

Patterns of peripheral blood B-cell subtypes are associated with treatment response in patients treated with immune checkpoint inhibitors: a prospective longitudinal pan-cancer study

Dominik A. Barth, Stefanie Stanzer, Jasmin A. Spiegelberg, Thomas Bauernhofer, Gudrun Absenger, Joanna Szkandera, Armin Gerger, Maria A. Smolle, Georg C. Hutterer, Sascha A. Ahyai, Tobias Madl, Florian Posch, Jakob M. Riedl, Christiane Klec, Philipp J. Jost, Julia Kargl, Martin H. Stradner, Martin Pichler

Angaben zur Veröffentlichung / Publication details:

Barth, Dominik A., Stefanie Stanzer, Jasmin A. Spiegelberg, Thomas Bauernhofer, Gudrun Absenger, Joanna Szkandera, Armin Gerger, et al. 2022. "Patterns of peripheral blood B-cell subtypes are associated with treatment response in patients treated with immune checkpoint inhibitors: a prospective longitudinal pan-cancer study." *Frontiers in Immunology* 13: 840207. <https://doi.org/10.3389/fimmu.2022.840207>.

Nutzungsbedingungen / Terms of use:

CC BY 4.0



Patterns of Peripheral Blood B-Cell Subtypes Are Associated With Treatment Response in Patients Treated With Immune Checkpoint Inhibitors: A Prospective Longitudinal Pan-Cancer Study

OPEN ACCESS

Edited by:

Isabelle Cremer,
Sorbonne Universités, France

Reviewed by:

Sophie Siberil,
INSERM U1138 Centre de Recherche
des Cordeliers (CRC), France

Philippe Rochigneux,
Institut Paoli-Calmettes (IPC), France

*Correspondence:

Martin Pichler
martin.pichler@medunigraz.at
Martin H. Stradner
martin.stradner@medunigraz.at

Specialty section:

This article was submitted to
Cancer Immunity
and Immunotherapy,
a section of the journal
Frontiers in Immunology

Received: 20 December 2021

Accepted: 10 March 2022

Published: 01 April 2022

Citation:

Barth DA, Stanzer S,
Spiegelberg JA, Bauernhofer T,
Absenger G, Szkandera J, Gerger A,
Smolle MA, Hutterer GC, Ahyai SA,
Madl T, Posch F, Riedl JM, Klec C,
Jost PJ, Kargl J, Stradner MH and
Pichler M (2022) Patterns of Peripheral
Blood B-Cell Subtypes Are Associated
With Treatment Response in
Patients Treated With Immune
Checkpoint Inhibitors:
A Prospective Longitudinal
Pan-Cancer Study.
Front. Immunol. 13:840207.
doi: 10.3389/fimmu.2022.840207

Dominik A. Barth¹, Stefanie Stanzer¹, Jasmin A. Spiegelberg¹, Thomas Bauernhofer¹,
Gudrun Absenger¹, Joanna Szkandera¹, Armin Gerger¹, Maria A. Smolle²,
Georg C. Hutterer³, Sascha A. Ahyai³, Tobias Madl^{4,5}, Florian Posch⁶, Jakob M. Riedl¹,
Christiane Klec¹, Philipp J. Jost¹, Julia Kargl^{6,7}, Martin H. Stradner^{8*} and Martin Pichler^{1,9*}

¹ Division of Oncology, Department of Internal Medicine, Medical University of Graz, Graz, Austria, ² Department of Orthopaedics and Trauma, Medical University of Graz, Graz, Austria, ³ Department of Urology, Medical University of Graz, Graz, Austria, ⁴ Gottfried Schatz Research Center, Molecular Biology and Biochemistry, Medical University of Graz, Graz, Austria, ⁵ BioTechMed-Graz, Graz, Austria, ⁶ Division of Hematology, Department of Internal Medicine, Medical University of Graz, Graz, Austria, ⁷ Otto Loewi Research Center, Division of Pharmacology, Medical University of Graz, Graz, Austria, ⁸ Division of Rheumatology and Immunology, Department of Internal Medicine, Medical University of Graz, Graz, Austria, ⁹ Department of Experimental Therapeutics, The University of Texas MD Anderson Cancer Center, Houston, TX, United States

Background: Immune checkpoint inhibitors (ICIs) have revolutionized systemic anti-tumor treatments across different types of cancer. Nevertheless, predictive biomarkers regarding treatment response are not routinely established yet. Apart from T-lymphocytes, the humoral immunity of B-lymphocytes is studied to a substantially lesser extent in the respective setting. Thus, the aim of this study was to evaluate peripheral blood B-cell subtypes as potential predictors of ICI treatment response.

Methods: Thirty-nine cancer patients receiving ICI therapy were included into this prospective single-center cohort study. All had a first blood draw at the date before treatment initiation and a second at the time of first response evaluation (after 8–12 weeks). Seven different B-cell subtypes were quantified by fluorescence-activated cell sorting (FACS). Disease control- (DCR) and objective response rate (ORR) were co-primary study endpoints.

Results: Overall, DCR was 48.7% and ORR was 25.6%, respectively. At baseline, there was no significant association of any B-cell subtype with neither DCR nor ORR. At the first response evaluation, an increase in the frequency of CD21⁺ B-cells was a statistically significant negative predictor of response, both regarding DCR (OR=0.05, 95%CI=0.00–0.67, $p=0.024$) and ORR (OR=0.09, 95%CI=0.01–0.96, $p=0.046$). An increase of the frequency of switched memory B-cells was significantly associated with reduced odds for

DCR (OR=0.06, 95%CI=0.01-0.70, $p=0.025$). Patients with an increased frequency of naïve B-cells were more likely to benefit from ICI therapy as indicated by an improved DCR (OR=12.31, 95%CI=1.13-134.22, $p=0.039$).

Conclusion: In this study, certain B-cell subpopulations were associated with ICI treatment response in various human cancer types.

Keywords: B cells, cancer, immune checkpoint inhibitor therapy, lymphocytes, response

1 INTRODUCTION

The recent introduction of immune checkpoint inhibitors (ICIs) as systemic treatment modality of solid malignancies has significantly improved patients' survival outcomes across a variety of human cancer types (1). By monoclonal antibody-mediated blocking of certain IC-molecules, mainly the programmed cell death protein 1 (PD-1), programmed cell death ligand 1 (PD-L1), and cytotoxic T-lymphocyte-associated protein 4 (CTLA4), cancer cells immune evasion can be reprogrammed and an immune response against cancer cells can be induced (2).

Although a substantial number of patients respond with long-lasting remissions or even complete remissions (CRs) upon ICI treatment initiation, a significant proportion of patients does not respond to ICI treatment at all (2). Depending on cancer entity, predictive biomarkers, such as the PD-(L)1 expression and related scores are routinely used as a potential decision tool and as predictive biomarkers in clinical practice in certain types of cancer (3, 4), yet their predictive ability is nevertheless inaccurate in most treatment settings. Besides sometimes complicated and dangerous side effects, these novel agents carry substantial treatment costs, which warrant the ongoing search for predictive biomarkers all the more, to better select appropriate candidates for ICI therapy (5–7).

To date, the crucial role of T-lymphocytes in the mode of action of ICIs in cancer therapy is well known (8–12). Nonetheless, recent evidence suggests a major role of B-lymphocytes and various B-cell subtypes in ICI-based cancer treatment (13). In addition, B-cells forming tertiary lymphoid structures (TLS) in the tumor microenvironment may also play an important role for an efficient anti-cancer immune response upon ICI therapy (14). However, a better functional understanding of different B-cell populations in ICI treatment, as well as prospective clinical studies evaluating B-cells as potential biomarkers for treatment efficacy in cancer patients undergoing ICI therapy are in general scarce. For instance, in cancer the relevance of CD21^{low} B-cells, a subset which linked to several autoimmune diseases (15) and are considered an exhausted and anergic B-cell subpopulation (15–19), undefined yet. Recently, the frequency of peripheral blood B-cells was associated with response rates in a retrospective cohort study, albeit an analysis of the B-cell subtypes involved was missing (20). As such, the potential value of peripheral blood B-cells as an easily accessible blood-based biomarker for the prediction of treatment response currently remains undefined. The aim of this

study was to prospectively evaluate the potential predictive value of peripheral blood B-cells for ICI treatment efficacy and to monitor the change of B-cell population levels during ICI treatment in a human pan-cancer setting.

2 METHODS

Forty-five consecutive cancer patients who were treated at the Division of Oncology, Department of Internal Medicine, Medical University of Graz and received ICI therapy between January 2017 and December 2020, were included into this prospective longitudinal biomarker study. Patients of all cancer types who were older than 18 years of age and who received mono- or combination ICI were included in the study. Patients with preexisting autoimmune diseases were excluded from the study.

All patients underwent a first blood draw at the date before treatment initiation and a second one 8–12 weeks after initiation of ICI treatment at the time of the first response evaluation. All patients were seen by an experienced oncologist before each treatment administration and they were evaluated for treatment response every 8–12 weeks by CT- or MRI-scans as appropriate, considering RECIST version 1.1 criteria. All blood samples were delivered in a standardized manner after collection to the Laboratory of the Division of Rheumatology and Immunology, Department of Internal Medicine, Medical University of Graz. All assays were calibrated and standardized for routine clinical sample testing.

The absolute number of CD19⁺ B cells was determined by flow cytometry as previously described (21). B-cell subsets were analyzed in peripheral blood, mononuclear cells isolated from lithium heparin blood by Ficoll gradient density centrifugation. One million cells were incubated with antibodies against CD19, IgD, CD24, CD38, CD27, CD86, CD21, and IgM (Miltenyi Biotec, Bergisch Gladbach, Germany) (**Supplementary Table 1**). Samples were measured using a FACS Canto II flow cytometer (BD Biosciences, Franklin Lakes, NJ, USA). Data were analyzed using the FACSDiva software (BD Biosciences) using a gating strategy as published previously (22) (**Supplementary Figure 1**).

B-cell subsets were classified according to the EUROclass gating strategy by Wehr et al. (23) as follows: CD21[−] (CD19⁺ CD38[−] CD21[−]), unswitched memory B cells (or marginal zone) B-cells (CD19⁺ CD27⁺ IgD⁺ IgM⁺), naïve B-cells (CD19⁺ CD27[−] IgD⁺ IgM⁺), transitional B-cells (CD19⁺ CD38⁺⁺ IgM⁺⁺), switched memory B-cells (CD19⁺ CD27⁺ IgD[−] IgM[−]),

regulatory B-cells (CD19⁺ CD24⁺ CD38⁺⁺) and plasmablasts (CD19⁺ CD38⁺⁺ IgM⁻). Regulatory B-cells were defined as proposed by Das et al. (24) and Blair et al. (25).

2.1 Statistical Analysis

Co-primary endpoints of this study were disease control rate (DCR), defined as the rate of patients who experienced CR, partial remission (PR) or stable disease (SD), and objective response rate (ORR), defined as the rate of patients who experienced CR or PR.

To assess for the association of clinico-pathological parameters with the B-cell measurements, the Mann-Whitney-U-test and the Kruskal-Wallis-test were used where appropriate. At baseline, uni- and multivariable logistic regression models were performed to assess whether different B-cell subpopulations might predict for treatment responses, whereby odds ratios (ORs) and 95% confidence intervals (CIs) are reported. Throughout the analysis, ORs for different B-cell subtypes were calculated and reported as per 1000, 100 or 10 units increase, as appropriate. Due to the hypothesis-generating character of our study, multivariable analyses were only adjusted for tumor type. To avoid perfect prediction of the outcome in the multivariable logistic regression analysis at the second blood draw, tumor types for adjustment were summarized as Non-Small Cell Lung Cancer (NSCLC), genitourinary [Renal cell carcinoma (RCC) and urothelial carcinoma (UC)], gastrointestinal (CRC, gastric and cholangiocellular carcinoma) and head and neck cancer. Increase vs. decrease or no change, absolute change (2nd blood draw - 1st blood draw) and relative change [(2nd blood draw - 1st blood draw)/1st blood draw*100] were considered in the analysis of changes in B-cell levels after 8-12 weeks of ICI treatment. Changes of B-cell levels were assessed using the Wilcoxon-Sign-Rank Test. A two-sided *p*-value of <0.05 was considered statistically significant in all analyses.

All statistical analyses were performed using Stata for Windows Version 16.1 (StataCorp LP, Collage Station, TX, USA). Box plots were created using GraphPad Prism Version 9.1.2 for Windows (GraphPad Software, San Diego, California, USA).

2.2 Ethics

Written informed consent was obtained from each patient included into this study. This study was approved by the local ethics committee of the Medical University of Graz (29-593 ex 16/17).

3 RESULTS

Overall, 45 patients treated with ICIs were included into this prospective single-center cohort study. One patient was excluded due to loss of follow-up. Five patients did not have B-cell FACS at treatment initiation and were subsequently excluded from analyses. Thus, 39 patients were included into the final analysis.

3.1 Baseline Characteristics

Twenty-seven (69%) patients were male, and the median age was 64 years. Fourteen (36%) patients had histologically verified

NSCLC, 9 (23%) patients had RCC, 7 (18%) patients had UC of the urinary bladder, 4 (10%) patients had head and neck squamous cell carcinoma and one (3%) patient each had gastric or cholangiocellular carcinoma. More than half of the patients (51%) were treated either with pembrolizumab (51%) or nivolumab (43%), whereas one patient each received treatment with atezolizumab or a combination therapy of nivolumab plus ipilimumab. More than half (51%) of the study population were in a 2nd line treatment setting, 14 (36%) patients received 1st line treatment and 5 (13%) patients received immunotherapy as a 3rd line treatment (**Supplementary Table 2**).

PR was achieved in 10 (26%) patients. No patient experienced CR during ICI treatment. Nine (23%) patients had SD and 20 (51%) patients had progressive disease (PD) at the time of first response evaluation after 8-12 weeks of ICI treatment. In the overall cohort, DCR was 48.7% and ORR was 25.6%. Due to the limited sample size and the hypothesis-generating character of the study, multivariable analyses were only adjusted to tumor type in all further analysis. However, the clinico-pathological parameters including age, sex, tumor type, treatment line, treatment modality, and histology were neither associated with DCR nor ORR in the univariable logistic regression model (all *p*>0.05, data not shown).

3.2 Association of Response-Rates With B-Cell Levels at Baseline

Summary measures of the B-cell subpopulations at baseline and respective associations with clinico-pathological parameters are shown in **Table 1**. Interestingly, absolute B-cell counts were significantly higher in female (median 6826 cells/μl; IQR 1369-10,018 cells/μl) than male patients (median 3053 cells/μl; IQR 639-5032 cells/μl) [*p*=0.0452] and the B-cell frequency was significantly higher in smokers (median 3.25%) vs. never-smokers (median 1.80%) [*p*=0.0119]. Apart from this observation, no differences in the B-cell distribution depending on clinico-pathological parameters were recorded.

Except for plasmablasts, which were significantly increased in responders, as indicated by DCR (*p*=0.048), there was no difference in the distribution of B-cell subtypes in responders and non-responders, as indicated by DCR and ORR (**Figures 1A-D**).

At baseline, there was no significant association, neither with DCR nor ORR of total B-cell, CD21⁻ B-cells, unswitched memory B-cells, transitional zone B-cells, naïve B-cells, switched memory B-cells, CD24⁺CD38⁺⁺ regulatory B-cells and plasmablasts measurements in both, univariable logistic regression models, as well as in the multivariable analysis adjusted for tumor entity (**Table 2**).

3.3 Change of B-Cell Counts and Frequency During ICI-Treatment

Twenty-seven (69.2%) patients had a second follow-up blood draw at the time point of first response evaluation after the start of ICI treatment, while 12 (30.8%) patients either dropped out of the study before having a second blood draw due to PD, death, unfitness for further treatment, or they had received no ICI therapy within 8 weeks for any other reason.

TABLE 1 | Association of B-cell subpopulations with clinico-pathological parameters.

		Median [IQR]	p-values				
			Age (>65yrs.)	Gender	Smoker	Tumor type	Treatment line
Lymphocytes	absolute count	145,922 [93,985 - 190,337]	0.6642	0.3745	0.8786	0.3338	0.5102
B-cells - total	absolute count	3725 [928 - 6,726]	0.4746	0.0452	0.0612	0.1768	0.6281
	% lymphocytes	2 [1.19 - 5.33]	0.4619	0.7348	0.0119	0.1249	0.5249
CD21-negative	absolute count	860 [246 - 2149]	1.0000	0.2098	0.0696	0.1477	0.7930
	% B cells	27 [21.4 - 48.4]	0.0642	0.4044	0.4234	0.5512	0.3513
unswitched memory B-cells	absolute count	241 [56 - 367]	0.8615	0.4048	0.1640	0.4200	0.8512
	% B cells	6.45 [4.31 - 9.18]	0.6043	0.1782	0.4780	0.3154	0.3909
Transitional Zone B-cells	absolute count	45 [12 - 193]	0.4876	0.9461	0.4735	0.3978	0.3483
	% B cells	1.55 [0.64 - 4.95]	0.6091	0.5732	0.8951	0.6548	0.1580
Naive B-cells	absolute count	1,818 [590 - 4,095]	0.3766	0.0606	0.2138	0.4098	0.6164
	% B cells	59.2 [48.1 - 78.3]	0.2429	0.3745	0.7919	0.5709	0.0664
Switched memory B-cells	absolute count	419 [179 - 915]	0.7421	0.0926	0.0934	0.2321	0.6357
	% B cells	16.7 [8.26 - 24.2]	0.5704	0.7584	0.8950	0.4727	0.1916
CD24⁺CD38⁺⁺ Regulatory B-cells	absolute count	0 [0 - 0] Maximum 2	0.5198	1.0000	0.3521	0.1714	0.8889
	% B cells	0 [0 - 0] Maximum 0.33	0.4532	0.8743	0.1715	0.1329	0.9366
Plasmablasts	absolute count	78 [19 - 266]	0.9275	0.0844	0.4737	0.2484	0.7873
	% B cells	2.97 [1.48 - 5.49]	0.3615	0.2578	0.5499	0.6135	0.9364

IQR, interquartile range; significant values are highlighted in bold.

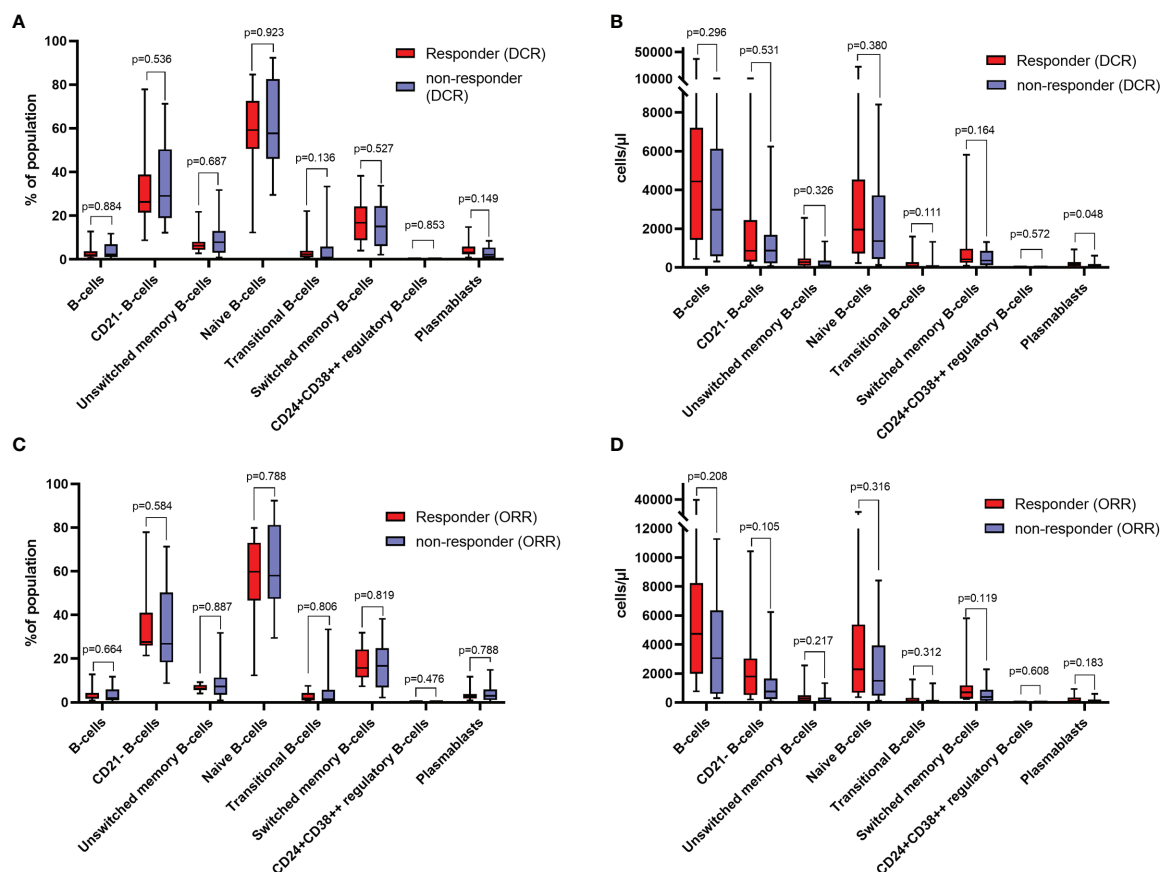


FIGURE 1 | (A) Relative distribution of different B-cell subtypes in responders and non-responders as indicated by disease control rate (DCR) at baseline. (B) absolute counts of different B-cell subtypes in responders and non-responders as indicated by DCR at baseline. (C) Relative distribution of different B-cell subtypes in responders and non-responders as indicated by objective response rate (ORR) at baseline. (D) absolute counts of different B-cell subtypes in responders and non-responders as indicated by ORR at baseline. Values in (A, C) are percentages of total lymphocytes (for B-cells), and percentages of total B-cells (for B-cell subsets). All groups compared by Mann-Whitney-U-test.

TABLE 2 | B-cell measurements and associations with disease control rate (DCR) and objective response rate (ORR) at baseline.

Variable		Disease Control Rate				Objective Response Rate			
		Univariable Analysis		Multivariable Analysis		Univariable Analysis		Multivariable Analysis	
		OR (95%CI)	p-value	OR (95%CI)	p-value	OR (95%CI)	p-value	OR (95%CI)	p-value
Lymphocytes	count*	1.01 (0.99-1.02)	0.067	1.01 (1.00-1.02)	0.037	1.01 (1.00-1.02)	0.039	1.01 (1.00-1.02)	0.045
B-cells – total	count*	1.08 (0.93-1.26)	0.300	1.15 (0.92-1.43)	0.222	1.10 (0.95-1.28)	0.189	1.10 (0.95-1.29)	0.200
	% lymph	0.96 (0.79-1.17)	0.679	0.99 (0.80-1.23)	0.925	1.02 (0.80-1.31)	0.853	1.00 (0.80-1.25)	0.984
CD21⁺ B-cells	count*	1.13 (0.81-1.59)	0.462	1.19 (0.81-1.76)	0.374	1.34 (0.93-1.96)	0.117	1.36 (0.92-2.02)	0.124
	% B***	0.87 (0.60-1.25)	0.441	0.86 (0.58-1.28)	0.463	1.05 (0.70-1.58)	0.803	1.05 (0.70-1.60)	0.793
Unswitched memory B-cells	count**	1.06 (0.92-1.22)	0.445	1.06 (0.91-1.23)	0.462	1.08 (0.94-1.24)	0.254	1.08 (0.94-1.24)	0.260
	% B	0.95 (0.85-1.05)	0.300	0.92 (0.83-1.03)	0.174	0.94 (0.81-1.08)	0.359	0.92 (0.78-1.08)	0.317
Transitional Zone B-cells	count**	1.05 (0.88-1.26)	0.580	1.06 (0.87-1.29)	0.539	1.07 (0.89-1.28)	0.459	1.07 (0.89-1.29)	0.446
	% B	1.00 (0.91-1.11)	0.945	0.99 (0.99-1.1)	0.916	0.94 (0.78-1.11)	0.448	0.94 (0.79-1.12)	0.456
Naïve B-cells	count*	1.08 (0.91-1.28)	0.372	1.11 (0.90-1.39)	0.333	1.11 (0.94-1.30)	0.217	1.11 (0.94-1.30)	0.218
	% B***	0.92 (0.65-1.30)	0.639	0.92 (0.64-1.32)	0.662	0.89 (0.60-1.32)	0.569	0.90 (0.60-1.35)	0.625
Switched memory B-cells	count**	1.08 (0.88-1.23)	0.210	1.14 (0.96-1.35)	0.139	1.08 (0.97-1.20)	0.161	1.07 (0.89-1.19)	0.177
	% B***	1.27 (0.65-2.49)	0.489	1.21 (0.60-2.45)	0.600	1.07 (0.50-2.30)	0.849	1.03 (0.46-2.31)	0.942
CD24⁺CD38⁺⁺ Regulatory B-cells	count	1.72 (0.42-7.13)	0.453	1.50 (0.34-6.72)	0.596	0.43 (0.05-3.41)	0.422	0.37 (0.04-3.26)	0.370
	% B	NA		NA		NA		NA	
Plasmablasts	count**	1.23 (0.87-1.73)	0.245	1.41 (0.89-2.24)	0.146	1.26 (0.90-1.76)	0.174	1.29 (0.91-1.85)	0.157
	% B	1.18 (0.94-1.48)	0.148	1.20 (0.94-1.52)	0.148	0.96 (0.76-1.21)	0.731	0.96 (0.76-1.23)	0.756

*per 1000 Unit increase, **per 100 Unit increase, ***per 10 Unit increase; % lymph – percent of total lymphocytes; % B – percent of total B-cells; NA – not applicable, significant values are highlighted in bold.

Regarding absolute counts at the first time of response evaluation, 11 (41%) patients had an increase in lymphocytes, 8 (30%) patients had an increase in total B-cells, 10 (37%) patients had an increase in CD21⁺ B-cells, 7 (26%) patients had an increase in unswitched memory B-cells, 10 (37%) patients had an increase in naïve B-cells, 13 (48%) patients had an increase in transitional zone B-cells, 9 (33%) patients had an increase in switched memory B-cells, and 15 (56%) patients had an increase in plasmablast counts. No individual showed an increase in CD24⁺CD38⁺⁺ regulatory B-cell counts. Significant changes in absolute counts of total B-cells (median change -789 cells/ μ l, $p=0.0422$), unswitched memory B-cells (median change -76 cells/ μ l, $p=0.0237$) and switched memory B-cells (median change -140 cells/ μ l, $p=0.0463$) could be observed, which were significantly lower at the time of response evaluation of ICI treatment in the overall cohort. However, there was no significant change of lymphocyte counts ($p=0.3997$), CD21⁺ B-cells ($p=0.0815$), naïve B-cells ($p=0.1775$), transitional zone B-cells ($p=0.9859$), CD24⁺CD38⁺⁺ regulatory B-cells ($p=0.6406$), and plasmablasts ($p=0.8919$) (Figure 2B).

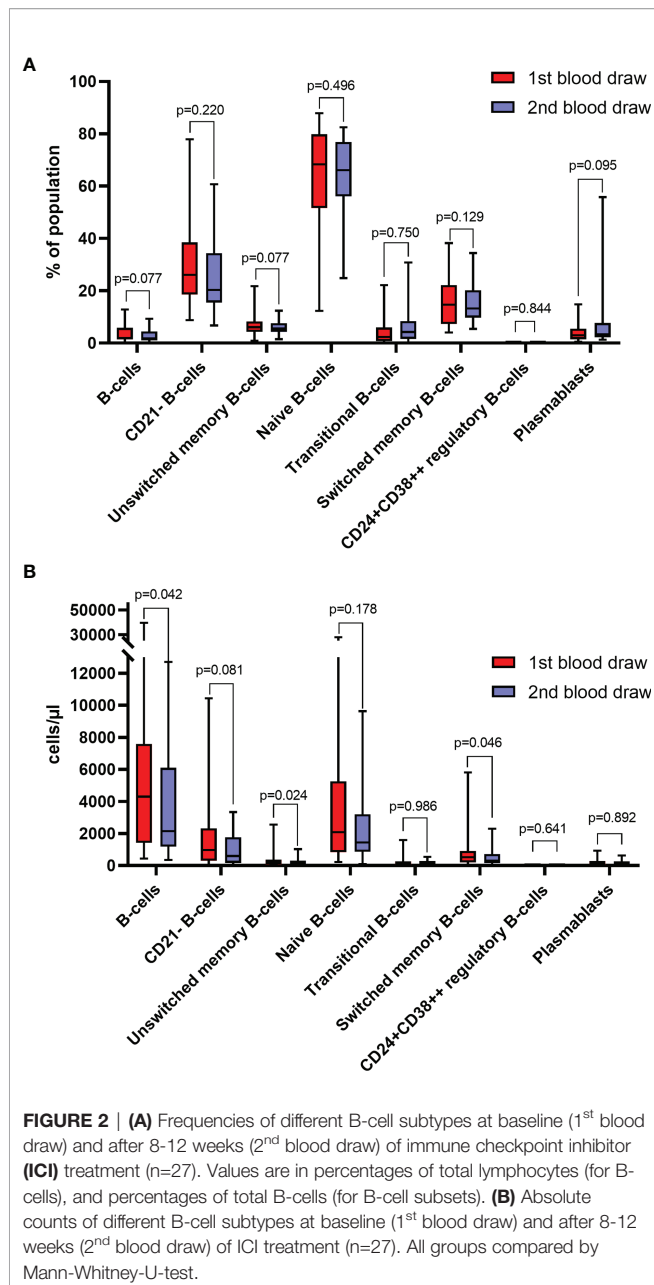
Regarding the frequency of B-cell subpopulations, 10 (37%) and 12 (44%) patients showed an increase in total B-cells and CD21⁺ B-cells, respectively. Fourteen (52%) patients each had an increase in unswitched memory B-cells, naïve B-cells, or switched memory B-cells, whereas 18 (67%) patients each showed increased frequencies transitional zone B-cells or plasmablasts after ICI treatment. No single patient had an increase in CD24⁺CD38⁺⁺ regulatory B-cell frequency. Yet, no significant changes in the frequencies of total B-cells ($p=0.0772$), CD21⁺ B-cells ($p=0.2198$), unswitched memory B-cells ($p=0.5149$), naïve B-cells ($p=0.7496$), transitional zone B-cells ($p=0.1286$), switched memory B-cells ($p=0.9247$), CD24⁺CD38⁺⁺ regulatory B-cells ($p=0.6406$) and plasmablasts ($p=0.0954$) were observed (Figure 2A).

3.3.1 Association of Changes in B-Cells With Response Rates

Patients with an increase in the frequency of naïve B-cells showed significantly greater odds for DCR in both, univariable (OR=7.00, 95%CI=1.10-44.61, $p=0.039$) and multivariable (OR=12.31, 95%CI=1.13-134.22, $p=0.039$) analyses. There was no statistically significant relationship between the frequency of naïve B-cells and treatment response, as indicated by ORR in uni- and multivariable analyses (Table 3). Yet, although a higher absolute change was not significantly associated with DCR and ORR in the univariable model, multivariable adjustment for tumor entity revealed a significant relationship of an absolute increase in the frequency of naïve B-cells with DCR (per 10-unit increase: OR=1.94, 95%CI=1.05-3.59, $p=0.035$) and ORR (per 10-unit increase: OR=2.15, 95%CI=1.07-4.34, $p=0.033$) (Table 4). There was no significant association with DCR and ORR when considering relative changes of naïve B-cell measurements (Supplementary Table 3).

Conversely, an increase of the frequency of switched memory B-cells was significantly associated with reduced odds for DCR in the univariable (OR=0.06, 95%CI=0.01-0.62, $p=0.018$) and multivariable (OR=0.06, 95%CI=0.01-0.70, $p=0.025$) logistic regression models. When considering ORR, the results were trending towards the same direction in both, the uni- and multivariable model (Table 3). However, there were no significant associations with neither DCR nor ORR when considering absolute and relative changes of the frequency of switched memory B-cells (Table 4 and Supplementary Table 3).

Patients who had an increase in the frequency of CD21⁺ B-cells during ICI treatment were less likely to respond to ICI treatment, as indicated by DCR in univariable (OR=0.11, 95%CI=0.02-0.72, $p=0.021$) and multivariable analyses adjusted for tumor type (OR=0.05, 95%CI=0.00-0.67, $p=0.024$). Concerning



ORR, an increase in the frequency of CD21⁺ B-cells was not significantly associated with response rate in the univariable model (OR=0.18, 95%CI=0.03-1.09, $p=0.061$), while it was a significant negative predictor for response when adjusted for tumor entity (OR=0.09, 95%CI=0.01-0.96, $p=0.046$) (**Table 3** and **Figure 3**). Similar results could be observed when considering absolute changes of the frequency of CD21⁺ B-cells between the first and second blood draw. An increase was significantly negatively linked to DCR in uni- (per 10-unit increase: OR=0.32, 95%CI=0.11-0.95, $p=0.040$) and multivariable analyses (per 10-unit increase: OR=0.19, 95%CI=0.04-0.81, $p=0.025$). Concerning ORR, an absolute change was not associated with ORR in the univariable analysis, but was

a significant predictor of response when adjusted for tumor entity (per 10-unit increase: OR=0.34, 95%CI=0.12-0.93, $p=0.036$) (**Table 4**). A relative change was numerically linked to DCR in uni- and multivariable analyses, yet the association was not statistically significant. Considering ORR, although close to significance, a relative change in the frequency of CD21⁺ B-cells was not associated with ORR in the univariable analysis (OR=0.09, 95%CI=0.01-1.25, $p=0.073$), but reached statistical significance in the multivariable logistic regression model (OR=0.21, 95%CI=0.00-0.71.9, $p=0.032$).

4 DISCUSSION

There is evidence that B-cells and TLS in the tumor microenvironment might be linked to ICI treatment response (14). However, the role of peripheral blood B-cells as potential biomarkers for treatment response has yet to be defined. Within our recent study, we prospectively evaluated the potential predictive value, as well as the changes over time of different B-cell subtypes in the peripheral blood of cancer patients undergoing ICI therapy. While we did not observe a statistically significant association of any B-cell subtype with treatment response at baseline, after 8-12 weeks of ICI treatment, we observed an increase of the frequency of CD21⁺ B-cells to be consistently associated with decreased odds for treatment response as indicated by both, DCR and ORR. Moreover, an increase of the frequency of naïve B-cells was significantly associated with increased odds for response, as indicated by DCR. Lastly, an increase in the frequency of switched memory B-cells was significantly negatively associated with treatment response indicated by DCR, whereas there was at least a numerical association with ORR.

The role of T-cells in cancer immunity and especially in the treatment with ICIs is well known (8, 9), however, although the importance of B-cells in forming anti-cancer immune reactions has been gradually revealed in recent years, a much better and detailed understanding of the role of different B-cell populations in cancer is still required (13). By their ability to present tumor antigens, to activate T-lymphocytes and a subsequential cytotoxic T-cell response, as well as to produce anti-tumor antibodies and cytokines, intra-tumoral B-cells and B-cells in regional tumor-draining lymph nodes support an effective anti-tumor immune response (13, 26, 27).

Recent evidence suggests that B-cell markers might be increased in tumors of patients who respond to ICI treatment (28). This was, for example, demonstrated by Helmink et al. (14) in small individual cohorts of melanoma- and RCC-patients, as well as in a TCGA (The Cancer Genome Atlas) RCC cohort. Additionally, tumor-infiltrating B-cells were previously linked to both longer survival and tumor stage in NSCLC (29, 30). Taken together, this suggests that the role of B-cells in cancer immunity might be independent of cancer entity, thus highlighting their importance across various cancer types. Furthermore, it does support the pan-cancer approach when evaluating the impact on peripheral blood B-cells on ICI treatment response in our present prospective study.

TABLE 3 | Increase vs. decrease or no change (reference) after 8-12 weeks of ICI treatment.

Variable		Disease Control Rate				Objective Response Rate			
		Univariable Analysis		Multivariable Analysis		Univariable Analysis		Multivariable Analysis	
		OR (95%CI)	p-value	OR (95%CI)	p-value	OR (95%CI)	p-value	OR (95%CI)	p-value
Lymphocytes	count	1.60 (0.30-8.50)	0.581	1.64 (0.22-12.30)	0.628	0.95 (0.19-4.68)	0.952	1.29 (0.21-7.84)	0.785
B-cells – total	count	1.75 (0.28-11.15)	0.554	1.06 (0.14-7.85)	0.951	4.67 (0.80-27.10)	0.086	5.07 (0.75-34.20)	0.095
	% lymph	0.63 (0.12-3.22)	0.574	0.28 (0.04-2.13)	0.220	1.22 (0.25-6.11)	0.807	0.91 (0.15-5.33)	0.913
CD21- B-cells	count	0.63 (0.12-3.22)	0.574	0.23 (0.28-1.89)	0.170	1.22 (0.25-6.11)	0.807	0.88 (0.15-5.24)	0.891
	% B	0.11 (0.02-0.72)	0.021	0.05 (0.00-0.67)	0.024	0.18 (0.03-1.09)	0.061	0.09 (0.01-0.96)	0.046
Unswitched memory B-cells	count	1.35 (0.21-8.82)	0.757	0.74 (0.09-5.76)	0.771	3.11 (0.53-18.38)	0.210	3.04 (0.45-20.37)	0.253
	% B	0.40 (0.08-2.12)	0.282	0.30 (0.04-2.04)	0.217	0.47 (0.10-2.29)	0.348	0.39 (0.07-2.10)	0.271
Transitional Zone B-cells	count	1.25 (0.25-6.24)	0.785	1.15 (0.16-8.11)	0.886	4.28 (0.80-22.93)	0.090	3.88 (0.61-24.74)	0.151
	% B	2.08 (0.39-11.06)	0.390	2.52 (0.39-16.29)	0.330	2.80 (0.45-17.38)	0.269	3.10 (0.45-21.21)	0.248
Naive B-cells	count	2.80 (0.45-17.38)	0.269	2.86 (0.40-20.79)	0.298	4.88 (0.90-26.42)	0.066	5.63 (0.93-34.05)	0.060
	% B	7.00 (1.10-44.61)	0.039	12.31 (1.13-134.22)	0.039	3.33 (0.63-17.57)	0.156	4.41 (0.70-27.74)	0.113
Switched memory B-cells	count	1.00 (0.18-5.46)	1.000	0.54 (0.08-3.87)	0.541	1.60 (0.31-8.25)	0.574	1.20 (0.20-7.16)	0.845
	% B	0.06 (0.01-0.62)	0.018	0.06 (0.01-0.70)	0.025	0.23 (0.04-1.25)	0.090	0.18 (0.03-1.18)	0.073
CD24*CD38** Regulatory B-cells	count	NA		NA		NA		NA	
	% B	NA		NA		NA		NA	
Plasmablasts	count	1.00 (0.20-5.00)	1.000	0.34 (0.03-4.15)	0.395	1.33 (0.27-6.50)	0.722	0.86 (0.12-6.19)	0.878
	% B	1.00 (0.18-5.46)	1.000	0.92 (0.12-7.03)	0.938	1.27 (0.24-6.82)	0.778	0.99 (0.16-6.07)	0.989

% lymph – percent of total lymphocytes; % B – percent of total B-cells; NA – not applicable, significant values are highlighted in bold.

TABLE 4 | Absolute changes of B-cells after 8-12 weeks of ICI treatment and associations with disease control rate (DCR) and objective response rate (ORR).

Variable		Disease Control Rate				Objective Response Rate			
		Univariable Analysis		Multivariable Analysis		Univariable Analysis		Multivariable Analysis	
		OR (95%CI)	p-value	OR (95%CI)	p-value	OR (95%CI)	p-value	OR (95%CI)	p-value
Lymphocytes	count*	1.00 (0.99-1.01)	0.630	1.00 (0.99-1.01)	0.755	1.00 (0.99-1.01)	0.748	1.00 (0.99-1.01)	0.635
B-cells – total	count*	0.99 (0.86-1.14)	0.907	0.96 (0.78-1.19)	0.724	0.98 (0.86-1.12)	0.776	0.96 (0.84-1.11)	0.609
	% B	1.09 (0.81-1.45)	0.577	0.98 (0.69-1.40)	0.931	1.13 (0.84-1.53)	0.416	1.07 (0.77-1.49)	0.691
CD21- B-cells	count**	0.99 (0.95-1.04)	0.778	0.98 (0.92-1.04)	0.514	0.99 (0.94-1.03)	0.470	0.98 (0.94-1.02)	0.340
	% B***	0.32 (0.11-0.95)	0.040	0.19 (0.04-0.81)	0.025	0.49 (0.21-1.19)	0.116	0.34 (0.12-0.93)	0.036
Unswitched memory B-cells	count**	0.96 (0.78-1.20)	0.730	0.94 (0.71-1.25)	0.663	1.00 (0.83-1.21)	0.999	0.98 (0.81-1.19)	0.822
	% B	0.95 (0.75-1.20)	0.655	0.91 (0.70-1.19)	0.494	0.99 (0.79-1.23)	0.898	0.96 (0.76-1.21)	0.709
Transitional Zone B-cells	count**	1.08 (0.86-1.35)	0.525	1.05 (0.80-1.37)	0.742	1.12 (0.85-1.47)	0.427	1.11 (0.83-1.50)	0.484
	% B***	2.47 (0.58-10.48)	0.220	2.27 (0.54-9.61)	0.265	3.95 (0.76-20.53)	0.102	4.38 (0.88-21.80)	0.071
Naive B-cells	count*	1.01 (0.84-1.21)	0.931	1.00 (0.80-1.26)	0.976	0.99 (0.82-1.18)	0.879	0.97 (0.80-1.17)	0.738
	% B***	1.54 (0.85-2.79)	0.156	1.94 (1.05-3.59)	0.035	1.67 (0.87-3.20)	0.122	2.15 (1.07-4.34)	0.033
Switched memory B-cells	count**	0.97 (0.87-1.09)	0.619	0.96 (0.83-1.10)	0.528	1.00 (0.91-1.10)	0.970	0.99 (0.90-1.09)	0.886
	% B***	0.30 (0.07-1.32)	0.110	0.26 (0.05-1.38)	0.113	0.81 (0.30-2.23)	0.688	0.64 (0.21-1.93)	0.426
CD24*CD38** Regulatory B-cells	count	NA		NA		NA		NA	
	% B	NA		NA		NA		NA	
Plasmablasts	count**	1.21 (0.81-1.82)	0.349	1.08 (0.63-1.87)	0.780	1.05 (0.70-1.56)	0.827	0.94 (0.57-1.54)	0.795
	% B	1.09 (0.88-1.34)	0.446	1.09 (0.84-1.41)	0.514	0.98 (0.88-1.08)	0.622	0.97 (0.87-1.08)	0.590

% B – percent of total B-cells, *per 1000 unit, **per 100 unit, ***per 10 unit increase; NA – not applicable, significant values are highlighted in bold.

In addition, Helmink et al. (14), observed that B-cells in the tumor microenvironment are primarily localized in TLS and a higher density of B-cells and TLS was associated with response. Mass cytometry in tumor and blood samples in a small sample study (n=10) revealed that non-responders had a higher frequency of naïve B-cells. Moreover, intra-tumoral CXCR3⁺ switched memory B-cells were increased in responders vs. non-responders (14). Since we observed a decrease of the frequency of switched memory B-cells in peripheral blood to be associated with response in our cohort, we hypothesized that switched memory B-cells might be drawn to and are subsequently enriched in the tumor microenvironment and

TLS of responders during ICI treatment, thus resulting in a decrease in peripheral blood. Eventually, switched memory B-cells could differentiate into plasma cells (31) in the tumor microenvironment, thereby supporting an anti-cancer immune reaction. Interestingly, plasma cells were also enriched in responders in the study of Helmink et al. (14). Yet, further longitudinal studies are warranted to better evaluate this hypothesis.

Regarding our observation of naïve B-cells with ICI-response, the number of naïve-like B-cells was previously shown to be higher in tumor samples of patients responding to ICI therapy before treatment initiation (32). In line with these results, in the

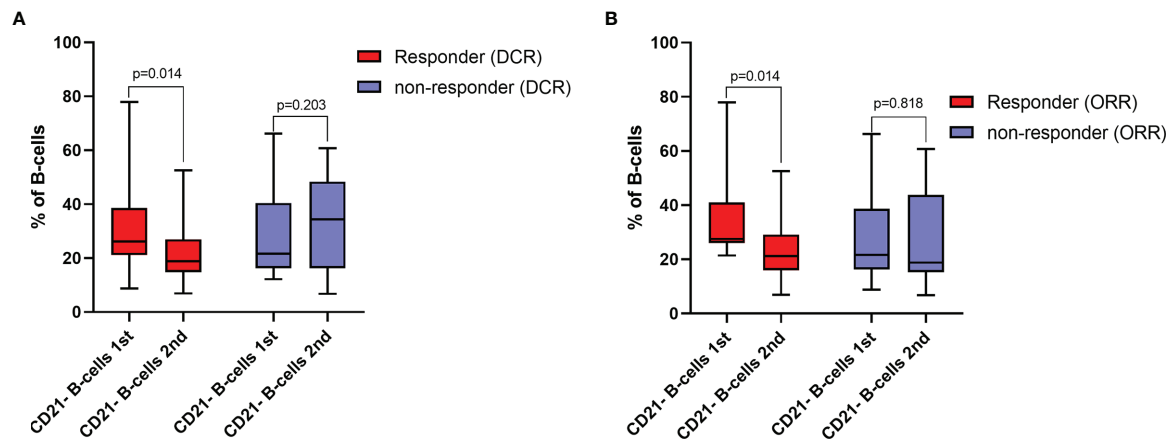


FIGURE 3 | (A) Frequencies of CD21⁺ B-cells at baseline (1st blood draw) and after 8-12 weeks (2nd blood draw) of immune checkpoint inhibitor (ICI) treatment in responders and non-responders for disease control rate (DCR). Values are in percentages of total B-cells. **(B)** frequencies of CD21⁺ B-cells at baseline (1st blood draw) and after 8-12 weeks (2nd blood draw) of ICI treatment in responders and non-responders for objective response rate (ORR). Values are in percentages of total B-cells. All groups compared by paired Sign-Rank-Test.

present study the increase of naïve B-cells during ICI treatment was associated with increased odds of response. Conversely, naïve B-cells have also been reported to be increased in tumors of non-responders to ICI treatment (14), thus definitive conclusions cannot be drawn and further studies are needed to clarify the role of naïve B-cells in ICI response.

Lastly, CD21^{low} B-cells that are linked to several autoimmune diseases, including rheumatoid arthritis, systemic lupus erythematosus, Sjogren's syndrome or common variable immunodeficiency (15), were previously reported in a context of B-cell exhaustion and might represent an anergic B-cell population (15–19). Yet, their role in cancer and especially under ICI treatment is still undefined. To the best of our knowledge, the present study is the first report showing a consistent significant association of CD21⁺ B-cells in the blood of cancer patients with ICI treatment efficacy. Considering our results of an increased frequency of CD21⁺ B-cells being independently associated with decreased odds of response, an increase in peripheral blood CD21⁺ B-cells during ICI treatment may resemble B-cell exhaustion over the course of treatment and could thus be linked to the failure of ICI therapy.

A recent retrospective study found pretreatment peripheral B-cells to be significantly decreased in patients showing response to ICI treatment in a pan-cancer cohort including 75 patients (20), which we could not validate in our prospective study. However, both uni- and multivariable regression models were not implemented in the cited study and an analysis of different B-cell populations was not conducted, which should be mentioned as an important limitation (20).

Some limitations of our present study have to be noted. Firstly, due to the limited sample size, our study might be underpowered to detect smaller differences in the distribution and association of B-cell subtypes with the clinical endpoints. However, we did observe significant signals of CD21⁺, naïve and

switched memory B-cells which presence in the tumor microenvironment has been previously linked to ICI treatment response. Thus, our data might be important for further planned larger scaled studies. Moreover, the clinical relevance of smaller changes may at least be questionable. Secondly, selection bias cannot be entirely excluded since the study was performed at a single center. Thirdly, due to the pan-cancer study design, follow-up protocols and ICI treatment dosing schemes may vary depending on cancer entity. Fourthly, PD-L1 expression status was missing for most patients since in most cases it was routinely assessed only in NSCLC. Fifth, we might have lost single plasmablasts due to our conservative CD19 gating.

In conclusion, different B-cell subtypes, notably CD21⁺, naïve and switched memory B-cells in the peripheral blood of cancer patients might represent potential novel biomarkers regarding treatment response during ICI therapy.

DATA AVAILABILITY STATEMENT

The datasets presented in this article are not readily available because The dataset for this study is not publicly available by request of the local ethic committee in order to protect the anonymity of the patients. Requests to access the datasets should be directed to martin.pichler@medunigraz.at.

ETHICS STATEMENT

The studies involving human participants were reviewed and approved by ethics committee of the Medical University of Graz (29-593 ex 16/17). The patients/participants provided their written informed consent to participate in this study.

AUTHOR CONTRIBUTIONS

MP, MSt and DB contributed to conception and design of the study. SS organized the database. DB performed the statistical analysis. All authors contributed to the interpretation to the results. DB wrote the first draft of the manuscript. All authors contributed to manuscript revision, read, and approved the submitted version.

FUNDING

This work was supported by the Austrian Science Fund (FWF; Hertha-Firnberg Grant T1112-B, to CK). TM was supported by Austrian Science Fund (FWF) grants P28854, I3792, DK-MCD W1226, DOC-130; Austrian Research Promotion Agency (FFG) Grants 864690 and 870454; the Integrative Metabolism Research Center Graz; Austrian Infrastructure Program 2016/2017, the Styrian

Government (Zukunftsfonds, doc.funds program), the City of Graz, and BioTechMed-Graz (Flagship project DYNIMO).

SUPPLEMENTARY MATERIAL

The Supplementary Material for this article can be found online at: <https://www.frontiersin.org/articles/10.3389/fimmu.2022.840207/full#supplementary-material>

Supplementary Figure 1 | Gating strategy of B-cell subsets.

Supplementary Table 1 | Antigens, clone and dye of the antibodies used.

Supplementary Table 2 | Descriptive table of the study population.

Supplementary Table 3 | Relative Changes per 100% increase of B-cells after 8–12 weeks of ICI treatment and associations with disease control rate (DCR) and objective response rate (ORR). % lymph – percent of total lymphocytes; % B – percent of total B-cells; NA – not applicable.

REFERENCES

- Havel JJ, Chowell D, Chan TA. The Evolving Landscape of Biomarkers for Checkpoint Inhibitor Immunotherapy. *Nat Rev* (2019) 19(3):133–50. doi: 10.1038/s41568-019-0116-x
- Jenkins RW, Barbie DA, Flaherty KT. Mechanisms of Resistance to Immune Checkpoint Inhibitors. *Br J Cancer* (2018) 118(1):9–16. doi: 10.1038/bjc.2017.434
- Darvin P, Toor SM, Sasidharan Nair V, Elkord E. Immune Checkpoint Inhibitors: Recent Progress and Potential Biomarkers. *Exp Mol Med* (2018) 50(12):1–11. doi: 10.1038/s12276-018-0191-1
- Cramer JD, Burtneis B, Ferris RL. Immunotherapy for Head and Neck Cancer: Recent Advances and Future Directions. *Oral Oncol* (2019) 99:104460–0. doi: 10.1016/j.oraloncology.2019.104460
- Smolle MA, Calin HN, Pichler M, Calin GA. Noncoding RNAs and Immune Checkpoints—Clinical Implications as Cancer Therapeutics. *FEBS J* (2017) 284(13):1952–66. doi: 10.1111/febs.14030
- Smolle MA, Prinz F, Calin GA, Pichler M. Current Concepts of non-Coding RNA Regulation of Immune Checkpoints in Cancer. *Mol Aspects Med* (2019) 70:117–26. doi: 10.1016/j.mam.2019.09.007
- Pichler M, Steyrer J. Cost-Effectiveness Analysis of the Use of Immunotherapy in Metastatic Solid Tumours in Austria by Applying the ESMO-Magnitude of Clinical Benefit Scale (ESMO-MCBS) Version 1.1. *ESMO Open* (2021) 6(4):100198. doi: 10.1016/j.esmoop.2021.100198
- Waldman AD, Fritz JM, Lenardo MJ. A Guide to Cancer Immunotherapy: From T Cell Basic Science to Clinical Practice. *Nat Rev* (2020) 20(11):651–68. doi: 10.1038/s41577-020-0306-5
- Paijens ST, Vledder A, de Bruyn M, Nijman HW. Tumor-Infiltrating Lymphocytes in the Immunotherapy Era. *Cell Mol Immunol* (2021) 18(4):842–59. doi: 10.1038/s41423-020-00565-9
- Sun C, Mezzadra R, Schumacher TN. Regulation and Function of the PD-L1 Checkpoint. *Immunity* (2018) 48(3):434–52. doi: 10.1016/j.immuni.2018.03.014
- Togashi Y, Shitara K, Nishikawa H. Regulatory T Cells in Cancer Immunosuppression — Implications for Anticancer Therapy. *Nat Rev Clin Oncol* (2019) 16(6):356–71. doi: 10.1038/s41571-019-0175-7
- Taheri M, Barth DA, Kargl J, Rezaei O, Ghafouri-Fard S, Pichler M. Emerging Role of Non-Coding RNAs in Regulation of T-Lymphocyte Function. *Front Immunol* (2021) 12:756042. doi: 10.3389/fimmu.2021.756042
- Largeot A, Pagano G, Gonder S, Moussay E, Paggetti J. The B-Side of Cancer Immunotherapy: The Underrated Tune. *Cells* (2019) 8(5):449. doi: 10.3390/cells8050449
- Helmkink BA, Reddy SM, Gao J, Zhang S, Basar R, Thakur R. B Cells and Tertiary Lymphoid Structures Promote Immunotherapy Response. *Nature* (2020) 577(7791):549–55. doi: 10.1038/s41586-019-1922-8
- Thorarinnsson K, Camponeschi A, Gjertsson I, Mårtensson I-L. CD21^{low} B Cells: A Snapshot of a Unique B Cell Subset in Health and Disease. *Scand J Immunol* (2015) 82(3):254–61. doi: 10.1111/sji.12339
- Isnardi I, Ng Y-S, Menard L, Meyers G, Saadoun D, Srdanovic I. Complement Receptor 2/CD21^{low} Human Naive B Cells Contain Mostly Autoreactive Unresponsive Clones. *Blood* (2010) 115(24):5026–36. doi: 10.1182/blood-2009-09-243071
- Charles ED, Brunetti C, Marukian S, Ritola KD, Talal AH, Marks K. Clonal B Cells in Patients With Hepatitis C Virus-Associated Mixed Cryoglobulinemia Contain an Expanded Anergic CD21^{low} B-Cell Subset. *Blood* (2011) 117(20):5425–37. doi: 10.1182/blood-2010-10-312942
- Visentini M, Cagliuso M, Conti V, Carbonari M, Casato M, Fiorilli M. The VH1-69-Expressing Marginal Zone B Cells Expanded in HCV-Associated Mixed Cryoglobulinemia Display Proliferative Anergy Irrespective of CD21^{low} Phenotype. *Blood* (2011) 118(12):3440–1. doi: 10.1182/blood-2011-05-353821
- Rakhmanov M, Keller B, Gutenberger S, Foerster C, Hoenig M, Driessen G. Circulating CD21^{low} B Cells in Common Variable Immunodeficiency Resemble Tissue Homing, Innate-Like B Cells. *Proc Natl Acad Sci USA* (2009) 106(32):13451–6. doi: 10.1073/pnas.0901984106
- Yuan S, Liu Y, Till B, Song Y, Wang Z. Pretreatment Peripheral B Cells Are Associated With Tumor Response to Anti-PD-1-Based Immunotherapy. *Front Immunol* (2020) 11:563653. doi: 10.3389/fimmu.2020.563653
- Rupp J, Dreö B, Gütl K, Fessler J, Moser A, Haditsch B. T Cell Phenotyping in Individuals Hospitalized With COVID-19. *J Immunol* (2021) 206(7):1478. doi: 10.4049/jimmunol.2001034
- Schulz E, Hodl I, Forstner P, Hatzl S, Sareban N, Moritz M. CD19+IgD+CD27^{low} Naïve B Cells as Predictors of Humoral Response to COVID 19 mRNA Vaccination in Immunocompromised Patients. *Front Immunol* (2021) 12:5245. doi: 10.3389/fimmu.2021.803742
- Wehr C, Kivioja T, Schmitt C, Ferry B, Witte T, Eren E. The EUROclass Trial: Defining Subgroups in Common Variable Immunodeficiency. *Blood* (2008) 111(1):77–85. doi: 10.1182/blood-2007-06-091744
- Das A, Ellis G, Pallant C, Lopes AR, Khanna P, Peppas D. IL-10-Producing Regulatory B Cells in the Pathogenesis of Chronic Hepatitis B Virus Infection. *J Immunol Baltim Md 1950* (2012) 189(8):3925–35. doi: 10.4049/jimmunol.1103139
- Blair PA, Noreña LY, Flores-Borja F, Rawlings DJ, Isenberg DA, Ehrenstein MR. CD19⁺CD24^{hi}CD38^{hi} B Cells Exhibit Regulatory Capacity in Healthy Individuals But are Functionally Impaired in Systemic Lupus Erythematosus Patients. *Immunity* (2010) 32(1):129–40. doi: 10.1016/j.immuni.2009.11.009
- Sharonov GV, Serebrovskaya EO, Yuzhakova DV, Britanova OV, Chudakov DM. B Cells, Plasma Cells and Antibody Repertoires in the Tumour

- Microenvironment. *Nat Rev Immunol* (2020) 20(5):294–307. doi: 10.1038/s41577-019-0257-x
27. Kinker GS, Vitiello GAF, Ferreira WAS, Chaves AS, Cordeiro de Lima VC, Medina TdaS. B Cell Orchestration of Anti-Tumor Immune Responses: A Matter of Cell Localization and Communication. *Front Cell Dev Biol* (2021) 9:1282. doi: 10.3389/fcell.2021.678127
 28. Amaria RN, Reddy SM, Tawbi HA, Davies MA, Ross MI, Glitza IC. Neoadjuvant Immune Checkpoint Blockade in High-Risk Resectable Melanoma. *Nat Med* (2018) 24(11):1649–54. doi: 10.1038/s41591-018-0197-1
 29. Germain C, Gnjatich S, Tamzalit F, Knockaert S, Remark R, Goc J. Presence of B Cells in Tertiary Lymphoid Structures Is Associated With a Protective Immunity in Patients With Lung Cancer. *Am J Respir Crit Care Med* (2014) 189(7):832–44. doi: 10.1164/rccm.201309-1611OC
 30. Wang S-S, Liu W, Ly D, Xu H, Qu L, Zhang L. Tumor-Infiltrating B Cells: Their Role and Application in Anti-Tumor Immunity in Lung Cancer. *Cell Mol Immunol* (2019) 16(1):6–18. doi: 10.1038/s41423-018-0027-x
 31. Dogan I, Bertocci B, Vilmon V, Delbos F, Mégret J, Storck S. Multiple Layers of B Cell Memory With Different Effector Functions. *Nat Immunol* (2009) 10(12):1292–9. doi: 10.1038/ni.1814
 32. Griss J, Bauer W, Wagner C, Simon M, Chen M, Grabmeier-Pfistershammer K. B Cells Sustain Inflammation and Predict Response to Immune Checkpoint

Blockade in Human Melanoma. *Nat Commun* (2019) 10(1):4186. doi: 10.1038/s41467-019-12160-2

Conflict of Interest: The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Publisher's Note: All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

Copyright © 2022 Barth, Stanzer, Spiegelberg, Bauernhofer, Absenger, Szkandera, Gerger, Smolle, Hutterer, Ahyai, Madl, Posch, Riedl, Klec, Jost, Kargl, Stradner and Pichler. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.