

Modelica Day 2015

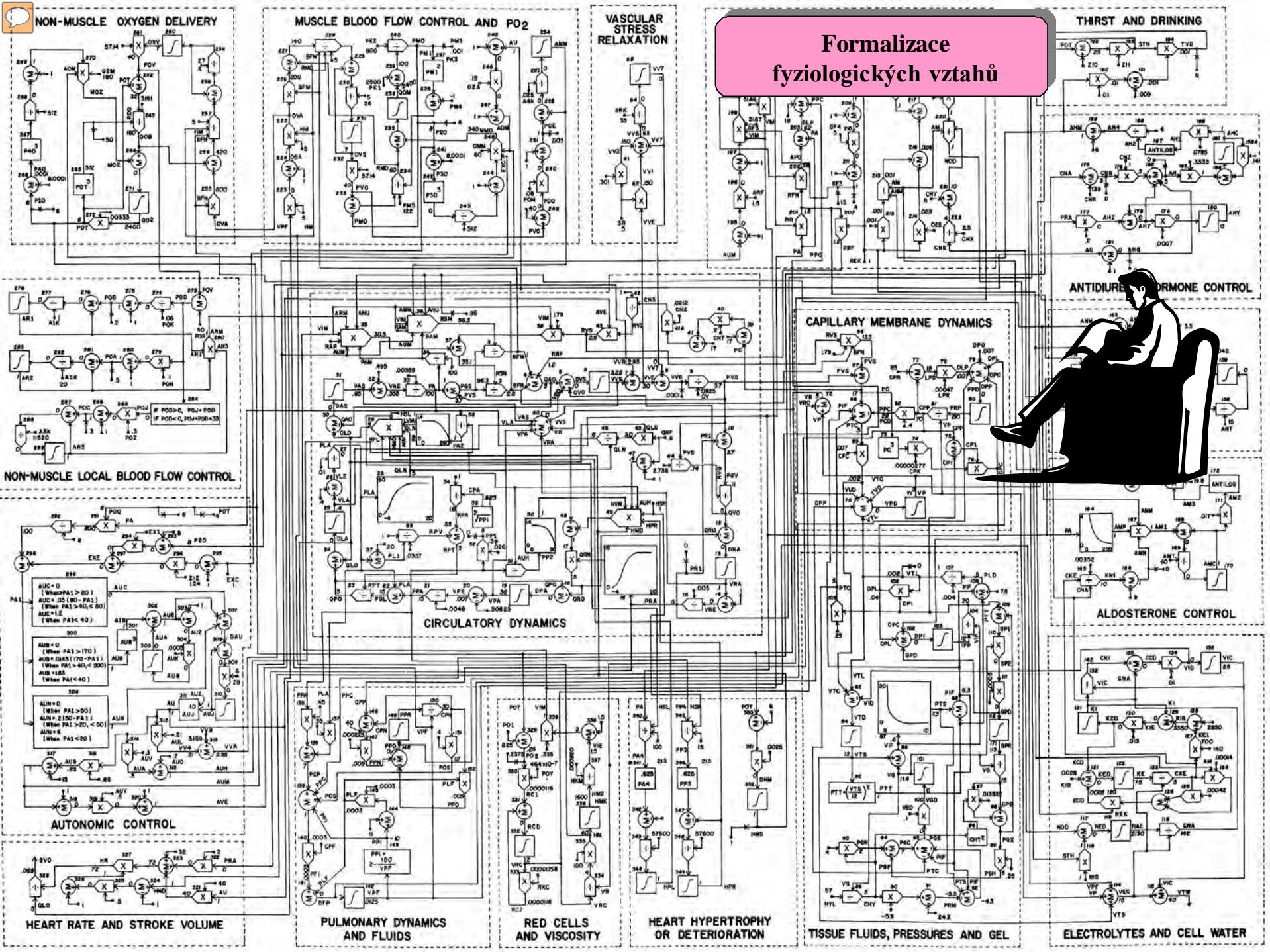
Jiří Kofránek

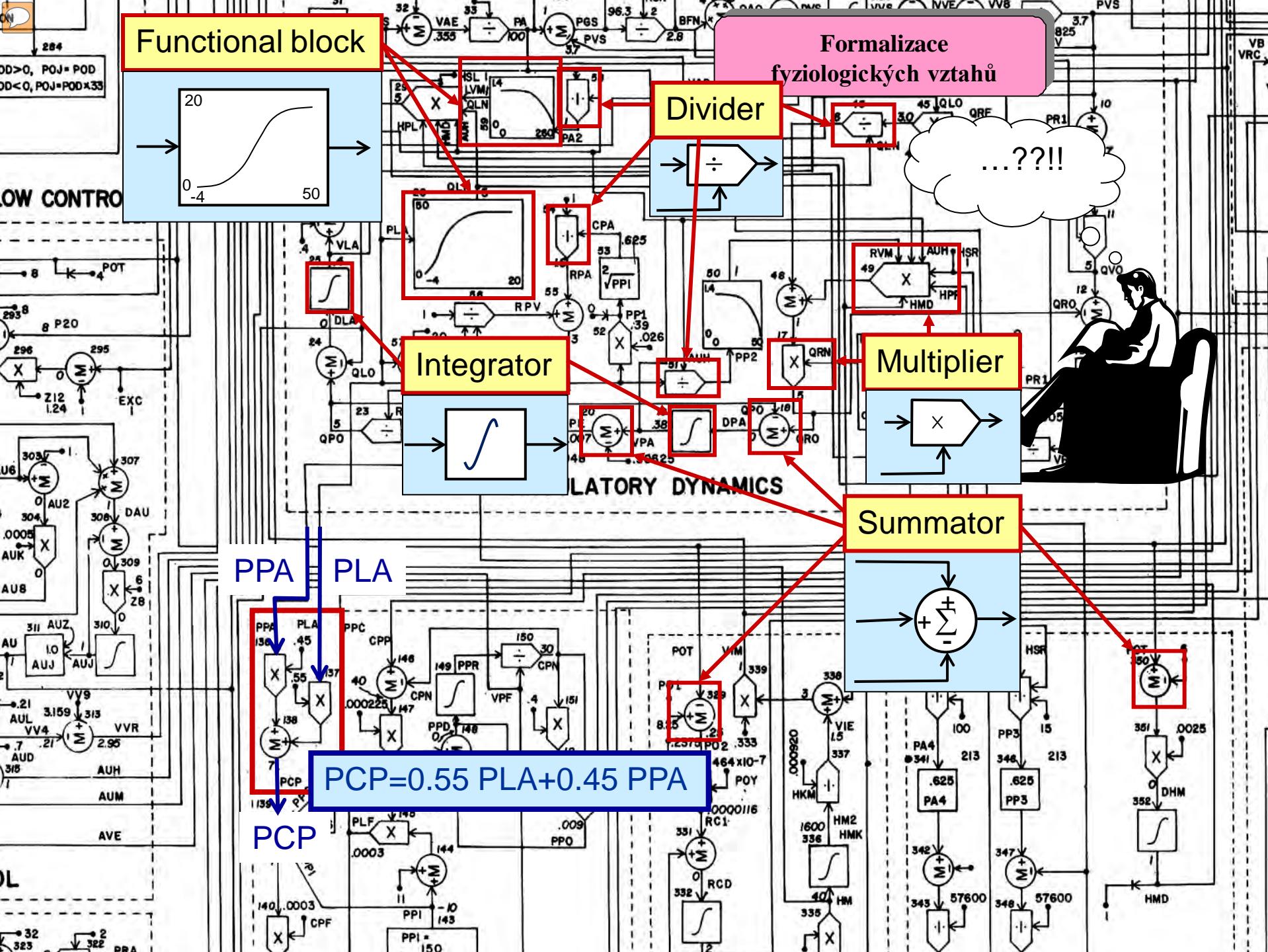


Naše cesta k Modelice

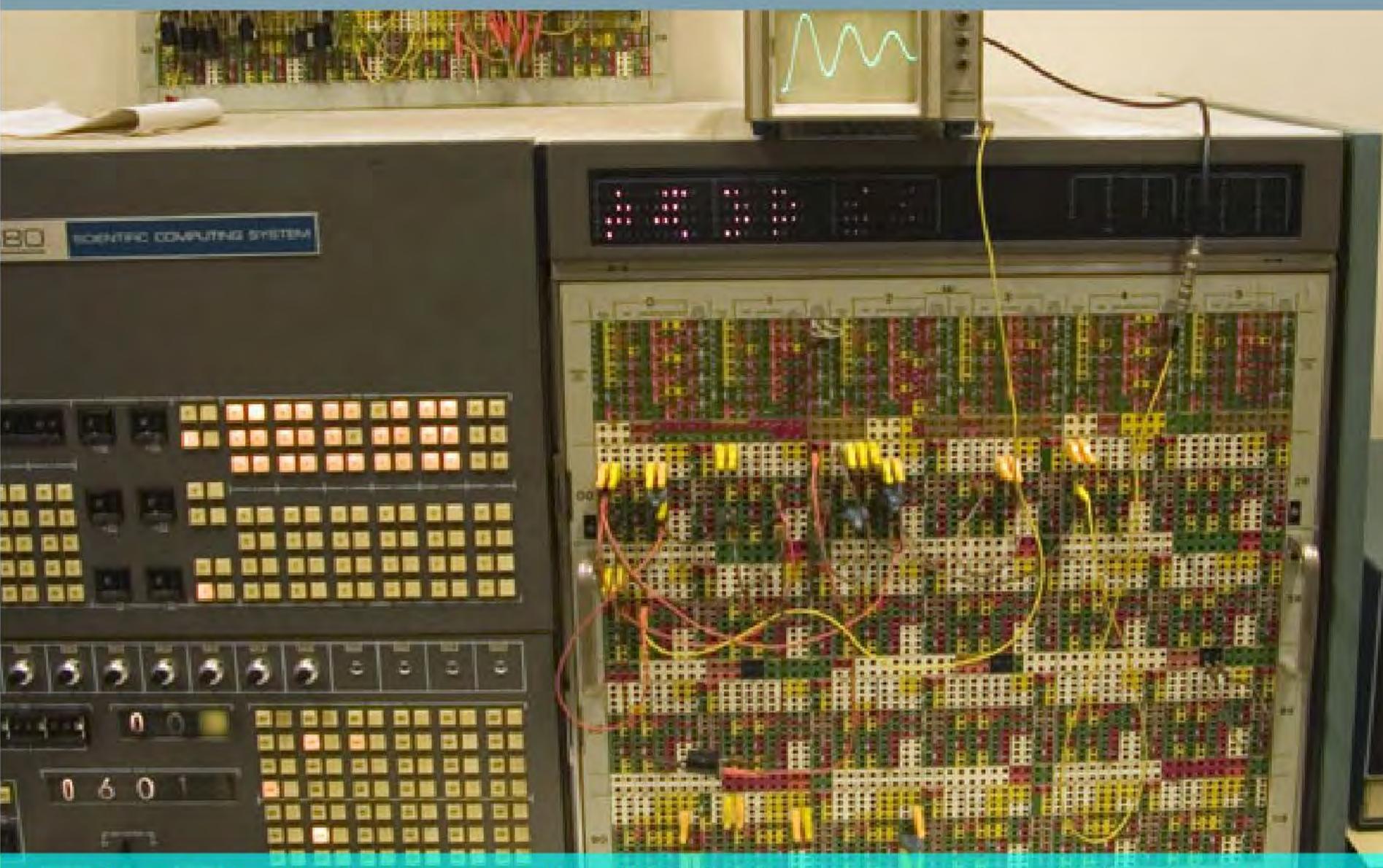
Jiří Kofránek

Formalizace fyziologických vztahů

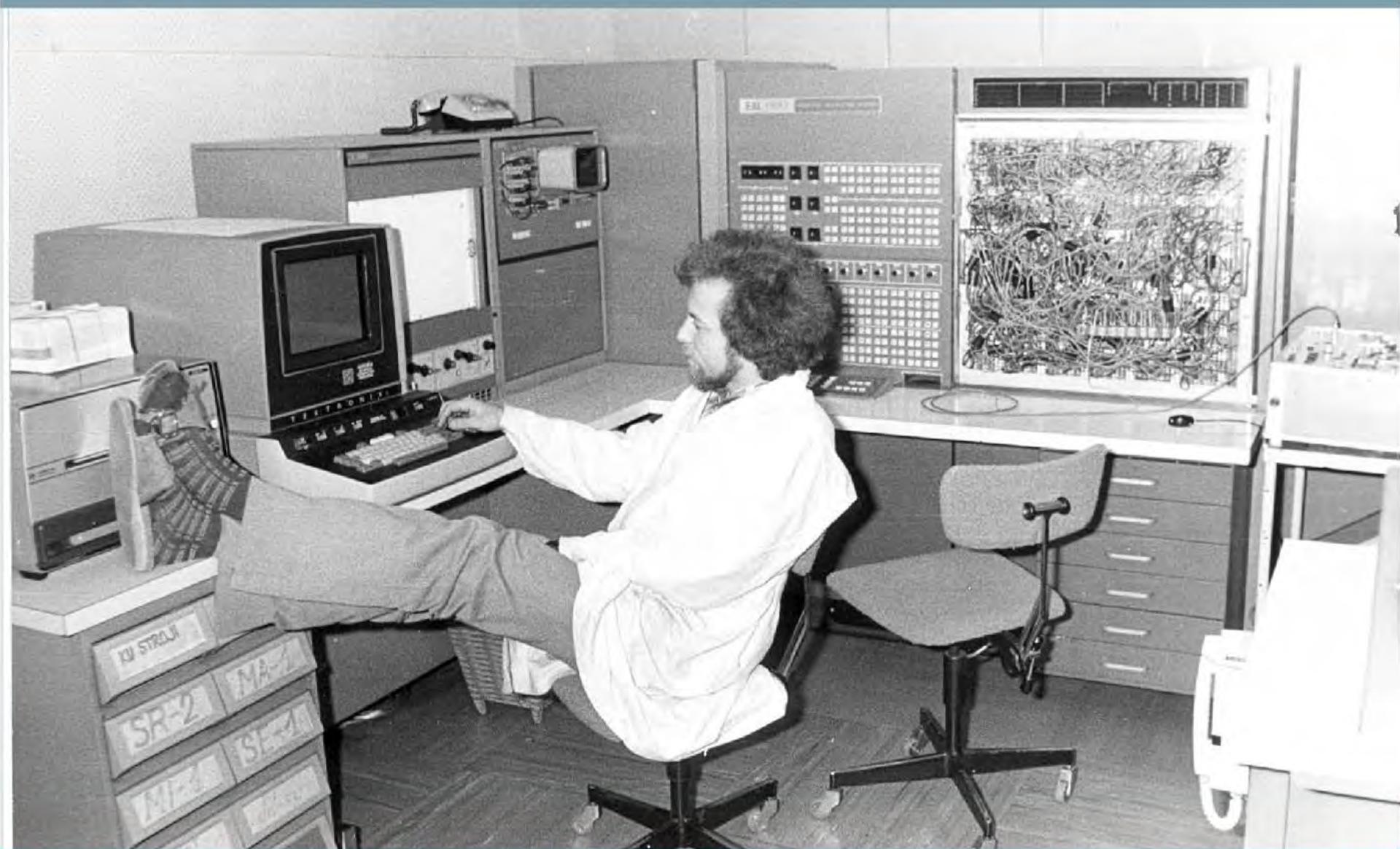




Nástoje pro modelování - specializovaná softwarová prostředí



Analogové a hybridní počítače: fyzické propojování výpočetních prvků kablíky



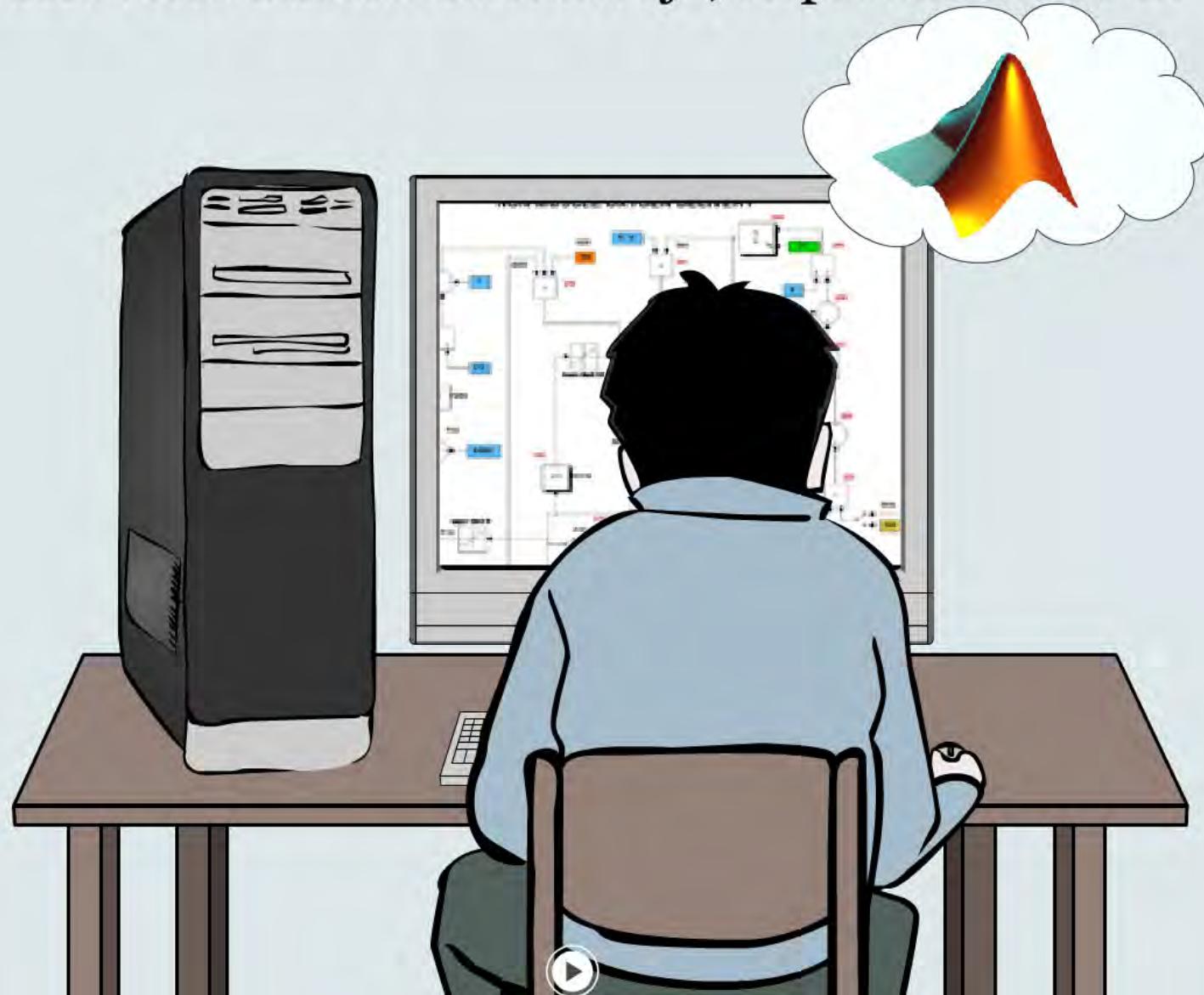
Analogové a hybridní počítače: fyzické propojování výpočetních prvků kablíky

Číslcové počítače: modely ve standardních programovacích jazycích (FORTRAN, PASCAL, C++ ...)

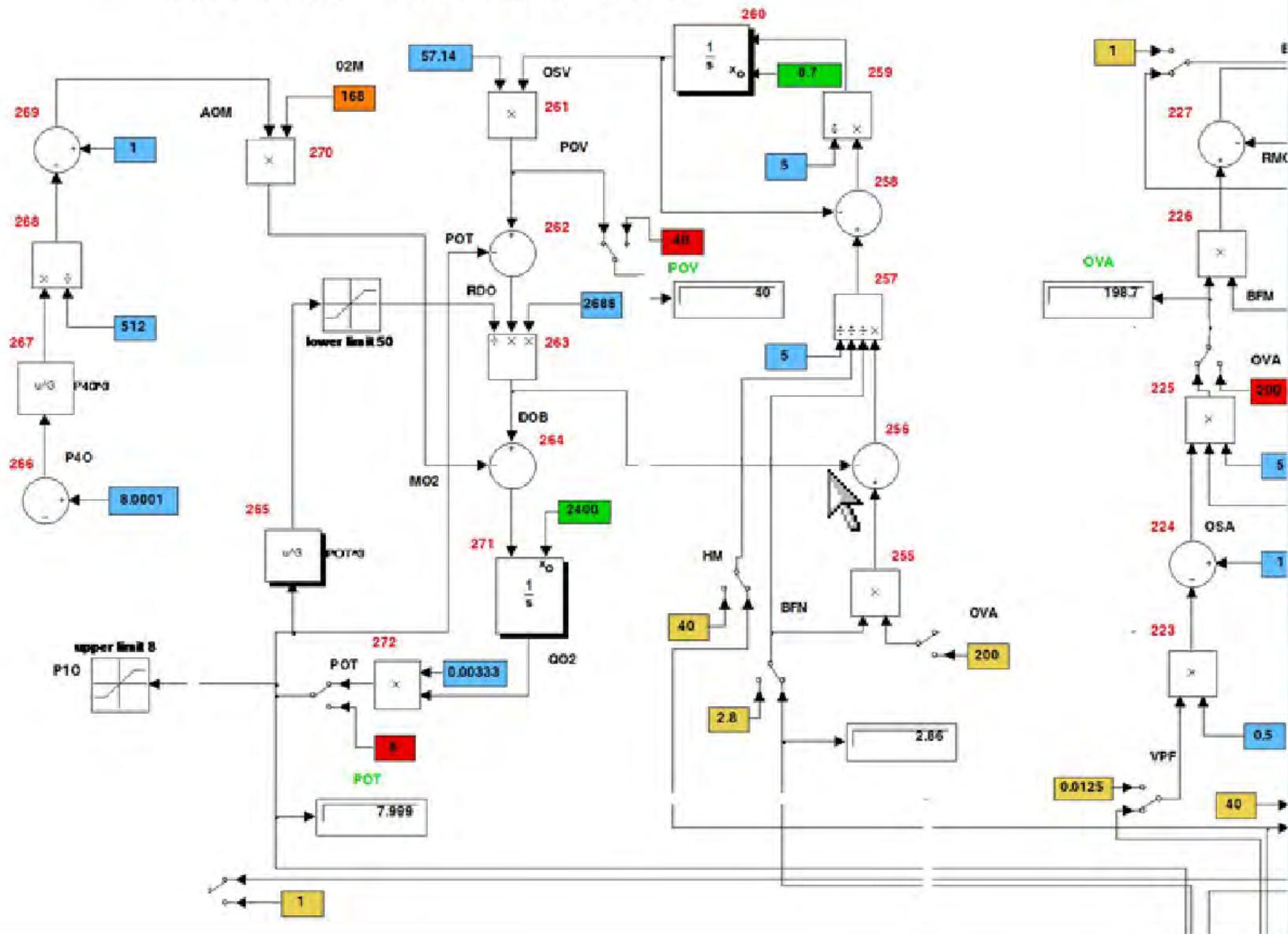


Analogové a hybridní počítače: fyzické propojování výpočetních prvků kablíky

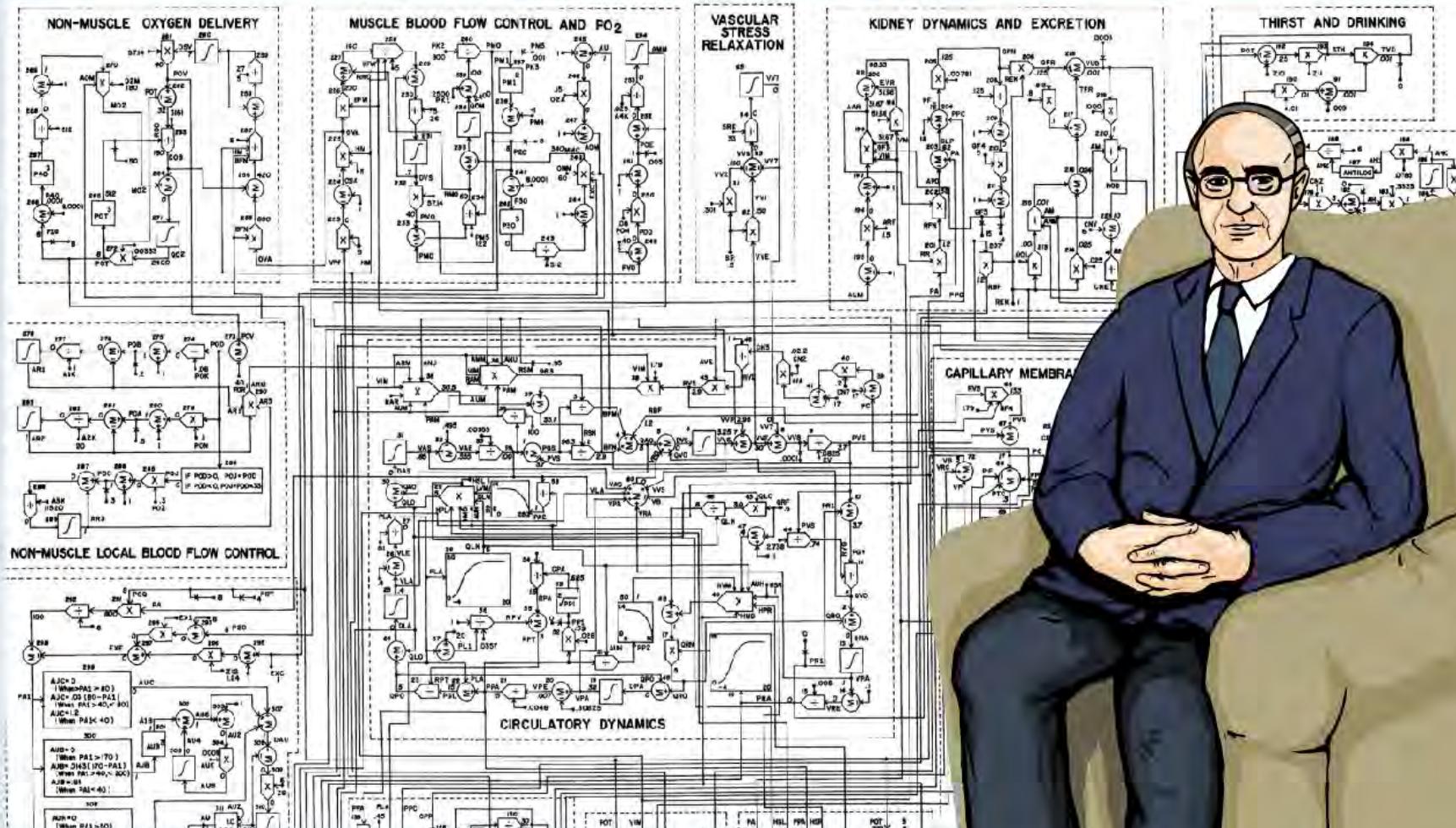
Specializované simulační nástroje, např. SIMULINK



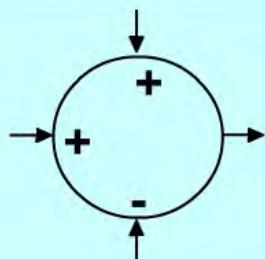
NON-MUSCLE OXYGEN DELIVERY



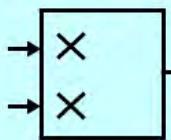
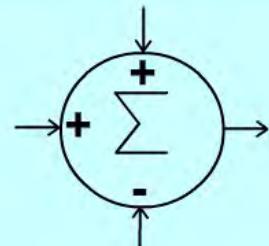
Guytonův diagram v Simulinku



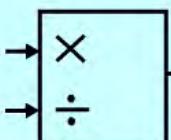
Guytonův diagram v Simulinku



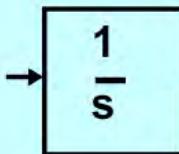
Sumátor



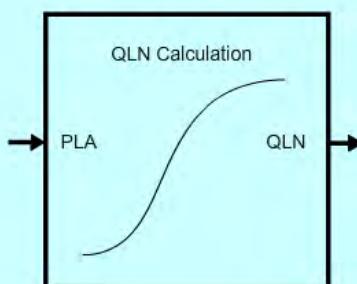
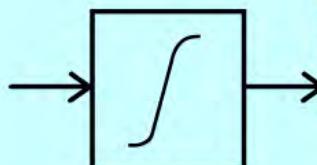
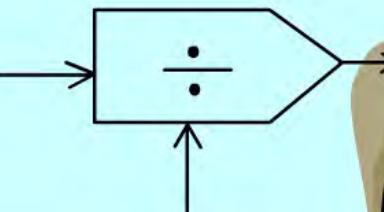
Násobička



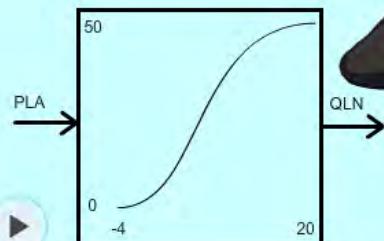
Dělička



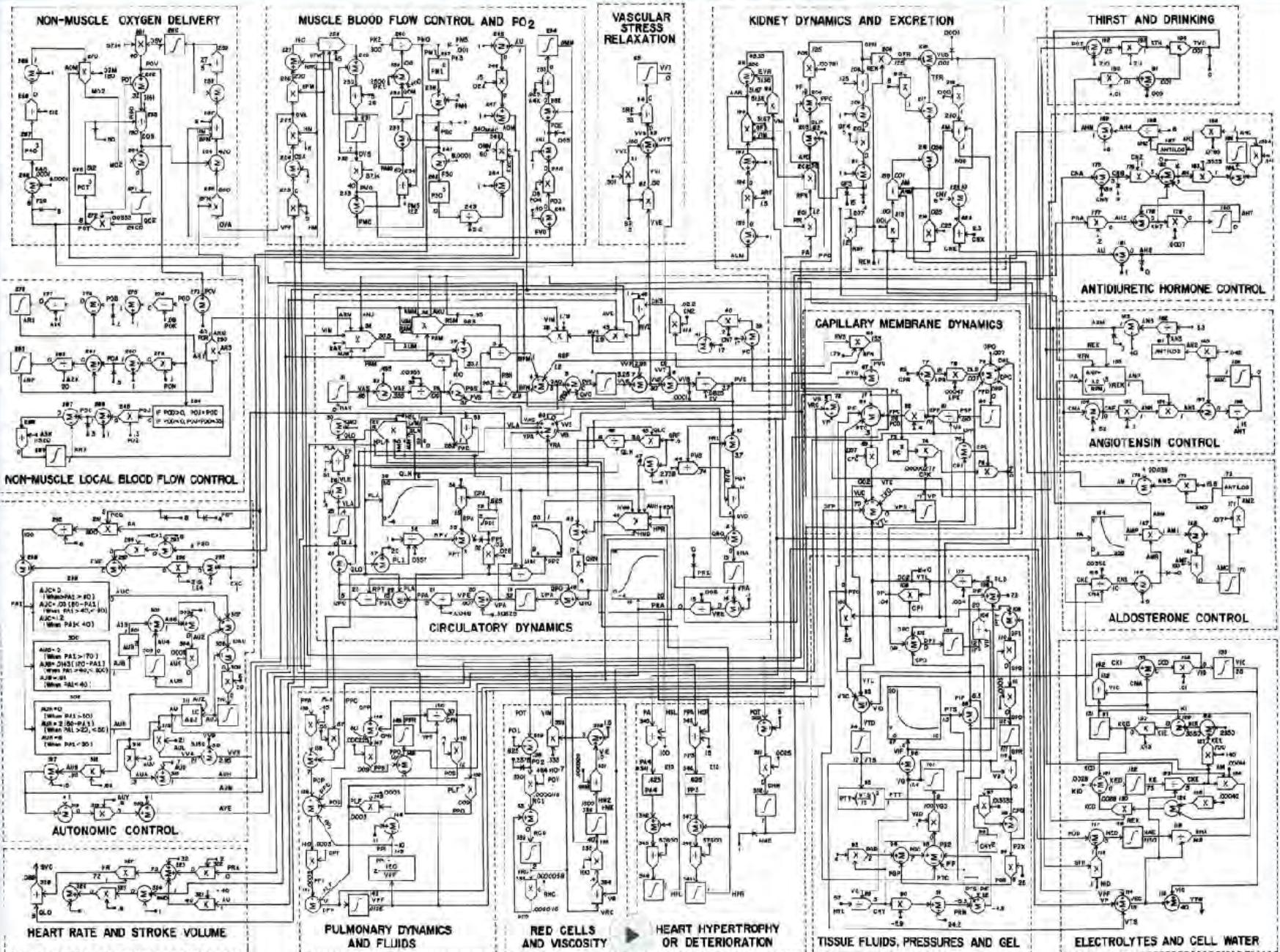
Integrátor



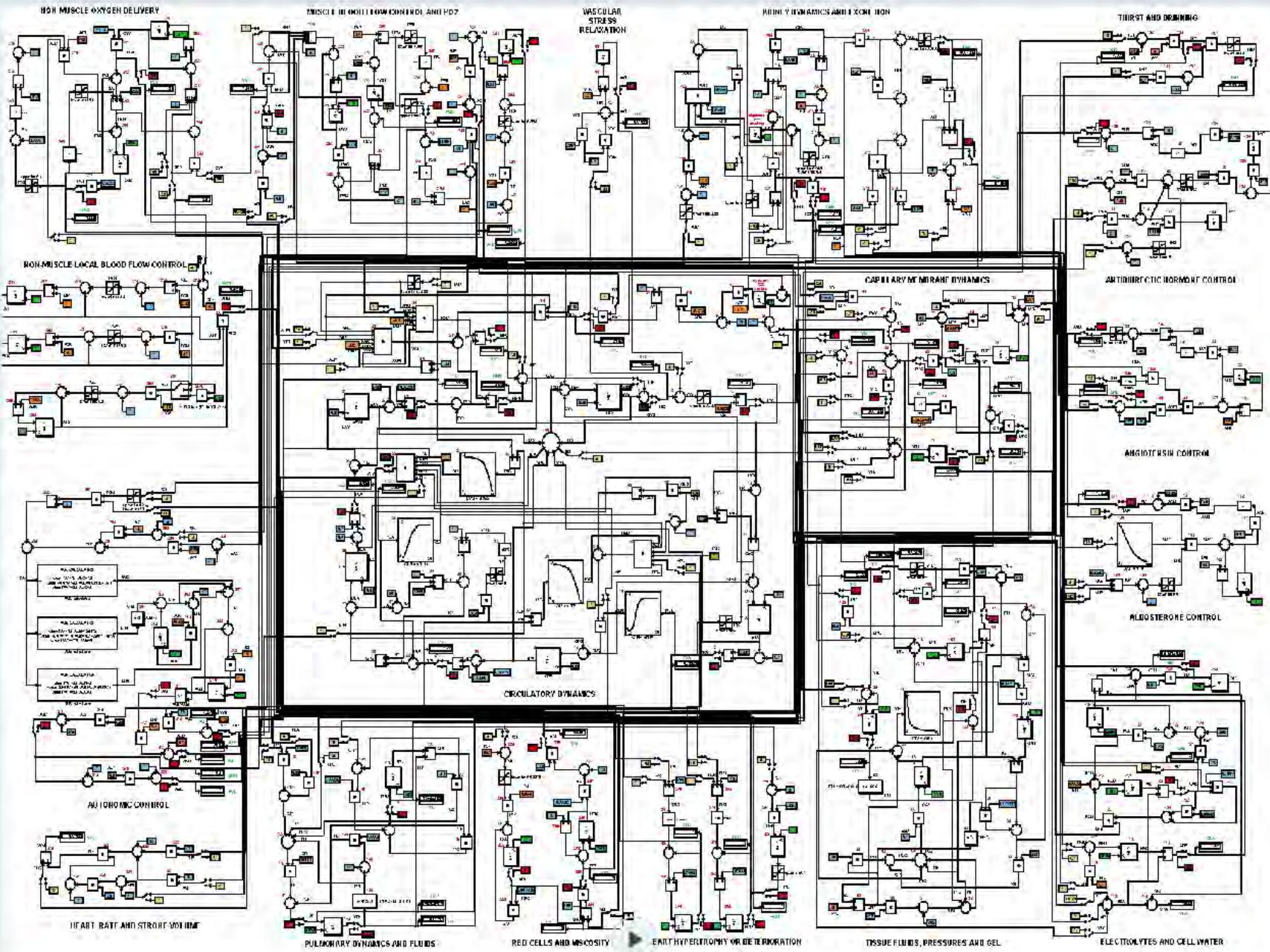
**Funkční
blok**



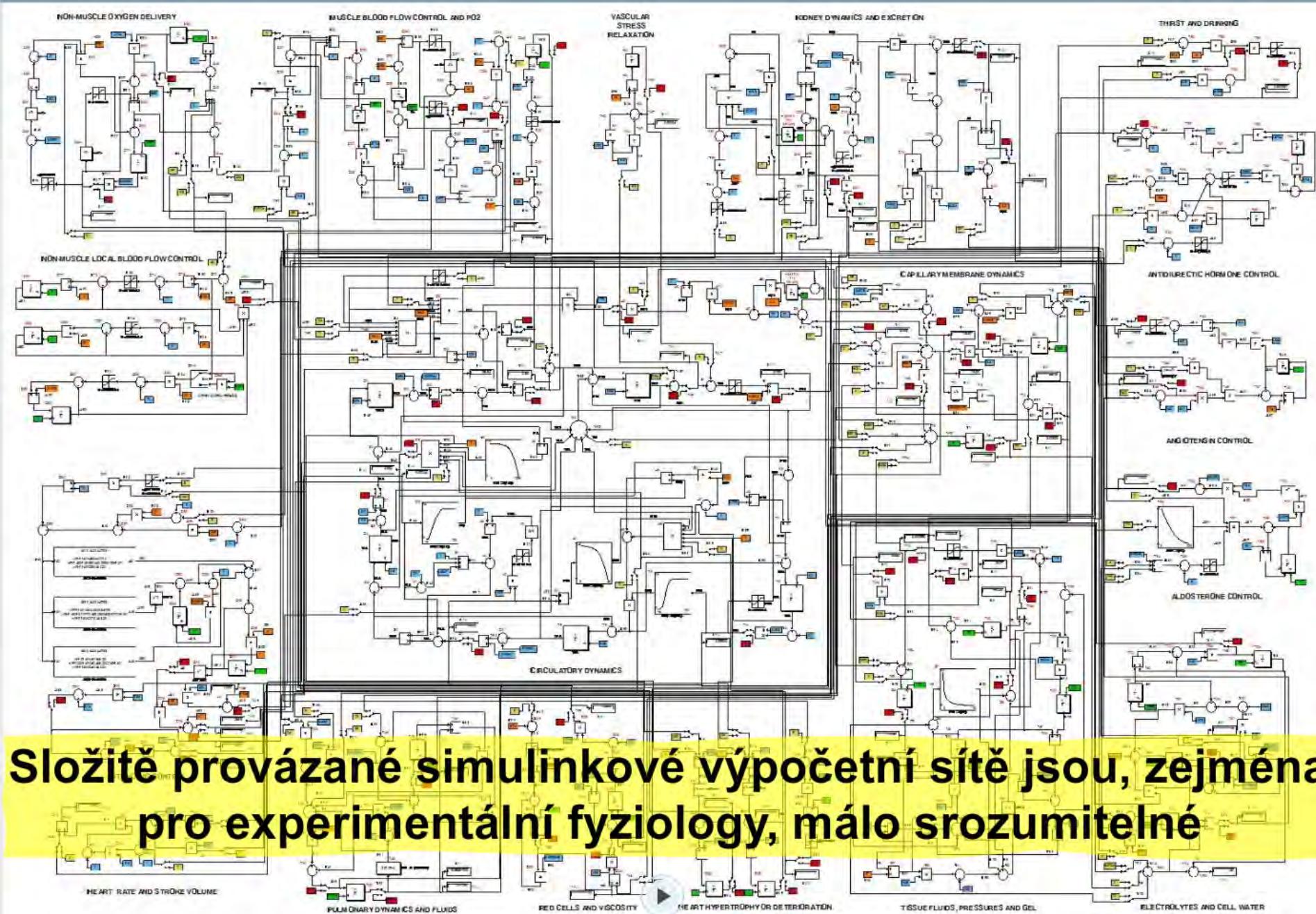
Guytonův diagram v Simulinku



Guytonův diagram v Simulinku



Simulinková knihovna Physiolibrary

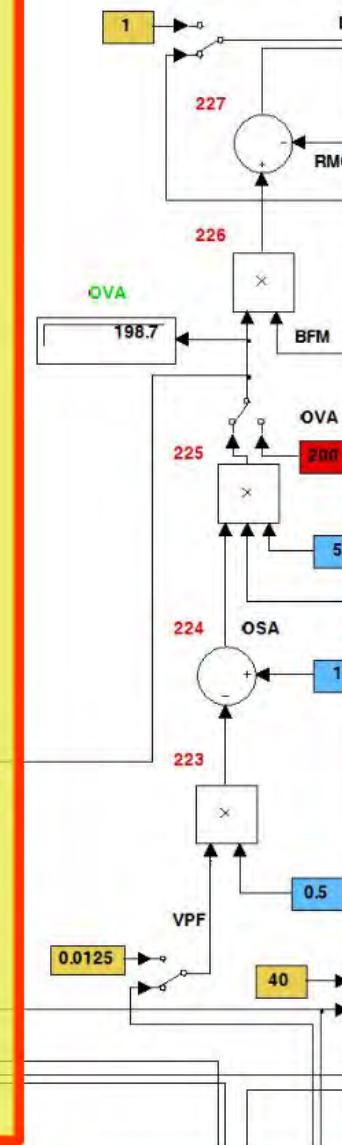


Složitě provázané simulinkové výpočetní sítě jsou, zejména pro experimentální fyziologie, málo srozumitelné

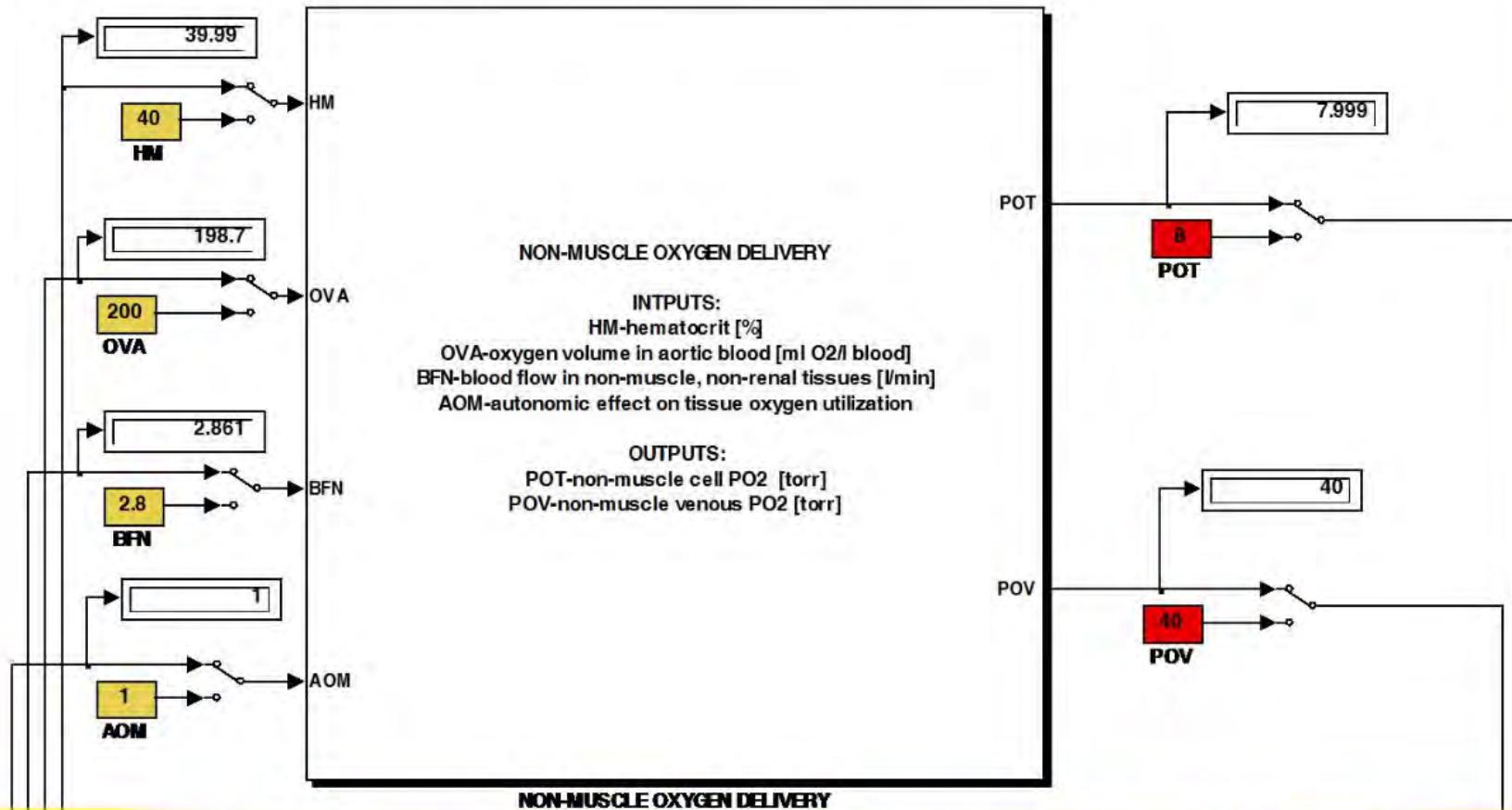
NON-MUSCLE OXYGEN DELIVERY



Proto se vyplatí jednotlivé výpočetní prvky agregovat do bloků a jejich vnitřní propojení před uživatelem skrýt

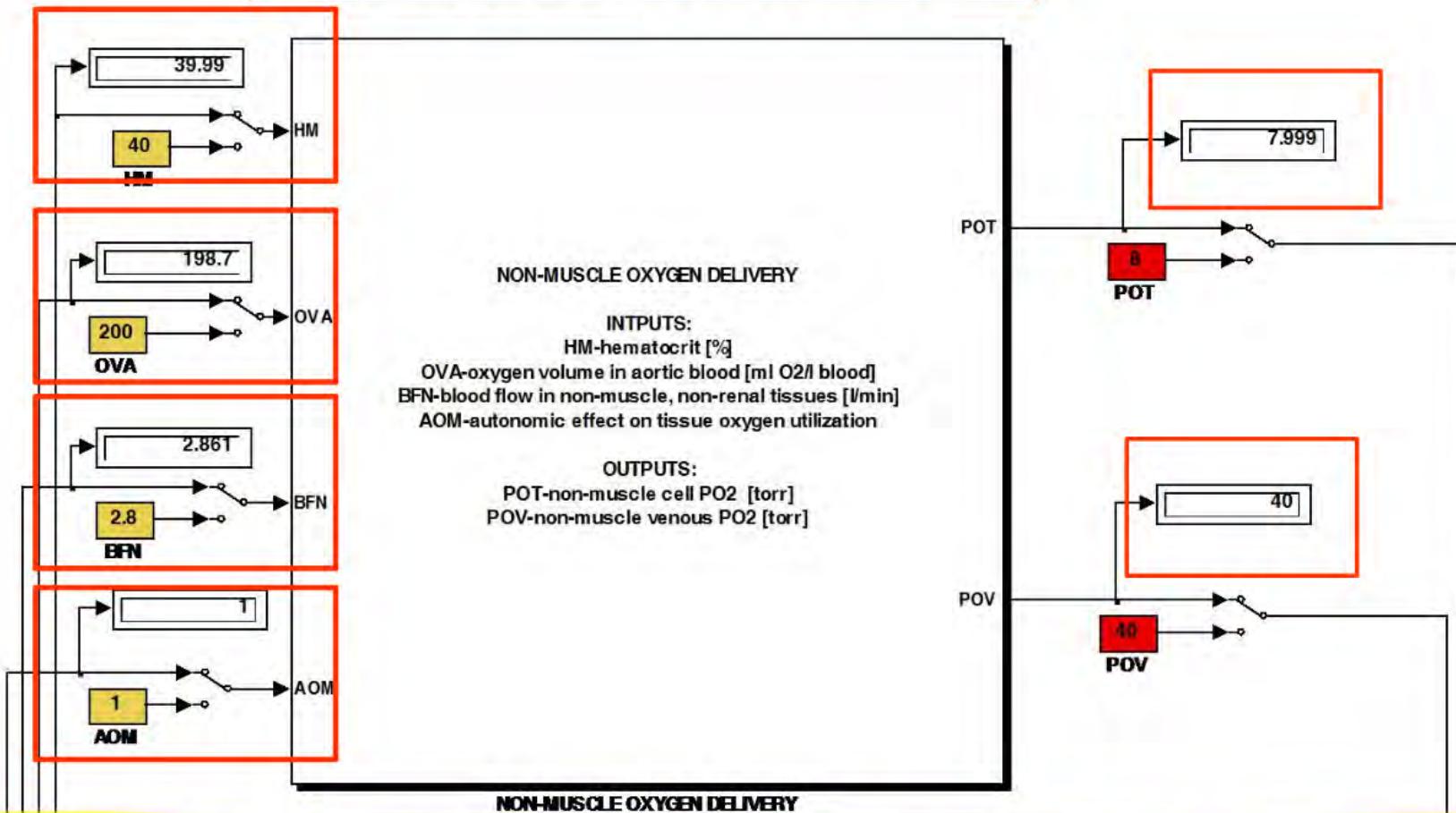


NON-MUSCLE OXYGEN DELIVERY



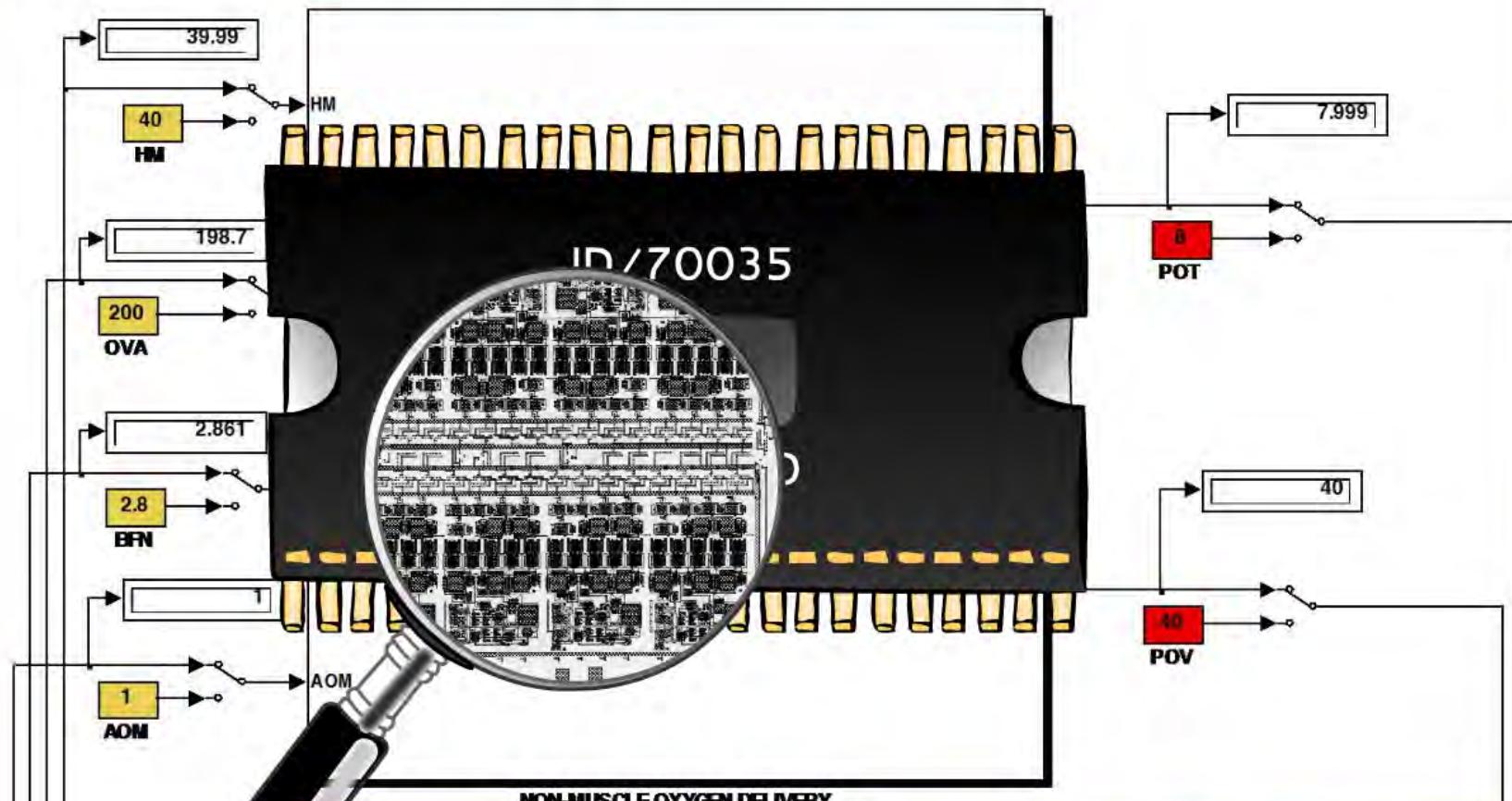
**Agregovaný simulační blok s definovanými vstupy
a výstupy je pro fyziology mnohem srozumitelnější**

NON-MUSCLE OXYGEN DELIVERY



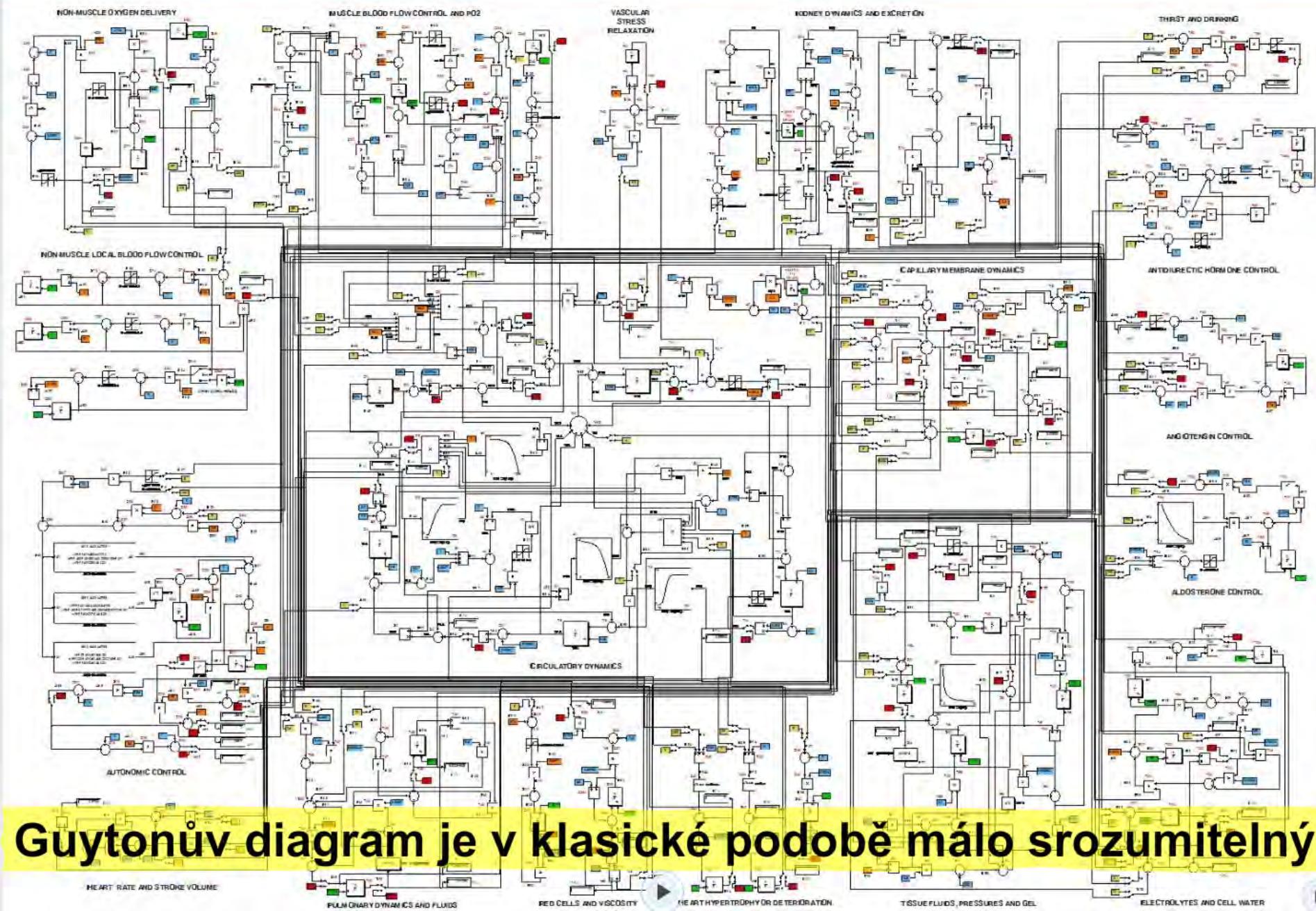
V simulačním nástroji "Simulink" můžeme testovat chování agregovaného simulačního bloku

NON-MUSCLE OXYGEN DELIVERY

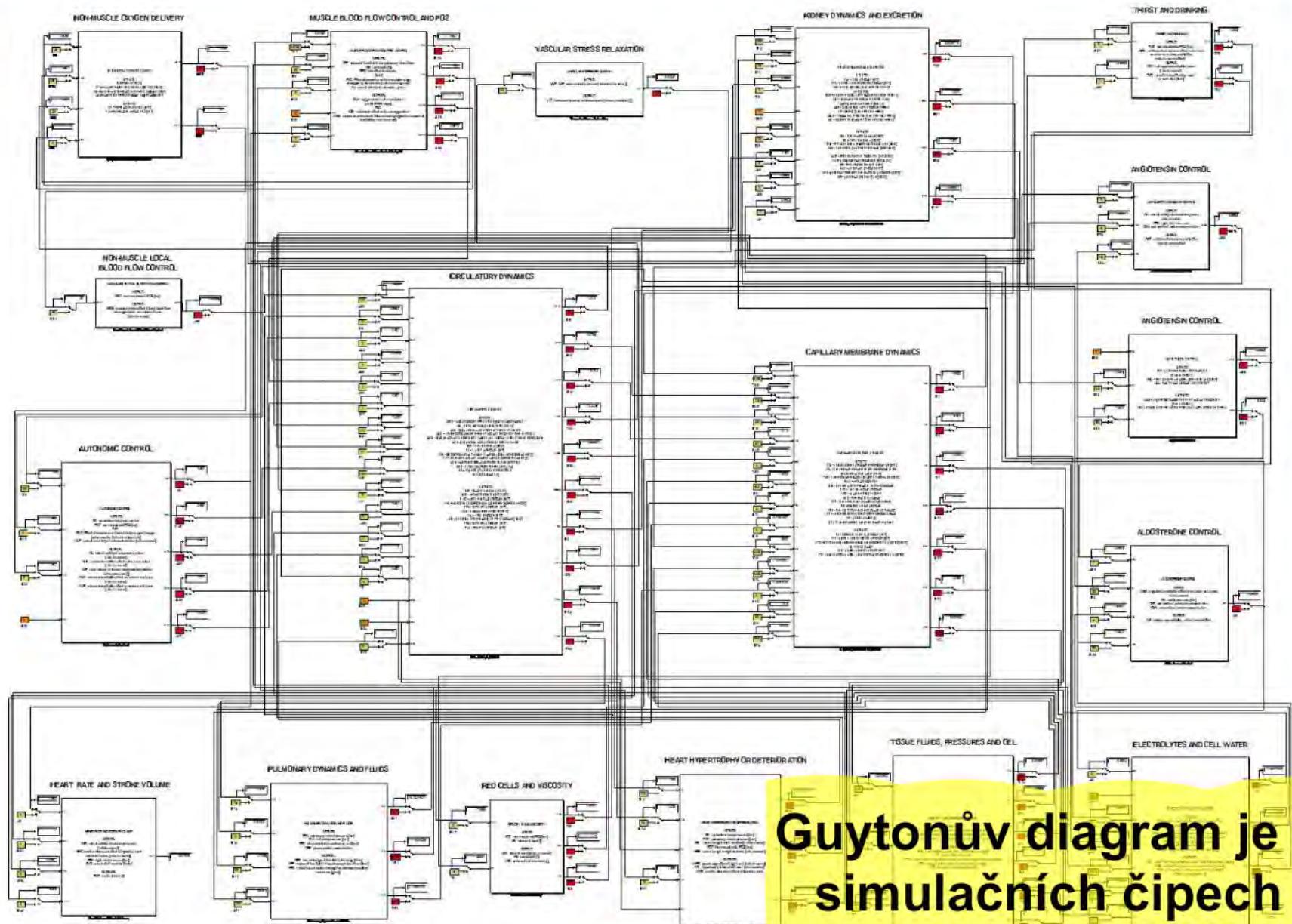


"Simulační čip" (obdobně jako elektronický čip) skrývá před uživatelem svoji vnitřní strukturu (rovnice)

Simulinková knihovna Physiolibrary

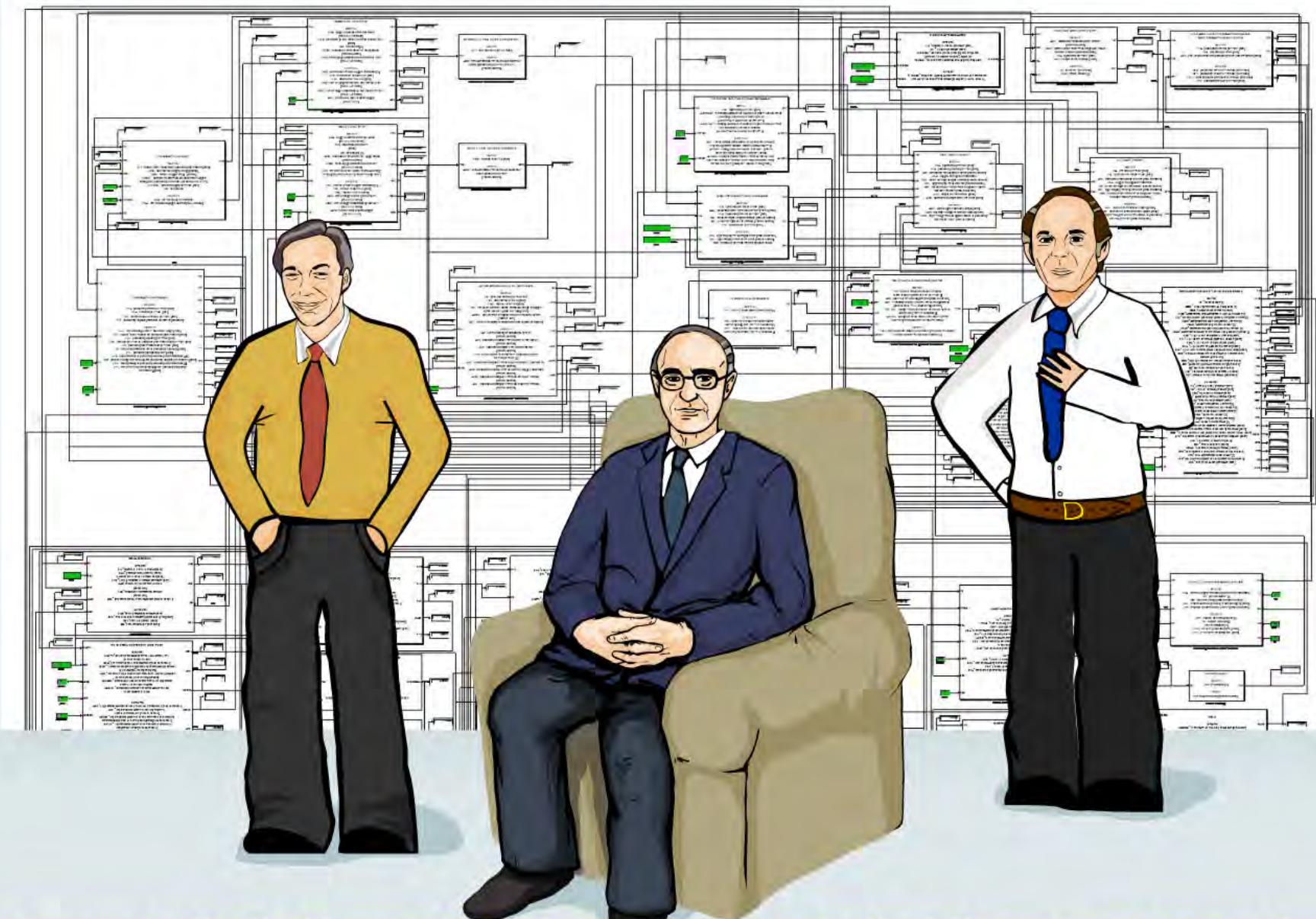


Simulinková knihovna Physiolibrary



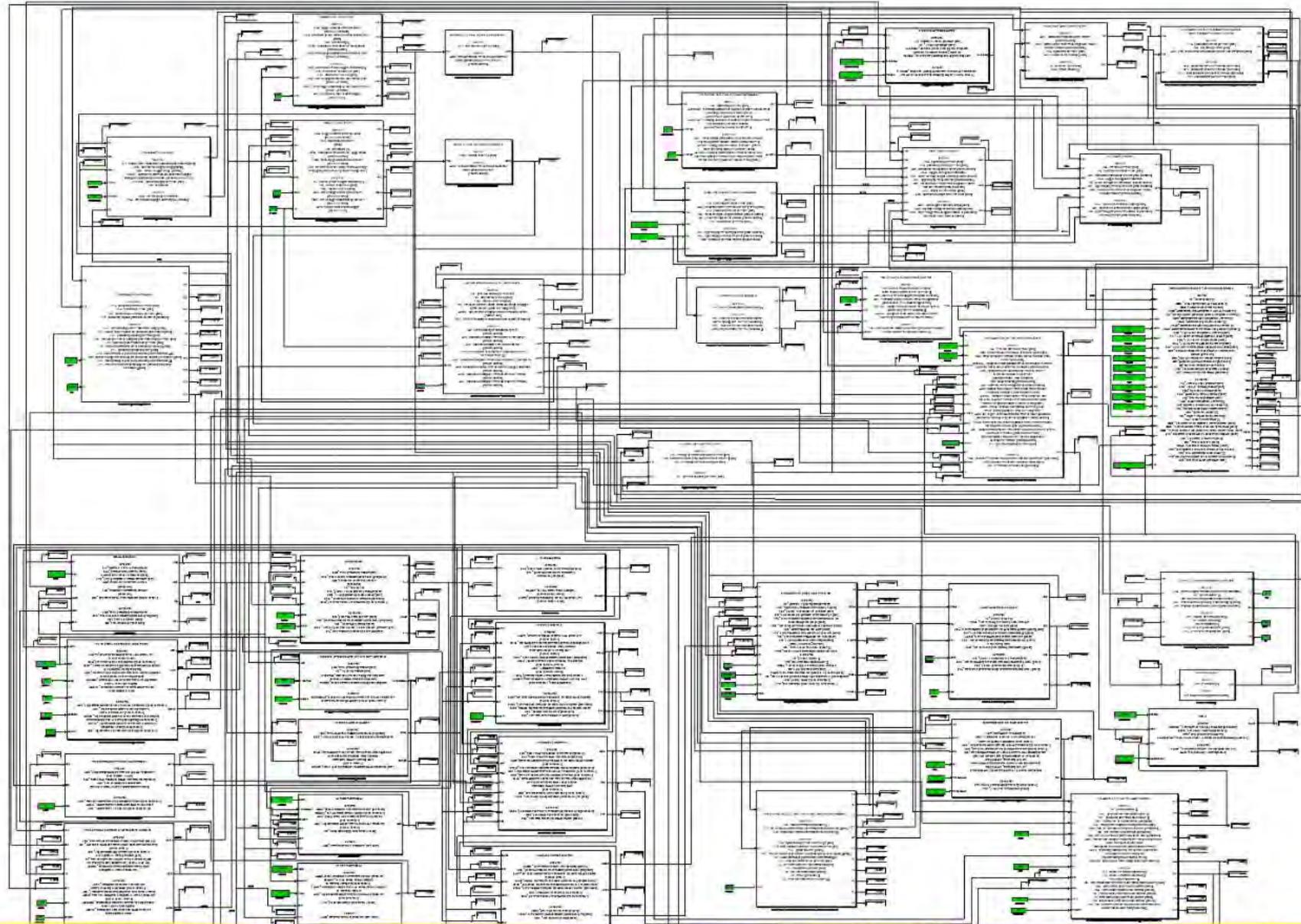
Guytonův diagram je v
simulačních čipech
srozumitelnější

Simulinková knihovna Physiolibrary



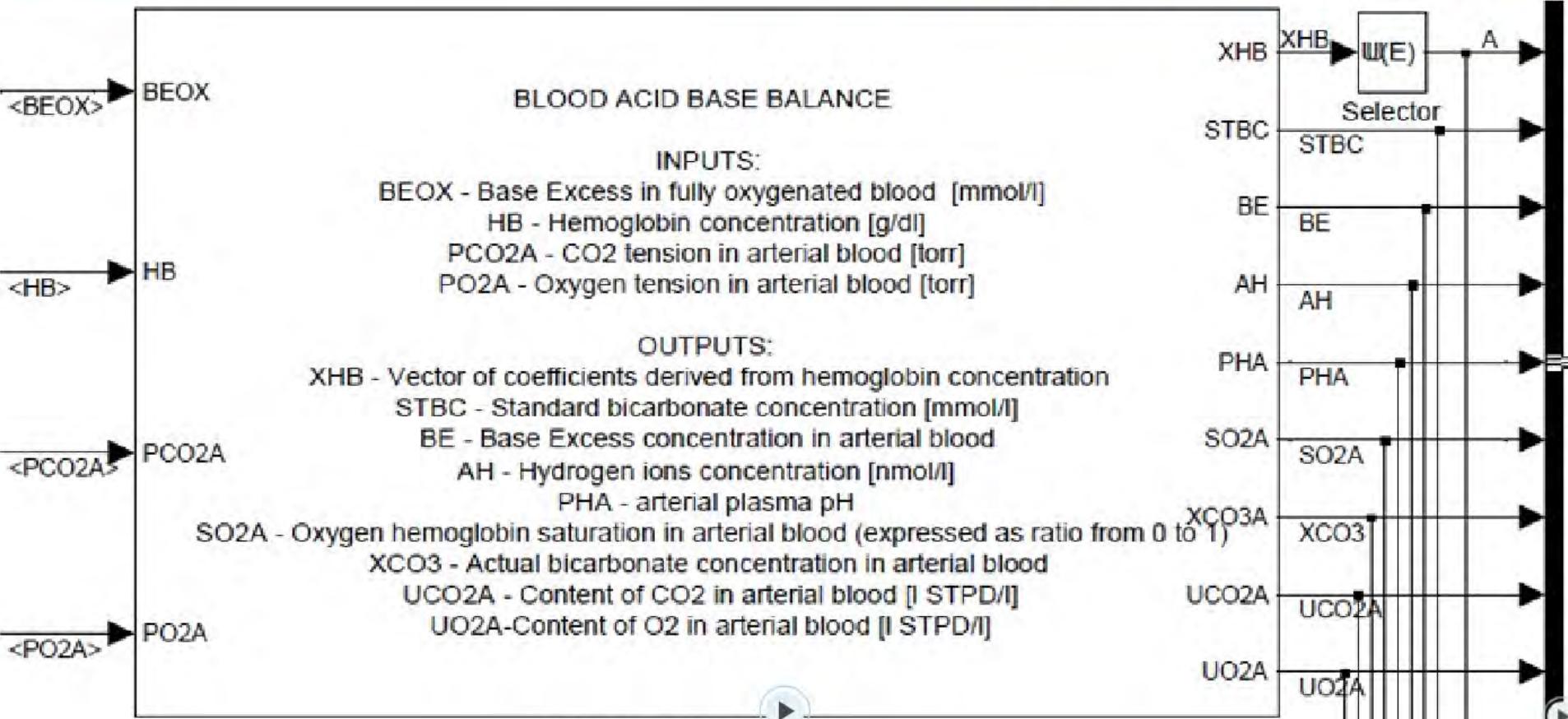
Pro složitější modely jsou "simulační čipy" nutností

Simulinková knihovna Physiolibrary

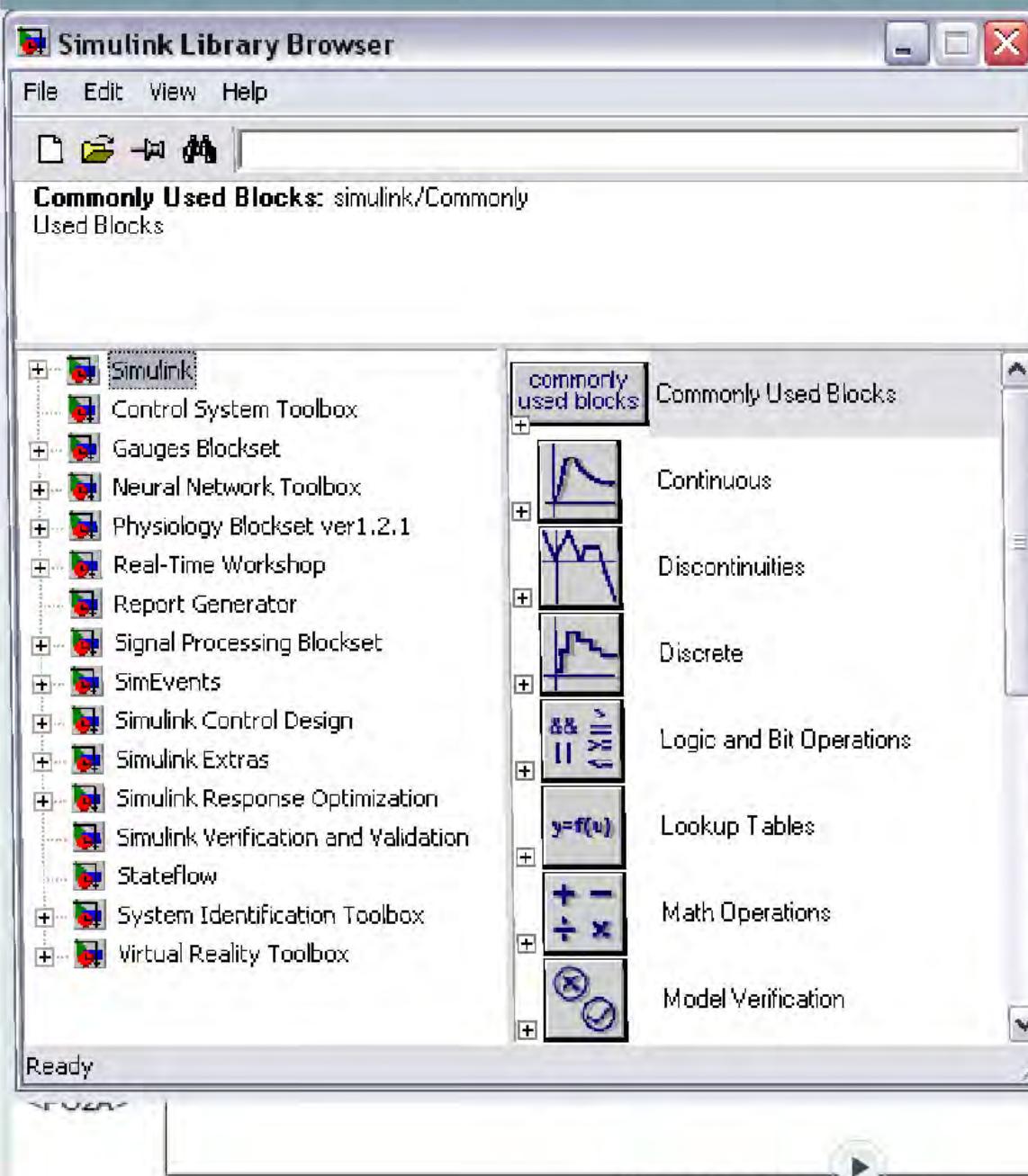


Pro složitější modely jsou "simulační čipy" nutností

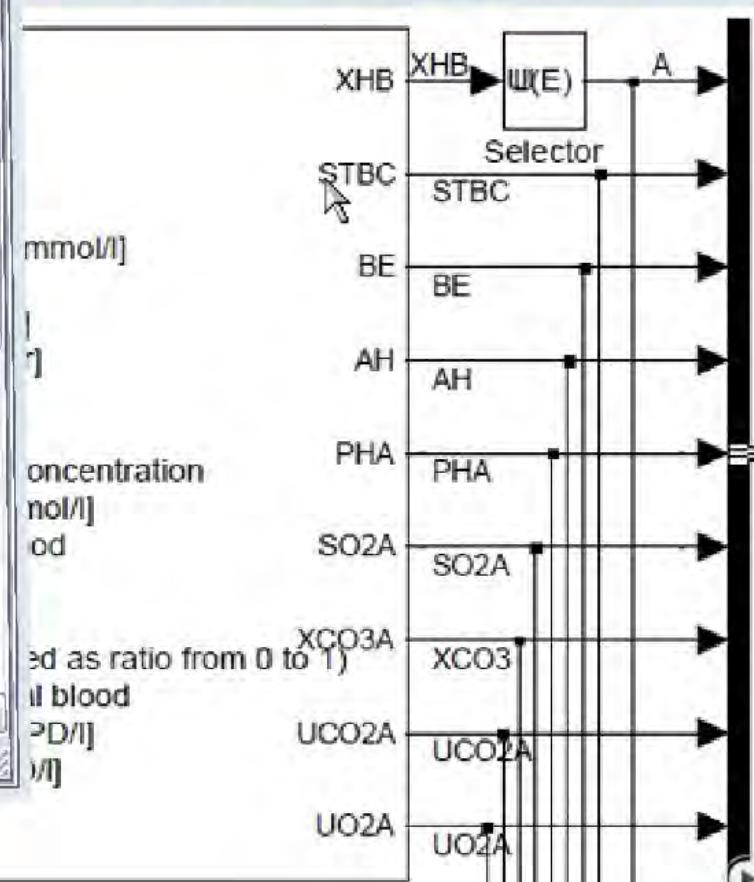
Jednou už vytvořené
 "Simulačních čipy"
 jsou znovupoužitelné!
 Dají se využívat v
 různých modelech



Simulinková knihovna Physiolibrary



Ve veřejně přístupné knihovně
PHYSIOLIBRARY je nabídka "simulačních čipů" ke stažení



Simulinková knihovna Physiolibrary

Help

File Edit View Go Favorites Desktop Window Help

Help Navigator

Contents Index Search Demos

MATLAB

- MATLAB Compiler
- MATLAB Report Generator
- MATLAB Web Server
- Bioinformatics Toolbox
- Control System Toolbox
- Curve Fitting Toolbox
- Datafeed Toolbox
- Financial Toolbox

R o a d m a p

Physiology blockset

Blood gases models

- Lungs & Tissues - Model of blood gases exchange. Respiration and O₂ / CO₂ metabolism.
- "Bloods Model" - The vascular model of blood gases exchange. Respiration and O₂ / CO₂ metabolism.

BLOOD ACID BASE BALANCE

INPUTS:

- BEOX - Base Excess in fully oxygenated blood [mmol/l]
- HB - Hemoglobin concentration [g/dl]
- PCO₂A - CO₂ tension in arterial blood [torr]
- PO₂A - Oxygen tension in arterial blood [torr]

OUTPUTS:

- XHB - Vector of coefficients derived from hemoglobin concentration
- STBC - Standard bicarbonate concentration [mmol/l]
- BE - Base Excess concentration in arterial blood
- AH - Hydrogen ions concentration [nmol/l]
- PHA - arterial plasma pH
- SO₂A - Oxygen hemoglobin saturation in arterial blood (expressed as ratio from 0 to 1)
- XCO₃A - Actual bicarbonate concentration in arterial blood
- UCO₂A - Content of CO₂ in arterial blood [l STPD/l]
- UO₂A - Content of O₂ in arterial blood [l STPD/l]

Block Diagram:

```
graph LR; XHB --> W((W(E))); Selector[Selector] --- BE; Selector --- AH; Selector --- PHA; Selector --- SO2A; Selector --- XCO3A; Selector --- UCO2A; Selector --- UO2A; W --> A
```

Ve veřejně přístupné knihovně PHYSIOLIBRARY je nabídka "simulačních čipů" ke stažení

Simulinková knihovna Physiolibrary

Help

File Edit View Go Favorites Desktop Window Help

Help Navigator x

Contents Index Search Demos

MATLAB MATLAB Compiler MATLAB Report Generator MATLAB Web Server Bioinformatics Toolbox Control System Toolbox Curve Fitting Toolbox Datafeed Toolbox Financial Toolbox Genetic Algorithm and Direct Search Toolbox Image Processing Toolbox Neural Network Toolbox Optimization Toolbox Signal Processing Toolbox Spline Toolbox Statistics Toolbox Symbolic Math Toolbox System Identification Toolbox Simulink Real-Time Workshop Simulink Accelerator Simulink Control Design Simulink Report Generator Simulink Response Optimization Simulink Verification and Validation Stateflow Gauges Blockset Physiology Blockset Blood Gases GOLEM Kidney Product Page (Web) Signal Processing Blockset

← → C M

Title: Simulink

R o a d m a p

Physiology blockset

Blood gases models

- [Lungs & Tissues](#) - Model of blood gases exchange. Respiration and O₂ / CO₂ metabolism.
- ["Bloody Mary"](#) - The varying scale model of blood gases exchange. Respiration and O₂ / CO₂ metabolism. The varying structure scale during the model running. The loop cutting. Etc.
- Help for [simulation chips used in blood gases](#) chapter.

All additional informations are available at <http://patf-biofyb.fjfi.cuni.cz>.

Golem model

- [Golem](#) - The large physiological model of inner body environment. This model is one part of project Physiome and actual structure is presented on web: www.physiome.cz.
- Help for [simulation chips used in golem model](#).

SIMA model

- [Cardiovascular model](#)

Kidney model

- [Kidney](#) - Model of the kidney.
- Help for [simulation chip used in kidney model](#).

Každý nabízený simulační blok musí být dobře dokumentován

Simulinková knihovna Physiolibrary

Help

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Help Navigator

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- * MATLAB Compiler
- * MATLAB Report Generator
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- * Financial Toolbox
- * Genetic Algorithm and Direct Search Toolbox
- * Image Processing Toolbox
- * Neural Network Toolbox
- * Optimization Toolbox
- * Signal Processing Toolbox
- * Spline Toolbox
- * Statistics Toolbox
- * Symbolic Math Toolbox
- * System Identification Toolbox
- * Simulink
- * Real-Time Workshop
- * Simulink Accelerator
- * Simulink Control Design
- * Simulink Report Generator
- * Simulink Response Optimization
- * Simulink Verification and Validation
- * Stateflow
- * Gauges Blockset
- * Physiology Blockset
 - Blood Gases
 - GOLEM
 - Kidney
 - Product Page (Web)
- * Signal Processing Blockset
- * Support and Web Services

Title: How Simulink Works (Using Simulink)

Physiology blockset - SIMA, Cardiovascular model

SIMA - Cardiovascular model

SIMA CM is an implementation of cardiovascular model according to description in [Applied Mathematical Models in Human physiology](#), Chapter 6. It's assembled from building blocks found in SIMA simulator part of library.

Current version in Physiolibrary serves as a demonstration and provides three outputs:

- * Atrial pressure [mmHg]
- * Left ventricular pressure [mmHg]
- * Left ventricular volume [ml]

Number of outputs can be changed easily.

Každý nabízený simulační blok musí být dobře dokumentován



MODELLING THE RESPIRATORY CONTROL SYSTEM IN HUMAN SUBJECTS FOR EXERCISE CONDITIONS

In our model, blood pH is calculated using a model developed by researchers at the University of Prague (Kofranek et al., 2007). The calculation can be regarded as a computer implementation of the Siggaard-Andersen nomogram which was explained in



A THESIS

SUBMITTED TO THE FACULTY OF ENGINEERING

OF THE UNIVERSITY OF GLASGOW

FOR THE DEGREE OF

DOCTOR OF PHILOSOPHY

**Knihovna
"simulačních čipů"
PHYSIOLIBRARY
je využívána**

Window Help

X

Title: ref. GOLEM

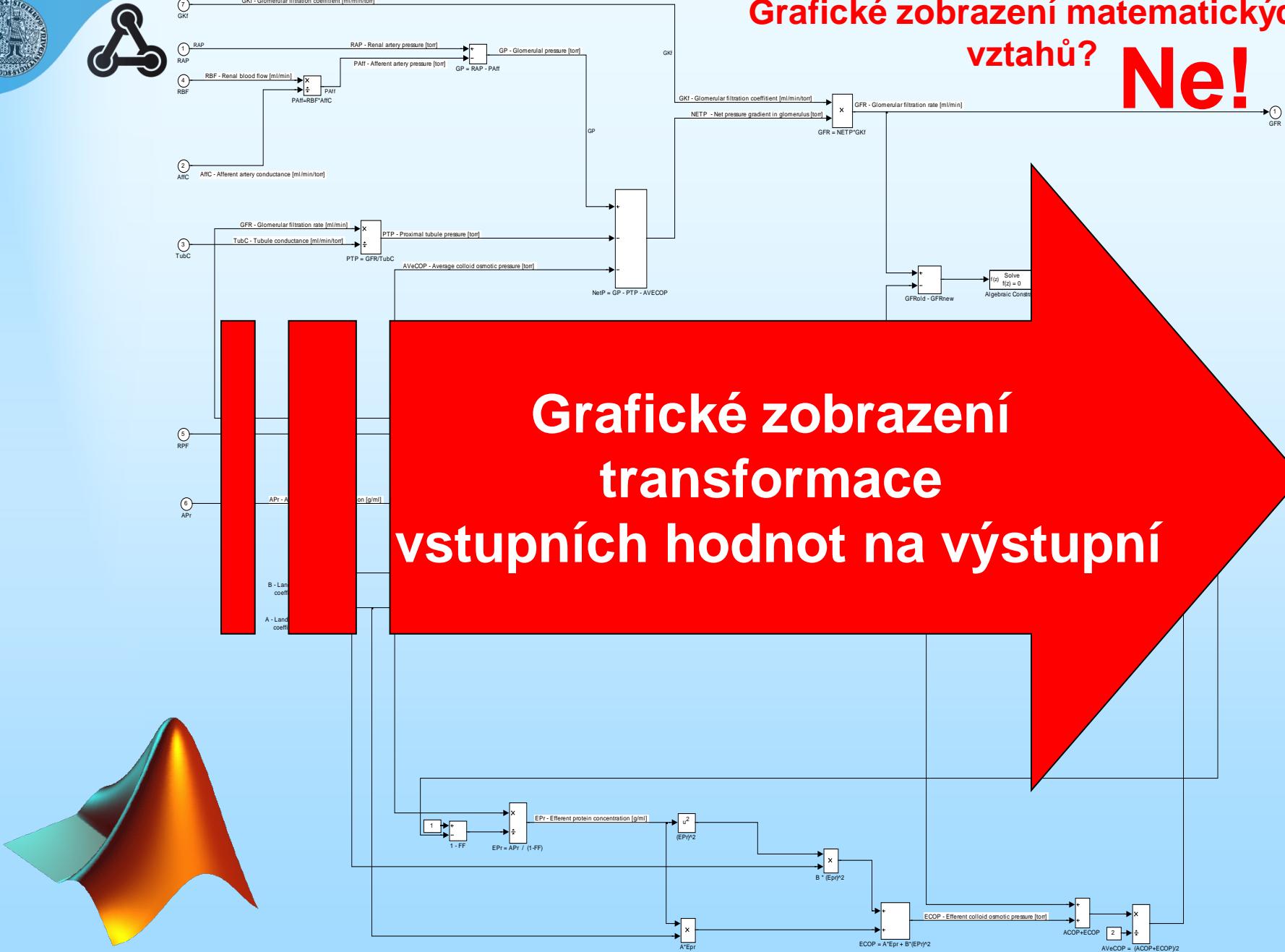
Help for Physiological Blockset - (blocks references)

Golem library - Alphabetical List of blocks references

GOLEM Library

- Acid Base Metabolic Balance
- Blood Acid Base Balance
- Blood Glucose Control
- Body Fluid Volume Balance
- Ca & Mg Balance
- Cardiovascular Block
- Chloride Balance
- Controller of Renal Function
- Diuresis & Urine Osmolarity
- Interstitial Pressure and Lymph Flow Rate
- Na & K Balance
- O₂ & CO₂ Exchange

Součástí knihovny
PHYSIOLIBRARY je i
model, který byl
podkladem výukového
simulátoru **GOLEM**

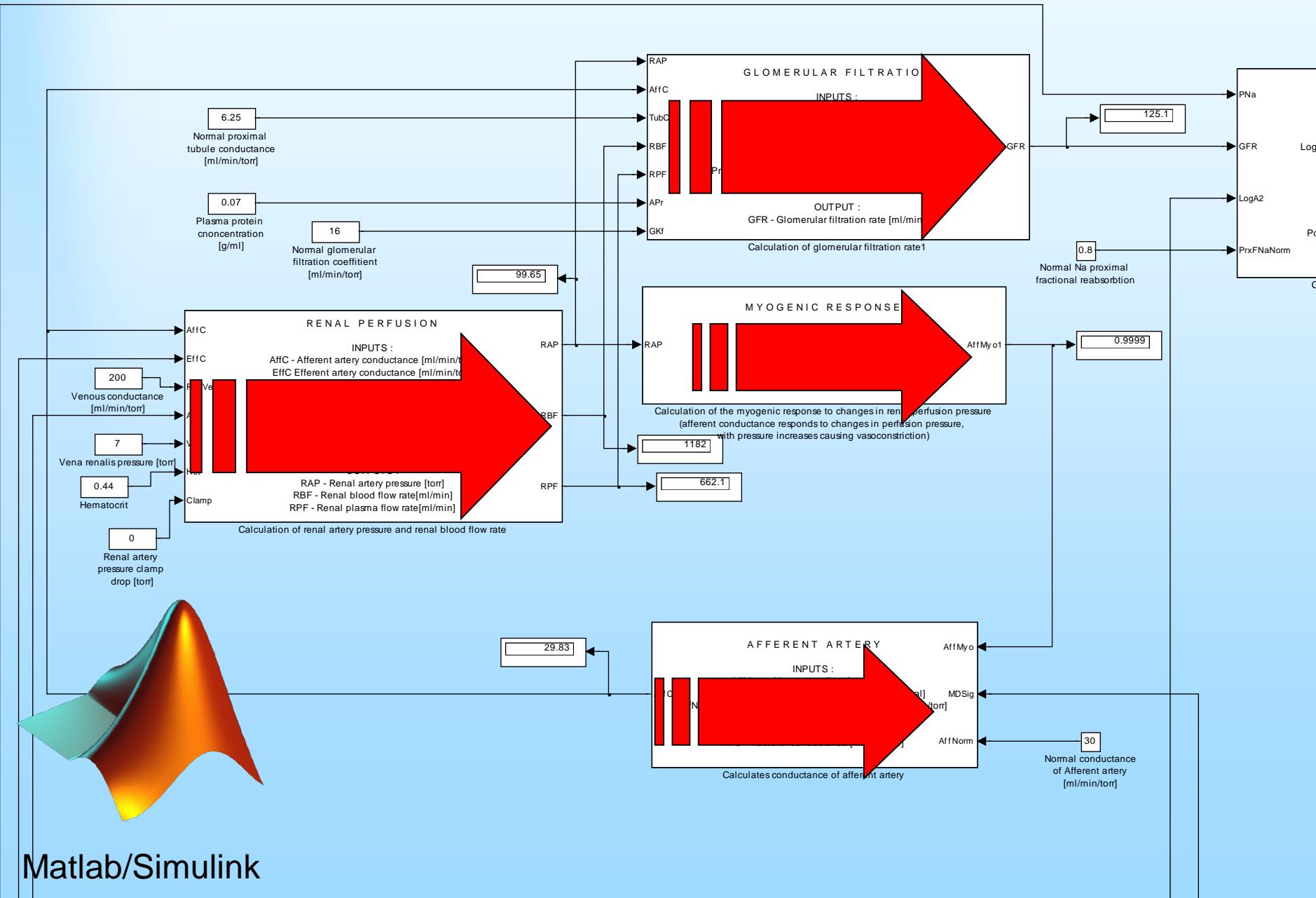


Matlab/Simulink

Grafické zobrazení matematických vztahů? Ne!



Softwarové nástroje pro tvorbu modelů



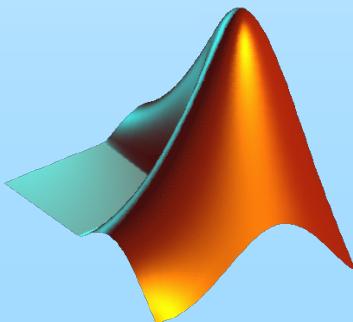


Kauzální modelovací nástroje

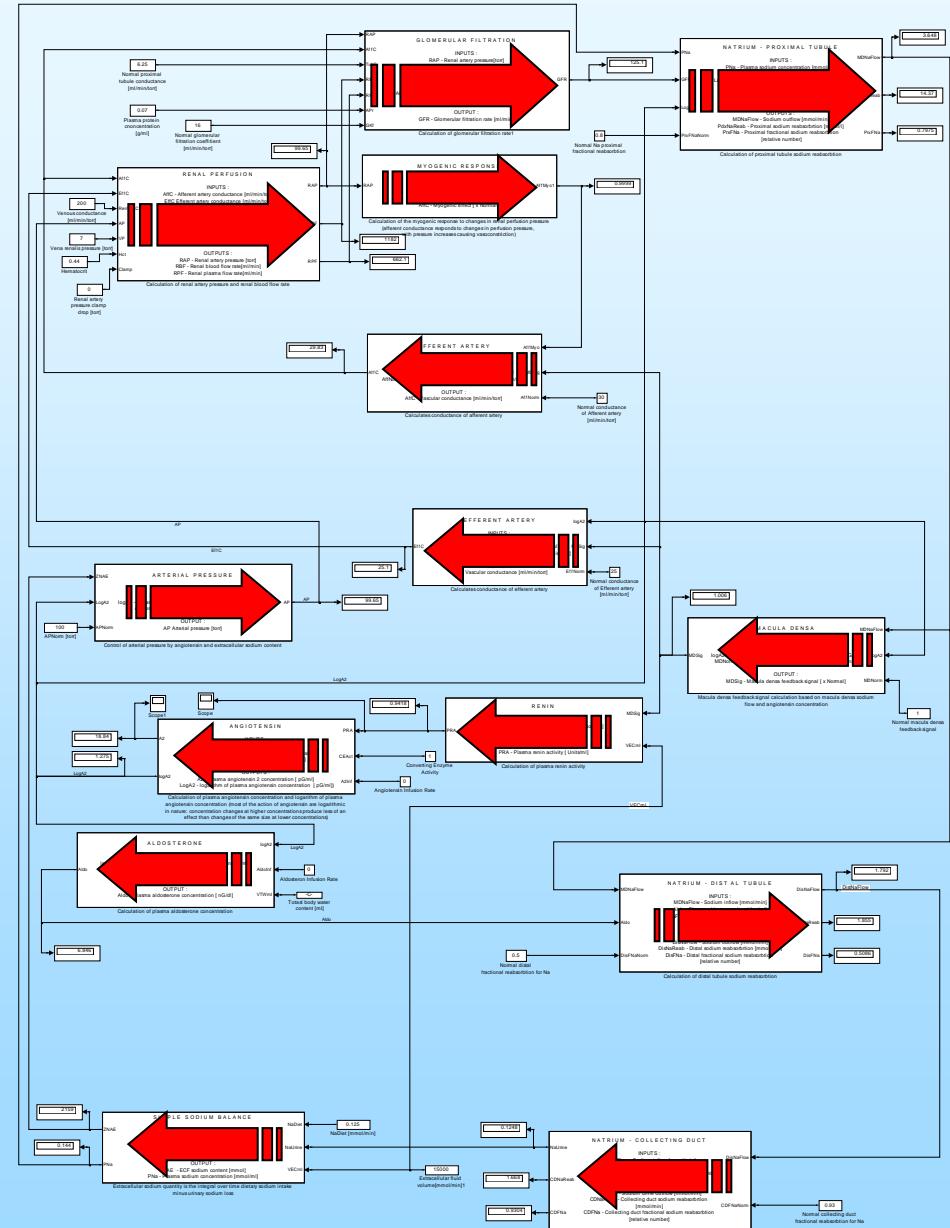
Je jednoznačně
definován
postup výpočtu

Kauzální modelování

Model v Simulinku
vyjadřuje spíše
způsob výpočtu než
strukturu modelované
reality



Matlab/Simulink





Modelica pro simulace



Guytonův model po 40 letech

Model HumMod (QHP - Quantitative Human Physiology)

The screenshot shows a Mozilla Firefox browser window with the following details:

- Title Bar:** HumMod | Welcome - Mozilla Firefox
- Menu Bar:** Soubor, Úpravy, Zobrazení, Historie, Záložky, Nástroje, Nápověda
- Address Bar:** http://hummod.org/
- Toolbar:** DAEMON Tools Lite, Astroburn Lite, Ostrava (Mosnov) +6 °C, Radio player
- Bottom Bar:** Edit, Post to Blog

The main content area displays the following information:

HumMod (hyoo—mod)—noun:

a better understanding of the human body; a mathematical model of integrated human physiology; a framework for developing physiology models and expanding current modeling practices.

Common Usage:

- Send astronauts to Mars
- Understand burn victim physiology
- Articulate clinical scenarios

Get Started

At the bottom of the page, there is a copyright notice:

©2010 — The University of Mississippi Medical Center
Supported by: NSF EPSCoR, NASA, and NHLBI
Icons By: [tiny icons](#)

At the very bottom left, it says "Hotovo".

***Model HumMod (Quantitative Human Physiology) je patrně dnes
nejrozsáhlejším modelem integrativní fyziologie člověka***

The screenshot shows the homepage of the HumMod website. At the top, there is a banner with the text "Model HumMod (Quantitative Human Physiology) je patrně dnes nejrozsáhlejším modelem integrativní fyziologie člověka". Below the banner is a standard browser header bar with icons for DAEMON Tools, DAEMON Tools Lite, Astroburn Lite, weather information for Ostrava (Mosnov) at +6 °C, and a radio player. The main content area has a light blue background. It features a large, bold title "HumMod" followed by its definition "(hyoo—mod)—noun:". Below the title is a descriptive paragraph: "a better understanding of the human body; a mathematical model of integrated human physiology; a framework for developing physiology models and expanding current modeling practices." To the left of this text is a section titled "Common Usage:" with three bullet points: "Send astronauts to Mars", "Understand burn victim physiology", and "Articulate clinical scenarios". To the right of the main text is a green rectangular button with the text "Get Started". At the bottom of the page, there is a footer with copyright information: "©2010 — The University of Mississippi Medical Center Supported by: [NSF EPSCoR](#), [NASA](#), and [NIH](#)". There are also links for "Icons By: [DryIcons](#)" and "Hotovo".

HumMod

(hyoo—mod)—noun:

a better understanding of the human body; a mathematical model of integrated human physiology; a framework for developing physiology models and expanding current modeling practices.

Common Usage:

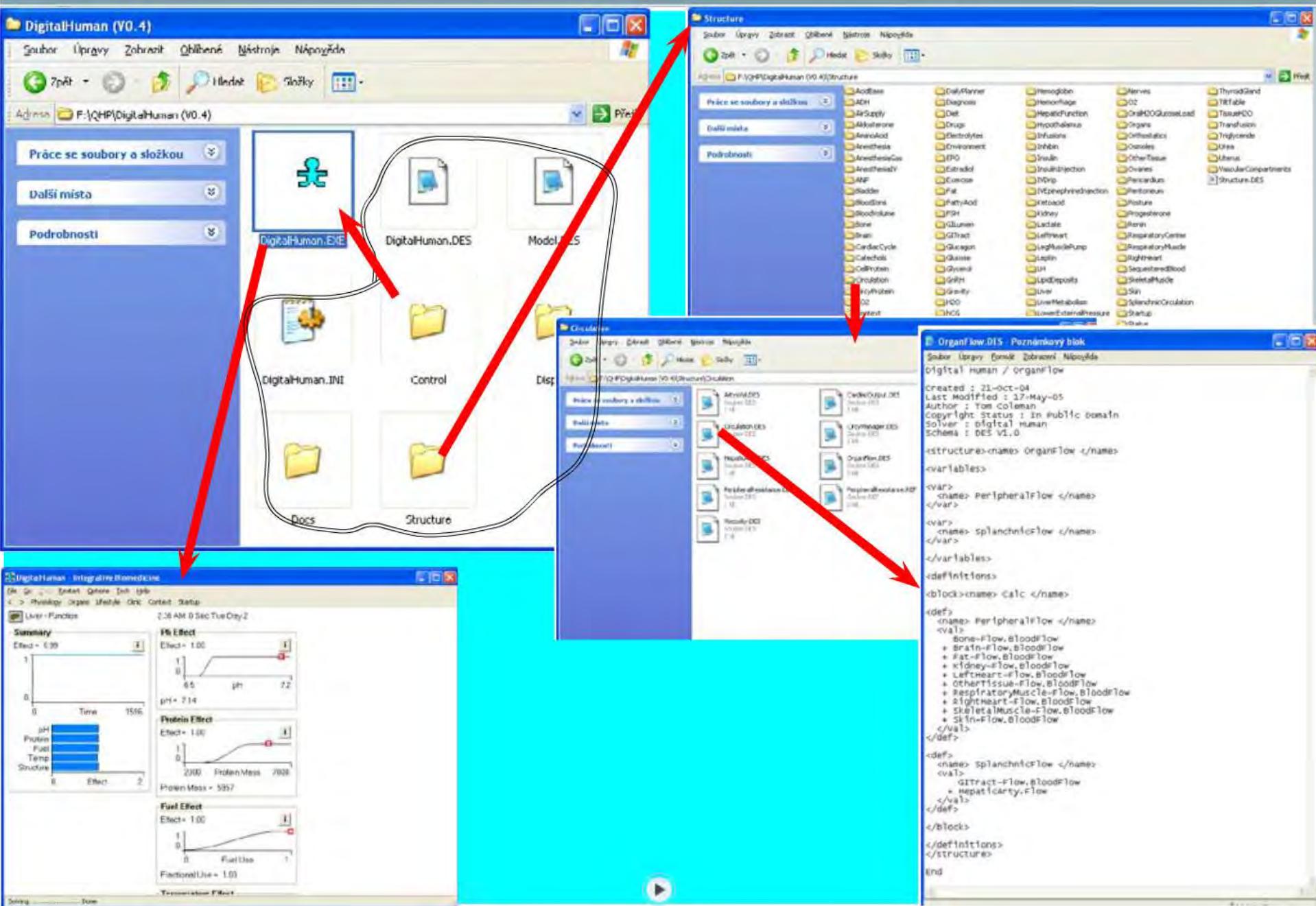
- Send astronauts to [Mars](#)
- Understand burn victim physiology
- Articulate clinical scenarios

[Get Started](#)

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Supported by: [NSF EPSCoR](#), [NASA](#), and [NIH](#)
Icons By: [DryIcons](#)

Hotovo

Model HumMod (QHP - Quantitative Human Physiology)



Model HumMod (QHP - Quantitative Human Physiology)

« Hummod » version_1.5

Structure » BloodVolume »

BloodVolume - Notepad

File Edit Format View Help

```
<?include [EPO]Delay.DES ?>
<Structure><name> BloodVolume </name>
<definitions>
<block><name> Params </name>
  <call> RBCClearance.Parms </call>
  <call> [EPO]Delay.Parms </call>
</block>
<block><name> calcvol </name>
  <call> RBCVol.calcvol </call>
  <call> RBCSolids.Calcvol </call>
  <call> RBCH2O.calcvol </call>
  <call> PlasmaVol.calcvol </call>
  <call> BloodVol.calcvol </call>
</block>
<block><name> calcv0 </name>
  <call> BloodVol.calcv0 </call>
</block>
<block><name> Dervs </name>
  <call> RBCSecretion.Dervs </call>
  <call> RBCClearance.Dervs </call>
  <call> [EPO]Delay.Dervs </call>
</block>
```

Organize

Favorite Links

- Documents
- Pictures
- Music
- More »

Folders

- Computers
- OS (C:)
- KOFRA
- HumMod
- version_1.5
- Structure
- BloodVolume

Matematické vztahy
ve zdrojovém textu
modelu



Ln 1, Col 1



HumMod

Version 1.2.1

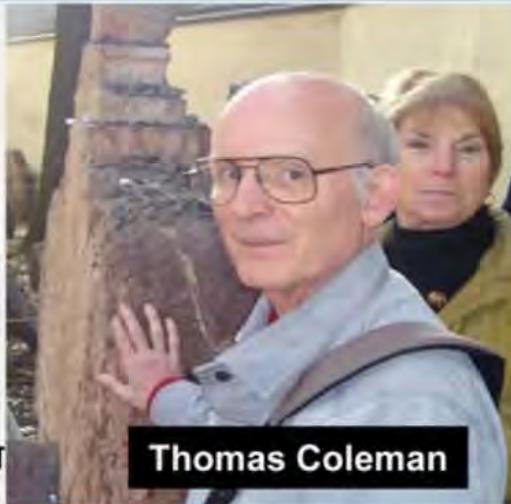
Build : Apr 20 2010

University of Mississippi Medical Center
Jackson, MS 39216 USA

Dr. Tom Coleman
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www.hummod.org



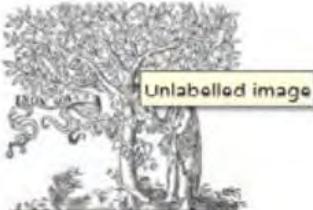
Thomas Coleman

*Thomas Coleman -
spoluautor původního
Guytonova modelu
a nyní hlavní architekt
modelu HumMod na
návštěvě v Praze*

Type:	File Folder (.5)
Location:	D:\Hummod
Size:	18,1 MB (19 003 011 bytes)
Size on disk:	59,6 MB (62 525 440 bytes)
Contains:	14 115 Files, 9 674 Folders

SAPHIR: a physiome core model of body homeostasis and blood pressure

By S. RANDALL TURK with SIMULINK has recently been published (Kofranek & Rusz 2007),
not to mention the current very elaborate and comprehensive Quantitative
Human Physiology model at the University of Mississippi, directly descended
from the early Guyton models by the colleagues of Guyton (Abram *et al.* 2007).
The originality of our core model implementation is our commitment to provide
documentation for each basic module, interactive modification of any aspect of the
model parameters or equations and, especially, the possibility of contributing new
detailed local sub-modules to complement or replace selected ‘basic modules’.
Crucial to this approach is an underlying computational toolbox that is
sufficiently robust to handle the wide range of spatial and temporal scales, and
flexible enough to accept sub-modules in a variety of formalisms, including not
only ODE/PDE dynamical systems models but also discrete modelling methods
(e.g. renale),



ELSEVIER

journal homepage: www.intl.elsevierhealth.com/journals/cmpb

Graphical simulation environments for modelling and simulation of integrative physiology

Violeta Mangourova^{a,*}, John Ringwood^a, Bruce Van Vliet^b

^a Dept. of Electronic Engineering, NUI Maynooth, Ireland

^b Biomedical Sciences Division, Memorial University, St. John's, Newfoundland, Canada

ARTICLE IN

Article history:

Received 27 April 2009

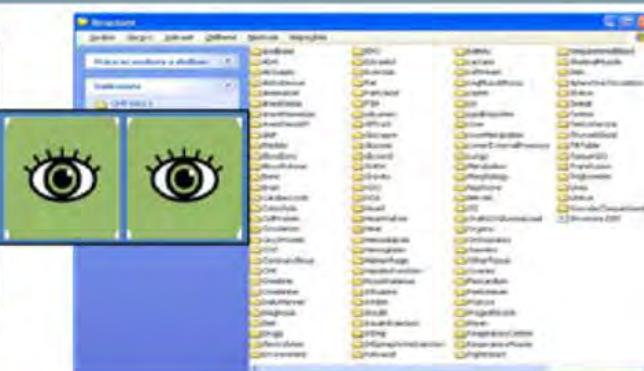
Makes the wider interactions in the model difficult to visualise. Coleman's extension of Guyton's work [5] via the QHP (Quantitative Human Physiology) model was available only in executable form (for teaching purposes) until recently, when an eXtensible Markup Language (XML) version of the model was released [6,7]. This extended model, however, is difficult to read and visualise, since no diagrammatic form exists.

Model HumMod (QHP - Quantitative Human Physiology)

Mozilla Firefox

Soubor Úpravy Zobrazit Historie Záhlby Yahoo! Nástroje Nápověda

CO2, Hemoglobin, AcidBase, ADH, AirSupply, Aldosterone, AminoAcid, if ANESTHESIA then Anesthesia, if not ANESTHESIA then NoAnesthesia, if ANP then DetailedANP, if not ANP then BasicANP, Bladder, BloodVolume, BloodLions, Bone, Brain, CardiacCycle, Catechols, CellProtein, Circulation, CervyProtein, CoronaryStatus, CPR, Creatine, Creature, DailyPlanner, Diagnosis, Diet, Drugs, Electrolytes, Environment, EPO, Estradiol, Exercise, Fat, FattyAcid, FSH, GITLumen, GITTract, Glucagon, Glucose, Glycerol, GnRH, Gravity, H2O, hCG, Heart, HeartValves, Heat, HemoDialysis, Hemorrhage, HepaticFunction, Hypothalamus, Inhibit, Infusions, Insulin, InsulinInjection, IVDrp, IVEpinephrineInjection, Ketoacid, Kidney, Lactate, LeftHeart, LegMusclePump, Leptin, LH, LipidDeposits, Liver, LiverMetabolism, LowerExternalPressure, Lungs, Metabolism, Morphology, Nephrons, Nerves, O2, OralH2OGlucoseLoad, Organs, Orthostatics, Osmoles, OtherTissue, Pericardium, Peritoneum, Posture, Progesterone, Renin, RespiratoryCenter, RespiratoryMuscle, RightHeart, SequesteredBlood, SkeletalMuscle, Skin, SplanchnicC, Testes, Testosterone, ThyroidGland, TktTable, TissueH2O, Transfusion, Triglyceride, VascularCompartments



QHPView

VascularCompartments

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VascularCompartments/VascularCompartments

(ref VascularCompartments.REF) (ref Ventricles.REF)
LeftAtrium, PulmArty, PulmCapys, PulmVeins, RightAtrium, SplanchnicVeins, LeftVentricle, RightVentricle,

[structure: VascularCompartments]
[definitions]
[block: CalcPressure(1)
RightAtrium.CalcPressure
LeftAtrium.CalcPressure
SystemicArtsy.CalcPressure
[block: CalcPressure(2)
SystemicVeins.CalcPressure
SplanchnicVeins.CalcPressure
[block: Dervs]
RightVentricle.Dervs
SystemicVeins.Dervs
SplanchnicVeins.Dervs
RightAtrium.Dervs
PulmArty.Dervs
PulmCapys.Dervs
PulmVeins.Dervs
LeftVentricle.Valves LeftVentricle RightVentricle-

VascularCompartments/SplanchnicVeins.DES

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VascularCompartments.REF (ref Ventricles.REF)

structure: SplanchnicVeins]
[variables]
[var: Inflow]
[var: Outflow]
[parm: V0 = 300.0]
[var: StressedVol]
[var: Pressure]
[var: ExternalPressure]
[parm: Compliance = 62.5]
[parm: Conductance = 178]
[equations]
der(Change) = Vol
initial equation: Vol = 300.0
errorLimit: 10.0
[definitions]
[block: CalcPressure]
StressedVol = (Vol - V0) MAX 0.0
ExternalPressure = 0.0
Pressure = (StressedVol / Compliance) + ExternalPressure
[block: Dervs]
DxMax = if Conductance > 0.0 then 0.5 * Compliance / Conductance else INFINITE
Inflow = OrganFlow.SplanchnicFlow
Outflow = Conductance * (Pressure - RightAtrium.Pressure)
Change = Inflow - Outflow

Equations



Physiolibrary x www.physiolibrary.org

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Physiolibrary

Physiolibrary is a free open-source Modelica library designed for modeling human physiology. This library contains basic physical laws governing human physiology, usable for cardiovascular circulation, metabolic processes, nutrient distribution, thermoregulation, gases transport, electrolyte regulation, water distribution, hormonal regulation and pharmacological regulation.

Library description

Our laboratory have a long tradition building physiological libraries, starting with Physiolibrary in Matlab/Simulink environment. The origin of this Modelica Physiolibrary was in the first version of our HumMod Golem Edition model implementation, where it was called HumMod Library. As the successors of Guyton's Medical Physiology School write, the original HumMod model is "The best, most complete, mathematical model of human physiology ever created". In cooperation with this group we are now developing together the new complex integrative model of physiology called [Physiomodel](#) based on Physiolibrary and HumMod.

We are also developing many types of smaller physiological models for use in medical education, so it was essential to separate this library from our Modelica model implementations. Our Physiolibrary contains only carefully-chosen elementary physiological laws, which are the basis of more complex physiological processes. For example from only three type of blocks (ChemicalReaction, Substance and MolarConservationMass) it is possible to compose the allosteric transitions or the Michaelis-Menten equation.

Library contains also the icons for higher level (HumMod) subsystem implementations:

HumMod-hummod-s...z... Zobrazit všechny stažené soubory...



Physiomodel

Model of physiology in Modelica based on [HumMod](#) and [Physiolibrary](#)

.zip

```
graph TD; A[nutrientsAndMetabolites] --> B[heat]; A --> C[water]; A --> D[proteins]; A --> E[gases]; A --> F[status]; B --> G[cardioVasculature]; B --> H[nerves]; C --> I[hormones]; C --> J[electrolytes]; E --> K[pH]; E --> L[O2]; F --> M[setup];
```

Physiomodel maintained by [physiology](#)

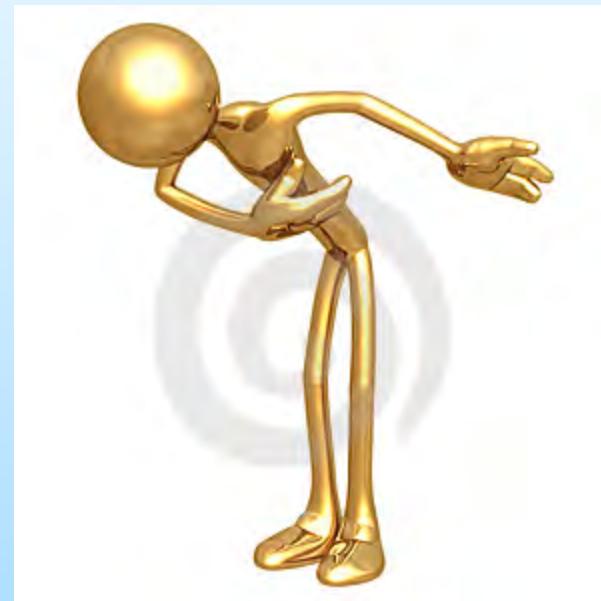
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HumMod-hummod-s... Zobrazit všechny stažené soubory...

Modelica pro simulace



Děkuji za pozornost



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Modelica pro simulace