# LEADING ARTICLE



# *Clostridium* Bacteria: Harnessing Tumour Necrosis for Targeted Gene Delivery

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# Abstract

Necrosis is a common feature of solid tumours that offers a unique opportunity for targeted cancer therapy as it is absent from normal healthy tissues. Tumour necrosis provides an ideal environment for germination of the anaerobic bacterium *Clostridium* from endospores, resulting in tumour-specific colonisation. Two main species, *Clostridium novvi*-NT and *Clostridium sporogenes*, are at the forefront of this therapy, showing promise in preclinical models. However, anti-tumour activity is modest when used as a single agent, encouraging development of *Clostridium* as a tumour-selective gene delivery system. Various methods, such as allele-coupled exchange and CRISPR-cas9 technology, can facilitate the genetic modification of *Clostridium*, allowing chromosomal integration of transgenes to ensure long-term stability of expression. Strains of Clostridium can be engineered to express prodrug-activating enzymes, resulting in the generation of active drug selectively in the tumour microenvironment (a concept termed Clostridium-directed enzyme prodrug therapy). More recently, Clostridium strains have been investigated in the context of cancer immunotherapy, either in combination with immune checkpoint inhibitors or with engineered strains expressing immunomodulatory molecules such as IL-2 and TNF- $\alpha$ . Localised expression of these molecules using tumour-targeting *Clostridium* strains has the potential to improve delivery and reduce systemic toxicity. In summary, *Clostridium* species represent a promising platform for cancer therapy, with potential for localised gene delivery and immunomodulation selectively within the tumour microenvironment. The ongoing clinical progress being made with C. novyi-NT, in addition to developments in genetic modification techniques and non-invasive imaging capabilities, are expected to further progress Clostridium as an option for cancer treatment.

## **Key Points**

*Clostridium* are a genus of anaerobic bacteria that can colonise tumour necrosis.

Strains of *Clostridium* can be engineered to express prodrug activating enzymes and immunomodulatory molecules selectively within the tumour microenvironment.

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# 1 Clostridium Species

*Clostridium* is a genus of Gram-positive obligate anaerobic bacteria [1]. In aerobic environments they exist as inert endospores, germinating into metabolically active vegetative cells in the absence of oxygen. This genus includes some well-known pathogenic species, such as *Clostridium tetani* and *Clostridium botulinum*, but most are generally non-pathogenic and reside in the subsoil [2], or the intestinal tract of animals and humans [3]. *Clostridium* species have been widely used in biotechnology to produce biofuels, such as ethanol [4] and butanol [5], but also have therapeutic applications due to their anaerobic nature. For example, areas of tumour necrosis are known to support the selective germination and growth of anaerobic bacteria such as *Clostridium* (Fig. 1) [6].

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# 2 The Use of Clostridium in Cancer Therapy

Areas of tumour necrosis are commonly observed in the core regions of solid tumours due to insufficient vascularisation and subsequent metabolic stresses such as hypoxia and nutrient deprivation. Tumour necrosis is associated with poor prognosis in a wide range of cancer types and can be prevalent at end-stage disease [7]. Being completely absent from normal healthy tissues, tumour necrosis ensures that germination and growth of *Clostridium* is targeted selectively to the tumour microenvironment. Two main species of *Clostridium* are frequently used in this context, namely *Clostridium novyi* and *Clostridium sporogenes*.

The most clinically advanced *Clostridium* species used for cancer therapy is *C. novyi*-NT, an attenuated variant of *C. novyi* with the lethal  $\alpha$ -toxin gene removed. *C. novyi*-NT was first identified as a promising species from a panel of 26 anaerobic bacteria that were screened for their ability to colonise and grow within necrotic tumours [8]. In preclinical mouse (CT26) and rabbit (VX2) models, intravenously injected endospores of *C. novyi*-NT were shown to germinate in necrotic tumour areas, producing complete responses in approximately 30% of treated animals [8,9]. This effect was immune-mediated, with cured animals developing long-term cellular immunity to the original tumours [9]. However, in a subsequent dose-escalation study in dogs with spontaneously occurring tumours, intravenous injection of C. novyi-NT endospores resulted in significant toxicities that necessitated the study to be discontinued [10]. This included anaemia, thought to be a result of tumour-related haemorrhage, and abscesses that required extensive surgical debridement. A follow-up study evaluated direct intratumoural injection, where 37.5% of treated dogs had objective responses (three complete and three partial responses in 16 animals) [11]. Based on the safety and efficacy observed in this study, a phase I clinical trial was initiated in patients with treatment-refractory solid tumours (NCT01924689). Radiological and clinical signs of tumour lysis were observed in 42% of treated patients with injected lesions, with stable disease being the best overall response [12]. However, this was associated with some significant toxicities, including sepsis and gas gangrene. All patients who experienced these dose-limiting toxicities eventually recovered.



**Fig. 1** Gram-twort staining of a CT26 mouse tumour colonised by *C. sporogenes* (purple rods), demonstrating the selectivity of germination to necrotic areas

Whilst promising, intratumoural injection of C. novvi-NT endospores limits the ability to target tumours beyond the reach of percutaneous injection. The use of C. sporogenes could allow for successful intravenous delivery of endospores. The first clinical trials for the use of C. sporogenes (then named C. butyricum M55) on cancer patients were initiated in the 1960s and 1970s [13,14]. In general, intravenous administration of endospores was well tolerated, with patients mainly experiencing fever and leukocytosis that was controllable by antibiotics [13]. Tumours in these patients frequently became liquified (indicative of oncolysis) and required drainage, but complete tumour regression was not observed [13]. More recently, C. sporogenes has been attenuated to enhance clinical safety, after identification of a nine gene cluster with high sequence similarity to the streptolysin S (SLS) operon of Streptococcus pyrogenes [15]. This haemolysis operon was deleted from the genome using CRISPR-cas9 technology, with the resulting strain (C. sporogenes-NT) now free from haemolytic activity and thus better positioned for successful clinical translation [16].

# 3 Clostridium as a Vector for Solid Tumours

Initially, *Clostridium* species demonstrated modest antitumour activity when used as a single agent. However, tumour regrowth eventually occurred from an outer rim of viable oxygenated cells where *Clostridium* could neither germinate nor penetrate. This led to the idea that engineered strains of *Clostridium* could be used as a tumour-selective gene delivery system [17]. In early studies, the enzyme

Fig. 2 Schematic representation of the key advantages in using Clostridium species as vectors for solid tumours. A Genomic autonomy minimises the risk of recombination with the host genome, B flagella provide motility to promote intratumoural spread, C genomic size allows for larger gene inserts at more than one loci, D control of proliferation is possible with generic antibiotics, and E Clostridium endospores are not immunogenic, allowing for repeated intravenous (IV) injections

cytosine deaminase (CD) was cloned into a clostridial expression vector and transformed into Clostridium beijerinckii [17]. CD can be used to convert the non-toxic prodrug 5-fluorocytosine (5-FC) into the active chemotherapy agent 5-fluorouracil (5-FU) [18]. Significant levels of CD in the bacterial supernatant of transformed C. beijerinckii cultures increased the sensitivity of tumour cells to 5-fluorocytosine by up to 500-fold, indicating successful expression and secretion of functional CD from C. beijerinckii [17]. Later, these findings were confirmed in an in vivo setting using a strain of C. beijerinckii engineered to express a nitroreductase enzyme from Escherichia coli, NfsB [19]. Nitroreductase activity was detectable in ten out of ten tumours for 5 days after intravenous injection of NfsB-expressing C. beijerinckii endospores, with no nitroreductase protein observed in normal tissues, demonstrating proof-of-principle for selective gene delivery to the tumour using Clostridium [19].

There are many advantages to using *Clostridium* species as a vector for solid tumours over other vector options (Fig. 2). For example, *Clostridium* endospores are not immunogenic [20], allowing for repeated systemic administration of endospores without neutralisation by circulating antibodies or the complement system. The ability to perform repeated intravenous injections of *Clostridium* endospores was first demonstrated in early human studies [14]. These findings were later confirmed in preclinical models, where multiple consecutive treatment cycles of *Clostridium* infection were possible in between periods of antibiotic treatment to clear the infection [21]. Another advantage to the use of *Clostridium* as a vector is their genomic size and autonomy.



Larger gene inserts at more than one loci are feasible, and infection and lysis of the host cells is not required for successful bacterial replication. As a consequence, there is minimal risk of recombination with the host genome and no loss of the transgene or any intracellular co-factors to the extracellular space. Germinated *Clostridium* are also flagellated, providing motility that can facilitate intratumoural spread. Importantly, control of proliferation is possible using antibiotics, either at the conclusion of treatment or if an adverse event occurs. Sensitivity to generic antibiotics such as metronidazole is an attractive safety feature that can help to alleviate safety concerns around the use of replicating biological vectors in humans.

# 4 Methods for Genetic Modification of *Clostridium*

To successfully develop *Clostridium* for use as a vector, methods that allow for construction of stable recombinant strains are essential. Initially, Clostridium species were successfully engineered to express genes from autonomous plasmids. Saccharolytic Clostridium strains such as C. beijerinckii were the first to be engineered for plasmid-based expression of various transgenes (covered in Sect. 3), but colonisation of tumours using these strains was low. Proteolytic Clostridium strains were much better at colonising tumour tissue but were initially overlooked due to their resistance to genetic modification. Eventually, a transformation method was developed for the proteolytic strain C. sporogenes that involved use of a polyethylene glycol based transfection buffer [22], but this method had low efficiency and reproducibility. Subsequent transformation protocols based on conjugation or electroporation were then developed for C. sporogenes that were highly successful [21,23], allowing for tumour-specific expression of a transgene following colonisation by Clostridium. However, plasmid-based expression of genes can result in segregational instability, in which the proportion of plasmid-containing cells within a population decreases over multiple cycles of replication [24,25].

More recently, a reliable method for the stable integration of expression constructs into the chromosome of *Clostridium* was developed, termed Allele-Coupled Exchange (ACE) [26]. ACE is based on a double-crossover homologous recombination event that transfers a region of a plasmid into a specific site in the *Clostridium* genome. Each gene sequence is usually codon optimised to ensure efficient expression in the *Clostridium* host. Formation of the desirable double-crossover cells is coupled to an associated selectable phenotype that allows these cells to be isolated. *Clostridium* strains generated using ACE have the transgene inserted into the *pyrE* locus. The *pyrE* gene is essential for Clostridium to synthesise uracil, a nucleobase that plays a role in the manufacture of proteins. By inactivating *pyrE* during integration, recombinant strains are rendered uracil auxotrophs, disabled from undesirable growth in the environment. Consequently, ACE does not require the use of antibiotic resistance markers. In addition, integration of the transgene into the chromosome provides complete segregational stability with no mechanism for horizontal gene transfer. Importantly, ACE allows for generation of Clostridium endospore preparations that are compatible with clinical regulatory requirements. Several recombinant strains of C. sporogenes have been generated using ACE technology, with functional expression of the desired transgene (a bacterial nitroreductase enzyme from Neisseria meningitides, NmeNTR) demonstrated [27,28]. However, ACE can be time consuming and large gene inserts ( $\geq 28$  kb) can decrease transformation efficiency [26].

The development of CRISPR-Cas9 genome editing technology has revolutionised the ability to engineer recombinant strains of Clostridium [29,30]. CRISPR-Cas9 allows genetic material to be added, modified, or deleted at precise locations in the genome using short 'guide' sequences of RNA that direct the Cas nuclease to its target. Here, doublestranded breaks are created in the targeted DNA sequence and one of the two main DNA repair pathways can then be exploited to render genes non-functional (non-homologous end joining) or to insert new genes (homology-directed repair). CRISPR-Cas9 has the advantage of being simple, quick and cheap to perform in *Clostridium* compared with other techniques. It has been performed successfully in both C. novyi-NT (genomic modification) [31] and C. sporogenes (gene insertion and gene deletion) [16,32], although it appears that design of optimal constructs and guide RNAs is strain-specific. The use of CRISPR-Cas9 is not without its challenges, however. Expression of Cas9 in bacteria can be toxic when combined with guide RNAs that have specific motifs at their protospacer adjacent motif (PAM)-proximal end [33]. These Cas9 complexes have been shown to bind at off-target positions within the bacterial chromosome, silencing essential genes and causing toxicity [34]. Alternative endonucleases (e.g. Cas12a, also known as Cpf1) have since been developed to overcome limitations of Cas9 and may prove more advantageous for genomic modification of *Clostridium* in future [35–37].

# 5 *Clostridium*-Directed Enzyme Prodrug Therapy

*Clostridium* has been used as a tumour-selective gene delivery vector primarily in the context of *Clostridium*-directed enzyme prodrug therapy (CDEPT). CDEPT is a cancer gene therapy approach whereby an exogenous

therapeutic enzyme expressed by *Clostridium* can metabolise otherwise inert prodrugs into active, cytotoxic metabolites (Fig. 3). Localised activation of prodrug in this way can create an improved therapeutic index and, in principle, produce continuous active drug concentrations within the tumour microenvironment. Depending on the tissue penetration capacity of the active metabolite(s), cytotoxic products may diffuse into and kill adjacent cells within the tumour; a phenomenon termed the 'bystander effect'. A robust bystander effect would allow for cell killing outside of the necrotic boundaries, beyond where *Clostridium* germination is possible.

#### (i) Enzyme-prodrug combinations

The therapeutic success of CDEPT relies partly on the properties of the enzyme/prodrug combination used. The prodrug activating enzyme should have a high  $k_{cat}$  (catalytic rate) and low  $K_m$  (substrate concentration required for effective catalysis to occur) to maximise production of the active metabolite. The prodrug itself should be a stable, water soluble, systemic agent that remains inert in normal tissues, with good extravascular penetration into the tumour. The physiochemical properties of the active prodrug metabolites should allow for good cellular uptake and the ability to diffuse throughout the tumour microenvironment to generate bystander cell killing. A range of enzyme/prodrug combinations have been studied in the context of CDEPT, although none have been evaluated in clinical trial to date.

Proteolytic *Clostridium* strains engineered for CDEPT were the first to demonstrate beneficial effects on tumour

growth in vivo after administration of prodrug, likely due to their improved ability to colonise tumours. Initial studies were performed using the CD/5-FC enzyme/prodrug combination. Intravenous injection of a strain *C. sporogenes* engineered to express CD resulted in expression of active CD exclusively in the tumour [22]. The combination of these CD-expressing spores and prodrug 5-FC produced a larger tumour growth delay than that produced by 5-FU (the active metabolite of 5-FC) when given at maximum tolerated dose [22]. However, the cell cycle selectivity of 5-FU [38] is a major drawback to this particular enzyme/prodrug combination.

Bacterial nitroreductase (NTR) enzymes metabolise a diverse range of nitroheterocyclic substrates [39] and provide an ideal alternative to CD for expression in Clostridium. Traditionally, NTRs have been used to convert the non-toxic prodrug CB1954 to the active metabolites that cause DNA mono adducts and crosslinks [40]. The NTR/ CB1954 combination is not cell cycle specific, as killing has been observed in both cycling and quiescent cells [41]. Clostridium has been engineered to express various bacterial NTRs for use in combination with CB1954. For example, intravenously injected spores of C. sporogenes expressing an NTR from Haemophilus influenzae produced significant in vivo anti-tumour efficacy when combined with administration of CB1954 (P < 0.0001when compared with untreated control animals) [21]. Interestingly, bacteria were cleared from the animals in this study using antibiotics between treatment cycles, with no reduction in numbers of germinated bacteria occurring after repeated administration of spores. This was in line with previous observations suggesting that repeated spore

Fig. 3 Schematic representation of Clostridium-directed enzyme prodrug therapy (CDEPT) using the enzyme/prodrug combination of nitroreductase/ PR-104A as an example. (1) Inert prodrugs can be administered systemically and diffuse into the tumour. (2) Exogenous therapeutic enzyme expressed by Clostridium can metabolise the inert prodrug into active, cytotoxic metabolites. (3) These metabolites can be designed to diffuse out of necrotic regions into the outer rim of viable tumour (metabolite 'bystander' effect)



administration is not immunogenic to the host [20,42]. More recently, a novel nitroreductase from Neisseria meningitidis (NmeNTR) was expressed from a strain of C. sporogenes that is potentially suitable for clinical trial due to the absence of antibiotic resistance markers [27]. NmeNTR-expressing C. sporogenes in combination with CB1954 produced a significant growth delay (P < 0.001when compared with untreated control animals), with 4/16 tumours undergoing maintained complete regressions [27]. In contrast, CB1954 administered alone had no effect on tumour volume, and a small, non-significant growth delay was observed when NmeNTR-expressing C. sporogenes spores were administered as a single agent [27]. Whilst encouraging, the dose of CB154 used in these studies is in excess of that which can be safely achieved in humans [43]. In addition, the NTRs described produce the 4-hydroxylamine metabolite of CB1954 exclusively [44], resulting in a limited bystander cell killing.

As an alternative to CB1954, the prodrug PR-104A (or its phosphate ester 'pre-prodrug' PR-104) has been proposed for use in combination with NTR due to its improved potency and potential for bystander cell killing [45]. PR-104 has been used in pre-clinical models of E. coli NTR-armed CDEPT with considerable success [23]. This NTR/prodrug combination produced superior antitumour activity (tumour growth delay > 13 days, P < 0.05) compared with NTR-expressing spores or PR-104 alone [23]. The NmeNTR strain of C. sporogenes described above also demonstrated a significant improvement in survival in combination with PR-104, an increase of 20 days compared with untreated control animals (P < 0.001) [28]. The superior activity of the NmeNTR/PR-104 combination was confirmed by a 100% overall response rate as determined by RECIST 1.1 criteria [28]. However, PR-104A has been identified as a substrate for aldo-keto reductase 1C3 (AKR1C3) [46], a human enzyme involved in steroid hormone and prostaglandin regulation [47]. Reduction of PR-104A by AKR1C3 is thought to contribute to the doselimiting toxicities observed in humans that significantly restricts exposure [48]. Fortunately, novel analogues of PR-104A are being developed that are resistant to metabolism by AKR1C3 but still have significant activity with NTR enzymes [49–51].

#### (i) Non-invasive imaging capability

For decades, reporter genes such as  $\beta$ -galactosidase and *Renilla* luciferase have been used to monitor transgene expression in tissue biopsy samples. However, these techniques are invasive and time consuming and are therefore not suitable for repeated clinical evaluation. With

tomographic scanning becoming more routine, the ability to non-invasively image certain reporter genes is now possible. This would provide a means to accurately determine *Clostridium* distribution and replication in real-time, particularly after systemic administration, and may assist with expediting clinical development of *Clostridium* vectors.

CD and NTR enzymes are most commonly imaged in vitro and in vivo using red or near-infrared fluorescent probes [52–54]. However, there are issues with light penetration through tissue, as it has been predicted that fluorescent light cannot penetrate beyond the depth of several centimetres [55]. The recent demonstration that NTR enzymes can metabolise positron emission tomography (PET) imaging agents already in clinical development for imaging of hypoxia is a significant advantage to the use of this enzyme in the CDEPT context over CD. <sup>18</sup>F-labelled 2-nitroimidazole compounds can detect hypoxic cells following an oxygen-sensitive single electron reduction step, leading to further reduction to the reactive products that can covalently bind to cell components and accumulate inside the cell [56]. NTR was hypothesised to bypass the oxygen-sensitive step to produce the reactive products directly by two-electron reduction [39]. The hypoxia PET probe <sup>18</sup>F-HX4 was first used to detect expression of an NTR from E. coli, where a significantly higher tumourto-blood ratio was observed for tumours over-expressing NTR compared with corresponding parental tumour [57]. These results were confirmed in a separate study using <sup>18</sup>F-FMISO to detect NTR expression, including in small metastatic lesions [58]. However, the innate background signal from hypoxic tissue may somewhat limit the utility of these probes for NTR imaging. NTR-selective analogues of these 2-NI PET probes are being developed to overcome this issue [59].

#### (iii) Increased sensitivity to antibiotics

An advantage to NTR as a transgene is its catalytic activity across a range of antibiotic and anti-infective substrates, providing an additional safety feature which may be useful for clinical development. The hypersensitivity to these substrates that is provided by the insertion of NTR could, in theory, assist in eliminating *Clostridium* infection at the end of treatment or if an adverse event occurs. Metronidazole (Flagyl<sup>TM</sup>) is most commonly associated with the treatment of anaerobic bacterial infections [60], and expression of NTR has been shown to render host cells more sensitive to this antibiotic [28,61]. NTR has also shown activity with other anti-infective substrates such as tinidazole, nitrofurantoin, nimorazole and misonidazole, offering some alternative options in case of antibiotic resistance [28].

# 6 Clostridium in the Context of Immunotherapy

Cancer immunotherapies have been widely hailed as a breakthrough for cancer treatment in the last decade, epitomised by the unprecedented results observed with immune checkpoint inhibitors (ICIs). This paradigm shift towards harnessing an anti-tumour immune response as a treatment strategy has led to a renaissance for tumour-targeting bacteria such as *Clostridium*. Strains of *Clostridium* can be engineered to express immunomodulatory molecules (Fig. 4), resulting in localised delivery to the tumour microenvironment whilst potentially minimising systemic toxicity, thus improving therapeutic index.

# (i) Interleukin 2 and Tumour Necrosis Factor alpha

Interleukin 2 (IL-2) is a key cytokine with pleiotropic effects on the immune system, including the differentiation and activation of T cells [62]. High doses of recombinant IL-2 can induce tumour regression in patients [63,64], lead-ing to its approval by the FDA for the treatment of metastatic melanoma and metastatic renal cell carcinoma. However, systemic administration of this molecule can cause significant dose-limiting toxicities, including high serum cytokine levels that appear to be directly or indirectly toxic to multiple organs [65]. This makes IL-2 an ideal candidate for tumour selective *Clostridium*-mediated expression. Initial studies used *C. acetobutylicum* DSM792 for plasmid-based expression of rat IL-2 at therapeutically relevant concentrations

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[66]. More recently, *C. sporogenes*-NT was used for plasmid-based expression of mouse IL-2 [16].

Tumour necrosis factor alpha (TNF- $\alpha$ ) is a major regular of inflammatory responses and can be a potent mediator of tumour regression <sup>67</sup>. Similar to IL-2, systemic administration of this cytokine is not possible as it results in serious dose-limiting toxicities that are associated with sepsis [68]. Therefore, a strain of C. acetobutylicum DSM792 was engineered to express mouse TNF- $\alpha$  from a plasmid in an attempt to improve tumour selectivity [69]. Biological activity of the secreted TNF- $\alpha$  was confirmed in a bioassay using WEH164 clone 13 cells (which are highly sensitive to the cytotoxic effect of TNF- $\alpha$ ), and the plasmid used to derive TNF- $\alpha$  expression was segregationally stable [69]. Whilst the above studies report expression and secretion of a biologically active cytokine from Clostridium, none of the engineered strains were tested for therapeutic efficacy in vivo. This next step will be crucial for progression of cytokineexpressing strains towards clinical trial.

(ii) Nanobodies

ICIs target inhibitory receptors (e.g., CTLA-4, PD-1) and ligands (e.g., PD-L1) expressed on T-cells, antigen presenting cells and tumour cells to stimulate the immune system and elicit an anti-tumour response. Diseases with historically poor long-term survival (e.g., advanced melanoma, advanced non-small cell lung cancer) are now experiencing significantly improved therapeutic outcomes [70,71]. Unfortunately, ICIs can be associated with systemic autoimmunerelated adverse events of varying severity, often requiring

Fig. 4 Schematic representation of how *Clostridium* can be utilised in the context of cancer immunotherapy. (1) *Clostridium* strains can be engineered to express immunomodulatory molecules selectively within the tumour microenvironment, improving therapeutic index. (2) Transgene-free *Clostridium* can be used to promote immune cell infiltration into the tumour microenvironment, potentially sensitising tumours to immunotherapy treatment



treatment cessation and clinical intervention, thereby limiting therapeutic utility [72]. In addition, some ICIs engage targets deep within the tumour microenvironment; these large macromolecules (146–149 kDa) may experience restricted diffusion into tumour tissue [73]. *Clostridium* strains offer a unique opportunity to generate high local concentrations of ICI, potentially improving the selectivity and therapeutic efficacy of ICIs in certain patients.

In the past, *C. novyi*-NT and *C. sporogenes* have been engineered to produce the variable domain of a heavy-chain antibody (VHH antibody or nanobody) specific for human HIF-1 $\alpha$  [74]. The antibodies isolated from these strains retained their binding capacity and specificity for the HIF-1 $\alpha$ antigen. It is reasonable to assume that production of VHH antibodies against more immunologically relevant molecules (e.g. CTLA-4, PD-1 and PD-L1) would be possible in the future, analogous to the strain of *E. coli* Nissle 1917 that produces VHH antibodies against PD-L1 and CTLA-4 [75].

(iii) Clostridium in combination with immune checkpoint inhibitors

Whilst checkpoint blockade has revolutionised cancer therapy, only a minority of patients are able to achieve durable, long-term remissions. Responsive patients often have pre-existing immune cell infiltrate within the tumour microenvironment. As an alternative to immunomodulatoryexpressing *Clostridium* vectors, selective colonisation of the tumour by transgene-free *Clostridium* strains could alter the local immune microenvironment to one that is more immunogenic through the innate and adaptive immune response to infection (Fig. 4). This approach has the potential to sensitise tumours to subsequent treatment with immunotherapy and improve rates of response, particularly in tumours that are inherently immunosuppressive and thus largely resistant to immunotherapy at present (e.g. pancreatic cancer and glioblastoma [76,77]).

Early clinical data with *C. novyi*-NT demonstrated that intratumoural injection of endospores produced a transient systemic cytokine response and tumour-specific T-cell responses [12]. These data supported an additional phase Ib dose escalation study to assess the safety and potential synergistic effects of *C. novyi*-NT in combination with pembrolizumab in advanced solid tumours (NCT03435952). Initial results indicate that this combination is well tolerated in the 16 treatment-refractory patients recruited thus far, with a confirmed overall objective response rate of 25% (made up of three partial responses and one complete response) [78]. Currently there is no confirmation of tumour colonisation in responding patients or information on host immune responses. This proof-of-principle study is ongoing to identify the optimal dose for phase II.

# 7 Conclusion

*Clostridium* species are a promising and versatile vector system for use in cancer therapy, with many advantages over traditional vector platforms. Recent developments in methods for the genetic modification of *Clostridium* have allowed for expression of prodrug-activating genes and immunotherapeutics selectively within the tumour micro-environment. However, clinical development has been slow thus far, likely due to challenging and complex regulatory requirements. The ongoing clinical progress being made with *C. novyi*-NT, in addition to the potential for non-invasive monitoring of transgene expression through PET imaging, should help to demonstrate the feasibility and relative safety of *Clostridium* vectors and hopefully lead to additional clinical studies in this area.

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