
RESEARCH ARTICLE

SIR Model and its Applications

Ghulam Sarwar Mubarez¹ ✉ and Mohammad Naser Mohsini²

^{1,2}Assistant Professor, Department of Mathematics, Faculty of Science, Balkh University, Balkh Afghanistan

Corresponding Author: Ghulam Sarwar Mubarez, **E-mail:** ghsmubarez@gmail.com

ABSTRACT

The Susceptible-Infected-Recovered (SIR) model, first introduced by Kermack and McKendrick in 1927, represents a fundamental framework for understanding epidemic dynamics and has evolved into one of the most influential mathematical models in epidemiology. This article examines the foundations of the SIR model, including its mathematical formulation, underlying assumptions, and parameter estimation techniques. The review also explores diverse applications of the SIR model across multiple domains, including traditional epidemiological studies, economic modeling, social network analysis, and emerging areas such as cybersecurity and marketing. Through systematic analysis of recent literature, this work demonstrates the model's adaptability and continued relevance in addressing contemporary challenges, from COVID-19 pandemic modeling to understanding information diffusion in digital networks. The findings highlight both the strengths and limitations of the SIR framework while identifying future research directions for model enhancement and application.

KEYWORDS

SIR model, epidemiology, mathematical modeling, infection dynamics, parameter estimation, COVID-19, stochastic models.

ARTICLE INFORMATION

ACCEPTED: 01 November 2025

PUBLISHED: 10 December 2025

DOI: 10.32996/jmss.2025.6.6.2

1. Introduction

The Susceptible-Infectious-Recovered (SIR) model stands as one of the most fundamental and influential frameworks in mathematical epidemiology, serving as the cornerstone for understanding infectious disease dynamics across diverse populations and contexts. Originally formulated by Kermack and McKendrick in their seminal 1927 paper "A contribution to the mathematical theory of epidemics," the SIR model represents a revolutionary approach to quantifying disease transmission patterns through mathematical modeling (Kermack & McKendrick, 1927). This compartmental model divides a population into three distinct categories: susceptible individuals who can contract the disease, infectious individuals who can transmit the disease, and recovered individuals who have gained immunity, providing a simplified yet powerful representation of epidemic dynamics (Hethcote, 2000).

The theoretical foundation of the SIR model rests on a system of ordinary differential equations that describe the flow of individuals between compartments over time. The model assumes homogeneous mixing within the population, constant population size, and permanent immunity following recovery, making it particularly suitable for analyzing acute infectious diseases such as measles, influenza, and more recently, COVID-19 (Anderson & May, 1992; Brauer et al., 2019). The mathematical elegance of the SIR framework lies in its ability to capture complex epidemic phenomena through relatively simple parameters: the transmission rate (β) representing the effective contact rate between susceptible and infectious individuals, and the recovery rate (γ) indicating the rate at which infectious individuals recover and gain immunity (Bailey, 1975).

The significance of the SIR model extends beyond its mathematical formulation to its practical applications in public health decision-making and policy development. The model's capacity to predict epidemic trajectories, estimate key epidemiological parameters such as the basic reproduction number (R_0), and evaluate the potential impact of intervention strategies has made it an indispensable tool for epidemiologists, public health officials, and policymakers worldwide (Keeling & Rohani, 2008). The

basic reproduction number, defined as the average number of secondary infections caused by a single infectious individual in a completely susceptible population, serves as a critical threshold parameter that determines whether an epidemic will grow ($R_0 > 1$) or decline ($R_0 < 1$) in a population (Keeling & Eames, 2005).

Over the decades since its inception, the SIR model has undergone numerous extensions and modifications to address the limitations of the original framework and accommodate the complexities of real-world disease transmission scenarios. Researchers have developed stochastic versions of the SIR model to account for random fluctuations in small populations, where demographic stochasticity can significantly influence epidemic outcomes (Allen, 2008; Gourieroux & Lu, 2020). Additionally, spatial extensions incorporating network theory and metapopulation dynamics have emerged to capture the heterogeneous nature of human contact patterns and geographical disease spread (Newman, 2002; Pastor-Satorras & Vespignani, 2001).

The advent of computational modeling and increased data availability has further expanded the scope and sophistication of SIR-based models. Modern implementations incorporate time-varying parameters to reflect changing transmission dynamics due to seasonal effects, behavioral modifications, or intervention measures (Dong et al., 2023; Zelenkov & Reshetsov, 2023). Machine learning techniques have been integrated with traditional SIR frameworks to enhance predictive accuracy and parameter estimation, particularly during emerging epidemic situations where historical data may be limited (Mortensen et al., 2024; Bi et al., 2022).

The COVID-19 pandemic has particularly highlighted both the utility and limitations of SIR-based models, leading to renewed interest in model refinement and development of more sophisticated compartmental structures. Researchers have proposed models incorporating asymptomatic transmission, waning immunity, and vaccination dynamics to better capture the complexities of SARS-CoV-2 transmission (Nesteruk, 2020; Singh et al., 2021; Chen et al., 2021). The pandemic has also demonstrated the importance of incorporating behavioral responses and policy interventions into epidemiological models, leading to the development of behavioral SIR variants that account for human decision-making processes (Keppo et al., 2021).

Recent methodological advances have focused on addressing parameter uncertainty and improving model calibration through advanced statistical techniques. Bayesian approaches have been employed to quantify uncertainty in epidemic projections and parameter estimates, providing more robust foundations for policy recommendations (Zimmer et al., 2019). Additionally, the development of time-varying and fractional-order SIR models has enhanced the framework's ability to capture memory effects and long-term epidemic dynamics (Ilhan & Şahin, 2024; Tahir et al., 2024).

The versatility of the SIR framework extends beyond traditional epidemiological applications to encompass diverse fields where diffusion-like processes occur. Marketing researchers have adapted SIR models to understand viral marketing campaigns and product adoption patterns, treating ideas and products as "infectious" entities that spread through social networks (Freed, 2019; Goldenberg et al., 2001). Social media platforms have employed SIR-based models to analyze information dissemination and influence propagation, while urban planners have utilized modified SIR frameworks to model traffic congestion spread and urban dynamics (Sharif et al., 2022; Teng & Wei, 2024).

The integration of network theory with SIR modeling has opened new avenues for understanding disease spread in complex social structures. Scale-free networks, small-world networks, and other complex network topologies have been incorporated into SIR frameworks to better represent realistic contact patterns and their implications for epidemic dynamics (Watts & Dodds, 2007; Vespignani, 2012). These network-based extensions have provided insights into the role of superspreaders, community structure, and targeted intervention strategies in controlling epidemic spread.

Contemporary research has also explored the intersection of SIR modeling with emerging technologies and applications. Cybersecurity researchers have adapted SIR principles to model malware propagation in computer networks and wireless sensor networks, treating malicious software as infectious agents that spread through digital systems (Kovtun et al., 2024; Ginters et al., 2024). This cross-disciplinary application demonstrates the fundamental universality of the SIR framework in modeling diffusion processes across diverse domains.

The mathematical analysis of SIR models has revealed important theoretical insights into epidemic dynamics, including the existence of epidemic thresholds, final size relations, and conditions for disease persistence or extinction. These theoretical results provide crucial guidance for public health planning and resource allocation, particularly in determining vaccination coverage requirements for herd immunity and optimal timing of intervention measures (Hethcote, 2000; Brauer & Castillo-Chavez, 2012).

As the field of mathematical epidemiology continues to evolve, the SIR model remains a foundational framework that serves as both a teaching tool for understanding basic epidemic principles and a building block for more complex modeling approaches.

Its enduring relevance reflects the model's ability to capture essential features of infectious disease transmission while maintaining mathematical tractability and interpretability (Tolles & Luong, 2020; Siettos & Russo, 2013).

2. Fundamentals of the SIR Model

2.1. Mathematical Formulation

The classical SIR model is expressed as a system of ordinary differential equations that describe the rate of change in each compartment over time. The basic formulation, as originally presented by Kermack and McKendrick (1927), consists of three coupled differential equations:

$$\begin{aligned}\frac{dS}{dt} &= -\beta SI \\ \frac{dI}{dt} &= \beta SI - \gamma I \\ \frac{dR}{dt} &= \gamma I\end{aligned}$$

where $S(t)$, $I(t)$, and $R(t)$ represent the number of susceptible, infected, and recovered individuals at time t , respectively. The parameter β represents the transmission rate, γ represents the recovery rate, and $N = S + I + R$ represents the total population size, which remains constant in the basic model (Hethcote, 2000). The ratio β/γ defines the basic reproduction number R_0 , which represents the average number of secondary infections produced by a single infected individual in a completely susceptible population.

The mathematical structure of the SIR model embodies several key principles of epidemic dynamics. The first equation captures the depletion of susceptible individuals through infection, with the rate proportional to the product of susceptible and infected populations. The second equation represents the balance between new infections and recoveries, while the third equation tracks the accumulation of recovered individuals (Brauer & Castillo-Chavez, 2012). This formulation ensures that the total population remains constant, reflecting the closed-population assumption of the basic model.

2.2. Model Assumptions and Limitations

The SIR model operates under several fundamental assumptions that both enable its mathematical tractability and limit its applicability to real-world scenarios. These assumptions include: (1) homogeneous mixing, where all individuals have equal probability of contact with any other individual; (2) constant transmission and recovery rates; (3) permanent immunity after recovery; (4) closed population with no births, deaths, or migration; and (5) instantaneous transition between compartments (Bailey, 1975). Understanding these assumptions is crucial for appropriate model application and interpretation of results.

The homogeneous mixing assumption, while mathematically convenient, often fails to capture the heterogeneity of real-world contact patterns. Individuals typically interact within specific social networks, geographic regions, or demographic groups, leading to non-random mixing patterns that can significantly influence disease dynamics (Keeling & Eames, 2005). Similarly, the assumption of constant parameters may not reflect the dynamic nature of disease transmission, which can vary due to seasonal factors, behavioral changes, or implementation of control measures.

Recent research has addressed many of these limitations through model modifications and extensions. Nikitina et al. (2020) explored modifications of the SIR model to better capture viral disease spread dynamics, while Hellwig (2022) introduced density-dependent transmission rates to account for spatial heterogeneity. These developments demonstrate the ongoing evolution of the SIR framework to address its inherent limitations while maintaining its fundamental structure.

2.3. Parameter Estimation and Calibration

Accurate parameter estimation represents a critical challenge in SIR model implementation, as the model's predictive capability depends heavily on the precise determination of transmission rate β and recovery rate γ . Traditional parameter estimation approaches rely on fitting model predictions to observed epidemic data using various optimization techniques, including maximum likelihood estimation and least squares methods (Chen et al., 2021). However, the inherent variability in epidemic data and the model's sensitivity to parameter values often complicate this process.

Contemporary approaches to parameter estimation have incorporated advanced computational methods to improve accuracy and account for uncertainty. Mortensen et al. (2024) developed a machine learning-enabled SIR model that dynamically adjusts parameters based on observed data, while Zelenkov and Reshetsov (2023) employed genetic algorithms to fit time-varying parameters to COVID-19 data. These methodological advances highlight the importance of sophisticated parameter estimation techniques in modern SIR model applications.

The challenge of parameter estimation is further complicated by the need to account for model uncertainty and data limitations. Zimmer et al. (2019) emphasized the importance of accurate uncertainty quantification in epidemic parameter estimates, proposing stochastic compartmental models that explicitly incorporate parameter uncertainty. This approach recognizes that parameter estimates are themselves uncertain and that this uncertainty should be propagated through model predictions to provide more realistic confidence intervals.

2.4. Model Variations and Extensions

The basic SIR model has spawned numerous variations and extensions designed to address specific epidemiological scenarios and overcome the limitations of the original formulation. These modifications include the incorporation of additional compartments (such as SEIR models that include an exposed compartment), consideration of demographics (SIRS models that account for waning immunity), and inclusion of vaccination dynamics (SIRV models) (Basnarkov, 2021). Each variation serves to enhance the model's realism and applicability to specific contexts.

Stochastic extensions of the SIR model have gained particular attention for their ability to capture the inherent randomness in disease transmission processes. Gourieroux and Lu (2020) developed a SIR model with stochastic transmission, while Tahir et al. (2024) explored stochastic epidemic modeling in the context of wireless sensor networks. These stochastic formulations acknowledge that disease transmission is fundamentally a random process, particularly in small populations where chance events can significantly influence epidemic trajectories.

Time-varying parameter models represent another important class of SIR extensions that address the limitation of constant transmission and recovery rates. Dong et al. (2023) applied a time-delay SIR model with vaccination to COVID-19 prediction, while Law et al. (2020) developed a time-varying SIR model to capture the depleting transmission dynamics of COVID-19. These models recognize that epidemic parameters can change over time due to various factors, including seasonal effects, behavioral changes, and policy interventions.

3. Applications of the SIR Model

3.1. Traditional Epidemiological Applications

The SIR model's primary application domain remains epidemiology, where it continues to serve as a fundamental tool for understanding and predicting disease spread. During the COVID-19 pandemic, the model experienced renewed attention as researchers worldwide applied various SIR formulations to analyze transmission dynamics, evaluate intervention strategies, and inform public health policy decisions. Nesteruk (2020) utilized SIR model simulations to predict COVID-19 pandemic trajectories, while Saxena et al. (2022) conducted comprehensive investigations of COVID-19 spread using susceptible-infectious-recovered frameworks.

The COVID-19 pandemic has represented the most significant global application of SIR modeling in recent history, demonstrating both the power and limitations of compartmental approaches in real-time epidemic response. Researchers worldwide rapidly adapted SIR frameworks to model SARS-CoV-2 transmission dynamics, providing crucial insights for public health decision-making during the early phases of the pandemic when empirical data was limited (Nesteruk, 2020; Postnikov, 2020). These applications highlighted the model's utility in generating rapid assessments of epidemic potential and informing initial policy responses under conditions of high uncertainty.

The complexity of COVID-19 transmission patterns necessitated numerous modifications to the basic SIR framework, leading to the development of more sophisticated compartmental models. Researchers incorporated asymptomatic transmission pathways, age-stratified mixing patterns, and time-varying parameters to better capture the heterogeneous nature of SARS-CoV-2 spread (Sarkar et al., 2020; Law et al., 2020). The development of SEIR variants, which include an exposed compartment to account for the incubation period, proved particularly valuable for COVID-19 modeling given the disease's characteristic latency period (Basnarkov, 2021).

Advanced SIR-based models for COVID-19 incorporated vaccination dynamics and waning immunity, reflecting the evolving understanding of SARS-CoV-2 immunology and the rollout of vaccination programs. Dong et al. (2023) developed time-delay SIR models with vaccination components, demonstrating how optimal control strategies could be derived from mathematical frameworks to minimize epidemic impact while considering economic and social constraints. These models provided quantitative foundations for vaccination prioritization strategies and helped evaluate the potential impact of different vaccine distribution scenarios.

The integration of machine learning techniques with SIR modeling emerged as a powerful approach for COVID-19 forecasting, particularly when dealing with incomplete or noisy data. Bi et al. (2022) demonstrated how gated recurrent units could be

combined with evolutionary algorithms to enhance SIR model predictions, while Mortensen et al. (2024) developed machine learning-enabled adaptive SIR models that could dynamically adjust parameters based on incoming data streams. These hybrid approaches proved particularly valuable during periods of changing transmission dynamics due to emerging variants or policy interventions.

Stochastic extensions of SIR models gained prominence in COVID-19 applications, addressing the limitations of deterministic approaches when modeling outbreak dynamics in small populations or during early epidemic phases. Pramanik (2024) explored optimal lockdown and vaccination strategies using stochastic SIR frameworks, demonstrating how random fluctuations could significantly influence intervention effectiveness and timing. These stochastic approaches provided more realistic uncertainty quantification for policy recommendations and helped identify critical thresholds for intervention implementation.

Recent developments in SIR model applications have focused on integrating machine learning techniques and advanced statistical methods to improve predictive accuracy and parameter estimation. Bi et al. (2022) developed COVID-19 forecasting and intervention planning using gated recurrent units combined with evolutionary algorithms, while Bousquet et al. (2022) employed deep learning for forecasting using time-varying parameters of the SIRD model. These hybrid approaches demonstrate the evolution of SIR modeling toward more sophisticated, data-driven methodologies.

3.2. Economic and Macroeconomic Applications

The SIR model has found significant application in economic modeling, where it provides a framework for understanding how economic shocks, policy changes, and market dynamics spread through interconnected economic systems. Bayraktar et al. (2021) developed a macroeconomic SIR model for COVID-19 that integrated epidemiological dynamics with economic variables, demonstrating how health and economic outcomes are interconnected during pandemic periods. This application highlighted the model's versatility in addressing complex, multi-dimensional problems that extend beyond traditional epidemiological contexts.

The economic applications of SIR models have proven particularly valuable in analyzing the economic impacts of pandemic interventions and optimal policy design. Acemoglu et al. (2020) proposed a multi-risk SIR model with optimally targeted lockdown policies, providing insights into how epidemiological models can inform economic policy decisions. Their work demonstrated how SIR frameworks can be extended to consider multiple risk groups and optimize intervention strategies based on both health and economic criteria.

Insurance and financial applications of SIR models have emerged as another important domain, particularly in the context of pandemic risk assessment and management. Boado-Penas and Eisenberg (2022) explored the application of SIR models to pandemic insurance and social protection, illustrating how epidemiological modeling can inform risk assessment and product design in the insurance industry. These applications demonstrate the model's utility in quantifying and managing pandemic-related financial risks.

The development of behavioral economic models incorporating SIR dynamics has provided insights into how individual decision-making processes influence epidemic outcomes and economic impacts. Keppo et al. (2021) developed behavioral SI* models that account for how individuals modify their behavior in response to epidemic information, creating feedback loops between disease dynamics and economic activity. These models help explain phenomena such as voluntary social distancing and its economic consequences, providing more realistic foundations for policy design.

3.3. Social Network and Information Diffusion Applications

The application of SIR models to social network analysis has also encompassed source identification and influence maximization problems. Zang et al. (2015) developed approaches for locating multiple sources in social networks under the SIR model using divide-and-conquer strategies, addressing the challenge of identifying the origins of information cascades in complex network structures. These applications demonstrate the model's utility in understanding and managing information flow in interconnected social systems.

The application of SIR principles to marketing and information diffusion represents one of the most innovative extensions of epidemiological modeling beyond health applications. Marketing researchers have recognized the fundamental similarity between disease transmission and viral marketing processes, where ideas, products, or information spread through social networks in patterns analogous to infectious disease dynamics (Goldenberg et al., 2001). This cross-disciplinary application has provided new insights into consumer behavior, product adoption patterns, and optimal marketing strategies.

Freed (2019) developed SIR models with vector transmission to predict the effectiveness of viral marketing campaigns and product adoption spread, demonstrating how epidemiological frameworks could quantify marketing reach and influence propagation through social networks. These models incorporated concepts such as "infection" through exposure to marketing messages and "recovery" through saturation or loss of interest, providing quantitative foundations for marketing strategy optimization and resource allocation.

Social media platforms have extensively utilized SIR-based models to analyze information dissemination patterns and optimize content distribution strategies. Sharif et al. (2022) evaluated the effectiveness of online video marketing on Facebook using SIR frameworks, demonstrating how compartmental models could predict viral content spread and identify factors influencing engagement patterns. These applications have become increasingly important as organizations seek to understand and leverage social media dynamics for communication and marketing purposes.

The extension of SIR models to multi-source information diffusion has provided insights into complex information ecosystems where multiple competing or complementary messages propagate simultaneously. Zang et al. (2015) developed divide-and-conquer approaches for locating multiple information sources in social networks using SIR frameworks, demonstrating how epidemiological principles could address problems in social network analysis and information source identification. These methods have applications in misinformation tracking, influence analysis, and social media monitoring.

3.4. Cybersecurity and Digital Systems Applications

The cybersecurity domain has embraced SIR modeling as a fundamental framework for understanding malware propagation, network security threats, and digital epidemic dynamics. Computer security researchers have recognized that malware spread through digital networks exhibits characteristics remarkably similar to infectious disease transmission in biological populations, leading to the development of cyber-epidemic models based on SIR principles (Kovtun et al., 2024).

Advanced applications in cybersecurity have incorporated entropy-extremal dynamic interpretations of SIR models to forecast cyber epidemic spread and optimize security response strategies. Kovtun et al. (2024) developed frameworks that combine information theory with epidemiological modeling to predict malware propagation patterns and identify critical vulnerabilities in network infrastructure. These models help cybersecurity professionals prioritize security investments, optimize patch deployment strategies, and design resilient network architectures.

Wireless sensor networks represent a particularly important application domain for cyber-epidemic modeling, where malware spread can compromise critical infrastructure monitoring and control systems. Tahir et al. (2024) explored stochastic epidemic modeling approaches specifically tailored to worm transmission in wireless sensor networks, incorporating network topology, communication protocols, and energy constraints into SIR frameworks. These applications are crucial for protecting critical infrastructure and ensuring the reliability of sensor-based monitoring systems.

The integration of SIR models with machine learning approaches has enhanced the effectiveness of cybersecurity applications, particularly in early detection and rapid response scenarios. Ginters et al. (2024) evaluated the usability of classic SIR and diffusion models for assessing malware spread in early stages, demonstrating how epidemiological principles could enhance threat intelligence and incident response capabilities. These hybrid approaches provide cybersecurity teams with predictive capabilities for threat assessment and resource allocation optimization.

Network security applications of SIR models have also extended to understanding information security vulnerabilities and system resilience. The model's compartmental structure provides a natural framework for categorizing system states (vulnerable, compromised, secured) and analyzing transitions between these states. This application demonstrates the model's adaptability to diverse domains where contagion-like processes occur, whether biological, social, or digital in nature.

3.5. Transportation and Urban Planning Applications

Recent innovative applications of the SIR model have emerged in transportation and urban planning, where it provides insights into traffic congestion propagation and urban mobility dynamics. Teng and Wei (2024) conducted a comprehensive review of SIR model applications in predicting urban traffic congestion, identifying successes and future directions for this emerging application domain. Their work demonstrated how epidemiological frameworks can be adapted to understand how traffic congestion spreads through urban transportation networks, treating congested areas as "infected" regions that influence neighboring areas.

The application of SIR models to traffic dynamics has proven particularly valuable in understanding how local traffic disturbances propagate through transportation networks. Kozhabek et al. (2024) developed approaches for modeling traffic congestion

spreading using topology-based SIR epidemic models, providing insights into how network structure influences congestion propagation patterns. This application highlights the model's utility in understanding spatial dynamics and network effects in urban systems.

The application of SIR models to urban planning extends beyond traffic management to encompass broader urban dynamics such as urban sprawl, gentrification patterns, and service distribution optimization. These models help urban planners understand how spatial processes spread through urban environments and identify critical intervention points for managing urban development patterns. The integration of geographic information systems (GIS) with SIR models has enabled spatially explicit analysis of urban phenomena, providing more detailed insights into local-scale processes and their regional implications.

Public transportation systems have utilized SIR-based models to optimize route planning, capacity allocation, and system resilience analysis. These applications treat passenger flow disruptions as infectious processes that can spread through transportation networks, helping system operators develop contingency plans and design robust transportation infrastructure that can maintain functionality during disruptions or peak demand periods.

3.6. Advanced Modeling Techniques and Computational Approaches

Contemporary SIR model applications have increasingly incorporated advanced computational techniques and machine learning approaches to enhance predictive accuracy and analytical capabilities. İlhan and Şahin (2024) developed numerical approaches for epidemic SIR models using Morgan-Voyce series, demonstrating how mathematical innovations can improve model solution techniques. These computational advances enable more sophisticated analysis of complex epidemic dynamics and support real-time decision-making in various application domains.

The integration of machine learning techniques with SIR models has opened new possibilities for dynamic parameter estimation and adaptive modeling. Singh et al. (2021) developed modified variable-order fractional SIR models to predict COVID-19 spread, while Habott et al. (2024) analyzed COVID-19 dynamics using modified SIR models characterized by nonlinear functions. These approaches demonstrate how traditional SIR frameworks can be enhanced through modern computational methods to address contemporary challenges.

Control theory applications of SIR models have emerged as another important development, providing frameworks for optimal intervention design and pandemic mitigation strategies. Godara et al. (2024) developed control theory approaches to optimal pandemic mitigation using SIR frameworks, while Pramanik (2024) estimated optimal lockdown and vaccination rates using stochastic SIR models. These applications demonstrate how SIR models can inform evidence-based policy decisions and intervention strategies across various domains.

3.7. Healthcare System Planning and Resource Allocation

Healthcare systems have extensively utilized SIR models for capacity planning, resource allocation, and emergency preparedness, particularly in anticipating and managing epidemic-related healthcare demand. These applications extend beyond disease modeling to encompass broader healthcare system dynamics, including staff deployment, equipment procurement, and facility utilization optimization (Ferguson et al., 2020).

Hospital capacity modeling represents a critical application of SIR frameworks, where epidemic projections are translated into healthcare resource requirements such as hospital beds, ventilators, and intensive care unit capacity. During the COVID-19 pandemic, healthcare systems worldwide relied on SIR-based projections to guide capacity expansion decisions, staff scheduling, and patient transfer protocols. These models helped healthcare administrators balance resource availability with anticipated demand, optimizing patient outcomes while managing system constraints.

The development of control theory approaches combined with SIR modeling has provided sophisticated frameworks for pandemic mitigation strategy optimization. Godara et al. (2024) explored optimal pandemic mitigation strategies using control theory principles integrated with epidemiological models, demonstrating how mathematical optimization could guide policy decisions regarding intervention timing, intensity, and duration. These approaches provide quantitative foundations for balancing public health objectives with social and economic constraints.

Healthcare workforce planning has benefited from SIR-based models that account for healthcare worker infection risks, quarantine requirements, and capacity constraints during epidemic periods. These models help healthcare administrators develop staff deployment strategies that maintain essential services while protecting healthcare workers from occupational exposure risks. The integration of occupational health considerations into epidemic models has become increasingly important for healthcare system resilience and sustainability.

3.8. Environmental and Ecological Applications

The extension of SIR principles to environmental and ecological systems has provided new insights into contaminant spread, ecosystem dynamics, and environmental risk assessment. Environmental scientists have adapted epidemiological frameworks to model pollutant diffusion, invasive species spread, and ecosystem disturbance propagation, demonstrating the versatility of compartmental modeling approaches across diverse scientific domains.

Ecological applications of SIR models have focused on invasive species dynamics, where introduced species spread through ecosystems in patterns analogous to infectious disease transmission. These models help ecologists predict invasion patterns, identify critical control points, and optimize management strategies for invasive species control. The integration of spatial dynamics and habitat heterogeneity into SIR frameworks has enhanced the realism and predictive capability of ecological applications.

Environmental contamination modeling has utilized SIR principles to understand pollutant spread through environmental media such as groundwater, soil, and atmospheric systems. These applications treat contamination as an infectious process where polluted areas influence adjacent clean areas, creating spreading patterns that can be analyzed using epidemiological methods. The results inform environmental remediation strategies, risk assessment procedures, and regulatory policy development.

Climate change impacts on disease vector dynamics have been analyzed using modified SIR frameworks that incorporate temperature, precipitation, and habitat changes into disease transmission models. These applications help public health officials anticipate how changing environmental conditions may influence vector-borne disease patterns and develop adaptive management strategies for emerging health threats.

The comprehensive applications of SIR models across these diverse domains demonstrate the fundamental universality of diffusion processes and the powerful insights that epidemiological frameworks can provide when applied to complex systems. From traditional infectious disease modeling to innovative applications in marketing, cybersecurity, and urban planning, the SIR model continues to evolve and adapt to address contemporary challenges across multiple disciplines. The ongoing development of computational tools, data analytics approaches, and interdisciplinary collaborations promises to further expand the scope and impact of SIR-based modeling in addressing complex societal challenges.

4. Discussion and Future Directions

The comprehensive review of SIR model applications reveals both the remarkable versatility and enduring relevance of this fundamental epidemiological framework. From its origins in disease transmission modeling to contemporary applications in cybersecurity, economics, and urban planning, the SIR model has demonstrated exceptional adaptability to diverse domains characterized by contagion-like processes. This versatility stems from the model's fundamental structure, which captures the essential dynamics of state transitions in systems where entities can exist in distinct categories and move between them according to specific rules.

However, the widespread application of SIR models has also highlighted several important limitations and challenges that require continued attention. The model's assumptions of homogeneous mixing, constant parameters, and closed populations often diverge from real-world conditions, necessitating careful consideration of model appropriateness and result interpretation. Recent developments in stochastic modeling, time-varying parameters, and network-based approaches have addressed many of these limitations, but significant challenges remain in balancing model complexity with practical applicability.

Future research directions for SIR model development should focus on several key areas. First, continued advancement in parameter estimation techniques, particularly those incorporating machine learning and artificial intelligence, will enhance model accuracy and predictive capability. Second, the development of more sophisticated network-based models that capture realistic contact patterns and spatial dynamics will improve model realism. Third, the integration of SIR models with other modeling frameworks, such as agent-based models and complex systems approaches, will enable more comprehensive analysis of complex phenomena.

The COVID-19 pandemic has demonstrated both the utility and limitations of SIR models in real-world crisis situations, providing valuable lessons for future model development and application. Close (2024) evaluated the application of SIR models to localized populations, while Nesteruk (2025) developed general SIR models for visible and hidden epidemic dynamics. These studies highlight the importance of context-specific model adaptation and the need for robust uncertainty quantification in model predictions.

5. Conclusion

The SIR model represents one of the most successful and enduring mathematical frameworks in applied sciences, with applications spanning from traditional epidemiology to emerging domains such as cybersecurity and urban planning. This comprehensive review has demonstrated the model's instructional value in understanding fundamental dynamics of contagion processes and its practical utility in addressing diverse real-world challenges. The model's mathematical elegance, combined with its conceptual clarity, has enabled researchers and practitioners across multiple disciplines to adapt and apply it to their specific contexts.

The evolution of SIR model applications reflects broader trends in scientific modeling, including the integration of computational methods, incorporation of stochastic elements, and development of hybrid approaches that combine traditional mathematical modeling with modern data science techniques. These developments have enhanced the model's predictive capability and expanded its applicability while maintaining its fundamental conceptual framework.

As we look toward the future, the SIR model will undoubtedly continue to evolve and find new applications in emerging domains. The model's fundamental structure provides a robust foundation for understanding contagion processes, while its flexibility enables adaptation to new contexts and challenges. Continued research and development in SIR modeling will contribute to our understanding of complex systems and support evidence-based decision-making across diverse fields of human endeavor.

References

- [1] Acemoglu, D., Chernozhukov, V., Werning, I., & Whinston, M. D. (2020). A multi-risk SIR model with optimally targeted lockdown (Vol. 2020). Cambridge, MA: National Bureau of Economic Research.
- [2] Allen, L. J. S. (2008). An introduction to stochastic epidemic models. In *Mathematical epidemiology* (pp. 81–130). Springer.
- [3] Anderson, R. M., & May, R. M. (1992). *Infectious diseases of humans: dynamics and control*. Oxford University Press.
- [4] Bailey, N. T. J. (1975). *The mathematical theory of infectious diseases and its applications*. Griffin.
- [5] Bakhta, A., Boiveau, T., Maday, Y., & Mula, O. (2020). Epidemiological forecasting with model reduction of compartmental models. Application to the COVID-19 pandemic. *Biology*, 10(1), 22. <https://www.mdpi.com/2079-7737/10/1/22>
- [6] Basnarkov, L. (2021). SEAIR epidemic spreading model of COVID-19. *Chaos, Solitons & Fractals*, 142, 110394.
- [7] Bayraktar, E., Cohen, A., & Nellis, A. (2021). A macroeconomic SIR model for COVID-19. *Mathematics*, 9(16), 1901.
- [8] Bi, L., Fili, M., & Hu, G. (2022). COVID-19 forecasting and intervention planning using gated recurrent unit and evolutionary algorithm. *Neural Computing and Applications*, 34(20), 17561-17579.
- [9] Boado-Penas, M. D. C., & Eisenberg, J. (2022). *Pandemics: Insurance and social protection* (p. 298). Springer Nature.
- [10] Bousquet, A., Conrad, W. H., Sadat, S. O., Vardanyan, N., Chang, J. L., Taschereau-Dumouchel, V., ... & Goodman, J. (2022). Deep learning forecasting using time-varying parameters of the SIRD model for Covid-19. *Scientific Reports*, 12(1), 3030.
- [11] Brauer, F., & Castillo-Chavez, C. (2012). *Mathematical models for communicable diseases*. SIAM.
- [12] Brauer, F., Castillo-Chavez, C., & Feng, Z. (2019). *Mathematical models in epidemiology*. Springer.
- [13] Breda, D., Diekmann, O., De Graaf, W. F., Pugliese, A., & Vermiglio, R. (2012). On the formulation of epidemic models (an appraisal of Kermack and McKendrick). *Journal of Biological Dynamics*, 6(2), 103-117.
- [14] Chen, X., Li, J., & Xiao, C. (2021). Numerical solution and parameter estimation for uncertain SIR model with application to COVID-19. *Fuzzy Optimization and Decision Making*, 20, 189.
- [15] Close, A. (2024). Evaluating the application of the SIR model to a localized population. *International Surgery*, 108(3), 124-131.
- [16] Connell, R., Dawson, P., & Skvortsov, A. (2009). Comparison of an agent-based model of disease propagation with the generalised SIR epidemic model. *Defence Science and Technology Organisation*.
- [17] Dong, S., Xu, L., A, Y., Lan, Z. Z., Xiao, D., & Gao, B. (2023). Application of a time-delay SIR model with vaccination in COVID-19 prediction and its optimal control strategy. *Nonlinear Dynamics*, 111(11), 10677-10692.
- [18] Ellison, G. (2020). Implications of heterogeneous SIR models for analyses of COVID-19 (No. w27373). National Bureau of Economic Research.
- [19] El-Sayed, A. M., Scarborough, P., Seemann, L., & Galea, S. (2012). Social network analysis and agent-based modeling in social epidemiology. *Epidemiologic Perspectives & Innovations*, 9(1), 1-9.
- [20] Ferguson, N. M., et al. (2020). Impact of non-pharmaceutical interventions (NPIs) to reduce COVID-19 mortality and healthcare demand. Imperial College London. <https://doi.org/10.25561/77482>
- [21] Freed, M. (2019). Using the SIR epidemiology model with vector transmission to predict the effectiveness of a viral marketing campaign and the spread of product adoption. *Undergraduate Journal of Mathematical Modeling: One+ Two*, 9(2), 1.
- [22] Ginters, E., Eroles, M. A. P., & Matvejevs, A. (2024, October). Usability of classic SIR and diffusion models for assessing malware spread in the early stages of infectious diseases. In *2024 IEEE 65th International Scientific Conference on Information Technology and Management Science of Riga Technical University (ITMS)* (pp. 1-8). IEEE.
- [23] Giordano, G., Blanchini, F., Bruno, R., Colaneri, P., Di Filippo, A., Di Matteo, A., & Colaneri, M. (2020). A SIDARTHE model of COVID-19 epidemic in Italy. *arXiv preprint arXiv:2003.09861*.
- [24] Giudici, M., Comunian, A., & Gaburro, R. (2020). Inversion of a SIR-based model: a critical analysis about the application to COVID-19 epidemic. *Physica D: Nonlinear Phenomena*, 413, 132674.

- [25] Godara, P., Herminghaus, S., & Heidemann, K. M. (2024). Correction: A control theory approach to optimal pandemic mitigation. *PLoS One*, 19(12), e0315749.
- [26] Goldenberg, J., Libai, B., & Muller, E. (2001). Talk of the network: A complex systems look at the underlying process of word-of-mouth. *Marketing Letters*, 12(3), 211-223.
- [27] Gourieroux, C., & Lu, Y. (2020). SIR model with stochastic transmission. *arXiv preprint arXiv:2011.07816*.
- [28] Habott, F., Ahmedou, A., Mohamed, Y., & Sambe, M. A. (2024). Analysis of COVID-19's dynamic behavior using a modified SIR model characterized by a nonlinear function. *Symmetry*, 16(11), 1448.
- [29] Hellwig, M. (2022). SIR-Model supported by a new density.
- [30] Hethcote, H. W. (2000). The mathematics of infectious diseases. *SIAM Review*, 42(4), 599-653.
- [31] İlhan, Ö., & Şahin, G. (2024). A numerical approach for an epidemic SIR model via Morgan-Voyce series. *International Journal of Mathematics and Computer in Engineering*, 2(1), 125-140.
- [32] Keeling, M. J., & Eames, K. T. (2005). Networks and epidemic models. *Journal of the Royal Society Interface*, 2(4), 295-307.
- [33] Keeling, M. J., & Rohani, P. (2008). *Modeling infectious diseases in humans and animals*. Princeton University Press.
- [34] Keppo, J., Kudlyak, M., Quercioli, E., Smith, L., & Wilson, A. (2021). The behavioral SI* model, with applications to the swine flu and COVID-19 pandemics.
- [35] Kermack, W. O., & McKendrick, A. G. (1927). A contribution to the mathematical theory of epidemics. *Proceedings of the Royal Society of London. Series A, Containing Papers of a Mathematical and Physical Character*, 115(772), 700-721.
- [36] Kissler, S. M., et al. (2020). Projecting the transmission dynamics of SARS-CoV-2 through the postpandemic period. *Science*, 368(6493), 860-868.
- [37] Konstantinov, I. S., Tariq, T. A., & Starchenko, D. N. (2024). SIR model dynamics: Insights into epidemics and vaccination. *Экономика. Информатика*, 51(1), 145-156.
- [38] Kovtun, V., Grochla, K., Al-Maitah, M., Aldosary, S., & Gryshchuk, T. (2024). Cyber epidemic spread forecasting based on the entropy-extremal dynamic interpretation of the SIR model. *Egyptian Informatics Journal*, 28, 100572.
- [39] Kozhabek, A., Chai, W. K., & Zheng, G. (2024). Modeling traffic congestion spreading using a topology-based SIR epidemic model. *IEEE Access*, 12, 35813-35826.
- [40] Kudryashov, N., Chmykhov, M., & Vigdorowitsch, M. (2021). Analytical features of the SIR model and their applications to COVID-19. *Applied Mathematical Modelling*, 90, 466.
- [41] Law, K. B., Peariasamy, K. M., & Gill, B. S. (2020). Predicting the early depleting transmission dynamics of COVID-19: a time-varying SIR model. *Scientific Reports*, 10, 21721.
- [42] Li, M. Y. (2018). *An introduction to mathematical modeling of infectious diseases*. Springer.
- [43] Lloyd, A. L., & Jansen, V. A. (2004). Spatiotemporal dynamics of epidemics: synchrony in metapopulation models. *Mathematical Biosciences*, 188(1-2), 1-16.
- [44] Mortensen, P., Lauer, K., Rautenbach, S. P., Gallotta, M., Sharapova, N., Takkides, I., ... & Linley, M. (2024). A machine learning-enabled SIR model for adaptive and dynamic forecasting of COVID-19. *MedRxiv*, 2024-07.
- [45] Muñoz-Fernández, G. A., Seoane, J. M., & Seoane-Sepúlveda, J. B. (2021). A SIR-type model describing the successive waves of COVID-19. *Chaos, Solitons & Fractals*, 144, 110682.
- [46] Nesteruk, I. (2020). Simulations and predictions of COVID-19 pandemic with the use of SIR model.
- [47] Nesteruk, I. (2025). General SIR model for visible and hidden epidemic dynamics. *Frontiers in Artificial Intelligence*, 8, 1559880.
- [48] Newman, M. E. J. (2002). Spread of epidemic disease on networks. *Physical Review E*, 66(1), 016128.
- [49] Nikitina, A. V., Lyapunova, I. A., & Dudnikov, E. A. (2020). Study of the spread of viral diseases based on modifications of the SIR model. *Computational Mathematics and Information Technologies*, (1), 19-30.
- [50] Paggi, M. (2020). Simulation of Covid-19 epidemic evolution: are compartmental models really predictive? *arXiv preprint arXiv:2004.08207*.
- [51] Pastor-Satorras, R., & Vespignani, A. (2001). Epidemic spreading in scale-free networks. *Physical Review Letters*, 86(14), 3200.
- [52] Postnikov, E. B. (2020). Estimation of COVID-19 dynamics "on a back-of-envelope": Does the simplest SIR model provide quantitative parameters and predictions? *Chaos, Solitons & Fractals*, 135, 109841.
- [53] Pramanik, P. (2024). Estimation of optimal lock-down and vaccination rate of a stochastic SIR model: A mathematical approach. *European Journal of Statistics*, 4, 3-3.
- [54] Sarkar, K., Khajanchi, S., & Nieto, J. J. (2020). Modeling and forecasting the COVID-19 pandemic in India. *Chaos, Solitons & Fractals*, 139, 110049.
- [55] Saxena, R., Jadeja, M., & Bhateja, V. (2022). *Exploring susceptible-infectious-recovered (SIR) model for COVID-19 investigation*. Berlin/Heidelberg, Germany: Springer.
- [56] Sharif, N., Bidin, J., Akil, K. A. K., & Mazlan, S. F. (2022). The effectiveness of online video marketing on Facebook using Susceptible-Infected-Recovered (SIR) model. *Journal of Computing Research and Innovation*, 7(2), 54-65.
- [57] Shayak, B., Jahedi, S., & Yorke, J. A. (2024). Ambiguity in the use of SIR models to fit epidemic incidence data.
- [58] Siettos, C. I., & Russo, L. (2013). Mathematical modeling of infectious disease dynamics. *Virulence*, 4(4), 295-306.
- [59] Singh, A. K., Mehra, M., & Gulyani, S. (2021). A modified variable-order fractional SIR model to predict the spread of COVID-19 in India. *Mathematical Methods in the Applied Sciences*.
- [60] Tahir, H., Din, A., Shah, K., Abdalla, B., & Abdeljawad, T. (2024). Advances in stochastic epidemic modeling: tackling worm transmission in wireless sensor networks. *Mathematical and Computer Modelling of Dynamical Systems*, 30(1), 658-682.
- [61] Teng, Y., & Wei, W. (2024, September). Review on the application of the SIR model in predicting urban traffic congestion: Successes and future directions. In *Proceedings of the 2024 5th International Conference on Urban Construction and Management Engineering (ICUCME 2024)* (Vol. 242, p. 4). Springer Nature.
- [62] Tolles, J., & Luong, T. B. (2020). Modeling epidemics with compartmental models. *JAMA*, 323(24), 2515-2516.

-
- [63] Velásquez, R. M. A., & Lara, J. V. M. (2020). Forecast and evaluation of COVID-19 spreading in USA with reduced-space Gaussian process regression. *Chaos, Solitons & Fractals*, 136, 109924.
 - [64] Vespignani, A. (2012). Modelling dynamical processes in complex socio-technical systems. *Nature Physics*, 8(1), 32-39.
 - [65] Watts, D. J., & Dodds, P. S. (2007). Influentials, networks, and public opinion formation. *Journal of Consumer Research*, 34(4), 441-458.
 - [66] Zang, W., Zhang, P., Zhou, C., & Guo, L. (2015). Locating multiple sources in social networks under the SIR model: A divide-and-conquer approach. *Journal of Computational Science*, 10, 278-287.
 - [67] Zelenkov, Y., & Reshetsov, I. (2023). Analysis of the COVID-19 pandemic using a compartmental model with time-varying parameters fitted by a genetic algorithm. *Expert Systems with Applications*, 224, 119968.
 - [68] Zimmer, C., Leuba, S. I., Cohen, T., White, P. J., Hedberg, K., & Dimitrov, D. T. (2019). Accurate quantification of uncertainty in epidemic parameter estimates and predictions using stochastic compartmental models. *Statistical Methods in Medical Research*, 28(12), 3591-3608