (46.6)
0 (unhealthy)	11.8	0.2	11.8	0.3
1 (pre-stage)	45.2	3.4	44.3	3.5
2 (optimal)	43.1	96.5	44	96.2
FVC (L)	1,216 (52.8)	1,085 (47.2)	2,102 (52.6)	1,896 (47.4)
0 (severe airway obstruction)	3.9	3.9	6.2	7
1 (moderate airway obstruction)	14.4	20.6	17	22
2 (good lung function)	81.7	75.5	80.8	71
Global cognitive function	820 (54.9)	674 (45.1)	1,804 (53.0)	1,599 (47.0)
0 (cognitive frail)	1.1	0.6	3.4	3.8
1 (moderate cognition)	5.5	5.3	18.5	20.3
2 (good cognition)	93.4	94.1	78.1	75.9

Table C.7: Score categories at T4 on the healthy aging indicators for complete and imputed sample

	Complete sample		Imputed sample	
	Women, N(%)	Men, N(%)	Women, N(%)	Men, N(%)
Systolic blood pressure (mm Hg)	1,818 (52.8)	1,624 (47.2)	2,342 (52.5)	2,119 (47.5)
0 (hypertension)	37.2	40.9	38	41.3
1 (high – normal)	23.1	30.3	22.9	30.6
2 (optimal)	39.7	28.8	39.1	28.1
Random glucose ( $mmol * L^{-1}$ )	1,656 (52.2)	1,515 (47.8)	2,301 (52.4)	2,092 (47.6)
0 (diabetes)	6	7.3	6.1	7.4
1 (pre stage diabetes)	18.2	27.2	18.6	28
2 (no diabetes)	75.7	65.5	75.3	64.6
Creatinine	1,600 (52.2)	1,464 (47.8)	2,314 (53.4)	2,016 (46.6)
0 (unhealthy)	18.8	1.6	18.7	2
1 (pre-stage)	43.5	5.9	42.9	5.8
2 (optimal)	37.7	92.5	38.4	92.2
FVC (L)	1,217 (52.8)	1,086 (47.2)	2,102 (52.5)	1,900 (47.5)
0 (severe airway obstruction)	1.2	1.7	1.3	3.1
1 (moderate airway obstruction)	4.4	8	6	8.8
2 (good lung function)	94.5	90.3	92.8	88.2
Global cognitive function	1,712 (52.5)	1,552 (47.5)	1,804 (53.0)	1,599 (47.0)
0 (cognitive frail)	9.9	8.1	9	7.3
1 (moderate cognition)	35	38	32.2	35.7
2 (good cognition)	55.1	54	58.8	57

Table C.8: Respondent characteristics of the total sample at T1

	Women	Men
	n=3255(53%)	n=2858(47%)
<b>Socio-demographic characteristics</b>		
Mean age (SD)	46.2 (10)	46.8 (10)
<i>Age categories, %</i>		
26-35 yr	17.5	15.4
36-45 yr	32.1	31.2
46-55 yr	28.5	31.4
56-65 yr	21.8	22.1
<i>Educational level, %</i>		
Low	63.8	47.5
Medium	22	29.7
High	14.2	22.9
Employed (yes)	44.4	76
Marital status (married)	81.4	82.1
<i>Subjective health</i>		
Excellent / very good	22.9	28.3
Good / moderate	62.9	59.2
Poor	14.2	12.5
Alcohol consumption (>1 per week)	48.9	76.6
Sleep (<7 hours per night)	13.2	19
<i>Smoking status</i>		
Smoker	30.7	32.2
Ex-smoker	32.4	40.5
Never smoker	36.9	27.3
Physical exercise (no)	51.3	52.8
BMI ( $kg * m^{-2}$ ), mean (SD)	25.7 (4.2)	26.1 (3.2)
<b>Healthy Aging index indicators</b>		
Systolic blood pressure, mmHg	122.3 (17.2)	128.7 (15.7)
Random glucose, $mmol * L^{-1}$	5.3 (1.5)	5.5 (1.5)
Creatinine, $mmol * L^{-1}$	72.9 (13.0)	72.6 (12.8)
Lung function ,FVC L	3.9 (0.6)	5.3 (1.0)
Global cognitive function, z-scores *	0.03 (0.7)	0.03 (0.7)
Healthy Aging Index	8.3 (1.5)	8.5 (1.4)

\* Scores of T2 are presented due to small sample size at T1





## APPENDIX D

# Supplement to Chapter 5

### D.1 Supplementary figures

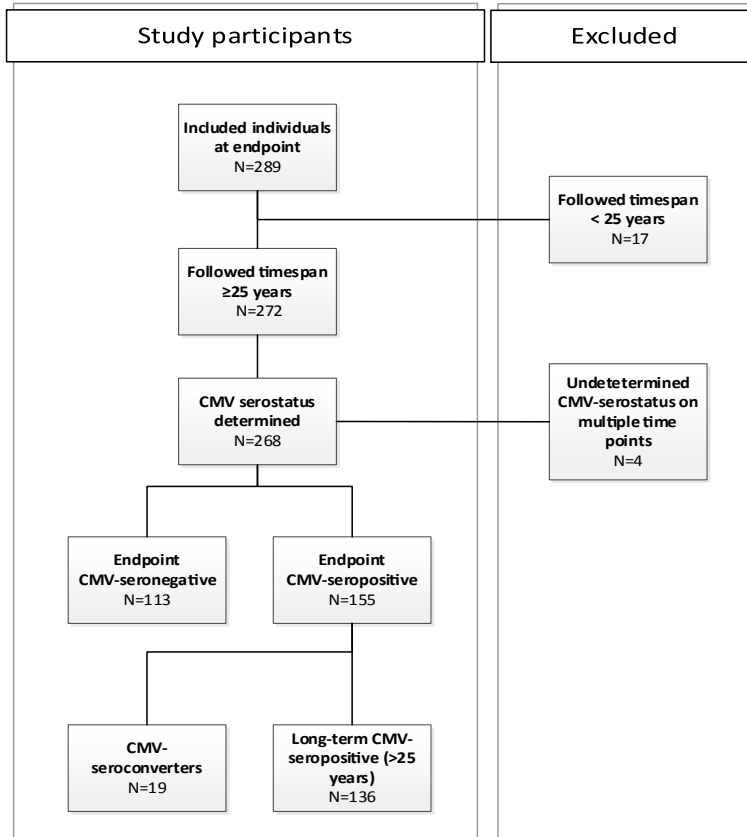


Figure D.1: Flow chart of selection of study subjects.

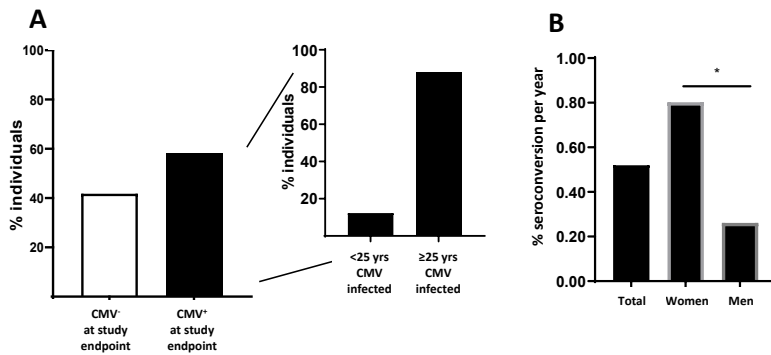


Figure D.2: CMV serostatus at study endpoint and seroconversion rate in the past 27 years. (a) Left panel: CMV serostatus at end point of all individuals. Right panel: duration of CMV infection in CMV+ individuals. (b) Seroconversion rate for total individuals and separated for women and men. Groups difference was investigated with the  $\chi^2$  test. Star indicates a P-value < 0.05.

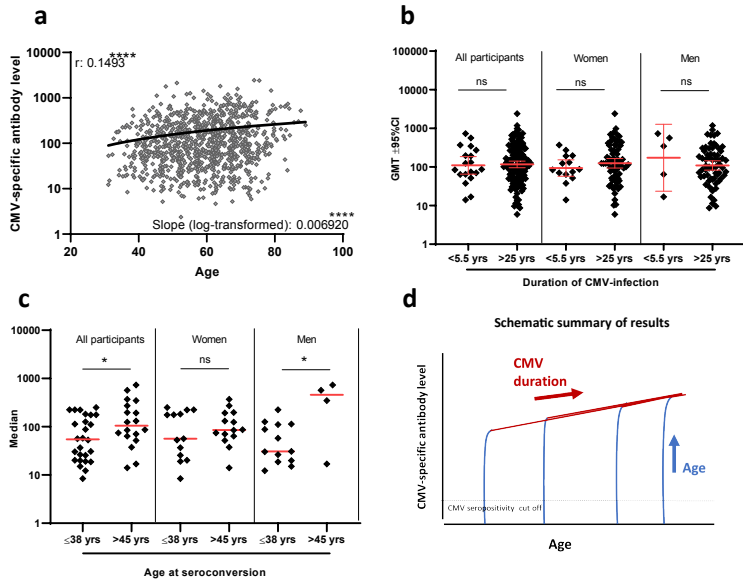


Figure D.3: CMV-specific antibody levels. (a) Results are summarized of the data of CMV-specific antibodies in both long-term CMV<sup>+</sup> individuals and CMV seroconverters. (b) Duration of CMV infection: CMV-specific antibody levels of recently seroconverted individuals (max  $<5.5$  year after CMV seroconversion,  $n = 19$ ) compared with those of long-term CMV<sup>+</sup> individuals ( $n=136$ ,  $> 25$  years CMV<sup>+</sup>), and similar comparison separately in women and men (middle and right panel, respectively). (c) Age at seroconversion: CMV-specific antibody levels of individuals that seroconverted at younger age ( $\leq 38$ yr of age,  $n = 26$ ) or older age ( $\geq 45$ yr of age,  $n = 18$ , mean age  $58.5 \pm 8.1$ , shortly after CMV seroconversion ( $< \max 5.5$  years)), with similar comparison separately in women and men (middle and right panel, respectively). (d) Schematic summary of results of CMV-specific antibody levels. GMT: geometric mean titer, 95% CI: 95% confidence interval.

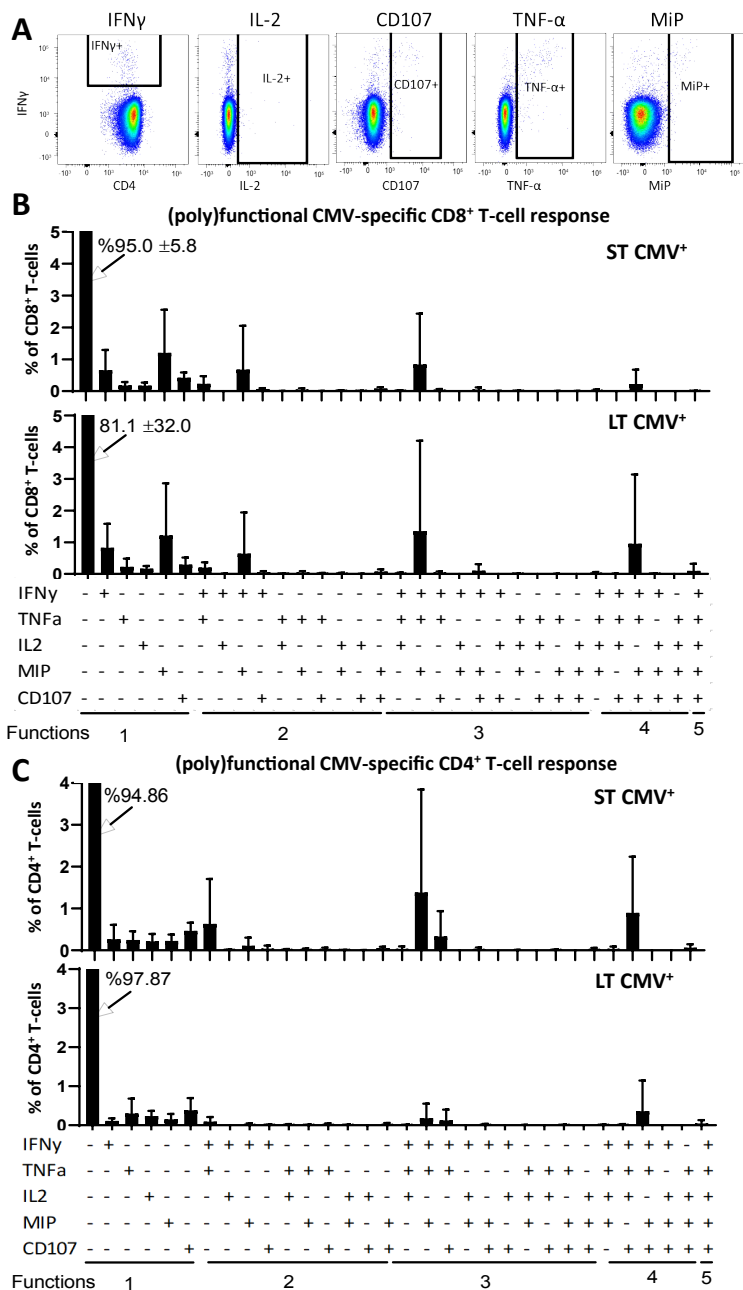


Figure D.4: (see next page)

Figure D.4 (*previous page*): Cytokine profiles after stimulation with CMV-specific peptide pools. (a) Representative flow cytometry plots in the CMV-specific T-cell stimulation assay after stimulation with CMV-specific peptides. (b,c) Percentage of cells that produce IFN $\gamma$ , TNF $\alpha$ , MIP-1 $\beta$ , CD107, IL-2, or a combination of these cytokines after in vitro stimulation of PBMCs for CD8 $^+$  T-cells (b) and (c) CD4 $^+$  T-cells.

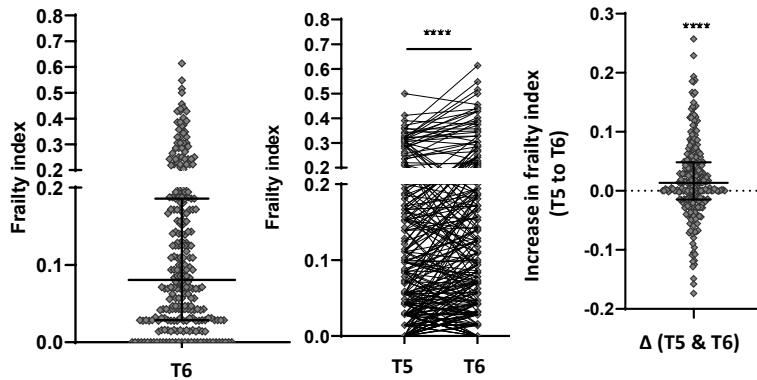


Figure D.5: Frailty index score at time point (T)6 (left panel) and the difference in frailty index score between T5 and T6 (middle and right panel).

## APPENDIX E

# Supplement to Chapter 6

### E.1 Supplementary figures and tables



Table E.1: Summary characteristics of the cytokines and chemokines in the cohort

Protein	Baseline concentration	Endpoint concentration	AUC
<b>C-reactive protein</b>			
CRP	1.03(0.93 – 1.14) * 10 <sup>6</sup>	1.22(1.11 – 1.34) * 10 <sup>6</sup>	1.56(1.43 – 1.70) * 10 <sup>6</sup>
<b>CC chemokines</b>			
CCL1/I-309	3.30(3.17 – 3.43)	3.34(3.22 – 3.46)	3.38(3.27 – 3.50)
CCL2/MCP-1	1.03(1.00 – 1.06) * 10 <sup>2</sup>	1.07(1.04 – 1.10) * 10 <sup>2</sup>	1.08(1.06 – 1.11) * 10 <sup>2</sup>
CCL5/RANTES	8.83(8.42 – 9.26) * 10 <sup>4</sup>	7.60(7.27 – 7.95) * 10 <sup>4</sup>	8.00(7.73 – 8.27) * 10 <sup>4</sup>
CCL11/Eotaxin	4.76(4.64 – 4.89)	4.97(4.84 – 5.11)	5.08(4.97 – 5.19)
CCL27/C-TACK	9.34(8.97 – 9.74) * 10 <sup>2</sup>	1.03(0.99 – 1.08) * 10 <sup>3</sup>	1.08(1.05 – 1.12) * 10 <sup>3</sup>
<b>CXC chemokines</b>			
CXCL9/MIG	3.08(2.95 – 3.23)	2.98(2.85 – 3.11)	3.15(3.05 – 3.26)
CXCL10/IP-10	2.37(2.29 – 2.46) * 10 <sup>2</sup>	2.83(2.73 – 2.93) * 10 <sup>2</sup>	3.01(2.92 – 3.10) * 10 <sup>2</sup>
CXCL11/I-TAC	6.07(5.64 – 6.53)	6.19(5.76 – 6.65)	6.98(6.59 – 7.39)
<b>Interleukins</b>			
IL-6	2.43(2.23 – 2.65)	2.31(2.13 – 2.51)	2.61(2.45 – 2.78)
IL-10	0.71(0.66 – 0.77)	0.66(0.62 – 0.72)	0.75(0.71 – 0.80)
<b>Soluble receptors</b>			
sCD14	2.15(2.10 – 2.21) * 10 <sup>6</sup>	2.21(2.15 – 2.27) * 10 <sup>6</sup>	2.27(2.22 – 2.32) * 10 <sup>6</sup>
sCD40L	4.14(3.90 – 4.40) * 10 <sup>2</sup>	3.69(3.48 – 3.91) * 10 <sup>2</sup>	4.05(3.87 – 4.24) * 10 <sup>2</sup>
sIL-6R	2.27(2.22 – 2.32) * 10 <sup>4</sup>	2.35(2.30 – 2.41) * 10 <sup>4</sup>	2.37(2.33 – 2.42) * 10 <sup>4</sup>
<b>Other</b>			
P selectin	7.95(7.43 – 8.50) * 10 <sup>4</sup>	6.80(6.40 – 7.22) * 10 <sup>4</sup>	8.04(7.73 – 8.36) * 10 <sup>4</sup>
sGP130	2.19(2.16 – 2.22) * 10 <sup>4</sup>	2.23(2.20 – 2.26) * 10 <sup>4</sup>	2.25(2.23 – 2.27) * 10 <sup>4</sup>
C5a	4.65(4.38 – 4.93) * 10 <sup>3</sup>	4.90(4.62 – 5.18) * 10 <sup>3</sup>	5.06(4.79 – 5.34) * 10 <sup>3</sup>
BDNF	1.85(1.79 – 1.91) * 10 <sup>4</sup>	1.69(1.63 – 1.77) * 10 <sup>4</sup>	1.75(1.69 – 1.81) * 10 <sup>4</sup>

*Note:*

Concentrations ( $pg * mL^{-1}$ ) and AUC values ( $years * pg * mL^{-1}$ ) are geometric mean values with 95% confidence intervals.

Table E.2: Associations between the inflammatory marker AUC levels and the change in frailty index score between the last two Doetinchem cohort study rounds

Inflammatory marker	n	P value	$\rho$	FDR	Selected
<b>Analyzed in men</b>					
CRP	52	0.03	0.43	0.47	No
CCL27/C-TACK	52	0.07	0.13	0.63	No
P selectin	52	0.11	-0.22	0.64	No
IL-6	47	0.28	-0.04	1.24	No
CXCL11/I-TAC	52	0.45	-0.07	1.63	No
sCD14	52	0.45	0.10	1.36	No
C5a	52	0.47	0.12	1.22	No
CCL11/Eotaxin	52	0.48	0.15	1.09	No
CXCL9/MIG	52	0.49	0.05	0.98	No
sGP130	47	0.60	-0.13	1.09	No
CXCL10/IP-10	52	0.64	-0.04	1.05	No
IL-10	47	0.70	-0.02	1.05	No
CCL1/I-309	52	0.77	0.10	1.07	No
CCL5/RANTES	52	0.82	0.09	1.06	No
sCD40L	52	0.87	-0.02	1.04	No
sIL-6R	52	0.89	-0.01	1.00	No
BDNF	52	0.95	0.01	1.00	No
CCL2/MCP-1	52	0.99	0.13	0.99	No
<b>Analyzed in women</b>					
sCD40L	58	0.02	-0.25	0.32	No
sCD14	58	0.05	0.31	0.41	No
IL-10	52	0.06	-0.36	0.38	No
IL-6	52	0.06	-0.37	0.29	No
CRP	58	0.09	0.18	0.33	No
P selectin	58	0.18	-0.14	0.54	No
CCL5/RANTES	58	0.18	-0.20	0.47	No
CCL11/Eotaxin	58	0.19	-0.16	0.42	No
BDNF	58	0.27	-0.23	0.54	No
CXCL9/MIG	58	0.33	-0.21	0.59	No
sIL-6R	58	0.36	0.06	0.59	No

CXCL11/I-TAC	58	0.37	-0.14	0.55	No
C5a	58	0.47	0.16	0.66	No
CCL1/I-309	58	0.70	-0.17	0.90	No
CCL2/MCP-1	58	0.79	-0.01	0.95	No
CXCL10/IP-10	58	0.80	0.04	0.90	No
sGP130	53	0.90	0.04	0.95	No
CCL27/C-TACK	58	1.00	0.00	1.00	No

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*Note:*

Permutation version of the Spearman test was used. Analyses are all stratified by initial frailty index score (in tertiles) and by age category (65-70yrs and 70-75yrs). FDR: False Discovery Rate

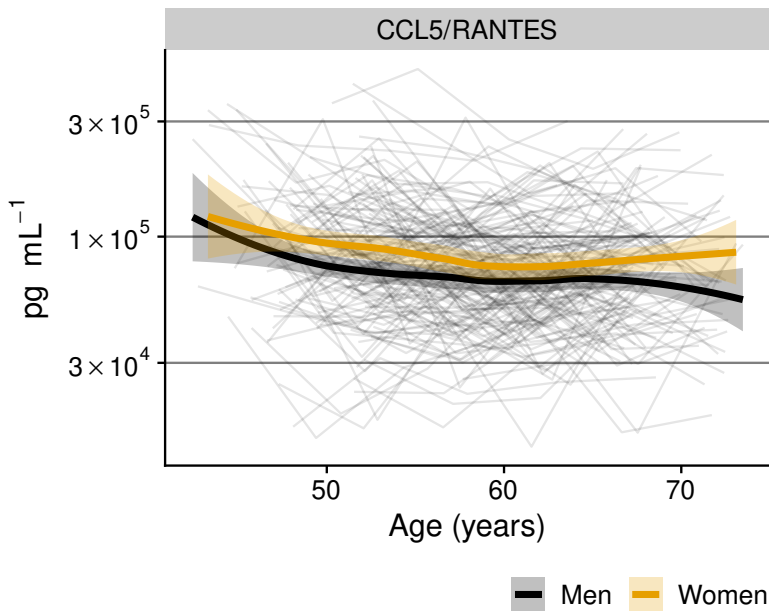


Figure E.1: The concentration of CCL5/RANTES over 20 years in men and women (showing that this is continuously higher in women).

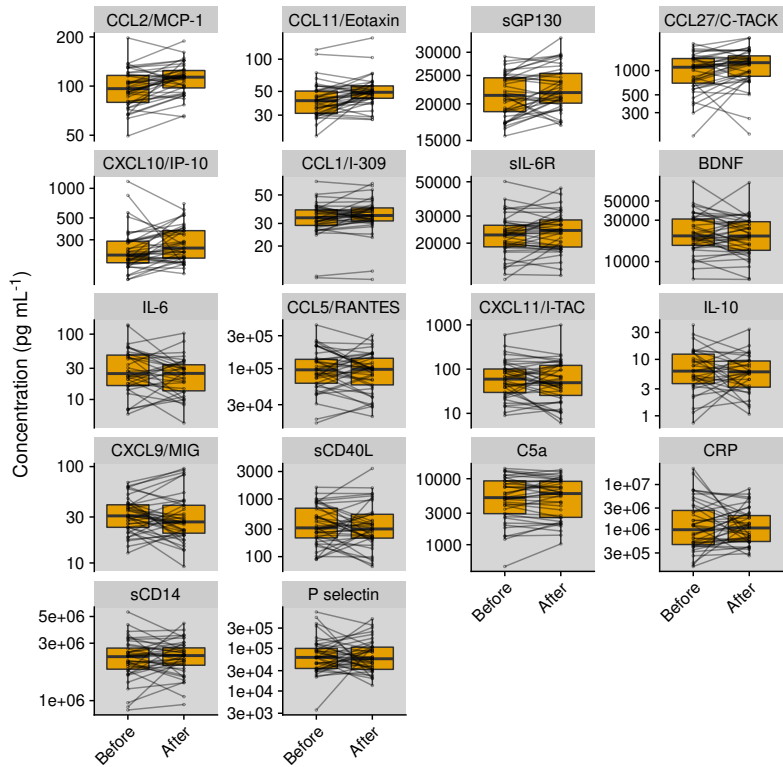


Figure E.2: Menopause is related to changes in inflammatory marker profile. Inflammatory marker concentrations are shown shortly before and shortly after menopause (average difference: 5.3 years) in women of whom data at both time-points are available ( $n=40/70$ ). Tiles of biomarkers in which an association was found with menopause are shown in a white background, others in a grey background.

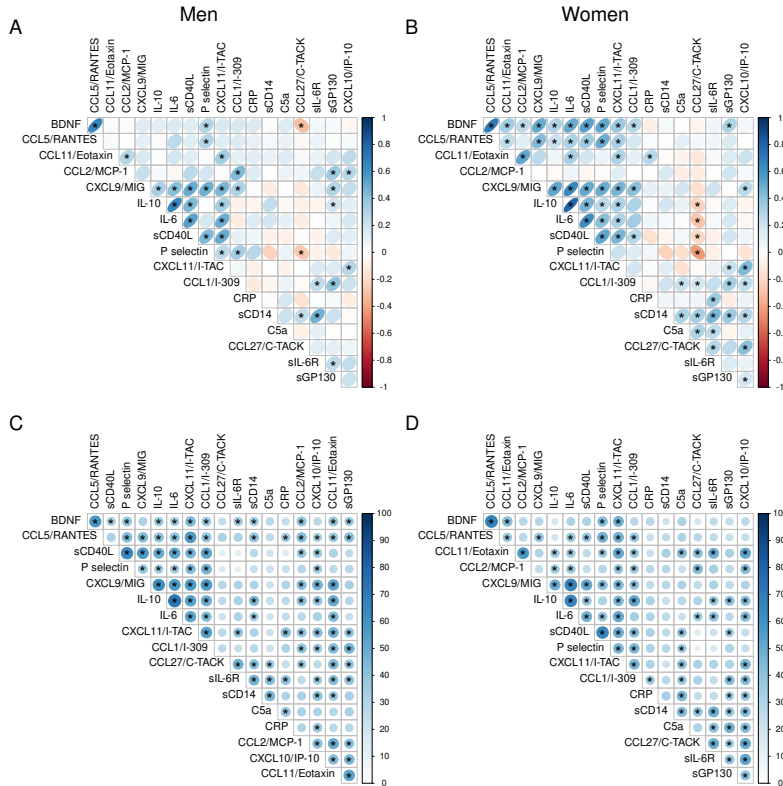


Figure E.3: Relationships between inflammatory markers shown as (A,B) correlation between pairs of inflammatory markers at study endpoint and (C,D) similarity between pairs of inflammatory marker trajectories during about 20years of follow-up. In (A,B) the direction and strength of the association is visualized with an oval shape and a color gradient. In (C,D) the blue gradient color and the size of the circles shows the percentage of participants of which a pair of biomarkers had the highest increase in concentration at the same moment in 20 years of follow-up. \* = an association between two inflammatory markers, with false discovery rate being set at a maximum of 15%. n=71 women, n=73 men.



# APPENDIX F

## Supplement to Chapter 7

### F.1 Supplementary figures and tables

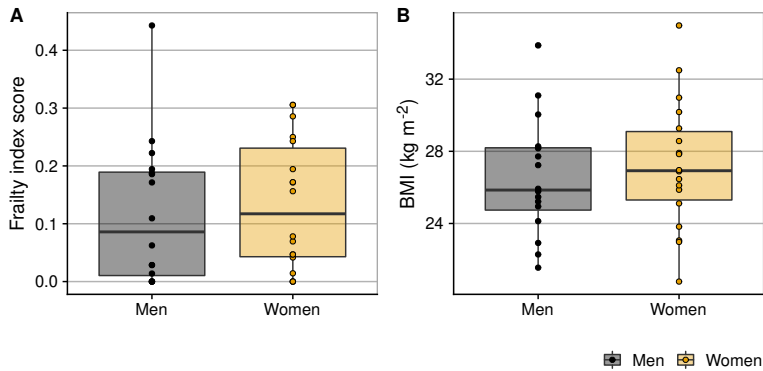


Figure F.1: Frailty index score (A) and BMI values (B) in men and women in the study population (n=16 men, n=18 women).



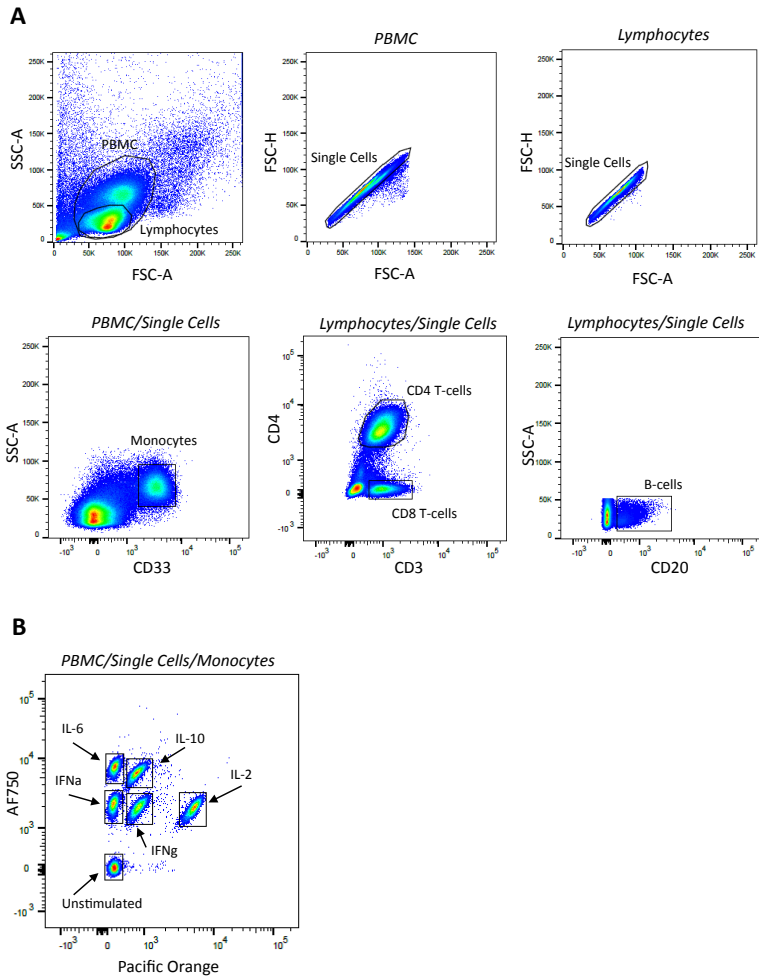


Figure F.2: (A) Phospho-flow cytometry gating strategy, showing gating of leukocytes and lymphocytes, single cells, monocytes, B cells, CD4 T cells and CD8 T cells. (B) Gating strategy to separate different stimulus conditions after barcoding and pooling samples. Gating shown is within monocytes and is representative for the strategy within the other investigated cell subsets.

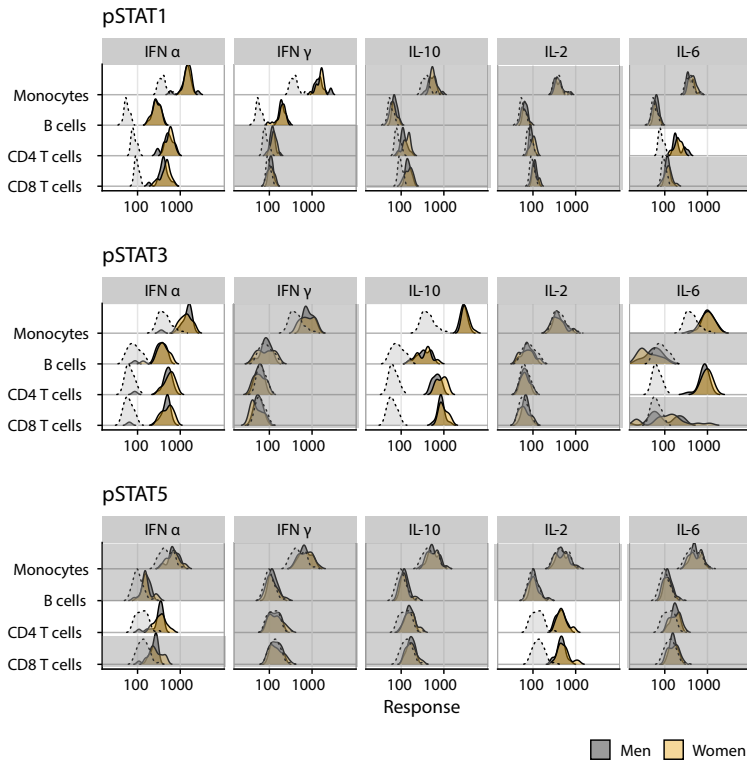


Figure F.3: Phosphorylated STAT (pSTAT1, pSTAT3, pSTAT5) expression at baseline (unstimulated, light-grey densities with dashed lines) and after stimulation in men (dark grey) and women (yellow) with IFN $\alpha$ , IFN $\gamma$ , IL-10, IL-2 or IL-6, in monocytes, B cells, CD4<sup>+</sup> T cells, CD8<sup>+</sup> T cells. Selected conditions for further analysis (median fold change >2) are shown with a white background, other plots (non-selected conditions) are presented with a dark grey background.

Table F.1: Selected phosflow conditions with fold change > 2

Stimulus	pSTAT	Fold change		
		Combined (n=32)	Men (n=16)	Women (n=18)
<b>Monocytes</b>				
IFN $\alpha$	pSTAT1	3.9	4.0	3.9
IFN $\gamma$	pSTAT1	3.9	3.9	3.8
IFN $\alpha$	pSTAT3	3.4	3.4	3.2
IL-10	pSTAT3	7.4	7.0	7.9
IL-6	pSTAT3	2.3	2.2	2.5
<b>B cells</b>				
IFN $\alpha$	pSTAT1	4.9	4.9	5.2
IFN $\gamma$	pSTAT1	3.5	3.3	3.6
IFN $\alpha$	pSTAT3	4.1	3.9	4.4
IL-10	pSTAT3	3.6	3.7	3.6
<b>CD4<sup>+</sup> T cells</b>				
IFN $\alpha$	pSTAT1	6.8	6.7	6.9
IL-6	pSTAT1	2.5	2.5	2.3
IFN $\alpha$	pSTAT3	8.1	7.4	9.0
IL-10	pSTAT3	11.9	10.7	13.4
IL-6	pSTAT3	13.9	12.9	14.8
IFN $\alpha$	pSTAT5	2.5	2.4	2.6
IL-2	pSTAT5	3.8	3.7	4.1
<b>CD8<sup>+</sup> T cells</b>				
IFN $\alpha$	pSTAT1	4.6	4.1	4.9
IFN $\alpha$	pSTAT3	7.3	6.9	8.3
IL-10	pSTAT3	15.2	12.6	16.0
IL-2	pSTAT5	3.6	3.2	4.1

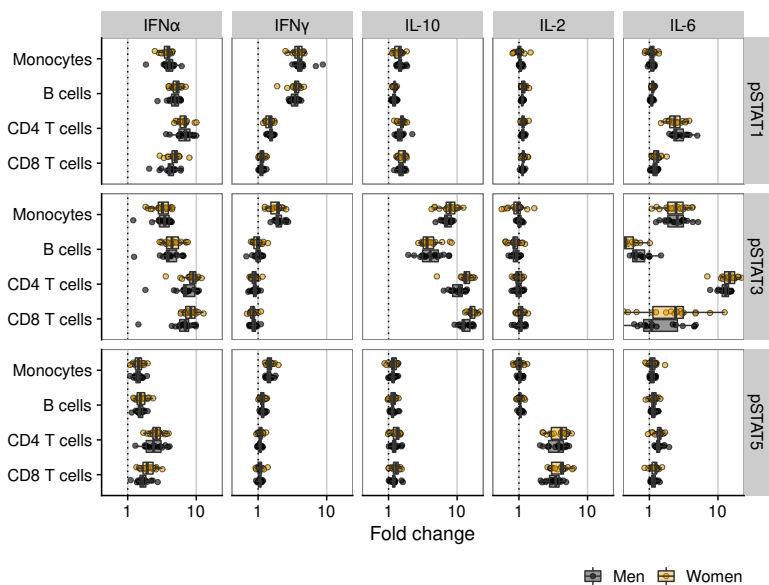


Figure F.4: Fold change (stimulus/baseline) in pSTAT levels (pSTAT1, pSTAT3, pSTAT5) after stimulation with IFN $\alpha$ , IFN $\gamma$ , IL-10, IL-2 or IL-6, in CD4<sup>+</sup> T cells, CD8<sup>+</sup> T cells, Monocytes, and B cells in men and women.

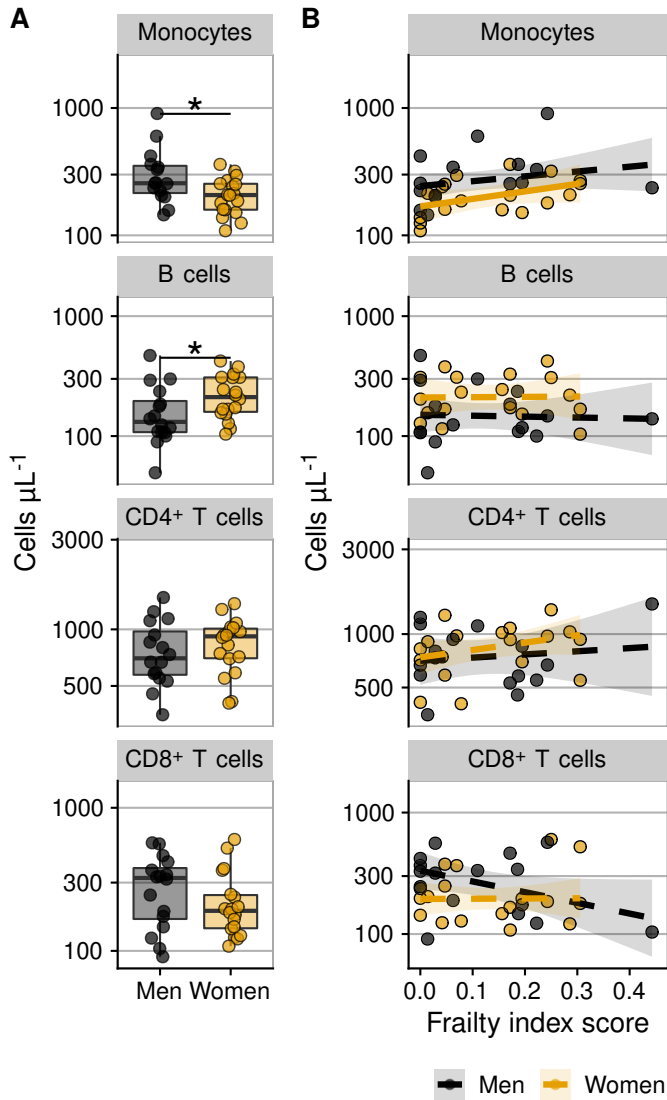


Figure F.5: (A) Association between sex and immune cell numbers. \*= association was found with false discovery rate <15%. (B) Association between immune cell numbers and frailty. Continuous trend line means association was found (with false discovery rate <15%), dashed line means that no association was found. Men: n=16, women n=18.

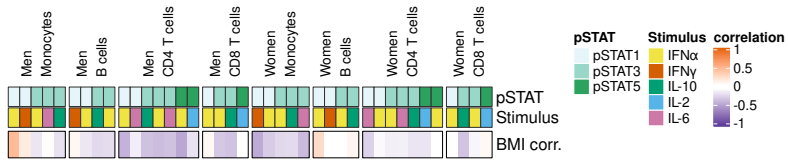


Figure F.6: Heatmap showing the relation between BMI and cellular response to cytokines detected by phosphorylation of STAT1, STAT3, and STAT5 (fold change with baseline levels) in monocytes, B cells, CD4<sup>+</sup> T cells and CD8<sup>+</sup> T cells of men and women. Every box displays the Spearman's  $\rho$  value, based on  $n=18$  women and  $n=16$  men.



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# Nederlandse samenvatting (Summary in Dutch)

Een goed werkend immuunsysteem is essentieel om gezond ouder te kunnen worden. Het is onmisbaar voor de bescherming tegen ziektekiemen, en voor het opruimen van dode cellen en afvalstoffen in het lichaam. Ook speelt het een belangrijke rol in het repareren van schade aan weefsels in het lichaam. Het immuunsysteem bestaat uit veel verschillende eiwitten, cellen en moleculen die op een zeer complexe manier met elkaar samenwerken en communiceren. Al deze verschillende immunologische cellen en stoffen houden elkaar in evenwicht. Bij een juiste balans worden schadelijke ziektekiemen snel geneutraliseerd, terwijl het eigen, gezonde weefsel niet wordt beschadigd. Als de balans tussen al deze cellen en stoffen van het immuunsysteem verstoord raakt, kan het zo zijn dat het immuunsysteem zijn werk slechter gaat vervullen, en mogelijk zelfs schade aanricht aan het eigen lichaam. Het slechter functioneren van het immuunsysteem komt vaker voor op oudere leeftijd, maar niet bij iedereen. De vraag is waarom dit bij sommige mensen wel gebeurt, en bij anderen niet. We kunnen een inzicht krijgen in het immuunsysteem doordat we verschillenden elementen en aspecten van dit systeem kunnen onderzoeken en meten. Zo kunnen we een aantal ‘immunologische biomarkers’ definiëren, die een beeld geven van het functioneren van het immuunsysteem.



Dit proefschrift is gericht op het identificeren van immunologische biomarkers van *frailty*. *Frailty* is een breed begrip en komt vaker voor naarmate men ouder wordt. Iemand is *frail* als diegene kwetsbaar is, een relatief hoge kans op ongunstige gebeurtenissen heeft zoals een val of een delirium, en minder goed van zulke ongunstige gebeurtenissen kan herstellen (Clegg et al., 2013; Kenneth Rockwood & Howlett, 2019; Walston et al., 2006). Er worden in dit proefschrift nieuwe inzichten gepresenteerd in hoe gemeten waardes van immunologische biomarkers veranderen met de (leef)tijd in een populatie van verouderende personen, en hoe deze veranderingen mogelijk samenhangen met de kans om *frail* te worden. Ook wordt onderzocht of de communicatie tussen immunologische eiwitten (cytokines) en immuuncellen is verstoord. Dit doen we door de reacties te meten die plaatsvinden binnenin immuuncellen wanneer ze met deze cytokines worden gestimuleerd, via een specifiek activatiemechanisme ('*JAK/STAT pathway*').

In **Hoofdstuk 2** beschrijven we de personen van wie er data worden gebruikt in dit proefschrift. Het zijn mensen uit de omgeving van Doetinchem die vanaf 1987 elke vijf jaar deel hebben genomen aan de 'Doetinchem Cohort Studie,' waarin veel van informatie over hun gezondheid is verzameld. Met allerlei onderzoeken zijn hun lichamelijk, cognitief en psychisch functioneren gemeten, en er zijn telkens meerdere bloedmonsters afgenomen. Verder ontwikkelen we in dit hoofdstuk een manier om *frailty* in een getal uit te drukken, door een '*frailty index score*' (K. Rockwood et al., 2011) samen te stellen in het Doetinchem Cohort. Deze score bestaat uit 36 verschillende kenmerken van ongezondheid welke bij de proefpersonen waren gemeten, of met een vragenlijst uitgevraagd. Analyses lieten zien dat hogere scores voor *frailty* gecorreleerd waren met kalenderleeftijd, zoals verwacht. Ook hingen hogere scores samen met een hoger sterfterisico, ongeacht de kalenderleeftijd. Dit komt overeen met wat gevonden is in andere studies waarin ook een *frailty index* was ontwikkeld en gebruikt (Collerton et al., 2012; K. Rockwood et al., 2011; Schoufour et al., 2017; Searle et al., 2008).

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In het bloed van 289 personen uit de Doetinchem cohort studie bepaalden we concentraties van leukocyten (witte bloedcellen). Leukocyten kunnen op basis van hun afstamming vanuit de stamcellen worden onderverdeeld in *myeloïde* en *lymfoïde* cellen. In **Hoofdstuk 2** zagen we in mannen en vrouwen met een hoge score voor *frailty* een groter aantal myeloïde cellen. Ook zagen we in dezelfde groep een chronisch hogere concentratie van de ontstekingsmarker CRP (*'C-reactive protein'*) over een tijdsspanne van 20 jaar. Dit suggereert dat chronische inflammatie (ontsteking) een rol speelt in het proces waardoor mensen *frail* worden op oudere leeftijd.

Een grote groep verschillende immuuncellen vormt samen het *'cellulaire immuunprofiel'*. Bij het meer in detail bestuderen hiervan vielen verschillen tussen mannen en vrouwen op: vrouwen hadden over het algemeen meer B – en T cellen, maar minder monocytten in het bloed (**Hoofdstuk 3**). Ook zagen we dat mannen en vrouwen een verschillend profiel hadden dat met *frailty* samenhang (**Hoofdstuk 3**). Dit profiel liet zien dat verschillen in aantallen immuuncellen mogelijk een grotere rol spelen bij vrouwen dan bij mannen in de processen die tot *frailty* leiden. Kenmerkend in dit profiel waren de hogere aantallen myeloïde cellen, zoals neutrofielen en monocytten. Deze myeloïde cellen hadden een sterkere associatie met *frailty* dan T- of B cellen; de associatie met *frailty* was ook sterker dan die tussen *frailty* en cellen waarvan bekend is dat de concentraties veranderen met hogere leeftijd, zoals lagere aantallen 'naïeve' CD8<sup>+</sup> T cellen maar juist hogere aantallen 'inflammatie regulerende' T cellen en hogere aantallen 'geheugen' T cellen, dat wil zeggen 'rijpe' cellen die zich niet verder differentiëren of delen. Deze veranderingen in het aantal T cellen met leeftijd hoeven dus niet te betekenen dat iemand *frail* is.

In **Hoofdstuk 4** worden verouderingspatronen in de algemene populatie in Nederland geanalyseerd, waarvoor we de Doetinchem studie als representatief hebben genomen. Dit deden we door een *'healthy aging index'* samen te stellen. De *healthy aging index* heeft minder variabelen dan de uitgebreide

*frailty index*, maar toch nog genoeg om een schatting te maken van de algehele lichamelijke gezondheid. Deze score kon gemeten worden over de tijd in de proefpersonen van het Doetinchem cohort. Hierdoor konden we met verschillende algoritmes verouderingspatronen in de algehele bevolking onderzoeken over meerdere jaren. Zoals verwacht, zagen we een afname in deze score, en dus tekenen van achteruitgang van lichamelijke functies, relatief vroeg in de levensloop, vanaf ongeveer 45 jaar. In mannen identificeerden we twee verschillende verouderingstrajecten, één waarin de veroudering min of meer ‘geleidelijk’ plaatsvond, en een kleinere groep waarin deze veroudering relatief snel plaatsvond. In vrouwen werd één patroon ontdekt, dat grotendeels samenviel met dat van de snel verouderende mannen. Belangrijke factoren welke een risico vormden om tot de snel verouderende groep te horen, waren een hoge ‘*Body Mass Index*’ (BMI) en een lage sociaal economische status.

Het is bekend dat een chronische cytomegalovirus (CMV) infectie van invloed is op de aantallen immuuncellen in het bloed (met name de geheugen T cellen). Omdat het immuunsysteem constant zijn best moet doen om deze virusinfectie onder controle te houden, wordt vaak aangenomen dat een CMV infectie zorgt voor snellere veroudering. In **Hoofdstuk 5** deden we onderzoek naar de rol van dit virus, en we lieten hier zien dat besmet raken met CMV niet sterk samenhangt met *frailty*. Aangezien we bloedmonsters van de proefpersonen hadden tot 30 jaar terug in de tijd, konden we ook bepalen wanneer iemand geïnfecteerd was geraakt, en zo ook hoe lang iemand geïnfecteerd was geweest met het virus. We zagen geen sterke relatie tussen de duur van infectie en *frailty*. Verder zagen we inderdaad veel hogere aantallen rijpe T cellen in personen die geïnfecteerd waren met CMV, zoals verwacht (Wertheimer et al., 2014), maar opnieuw maakte het voor deze cel aantallen niet veel uit of iemand heel lang (>25 jaar) of juist kortgeleden geïnfecteerd was. De theorie dat cumulatieve blootstelling van het lichaam aan een chronische CMV infectie voor toenemende ‘stress’ zorgt op het immuunsysteem en daarmee voor snellere

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algemene veroudering, konden daarom niet met onze resultaten ondersteund worden.

In meerdere artikelen is beschreven dat er in oudere mannen en vrouwen vaak een chronische, laaggradige inflammatie optreedt, wat mogelijk komt door een minder adequate immuunrespons (Daniel Baylis et al., 2013; Claudio Franceschi et al., 2006). Bij het gedetailleerd bestuderen van indicatoren van chronische inflammatie (inflammatoire biomarkers), en hun verandering in concentratie over de tijd, zagen we dat markers gerelateerd aan  $\text{IFN}\gamma$  en aan bloedplaatjesactivatie sterk gerelateerd waren aan veranderingen van de meeste andere markers, en dus mogelijk een centrale plaats innemen in een inflammatoir biomarkerprofiel (**Hoofdstuk 6**). We zagen ook meerdere duidelijke verschillen tussen mannen vrouwen in de biomarker concentraties en veranderingen daarin. Deze verschillen werden minder duidelijk na de leeftijd van ongeveer 60 jaar. Mogelijk spelen hier hormonale veranderingen rond de menopauze een rol. De inflammatoire markers geassocieerd met *frailty* waren de markers gerelateerd aan de IL-6 signalerings ‘*pathway*’. In vrouwen zagen we ook dat met de leeftijd toenemende concentraties van sCD14, een marker van monocytten activatie, gerelateerd waren aan *frailty*. De meeste associaties verdwenen na het corrigeren voor BMI; dit laat zien dat BMI en overgewicht mogelijk belangrijke factoren kunnen zijn die het immuunmarker profiel dat gerelateerd is aan *frailty* mede bepalen.

Als laatste hebben we nog onderzocht of de functie van immuuncellen anders is in kwetsbare, ‘frail,’ ouderen vergeleken met gezonde ouderen (**Hoofdstuk 7**). Hier zagen we dat de immuuncellen van *frail* ouderen de signalen van specifieke immuunmarkers (‘cytokines’) minder goed verwerken en minder goede respons (reactie) hebben op die signalen. Dit zagen we in B cellen, in T cellen, en in monocytten. De responsen die we hier gemeten hadden waren ‘pSTAT’ responsen (pSTAT1, pSTAT3, en pSTAT5). Dit kan een teken zijn dat laaggradige inflammatie het immuunsysteem ‘uitput,’ met verminderde signalering via de zogenoemde ‘JAK/STAT pathway’ tot gevolg,

wat mogelijk dus leidt tot slechtere immuunresponsen in *frail* personen. Van belang is verder ook dat verminderde IL10-geïnduceerde pSTAT3 responsen werden gezien in zowel mannen die *frail* waren als in mannen met chronisch hogere levels van een ontstekingsmarker (CRP). Aangezien IL10 een immuunregulerende werking heeft, zou het dus kunnen zijn dat *frail* mannen minder goed immuunreacties kunnen ‘dempen.’

## Conclusie

Het is een cruciale vraag waarom sommige mensen gezond oud worden en anderen niet. Om deze vraag te beantwoorden is het zeer belangrijk te weten wat de rol van het immuunsysteem hierin is. Dit proefschrift is erop gericht om immunologische biomarkers te vinden welke gerelateerd zijn aan veroudering en, met name, *frailty* op oudere leeftijd. Specifiek keken we naar hoe verschillende immunologische biomarkers met elkaar in verband staan, hoe de concentraties van deze biomarkers veranderen over de tijd, en hoe deze veranderingen zijn gerelateerd aan klinisch relevante tekenen van immuunveroudering. Verder hebben we onderzocht welke immunologische mechanismen belangrijk zijn in het ontwikkelen van een sluimerende chronische ontsteking.

We vonden inderdaad een ‘immuunprofiel’ gerelateerd aan *frailty*. We zagen tekenen van ontregeling van immuuncellen uit de myeloïde cellijn. Ook zagen we verschillen tussen mannen en vrouwen; vrouwen lieten meer en ook sterkere associaties zien tussen cellulaire en moleculaire immuunmarker en *frailty*. Dit laat zien dat het van belang is om mannen en vrouwen apart te onderzoeken bij het bestuderen van veroudering. Gebaseerd op onze resultaten lijkt een chronische CMV infectie minder dan verwacht *frailty* te beïnvloeden. Ten slotte zagen we dat het verwerken van signalen van cytokines mogelijk verstoord is in immuuncellen van *frail* ouderen. Mogelijk kunnen deze resultaten houvast bieden om in de toekomst een behandeling te

vinden tegen veroudering van het immuunsysteem. Een belangrijke hypothese van ons was dat laaggradige inflammatie vooraf gaat aan het ontstaan van *frailty*. Hoewel we geen definitieve conclusies kunnen trekken, vonden we inderdaad aanwijzingen welke deze hypothese ondersteunen.

De resultaten uit dit proefschrift laten zien hoe het immuunsysteem betrokken is in het zeer complexe, multifactoriële verouderingsproces. Nieuwe immuunmarkers zijn hier ontdekt, welke verschillen tussen mannen en vrouwen en een aanvulling zijn op de al bekende klinische markers gerelateerd aan veroudering. We hopen daarom dat dit proefschrift een opstap is voor vervolgstudies om nieuwe technieken toe te passen in cohorten waarin proefpersonen langdurig gevolgd zijn, om zo uiteindelijk steeds beter te kunnen voorspellen welke personen gezond verouderen en welke niet.



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