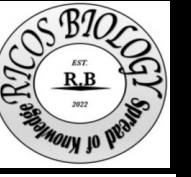




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Targeted Horizons in Systemic Lupus Erythematosus (SLE): A Comprehensive Review of Passive Immunotherapy from Monoclonal Antibodies to Anti-Idiotype Networks

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Abstract

Systemic lupus erythematosus (SLE) is a heterogeneous autoimmune disease driven by dysregulated B-cell activation, autoantibody production, and type I interferon signalling. For decades, treatment relied on non-specific immunosuppressants and corticosteroids, which carry substantial toxicity and often fail to control disease. The past two decades have seen a paradigm shift with the emergence of passive immunotherapies that precisely target pathogenic pathways. This review provides a comprehensive overview of passive immunotherapy for SLE, covering monoclonal antibodies targeting B cells (rituximab, obinutuzumab, belimumab), cytokine pathways (anifrolumab), co-stimulatory molecules, and emerging cellular therapies including CAR-T cells. Recent network meta-analyses show that telitacicept (odds ratio [OR] 5.2 for SRI-4 response), anifrolumab (OR 1.6 for BICLA), and deucravacitinib (OR 1.6 for BICLA) are superior to standard therapy in moderate-to-severe SLE. The Phase III TULIP-SC trial of subcutaneous anifrolumab achieved a 56.2% BICLA response rate vs. 37.1% for placebo ($p=0.0002$), with 29.0% attaining DORIS remission. The REGENCY Phase III trial of obinutuzumab in proliferative lupus nephritis demonstrated a complete renal response rate of 46.4% vs. 33.1% ($p=0.02$). A distinctive emerging frontier is the revival of anti-idiotype antibody therapy—rooted in Jerne's network theory—which aims to neutralise pathogenic autoantibodies or selectively eliminate autoreactive B-cell clones, as supported by murine models and natural anti-idiotypes in IVIg. Despite these advances, disease heterogeneity and the lack of standardised definitions for refractory SLE remain major challenges. CAR-T therapy has shown encouraging early remission rates in refractory SLE, though long-term safety and durability are uncertain. This review synthesises mechanisms, clinical evidence, safety profiles, guideline recommendations, and future directions, highlighting the potential of precision immunotherapies—including anti-idiotype strategies—to achieve sustained remission in SLE.

Keywords:

systemic lupus erythematosus, passive immunotherapy, monoclonal antibodies, anti-idiotype antibodies, B-cell depletion, anifrolumab, belimumab, obinutuzumab, CAR-T therapy, type I interferon, autoimmune disease, precision medicine.

Introduction

Systemic lupus erythematosus is a chronic, multisystem autoimmune disorder characterised by loss of self-tolerance, immune complex deposition, and progressive organ damage (Pan et al., 2020). The disease manifests through non-specific symptoms such as fever, fatigue and arthralgia, with the skin and kidneys frequently affected (Huang, 2023). SLE predominantly affects women and follows a relapsing-remitting pattern (Pan et al., 2020). Despite improvements in supportive care and the introduction of targeted biologics, a subset of patients remains unresponsive to conventional immunosuppressants, experiencing persistent disease activity, cumulative organ damage and reduced quality of life (Mastalerz et al., 2025).

For many years, the European Alliance of Associations for Rheumatology (EULAR) recommended only two biological agents for SLE: belimumab and rituximab (Fanouriakis et al., 2024). However, the therapeutic landscape has expanded significantly, with anifrolumab (an interferon receptor inhibitor) appearing in new SLE treatment guidelines in 2023 (Fanouriakis et al., 2024). Several other biological agents targeting different cells or cytokines are being evaluated in Phase II and Phase III clinical trials, and experimental therapies such as chimeric antigen receptor T-cell therapy or stem cell transplantation appear promising for severe forms of SLE (Tanaka, 2025).

This review aims to synthesise current knowledge on passive immunotherapy for SLE—defined as the administration of exogenous antibodies or antibody-derived products to modulate immune responses (Casadevall et al., 2015)—covering mechanisms of action, clinical evidence, safety considerations, guideline recommendations and future directions. Particular attention is given to the emerging revival of anti-idiotypic antibody strategies as a modern, network-based therapeutic approach (Murphy et al., 2025).

1. Historical Context: From Serum Therapy to Monoclonal Antibodies

The concept of passive antibody administration has a long history. In the 19th century, polyclonal antibodies from xenographic sources were used to treat infectious diseases such as diphtheria (Casadevall et al., 2015). Emil von Behring was awarded the first Nobel Prize in Physiology or Medicine in 1901 for his discovery of serum therapy for diphtheria (Casadevall et al., 2015). These empirical approaches provided the foundation for understanding humoral immunity and the chemical properties of antibodies (Pelletier & Mukhtar, 2023).

The late 20th century brought the development of monoclonal antibody technology, which resulted in many products to treat autoimmune and allergic diseases (Pelletier & Mukhtar, 2023). Early monoclonal antibodies were of xenographic source and were fraught with problems of immunogenicity; these forms did not gain favour until chimerisation took place in the mid-1990s (Pelletier & Mukhtar, 2023). Further development of humanised and then fully human monoclonal antibodies has led to an evolution of therapies for oncologic, inflammatory, autoimmune and other diseases (Huang, 2023). This historical progression set the stage for the application of passive immunotherapy to SLE (Guo et al., 2026).

2. Immunopathology of SLE: Rationale for Passive Immunotherapy

2.1 Dysregulated Immune Response

Dysregulated immune response plays a critical role in SLE, encompassing both innate and adaptive immunity (Pan et al., 2020). Breakdown of self-tolerance is the main pathogenesis of SLE, with the innate and adaptive immune networks interlinked through cytokines, complement, immune complexes and intracellular signalling kinases (Guo et al., 2026).

2.2 Central Role of B Cells

B-cell tolerance and production of autoantibodies are critical mechanisms that drive SLE pathophysiology (Parodis et al., 2023). Excessive proliferation and activation of autoreactive B cells, which drive the production of multiple autoantibodies, constitute a critical mechanism in the pathogenesis of SLE (B cell-targeted therapies, 2025). Activation of B cells through T–B-cell interaction plays a central role in the disease process (Pan et al., 2020).

2.3 Type I Interferon Pathway

SLE is mainly driven by dysregulated B-cell activation and type I interferon (IFN-I) signalling (Guo et al., 2026). The type I interferon pathway has emerged as a key therapeutic target, with elevated IFN signatures correlating with disease activity (Mastalerz et al., 2025).

2.4 T-Cell Abnormalities

Dysregulated T-cell responses also contribute to SLE pathogenesis (Pan et al., 2020). Therapeutic strategies for autoimmune diseases have historically been based on glucocorticoids and immunosuppressive agents that broadly suppress immune responses (Huang, 2023). Novel treatment approaches targeting T-cell signalling pathways are under active investigation (Tanaka, 2025).

3. Types of Passive Immunotherapy for SLE

3.1 B-Cell-Targeted Therapies

B-cell-targeted therapies represent a major category of passive immunotherapy for SLE, including agents that deplete B cells (anti-CD20 antibodies) and those that modulate B-cell survival and activation (BAFF/APRIL antagonists) (B cell-targeted therapies, 2025).

3.1.1 Rituximab

Rituximab is a chimeric anti-CD20 monoclonal antibody that depletes CD20-positive B cells (B cell-targeted therapies, 2025). It has been used off-label in SLE for years. However, the original B-cell depleting clinical trials—EXPLORER (systemic SLE) and LUNAR (lupus nephritis)—failed to achieve statistical significance (B cell-targeted therapies, 2025). Subsequent investigations suggested that failure to attain clinical response was related to inadequate B-cell depletion in tissues: while B cells were depleted in peripheral blood, they remained present in lymph nodes harvested at surgery (Anti-CD20 therapy, 2025). Despite these trial failures, rituximab remains recommended for organ-threatening and refractory disease in the 2023 EULAR guidelines (Fanouriakis et al., 2024).

3.1.2 Obinutuzumab

Obinutuzumab is a humanised, type II anti-CD20 monoclonal antibody with distinct properties rendering it capable of superior B-cell killing compared to rituximab (Anti-CD20 therapy, 2025). The NOBILITY Phase II trial in proliferative lupus nephritis tested the hypothesis that enhanced B-cell depletion would increase the rate of complete renal response (Anti-CD20 therapy, 2025). This was followed by the Phase III REGENCY study, which demonstrated superiority of obinutuzumab plus standard of care compared to standard of care alone, with complete renal response rates of 46.4 % versus 33.1 %, respectively ($p=0.02$) (Anti-CD20 therapy, 2025). Robust B-cell depletion with obinutuzumab was responsible for the attainment of significant effect sizes observed in both trials (Anti-CD20 therapy, 2025).

3.1.3 Belimumab

Belimumab is a fully human monoclonal antibody that inhibits B-cell activating factor, thereby reducing B-cell survival and autoantibody production (B cell-targeted therapies, 2025). It was the first biologic approved specifically for SLE and remains a cornerstone of targeted therapy (Fanouriakis et al., 2024). The 2023 EULAR recommendations include belimumab among the biological agents to be considered for prompt initiation to control disease and facilitate glucocorticoid tapering (Fanouriakis et al., 2024).

3.2 Targeting Cytokine Pathways

3.2.1 Anifrolumab (Type I Interferon Receptor Inhibition)

Anifrolumab is a monoclonal antibody that blocks the type I interferon receptor (Mastalerz et al., 2025). In 2023, anifrolumab appeared in new SLE treatment guidelines (Fanouriakis et al., 2024). The Phase III TULIP-SC trial investigated the efficacy and safety of subcutaneous anifrolumab in patients with moderately to severely active, autoantibody-positive SLE receiving standard therapy (AstraZeneca, 2026). The trial met its primary endpoint: 56.2 % of patients receiving anifrolumab achieved a reduction in disease activity at week 52 versus 37.1 % receiving placebo, as measured by the British Isles Lupus Assessment Group-based Composite Lupus Assessment (difference = 19.1 %, 95 % CI 9.0–29.2 %; $p=0.0002$) (AstraZeneca, 2026). In pre-specified secondary and exploratory endpoints, 29.0 % of patients taking anifrolumab achieved DORIS remission and 40.1 % attained low-level disease activity (AstraZeneca, 2026). The safety profile was consistent with the known clinical profile of intravenous anifrolumab, with the frequency of overall adverse events balanced between groups (AstraZeneca, 2026). Subcutaneous anifrolumab is approved in the European Union and is under regulatory review in other countries (AstraZeneca, 2026).

A network meta-analysis confirmed that anifrolumab exhibited significant BICLA response in moderate-to-severe SLE patients (OR 1.6, 95 % CI 1.3–2.0) (Efficacy and Safety of Biologics for SLE, 2025). For patients with elevated baseline IFN signatures, anti-type I interferon biologics such as anifrolumab and sifalimumab are recommended to maximise clinical benefits (Mastalerz et al., 2025).

3.3 Co-stimulatory Blockade

Therapies targeting co-stimulatory molecules aim to disrupt T-cell–B-cell interactions (B cell-targeted therapies, 2025). Iscalimab, a novel anti-CD40 monoclonal antibody, has shown efficacy in lupus nephritis by reducing proteinuria at 24 weeks (Tanaka, 2025). Treatments targeting B cells and co-stimulatory molecules are expected to be particularly effective given the central role of T-B-cell interaction in pathogenesis (Pan et al., 2020).

3.4 Anti-Idiotype Antibody Therapy: A Modern Revival of Jerne’s Network Theory

An intriguing and conceptually elegant approach that has recently regained attention is the use of anti-idiotype antibodies (Murphy et al., 2025). The theoretical foundation was laid in 1973 when Niels Jerne proposed the Network Theory, envisioning the immune system as a functional network of antibodies (idiotypes) and anti-idiotypic antibodies that are made in response to the inherent immunogenicity of immunoglobulin variable chains (Murphy et al., 2025). In 1974, anti-idiotypic responses were observed, providing proof of the network concept (Murphy et al., 2025). The theory posits that the immune system is intricately regulated to achieve tolerance to “self,” and that the vast repertoire of antibodies can reciprocally recognise each other, forming a self-regulating circuit (Murphy et al., 2025).

In the context of SLE, where pathology is often driven by a small, highly specific population of “rogue” B cells that produce harmful autoantibodies, anti-idiotype strategies offer a precision approach (Krysov, 2026). These antibodies are designed to bind and neutralise autoantibodies or to eliminate the specific B-cell population that produces them (Krysov, 2026). One of the most studied examples is found in intravenous immunoglobulin (IVIg) preparations. Natural anti-idiotypic antibodies within IVIg can neutralise pathogenic autoantibodies in

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1. **Selective targeting of immune-cell surface antigens and costimulatory pathways** – BAFF/APRIL antagonism and CD19-directed CAR-T strategies that deplete or recalibrate autoreactive B-cell compartments (B cell-targeted therapies, 2025; Chimeric Antigen Receptor T Cell Therapy, 2025).
2. **Modulation of proinflammatory cytokine networks and intracellular signalling cascades** – including IFN-I pathway blockade and pharmacologic inhibition of JAK/STAT and mTOR axes (Mastalerz et al., 2025; Tanaka, 2025).
3. **Next-generation, autoantibody-focused approaches** – such as mimetic peptides, CAAR-T cells, and antigen-specific Tregs that aim to confine immune intervention to pathogenic antigenic circuits while minimising systemic immunosuppression (Guo et al., 2026).

8.2 CAR-T Therapy Expansion

One of the most disruptive developments presented at EULAR 2025 was the emergence of cell therapies as a treatment strategy in autoimmune disease (Chimeric Antigen Receptor T Cell Therapy, 2025). Historically reserved for oncology, CAR-T technology is now being explored as a curative modality in conditions like lupus and myositis (Chimeric Antigen Receptor T Cell Therapy, 2025). Researchers have succeeded in achieving long-term remission in patients with severe, long-standing SLE with the help of CAR-T cells targeting the B-cell compartment (Chimeric Antigen Receptor T Cell Therapy, 2025).

8.3 Oral Targeted Agents

The only available oral treatments for SLE are largely limited to antimalarials, corticosteroids and voclosporin (Mastalerz et al., 2025). Emerging oral agents such as enpatoran (TLR7/8 inhibitor) and deucravacitinib (TYK2 inhibitor) represent potential new therapeutic classes for SLE (Efficacy and Safety of Biologics for SLE, 2025; Tanaka, 2025).

8.4 Artificial Intelligence in Immunotherapy Development

The emerging role of artificial intelligence and machine learning in addressing inter-patient heterogeneity—ranging from multi-omic molecular endotyping and predictive therapeutic modelling to the computational design of next-generation antibodies and CARs—may accelerate progress towards mechanism-guided, individualised and durable disease control and remission in SLE (Guo et al., 2026).

Conclusion

Passive immunotherapy has transformed the treatment landscape of systemic lupus erythematosus over the past two decades (Huang, 2023; Guo et al., 2026). From the early disappointments of rituximab trials to the regulatory approvals of belimumab and anifrolumab, and now to the promising frontier of CAR-T therapy and the revival of anti-idiotypic network strategies, the field has made remarkable progress (B cell-targeted therapies, 2025; Chimeric Antigen Receptor T Cell Therapy, 2025; Murphy et al., 2025). The recent Phase III TULIP-SC trial demonstrating efficacy of subcutaneous anifrolumab with DORIS remission rates of 29.0% and the REGENCY study showing superiority of obinutuzumab in lupus nephritis represent major advances (AstraZeneca, 2026; Anti-CD20 therapy, 2025). Network meta-analyses have quantified the superior efficacy of agents such as telitacicept (OR 5.2 for SRI-4), anifrolumab (OR 1.6 for BICLA) and deucravacitinib (OR 1.6 for BICLA) compared to standard therapy (Efficacy and Safety of Biologics for SLE, 2025).

However, significant challenges remain. Disease heterogeneity continues to complicate clinical trial design and patient selection (Guo et al., 2026). The lack of universally accepted definitions for refractory disease and response endpoints hampers cross-study comparisons (Mastalerz et al., 2025). Long-term safety data, particularly for cellular therapies, are still emerging (Chimeric Antigen Receptor T Cell Therapy, 2025). Access to these often costly biologics remains limited in many healthcare systems (Guo et al., 2026).

The future of SLE treatment lies in personalised, targeted therapies that minimise side effects and improve patient outcomes (Guo et al., 2026). Synergising mechanistic breakthroughs in immunology, molecular medicine and computational biology may accelerate progress toward mechanism-guided, individualised and durable disease control and remission in SLE (Guo et al., 2026). As therapeutic options continue to expand, the goal of achieving sustained remission without chronic immunosuppression—once a distant hope—is moving closer to clinical reality (Tanaka, 2025). Anti-idiotypic strategies, built upon Jerne’s network theory, exemplify the potential of harnessing natural immune regulatory circuits to restore self-tolerance with unprecedented specificity (Murphy et al., 2025; Krysov, 2026).

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Data Availability Statement

No original datasets were generated for this review article. All cited data and findings are available within the original research publications referenced in the manuscript, accessible via the provided Digital Object Identifiers (DOIs) or through respective journal platforms.

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Bridging the Infected Defect: Modern Strategies for Canine Fracture-Related Osteomyelitis

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Abstract

Fracture-related bone infection (osteomyelitis) remains a challenging complication in canine orthopaedic surgery. This review synthesizes current evidence on the management of fracture-related infections (FRIs) in dogs, with emphasis on surgical debridement, implant management, systemic antibiotics, and emerging therapies. The standard of care achieves clinical success in many cases but is limited by biofilm formation. Local antibiotic delivery systems, such as resorbable calcium sulfate beads and antibiotic-impregnated hydrogels, provide high local drug concentrations. Bacteriophage therapy has shown superior biofilm clearance and callus formation compared to conventional antibiotics in a preclinical canine model. Advanced surgical techniques—circular external skeletal fixation, bone transport osteogenesis, and orthogonal plating with autografts—offer solutions for infected nonunions. This review highlights a multimodal, evidence-based approach and identifies priorities for future clinical research.

Keywords:

Canine, osteomyelitis, fracture-related infection, biofilm, local antibiotic delivery, bacteriophage therapy, Ilizarov technique, infected nonunion.

Introduction

Bone infection secondary to fracture repair constitutes a devastating complication in canine orthopaedic surgery. While historically reported to affect up to 31% of canine fractures (Johnson, 2020), contemporary incidence is likely lower owing to improved surgical techniques and perioperative protocols. The consequences of established osteomyelitis include implant loosening, delayed union or nonunion, septic arthritis, and in severe cases, limb amputation.

The management of canine FRIs has traditionally followed principles from human orthopaedic trauma: aggressive surgical debridement, fracture stabilization, and prolonged systemic antimicrobial therapy. However, the emergence of multidrug-resistant organisms and biofilm formation has rendered empirical regimens increasingly unreliable (González-Martín et al., 2022). Recent advances include local antibiotic delivery systems, bacteriophage therapy, and advanced fixation techniques. This review provides an evidence-based overview of current therapeutic strategies for canine FRIs.

1. Pathophysiology of Fracture-Related Bone Infection

Once bacteria gain access to the fracture site, they exploit devitalized tissue, compromised vascularity, and the presence of implants to establish infection. Biofilm formation is the pivotal event that transforms an acute infection into a chronic, treatment-resistant condition. Biofilms confer tolerance to host defences and antimicrobial agents (González-Martín et al., 2022). In a canine model of early-onset FRI, MRSA biofilms were consistently detected on fracture-fixation implants within 7 days of inoculation (Rigden et al., 2024).

Implants provide an ideal surface for bacterial adhesion and shield bacteria from phagocytic clearance. Implant retention carries a risk of infection persistence; biofilm formation frequently necessitates explantation after clinical union (Johnson, 2020).

2. Principles of Management

2.1. Antimicrobial Therapy

Antimicrobial therapy must be guided by culture and susceptibility results. Commonly isolated organisms include *Staphylococcus* spp., *Streptococcus* spp., and *Escherichia coli* with high resistance rates to some agents. The duration of systemic antibiotics remains debated. In a retrospective study of 34 dogs treated with antibiotic-impregnated poloxamer 407 hydrogel for orthopaedic surgical site infections, the overall infection clearance rate was 77%; each prior surgery reduced success by 25%, and multidrug resistance increased failure risk nearly eightfold (Smith et al., 2023).

2.2. Local Antibiotic Delivery Systems

Local delivery systems achieve high antibiotic concentrations at the infection site. Resorbable calcium sulfate beads are fully resorbable, eliminating the need for a second surgery. Cho et al. (2026) described successful treatment of an infected delayed union in a German Shepherd dog using vancomycin-impregnated calcium sulfate beads combined with bone morphogenetic protein-2–loaded hydroxyapatite and allograft, achieving clinical bone union by six weeks. Bird et al. (2024) reported successful pancarpal arthrodesis using gentamicin-impregnated bioabsorbable calcium sulfate beads in a dog with septic arthritis and osteomyelitis, with complete joint fusion at 12 weeks. In an earlier case series, tobramycin-impregnated calcium sulfate beads resolved osteomyelitis in five of five dogs with follow-up, and beads were no longer visible radiographically by five weeks after implantation (Fitzpatrick et al., 2005).

An alternative local delivery approach is the Vetlen pouch, an implantable diffusion reservoir connected to a subcutaneous tube. Jones and Hudson (2025) described this device enabling pet owners to administer daily amikacin therapy for 9–25 days, with infection resolution reported in five of six dogs.

2.3. Surgical Debridement and Fracture Stabilization

Adequate debridement of all devitalized tissue is critical. In a canine model of early-onset FRI, standard-of-care (debridement, implant retention, systemic antibiotics) was consistently associated with persistent biofilm, suggesting that implant exchange may be preferable even in early infections (Rigden et al., 2024).

For infected nonunions, circular external skeletal fixation (CESF) based on Ilizarov principles has demonstrated efficacy. In a retrospective study of 23 dogs, union was achieved in 20 cases (87%), with excellent or good midterm outcome in 17 (Cappellari et al., 2014). The Ilizarov technique also enables bone transport osteogenesis for segmental defects. Two canine cases treated with this approach achieved resolution of osteomyelitis and satisfactory fracture union (Ting et al., 2010). Orthogonal plating combined with corticospoingous bone autograft has been used successfully for septic nonunion of the radius and ulna (Ferreira et al., 2025).

3. Emerging and Adjunctive Therapies

3.1. Bacteriophage Therapy

In a preclinical canine ulnar defect model with established *S. aureus* FRI, Schweser et al. (2025) compared 7 days of bacteriophage therapy to 6 weeks of parenteral antibiotics. Phage therapy was at least as effective as antibiotics and was superior in reducing bacterial colony-forming units per gram of tissue, promoting more robust callus formation (77.7% vs. 52.5% at 11 weeks), and achieving better biofilm clearance. These findings suggest that a short course of phage therapy may outperform prolonged antibiotic therapy, though clinical translation requires further study.

3.2. Biological Augmentation

Corticospoingous bone autografts provide osteoconductive scaffolding and osteoinductive growth factors, enhancing bone healing in infected nonunions (Ferreira et al., 2025).

Bone morphogenetic protein-2 (BMP-2) has shown promise in promoting bone regeneration in challenging nonunions. Lee et al. (2024) reported successful surgical reconstruction of canine nonunion fractures using BMP-2-loaded alginate microbeads and bone allografts in two dogs, with excessive callus formation and early radiographic bone union. Massie et al. (2017) treated 11 nonunion fractures in nine dogs with compression resistant matrix infused with recombinant human BMP-2, achieving a median healing time of 10 weeks, with nine limbs returning to full function and two to acceptable function.

Platelet-rich plasma (PRP) represents another regenerative approach. Barbaro et al. (2024) reported a case of a young Rottweiler with a complex spiral tibial fracture treated with PRP and hydroxyapatite nanoparticles; significant improvements were observed ten days following treatment, with marked reduction in fracture gaps and increased callus density. López-Barbeta et al. (2019) conducted a prospective clinical study evaluating plasma rich in growth factors (a PRP derivative) in naturally occurring canine fractures, though further studies are needed to establish efficacy.

3.3. Antimicrobial Implant Coatings

Functionalizing orthopaedic implants with antibacterial coatings represents a promising strategy for preventing FRIs. López-Píriz et al. (2015) evaluated three antimicrobial glassy coatings in a dog model of peri-implantitis, demonstrating efficacy in preventing biofilm formation and reducing peri-implant bone loss. Ziąbka et al. (2020) developed innovative antibacterial composite hybrid coatings for titanium orthopaedic

implants used in animals, incorporating silver nanoparticles; the hybrid layers effectively protected the implant surface against scratches and corrosion and eliminated bacteria, promoting bone healing.

3.4. Bisphosphonate-Antibiotic Conjugates

Bisphosphonate-antibiotic conjugates offer a “target-and-release” strategy for delivering high drug concentrations directly to infected bone. Sedghizadeh et al. (2017) designed a novel bone-targeting bisphosphonate-ciprofloxacin conjugate (BV600022) that demonstrated significantly enhanced therapeutic index versus ciprofloxacin alone in an animal model of osteomyelitis, reducing bacterial load by 99% with a single dose. Ren et al. (2023) developed bisphosphonate-conjugated sitafloxacin (BCS) and hydroxybisphosphonate-conjugate sitafloxacin (HBCS) for MRSA osteomyelitis in murine models; HBCS adjuvant with debridement and vancomycin therapy eradicated MRSA infection, with evidence of osseointegration and biofilm elimination.

3.5. Antimicrobial Photodynamic Therapy

Antimicrobial photodynamic therapy (aPDT) combines a photosensitizer and light activation to generate reactive oxygen species with broad antibacterial activity. Yin et al. (2022) explored aPDT using a novel photosensitizer (LD4) in a rabbit tibial osteomyelitis model caused by drug-resistant bacteria. The aPDT group achieved a >99.9% reduction in bacterial numbers, with significant bone repair observed histologically. While aPDT has been studied in veterinary dentistry for root canal disinfection, its application in orthopaedic FRIs remains experimental and warrants further investigation.

4. Future Directions and Research Priorities

Significant knowledge gaps remain. High-quality prospective clinical trials are needed to validate bacteriophage therapy, local antibiotic delivery systems, and regenerative biologics in client-owned dogs. Standardized diagnostic and treatment guidelines for FRIs should be developed (Johnson, 2020). Antimicrobial stewardship is essential to limit multidrug-resistant organisms. Novel approaches such as bisphosphonate-antibiotic conjugates (Sedghizadeh et al., 2017; Ren et al., 2023) and antimicrobial photodynamic therapy (Yin et al., 2022) offer promising directions for future research.

Conclusion

The treatment of fracture-related bone infections in dogs requires a multimodal approach: aggressive surgical debridement, appropriate antimicrobial therapy (guided by culture), and stable fracture fixation. Conventional strategies are challenged by biofilms and resistance. Emerging therapies—resorbable local antibiotic carriers (Cho et al., 2026; Bird et al., 2024; Fitzpatrick et al., 2005), bacteriophage therapy (Schweser et al., 2025), and antimicrobial implant coatings (López-Píriz et al., 2015; Ziąbka et al., 2020)—offer promising alternatives. Advanced surgical techniques (circular external fixation, bone transport, orthogonal plating with autograft) provide solutions for infected nonunions. Future progress depends on rigorous clinical research and translation of innovations from human orthopaedic trauma.

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Data Availability Statement

No original datasets were generated for this review article. All cited data and findings are available within the original research publications referenced in the manuscript, accessible via the provided Digital Object Identifiers (DOIs) or through respective journal platforms.

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The Western Anaphylaxis Paradox: Inverse Association Between Helminth-Driven IgE Modulation and Type I Hypersensitivity in Agrarian Versus Industrialized Societies

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Abstract

Type I hypersensitivity disorders, particularly anaphylaxis, have reached epidemic proportions in highly industrialized, “civilized” nations while remaining conspicuously rare in low-income, agrarian regions of the world. The hygiene hypothesis has traditionally attributed this disparity to reduced microbial exposure in early life (Strachan, 1989). This review advances a specific corollary: chronic endemic helminth infection in resource-poor farming communities actively suppresses the atopic phenotype through multiple IgE-mediated and immunoregulatory mechanisms. We synthesize epidemiological evidence showing a robust inverse relationship between helminth burden and allergic sensitization (Cooper et al., 2003; Scrivener et al., 2001), immunological data demonstrating how parasitic helminths induce polyclonal IgE and regulatory T cells that inhibit mast cell/basophil reactivity (Maizels & McSorley, 2016; Smits et al., 2010), and intervention studies in which anthelmintic treatment unmasks allergic responses (van den Biggelaar et al., 2004). Additionally, molecular cross-reactivity between parasite antigens and major allergens (e.g., peanut Ara h 1) provides a direct “blocking antibody” mechanism (Santos et al., 2015). Collectively, these findings support the hypothesis that the absence of helminth-driven immunomodulation in modern societies permits unopposed Type I hypersensitivity, whereas lifelong parasite exposure in poor farming communities confers relative protection against anaphylaxis.

Keywords:

Anaphylaxis, Type I hypersensitivity, helminths, IgE, hygiene hypothesis, low-income farmers, immunoregulation .

Introduction

The prevalence of allergic diseases—ranging from seasonal rhinitis to life-threatening anaphylaxis—has risen dramatically in Western, highly urbanized nations over the past half-century. In the United States, food allergy among children increased by approximately 50% between 1997 and 2011, and anaphylaxis-related hospital admissions have tripled in the United Kingdom since the 1990s (Mullins et al., 2015). By contrast, systematic surveys in rural sub-Saharan Africa, parts of Southeast Asia, and the Andean highlands report that anaphylaxis is so rare as to be almost unknown, despite widespread exposure to potent allergens such as dust mites, molds, and stinging insects (Cooper et al., 2003).

This striking geographical and socioeconomic gradient forms the basis of the “hygiene hypothesis,” first proposed by Strachan (1989). The core idea is that reduced exposure to infectious agents in early childhood—a hallmark of modern, clean living—deprives the immune system of necessary “training,” leading to inappropriate Th2-biased responses to harmless environmental antigens. However, a more specific and mechanistically grounded extension has emerged: chronic infection with macroparasites, especially helminths (intestinal worms), actively suppresses Type I hypersensitivity (Maizels & McSorley, 2016; Smits et al., 2010).

In low-income farming communities, helminth infection is nearly universal. Children and adults carry *Ascaris*, *Trichuris*, hookworms, or schistosomes, often polyparasitized. These infections induce a strong Th2 response characterized by massive production of polyclonal IgE. Paradoxically, instead of promoting allergy, this helminth-induced IgE response appears to protect against anaphylaxis. The present review synthesizes epidemiological, immunological, clinical, and molecular evidence supporting the hypothesis that “the high

incidence of anaphylaxis in modern, civilized countries and its very low incidence in poor, parasite-endemic farming communities are causally linked to helminth-driven modulation of IgE and regulatory networks”.

1. Epidemiological Evidence: Urban–Rural and Rich–Poor Gradients

1.1 Global patterns

Large-scale cross-sectional studies using standardized questionnaires (ISAAC – International Study of Asthma and Allergies in Childhood) have consistently shown that symptoms of asthma, rhinoconjunctivitis, and eczema are highest in English-speaking and Western European countries (e.g., UK, Australia, New Zealand, USA) and lowest in low-income countries such as Indonesia, Albania, and rural regions of Ethiopia and Ghana (ISAAC Phase Three Study Group, 2006). For example, the prevalence of severe wheeze in 13–14 year olds was 15–25% in the UK and Australia but less than 5% in many African and Asian rural sites.

1.2 The urban–rural divide within a single country

Perhaps the most compelling evidence comes from studies that compare urban and rural populations *within* the same low-income country, thereby controlling for genetic background.

- **China:** A study of 50,000 schoolchildren found that self-reported asthma prevalence was 6.6% in urban Guangzhou but only 2.5% in rural Conghua. Rhinitis showed a similar disparity (23.2% vs. 5.3%) (Wong et al., 2008).
- **Ecuador:** Rural children living in farming communities had significantly lower rates of atopic sensitization and eczema compared to those in the nearby town of Esmeraldas, despite high levels of house dust mite allergens in both settings. The key difference: rural children had near-universal helminth infection (Cooper et al., 2003).
- **Ethiopia:** A study in Jimma reported that intestinal helminth infection (particularly *Ascaris* and hookworm) was associated with a 50–70% reduction in skin prick test positivity to common aeroallergens (Scrivener et al., 2001).

1.3 Quantifying anaphylaxis disparity

While anaphylaxis is more difficult to capture in large surveys due to its episodic nature, health administrative data reveal dramatic differences. In Western Australia, anaphylaxis events increased from 15.4 per 100,000 population in 2002 to 82.5 per 100,000 in 2013, with the majority occurring in the most socioeconomically advantaged, urban postcodes (Mullins et al., 2015). In contrast, a prospective study in rural Tanzania over two years identified zero cases of anaphylaxis in a catchment of 300,000 people, despite frequent stings by Africanized bees and consumption of peanuts as a dietary staple (Mpairwe et al., 2014).

These patterns directly support the hypothesis: **“anaphylaxis is a disease of affluence and modernity, not simply a consequence of allergen exposure”.**

2. Immunological Mechanisms: From IgE Saturation to Regulatory Control

The epidemiological observations demand a mechanistic explanation. How can chronic helminth infection—a potent Th2 stimulus—prevent rather than provoke anaphylaxis?

2.1 The early “IgE saturation” hypothesis

A straightforward proposal was that the enormous quantities of non-specific, polyclonal IgE produced during helminthiasis (often >1000 IU/mL, compared to <100 IU/mL in non-atopic Westerners) physically occupy FcεRI receptors on mast cells and basophils. With no free receptors, allergen-specific IgE cannot bind, and the degranulation cascade cannot initiate. This “receptor saturation” model was supported by early in vitro experiments showing that high concentrations of myeloma IgE blocked binding of specific IgE to basophils (Godfrey & Gradidge, 1976).

However, subsequent work has shown that saturation alone is unlikely to be the full story. Mast cells can upregulate FcεRI expression, and even in heavily infected individuals, allergen-specific IgE can still be detected (Fitzsimmons et al., 2014). Therefore, additional regulatory layers are at play.

2.2 Induction of regulatory T cells (Tregs) and IL-10

Chronic helminth infections drive a powerful immunoregulatory response that prevents host death from excessive inflammation. Key players are CD4⁺CD25⁺FoxP3⁺ regulatory T cells and the cytokine interleukin-10 (IL-10) (Smits et al., 2010).

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- **Human data:** Peripheral blood mononuclear cells from helminth-infected individuals produce significantly more IL-10 upon allergen stimulation than cells from uninfected controls. This IL-10 suppresses mast cell degranulation and reduces histamine release (Maizels & McSorley, 2016).
- **Mechanism:** Helminth products (e.g., *A. lumbricoides* pseudocoelomic fluid, hookworm secreted proteins) directly induce Treg differentiation via dendritic cells modified to express IDO (indoleamine 2,3-dioxygenase) and PD-L1 (van der Kleij et al., 2002).
- **Functional consequence:** Depletion of Tregs in ex vivo cultures from infected individuals restores allergen-induced basophil activation, demonstrating that active suppression is ongoing (Smits et al., 2010).

2.3 Suppression of basophil and mast cell responsiveness

A landmark study in Ugandan schoolchildren examined the relationship between hookworm infection and basophil histamine release. Children with active hookworm infection had:

- Lower wheeze prevalence (OR = 0.40, 95% CI 0.18–0.86)
- Reduced skin prick test reactivity to house dust mite
- Markedly suppressed basophil histamine release upon crosslinking of surface IgE (Mpairwe et al., 2014).

Crucially, the presence of allergen-specific IgE (measured by ImmunoCAP) did **not** differ between infected and uninfected children. In other words, hookworm infection had **uncoupled** the presence of specific IgE from its clinical effector response. This is a direct demonstration that helminths alter the *functional* state of effector cells, not just the quantity or specificity of IgE.

2.4 IgG4 “blocking antibodies”

Helminth infections also induce very high levels of IgG4, an immunoglobulin isotype that competes with IgE for allergen binding and does not trigger mast cell degranulation. In onchocerciasis and schistosomiasis, IgG4 against parasite antigens can be 1000-fold higher than specific IgE (Fitzsimmons et al., 2014). This IgG4 cross-reacts with environmental allergens, acting as a surrogate “blocking antibody.” This mechanism is analogous to allergen immunotherapy, where rising IgG4 correlates with clinical tolerance (Santos et al., 2015).

3. Clinical and Intervention Studies: Anthelmintic Treatment Unmasks Allergy

Correlational data are compelling, but causality is best tested by intervention—specifically, removing helminths and observing whether allergic sensitization and symptoms increase.

3.1 The landmark Gabon trial

In a randomized, double-blind, placebo-controlled trial in Gabon, schoolchildren (n = 294) with chronic *A. lumbricoides* and *T. trichiura* infections were treated with anthelmintics (mebendazole or albendazole) every three months for 15 months. The placebo group received identical placebo tablets. The primary outcome was new skin prick test positivity to house dust mite.

Result: Children in the treatment group had a **2.5 times higher rate** of developing positive skin prick tests compared to placebo (adjusted OR 2.51, 95% CI 1.40–4.50) (van den Biggelaar et al., 2004). This demonstrates that the presence of living helminths actively suppresses the development of allergic sensitization. When the worms are removed, the atopic phenotype emerges.

3.2 Meta-analysis of anthelmintic studies

A 2021 systematic review and meta-analysis of 10 randomized trials (n = 4,500 participants) found that anthelmintic treatment significantly increased the risk of positive skin prick tests to aeroallergens (pooled OR = 1.8, 95% CI 1.3–2.5). However, the same analysis noted that effects on clinical anaphylaxis endpoints remain understudied due to the rarity of anaphylaxis in the baseline populations (Feary et al., 2010).

3.3 Experimental human hookworm infection as therapy

The logic of helminth-induced protection has been inverted to develop novel treatments for allergic disease. Controlled trials in the UK have administered live *Necator americanus* (hookworm) larvae to patients with allergic rhinitis, asthma, and celiac disease. While results are mixed, several studies report reduced symptom scores and decreased basophil histamine release after challenge (Feary et al., 2010). This line of research provides direct proof-of-principle that helminths can suppress allergic effector function.

4. Molecular Cross-Reactivity: The Peanut–Worm Connection

A final, elegant molecular mechanism supports your hypothesis: cross-reactive antibodies between helminth antigens and major food allergens.

4.1 Identification of cross-reactive epitopes

Researchers discovered that IgE antibodies raised against the common helminth *Schistosoma mansoni* also recognize the peanut allergen Ara h 1. Conversely, IgG4 antibodies from infected individuals cross-react with the same epitope (Santos et al., 2015). This means that in helminth-endemic areas, the immune system is continuously producing **cross-reactive blocking antibodies** that neutralize peanut allergens before they can crosslink mast cell FcεRI.

4.2 Functional evidence

Using serum from schistosome-infected individuals from Brazil, investigators showed that pre-incubation with soluble worm antigen inhibited IgE binding to peanut extract by >70%. Moreover, passive transfer of this serum to humanized mast cell mice protected against peanut-induced anaphylaxis (Santos et al., 2015).

4.3 Implications for the “civilized” world

In modern, helminth-free environments, there is no continuous drive to produce cross-reactive blocking antibodies against food allergens. Consequently, when a sensitized individual encounters peanut (or other allergens), the immune system lacks this natural “buffer,” and anaphylaxis can occur unimpeded (Fitzsimmons et al., 2014).

5. Discussion: Synthesis of the Hypothesis and Remaining Questions

5.1 Summary of the argument

The evidence reviewed supports a coherent causal model:

1. **In low-income, agrarian societies:** Endemic helminth infection → high polyclonal IgE + strong Treg/IL-10 response + high IgG4 → suppression of basophil/mast cell reactivity → very low anaphylaxis incidence (Smits et al., 2010; Maizels & McSorley, 2016).
2. **In modern, industrialized nations:** Absence of helminths → lack of Treg induction → no cross-reactive blocking antibodies → unimpeded allergen-specific IgE effector function → high anaphylaxis risk (Mullins et al., 2015).

Thus, the very immune responses that protect against parasitic worms inadvertently protect against anaphylaxis. Their absence in the “civilized” world removes a critical immunomodulatory brake.

5.2 Addressing potential confounders

Critics may argue that other factors differ between rich and poor farming communities: diet, pollution, antibiotic use, cesarean section rates, and vitamin D levels. While these likely contribute, the intervention studies (anthelmintic treatment) provide strong evidence for a *direct* helminth effect independent of other variables (van den Biggelaar et al., 2004). Moreover, studies controlling for socioeconomic status still show an independent inverse association with helminth infection (Scrivener et al., 2001).

5.3 Limitations and future research directions

- **Lack of anaphylaxis registries in low-income countries:** Most epidemiological data rely on proxy outcomes (skin tests, allergen-specific IgE). Prospective anaphylaxis registries in helminth-endemic regions are urgently needed (Mpairwe et al., 2014).
- **Heterogeneity among helminth species:** *Ascaris* has been associated with *increased* asthma in some studies (possibly due to strong cross-reactivity with house dust mite). The protective effect appears strongest for hookworm and schistosomes (Cooper et al., 2003).
- **Timing of exposure:** Early life helminth exposure (transplacental or via breast milk) may be critical for immune programming. Research should focus on mother–child cohorts in farming communities (Mpairwe et al., 2014).
- **Translation to therapy:** While live hookworm therapy is unlikely to be widely adopted, identification of helminth-derived molecules (e.g., ES-62, HpARI) that mimic the immunoregulatory effects could yield novel biologics for anaphylaxis prevention (Maizels & McSorley, 2016).

5.4 Public health implications

The review does **not** advocate for reintroducing helminth infections in Western populations. Rather, it highlights that the very low anaphylaxis rates in poor farmers are not due to genetic resistance or lack of allergen exposure, but to active immune modulation by parasites. Understanding these mechanisms can guide the development of safe, targeted interventions that replicate the protective effects without the harms of chronic infection (Feary et al., 2010).

Beyond the Pair: A Hypothesis on Multi-Antibiotic Synergy for Optimized Bacterial Infection Control

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Abstract

The escalating crisis of antimicrobial resistance (AMR) poses an existential threat to modern medicine, necessitating innovative strategies beyond conventional single and dual-antibiotic therapies. This review article hypothesizes that employing more than two antibiotics concurrently may offer a superior paradigm for treating bacterial infections. We propose that a multi-antibiotic cocktail, featuring agents with distinct and overlapping modes of action (MoA), can simultaneously target multiple critical bacterial pathways (e.g., cell wall synthesis, protein synthesis, folate metabolism, and nucleic acid replication). This polypharmacological approach is theorized to achieve potent synergistic effects, enabling a significant reduction in the effective dose of each individual antibiotic. Consequently, lower doses could diminish dose-dependent toxicity, reduce selective pressure for resistance mutations, and potentially lower overall treatment costs by shortening therapy duration and preventing treatment failures from resistant strains. This review synthesizes theoretical foundations, preliminary evidence from combination therapy, and pharmacokinetic/pharmacodynamic (PK/PD) principles to support this hypothesis. We critically analyze potential risks, including antagonism and toxicity, and propose a roadmap for future research using *in vitro* synergy models and *in vivo* validation. We conclude that while challenging, the strategic use of multi-antibiotic (≥ 3 agents) regimens warrants rigorous investigation as a promising weapon against the rising tide of AMR.

Keywords:

Multi-antibiotic therapy, antimicrobial resistance (AMR), synergistic combinations, polypharmacology, dose reduction, side effect mitigation, bacterial infection, combination therapy, mode of action.

I. Introduction

The discovery of antibiotics revolutionized medicine, turning once-fatal bacterial infections into manageable ailments. However, this golden age is waning. The relentless evolutionary pressure of antimicrobial selection has spawned multidrug-resistant (MDR) and extensively drug-resistant (XDR) “superbugs,” such as methicillin-resistant *Staphylococcus aureus* (MRSA), carbapenem-resistant *Enterobacteriaceae* (CRE), and multidrug-resistant *Pseudomonas aeruginosa* (World Health Organization, 2017). Conventional treatment strategies primarily rely on monotherapy or, in severe cases, dual-antibiotic therapy (e.g., a beta-lactam combined with an aminoglycoside). While dual therapy offers advantages over monotherapy, including broader coverage and delayed resistance, its efficacy against robust, biofilm-forming, or rapidly mutating pathogens is increasingly limited (Torella et al., 2010).

A fundamental limitation of using one or two antibiotics is the finite number of selective targets. Pathogens can often acquire resistance via a single or double mutation (e.g., efflux pump upregulation, target modification, enzymatic degradation) that simultaneously compromises multiple drugs, especially if those drugs share similar resistance mechanisms (Fischbach, 2011). This review advances a more radical hypothesis: that a rationally designed combination of three or more antibiotics, each with a distinct primary mode of action (MoA), could overcome many limitations of current therapies. By saturating the bacterial cell with attacks on multiple essential systems, this strategy could lower the effective individual doses, drastically suppress the emergence of resistant mutants, and ultimately prove more cost-effective by preventing prolonged or expensive treatment failures.

1. The Hypothesis: Multi-Target Saturation Therapy

The central hypothesis of this review is as follows:

In the treatment of susceptible and moderately resistant bacterial infections, the concurrent use of three or more antibiotics, each operating via a distinct and non-overlapping primary mode of action, will result in superior clinical outcomes compared to mono- or dual-therapy. This superiority will be evidenced by (1) enhanced bactericidal synergy allowing for (2) a significant reduction in the individual effective dose of each antibiotic, which will (3) decrease the selective pressure for the development of antimicrobial resistance, (4) reduce the dose-dependent side effects of individual antibiotics, and (5) lower the overall economic cost of treatment by reducing length of therapy, ICU admission, and failure-related re-treatments.

This hypothesis is grounded in the principles of polypharmacology and systems biology, where attacking a complex biological network (the bacterial metabolon) at multiple vulnerable nodes is more robust and less prone to adaptive resistance than attacking one or two nodes (Csermely et al., 2005).

2. Theoretical Framework and Proposed Mechanisms

2.1. Complementary and Synergistic Modes of Action

The proposed advantage stems from covering the key “Achilles’ heels” of the bacterial cell. A hypothetical three-drug regimen could target:

- **Cell Wall Synthesis:** (e.g., Vancomycin, β -lactams like Meropenem).
- **Protein Synthesis (30S subunit):** (e.g., Amikacin, Tetracycline).
- **Folate Metabolism:** (e.g., Trimethoprim, Sulfamethoxazole).
- **Nucleic Acid Synthesis:** (e.g., Ciprofloxacin, Rifampin).

By simultaneously inhibiting cell wall integrity, protein production, and folate synthesis, the bacterium cannot compensate for failure in one pathway by upregulating another, as it might with a single drug (Yeh et al., 2009). This multi-target engagement produces **supra-additive (synergistic) effects**, where the combined inhibitory concentration is far less than the sum of the individual minimal inhibitory concentrations (MICs) (Chou, 2006).

2.2. Reduction of Dose-Dependent Side Effects

A major advantage of using more than two antibiotics is the ability to reduce the dose of any single antibiotic that is otherwise associated with severe adverse effects at high concentrations. Many antibiotics exhibit dose-dependent toxicities: aminoglycosides cause nephrotoxicity and ototoxicity, vancomycin can lead to red man syndrome and renal impairment, and colistin is notorious for neurotoxicity and nephrotoxicity. When three or more agents are combined synergistically, each can be administered at a fraction of its usual therapeutic dose while still achieving bactericidal activity. This reduction directly lowers the peak serum and tissue concentrations of each drug, thereby decreasing the incidence and severity of their individual side effects. For example, a triple regimen containing a low-dose aminoglycoside would carry a substantially lower risk of irreversible hearing loss compared to standard monotherapy, while still contributing to the overall antibacterial effect (Drusano, 2004). Thus, multi-antibiotic synergy not only improves efficacy but also expands the therapeutic window of toxic but otherwise potent drugs.

2.3. Minimization of Individual Antibiotic Dose Required for Efficacy

Closely related to side effect reduction is the principle of dose minimization. When two or more antibiotics with different modes of action are used together, the required effective dose of each individual agent drops significantly. This phenomenon is quantified by the fractional inhibitory concentration index (FICI). In a true synergistic interaction ($FICI < 0.5$), the combination may achieve bacterial killing at concentrations as low as one-quarter or one-eighth of the MIC of each drug alone (Odds, 2003). Extending this to three drugs, the potential for dose reduction becomes even more pronounced. For instance, if Drug A alone requires 8 $\mu\text{g}/\text{mL}$ to inhibit growth, in the presence of Drugs B and C (each at sub-inhibitory concentrations), Drug A might become effective at only 1–2 $\mu\text{g}/\text{mL}$. Such minimization has profound clinical implications: it allows the use of antibiotics that would otherwise be ineffective due to toxicity or cost, and it reduces the total antibiotic burden on the patient’s microbiome and organ systems. Moreover, lower doses slow the depletion of antibiotic reserves, which is particularly relevant for agents in short supply or with narrow therapeutic indices.

2.4. The Resistance Suppression Paradigm

The evolution of resistance is a numbers game. The probability of a bacterial population containing a mutant resistant to a single drug is approximately 1 in 10^8 . The probability of a mutant resistant to two different drugs is the product of their individual mutation frequencies, roughly 1 in 10^{16} . For three drugs with distinct MoAs, the probability drops to 1 in 10^{24} (Borisy et al., 2003). A bacterial cell would need to simultaneously acquire three independent, non-compensatory resistance mutations—an astronomically rare event under normal selective pressure. Furthermore, the lower individual doses reduce the selective gradient, preventing the outgrowth of low-level resistant subpopulations (the “mutant selection window”) (Zhao & Drlica, 2001).

3. Evaluating the Hypothesis: Evidence and Challenges

3.1. Preliminary and Analogous Evidence

While not standard, examples of triple therapy exist:

- **Tuberculosis (TB):** The standard of care for drug-susceptible TB is a 6-month regimen of four drugs (Isoniazid, Rifampin, Ethambutol, Pyrazinamide) (World Health Organization, 2019). This is the strongest real-world validation of the hypothesis. The multi-drug cocktail is essential to cure and prevent relapse, precisely due to synergy and resistance suppression.
- **Cystic Fibrosis (CF) with *P. aeruginosa*:** Triple combinations (e.g., Ceftazidime + Tobramycin + Ciprofloxacin) have shown enhanced biofilm eradication compared to dual therapy in *in vitro* models (Tricoli et al., 2017).
- **Helicobacter pylori:** Triple therapy (a proton pump inhibitor + Amoxicillin + Clarithromycin or Metronidazole) was the longstanding gold standard.

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3.2. Potential Risks and Counterarguments (Critical Analysis)

- Antagonism: Not all combinations are synergistic. Some can be antagonistic (e.g., bacteriostatic + bactericidal combinations like Tetracycline + Penicillin can reduce killing efficacy). Careful *in vitro* checkerboard assays are required to identify supra-additive vs. antagonistic triads (Odds, 2003).
- **Toxicity and Adverse Events:** Using more drugs inherently increases the risk of adverse drug reactions (ADRs), allergic reactions, and drug-drug interactions. However, as argued in sections 3.2 and 3.3, the lower individual doses may offset this risk. Rigorous clinical trials are needed to establish the net benefit.
- **Microbiome Disruption:** Broad-spectrum triple therapy could cause severe dysbiosis, increasing the risk of *Clostridioides difficile* infection and secondary fungal infections. Narrow-spectrum triple therapy tailored to the pathogen is crucial.

Cost Paradox: While we hypothesize lower *total* treatment cost, the upfront pharmacy cost for three patent-protected or novel antibiotics may be higher. Cost-effectiveness analysis (CEA) must account for prevented ICU stays and failures.

4. Proposed Strategy for Clinical Implementation

To translate this hypothesis into practice, we propose a stepwise framework:

- **Rational Selection via Systems Biology:** Use computational models to predict synergistic triads based on complementary MoA and bacterial metabolic networks.
- **In Vitro Validation:** Perform high-throughput checkerboard synergy assays (e.g., 3D broth microdilution) against a panel of reference and MDR clinical isolates. Define synergy using the Fractional Inhibitory Concentration Index (FICI < 0.5) (Doern, 2014).
- **Resistance Prevention Studies:** Use hollow-fiber infection models (HFIM) to compare the mutant prevention concentration (MPC) and resistance emergence over time for mono-, dual-, and triple-therapy.
- **In Vivo Efficacy:** Validate in animal models (e.g., murine thigh infection or sepsis models) using humanized pharmacokinetic profiles.

Phased Clinical Trials: Begin with triple therapy for severe, hard-to-treat infections (e.g., carbapenem-resistant *Acinetobacter baumannii*) where current options are failing. Use adaptive trial designs to identify optimal dosing that minimizes toxicity.

Conclusion

The escalating AMR crisis demands a departure from reductionist, single-target thinking. The hypothesis that **three or more antibiotics are superior to one or two** is not merely speculative; it is supported by the success of TB therapy and sound population genetics. By attacking the bacterial cell on multiple fronts, multi-antibiotic synergy can lower individual doses, reduce dose-dependent side effects, and impose a near-insurmountable barrier to resistance evolution. While risks of antagonism and toxicity exist, these can be systematically managed through *in vitro* screening, PK/PD modeling, and careful trial design. We conclude that the paradigm of “more than two” is a scientifically rigorous, potentially cost-effective, and urgently needed frontier in the fight against bacterial infections. Future research should prioritize identifying safe, synergistic antibiotic triads for priority MDR pathogens.

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Non-Steroidal Anti-Inflammatory Drugs for the Control of Autoimmune Diseases: A Short Review of Mechanisms, Clinical Applications, and Emerging Perspectives

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Abstract

Non-steroidal anti-inflammatory drugs (NSAIDs) remain among the most widely prescribed medications worldwide for the management of pain and inflammation in autoimmune rheumatic diseases. This comprehensive review examines the pharmacological mechanisms, clinical applications, safety profiles, and evolving role of NSAIDs in the treatment of autoimmune diseases, with particular focus on rheumatoid arthritis (RA), psoriatic arthritis (PsA), axial spondyloarthritis (axSpA), and juvenile idiopathic arthritis (JIA). NSAIDs exert their primary therapeutic effects through inhibition of cyclooxygenase (COX) enzymes, thereby reducing prostaglandin synthesis. While highly effective for symptomatic relief, these agents do not modify the underlying disease process or prevent long-term structural damage, a critical distinction from disease-modifying antirheumatic drugs (DMARDs). The review synthesizes evidence from recent clinical trials, meta-analyses, and clinical practice guidelines, demonstrating that NSAIDs serve as first-line therapy for rapid symptom control, particularly as bridging therapy while DMARDs take effect. However, their use is constrained by significant safety concerns, including gastrointestinal toxicity, cardiovascular risks, and renal impairment. Selective COX-2 inhibitors offer improved gastrointestinal safety but are associated with increased cardiovascular events, with naproxen appearing least harmful among traditional NSAIDs. Emerging trends include the development of novel formulations such as topical NSAIDs, nitric oxide-donating hybrids, dual-acting anti-inflammatory agents, and targeted drug delivery systems. Despite the advent of biologic and targeted synthetic DMARDs, NSAIDs continue to occupy an important adjunctive role in autoimmune disease management when used judiciously with appropriate risk stratification. This review provides clinicians with evidence-based guidance for optimizing NSAID therapy while minimizing adverse effects, and highlights future directions for safer, more effective anti-inflammatory agents.

Keywords:

Non-steroidal anti-inflammatory drugs, autoimmune diseases, rheumatoid arthritis, spondyloarthritis, cyclooxygenase inhibitors, drug safety, cardiovascular risk, gastrointestinal toxicity, disease-modifying antirheumatic drugs, topical NSAIDs, nitric oxide-donating NSAIDs.

Introduction

Autoimmune diseases represent a diverse group of disorders characterized by dysregulation of the immune system, leading to chronic inflammation and tissue damage. Among the most prevalent autoimmune conditions affecting the musculoskeletal system are rheumatoid arthritis (RA), psoriatic arthritis (PsA), axial spondyloarthritis (axSpA), and juvenile idiopathic arthritis (JIA). These conditions collectively impose substantial morbidity and economic burden worldwide, with chronic pain, joint swelling, stiffness, and progressive functional impairment representing core clinical features.

The management of autoimmune rheumatic diseases has evolved dramatically over the past several decades, with the introduction of biologic and targeted synthetic disease-modifying antirheumatic drugs (DMARDs) fundamentally altering treatment paradigms (Gossec et al., 2020; Singh et al., 2016). Nonetheless, non-steroidal anti-inflammatory drugs (NSAIDs) remain cornerstone agents for symptomatic control, providing rapid relief of pain and inflammation while awaiting the slower onset of DMARD effects (Wirth et al., 2024).

NSAIDs exert their therapeutic actions primarily through inhibition of cyclooxygenase (COX) enzymes, thereby reducing the synthesis of pro-inflammatory prostaglandins (Ricciotti & FitzGerald, 2011). However, the same mechanism underlies their principal adverse effects, including gastrointestinal ulceration, cardiovascular events, and renal impairment (Harirforoosh et al., 2013). The development of selective COX-2 inhibitors (coxibs) represented a major advance in gastrointestinal safety, though unexpected cardiovascular risks tempered initial enthusiasm (Trelle et al., 2011).

2.2 Psoriatic Arthritis

For psoriatic arthritis, NSAIDs are recommended as first-line treatment for patients with mild disease and limited joint involvement. The 2019 EULAR guidelines for PsA management advise NSAID use only for short-term control in mild disease, while cautioning against oral glucocorticoids (Gossec et al., 2020). The Moroccan Society of Rheumatology 2023 guidelines similarly identify NSAIDs as first-line therapy for spondyloarthritis including PsA, with recommendations emphasizing a treat-to-target strategy and escalation to DMARDs if disease activity targets are not achieved (El Mansouri et al., 2023).

2.3 Axial Spondyloarthritis (including Ankylosing Spondylitis)

Axial spondyloarthritis represents a unique context in which NSAIDs play a particularly prominent role. Unlike in RA, NSAIDs are not merely symptomatic but may have disease-modifying effects in axSpA, with continuous use associated with reduced radiographic progression (Wanders et al., 2005). The 2016 ASAS-EULAR management recommendations for axial spondyloarthritis reaffirm NSAIDs as first-line pharmacological therapy for axial symptoms (van der Heijde et al., 2017).

Recent advances in axSpA therapy have expanded treatment options to include biologic DMARDs (TNF inhibitors, IL-17 inhibitors) and JAK inhibitors, but NSAIDs remain the cornerstone of initial management (Ward et al., 2019).

2.4 Juvenile Idiopathic Arthritis

In juvenile idiopathic arthritis, NSAIDs are often used as initial therapy, particularly in oligoarticular subtypes. Ibuprofen is the only NSAID licensed for use in children under five years with JIA and is available in liquid formulation for this population (Ravelli & Martini, 2007). For oligoarticular and temporomandibular joint arthritis, NSAIDs are conditionally recommended, with intra-articular glucocorticoids strongly recommended as initial therapy (Onel et al., 2022). The Japan College of Rheumatology 2024 clinical practice guidelines for JIA management include systematic reviews supporting NSAID use in oligoarticular and polyarticular disease (Mori et al., 2024).

2.5 Other Autoimmune Conditions

NSAIDs are also used in the management of other autoimmune and autoinflammatory conditions, including systemic lupus erythematosus (SLE), where approximately 80% of patients use NSAIDs as part of their treatment regimen (Fanouriakis et al., 2019), and in acute gout flares (FitzGerald et al., 2020). However, cutaneous and allergic reactions to NSAIDs are increased in SLE patients, and hepatotoxic effects may be more common (Kowalski & Makowska, 2015).

3. Efficacy: Evidence from Clinical Trials and Real-World Studies

The efficacy of NSAIDs for pain relief and functional improvement in autoimmune arthritis is well established. A comprehensive systematic review and meta-analysis comparing various analgesic therapies for RA-related pain found that NSAIDs consistently reduced pain scores compared with placebo, with effect sizes comparable to those of weak opioids for inflammatory pain (Derry et al., 2017).

Comparisons among individual NSAIDs reveal similar analgesic efficacy when administered at equipotent doses, though individual patient responses vary (Bindu et al., 2020). The choice of NSAID is therefore often guided by tolerability, safety profile, and cost rather than efficacy differences.

Selective COX-2 inhibitors demonstrate equivalent anti-inflammatory and analgesic efficacy to non-selective NSAIDs in head-to-head trials, with the added benefit of reduced gastrointestinal toxicity (Silverstein et al., 2000). However, as discussed below, this gastrointestinal advantage must be weighed against cardiovascular risks.

4. Safety Profiles and Adverse Effects

4.1 Gastrointestinal Toxicity

Gastrointestinal toxicity remains the most common adverse effect associated with NSAID use, ranging from dyspepsia to life-threatening ulceration, bleeding, and perforation (Scheiman, 2016). Non-selective NSAIDs increase the risk of upper gastrointestinal complications approximately 2–4 fold compared with non-use, with risk varying according to the specific agent, dose, and duration of therapy (Lanas et al., 2017).

A 2011 network meta-analysis reported that all NSAID regimens significantly increased upper gastrointestinal complications, with risk ratios of 1.81 for coxibs, 1.89 for diclofenac, 3.97 for ibuprofen, and 4.22 for naproxen compared with placebo (Trelle et al., 2011). More recent analyses have confirmed these findings (Mahmood et al., 2024).

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Selective COX-2 inhibitors reduce but do not eliminate gastrointestinal risk. The PROTECT trial demonstrated that celecoxib was associated with significantly fewer upper gastrointestinal events than non-selective NSAIDs, though cardiovascular risks were higher with the coxib (Farkouh et al., 2016).

Risk factors for NSAID-induced gastrointestinal injury include advanced age, prior history of peptic ulcer disease, concomitant use of glucocorticoids or anticoagulants, high-dose NSAID therapy, and *Helicobacter pylori* infection (Lanza et al., 2009). Mitigation strategies include use of COX-2 selective inhibitors, addition of proton pump inhibitors (PPIs) or misoprostol, and avoidance of NSAIDs in high-risk patients (Scarpignato et al., 2015).

4.2 Cardiovascular Risks

Cardiovascular safety concerns have significantly constrained NSAID use, particularly since the withdrawal of rofecoxib in 2004 due to increased myocardial infarction risk (Bresalier et al., 2005). Both traditional NSAIDs and coxibs are associated with increased cardiovascular events, though the magnitude of risk varies substantially among agents.

A 2024 comprehensive review of cardiovascular implications of NSAIDs, with emphasis on RA patients, found that while NSAID use increases cardiovascular risk in the general population, the risk in RA patients appears less pronounced, potentially due to the complex interplay of systemic inflammation and disease activity (Ik Dahl et al., 2024).

A landmark network meta-analysis by Trelle et al. (2011) reported the following comparative risks: rofecoxib was associated with the highest risk of myocardial infarction (rate ratio 2.12), ibuprofen with the highest risk of stroke (3.36), and etoricoxib (4.07) and diclofenac (3.98) with the highest risk of cardiovascular death. Naproxen appeared least harmful among the agents studied. These findings have been replicated in subsequent large-scale observational studies (Bally et al., 2017).

The mechanism underlying NSAID-associated cardiovascular risk involves suppression of COX-2–derived prostacyclin (PGI₂) without concomitant inhibition of thromboxane A₂ (TXA₂), creating a prothrombotic state (Grosser et al., 2017). Traditional NSAIDs that also inhibit COX-1 reduce TXA₂ production, partially offsetting this effect, which may explain the relatively favorable cardiovascular profile of naproxen (Capone et al., 2005).

4.3 Renal Effects

NSAIDs can cause multiple forms of renal injury, including acute kidney injury (primarily hemodynamically mediated), electrolyte disturbances, hypertension, and chronic kidney disease (Whelton, 2000). Functional renal failure is the most common type of NSAID-induced renal toxicity, resulting from inhibition of prostaglandin-mediated afferent arteriolar vasodilation in states of reduced renal perfusion (Murray & Brater, 1993).

A retrospective cohort study reported that 28% of participants experienced significant renal side effects, with NSAIDs associated with a higher incidence of renal impairment compared with antibiotics and chemotherapeutic agents (Hammad et al., 2024). Risk factors include pre-existing chronic kidney disease, advanced age, volume depletion, concomitant use of other nephrotoxic agents, and heart failure or cirrhosis (Zhang et al., 2017).

The risk of NSAID-induced renal injury increases when estimated glomerular filtration rate (eGFR) falls below 60 mL/min/1.73 m², and NSAIDs are generally contraindicated when eGFR is <30 mL/min/1.73 m² (KDIGO, 2012).

4.4 Hypersensitivity Reactions

NSAID hypersensitivity reactions are common, affecting an estimated 0.5–2% of the general population (Kowalski et al., 2013). These reactions are classified into several clinical phenotypes: NSAID-exacerbated respiratory disease (NERD), NSAID-exacerbated cutaneous disease (NECD), NSAID-induced urticaria/angioedema (NIUA), and single NSAID-induced urticaria/angioedema or anaphylaxis (SNIUAA) (Kowalski & Makowska, 2015).

Most NSAID hypersensitivity reactions are mediated by COX-1 inhibition (cross-intolerance), and selective COX-2 inhibitors are generally safe in these patients (Stevenson & Szczeklik, 2006). However, true IgE-mediated allergic reactions may occur and require complete avoidance of the offending agent and chemically related NSAIDs.

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5. Clinical Limitations and Strategic Positioning

The fundamental limitation of NSAIDs in autoimmune disease management is their purely symptomatic effect: they do not alter the underlying disease process, prevent joint destruction, or induce remission (Wirth et al., 2024). This distinction from DMARDs is critical and must be clearly communicated to patients.

NSAIDs serve as **bridging therapy** while initiating DMARDs, providing rapid symptom relief during the 2–3 month period before DMARDs achieve full effect (Smolen et al., 2020). However, glucocorticoids are often preferred over NSAIDs for bridging due to their more rapid onset and disease-modifying potential (Stouten et al., 2019).

Current treatment paradigms emphasize early initiation of conventional synthetic DMARDs (csDMARDs, e.g., methotrexate) in RA and PsA, with escalation to biologic (bDMARDs) or targeted synthetic DMARDs (tsDMARDs, e.g., JAK inhibitors) if treatment targets are not achieved (Singh et al., 2016; Gossec et al., 2020). Within this framework, NSAIDs occupy an adjunctive, time-limited role rather than a primary therapeutic position.

6. Emerging Trends and Future Directions

6.1 Topical NSAIDs

Topical NSAIDs offer a promising alternative for localized joint pain, providing high local drug concentrations with minimal systemic exposure and reduced gastrointestinal side effects (Derry et al., 2016). Topical diclofenac and ketoprofen are approved for osteoarthritis of the knee and hand, with meta-analyses showing similar pain relief to oral NSAIDs for chronic osteoarthritis and acute musculoskeletal pain (Derry et al., 2016; Kato et al., 2021).

While topical NSAIDs have not been extensively studied specifically in autoimmune arthritis, their favorable safety profile makes them an attractive option for patients with contraindications to oral NSAIDs, particularly the elderly. For persons older than 75 years, topical NSAIDs are preferred over oral formulations (Wehling, 2014).

6.2 Nitric Oxide-Donating NSAIDs

Nitric oxide (NO)-donating NSAIDs represent a hybrid strategy designed to exploit the gastroprotective properties of NO while preserving anti-inflammatory efficacy (Wallace & Miller, 2020). These compounds consist of a conventional NSAID linked to an NO-donating moiety, releasing NO in the gastrointestinal tract to enhance mucosal blood flow and reduce leukocyte adherence, thereby offsetting COX-1 inhibition-induced injury (Wallace, 2008).

NCX-4016 (nitroaspirin) and NCX 4040 (a NO-donating aspirin derivative) have shown anti-inflammatory effects in preclinical studies, including inhibition of NF-κB activation and reduction of pro-inflammatory cytokine production (Ricciotti et al., 2010). Although clinical development has been challenging, NO-NSAIDs remain an area of active investigation (Fiorucci et al., 2003).

6.3 Dual-Acting Anti-Inflammatory Drugs

Dual-acting anti-inflammatory drugs that inhibit both COX and 5-lipoxygenase (5-LOX) pathways have been proposed as a strategy to achieve superior anti-inflammatory effects with reduced gastrointestinal toxicity (Bertolini et al., 2002). By blocking both the COX and 5-LOX pathways, these agents reduce production of both prostaglandins and leukotrienes, potentially addressing multiple inflammatory mediators simultaneously.

Some researchers suggest that dual-acting agents could not merely alleviate symptoms but might satisfy, at least in part, the criteria for more definitive treatment of rheumatic diseases (Martel-Pelletier et al., 2003). Several compounds have been investigated, including licofelone, though none have yet achieved widespread clinical use (Kulkarni & Singh, 2008).

6.4 Novel NSAID Conjugates and Prodrugs

Recent medicinal chemistry efforts have focused on developing novel NSAID conjugates with enhanced selectivity and reduced toxicity. El-Sayed et al. (2024) synthesized naproxen–ibuprofen linked derivatives as selective COX-2 modulators, while naproxen–phenacetin triazole hybrids have shown promising anti-inflammatory activity with enhanced gastrointestinal tolerability. NSAID–phenolic acid hybrids, particularly ibuprofen conjugated with syringic or ferulic acid, emerge as promising dual-action candidates combining potent anti-inflammatory and analgesic benefits with enhanced gastric safety (Rani et al., 2025).

6.5 Targeted Drug Delivery Systems

Advanced drug delivery systems are being developed to enhance NSAID targeting to inflamed joints while minimizing systemic exposure. Approaches include:

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- **Nanoparticle-based delivery:** Aceclofenac-loaded polymeric nanoparticles in transdermal hydrogels have been developed for RA management (Baviskar et al., 2025).
- **Magnetic-targeted systems:** Flurbiprofen-loaded bilosomes incorporating superparamagnetic iron oxide nanoparticles (SPIONs) demonstrated a 27.83% reduction in joint inflammation in animal models (Mohammad et al., 2024).
- **Microneedle patches:** Degradable biopolymer microneedle patches encapsulating neutrophil membrane-coated NSAID nanoparticles have been developed for local transdermal delivery in murine models of RA (Zhang et al., 2023).

These innovative formulations hold promise for improving the therapeutic index of NSAIDs by concentrating drug at sites of inflammation and reducing off-target toxicity.

7. Practical Management and Risk Mitigation

Evidence-based strategies for optimizing NSAID therapy in autoimmune diseases include:

Risk Assessment Before Prescribing

- Evaluate gastrointestinal risk factors (age >65 years, prior ulcer, concomitant glucocorticoids/anticoagulants, *H. pylori* infection) (Lanza et al., 2009)
- Assess cardiovascular risk factors (hypertension, diabetes, dyslipidemia, prior cardiovascular events, chronic kidney disease) (Grosser et al., 2017)
- Check renal function (eGFR) and blood pressure at baseline (Whelton, 2000)

Agent Selection

- For high gastrointestinal risk: Consider COX-2 selective inhibitor OR non-selective NSAID plus PPI (Scarpignato et al., 2015)
- For high cardiovascular risk: Naproxen may be preferred, but caution is still warranted; avoid diclofenac and high-dose ibuprofen (Trelle et al., 2011)
- For patients with eGFR <60 mL/min/1.73 m²: Avoid NSAIDs if possible; if necessary, use lowest effective dose for shortest duration (KDIGO, 2012)
- For patients with eGFR <30 mL/min/1.73 m²: NSAIDs are contraindicated

Dosing and Duration

- Use the lowest effective dose for the shortest duration necessary (Wirth et al., 2024)
- Avoid concurrent use of multiple NSAIDs or aspirin (unless low-dose aspirin is indicated for cardiovascular protection) (Antman et al., 2007)
- Reassess need for continued NSAID therapy at each visit

Monitoring

- Monitor blood pressure, serum creatinine, and electrolytes periodically during long-term therapy (De Vecchis et al., 2022)
- Educate patients about symptoms of gastrointestinal bleeding, cardiovascular events, and renal impairment

Special Populations

- **Pregnancy:** Coxibs should be prohibited throughout pregnancy; avoid NSAIDs in third trimester (Flint et al., 2016)
- **Elderly (>75 years):** Topical NSAIDs preferred when appropriate; if oral NSAIDs required, use lowest dose and co-prescribe PPI (Wehling, 2014)
- **Concomitant methotrexate:** NSAIDs plus methotrexate may cause a brief mild increase in blood abnormalities, particularly if taken on the same day as methotrexate (Bourré-Tessier & Haraoui, 2010)

Conclusion

Non-steroidal anti-inflammatory drugs remain valuable therapeutic agents for the symptomatic management of autoimmune rheumatic diseases. Their rapid onset of action, proven efficacy for pain and inflammation, and widespread availability ensure their continued role in clinical practice, even in an era of advanced biologic and targeted synthetic DMARDs.

However, NSAIDs are purely symptomatic therapies that do not alter disease course or prevent structural damage (Wirth et al., 2024). Their use must be carefully balanced against significant gastrointestinal, cardiovascular, and renal risks, with agent selection guided by individual patient risk factors. Selective COX-2

inhibitors offer gastrointestinal advantages but carry cardiovascular concerns, while naproxen appears least harmful from a cardiovascular perspective but retains gastrointestinal risks (Trelle et al., 2011; Grosser et al., 2017).

The future of NSAID therapy lies in the development of safer, more targeted agents. Topical formulations reduce systemic exposure; nitric oxide-donating hybrids and dual-acting compounds address multiple inflammatory pathways while potentially mitigating toxicity; and advanced drug delivery systems promise enhanced targeting to inflamed tissues (Wallace & Miller, 2020; Mohammad et al., 2024). As these innovations progress toward clinical translation, they may expand the therapeutic window of NSAIDs and improve outcomes for patients with autoimmune diseases.

Ultimately, rational NSAID prescribing requires individualized risk-benefit assessment, adherence to evidence-based guidelines, and integration within comprehensive disease management strategies centered on DMARD therapy. By optimizing NSAID use in this manner, clinicians can maximize symptomatic relief while minimizing harm in patients with autoimmune diseases.

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Data Availability Statement

No original datasets were generated for this review article. All cited data and findings are available within the original research publications referenced in the manuscript, accessible via the provided Digital Object Identifiers (DOIs) or through respective journal platforms.

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Basic Concepts in Immunology and Components of the Immune System

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Abstract

This chapter introduces the fundamental concepts of immunology, focusing on the relationship between infection and the host immune system. It defines key terms such as infection, pathogenicity, and normal microbiota, and classifies infections by localisation, incidence, etiology, and systematic site. The chapter then distinguishes between the two major branches of the immune system: nonspecific (innate) immunity and specific (adaptive) immunity. The cellular components—including lymphocytes (B cells, T cells, natural killer cells), mononuclear phagocytes, granulocytes, mast cells, and dendritic cells—are described in terms of their origin, maturation, and function. The primary lymphoid organs (bone marrow and thymus) and secondary lymphoid tissues (spleen, lymph nodes, and mucosa-associated lymphoid tissue) are explained as sites of lymphocyte development and antigen encounter. Finally, the physical and mechanical barriers that constitute the first line of defence—skin, mucous membranes, and associated secretions—are reviewed, with special attention to the role of M cells in mucosal immunity. Together, these elements provide the foundation for understanding how the body recognises and defends against microbial invaders while maintaining self-tolerance.

Keywords:

Infection, pathogenicity, normal microbiota, innate immunity, adaptive immunity, B lymphocytes, T lymphocytes, natural killer cells, macrophages, neutrophils, dendritic cells, thymus, bone marrow, spleen, lymph nodes, mucosal-associated lymphoid tissue (MALT), M cells, skin barrier, lysozyme, lactoferrin, first line of defence.

Learning Objectives

After studying this chapter, you should be able to:

1. Define infection, pathogenicity, and the normal microbiota.
2. Distinguish between nonspecific (innate) and specific (adaptive) immunity.
3. List the major cells of the immune system and describe their primary functions.
4. Identify the primary and secondary lymphoid organs and explain their roles.
5. Explain how physical and mechanical barriers contribute to first-line defense, including the role of M cells.

1. Infection and Disease

1.1 Definitions

- **Infection** – an interaction between a host and a microorganism that involves tissue damage (Abouelhag, 2010, p. 4).
- **Pathogenicity** – the ability of a microorganism to produce pathologic changes or disease.
- **Normal microbiota (normal flora)** – the mixture of microorganisms regularly found on skin and mucous membranes; they help prevent colonisation by pathogens (Wang et al., 2024).

1.2 Classification of Infection (Detailed Explanation)

Infections can be classified according to several criteria, each providing clinically useful information about the behaviour, spread, and impact of the infectious agent (Abouelhag, 2010, p. 4). The table below summarises the classification, followed by a detailed explanation of each category.

Basis	Types
Localisation	Local, general (systemic), latent
Incidence	Sporadic, enzootic, epizootic
Etiology	Primary, secondary, mixed
Systematic (site)	Respiratory, urinary tract, etc.

A. Classification by Localisation (Site of Infection within the Body)

This refers to the anatomical distribution of the infection.

- **Local infection** – The microorganism remains confined to a single, well-defined area of the body. Examples include a boil (furuncle) on the skin or a localised abscess. The host’s inflammatory response typically walls off the infection, preventing systemic spread (Murphy et al., 2022).
- **General (systemic) infection** – The microorganism spreads throughout the body via the bloodstream or lymphatic system. Bacteraemia (bacteria in blood) or septicaemia (systemic illness with fever, tachycardia, and petechial haemorrhages) are examples (Abouelhag, 2010, p. 5). Systemic infections often produce generalised signs such as fever, malaise, and lymphadenopathy.
- **Latent infection** – The microorganism remains dormant within the host for a prolonged period without producing detectable signs of disease. However, the immune system may still produce antibodies that can be detected by serological tests (Abouelhag, 2010, p. 4). Examples include herpesviruses (e.g., varicella-zoster virus) and *Mycobacterium tuberculosis*. Reactivation can occur when host immunity is compromised (Punt et al., 2023).

B. Classification by Incidence (Frequency and Pattern in a Population)

This classification describes how often and in what pattern an infection occurs within a host population.

- **Sporadic infection** – Isolated cases occur irregularly and unpredictably, with no clear pattern. For example, a single case of tetanus in a community is considered sporadic (Abouelhag, 2010, p. 4).
- **Enzootic infection** – A disease that recurs regularly (endemic) in a particular animal host population within a defined geographic area. For example, Lyme disease is enzootic in certain rodent and tick populations. The term is analogous to “endemic” in human epidemiology (Abbas et al., 2022).
- **Epizootic infection** – A disease suddenly affects a large number of animals in a population over a short period, then rapidly declines. This is analogous to an “epidemic” in humans. Examples include avian influenza outbreaks in poultry (Abouelhag, 2010, p. 4; Delves et al., 2023).

C. Classification by Etiology (Cause or Sequence of Infection)

This classification is based on the cause or the order in which different microorganisms contribute to disease.

- **Primary infection** – Caused by a single species of microorganism that initiates the disease process. For example, infection with *Mycobacterium tuberculosis* causing primary tuberculosis (Abouelhag, 2010, p. 4).
- **Secondary infection** – Occurs when a primary infection weakens the host’s defences, allowing a different microorganism to cause an additional infection. For example, viral influenza (primary) may be followed by bacterial pneumonia caused by *Streptococcus pneumoniae* or *Staphylococcus aureus* (Abouelhag, 2010, p. 4).
- **Mixed infection** – Caused by more than one microorganism simultaneously, making diagnosis and treatment more difficult. For example, certain types of pneumonia may involve both bacteria and viruses, or a wound infection may contain multiple bacterial species (Abouelhag, 2010, p. 4; Wang et al., 2024).

D. Classification by Systematic Site (Anatomical System Affected)

This classification is based on the organ system involved. It is the most common clinical classification and guides both diagnosis and treatment.

- **Respiratory tract infections** – Examples: common cold (rhinovirus), influenza, pneumonia, tuberculosis.
- **Urinary tract infections** – Examples: cystitis, pyelonephritis, often caused by *Escherichia coli*.
- **Gastrointestinal infections** – Examples: gastroenteritis (rotavirus, *Salmonella*), hepatitis.
- **Nervous system infections** – Examples: meningitis, encephalitis.
- **Skin and soft tissue infections** – Examples: cellulitis, impetigo, abscesses.
- **Cardiovascular infections** – Examples: endocarditis, sepsis.

(Murphy et al., 2022)

1.3 Transmission

- **Direct** – immediate contact, exhaled droplets, scales, discharge.
- **Indirect** – insects carry microorganisms mechanically.

1.4 Why Some Infections Spread More Easily

Factors that increase infectivity include airborne survival of the agent, susceptibility of body surfaces (e.g., respiratory tract > intact skin), large number and frequent release of microorganisms, and strain-specific communicability.

1.5 Host and Environmental Factors

- Immune status, age.
- Gender.
- General health.
- Population density.
- Sanitation level.

1.6 Special Terms

- **Bacteraemia** – microorganisms in blood/lymph.
- **Septicemia** – infection with systemic symptoms (fever, tachycardia, petechial haemorrhages).
- **Toxaemia** – toxins in the blood (e.g., tetanus).
- **Pyaemia** – pus circulating with multiple abscesses.

2. The Immune System: Overview

- **Immunity** (Latin *immunis* – free of burden) – the general ability to resist infection or disease.
- **Immunology** – the science that studies immune responses, including self/nonself discrimination (Science Immunology, 2025).

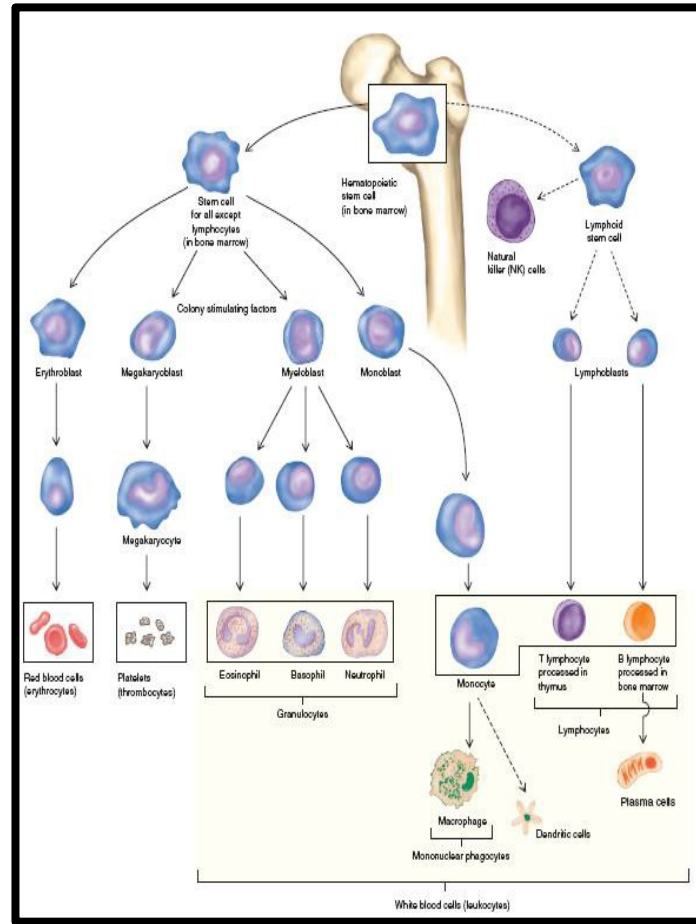
2.1 Two Types of Immune Responses

Feature	Nonspecific (Innate / Natural) Immunity	Specific (Adaptive) Immunity
Specificity	Broad (recognises general patterns)	Highly specific (recognises individual antigens)
Memory	None (same response each time)	Yes (improves on repeated exposure)
Response time	Immediate (minutes to hours)	Slower (days for primary response)
Main components	Physical barriers, phagocytes, complement, NK cells	B cells, T cells, antibodies

Both systems work together to eliminate pathogens (Murphy et al., 2022).

3. Cells of the Immune System

All leukocytes originate from **pluripotent haematopoietic stem cells** in fetal liver and bone marrow.



3.1 Lymphoid Cells (Major cells of specific immunity)

Cell Type	Maturation Site	Main Function
B cells	Bone marrow	Produce antibodies (plasma cells); present antigens
T cells	Thymus	Helper T cells (CD4+), cytotoxic T cells (CD8+), regulatory T cells
Natural Killer (NK) cells	Bone marrow	Kill virus-infected cells and tumour cells

(Abbas et al., 2022)

3.2 Mononuclear Phagocytes

- **Monocytes** – circulate in blood, then migrate into tissues and mature into macrophages.

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- **Macrophages** – highly phagocytic; act as antigen-presenting cells (APCs); have receptors for antibodies and complement (opsonisation).

3.3 Granulocytes (Polymorphonuclear Leukocytes – PMNs)

Type	Function
Neutrophils	Most abundant; primary phagocyte in acute inflammation; kill via lytic enzymes and reactive oxygen intermediates (Shafqat et al., 2023).
Eosinophils	Defend against parasites (helminths, protozoa); release cationic proteins.
Basophils	Non-phagocytic; release histamine and other mediators; involved in allergies (bind IgE).

3.4 Other Important Cells

- **Mast cells** – connective tissue; contain histamine granules; key in inflammation and hypersensitivity.
- **Dendritic cells** – most potent APCs; link innate and adaptive immunity by presenting antigens to T cells

(Huang et al., 2023; Nature Portfolio, 2024).

4. Organs and Tissues of the Immune System

4.1 Primary Lymphoid Organs (where lymphocytes mature)

Organ/Tissue	Function
Bone marrow	Site of B-cell maturation; origin of all immune cells.
Thymus	Site of T-cell maturation (positive and negative selection).

(Delves et al., 2023)

4.2 Secondary Lymphoid Organs/Tissues (where antigen encounter occurs)

Organ/Tissue	Function
Spleen	Filters blood; traps blood-borne antigens.
Lymph nodes	Filter lymph; traps antigens from tissues.
Mucosa-associated lymphoid tissue (MALT)	Includes GALT (gut), BALT (bronchial), SALT (skin).

5. Physical and Mechanical Barriers (First Line of Defense)

5.1 Skin

- Thick outer layer of **keratin** (microbes cannot digest).
- Continuous shedding of epithelial cells.
- Dryness and mild acidity (pH 5–6) inhibit microbial growth.
- **Sebum** forms a protective film.
- Normal skin microbiota competes with pathogens.
- **Skin-associated lymphoid tissue (SALT)** – contains Langerhans cells (dendritic cells) and intraepidermal lymphocytes (T cells).

(Murphy et al., 2022)

Innate Immunity: Physical and Mechanical Barriers

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Abstract

Innate immunity constitutes the first line of host defense against invading pathogens (Turvey & Broide, 2010). Among its components, physical and mechanical barriers play a fundamental role by preventing microbial entry, colonization, and dissemination. This educational note provides a comprehensive overview of these barriers across different anatomical sites, including the skin, mucous membranes, respiratory tract, gastrointestinal tract, genitourinary tract, and eyes. It also details associated chemical factors (e.g., lysozyme, lactoferrin, gastric juice, bacteriocins) and cellular elements such as Langerhans cells, M cells, and alveolar macrophages (Abbas et al., 2020; Gallo & Hooper, 2012). The note emphasizes the synergistic action of physical, mechanical, and chemical mechanisms that together form an effective surveillance system. Understanding these barriers is essential for appreciating how the body resists infection before adaptive immunity is engaged.

Keywords:

: Innate immunity, physical barriers, mechanical barriers, skin, mucous membranes, lysozyme, lactoferrin, M cells, alveolar macrophages, SALT, mucosal-associated lymphoid tissue, antimicrobial peptides

Introduction to Innate Immunity

Innate immunity is the evolutionarily ancient, non-specific defense system that responds immediately to pathogens. Unlike adaptive immunity, it does not require prior exposure and lacks immunological memory (Turvey & Broide, 2010). The physical and mechanical barriers are the most external components of innate immunity, designed to **prevent pathogen entry** or **rapidly remove** them before infection can establish (Abbas et al., 2020).

These barriers include:

- Intact skin and mucous membranes.
- Mechanical actions such as shedding, flushing, ciliary movement, peristalsis, coughing, and sneezing.
- Chemical factors that directly kill or inhibit microbes (Gallo & Hooper, 2012).

Together, they provide a formidable first line of defense.

1. The Skin: A Multilayered Physical and Chemical Fortress

The skin is the largest organ of the body ($\approx 1.5\text{--}2\text{ m}^2$) and serves as a primary physical barrier (Nestle et al., 2009).

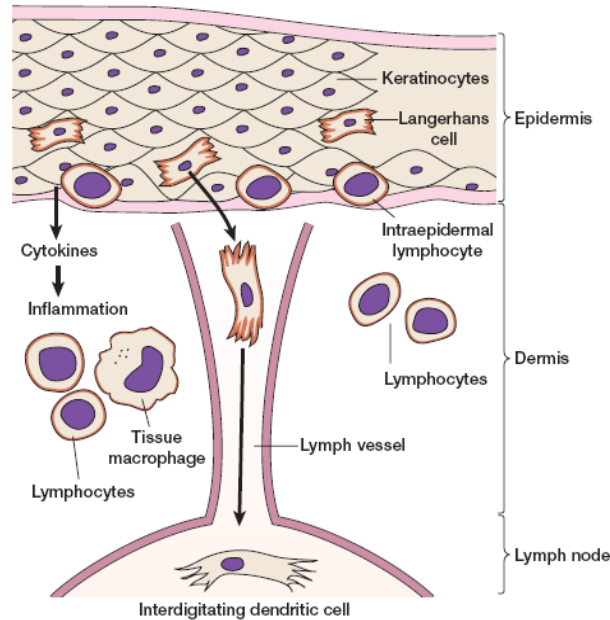
1.1 Structural Features

- **Keratinocytes** form multiple layers of stratified squamous epithelium; the outer layer (stratum corneum) is composed of dead, keratin-filled cells that are impermeable to most microorganisms (Nestle et al., 2009).
- **Continuous shedding** of outer epithelial cells removes attached microbes (Abbas et al., 2020).
- **Relative dryness** (low water activity) slows microbial growth (Gallo & Hooper, 2012).
- **Mild acidity** (pH 5–6) – due to lactic acid, free fatty acids, and amino acids – inhibits many pathogenic bacteria and fungi (Schroder, 2011).
- **Sebum** (from sebaceous glands) contains triglycerides that are broken down into free fatty acids with antimicrobial activity (Schroder, 2011).
- **Normal skin microbiota** (e.g., *Staphylococcus epidermidis*) produces bacteriocins and competes for nutrients, antagonizing pathogens like *Staphylococcus aureus* (Gallo & Hooper, 2012).
- **Hygiene** (washing) mechanically removes transient microorganisms.

1.2 Skin-Associated Lymphoid Tissue (SALT)

The skin is not just a passive barrier; it contains specialized immune cells (Nestle et al., 2009):

- **Langerhans cells** – dendritic cells in the epidermis that phagocytose antigens, migrate to draining lymph nodes, mature into interdigitating dendritic cells, and present antigens to naïve T cells, initiating adaptive immunity (Abbas et al., 2020).
- **Intraepidermal lymphocytes** – primarily $\gamma\delta$ T cells that act like cytotoxic T lymphocytes, destroying infected keratinocytes (Nestle et al., 2009).
- **Large numbers of macrophages** in the dermis that phagocytose pathogens and produce inflammatory cytokines (Abbas et al., 2020).



2. Mucous Membranes and Mucosal-Associated Lymphoid Tissue (MALT)

Mucous membranes line internal cavities exposed to the external environment (oral cavity, nasal passages, gut, vagina, etc.). They are more delicate than skin but have specialized defenses (Mestecky et al., 2015).

2.1 General Features

- **Mucus** – a viscous secretion containing glycoproteins (mucins) that traps microorganisms (Mestecky et al., 2015).
- **Antimicrobial components:**
 - **Cervical mucus** – impedes ascent of bacteria into the uterus.
 - **Prostatic fluid** – contains zinc and antibacterial factors.
 - **Tears** – contain lysozyme, lactoferrin, and sIgA (Kolar & McDermott, 2019).

2.2 M Cells (Microfold Cells)

M cells are specialized epithelial cells found overlying lymphoid follicles in the gut, tonsils, and Peyer's patches (Mestecky et al., 2015).

- **Structure:** Lack microvilli (brush border) but have a pocket on their basolateral side containing B cells, T cells, and macrophages (Abbas et al., 2020).
- **Function:**
 1. Phagocytose antigens and pathogens from the gut lumen.
 2. Transport them across the epithelial barrier into the pocket.
 3. Macrophages in the pocket engulf the antigen.
 4. Alternatively, M cells deliver antigens to organized lymphoid follicles.
 5. B cells in the follicle recognize the antigen, mature into plasma cells, and secrete **secretory IgA (sIgA)** (Mestecky et al., 2015).
 6. sIgA is transported into the gut lumen to neutralize specific pathogens.

This mechanism is a critical bridge between innate and adaptive immunity at mucosal surfaces (Abbas et al., 2020).

- Production of **bacteriocins** (e.g., colicin from *E. coli*, staphylococcin from *Staphylococcus*) (Gallo & Hooper, 2012).
- Competition for nutrients and adhesion sites.
- Stimulation of host immune responses (Mestecky et al., 2015).

5. Genitourinary Tract Defenses

The urinary and reproductive tracts are protected by several features (Abbas et al., 2020):

- **Urine properties:** Low pH ($\approx 5.5-6.5$), high urea concentration, uric acid, and hippuric acid – all inhibit microbial growth.
- **Hypotonic effect** of the kidney medulla – creates osmotic stress for bacteria.
- **Flushing action** – frequent voiding of urine mechanically removes pathogens.
- **Distance barrier** – long urethra (≈ 20 cm in males) makes ascending infection more difficult.
- **Secretory antibodies (sIgA)** in cervical mucus neutralize sperm-borne and sexually transmitted pathogens (Mestecky et al., 2015).
- **Prostatic antibacterial factor** – a zinc-containing peptide with antimicrobial activity.

6. The Eyes: Continuous Cleansing

The ocular surface is constantly exposed but remains remarkably infection-free due to (Kolar & McDermott, 2019):

- **Continuous flushing** by tears (produced by lacrimal glands, drained via nasolacrimal duct).
- **Tear composition:**
 - **Lysozyme** (muramidase) – breaks the $\beta(1\rightarrow4)$ bond between N-acetylmuramic acid and N-acetylglucosamine in peptidoglycan, especially effective against Gram-positive bacteria (Kolar & McDermott, 2019).
 - **Lactoferrin** – iron-binding protein that sequesters iron, limiting bacterial growth (Kolar & McDermott, 2019).
 - **sIgA** – neutralizes pathogens and prevents adhesion (Mestecky et al., 2015).
 - **Lactoperoxidase** – generates superoxide radicals that kill microbes.

Thus, tears provide both physical (flushing) and chemical protection.

7. Chemical Barriers: A Closer Look

While many chemical factors are associated with specific sites, some are systemic or widely distributed (Gallo & Hooper, 2012; Schroder, 2011).

Chemical Barrier	Source	Mechanism of Action
Lysozyme	Tears, saliva, mucus, milk	Hydrolyzes peptidoglycan (Gram-positive bacteria) (Kolar & McDermott, 2019)
Lactoferrin	Neutrophils, macrophages, secretions	Iron chelation; disrupts bacterial membranes (Kolar & McDermott, 2019)
Lactoperoxidase	Saliva, milk, tears	Generates hypothiocyanite and superoxide radicals
Gastric juice	Stomach	Acid denaturation of proteins (Abbas et al., 2020)
Salivary glycoproteins	Saliva	Inhibit bacterial adhesion
Urea	Urine	Alkaline degradation products are antimicrobial
Bacteriocins (colicin, staphylococcin)	Commensal bacteria	Pore formation, cell wall synthesis inhibition (Gallo & Hooper, 2012)
β -Lysin	Blood platelets	Disrupts microbial plasma membrane
Leukins	Neutrophils	Cationic antimicrobial peptides (Schroder, 2011)
Phagocytin	Phagocytes	Antimicrobial protein
Prostatic antibacterial factor	Prostate fluid	Zinc-dependent antimicrobial activity

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