

Brief Report

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Brief Report

Hemodiafiltration May Be Associated with Senescence-Related Phenotypic Alterations of Lymphocytes Which May Predict Mortality in Dialysis Patients

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Abstract: Introduction: Senescence-resembling alterations on lymphocytes of dialysis patients have been widely described. However, pathophysiology behind these phenomena has not been clarified. In this study, we examine the impact of dialysis prescription on T and B lymphocytes, in dialysis patients. Methods: T and B cell subsets were determined with flow cytometry in 36 hemodialysis and 26 hemodiafiltration patients, according to the expression of CD45RA, CCR7, CD31, CD28, CD57 and PD1 for T cells, and IgD and CD27 for B cells. Immune phenotype was associated with dialysis modality, hemofiltration volume, and mortality. Results: Compared to hemodialysis, hemodiafiltration patients had significantly decreased percentage of CD4+CD28-CD57- T cells [3.8(2.4-5.3) vs 2.1(1.3-3.3)%, respectively, $P=0.002$] and exhausted CD4+ T cells [14.1(8.9-19.4) vs 8.5(6.8-11.7)% , respectively, $P=0.005$]. Additionally, hemofiltration volume was negatively correlated to CD8+ EMRA T cells ($r=-0.46$, $P=0.03$). Finally, increased exhausted CD4+ T cells percentage was associated with increased all-cause mortality in dialysis patients, independently of age. Conclusion: Hemodiafiltration, especially with high hemofiltration volume, may have beneficial effects on senescence-related immune phenotype. Immune phenotype may also be a predicting factor for mortality in dialysis patients.

Keywords: ESRD; dialysis; hemodiafiltration; Immune senescence; immune exhaustion

1. Introduction

Immune alterations in End-Stage Kidney Disease (ESKD) have been widely described and involve both innate and acquired immunity [1]. These disturbances are closely related to persistent and chronic activation of inflammatory response, and are implicated in several clinical complications of ESKD, such as susceptibility to serious infections, inadequate response to vaccination, increased malignancy incidence, frailty, and nutritional disturbances, and above all increased cardiovascular risk and mortality [2–5].

In the setting of ESKD, acquired immunity alterations have been more extensively studied. Several studies show that lymphocytes from patients with ESKD exhibit phenotypic and functional characteristics of older individuals, thus resembling prematurely senescent cells. These include

lymphopenia with inverted CD4+/CD8+ ratio, elimination of naïve and low differentiation subpopulations, increased cytotoxicity, decrease of regulatory T cells and imbalance among Th responses [6–8].

Several pathophysiological mechanisms have been proposed to elicit these alterations. Premature thymic involution and cytokine dysregulation, particularly decrease of IL-7 and increase of TNF- α have been associated with naïve T cells elimination and senescent CD28- T cells accumulation [9,10]. Moreover, dialysis related factors including dialyzers biocompatibility, water contamination, frequent catheter infection and volume and sodium overload appear to play a major role [11,12]. Most importantly, increased levels of uremic toxins, mainly produced by impaired intestinal flora are poorly removed by dialysis and contribute to inflammatory activation [13].

Conventional hemodialysis (HD) is the most widespread method of kidney replacement therapy and relies on solute clearance with diffusion. However, its capacity to remove larger molecules is limited due to the relatively low speed of diffusion of these molecules through the dialyzer membrane. Online hemodiafiltration (HDF) provides an additional solute clearance with convection, which is achieved by the removal of large fluid volumes through a membrane of high hydraulic permeability (“high flux” membrane). The fluid removed is substituted with sterile dialysate of optimal electrolyte composition, infused directly into the bloodstream of the patient, hence, ultra-pure water is needed [14]. Online HDF improves removal of middle molecular weight solutes. Most importantly, appears to have a beneficial effect in all-cause and cardiovascular mortality, in oxidative stress and hospitalization rates and, also, improve the patients’ quality of life [15,16]. Moreover, a reduction of inflammatory markers has been associated to HDF [17].

Despite the wide range of clinical associations that immune alterations have on ESKD patients, there are no data available to the impact of dialysis prescription on immune phenotype. The present study aims to investigate for potential discrepancies in lymphocytic phenotypes, between ESKD patients, on two main dialysis modalities, HD and HDF, examine the role of hemofiltration volume (HFV), and explore the impact of immune phenotype to mortality.

2. Results

2.1. Patients’ Characteristics

Patients’ demographics are shown in table 1. Patients were equally distributed within the three age groups. Sex distribution did not differ among the three age groups (P=0.26), however, patients in the 20-40 year age group had a significantly lower body weight and BMI (P=0.03) . A 30% of the patients had a history immunosuppressive treatment, either due to their primary cause of ESKD, or after kidney transplantation. However, the vast majority of the patients had ceased any immunosuppressive treatment for a considerable time period before study initiation (Table 1). Dialysis prescription differed among the groups, as more patients were on conventional HD in the older group 61-80 years (P=0.02), without significant difference in the mean duration of each dialysis session between the groups (P=0.14). Vintage of dialysis differed significantly among the groups, as patients in the middle age group 41-60 years, had a mean vintage of 101 \pm 81months, P=0.01 in comparison to the young patients group (45 \pm 48 months) or the older patients group (67 \pm 44 months).

Table 1. Patients’ baseline characteristics.

	20-40	41-60	61-80	Total
N	21	21	20	62
Mean age \pm sd, years	30.5 \pm 6.3	52.2 \pm 5.6	71.2 \pm 6.4	51.3 \pm 17.7
Sex, male/female	14/7	14/7	9/11	37/25
Mean weight \pm sd, kg	61 \pm 11.5	81.3 \pm 19.2	71.1 \pm 12.5	71.3 \pm 17
Mean BMI \pm sd, kg/m ²	21 \pm 3.2	25.9 \pm 3.9	25.5 \pm 3.6	24.5 \pm 4.1
<i>Cause of ESRD</i>				
Primary glomerulonephritis (%)	3(14.3)	10(47.6)	4(20)	17(27.4)

Hypertension	1(4.8)	2(9.5)	0(0)	3(4.8)
Obstructive nephropathy	5(23.8)	1(4.8)	2(10)	8(12.9)
Polycystic kidney disease	1(4.8)	2(9.5)	4(20)	7(11.2)
Other	6(28.6)	1(4.8)	1(5)	7(11.2)
Unknown	5(23.8)	5(23.8)	9(45)	19(30.6)
<i>Dialysis characteristics</i>				
Classic hemodialysis	11	9	16	36
Online hemodiafiltration	10	12	4	26
Mean vintage \pm sd, months	45 \pm 48	101 \pm 81	67 \pm 44	72 \pm 64
Kt/V \pm sd	1.29 \pm 0.05	1.27 \pm 0.06	1.23 \pm 0.04	1.27 \pm 0.07
Mean dialysis duration \pm sd, Min	240 \pm 14	246 \pm 11	234 \pm 13	240 \pm 14
<i>Comorbidities</i>				
Hypertension	11	11	10	32
Cardiovascular disease	2	11	8	21
Secondary hyperparathyroidism	11	12	11	34
Anemia	18	15	18	51
History of kidney transplantation (%)	4(19)	6(28.6)	0(0)	10(16.1)
Prior immunosuppression (%)	5(23.8)	11(52.3)	3(15)	19(30)
Median time since immunosuppression cessation(IQR), months	75(43-148)	45(36-63)	48(-)	48(40-96)
<i>Laboratory values, median (IQR)</i>				
Hemoglobin, g/dl	11.6(10.8-11.9)	11.2(10.4-11.7)	10.5(10-11.1)	11(10.5-11.6)
Urea, mg/dl	130(96-149)	145(117-154)	121(112-140)	127(110-149)
Creatinine, mg/dl	9.7(7.8-11.8)	9.1(7.1-10.9)	8.4(5.8-9.5)	9.1(7.1-10.8)
Calcium, mg/dl	9.1(8.8-9.2)	8.9(8.8-9.5)	9.2(8.7-9.6)	9.1(8.8-9.3)
Phosphorus, mg/dl	4.6(4-5.5)	4.6(3.3-5)	4(3.8-4.5)	4.4(3.8-5.1)
Parathormone, pg/ml	162(89-391)	268(145-376)	197(98-244)	206(106-353)
Albumin, g/dl	4.3(3.9-4.4)	4.2(3.9-4.3)	3.9(3.8-4.1)	4.1(3.9-4.3)
C reactive protein (mg/l)	1.7(1.2-3.7)	2.5(1.4-7.4)	3.7(1.8-7.4)	2.4(1.4-4.7)

2.2. Comparison of Immune Phenotype between HD and HDF Patients

Online HDF was applied in 26/62(42%) patients. Age of patients did not differ significantly between the two groups [57(38-71) vs 49(31-58)years for HD and HDF respectively, $P=0.08$]. Patients on HDF had significantly higher proportions of low differentiation CD4+CD28+CD57- and decreased proportions of highly differentiated CD4+CD28- lymphocytes. However, within the compartment of CD28- T cells, the difference between the two patient groups was restrained only in CD28-CD57-subset which was increased in HD patients, while terminally differentiated CD28-CD57+ T cells did not differ between the two groups (Figure 1). In the CD8+ compartment, proportions of CD28+CD57- T cells and total CD28- T cells did not differ between the two subgroups, whereas CD8+CD28-CD57- T cells were significantly reduced in the HDF group. Absolute numbers and proportions of T cells according to expression of CD28 and CD57 are shown in Supplemental Table 1.

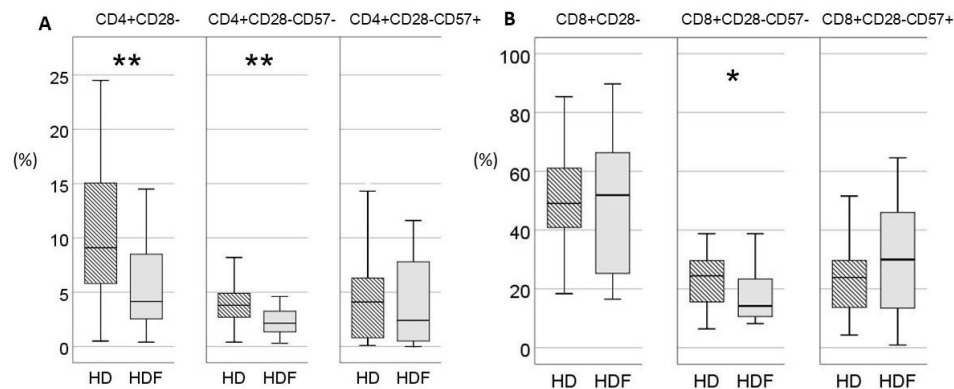


Figure 1. (A) Differences in expression of CD28 and CD57 surface markers on CD4+ T cells between patients on conventional hemodialysis (HD) and online hemodiafiltration (HDF). (B) Differences in expression of CD28 and CD57 surface markers on CD8+ T cells between patients on HD and HDF. T cell subsets are given as percentages. * $p < 0.05$, ** $p < 0.01$.

To adjust for age, the two patient cohorts were divided into three age groups and ANCOVA was performed. Patients on HDF had increased proportions of CD4+CD28+CD57- ($P=0.03$, for modality), and decreased proportions of CD4+CD28- ($P=0.02$, for modality) and CD4+CD28-CD57- ($P=0.02$, for modality), independently of age ($P=0.37$, $P=0.41$ and $P=0.88$ for the three subsets respectively), with no significant interaction between age group and modality ($P=0.9$, $P=0.88$ and $P=0.9$, respectively). Proportions of CD4+CD28-CD57+ T cells remained non-significant for dialysis modality and age group (Figure 2). In contrast, CD8+ subsets were affected by age in ANCOVA ($P=0.003$, $P=0.006$ and $P=0.01$ for CD28+CD57-, CD28- and CD28-CD57- respectively), while neither dialysis modality nor age were significant in ANCOVA for CD8+CD28-CD57- T cells proportion ($P=0.06$, $P=0.61$ and $P=0.358$ for modality, age and interaction, respectively)(Figure 2).

Moreover, there was a significant difference in percentage of exhausted CD4+PD1+ cells [14.1(8.9-19.4) vs 8.5(6.8-11.7)% for HD and HDF respectively, $P=0.005$], whereas the percentage of CD8+PD1+ cells did not differ significantly between the two groups [33.8(12.7-52.2) vs 17.4(8.5-41)% for HD and HDF respectively, $P=0.06$]. The impact of age in ANCOVA was not significant neither in the CD4+PD1+ ($P=0.42$) nor in CD8+PD1+ T cells percentages ($P=0.82$). The significance of dialysis modality was retained after adjusting for age only for CD4+PD1+ T cell percentage (Figure 2).

Counts and percentages of naïve and memory T and B cells did not differ between the two groups. Moreover, levels of urea, creatinine, and phosphorous levels did not correlate with any of the studied T and B cells subsets (data not shown).

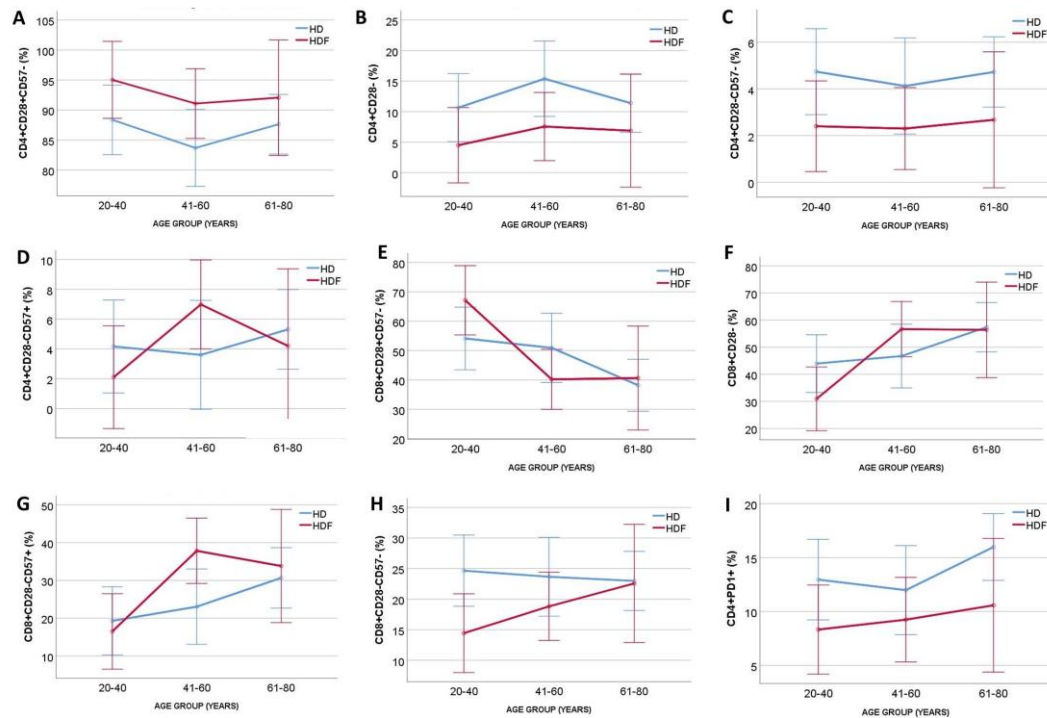


Figure 2. ANCOVA for modality and age group for different T cell subsets percentages (A) Low differentiated CD4+CD28+CD57- T cells (B) Highly differentiated CD4+CD28- T cells (C) Highly differentiated CD4+CD28-CD57- T cells (D) Senescent CD4+CD28-CD57+ T cells (E) Low differentiated CD8+CD28+CD57- T cells (F) Highly differentiated CD8+CD28- T cells (G) Highly differentiated CD8+CD28-CD57+ T cells (H) Senescent CD8+CD28-CD57+ T cells (I) Exhausted CD4+ T cells.

2.3. Effect of HFV

Correlation of T cells subsets with hemofiltration volume in patients on HDF revealed a significant positive association between HFV and central memory CD4+ ($r=0.46$, $P=0.03$) and CD8+ cell counts ($r=0.51$, $P=0.01$) and a significant negative correlation with CD8+ EMRA T cells count ($r=-0.46$, $P=0.03$). Moreover, a positive correlation between HFV with total B cell count was observed ($r=0.46$, $P=0.03$). Out of the B cell subpopulations HFV mainly affected naïve ($r=0.53$, $P=0.008$) and switched memory B cells ($r=0.5$, $P=0.02$), whereas its association with IgD-CD27- B cells ($r=0.41$, $P=0.05$) was marginal. (Figure 3). In univariate regression model the effect of HFV remained significant for all the above subsets except for IgD-CD27- B cells (Table 2).

Multivariate analysis performed for each lymphocyte population, including age and HFV, showed that HFV was an independent factor for CD8+ central memory and EMRA T cell population, whereas age was the main factor affecting CM CD4+ , total B and naïve B cell count (Table 2).

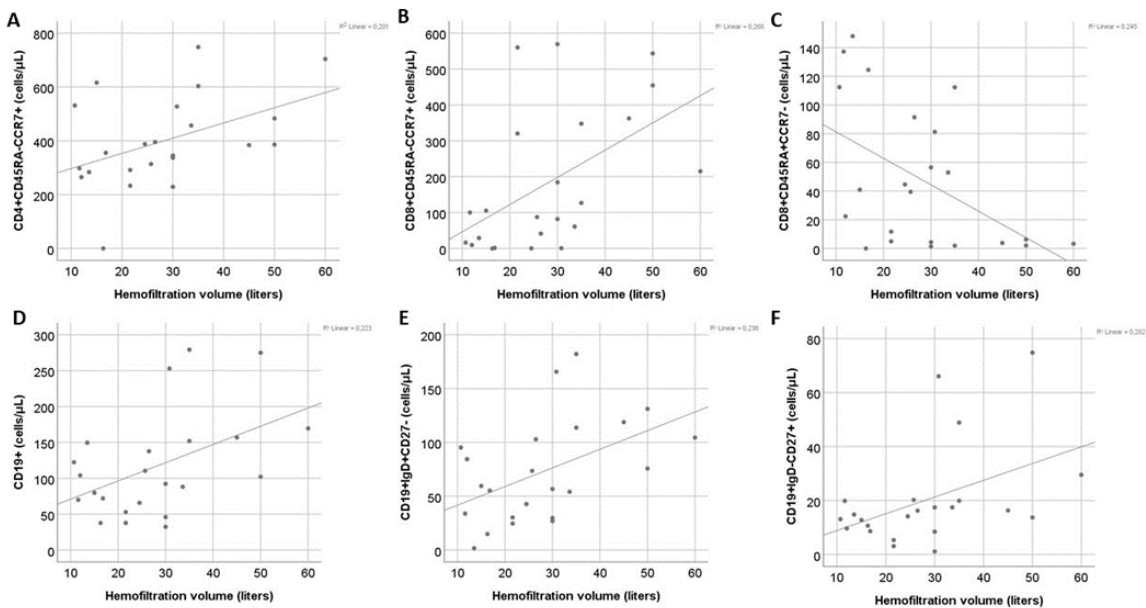


Figure 3. Linear correlation of hemofiltration volume to different T cell and B cell subsets. A. Central memory CD4+ T cells B. Central memory CD8+ T cells. C. Effector memory CD8+ T cells re-expressing CD45RA (EMRA). D. Total B cells. E. Naïve B cells. F. Switched memory B cells.

Table 2. Univariate linear regression of T and B cell subsets for hemofiltration volume and multivariate regression of T and B cell subsets for hemofiltration volume and age.

Lymphocyte subsets (cells/μL)	Univariate Linear Regression				Multivariate Linear Regression					
	Hemofiltration volume				Hemofiltration volume			Age (years)		
	B	R²	95% CI	P	PC	95% CI	P	PC	95% CI	P
CD4+CD45RA-CCR7+	0.45	0.2	0.5, 10.8	0.03	0.33	-1.1, 9.0	0.12	-0.43	-9.8, -0.2	0.04
CD8+CD45RA-CCR7+	0.52	0.26	1.8, 13.1	0.01	0.48	1.1, 13.3	0.02	-0.05	-6, 5	0.81
CD8+CD45RA+CCR7-	-0.49	0.24	-3.2, -0.3	0.02	-0.56	-3.6, -0.6	0.006	-0.32	-2.4, 0.3	0.13
CD19+	0.47	0.22	0.3, 4.5	0.02	0.36	-0.3, 3.8	0.09	-0.42	-3.9, -0.03	0.04
CD19+IgD+CD27-	0.48	0.23	0.3, 3.2	0.02	0.38	-0.2, 2.6	0.08	0.48	-2.8, -0.2	0.02
CD19+IgD-CD27+	0.45	0.20	-0.05, 1.1	0.03	0.39	0.05, 1.1	0.07	-0.16	-0.7, 0.35	0.47
CD19+IgD-CD27-	0.12	0.01	-0.3, 1.3	0.30	-0.07	-1.1, 0.8	0.75	-0.11	-1.2, 0.6	0.61

2.4. Effect of vintage

Dialysis Vintage was negatively correlated with the total count of T cells and CD4+ T cells ($r=-0.35$, $P=0.005$ and $r=-0.46$, $P<0.001$, respectively), while total CD8+ T cells were not affected. The effect of vintage was evident only in low differentiation CD4+ T cell subsets ($r=-0.51$, $P<0.001$, $r=-0.38$, $P=0.002$ and $r=-0.31$, $P=0.01$ for naïve, RTEs and CM CD4+ T cells respectively). Moreover, there was a significant negative correlation with CM CD8+ T cells ($r=-0.29$, $P=0.02$). Finally, total B cell and memory B cell count were also negatively correlated with dialysis vintage ($r=-0.26$, $P=0.04$ and $r=-0.25$, $P=0.04$) (Figure 4). However, in multivariate analysis for vintage and age the effect of vintage remained significant only for total CD4+ and naïve CD4+ count, whereas age was the main predictive factor for CD4+ RTEs and B cell subsets (Table 3). Total lymphocytes and CM cells, in both CD4+ and CD8+ compartments, were not affected neither by vintage nor age in the multivariate model.

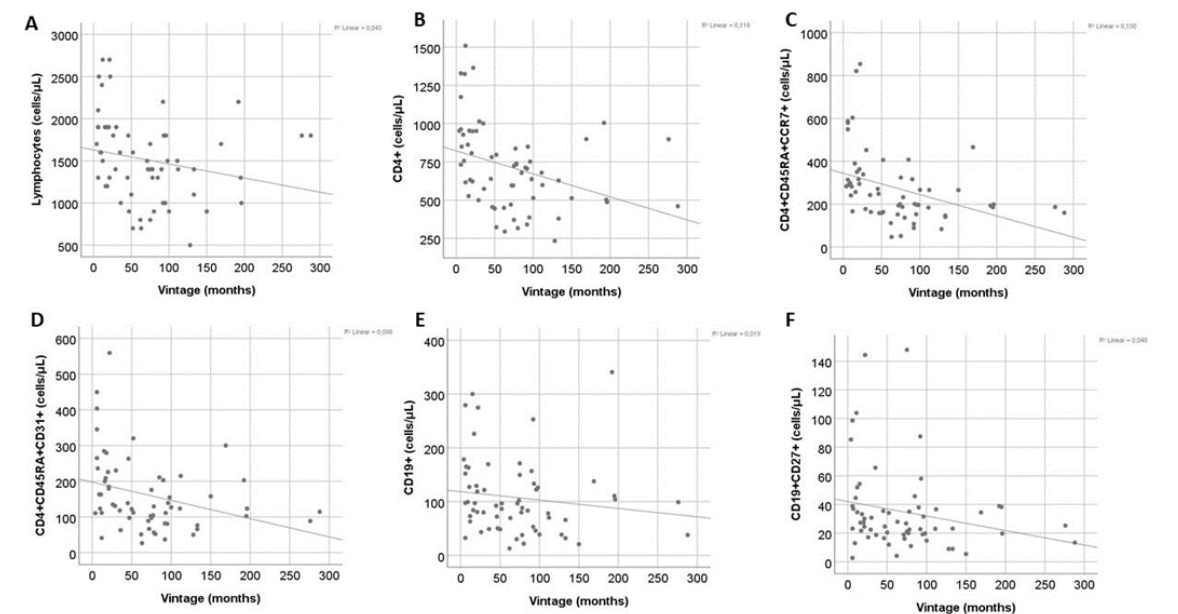


Figure 4. Linear correlation of dialysis vintage to different T cell and B cell subsets. A. Total lymphocytes B. Total CD4+ T cells. C. Naïve CD4+ T cells. D. CD4+ recent thymic emigrants E. Total B cells. F. Memory B cells.

Table 3. Univariate linear regression of T and B cell subsets for dialysis vintage and multivariate regression of T and B cell subsets for dialysis vintage and age.

Lymphocyte subsets (cells/μL)	Univariate Linear Regression					Multivariate Linear Regression					
	Vintage (months)					Vintage (months)			Age (years)		
	BC	R²	95% CI	P	PC	95% CI	P	PC	95% CI	P	
Total Lymphs	-0.21	0.04	-3.6, 0.3	0.09	-0.17	-3.3, 0.6	0.18	-0.16	-12.0, 2.5	0.19	
CD4+	-0.34	0.12	-2.6, -0.4	0.006	-0.30	-2.3, -0.2	0.02	-0.22	-7.1, 0.4	0.08	
CD4+CD45RA+CD31+	-0.31	0.98	-0.9, -0.1	0.01	-0.23	-0.7, 0.02	0.06	-0.46	-3.9, -1.3	<0.001	
CD4+CD45RA+CCR7+	-0.38	0.15	-1.6, -0.3	0.002	-0.35	-1.5, -0.2	0.006	-0.17	-3.8, 0.8	0.19	
CD4+CD45RA-CCR7+	-0.17	0.03	-1.2, 0.2	0.17	-0.14	-1.2, 0.3	0.26	-0.11	-4.0, 1.5	0.37	
CD8+CD45RA-CCR7+	-0.13	0.02	-1.2, -0.4	0.32	-0.08	-1.0, 0.5	0.53	-0.20	-5.3, 0.6	0.12	
CD19+	-0.14	0.02	-0.4, 0.1	0.28	-0.01	-0.2, 0.2	0.88	-0.52	-3.0, -1.2	<0.001	
CD19+CD27+	-0.22	0.05	-0.2, 0.02	0.08	-0.15	-0.1, 0.04	0.22	-0.29	-0.8, -0.06	0.02	

2.5. Immune phenotype may be related to increased mortality

All patients were followed up prospectively after phenotypic evaluation for at least 24 months and deaths were recorded. A 13% (8/62) of patients died during the follow-up period. Deceased patients had significantly higher proportion of exhausted CD4+PD1+ T cells in comparison to survivors [18.8(14-25) vs 11.5(7-16.9)% for respectively, P=0.006] and decreased count of total B cells [40(24-115) vs 97(70-150)cells/μL, respectively, P=0.02]. This difference affected mainly naïve and IgM memory B cells (Figure 5A). Kaplan-Meier survival analysis for patients with proportion of CD4+PD1+ above and below median was significant (P=0.002) (Figure 5B). Moreover, significant effect of exhausted CD4+PD1+ T cells proportion remained in Cox Regression either when examined alone as a continuous variable (P=0.003, 95%CI: 1.0, 1.2), or in combination with age (CD4+PD1+ proportion: P=0.03, 95% CI: 1.01, 1.23, age: P=0.02, 95% CI: 1.01, 1.44).

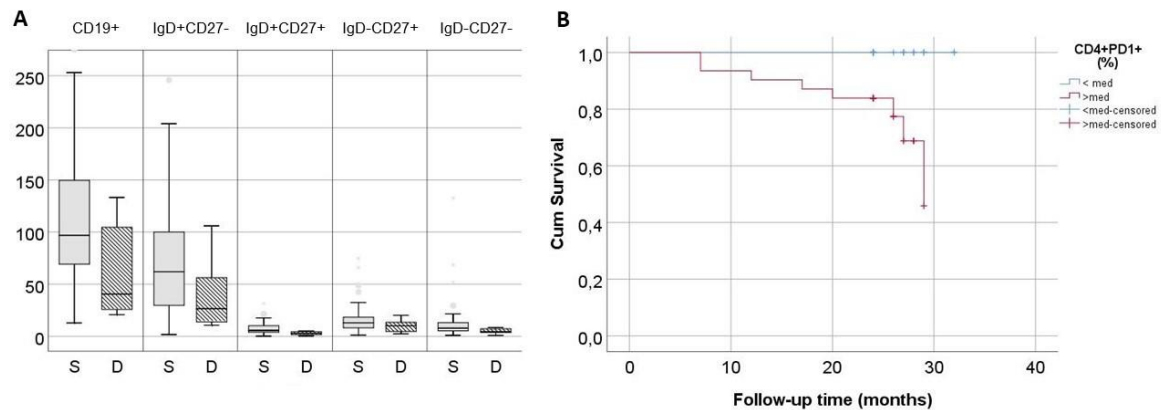


Figure 5. (A) Comparison of B cell subsets between survivors and deceased patients. (B) Kaplan-Meier survival analysis of dialysis patients according to percentage of exhausted CD4+ T cells. S: survivors, D: deceased.

3. Discussion

Alterations of acquired immunity have been studied to a certain extent in ESKD. However, the effect of dialysis modality and prescription has not been studied to date. Scope of the present study was to examine differences of senescence-related phenotype of T and B lymphocytes between patients on conventional HD or online HDF, determine the effect of hemofiltration volume and dialysis vintage and investigate the impact of T and B cells subsets to mortality.

Dialysis modality appeared to have significant effect on T cells phenotype, in terms of the expression of CD28 and PD1 molecules on CD4+ and CD8+ T lymphocytes. Patients on HDF had an improved phenotype with higher percentage of low differentiation CD28+ and lower percentage of senescent CD28-CD57- T cells. CD28, a costimulatory molecule is downregulated after multiple activations of a T cell clone [18], or as a result of cytokine action in absence of antigenic stimulation [19–21], and has been documented in many chronic inflammatory conditions, included ESKD [8, 22–23]. High percentages of CD4+CD28- T cells have been associated to instable angina and to increased risk for relapse of acute coronary event or stroke [24–28]. Moreover, these cells have been found within atheromatous plaque in animals and have been associated to vascular smooth muscle cell apoptosis and increased risk of plaque rupture [29]. CD8+CD28- T cells were also found to be increased in dialysis patients with cardiovascular disease [23].

Significant correlations between T cell phenotype with the hemofiltration volume were found in HDF patients. Our analysis revealed a clear positive effect of increased hemofiltration volume on central memory T cells, both of CD4+ and CD8+ compartment, while hemofiltration volume was negatively correlated with CD8+ EMRA T cells, a terminally differentiated subset with strong senescent characteristics [30]. Interestingly, CD8+ EMRA cell population has been positively associated with a protein bound uremic toxin, p-cresyl sulfate [31]. Hence, increased clearance of protein-bound uremic toxins, potentially achieved with HDF may contribute to the improvement of the immune phenotype of dialysis patients. Finally, a potential positive effect was also found on B cell compartment, however, in multivariate analysis, age was the main determinant of B cell subsets.

Dialysis vintage was also associated with several immunophenotype alterations. Vintage was the main determinant of total CD4+ and naïve CD4+ T cells, whereas age of CD4+ RTE and B cells in the multivariate model, despite vintage being significant in the linear correlation analysis. In 2011, Borges et al. reported a positive association of CD4+ apoptotic cells with dialysis vintage, a finding consistent to the results of the present study [32]. Central memory T cell count had a significant negative correlation to vintage, which, however, was not retained in multivariate analysis. Results from previous studies are controversial, as a recent large multicenter study supported the effect dialysis vintage on the percentage of central memory CD4+ cells [31], while other investigators claimed no significant correlation [22]. In consistency with our results, Stefanidis et al, have recently reported a gradual reduction in T lymphocyte telomere length during dialysis vintage [33].

Finally, according to the findings of the present study, immune phenotype might predict all-cause mortality in dialysis patients. Literature on the role of immune phenotype on dialysis patients' survival is, to date, poor. One study, published in 2020, associated decreased levels of low-differentiation T cell subsets to increased all-cause mortality in dialysis patients [34]. These findings were not confirmed by the present study, potentially due to lower number of patients and shorter follow-up time. B cells count $< 100/\mu\text{L}$ were also associated to increased cardiovascular and all-cause mortality in dialysis patients, according to a study of Molina et al [35]. The present study adds one more potential predictive factor, associated with mortality in this patients' group, the percentage of exhausted CD4⁺ T cells.

Limitations of the study include the relatively low number of patients enrolled and the limited follow-up interval. Moreover, patients were not initiated in a randomized way in HD or HDF before initiation of the study. Thus, more studies are needed to confirm our results.

In conclusion, HDF is potentially beneficial to patients as it was associated with improved immune phenotype in terms of CD28 and PD1 expression on T cells. An additional benefit was observed in high hemofiltration volumes, which were associated to decreased count of terminally differentiated CD8⁺ EMRA T cells. Moreover, dialysis vintage affected total and naïve CD4⁺ T cell counts, the latter, however, being an unmodifiable factor. Finally, exhausted CD4⁺ T cells might be a predictive marker for all-cause mortality in dialysis patients.

4. Materials and Methods

4.1. Patients

In this prospective, observational study, a total of 62 adult patients on chronic maintenance HD were included and followed up for at least two years. All the patients received a thrice-weekly dialysis program with HD or online HDF for at least one year, with $\text{Kt/V} > 1.2$. Patients with active malignancy, hematological or autoimmune diseases, recent infection, or vaccination (< 3 months), or patients who had received immunosuppressive treatment one year before enrollment were excluded from the study. As diabetes mellitus has been shown to provoke alterations to cellular immunity, diabetic patients were deemed not eligible for the study [36-37]. Moreover, patients who changed from HD to HDF or vice versa, for any reason, were also excluded. As many lymphocyte subpopulations are affected by ageing, though not in a linear way, we decided to stratify patients in three age groups, 20-40, 41-60 and 61-80 years. Patients were followed up for two years and cardiovascular events, de novo cancer diagnosis and deaths were recorded. The study was approved by the Institutional Review Board of the Medical School of the Aristotle University of Thessaloniki (ref No 2273/15-12-2020) and was conducted according to the principles of the Declaration of Helsinki. All study participants signed an informed consent prior to enrollment.

4.2. Flow Cytometry

Flow cytometry was performed in fresh total blood, collected immediately before the initiation of a mid-week dialysis session, processed no more than 12 hours after collection. Specimens were kept in the refrigerator at 4°C until examination. Proportions of CD4⁺, CD8⁺, and B lymphocytes subsets were determined using a cell counter (Navios Flow Cytometer, Beckman Coulter), according to the manufacturer's recommendations, as described previously [38]. For each sample four different panels of markers were prepared. Briefly, the lymphocytes were stained with conjugated antibodies for CD45, CD3, CD4, CD8, CD45RA, CCR7, CD28, CD31, CD57, PD1, CD19, CD27 and IgD, in four different tubes, as described in Supplemental Table 2, and lymphocyte subsets were defined as shown in Supplemental Table 3. Gating strategy is shown on Supplemental Figure 1.

4.3. Statistical Analysis

Statistical analysis was performed with SPSS.25 for Windows (IBM, Armonk, NY, USA). Continuous variables are reported as median(25th–75th percentile). Differences between groups were evaluated using Chi-square test for categorical variables. Comparisons of continuous variables were

evaluated with Mann–Whitney U test for comparison between two groups or Kruskal–Wallis test for comparisons among three groups. Analysis of Covariance (ANCOVA) was used to evaluate the effect of age in comparisons among subgroups. Spearman Rank Correlation Coefficient was used to estimate correlation between continuous variables. Multivariate linear regression was used for the significance of multiple continuous variables. For clinical outcomes of patients, Kaplan–Meier and Cox regression model analysis were performed. A p-value < 0.05 was considered statistically significant.

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Informed Consent Statement: Informed consent was obtained from all subjects involved in the study.

Data Availability Statement: Data will be available from the authors upon request.

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