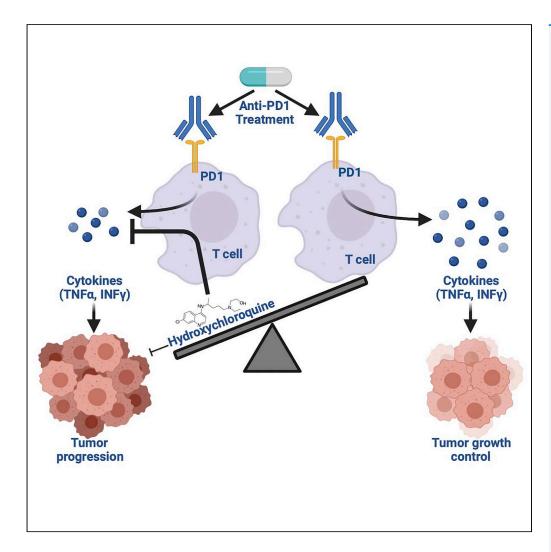
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tim.greten@nih.gov

HIGHLIGHTS

HCQ inhibits both antigen-specific and nonspecific CD4⁺ and CD8⁺ T function

HCQ inhibits tumor cell growth *in vitro*

HCQ impairs anti-PD1 therapy

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Hydroxychloroquine can impair tumor response to anti-PD1 in subcutaneous mouse models

Simon Wabitsch,¹ John C. McVey,^{1,2} Chi Ma,¹ Benjamin Ruf,¹ Olena Kamenyeva,³ Justin D. McCallen,¹ Laurence P. Diggs,¹ Bernd Heinrich,¹ and Tim F. Greten^{1,4,*}

SUMMARY

Hydroxychloroquine (HCQ) is a well-known anti-inflammatory drug but is also known as an anti-inflammatory drug. Here, we evaluate the influence of HCQ treatment on the effect of anti-PD1 tumor immunotherapy. Anti-PD1 therapy-sensitive tumor lines MC38, CT26, and RIL-175 were used to investigate the impact of HCQ on anti-PD1 therapy efficacy. *In vitro* assays demonstrated that HCQ directly inhibited tumor cell growth in all the tested tumor cell lines. HCQ treatment impaired both antigen-specific and nonspecific T-cell production of TNF α and IFN γ *in vitro* and *in vivo*. Importantly, in all the three tumor models, HCQ treatment significantly impaired the response to anti-PD1 treatment, accompanying diminished *in vivo* T-cell activation and reduced tumor-infiltrating, antigen-specific CD8⁺ T cells. This study shows that HCQ treatment can result in immunotherapy failure due to its immunosuppressive effects that offset both increased MHC-I expression by tumor cell and direct cytotoxicity.

INTRODUCTION

Hydroxychloroquine (HCQ) is an established anti-malarial medication and a pH-dependent autophagy inhibitor. Recently, its anti-tumor function has been described wherein HCQ treatment reversed immune evasion of pancreatic ductal adenocarcinoma by upregulating MHC-I surface expression on tumor cells and promoted tumor recognition by cytotoxic CD8⁺ T cells, resulting in responsiveness to anti-CTLA4 + anti-PD1 therapy (Yamamoto et al., 2020). Furthermore, IL-2 alone is an efficient immunotherapy in an advanced murine metastatic liver tumor model. IL-2 inhibits tumor growth by enhancing immune cell proliferation and infiltration in the liver and spleen; however, the anti-tumor effects of IL-2 immunotherapy significantly heighten when coupled with administration of autophagy inhibitor, chloroquine (CQ) (Liang et al., 2012). Similarly, in renal cell carcinoma, CQ is used to improve IL-2-mediated anti-tumor immunity by enhancing dendritic cell, T-cell, and natural killer cell function and limiting ATP production through inhibition of oxidative phosphorylation and promotion of apoptosis (Lotze et al., 2012). More importantly, an interaction between autophagy and immune checkpoints has been identified. In both in vitro melanoma and ovarian cancer models and using in vivo NSG (Non-obese diabetic scid gamma) mouse models, PD-L1 attenuated autophagy by blunting the ability of autophagy inhibitors to limit tumor proliferation (Clark et al., 2016). Furthermore, PD-L1 attenuation elevated mammalian target of rapamycin complex 1 (mTORC1) activity and enhanced the anti-proliferative effects of the mTORC1 inhibitor rapamycin, an autophagy inhibitor (Clark et al., 2016). These findings suggest that HCQ may enhance ICI therapy against cancer.

However, HCQ is known to cause immunosuppression (Hu et al., 2017). Indeed, it is widely used for treating autoimmune diseases such as systemic lupus erythematosus and rheumatoid arthritis. The mechanisms of HCQ-induced immunosuppression in these diseases are well known (Jeong and Jue, 1997; Mok, 2017; Weber and Levitz, 2000; Zhu et al., 1993). In the context of cancer, autophagy limits immune cell-mediated cytotoxicity (Rao et al., 2014), and the loss of autophagy genes promote tumor inflammation (Degenhardt et al., 2006; Guo et al., 2013). Most immune checkpoint inhibitors (ICIs) under investigation block one of three immune checkpoint molecules (PD-1, PD-L1, or CTLA4) on cancer or immune cells. This way, ICIs unleash the adaptive immune system and lead to an effective anti-tumor immune response (Wei et al., 2018). This mechanism may be impaired by the immunosuppressive actions of HCQ.

Of note, HCQ has gained a significant amount of interest during the coronavirus disease 2019 (COVID-19) health crisis (Gautret et al., 2020; Liu et al., 2020). Early studies during the pandemic suggested that HCQ can decrease

¹Gastrointestinal Malignancy Section, Thoracic and GI Malignancies Branch, Center for Cancer Research, National Cancer Institute, National Institutes of Health, 9000 Rockville Pike, 10/3B43, Bethesda, MD 20892, USA

²Cleveland Clinic Lerner College of Medicine at Case Western Reserve University, Cleveland, OH, USA

³Biological Imaging Section, Research Technology Branch, National Institute of Allergy and Infectious Diseases, Bethesda, MD, USA

⁴Lead contact

*Correspondence: tim.greten@nih.gov

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viral fusion and release of viral RNA from the endosome, thus potentially lowering viral spread within the patient (Al-Bari, 2015; Liu et al., 2020). However, a recent randomized trial failed to show prophylactic effects in patients exposed to severe acute respiratory syndrome coronavirus-2. Nevertheless, it is still used to treat COVID-19, along with many immunologic and rheumatologic diseases (Guan et al., 2020; Zhou et al., 2020).

This study addresses the timely and clinically relevant question of whether HCQ treatment may affect anti-PD1 treatment efficacy. We show that HCQ treatment impairs tumor cell growth *in vitro*. In contrast, HCQ impaired the response of mouse tumor models to anti-PD1 treatment. These findings suggest that the immunosuppressive effects of HCQ counteract both the upregulated presentation of tumor antigens and its direct anti-tumor effects.

RESULTS

HCQ treatment impairs T-cell function under tumor-free conditions

Anti-PD1 activates T-cell responses in patients with cancer, so we studied the effect of HCQ on T-cell function. Splenocytes from naive mice were stimulated with PMA (Phorbol myristate acetate)/ionomycin in the presence of HCQ for 4 hr to test the immediate effects of HCQ on T-cell function (Figures 1A–1D). IFN_Y and TNF^{α} production by T cells were measured by intracellular flow cytometry. HCQ caused a dose-dependent reduction of IFN_Y and TNF^{α} production in both CD4⁺ and CD8⁺ T cells (Figures 1A–1D), exemplary gating strategy shown in Figure S1A). At the 20µM concentration, we observed a strong inhibition of T-cell cyto-kine production. Next, we studied the effect of 2 hr (h) of HCQ pretreatment on T-cell stimulation (Figures 1E–1H). Again, HCQ pretreatment reduced TNF^{α} production in CD4⁺ T cells (Figure 1F) and CD8⁺ T cells (Figures 1E and 1G). We repeated these experiments using murine peripheral blood mononuclear cells (PBMCs). Again, we observed a consistent inhibition of IFN_Y and TNF^{α} production by CD4⁺ and CD8⁺ T cells when PBMCs were stimulated and simultaneously treated with HCQ (Figures S1G–S1J).

ELISA (Enzyme-linked immunosorbent assay) analysis of IFN γ and TNF α production by stimulated splenocytes treated with HCQ or pre-treated with HCQ confirmed the decrease in IFN γ and TNF α in both experimental settings (Figures 1I and 1J).

Next, we studied the effect of HCQ on antigen-specific T-cell responses. To test the MHC-I-restricted CD8⁺ T-cell response, we used OT-1 cells, whereas OT-2 cells were used to test MHC-II-restricted CD4⁺ T-cell response. CFSE (Carboxyflourescein succinimidyl ester)-labeled OT-1 or OT-2 cells were stimulated *in vitro* with their cognate peptide, and we measured cell proliferation. Consistently, HCQ reduced the proliferation of both OT-1 and OT-2 T cells, and the effect was more prominent in CD4⁺ OT-2 T cells (Figures 1K and 1L). Also, the concentration of IFN γ in the supernatant of the stimulated OT-1 and OT-2 splenocytes was lower upon HCQ treatment (Figure 1M).

To confirm HCQ-induced T-cell inhibition occurs *in vivo*, we treated naive C57BL/6 mice with HCQ at a dose suggested for treating patients with COVID-19 (Figure 1N). Splenocytes isolated from HCQ treated or control mice were stimulated with PMA/ionomycin followed by intracellular cytokine flow cytometry analysis. Similarly, the production of IFN γ and TNF α in CD8⁺ T cells and TNF α in CD4⁺ T cells (Figures 10–1Q) were significantly reduced. IFN γ production by CD4⁺ T cells was not affected (Figure 1R). Our results demonstrate that HCQ effectively impaired T-cell function, which has a potential negative impact on anti-PD1 therapy.

HCQ treatment directly inhibits tumor cell growth

Next, we studied the effect of HCQ treatment on the *in vitro* cell growth of murine and human tumor cell lines. Using an MTT (3-(4,5-dimethylthiazol-2-yl)-5-(3-carboxymethoxyphenyl)-2-(4-sulfophenyl)-2H-tetrazolium) assay, we observed dose-dependent growth inhibition of murine colorectal carcinoma cell lines, CT26 and MC38, murine hepatocellular carcinoma cell line, RIL-175, murine cholangiocarcinoma cell line, SB1, as well as human cholangiocarcinoma cell lines, EGI-1 and TFK-1 (Figures 2A–2F). Consistent with this finding, Ki67 expression was impaired in these cell lines after HCQ treatment (Figures S2A–S2E). Next, we studied the effect of HCQ treatment on molecules associated with tumor immune evasion.

Different tumor cell lines were treated with HCQ for 48hr, and MHC-I, PD-L1, and PD-L2 were tested by flow cytometry. Interestingly, MHC-I was upregulated in CT26, SB1, and MC38 but not RIL-175 as measured by

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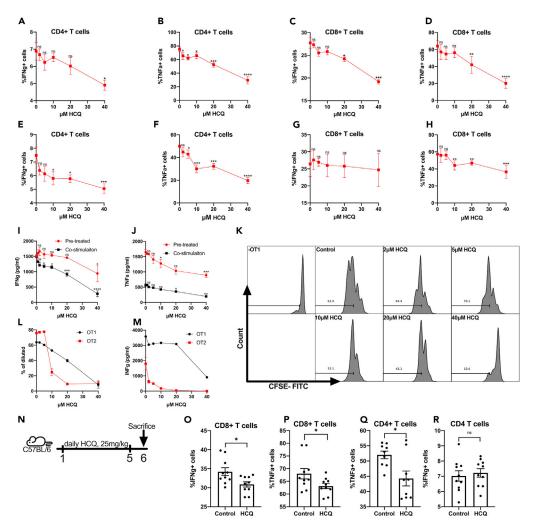


Figure 1. HCQ treatment impairs T-cell function

 $(A-D) \mbox{ Percentage of IFN} \gamma^+ (A) \mbox{ and TNF} \alpha^+ (B) \mbox{ in CD4}^+ T \mbox{ cells and IFN} \gamma^+ (C) \mbox{ and TNF} \alpha^+ (D) \mbox{ in CD8}^+ T \mbox{ cells } (n = 7 \mbox{ per group}). \mbox{ Splenocytes were stimulated with PMA and ionomycin (2 \mu L/ml) \mbox{ and simultaneously treated with HCQ for 4h. Data represent mean <math>\pm$ SEM. ns = non-significant; *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001; student's t-test. (E-H) Percentage of IFN} (E) \mbox{ and TNF} \alpha^+ (F) \mbox{ in CD4}^+ T \mbox{ cells and IFN} \gamma^+ (G) \mbox{ and TNF} \alpha^+ (H) \mbox{ in CD8}^+ T \mbox{ cells } (n = 4-8 \mbox{ per group}). \mbox{ Splenocytes were treated with HCQ for 2h, washed, and then stimulated with PMA and ionomycin (2 \mu L/ml) for 4h. Data represent mean \pm SEM. ns = non-significant; *p < 0.05, **p < 0.01, ***p < 0.001, ****p < 0.0001; student's t-test. (I and J) IFN (J) and TNF \alpha (K) concentration (pg/mL) \mbox{ in the supernatant of splenocytes treated with HCQ for 2h, washed and then stimulated with PMA and ionomycin (2 \mu L/ml) for 4 h (pre-treatment) or stimulated with PMA and ionomycin (2 \mu L/ml), and simultaneously treated with HCQ for 4 h (co-stimulation) (n = 3 \mbox{ per group}). Data represent mean \pm SEM. ns = non-significant; *p < 0.0001; student's t-test.

(K) Representative FLOW histogram of the effect of HCQ treatment on OT-1 proliferation (gating strategy: $CD3^+CD8^+CFSE^+$) in the CFSE assay.

(L) Dilution of CFSE-labeled OT-1 and OT-2 splenic cells after HCQ treatment for 48 hr (OT-1, n = 5) and 72 hr (OT-2, n = 5). Cells were stimulated with OVA I or -II peptide. Data represent mean \pm SEM.

(M) IFN γ concentration (pg/mL) in the supernatant of OT-1 and OT-2 splenic cells after HCQ treatment for 48 hr (OT-1) and 72 hr (OT-II). Cells were stimulated with OVA I or -II peptide (n = 3 per group). Data represent mean \pm SEM; student's t-test.

(N) Experimental setup: Mouse model of healthy C57BL/6 mice treated with HCQ by daily gavage and received anti-PD-1 (200μ g/mouse at indicated time points).

(O-R) Percentage of IFN γ^+ (O) and TNF α^+ (P) in CD8⁺ T cells and TNF α^+ (Q) and IFN γ^+ (R) in CD4⁺ T cells (n = 10 per group) after *in vivo* treatment of mice with HCQ. Splenocytes were stimulated *ex vivo* stimulation for 4 h. Two groups are shown: control treatment (control) vs HCQ treatment. Data represent mean \pm SEM. ns = non-significant; *p < 0.05; student's t-test.

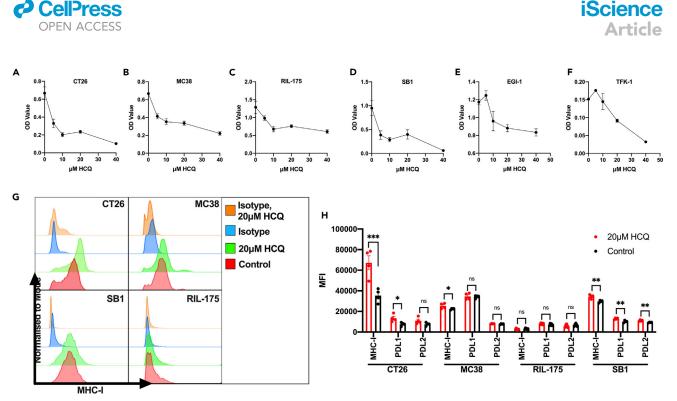


Figure 2. The effects of HCQ treatment on tumor cells

(A–F) OD value of CT26 (A), MC38 (B), RIL-175 (C), SB1 (D), EGI-1 (E), and TFK-1 (F) after HCQ treatment for 72 hr (n = 5–6 per group). MTT assay was used to measure cell viability. Data represent mean ± SEM.

(G).Representative histograms plot of surface MHC-I expression on tumor cells (CT26, MC38, SB1, RIL-175) after 48 hr of 20µM HCQ and control measured by flow cytometry.

(H).Cell surface expression of MHC-1, PD-L1, and PD-L2 on tumor cells after 20μ M HCQ treatment or control for 48 hr measured by flow cytometry (n = 3–5 per group). Data represent mean \pm SEM; ns = non-significant; *p < 0.05; **p < 0.01, ***p < 0.001, ****p < 0.001; student's t-test.

median fluorescence intensity (MFI) (Figures 2G and 2I). The MHC-I upregulation in HCQ-treated CT26, SB1, and MC38 was confirmed by immunofluorescence staining (Figures S2F–S2J). As expected, autophagy inhibition due to HCQ treatment was confirmed by increased LAMP-1 staining(Mauthe et al., 2018) (Figures S2F–S2J). Additionally, our results show that together with surface MHC-I upregulation, CT26 and SB1 cell lines have increased surface expression of PD-L1 and PD-L2 as measured by MFI (Figure 2I). Similar increases of surface MHC-I expression measured by MFI were observed in human cell lines, EGI-1 and TFK-1 (Figures S2K and S2L). The direct anti-tumor function of HCQ and the upregulation of surface MHC-I, PDL1, and PDL2 suggest the potential anti-tumor benefits of HCQ treatment when combined with anti-PD1 therapy.

HCQ treatment impairs anti-PD1 response in tumor-bearing mice

The counteracting effects of HCQ between tumor cells and T cells prompted us to investigate whether HCQ treatment impacts anti-PD1 efficacy in anti-PD1-sensitive tumors. Three tumor-bearing mouse models including MC38 and RIL-175 in C57BL/6 mice and CT26 in BALB/c mice were chosen based on previous publications and our results (Brown et al., 2018; Yu et al., 2019; Homet Moreno et al., 2016). Tumor cells were injected subcutaneously, and HCQ treatment was started when tumors were established (5 days). Mice received three doses of anti-PD1 (day 7, 10, and 15), and tumor growth was followed until day 19 (Figure 3A). Consistent with previous reports, anti-PD1 treatment effectively impaired tumor growth in all three subcutaneous tumor models (Figures 3B–3G). A single treatment of HCQ promoted tumor size growth of subcutaneous MC38 tumor but had no significant effect on subcutaneous CT-26 or RIL-175 tumor (Figure 3G). Importantly, in all three models, HCQ and anti-PD1 combination treatment led to higher tumor weights compared to anti-PD1 single treatment (Figures 3E–3G). Using three tumor lines in two mouse strains, our results clearly demonstrate that HCQ treatment can impair the efficacy of anti-PD1 therapy.

HCQ treatment impairs anti-PD1-enhanced anti-tumor immune responses

Since HCQ treatment has direct cell growth inhibitory function against MC38, CT26, and RIL-175 (Figures 2A–2C), the impaired anti-PD1 treatment efficacy in combination therapy is likely due to the T-cell function

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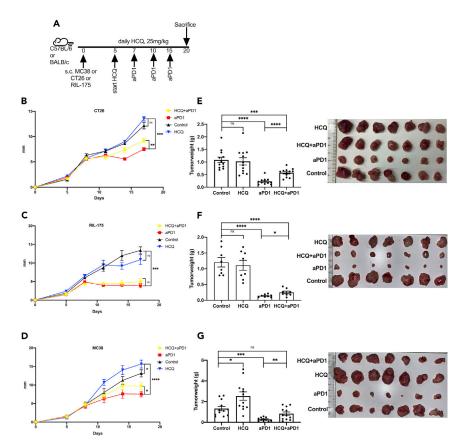


Figure 3. HCQ impairs anti-PD1 treatment response in mice in subcutaneous tumors

(A) Experimental setup: Mouse model of C57BL/6 (MC38 and RIL-175 tumor cells) and BALB/c (CT26 tumor cells) mice with subcutaneous injection of 10^6 tumor cells. HCQ and anti-PD1 treatment (aPD1, 200μ g/mouse at indicated time points).

(B–D).Tumor growth of subcutaneous CT26 (B, n = 12), RIL-175 (C, n = 8), and MC38 (D, n = 12) tumors over time. Four groups are shown: control treatment (control) vs HCQ treatment (HCQ) vs anti-PD1 treatment (aPD1) vs combination (HCQ + aPD1). Tumor growth is shown as largest tumor diameter in mm. Data represent mean \pm SEM; ns = non-significant; *p < 0.05; **p < 0.01, ***p < 0.001, ****p < 0.0001; student's t-test.

(E-G) Weight of CT26 (E, n = 12 per group), RIL-175 (F, n = 8), MC38 (G, n = 12) and subcutaneous tumors comparing control treatment (control) vs HCQ treatment (HCQ) vs anti-PD1 treatment (aPD1) vs combination (HCQ + aPD1). Representative pictures of subcutaneous CT26, RIL-175, and MC38 tumors. Data represent mean \pm SEM; ns = non-significant; *p < 0.05; **p < 0.01, ***p < 0.001, ***p < 0.001; student's t-test.

inhibition as suggested in Figure 1. To support this hypothesis, we isolated and analyzed splenocytes from MC38 tumor-bearing mice treated with anti-PD1 \pm HCQ. The expression of activation marker CD69 on both CD4⁺ and CD8⁺ T cells was greatly increased in the anti-PD1 treatment group (Figures 4A and 4B, exemplary gating strategy is shown in Figure S1B). The addition of HCQ treatment completely blocked the CD69 increase when compared to combination treatment. T-cell function was further analyzed using intercellular cytokine flow cytometry analysis after *ex vivo* PMA/ionomycin stimulation. Upon anti-PD1 treatment, both CD4⁺ and CD8⁺ T cells produced significantly more IFN γ and TNF α (Figures 4C–4F). Anti-PD1 in combination with HCQ did not reduce IFN γ and TNF α production in CD8⁺ T cells (Figures 4C and 4D). In CD4⁺ T cells, the combination therapy decreased IFN γ but not TNF α production (Figures 4E and 4F).

Tumor-infiltrating lymphocytes (TILs) were analyzed in CT26 tumors, and anti-PD1 treatment caused a robust increase of CD4⁺ and CD8⁺ TILs (Figures 5A and 5B). Additionally, HCQ treatment greatly reduced the anti-PD1 treatment-dependent accumulation of CD4⁺ and CD8⁺ TILs. Furthermore, we used an AH-1 tetramer to quantify tumor antigen-specific CD8⁺ T cells in CT26 tumors³⁴. Again, the absolute number of AH-1 tetramer⁺ CD8⁺ TIL cells was significantly higher with anti-PD1 alone treatment but dropped to





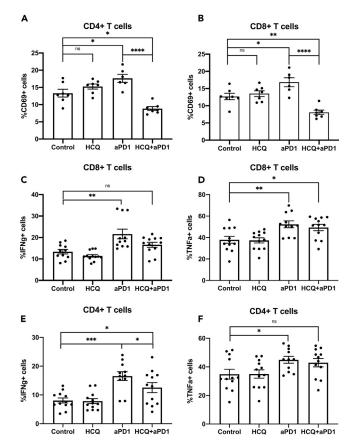


Figure 4. HCQ treatment impairs T-cell activation in tumor-bearing mice

(A and B) Percentage of splenic CD69⁺CD4⁺ (A) and CD8⁺ (B) T cells of MC38 tumor-bearing mice after HCQ and anti-PD1 treatment (n = 7–8 per group). Experimental setup is shown in Figure 3A and group setup as in Figure 3B. Data represent mean \pm SEM; ns = non-significant; *p < 0.05; **p < 0.01; ***p < 0.001; student's t-test.

(C–F) Percentage of splenic IFN γ ⁺ (C) and TNF α ⁺ (D) and CD8⁺ IFN γ ⁺ (E) and TNF α ⁺ (F) CD4⁺ T cells of MC38 tumorbearing mice after HCQ and anti-PD1 treatment (n = 11–12 per group). Experimental setup is shown in Figure 3A and group setup as in Figure 3B. Data represent mean \pm SEM; ns = non-significant; *p < 0.05; **p < 0.01; ***p < 0.001; student's t-test.

baseline in the combination group (Figures 5C and 5D). Together, our results suggest that HCQ impairs the anti-PD1 treatment-induced anti-tumor T-cell response and subsequently leads to reduced anti-tumor effects.

DISCUSSION

HCQ is an immunosuppressive drug utilized in treating various autoimmune diseases, such as systemic lupus erythematosus and rheumatoid arthritis (Lee et al., 2011; Schrezenmeier and Dörner, 2020). In the context of autoimmune disease, HCQ inhibits immune function through many mechanisms such as alterations in autophagy and lysosomal pH (Hu et al., 2017). For example, autophagy and lysosomal function in dendritic cells is an important mechanism by which MHC-II antigens are processed and presented. The blockage of autophagy and rising lysosomal pH impairs MHC-II presentation and thus the function of CD4+ T cells (Schmidt et al., 2017). In the context of cancer, MHC-II antigen presentation and CD4+ T cells response are an important component of anti-tumor immunity, as well as response to ICIs. Furthermore, HCQ can inhibit the secretion of important cytokines such as IFN γ , TNF α , and IL-6 (Alspach et al., 2019). These studies suggest that the use of HCQ may inhibit anti-tumor response, especially when using ICIs.

Autophagy inhibition is being increasingly recognized as an important anti-tumor approach to improve traditional chemotherapy (Boone et al., 2015; Briceño et al., 2007). Two recent reports showed that CQ improves the efficacy of chemotherapy by inhibition of autophagy (Kinsey et al., 2019; Bryant et al., 2019). Both

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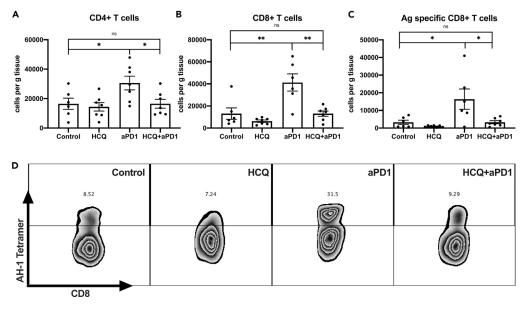


Figure 5. Effects of HCQ treatment on tumor-infiltrating lymphocytes

(A and B) Tumor-infiltrating CD4+ (A) and CD8+ (B) T cells per gram of CT26 tumor after HCQ and anti-PD1 treatment (n = 6-7 per group). Experimental setup is shown in Figure 3A and group setup as in Figure 3B. Data represent mean \pm SEM; ns = non-significant; *p < 0.05; student's t-test.

(C) Tumor-infiltrating AH-1+CD8+ T cells per gram of CT26 tumor after HCQ and anti-PD1 treatment (n = 6–7 per group). Experimental setup is shown in Figure 3A and group setup as in Figure 3B. Data represent mean \pm SEM; ns = non-significant; *p < 0.05; student's t-test.

(D) Representative contour plot and frequencies of tumor-infiltrating antigen-specific AH-1+CD8+ cells in CT26 tumors after HCQ and anti-PD1 treatment (gating strategy: CD3⁺CD8⁺AH1⁺). Experimental setup is shown in Figure 3A and group setup as in Figure 3B.

of these studies investigated the effects of CQ in KRAS-mutated pancreatic ductal adenocarcinoma and found that CQ had synergistic effects with KRAS pathway inhibitors. Other studies show that CQ and HCQ can be used to improve chemotherapy by inhibiting autophagy and raising lysosomal pH (Goldberg et al., 2012; Chen et al., 2018).

While there is compelling evidence to suggest that HCQ and CQ can improve the efficacy of chemotherapy, little is known about the immunosuppressive properties of HCQ and CQ in relation to immunotherapies such as ICIs. Studies in mice have shown that CQ and HCQ can induce a shift from immunosuppressive M2-tumor-associated macrophages (TAMs) to immune-promoting M1-TAMs, which may sensitize glioblastoma to anti-PD1 treatment (Hsu et al., 2020; Li et al., 2018). A recent report showed that inhibiting autophagy reversed the downregulation of MHC-I surface expression on pancreatic cancer cells. Moreover, CQ sensitized otherwise ICI therapy-insensitive pancreas cancer to dual treatment with anti-PD1 and anti-CTLA4 (Yamamoto et al., 2020), suggesting that CQ can potentially improve ICIs. However, the study did not test the direct effect of CQ on anti-tumor immunity and did not address its influence on ICI therapy-sensitive cancer. Of note, the study observed that CQ failed to boost anti-PD1 monotherapy against pancreatic cancer. Other reports using tumor models without downregulated MHC-I expression showed that genetic inhibition of autophagy or the use of HCQ had no improved function of T cells in tumor-bearing mice (Starobinets et al., 2016). Taken together, these studies paint an unclear picture of how the immunosuppressive drugs, HCQ and CQ, affect the adaptive immune response in the context of ICIs.

At a time when there is increasing interest in HCQ for the treatment of cancer, as well as COVID-19, this study addresses whether HCQ affects anti-PD1 therapy. Our results show that HCQ diminishes the anti-tumor effects of anti-PD1 in solid tumor mouse models. We demonstrate that HCQ impairs the production of proinflammatory cytokines in mice bearing various tumor types. We also show that HCQ reduces the number of macrophages and dendritic cells, which likely impairs a robust T-cell response. Finally, we found that MHC-I surface expression was only increased in a subset of tumor cell types, indicating significant heterogeneity within different tumor cell lines. Of note, HCQ-treated tumors showed accelerated MC38 tumor





growth compared to control. Taken together, these results suggest that the immunosuppressive properties of HCQ outweigh its anti-tumor properties in tumor-bearing mice treated with anti-PD1.

These results suggest that HCQ should not be used as an autophagy inhibitor in combination with anti-PD1 in patients with cancer. Furthermore, patients with COVID-19 treated with ICIs should be excluded from the administration of HCQ.

Limitations of the study

Subcutaneous mouse models are useful in elucidating *in vivo* treatment effects, yet orthotopic models provide in-depth analysis regarding the organ-specific tumor microenvironment. Our study demonstrates that HCQ impairs anti-PD1 treatment through an associated decrease in tumor-infiltrating, antigen-specific lymphocytes. *In vitro* and *in vivo* models suggest that CD4⁺ and CD8⁺ T cells produce proinflammatory cytokines, yet we do not demonstrate definite molecular mechanisms describing the immune suppressive effects of HCQ. The *in vivo* data are consistent between three tumor cell lines and two mouse strains, yet the absence of human data limits the study's immediate translatability. Finally, HCQ is being explored as an adjuvant to several chemotherapy regimens. The anti-tumor effects of HCQ through autophagy inhibition are apparent, and future studies are needed to successfully utilize HCQ, while also mitigating its immune suppression functions.

Resource availability

Lead contact

Further requests should be directed to and will be fulfilled by the Lead Contact: Tim Greten (tim.greten@ nih.gov).

Material availability

No new unique reagents were generated in this study.

Data and code availability

No unpublished custom code, software, algorithm, or data sets were used in this study.

METHODS

All methods can be found in the accompanying Transparent methods supplemental file.

SUPPLEMENTAL INFORMATION

Supplemental information can be found online at https://doi.org/10.1016/j.isci.2020.101990.

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AUTHOR CONTRIBUTIONS

S.W., C.M., J.D.M., and T.F.G. contributed in project design; S.W., B.R., O.K., and J.D.M. completed all experiments; S.W., J.C.M., C.M., B.R., L.D., B.H., and T.F.G. contributed in data analysis and interpretation; S.W., J.C.M., and T.F.G. wrote the original draft; and all authors contributed in reviewing and editing.

DECLARATION OF INTERESTS

The authors declare no potential conflicts of interest.

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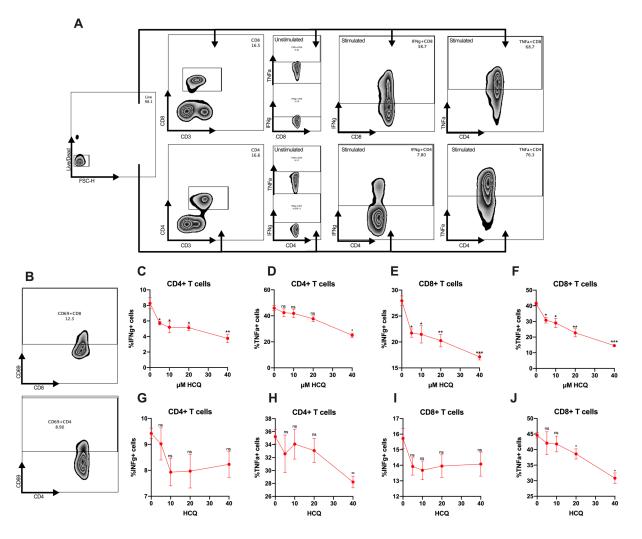
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Supplemental Information

Hydroxychloroquine can impair tumor response

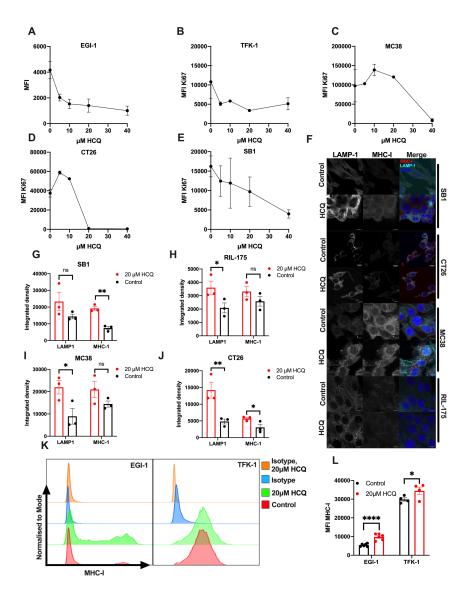
to anti-PD1 in subcutaneous mouse models

Simon Wabitsch, John C. McVey, Chi Ma, Benjamin Ruf, Olena Kamenyeva, Justin D. McCallen, Laurence P. Diggs, Bernd Heinrich, and Tim F. Greten



Supplemental Figure 1: HCQ impairs PBMCs T cell function. Related to Figure 1.

- A. Representative contour plot of the gating strategy for CD8⁺ (upper row) and CD4⁺ (lower row) T cells and gating for IFN γ^+ and TNF α^+ of unstimulated and stimulated splenocytes.
- **B**. Representative contour plot of the gating strategy for CD69⁺ in CD8⁺ (upper row) and CD4⁺ (lower row) T cells (gating on CD4+ and CD8+ T cells from Fig. S1A).
- **C-F.** Percentage of IFNγ⁺ (**C**) and TNFα⁺ (**D**) in CD4⁺ T cells and IFNγ⁺ (**E**) and TNFα⁺ (**F**) in CD8⁺ T cell (n=4 per group). PBMCs were stimulated with PMA and ionomycin (2µL/ml) and simultaneously treated with HCQ for 4h. Data represent mean ± SEM. ns= non-significant; *p<0.05, **p<0.01, ***p<0.001; student's t-test.
- **G-J.** Percentage of IFNγ⁺ (**E**) and TNFα⁺ (**F**) in CD4⁺T cells and IFNγ⁺ (**G**) and TNFα⁺ (**H**) in CD8⁺T cells (n=4 per group). PBMCs were treated with HCQ for 2h, washed and then stimulated with PMA and ionomycin (2µL/ml) for 4h. Data represent mean ± SEM. ns= non-significant; *p<0.05, **p<0.01, ****p<0.001, ****p<0.0001; student's t-test.



Supplementary Figure 2: HCQ inhibits tumor cell proliferation and upregulates MHC-1. Related to Figure 2.

- A-E. Expression of Ki67 in EGI-1 (A), TFK-1 (B), MC38 (C), CT26 (D) and SB1 (E) tumor cells after HCQ treatment for 48h. Ki67 was measured by flow cytometry (n=3 per cell line). Data represent mean ± SEM.
- **F**. Representative images of SB1, CT26, MC38 and RIL-175 tumor cells after 20μM HCQ treatment for 48h or control. Tumor cells were stained with LAMP-1 (column 1), MHC-I (column 2) and merge (column 3). Scale bar: 10μm.
- **G-J.** Median integrated density per cell of LAMP-1 and MHC-1 staining of SB1 (H), RIL-175 (I), MC38 (J) and CT26 (K) cells after 24h of HCQ treatment and control (n=3 high power field). Data represent mean ± SEM; ns= non-significant; *p<0.05, **p<0.01; student's t-test.
- **K.** Representative FLOW plot of MHC-I expression on tumor cells (EGI-1 and TFK-1) after 48h of 20μM HCQ and control.
- L. MFI of cell surface expression of MHC-1 on EGI-1 and TFK-1 tumor cells after 20µM HCQ treatment or control for 48h measured by flow cytometry (n=5 per group). Data represent mean ± SEM; ns= non-significant; ***p<0.001, ****p<0.0001; student's t-test.

Transparent Methods

Cell Lines

Two murine colorectal carcinoma cell lines CT26(Brattain et al., 1980) and MC38(Corbett et al., 1975), one murine hepatocellular carcinoma cell line RIL-175(Yu et al., 2019), one murine cholangiocarcinoma cell lines SB1 (Rizvi et al., 2018) and two human cholangiocarcinoma cell lines TFK-1(Saijyo et al., 1995) and EGI-1 (Wolf et al., 1985) were used in this study.

Mouse strains and reagents

Female C57BL/6 and BALB/c mice were purchased from the Charles River Laboratory. OT-1 (C57BL/6-Tg(TcraTcrb)1100MJb/J) and OT-2 (B6.Cg-Tg(TcraTcrb)425Cbn/J) mice were purchased from the Jackson Laboratory. Mice were used at age 8-12 weeks for experiments. All experiments were conducted according to local institution guidelines and approved by the Animal Care and Use Committee of the National Institutes of Health, Bethesda, USA. HCQ was obtained from the NIH clinical center pharmacy.

Animal studies

Subcutaneous injections were performed in 8-week-old C57BL/6 and BALB/c female mice. 1x10⁶ CT26 or MC38 or RIL-175 cells were re-suspended in 100µL of PBS and the tumor cell suspension was injected in the lateral flank. All mice were randomized prior to initiation of treatment. Mice (n=5-8 per group) were then treated 5 days after injection with either vehicle control (saline), daily gavage of 25mg/kg HCQ or 200µg anti-PD-1 (clone 29F.1A12, BioXCell) on days 7, 10, 15. One group received combined treatment with both HCQ and anti-PD1. Tumor size was measured by diameter and mice were sacrificed at day 20 post tumor cell injection. Tumors were then excised and weighed. Tumor weights were compared between groups and lymphocytes were isolated from spleens and tumors for further flowcytometry analysis. Numbers of infiltrating lymphocytes were counted by flow cytometry and divided by tumor weight (g). For the immune profiling experiments, non-tumor bearing mice (n=5 per group) were treated with vehicle control or daily gavage of 25mg/kg HCQ. Mice were sacrificed at day 6 and splenic lymphocytes were isolated.

In vitro T cell activation assay

Splenocytes were isolated from C57BL/6 mice and $2x10^5$ cells were seeded into 96-well plate in T cell media. Cells were stimulated with 2μ L/ml of leukocyte activation cocktail (containing Phorbol 12-Myristate 13-Acetate [PMA], ionomycin and GolgiPlugTM [Brefeldin A], BD Bioscience, 51-2042E) and treated with 1, 2, 5, 10, 20 and 40 μ M HCQ for 4h during stimulation and 2h prior stimulation. PBMCs were isolated and $1x10^5$ cells were seeded into a 96-well plate, stimulated as splenocytes and treated with 5, 10, 20 and 40 μ M HCQ for 4h during stimulation. The levels of IFN γ and tumor necrosis factor alpha (TNF α) of CD4⁺ and CD8⁺ cells were analyzed by flow cytometry.

In vitro T cell proliferation assay

OT-1 or OT-2 splenic T cells were CFSE-labeled and seeded into 96-well plates. Total of $2x10^5$ cells per well were stimulated with 1µg/ml OVA-1 or OVA-2 peptide (Invivogen) and treated with 0.5, 1, 2, 5, 10, 20 and 40 µM HCQ in T cell media (RPMI, 10% FCS, 0.05mM 2-Mercaptoethanol, 2.5 µL/L IL-2) for 48h (OT-1) and 72h (OT-2). Co-cultures experiments were performed in triplicate for each condition. Proliferation of CFSE+CD8-OT-1 and CFSE+CD4-OT-2 cells was measured by flow cytometry (Gamrekelashvili et al., 2013).

ELISA

IFN γ and TNF α concentration in the supernatant of stimulated splenocytes (2µL/ml of leukocyte activation cocktail containing PMA and ionomycin; Biolegend, 423301) was determined using mouse IFN γ (abcam, ab100689) and TNF α (abcam ab 100747) KIT. Cells were treated with 1, 2, 5, 10, 20 and 40 µM HCQ for 4h during stimulation or 2h prior stimulation. IFN γ supernatant concentration of OT-1 or OT-2 splenic T cells was measured after treatment with 1, 2, 5, 10, 20 and 40 µM HCQ and stimulation with OVA-1 and OVA-2 peptide for 48h (OT-1) and 72h (OT-2).

In-vitro tumor cell viability assay and antigen profiling

A total of 5000 CT26, MC38, and SB1 cells per well and 10000 TFK-1 and EGI-1 cells were seeded overnight into 96-well plate in 200µL media (RPMI, 10% FBS, 1%Pen/Strep). The media was removed, and

cells were treated with 5, 10, 20 and 40 μ M HCQ for 72h. Cell viability was measured with the MTT Cell Proliferation Kit (abcam, ab211091) according to manufacturer's instructions. For intracellular and extracellular staining of tumor cells, $3x10^5$ cells per well were seeded overnight. Treatment with 20 μ M HCQ was performed for 24h and analyzed using flowcytometry.

Immunostaining and confocal microscopy

Cell lines were cultured for 48h on coverslips in a 6-well plate, washed with PBS and treated with 20µM HCQ in media for 24h. Cells were fixed with paraformaldehyde for 15min at room temperature and permeabilized with 0.1% Saponin (EMD Millipore Corp.) for 30min at RT. Blocking was performed with PBS containing 20% FCS (Sigma) for 30 min and incubated with primary antibodies overnight at 4°C. The following antibodies were used for immunofluorescence: murine anti-MHC-I (clone ERMP42, abcam), murine anit-LAMP-1 (clone C54H11, Cell Signaling). Confocal microscopy was performed using Leica DMi8 inverted 5 channel confocal microscope equipped with ultra-sensitive hybrid detectors (HyDs, Leica Microsystems) and full range of visible lasers. Diode laser for 405 nm and HeNe lasers for 594 and 633 nm excitation were used at minimal laser power (0.05-1%). Z stacks of single cells (Z=10–15 µm) were collected using 63X objective (Leica Microsystems). Post-acquisition image processing was performed using Imaris software (Imaris version 9.5.1, Bitplane AG). Median integrated density per cell was determined by ImageJ.

Flow cytometry

Splenocytes and tumor infiltrating mononuclear cells were prepared. Briefly, tumors were removed immediately after mice were sacrificed. After homogenization using gentleMACS Octo Dissociator, debris was removed by passing through a 100µm and 40µm cell strainer (Fisherbrand). Tumor infiltrating cells were isolated by cell separation media (Cedarlane) centrifugation (2000xg, 20min). Cells were surfacelabeled with indicated antibodies for 30 minutes at 4°C. Intracellular staining using a Foxp3/transcription factor staining buffer set or Fixation/Permeabilization Solution Kit (both eBioscience) was used according the manufacturer's instructions. Flow cytometry was performed on CytoFLEX LX platforms and results were analyzed using FlowJo software version 10.4.2 (BD Biosciences). Dead cells were excluded by using live/dead fixable near-IR dead cell staining kit (ThermoFisher scientific). The following antibodies were used for flow cytometry analysis: anti-CD3-BV605 (clone 17A2, Biolegend), CD8-Alexa Fluor 700 (clone 53-6.7, Biolegend), anti-CD8-PB (clone 53-6.7, Biolegend), anti-CD69-BV421 (clone H1.2F3, Biolegend), anti-PD1-FITC (clone RMP-1, Invitrogen and clone J43, Invitrogen), anti-CTLA4-PE (clone UC10-4B9, Biolegend), anti-CD4-Alexa Fluor 700 (clone GK1.5, Biolegend), anti-IFNy-APC (clone XMG1.2, Biolegend), anti-TNFα-PE (clone MP6-XT22, Biolegend), H-2L(d) MuLV gp70 Tetramer SPSYVYHQF APC conjugated (NIH Tetramer Core Facility), anti-PD-L1-PerCP/Cy5.5 (clone 10F.9G2, Biolegend) anti-PD-L2-BV421 (clone TY25, Biolegend), anti-HU-HLA-APC (clone W6/32, Invitrogen), anti-H2Kb/H2DB-APC (clone 28-8-6, Biolegnd).

Statistical analysis and graphical design

Sample size for animal studies were guided by previous study in our laboratory in which similar or identical liver tumor models were used. For all models, randomization and blinding were performed. Statistical analysis was performed with GraphPad Prism 8 (GraphPad Software). Significance of the difference between groups was calculated by Student's unpaired t-test, one-way ANOVA (Tukey's and Bonferroni's multiple comparison test). P<0.05 was considered as statistically significant.

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