



Review article

Walking the tightrope: Balancing immune checkpoints and their inhibitors in host defence against bacterial infections

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Abstract

Immune checkpoints are one of the mechanisms that maintain the balance between immunotolerance and immunopathology. These mechanisms are often exploited by tumour cells. Thanks to extensive global testing of anticancer drugs, a revolutionary cancer therapy based on immune checkpoint inhibitors (ICIs) has been developed. Some pathogens exploit regulatory checkpoint interactions, contributing to the establishment of hidden, long-term, or persistent infections in which the host is unable to eliminate the pathogen. However, a structured overview of immune checkpoint involvement in bacterial infections remains underrepresented in the current scientific literature. We conducted a literature review focusing on the documented role of selected immune checkpoints (specifically PD-1, PD-L1, TIM-3, LAG-3, CTLA-4, and TIGIT) during infections of humans and animals with clinically relevant bacterial pathogens from the Actinobacteria, Chlamydiae, Firmicutes, Proteobacteria, and Spirochaetae phyla. The current state of knowledge suggests that applied research into immune checkpoints and the controlled use of ICIs has great potential to improve the diagnosis, treatment, and prognosis of serious human bacterial infections.

Keywords: Bacteria; ICIs; Immune checkpoint; Infection; Therapy

Highlights:

- Immune checkpoints play a significant role during bacterial infections.
- Pathogenic bacteria exploit immune checkpoints in order to evade host immunity.
- Checkpoint inhibitors can improve multiple aspects of infectious disease management.

Introduction

A fundamental property of the immune system is to correctly distinguish between self and non-self and to develop the appropriate immune response against invading pathogens. During infection, regulation of the inflammatory response is crucial to prevent excessive tissue damage due to host defence against pathogens. Prolonged immune activation is energetically demanding; timely control of the immune response and return to tissue homeostasis is beneficial for the host organism (Paludan et al., 2021). This delicate balance between tolerance and immunopathology is preserved by various pathways including anti-inflammatory cytokines, regulatory T cells, M2 macrophages and last but not least, by specific inhibitory immune checkpoints (IC), such as PD-1, PD-L1, CTLA-4, TIM-3, LAG-3, and TIGIT (Fig. 1) (Pauken and Wherry, 2015).

Programmed cell death protein 1 (**PD-1**; CD279) is a molecule that inhibits some lymphocyte functions, such as cytokine production, cell proliferation, and cytotoxicity. The interaction of PD-1 with its ligands PD-L1 (programmed

death-ligand 1; CD274) and PD-L2 (programmed death-ligand 2; CD273) is important in the control of peripheral T lymphocyte activation in infectious diseases, since it limits the response of subpopulations of effector cells to avoid tissue damage (Sampedro-Nunez et al., 2018). T cell immunoglobulin and mucin domain-containing protein 3 (**TIM-3**; CD366) can have dual and context-dependent effects on the immune system through multiple ligands comprising galectin 9, phosphatidylserine, HMGB1 alarmin, and CEACAM1 (Gorman and Colgan, 2014; Wolf et al., 2020). Lymphocyte-activation gene 3 (**LAG-3**; CD223) binds to its canonical ligand MHC class II with higher affinity than CD4, thus compromising CD4+ T cell activation (Huard et al., 1997). The coinhibitory molecule cytotoxic T-lymphocyte-associated protein 4 (**CTLA-4**; CD152) competes with the costimulatory molecule CD28 for the binding to CD80 and CD86 on antigen-presenting cells (Berg and Zavazava, 2008). T cell immunoreceptor with Ig and ITIM domains (TIGIT) regulates T cell and NK cell-based immunity via high-affinity binding to CD155 and lower-affinity binding to CD112 on T and NK cells (Tang et al., 2023).

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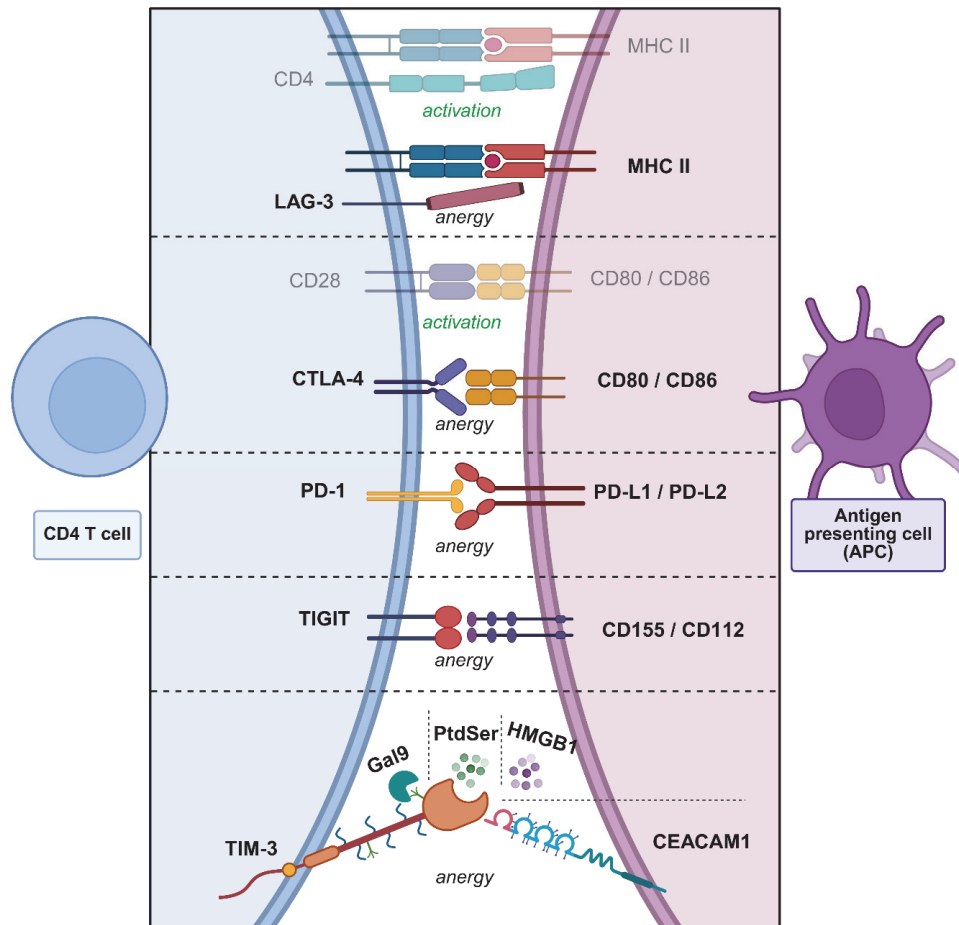


Fig. 1. Inhibitory immune checkpoints in immunological synapse. Interactions between key inhibitory immune checkpoints on a CD4 T cell and their ligands on an antigen presenting cell (APC) are highlighted in bold. T cell receptors such as LAG-3, CTLA-4, PD-1, TIGIT, and TIM-3 bind their corresponding ligands (MHC-II, CD80/CD86, PD L1/PD L2, CD155/CD112, and CEACAM1/HMGB1/PtdSer/Gal9) on the APC, leading to T cell anergy. Interactions resulting in cell activation, such as CD4–MHC II and CD28–CD80/CD86, are shown in grey and are diminished by the anergic effect. List of abbreviations: MHC II: major histocompatibility complex class II; LAG-3: lymphocyte-activation gene 3; CTLA-4: cytotoxic T-lymphocyte-associated protein 4; PD-1: programmed cell death protein 1, PD-L1: programmed death-ligand 1; PD-L2: programmed death-ligand 2; TIGIT: T cell immunoreceptor with Ig and ITIM domains; TIM-3: T cell immunoglobulin and mucin domain-containing protein 3; Gal9: galectin-9; PtdSer: phosphatidylserine; HMGB1: high-mobility group box 1 alarmin; CEACAM1: carcinoembryonic antigen-related cell adhesion molecule 1. Created in BioRender (Kovaříková, 2025).

Besides the traditional and well-described system of membrane-bound immune checkpoints and their ligands depicted in Fig. 1, soluble forms of these molecules also exist, produced through the alternative splicing of mRNA or the cleavage of membrane-bound checkpoint proteins. Soluble checkpoints can be present in body fluids, including plasma and cerebrospinal fluid, and also in extracellular vesicles. The functional significance of soluble immune checkpoints and their roles within both (patho)physiology and disease context have not yet been fully clarified (Joseph et al., 2022; Pitts et al., 2024).

Checkpoint molecules play an important role in two phases of the immune response: i) in immunological synapsis during antigen presentation where they prevent T cell hyperreactivity (Francisco et al., 2009); ii) during execution of immune cell effector functions in order to reduce collateral tissue damage. Tumour cells notoriously hijack these mechanisms to avoid the immune response and promote their own survival and progression. This insight led to the development of revolutionary cancer therapies based on immune checkpoint inhibitors

(ICIs). Currently, multiple ICIs have either been approved or are undergoing clinical trials for treatment of various cancer types (Lee et al., 2022). The impact of ICIs on survival and prognosis across cancers is significant. In advanced melanoma, the combination of nivolumab (PD-1 inhibitor) and ipilimumab (CTLA-4 inhibitor) has been shown to increase the median overall survival to 72 months, compared with the previously used chemotherapeutic dacarbazine, which achieved a median survival of only 9.1 months (Robert et al., 2011; Wolchok et al., 2022).

Proper regulation of checkpoint immune response is crucial for optimal coping with infections. While there has been a breakthrough in the targeting of immune checkpoints in cancer, increasing evidence indicates that some pathogens exploit inhibitory checkpoint interactions, contributing to the chronicity of the viral, bacterial, and parasitic infections. In this review, we focus on examples of how important the checkpoints are during the infection of host organism with selected clinically relevant bacterial pathogens, and simultaneously highlight their potential as therapeutic targets.

The role of inhibitory immune checkpoints in selected bacterial infections

Actinobacteria

Mycobacterium spp.

It is well established that multiple immune checkpoints, including PD-1/PD-L1, TIM-3, and LAG-3, play an important role in the immune response to mycobacterial infection. Chronic mycobacterial infection can progressively impair antigen-specific T cell responses, resulting in a phenotype of exhausted T cells and inversely increased mycobacterial burden. In infected mice, TIM-3 neutralization via antibody or genetic knockout effectively reduces bacilli burden and prolongs survival (Jayaraman et al., 2016). Pan et al. (2022) observed elevation of PD-L1 in lung tissue from patients with pulmonary tuberculosis. Sporadic case reports mention a positive effect of anti-PD-1 therapeutics in resolving mycobacterial infection (Ishii et al., 2018; Liu et al., 2023). LAG-3 is upregulated during mycobacterial infection in non-human primates and its overexpression is characteristic for both pulmonary and extrapulmonary tuberculosis (Pan et al., 2024; Phillips et al., 2017). TIGIT is downregulated on NK cells from patients with latent tuberculosis (TB) (Harris et al., 2020), whereas in active TB patients, TIGIT expression on CD8+ T cells is significantly upregulated and positively correlates with disease severity. Moreover, TIGIT blockade reduces lung bacterial burden in *M. tuberculosis*-infected mice (Zhou et al., 2025).

It seems that a certain amount of immune checkpoint activation is necessary for optimal control of mycobacterial infection, since disruption of PD-1/PD-L1 signalling in *in vivo* animal models and human 3D model leads to increased *M. tuberculosis* burdens and greater vulnerability of the hosts (Barber et al., 2011; Kauffman et al., 2021; Tezera et al., 2020). Indeed, many studies acknowledge the association between the use of ICIs and increased risk of acute mycobacterial infections or reactivation of latent tuberculosis; reviewed e. g. in Anand et al. (2020), and case reports continue to accumulate (Anidi et al., 2024; Inada et al., 2025; Yamaba et al., 2022).

In patients infected with *M. leprae*, increased expression of LAG-3 and PD-1/PD-L1 in skin lesions and peripheral blood cells contributes to the hyporesponsive immune profile, which is characteristic of leprosy patients (de Castro et al., 2025; Tarique et al., 2023).

Chlamydiae

Chlamydiaceae

Chlamydia infection represents a case in which the adaptive immune system is unable to protect against reinfection. This can be partially caused by type I interferon-induced upregulation of PD-L1 in the genital tract following *Chlamydia trachomatis* murine infection (Fankhauser and Starnbach, 2014; Reinhold-Larsson and Starnbach, 2025). On the other hand, an animal model showed that PD-L1 and TIM-3 signalling can reduce genital tract pathologies without significant suppression of immunity against *C. muridarum* infection (Peng et al., 2011). In the case of lung inflammation caused by *Chlamydomydia pneumoniae*, the early targeting of abundantly upregulated PD-L1 has no effect on pathogen load but prevents infection-induced airway hyperresponsiveness (Starkey et al., 2016). Finally, in *C. psittaci*-infected mice, TIM-3 expression is upregulated, and its blockade leads to more effective pathogen clearance, diminishes lung pathology, and improves Th1 and Th-17 mediated immune response (Li et al., 2023b).

Firmicutes

Streptococcus spp.

Multiple inhibitory immune checkpoint proteins, including PD-L1 and LAG-3, were upregulated at the maternal-fetal interface exposed to *Streptococcus agalactiae* experimental infection of *Macaca nemestrina* primates (Manuel et al., 2023). PD-1(-/-) mice, as well as wild-type mice treated with a PD-1 blocking antibody, exhibit significantly increased survival against lethal *Streptococcus pneumoniae* infection, since PD-1 expressed by B cells seems to play the central role in regulation of protection against *S. pneumoniae* (McKay et al., 2015). Moreover, blockade of the PD-1/PD-L1 pathway mitigates the symptoms of murine meningitis caused by *S. pneumoniae* (Ma et al., 2023).

Staphylococcus aureus

The PD-1/PD-L1 axis is strongly involved in the immune response to the common bacterium *Staphylococcus aureus* throughout life. Early in infancy, PD-L1 blockade during *S. aureus* stimulation reduces the induction of neonatal Tregs (Rabe et al., 2014). Likewise, both neonatal and adult CD4+ T cells upregulate PD-1 in response to *S. aureus* infection and the blockade enhances early *S. aureus*-initiated responses (Majer et al., 2023). Since the PD-1/PD-L1 axis contributes to immune dysfunction after burn injury, anti-PD-L1 treatment improves bacterial clearance and the survival of mice following post-burn systemic infection (Patil et al., 2018). Orthopaedic implant-associated *S. aureus* osteomyelitis is characterized by an immunosuppressive profile with upregulated PD-1 and PD-L1, which suppresses macrophage bactericidal function. PD-1/PD-L1 inhibitors not only reduce bacterial load in bone tissue but also promote healing and support memory T cells' development (Li et al., 2023a). Interestingly, a significant number of *S. aureus*-specific Tregs expressing TIGIT were found in healthy individuals, which may hinder vaccine responses (Clegg et al., 2024).

Listeria monocytogenes

The role of PD-1/PD-L1 axis during *Listeria monocytogenes* infection is unclear. Whereas PD-1 deficiency increases the resistance of mice to *L. monocytogenes* infection (Yao et al., 2009), some authors report a potential costimulatory effect of PD-L1 towards *Listeria* antigen-specific adaptive responses (Seo et al., 2008; Xu et al., 2013). However, TIM-3 helps *L. monocytogenes* to evade the immune system by inhibiting macrophage-mediated phagocytosis *in vitro* and MHC-I-restricted antigen presentation by murine macrophages *in vivo* (Wang et al., 2017b; 2020).

Proteobacteria

Brucella spp.

Chronic infection of mice with *Brucella melitensis* leads, among other effects, to CD8+ T cell exhaustion with upregulation of PD-1 and LAG-3 (Durward-Diioia et al., 2015). Human brucellosis is accompanied by the upregulation of PD-1 during both acute and chronic infection, and CTLA-4+ Tregs in chronic disease (Sun et al., 2021). Moreover, a vaccine containing CTLA-4 together with *Brucella* surface proteins had higher antigenicity and immunogenicity (Guo et al., 2023). Also, *Brucella* antigen Omp25 significantly upregulates PD-1 in monocytes/macrophages and negatively regulates IL-12. PD-1 blockade restores IL-12 production and promotes Th1 responses (Cui et al., 2017).

Rickettsiales

PD-1 influences immune responses in different *Rickettsia* infections: patients infected with *R. rickettsii* had a higher percentage of CD8+PD-1+ cells, and patients infected with *R. typhi* had a higher number of cells corresponding to CD4+PD-1+ cells (Dzul-Rosado et al., 2023). Also, patients infected with rickettsial agent, *Orientia tsusugamushi*, the causative agent of scrub typhus, significantly upregulate PD-1, LAG-3, and TIM-3 on circulating NKT cells, resulting in their functional impairment (Kang et al., 2018). On the other hand, the blocking or silencing of TIM-3 signalling reduces intracellular killing and thus boosts rickettsial burdens in both *in vitro* and *in vivo* models of early phase *R. heilongjiangensis* infection (Yang et al., 2016).

Enhanced expression of PD-1 and LAG-3 on T cells following *Anaplasma marginale* infection in cattle contributes to the exhaustion of specific T cells and potentially leads to the persistence of this rickettsia in organism (Okagawa et al., 2016).

Bordetella pertussis

In contrast to *Bordetella pertussis* strains from the era before the whole-cell pertussis vaccines were introduced, recently circulating strains of *B. pertussis* induce higher levels of PD-L1 in a TLR2-dependent manner on human monocyte-derived dendritic cells. Emerging *B. pertussis* strains have thus evolved to induce a more regulatory response, which can be a pathogen strategy to increase its spreading in highly vaccinated population (Hovingh et al., 2017).

Neisseria gonorrhoeae

Neisseria gonorrhoeae is able to suppress the protective immune response at different levels, including polarization of macrophages into regulatory M2b phenotype and upregulation of PD-L1 on macrophages and dendritic cells upon cocultivation with gonococci (Ortiz et al., 2015; Zhu et al., 2012).

Coxiella burnetii

During the acute phase of Q fever caused by *Coxiella burnetii*, PD-L1 expression on T cells is upregulated, resulting in a decreased number of memory T cells (Ka et al., 2015). In chronic Q fever endocarditis, overexpression of PD-1 is observed in intermediate monocytes and CD4+ T cells (Ka et al., 2014).

Pseudomonas aeruginosa

Infection with *Pseudomonas aeruginosa* is common during immunopathologic conditions (e.g., extensive injuries, surgical traumas, cystic fibrosis). Blockade of PD-1/PD-L1 axis protects experimental mice against *P. aeruginosa* infection after burn injury, fracture surgery, or exposure to air pollutants (Luo et al., 2024; Patil et al., 2018; Zhang et al., 2019). In case of *P. aeruginosa* colonization of cystic-fibrosis-affected respiratory tract, PD-1/PD-L1 blocking assays improved the immune response and T cell effectivity (Averdano-Ortiz et al., 2019). In contrast, TIM-3/galectin-9 signaling is beneficial for the neutrophil-mediated killing of *P. aeruginosa* (Vega-Carrascal et al., 2014).

Haemophilus spp.

In cases of *Haemophilus* colonization of the lower respiratory tract, PD-1 and CTLA-4 pathways seem to play a complex regulatory role in controlling this opportunistic pathogen: patients with chronic obstructive lung disease produce more PD-1 and CTLA-4 in the airways, which makes them more susceptible to *H. influenzae* infections (Kalathil et al., 2014). In concordance, anti-PD-L1 antibodies enhance antibody production and

improved the bacterial clearance of *H. influenzae* in a mouse model (Yoshinaga et al., 2025), and *Haemophilus* loads positively correlate with PD-L1 expression in mixed granulocytic childhood asthma (Kim et al., 2024). CTLA-4 positive Tregs contribute to the persistence of *H. ducreyi* genital ulcers (Li et al., 2010).

Salmonella spp.

According to current knowledge, multiple immune checkpoints play a significant role in the course of *Salmonella* infections. The infection of intestinal epithelial cells with *Salmonella enterica* serovar Typhimurium induces PD-L1 in both human and rodent cells, and this effect is potentiated by gamma interferon (Sahler et al., 2018). Also, *Salmonella*-infected B cells express more PD-L1, which leads to an impaired CTL response during persistent *Salmonella* infection (Lopez-Medina et al., 2015). PD-L1 upregulation is associated with *Salmonella* pathogenicity island 2 effector molecule SseL (Chopra et al., 2025). Effector responses during *Salmonella* Typhimurium infection are further inhibited by IL-10, originating from LAG-3-expressing the subpopulation of plasma cells (also co-expressing CD138, PD-L1 and PD-L2). This regulation leads to higher bacterial loads in murine tissues and the reduced survival of mice (Lino et al., 2018). Whereas TIM-3/galectin-9 interaction is important for the effective activation and bactericidal activity of intestinal macrophages during *S. Typhimurium* infection (Yu et al., 2018), TIM-3 antibody blockade restores Th1 responses dampened by iron overdose and improves bacterial control (Pfeifhofer-Obermair et al., 2021). Finally, antibody blockade of TIGIT in murine *in vivo* model is able to enhance immunity and improve the clearance of *S. Typhimurium* (McCulloch et al., 2024).

Helicobacter pylori

Infection with *Helicobacter pylori* is a well-established risk factor for the development of gastric cancer, where inhibitory immune checkpoints play a significant role. Whereas TIM-3 overexpression leads to reduced *H. pylori*-associated proinflammatory immune response in RAW264.7 macrophages *in vitro* (Wang et al., 2017a), the association between *H. pylori* and the PD-1/PD-L1 axis is in the spotlight, since *H. pylori* significantly increases PD-L1 expression in organoid cultures and gastric epithelium via Hedgehog signalling pathway, creating premalignant lesions (Holokai et al., 2019; Koh et al., 2021). Indeed, clearance of *H. pylori* is negatively affected by enhanced PD-L1 expression on gastric epithelial cells (Beswick et al., 2007; Das et al., 2006), and more importantly, *H. pylori* infection may reduce the efficacy of PD-1/PD-L1 blockade therapy of cancer (Oster et al., 2022; Shatila et al., 2025; Yang et al., 2025).

Spirochaetae

Spirochaetaceae

In patients infected with *Treponema pallidum*, a notably higher percentage of follicular helper T cells expressing PD-1 is present in the stage of latent syphilis, characterized by no clinical symptoms and high antibody titres against *Treponema pallidum* (Shen et al., 2023). The exposure of dendritic cells and macrophages to live *Borrelia burgdorferi* spirochetes drives the expression of immunoregulatory molecules, including PD-L1, LAG-3, and TIM-3 (Gutierrez-Hoffmann et al., 2023; Helble et al., 2022). PD-1 knockout mice showed no differences in bacterial loads but the absence of PD-1 led to increased T cell expansion and accumulation in joints and draining lymph nodes of infected mice (Helble et al., 2022).

An overview of the literature addressing the role of checkpoints in bacterial infections is presented in Table 1.

Table 1. Summary of the current knowledge about the involvement of immune checkpoints in the course of bacterial infections

Pathogen	PD-1/ PD-L1	TIM-3	LAG-3	CTLA-4	TIGIT	Reference
<i>Mycobacterium</i> spp.	•	•	•		•	Anand et al., 2020; Anidi et al., 2024; Barber et al., 2011; de Castro et al., 2025; Harris et al., 2020; Inada et al., 2025; Ishii et al., 2018; Jayaraman et al., 2016; Kauffman et al., 2021; Liu et al., 2023; Pan et al., 2022, 2024; Phillips et al., 2017; Tarique et al., 2023; Tezera et al., 2020; Yamaba et al., 2022; Zhou et al., 2025
Chlamydiaceae	•	•				Fankhauser and Starnbach, 2014; Li et al., 2023b; Peng et al., 2011; Reinhold-Larsson and Starnbach, 2025; Starkey et al., 2016
<i>Streptococcus</i> spp.	•		•			Ma et al., 2023; Manuel et al., 2023; McKay et al., 2015
<i>Staphylococcus aureus</i>	•				•	Clegg et al., 2024; Li et al., 2023a; Majer et al., 2023; Patil et al., 2018; Rabe et al., 2014
<i>Listeria monocytogenes</i>	•	•				Seo et al., 2008; Wang et al., 2017b, 2020; Xu et al., 2013; Yao et al., 2009
<i>Brucella</i> spp.	•	•		•		Cui et al., 2017; Durward-Diioia et al., 2015; Guo et al., 2023; Sun et al., 2021
Rickettsiales	•	•	•			Dzul-Rosado et al., 2023; Kang et al., 2018; Okagawa et al., 2016; Yang et al., 2016
<i>Bordetella pertussis</i>	•					Hovingh et al., 2017
<i>Neisseria gonorrhoeae</i>	•					Ortiz et al., 2015; Zhu et al., 2012
<i>Coxiella burnetii</i>	•					Ka et al., 2014, 2015
<i>Pseudomonas aeruginosa</i>	•	•				Avendano-Ortiz et al., 2019; Luo et al., 2024; Patil et al., 2018; Vega-Carrascal et al., 2014; Zhang et al., 2019
<i>Haemophilus</i> spp.	•			•		Kalathil et al., 2014; Kim et al., 2024; Li et al., 2010; Yoshinaga et al., 2025
<i>Salmonella</i> spp.	•	•	•		•	Chopra et al., 2025; McCulloch et al., 2024; Lino et al., 2018; Lopez-Medina et al., 2015; Pfeifhofer-Obermair et al., 2021; Sahler et al., 2018; Yu et al., 2018
<i>Helicobacter pylori</i>	•	•				Beswick et al., 2007; Das et al., 2006; Holokai et al., 2019; Koh et al., 2021; Oster et al., 2022; Shatila et al., 2025; Wang et al., 2017a; Yang et al., 2025
Spirochaetaceae	•	•	•			Gutierrez-Hoffmann et al., 2023; Helble et al., 2022; Shen et al., 2023

Discussion and future prospects

When exploring the relationship between bacteria colonizing the host and inhibitory immune checkpoints, it becomes evident that most existing literature focuses on their role in cancer immunotherapy and the interaction of ICIs with host microbiome. Indeed, gut microbiota has been identified as a key driver of ICIs efficacy against cancer (reviewed elsewhere, e.g., in Chen et al., 2025, Perl et al., 2025 and Vig and Dubey, 2025). Many studies have also explored how ICIs therapy may lead to bacteria-associated side effects, starting from *Clostridium difficile* infection during ICIs-induced colitis (Vuillamy et al., 2023) through pathogenic colonization of the genitourinary tract and skin (Ross et al., 2022), and ending with the reactivation of latent infections such as tuberculosis (Inada et al., 2025). Despite the extensive research on immune checkpoints in oncology, microbiome interactions, and experimental viral infections, their specific role in bacterial infections has received limited scientific attention.

To understand the potential role of immune checkpoints in bacterial infections, it is important to note that during the

coevolution of bacteria and their hosts, the pathogens have evolved several immunity-based strategies that enable them to evade host immunity and establish hidden, long-term, or persistent infection with the host unable to eliminate the pathogen. Persistent antigenic stimulation, observed both in cancer and chronic infections, can lead to progressive T cell exhaustion, characterized by diminished effector functions and the increased expression of inhibitory immune checkpoints. This fact suggests that immune checkpoints may play a crucial role in determining the course of certain bacterial infections and therefore opens the possibility of their therapeutic use in human medicine.

While antibiotics remain the primary treatment for bacterial infections, their effectiveness is not universal – particularly in chronic or biofilm-associated conditions. Identifying strategies to enhance antibiotic treatment effectivity is therefore beneficial. Although some studies have shown that concomitant use of antibiotics reduces the efficacy of ICIs antitumour therapy (Wu et al., 2021), recent evidence suggests that immune checkpoint inhibition in certain conditions may enhance antimicrobial efficacy. For instance, PD-1/PD-L1 blockade significantly improved outcomes in a murine model

of *Staphylococcus aureus* osteomyelitis, acting as a potent adjuvant to antibiotic therapy and promoting bacterial clearance (Li et al., 2023a).

Another promising application of immune checkpoint modulation lies in enhancing vaccine responsiveness. In some healthy individuals, immune responses may be suppressed due to natural background checkpoint expression that can dampen the activation and proliferation of immune cells, resulting in suboptimal immune response following vaccination (Clegg et al., 2024). This pre-exposure immunosuppression can potentially be overcome through the administration of immune checkpoint inhibitor prior to vaccination.

In the discussion of using ICIs as immunomodulatory tools in human medicine, it is necessary to consider the potential risks associated with off-target effects of conventional ICIs, given the broad functional roles that checkpoints play in the body. However, evidence suggests that certain immune checkpoints may be pathogen specific. For instance, CD84 has been identified as a putative tuberculosis-specific inhibitory receptor (Zheng et al., 2022). The discovery of new, infection-specific checkpoints may bring fresh impetus to the treatment of chronic infections.

The potential of immune checkpoints and ICIs extends beyond immune modulation. Soluble immune checkpoints, which are elevated during infectious states like tuberculosis and bacterial sepsis, correlate with common laboratory markers and treatment responses. Moreover, elevated serum concentrations of some soluble inhibitory immune checkpoints correlate with higher mortality in bacterial sepsis. Taken together, certain soluble immune checkpoints may serve as potential biomarkers for disease surveillance and prognosis (Chen et al., 2022; Mearelli et al., 2025).

Conclusion

Thanks to extensive global testing of anticancer drugs, we have a large amount of data on ICIs, which could also be transferred to other areas of human medicine – although we are aware of the challenges posed by the application of ICIs. In addition, there are many infections whose relationship to immune checkpoints remains poorly understood or unexplored. Taken together, modulation of immune response via immune checkpoints can be an emerging tool for the improvement of diagnosis, therapy, and determining the prognosis of important human infections. For this reason, we believe that future research should be directed towards this area.

Author contributions

H. Langhansová conceived and designed the study. All authors performed the literature search. The first draft of the manuscript was written by H. Langhansová and A. Palounková, the figure was designed by A. Kovaříková and A. Palounková. All authors edited several drafts of the manuscript and read and approved the final version.

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Ethical aspects and conflict of interest

The authors have no conflict of interest to declare.

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