

Association between Immunosenescence Phenotypes and pre-frailty in Older Subjects

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Published in:

The Journals of Gerontology. Series A, Biological Sciences and Medical Sciences

DOI:

[10.1093/gerona/gly135](https://doi.org/10.1093/gerona/gly135)

Publication date:

2019

License:

Unspecified

Document Version:

Accepted author manuscript

[Link to publication](#)

Citation for published version (APA):

Dinh, H. C., Bautmans, I., Beyer, I., Mets, T., Onyema, O. O., Forti, L. N., ... Jansen, B. (2019). Association between Immunosenescence Phenotypes and pre-frailty in Older Subjects: Does Cytomegalovirus Play a Role? *The Journals of Gerontology. Series A, Biological Sciences and Medical Sciences*, 74(4), 480-488. <https://doi.org/10.1093/gerona/gly135>

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1 **Abstract**

2 Frailty is highly prevalent in old age and confers an important mortality risk. Although the causes
3 of frailty are multiple, immuno-senescence (IS) - predominantly driven by cytomegalovirus (CMV)
4 - has been implicated in the pathophysiology of the syndrome. Thus far, research examining the
5 association between IS and frailty is sparse and equivocal. Therefore, we aimed to clarify the
6 impact of CMV on IS and its relevance to the frailty concept. 173 persons aged 80 to 99 years were
7 enrolled. Pre-frailty was defined according to Fried's criteria. Anti-CMV IgG and serum IL-6 were
8 measured using Architect iSystem and Luminex, respectively. T-cell phenotypes were determined
9 using flow cytometry. The prevalence of pre-frailty was 52.6%, increased with age ($p=0.001$), and
10 was greater in men than women ($p=0.044$). No relationship was found between pre-frailty and
11 positive CMV serology. CMV-seropositivity was significantly associated with less naïve cells,
12 more memory and senescence-prone phenotypes (all $p<0.001$). Further, high IL-6 levels, more
13 memory, and less naïve T-cells were separately associated with pre-frailty (all $p<0.05$) in CMV-
14 negative, but not positive, subjects. After adjusting for potential confounders, however, only IL-6
15 was predictive of pre-frailty. We conclude that the presence of pre-frailty is independent from
16 CMV infection in very old subjects.

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21 **Key words:** Senescence, Inflammation, Lymphocytes, Robust

1 **Introduction**

2 Frailty is a complex geriatric syndrome that results from a decreased physiologic reserve in
3 multiple organ systems, to the extent that minor stress will put a number of physiological systems
4 beyond the threshold of symptomatic clinical failure (1). It is very prevalent among older people -
5 with estimated prevalence of up to one-third of those aged 80 years and over - and is associated
6 with an increased risk of disability, falls, morbidity, hospitalization, institutionalization and death
7 (2-4). Given the expanding older adult population, the numbers of frail older people will increase,
8 particularly as the current numbers of the oldest old are predicted to triple over the next 30 years
9 (<http://www.un.org/esa/population/publications/worldageing19502050/>).

10 A pre-frail state has been described by several researchers as an incomplete physical frailty
11 phenotype (5-7) and pre-frail older adults have more than twice the risk of becoming frail than
12 robust ones (8). In a recent meta-analysis pre-frailty was shown to be a high risk factor for mortality
13 (OR=1.761 [1.359, 2.282], HR/RR=1.466 [1.323, 1.624]), disability in basic (OR=1.855 [1.347,
14 2.556], HR/RR=1.587 [1.442, 1.747]) and instrumental (OR=2.302 [1.947, 2.721]) activities of
15 daily life, physical limitations (OR=1.813 [1.412, 2.328], HR/RR=1.484 [1.328, 1.658]), falls
16 (HR/RR=1.167 [1.049, 1.299]), and hospitalization (OR=1.527 [1.191, 1.959], HR/RR=1.148
17 [1.063, 1.239]) (4). Notwithstanding, it has been shown that frailty is a highly dynamic condition
18 that can revert, particularly in pre-frail individuals (9). Scientific evidence suggests that pre-frail
19 older adults respond more successfully to physical interventions than those who have already
20 moved to a frail state (10). Therefore, pre-frail older persons can be considered as an important
21 target group to counter frailty. However, the clinical and physiological profiles of pre-frail older
22 adults are scarcely described in literature, especially for the oldest old.

1 Although the pathophysiology of (pre-)frailty needs further elucidation, given its complex and
2 multifactorial nature, there is growing evidence for the involvement of immuno-senescence (IS)
3 and its associated conditions in the development of the syndrome (11,12). Inflammaging (13,14)
4 and Immune Risk Profile (IRP) (15) are two recent concepts regarding IS that are increasingly
5 being recognized to be, at least in part, the cause of increased susceptibility to frailty and death in
6 older subjects. Inflammaging refers to a chronic low grade inflammatory profile (CLIP) with
7 advancing age, and emerging studies have shown that this heightened inflammatory state may play
8 a central role in the pathogenesis of pre-frailty and frailty, either by promoting protein degradation,
9 or through its deregulation of other metabolic pathways (16). On the other hand, IRP is
10 characterized by a shift in T-cell sub-population types manifested by decreased CD4+/CD8+ T-
11 cell ratio, lower numbers and proportions of naïve and early-differentiated T-cells (defined as cells
12 expressing the costimulatory molecule CD28 and lacking the cell surface receptor CD57), with a
13 concomitant accumulation of highly differentiated memory and senescent T-cells, identified by the
14 expression of CD57 and/or absence of CD28. IRP is strongly associated with seropositivity to
15 chronic viral infections such as cytomegalovirus (CMV), suggesting that CMV infection may be a
16 driving force behind the shifts in T-cell subsets. Indeed, age-related increase of memory CD8+ T-
17 cells, is paralleled by an increase in the proportion of CMV epitope-specific T-cells. Khan et al.
18 portrayed that individual CMV epitope-specific CD8+ T-cells could represent up to 23% of the
19 total CD8+ T-cells in older adults with CMV infection (17,18). This clonal expansion of CMV-
20 specific CD8+ T-cells is thought to exacerbate human T-cell IS, and thereby increase the
21 susceptibility to inflammatory processes (19). Also, high levels of CMV IgG antibodies have been
22 inconsistently reported to be associated with an increased risk of pre-frailty. In women under 80
23 years of age, Wang et al. reported an increased prevalence of pre-frailty in those with high CMV
24 antibody concentrations compared to CMV-seronegative women (20). Therefore, cellular

1 mechanisms - in concert with alterations in inflammatory processes - may be implicated in the (pre-
2)frailty syndrome.

3 Our understanding of the effects of multiple deregulations in the T-cell pool in mediating frailty
4 with advancing age is imperfect. In a study of community-dwelling adults aged 55 years and over,
5 frailty and pre-frailty were predicted by the frequency of terminal effector CD8⁺ T cells (21).
6 However, in a large population based study on persons older than 85, an inverse relationship of
7 memory/naïve CD8 T-cell ratio with pre-frailty was observed, that was contrary to expectation
8 (22). Additionally, in a large population-based study of 724 community dwelling women, Schmaltz
9 and colleagues (12) could not confirm the results of Wang et al. (20) indicating an increased
10 prevalence of pre-frailty in subjects with higher levels of CMV antibodies.

11 In light of this ongoing controversy and limited available data, we sought to clarify the impact of
12 CMV on the relationship between IS phenotypes and pre-frailty in community-dwelling older
13 subjects.

14

15 **Method**

16 **Participants and study design**

17 The BrUssels sTudy on The Early pRedictors of FraiLty (BUTTERFLY) is an ongoing
18 longitudinal study - organized by the Vrije Universiteit Brussel, Universiteit Ziekenhuis Brussel,
19 and Universiteit Gent - designed to identify the determinants for active and healthy aging and for
20 early stages of frailty in the oldest old. Apparently healthy older individuals (≥ 80 years old) who
21 presented no acute pathology, able to walk, and living independently in the community were
22 recruited for this observational study. Recruitment was done - between February 2015 and February

1 2017 - by advertisement through day centres, health insurance companies, seniors associations,
2 general practitioners, municipalities, and other public places. Participants were excluded if they
3 met any of the following criteria: acute pathology, cognitive impairment (unable to understand
4 instructions and/or mini mental state examination score < 24/30); diagnosis of cancer during the
5 past 6 months; undergone surgery, radiotherapy, or chemotherapy within the past 6 months or
6 scheduled in the near future. When eligible, the subjects were examined by a team of MDs and
7 researchers to determine whether they portrayed any sign of frailty. Frailty was operationalized
8 using 3 well known definitions: Fried's frailty phenotype focusing mostly on physical frailty, the
9 Groningen Frailty Indicator with a mainly psychosocial approach, and the Rockwood Frailty Index,
10 which focuses on the medical aspects of frailty. Potential participants were excluded if they were
11 identified as frail based on the Fried's criteria. This paper was based on the baseline data of the
12 first 173 included subjects - 81 women and 92 men - and, for the purpose of the present report,
13 only physical frailty was considered since Fried's frailty index is the only one that identifies pre-
14 frail subjects. The study protocol was approved by the local ethics committee in accordance with
15 the Declaration of Helsinki and each participant gave a written informed consent.

16

17 **Flow cytometry analysis**

18 Venous blood specimens were collected in the morning for serum (stored at -80°C until analysis)
19 and for EDTA anticoagulated blood. Peripheral blood leucocytes were recovered as described
20 previously (23). Briefly, EDTA blood was exposed to lysis buffer for 10 min. After lysing the red
21 blood cells, the blood leucocytes were centrifuged at 2,800 rpm for 4 min. Thereafter, the cells
22 were isolated, washed twice in PBS containing 1% BSA at 2,800 rpm for 3 min, and re-suspended
23 in 200µl PBS containing 1% BSA.

1 Antibodies were initially titrated to determine the optimal conditions for flow cytometry analysis
2 before staining. About 5×10^5 cells were stained with 3 μ L each of PE-CY5-labeled anti-CD8
3 (Becton Dickinson, San Jose, CA, USA), PE-CY7-labeled anti-CD3 (Biolegend, San Diego, CA,
4 USA), FITC-labelled anti-CD28 (Biolegend, San Diego, CA, USA), and PE-labelled anti-CD57
5 (Biolegend, San Diego, CA, USA). After 20 min incubation at room temperature in the dark, cells
6 were washed at 2,800 rpm for 3 min, and 500 μ L of FACS flow solution (Becton Dickinson, San
7 Jose, CA, USA) were added.

8 The labelled samples were analyzed with a Coulter FC 500 flow cytometer (Beckman Coulter,
9 Fullerton, CA, USA). Data acquisition was performed using the Coulter CXP software (Epics).
10 The lymphocyte subpopulation was gated according to size and granularity in the forward vs. side
11 scattergram, and as such, dead cells were excluded. Fluorescence-minus-one controls were used to
12 distinguish positive from negative events and the various lymphocyte clusters were identified
13 according to their expression or non-expression of a combination of surface markers (see
14 Supplementary Figure 1). As CD3⁺ T-cells almost exclusively express CD4 or CD8 (24) and
15 because in a previous study - we found that at least 95% of CD3⁺ CD8⁻ cells from our subjects
16 were CD4⁺ (23) - we considered the CD8⁻ T-cells to be largely CD4⁺ T-cells (23). In this
17 perspective, CD8⁻/CD8⁺ T-cell ratio could represent an imperfect but acceptable approximation
18 of CD4⁺/CD8⁺ T-cell ratio in our setup.

19

20 **Serum CMV IgG and IL-6 determination**

21 Serum levels of CMV IgG were measured by a chemiluminescent microparticle immunoassay on
22 the ARCHITECT iSystem (Abbott Diagnostics, Abbott Park, Ireland) with an assay sensitivity and
23 specificity of 100% and 99%, respectively. Assays were regarded as positive if they had
24 concentrations of 6.0 arbitrary units (AU)/mL or greater and negative if they had concentrations of

1 less than 6.0 AU/mL. The detection limit of 6 AU/mL was based on the indications from the
2 manufacturer of the CMV IgG kit. The intra-assay and inter-assay coefficients of variation ranged
3 from 4.39% to 5.67% and from 4.87% to 6.17%, respectively. The serum levels of IL-6 were
4 determined using an IL-6 ultrasensitive singleplex Bead kit (Lifetechnologies USA). For IL-6
5 determination, the limit of detection, intra-assay and inter-assay coefficients of variation were <
6 0.05 pg/mL, 7.59%, and 9.99% respectively. All reagents were applied according to the
7 manufacturers' instructions.

8

9 **Frailty indicators**

10 Fried et al. (25) developed an operational definition of frailty containing five criteria: weight loss,
11 exhaustion, physical activity, gait speed, and grip strength. Each item is dichotomized and a total
12 score of 0 means robustness, a score of 1-2 refers to pre-frailty, while a score of 3 or more signifies
13 the presence of frailty (25). This construct of frailty is widely used, with the originally proposed
14 measures, as well as in modified constructs. Inspired by the operational definition of Fried, our
15 approach was based on four frailty characteristics suggested in previous research: weight loss,
16 exhaustion, gait speed, and grip strength (5). Weight loss was evaluated by the self-reported
17 question: *'In the last six months, have you lost more than 4,5kg unintentionally?'* which was
18 answered by yes (1) or no (0). Exhaustion was measured similarly to the original Fried phenotype,
19 questioning two statements from the CES-D Depression Scale (26): *'I felt that everything I did was*
20 *an effort'* and *'I could not get going'*. The participants were asked: *'How often in the last week did*
21 *you feel this way?'* and were scored 0 for rarely or none of the time, 1 for some or a little of the
22 time, 2 for a moderate amount of time, or 3 for most of the time. When participants scored a 2 or 3
23 on either of the two statements, they received a point on the frailty scale for exhaustion. Gait speed
24 was measured by timing the walked distance of 4.5m and was stratified for gender and height, as

1 proposed by Fried (25). Participants were scored a point for slow walking if their walking time was
2 ≥ 7 seconds in men ≤ 173 cm and women ≤ 159 cm, and if their time was ≥ 6 seconds in men $>$
3 173 cm and women > 159 cm. Grip strength was performed using the Martin Vigorimeter, a reliable
4 and practical instrument which measures handgrip strength in kPa (27). Cut-offs were 42kPa for
5 women, and 71kPa for men. Participants showing a lower grip strength received a point for this
6 item (28). The frailty scale contained 4 items and in analogy with previous research, the following
7 scoring system was put forward to assign the level of frailty: a score of 0/4 signifies robustness, 1-
8 2/4 points means pre-frailty and with a score of 3 or 4/4 one is considered frail (29).

9 Measurements of height and weight were taken and body mass index (weight (kg)/ height² (m²))
10 was calculated.

11 Medical history

12 Participants were asked whether a doctor had ever told them that they had any of the following
13 conditions: hypertension, ischemic heart disease, heart failure, peripheral vascular disorders,
14 cerebrovascular disorders, thyroid disorders, diabetes mellitus, cancer, respiratory disorders,
15 musculoskeletal conditions, osteoporosis, eye disorders, falls, skin disorders, kidney problems,
16 problems with urination, depression or anxiety.

17

18 **Statistical Analyses**

19 Statistical analysis was performed using IBM SPSS version 22.0. Data were tested for normality
20 using the Kolmogorov-Smirnov goodness of fit test. Most of the data were not normally distributed
21 even after log-transformation and as such, nonparametric tests were applied during analysis. The
22 Wilcoxon's Signed Rank test, Kruskal-Wallis and Mann-Whitney U tests were used for continuous

1 variables. Comparisons between categorical variables were performed using the chi-square test or
2 Fisher exact test, where appropriate. Spearman's rank correlations were used to determine
3 associations between participants' characteristics and CMV titers. Also, because the relationship
4 between T-cell differentiation markers and the presence of pre-frailty differed significantly by
5 CMV serostatus (p for interaction terms < 0.05), data were analyzed for CMV-seropositive and
6 CMV-negative subjects separately. A binary logistic regression was applied to explore the
7 relationship between IL-6 and T-cell differentiation markers and the risk of pre-frailty. Participants
8 were classified into three groups - of about the same number of subjects - according to the levels
9 of IL-6 as Low, < 1.4 pg/mL; Intermediate, 1.4 to 2.5 pg/mL and High, > 2.5 pg/mL and the Low
10 group was the reference group. This concentration range was chosen based on findings by other
11 authors (30) indicating that subjects aged 65 years and older are at higher risk of functional decline
12 if they have circulating levels of IL-6 greater than 2.5 pg/mL. Collinearity was assessed with the
13 variance inflation factor, and the naïve/early-differentiated phenotypes were removed due to a
14 significant collinearity with their more differentiated counterparts. Analyses were carried out with
15 and without adjustment for the potential confounders: age, sex, BMI, heart failure, use of anti-
16 inflammatory drugs, and smoking habits. Statistical significance was set *a priori* at two-sided $p <$
17 0.05 .

18

19 **Results**

20 **Descriptive statistics**

21 As portrayed in Supplementary Table 1, the overall prevalence of pre-frailty was 52.6%. Pre-frail
22 subjects were significantly older than robust individuals ($p = 0.001$) and the prevalence of pre-

1 frailty was greater in men than women ($p = 0.044$). Pre-frailty was associated with increased BMI
2 ($p = 0.019$) and low CD8⁺ counts ($p = 0.021$). The IgG for CMV was positive in 92 (53.2 %) subjects and no direct association was found between CMV seropositivity or CMV titer and pre-
3 frailty. However, more pre-frail subjects tended to have a history of heart failure compared to
4 robust ($p < 0.05$, supplementary table 2).

6

7 **IS phenotypes according to pre-frailty and CMV serostatus**

8 Table 1 shows the IS phenotypes according to pre-frailty and CMV serostatus. No significant
9 difference was found in the percentage of T-cell differentiation markers or CD8⁻/CD8⁺ T-cell
10 ratio between pre-frail and robust individuals. The pro-inflammatory cytokine IL-6 was
11 significantly higher in pre-frail subjects compared to robust ($p < 0.001$). The CMV-seropositive
12 group was characterized by a significantly higher proportion of highly differentiated memory and
13 senescence-like phenotypes, in both the CD8⁺ and the CD8⁻ sub-populations of T-cells (all $p <$
14 0.001). On the other hand, CD28⁺CD57⁻ expressing cells (mainly representing the naïve
15 phenotype) in both lineage markers of the lymphocyte subset as well as CD8⁻/CD8⁺ T-cell ratio
16 were significantly higher in subjects without CMV compared to their CMV-seropositive
17 counterparts (all $p < 0.001$). No significant difference was found in IL-6 levels with respect to
18 CMV serostatus.

19

20 **Association between IS phenotypes and pre-frailty stratified by CMV serostatus**

21 Considering the significant impact of CMV on the proportion of various T-cell subsets, we
22 investigated the T-cell differentiation phenotypes according to pre-frailty status and separately in

1 CMV-seropositive and CMV-negative subjects (see Table 2). In the CMV-seronegative
2 population, we found a significantly higher proportion of the highly differentiated memory
3 phenotypes and a lower proportion of the naïve cell subset - in the CD8⁻ compartment - in pre-
4 frail subjects compared to robust (all $p < 0.05$, see Table 2). A similar trend was found for the
5 CD8⁺ compartment. Also, pre-frailty was associated with higher levels of IL-6 ($p < 0.001$) in CMV-
6 negative subjects. No significant difference was recorded concerning the percentages of T-cell
7 phenotypes, IL-6 or CD8⁻/CD8⁺ T-cell ratio between the robust and pre-frail groups in the
8 seropositive CMV population.

9

10 **Association between IS phenotypes and CD8⁻/CD8⁺ T-cell ratio category**

11 We further investigated the association between T-cell subsets and the CD8⁻/CD8⁺ T-cell ratio
12 category in the whole cohort (see Figure 1) as well as by CMV serostatus (see Figure 2 and
13 Supplementary Figure 2). 15 (8.7%), 122 (70.5%) and 36 (20.8%) subjects had a ratio < 1 , ratio =
14 1 to 4 and ratio > 4 , respectively. The frequency of cells expressing the highly differentiated
15 memory phenotype was significantly higher in the ratio < 1 group compared to the other groups,
16 both in the CD8⁻ and CD8⁺ sub-populations of T-cells (all $p < 0.01$). Also, the ratio < 1 group was
17 characterized by a significantly higher proportion of the senescence-like phenotypes (all $p < 0.05$)
18 compared to the other groups, in the CD8⁻ pool. On the other hand, the proportion of cells
19 expressing the predominantly naïve phenotype was significantly higher in the ratio > 4 group
20 compared to the other groups ($p < 0.001$). Figure 2 and Supplementary Figure 1 show the
21 distribution of various T-cell sub-populations according to the CD8⁻/CD8⁺ ratio categories and by
22 CMV serostatus. For the CMV-negative group, we found a significantly higher percentage of the
23 highly differentiated memory and senescence-like phenotypes in the ratio < 1 group compared to

1 the other groups (all $p < 0.05$). Contrary wise, the frequency of the naïve phenotypes was
2 significantly higher in the ratio > 4 group compared to the other groups (all $p < 0.01$, see Figure 2).
3 For the CMV-positive group, we found a significantly higher percentage of the highly
4 differentiated memory cells and lower percentage CD8+CD28+CD57+ cell in the ratio < 1 group
5 compared to the other groups (all $p < 0.05$). The percentages of the other differentiation phenotypes
6 did not differ with respect to CD8-/CD8+ T-cell ratio among the CMV-seropositive individuals
7 (see Supplementary Figure 1).

8

9 **Association between IS phenotypes and pre-frailty stratified by CD8-/CD8+ T-cell ratio** 10 **category**

11 The association between IS phenotypes and pre-frailty was not consistent among the various
12 categories of CD8-/CD8+ T-cell ratio (see supplementary Table 3). In the CD8- compartment of
13 T-cells, we found a significantly higher proportion of the highly differentiated memory and
14 senescence-like phenotypes and lower proportion of the naïve phenotype in pre-frail compared to
15 robust subjects in the CD8-/CD8+ T-cell ratio > 4 group (all $p < 0.05$). A similar trend was found
16 in the CD8+ subset. It is noteworthy that more than two thirds (69.4%) of the subjects in the
17 CD8-/CD8+ T-cell ratio > 4 group was CMV-negative.

18

19 **Predictors of prefrailty**

20 Finally, logistic regression was used to determine predictors of the risk of pre-frailty in CMV-
21 seropositive and CMV-negative subjects separately. Since significant correlations were found
22 among the T-cell phenotypes within the CD8- and CD8+ T-cell compartments, we entered just the

1 senescence-prone CD57+ phenotype of each T-cell compartment in the regression analyses. When
2 parameters associated with inflammation, senescence and IRP were entered into the model -
3 corrected for age, sex, BMI, history of heart failure, use of anti-inflammatory drugs, and smoking
4 habits - only changes in IL-6 predicted the risk of pre-frailty, and this was seen only in the CMV-
5 negative group (see Table 3). In a separate analysis, we did include the CD28- subset in the
6 regression analysis. However, the frequency of CD28- T-cells was not predictive of the risk of
7 pre-frailty. Moreover, the inclusion of CD28- phenotype did not influence the predictive ability of
8 IL-6 in identifying pre-frailty (data not shown). We also performed the regression analysis
9 modeling IL-6 as continuous variable and without stratification for CMV - i.e. by including CMV
10 serostatus as a covariate – and the results remained substantially unchanged, portraying only IL-6
11 as a probable predictor of the risk of pre-frailty (data not shown).

12

13 **Discussion**

14 Exploring baseline data from the longitudinal BUTTERFLY study, we investigated the impact of
15 CMV on IS and its relevance to the frailty concept in a very old population. The findings of the
16 present study indicate that pre-frailty does not require the CMV infection as a necessary factor for
17 its development in very old subjects. More so, our study put in doubt the predominant role of the
18 CMV infection on the inflammatory profile of very old persons.

19 In this study on people older than 80 years, IgG for CMV was positive in 92 (53.2 %) subjects.
20 Higher CMV prevalence have often been described in older adults. In a study of 549 community-
21 dwelling persons aged 80 and older in Belgium – where the current study was performed - Mathei
22 et al. (31) reported 74% positive CMV serology. However, in their study, they included patients
23 unlike the apparently healthy population of the present study. Further, a Finnish study (32) showed

1 that CMV seroprevalence was higher in Helsinki compared to a rural area in the southwest of the
2 country (70.7% versus 56.3%, respectively). Therefore, it is conceivable that the overall CMV
3 seropositivity can change over time as a result of changes in health status, age, and socio-economic
4 situation (33).

5 We found no significant relationship between CMV seropositivity and the pro-inflammatory
6 cytokine IL-6. Although positivity for IgG class anti-CMV antibodies cannot distinguish between
7 participants with persistent and those with resolved infection (34), evidence for frequent age-
8 related reactivation and increased viral load of CMV in individuals with positive CMV serology
9 has been reported (35,36). More so, data indicating an age-related prevalence of CMV infection –
10 15% vs 63%, for subjects < 20 years and those > 60 years, respectively – in CMV seropositive
11 individuals has been reported (37). In this light, our observation put in doubt the predominant role
12 of the CMV infection on the inflammatory profile of very old persons.

13 The data portray that pre-frailty does not require the CMV infection as a necessary factor for its
14 development in very old subjects. This finding supports the relatively few published works putting
15 in doubt the predominant role of the CMV infection in frailty states (22,31). In a large population-
16 based study on persons older than 85 years in England no evidence was found to support the
17 association of CMV seropositivity with pre-frailty or frailty (22). More strikingly, in another
18 population-based study in the oldest old in Belgium a negative association between positive CMV
19 serology and frailty states was reported (31). These findings, regarding CMV-serostatus and frailty
20 states in very old subjects, might reflect a survival effect as was proposed by Adriaensen et al. (38).
21 From this perspective, individuals susceptible to the long-term deleterious effects of CMV
22 exposure are more likely to die at a younger age (20) and thus be under-represented in a cohort of
23 very old people like ours. Accordingly, Derhovanessian and colleagues (39) found that CMV-

1 infected offspring from long-lived families had significantly lower levels of pro-inflammatory
2 parameters than did their age-matched CMV-infected controls, hypothetically reflecting a better
3 immunological control of the virus - with less contributing factors to frailty status - in the siblings
4 of long-lived families (39). A better immunological control would imply less reactivation of the
5 virus and perhaps up-regulation of the anti-inflammatory pathway (40). This reasoning might
6 explain the lack of difference in IL-6 levels with respect to CMV serostatus and the absence of a
7 relationship between IL-6 and pre-frailty among the CMV-seropositive subjects, in the perspective
8 that the frailty status might depend on the balance between pro- and anti-inflammatory cytokines
9 (41).

10 Pre-frailty was clearly associated with increased IL-6 independent of age, sex, BMI, history of
11 heart failure, use of anti-inflammatory drugs, and smoking habits, in CMV-negative subjects,
12 indicating that inflammatory regulators other than CMV are involved. This observation is
13 consistent with the consensus that pro-inflammatory parameters, particularly IL-6, may inhibit the
14 synthesis of IGF-1 and induce - through its catabolic effects on muscles - skeletal muscle mass loss
15 (42,43). In this light, subjects may become less active, and express physical characteristics of frailty
16 including low muscle strength, exhaustion, reduced physical activity and unintentional weight loss.
17 In agreement with the present study, many reports in both cross-sectional and longitudinal studies
18 have consistently found elevated levels of various inflammatory markers among pre-frail as well
19 as frail individuals (21,22,44). Considering the well-established burden of CLIP in the elderly, it is
20 reasonable to think that CLIP would maintain and reinforce the frailty syndrome in older subjects.
21 Notwithstanding, the absence of association between IL-6 and pre-frailty in CMV seropositive
22 individuals deserves further investigation.

1 In our cohort of older adults, 15 (8.7%) subjects had a CD8⁻/CD8⁺ ratio <1, which was associated
2 with CMV-seropositivity. Large increases in the proportion of memory and senescence-like
3 phenotypes and decrease in naïve cell phenotypes were significantly associated with CMV
4 seropositivity. Similar associations were found between T-cell subtypes and CD8⁻/CD8⁺ ratio <1,
5 albeit in CMV negative subjects. Loss of CD28 marker and increase in the CD57 marker on T-
6 cells of very old subjects with IRP was reported for Swedish OCTO and NONA cohorts (45).
7 However, prudence should be exercised when drawing conclusions from the present results, since
8 our CD8⁻/CD8⁺ parameter is a surrogate, which might differ from the originally used CD4⁺/CD8⁺
9 ratio.

10 Pre-frail subjects were more prone to have a history of heart failure compared to robust. This
11 finding corroborates results from other authors indicating an increased risk of heart failure
12 diagnosis in community-dwelling individuals with moderate and severe frailty (46,47). Although
13 it is not clear how heart failure and pre-frailty may be linked, both phenomena are associated with
14 inflammation (48). There is emerging evidence of NLRP3 activation in heart failure patients, with
15 resulting inflammation (49). On the other hand, aged mice deficient in the NLRP3 inflammasome
16 exhibit enhanced walk distance and running time as compared to their wild-type controls,
17 suggesting that NLRP3 may enhance inflammation and thereby lead to pre-frailty (50). Whether
18 the NLRP3 inflammasome may represent a common pathway by which pre-frailty and heart failure
19 interact requires future investigation.

20

21 **Limitations**

1 The findings of the present study should be interpreted within its limitations. First, this was a cross-
2 sectional study, which precludes causal relationships. Therefore, caution should be exercised when
3 making inferences about temporality. Second, since our intent was to focus on markers of IS with
4 regards to pre-frailty, we did not investigate psycho-social factors, which are modifying factors for
5 the development of pre-frailty. The authors also acknowledge the limitation that the selection of
6 apparently healthy individuals might have masked other pre-frailty related patterns. More so, given
7 the small sample size in some of the subsets of the population, it is possible that our study was not
8 sufficiently powered to detect small differences between groups. Despite some limitations, this
9 study adds a highly needed element in the context of frailty and associated characteristics. Our
10 attempt to simultaneously investigate CMV, inflammatory, and IS markers in the same cohort
11 offers insights into the impact of CMV on IS phenotypes and their relevance in the setting of pre-
12 frailty, thus extending current knowledge on the frailty concept. Another strong point is that this
13 study was performed in very old subjects with a distinctly different physiologic profile compared
14 to the relatively younger adult participants in most literature reports. Our finding of no association
15 between pre-frailty and CMV-seropositivity, makes our study complementary to previous studies
16 in younger populations. Moreover, the observation that subjects' CMV serostatus may define the
17 association between IS phenotypes and pre-frailty could act as a guide for future research
18 concerning IS phenotypes and their association with frailty status.

19

20 **Conclusions**

21 The findings of the present study indicate that the presence of pre-frailty is independent of CMV
22 infection in very old subjects. Further, higher concentrations of the inflammatory cytokine IL-6
23 was predictive of pre-frailty in CMV-negative but not in CMV-seropositive subjects. Whether IL-

1 6 might facilitate the identification of people at risk of developing pre-frailty - at least for subjects
2 without CMV - deserves further study.

3 **Funding**

4 This study was funded by a grant from the People's Committee of Hochiminh City, Vietnam (grant
5 number 35-QĐ/BTCTU) to [Hung Cao Dinh] and an "Interdisciplinary Research Program" grant
6 from the research council of the Vrije Universiteit Brussel (grant number IRP3).

7

8 **Conflict of interest**

9 All authors certify that they comply with the ethical guidelines for publishing in the Journal of
10 Gerontology: Biological Sciences. None of the authors have any conflict of interest with any entity
11 with regard to this study. The authors have no other conflict of interest to declare.

12

13 **Acknowledgment**

14 The authors would like to thank all members of the Gerontopole Brussels Study group, comprising
15 Ivan Bautmans, Dominique Verté, Ingo Beyer, Mirko Petrovic, Liesbeth De Donder, Tinie Kardol,
16 Gina Rossi, Peter Clarys, Aldo Scafoglieri, Eric Cattrysse, Paul de Hert, and Bart Jansen for their
17 inputs in the conception of the project.

18

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43

1 **Captions for Tables**

2 **Table 1.** IS phenotypes according to pre-frailty and CMV serostatus

3 **Table 2.** Association between IS phenotypes and pre-frailty stratified by CMV serostatus

4 **Table 3.** Odds ratio (95% CI) of cross-sectional logistic regression analyses of the association
5 between inflammatory and senescence parameters and prevalent pre-frailty.

6 **Supplementary Table 1.** Characteristics of the cohort according to frailty and CMV serostatus

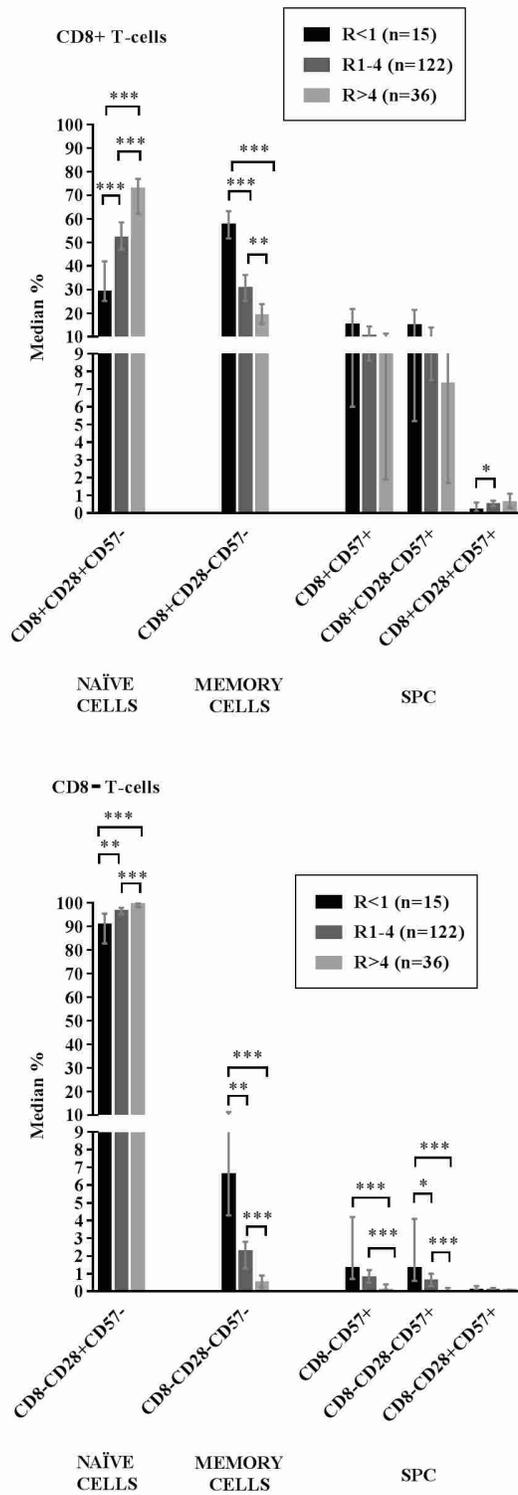
7 **Supplementary Table 2.** Overview of participants' comorbidities according to frailty status

8 **Supplementary Table 3.** Association between IS phenotypes and pre-frailty stratified by
9 CD8⁻/CD8⁺ category

10

1 Captions for Figures

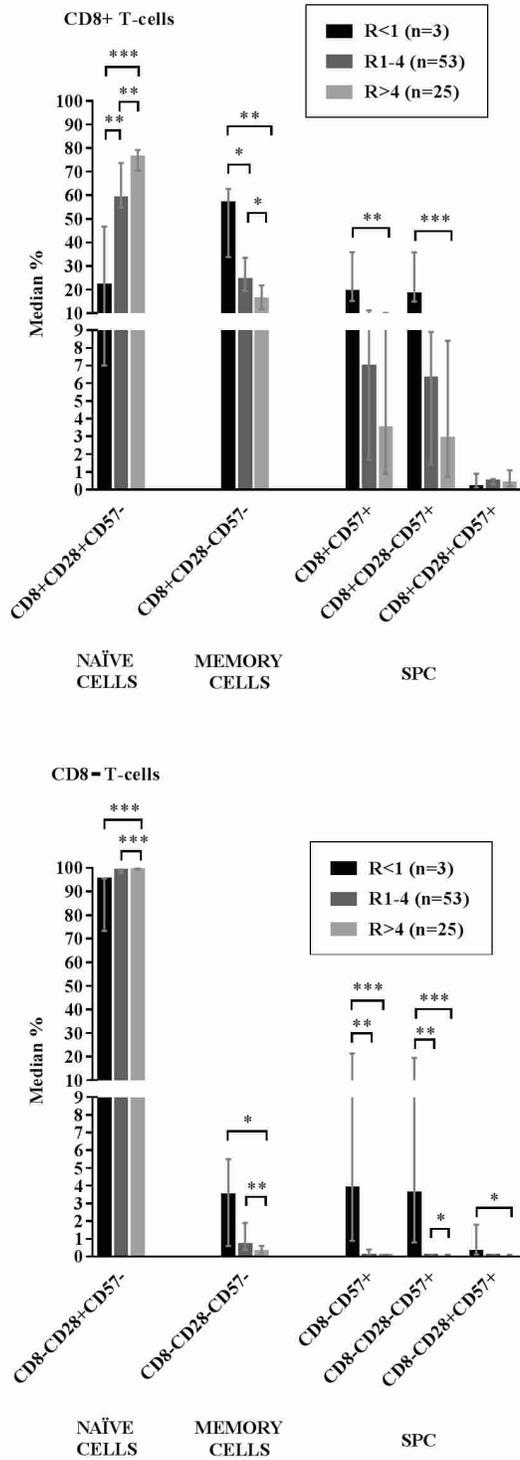
Figure 1. Association between T-cell differentiation markers and CD8⁻/CD8⁺ T-cell ratio in the cohort.



Note: data represent median percentage of cells within the CD3⁺CD8⁺ or CD3⁺CD8⁻ T-cell subsets. SPC, senescence-prone cells. The error bars represent 95% confidence intervals. *p<0.05, **p<0.01, ***p<0.001.

1 **Figure 1.** Association between T-cell differentiation markers and CD8⁻/CD8⁺ T-cell ratio in the
2 cohort.
3 Data represent median percentage of cells within the CD3⁺CD8⁺ and CD3⁺CD8⁻ T-cell subsets.
4 SPC, senescence-prone cells. The error bars represent 95% confidence intervals. *p < 0.05, **p <
5 0.01, ***p < 0.001
6

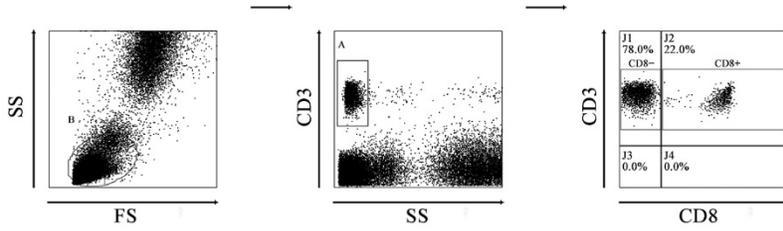
Figure 2. Association between T-cell differentiation markers and CD8⁻ /CD8⁺ T-cell ratio in CMV-seronegative subjects.



Note: data represent median percentage of cells within the CD3⁺CD8⁺ or CD3⁺CD8⁻ T-cell subsets. SPC, senescence-prone cells. The error bars represent 95% confidence intervals. *p<0.05, **p<0.01, ***p<0.001.

1 **Figure 2.** Association between T-cell differentiation markers and CD8⁻/CD8⁺ T-cell ratio in
2 CMV-seronegative subjects.
3 Data represent median percentage of cells within the CD3⁺CD8⁺ and CD3⁺CD8⁻ T-cell subsets.
4 SPC, senescence-prone cells. The error bars represent 95% confidence intervals. *p < 0.05, **p <
5 0.01, ***p < 0.001
6

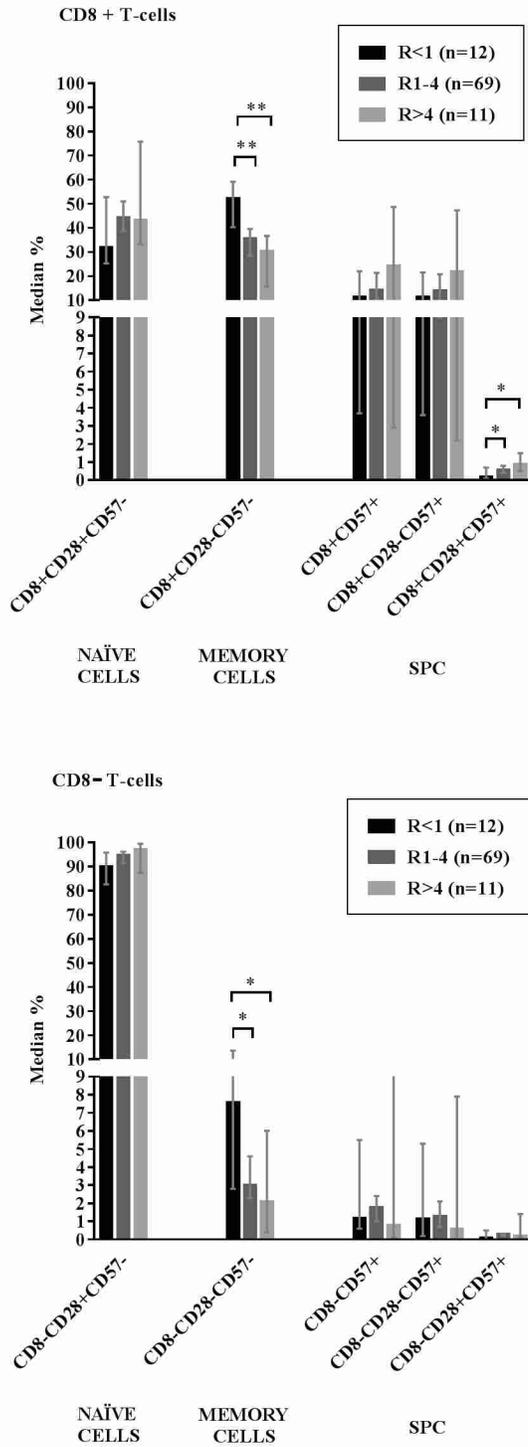
Supplementary figure 1. Representative dot plots for the delineation of the different sub-populations by flow cytometry.



By combining side scatter (SS) versus forward scatter (FS) and anti-CD3 conjugated fluorochrome fluorescence versus SS plots, CD3+ cells were identified as cells that were both in gates A and B. CD3+ cells were further sub-divided based on the expression or non-expression of CD8.

- 1
- 2 **Supplementary Figure 1.** Representative dot plots for the delineation of the different sub-
- 3 populations by flow cytometry.
- 4 By combining side scatter (SS) versus forward scatter (FS) and anti-CD3 conjugated fluorochrome
- 5 fluorescence versus SS plots, CD3+ cells were identified as cells that were both in gates A and B.
- 6 CD3+ cells were further sub-divided based on the expression or non-expression of CD8.
- 7

Supplementary Figure 2. Association between T-cell differentiation markers and CD8⁻/CD8⁺ T-cell ratio in CMV-seropositive subjects.



Note: data represent median percentage of cells within the CD3⁺CD8⁺ or CD3⁺CD8⁻ T-cell subsets. SPC, senescence-prone cells. The error bars represent 95% confidence intervals. *p < 0.05, **p < 0.01.

1 **Supplementary Figure 2**

2 Association between T-cell differentiation markers and CD8⁻/CD8⁺ T-cell ratio in CMV-
3 seropositive subjects.

4 Data represent median percentage of cells within the CD3⁺CD8⁺ and CD3⁺CD8⁻ T-cell subsets.

5 SPC, senescence-prone cells. The error bars represent 95% confidence intervals. *p < 0.05, **p <

6 0.01.

Table 1. IS phenotypes according to pre-frailty and CMV serostatus

Parameter	Robust (n=82)	Pre-frail (n=91)	CMV+ (n=92)	CMV- (n=81)
T-cell subset				
CD8+ T-cells				
CD8+CD28+CD57- (naïve)	51.05 (40.88)	55.50 (34.80)	42.80 (28.33)	68.30 (29.25) ^{b**}
CD8+CD28-CD57- (memory)	29.65 (31.12)	31.40 (24.80)	36.60 (27.03)	22.50 (21.65) ^{b**}
CD8+CD28-CD57+ (SPC)	9.25 (20.68)	8.40 (17.80)	14.70 (19.00)	6.30 (11.90) ^{b**}
CD8+CD28+CD57+ (SPC)	0.50 (0.70)	0.50 (1.10)	0.60 (0.95)	0.50 (0.90)
CD8- T-cells				
CD8-CD28+CD57- (naïve)	96.95 (8.9)	97.40 (6.10)	94.80 (9.18)	99.20 (2.60) ^{b**}
CD8-CD28-CD57- (memory)	1.85 (4.73)	2.00 (4.20)	3.05 (6.00)	0.50 (1.80) ^{b**}
CD8-CD28-CD57+ (SPC)	0.55 (2.73)	0.30 (1.80)	1.15 (2.68)	0.00 (0.30) ^{b**}
CD8-CD28+CD57+ (SPC)	0.10 (0.30)	0.10 (0.30)	0.20 (0.30)	0.10 (0.10) ^{b**}
Inflammatory marker				
IL-6 (pg/mL)	1.50 (1.17)	2.06(2.30) ^{a**}	1.78 (1.90)	1.65 (1.79)
IRP marker				
CD8-/CD8+ ratio (n (%))				
<1	8 (9.76)	7 (7.69)	12 (13.04)	3 (3.70) ^{b**}
1-4	60 (73.17)	62 (68.13)	69 (75.00)	53 (65.43)
>4	14 (17.07)	22 (24.18)	11 (11.96)	25 (30.87)

Note: The values denote median (Interquartile range), unless otherwise stated; IS= immunosenescence, CMV cytomegalovirus; IRP = immune risk profile; SPC= senescence-prone cells; ^a difference between robust and pre-frailty; ^b difference between CMV+ and CMV-. *p < 0.05; **p < 0.01.

1

2

Table 2. Association between IS phenotypes and pre-frailty stratified by CMV serostatus.

Parameter	CMV+ (n=92)		CMV- (n=81)	
	Robust (n=47)	Pre-frail (n=45)	Robust (n=35)	Pre-frail (n=46)
T-cell subset				
CD8+ T-cells				
CD8+CD28+CD57- (naïve)	41.70 (28.00)	49.80 (33.30)	72.60 (34.00)	60.00 (29.13)*
CD8+CD28-CD57- (memory)	37.40 (21.10)	33.70 (22.75)	19.50 (25.40)	23.90 (25.90)
CD8+CD28-CD57+ (SPC)	16.40 (16.10)	12.80 (19.35)	1.70 (10.60)	7.50 (12.58)
CD8+CD28+CD57+ (SPC)	0.60 (0.60)	0.60 (1.20)	0.40 (0.70)	0.50 (1.13)
CD8- T-cells				
CD8-CD28+CD57- (naïve)	92.50 (10.10)	95.40 (8.80)	99.50 (0.70)	99.00 (4.00)*
CD8-CD28-CD57- (memory)	4.00 (7.00)	2.80 (5.35)	0.40 (0.50)	0.90 (2.83)**
CD8-CD28-CD57+ (SPC)	1.30 (4.20)	0.70 (2.25)	0.00 (0.20)	0.10 (0.45)
CD8-CD28+CD57+ (SPC)	0.30 (0.30)	0.20 (0.30)	0.10 (0.10)	0.10 (0.10)
Inflammatory marker				
IL-6 (pg/mL)	1.56 (1.82)	2.06 (2.25)	1.28 (1.07)	2.20 (2.33)**
IRP marker				
CD8-/CD8+ ratio (n (%))				
<1	6 (12.77)	6 (13.33)	2 (5.71)	1 (2.17)
1-4	38 (80.85)	31 (68.89)	22 (62.86)	31 (67.39)
>4	3 (6.38)	8 (17.78)	11 (31.43)	14 (30.44)

Note: The values denote median (Interquartile range), unless otherwise stated; IS= immunosenescence; CMV = cytomegalovirus; IRP = immune risk profile; SPC = senescence-prone cells; Phenotype frequencies were expressed as percentages within the CD3+CD8+ or CD3+CD8- T-cells; *p < 0.05, **p < 0.01: difference between robust and pre-frail.

Table 3. Odds ratio (95% CI) of cross-sectional logistic regression analyses of the association between inflammatory and senescence parameters and prevalent pre-frailty

Parameter	Unadjusted Model				Adjusted Model			
	CMV+ (n=92)		CMV- (n=81)		CMV+ (n=92)		CMV- (n=81)	
	OR (95% CI)	p value	OR (95% CI)	p value	OR (95% CI)	p value	OR (95% CI)	p value
IL-6								
Low	reference		reference		reference		reference	
Intermediate	2.12 (0.66-6.76)	0.206	5.47 (1.40 -21.39)	0.015	2.99 (0.84-10.68)	0.091	5.98 (1.33-26.86)	0.020
High	2.04 (0.64-6.54)	0.231	17.29 (3.81-78.39)	<0.001	2.08 (0.59-7.29)	0.252	18.38 (3.44-98.21)	0.001
CD8-/CD8+ ratio	1.23 (0.91-1.67)	0.178	1.10 (0.81-1.50)	0.548	1.27 (0.91-1.78)	0.166	1.12 (0.79 -1.58)	0.522
CD8+CD57+	1.00 (0.96 -1.04)	0.923	1.02 (0.96-1.09)	0.544	1.00 (0.96-1.05)	0.915	1.03 (0.96-1.10)	0.413
CD8-CD57+	0.93 (0.83-1.05)	0.259	0.86 (0.52-1.44)	0.569	0.93 (0.81-1.06)	0.272	0.80 (0.44-1.45)	0.465
age	1.18 (1.01-1.39)	0.043	1.30 (1.05-1.62)	0.018	1.18 (1.00-1.40)	0.050	1.33 (1.05-1.68)	0.017
sex	0.48 (0.19 -1.21)	0.119	0.44 (0.14 -1.38)	0.159	0.35 (0.11-1.10)	0.073	0.27 (0.06-1.11)	0.069
body mass index	-		-		1.00 (0.86- 1.17)	0.998	1.19 (0.98- 1.43)	0.075
smoking	-		-		0.57 (0.17- 1.93)	0.370	0.78 (0.21- 2.96)	0.714
heart failure	-		-		7.33 (1.12- 47.99)	0.038	11.63 (0.40- 334.53)	0.152
use of anti-inflammatory drugs.	-		-		0.53 (0.20- 1.43)	0.208	0.39 (0.11- 1.34)	0.134

Note: Unless otherwise specified, data are presented as odds ratio (OR) and 95% confidence interval (CI); CMV = cytomegalovirus; IL-6 = interleukin 6; Low (< 1.4 pg/mL), Intermediate (1.4-2.5 pg/mL), High (> 2.5 pg/mL). Adjusted model: body mass index, smoking, heart failure, and use of anti-inflammatory drugs.