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Limited T cell response to SARS-CoV-2 mRNA vaccine among patients with cancer receiving different cancer treatments

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1. Introduction

Coronavirus disease 2019 (COVID-19) pandemic has led to more than 245 million cases and 5 million of deaths globally since the end of 2019 [1]. There has been an unprecedented global effort to develop vaccines against COVID-19, which has been a crucial step to control the pandemic. In general population, vaccines efficacy ranges between 60% and 94% and presents a good safety profile [2–7].

Patients with cancer (PC) are at high risk of acquiring COVID-19 due to their immunosuppression and higher exposition rate with frequent hospital visits. In addition, they can develop more serious complications with COVID-19 infection [8–10]. For these reasons, COVID-19 vaccination is highly recommended in these patients [11,12].

Data about the efficacy and safety of vaccines in PC comes from heterogeneous prospective studies [13–22]. Many of them include patients with solid and hematological tumors, with different types of cancer treatments and/or using different vaccines. Due to its heterogeneity, it is difficult to elucidate the specific influence of different cancer treatments on the vaccine’s humoral and cellular responses [13,17,22].

It is widely agreed that humoral response to vaccine is poor after the first administration but reaches similar rates of positive seroconversion after the second dose, when compared with a healthy population. However, antibody median titers are usually lower than in health care workers (HCW). It is still not clear how important could be this for a long-term protection [23].

Besides seroconversion, T cell response in this immunosuppressed population has emerged as an important tool to address the immunogenicity against COVID-19, and several studies even described a lower T cell response after COVID-19 infection, including CD8+ T cells and CD4+ T cells [24,25], and this could be correlated with COVID-19 severity [24]. Furthermore, T cell response could be crucial to protect against the disease with new viral variants, as recently demonstrated [26]. However, there is a paucity of data about vaccine T cell response in PC, especially by considering the possible influence of different
anticancer therapies in the immune mechanisms of T cell response. Moreover, this cellular response to vaccine could be altered in the presence of cross-reactive immunity, an important issue since pre-existing T cell response to SARS-CoV-2 has been observed in 30%−60% of unexposed individuals [27,28].

To accordingly elucidate the rate of both humoral and cellular response according to pre-existing T cell response and type of anticancer therapy, we evaluated the efficacy and safety of mRNA-1273 SARS-CoV-2 vaccination in 3 prospectively selected cohorts of patients with solid tumors treated with broadly used cancer treatments such as cytotoxic chemotherapy (CT), immunotherapy (IT), and cyclin-dependent kinases 4 and 6 inhibitors (CDK4/6i), using as control a cohort of HCW.

2. Methods

We conducted a prospective study at the Ramon y Cajal University Hospital in Madrid. Consecutive PC of three different types of therapy, aged 18 years or older visiting our Oncology Department previously to be vaccinated (from April 21 to May 13) were offered to participate in the study. As mentioned, three types of participants were enrolled: individuals with solid tumours treated with CDK4/6i (cohort A), those receiving IT (cohort B), and patients who had received or were receiving CT (cohort C). In addition, HCWs were enrolled to serve as control group. In all cohorts, patients with a history of clinical COVID-19 or anti-N baseline positive serology against SARS-CoV-2 were excluded from the study.

After inclusion, participants received two mRNA-1273 vaccinations intramuscularly, 28 days apart. The vaccination program for PC was designed by the public healthcare system authorities, and we could not control the timing of the vaccine administration within the anticancer therapy schedule. The primary objective was to analyse and compare the rate and magnitude of T cell response at least 25 days after the second dose of vaccine, whereas humoral response defined as specific serologic response was used as secondary objective.

Three blood samples were obtained from each patient: before the administration of the first vaccine dose (baseline, BL), at least 21 days after the administration of the first vaccine dose (1D), and three to four weeks after the second vaccine dose (2D). For the different analysis, 30 mL of venous blood were obtained in EDTA tubes and processed within 2 h after the collection. After the centrifugation, the plasma fraction was stored to −80 °C, while the cellular fraction was diluted in phosphate-buffered saline (PBS) followed by a Ficoll density gradient centrifugation for the isolation of PBMCs, which were subsequently washed and frozen with foetal bovine serum (FBS) and dimethyl sulfoxide (DMSO) 8%. Initially, antibodies to the SARS-CoV-2 nucleocapsid (COVID-19-SARS-CoV-2 IgG ELISA, Demeditech, Germany; positivity threshold 11 relative units (RU)/mL) was performed to identify those with previous infection. In the three consecutive samples, specific antibodies to SARS-CoV-2 Spike protein (SARS-CoV-2 IgG II Quant Alinity; Abbott, Maidenhead, UK; positivity threshold 50 arbitrary units (AU)/mL) was measured. Following WHO recommendation for standardization of anti-SARS-CoV-2 immunoglobulin determination, we converted our antibody levels units to BAU/mL (binding antibody units, conversion factor 0.142) [29]. Consequently, for specific antibodies to SARS-CoV-2 Spike protein positivity threshold is equivalent to 7.14 BAU/mL.

Cellular immune response was assessed at baseline and after complete vaccination (2D time point). Briefly, SARS-CoV-2-specific CD4+ and CD8+ T cells were measured using in vitro stimulation with SARS-CoV-2 peptide pools of viral proteins encompassing the spike (S), membrane (M), and nucleocapsid (N), followed by quantitation of CD4+ and CD8+ T cell specific interferon (IFN)-γ in live cell flow cytometry, using peripheral blood mononuclear cell (PBMC) samples from all subjects. It was considered reactive if the proportion of positive cells in stimulated wells was at least 2-fold higher in comparison with the negative control wells (unstimulated).

Patient reported outcomes were collected after both the first and the second vaccine doses. Adverse events were recorded by the participants using self-completed questionnaires after each vaccine dose.

The study was approved by our institutional ethics committee (Ramon y Cajal University Hospital Ethic Committee, EC 412/21). All the participants provided written informed consent before their inclusion in the study. The study was conducted following the Good

| Table 1 | Patient characteristics. |
|---------|------------------------|
|         | HCW (n = 36) | PC (n = 78) |
| Ageb    | 46.7 (39.9−58.8) | 58.5 (51.2−68.7) |
| Female  | 69.4%        | 73.1%        |
| Hypertension | 11.1%    | 33.3%a       |
| Diabetes | 5.6%         | 11.5%        |
| Dyslipidaemia | 25%      | 24.4%        |
| Cardiopathy | 5.6%     | 11.5%        |
| Pneumopathy | 5.6%      | 6.4%         |
| Thrombotic event | −        | 7.8%         |
| Smoking | 50%          | 44.9%        |
| Steroids | −            | 9%           |
| Anticoagulation | −        | 6.4%         |

a Results with p < 0.05 compared to controls.
b Expressed as median (IQ25−75).
Clinical Practice guidelines and the World Medical Association Declaration of Helsinki.

2.1. Statistical analysis

Continuous variables were expressed as the median and interquartile range (IQ25-75) and categorical variables by frequencies and proportions. Comparisons between groups were performed using two-tailed statistical tests, chi-square or Fisher’s exact tests for categorical variables, and Mann–Whitney test or 1-way analysis of variance (Kruskal–Wallis test) with Dunn’s correction for multiple comparisons, as appropriate. Paired samples were compared using Wilcoxon-signed rank test. Statistical significance was defined as two-sided p values <0.05. Statistical analysis was performed with STATA version 15 software, and GraphPadPrism version 8 for figures.

3. Results

3.1. Participants

Overall, 131 patients were enrolled in the study, but 17 patients were excluded at BL for serological evidence of previous SARS-CoV-2 infection. Therefore, 114 participants were the final sample of the study, 36 HCW and 78 PC. PC were subdivided in three groups: 26 patients treated with CDK4/6i, 20 patients treated with IT and 32 patients treated with CT.

Characteristics of enrolled individuals are shown in Table 1. Specifically, cancer subtypes and the different anticancer treatments administered are available in Supplementary Table S1. As it can be observed, PC were older and had more comorbidities compared to HCWs. It should be noted that 9% and 6.4% of PC were receiving treatment with steroids (more than 10 mg daily of prednisone or equivalent) and anticoagulation therapy (with new oral anticoagulants or low-molecular-weight heparin), respectively.

3.2. Humoral response (Table 2, Fig. 1)

Following vaccination, the rate specific anti-S serologic positivity (anti-S IgG cut-off point at 7.14 BAU/mL) was significantly higher in HCW compared to PC after 1D (100% versus 83.8%; p = 0.04), but similar after 2D (100% versus 95.8%; p = 0.5), but this difference was driven by PC treated with CT after 1D (100% versus 64.5%; p = 0.001) with no differences after 2D. However, the rate of humoral response of CDK4/6i and IT treated cohorts were similar compared to HCW at any timepoint. Regarding the magnitude of response, anti-S IgG levels were 1.15-fold higher in HCW compared to PC; and 2.88-fold higher when compared to PC treated with CT. No statistically significant differences were observed between PC treated with CDK4/6i and IT compared to HCW.

Considering a cut-off point at 300 BAU/mL, serologic positivity after 2D was similar between PC and HCW (97% versus 89%; p = 0.27). Only PC treated with CT showed significant lower rate of humoral response compared to HCW (97% versus 76%; p = 0.018).

3.3. Cellular response (Table 2, Fig. 2)

At baseline, pre-existing CD4+ T cell response was found in 36% of HCW and 17% of PC (p = 0.03), and pre-existing CD8+ T cell response in 31% and 23% (p = 0.5), and again the lower proportion of cross-reactivity was found in those treated with CT (p = 0.01 for CD4+ T cells).

Globally, the rate of cellular response after 2D was significantly lower in PC than in HCW for both anti-S CD4 (91.7% versus 59.7%; p < 0.001) and CD8 (94.4% versus 55.6%; p < 0.001) T cells. This effect was consistent among all PC subgroups (CDK4/6i, IT and CT) for both anti-S CD4 and CD8 T cells (p < 0.05). Furthermore, the magnitude of response measured by the percentage of IFN-g producing CD4+ and CD8+ T cells was also significantly lower in PC in comparison with HCW, showing a decreasing quantity from CDK4/6i to IT to CT.

| Table 2 | Results of immunogenicity of SARS-CoV-2 vaccine. |
|---------|-------------------------------------------------|
|         | HCW | PC | CDK4/6i | IT | CT |
| Positive serology (%) | | | | | |
| After 1D | 100% | 83.8% | 100% | 94.4% | 64.5% |
| After 2D | 100% | 95.8% | 100% | 100% | 89.7% |
| Anti-S IgG (BAU/mL) | | | | | |
| After 1D | 142 | 91 | 367 | 78 | 38 |
| After 2D | 1766 | 1540 | 3427 | 2034 | 613 |
| Anti-S CD4 T cell response (%) | | | | | |
| BL | 36.1% | 16.7% | 23.1% | 20% | 9.4% |
| After 2D | 91.7% | 59.7% | 69.2% | 58.8% | 51.7% |
| Anti-S CD8 T cell response | | | | | |
| BL | 30.6% | 23.1% | 34.6% | 25% | 12.5% |
| After 2D | 94.4% | 55.6% | 69.2% | 64.7% | 37.9% |

a Results with p < 0.05 compared with HCW.
b Expressed as median.
6i treated patients, to those with immunotherapy and to chemotherapy cohort.

We did not find any predictive factor for this poor cellular response among the baseline patient characteristics.

3.4. Patients without cross-reactivity at baseline (Fig. 3)

To avoid the possible interference of pre-existing T cell response in the rate and magnitude of response, especially among HCWs, we repeated the humoral and cellular response analysis after withdrawing those participants with T cell cross-reactivity. In these 82 patients (22 healthy volunteers, 17 patients treated with CDK4/6i, 15 with IT and 28 with CT), humoral response was similar between controls and PC after full vaccination. It should be noted that CDK4/6i patients reached significantly higher IgG levels after 1D compared with controls (p = 0.02); and CT patients had notably lower IgG titres after 1D (p = 0.02) and 2D (p = 0.03) compared to controls. In addition, the rate of positive serology in CT patients was particularly lower than
controls both after 1D (64.3%; p = 0.007) and 2D (88.5%; p = 0.1).

At a cut-off point of 300 BAU/mL, we found that PC treated with CDK4/6i reached higher rate of positive serology after 1D compared with controls (56%; p = 0.02), without significant differences after 2D. Additionally, CT patients showed lower humoral response after 2D compared to controls (76%; p = 0.02).

Regarding cellular response, PC achieved lower CD4 T cell response (50.9% versus 95.5%, p < 0.001) compared with HCW. These results were consistent within the subgroups of patients treated with CDK4/6i (64.7%; p = 0.02), IT (41.7%; p = 0.001) and CT (46.2%; p < 0.001). Comparable results were found for CD8 T cell response compared with controls (45.5% versus 95.5%, p < 0.001) and among the predefined subgroups of CDK4/6i (58.8%; p = 0.007), IT (58.3%; p = 0.14) and CT (30.8%; p < 0.001). Again, the magnitude of T cell response was also significantly blunted in the different groups of PC, especially for CD4+ T cell response after vaccine (Fig. 3).

3.5. Safety (Table 3, Fig. 4)

Overall, PC reported notably more adverse events than HCW (96.6% versus 69.2%, p < 0.001). The proportion of patients experiencing local pain, fatigue, headache, and chills was significantly higher in PC compared with HCW (p < 0.05). Symptomatic treatment for adverse events was equally administered in PC (44.2%) and controls (42.3%). None of the patients or controls experienced serious adverse events or needed hospitalization secondary to vaccination.

3.6. Follow up

Our study also included a follow-up period of 6 months to detect symptomatic COVID-19 cases. After a scheduled phone contacts during this period, only two cases
were observed. None of them required hospitalization. Seven patients died during this period without evidence of COVID-19 infection.

4. Discussion

The ONCOVac study was designed to prospectively evaluate the serologic status, immunogenicity, and safety of mRNA-1273 vaccine in three cohorts of advanced PC under different oncologic treatments, ruling out the possibility of previous undiagnosed SARS-CoV-2 infection and the existence of pre-existing T cell immunity. Our data demonstrated that T cell response is significantly blunted in percentage and magnitude in all the PC, a striking finding that could confirm the special risk situation for these patients.

The different cohorts were carefully selected based on the widespread use of these cancer therapies in oncology, and their different mechanisms of action, which could potentially influence on the immune system and thus, on vaccination efficacy. Both CDK4/6i and CT constitute immunosuppressive therapies causing mainly neutropenia through reversible bone marrow suppression by cell cycle arrest or depletion of precursor myeloid cells by apoptotic death. On the contrary, immune checkpoint inhibitors (ICI) restore an exhausted immunity and enhance cytotoxic immune response which could, hypothetically, increase the rate of response to COVID-19 vaccines.

In the entire cohort, our results showed lower seroconversion rates compared with HCW after the first dose, without differences after the second dose. These results agree with previous published studies addressing this question and confirm that the humoral response pattern in PC is gradual and slower than in a noncancer population, especially for those with haematologic malignancies. After two doses of COVID-19 vaccines, the majority of patients become seropositive [23,30,31]. This difference after the first dose was mainly driven by a lower humoral response in chemotherapy patients, while CDK4/6i and IT responded similarly to HCW at any time point. In fact, patients receiving CDK4/6i or IT had antibody titers 6- and 3-fold higher than those patients who received CT, respectively. This is also in line with available data showing that among patients receiving CT, there is a higher proportion of patients with a weak or lack of response after the first vaccine dose [32]. Also, consistent with our results, published data with CDK4/6i treated patients confirmed that SARS-CoV-2 neutralizing antibody titers were similar to HCW after the first dose of vaccine in patients with breast cancer [33].

Notably, we observed a good humoral response in those patients receiving immunotherapy, although contradictory data have been published on the influence of IT on vaccine efficacy in terms of humoral response. A recent study documented only 25% of appropriate neutralizing antibodies titers in PC under IT compared with 65.7% of HCW after the first dose [19]. Conversely, the VOICE trial described seroconversion rates of 99.3% and 100% among patients receiving IT and IT-based treatment, respectively [34].

Our study was able to evaluate the anti-S T cell response after two doses of vaccine, important information since few data about cellular response are currently available. We observed a very limited response in the rate and magnitude of CD4+ and CD8+ T cell response after two doses of the vaccine in PC comparing to noncancer population. Notably, we found that the overall rate of CD4+ T cell response was lower than expected in all the cohorts of PC compared with HCW (59.7% versus 91.7%), oscillating from 51.7% of cellular response for those patients receiving CT to 69.2% for those treated with CDK4/6i.

The lack of T cell response could be of vital importance in cancer population, since antigen-specific CD4+ T cell response plays an important role in antigen-specific B cell development, maturation and survival [35,36], and antigen-specific memory CD4+ and CD8+ T cells are likely to be less impacted by antibody escape mutations in variant viral strains [26,37].

This concerning data could be controversial for those patients receiving IT, considering the mechanism of action of these drugs, as well as data from a previous report showing an increased T cell immunity after SARS-CoV-2 infection [38]. Nevertheless, our results are in line with an exploratory finding of the CAPTURE trial, that highlighted a negative impact of IT on cellular immune responses [39]. The reason for this unexpected low T cell response to mRNA-1273 COVID-19 vaccine in this population is unknown and investigation in this issue should be encouraged.

In our study, for the first time, we also present data on the importance of pre-existing, cross-reactive, T cell immunity in PC. We showed that this cross-reactive immunity is also lower in patients with different types of cancer in comparison with HCW, and even lower in those receiving chemotherapy. We have previously demonstrated that cross-reactivity is associated with a better cellular response to the vaccine in noncancer patients [40]. Therefore, we were able to avoid the bias associated with the presence of this cross-reactivity immunity, showing that the differences between PC and patients without cancer in the rate and magnitude of T cell response were even higher after excluding these cases.

In terms of safety, PC presented significantly more adverse events than HCW (96.6% versus 69.2%, p < 0.001), mainly driven by local pain, fatigue, headache, and chills (p < 0.05). This contrasts with available
literature showing lower incidence of local (36% versus 52%) and systemic (25 versus 32%) symptoms for PC compared to healthy control after BNT162b2 vaccine [20]. On the other hand, in a study that specifically addressed the safety of IT, PC experienced similar systemic symptoms with the exception of myalgia [41]. In our opinion, this information should be interpreted with caution given the coexistence of cancer or treatment derived symptoms and heterogeneity between groups.

Our study has some limitations in addition to the small sample size. First, we did not include the analysis of neutralizing antibodies; although a good correlation between neutralizing antibodies and anti-spike antibodies has been observed [42]. Second, the cohorts were different in other factors such as age or sex, that could contribute to differences in the rate of immune response to vaccine. The median age of our control group is statistically younger, and age inversely correlates with the response to vaccines. The fact that 9% of PC were receiving concurrent steroids, which may affect the results is usual in the cancer patient population.

Additionally, there is still controversy about the antibody’s titer cutoff considered protective due to a lack of technique and assays harmonization across published studies, which make it difficult to establish a uniform threshold. In order to overcome this issue, we present the results using both, the SARS-CoV-2 IgG II Quant Alinity threshold (50 AU/mL, or 7,14 BAU/mL), and the cutoff more recently recommended as protective of 300 BAU/mL [43].

Finally, in the follow-up period we did not find high incidence of COVID-19 infections in this cohort and, therefore, we cannot link the absence or a weaker T cell response with a higher risk of COVID-19 infection.

In conclusion, we demonstrate that PC showed a blunted T cell response following mRNA vaccination, slightly modified by the type of cancer therapy and the pre-existing immunity. Although it seems that antigen-specific memory CD4+ and CD8+ T cells are an important weapon against variant viral strains, it remains to be determined if the specific T cell response can protect individuals against COVID-19. This new data together with previously published literature [43], reinforce the idea of administering a booster dose in this population.

CRediT author statement

Alfonso Cortés: Conceptualization, Project administration, Writing- Original draft preparation, Writing- Reviewing and Editing, Supervision. Jose Luis Casado: Conceptualization, Methodology, Writing- Reviewing and Editing, Supervision. Federico Longo: Conceptualization, Formal Analysis, Writing- Reviewing and Editing. Juan José Serrano: Conceptualization, Data curation, Writing- Reviewing and Editing. Cristina Saavedra: Writing- Reviewing and Editing. María Fernández: Resources. María Gion: Resources. Noelia Martínez Jaínez: Resources. Ainara Soria: Resources. Teresa Alonso: Resources. Inigo Martinez: Resources. Yolanda Lage: Resources.
Elena Lopez Miranda: Resources. Maria Eugenia Olmedo: Resources. Pablo Reguera: Resources. Pablo Gajate: Resources. Javier Molina: Resources. Eva Guerra: Resources. Raquel Fuentes: Resources. Beatriz Romero: Investigation. Mario J Rodríguez-Dominguez: Investigation. Alejandro Vallejo: Formal Analysis, Reviewing and Editing. Alfredo Carrato: Reviewing and Editing.

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**Conflict of interest statement**

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**Appendix A. Supplementary data**

Supplementary data to this article can be found online at https://doi.org/10.1016/j.ejca.2022.02.017.

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