



University of Idaho
College of Science



**Systems Medicine
of Infectious Diseases**
Research and Consulting Group

Adaptive Observers for Medical Digital Twins

A Framework for personalized medicine

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University of Idaho

Panel 1. A 52-year-old male with a history of stroke and hemiplegia is found confused in his prison cell.



Taken, and adapted with Gemini, from:

[1] Laubenbacher, R., Mehrad, B., Shmulevich, I., & Trayanova, N. (2024). **Digital twins in medicine.** Nature Computational Science, 4(3), 184-191.

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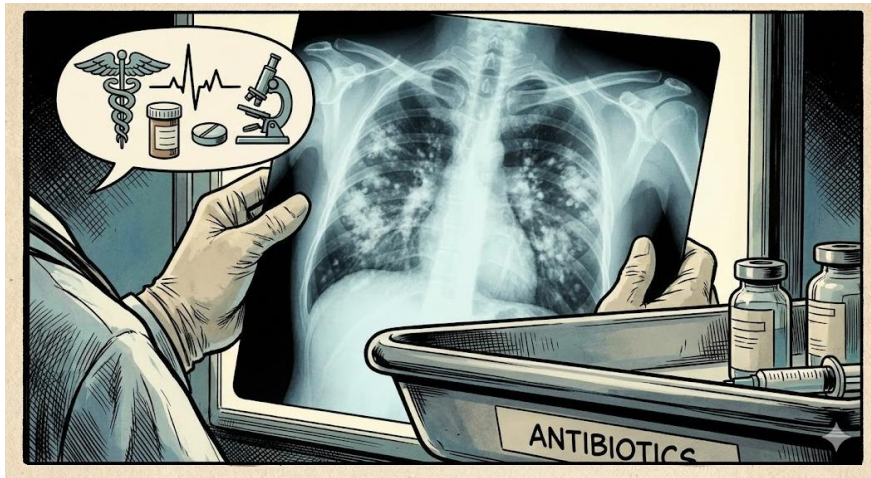
Panel 2. At ED, patient presents hypotension, mild renal insufficiency, and bilateral airspace disease on chest X-ray.



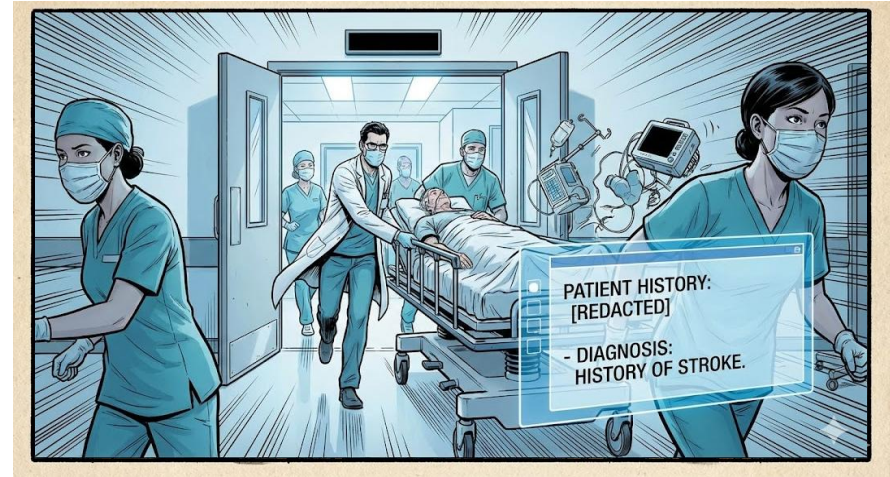
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Panel 3. Pneumonia diagnosed and empiric severe pathogen antibiotics started.



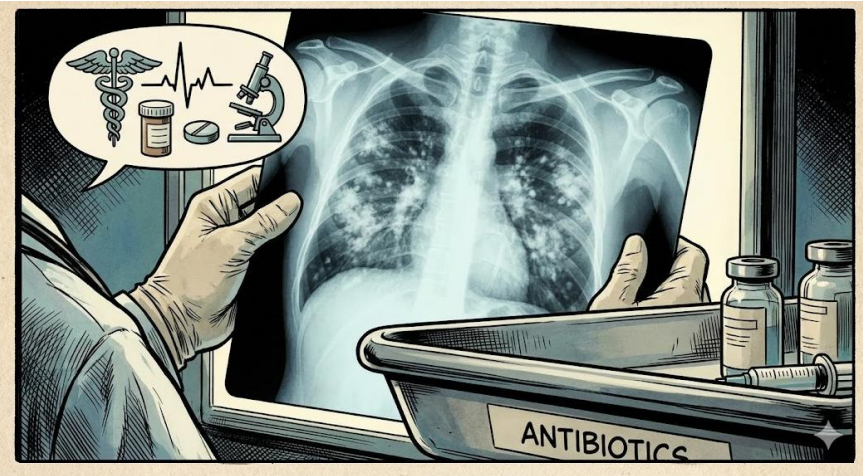
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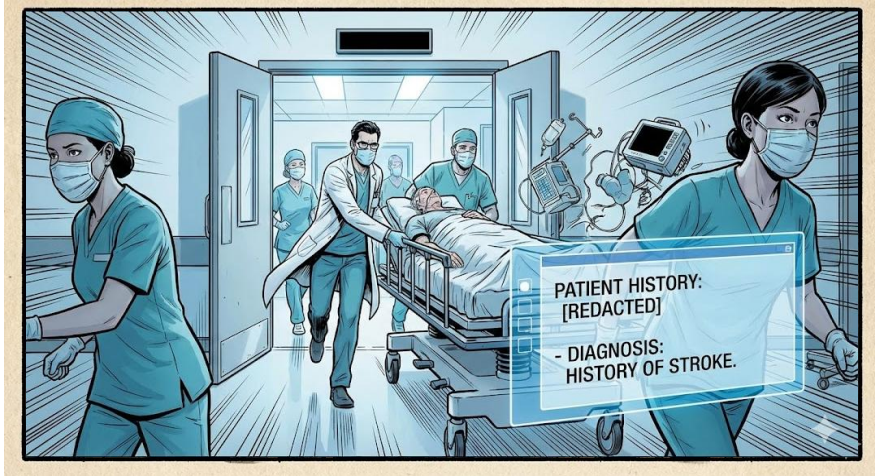


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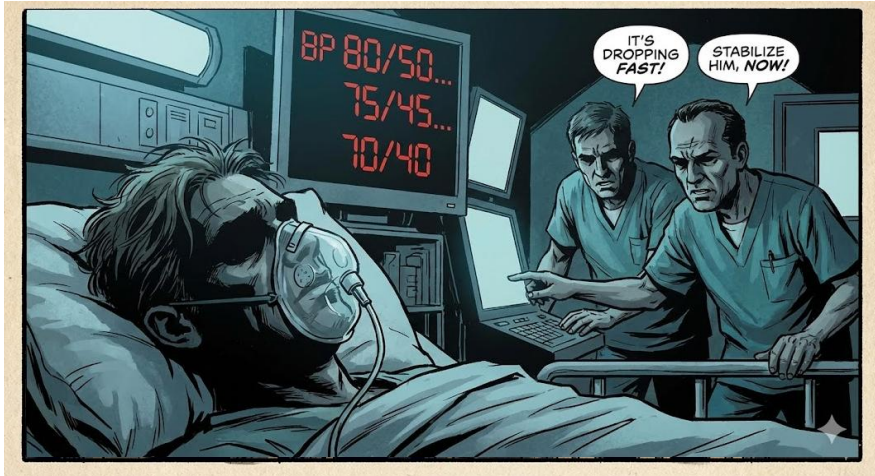


Suspected pathogen: **Klebsiella pneumoniae**.

Panel 2. At ED, patient presents hypotension, mild renal insufficiency, and bilateral airspace disease on chest X-ray.



Panel 4. 6 h later, recurrent hypotension requiring intravenous pressors and escalating supplemental oxygen.



Panel 5. 3 h after deterioration, requires endotracheal intubation and mechanical ventilation for ARDS-induced hypoxic respiratory failure.



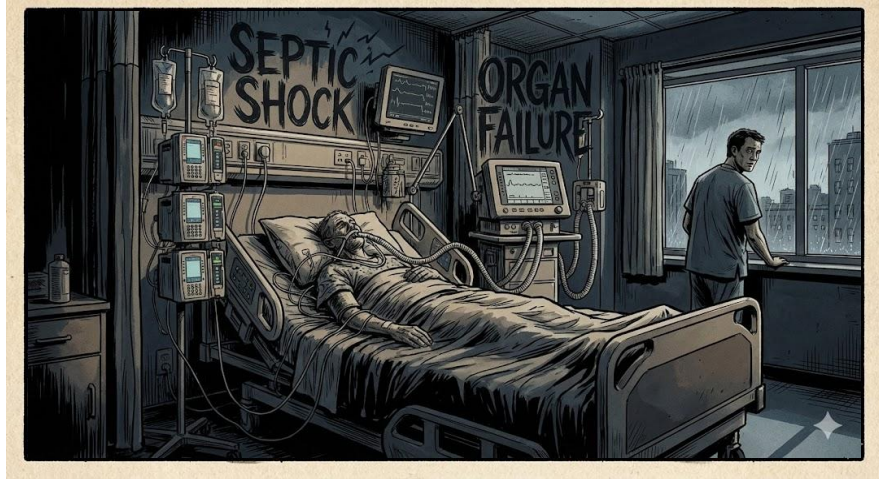
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Panel 6. Condition progress to refractory septic shock and acute kidney injury **despite receiving antibiotics to which the *Klebsiella pneumoniae* was sensitive.**



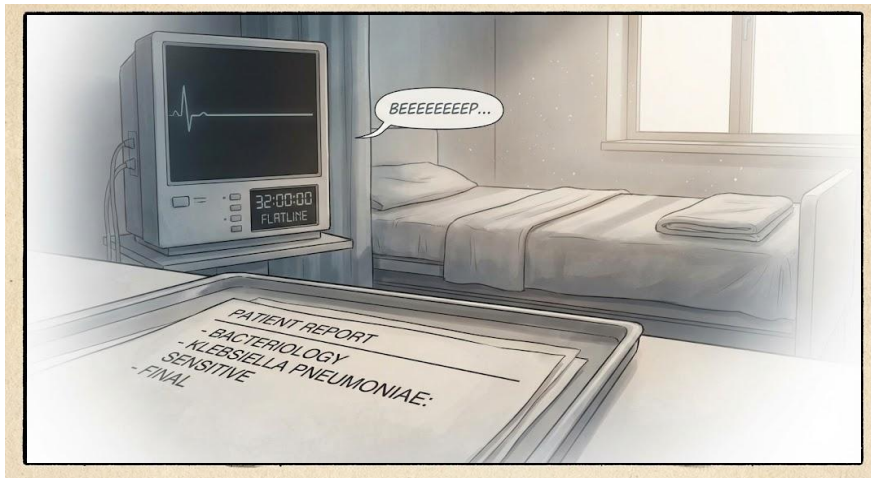
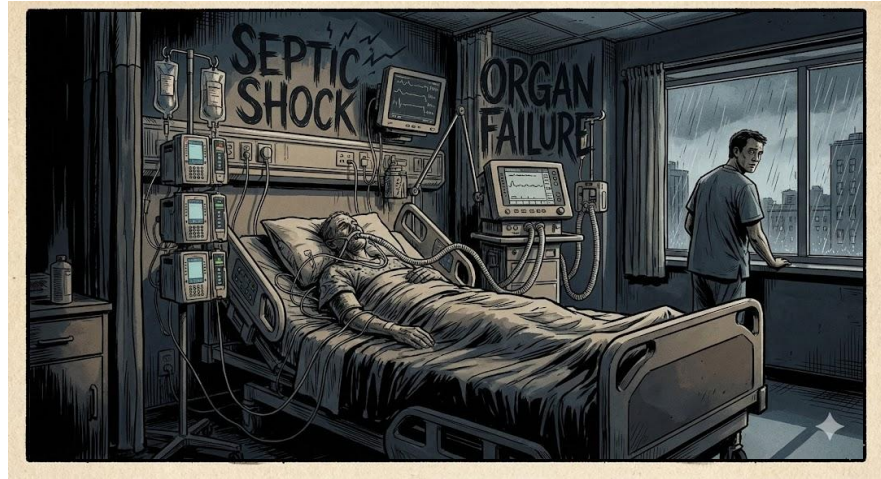
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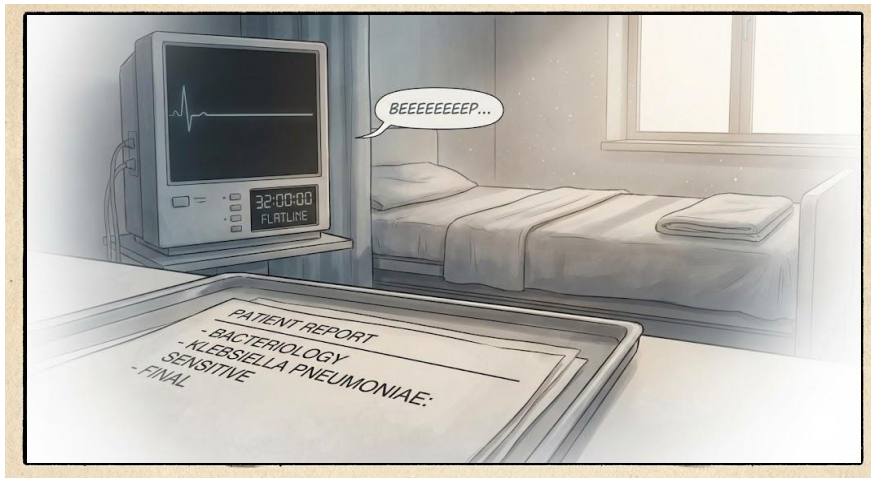
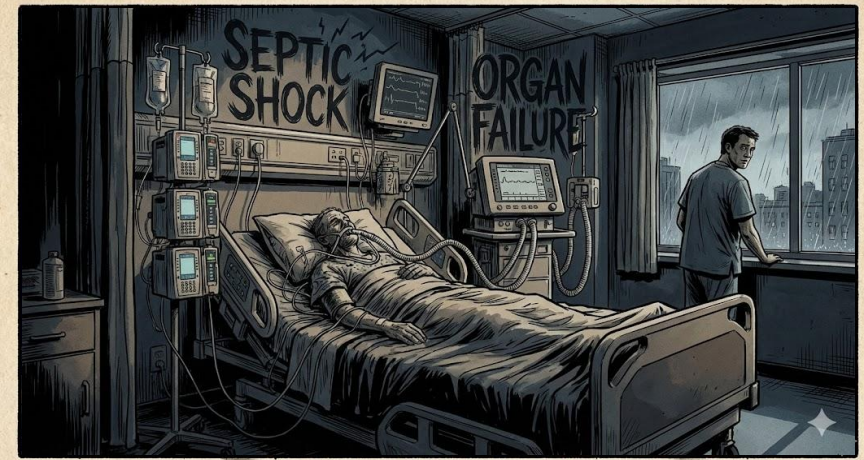
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Why we need **Digital Twins in medicine?**
This one represent the biggest promise and challenge in applying computational modeling at the level of individual patients, i.e., personalized medicine.

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1. A model of the physical twin.
2. An evolving set of data relating to the object, and
3. A means of dynamically updating or adjusting the model in accordance with data.

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A **digital twin** is an integrated multiphysics, multi-scale, probabilistic simulation of a vehicle or system that uses the best available physical models, sensor updates, flight history, etc., to mirror the life of its flying twin. The **digital twin** is ultra-realistic and may consider one or more important and interdependent vehicle systems, including propulsion and energy storage, life support, avionics, thermal protection, etc.

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A trade-off arises

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Medical Digital Twin [1,4]:

A computational model of the system to be twinned (heathy or sickness processes) that is connected to the physical twin in a bidirectional fashion over time, periodically recalibrated with patient data, and provides patient predictions over time to prevent or treat diseases.

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[4] **Foundational Research Gaps and Future Directions for Digital Twins** (National Academies of Sciences, Engineering and Medicine, 2023); <https://doi.org/10.17226/26894>.

Digital Twins in Medicine

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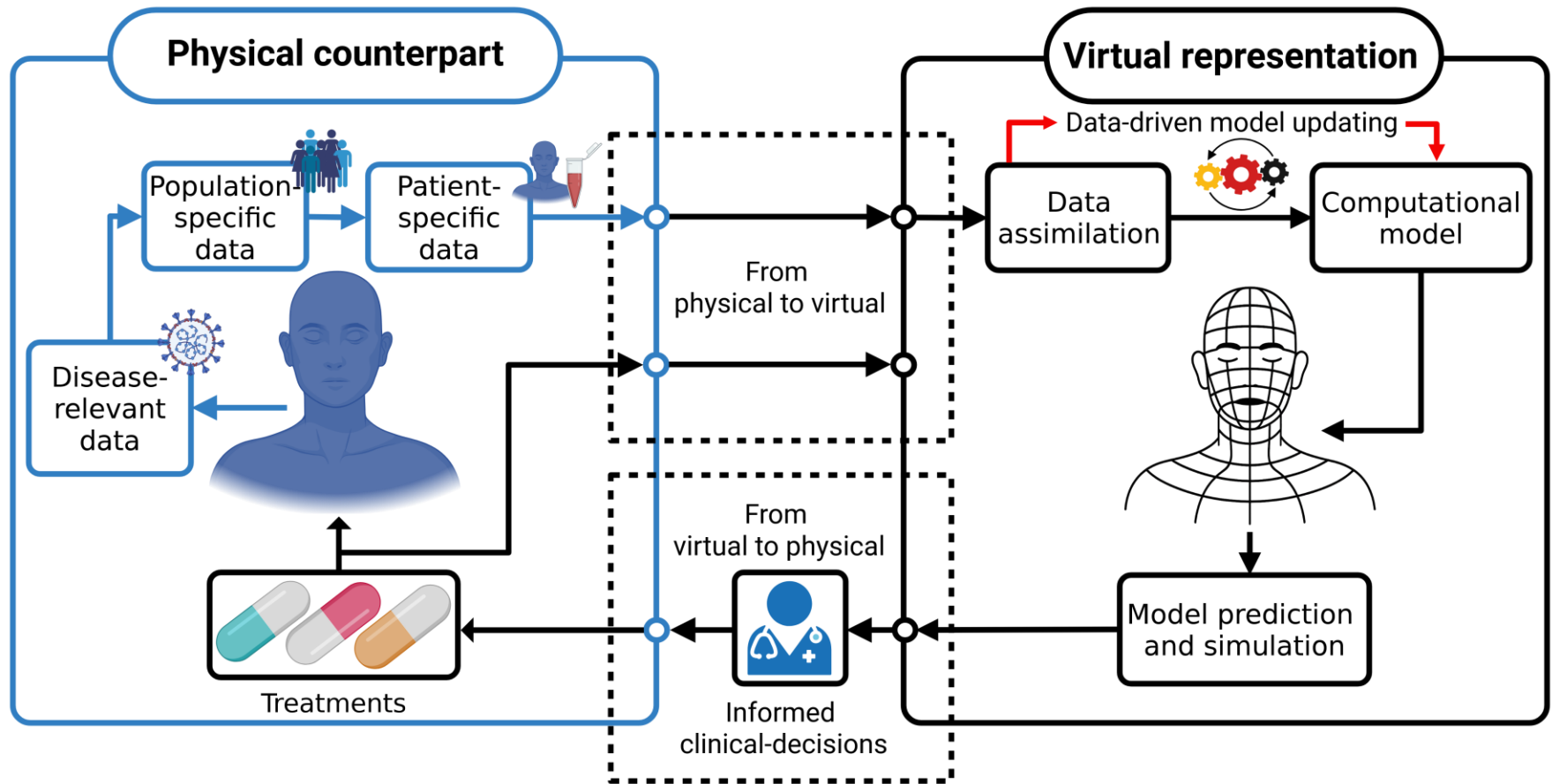


Figure 1.a. The standard Digital Twin Framework [5] .

Digital Twins in Medicine

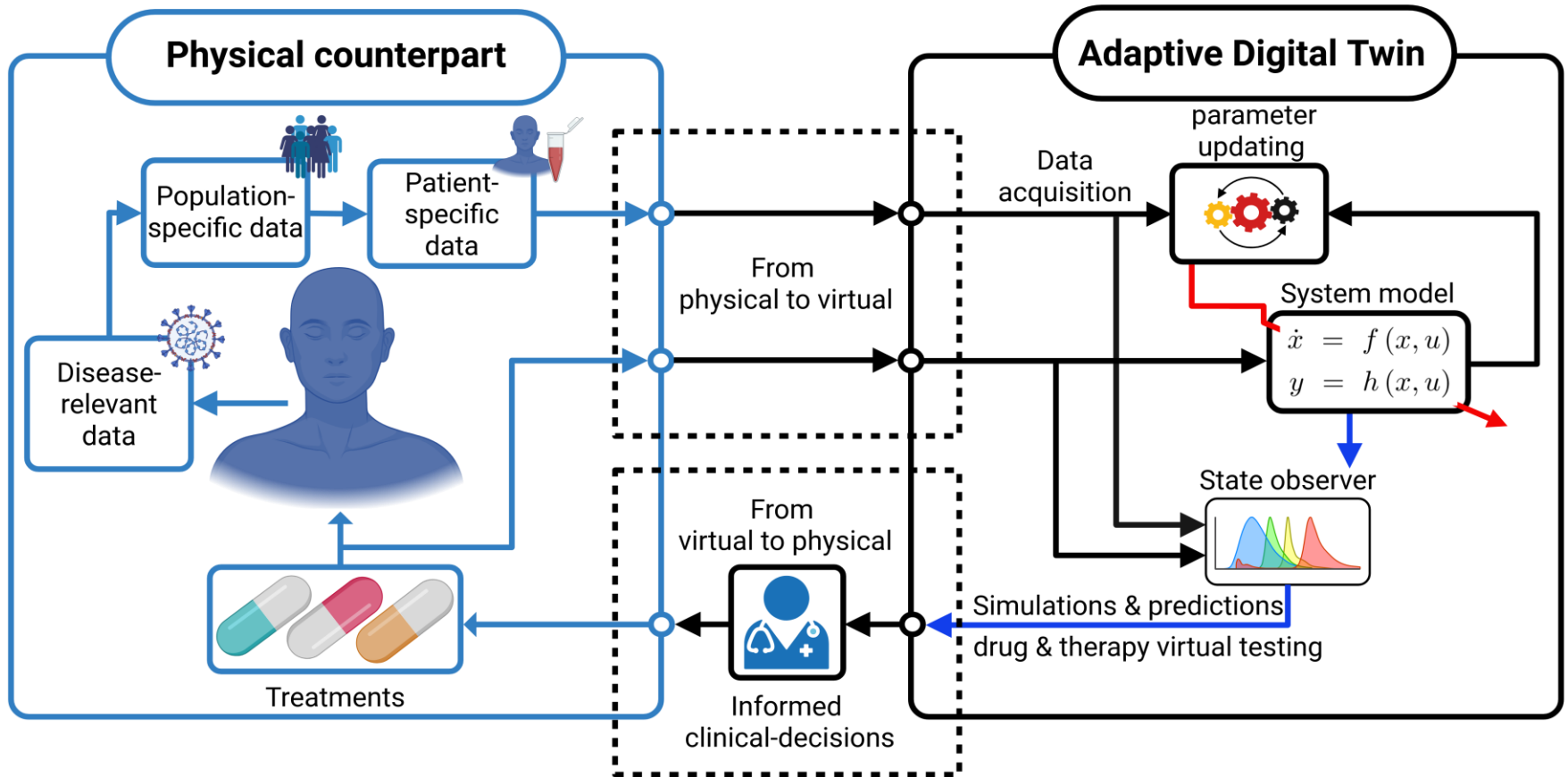


Figure 1.b. Adaptive Digital Twin Framework [5] .

Mathematical foundations for model parameter updating

Model Parameter Updating:

1. Adaptive Observers [6].
2. Hybrid Extended Kalman Filter [7].

Taken from:

[6] Besancon, G., de León-Morales, J., & Huerta-Guevara, O. (2006). *On adaptive observers for state affine systems*. International journal of Control, 79(06), 581-591.

[7] Simon, D. (2006). *Optimal state estimation: Kalman, H infinity, and nonlinear approaches*. John Wiley & Sons.

Mathematical foundations for model parameter updating

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Biomedical applications:

1. HIV mutation model [8].
2. Antimicrobial Resistance model [9] .

Taken from:

[8] Hernandez-Vargas, E. A., & Middleton, R. H. (2013). ***Modeling the three stages in HIV infection***. Journal of theoretical biology, 320, 33-40.

[9] Hernández-Vargas, E. A., Parra-Rojas, C., & Olaru, S. (2021, December). ***Switching logistic maps to design cycling approaches against antimicrobial resistance***. In 2021 60th IEEE conference on decision and control (CDC) (pp. 4248-4253). IEEE.

HIV mutation model

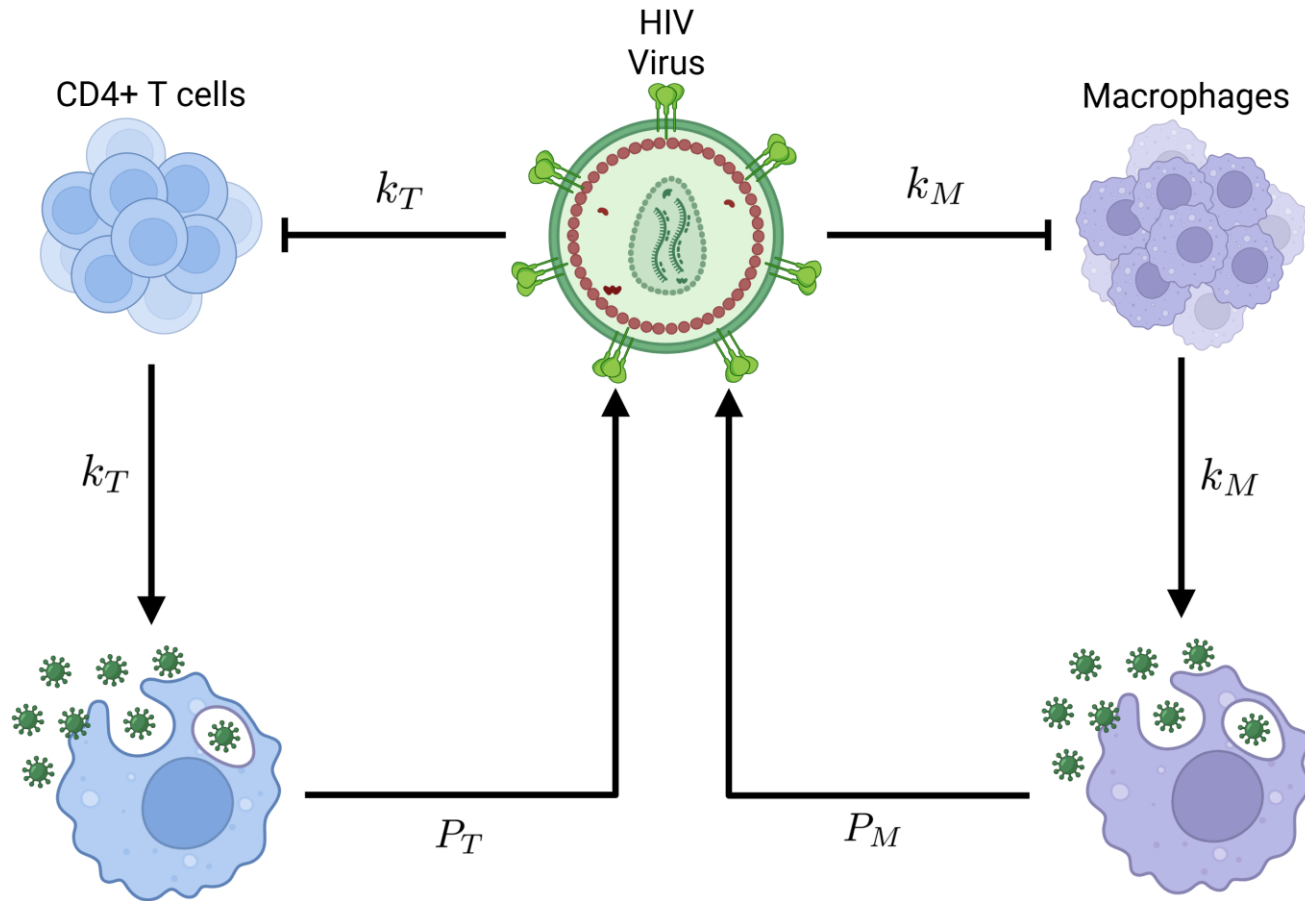


Figure 2. HIV infection scheme, taken from [8].

HIV mutation model

$$\dot{T} = s_T + \frac{\rho_T}{C_T + V_T} T V_T - \sum_{i=1}^n k_{T,\sigma}^i T V_i - \delta_T T$$

$$\dot{M} = s_M + \frac{\rho_M}{C_M + V_T} M V_T - \sum_{i=1}^n k_{M,\sigma}^i M V_i - \delta_M M$$

$$\dot{T}_i^* = k_{T,\sigma}^i T V_i + \sum_{j=1}^n \mu m_{i,j} V_j T - \delta_{T^*} T_i^*$$

$$\dot{M}_i^* = k_{M,\sigma}^i M V_i + \sum_{j=1}^n \mu m_{i,j} V_j M - \delta_{M^*} M_i^*$$

$$\dot{V}_i = p_{T,\sigma}^i T_i^* + p_{M,\sigma}^i M_i^* - \delta_V V_i.$$

HIV mutation model

$$\dot{T} = s_T + \frac{\rho_T}{C_T + V_T} T V_T - \sum_{i=1}^n k_{T,\sigma}^i T V_i - \delta_T T$$

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$$\dot{V}_i = p_{T,\sigma}^i T_i^* + p_{M,\sigma}^i M_i^* - \delta_V V_i. \quad (1)$$

Can we update model parameters to estimate infections rates and the number of infected cells from measurements?



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COVID-19



Our research group seeks new avenues for holistic understanding against infections at the meeting point of mathematics, machine learning, and infection biology. By integrating multi-scale experimental data, modeling, and the power of computational simulations we aim to:

COVID-19 Research

Tweets by @systmedicine

Esteban A. Hernandez-Vargas Retweeted
Jesus Alvarez
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Contact us at:

esteban@systemsmedicine.de

Are you enjoying this work so far?

**Contact jserenom@uidaho.edu for
the extended version of the slides**

References

- [1] Wright, L., & Davidson, S. (2020). How to tell the difference between a model and a digital twin. AMSES, 7, 1-13.
- [2] Hernandez-Vargas, E. A., & Middleton, R. H. (2013). Modeling the three stages in HIV infection. Journal of theoretical biology, 320, 33-40.
- [3] Rodríguez, A., et al. An adaptive observer for operation monitoring of anaerobic digestion wastewater treatment. J. Chem. Eng. 269 (2015): 186-193.
- [4] Hernandez-Vargas, E. A. (2019). Modeling and control of infectious diseases in the host: with MATLAB and R. Academic Press.