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Personal history of infections and immunotherapy: Unexpected links and possible therapeutic opportunities

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Abstract
The recent breakthroughs in the understanding of tumor immune biology have given rise to a new generation of immunotherapies, harnessing the immune system to eliminate tumors. As the typology and frequency of encountered infections are susceptible to shape the immune system, it could also impact the efficiency of immunotherapy. In this review, we report evidences for an indirect link between personal history of infection and different strategies of immunotherapy. In the current context of interest rise for personalized medicine, we discuss the potential medical applications of considering personal history of infection to design immunotherapeutic strategies.

Keywords
Immunotherapy; Infections; Personalized medicine; Cancer vaccine; Immune checkpoints; Adoptive cell transfe
Introduction

Genomic instability, common to all cancers,\(^1\) is a challenge for current cytotoxic therapies and represents the main cause of therapy failure and cancer relapse. Indeed, treatments generally target the dominant clones within the tumor and result in massive cell death. Nevertheless, it may paradoxically provide a selective pressure for the expansion of variant cells resisting to the drug used.\(^2\) Combinations of chemotherapies have been tested, and have shown encouraging results, but only delay emergence of resistance and rarely avoid it.\(^3\) This is why the research on cancer treatment has evolved from relatively nonspecific cytotoxic agents to selective, mechanism-based therapeutics. These ‘targeted therapies’ have been designed to interfere with specific molecular targets that are known to have a critical role in tumor growth or progression.\(^4\) For instance, treatments based on VEGF have increased the time before the cancer progresses as well as improving survival for patient with renal cell carcinoma.\(^5\)

Another key point in the recent evolution of cancer therapies is the observation that cancerous patients can be stabilized over metronomic treatments (chemo-, radio- or targeted therapies) through the induction of a long-term anti-cancer immune response or vaccine-like effect.\(^6\) To this extent, the recent breakthroughs in the understanding of tumor immune biology have given rise to a new generation of treatments which attempts to harness the exquisite power and specificity of the immune system for the treatment of malignancy. This cancer immunotherapy could allow overcoming immune tolerance to tumors and the production of adaptive anti-cancer responses able to deal with genomic instability.\(^7\)

Indeed, since the beginning of the 20th century, it has been hypothesized that immune system may play a role in cancer surveillance and prevention. The pioneering works of Dr William B. Coley have showed that stimulation of immune system by infections could be associated with tumor growth control.\(^8\) However, direct evidence for immunosurveillance of human cancers has been hard to prove experimentally and immunotherapy has mainly consisted in passive transfer of anti-cancer antibodies and T cells from donor to induce an immediate immune reaction against cancer.\(^9\) Recently, the clinical success of new immunotherapies, such as immune checkpoint inhibitors leading to complete tumor elimination in some patients with advanced cancers,\(^10\) supports the idea that immune system is a powerful weapon against cancer. Nevertheless, the successes of immunotherapy are extremely heterogeneous, sometimes associated with dramatic side-effects,\(^11\) and resistance has already been reported.\(^12\) Thus, identification of predictive biomarkers allowing selection of patients predisposed to respond to immunotherapy remains a challenge for immunologists and oncologists.

In addition to immnosurveillance of cancer, a prime function of immune system is the defense against infectious agents (i.e., viruses, bacteria, fungi, protozoans and metazoans that exploit other organisms, called hosts, to complete their life cycle). At an individual scale, humans are exposed to a high number of infections (through contact, ingestion or inhalation among other possibilities) which sometimes result in infraclinical symptoms.\(^13\) The community of organisms which have infected an individual during its life represents the personal history of infection. As cancer cells and infections interact with organisms’ defense mechanisms, typology and frequency of encountered infections are susceptible to shape the functioning of immune system, and consequently the efficiency of immunotherapy. Here, we review the indirect evidences suggesting that personal history of infections may impact, in a non-negligible way, many
mechanisms targeted by immunotherapeutic strategies and therefore may explain their contrasting successes as well as open new opportunities to improve them.

**Strategies envisioned for immunotherapy**

Today, several categories of immunotherapy exist which differ according to their target. First, we can distinguish immunotherapies designed to directly target tumor cells but which rarely result in a long-lasting anti-tumor immune response. It includes anti-tumor antibodies as well as cellular therapy, also known as adoptive cell transfer, aiming to select and/or to engineer tumor-specific immune cells that are thereafter administrated by local injection (i.e., intratumoral). A second category includes immunotherapies that stimulate the patient own immune system to elicit cancer cells elimination and an immune memory protective against tumor-associated antigens (TAAs). They are generally administrated by intravenous injection to generate a systemic immune response. They include antibodies directed against immune-modulating molecules present on immune cells such as CTLA-4, PD-1 and PDL-1, PDL-1 being also expressed by tumor cells themselves. At the interface between these two strategies, oncolytic therapies are based on viruses that show a tropism for cancer cells. This cytotoxic therapy stimulates both cancer cell destruction as well as anti-tumor immune response. Finally, immunotherapy also includes strategies based on immune-mechanisms, such as vaccines that either treat existing cancer or prevent development of a cancer.

**Cellular therapy**

**CAR T cells**

Adoptive transfer of T cells engineered to express chimeric antigen receptor (CAR) has emerged as a powerful immunotherapeutic strategy and is now the focus of numerous clinical trials. Especially, CAR-modified T cells targeting the B cell specific antigen CD19 have shown great promise in the treatment of chronic lymphocytic leukemia (CLL) and acute lymphocytic leukemia (ALL). The most notable toxicity of this treatment is the cytokine release syndrome (CRS), which is a systemic inflammatory process resulting from attack by CAR T cells of non-malignant cells that express targeted tumor antigen. In some patients, CD19-targeted T cells therapy has also been associated to neurological disturbances. A study has shown that CAR T cells are able to penetrate blood-brain barrier and they have been observed in the cerebrospinal fluid of the patient experiencing neurological toxicities. We can expect that neurological complications may be more frequent in patients with a high level of B cell infiltration in the brain. Indeed, the immune response produced by CAR T cells recognizing CD19 B cells could be associated with cytotoxicity to the surrounding tissues.

In addition to the natural variability between individuals, the level of cerebral infiltration by B cells could also be the result of past infections. Notably, acute malaria infection has been associated to a significant increase of CD19 B lymphocyte percentages in the brain of infected animals in a rodent malaria model. This increase is susceptible to protect against cerebral malaria, which is a frequent and deadly complication in children. It may be worth mentioning that certain viral brain infections (i.e., cytomegalovirus (CMV), West Nile Virus) may also be associated with an infiltration of immune cells and especially immune-regulatory B cells in the
brain, as suggested by an experimental study on mice.\textsuperscript{22} However, to our knowledge, the persistence of B cells within the brain after recovery of infection has not been investigated. If this was confirmed, neurological complications with CD19-targeted CAR T cells could be more common in patients with an history of malaria or such viral infections.

Another approach to generate CAR T cells is to use natural virus-specific T cells that have been modified to express antitumor CAR. These techniques generally lay on the dual-specificity of CAR T cells for infected cells and for tumor cells. In this manner, clonal expansion resulting from response to infection may benefit to tumor cell control and elimination. For example, EBV-specific T cells have been transduced with tumor-specific chimeric receptors that allow the recognition of G\textsubscript{D2}, a TAA present on several tumors of neural crest origin.\textsuperscript{23} The authors have observed that EBV-specific CAR T cells recognize EBV-infected targets through their conventional T-cell receptors and tumor targets through their chimeric receptors.\textsuperscript{24} Consequently, they are maintained by the presence of EBV-infected B cells and induce a higher anti-tumor activity which has increased survival in two mice tumor models: neuroblastoma and Hodgkin diseases. Influenza and CMV have also been used to generate dual-specific CAR T cells\textsuperscript{25,26} but their efficiencies in presence of the virus have not been tested.

We would like to propose here that the establishment of personal history of infection (especially by viruses and P. falciparum), before treatment administration, may help to select patients in order to prevent the apparition of dramatic adverse-effect, such as those observed with CD19-targeted CAR T cells. In addition, we can suspect that the use of virus-based CAR T cells in patients who have been already infected by the natural virus, could show a higher efficiency. For instance, EBV infects more than 80\% of the adult population.\textsuperscript{27} In these patients, the use of EBV-specific CAR T cells without simultaneous introduction of the virus, or with lower viral dose to avoid side-effects, could improve anti-tumor responses (Fig. 1). The generation of a cocktail of dual-specific CAR T cells, specific to persisting pathogens (EBV, CMV, herpesvirus) and engineered to target a panel of common TAAs, could improve immunotherapy efficiency and raises new perspectives for prevention.

**TILs**

Given the lack of well-defined TAAs for most human tumors, the use of naturally occurring tumor infiltrating lymphocytes (TILs) represents a good alternative to CAR T cells. In fact, TILs are able to recognize multiple TAAs that increase their ability to keep tumor growth in check. Nevertheless, insufficient migration\textsuperscript{28} and decreased anti-tumor function in the immunosuppressive tumor microenvironment\textsuperscript{29} are significant limitations to treat cancer patients with adoptive transfer of ex-vivo expanded TILs. However, the identification of cytotoxic tumor infiltrating T cells that recognize clonal mutations, shared by all tumor cells, hold promise for adoptive cellular therapy strategies in combination with the appropriate immunomodulatory agent.\textsuperscript{30}

Recently, tumor-specific T cells from patients have been engineered to carry a second T cell receptor (TCR) recognizing antigens from the bacteria Listeria monocytogenes.\textsuperscript{31} These polyclonal dual-specific T cells show a greater antitumor effect in presence of specific bacterial antigens, which are simultaneously injected at the tumor site. Indeed, the presence of bacterial antigens allows a greater clonal expansion of dual-specific T cells and decreases
immunosuppression in tumor micro-environment. However, patients who have already been exposed to the bacteria, or a closely related one, could show a stronger immune response against these antigens. This may result in a quick elimination of the bacterial antigens, neutralizing the positive impact on clonal expression and thus decreasing therapy efficiency. But, the consequences of this pre-exposition in term of treatment efficacy have not been investigated. Thus, within the context of personalized medicine rise, the choice of an additional infectious agent in adoptive TILs transfer should rely on the personal history of infections of the patient in order to maximize therapy efficiency.

Immune checkpoint inhibitors

**Anti-CTLA-4**

Cytotoxic T-lymphocyte–associated antigen 4 (CTLA-4) is a negative regulator of T cell activation. CTLA-4 acts by competing with the co-stimulatory molecule CD28 for binding to shared ligands CD80 and CD86 expressed by antigen-presenting cells. Antibodies that block CTLA-4, such as the ipilimumab, are known to increase T-cell activation and proliferation. Therefore, blocking CTLA-4 has the potential to enhance antitumor immune responses. Treatment with ipilimumab can cause significant tumor response in patients with metastatic melanoma. However, complete response has rarely been observed and immune-related adverse events, related to tolerance break and consequent auto-immunity, happen at high frequency.

Regarding the interactions between infections and CTLA4, a study has shown that the expression of CTLA-4 was depressed in peripheral blood mononuclear cells from tuberculosis patients, suggesting that people exposed to Mycobacterium tuberculosis could be more responsive to such immunotherapy. Nowadays, up to one-third of the world’s population is infected with M. tuberculosis and the majority of infected persons have a clinically latent infection. Even if the prevalence of tuberculosis is expected to be low in developed countries, where such immunotherapeutic strategy is available, this mechanism could be mimicked by TB vaccination (Bacillus Calmette-Guerin (BCG) is routinely administrated in many countries) or applied to infection with other mycobacteria. Interestingly, intravesical injection of BCG vaccine is approved for the treatment of non-invasive bladder cancer and its efficiency has been tested in combination with antiCTLA-4 antibodies in a murine model. It has been observed that mice treated with BCG and anti-CTLA-4 show an improved survival than mice receiving BCG only. Therefore, the use of ipilimumab in patients with a history of mycobacterial infection, where the target of ipilimumab is already under-expressed, could block the immunosuppressive action of remaining molecules and could therefore result in stronger immune activation (Fig. 2A). Thus, it would be intriguing to evaluate in clinical trials whether tuberculosis/vaccination status might have an impact on response to anti-CTLA-4 but also whether it would allow diminishing the dose of administration to avoid immuneregulation of immune system. Expression of PD-1, signed related adverse events (Fig. 2B).
Anti-PD-1 & anti-PD-L1
Programmed death 1 (PD-1) is a key immune-checkpoint receptor expressed by activated T cells that mediates downregulation of immune system. Expression of PD-1, signed the “exhausted phenotype” of effector T cells, has been first identified through studies of chronic viral infections in mice in which PD-1/PD-L1 was found to be an important negative regulatory feedback loop to ensure immune homeostasis. Cancer immunotherapies based on the administration of anti-PD-1 (such as nivolumab and pembrolizumab) or anti-PD-L1 (such as atezolizumab, avelumab or durvalumab) aim to inhibit the interaction between PD-L1, expressed by tumor cells but also by stromal cells, and PD1, expressed by T cells, and thus restore T-cell antitumor activity. Many investigators have addressed the question of whether the expression of PD-1 or PD-L1 might be a prognostic marker of treatments with immune checkpoint inhibitors. Although it is still a matter of debate for PD-L1, PD-1 expression within the tumor reflecting the presence of specific anti-tumor lymphocytes has been associated with a positive prognostic feature in many cancers. Importantly, the basal level of PD-1 expression by T cells could be shaped by personal history of infection. As mentioned above, the increase of PD-1 expression on antigen-experienced effector T cells during chronic infections, is a physiological mechanism to constrain the adaptive immune response and prevent auto-immunity. For instance, HIV infection has been associated with an increase of PD-1 expression on NK cells. The alteration of PD-1 expression in the context of acute infections has also attracted increasing attention. It has been reported that acute infections with M. tuberculosis and influenza virus result in a significant increase of PD-1 on NK cells and CD8 C T cells respectively. Moreover, evidences suggest that PD-1 expression could also be modified at long-term through immunological memory. In fact, memory CD8 C T cells excessively up-regulate PD-1 expression following vaccinia virus reinfection. Hence patients who have experienced a relatively high number of infections may also express high level of PD-1, in activated as well as memory immune cells, reflecting the immunosuppression of pre-existing T cell immunity. Contrarily to NK cells, CD8 C T cells are antigen-dependant and thus specific T cells could target cancer cells only if TAAs share similarities with infectious agents. This situation has already be suggested between influenza virus and lung cancer cells. In this context, we could expect that checkpoint inhibitors will present a lower efficacy in patients with a high basal level of PD-1. We also suggest that this effect may not be limited by clonal selection of cancer cells. In fact, cancer cells which harbor the ligand PD-L1 may have been favored because it allows them to escape immune destruction by the high number of immune cells expressing PD-1. If so, new strategies, aiming to increase immunotherapy efficiency and avoid dangerous side-effects, should consider adapting the dose of treatments with antagonists of the PD-1/PDL-1 interaction, according to the past history of infections of each patient.

Tumor-targeting monoclonal antibodies

Although developed to directly target malignant cells, it is now recognized that the therapeutic activity of tumor antigen-targeting monoclonal antibodies (mAb) is associated to their capacity to mobilize, through FcγR-dependent mechanisms, both innate and adaptive immune cells. However, tumor escape from mAb-induced tumor immune surveillance remains one of the
main clinical issues. As an example, anti-HER2/ErbB2 mAb (Trastuzumab) aims to reduce cell proliferation by blocking a receptor (ErbB2) of a growth factor highly expressed by many cancer cells. This therapy has substantially improved outcome of HER2-positive breast cancer and, given the importance of immune responses in trastuzumab therapy, therapy combined with immune stimulating agents have been used with success.

Recently, co-injection of polyinosinic-polycytidylic acid (PolyI:C) and CpG oligodeoxynucleotides (CpG) with trastuzumab shows higher efficiency than anti-ErbB2 therapy used alone. The authors suggest that these molecules may represent pertinent adjuvant stimulating innate responses including interferon expression which is required in trastuzumab therapy. PolyI:C is a synthetic viral-like double-stranded RNA (dsRNA) which bind to the innate immunity receptors TLR3, RIG-I and MDA-5. It has been used to mimic activation of dendritic cells by viral infections as most of viruses produce dsRNA at some point in their replication. CpG, a ligand for TLR9, mimics the presence of bacterial DNA and is also considered as a pathogen-associated molecular pattern that activates DC and B cells in particular.

Thus, whether a history of viral and/or bacterial infections could mimic the action of these molecules, and especially the increase expression of interferons, remains an interesting hypothesis. In accordance with this idea, it has been shown that interferon (a and g) levels were significantly higher in patients with chronic hepatitis B infection but also in patients who cleared the virus. Thus, one can suggest that personal history of hepatitis B infection might naturally boost anti-tumor responses in patients who received anti-HER2 therapy. In addition, over the last few years, an increasing number of works uncovered that innate immune cells (including monocytes, macrophages, and natural killer cells) display prolonged changes in their functionalities after infections or vaccination therapies, a phenomenon called “trained immunity”. According to this idea, innate immune cells could harbor an increased production of cytokines and other inflammatory mediators, including interferons, after recent infections. The increase in innate immune cells responsiveness may potentiate the adjuvant effect and thus the efficiency of anti-tumor targeting therapies.

**Oncolytic viruses**

Oncolytic viruses have emerged as a novel class of immune-based cancer therapy through their natural preference for cancerous cells and their capacity to trigger lysis of tumor cells as well as activation of anti-tumor immunity. Advances in molecular biology have also allowed the modification of other viruses to make them specific to neoplastic tissues and/or to combine them with immunogenic chemotherapy or immune checkpoint blockers to break tumor-induced immune tolerance. For instance, recombinant measles viruses have been intravenously injected to treat human patients with bone-marrow cancer. As for any non-oncolytic infections, the strength of the immune response to oncolytic viruses may be dependent of previous exposition. For instance, treatment based on the recombinant measles viruses led to a significant resolution of tumor only in the two patients who were measles-seronegative, suggesting that absence of prior exposure to measles virus has allowed
mounting an immune response strong enough to eliminate cancerous cells. This highlights that personal infection history could be crucial to identify which oncolytic agents can be envisioned in these strategies.

In addition, a genetically engineered herpes simplex virus, called T-VEC, has been recently approved to treat advanced melanoma. The treatment is administrated by intra-tumoral injection to seronegative patients. However, 67% of the population (0–49 years) had current herpes simplex virus-1 infection in 2012 (WHO). Thus, the proportion of patients who could receive the treatment could be lower than expected. Such considerations could be applied to other oncolytic therapy such as poxvirus vaccine in the treatment of prostate cancer or BCG therapy for bladder cancer. Thus, it suggests the necessity to investigate the potential of less common virus to design new oncolytic therapies.

**Prevention strategies**

**Prevention with infectious vaccines**

Several evidences show that vaccination against specific infectious agents could be used to prevent cancer. Protection against melanoma, lymphoma or leukemia has been reported 10 years after BCG, vaccinia or yellow fever vaccination. These findings might be explained by non-specific effects of these vaccines through the shifting of the immune response towards a Th1 profile or through cross reactivity. Indeed, vaccines may contain pathogen antigens with amino-acid sequences that are homologous with those of certain TAAs. By this cross-reactive effect, vaccination allows eliminating malignant cells as soon as they expressed the shared-antigens. For instance, a prior immunization with BCG vaccine, which has antigenic similarity with human endogenous retroviruses (HERV-KMEL) expressed in 95% of malignant melanocytes, has been associated with a better survival in patients with melanoma. Such considerations highlight the importance to consider vaccination as a component of personal history of infection and to study their potential for immunotherapeutic prevention. However, such studies may appear hardly conceivable because of methodological constrains such as cohort size and follow-up duration. Meta-analyses of several cohort studies and systematic reviews should nevertheless be considered to circumvent these limitations.

**Preventive cancer vaccines**

The potential of TAAs as targets for immunotherapy has already been largely investigated. Recently, it has been suggested that these antigens could be also candidates for preventive vaccines, which could hamper initiation of cancer cell proliferation. However, as it seems to be the case for therapy, the response to cancer vaccines could be conditioned by past history of infection. Here, we report examples of interaction between immune responses to infections and to cancer associated antigens.

**Mumps and MUC1 vaccine**

MUC1 is a normal epithelial cell antigen that was characterized as a TAA due to its abnormal expression in most epithelial adenocarcinomas, and the presence of anti-MUC1 antibodies in
patients with cancer has already been associated with a more favorable prognosis. This suggests that pre-existing antiMUC1 humoral response could help to control tumor growth or even prevent cancer proliferation. Interestingly, it has been described that mumps parotitis causes abnormal MUC1 expression on salivary gland and a negative correlation has early been observed between ovarian cancer and past history of mumps infection. Taken together, it has suggested that generation of anti-MUC1 immunity during mumps infection might protect against cancer development. Reinforcing such hypothesis, it has been reported that women who have experienced childhood mumps show a higher proportion of antiMUC1 antibodies which could increase humoral responses and immune recognition of MUC1. Accordingly, MUC1 has been studied as a target for antibody-based immunotherapies, some of which have been able to induce the elimination of ovarian cancer cells which express MUC1.

Multiple MUC1 vaccines for treatment of ovarian cancer are now in development and have given significant results. The elaboration of prophylactic vaccines has also been proposed and tested in a murine model of colon adenoma where it elicits a long term memory. However, mumps is a very common childhood disease suggesting that a large proportion of the population could have been exposed. We proposed that the consequences of a pre-existing immune memory developed against the infectious antigens should be investigated in the case of vaccines based on similar antigens. Indeed, it may help to better adapt the number of vaccine boosts which are necessary to reestablish a potent anti-tumor immune environment.

Varicella and vaccine against cyclin B1
Several tumors express high level of cyclin B1 such as lung, breast and prostate cancers, and it has been demonstrated that cyclin B1-specific immune responses may be important in cancer immunosurveillance. Indeed, a prophylactic vaccine against cyclin B1 can delay spontaneous cyclin B1C tumor growth in a murine lymphoma model and increases median survival. Interestingly, aberrant expression of cyclin B1-specific antibodies has also been reported in healthy individuals and infections seem to be a reasonably good candidate to explain the presence of cyclin B1-specific responses in healthy individuals. Indeed, a study has shown that cyclins are highly induced in varicella-zoster virus (VZV) infected cells. After the primary infection, Varicella-Zoster virus is able to establish a latent infection in the nervous system. The immune memory against the virus can persist for more than 20 years after the primo-infection and immune responses can be subsequently boosted either by endogenous re-exposure (silent reactivation of latent virus) or exogenous re-exposure (environmental). Thus, one can speculate that immune memory against VZV infection could boost immunosurveillance of cancer cells but could also interfere with the efficiency of cycline B1-based vaccines. In this context, the number of vaccine boosts should be determined in accordance with past history of VZV infection.

Personal history of infection and vaccine elaboration
There is a growing body of evidence showing that infections, both in childhood and adulthood, can provide a protection against the risk of developing malignancies. For example, exposures
to childhood diseases have been observed to reduce the risk of developing multiple cancers, including melanoma.  

Among childhood diseases, a particular attention has been paid to measles virus. It has been observed that specific MHC class I self peptide (Hsp90b<sub>570-578</sub>) are expressed both on melanoma cells and on measles-infected cells. Thus, the autoreactive CD8<sup>+</sup>T cells generated after measles infection could persist and enhance immunosurveillance of cancer cells, giving another example of cross-reactivity. During adulthood, history of acute infections (i.e., common colds and influenza) has been associated with a decreased of cancer risk in control-case study. This has been experimentally confirmed by a study showing that influenza infected mice harbor smaller lung tumor than naïve mice. They have also reported the overexpression of 5 TAAs (GAPDH, H4, MDH2, A2) both in infected cells and in tumor cells.

These evidences suggest that immune memory for antigens produced during influenza or measles infection has allowed the production of long-lasting antibodies against these specific TAAs which could be involved in cancer cell elimination. From these two examples, it seems that therapeutic potential of molecules, such as Hsp90b<sub>570-578</sub> and GAPDH, should be investigated. However, studies should not under-estimate the effects of previous natural infections, especially in the case of recurrent infections of fast-evolving virus (i.e., influenza virus), on vaccine efficiency.

Conclusions

Even if accumulating evidence suggests that infections may play a role in cancer risk, their influence in the responses to immunotherapy has been rarely investigated. Here, we report the evidences for indirect links between personal history of infections and different immunotherapeutic strategies. First, infections could be associated with a modified expression of the therapeutic target, such as increase of PD-1 following acute infections. Second, pre-exposition to additional infectious agents (as latent EBV infection in the case of EBV-specific CAR T cells) could increase cancer therapy efficiency. Finally, nononcogenic pathogens may induce long-lasting immunity against antigens shared with cancerous cells (i.e., mumps and antigens associated with ovarian cancer) and thus modify immunotherapy efficiency. Some immunotherapeutic strategies already mimic or rely on the effect of infection on immune system. It confirms that approaches, linking immunotherapy and personal history of infection, should be developed in a greater extent to consider the huge diversity of infectious agents, chronic but also acute, that may play a role.

While the potential interactions of infectious agents with immunotherapies cited here need to be experimentally tested and clarified, identifying the links between personal history of infection and immunotherapy could help clinicians to personalize their immunotherapeutic strategies, improving them or even envisioning new ones. First, it may help to adapt therapies through the determination of an optimal dose of administration. In addition, it may orient the clinicians’ choice toward the most adapted strategy, in accordance with the patient’s immune profile. Second, it could bring crucial insights to understand the diversity of responses to immunotherapy in cancer patients but also to forecast the apparition of adverse effects which
could be crucial to design safe treatments. Third, considering personal history of infection could help to design new oncolytic therapies based on less common viruses as it is expected that they will generate stronger immune responses. Finally, it may allow the elaboration of new preventive measures based on the knowledge of immune mechanisms involved in response to infections. It is worth mentioning that the role of vaccines to induce some of the immune mechanisms described in the context natural infections should be investigated as they represent a good opportunity of prevention. Retrospective analyses of data coming from immunotherapy trials could already give crucial clues to figure out the contribution of personal history of infection and vaccination on the heterogeneity of outcomes observed after the treatments.

**Competing interest**
The authors declare no competing interests.

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Figure 1
Influence of latent EBV infection on treatment with dual-specific CAR T cells. Latent and introduced virus could both be recognized by T-cell receptors and could increase CAR T cell clonal expansion. Thus, in patient with latent viral infection, the use of EBV-specific CAR T cells could result in stronger anti-tumor responses even without introduced EBV. TAA: tumor-associated antigens.
Figure 2

TB – CTLA-4 interaction hypothesis. (A) In patients with BCG vaccination or past history of tuberculosis (TB), CTLA-4 expression is down-regulated. For the same dose of ipilimumab (or other anti-CTLA-4 drugs), these patients will better respond to the treatment as more lymphocytes T (LT) will be targeted. (B) Proposition of clinical trial design to measure the impact of TB pre-exposure or BCG vaccination on treatment efficiency.
A

Patient without BCG vaccination or history of TB

Normal expression of CTLA-4
Variable responses to anti-CTLA-4

Patient with BCG vaccination or history of TB

Lower global CTLA-4 expression
Enhanced anti-tumor immunity

- LT expressing CTLA-4
- LT with depressed CTLA-4 expression
- Anti-CTLA-4 (ipilimumab)

B

Prospective study 1: Patients with ipilimumab 1\textsuperscript{st} lane

Impact of TB pre-exposure or BCG vaccination on response to treatment and toxicity

Prospective study 2: Patients with ipilimumab 1\textsuperscript{st} lane

Patients without TB pre-exposure or BCG treatment

Ipilimumab regular dosage

Patients with TB pre-exposure or BCG treatment

Ipilimumab regular dosage
Ipilimumab sub-optimal dosage

Comparison of treatment efficacy and toxicity