



BioGears Visualizer

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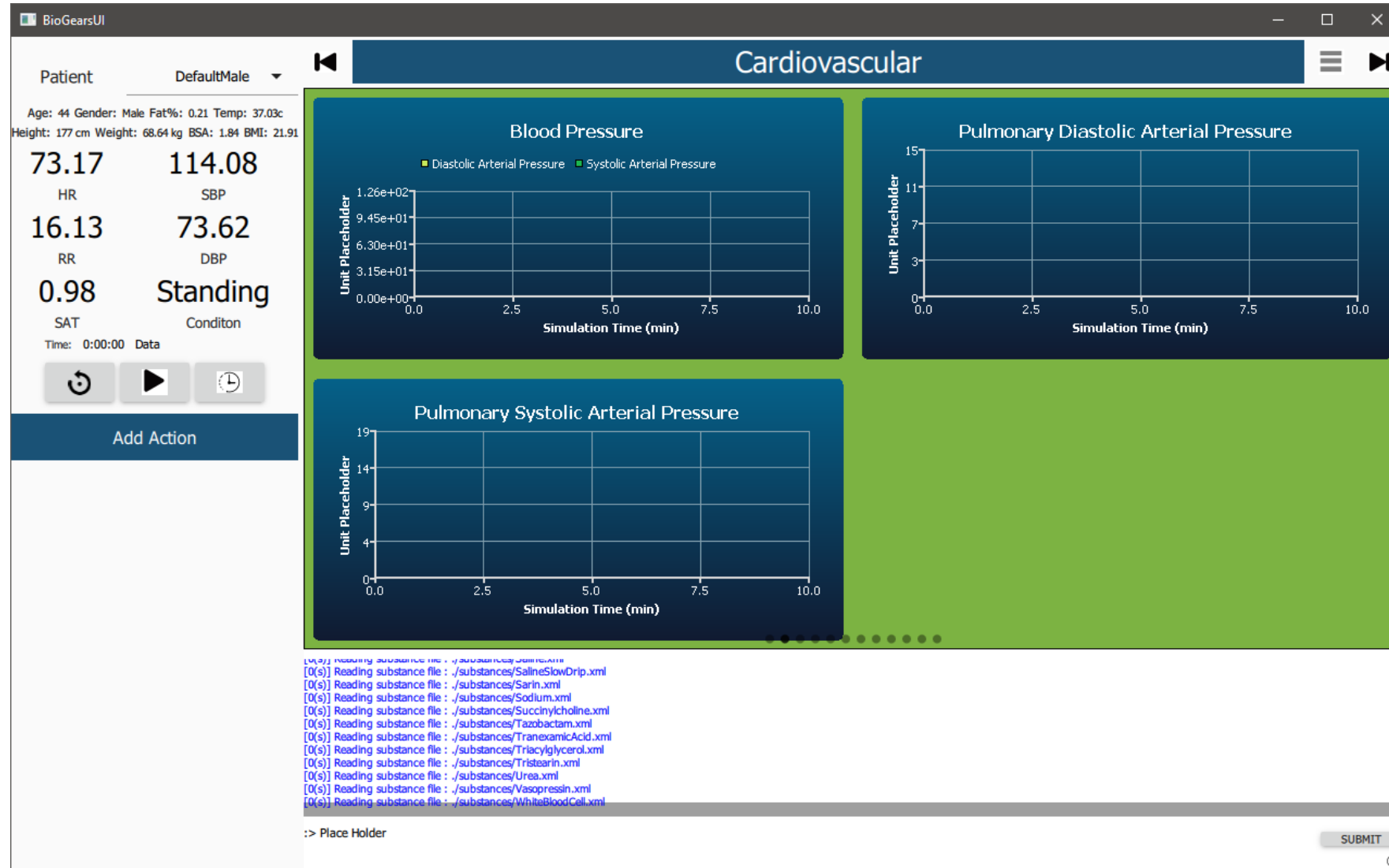
Streamlined UI for experimenting with Insults/Injuries on a patient.

- Timeline:
 - Now: 0.5 – Alpha Release
 - Custom Actions
 - Simulation Controls
 - Organized Data Visualization
 - December 2020: Version 1.0
 - Improved Theme
 - Curated Physiology Widgets
 - Patient State Creation
 - Scenario Timeline
 - Version 2.0:
 - Multi Window UI
 - LUA Based Scripting Engine
 - Parallel Simulations

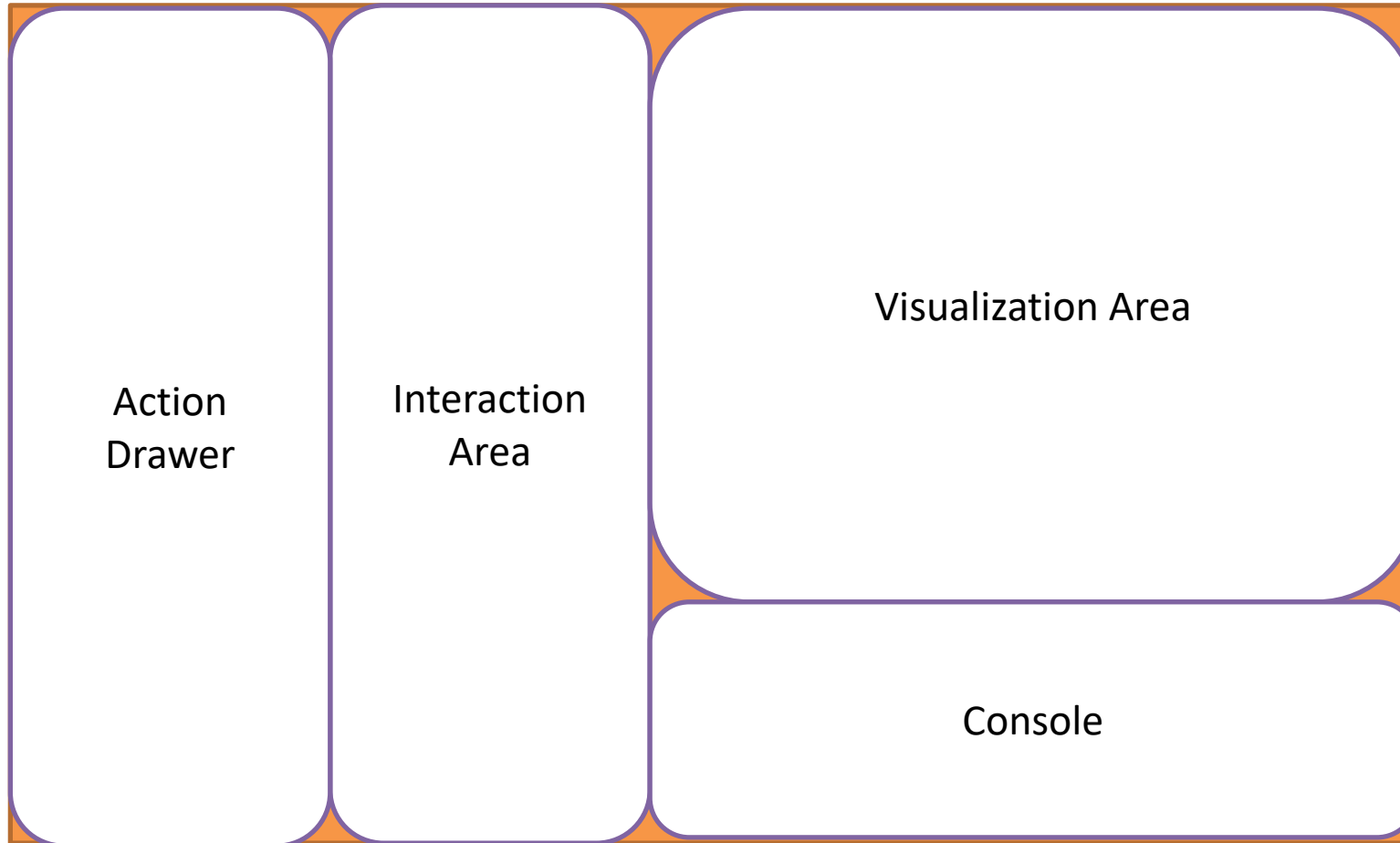


Alpha Release Walkthrough

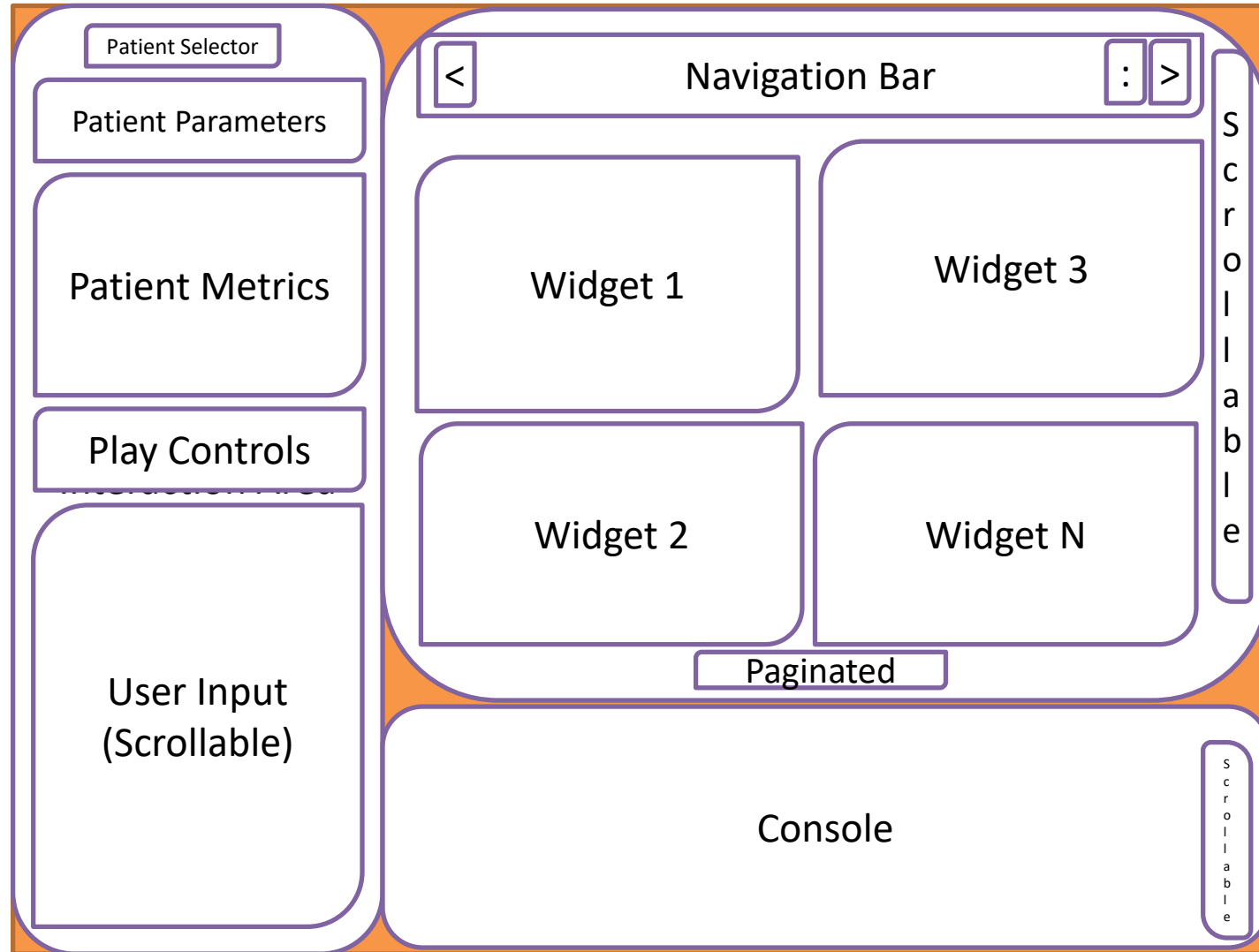
Current UI



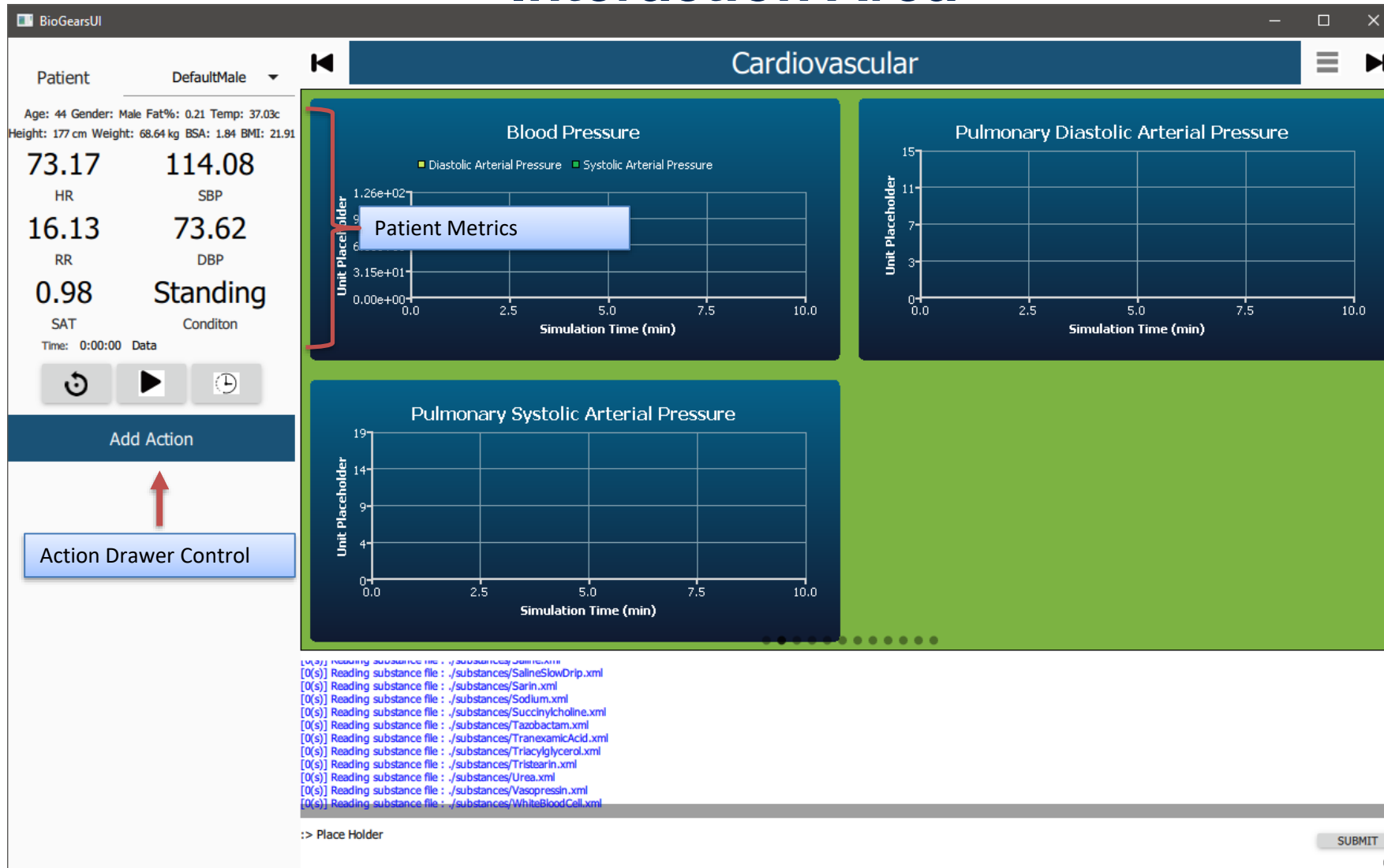
Current UI



UI Breakdown

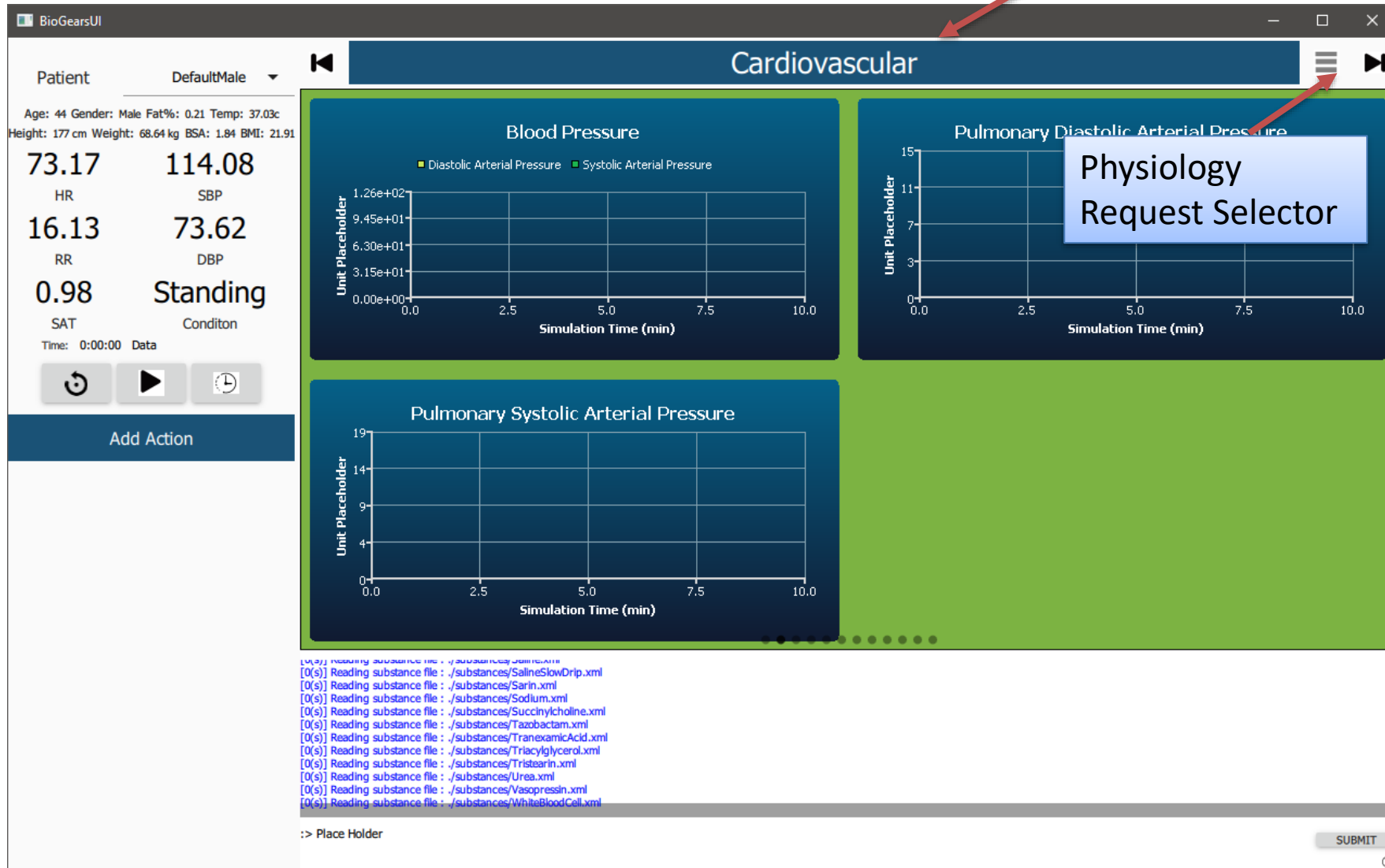


Interaction Area



Visualization Area

Current Physiology Category



Physiology Request Selection

Patient

DefaultMale

Age: 44 Gender: Male Fat%: 0.21 Temp: 37.03c

Height: 177 cm Weight: 68.64 kg BSA: 1.84 BMI: 21.91

73.17

HR

114.08

SBP

16.13

RR

73.62

DBP

0.98

SAT

Standing

Conditon

Time: 0:00:00 Data

↺

▶

🕒

Add Action

Blood Chemistry

Arterial Blood PH

Oxygen Saturation

Red Blood Cell Count

☒ arterialBloodPH
 ☐ arterialBloodPHBaseline
 ☐ bloodDensity
 ☐ bloodSpecificHeat
 ☐ bloodUreaNitrogenConcentration
 ☐ carbonDioxideSaturation
 ☐ carbonMonoxideSaturation
 ☐ hematocrit
 ☐ hemoglobinContent
 ☒ oxygenSaturation

--> Results Here

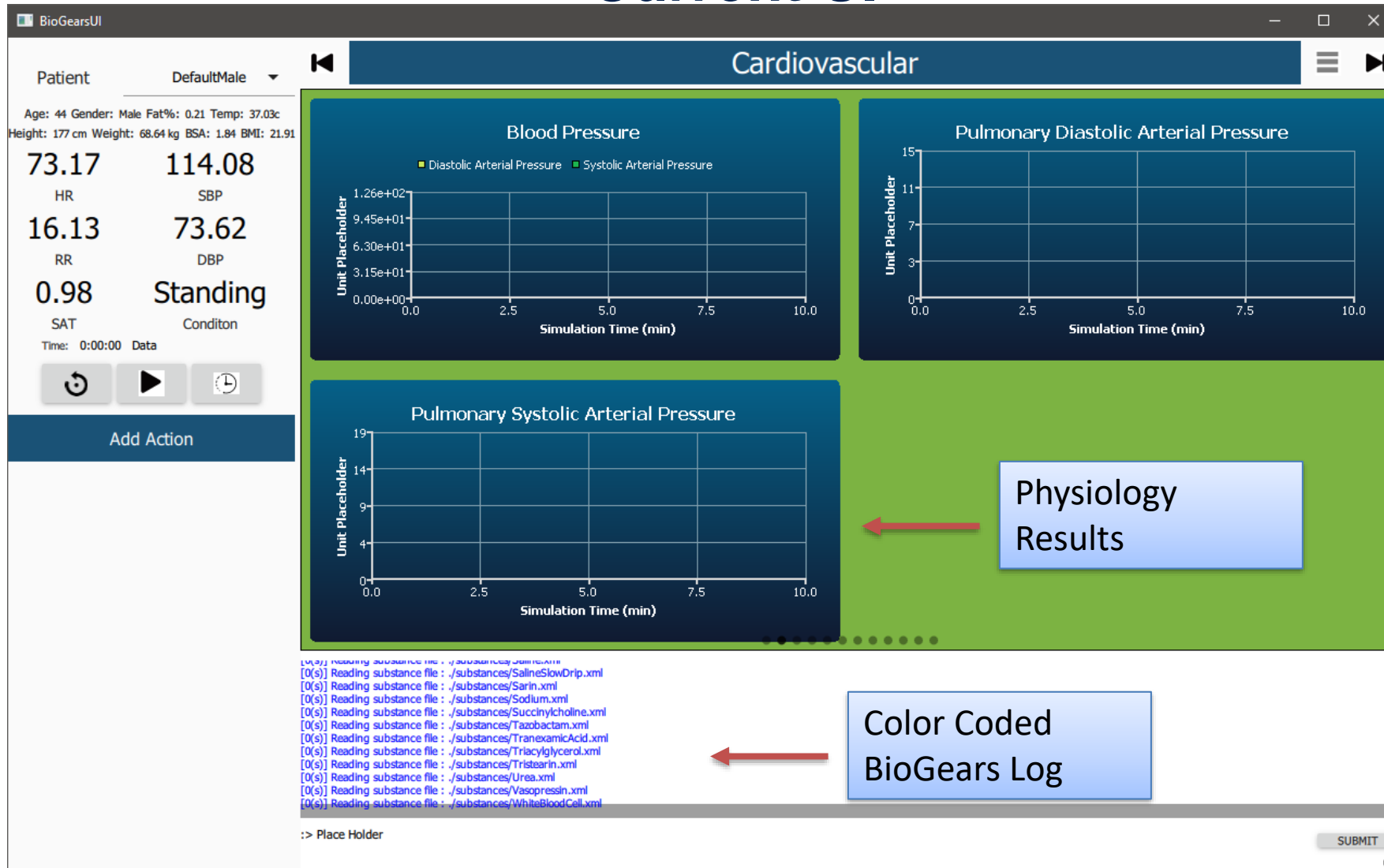
[0(s)] Reading substance file : ./substances/Acetoacetate.xml
[0(s)] Reading substance file : ./substances/Albumin.xml
[0(s)] Reading substance file : ./substances/Albuterol.xml
[0(s)] Reading substance file : ./substances/AminoAcids.xml
[0(s)] Reading substance file : ./substances/Antigen_A.xml
[0(s)] Reading substance file : ./substances/Antigen_B.xml
[0(s)] Reading substance file : ./substances/Bicarbonate.xml
[0(s)] Reading substance file : ./substances/Blood_ABNegative.xml
[0(s)] Reading substance file : ./substances/Blood_ABPositive.xml
[0(s)] Reading substance file : ./substances/Blood_ANegative.xml

::: Place Holder

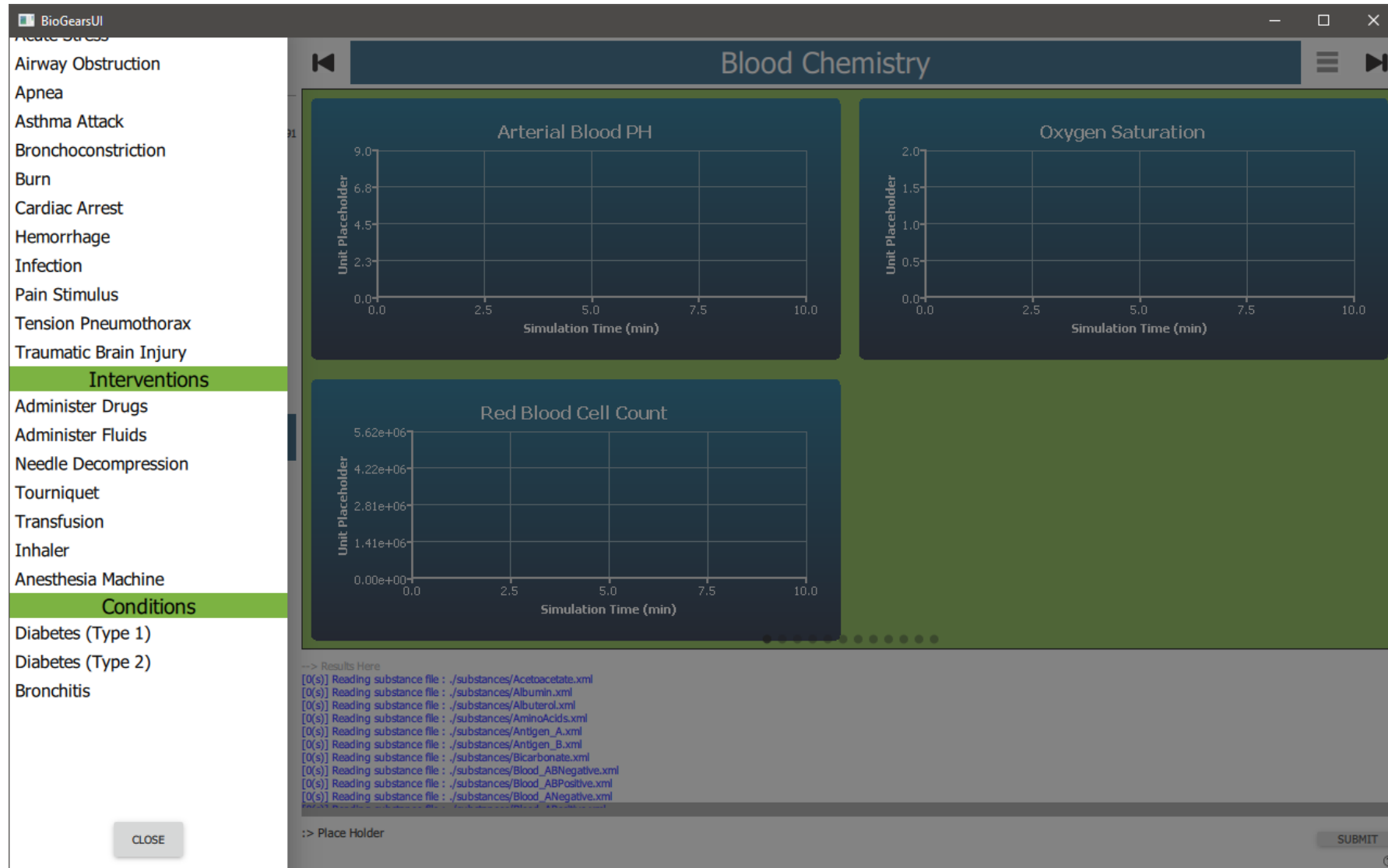
SUBMIT

Data Request Selector

Current UI



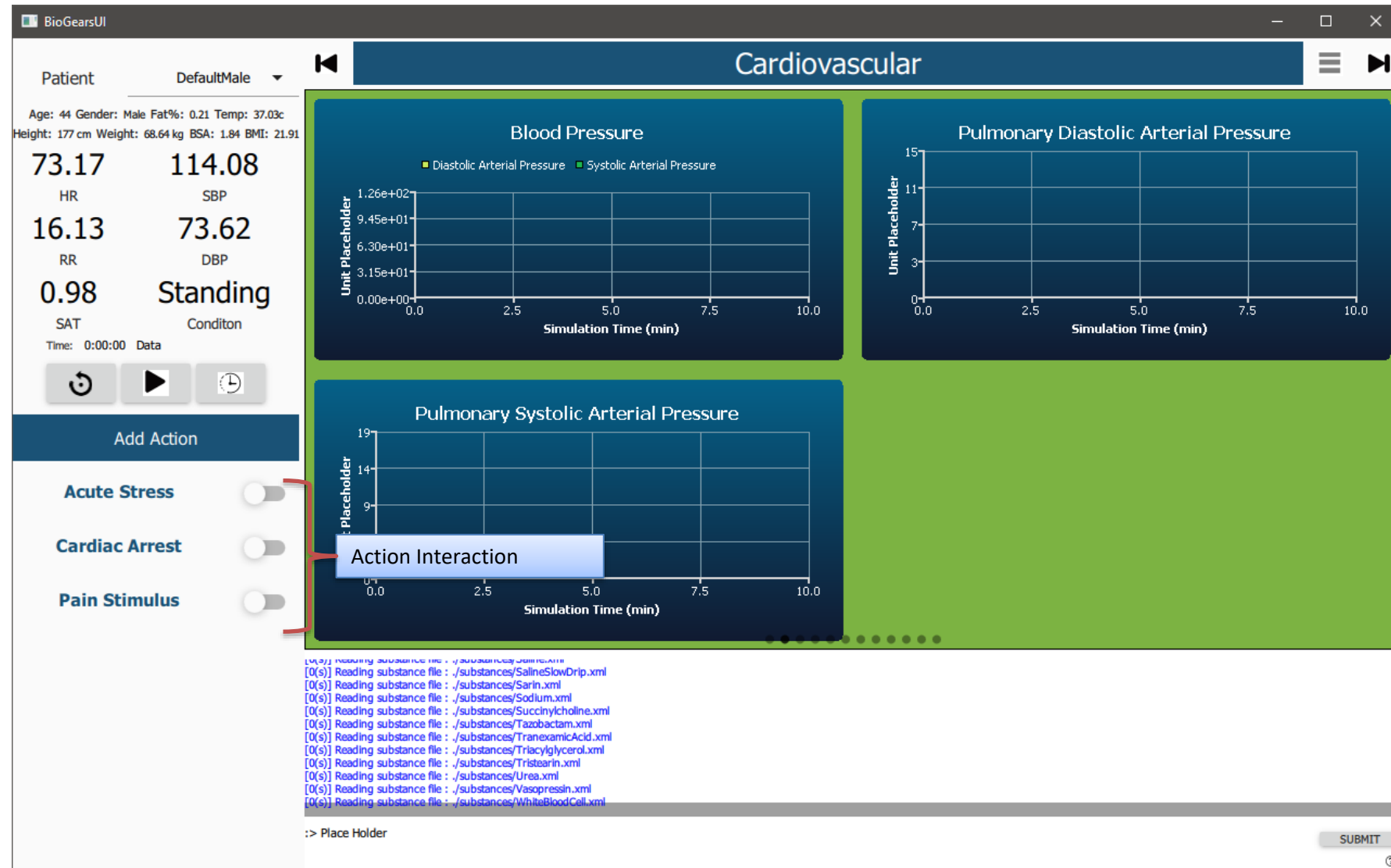
Action Selection



Action Creation

The screenshot displays the BioGearsUI Administer Fluids Editor. A dropdown menu is open, showing the following options: PiperacillinTazobactam, Plasma, RingersLactate, Saline, and SalineSlowDrip. The 'Plasma' option is currently selected. The background interface includes a 'Blood Chemistry' section with graphs for 'Arterial Blood PH' and 'Red Blood Cell Count', and a list of physiological parameters on the right, such as 'arterialBloodPH', 'arterialBloodPHBaseline', 'bloodDensity', 'bloodSpecificHeat', 'bloodUreaNitrogenConcentration', 'carbonDioxideSaturation', 'carbonMonoxideSaturation', 'hematocrit', 'hemoglobinContent', and 'oxygenSaturation'. The 'oxygenSaturation' parameter is checked. At the bottom, there is a 'Results Here' section with a list of substance files being read, including Acetate, Albumin, Albuterol, AminoAcids, Antigen_A, Antigen_B, Bicarbonate, Blood_ABNegative, Blood_ABPPositive, and Blood_ABNegative. A 'SUBMIT' button is visible in the bottom right corner.

Current UI



Data Request Organization

Vitals

- Blood Pressure
- Respiration Rate
- Heart Rate
- O2 Saturation
- Cardiac Output
- Blood Volume
- Stroke Volume
- Central Venous Pressure
- Tidal Volume

Cardiopulmonary

- Cerebral Perfusion Pressure
- Intracranial Pressure
- Systemic Vascular Resistance
- Pulse Pressure
- IE Ratio
- Total Pulmonary Ventilation
- Lung Volume
- Alveolar Ventilation
- Dead Space Ventilation
- Transpulmonary Pressure

Data Request Organization

Data Request Blood Chemistry

- Blood Gas Concentrations
- Combined CO₂ and O₂ Partial Pressure
- Blood pH
- Hematocrit
- Strong Ion Difference

Energy & Metabolism

- Core Temperature
- Sweat Rate
- Skin Temperature
- Total Metabolic Rate
- Stomach Contents
 - Make this a single plot w/ sub-requests for different nutrients (calcium, sodium, ...)
- Oxygen Consumption Rate
- CO₂ Production Rate

Data Request Organization

Drugs & Substances

- Active drugs
- Substances
- Nutrients

Fluid Levels

- Extravascular fluid volume
- Extracellular fluid volume
- Intracellular fluid volume
- Total body fluid



Visualizer Initial Release

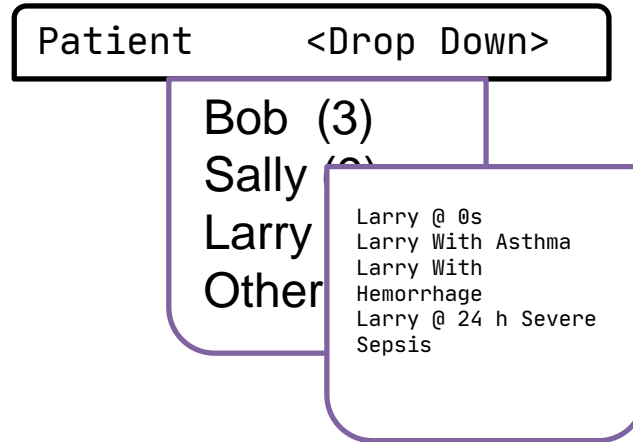
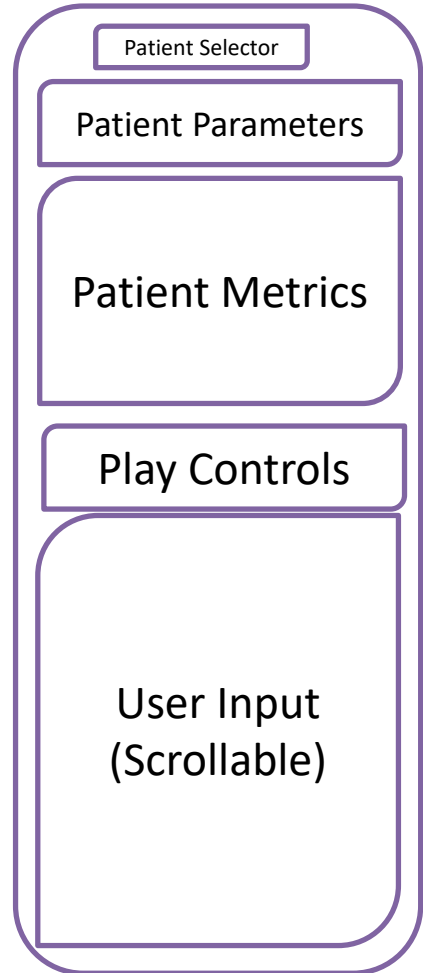
Release December 2020

Goals for initial binary release (v1.0)

- **Improved Interaction**
 - Ability to adjust variable actions in main UI
 - Distinguish between monetary actions and continual actions.
- **Improved Theme**
 - Uniform use of fonts across the UI
 - Common color scheme for metrics (Normal, Warning, Critical...)
 - Animated exercise state
 - Uniform scheme for widgets (Label font, legend position, line style...)
- **Curated Physiology Widgets**
 - Custom visualization to match common clinical interfaces (Patient Vitals, Metabolic Panels...)
- **Data Exports**
 - Patient States
 - Patient Files
 - Scenario Timelines
- **Scenario File Loading**

Interaction Improvements

Nest Patient Selection



- Future function File -> Save
- Decide Name
- Decide Description
- Write File to States
- In Subfolder
states/<patient_name>/

Purpose:

- Select initial conditions
- Switch state and restart

Initialization

- UI Parses all patients in patients/
- UI Parses all states in states/ and maps them to patient names

UX

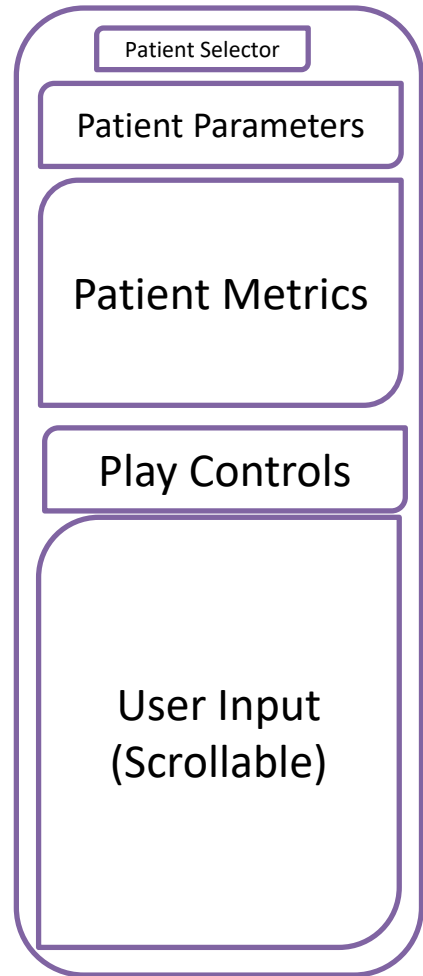
- Select drop down browse to patient
- Cascading menus allow selection of specific states for a common patient
- Mouse over text gives full description string from state file

On Click

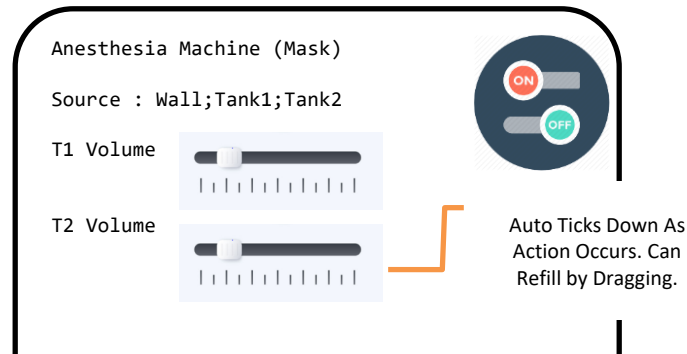
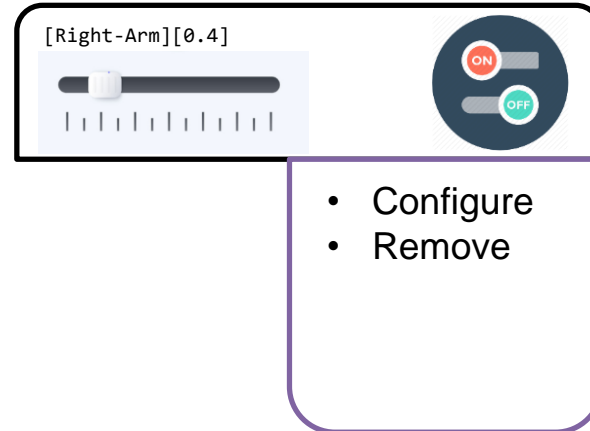
- Stop simulation
- Reset all visualization widgets
- Load state

Interaction Improvements

Customizable Action Values (Where Possible)



Severity Widget



Purpose:

- Each Action has its own custom widget
- User Input is scrollable to allow us to see different actions
- Optimized font size for data density while ensuring readability

UX:

Right Click

- Configure: display original configuration widget and adjust bound parameters.
- Remove: deletes widget from action parameters

Toggle

- On: apply Action with current value
- Off: reverse Action if possible
 - Timed actions can not be toggled off but will show off once they are no longer active

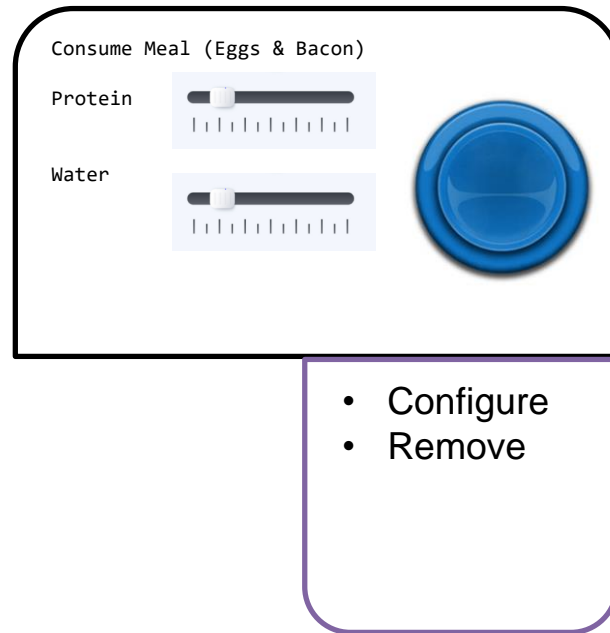
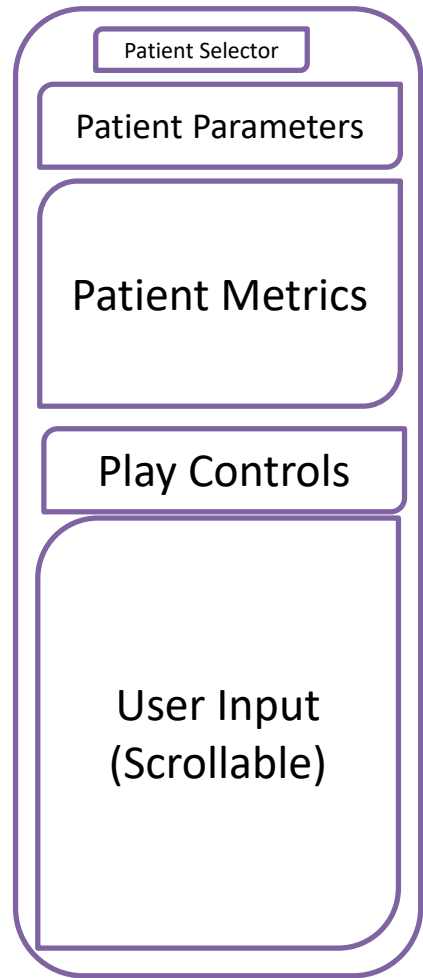
Fields

- Manipulate : changes Action live if active else just stores new values
 - Interaction model is different based on UI configurable actions.

*Some Actions are unique (anesthesia machine, pneumothorax)

Interaction Improvements

Momentary Actions



Purpose:

- Some actions are instantaneous and can not be toggled
 - Ex: Consume Meal

UX:

Right Click

- Configure: display original configuration widget and adjust bound parameters
- Remove: delete widget from action parameters

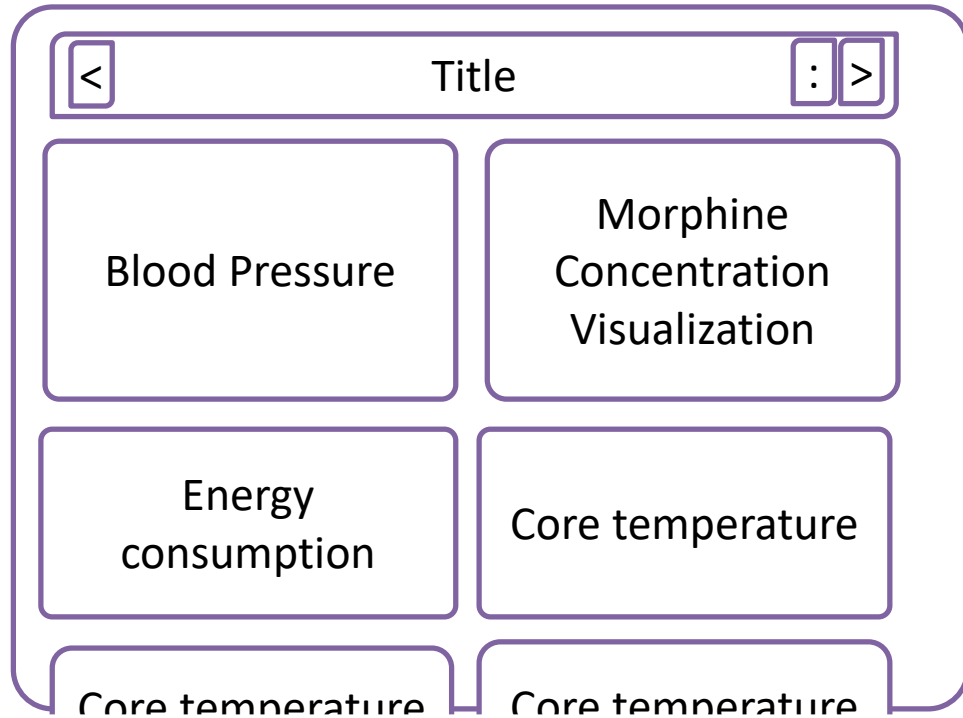
On Click

- Apply Action with current value

Fields

- Manipulate : changes Action live if active else just stores new vales
 - Interaction model is different based on UI configurable actions.

Visualization Improvements



Purpose: Paged Visualization Area of BioGears Metrics

Categories

- Categories Based on Data Request Organization

Controls

- **<** Previous Category (Loops)
- **>** Next Category (Loops)
- **:** Opens Selection Menu of Widgets Available on Current Page

Features:

- Dynamic Graphs
- Custom Visualizations
- Scrollable Area to Allow

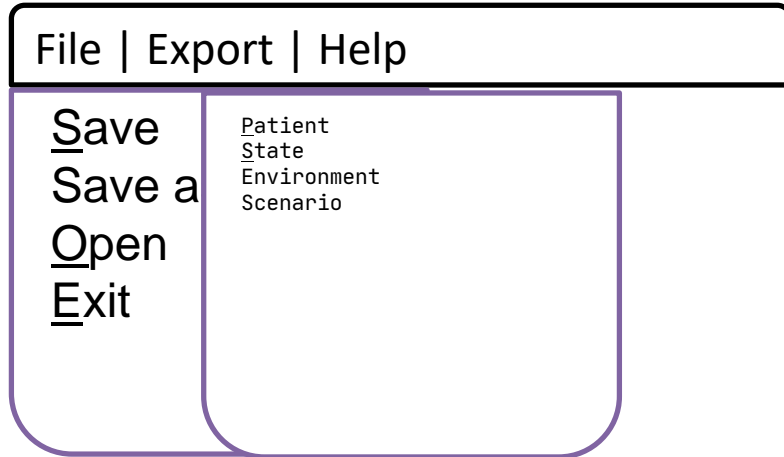
Custom Visualization

Moving forward we would like to produce several non plot-based visualizations.

Some ideas include:

- Patient Vitals
- Substance concentration visualizer (in the body)
- Pulmonary Activity
- Breathing Cycle
- Bladder volume

Data Exports



Purpose:

- Allows users to save work for later use
- Convenient way to create save points for long simulations and repeated demonstrations

UX

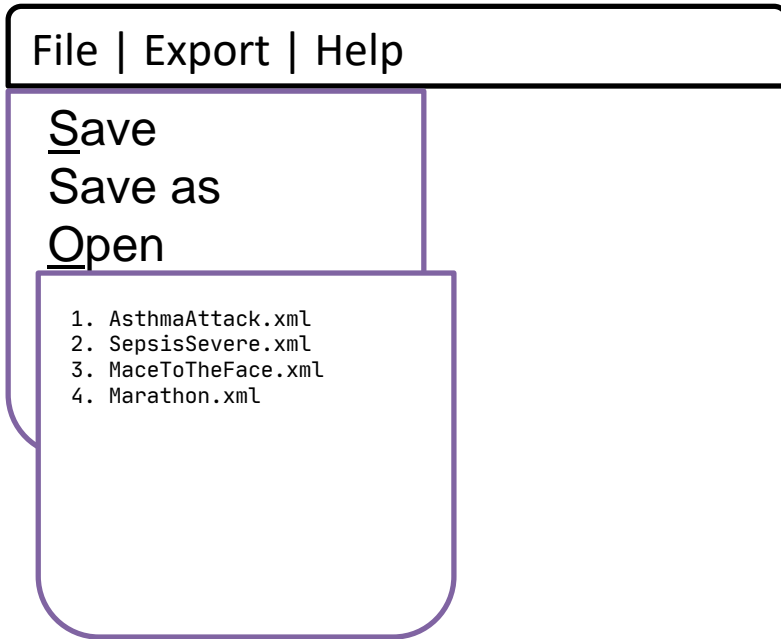
File

- Save – <patient_name>@<scenario_time>.xml
- Save As – User Chosen Filename
- Open – Open a Scenario File
- Exit – Exit Program

Export

- Patient – Export Patient to a File
- State - Same as Save
- Environment – Export env to a File
- Scenario – Export Current Scenario to a File

Scenario Open



Purpose:

- Allows user to quickly open a scenario file
- State opening handled in interaction area

UX:

- Jump list of previous 10 scenarios

Hot Keys

- Ctrl 0-9 (Load matching entry)
- Control + O (File Dialog)

On Click

- (Entry) Load shown scenario
- (Load) File Dialog for finding new scenario file

Scenario Timeline

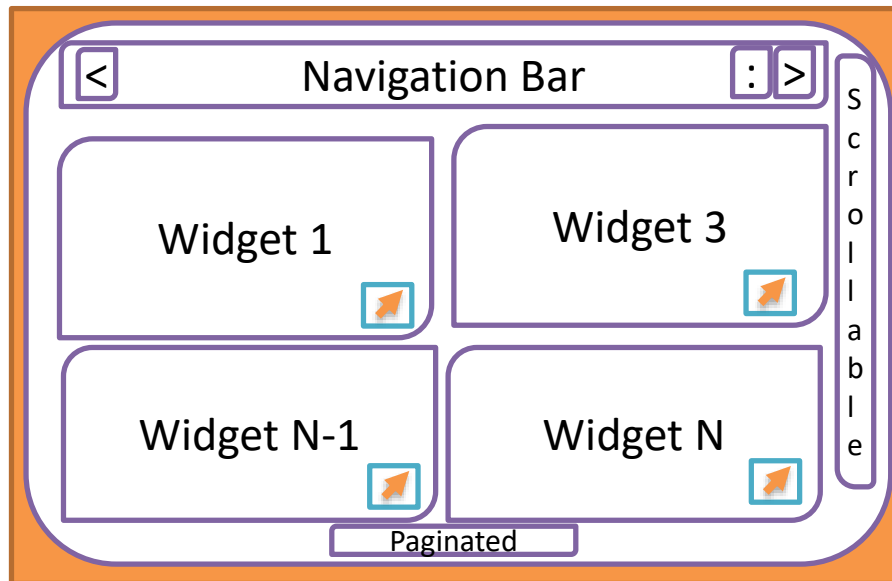
- In the current UI actions cannot be scheduled
- Instantaneous applications of actions is great for experimentation and demonstration, but bad for repeatable trials
- Scenario timeline would allow scheduling of events at a given simulation time
- Timelines could be imported and exported by CDM::ScenarioData format

Future Functionality

- Multi Window UI
 - Ability to detach widgets from main window to a sibling window
- LUA Based Scripting Engine
 - Command line interface to apply actions
 - Compound actions through LUA scriptable buttons
- Parallel Simulations
 - Scenario 2.0 batch runs where branch points launch multiple simulations
 - Ability to perform state generation
- cmd_bio Parity
 - Patient State Generation
 - Substance Data Generation
 - .config file support for batch runs.

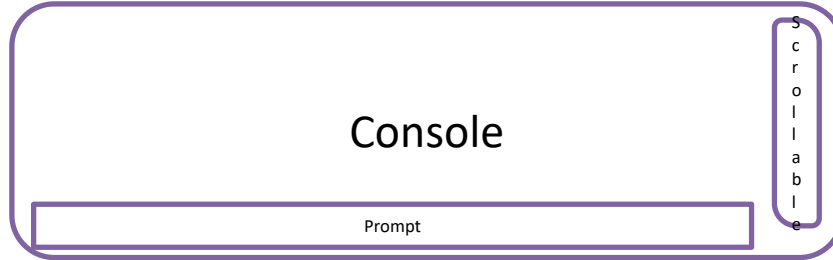
Multi UI Window

- Pop out windows for complex widgets
- Allows users to see multiple categories simultaneously

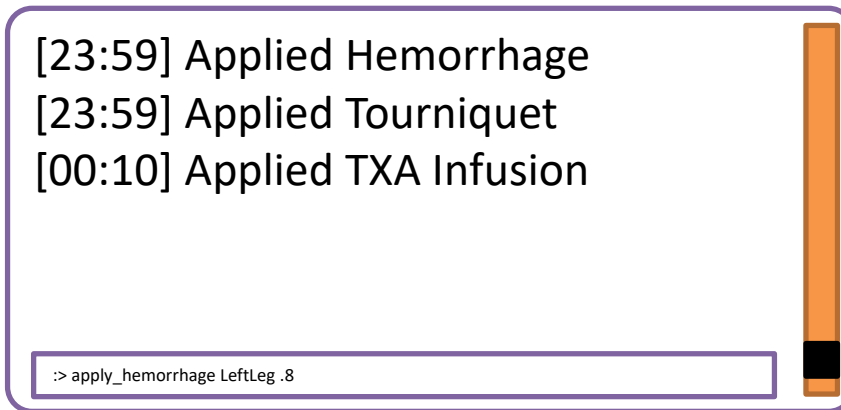


LUA Scripting

Breakdown



Example



Purpose: All CMD-like control of biogears actions and insults.

LUA engine will allow API access to current engine

Features

- Tab Complete
- Prompt History = 1000 inputs
- Prompt History Storage File = biogears_history
- Interactive help function

Example

- *hemorrhage LeftLeg .8*
- *pain_stimulus LeftLeg .5*
- *infuse_compound txa .5 1*
- Help hemorrhage

LUA Scripting (Stretch Goal)

- Custom Widgets using Data Request API
 - Advance features to assist in model validation
 - Generic plot support
 - Configurable Y-scale factors
 - Text field entry of compartment and physiology requests
- Custom Action using Exposed API
 - Advance feature to assist in action development
 - Editable text fields for lua scripting
 - Syntax verification before saving
 - Log messages reported in console area

CMD_BIO Parody

BioGears intends to eventually reach feature parity with our command line utilities.

- **Future Features:**

- Generate Patient States
 - Run stabilization for all patients in patient_dir and create a [patient@0s.xml](#) in states_dir
- Generate Substance Data
 - Read Substance.csv or excel master file and generate CDM files in substances/
- Run Parallel Simulations using Multi-Threading from a *.config file
 - Configure multiple batch runs with varying parameters
 - Support multiple patients and multiple timelines
 - Automatically run a configured number in parallel and save results to csv file
- Export Graph Plots to Image Format.
 - After a run give csv results file
 - Generate PNG images for presentations for proof of validation
 - Useful for presenters and for historical review
 - Plots will feature baseline support to show variation from expected or clinical data points
 - Complex feature which may take several iterations to implement fully