

Oral microbiota: Roles and treatment in radiation injury (Review)

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Abstract. While radiotherapy for malignant tumors increases the rate of local control for patients, it also inevitably causes damage to the surrounding normal tissues, resulting in different types and degrees of radiation injury (RI). RI has a significant negative impact on quality of life. However, there is no viable method for preventing and treating these complications. The oral microbiota plays a key role in the development of numerous categories of RI, such as radiation-induced brain injury, radiation-induced oral injury and radiation-induced lung injury. The aim of the present review is to clarify the potential mechanisms of RI and provide a comprehensive overview of the association between 10 distinct categories of RI and oral microbiota, thereby shedding light on novel clinical approaches for the prevention and treatment of RI based on modulating oral microbiota. This may open new avenues for targeted interventions.

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1. Introduction

One of the primary therapies for malignant tumors is radiotherapy, the effectiveness of which is closely correlated with

the radiation dose. Following radiotherapy for tumors, patients may experience various acute and chronic radiation injuries (RIs) in different sites of the body; a prime example is radiation-induced lung injury (RILI), which encompasses any lung toxicity induced by radiation therapy and manifests acutely as radiation pneumonitis and chronically as radiation pulmonary fibrosis. RI is the main factor that restricts the effectiveness of radiotherapy, as well as affecting the prognosis and quality of life of patients, as its severity increases rapidly with higher radiation doses (1). Nevertheless, currently, our understanding of the mechanisms underlying different RI types remains limited, and there is a lack of adequate clinical approaches for prevention/treatment. Therefore, there is a pressing need for novel and efficient solutions to prevent and treat RI.

Given the ongoing advancements in 16S ribosomal RNA sequencing and quantitative macro-genome detection technology, combined with decreasing costs, the investigation of the human microbiota as the 'second genome' has attracted widespread attention from medical researchers both domestically and internationally (2). Comprehensive analysis of the microbiota genome structure in various regions of the human body, such as the intestinal tract, oral cavity, respiratory tract, vagina and skin, has deepened our understanding of the diversity and functional abundance of microbiota that play a vital role in maintaining fundamental physiological functions and contribute to the development of associated diseases (3). The oral microbiota and its dysregulated metabolites have been shown to contribute significantly to the onset, progression, metastasis, diagnosis and prognosis of malignant tumors (4-8). It is worth noting that the oral microbiota is strongly linked to several categories of RIs. Therefore, the oral microbiota has demonstrated significant promise in the prevention and treatment of RI, prompting extensive exploration by researchers worldwide.

Within this review, the processes by which oral microbiota participates in 10 distinct categories of RI are delineated. Additionally, the impact of the oral microbiota on RI located at various sites is discussed. The aim of the present review is to present the most up-to-date medical evidence, paving an efficient path toward the prevention and treatment of RI.

2. Overview of oral microbiota and RI

The oral microbiota are a diverse group of microorganisms that include viruses, protozoa, fungi, archaea and bacteria. According to the Human Oral Microbiome Database (www.humanoralmicrobiome.org),

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homd.org), the oral cavity harbors 774 microbial species. These species, categorized at the phylum level, primarily consist of *Actinobacteria*, *Bacteroidetes*, *Firmicutes*, *Ascomycota*, *Spirochaetes* and *Saccharomycetes*. These species play a crucial role in defending against external pathogens and preserving oral health. Nevertheless, they can also induce oral disorders, such as dental caries, periodontal disease and oral malignancies (9-11). The most notable pathogens among these are *Porphyromonas gingivalis* (*P. gingivalis*) and *Fusobacterium nucleatum* (*F. nucleatum*), both gram-negative anaerobic bacteria (8). In addition, there is growing evidence that oral microbiota can affect distal organs through a variety of mechanisms. Specifically, oral microbiota can cause disease through direct translocation to distal organs, such as translocation to the lungs leading to aspiration pneumonia (12), and translocation to the gastrointestinal tract leading to gastrointestinal diseases and malignant tumors (13).

Another mechanism acts indirectly through the body's immune system and inflammatory responses leading to distal diseases, such as exacerbating atherosclerosis by inducing the apoptosis of smooth muscle cells and inhibiting the apoptosis of macrophages (14), and causing inflammation in the nerves resulting in Alzheimer's disease (15). Therefore, during radiation therapy, oral microbiota and their metabolites can reach various organs in the body through multiple pathways (Fig. 1) and participate in the development of RI through multiple mechanisms (Fig. 2) (16-21).

Radiotherapy, a cornerstone of oncological treatment, utilizes precisely targeted high-energy radiation to eradicate malignant tumor cells. This therapeutic intervention inevitably inflicts collateral damage on healthy tissues through three multifactorial mechanisms: Direct cellular injury, the radiation-associated bystander effect (22), and inflammatory cascades induced by activation of both innate and adaptive immune systems (1). These pathological processes collectively lead to RI (Fig. 1). These injuries can manifest as both acute and chronic adverse reactions. Acute adverse effects are prevalent, with the majority of patients experiencing mild to moderate fatigue, skin and mucous membrane damage in the irradiated area, fever, nausea and vomiting, and reduced peripheral blood counts following radiotherapy. These harmful processes substantially compromise long-term survival outcomes and quality of life, thereby acting as a dose-limiting factor that forces clinicians to balance tumor control against toxicity risks and restricts feasible radiation dose escalation, even when higher doses might theoretically improve survival (23).

3. Investigation of the association between oral microbiota and different categories of RI

Radiation-induced brain injury (RIB). RIB is a significant adverse reaction following radiotherapy for brain or head and neck malignant tumors. RIB includes damage to the hippocampus and white matter, vascular abnormalities, demyelination, localized neurological deficits and elevated intracranial pressure (24). RIB is primarily characterized by cognitive impairment and neurological inflammation (25). According to statistics, 50 to 90% of brain tumor patients with survival periods >6 months develop RIB (26). Previous research demonstrated that the connections between the oral

microbiota and the central nervous system, known as the oral-microbiota-brain axis (27), are linked to cognitive function and neuroinflammation, through which the oral microbiota and their metabolites interact with the immune system. For example, bacteria can secrete immunostimulants such as lipopolysaccharide (LPS) and peptidoglycan into the circulation, enabling them to enter the brain. LPS from Gram-negative bacteria in the oral cavity induces the expression of neuronal inflammatory proteins in SH-SY5Y and HMC3 cells (28). Oral microbiota components or some of their metabolites can even cross the blood-brain barrier directly, altering the inflammatory state of the central nervous system and leading to cognitive dysfunction through a variety of mechanisms (29). For example, Bulgart *et al* (30) demonstrated that *P. gingivalis* metabolite gingipain protease disrupts the processing of the transmembrane protein amyloid precursor protein, thereby affecting synaptic stability and neurite growth; and a more recent study showed that *P. gingivalis* can cause cognitive dysfunction by activating the p38 mitogen-activated protein kinase (MAPK) signaling pathway (31). Prior studies have established that excessive activation of microglia in the central nervous system is crucial for the development and advancement of radiation-induced brain damage (32,33), whereas oral microbiota can affect brain function via intracranial microglia. The oral microbiota can affect brain function via microglial activation. Chuang *et al* (34) revealed that outer membrane vesicles derived from *P. gingivalis* cause neurotoxicity and microglial activation.

Radiation-induced oral injury. Ultrafast division of oral mucosal cells makes them susceptible to radiation. Radiation-induced oral mucositis (RIOM) is the predominant adverse effect on the oral mucosa following head and neck radiotherapy. RIOM is characterized by congestion, edema, ulceration and discomfort in the oral mucosa. Statistics indicate that nearly all patients undergoing head and neck radiation experience RIOM, with over half of them developing severe oral mucositis (35,36).

The oral microbiota undergo changes during radiation therapy; for example, a clinical study by Nodit *et al* (37) systematically explored the dynamics of the oral microbiome during cancer RIOM. At radiation therapy completion, significant microbial shifts were observed, including increased abundance of *Fusobacterium* and *Streptococcus*, alongside decreased levels of *Rothia* and *Lautropia*. Numerous studies have shown that alterations in the oral microbiota after radiotherapy have the potential to be valuable in predicting the incidence and evaluating the severity of mucositis. Hou *et al* (38) investigated the characteristics of changes in the oropharyngeal mucosal microbiota of patients with nasopharyngeal carcinoma during radiotherapy, and no significant change was found in the bacterial α -diversity of the mucosal microbiota, whereas the abundance of *Prevotella*, *Clostridium*, *Mycobacterium*, *Porphyromonas* and *Aeromonas* showed significant dynamic changes in abundance, with their peaks coinciding with episodes of severe mucositis (38). By contrast, Zhu *et al* (39) found that patients with severe mucositis had significant microbial α -diversity and higher abundance of *Actinobacteria*, while Vesty *et al* (40) demonstrated that the presence of Gram-negative bacilli, *Clostridium innocuum*

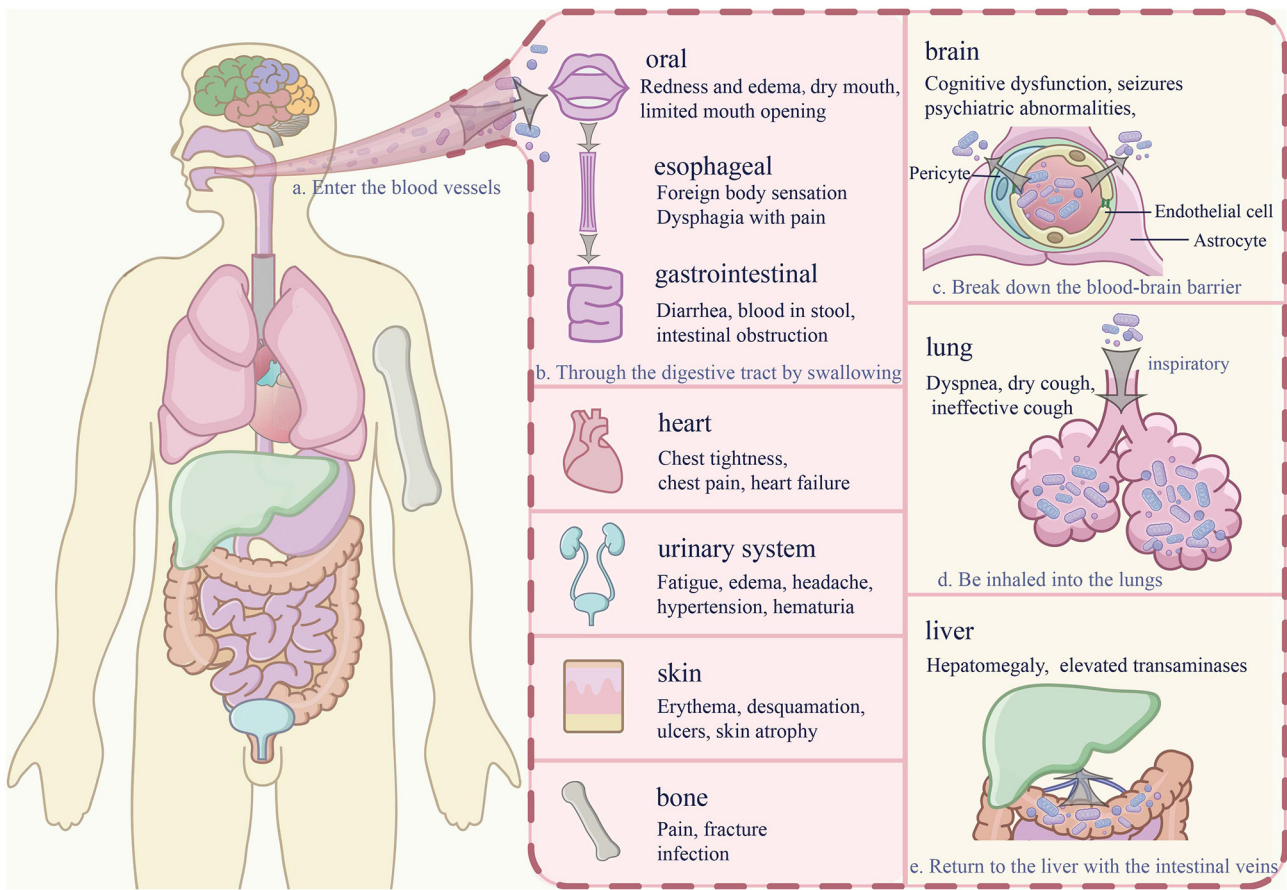


Figure 1. Multiple paths allow oral bacteria and their metabolites to reach different organs in the body. Oral bacteria and their metabolites can: (a) Enter into the blood vessels and travel to different organs through the bloodstream, (b) colonize the esophagus, stomach and intestines through the digestive tract via swallowing, (c) enter the brain by breaking through the blood-brain barrier, (d) enter the lungs through the saliva and (e) return to the liver through the intestinal veins in the intestinal tract.

and *Porphyromonas*, correlated with the severity of RIOM. In addition, a retrospective multicenter study of 326 patients by Nishii *et al* (41) revealed that higher levels of RIOM were significantly associated with increased incidence of oral candidiasis.

These clinical studies suggest that dysregulation of oral microbial ecology may exacerbate RIOM severity. There is also growing evidence that changes in the oral microbiota also influence the development and prognosis of mucositis through a range of mechanisms. Groeger and Meyle (42) found that *P. gingivalis* could mediate enhanced IL-33 secretion by epithelial cells, which in turn augmented the Th2 cytokine-mediated inflammatory response. Hirahara *et al* (43) demonstrated that the oral microbiota can cause excessive persistence of inflammation by inducing aberrant polarization of M1/M2 macrophages. Suárez *et al* (44) similarly found that Toll-like receptors (TLRs) initiate a pro-inflammatory response and upregulate antigen presentation by immune cells bound to LPS, exacerbating the oral inflammatory response. Experiments showed that the oral microbiota interacts with TLRs to regulate the immune system. Specifically, LPS from Gram-negative microbiota activate TLRs, subsequently activating the NF-κB pathway, driving the release of TNF-α and IL-6, and triggering an inflammatory cascade (17). Therefore, modulation of the oral microbiota is a potential target for combating (45).

Researchers have identified several protective agents within the microbiome against RIOM: Zhao *et al* (46) developed fulleranol, an antioxidant that inhibits apoptosis, maintains microbial homeostasis and protects against RIOM by scavenging radiation-induced excess reactive oxygen species and upregulating intrinsic enzyme-activated antioxidants. Topical Omega-3 nanoemulgel demonstrated a beneficial effect in preventing RIOM, mediated by its ability to regulate oral microbial dysbiosis; specifically, Morsy *et al* (47) found it significantly reduced the *Firmicutes/Bacteroidetes* ratio in patients with head and neck cancer undergoing radiotherapy.

Radiation-induced esophageal injury. As a common inflammatory reaction to thoracic radiotherapy, radiation-induced esophagitis (RE) is most often observed following radiotherapy for esophageal cancer, lung cancer, breast cancer, mediastinal malignancy, lymphoma and other thoracic cancers (48). Following deglutition, the oral microbiota translocate to the esophagus via the saliva, forming an esophageal microbiota that is compositionally analogous to its oral counterpart (49). Thus, in addition to acting indirectly through the immune system and inflammatory response, oral microbiota may be directly involved in the progression of radiation esophagitis by migrating into the esophagus. Research has demonstrated that Gram-negative microbiota exhibit predominance in the esophageal microbiota of individuals with esophagitis (50),

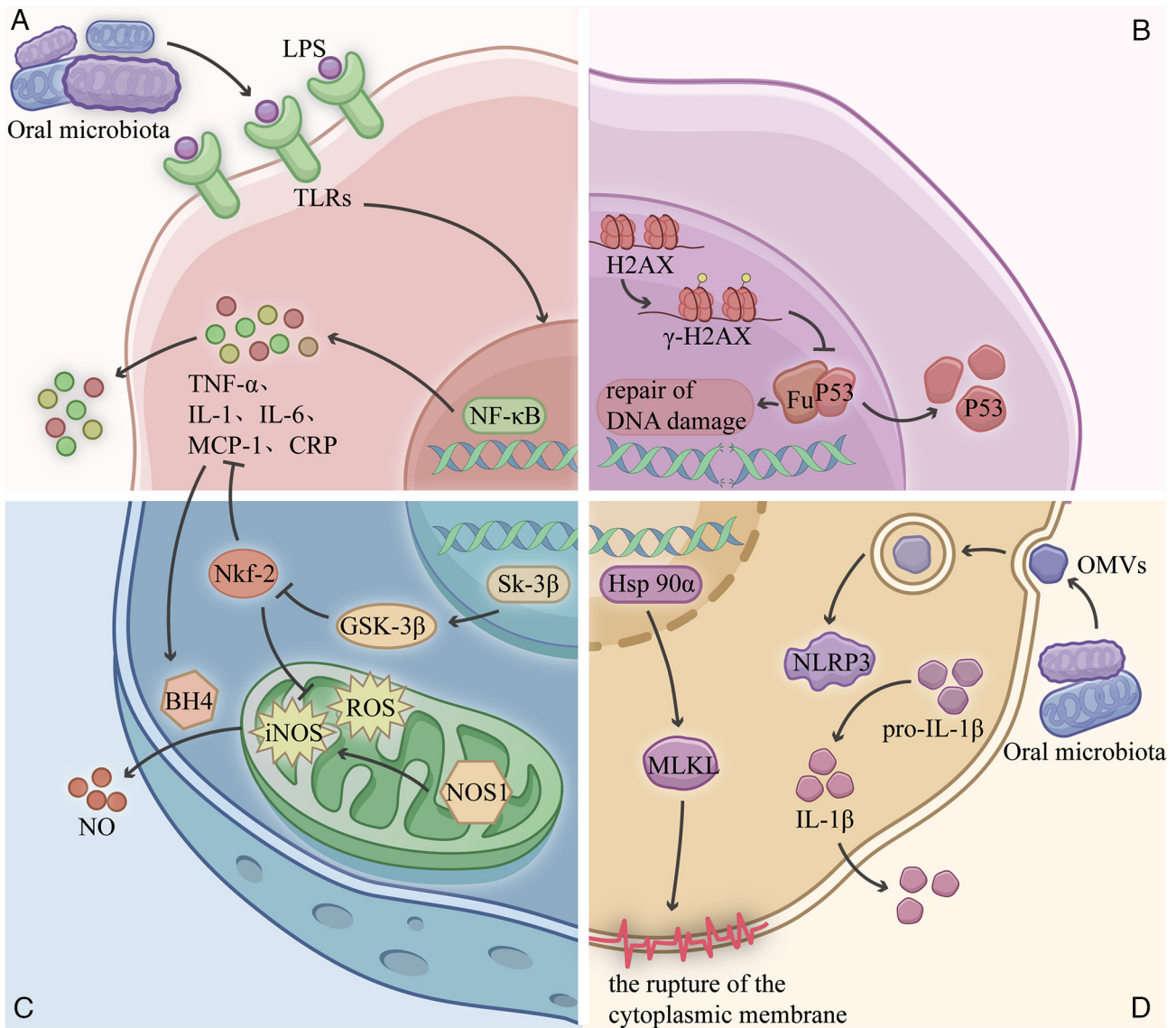


Figure 2. Molecular mechanisms of the oral microbiota that are involved in RI. (A) Initiation of the inflammatory response: The LPS of Gram-negative bacteria stimulates TLRs, which in turn trigger an inflammatory cascade response, release TNF- α , IL-6 and other inflammatory factors, and activate the NK- κ B pathway (16,17). (B) Involvement in DNA damage: The key proteins involved in DNA damage repair, Ku and p53, are expressed less frequently, while γ H2AX is expressed more frequently, in oral bacteria, suggesting both increased DNA damage and inadequate DNA repair (18). (C) Contribution to oxidative stress: The oral microbiota decrease Nrf-2 activity, worsen oxidative stress and increase inflammation via the GSK-3 β /BH4/eNOS/Nrf2 pathway. LPS also activates NOS1 and upregulates iNOS, and Nrf-2 controls the expression of antioxidant enzymes to protect cells and reduce oxidative stress (19). (D) Additional mechanisms: The oral microbiota can upregulate Hsp90 α and activate the tetrahydrobiopterin (MLKL), which causes cytoplasmic membrane rupture and cellular necrosis (20). Oral microbial OMVs can activate the NLRP3 inflammasome and increase IL-1 β production, aggravating cellular inflammation (21). MLKL, mixed lineage kinase domain-like pseudokinase; LPS, lipopolysaccharide; TLR, Toll-like receptor; NOS1, nitric oxide synthase 1; OMV, outer membrane vesicle.; ROS, reactive oxygen species; NLRP3, nucleotide-binding oligomerization domain-like receptor family pyrin domain-containing 3; GSK-3 β , glycogen synthase kinase 3 β ; Nrf-2, nuclear factor erythroid 2-related factor 2; γ H2AX, phosphorylated histone H2AX; MCP-1, monocyte chemoattractant protein-1; CRP, C-reactive protein; iNOS, inducible nitric oxide synthase.

mirroring the alterations in the oral microbiota observed in people with oral mucositis. Furthermore, it has been established that LPS present in Gram-negative microbiota stimulate TLRs to exacerbate inflammatory reactions. Notably, LPS induces nitric oxide synthase (NOS)1 upregulation and increases the production of inducible NOS, resulting in the relaxation of the lower esophageal sphincter (51). In addition, LPS impairs the rate of stomach emptying via COX-2 stimulation, consequently increasing gastric pressure (50) and thereby elevating esophagitis risk. Although no specific research has revealed a direct link between the oral microbiota and the onset of radiation

esophagitis, the aforementioned evidence suggests a possible association between the two.

RILI. RILI is a severe complication from radiation to the lung tissue, characterized by early radiation pneumonitis (RP) and late radiation pulmonary fibrosis, reported in 25% of treated cases (52). Clinically, RILI not only impacts the health and treatment outcomes of cancer patients, but may lead to life-threatening complications in severe cases. The oral cavity and the lungs are anatomically contiguous, allowing microbial translocation via salivary aspiration and subsequent inhalation

into the lungs. Furthermore, the organization of the microbiota in both the oral cavity and the lungs is similar (53). Multiple studies have demonstrated that the oral cavity and the lungs exert mutual influence (54,55). Thus, researchers proposed the oral-lung axis (56), with the oral microbiota significantly contributing to the pathogenesis of pulmonary inflammatory conditions.

The animal model described in the study by Benedyk *et al* (16) showed a significant elevation in the peripheral blood concentrations of TNF- α , IL-6, monocyte chemoattractant protein-1 and C-reactive protein following intratracheal administration of *P. gingivalis*. These inflammatory mediators promote airway inflammation by mediating the recruitment and activation of key immune cells, including neutrophils and monocytes, within the respiratory tract (16). Other researchers found that *P. gingivalis* and *F. nucleatum*, whose pathogenic mechanism is similar to that of *P. gingivalis*, can induce the expression of IL-6 and IL-8 in the bronchial epithelial cells, triggering inflammatory cascades, with IL-8 stimulating the production of the extracellular matrix by lung fibroblasts, a critical mechanism in fibrogenesis (57,58). Mechanistically, Feng *et al* (20) proved that *P. gingivalis* stimulates mixed lineage kinase domain-like pseudokinase-mediated necroptosis by increasing the expression of Hsp90 α , resulting in plasma membrane disruption. This directly triggers and indirectly spreads inflammatory responses. Concurrently, *F. nucleatum* can activate MAPKs and NF- κ B, which in turn stimulate matrix metalloproteinases that play a role in airway reconstruction (59). Beyond direct cellular effects, oral pathogens can synergistically interact with pulmonary microbiota to enhance pulmonary inflammatory responses. *P. gingivalis* or its virulence factors can potentiate the pathogenicity with *Streptococcus pneumoniae* and *Pseudomonas aeruginosa* to intensify the inflammatory disease response (57,60). Clinical corroboration comes from a clinical study by O'Dwyer *et al* (61), which found that oral microbial diversity predicted disease severity and death in patients with pulmonary fibrosis. Despite these compelling associations linking oral microbiota with pulmonary inflammation and pulmonary fibrosis, their involvement in RILI remains unclear.

Radiation-induced gastrointestinal injury (RIGI). Primary malignant tumors in the abdomen and pelvis mostly cause radiation-induced gastrointestinal damage. This damage includes radiation-induced enteritis (REn), gastric injury, duodenitis and proctitis. Among these complications, REn is the most prevalent and can be life-threatening for patients in severe cases (23,62). The mouth is the beginning of the digestive tract, and although these two loci are physiologically distinct, they are directly connected and can interact in a number of ways. Therefore, the microbiota in the mouth and intestines can mutually interact to establish an oral-intestinal axis (63,64). Yamazaki (65) established that the oral microbiota induced intestinal dysbiosis, including various inflammatory disorders, by inducing dysbiosis of the intestinal microbiota. Emerging evidence further implicates the oral microbiota in RIGI progression. For example, oral microbiota can lead to reduced expression of tight junction protein genes in intestinal tissues and disrupt intestinal

barrier function (65). Atarashi *et al* (66) found that *Klebsiella spp.* isolated from the oral cavity could colonize the mouse intestine and trigger Th1-mediated inflammatory responses. Dong *et al* (67) found that oral microbial transplantation (OMT) impaired the therapeutic efficacy of radiotherapy in colorectal malignant tumors. Furthermore, notably, it was found experimentally that colorectal cancer mice treated with radiation therapy in combination with metronidazole had reduced overall tumor load and REn (49). This suggests the potential of metronidazole in counteracting oral microbiota-mediated radiotherapy complications.

Radiation-induced heart injury. Due to the anatomical proximity of the heart, radiotherapy to the chest can result in radiation-induced heart disease (RIHD). This may include damage to coronary arteries or valves, obstruction of the conduction system, pericardial injury, myocardial injury, myocardial fibrosis and other related conditions (68,69). RIHD represents the primary cause of non-cancer-related mortality in thoracic radiotherapy recipients (70). Emerging evidence implicates periodontal pathogens in cardiovascular pathophysiology through multiple mechanisms.

In experimental models, Peron *et al* (71) demonstrated that *P. gingivalis* causes elevated numbers of DNA breaks, increased malondialdehyde levels, increased oxidized protein levels and enhanced macrophage infiltration in the myocardium, leading to adverse effects. Bijla *et al* (72) revealed that gingival protease released by *P. gingivalis* impairs autophagy by preventing the fusion of autophagosomes and lysosomes in cardiomyocytes, resulting in reduced cardiomyocyte viability and promoting apoptosis.

Radiation-induced skin injury (RISI). RISI manifests universally in radiotherapy recipients due to the high mitotic index of epidermal keratinocytes. RISI occurs to varying degrees after radiotherapy and can be categorized into acute and chronic injuries. Acute RISI includes dry and wet desquamation, skin necrosis, ulceration and hemorrhage, while chronic RISI comprises chronic ulceration, actinic keratosis, capillary dilatation, fibrosis and skin malignant tumors (73). Several studies have demonstrated microbiota alterations as a potential pathogenetic mechanism for radiation dermatitis (74,75). There is also a link between the oral microbiota and skin inflammation. Li and Yosipovitch (76) found that patients with atopic dermatitis had significantly reduced oral microbiota diversity, characterized by a decreased abundance of pro-inflammatory-associated microbiota compared with that in healthy individuals, with functional antagonism between these microbial communities, suggesting that skin inflammation can be controlled based on the modulation of the oral microbiota. Notably, as aforementioned, the oral microbiota can promote pro-inflammatory cytokines such as IL-1 to initiate an inflammatory cascade response. Janko *et al* (77) found that IL-1 or IL-1 receptor-deficient mice exhibited reduced inflammation and attenuated radiation dermatitis severity. Collectively, microbial-mediated therapy may be a viable option for RISI.

Radiation-induced bone injury (RIBI). Considering the characteristically high calcium concentration in bone tissue, it can absorb up to 30 to 40% more radiation than

the surrounding tissues when exposed. This elevated absorption makes bone particularly vulnerable to RI at equivalent exposure doses (78). RIBI includes several conditions, such as osteoporosis, osteomyelitis, pathological fracture, osteonecrosis and growth restriction (57). Recent studies have revealed that radiation-induced osteonecrosis of the jaw, which is caused by radiotherapy for head and neck malignant tumors, is associated with notable alterations in the oral microbiota (58). These changes interfere with bone regeneration by regulating particular metabolic pathways that enhance osteoclast activity, thereby exacerbating radiation-induced osteonecrosis (79). However, the precise mechanisms underlying how oral microbiota influence osteogenesis have not yet been fully elucidated. Evidence suggests that the oral microbiota imparts osteoimmunomodulatory effects and modulates bone resorption (80). For instance, Xu *et al* (81) demonstrated that *P. gingivalis* and *F. nucleatum* promote bone resorption by activating bone remodeling cells through pro-inflammatory cytokines and modulation of the key axis, the receptor activator of nuclear factor- κ B ligand-receptor activator of nuclear factor- κ B-osteoprotegerin axis. In an experimental model, *P. gingivalis* LPS activated various cell types, including monocytes, endothelial cells and epithelial cells, resulting in the release of pro-inflammatory mediators that initiate an immune-inflammatory response in host tissues. Kovtonyuk *et al* (82) proposed that microbiota drive inflammation in hematopoietic stem cells via the microbiota/IL-1/IL-1R1 pathway. These findings collectively suggest the mechanistic involvement of the oral microbiota in RIBI progression.

Radiation-induced urinary system injury. The kidney is highly susceptible to radiotherapy, and radiation-induced kidney injury (RIKI) may result from abdominal or paraspinal malignancies, or systemic radiotherapy. The acute phase of RIKI is marked by malaise, edema, headache, hypertensive crisis and cardiorenal syndrome, whereas the chronic phase is defined by hypertension, anemia and progressive renal atrophy (83). The bladder is a critical organ susceptible to pelvic tumor radiotherapy. Radiation-induced bladder injury manifests as hemorrhagic cystitis, dysuria and overactive bladder syndrome (84). Yuan *et al* (85) introduced the oral-genitourinary axis derived from epidemiological and clinical research, identifying oral microbiota such as *P. gingivalis*, *Streptococcus oralis* and *Campylobacter jejuni*, and periodontal pathogens as contributors to elevated instances of prostatic hyperplasia, prostatitis, bladder cancer and other bladder diseases, via inflammation and the Akt signaling pathway (86-88). An increasing number of researchers have undertaken comprehensive investigations into the oral microbiota involved in the advancement of kidney disorders. Tooth brushing and mastication facilitate the entry of oral microbiota into the bloodstream through ulcerated periodontal pockets, resulting in bacteremia. Proinflammatory mediators from microbiota in the systemic circulation exacerbate the inflammatory burden on renal tissues (89). Kajiwara *et al* (90) revealed that the LPS of *P. gingivalis* stimulates Mac-1/podoplanin-positive macrophages through the overexpression of vascular cell adhesion protein 1 and E-selectin in glomerular infiltration,

leading to renal inflammation characterized by glomerulosclerosis and tubulitis. Consequently, the oral microbiota may have a role in the etiology of RIKI.

Radiation-induced liver injury. Radiation-induced liver injury (RILI) is a common complication of radiotherapy for thoracic and abdominal neoplasms, particularly hepatocellular carcinoma (91). Clinical data reveal that ~60% of patients with liver cancer receive radiotherapy to varying degrees, and RILI significantly limits the clinical utility of this treatment (91). Researchers have proposed the oral-gut-liver axis following an in-depth examination of the association between the liver and oral microbiota (92). The connection between the liver and the oral cavity may occur via the gut, where impaired intestinal permeability allows the direct translocation of oral microbiota metabolites and inflammatory mediators from the oral cavity into the systemic circulation or to the liver. Upon entering the liver, the oral microbiota may promote the progression of liver disease via multiple mechanisms. For instance, vesicles from *P. gingivalis* can be released into the cytoplasm of hepatocytes to diminish phospho-glycogen synthase kinase 3 activation and inhibit gluconeogenesis via the insulin/insulin receptor substrate-1 receptor signaling pathway (68). The LPS of *P. gingivalis* prompts the release of galectin-3 from hepatocytes and macrophage-2 antigen-positive Kupffer cells, subsequently activating the TGF- β 1/TGF β R2 cascade, which upregulates α -smooth muscle actin transcription and facilitates the differentiation of hepatic stellate cells into myofibroblasts (93). Research has demonstrated that the oral microbiota provoke hepatic dysfunction and fibrosis by enhancing immune cell infiltration and inflammatory gene expression in the liver, thereby activating the pro-inflammatory NF- κ B pathway (94). However, to date, the comprehensive understanding of the causal association between the oral microbiota and RILI remains incomplete. This causal association between the oral microbiota and hepatic RI requires further elucidation.

4. Oral microbiota-based therapeutic approaches for RI

Oral microbiota-based approaches may be a useful method for the prevention and treatment of RI (Fig. 3). The homeostatic balance of oral microbiota contributes to systemic equilibrium. Healthy lifestyle choices, low-carbohydrate diets and good oral hygiene practices all help to maintain dynamic oral microbiota equilibrium and reduce the oral microbiota burden (95). By introducing beneficial bacterial species and substrates, probiotics, prebiotics and synbiotics help to reverse microbial dysbiosis (96). As aforementioned, *Streptococcus salivarius* K12, a probiotic, has been shown to restore the oral microbiota and cure radiographic oral mucositis (97). By reducing oral microbiota's load and obstructing pathogenic pathways, antibiotics and inhibitors that target harmful microbiota and their derivatives can successfully mitigate the pathogenic consequences of pathogens. Protease inhibitors, for instance, can reduce the inflammatory response and the *P. gingivalis* microbiota load (13). Notably, OMT has the ability to directly alter the microbiota ecosystem of the recipient, which makes it a potentially useful therapeutic strategy for RI.

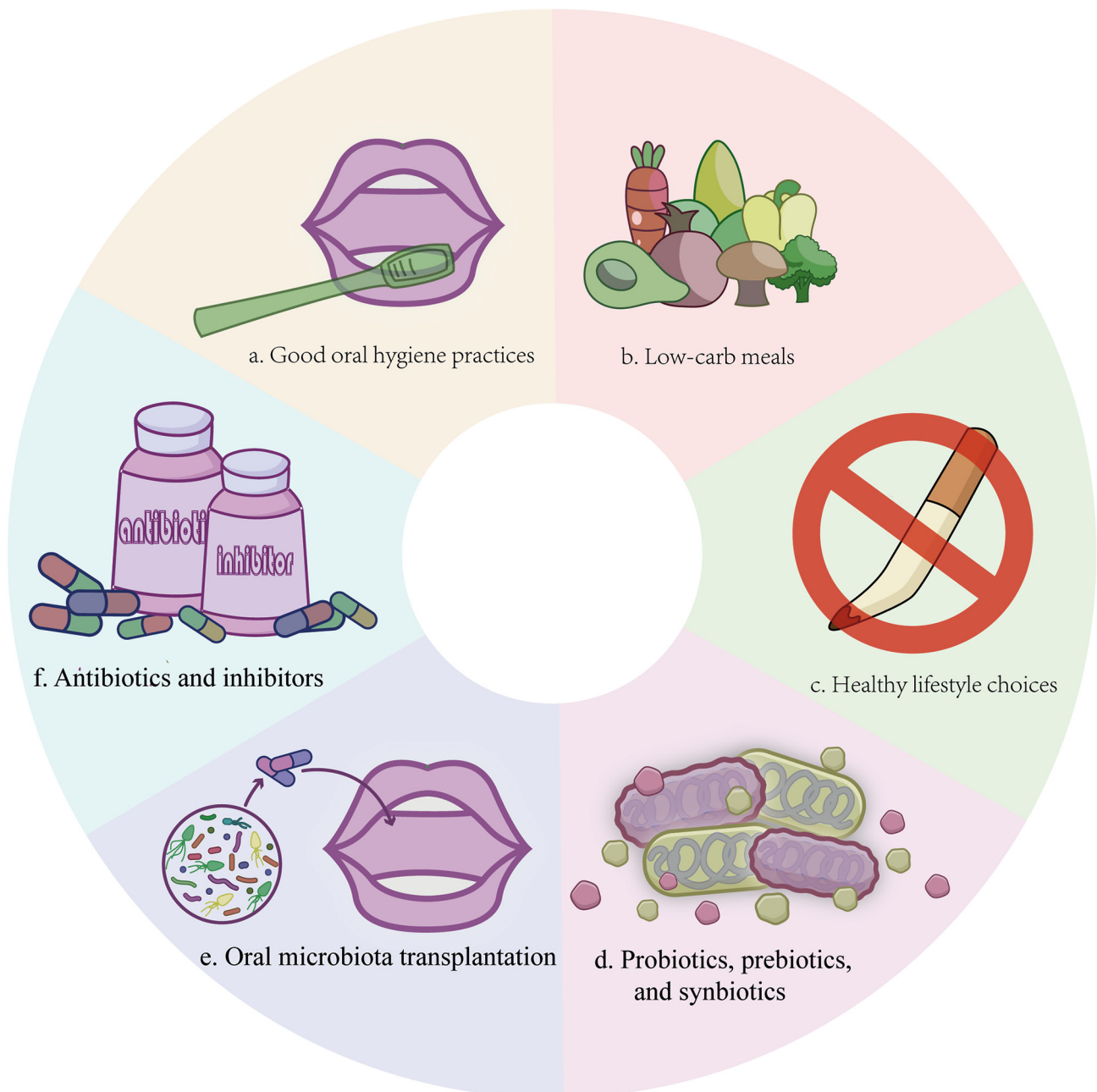


Figure 3. Oral microbiota-based therapeutic approaches for RI. Useful approaches include: (a) Good oral hygiene practices, (b) low-carbohydrate meals, (c) healthy lifestyle choices, (d) Probiotics, prebiotics and synbiotics, (e) oral microbiota transplantation, and (f) antibiotics and inhibitors.

Probiotics, antibiotics and inhibitors. Oral microbial dysbiosis can affect RI, and so probiotics, as a relatively predictable and safe measure to regulate microecology, are of interest to researchers.

First, despite the clinical trial by De Sanctis *et al* (98) demonstrating that *Lactobacillus brevis* CD2 lozenges failed to prevent RIOM in patients with head and neck cancer, strategic modulation of salivary microbiota homeostasis remains a compelling therapeutic avenue. Subsequently, a clinical trial by Peng *et al* (99) demonstrated that the probiotic *Streptococcus salivarius* K12 alleviated RIOM through suppression of opportunistic pathogens and reconstruction of murine oral flora. These findings are consistent with animal experiments conducted by Wang *et al* (97), which revealed that *Streptococcus salivarius* K12 ameliorate RIOM by suppressing oral anaerobic bacterial overgrowth. A clinical trial further

showed that topical application of *Streptococcus salivarius* K12 lozenges effectively decrease RIOM incidence, delay symptom onset and shorten disease duration (45). Moreover, nisin (a lantibiotic) was found to modify the oral microbiota composition in healthy rats and dogs (100,101). Animal studies by Gao *et al* (102) additionally demonstrated that nisin-producing *Lactococcus lactis* probiotics significantly lowered periodontal pathogen levels, attenuated alveolar bone loss and modulated host responses to oral/systemic inflammatory diseases, offering therapeutic implications for RI. In the context of RIB, Zhao *et al* (103) reported that nisin treatment significantly suppressed radiation-induced upregulation of pro-inflammatory cytokines (IL-1 β , IL-6 and TNF- α) in brain tissue affected by periodontal infections, suggesting therapeutic potential for RIB.

Furthermore, strategies targeting pathogenic bacteria through antibiotics or virulence factor inhibitors may mitigate pathogenicity via microbial load reduction and pathogenic pathway blockade. For instance, protease inhibitors have been shown to decrease *P. gingivalis* colonization and inflammatory responses (13). Notably, as previously mentioned, Dong *et al* (67) experimentally demonstrated that metronidazole combined with radiotherapy in mice with colorectal cancer resulted in reduced tumor burden and RE severity, implying therapeutic potential through modulation of oral microbiota-mediated interference with radiotherapeutic efficacy.

OMT. OMT, defined as the deliberate transfer of beneficial oral microbiota from healthy donors to patients, aims to establish microbial communities capable of modulating inflammatory responses (104). Accumulating evidence has demonstrated significant associations between oral microbiota dysbiosis and RI, prompting exploration of OMT as a therapeutic intervention for RI. In a murine model of radiation-induced head and neck injury, Xiao *et al* (105) administered OMT from healthy mice to irradiated mice with head and neck lesions, with mitigated oral mucositis demonstrating its potential as a novel treatment for RIOM. Clinically, Goloshchapov *et al* (106) recently reported successful OMT application in preventing chemotherapeutic oral mucositis. In this clinical trial, researchers showed that OMT effectively prevented oral mucositis following chemotherapy. This was achieved by administering repeated daily injections of donor saliva from healthy donors to patients for three chemotherapy cycles and before autologous hematopoietic cell transplantation (106). OMT can also be used to treat RP. AbdelMassih *et al* (107) hypothesized the therapeutic potential of OMT for low-grade inflammatory illnesses based on established links between oral ecological imbalance and pneumonia development. However, in investigations on radiotherapy for colorectal malignant tumors, scientists have discovered that OMT may have an adverse effect. Dong *et al* (67) discovered that OMT reduced the effectiveness of radiotherapy for colorectal malignant tumors. *et al* Verification of the use of this antibiotic for pretreatment of clinical colorectal malignant tumors radiation requires further investigation through clinical studies. Clinical translation warrants robust validation. Although OMT research has advanced, its clinical application remains experimental, mandating methodologically rigorous trials to establish therapeutic efficacy and safety.

Clinical translation program for treatment strategies. For current prevention and treatment strategies, future studies should prioritize expanding cohort sizes and diversifying research populations to validate the reliability and generalizability of current findings.

However, the mechanistic interplay between the oral microbiota and RI remains exploratory. Future investigations should focus on the following: i) Elucidating specific microbiota-radiation interaction mechanisms through establishment of oral microbiota-organ axis animal models; ii) deciphering molecular pathways mediating microbial effects on irradiated tissues across body regions; and

iii) characterizing microbiota-immune system crosstalk during radiation pathogenesis. This systematic approach will advance a comprehensive understanding of host-microbe dynamics to inform novel RI prevention and therapeutic strategies.

In addition, based on the existing research results, precise intervention strategies can be developed. For example, for targeted bacterial inhibition technologies, exploration is required of the development of effective intervention strategies against oral microbial pathogens, such as *P. gingivalis* and *F. nucleatum*, including precision-targeted inhibitors, antibiotics and oral cleansers. These strategies aim to reduce the number or pathogenicity of pathogenic bacteria in the oral cavity, thereby reducing the risk of RI. For immunomodulatory strategies, oral mucosa-targeted vaccines can be developed to induce specific IgA responses, and phage therapy can be explored to selectively remove drug-resistant pathogenic bacteria. Microecological remodeling techniques, such as OMT to rebuild healthy flora, are able to construct pH-responsive slow-release systems to maintain oral microenvironmental homeostasis.

Notably, research on the oral microbiota and RI across anatomical sites involves multiple disciplines, including dentistry, oncology, radiation oncology, immunology and systems-level expertise. Consequently, enhanced interdisciplinary collaboration is required to accelerate research progress in this domain. Through integration of cross-disciplinary resources and technical capabilities, researchers can more comprehensively elucidate the intrinsic mechanisms underlying this association and develop more effective preventive and therapeutic strategies.

In the future, a multimodal diagnosis and treatment system can also be established, integrating a rapid saliva flora detection chip, an artificial intelligence-assisted diagnostic system and a personalized flora regulation program to establish a closed-loop management system of early warning of the risk of RI, early intervention and efficacy assessment.

5. Patient management strategies for radiotherapy

The management of patients with malignant tumors throughout the course of their illness will be an indispensable medical treatment in the future. The radiotherapy-phase management system has sparked considerable deliberation among healthcare professionals, prompting the emergence of a multitude of management modalities to mitigate the incidence and severity of RI (83-85). Zhao *et al* (108) demonstrated that the bound healthcare cohort management model can effectively prevent RILI and improve the quality of life of patients. The core of the bound medical and nursing cohort management model lies in the establishment of a medical and nursing bound care team, with members performing their respective roles to provide health education, basic care, psychological intervention and continuity of care for patients. Wang *et al* (109) found that the application of the 4F (all-weather, all-process, all-system and all-service) nursing management model in patients who underwent nasopharyngeal carcinoma radiotherapy had a very good effect, helping to enhance treatment engagement, improve self-care ability, promote quality of life and reduce the incidence of radiotherapy-related adverse reactions. The

retrospective cohort study by Xiao *et al* (110) illustrated that the 4R (reduction, readiness, response and recovery) crisis management theory could significantly reduce the incidence and severity of RE in patients during radiotherapy for breast malignant tumors. These evidence-based management strategies provide actionable insights for RI patient care, effectively enhancing patient engagement and self-management capacity.

6. Summary and outlook

With marked advances in radiotherapy technology, its clinical applications in oncology have become increasingly widespread. However, there are still substantial challenges in the clinical management of RI, particularly regarding accurate diagnosis and evidence-based prevention and therapeutic strategies. The purpose of this review is to establish a systematic and effective microbiota-based intervention through the in-depth study of oral microbiota and different categories of RI. The present review describes the association between the two, and the potential prevention and treatment strategies. Nevertheless, critical questions persist regarding the precise mechanisms through which the oral microbiota modulate radiotherapy outcomes, while these proposed prevention and treatment strategies require rigorous clinical validation to assess their clinical feasibility and safety profiles.

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Authors' contributions

JS drafted the manuscript and contributed to writing the content. LX drafted the manuscript and revised the content. Both authors have read and approved the final manuscript. Data authentication is not applicable.

Ethics statement and consent to participate

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Competing interests

The authors declare that they have no competing interests.

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