



BioGears: New Models and Capabilities

The Advantages of Continuous Model Development

17 MARCH 2020

Outline

- New model motivation
- Inflammation
 - Infection / Sepsis
 - Burn
 - Antibiotic Pharmacodynamics
- Cerebral Model expansion
- Whole Blood
- Models in development

Motivation

- BioGears model development is driven by:
 - Demonstrated past performance to JPC-1
 - BioGears Follow-On contract
 - Collaboration with medical simulation community
 - Virtual Heroes Division (VHD) at ARA
 - United States Army Institute of Surgical Research (USAISR)
 - University of Washington—WWAMI Institute for Simulation in Healthcare (WISH)
 - Advanced Modular Manikin (AMM)
 - User Buy-In
 - Outreach on Github



BioGears Inflammatory Response Model

A dynamic system that mediates response to thermal injury and infection

Applications



Inflammatory Response

- WBC = White blood cell count
- T = Temperature
- RR = Respiration rate
- NO = Nitric oxide
- D = Tissue Damage

Pain Sub-model

- Epi = Epinephrine
- RR = Respiration rate

Hypovolemia Response

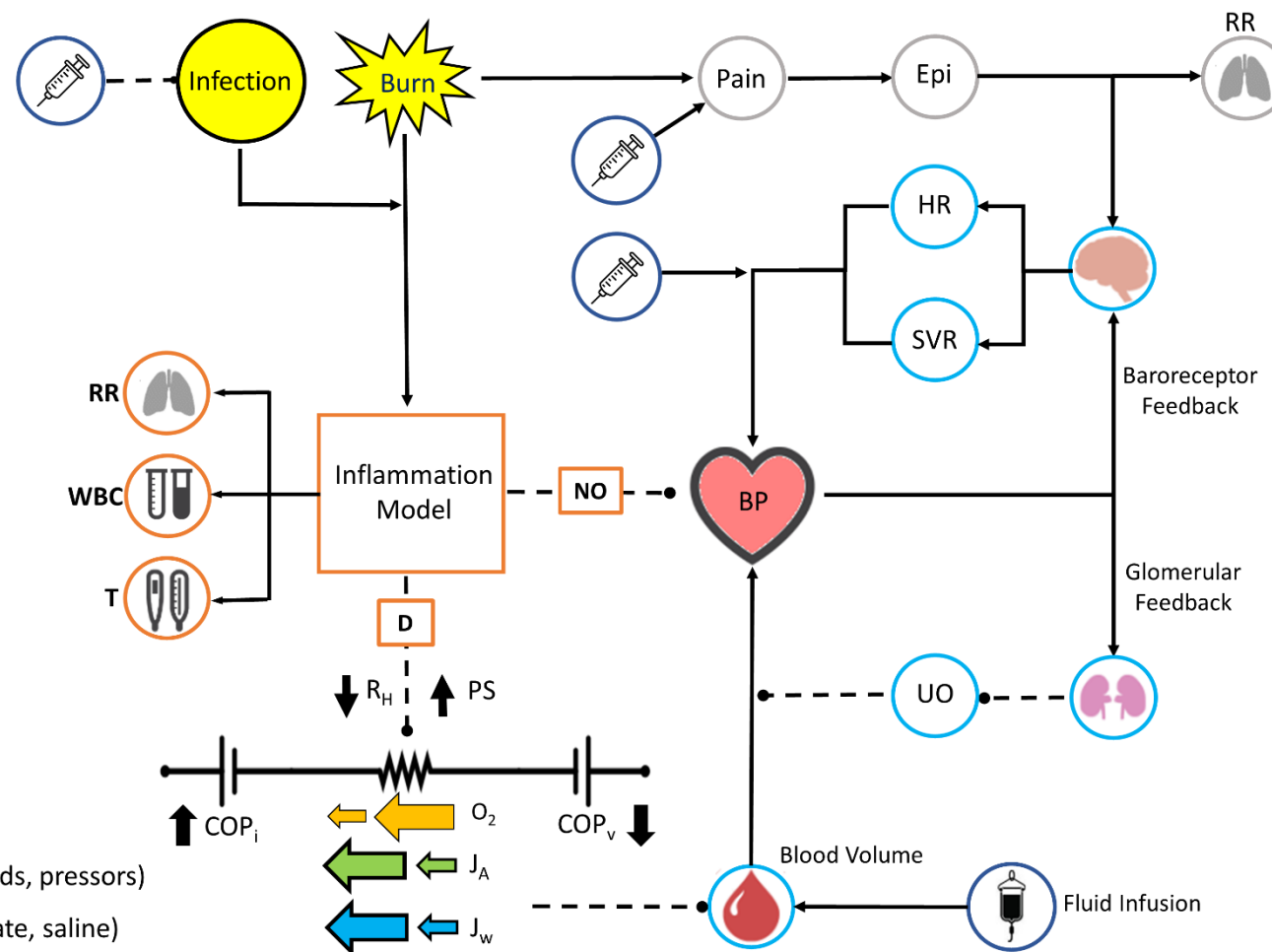
- HR = Heart rate
- SVR = Systemic vascular resistance
- UO = Urine output
- BP = Blood pressure

Microcirculation Sub-model

- $COP_{i/v}$ = Colloid osmotic pressure
- J_w = Fluid flux
- J_A = Albumin flux
- R_H = Hydraulic resistance
- PS = Solute permeability

Treatment Options

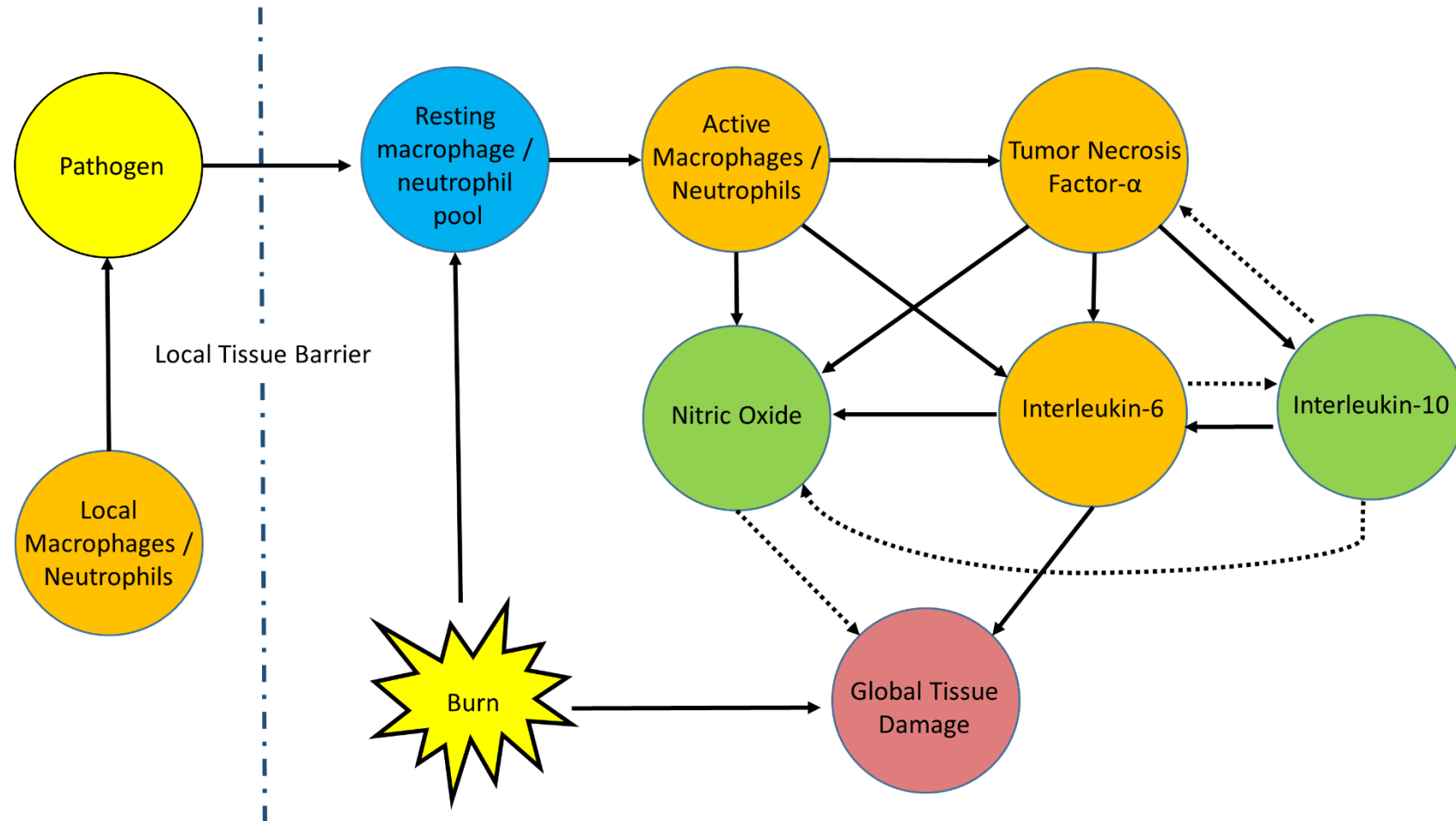
- Medication (antibiotics, pain meds, pressors)
- Fluid resuscitation (Ringer's lactate, saline)



Inflammation Model

Pathogen invasion and thermal trauma initiate the inflammatory response model

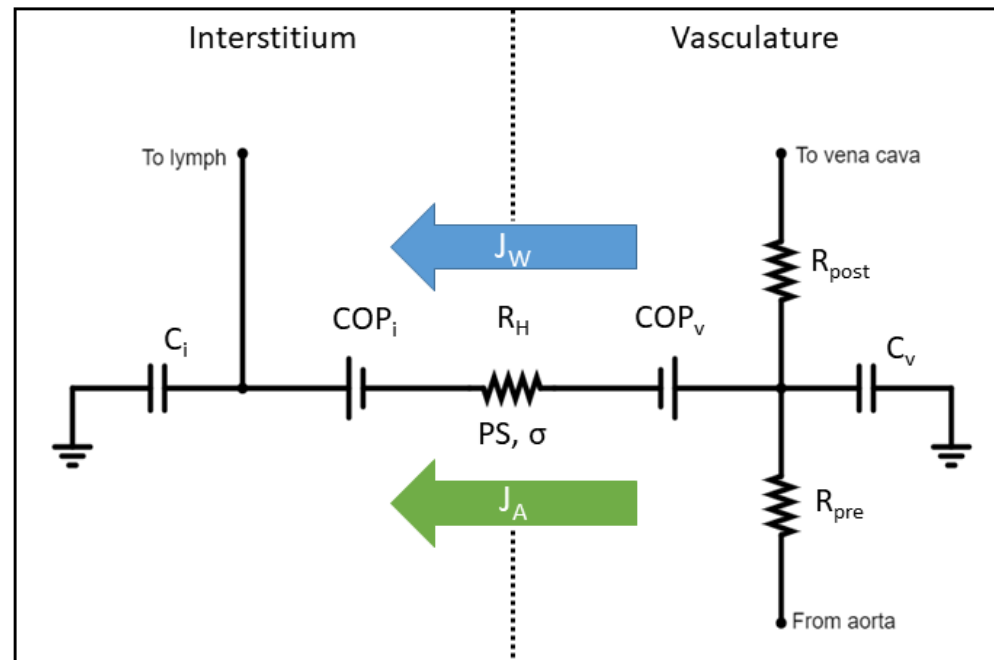
- Infection^[1-4]
 - Pathogen originates in tissue where it is combated by local immune mediators
 - Inflammation weakens tissue barrier integrity
 - Pathogen can enter circulation, leading to systemic inflammatory response (SIRS)
- Burn^[2,4]
 - Thermal trauma activates systemic inflammatory response
 - Inflicts initial hit on global tissue health



Microcirculation Model

Inflammatory-mediated tissue damage causes fluid shift and relative hypovolemia

- BioGears models the cardiovascular system as a fluid circuit with compartments representing distinct regions (e.g. muscle, skin, liver)
- Each vascular compartment communicates with an interstitial compartment, transferring fluid and substances
- Filtered fluids and substances are returned to the vasculature via a lymph pathway (amount returned = amount filtered at steady state)



σ = Membrane reflection coefficient

$C_{i/v}$ = Interstitial / vascular compliance

$R_{pre/post}$ = Pre / post-capillary resistance

- Query volumetric flux (J_W) from BioGears circuit solver
- For vascular (v) and interstitial (i) compartments:

1. Calculate total plasma protein concentration (C_{pp}) from albumin concentration (C_A) assuming:^[8]

$$C_{pp} = 1.6C_A$$

2. Calculate colloid osmotic pressure (COP)^[8]

$$COP = 2.1 \cdot C_{pp} + 0.18 \cdot C_{pp}^2 + 0.009 \cdot C_{pp}^3$$

3. Calculate albumin flux (J_A)^[9]

$$J_A = J_W \cdot (1 - \sigma) \cdot \frac{C_{A,v} - C_{A,i} \cdot \exp\left(-J_W \cdot \frac{(1 - \sigma)}{PS}\right)}{1 - \exp\left(-J_W \cdot \frac{(1 - \sigma)}{PS}\right)}$$

4. Update J_W (function of COP, hydrostatic pressure, and R_H)

* For more on lumped-parameter cardiopulmonary modeling, see refs [6, 7]

Antibiotic Pharmacodynamics Model

- Assume that antibiotics act by decreasing the net bacterial growth rate^[10-11]

$$k_{net} = k_{max} - \frac{(k_{max} - k_{min}) \cdot (C_u / MIC)^\delta}{(C_u / MIC)^\delta - k_{min} / k_{max}}$$

- k_{net} = Net bacteria growth rate
 - k_{max} = Bacteria growth rate in absence of antibiotic (set by model)
 - k_{min} = Minimum growth rate imposed by antibiotic ($< 0 \rightarrow$ bacteria death)
 - C_u = Free antibiotic concentration
 - MIC = Minimum inhibitory concentration (set in Infection Action)
 - δ = Shape parameter
- Noteworthy Features
 - $C_u = MIC \rightarrow k_{net} = 0$
 - Slow bacteria growth when $C_u < MIC$
 - Bacteria death when $C_u > MIC$
 - Users set k_{min} rather than an EC_{50}
 - This seems to be a more intuitive input

Demo: Action Initiation

- Infection Demonstration

- Progression of three levels of infection (“Mild”, “Moderate”, “Severe”)
- Antibiotic administration in a severely infected case
- Fluid and pressor administration in a septic case
- Setup:
 - Initiation requires a severity, bacteria MIC, and location (for future implementations when localized interactions are better modeled)

```
<Action xsi:type="InfectionData" Severity="Severe">
  <MinimumInhibitoryConcentration value="8.0" unit="mg/L"/>
  <Location>Skin</Location>
</Action>
```

- Note: Infection scenarios are *very* long.
 - BioGears library includes infection simulations saved at various points throughout simulation
 - These saved states can be loaded into new scenarios
 - Useful for cases in which latter stages of infection are main interest of simulation (such as two treatment simulations to be shown)

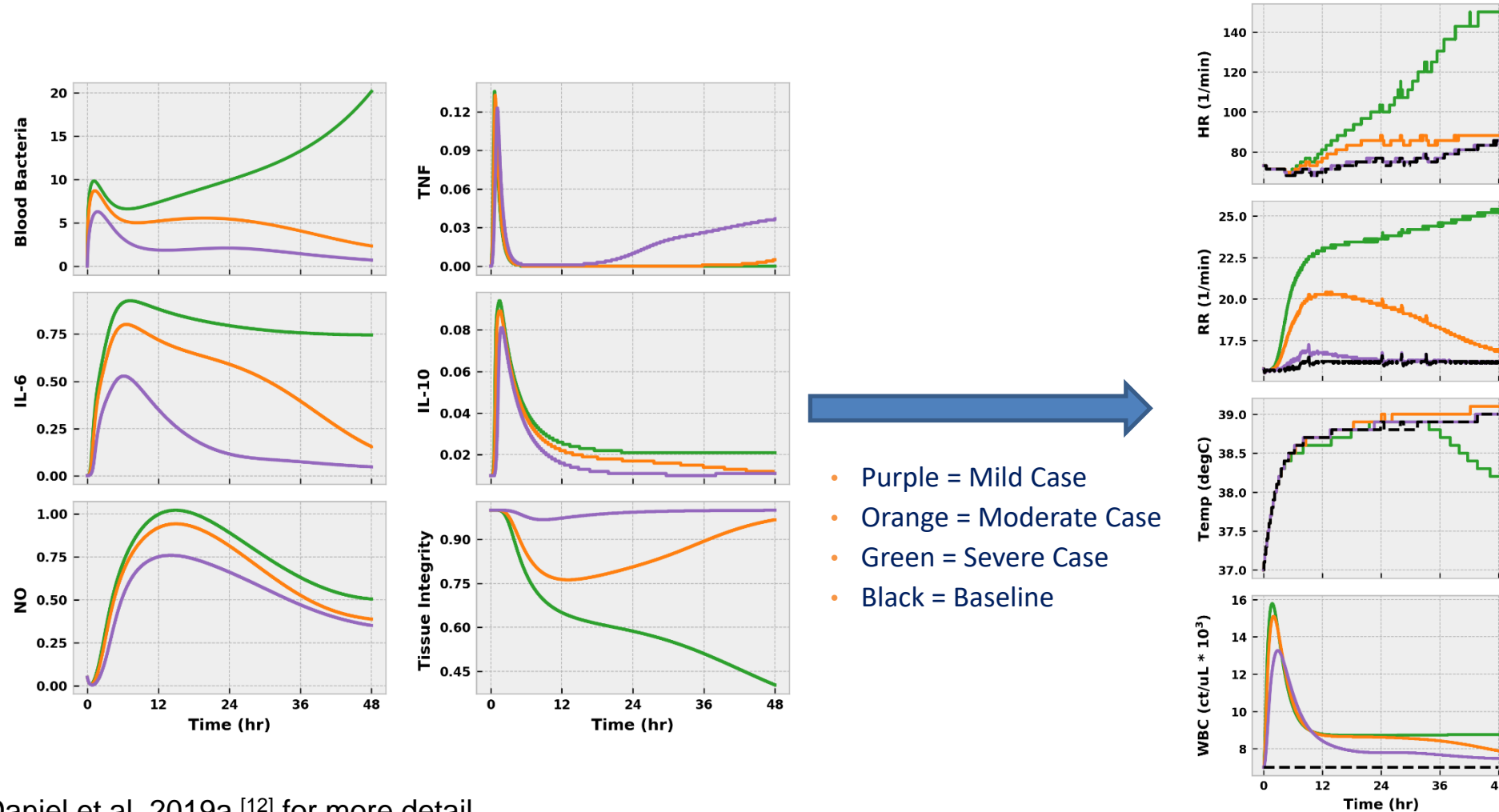
- Burn Wound Demonstration

- Progression of three burn wounds: 10%, 25%, and 40% TBSA
- Fluid resuscitation and pain management in the 25% TBSA case
- Setup:

```
<Action xsi:type="BurnWoundData">
  <TotalBodySurfaceArea value="0.25"/>
</Action>
```


Output: Infections of varying severities

Prolonged and excessive inflammation leads to symptoms of systemic inflammatory response syndrome (SIRS)*

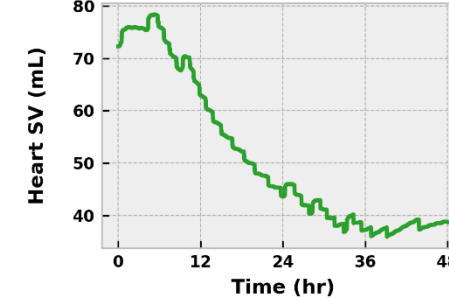
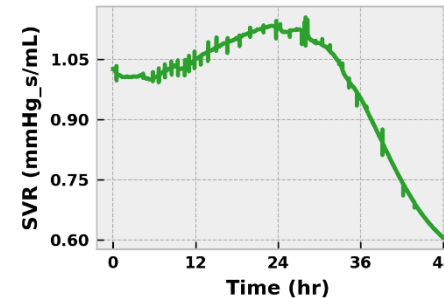
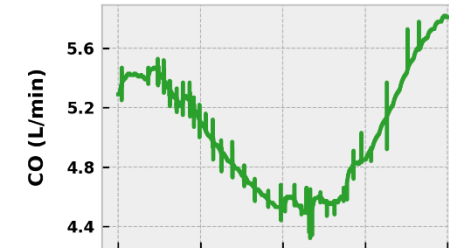
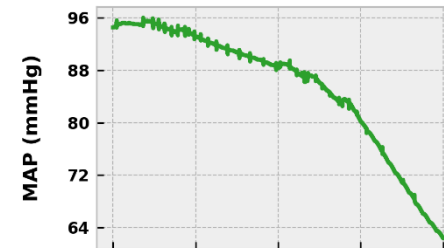
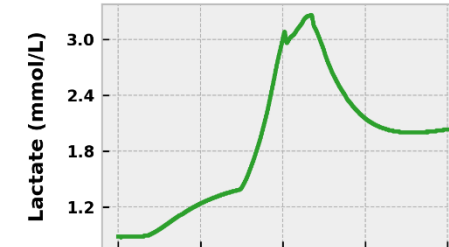
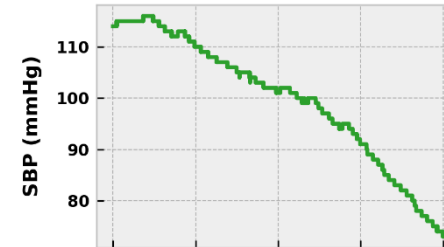
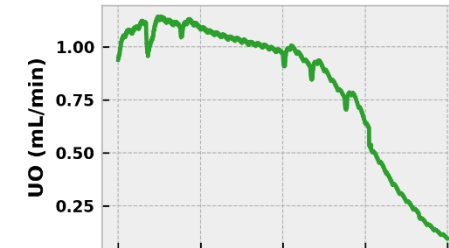
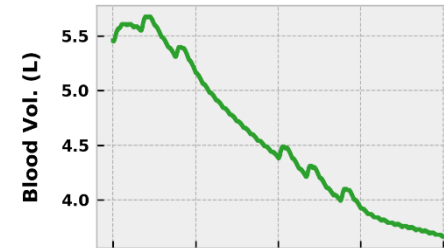


* See McDaniel et al. 2019a [12] for more detail

Output: Septic Infection

Clinically-relevant markers for sepsis and septic shock diagnosis according to Sepsis-3^[11] definitions*

- Sepsis
 - Systolic blood pressure < 100 mmHg
 - Respiration rate > 22 (previous slide)
 - Renal organ failure assessment score
 - UO < 500 mL/day (0.34 mL/min) = 3
 - UO < 200 mL/day (0.14 mL/min) = 4
- Septic Shock
 - MAP < 65 mmHg
 - Serum lactate > 2.0 mmol/L

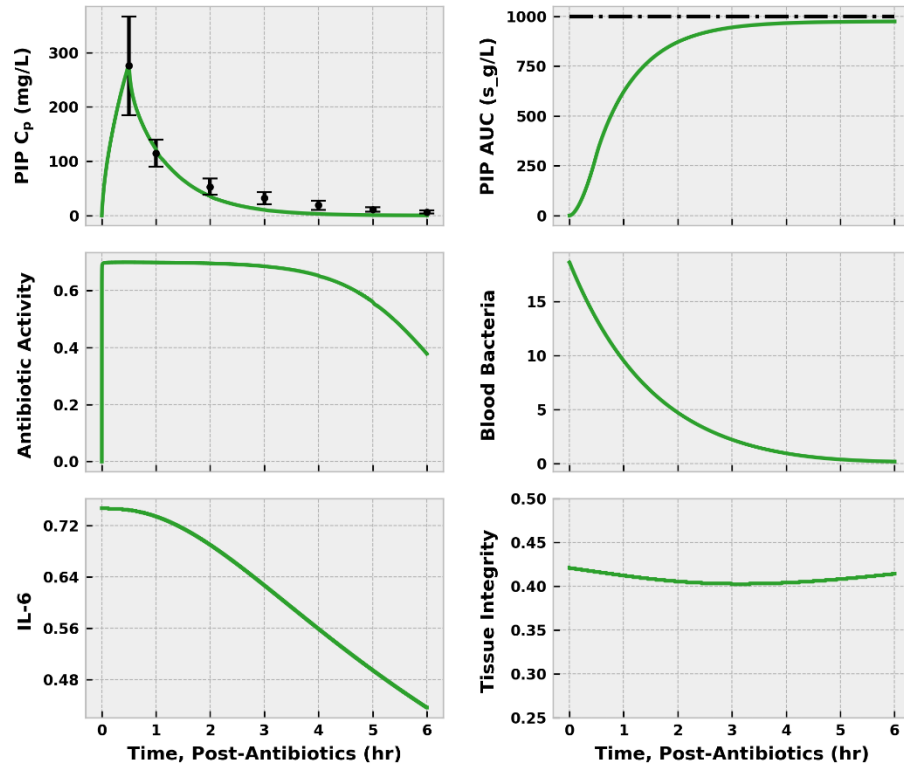


- SBP = Systolic blood pressure
- MAP = Mean arterial pressure
- SVR = Systemic vascular resistance
- UO = Urine output
- CO = Cardiac output
- SV = Stroke volume

* See McDaniel et al. 2019a^[12] for more detail

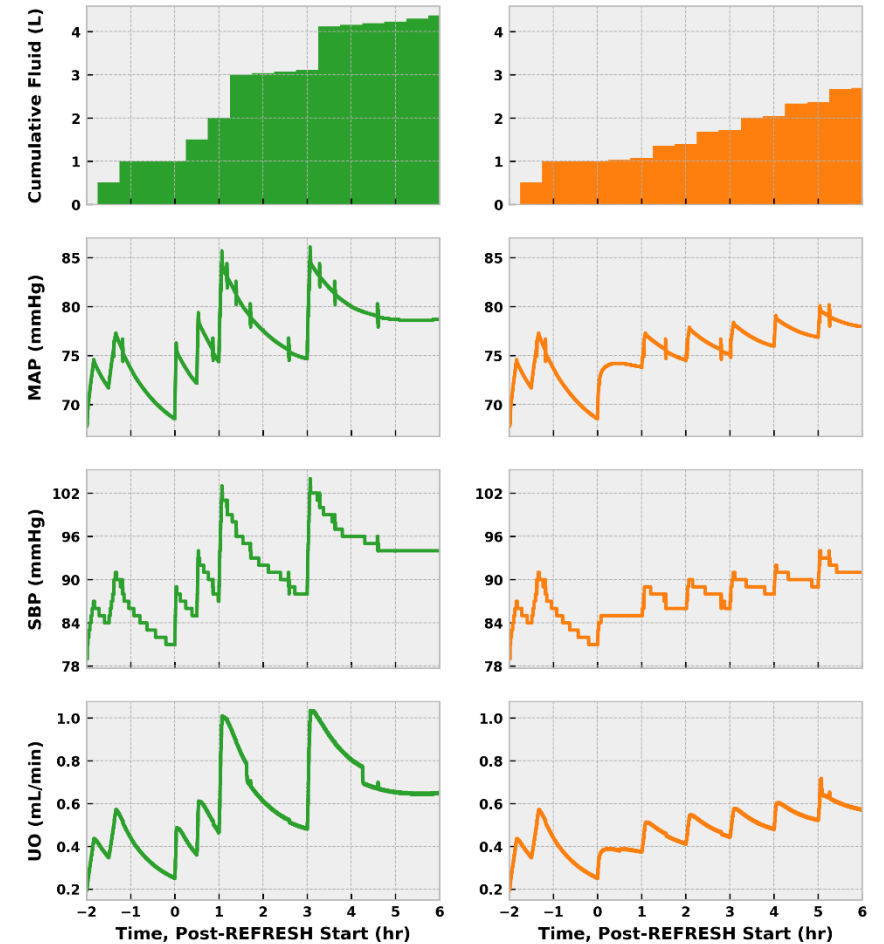
Output: Sepsis Treatment

Model demonstrates distinct outcomes depending on treatment strategy*



Administration of piperacillin/tazobactam. Plasma concentration (C_p) and area under curve (AUC) compared to values in literature^[14]. The antibiotic reduces the bacteria population, leading to reduction in inflammatory IL-6 population and improvement in tissue integrity.

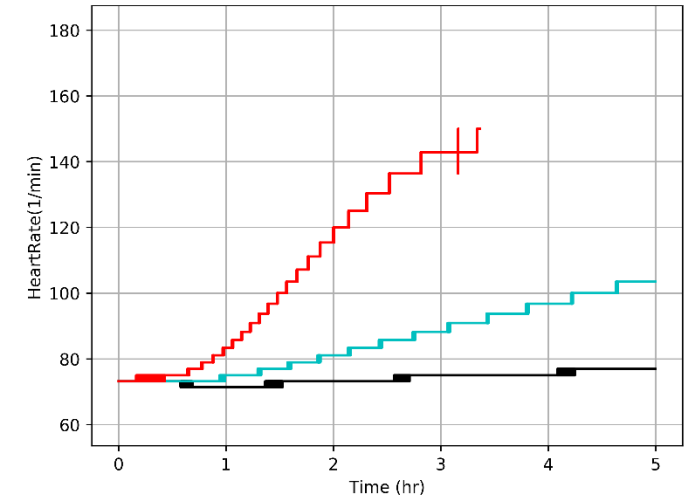
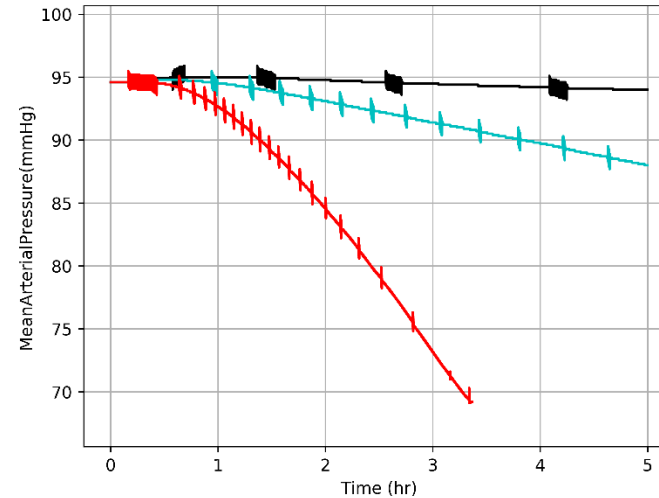
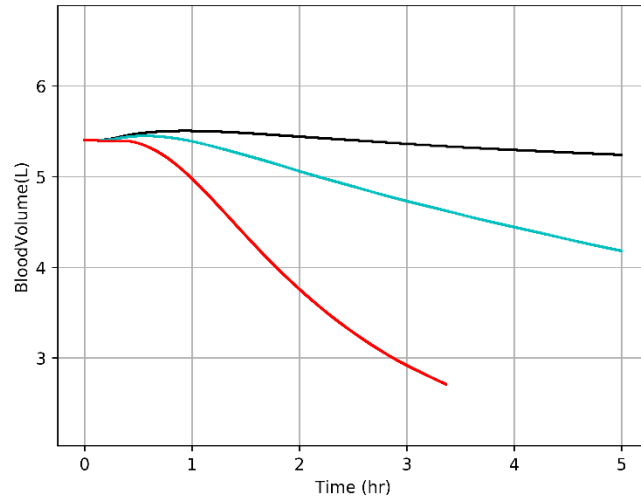
* See McDaniel et al. 2019a^[12] for more detail



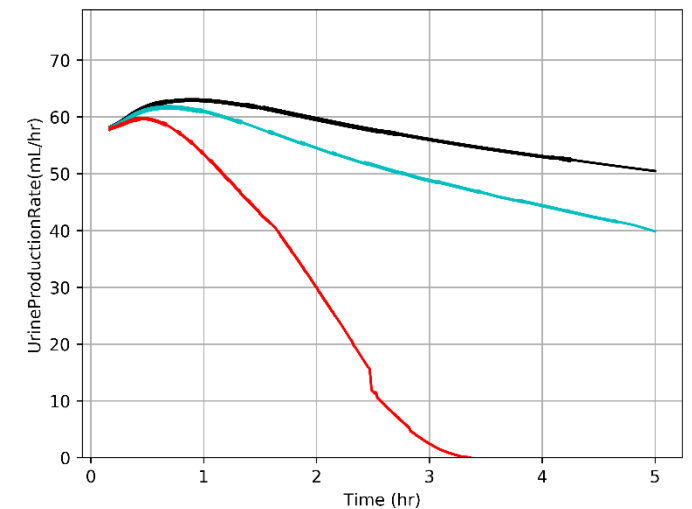
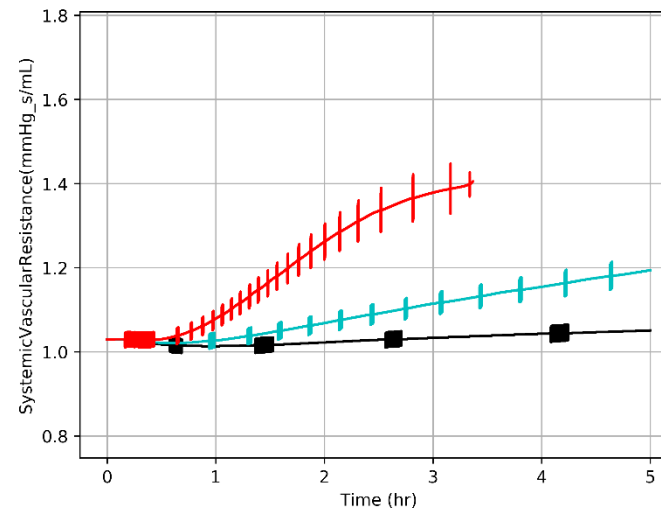
Comparing two treatment strategies described in MacDonald^[15-16]. Left: Early goal directed therapy. Right: Early pressors, reduced fluids

Output: Burns of Varying Severity

Large burns cause massive fluid shift that can lead to death*



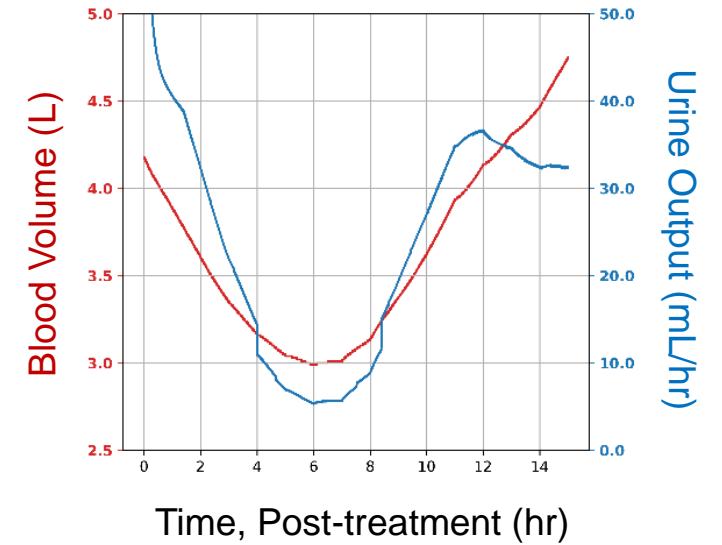
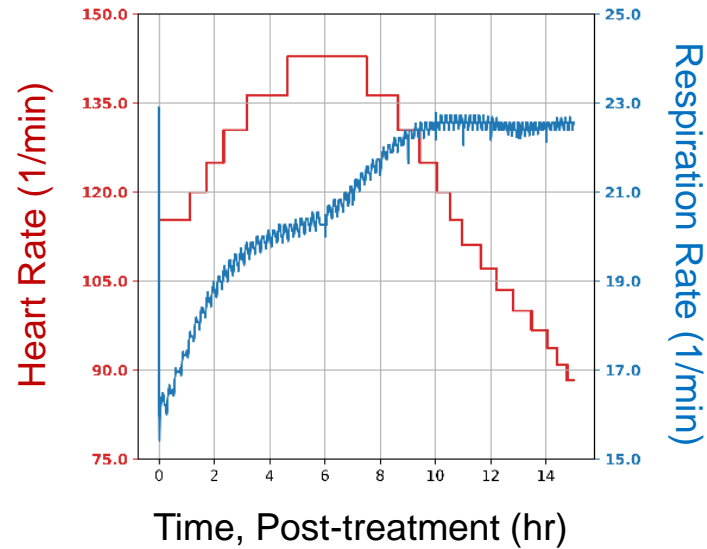
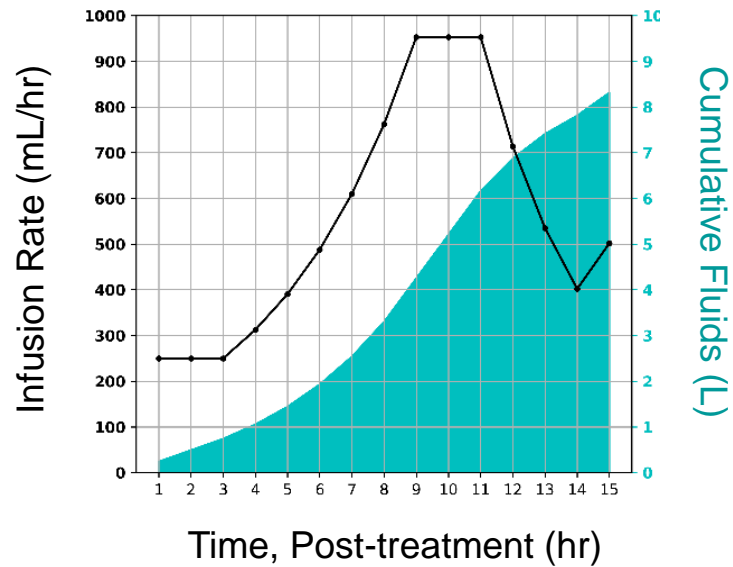
- Black = 10% TBSA (fraction of total body surface area)
- Blue = 25% TBSA
- Red = 40% TBSA



* See McDaniel et al. 2019b^[17] for more detail

Output: Burn Treatment

Burn size and volume resuscitation status (as indicated by UO) drive treatment*



A 25% TBSA burn treated with ketamine infusion (for pain) and administration of ringer's lactate solution (250 mL/hr initial infusion rate). Rate of fluids adjusted each hour according to resuscitation status. Treatment strategy derived from [18-20].

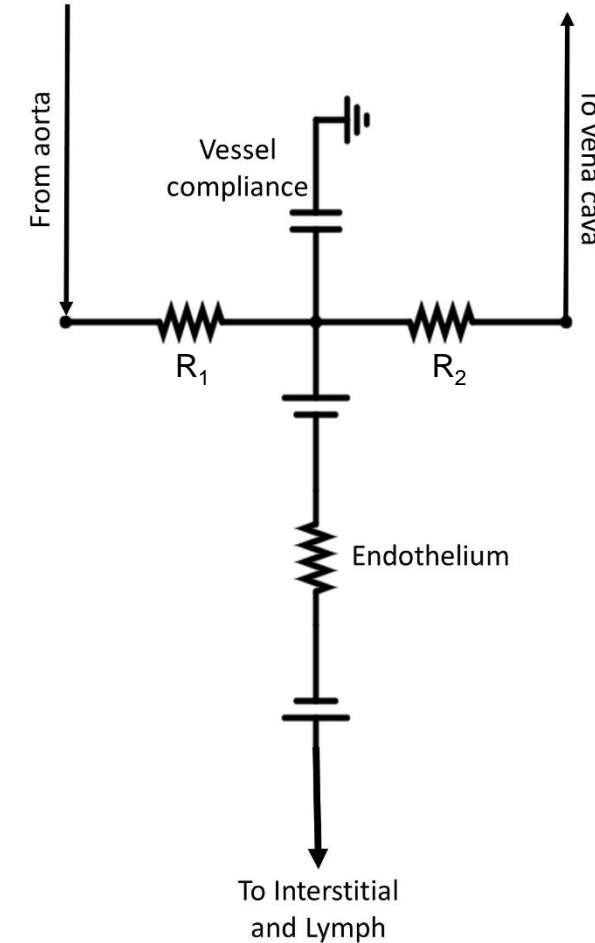
* See McDaniel et al. 2019b^[17] for more detail

Inflammation: Future Work

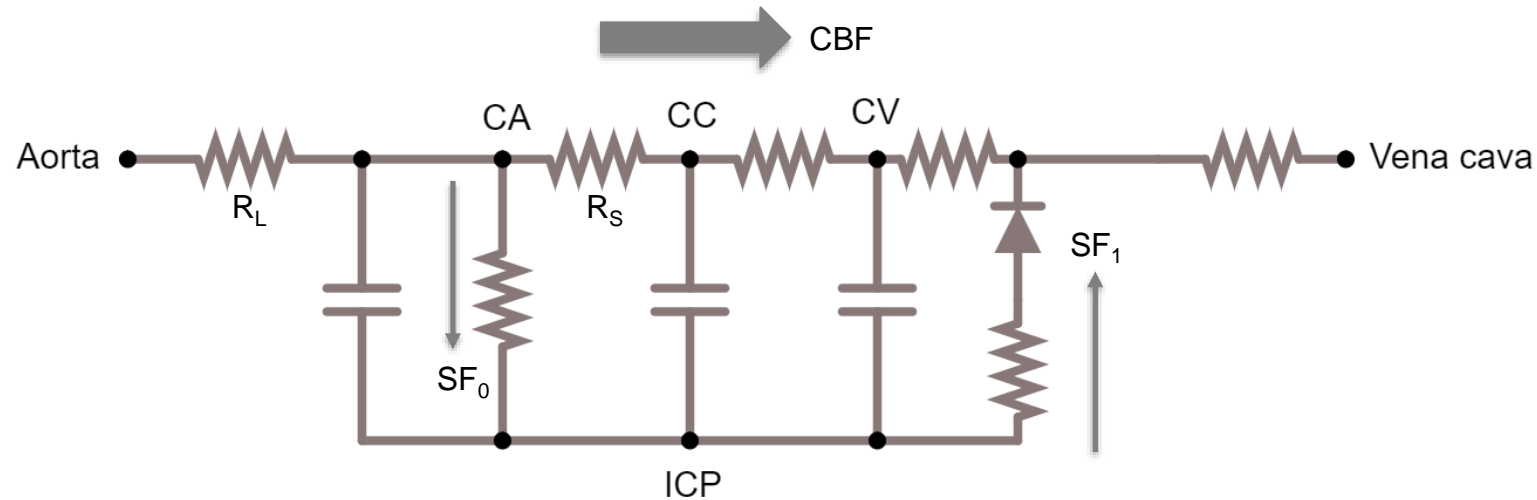
- Model variability
 - Patient inflammatory response parameters (e.g. up-regulation of IL-6 by TNF)
 - Bacteria characteristics (e.g. growth rate)
 - Initial infection levels (expand from “Mild”, “Moderate”, “Severe”)
- Model of acute lung injury (ALI) and acute respiratory distress syndrome (ARDS)
- Tune fluid loss during burn
- Coagulopathy
- Define inflammatory mediators as BioGears Substance types
 - Improved spatial resolution
 - Drug-pathogen interactions at compartment-level
- Link hemorrhage model to inflammation cascade

Cerebral Model Expansion

- Background
 - Cerebral circuit previously implemented a three-element *windkessel* model
 - Additional pathway for interstitial space
 - Traumatic Brain Injury action increased resistance on pre-capillary (R_1) and post-capillary (R_2) paths
 - No autoregulation
- Goals
 - Increase fidelity of circuit for more accurate blood and oxygen tracking
 - Introduce cerebral regulation to prioritize oxygen delivery to brain



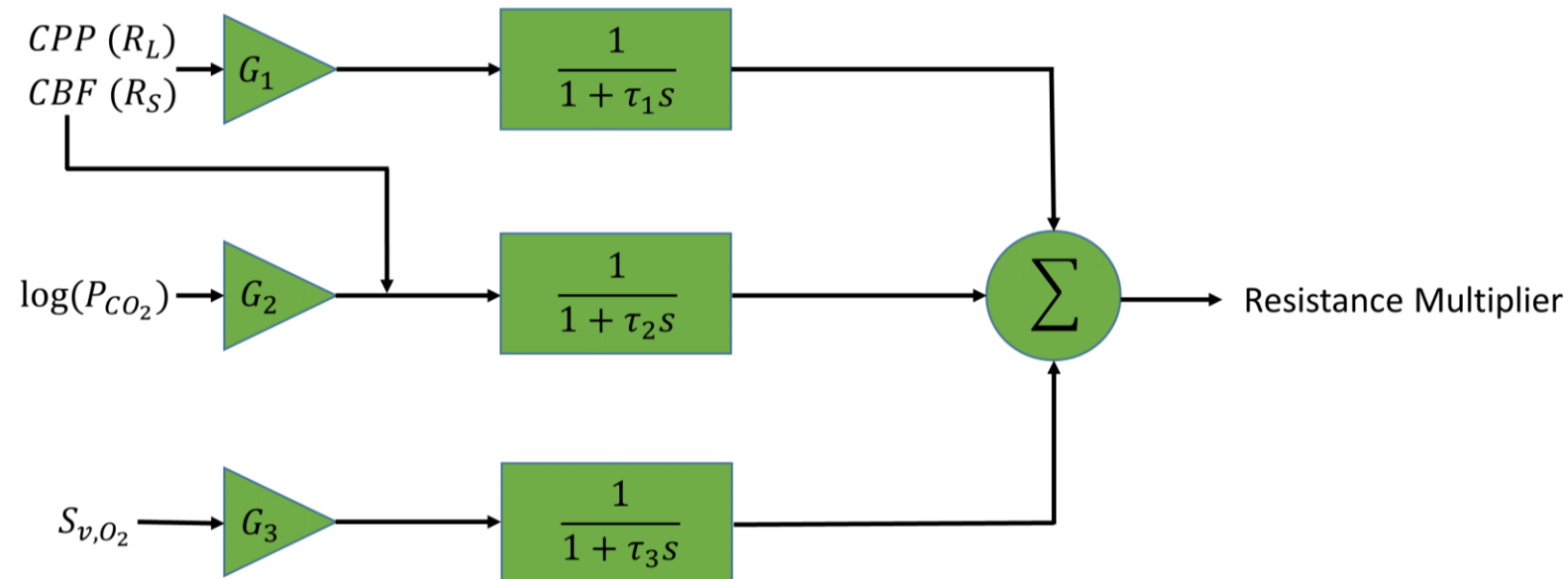
Updated Cerebral Circuit



- Discretize brain sub-circuit to include:
 - Cerebral arteries (CA)
 - Cerebral capillaries (CP)
 - Cerebral veins (CV)
- The volume of each region responds to pressure changes (compliance paths)
- Intracranial pressure (ICP) determined by fluid stored in brain
- Cerebral blood flow (CBF) calculated according to total pressure drop and resistance across sub-circuit
- Large (R_L) and small artery (R_S) resistances are subject to auto-regulatory control (next slide)
- Some fluid flows into interstitial space as spinal fluid (SF_0) and returns to vasculature (SF_1)
- Circuit design and baseline parameters adapted from [21]

Cerebral Autoregulation

- Model assumptions^[22-23]
 - R_L subject to first-order control via CPP
 - R_S subject to first-order control via CBF
 - R_L and R_S subject to first-order control via $\log(P_{CO_2}) \rightarrow$ Reflects pH dependence
 - The effects of CO_2 are attenuated by low CBF
 - R_L and R_S subject to first-order control via venous cerebral O_2 saturation
- Model outputs
 - R_L resistance multiplier: Bounds = [0.8, 1.2]
 - R_S resistance multiplier: Bounds = [0.75, 1.25]

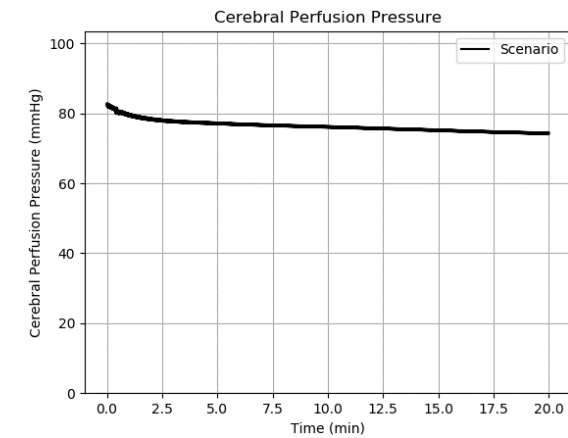
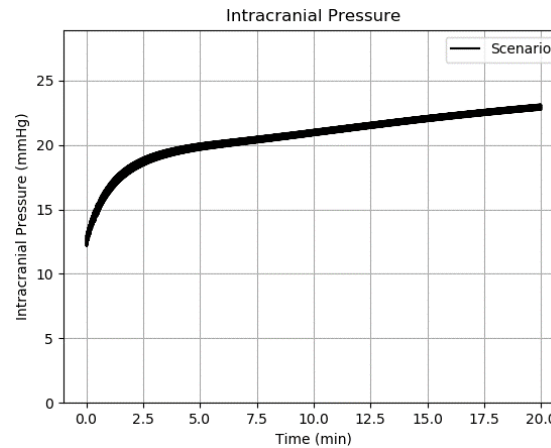
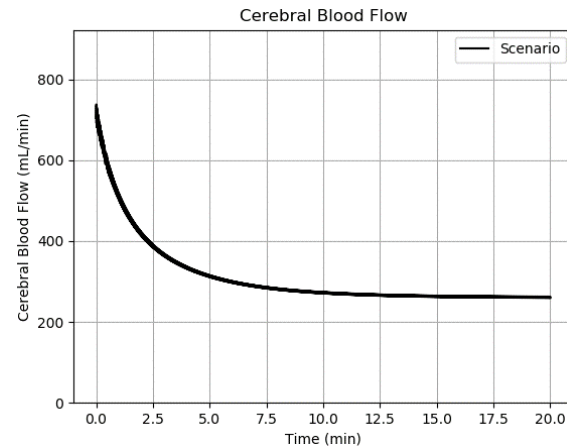


*Gains (G) and time constants (τ) obtained from [23]

Cerebral Model: Example Results

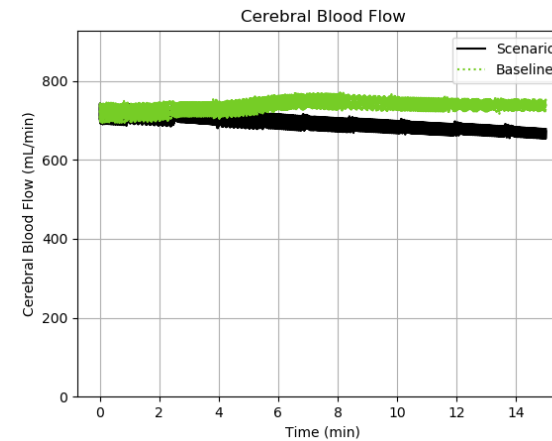
- Traumatic Brain Injury

- Severity Input = 0.75
- Results follow trends reported in [24]



- Effect of autoregulation

- Hemorrhage (initial bleeding rate = 100 mL/min) for 15 minutes
- Green = Cerebral autoregulation ON
- Black = Cerebral autoregulation OFF

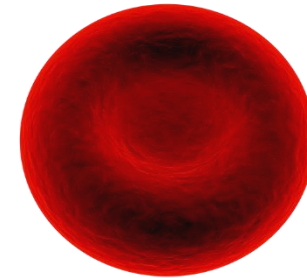


Cerebral Model: Future Work

- No short-term work planned
- Long-term
 - Improve blood-gas tracking in brain to increase fidelity of feedback model
 - Possible integration with ARA TBI project

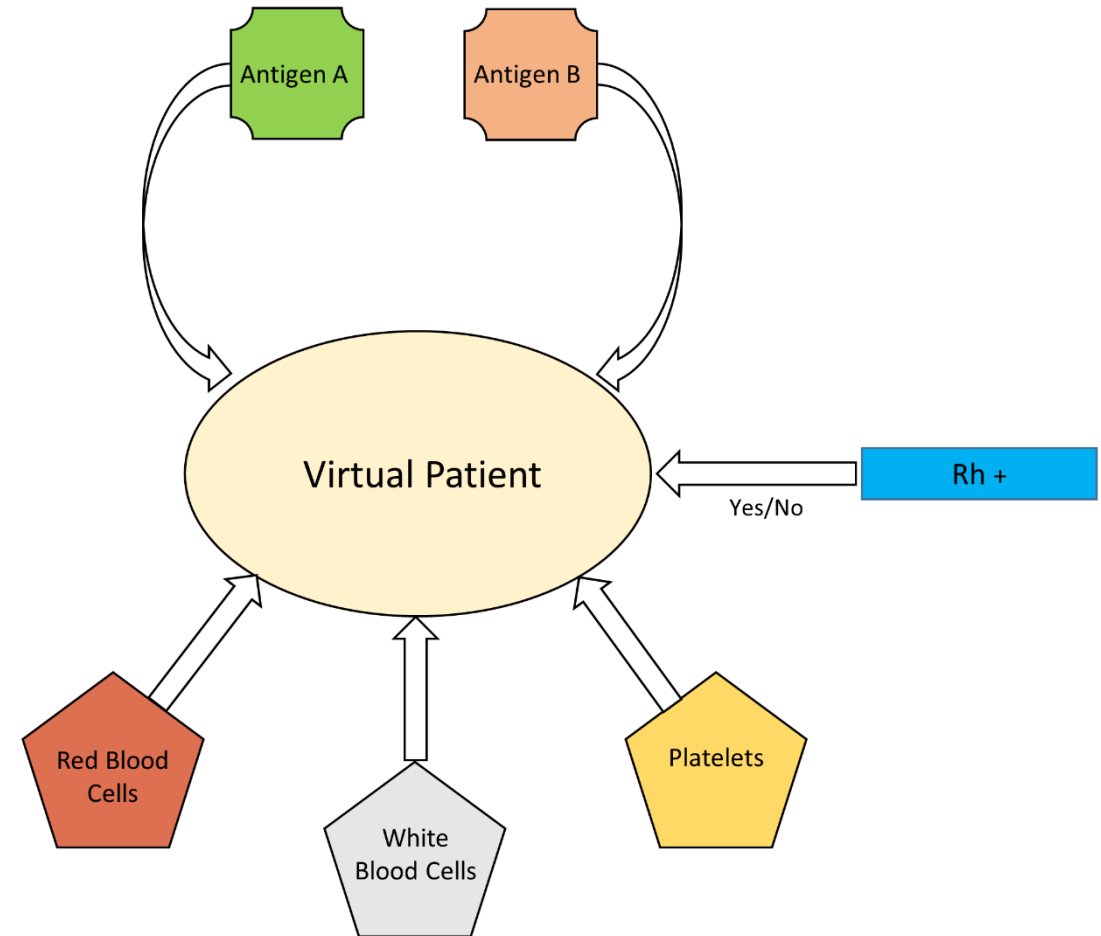
Whole Blood

- Background
 - Previous model
 - One substance compound: Blood
 - Constituent components
 - Hemoglobin
 - Oxyhemoglobin
 - Sodium
 - Albumin
 - Urea
 - Glucose
 - Triacylglycerols
 - Calcium
 - Creatinine
 - New model goals
 - Support ABO blood type definition for virtual patients
 - Support Rh factor definition for virtual patients
 - Distinguish between whole blood and plasma
 - Support transfusion action that checks for blood acceptor – donor compatibility



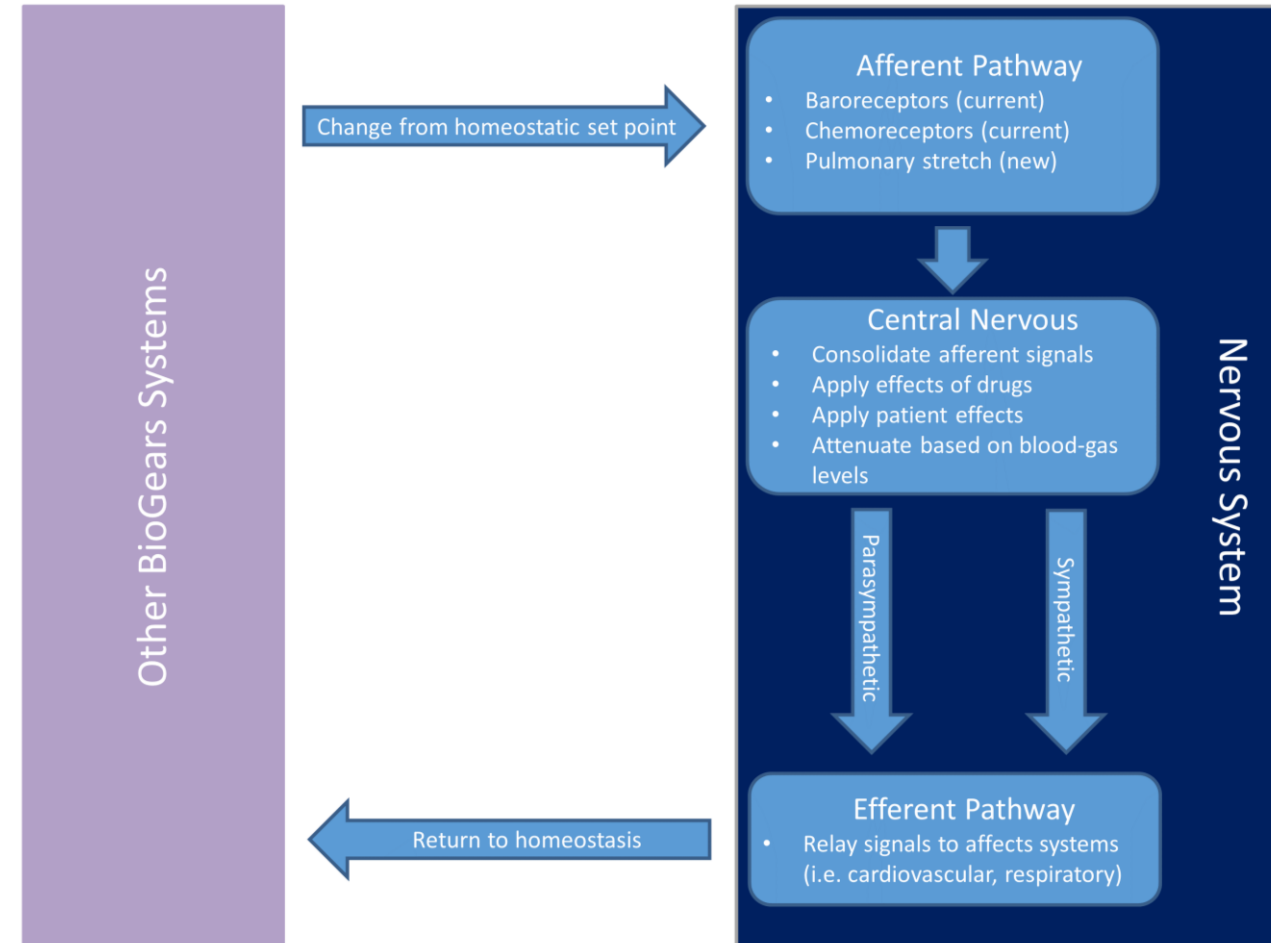
Whole Blood

- New Substance Definitions
 - Antigen A, B
 - Red Blood Cells, White Blood Cells, Platelets
 - Concentrations set according to standard literature values^[Boron]
- Rh Classification (Boolean)
 - Set to True (+) or False (-)
- No antibodies
 - Inferred by presence/absence of antigens
- Transfusion Action
 - Covered in detail in Multi-Trauma Presentation



Models in Development

- Advanced Stages: Integrated Nervous Model
 - Current
 - Feedback systems treated as separate entities
 - Response of one system does not inform the others
 - Proposed
 - Increase fidelity of existing baroreceptor and chemoreceptor models
 - Introduce new feedback mechanisms
 - Process all signals simultaneously



Models in Development

- Preliminary Stages—Long term goals
 - Updated Nutrient Kinetics model
 - Updated Insulin / Glucose regulation model

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