

A SECIR MODEL BASED ON INTEGRO-DIFFERENTIAL EQUATIONS FOR EPIDEMIC OUTBREAKS

An extension of age-of-infection models

Anna Wendler

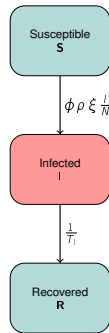
Joint work with *Lena Plötzke* and *Martin J. Kühn*

SIR model based on ordinary differential equations (ODE)

$$S'(t) = -\frac{S(t)}{N} \phi(t) \rho(t) \xi(t) I(t)$$

$$I'(t) = \frac{S(t)}{N} \phi(t) \rho(t) \xi(t) I(t) - \frac{1}{T_I} I(t)$$

$$R'(t) = \frac{1}{T_I} I(t)$$



$\phi(t)$: Number of contacts at time t

$\rho(t)$: Transmission probability at time t

$\xi(t)$: Proportion of infected individuals that are not isolated at time t

T_I : Mean stay time in compartment I

- Simple ODE models are restricted to exponentially distributed stay times in compartments
⇒ Unrealistic assumption¹
 - Choice of transition distribution impacts disease dynamics, in particular at change points
- ⇒ Need for flexible choice of transition distributions
- ⇒ Use model based on integro-differential equations (IDE)

¹Wearing et al., PLOS Medicine (2005), <https://doi.org/10.1371/journal.pmed.0020174>.

$$S'(t) = -\frac{S(t)}{N} \phi(t) \rho(t) \int_{-\infty}^t \xi(t, t-x) \underbrace{\gamma_i^R(t-x) \underbrace{\sigma_S^I(x)}_{\text{New infections at time } x}}_{\text{Number of individuals that are still infected at time } t} dx$$

$\sigma_S^I(x)$: Number of individuals transitioning from S to I at time x

$\gamma_i^R(\tau)$: Mean proportion of individuals that are still infected at infection age τ

$\phi(t)$: Number of contacts at time t

$\rho(t)$: Transmission probability at time t

$\xi(t, \tau)$: Proportion of infected individuals that are not isolated at time t and infection age τ

$$\begin{aligned} S'(t) &= -\frac{S(t)}{N} \phi(t) \rho(t) \int_{-\infty}^t \xi(t, t-x) \gamma_I^R(t-x) \sigma_S^I(x) dx \\ &= -S(t) \varphi(t) \end{aligned}$$

$\sigma_S^I(x)$: Number of individuals transitioning from S to I at time x

$\gamma_I^R(\tau)$: Mean proportion of individuals that are still infected at infection age τ

$\phi(t)$: Number of contacts at time t

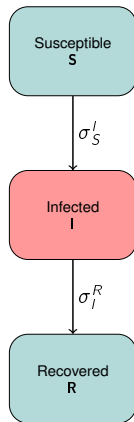
$\rho(t)$: Transmission probability at time t

$\xi(t, \tau)$: Proportion of infected individuals that are not isolated at time t and infection age τ

$$S'(t) = -\frac{S(t)}{N} \phi(t) \rho(t) \int_{-\infty}^t \xi(t, t-x) \gamma_i^R(t-x) \sigma_S^I(x) dx$$

$$I(t) = \int_{-\infty}^t \gamma_i^R(t-x) \sigma_S^I(x) dx$$

$$R(t) = \int_{-\infty}^t (1 - \gamma_i^R(t-x)) \sigma_S^I(x) dx = \int_{-\infty}^t \sigma_i^R(x) dx$$

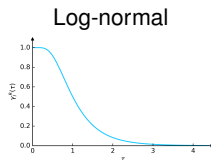
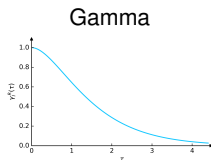
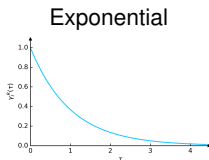


$$S'(t) = -\frac{S(t)}{N} \phi(t) \rho(t) \int_{-\infty}^t \xi(t, t-x) \gamma_i^R(t-x) \sigma_S^I(x) dx$$

$$I(t) = \int_{-\infty}^t \gamma_i^R(t-x) \sigma_S^I(x) dx$$

$$R(t) = \int_{-\infty}^t (1 - \gamma_i^R(t-x)) \sigma_S^I(x) dx$$

Flexible choice of transition distributions:



$$\sigma_S^I(x) = -S'(x) = S(x)\varphi(x)$$

$$\begin{aligned}\sigma_I^R(x) &= \frac{d}{dx} \left(\int_{-\infty}^x (1 - \gamma_I^R(x-u)) \sigma_S^I(u) du \right) \\ &= - \int_{-\infty}^x \gamma_I^{R'}(x-u) \sigma_S^I(u) du\end{aligned}$$

$$\begin{aligned} S'(t) &= -S(t) \varphi(t) \\ &= -\frac{S(t)}{N} \phi(t) \rho(t) \int_{-\infty}^t \xi(t, t-x) \gamma_l^R(t-x) \sigma_S^l(x) dx \end{aligned}$$

Non-standard finite difference scheme¹:

$$\widehat{S}'(t_{n+1}) = -\widehat{\sigma}_S^l(t_{n+1}) = -\widehat{S}(t_{n+1}) \widehat{\varphi}(t_n)$$

¹Messina et al., *Journal of Computational Dynamics* (2022), <http://dx.doi.org/10.3934/jcd.2021029>.

Backwards difference scheme for \widehat{S}' :

$$\frac{\widehat{S}(t_{n+1}) - \widehat{S}(t_n)}{\Delta t} = -\widehat{S}(t_{n+1})\widehat{\varphi}(t_n)$$
$$\iff \widehat{S}(t_{n+1}) = \frac{\widehat{S}(t_n)}{1 + \Delta t \widehat{\varphi}(t_n)}$$

Non-standard rectangular rule to approximate integral:

$$\widehat{\varphi}(t_{n+1}) = \frac{1}{N} \phi(t_{n+1}) \rho(t_{n+1}) \Delta t \sum_{i=a}^n \xi(t_{n+1}, t_{n+1-i}) \gamma_i^R(t_{n+1-i}) \widehat{\sigma}_S^l(t_{i+1})$$

Flows:

$$\hat{\sigma}_S^I(t_{n+1}) = \hat{S}(t_{n+1}) \hat{\varphi}(t_n)$$

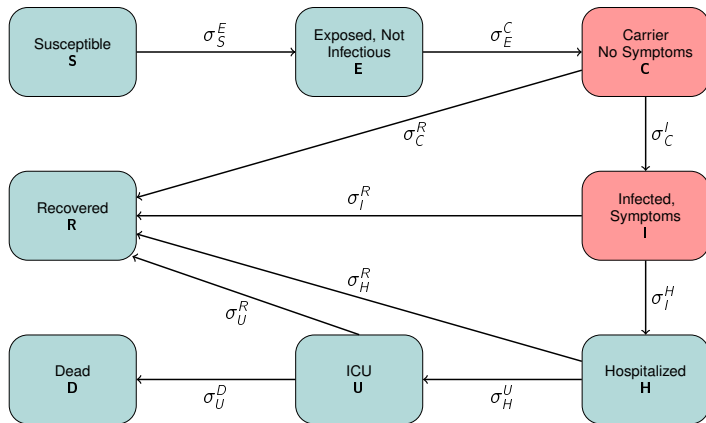
$$\hat{\sigma}_I^R(t_{n+1}) = -\Delta t \sum_{i=a}^n \gamma_i^{R'}(t_{n+1-i}) \hat{\sigma}_S^I(t_{i+1})$$

Remaining compartments:

$$\hat{I}(t_{n+1}) = \hat{I}(t_n) + \Delta t \hat{\sigma}_S^I(t_{n+1}) - \Delta t \hat{\sigma}_I^R(t_{n+1})$$

$$\hat{R}(t_{n+1}) = \hat{R}(t_n) + \Delta t \hat{\sigma}_I^R(t_{n+1})$$

Extension to IDE-SECIR model^{1,2}



¹ Kühn et al., *Mathematical Biosciences* (2021), <https://doi.org/10.1016/j.mbs.2021.108648>.

² Wendler et al., in preparation.

Theorem¹

Let $\Delta t > 0$ be arbitrary but fixed. Let $\frac{d}{d\tau} \gamma_{z_1}^{z_2}(\tau)$ be approximated using a backwards difference scheme and let there hold

$$\widehat{S}(t_0) + \widehat{E}(t_0) + \widehat{C}(t_0) + \widehat{I}(t_0) + \widehat{H}(t_0) + \widehat{U}(t_0) + \widehat{R}(t_0) + \widehat{D}(t_0) = N.$$

Then the following holds.

- For all $n \in \mathbb{N}$ there holds

$$\widehat{S}(t_n) + \widehat{E}(t_n) + \widehat{C}(t_n) + \widehat{I}(t_n) + \widehat{H}(t_n) + \widehat{U}(t_n) + \widehat{R}(t_n) + \widehat{D}(t_n) = N.$$

- The sequence $(\widehat{S}(t_n))_{n \in \mathbb{N}_0}$ is monotonically decreasing and

$$\lim_{n \rightarrow \infty} \widehat{S}(t_n) = \widehat{S}_\infty(\Delta t).$$

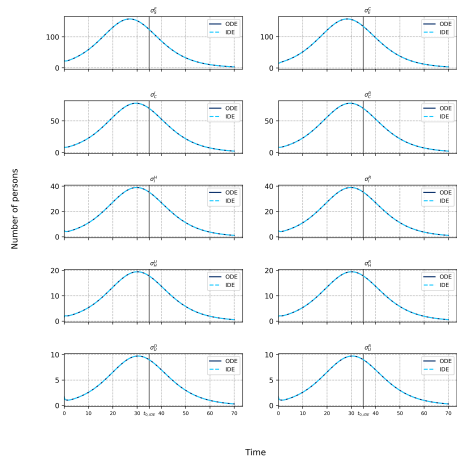
- The sequences $(\widehat{\sigma}_{z_1}^{z_2}(t_n))_{n \in \mathbb{N}_0}$ are bounded and

$$\lim_{n \rightarrow \infty} \widehat{\sigma}_{z_1}^{z_2}(t_n) = 0.$$

¹Wendler et al., in preparation.

Comparison of IDE and ODE model

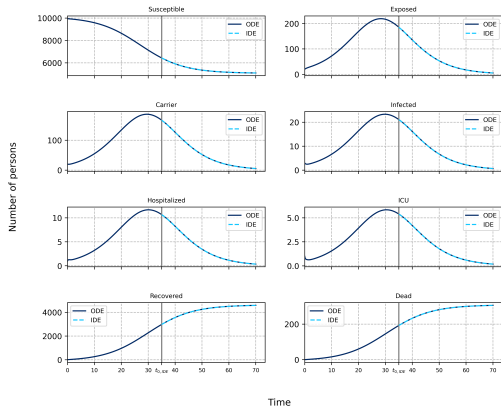
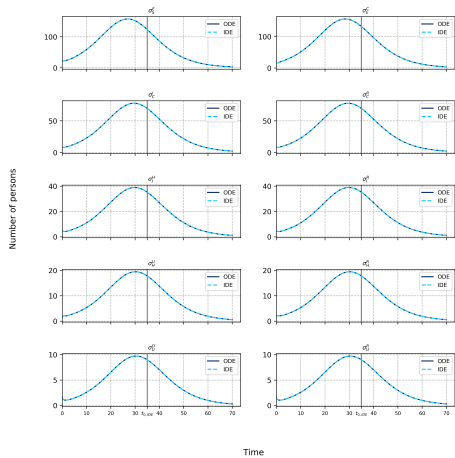
- With appropriate choice of parameters: IDE model reduces to ODE model



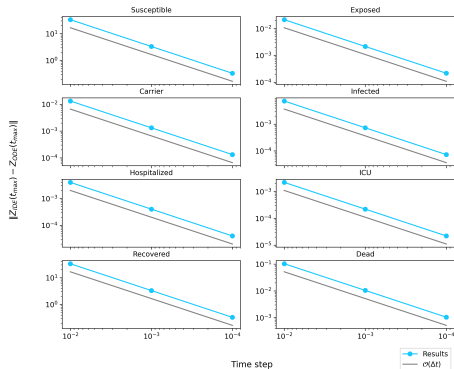
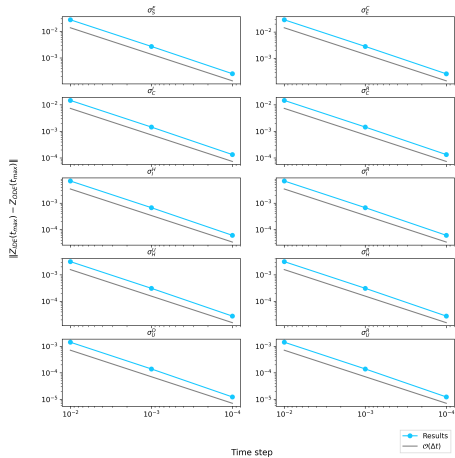
- To validate our discretization:

- Solve ODE model with Runge-Kutta using small time step
⇒ Define this as ground truth
- Compute initial flows for IDE model until $t_{0,IDE}$ based on ODE results
- Simulate IDE model and compare with ODE results

Comparison of IDE and ODE model

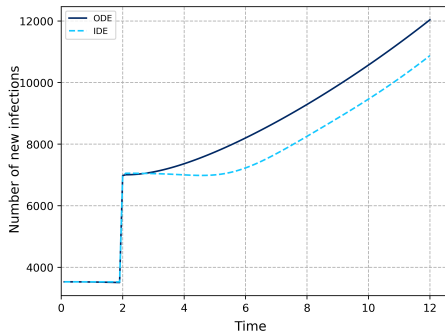
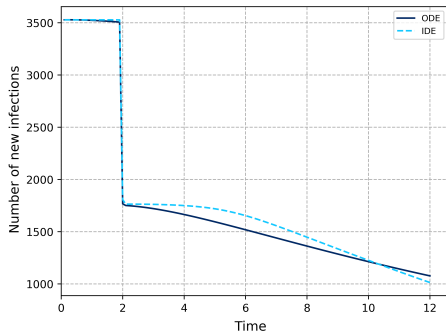


Convergence of IDE model



Behavior at change points

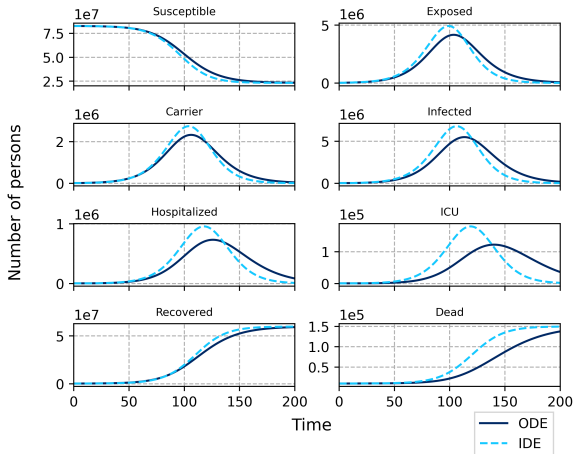
- In IDE model: Log-normally distributed stay times¹
- Change contact rate to model implementation and cancelation of NPIs



- IDE model reacts slower to change in contact rate than ODE model

¹Kerr et al., PLOS Computational Biology (2021), <https://doi.org/10.1371/journal.pcbi.1009149>.

Epidemic peak behavior for compartments



- Higher maximum in Hospitalized and ICU compartments when using IDE model

Conclusion



- Using IDE model allows flexible choice of transition distributions
- Application of non-standard numerical scheme
 - Conserves important properties of disease dynamics
 - Converges linearly
- Choice of distributions has significant impact on disease dynamics

Thank you for your attention!



GitHub repository: <https://github.com/SciCompMod/memilio>

$$S'(t) = -\frac{S(t)}{N - D(t)} \phi(t) \rho(t) \\ \int_{-\infty}^t \xi_C(t, t-x) (\mu_C^I \gamma_C^I(t-x) + (1 - \mu_C^I) \gamma_C^R(t-x)) \sigma_E^C(x) \\ + \xi_I(t, t-x) (\mu_I^H \gamma_I^H(t-x) + (1 - \mu_I^H) \gamma_I^R(t-x)) \sigma_C^I(x) dx$$

IDE-SECIR model

$$E(t) = \int_{-\infty}^t \gamma_E^C(t-x) \sigma_S^E(x) dx$$

$$C(t) = \int_{-\infty}^t (\mu_C^I \gamma_C^I(t-x) + (1 - \mu_C^I) \gamma_C^R(t-x)) \sigma_E^C(x) dx$$

$$I(t) = \int_{-\infty}^t (\mu_I^H \gamma_I^H(t-x) + (1 - \mu_I^H) \gamma_I^R(t-x)) \sigma_C^I(x) dx$$

$$H(t) = \int_{-\infty}^t (\mu_H^U \gamma_H^U(t-x) + (1 - \mu_H^U) \gamma_H^R(t-x)) \sigma_I^H(x) dx$$

$$U(t) = \int_{-\infty}^t (\mu_U^D \gamma_U^D(t-x) + (1 - \mu_U^D) \gamma_U^R(t-x)) \sigma_H^U(x) dx$$

$$R(t) = \int_{-\infty}^t \sigma_C^R(x) + \sigma_I^R(x) + \sigma_H^R(x) + \sigma_U^R(x) dx$$

$$D(t) = \int_{-\infty}^t \sigma_U^D(x) dx.$$

$$\sigma_S^E(t) = -S'(t) = S(t)\varphi(t)$$

$$\sigma_C^I(t) = -\int_{-\infty}^t \gamma_C^{I'}(t-x) \mu_C^I \sigma_E^C(x) dx$$

$$\sigma_I^H(t) = -\int_{-\infty}^t \gamma_I^{H'}(t-x) \mu_I^H \sigma_C^I(x) dx$$

$$\sigma_H^U(t) = -\int_{-\infty}^t \gamma_H^{U'}(t-x) \mu_H^U \sigma_I^H(x) dx$$

$$\sigma_U^D(t) = -\int_{-\infty}^t \gamma_U^{D'}(t-x) \mu_U^D \sigma_H^U(x) dx$$

$$\sigma_E^C(t) = -\int_{-\infty}^t \gamma_E^{C'}(t-x) \sigma_S^E(x) dx$$

$$\sigma_C^R(t) = -\int_{-\infty}^t \gamma_C^{R'}(t-x) (1 - \mu_C^I) \sigma_E^C(x) dx$$

$$\sigma_I^R(t) = -\int_{-\infty}^t \gamma_I^{R'}(t-x) (1 - \mu_I^H) \sigma_C^I(x) dx$$

$$\sigma_H^R(t) = -\int_{-\infty}^t \gamma_H^{R'}(t-x) (1 - \mu_H^U) \sigma_I^H(x) dx$$

$$\sigma_U^R(t) = -\int_{-\infty}^t \gamma_U^{R'}(t-x) (1 - \mu_U^D) \sigma_H^U(x) dx$$