

Basic Tracer Kinetic Concepts

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Steady state of the system

- i.e. the physiologic parameter is constant during the measurement
- Examples: flow (ml/s), perfusion (ml/g/s), CMRO₂ (mmol/g/s), glucose uptake (mmol/g/s)
- Consider: duration of the measurement in relation to the a spontaneous change of the parameter or timing of a perturbation of the parameter

Steady state of the system

- Exceptions: the physiologic parameter oscillates relative fast compared to the duration of the measurement
- Note: steady state not necessary implies that fluxes or concentration is constant in time

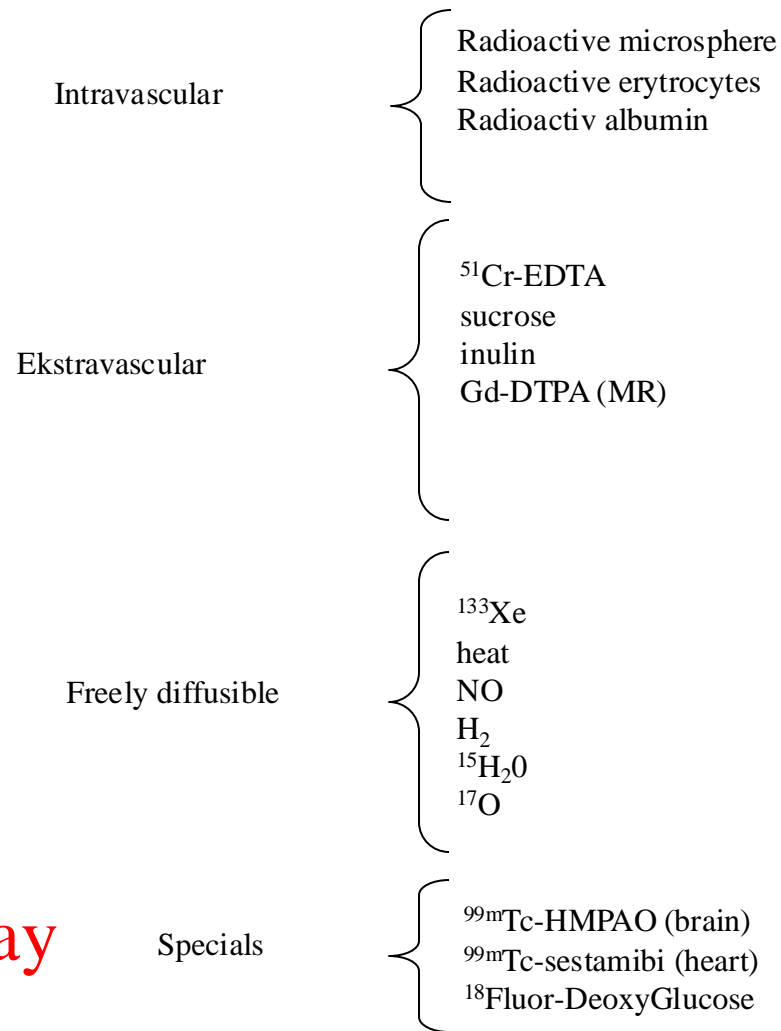


Tracers and indicators

- Tracers: labelled substances, behaves physically and chemically like the modersubstances;
- e.g. H_2^{15}O , $^{17}\text{O}_2$, ^{57}Co -vitB12, ^{131}I -thyroxin
- Or behaves nearly like the modersubstance
- e.g. ^{18}F FDG , ^{125}I -albumin, ^{131}I -insulin
- Indicators: not necessary related to a modersubstance
- e.g. contrast agents – x-rays – SPECT ($^{99\text{m}}\text{Tc}$ -HMPAO, $^{99\text{m}}\text{Tc}$ -sestamibi), - MRI (Gd-DTPA, Mn-DPDP)
- Law of conservation: mass balance
- **Note: tracers can be intravascular, extracellular, free difussible, bound to a receptor or behave in a more specific way**

Should not disturb the system we are studying

•Note:
tracers can be
intravascular,
extracellular,
free diffusible,
or behave in a
more specific way



Linearity of a system



**$RF(t)$: response function or more correctly
: The impulse response function**

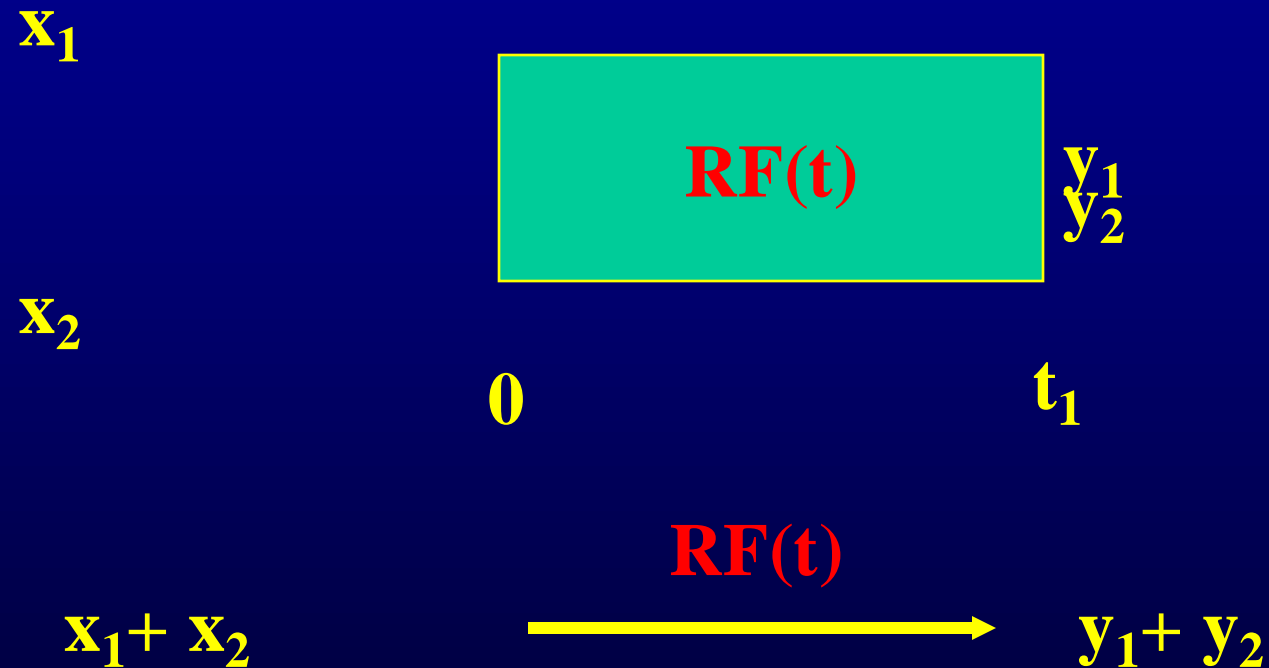
Linearity of a system



scaling



Linearity of a system



Principle of superposition

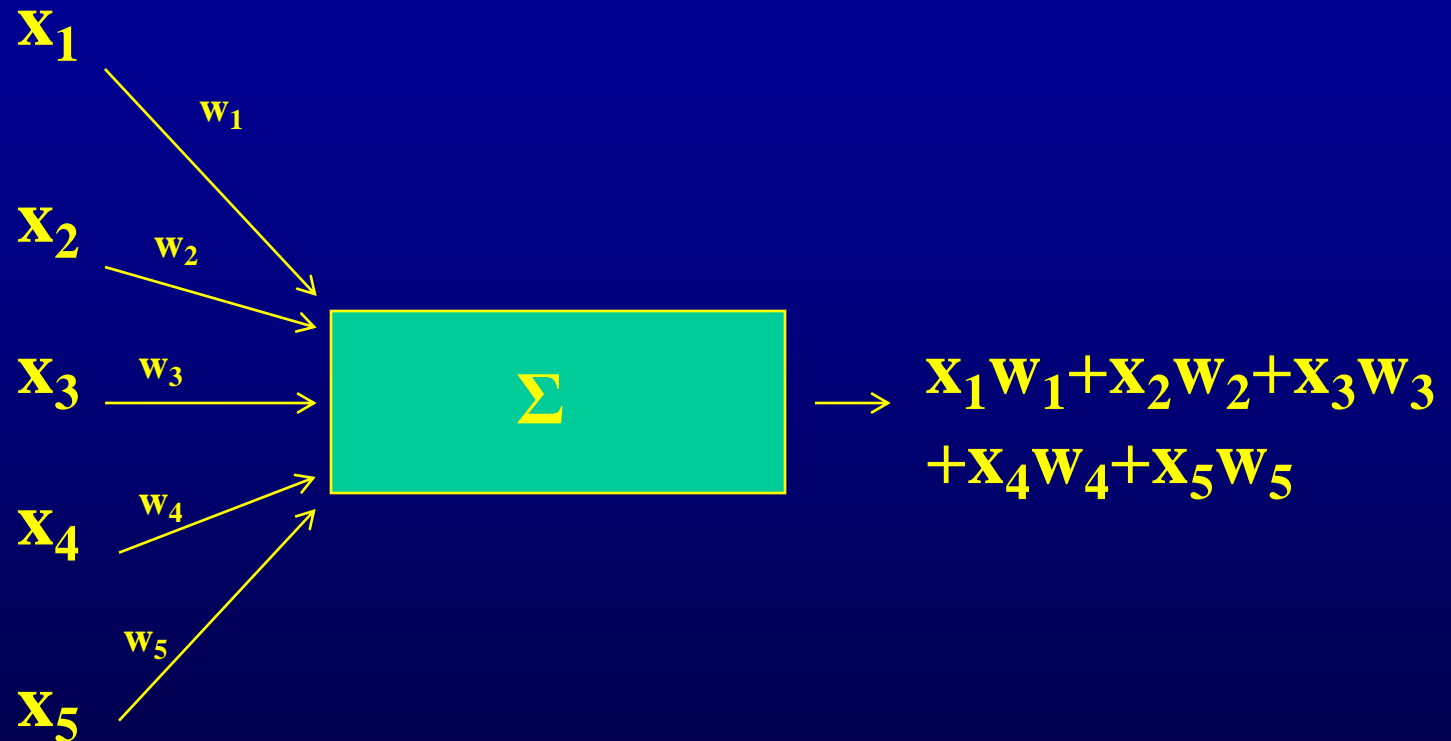
Examples

$$3 + 4 \frac{3}{4}$$

x 2

$$6 \frac{14}{8}$$

Examples



Examples

$$\log_3 3 + \log_4 4$$

log

$$\log_3 \log_4 (3+4)$$

$$\log_3 3 + \log_4 4 \neq \log_3 (3+4)$$

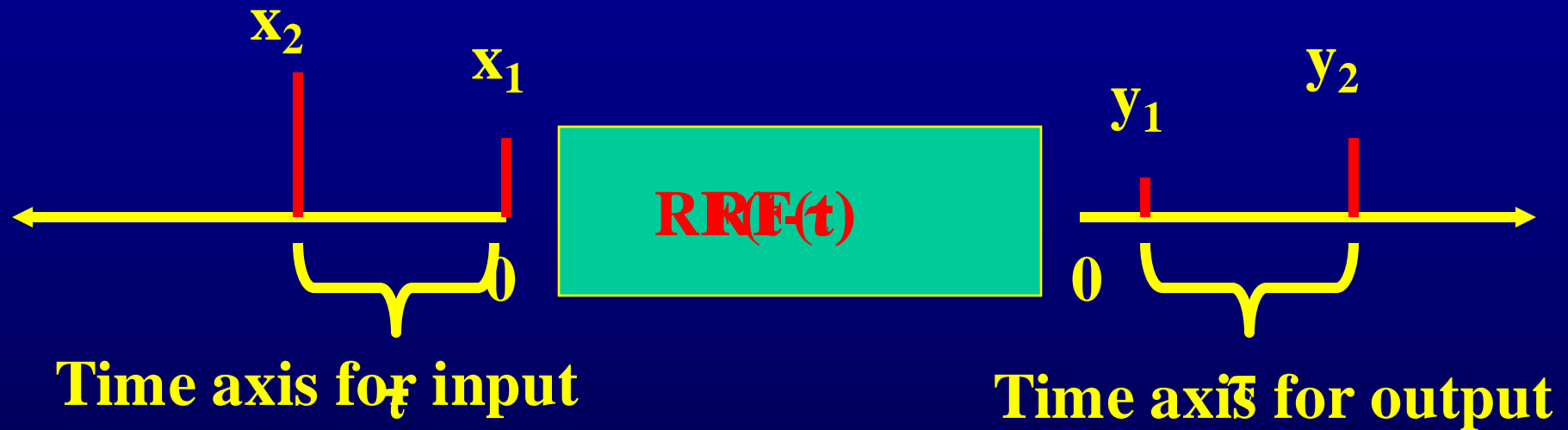
Linearity of a system

$$\mathbf{a} \mathbf{x}_1 + \mathbf{b} \mathbf{x}_2 \xrightarrow{\text{RF}(t)} \mathbf{a} \mathbf{y}_1 + \mathbf{b} \mathbf{y}_2$$

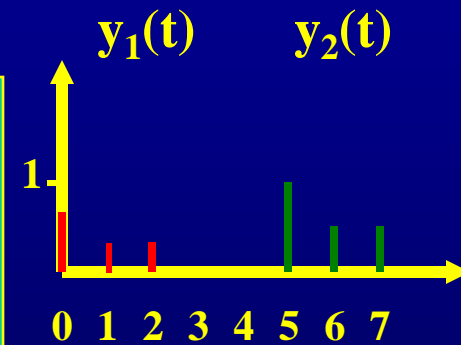
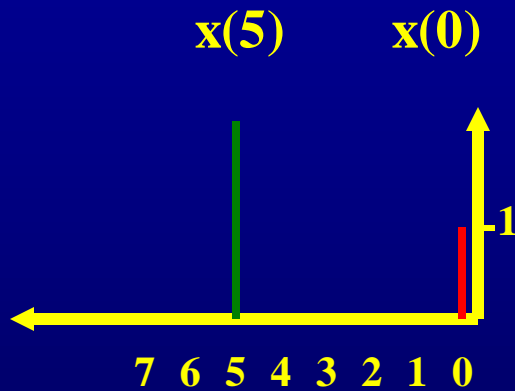
Time invariance of a system



Specification of time



Example



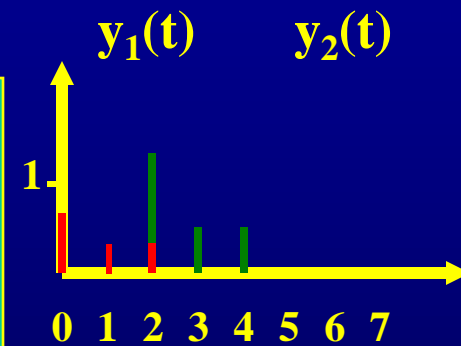
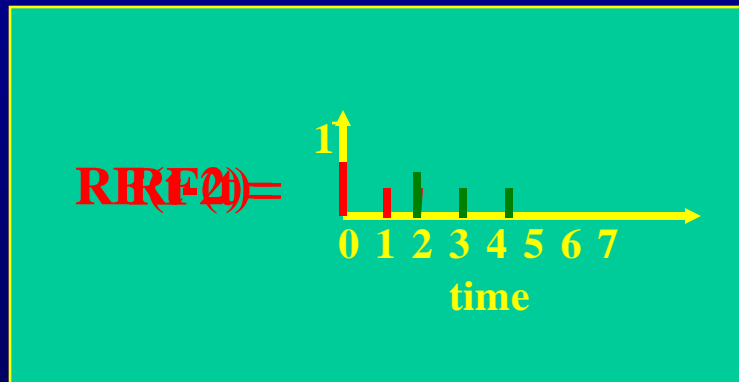
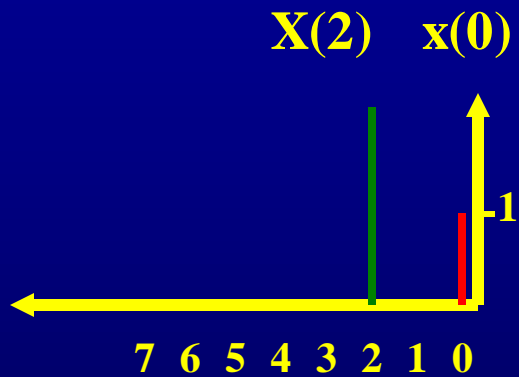
$$y_1(t) = x(0) \cdot RF(t)$$

$$y_2(t) = x(5) \cdot RF(t)$$

$$y_2(t) = x(5) \cdot RF(t-5)$$

Does not work !!!

Example



$$y_1(t) = x(0) \cdot RF(t)$$

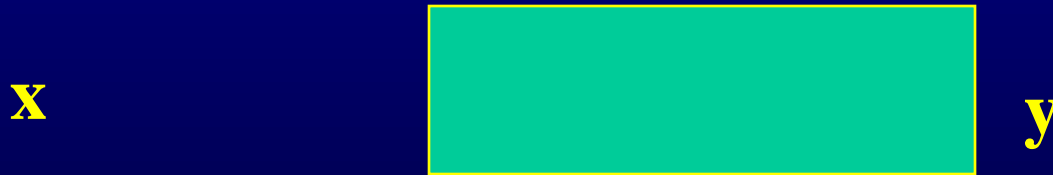
$$y_2(t) = x(2) \cdot RF(t)$$

$$y_2(t) = x(2) \cdot RF(t-2)$$

Does not work !!!

Causality of a system

Output is only observed after an input has enter the system



Causality of a system

Output is only observed after an input has enter the system



Can a biological system
behave like such a system?
Describe in words how a biological
system could interact with a
instantaneous tracer input
!

Linearity of a imaging system?

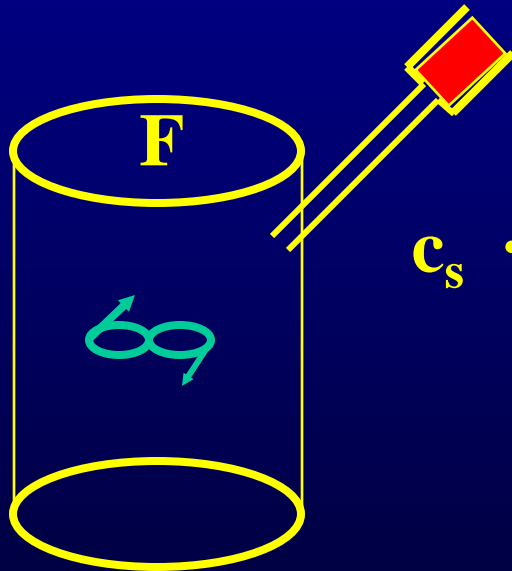
!

Break

Indicator-dilution methods

Constant Infusion (Stewart principle)

The aim : to measure the flow of an organ or a vessel or a pipeline



$$[F] = \text{ml/s}$$

$$c_s \cdot F_s = j_{\text{in}} \quad \text{flux !!!!}$$

$$[c_s] = \text{mmol/ml}$$

$$[F_s] = \text{ml/s}$$

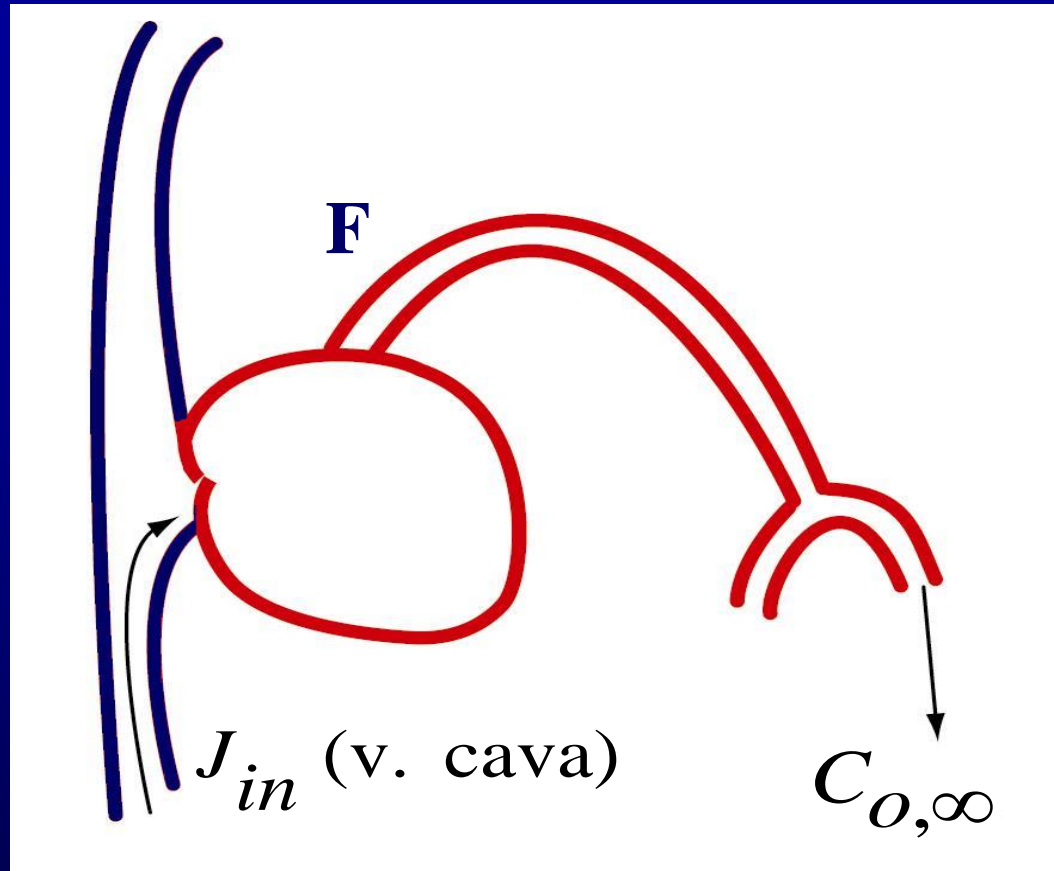
$$[j_{\text{in}}] = \text{mmol/s}$$

$$\mathbf{c}_s \cdot \mathbf{F}_s = (\mathbf{F} + \mathbf{F}_s) \cdot \mathbf{c}_\infty$$

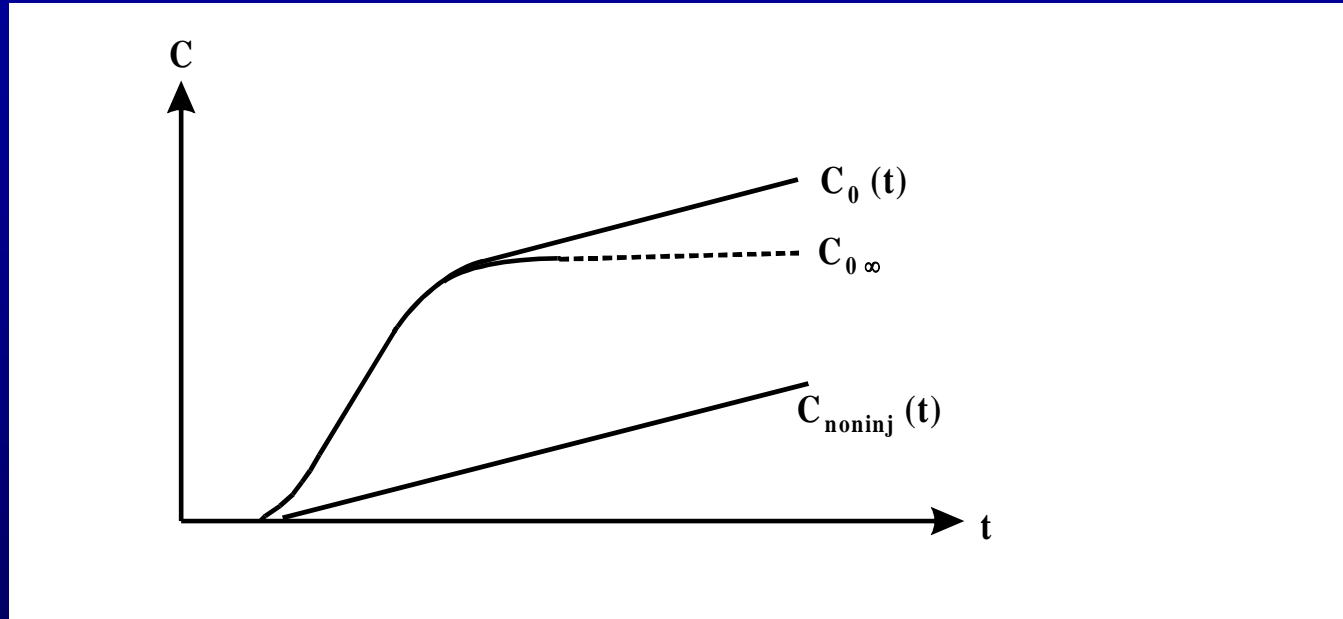
$$\mathbf{F}_s \ll \mathbf{F} \Rightarrow \quad \mathbf{F} = \mathbf{F}_s \cdot \mathbf{c}_s / \mathbf{c}_\infty$$

$$\mathbf{F} = \mathbf{j}_{\text{in}} / \mathbf{c}_\infty$$

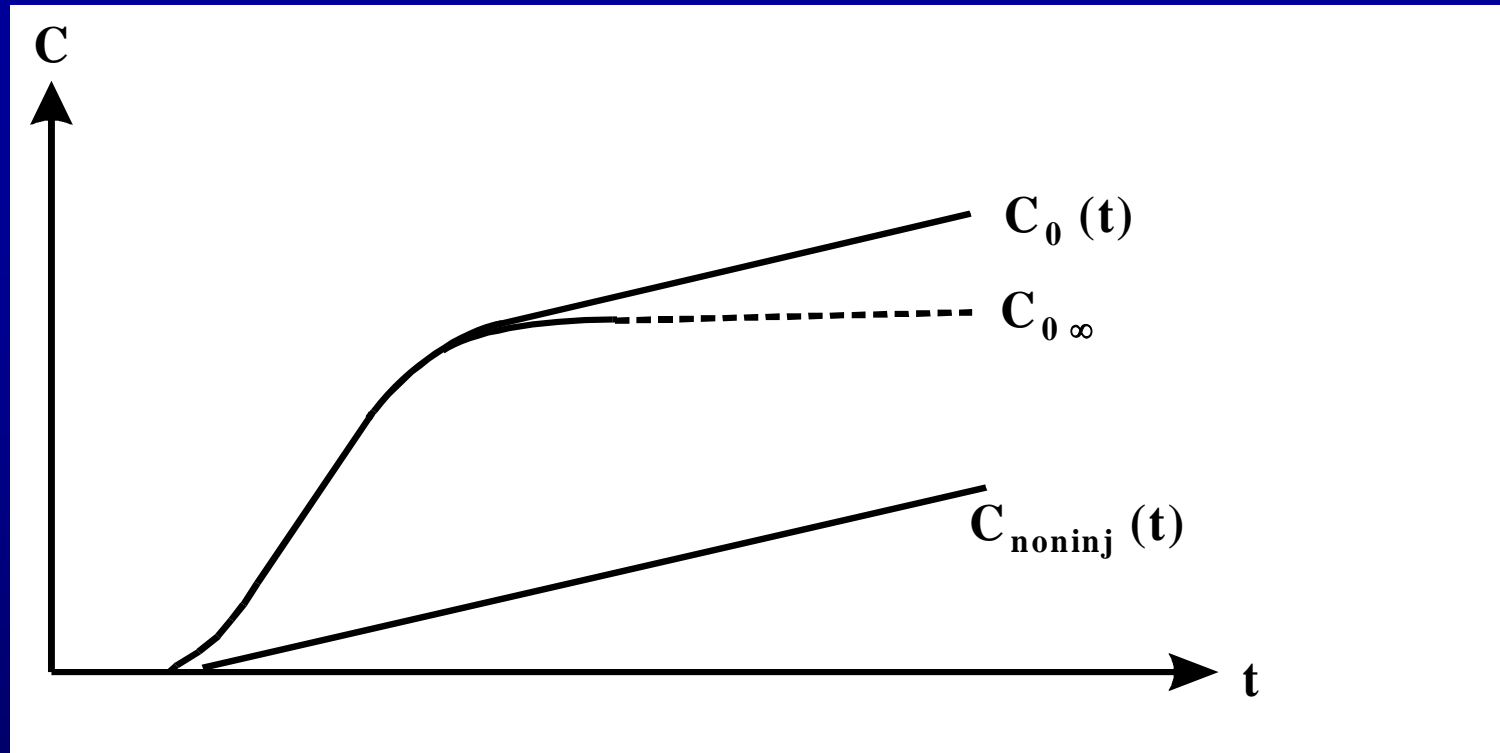
Examples and recirculation



Stewart's principle: Continuously infusion in vena cava, and outlet concentration measurement from a peripheral artery.



Measurement of concentration at the outlet and the "noninj" side.

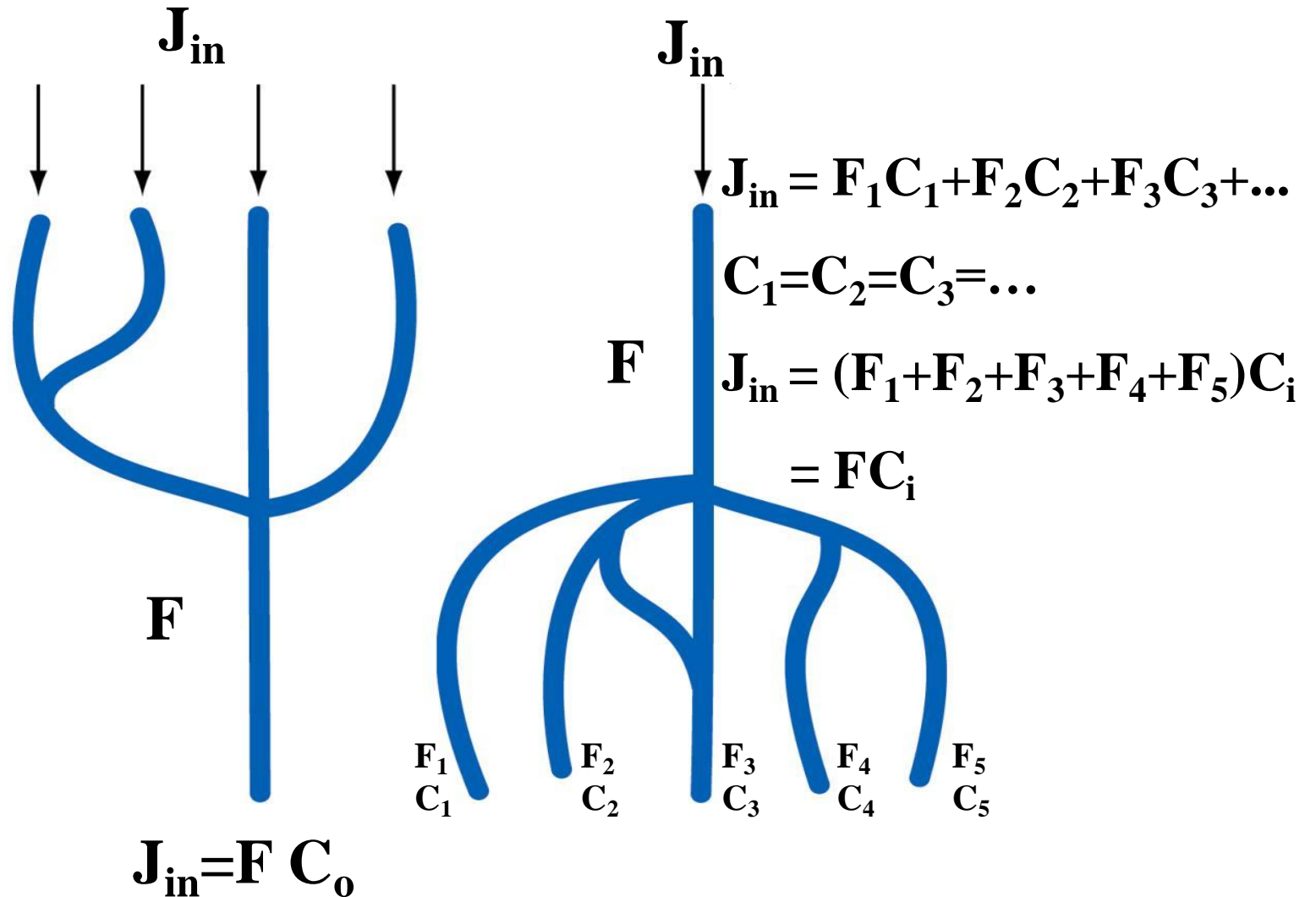


$$J_o = J_{in} + FC_{noninj}(t) \quad \text{D}$$

$$FC_o(t) = J_{in} + FC_{noninj}(t) \quad \text{U}$$

$$F = \frac{J_{in}}{C_o(t) - C_{noninj}(t)}$$

Bolus Fraction principle - Sapirsteins principle

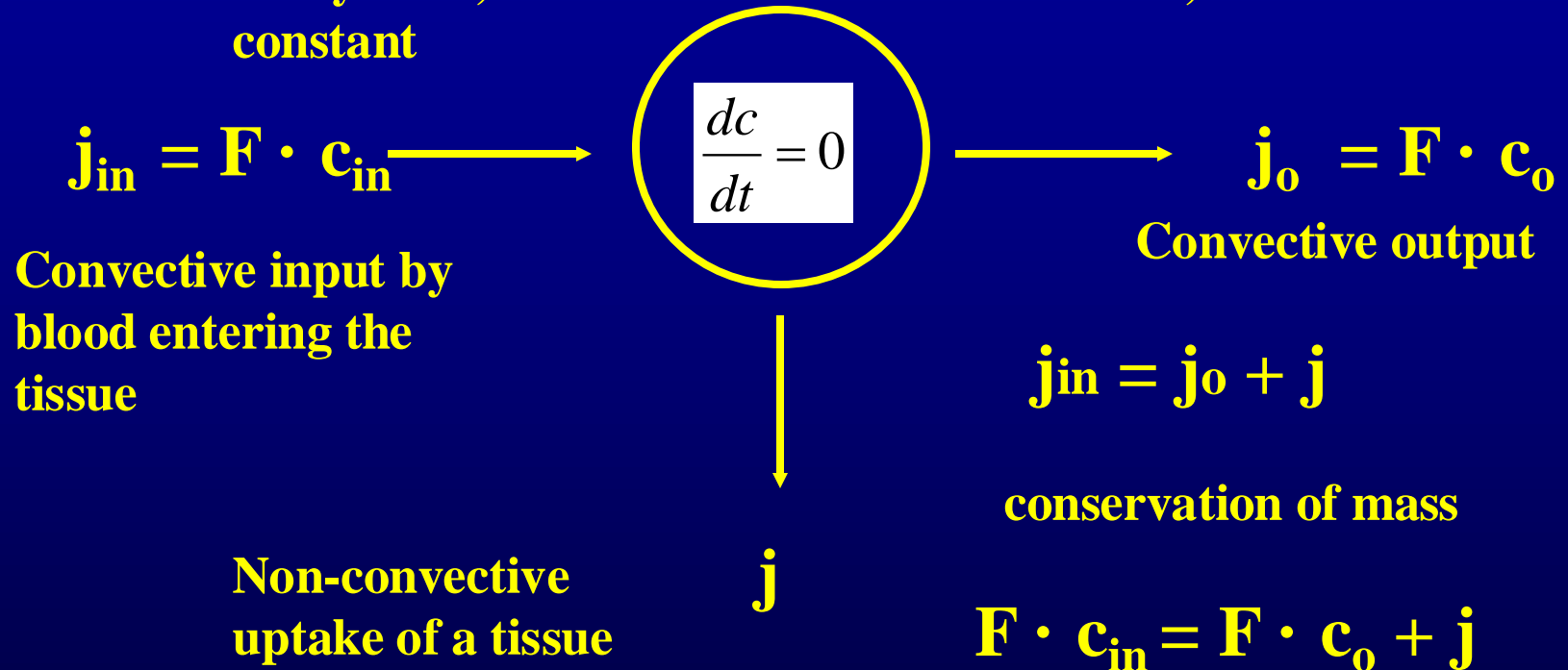


Fick's-principle

The conservation of matter

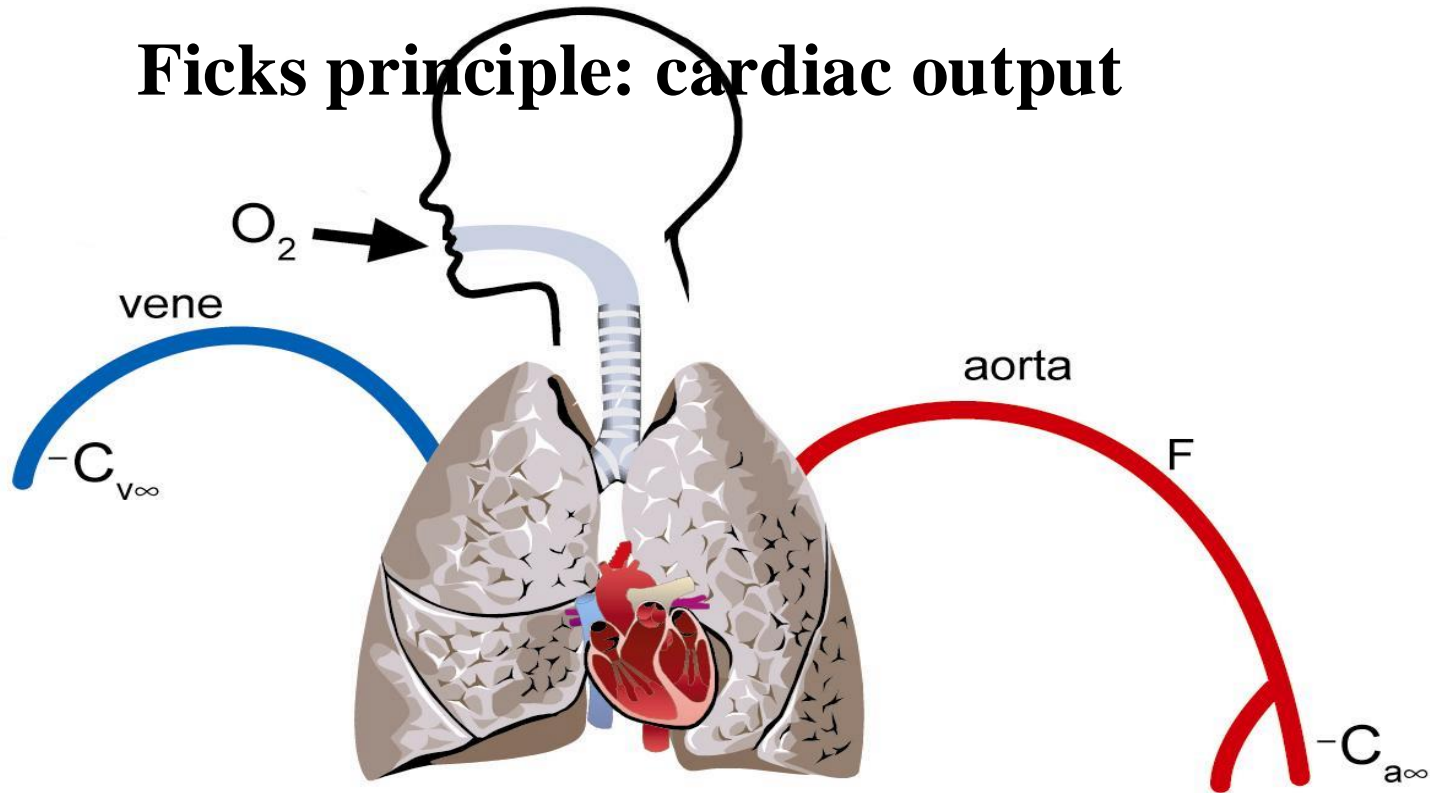
The principle of Fick

Stady state; Concentration here is constant , fluxes are here constant



$$F = \dot{j} / (c_{in} - c_o)$$

Ficks principle: cardiac output



$$J_a = J_{O_2} + J_v$$

$$F \cdot C_{a\infty} = J_{O_2} + F \cdot C_{v\infty} \Rightarrow$$

$$F = \frac{J_{O_2}}{C_{a\infty} - C_{v\infty}}$$

Cerebral metabolic rate of oxygen $CMRO_2$

- Ficks formel

Bloodsample

$$CMRO_2 = 4 \cdot [Hgb] \cdot CBF \cdot (S_a O_2 - S_v O_2)$$

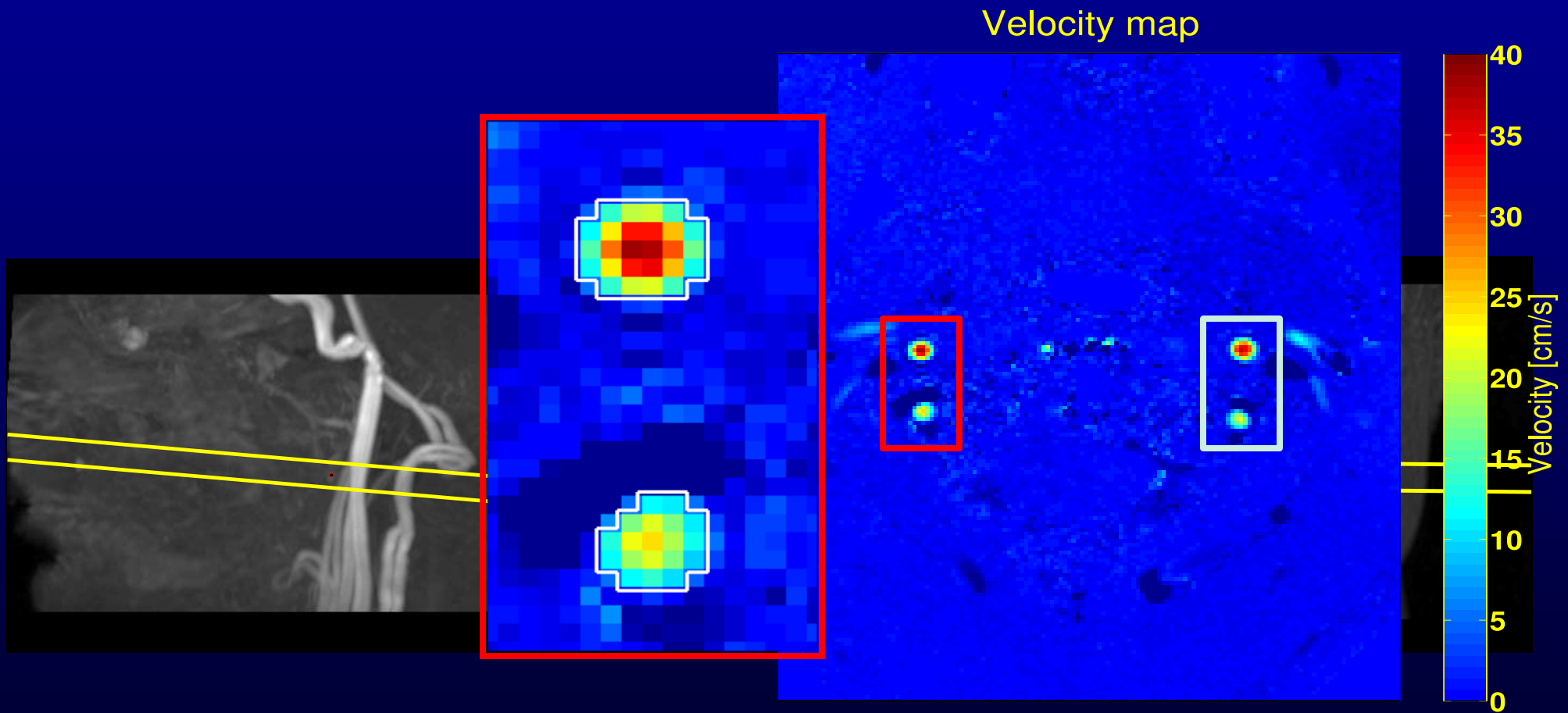
MRI phase contrast mapping

Puls-oximetri (A-cath)

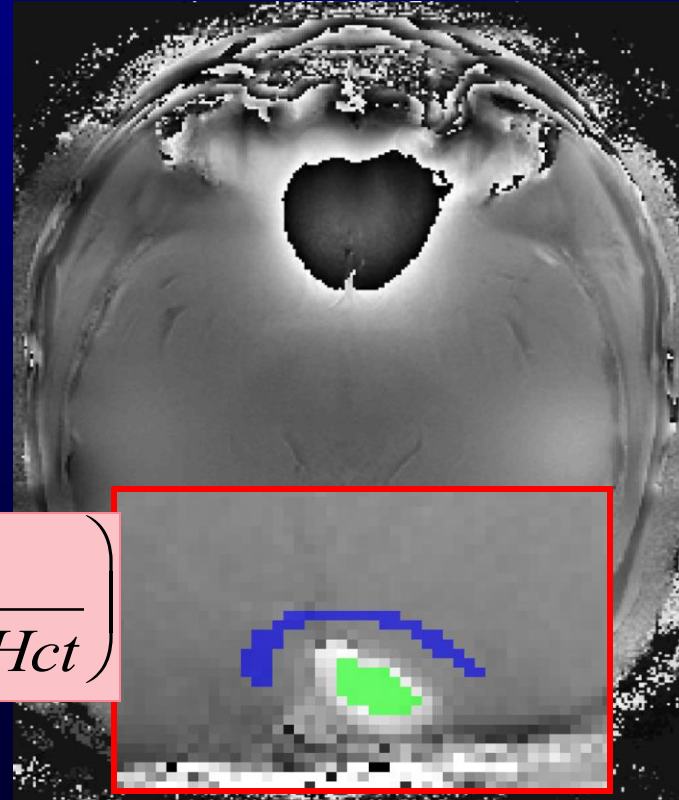
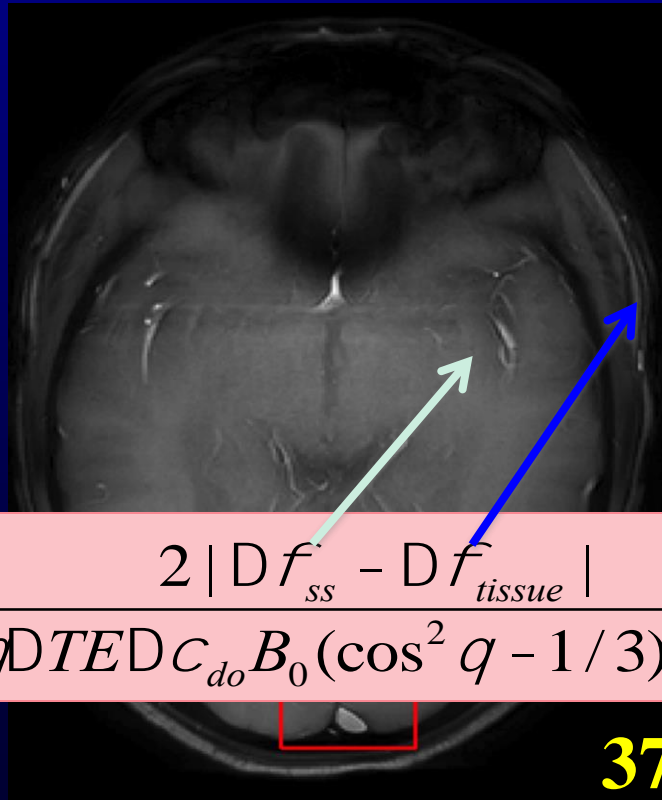
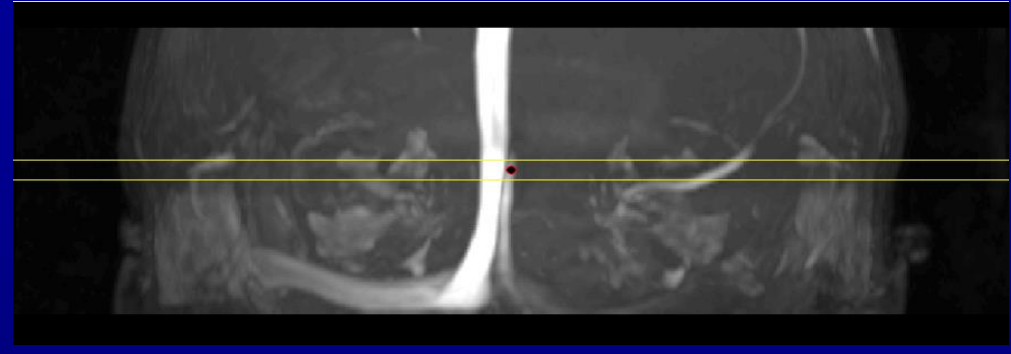
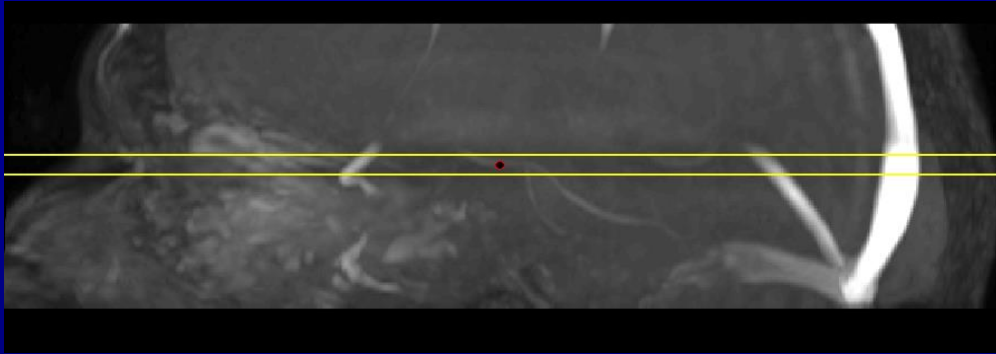
MRI susceptibility-based oximetry from Saggital sinus: venous blood from brain

CBF – Fase kontrast MRI

- Velocity through plane (orthogonal the arteries) and area



Susceptibility based oximetry

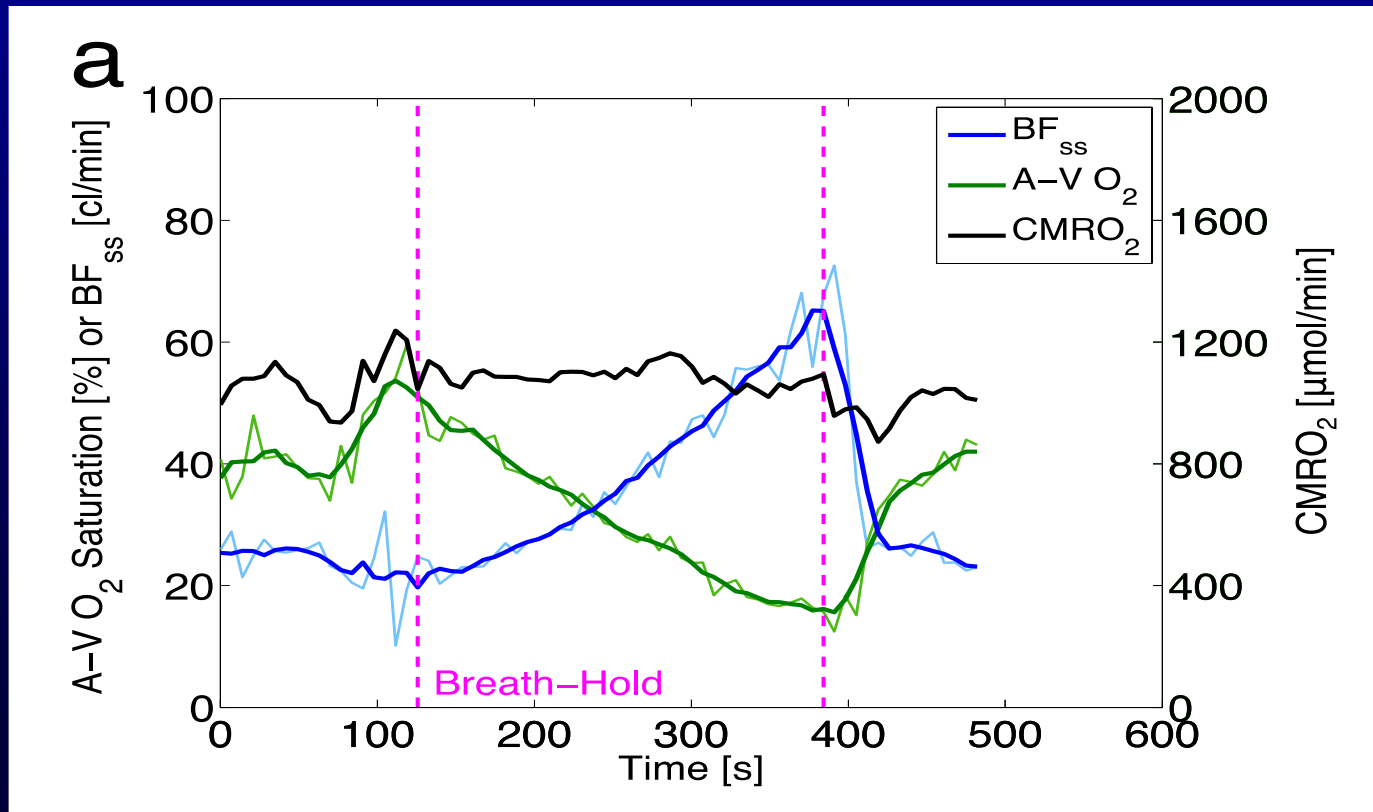


$$SvO_2 = \left(1 - \frac{2 |Df_{ss} - Df_{tissue}|}{gDTEDC_{do} B_0 (\cos^2 q - 1/3) Hct} \right)$$

Breathhold: $CMRO_2$

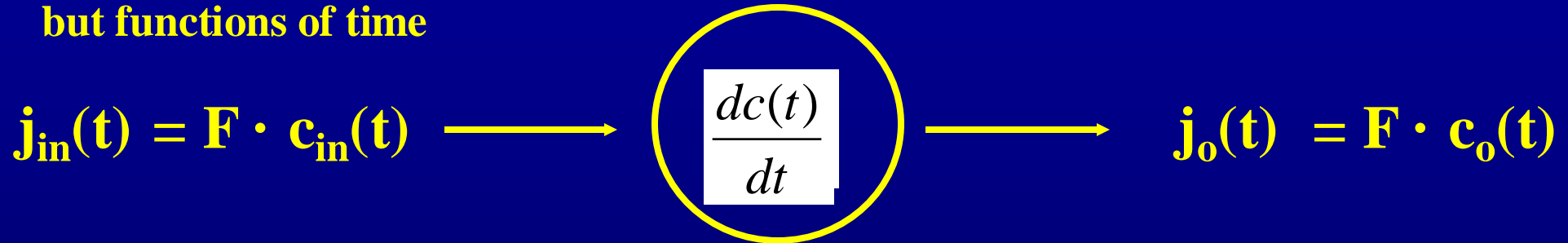
- $CMRO_2 = 4 \cdot [Hgb] \cdot BF_{ss} \cdot (S_aO_2 - S_vO_2)$
- Blood-flow in sagittal sinus (BF_{ss})
- Arteriovenous oxygen-difference ($A-V O_2$)

Vestergaard MB, Larsson HBW. Cerebral metabolism and vascular reactivity during breath-hold and hypoxic challenge in freedivers and healthy controls. *J Cereb Blood Flow Metab* 2017 .



Extending the principle of Fick

The fluxes are not constant, but functions of time The concentration here is not constant



conservation of mass

$$v \frac{dc(t)}{dt} = F \cdot c_{in}(t) - F \cdot c_o(t) - j(t)$$

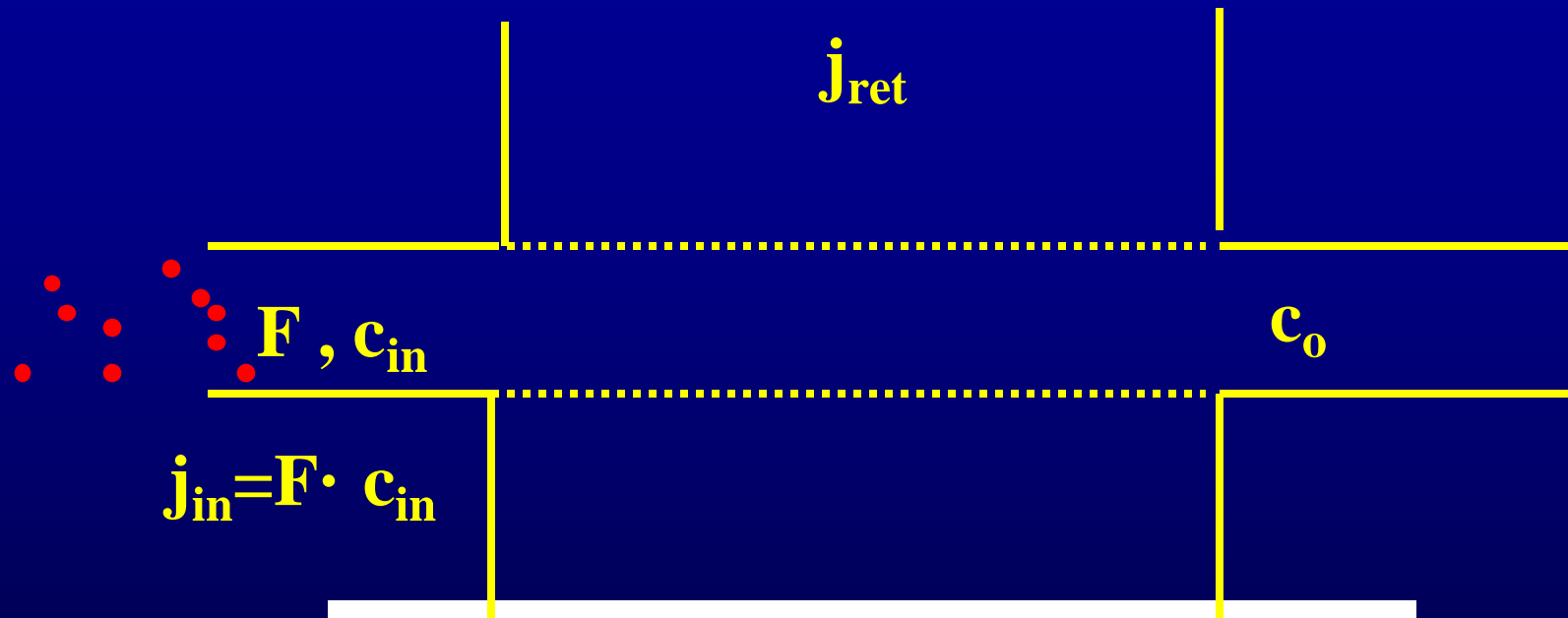
$$j_{in}(t) \neq j_o(t) + j(t)$$

$$j(t) = K_i \cdot c(t)$$

$$v \frac{dc(t)}{dt} = F \cdot c_{in}(t) - F \cdot c_o(t) - K_i \cdot c(t)$$

Break

Extraction fraction

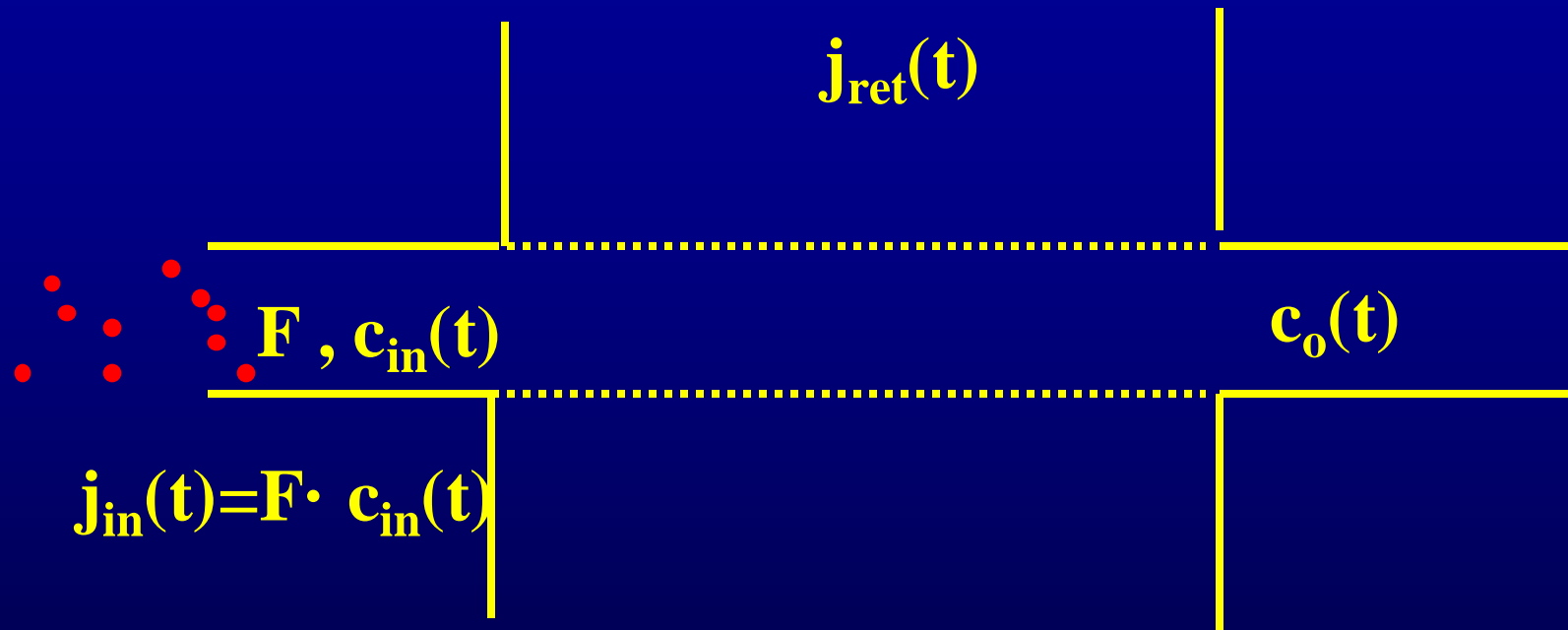


Extraction:

$$E = \frac{\dot{j}_{ret}}{\dot{j}_{in}} = \frac{F \cdot c_{in} - F \cdot c_o}{F \cdot c_{in}} = \frac{c_{in} - c_o}{c_{in}}$$

The transmitted fraction = 1-E

Extraction fraction

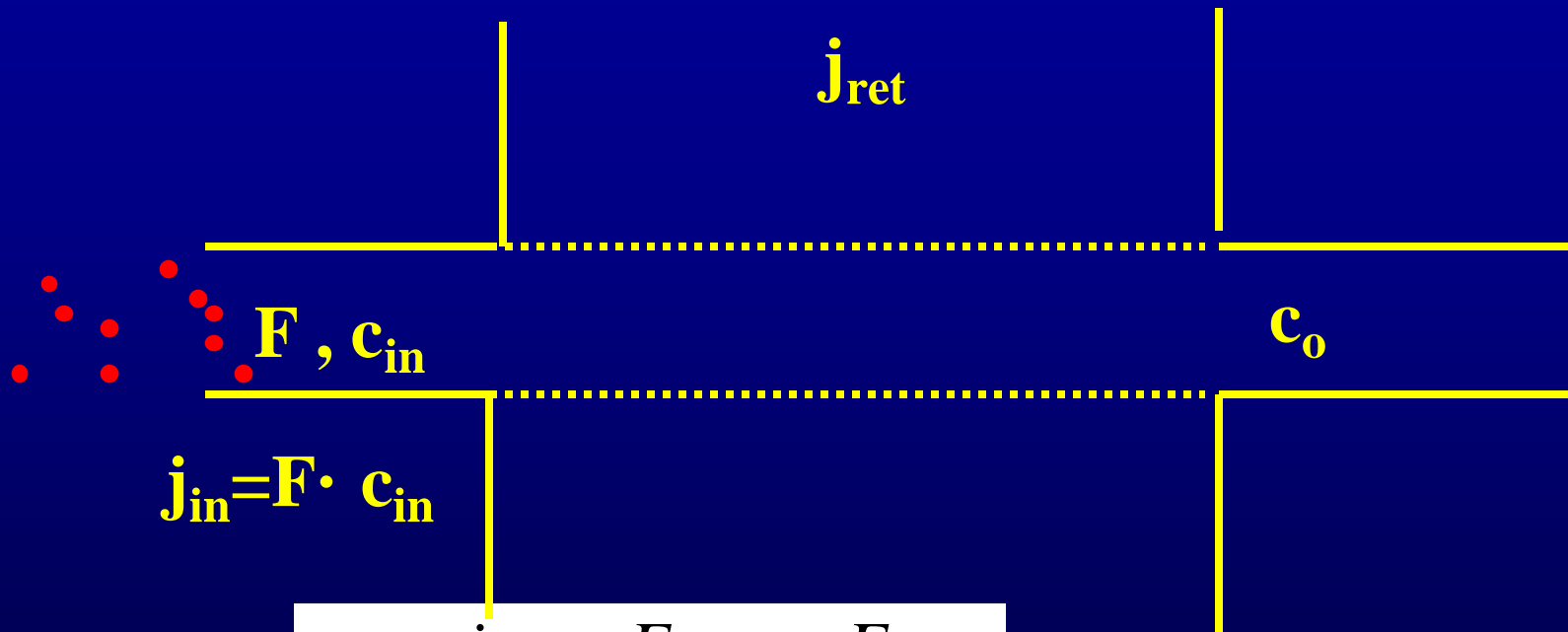


Extraction:

$$E = \frac{j_{ret}(t)}{j_{in}(t)} = \frac{F \cdot c_{in}(t) - F \cdot c_o(t)}{F \cdot c_{in}(t)} = \frac{c_{in}(t) - c_o(t)}{c_{in}(t)}$$

E constant ?

Clearance



clearance:

$$Cl = \frac{j_{ret}}{c_{ref}} = \frac{F \cdot c_{in} - F \cdot c_o}{c_{ref}}$$

$[Cl] = \text{ml/s}$

Clearance

It is a fictive flow: the volume of reference fluid containing the indicator amount taken up or cleared per unit time

Clearance



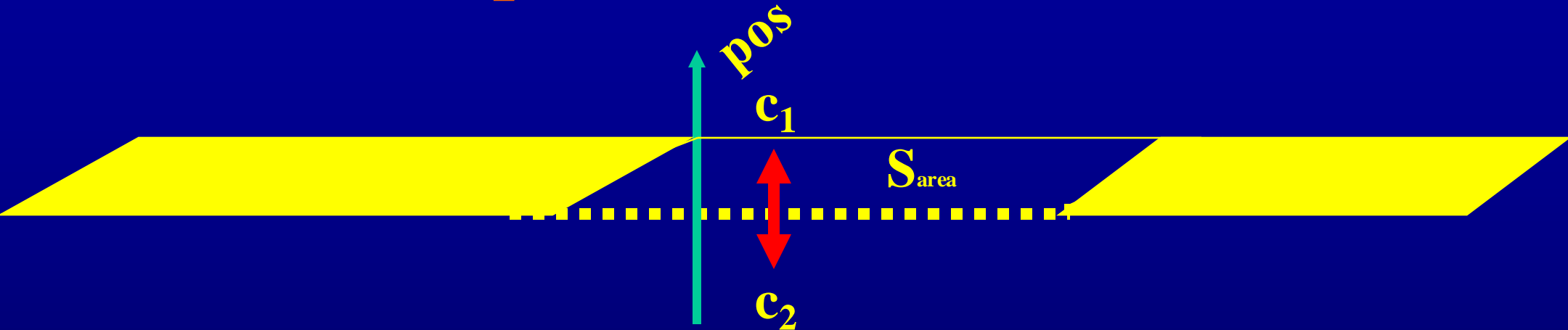
clearance:

$$Cl = \frac{j_{ret}}{c_{ref}} = \frac{j_{ret}}{c_{in}} = \frac{F \cdot c_{in} - F \cdot c_o}{c_{in}} = F \cdot E$$

$$j_{ret} = Cl \cdot c_{in} = K_i \cdot c_{in} = F \cdot E \cdot c_{in}$$

Break

Transport over a membrane



$$j \quad ? \quad c_2 - c_1$$

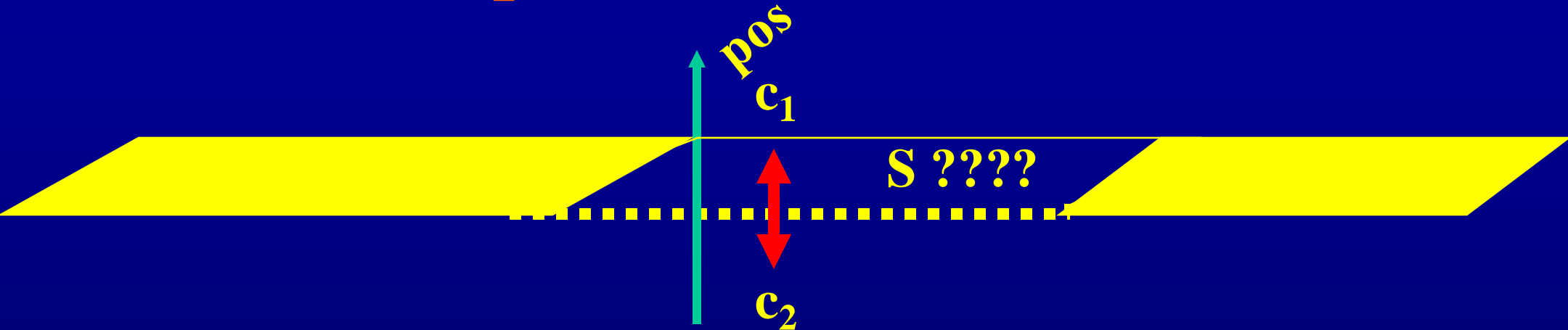
$$j/S = P(c_2 - c_1)$$

$$j/S = -P(c_1 - c_2)$$

$$\text{mol/cm}^2/\text{s} = [P] \text{ mol/ml}$$

$$[P] = \text{cm/s}$$

Transport over a membrane



$$j = -PS(c_1 - c_2)$$

S = surface area

$$\text{mol/s} = [PS] \text{ mol/ml}$$

$$[PS] = \text{cm}^3/\text{s} = \text{ml/s}$$

$$[PS] = \text{cm}^3/\text{g/s} = \text{ml/g/s}$$

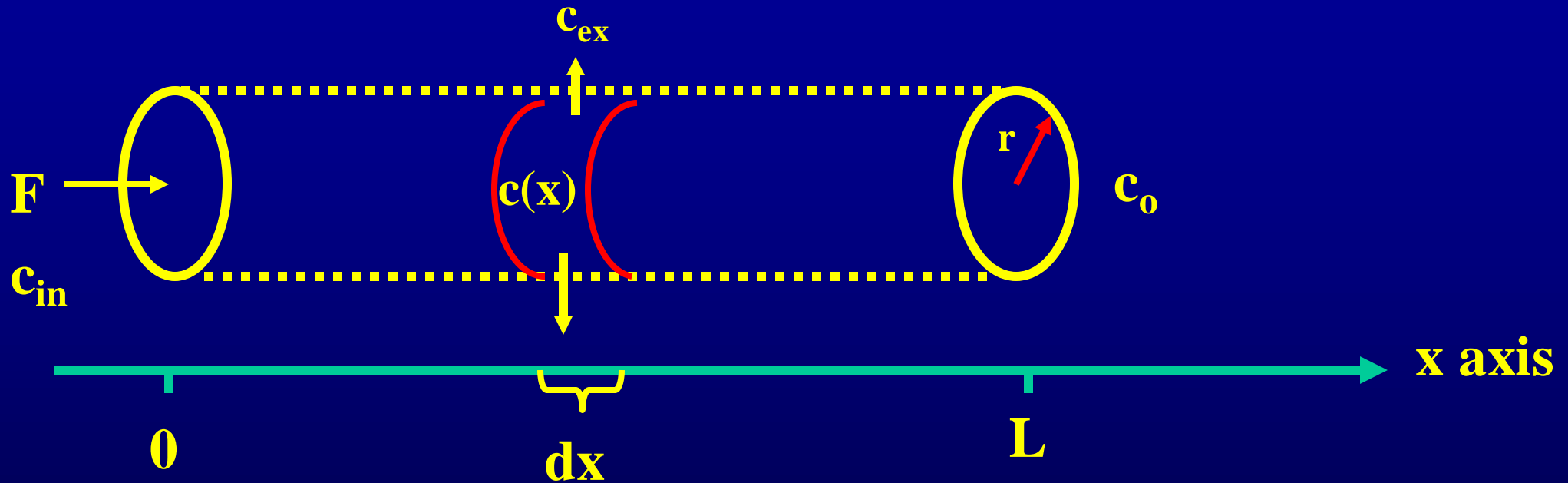
Crone (1963) & Renkin (1959) equation

Transport over the capillary membrane



$$c_o = c_{in} \exp(-PS/F)$$

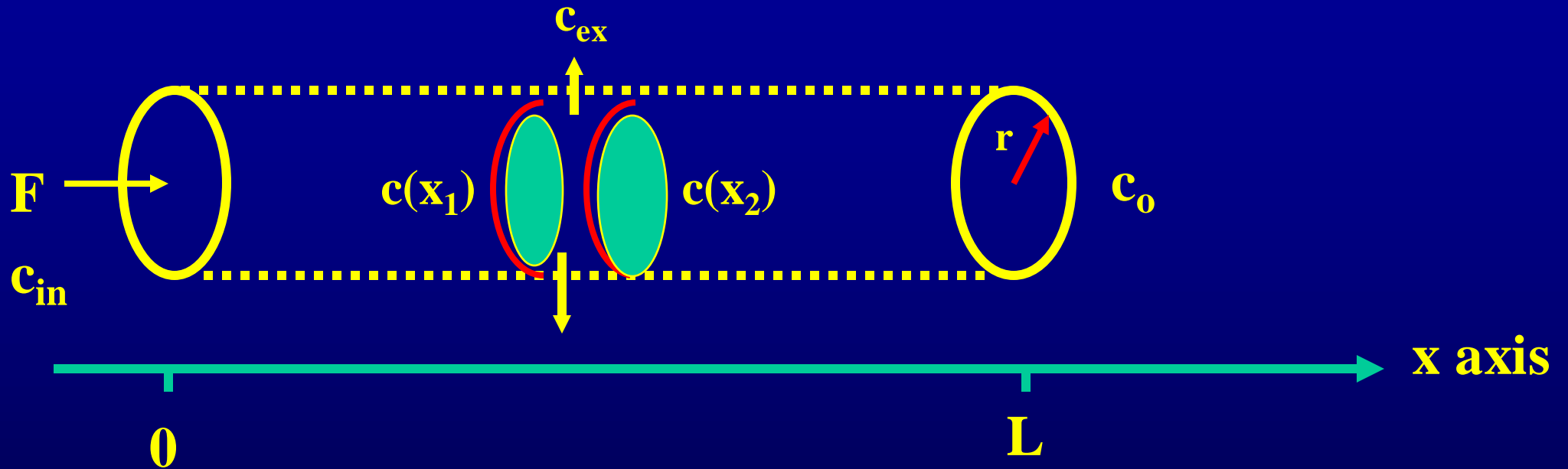
Crone (1963) & Renkin (1959) equation



The flux out
of dx :

$$dj = -\frac{dS}{S} PS(c_{ex} - c(x)) = \frac{2\pi r dx}{2\pi r L} PS c(x)$$

Crone (1963) & Renkin (1959) equation



The loss inside the capillary:

$$dj = -F(c(x_2) - c(x_1))$$

Fick's
principle

$$dj = -Fdc(x)$$

Crone (1963) & Renkin (1959) equation

Transport over the capillary membrane

$$\left. \begin{aligned} dj &= \frac{2\pi r dx}{2\pi r L} PS c(x) \\ dj &= -F dc(x) \end{aligned} \right\} \Rightarrow \frac{dc(x)}{c(x)} = -\frac{PS}{LF} dx$$

$$\int_{c_{in}}^{c_o} \frac{dc(x)}{c(x)} = -\int_0^L \frac{PS}{LF} dx$$

$$\ln \frac{c_o}{c_{in}} = -\frac{PSL}{LF}$$

$$c_o = c_{in} \exp(-PS/F)$$

Crone (1963) & Renkin (1959) equation

$$c_0 = c_i e^{-\frac{PS}{F}} \Rightarrow \frac{c_0}{c_i} = e^{-\frac{PS}{F}}$$

$$1 - \frac{c_0}{c_i} = 1 - e^{-\frac{PS}{F}} \Rightarrow \frac{c_i - c_0}{c_i} = 1 - e^{-\frac{PS}{F}}$$

$$E = 1 - e^{-\frac{PS}{F}} \wedge Cl = FE \Rightarrow Cl = K_i = F(1 - e^{-\frac{PS}{F}})$$

Accumulation of tracer in tissue can be
Flow Limited or Diffusion Limited

Flow limited : PS/F is large

$$\mathbf{E = 1 - \exp(-PS/F) \quad E \rightarrow 1 \text{ for } PS/F \rightarrow \infty}$$

$$\mathbf{CI = F E \rightarrow F}$$

Accumulation of tracer in tissue can be
Flow Limited or Diffusion Limited

Diffusion limited : PS/F is small

$$\mathbf{E = 1 - \exp(-PS/F)} \quad \mathbf{E \rightarrow 0 \text{ for } PS/F \rightarrow 0}$$

$$\mathbf{E = 1 - \exp(-PS/F) \approx 1 - (1 - PS/F) = PS/F}$$

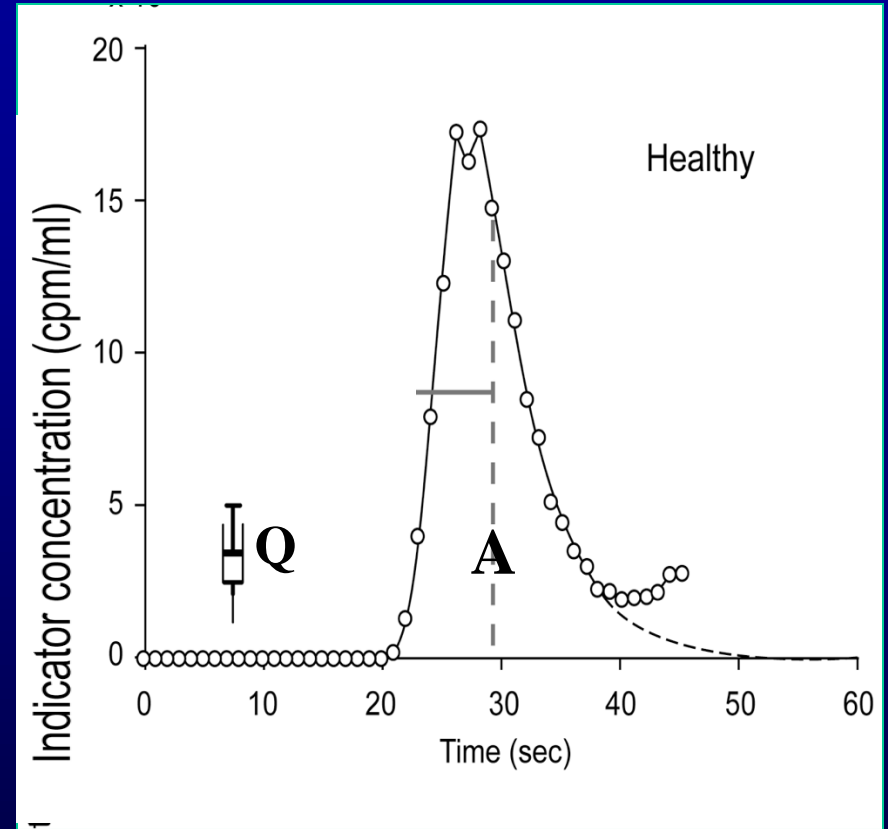
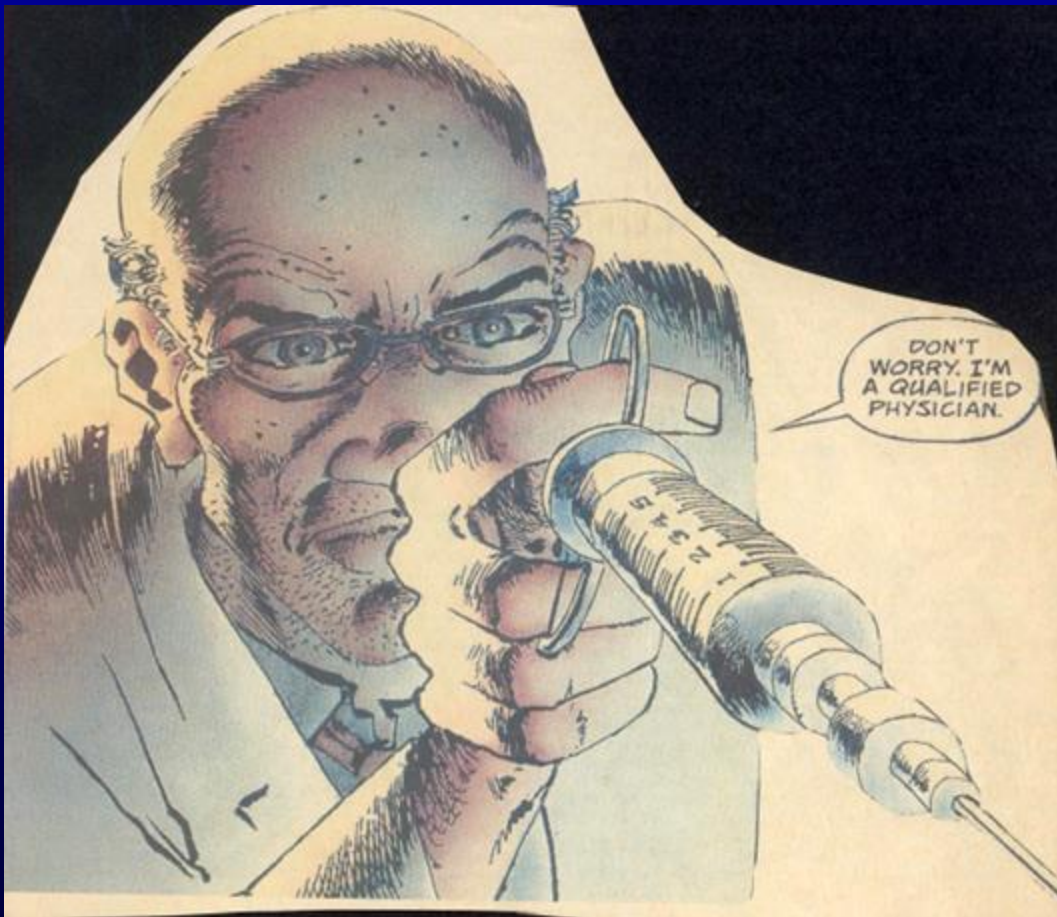
$$\mathbf{CI = F E \rightarrow PS}$$

Break

Indicator-technique

Stewart-Henriques-Hamilton

Bolus injection

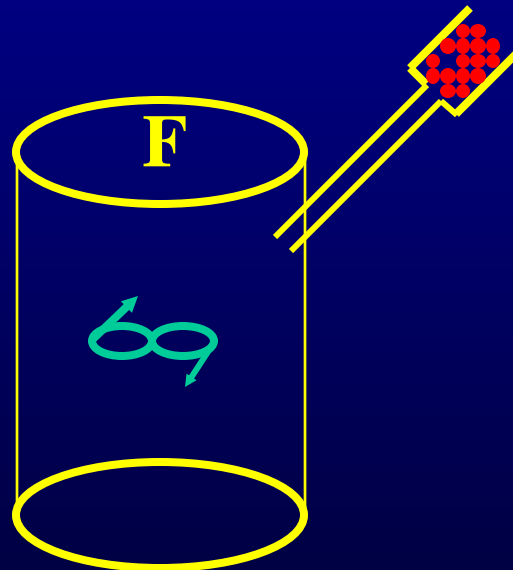


$$CO = Q/A$$

Indicator-dilution methods continued

Bolus injection (Henriques-Hamilton-Bergner principle)

The aim : to measure the flow of an organ or a vessel or a pipeline

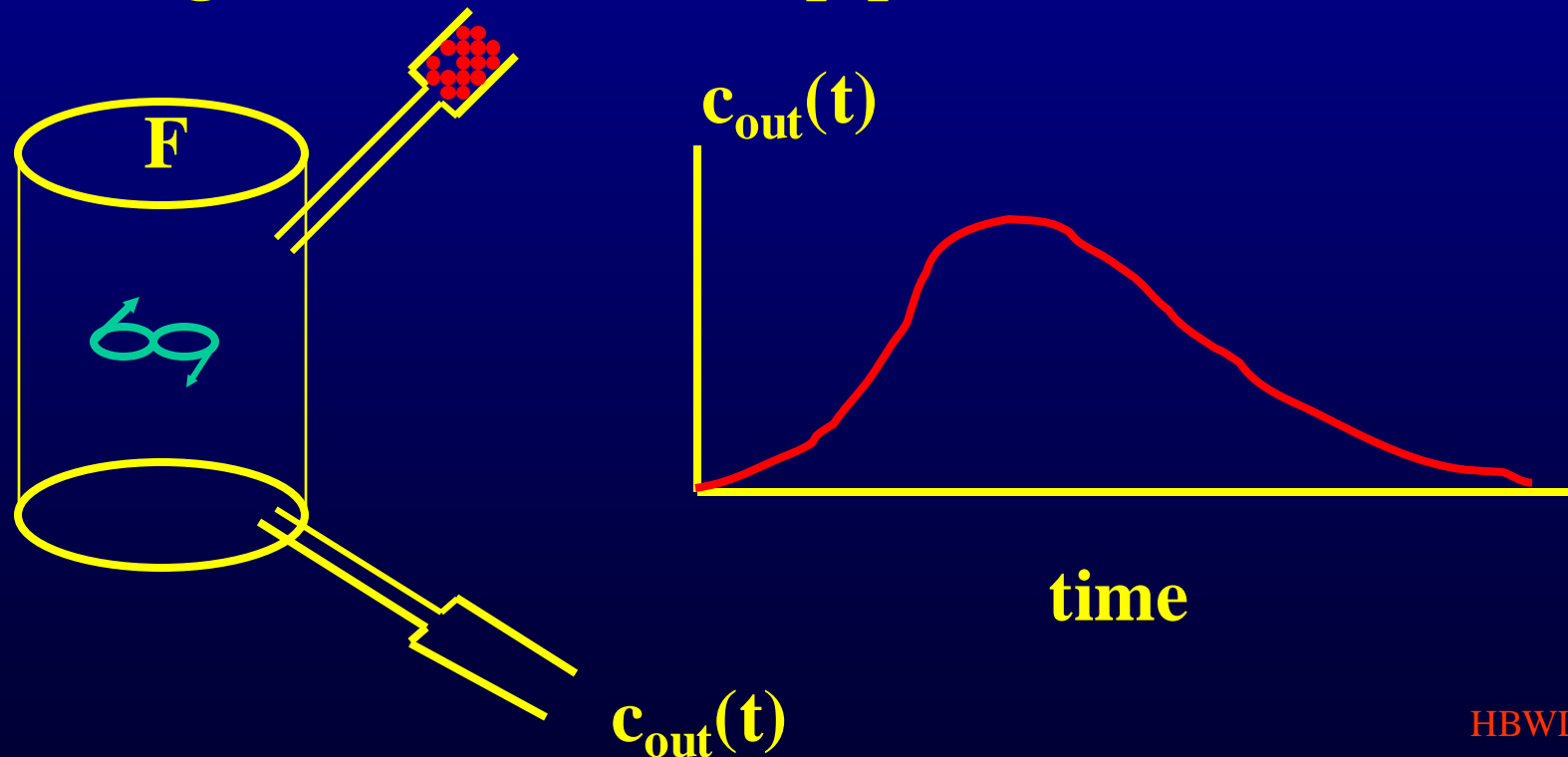


**Injection of bolus Q_0 ,
a known amount of
tracer**

Indicator-dilution methods continued

Bolus injection (Henriques-Hamilton-Bergner principle)

The aim : to measure the flow of an organ or a vessel or a pipeline



Indicator-dilution methods continued

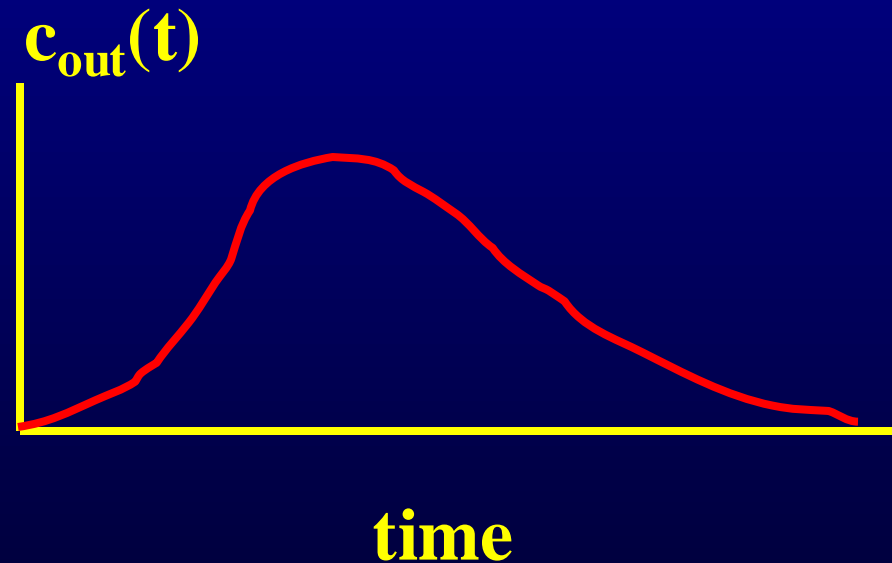
Bolus injection (Henriques-Hamilton-Bergner principle)

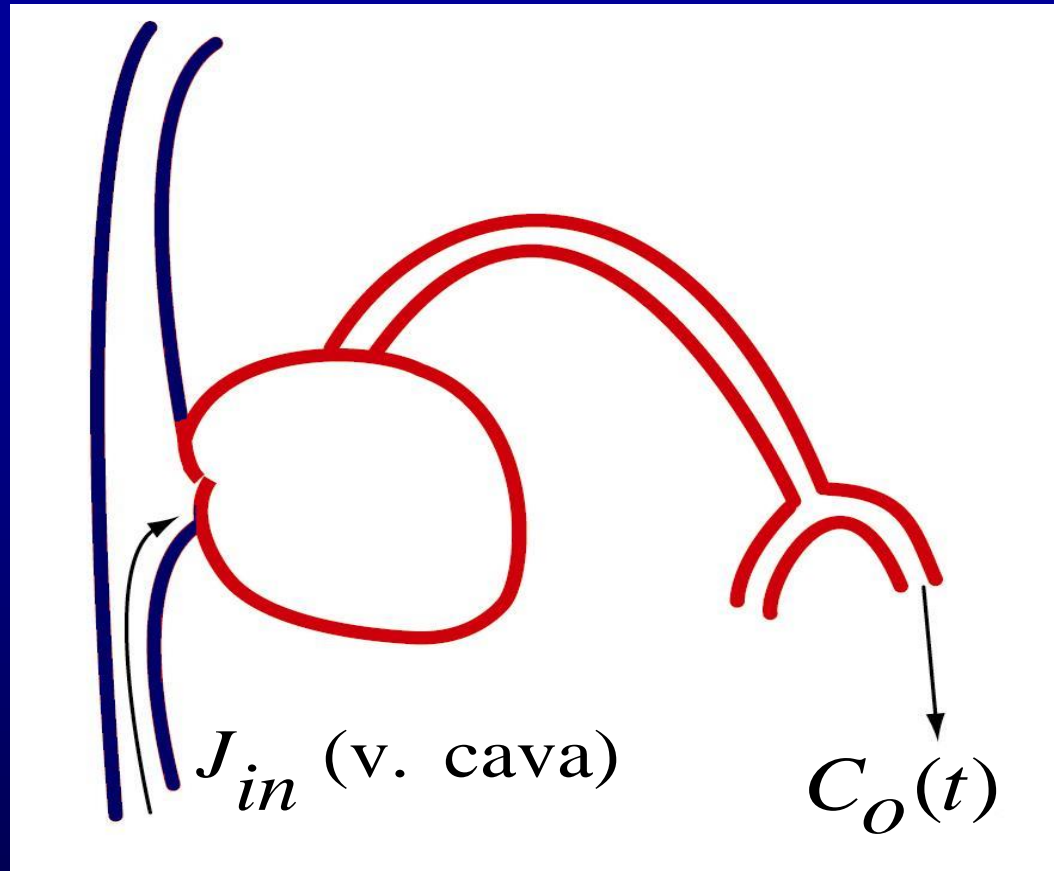
The aim : to measure the flow of an organ or a vessel or a pipeline

$$dQ(t) = F \cdot c_{out}(t) \cdot dt$$

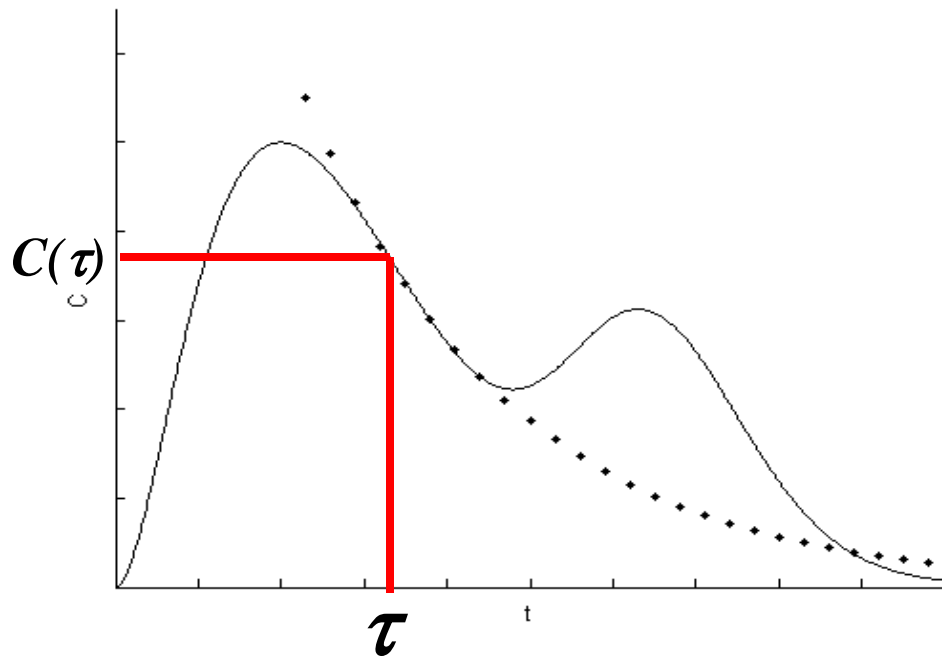
$$\int_0^{\infty} Q_0 = \int_0^{\infty} F \cdot c_{out}(t) \cdot dt$$

$$F = \frac{Q_0}{\int_0^{\infty} c_{out}(t) dt}$$





Bolus injection in vena cava/periferal vein, and outlet concentration measurement from a peripheral artery.



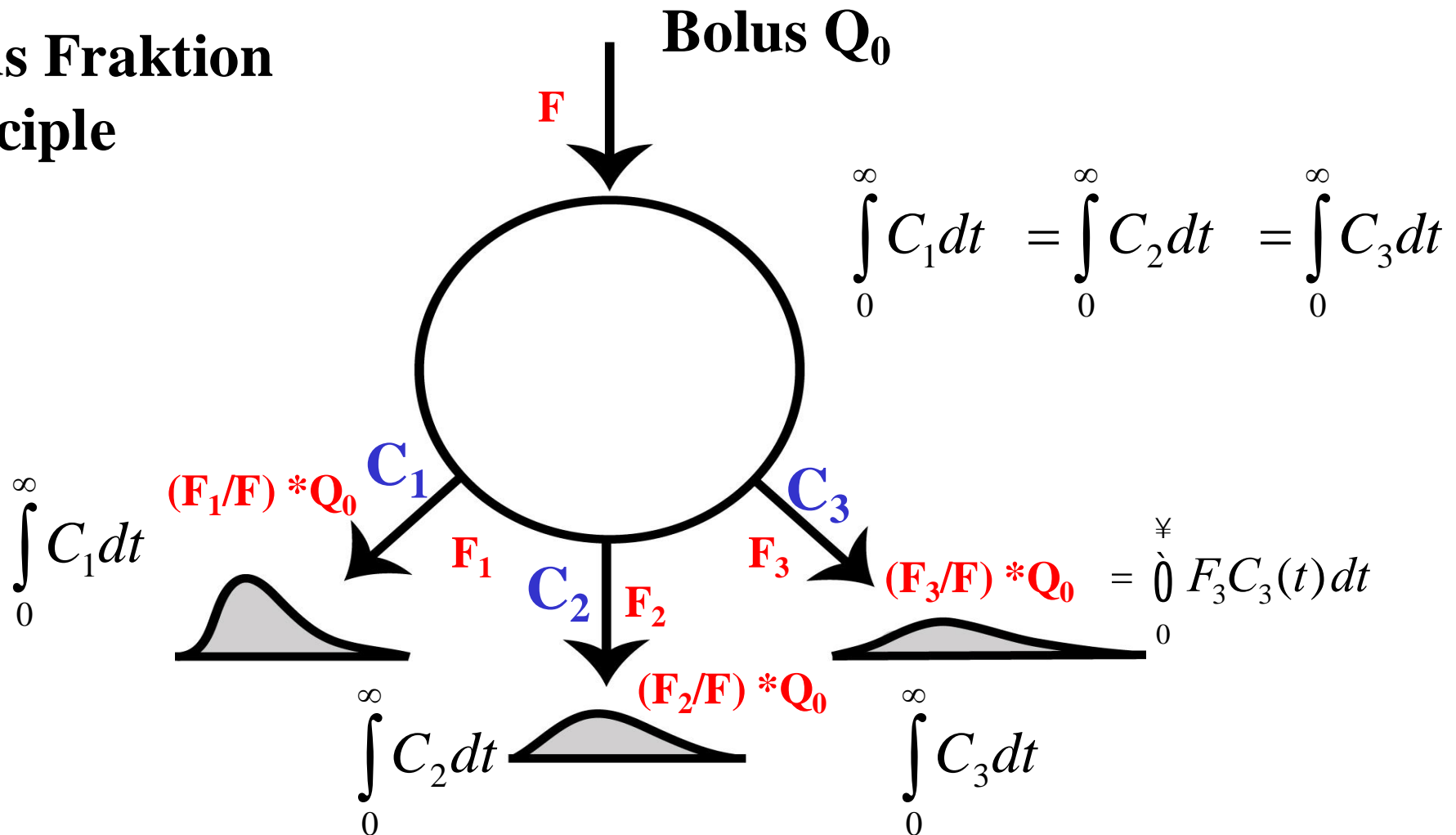
$$\int_0^{\infty} C_o(t) dt = \int_0^{\tau} C_o(t) dt + \int_{\tau}^{\infty} C(\tau) e^{-k(t-\tau)} dt =$$

$$\int_0^{\tau} C_o(t) dt + \frac{C(\tau)}{-k} \left[e^{-k(t-\tau)} \right]_{\tau}^{\infty} =$$

$$\int_0^{\tau} C_o(t) dt + \frac{C(\tau)}{k} \Rightarrow$$

$$F = \frac{Q_o}{\int_0^{\tau} C_o(t) dt + \frac{C(\tau)}{k}}$$

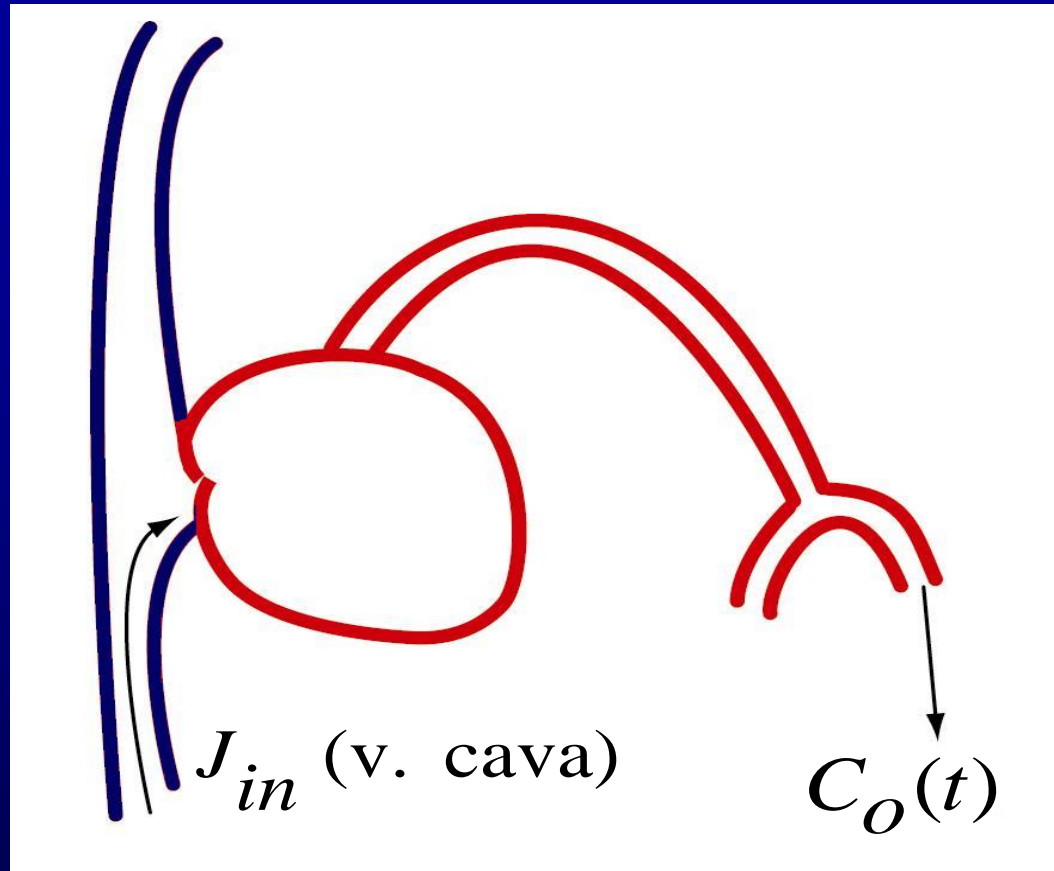
Bolus Fraction principle



Equal area rule. The shape is different but the areas of the different outlets are equal. This allows us to choose freely the most appropriate sampling point with regards the outlet concentration measurement.

$$(\mathbf{F}_i/\mathbf{F}) * \mathbf{Q}_0 = \int_0^{\infty} F_i C_i(t) dt$$

$$F = \frac{Q_0}{\int_0^{\infty} C_i(t) dt}$$

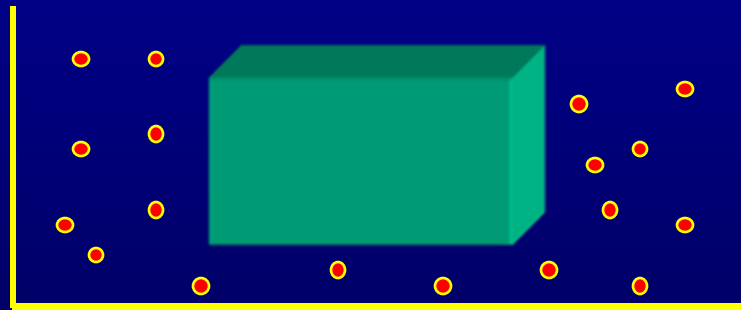


So the outlet concentration can be measured from a convenient artery

Bolus injection in vena cava/periferal vein, and outlet concentration measurement from a peripheral artery.

The volume of distribution: V_d

A tissue element



Incubation with a reference fluid

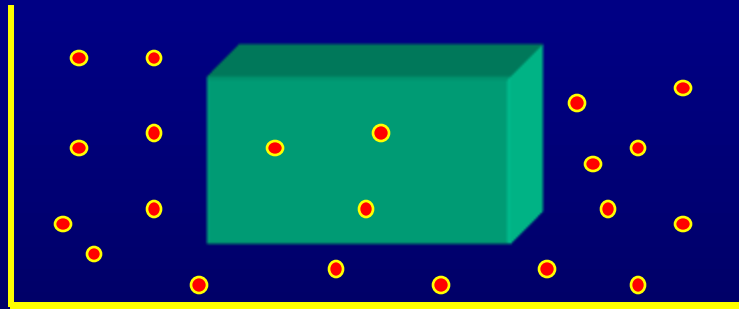
with a concentration c_{ref}

$$V_d \equiv Q/c_{\text{ref}}$$

$$[V_d] = \text{mmol}/\text{mmol}/\text{ml} = \text{ml}$$

The volume of distribution: V_d

A tissue element



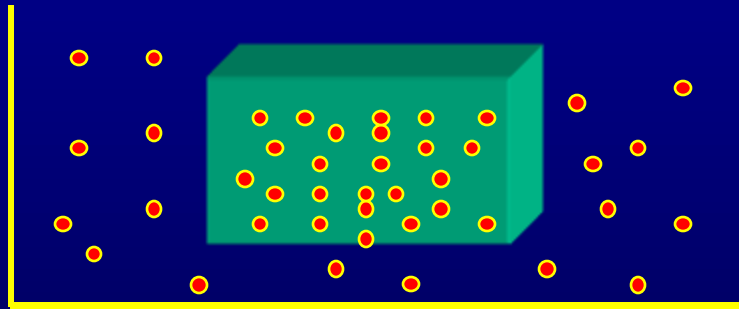
$$V_d \equiv Q/c_{\text{ref}}$$

V_d larger or smaller

**than the real volume of
the tissue ?**

The volume of distribution: V_d

A tissue element



$$V_d \equiv Q/c_{\text{ref}}$$

V_d larger or smaller

**than the real volume of
the tissue ?**

The volume of distribution: V_d

$$V_d \equiv Q/c_{\text{ref}}$$

It is the volume of the reference fluid which contains the amount Q

The partition coefficient $\lambda \equiv V_d/W$ or V_d/V

**W is either the (real) mass of the tissue : $[\lambda] = \text{ml/g}$
or**

V is the (real) volume of the tissue : $[\lambda] = \text{ml/ml}$

The partition coefficient λ

$$c_{\text{tissue}} = Q/W \qquad c_{\text{tissue}} = Q/V$$

Where W is either the real mass of tissue: $[c_{\text{tissue}}]=\text{mmol/g}$

Or

V is the (real) volume of the tissue: $[c_{\text{tissue}}]=\text{mmol/ml}$

$$\lambda \equiv \frac{V_d}{W} = \frac{Q}{c_{\text{ref}} \cdot W} = \frac{c_{\text{tissue}}}{c_{\text{ref}}}$$

Examples

- Plasma concentration is 200 ng/ml
- Total amount of substance 10 mg
- Volume of distribution is $10\text{mg}/200 \text{ ng/ml} = 50 \text{ L}$

Examples

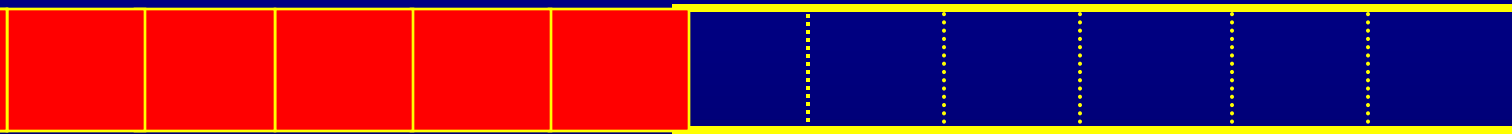
- Regional tissue concentration: 100 kBq/cm³
- Plasma concentration: 5 kBq/ml
- Volume of distribution:
 $(100 \text{ kBq/cm}^3) / (5 \text{ kBq/ml}) = 20 \text{ ml/cm}^3$

That 20 ml plasma would be required to account for the tracer in just 1 cm³ of tissue

Break

Mean transit time

The simplicity of this concept



$$V_d = 6 \text{ ml}$$

$$\text{A flow } F = 1 \text{ ml/s}$$

What is the (mean) transit time of the tracer in this compartment ?

$$\bar{t} = V_d / F = 6 \text{ ml} / 1 \text{ ml/s} = 6 \text{ s}$$

Mean transit time

$$\bar{t} = V_d / F$$

$$\lambda = V_d / W$$

$$f = F / W$$

$$\bar{t} = \lambda / f$$

Mean transit time

The definition

$$\bar{t} = \frac{1}{Q_0} (t_1 \cdot \Delta Q_1 + t_2 \cdot \Delta Q_2 + t_3 \cdot \Delta Q_3 + \dots + t_i \cdot \Delta Q_i + \dots) \wedge Q_0 = \sum_i \Delta Q_i$$

$$\bar{t} = \frac{1}{Q_0} \sum_i t_i \cdot \Delta Q_i = \sum_i t_i \cdot \frac{\Delta Q_i}{Q_0} = \sum_i t_i \cdot \frac{\Delta Q_i}{Q_0 \cdot \Delta t} \xrightarrow{\text{lim}} \int_0^{\infty} t \cdot \frac{dQ(t)}{Q_0 \cdot dt} \cdot dt$$

Define the frequency function of transit times:

$$h(t) \equiv \frac{dQ(t)}{Q_0 \cdot dt}$$

$$[h(t)] = 1/s$$

The frequency function

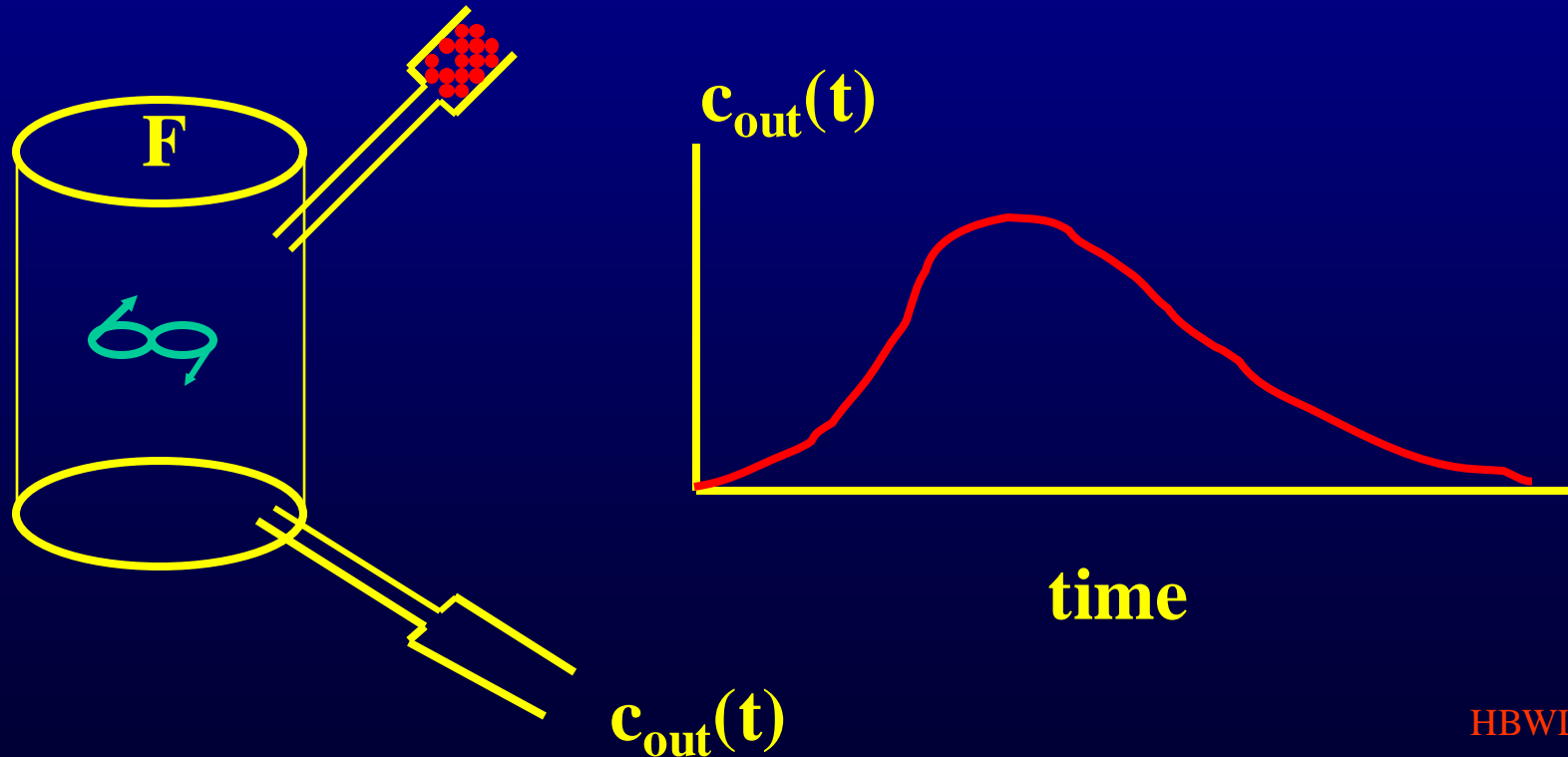
$$h(t) \equiv \frac{dQ(t)}{Q_0 \cdot dt}$$

In words : It is the fraction of the dose given as an impuls (a delta function), which leaves the system per unit time !!!! , at time t, (and therefore a function of time)

$$\bar{t} = \int_0^{\infty} t \cdot h(t) \cdot dt$$

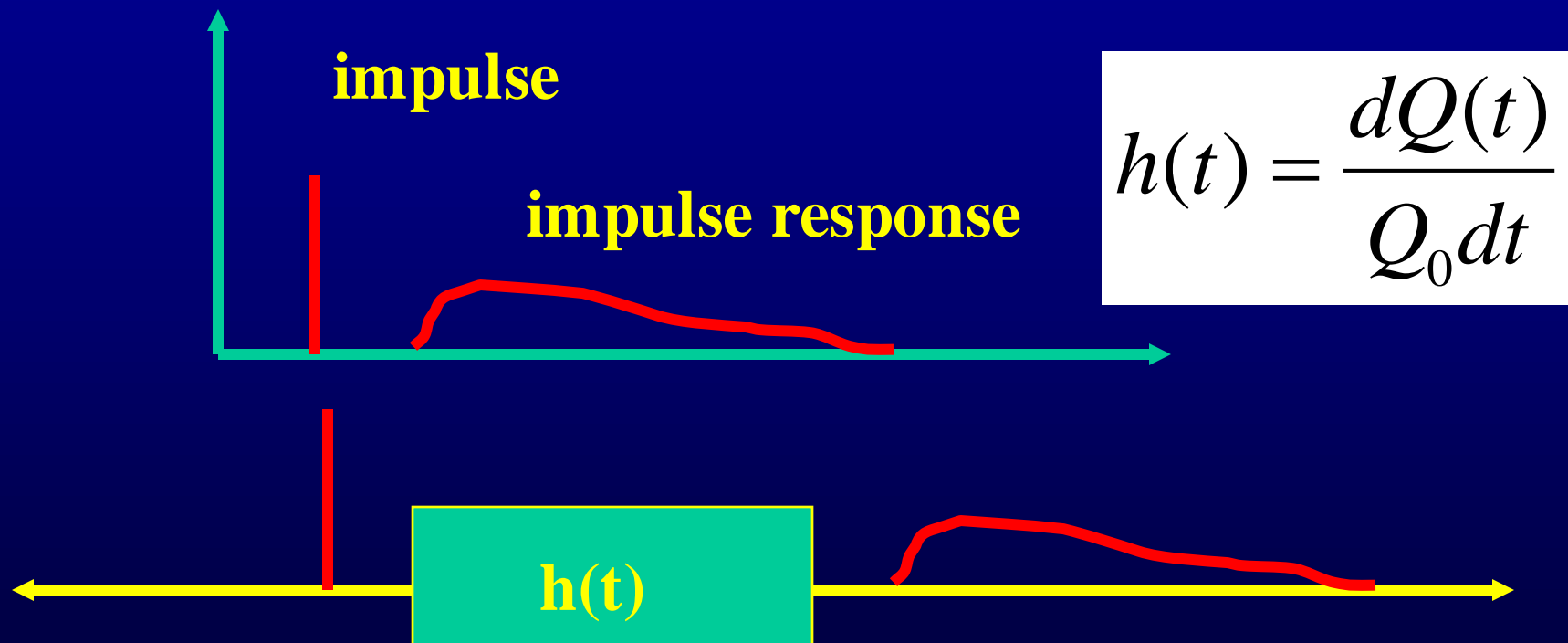
Finding $h(t)$

$$h(t) = \frac{c_{out}(t)}{\int_0^{\infty} c_{out}(t) \cdot dt} \cdot \frac{F \cdot c_{in}}{\bar{t}} = \int_0^{\infty} t \cdot h(t) \cdot dt$$



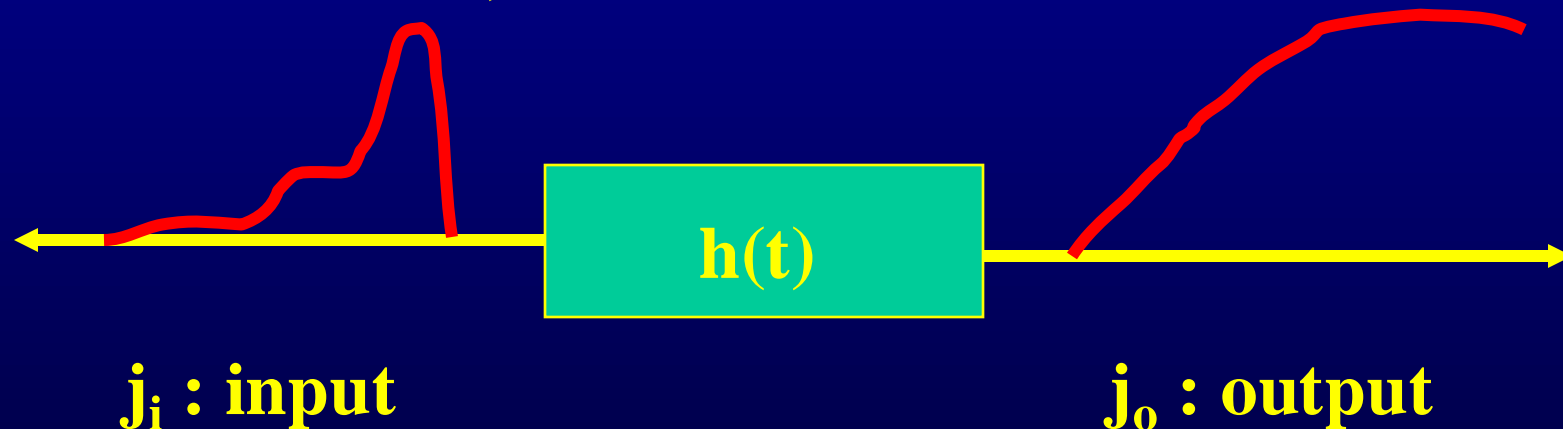
Break

The impulse response of a inlet and outlet system (artery – vein system)

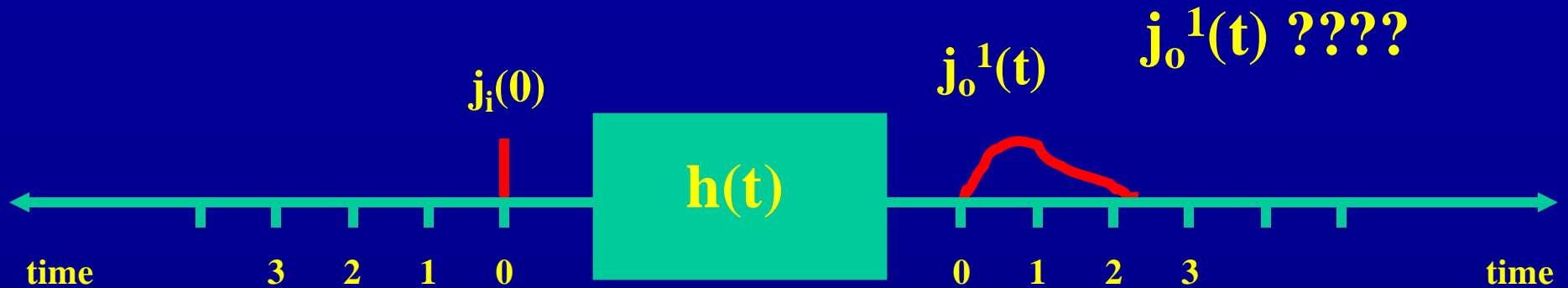


Why is $h(t)$ interesting ?

Because it relates input to an output in the case of the input not being a bolus (a deltafunction) !



$$j_o(t) = j_i(t) \otimes h(t) = \int_0^t j_i(\tau) h(t - \tau) d\tau$$

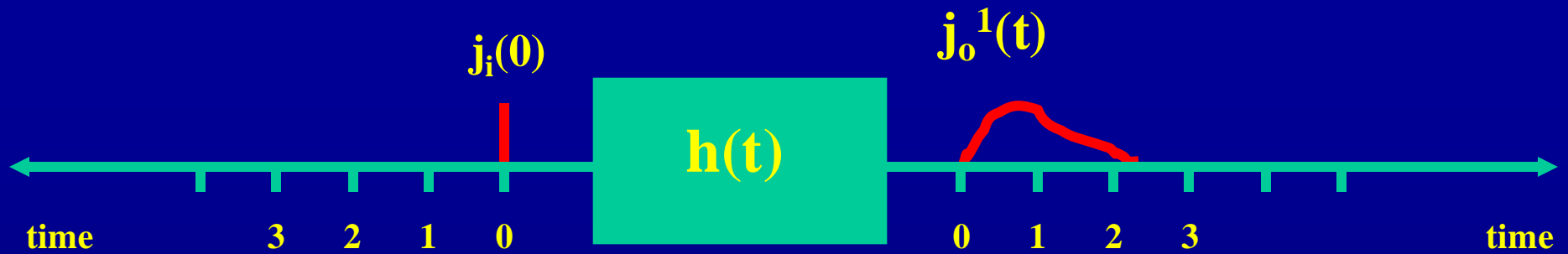


e.g $j_i(0) = 1 \text{ mmol} / 0.01\text{s}$

$$j_o^1(t) = j_i(0) \Delta\tau h(t)$$



**Flux (number pr unit time - as a function of time)
leaving the system due to an input at time zero**

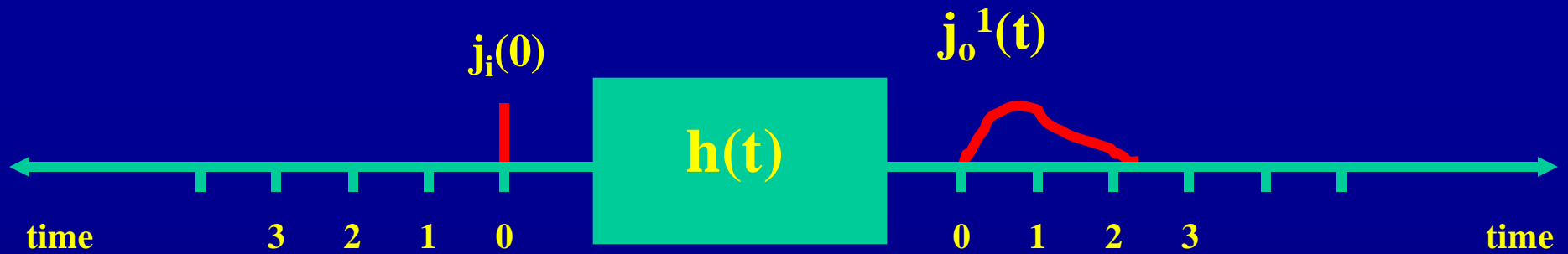


e.g $j_i(0) = 1 \text{ mmol} / 0.01\text{s}$

$$j_o^1(t) = j_i(0) \Delta\tau h(t)$$



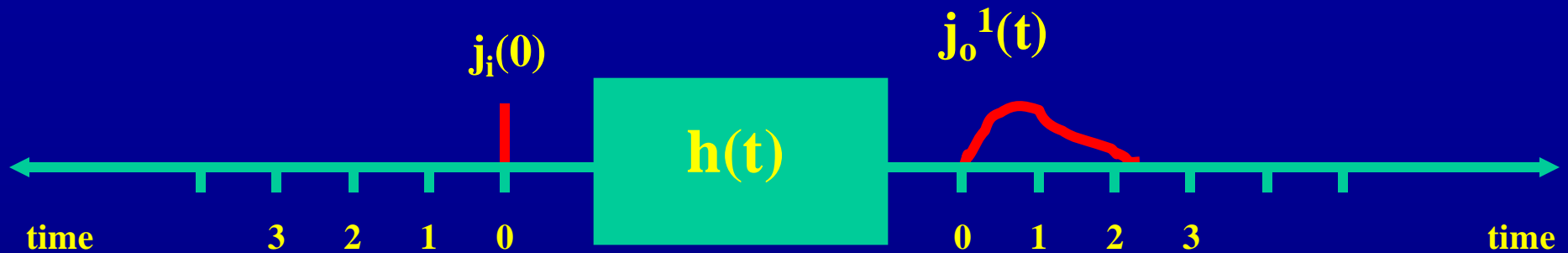
**Flux entering
the system at
time zero**



e.g $j_i(0) = 1 \text{ mmol} / 0.01\text{s}$

$$j_o^1(t) = j_i(0) \Delta\tau h(t)$$

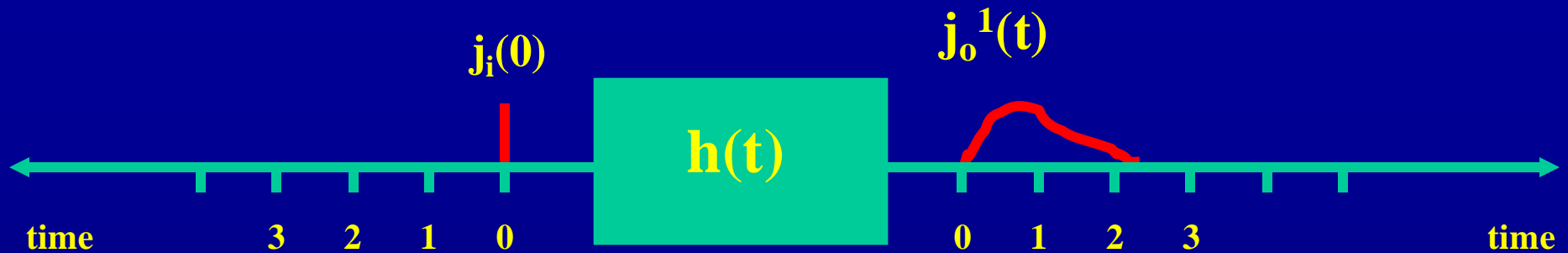
A small time
interval



e.g $j_i(0) = 1 \text{ mmol} / 0.01\text{s}$

$$j_o^1(t) = \underbrace{j_i(0) \Delta\tau}_{\text{Amount of tracer entering at } t=0} h(t)$$

The amount (the number) of tracer entering the system at time zero

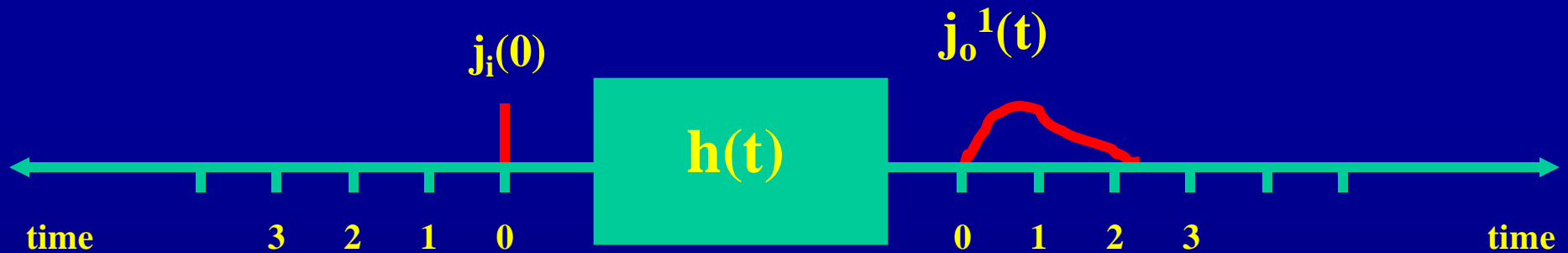


e.g $j_i(0) = 1 \text{ mmol} / 0.01\text{s}$

$$j_o^1(t) = j_i(0) \Delta\tau h(t)$$

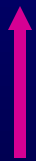


The impulse response function: the fractional amount (the number) pr. unit time - leaving the system as a function of time

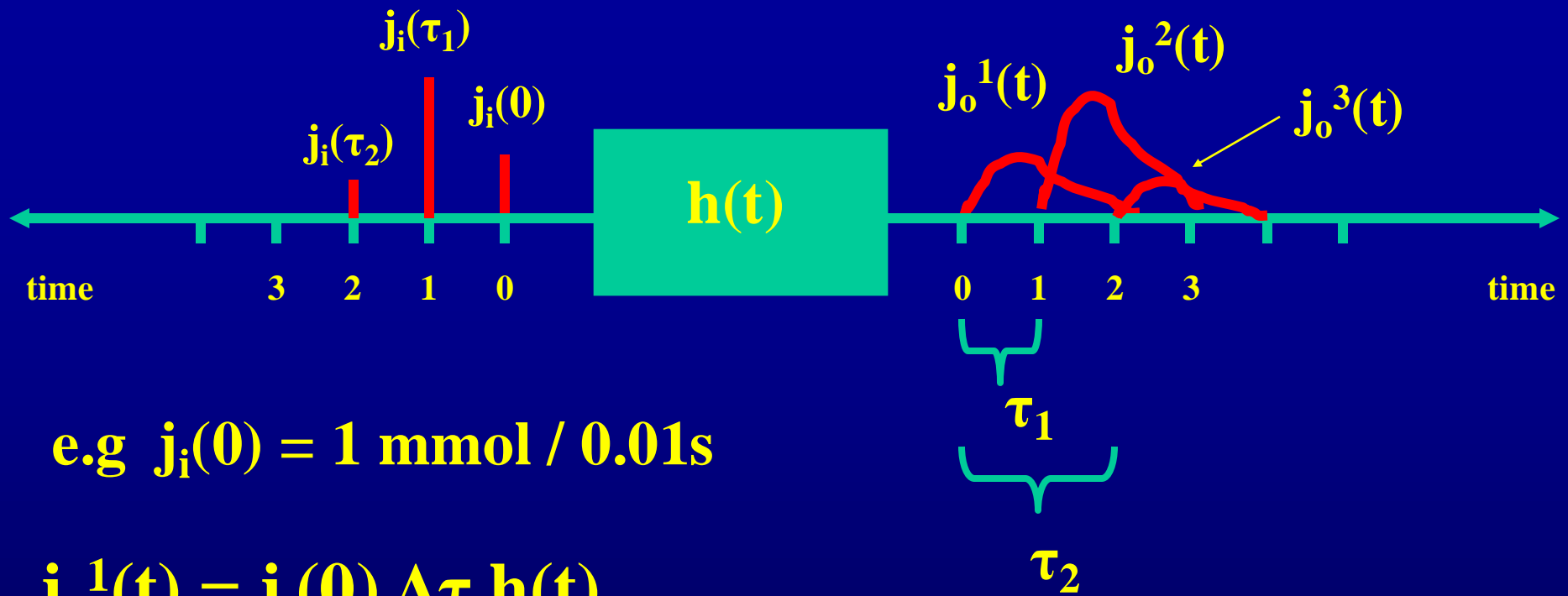


e.g $j_i(0) = 1 \text{ mmol} / 0.01\text{s}$

$$j_o^1(t) = j_i(0) \Delta\tau h(t)$$



**Flux (number pr unit time - as a function of time)
leaving the system due to an input at time zero**

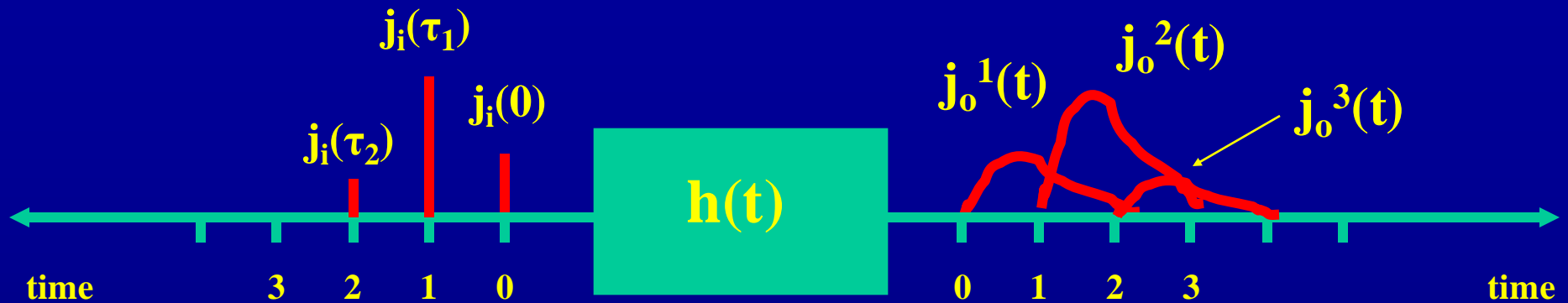


e.g $j_i(0) = 1 \text{ mmol} / 0.01\text{s}$

$$j_o^1(t) = j_i(0) \Delta\tau h(t)$$

$$j_o^2(t) = j_i(\tau_1) \Delta\tau h(t - \tau_1) = j_i(\tau_1) h(t - \tau_1) \Delta\tau$$

$$j_o^3(t) = j_i(\tau_2) h(t - \tau_2) \Delta\tau$$



$$j_o^1(t) = j_i(0) \Delta\tau h(t)$$

$$j_o^2(t) = j_i(\tau_1) h(t - \tau_1) \Delta\tau$$

$$j_o^3(t) = j_i(\tau_2) h(t - \tau_2) \Delta\tau$$

Total flux $j_o(t) = j_o^1(t) + j_o^2(t) + j_o^3(t) =$

$$j_i(0) h(t-0) \Delta\tau + j_i(\tau_1) h(t - \tau_1) \Delta\tau + j_i(\tau_2) h(t - \tau_2) \Delta\tau$$

$$j_0(t) = \sum_0^N j_i(\tau_n) h(t - \tau_n) \Delta \tau$$

$$\Delta \tau \rightarrow 0 \Rightarrow j_0(t) = \int_0^t j_i(\tau) h(t - \tau) d\tau$$

$$j_o(t) = j_i(t) \otimes h(t)$$

$$j_o(t) = F c_o(t)$$

$$j_i(t) = F c_i(t)$$

$$c_o(t) = c_i(t) \otimes h(t)$$

Break



$h(t)$ & $H(t)$!!

$h(t)$ is an analogous to a probability density function

The first moment of this function corresponds to the mean value or expectation value

$$\int_0^{+\infty} h(t) dt = \frac{1}{\int_0^{\infty} c_o(t) dt} \int_0^{\infty} c_o(t) dt = 1$$

$$\bar{x} = E(x) = \int_0^{\infty} x p(x) dx$$

$$\bar{t} = \int_0^{\infty} t h(t) dt$$

Mean transit time

The definition

$$\bar{t} = \frac{1}{Q_0} (t_1 \cdot \Delta Q_1 + t_2 \cdot \Delta Q_2 + t_3 \cdot \Delta Q_3 + \dots + t_i \cdot \Delta Q_i + \dots) \wedge Q_0 = \sum_i \Delta Q_i$$

$$\bar{t} = \frac{1}{Q_0} \sum_i t_i \cdot \Delta Q_i = \sum_i t_i \cdot \frac{\Delta Q_i}{Q_0} = \sum_i t_i \cdot \frac{\Delta Q_i}{Q_0 \cdot \Delta t} \xrightarrow{\text{lim}} \int_0^{\infty} t \cdot \frac{dQ(t)}{Q_0 \cdot dt} \cdot dt$$

Define the frequency function of transit times:

$$h(t) \equiv \frac{dQ(t)}{Q_0 \cdot dt}$$

$$[h(t)] = 1/s$$

$h(t)$ & $H(t)$!!

$$h(t) = \frac{dQ(t)}{Q_0 dt} = \frac{c_o(t)}{\int_0^{\infty} c_o(\tau) d\tau}$$

The fraction that leaves the system as a function of time pr unit time after a bolus inj

$$h(t) dt = \frac{dQ(t)}{Q_0}$$

The fraction that leaves the system as a function of time in a short time interval ^{HBWL}

$h(t)$ & $H(t)$!!

$$\int_0^{t_1} h(t) dt = \frac{1}{Q_0} \int_0^{t_1} dQ(t) = \frac{1}{Q_0} (Q(t_1) - Q(0))$$

**The fraction
having left the
system in the
time interval
 $0:t_1$ (after a
bolus injection)**

$$\int_{t_1}^{t_2} h(t) dt = \frac{1}{Q_0} \int_{t_1}^{t_2} dQ(t) = \frac{1}{Q_0} (Q(t_2) - Q(t_1))$$

**The fraction
having left the
system in the
time interval
 $t_1:t_2$**

$h(t)$ & $H(t)$!!

$$H(t) \equiv \int_0^t h(\tau) d\tau = \frac{1}{Q_0} \int_0^t dQ(\tau) = \frac{Q(t)}{Q_0}$$

The fraction remaining in the system at time t after a bolus inj

$$1 - H(t) = 1 - \int_0^t h(\tau) d\tau$$

The fraction having left the system in the time interval $0:t$ (after a bolus injection)

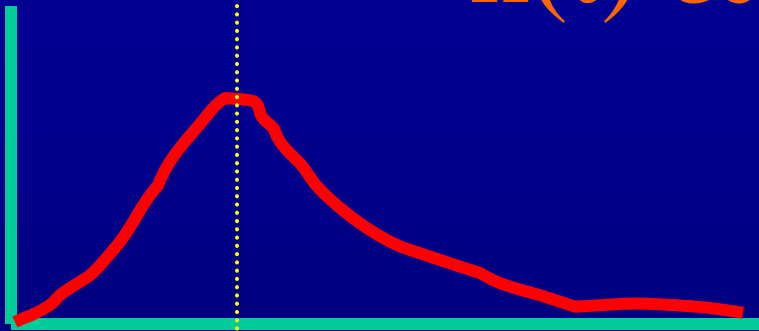
$h(t)$ & $H(t)$!!

**The residue impulse
response function**

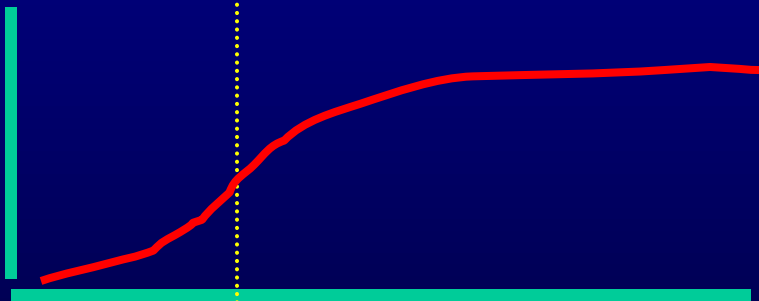
$$1 - H(t)$$

$h(t)$ & $H(t)$!!

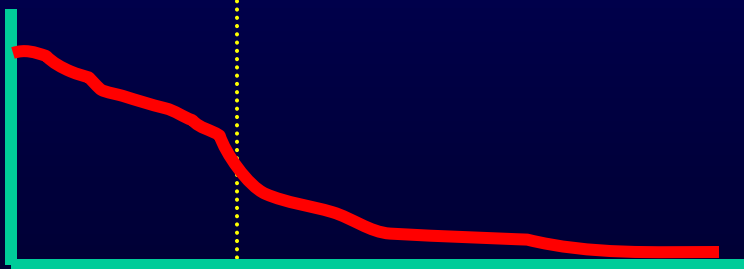
$h(t)$



$H(t)$



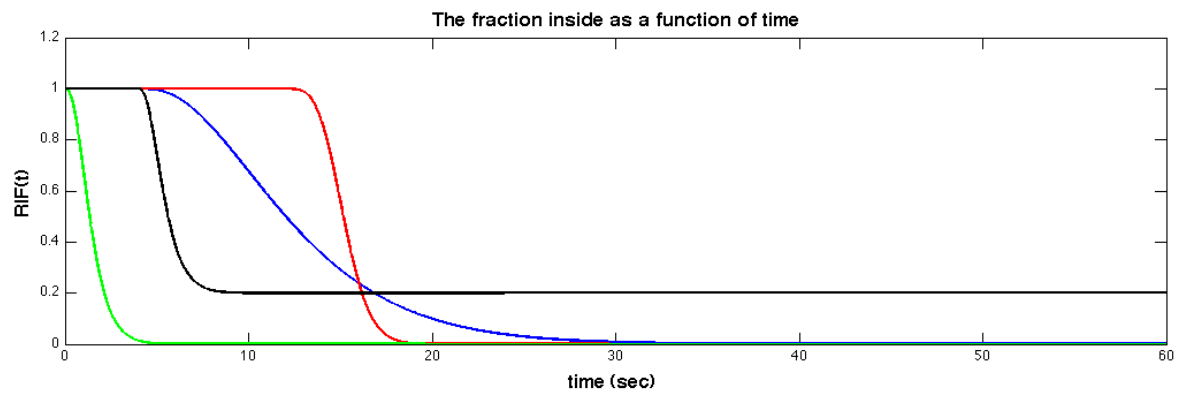
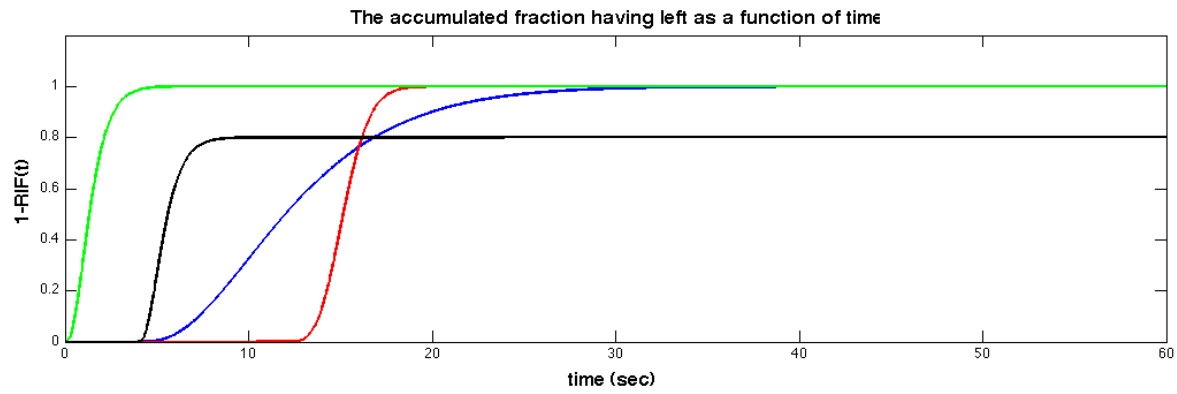
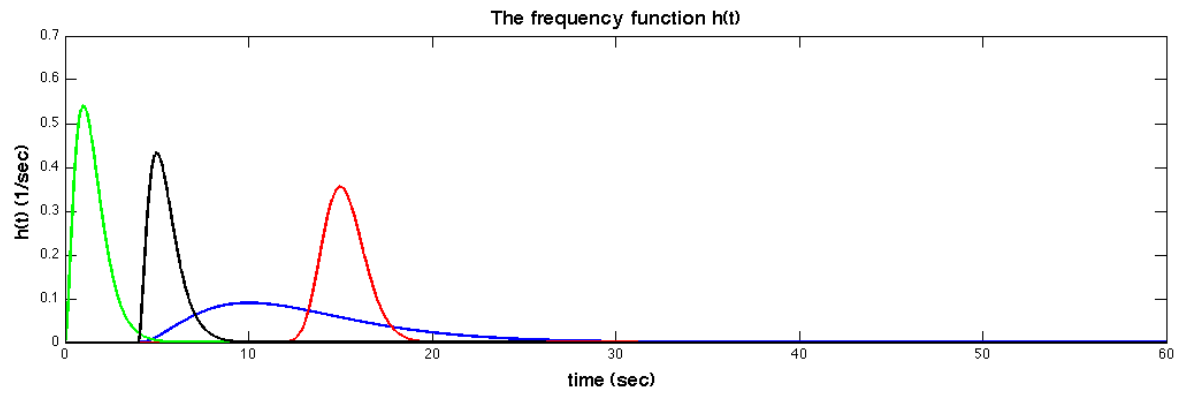
$1-H(t)$



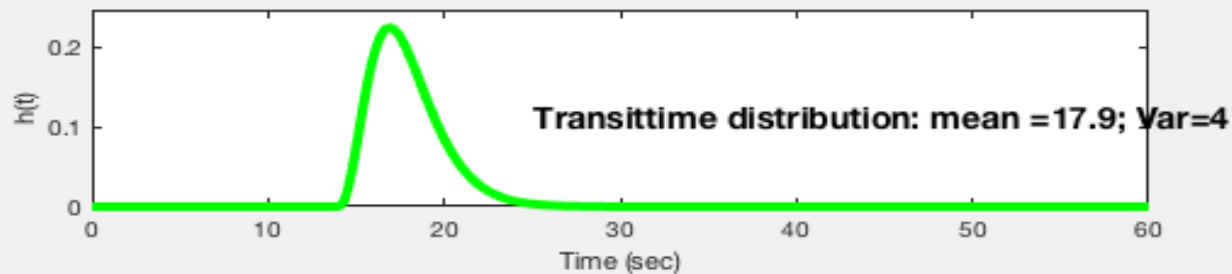
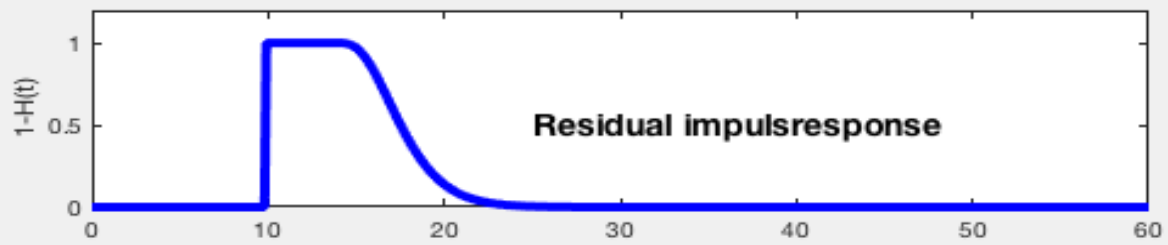
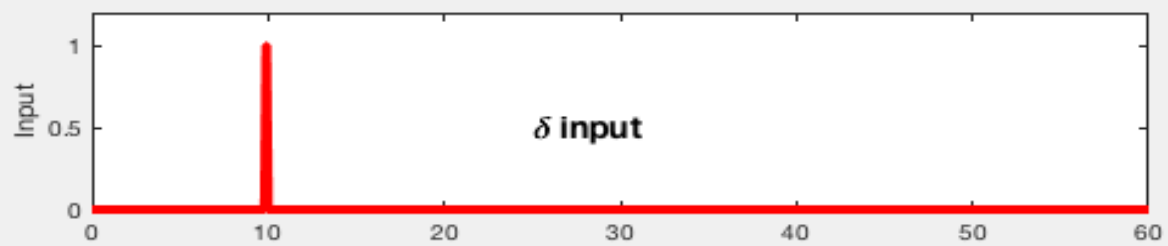
$$h(t) = \frac{dH(t)}{dt}$$

$$H(t) = \int_0^t h(\tau) d\tau$$

$$1 = H(t) + (1 - H(t))$$



CTH modeling



input



tissue



output

$h(t)$ & $H(t)$!!

$$\bar{t} = \int_0^{\infty} [1 - H(t)] dt$$

$$\bar{t} = \frac{1}{Q_0} (t_1 \cdot \Delta Q_1 + t_2 \cdot \Delta Q_2 + t_3 \cdot \Delta Q_3 + \dots + t_i \cdot \Delta Q_i + \dots)$$

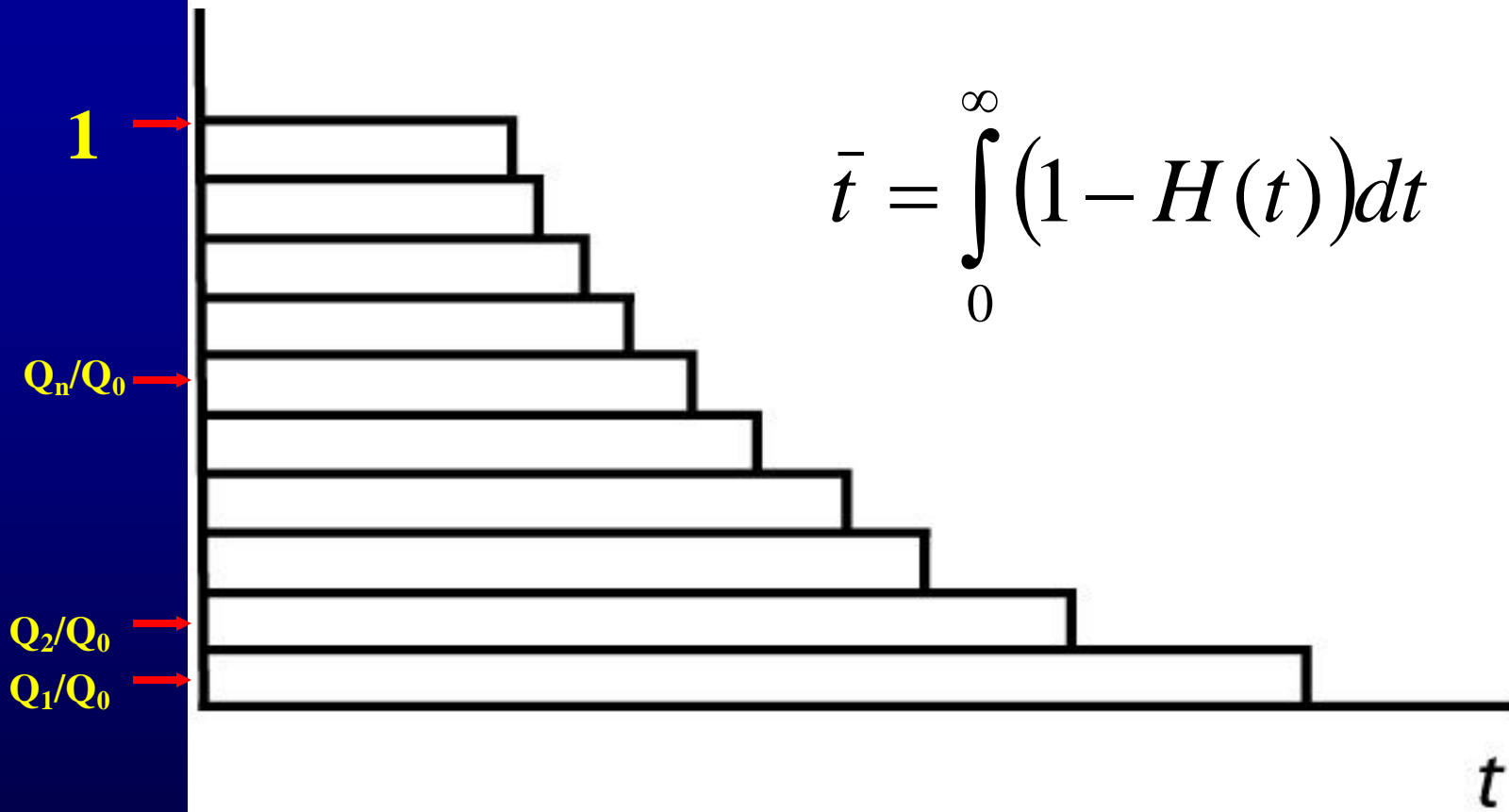


Illustration of transittimes and mean transittime estimated by residual measurement.

Mean transit time

The definition

$$\bar{t} = \frac{1}{Q_0} (t_1 \cdot \Delta Q_1 + t_2 \cdot \Delta Q_2 + t_3 \cdot \Delta Q_3 + \dots + t_i \cdot \Delta Q_i + \dots) \wedge Q_0 = \sum_i \Delta Q_i$$

$$\bar{t} = \frac{1}{Q_0} \sum_i t_i \cdot \Delta Q_i = \sum_i t_i \cdot \frac{\Delta Q_i}{Q_0} = \sum_i t_i \cdot \frac{\Delta Q_i}{Q_0 \cdot \Delta t} \xrightarrow{\text{lim}} \int_0^{\infty} t \cdot \frac{dQ(t)}{Q_0 \cdot dt} \cdot dt$$

Define the frequency function of transit times:

$$h(t) \equiv \frac{dQ(t)}{Q_0 \cdot dt}$$

$$[h(t)] = 1/s$$

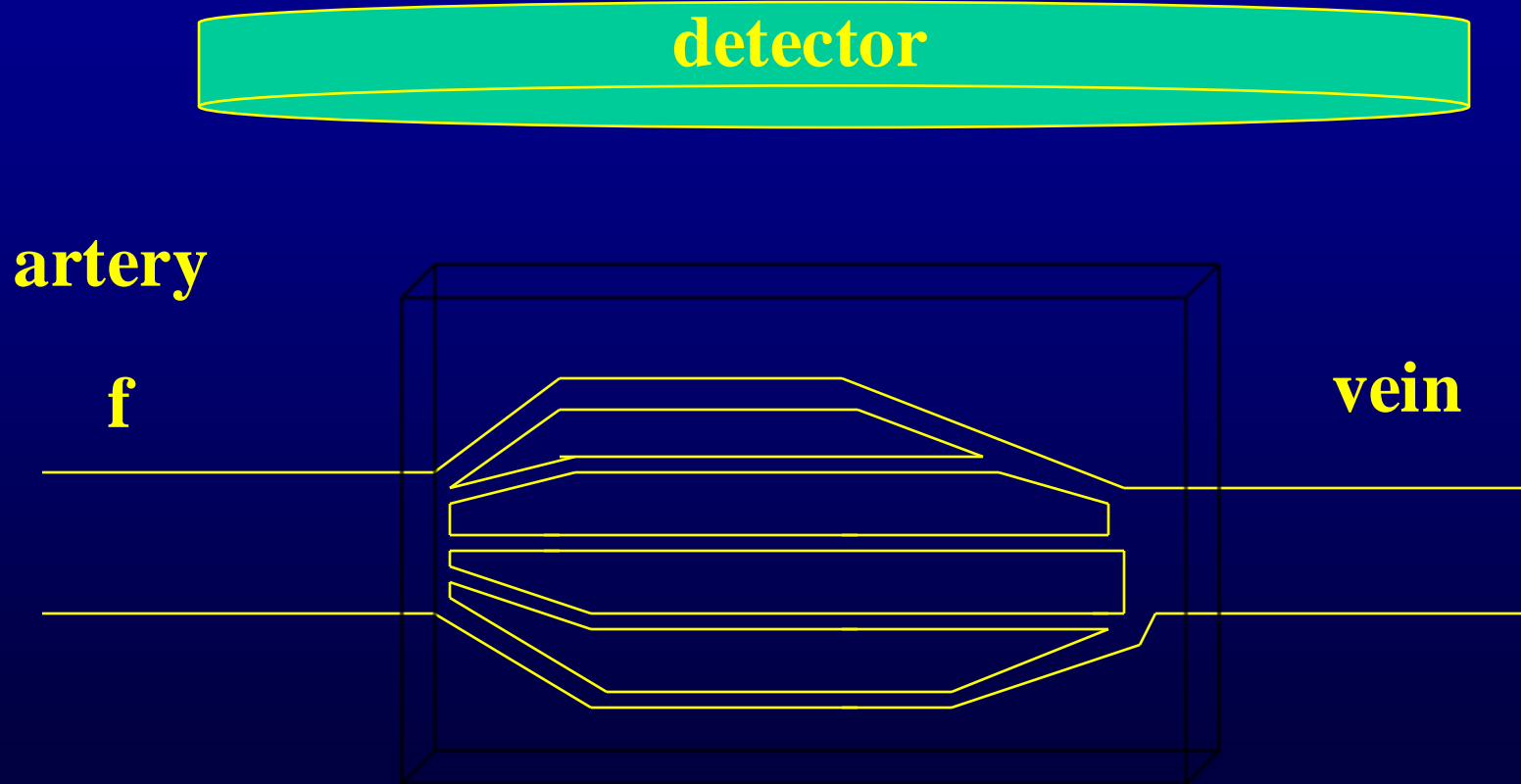
Break

Residue detection in CT-PET- SPECT- MRI

**The residue impulse response function:
The fraction remaining in the tissue at
time t after a brief (delta) input**

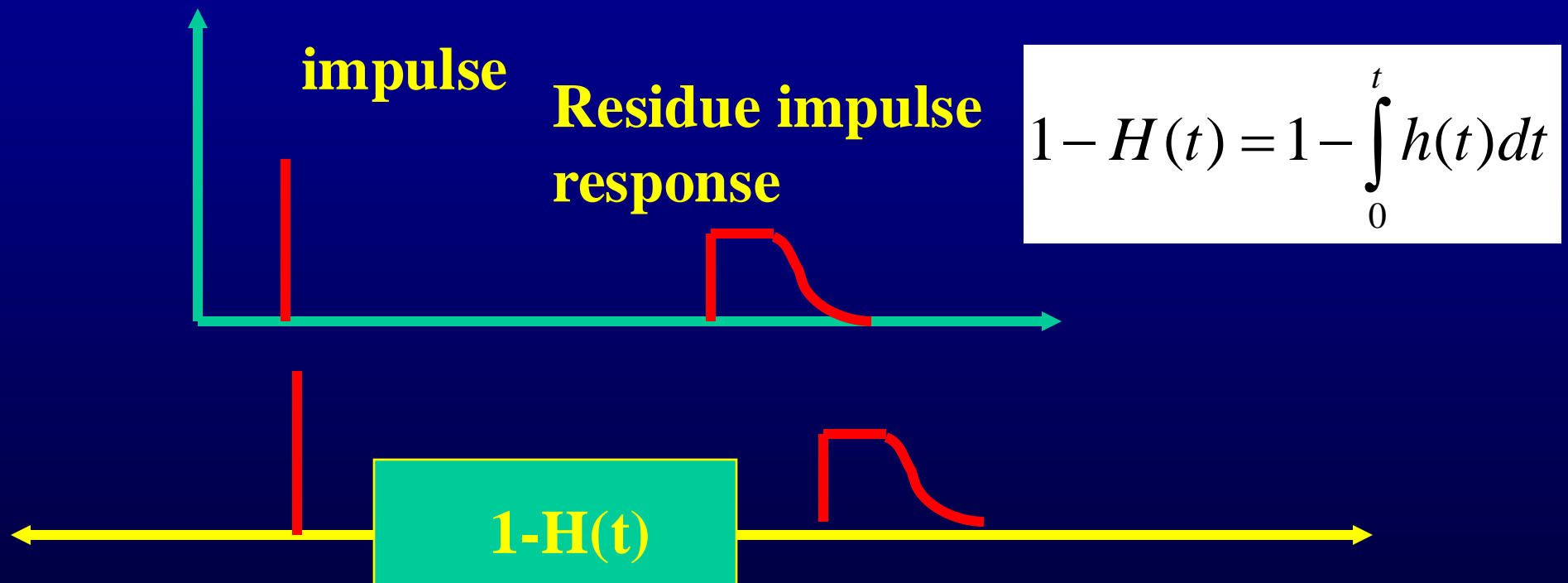
$$1 - H(t)$$

Measuring perfusion by an external registration: CT, SPECT, PET, MRI



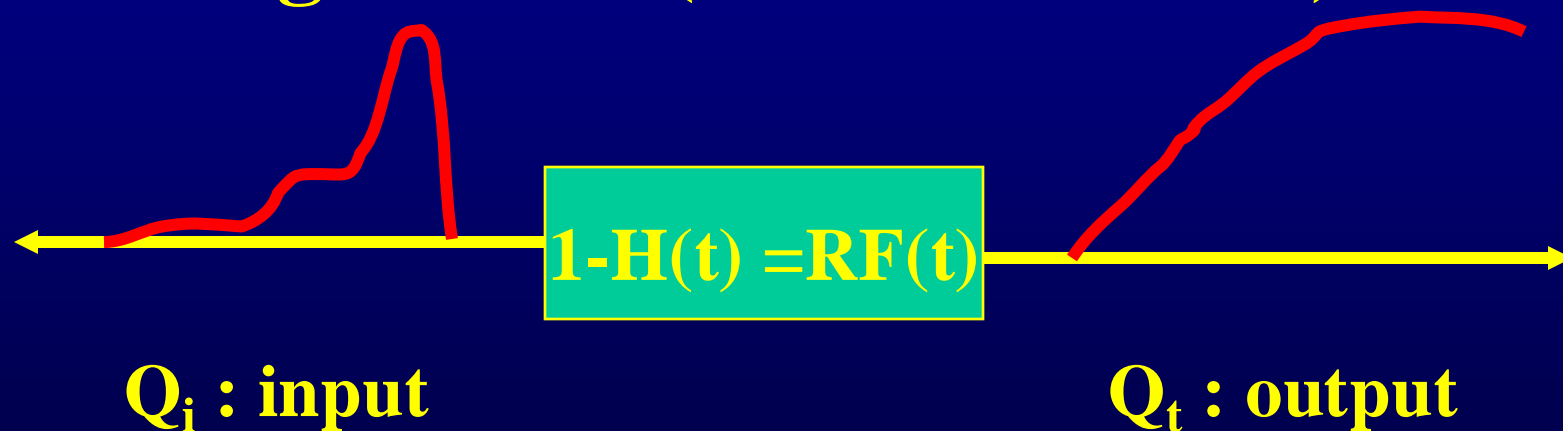
f: flow or perfusion [ml/min /100g]

The impulse response as measured by an external measuring system



Why is $1-H(t)$ interesting ?

Because it relates the input to the tracer amount in tissue in the case of the input not being a bolus (a deltafunction) !



$$C_t(t) = f C_a(t) \otimes RF(t) = f \int_0^t C_i(\tau) RF(t - \tau) d\tau$$



$$Q_i(0) = F C_i(0) \Delta\tau$$



The number which enters the system at time zero



$$Q_i(0) = F C_i(0) \Delta\tau$$



The total perfusion (Flow)



$$Q_i(0) = F C_i(0) \Delta\tau$$



The concentration of the tracer at the inlet at time zero



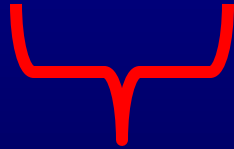
$$Q_i(0) = F C_i(0) \Delta\tau$$



Infinitively small time interval



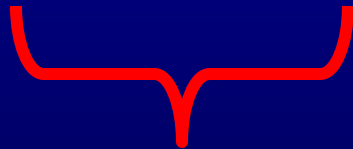
$$Q_i(0) = F C_i(0) \Delta\tau$$



The flux which enters the system at time zero



$$Q_i(0) = F C_i(0) \Delta\tau$$



The number which enters the system at time zero



$$Q_i(0) = F C_i(0) \Delta\tau$$

$$Q_t^1(t) = Q_i(0) RF(t)$$



The number (amount) of tracer in tissue as a function of time due to an input at time zero



$$Q_i(0) = F C_i(0) \Delta\tau$$

$$Q_t^1(t) = Q_i(0) RF(t)$$

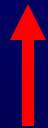


The relative number (amount) of tracer in tissue as a function of time due to an input at time zero



$$Q_i(0) = F C_i(0) \Delta\tau$$

$$Q_t^1(t) = Q_i(0) RF(t)$$



The number (amount) of tracer which enters the tissue at time zero



$$Q_i(0) = F C_i(0) \Delta\tau$$

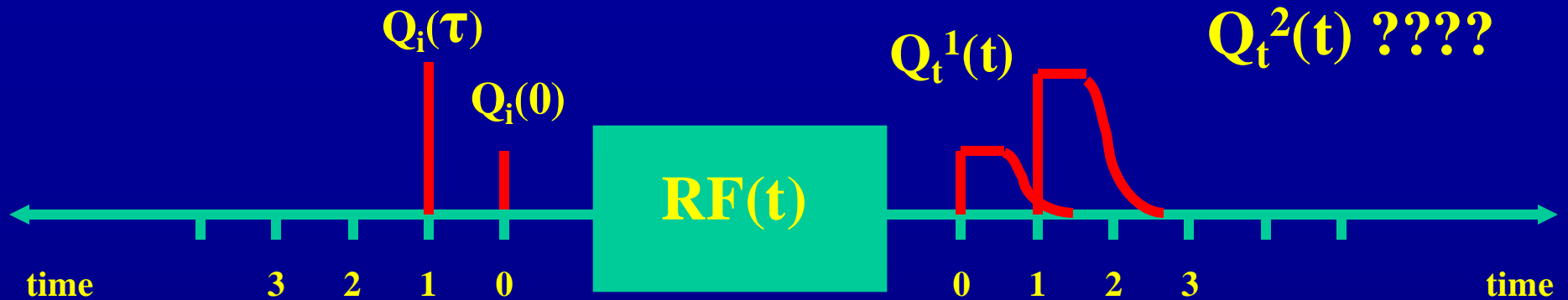
$$Q_t^1(t) = \underbrace{Q_i(0)}_{\text{input}} RF(t)$$

The number (amount) of tracer in tissue as a function of time due to an input at time zero



$$Q_i(0) = F C_i(0) \Delta\tau$$

$$Q_t^1(t) = Q_i(0) RF(t) = F C_i(0) \Delta\tau RF(t)$$

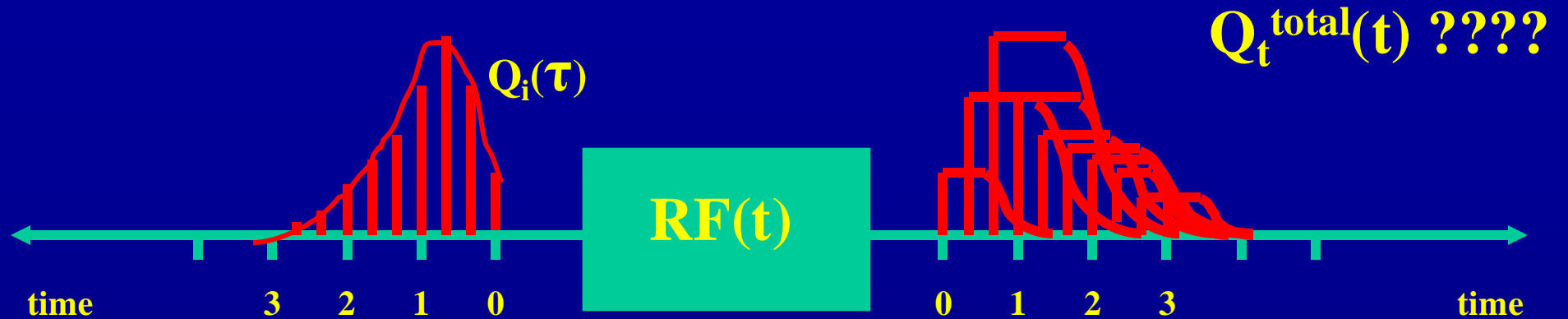


$$Q_i(0) = F C_i(0) \Delta\tau$$

$$Q_t^1(t) = Q_i(0) RF(t) = F C_i(0) \Delta\tau RF(t)$$

$$Q_t^2(t) = Q_i(\tau) RF(t-\tau) = F C_i(\tau) \Delta\tau RF(t-\tau)$$

Total amount in tissue at time t: $Q_t^{\text{total}}(t) = Q_t^1(t) + Q_t^2(t)$



Total amount in tissue at time t:

$$Q_t^{\text{total}}(t) = Q_t^1(t) + Q_t^2(t) + Q_t^3(t) + Q_t^4(t) + \dots \Rightarrow$$

$$Q_t^{\text{total}}(t) = \sum F C_i(\tau) RF(t - \tau) \Delta \tau$$

$$Q_t^{\text{total}}(t) = \int_0^t F C_i(\tau) RF(t - \tau) d\tau$$

$$Q_t^{total}(t) = \int_0^t F C_i(\tau) RF(t - \tau) d\tau$$

$$Q_t^{total}(t) = C_t(t) \textit{Weight}$$

$$C_t(t) = \frac{F}{W} \int_0^t C_i(\tau) RF(t - \tau) d\tau$$

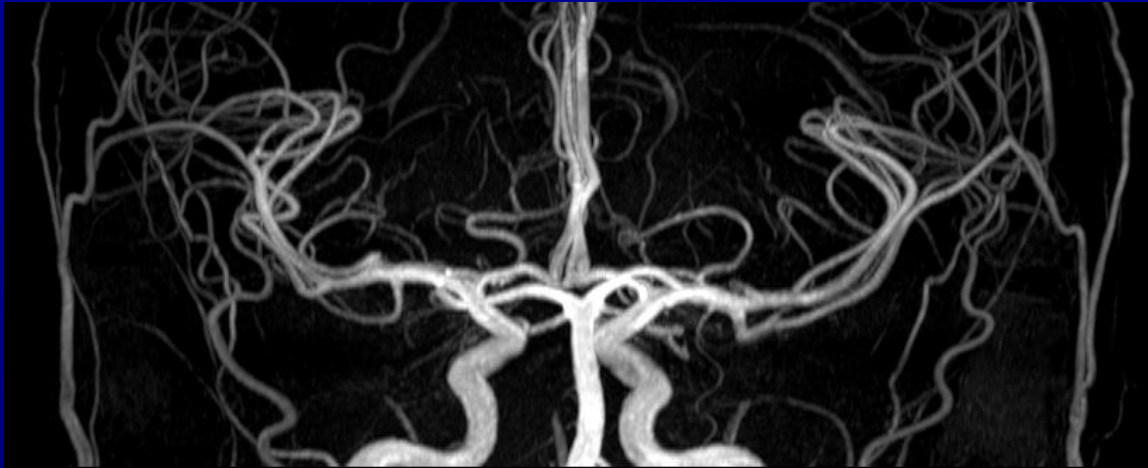
$$C_t(t) = f \int_0^t C_i(\tau) RF(t - \tau) d\tau$$

Convolution from the MATLAB point

- MATLAB

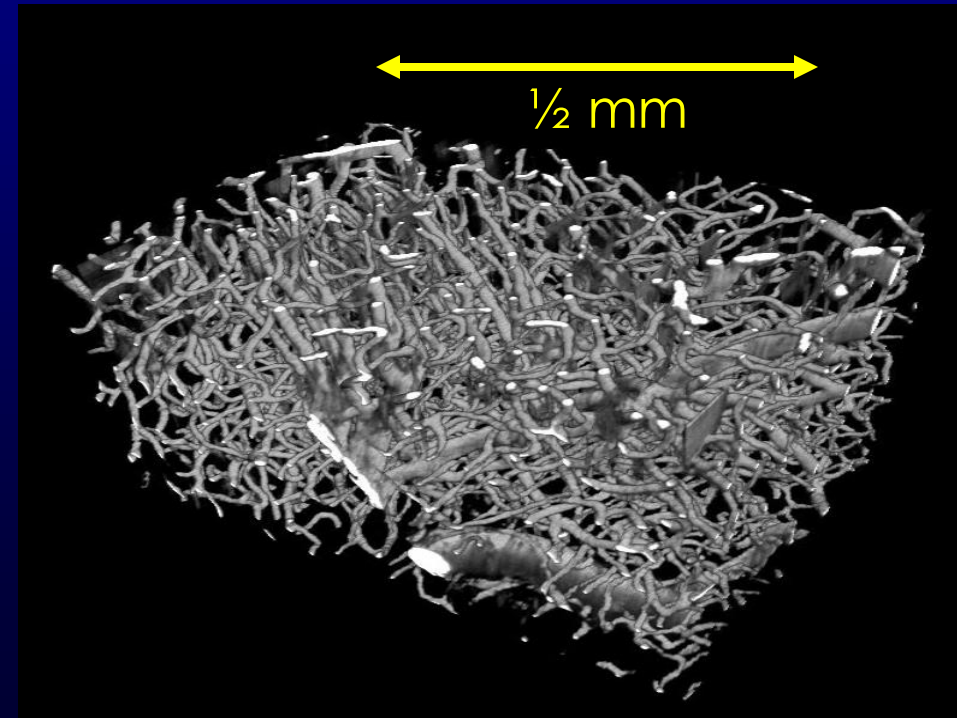
Break

What is perfusion?



Large vessels : flow

Perfusion: related to the microvascular system ~ the capillaries

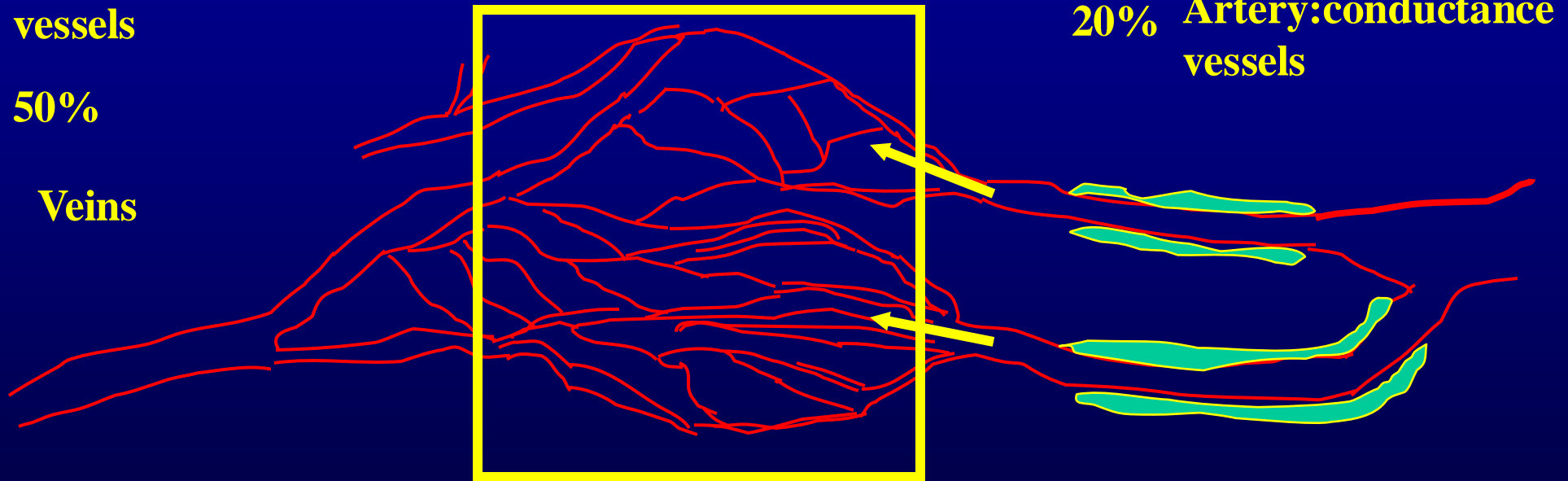


The vascular system of the brain and perfusion

Venules: capacity vessels

50%

Veins

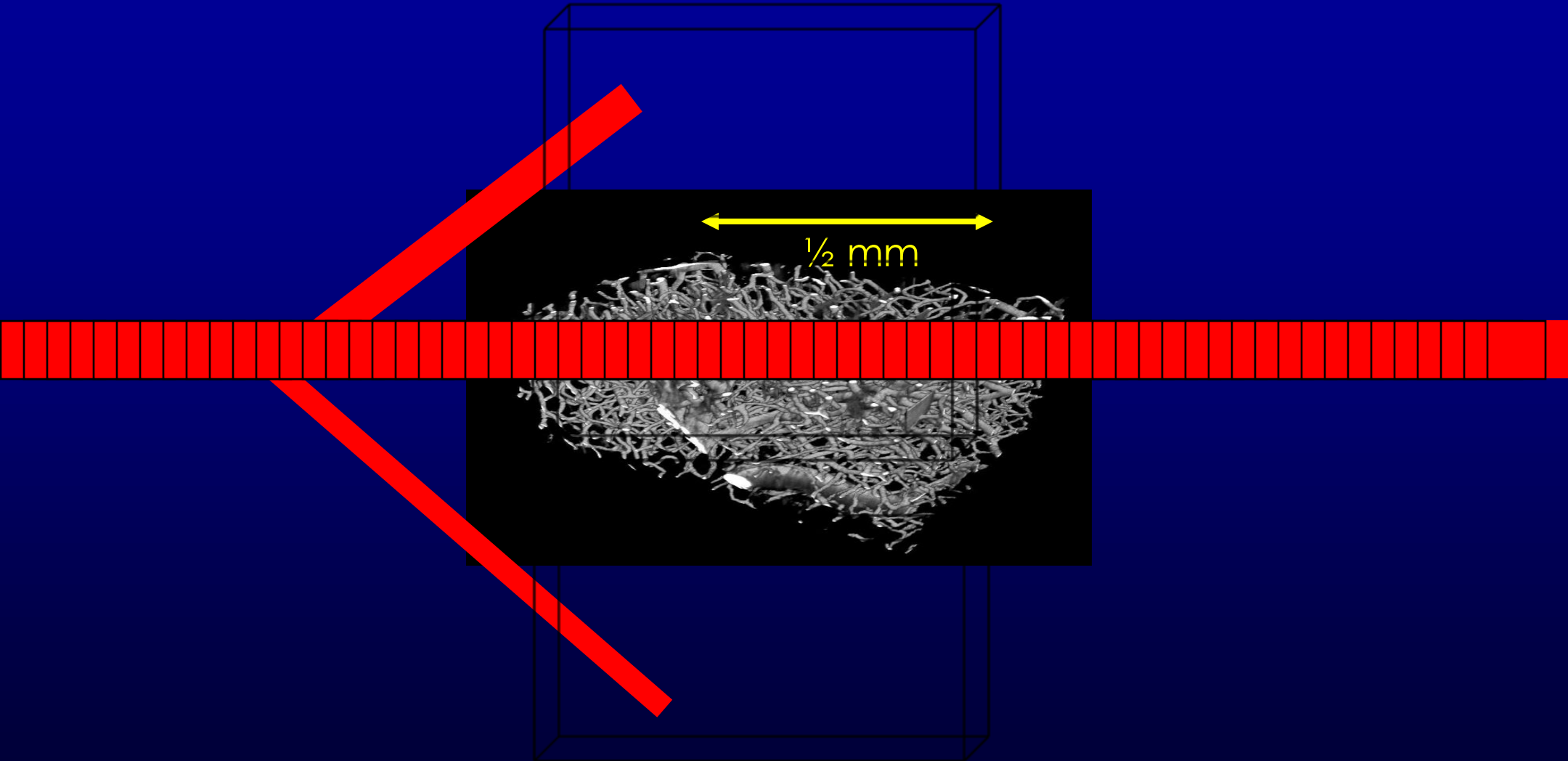


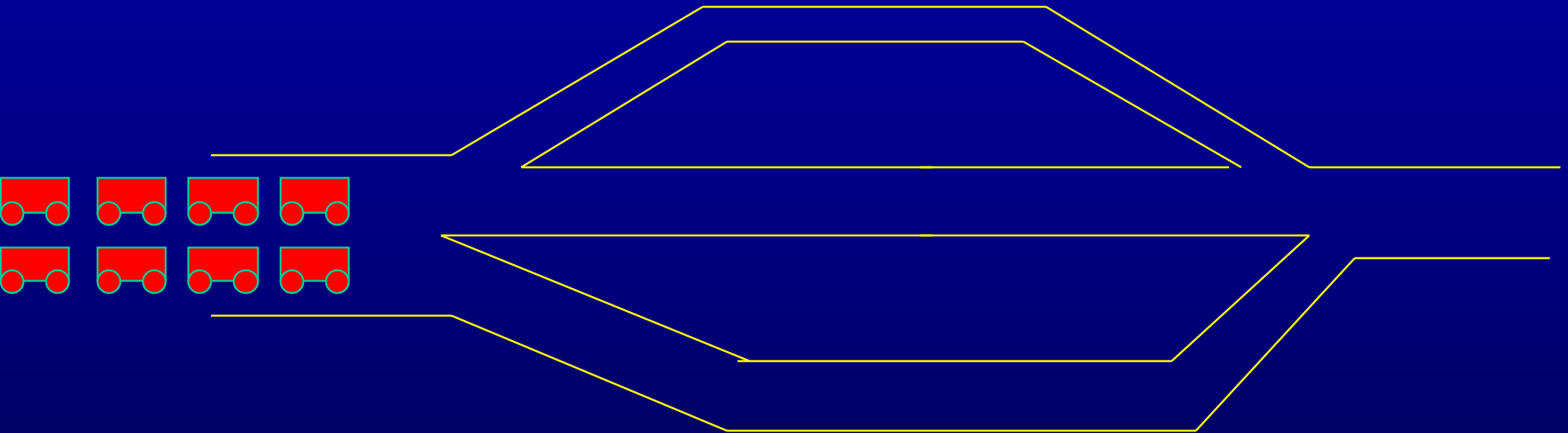
20% Artery: conductance vessels

30% Capillaries: exchange vessels ~ transport ~ diffusion

Arteriole: resistance vessels

Perfusion metrics in imaging: ml/min/100g or ml/min/100ml

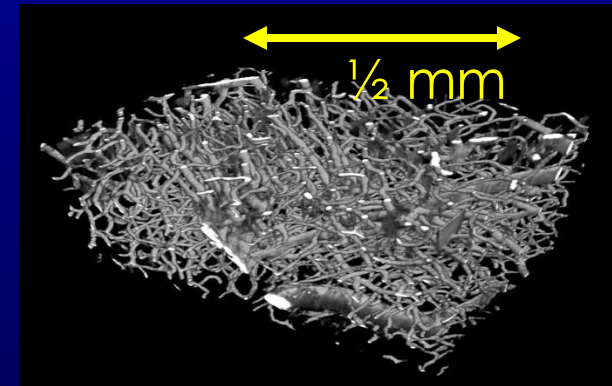




**Number of transport (ml) vehicles entering
100 ml tissue pr. time unit::
20 - 80 ml/min/100 ml tissue volume**

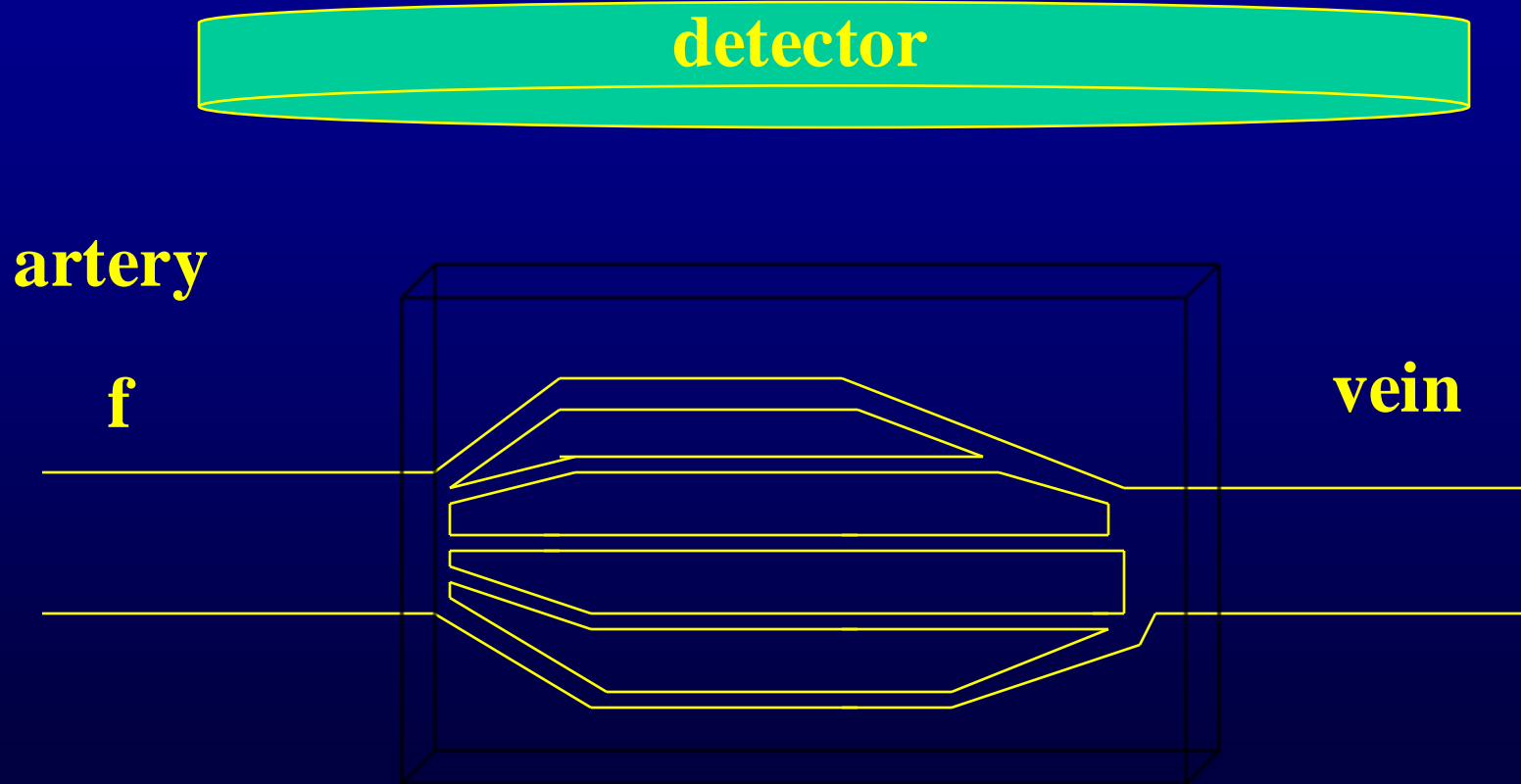
Important metrics

- **Perfusion:** – f [ml/min/100g] or [ml/min/100ml]
- **Brain Perfusion ('flow') :** Cerebral blood flow **CBF** [ml/100g/min]
- **Cerebral blood volume: CBV** [ml/100g]



- **Mean transit time: MTT** [s]
- **Blood brain permeability: PS product** [ml/100g/min]

Measuring perfusion by an external registration: CT, SPECT, PET, MRI



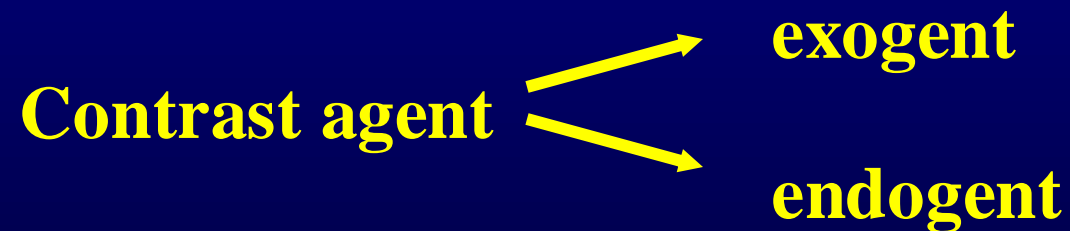
f: perfusion in [ml/min /100g]

How can it be measured ?

Add a contrast agent carried by the blood to the tissue

How can it be measured ?

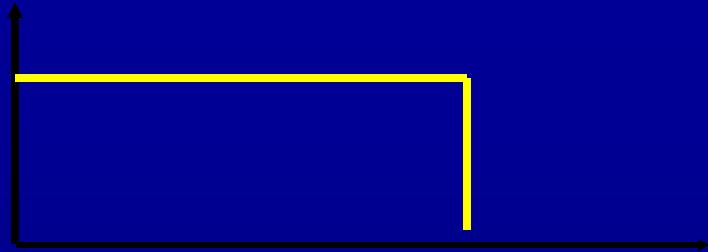
Add a contrast agent carried by the blood to the tissue



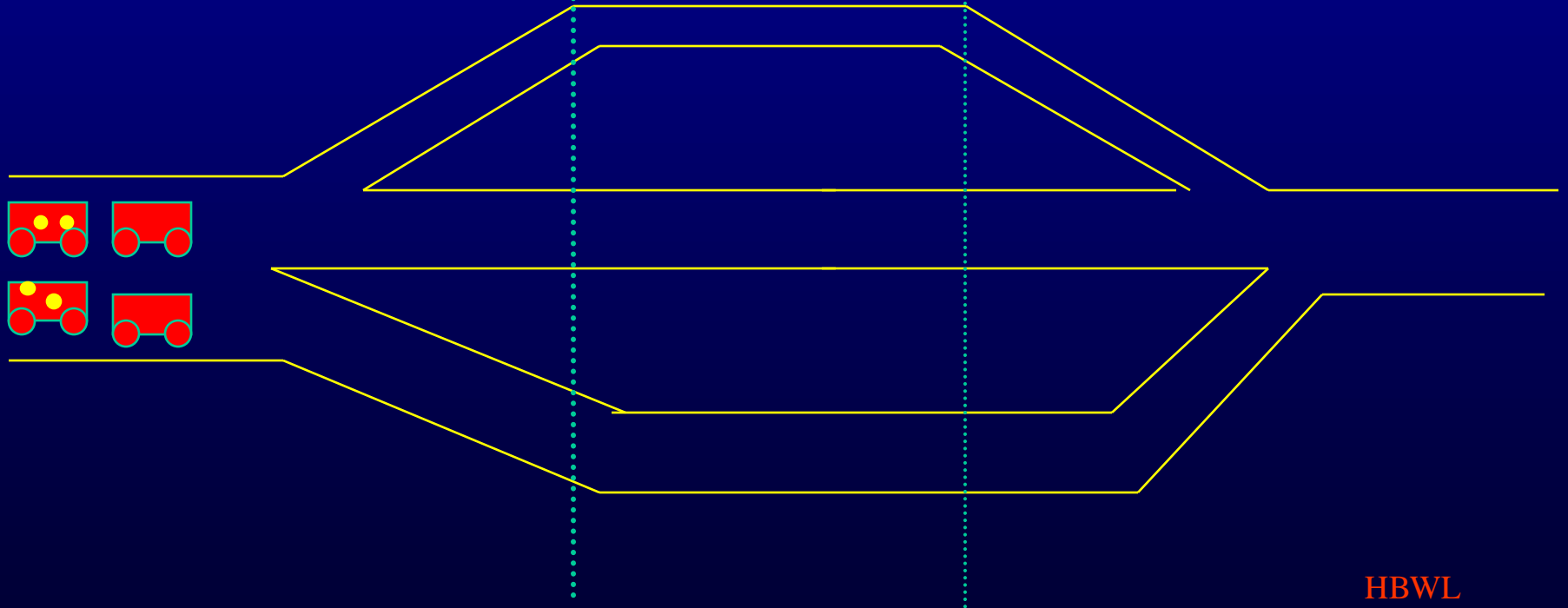
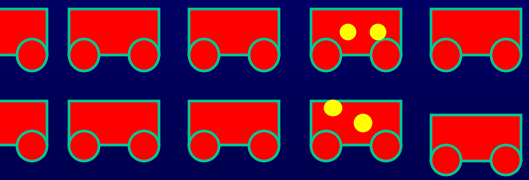
The complicated part: Single bolus injection and external registration



Signal $\approx C_{tis}$



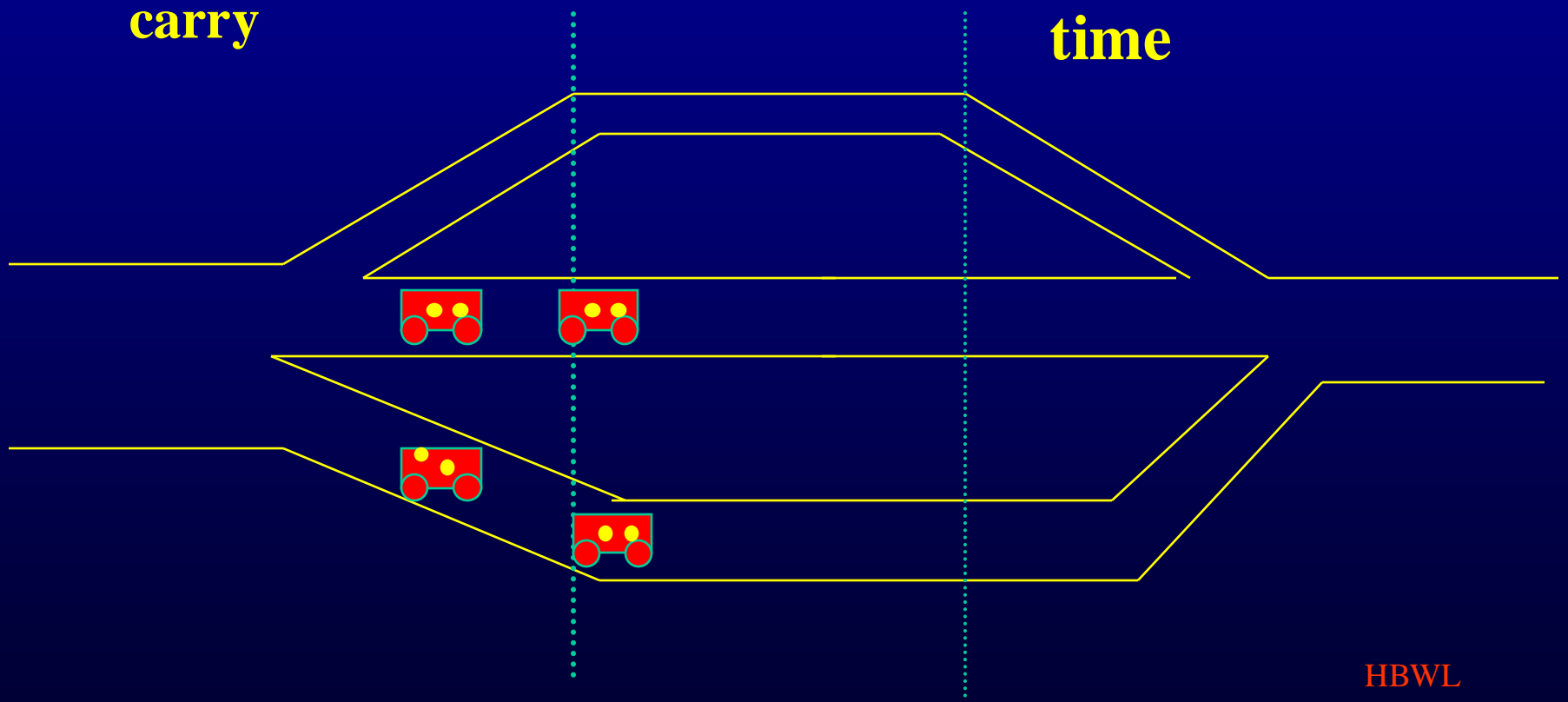
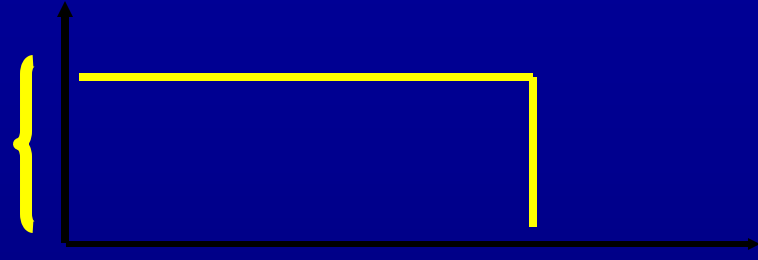
time



$$\text{Signal} \approx C_{\text{tis}}(0) : f C_a(0)$$

Perfusion (f) = vehicles/min
ml/min x

Conc (C_a) = the cargo they carry
mmol/ml carry



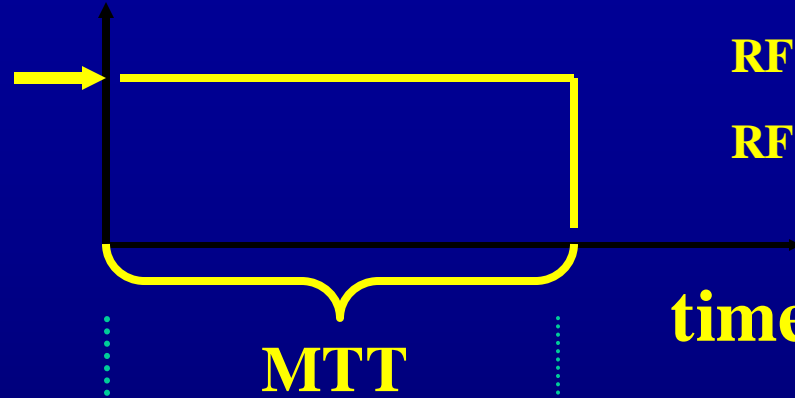
$$C_{tis}(t) = f C_a(0) \Delta t \mathbf{RF}(t)$$

$$C_{tis}(0) \doteq f C_a(0) \Delta t$$

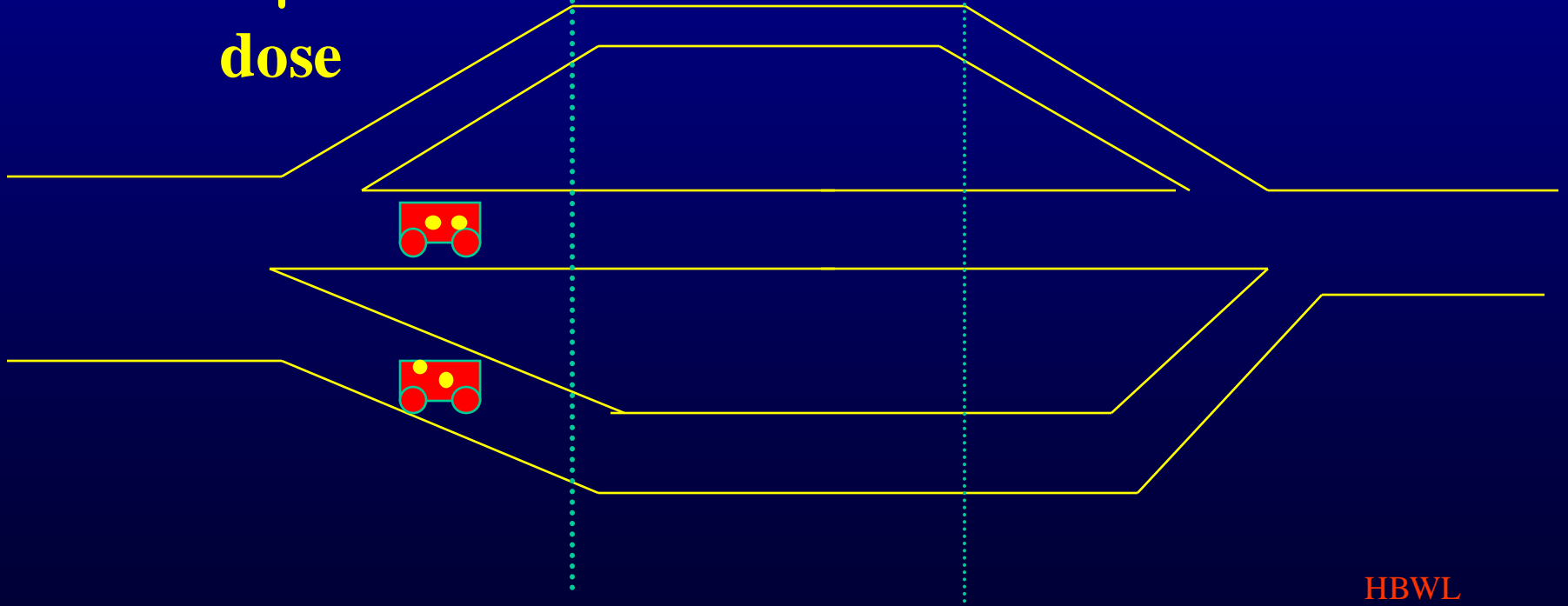
$f C_a(0) \Delta t$
flux
dose

$\mathbf{RF}(t) = 1$ for $t < \mathbf{MTT}$

$\mathbf{RF}(t) = 0$ for $t > \mathbf{MTT}$



time



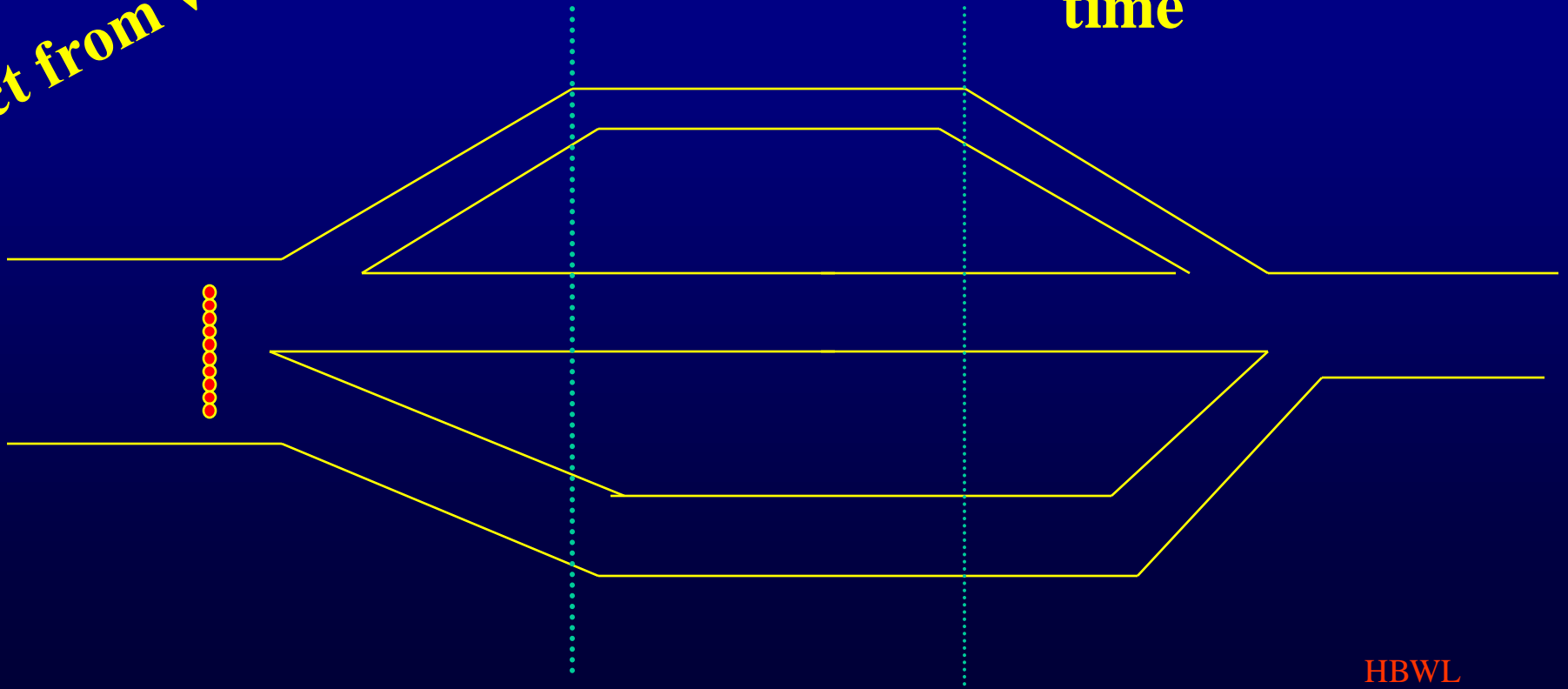
$$C_{tis}(t) = f C_a(0) \Delta t RF(t)$$

$$C_{tis}(0) = f C_a(0) \Delta t \rightarrow$$



time

Abstract from vehicles



Summing up: direct short bolus

Measure the
tissue conc

Measure the input conc i.e.
input function

Scanner signal : $C_{tis}(t) = f C_a(0) \Delta t RF(t)$

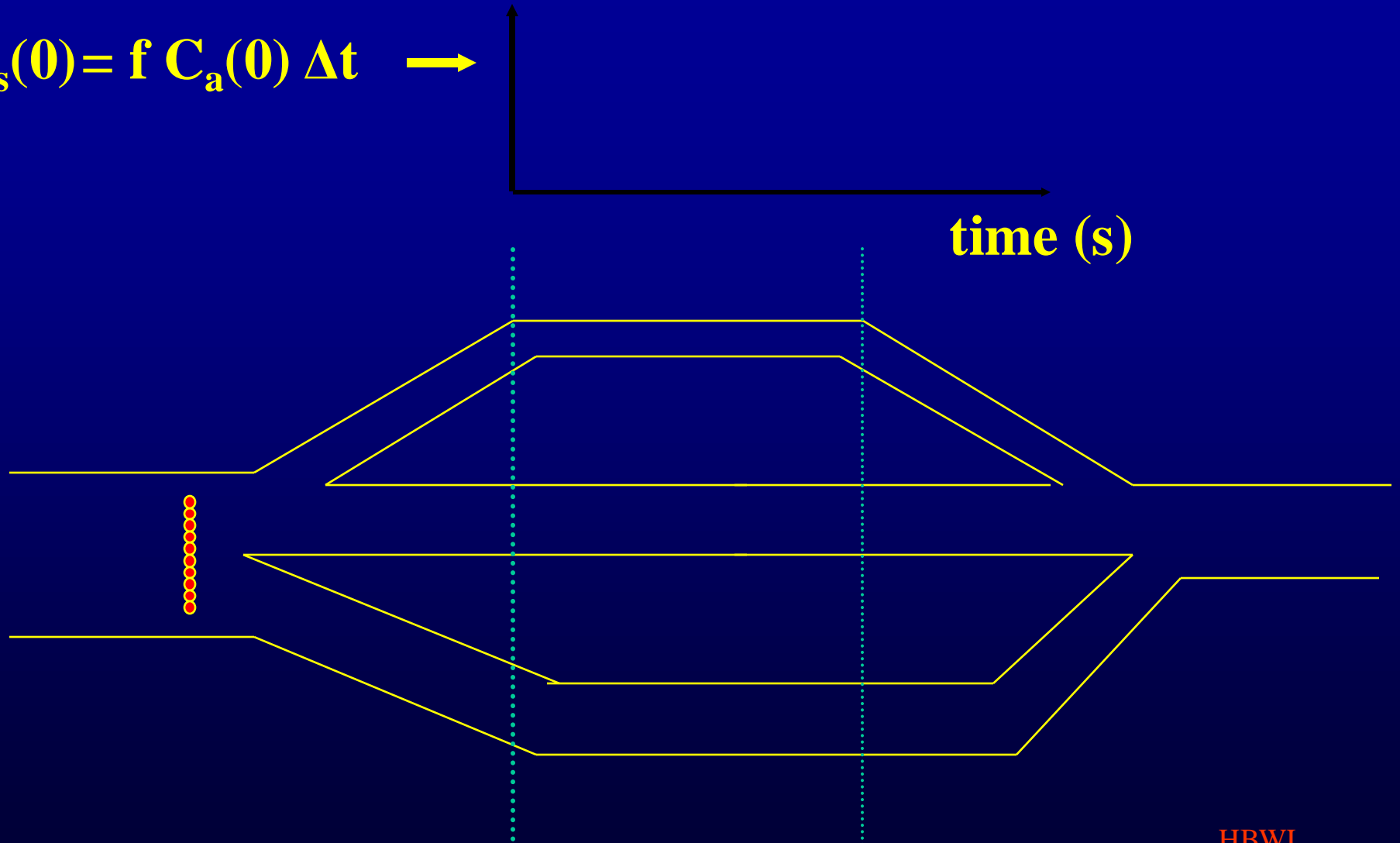
Estimate f and $RF(t)$

Different perfusion tracers behaves differently



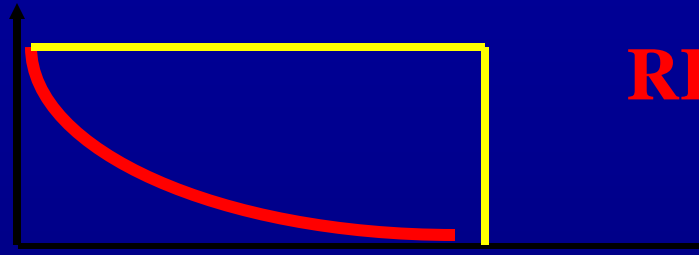
$$C_{tis}(t) = f C_a(0) RF(t)$$

$$C_{tis}(0) = f C_a(0) \Delta t \rightarrow$$



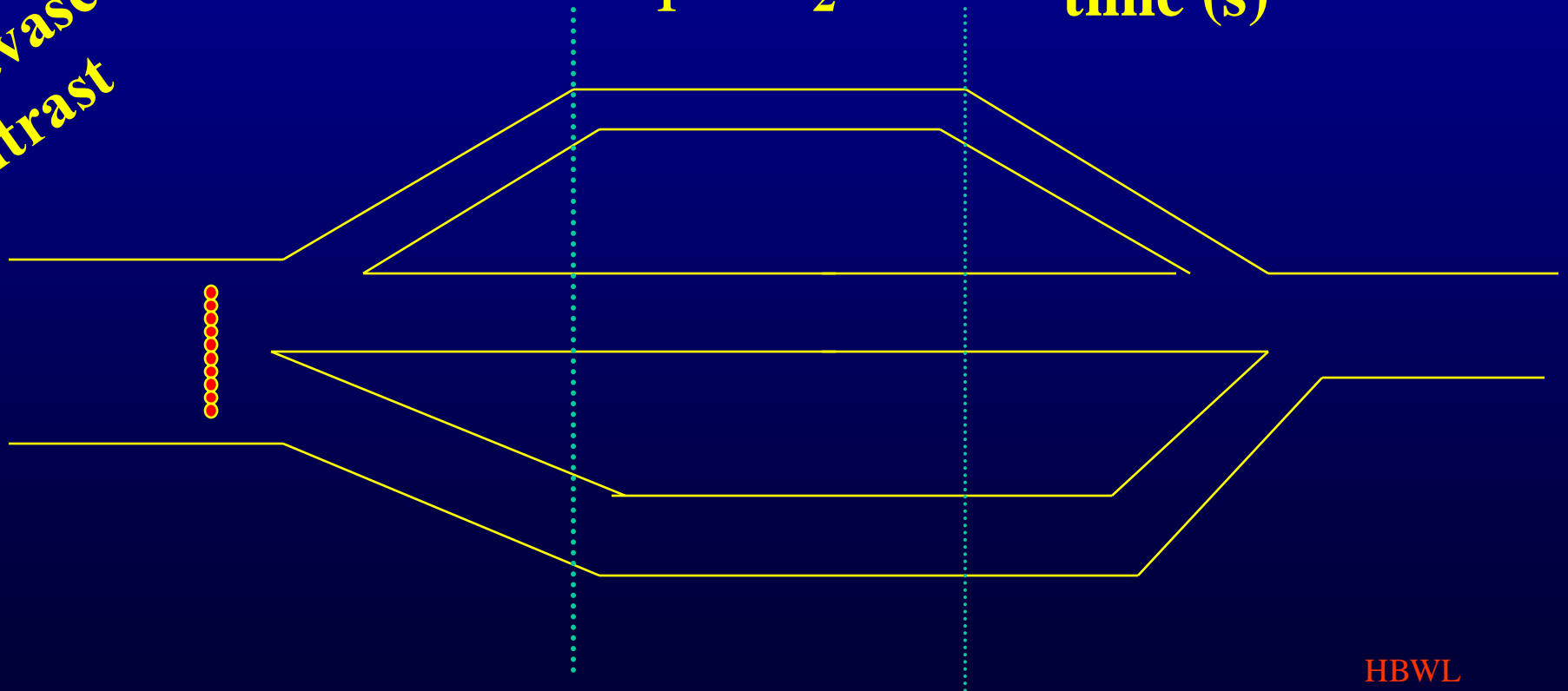
$$C_{tis}(t) = f C_a(0) \Delta t RF(t)$$

$$C_{tis}(0) = f C_a(0) \Delta t \rightarrow$$



$$RF(t) = e^{-k_2 t}$$

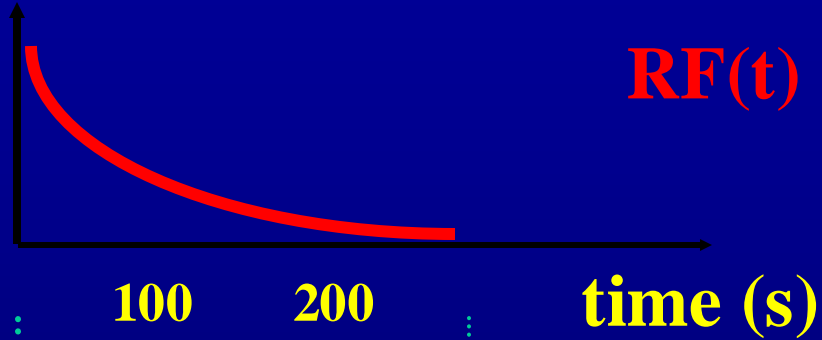
**Intravascular
contrast**



$$C_{tis}(t) = f C_a(0) \Delta t RF(t)$$

$$C_{tis}(0) = f C_a(0) \Delta t \rightarrow$$

$$RF(t) = e^{-k_2 t}$$



Freely
difusional
contrast agent

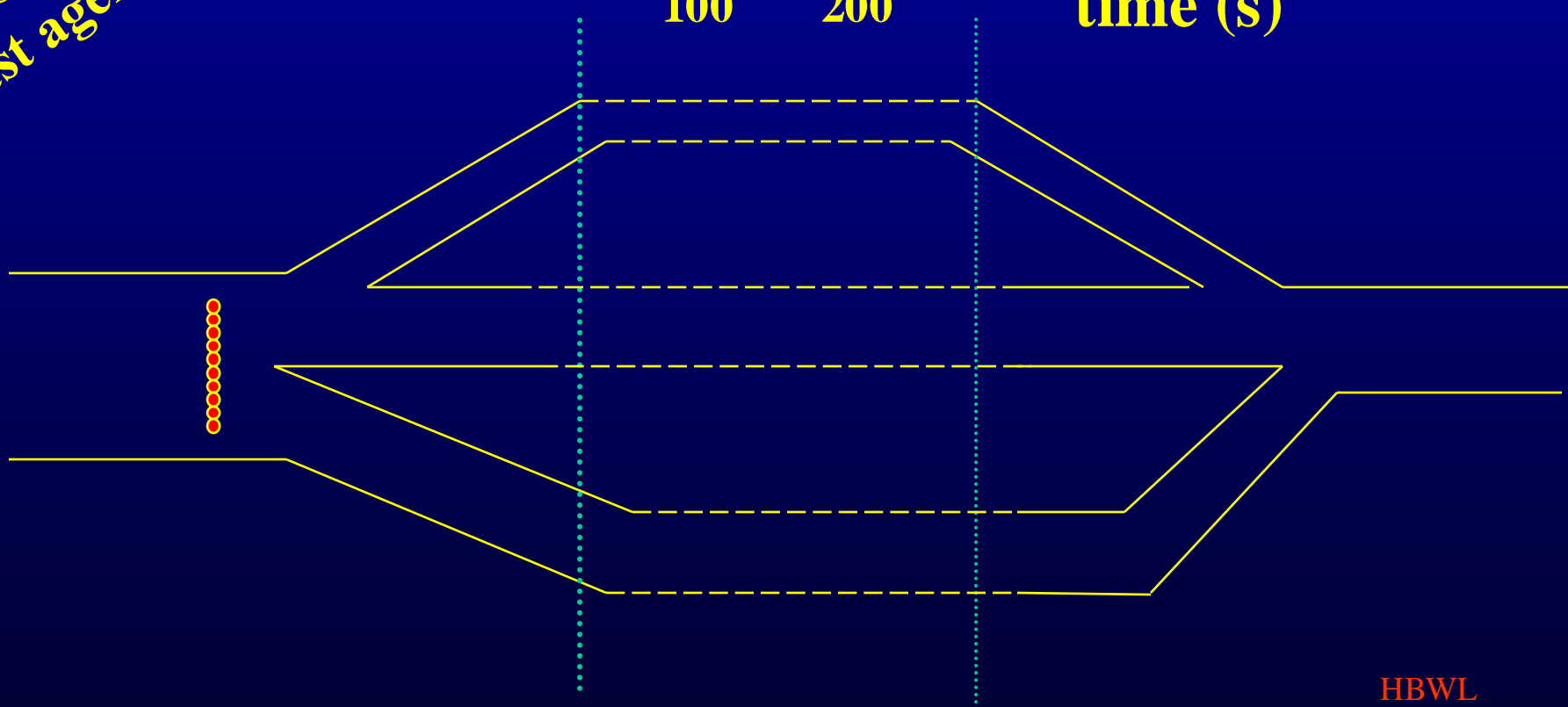
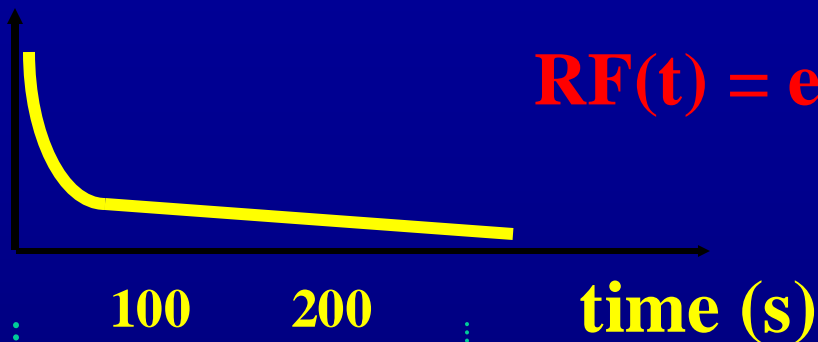


$$C_{tis}(t) = f C_a(0) \Delta t RF(t)$$

$$C_{tis}(0) = f C_a(0) \Delta t \rightarrow$$

$$RF(t) = e^{-k_2 t} + e^{-k_3 t}$$

Extravascular
contrast agent



The residue impulse response function $RF(t)$

$RF(t)$: the fraction of the injected dose remaining in the tissue (voxel) as a function of time

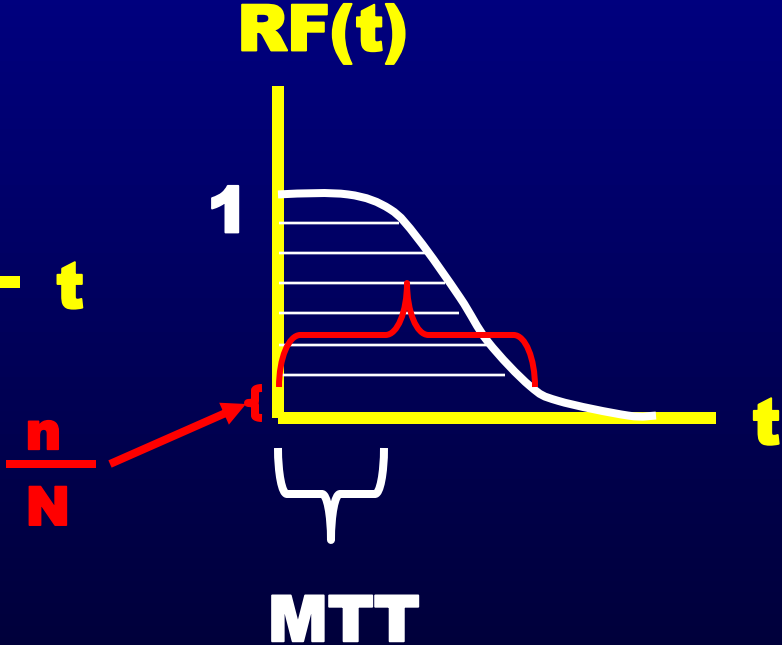
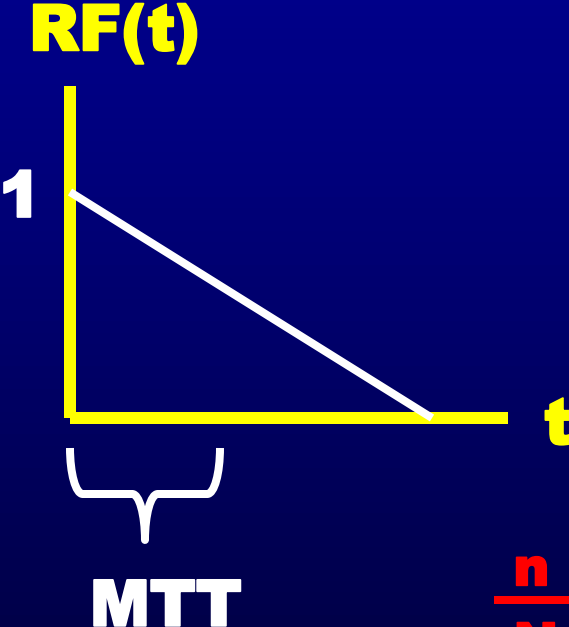
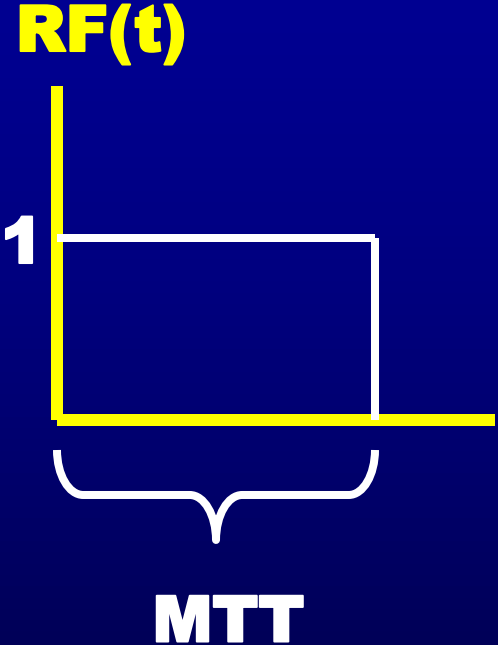
Mean transit time : MTT

$$MTT = \int_0^{\infty} RF(t)$$

Mean transit time : MTT

$$MTT = \int_0^{\infty} RF(t) dt$$

$$MTT = \sum \frac{n}{N} t$$



Generally

Perfusion: f

Distribution vol: V_d

Mean transit time: MTT

$$f = \frac{V_d}{MTT}$$

For an intravascular contrast agent, e.g. in brain MRI we have:

Brain perfusion: CBF

Brain blood volume: CBV

Mean transit time: MTT

$$CBF = \frac{CBV}{MTT}$$

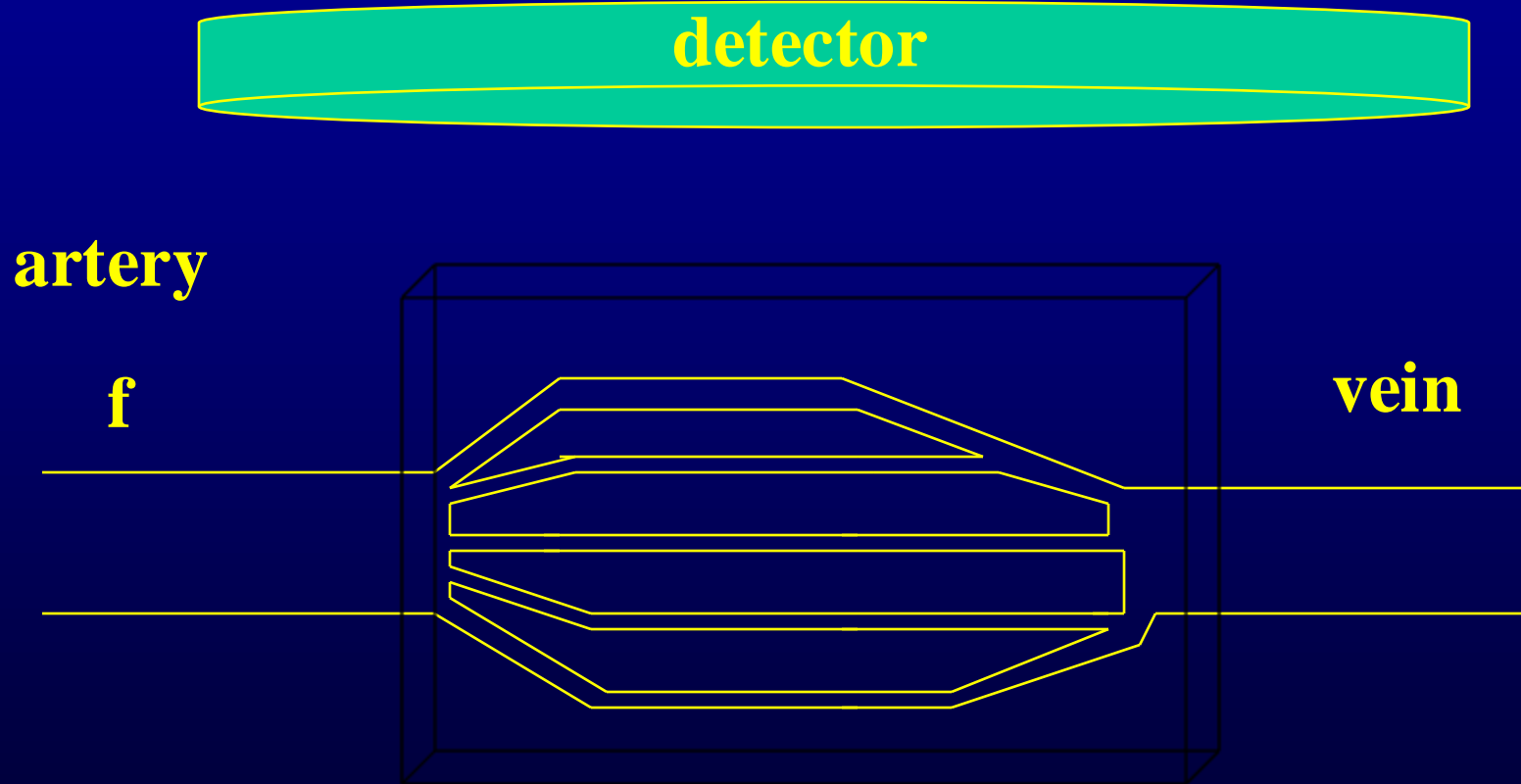
The really complicated part: Deconvolution



We cannot apply a bolus directly in the tissue !



Measuring perfusion by an external registration: CT, SPECT, PET, MRI



f: flow or perfusion [ml/min /100g]

The final step

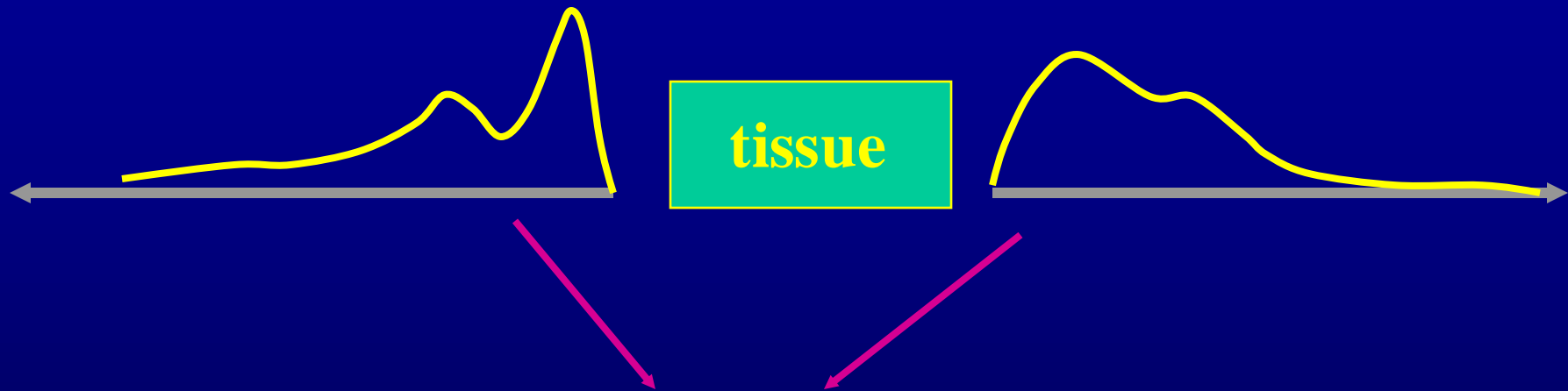
We cannot apply a bolus directly in the tissue !

Input :

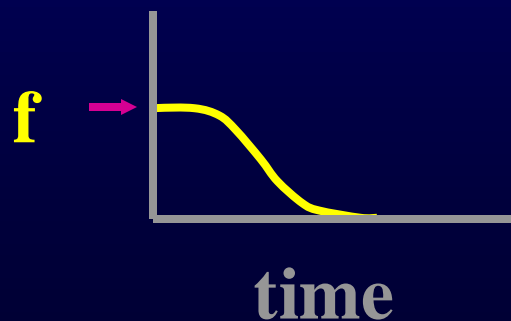
$$C_a(t)$$

Tissue enhancement :

$$C_{tis}(t) = \int_0^{\infty} f C_a(\tau) RF(t - \tau) d\tau$$



Deconvolution :
find f $RF(t)$



Input : $C_a(t)$

Tissue enhancement :

$$C_{tis}(t) = f C_a(0) RF(t) \Delta t$$



Input : $C_a(t)$

Tissue enhancement :

$$C_{tis}(t) = ?$$



Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = ?$$



If the linearity of the system exist

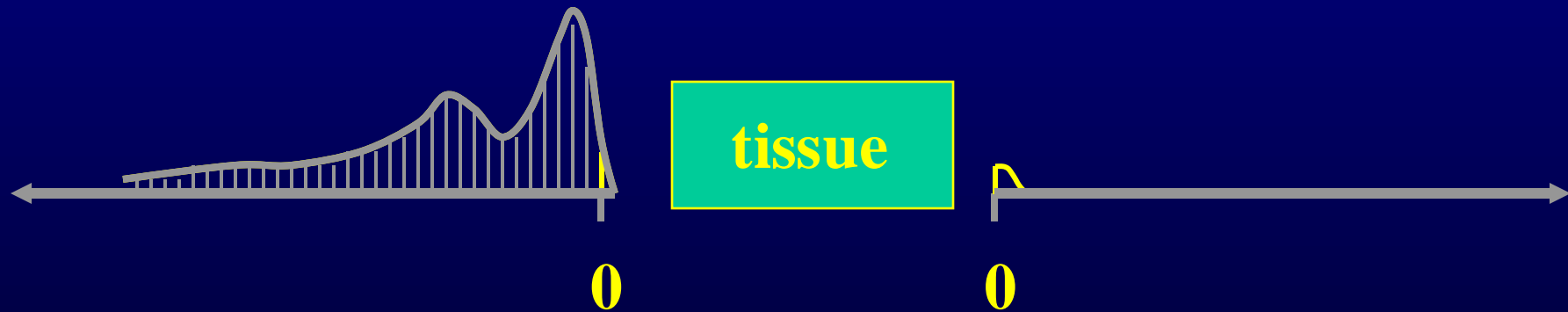


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(0) RF(t - 0) \Delta\tau$$

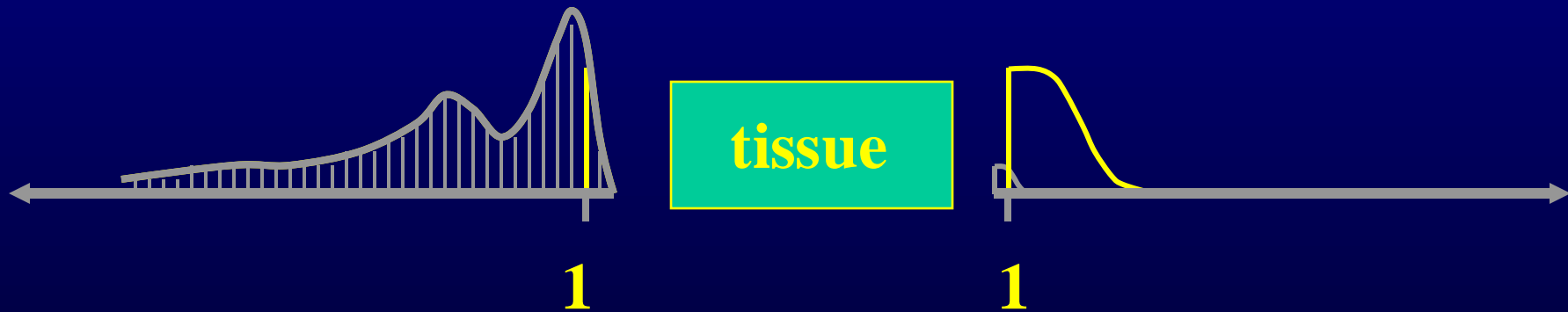


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(1) RF(t - 1) \Delta\tau$$

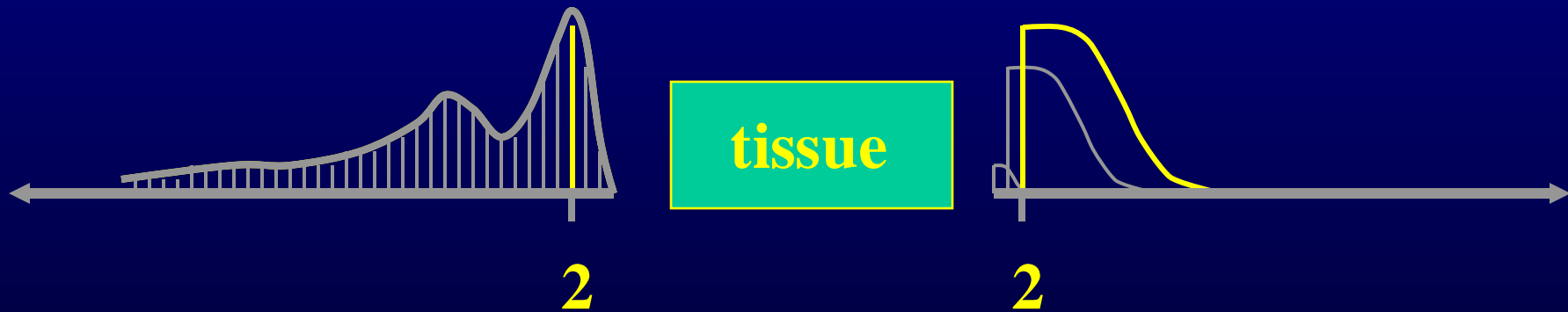


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(t) RF(t - t) \Delta\tau$$

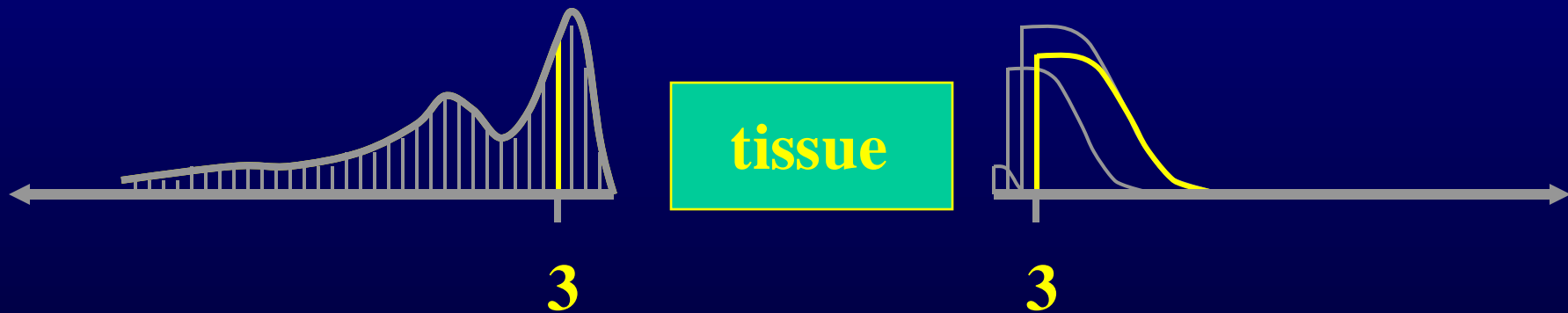


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(\tau) RF(t - \tau) \Delta\tau$$

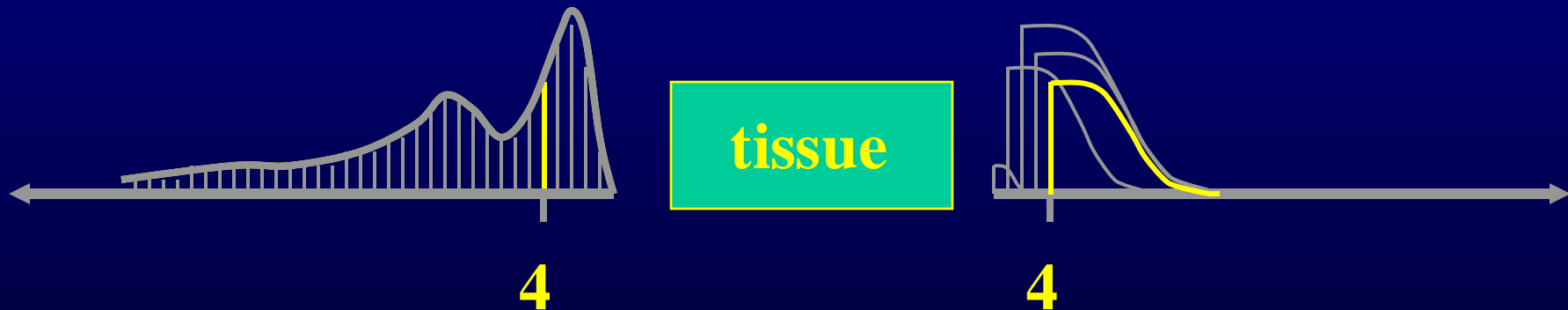


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(4) RF(t - 4) \Delta\tau$$



Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(5) RF(t - 5) \Delta\tau$$

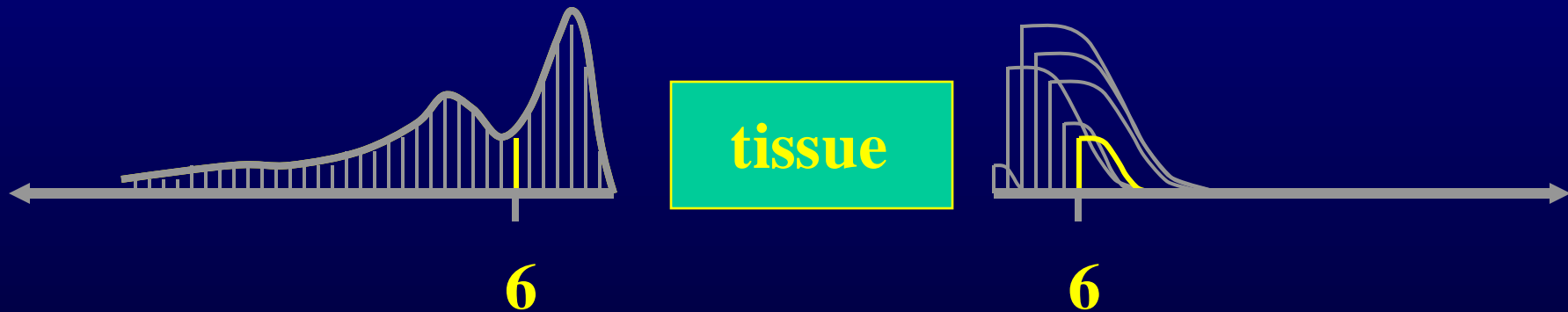


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(\tau) RF(t - \tau) \Delta\tau$$

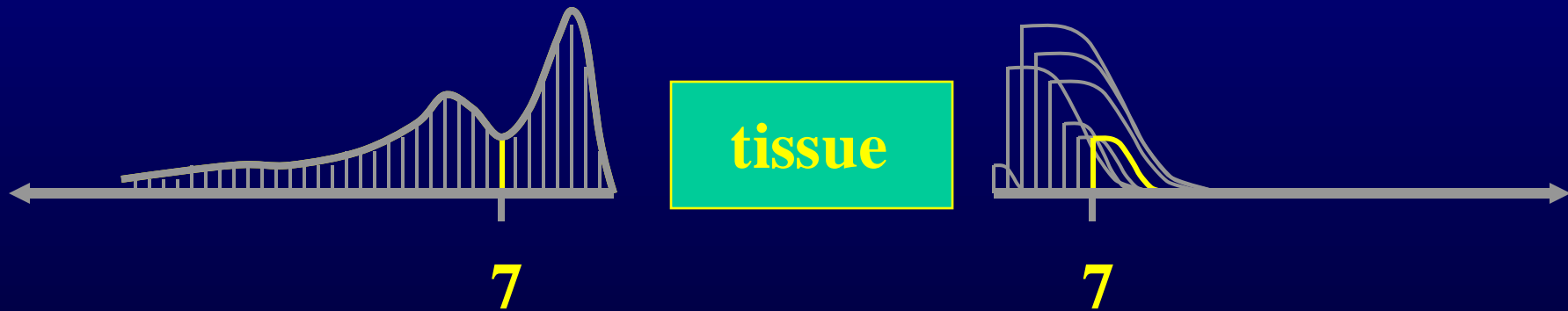


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(\tau) RF(t - \tau) \Delta\tau$$

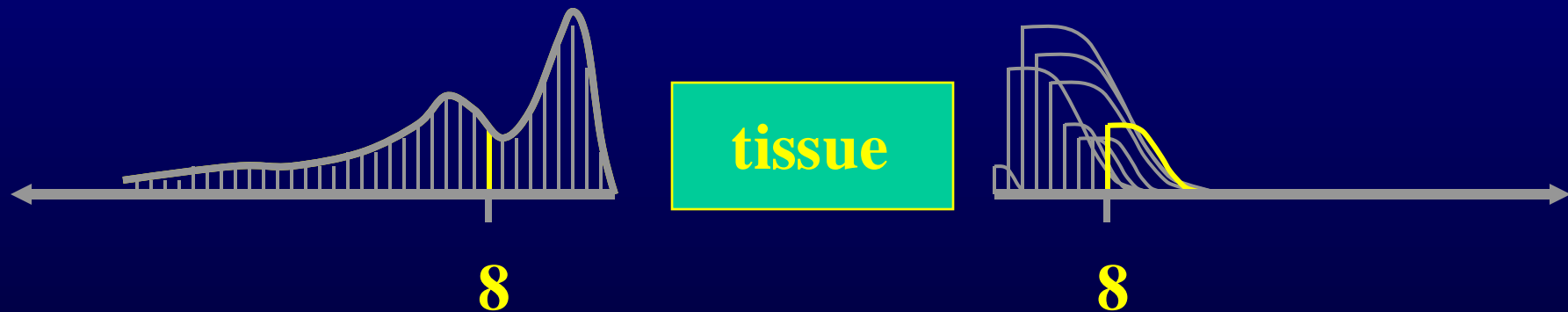


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(\delta) RF(t - \delta) \Delta\tau$$

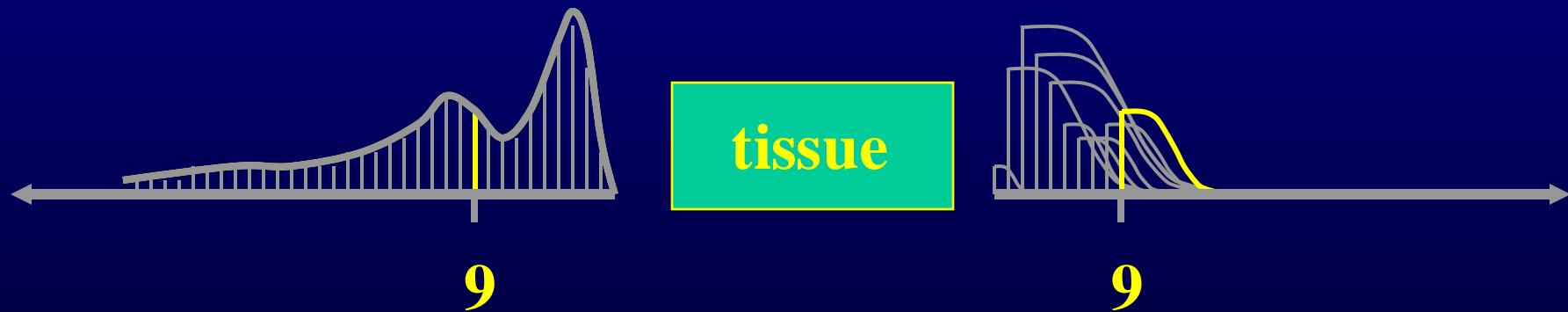


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(\theta) RF(t - \theta) \Delta\tau$$

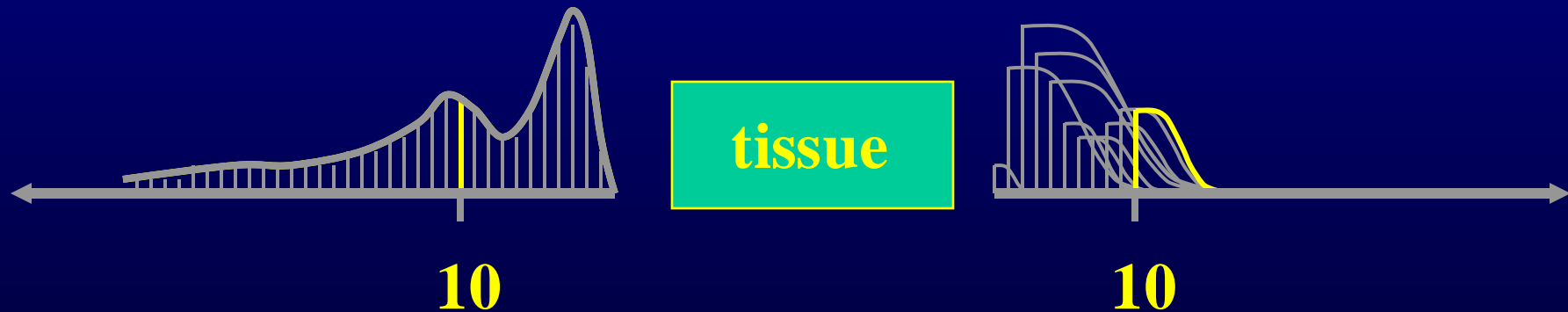


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(10) RF(t - 10) \Delta\tau$$

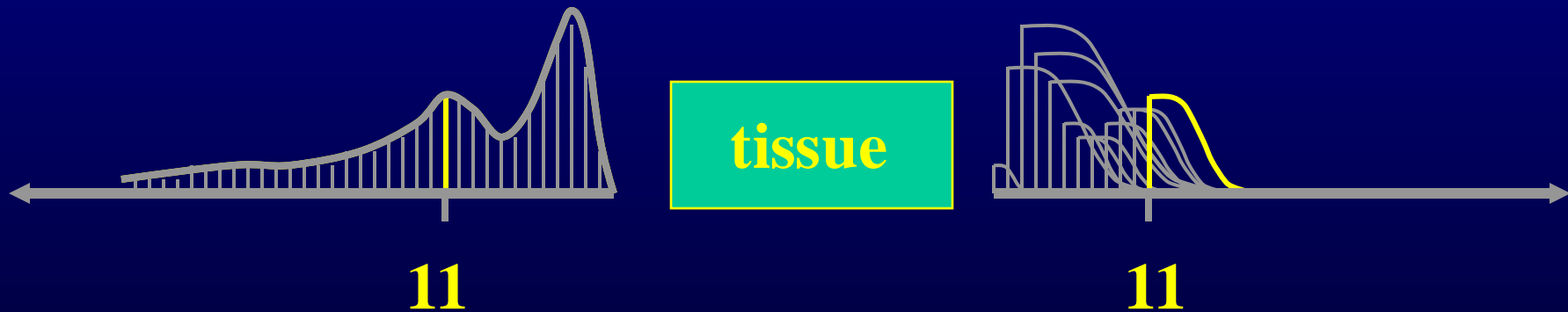


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(t) RF(t - t) \Delta\tau$$

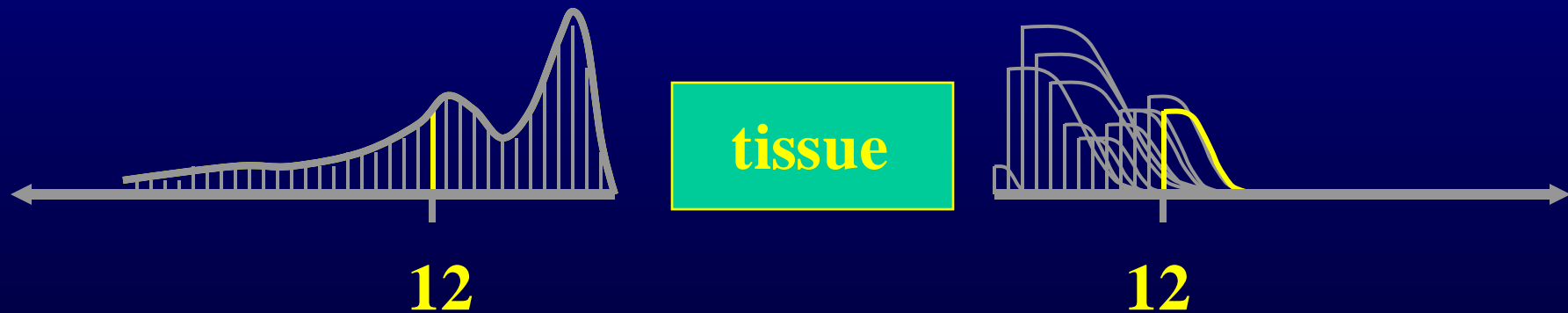


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(12) RF(t - 12) \Delta\tau$$

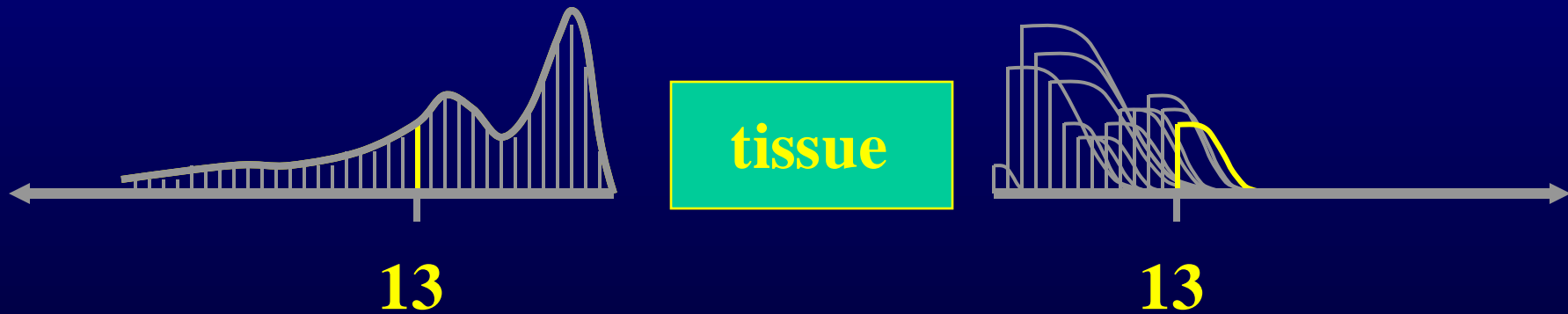


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(13) RF(t - 13) \Delta\tau$$

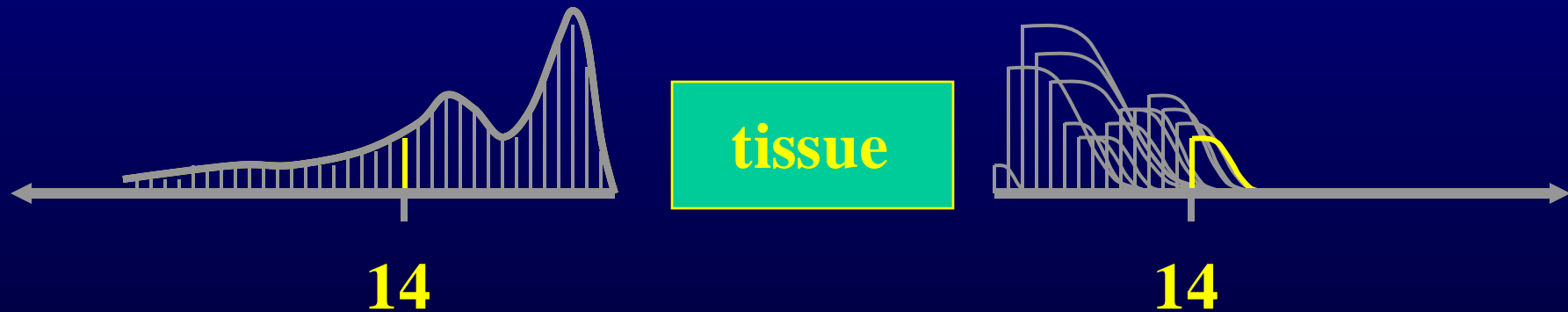


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(14) RF(t - 14) \Delta\tau$$

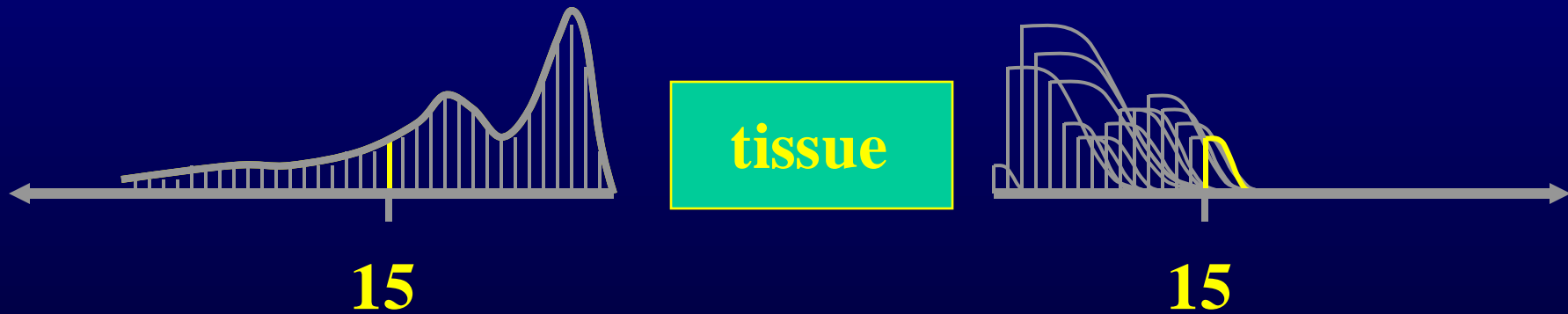


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(15) RF(t - 15) \Delta\tau$$

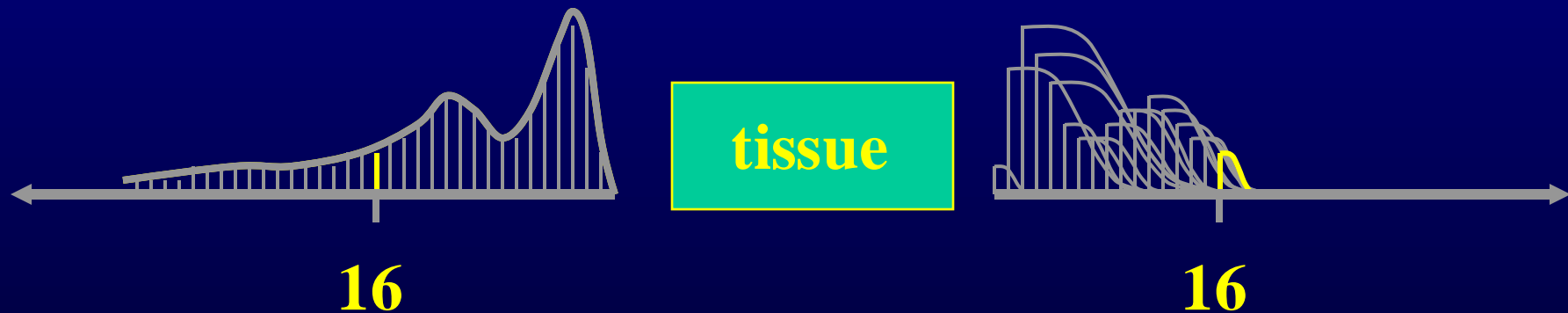


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(16) RF(t - 16) \Delta\tau$$

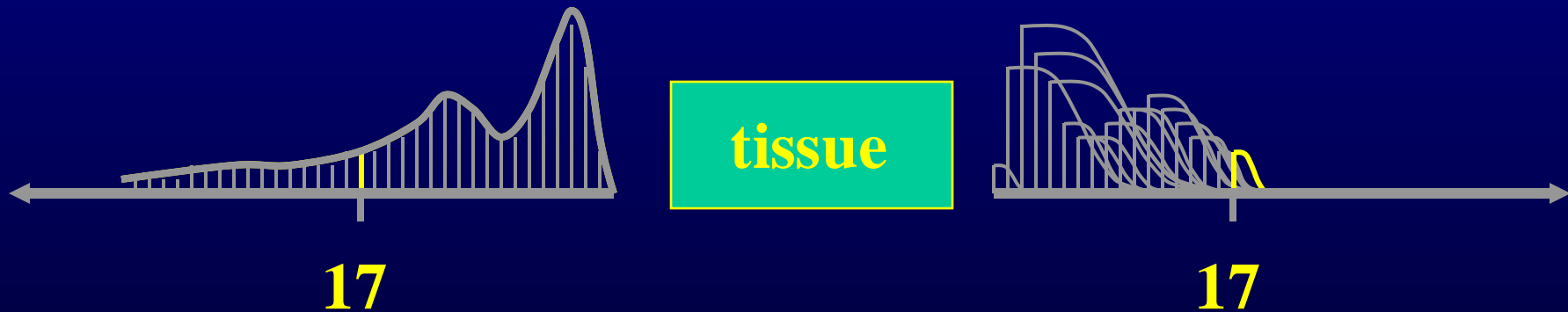


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(17) RF(t - 17) \Delta\tau$$

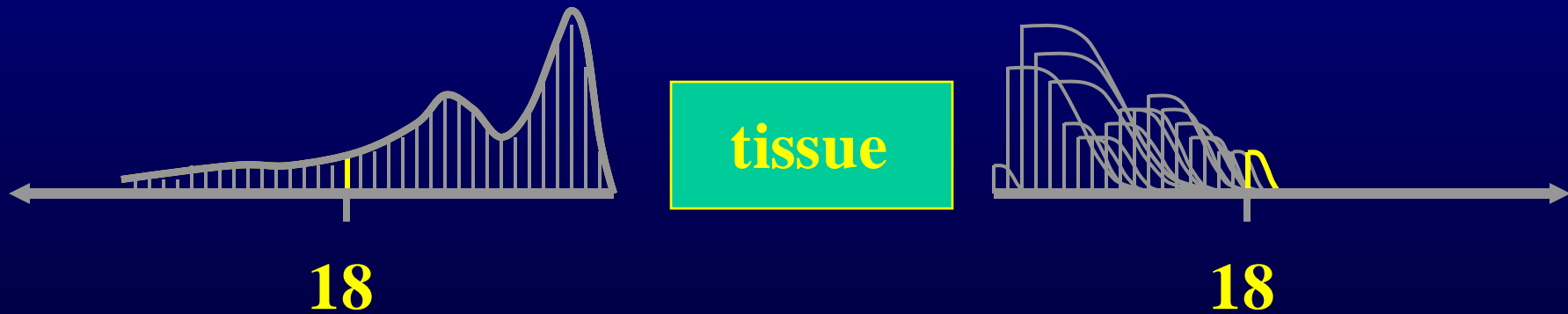


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(18) RF(t - 18) \Delta\tau$$

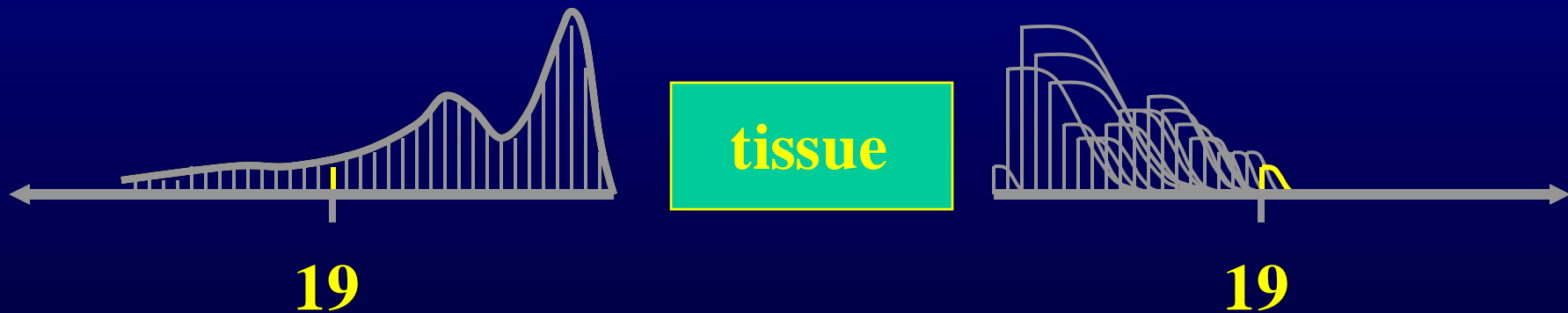


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(19) RF(t - 19) \Delta\tau$$

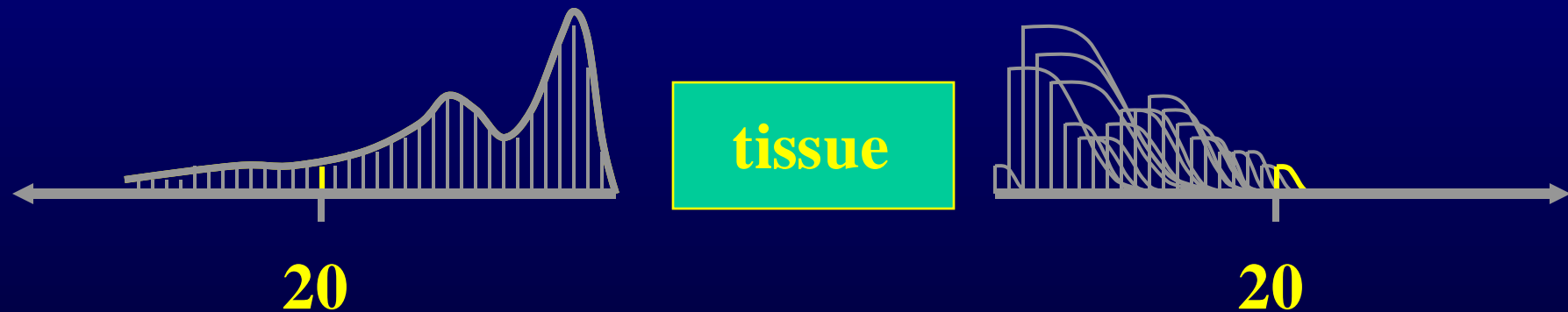


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(20) RF(t - 20) \Delta\tau$$

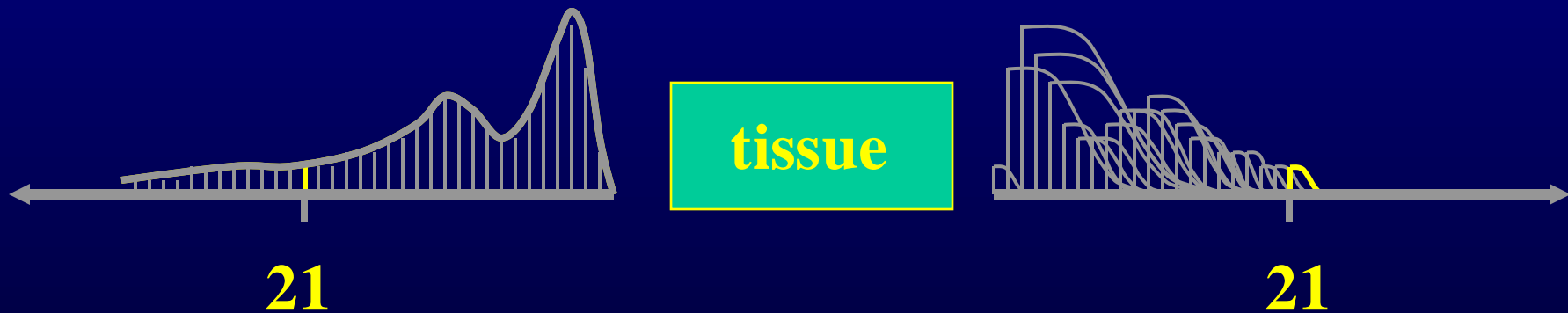


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(t) RF(t - \tau) \Delta\tau$$

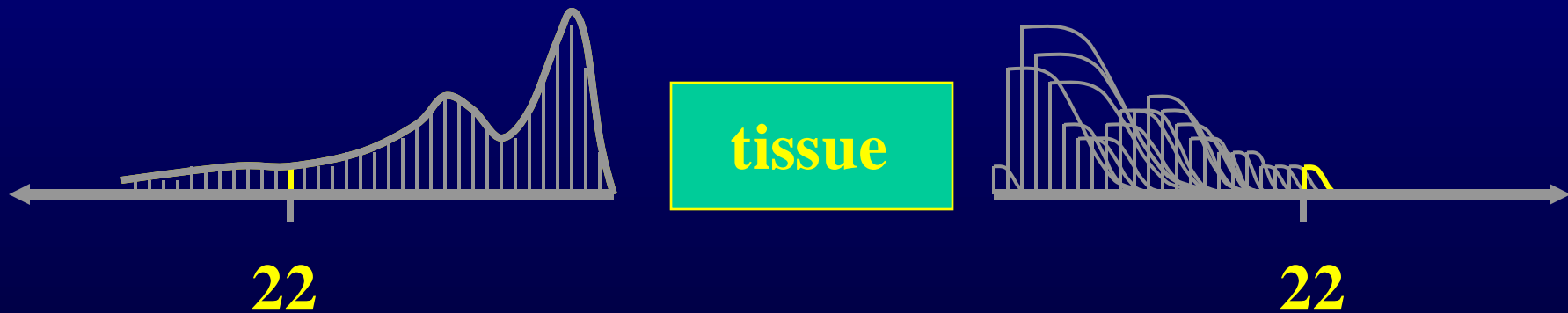


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(t) RF(t - t) \Delta\tau$$



Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(23) RF(t - 23) \Delta\tau$$

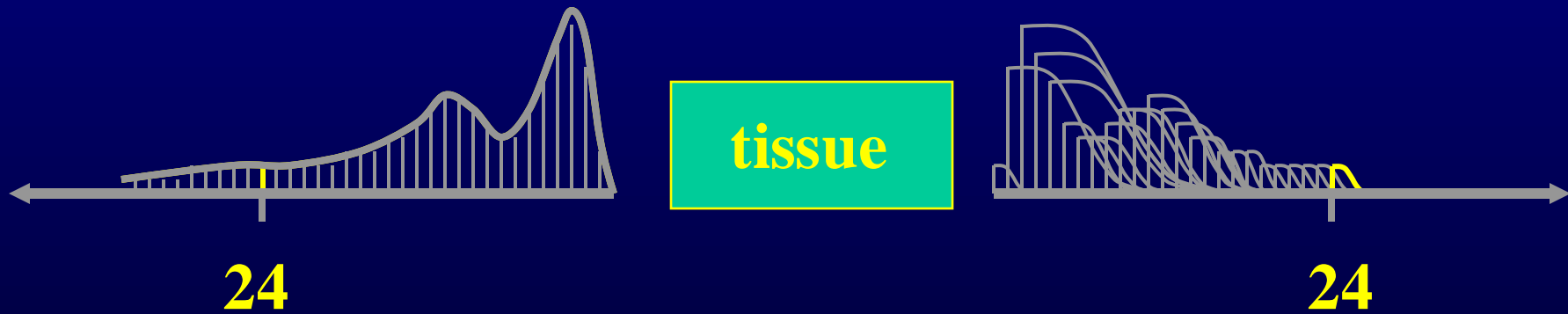


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(24) RF(t - 24) \Delta\tau$$

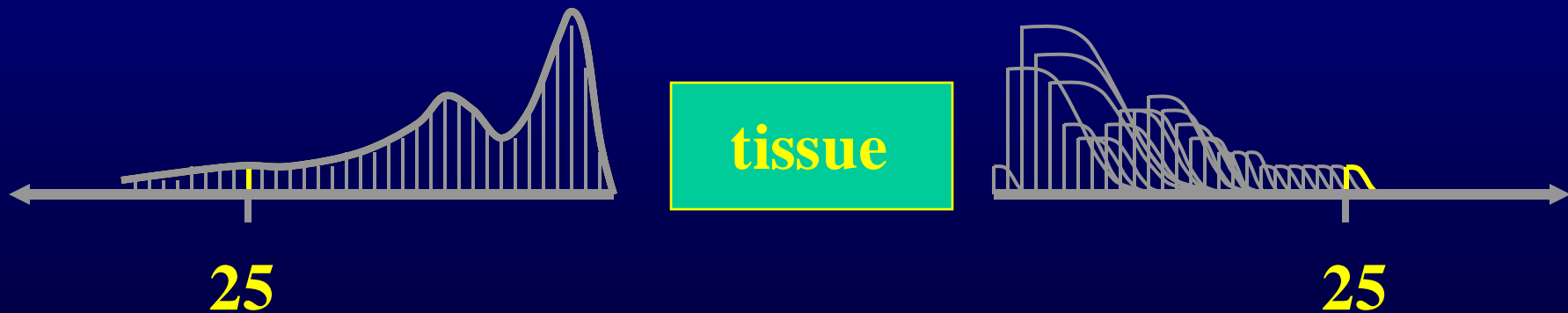


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(25) RF(t - 25) \Delta\tau$$

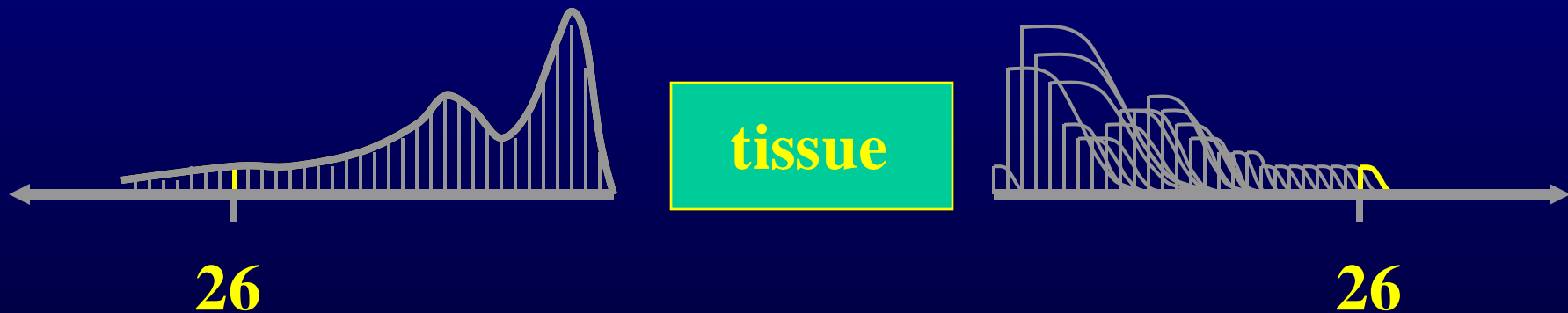


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(26) RF(t - 26) \Delta\tau$$

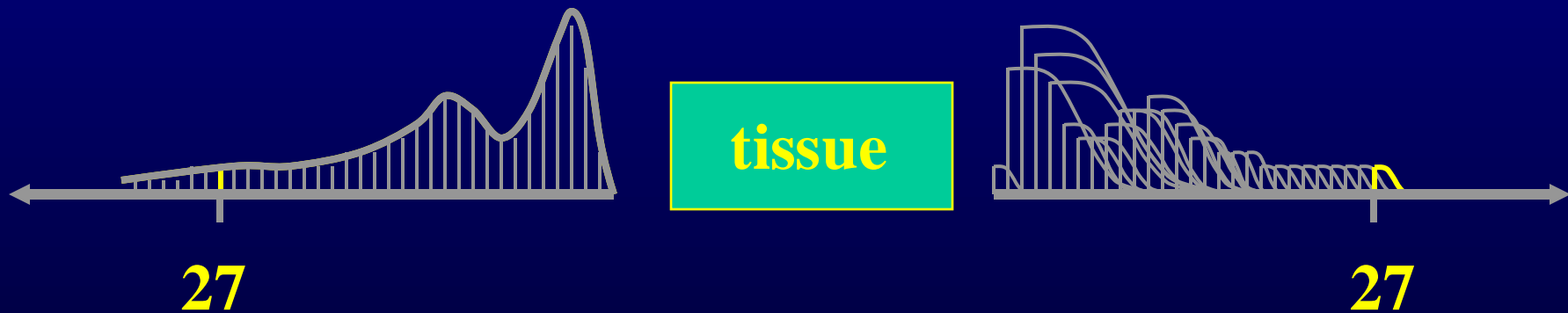


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(27) RF(t - 27) \Delta\tau$$

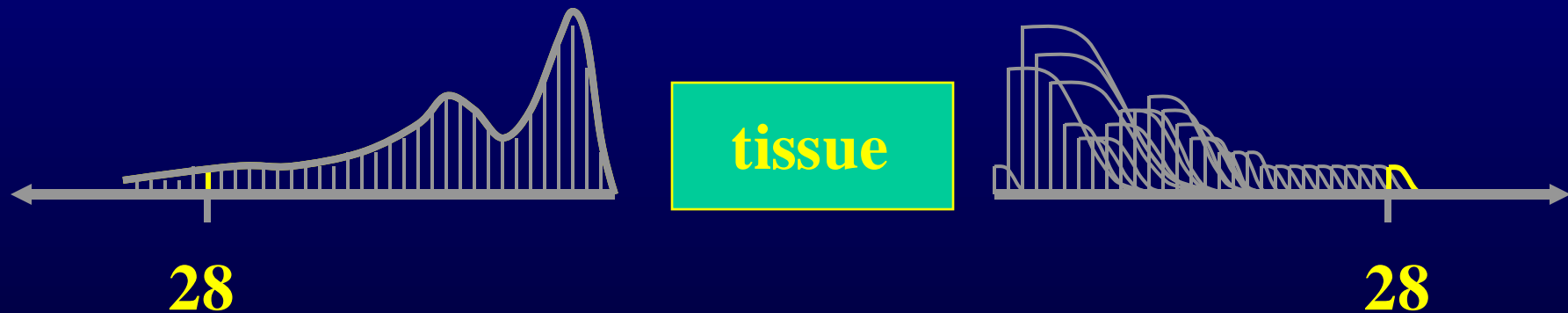


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(28) RF(t - 28) \Delta\tau$$

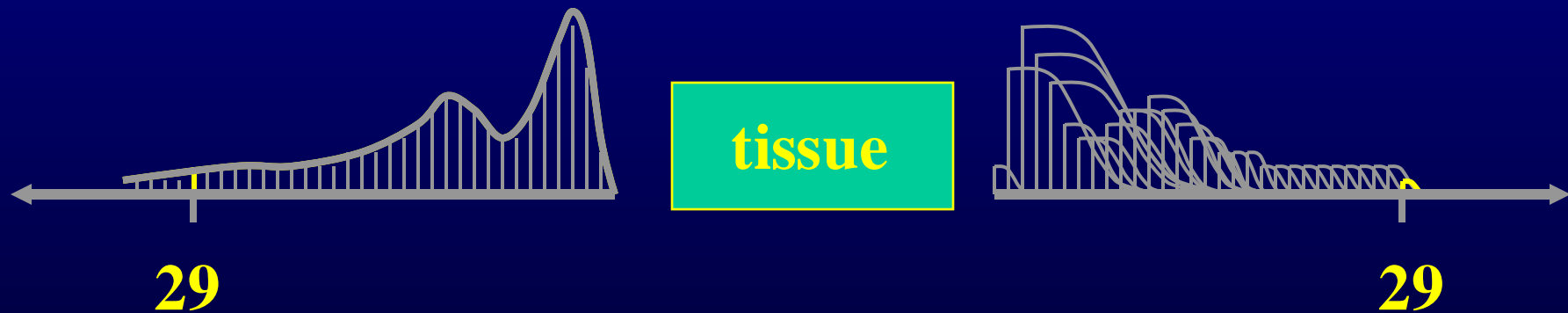


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(29) RF(t - 29) \Delta\tau$$

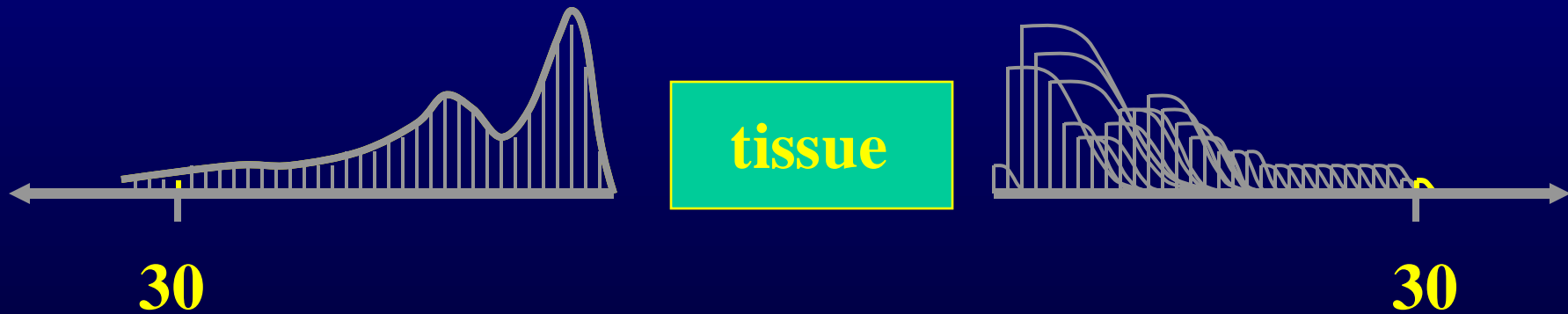


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(30) RF(t - 30) \Delta\tau$$

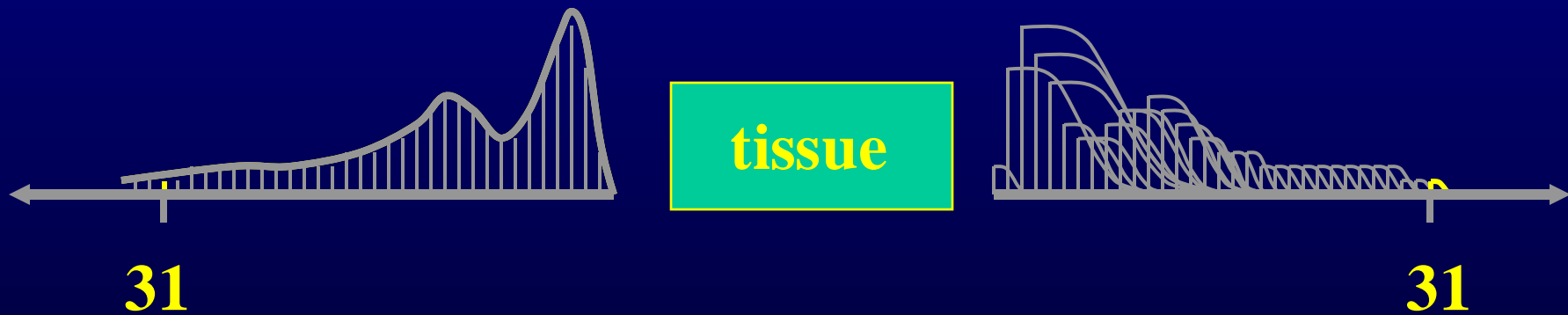


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(31) RF(t - 31) \Delta\tau$$

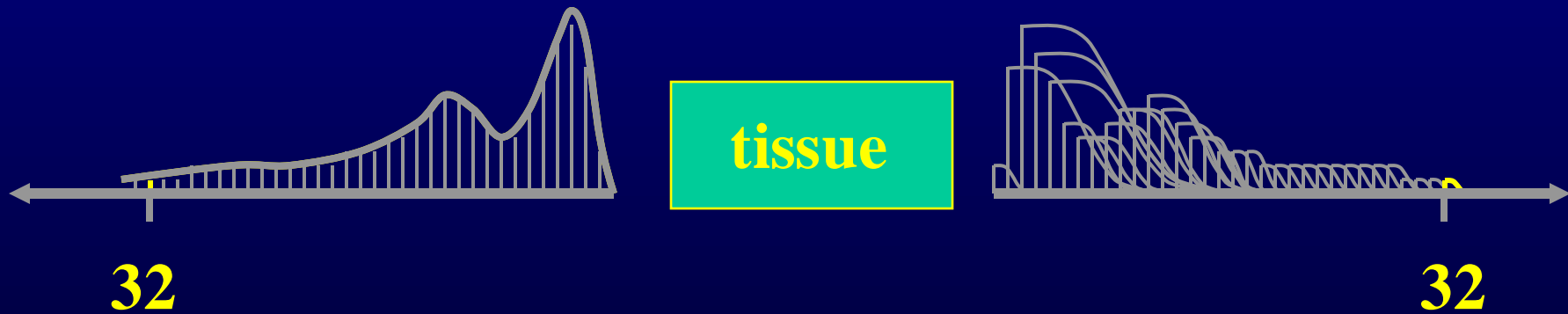


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(32) RF(t - 32) \Delta\tau$$

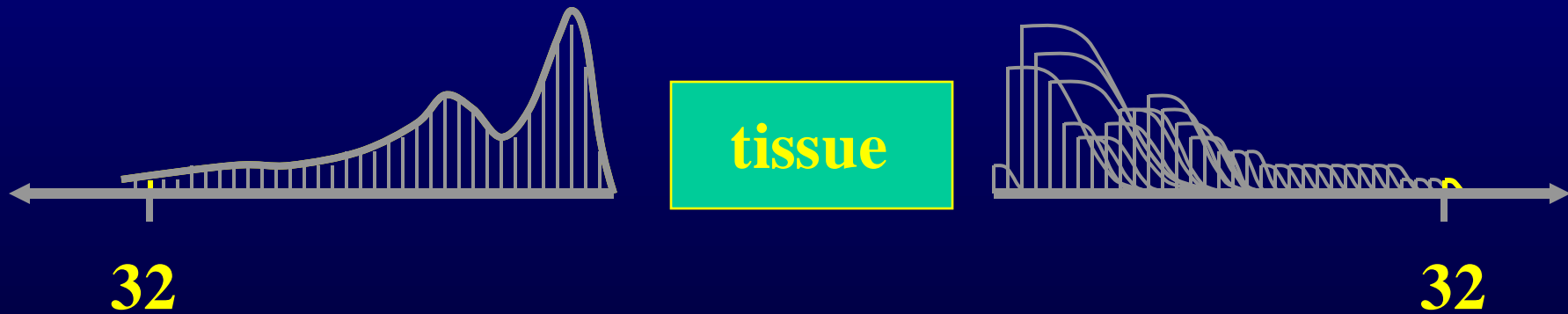


Input : $C_a(t)$

**composed of many
small input**

Tissue enhancement :

$$C_{tis}(t) = f C_a(32) RF(t - 32) \Delta\tau$$

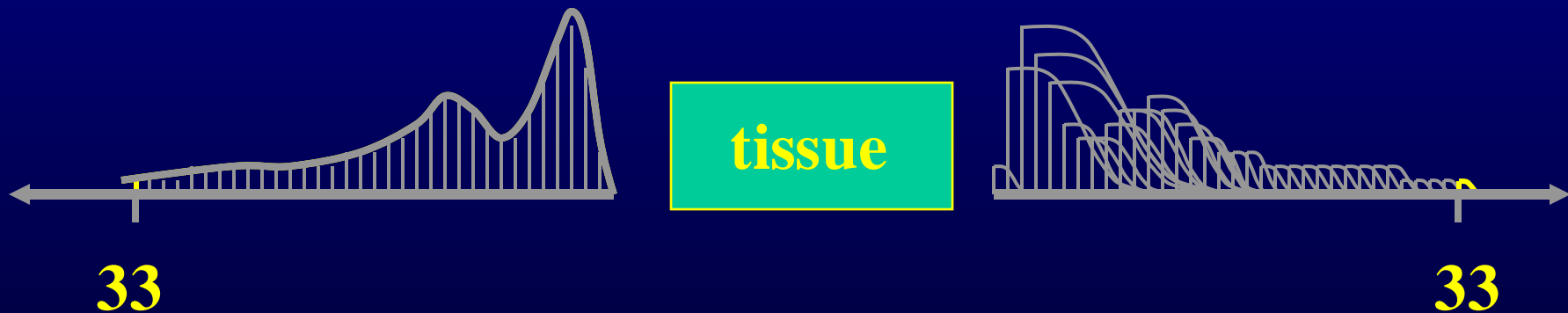


Input : $C_a(t)$

**composed of many
small input**

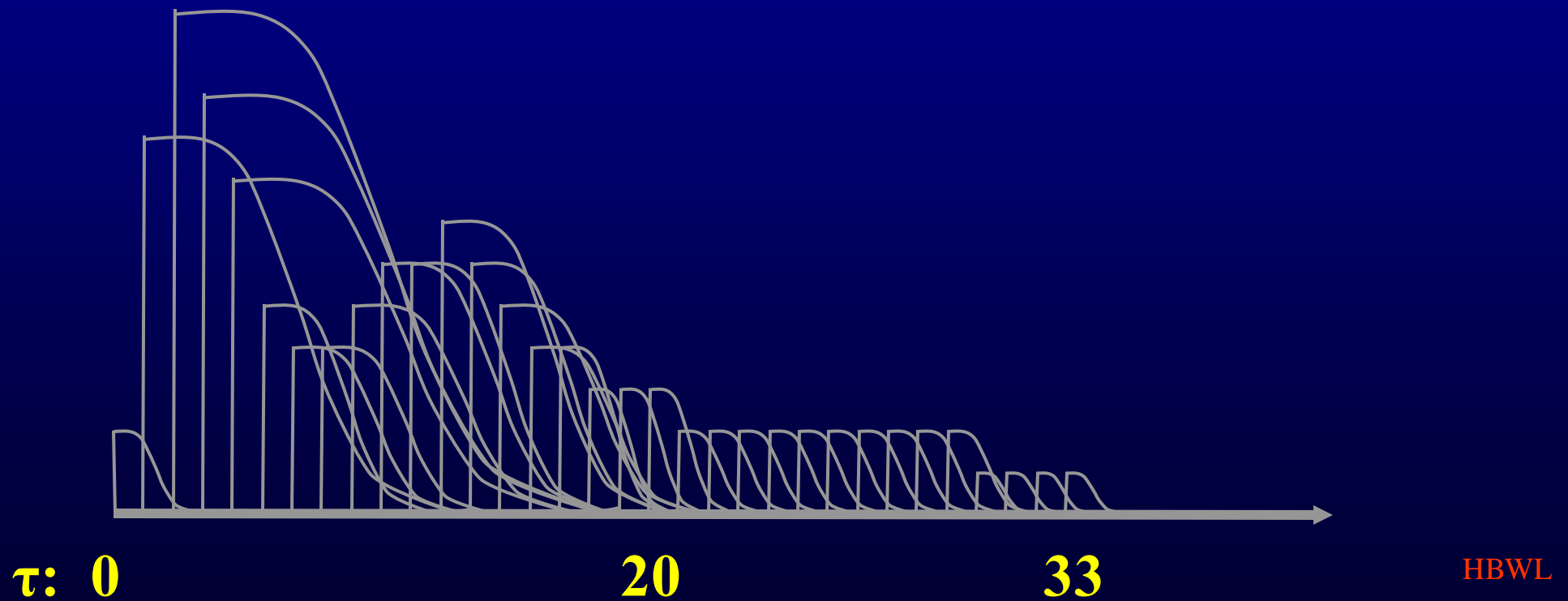
Tissue enhancement :

$$C_{tis}(t) = f C_a(33) RF(t - 33) \Delta\tau$$



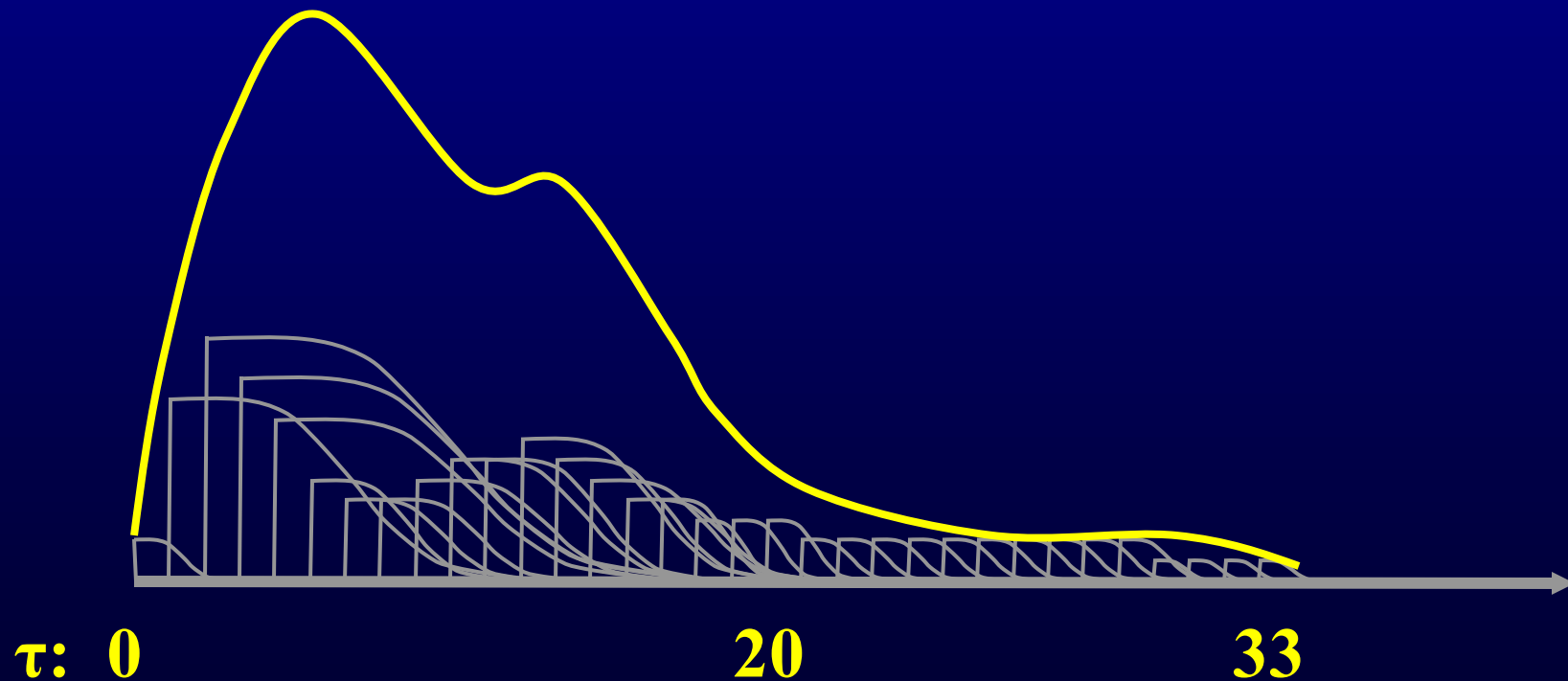
Tissue enhancement :

$$\int C_a(\tau) RF(t - \tau) \Delta\tau ; \tau = 0:33$$



Total tissue enhancement :

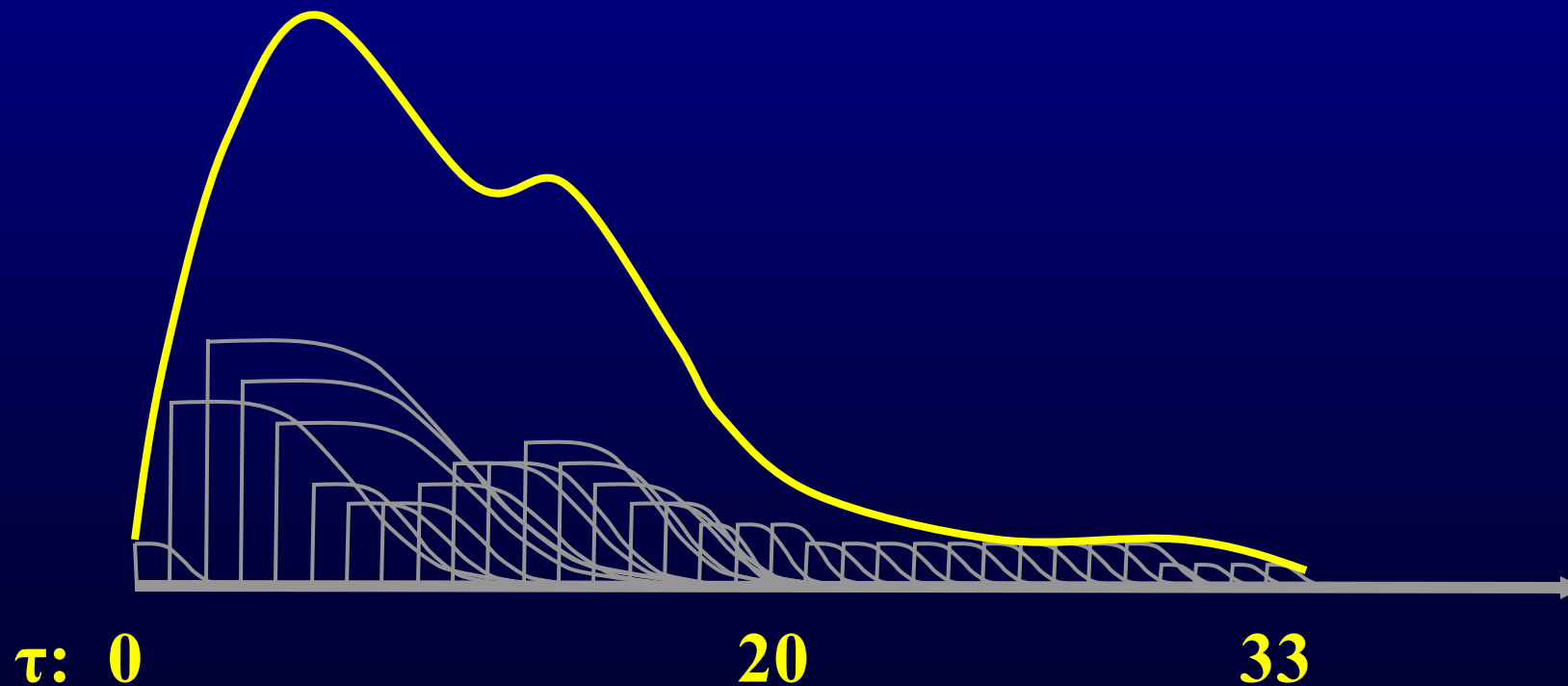
$$C_{tis}(t) = \sum f C_a(\tau) RF(t - \tau) \Delta\tau ; \tau = 0:33$$



Total tissue enhancement :

$$C_{tis}(t) = \int_0^t C_a(\tau) RF(t - \tau) d\tau ; \tau = 0:t$$

The convolution integral

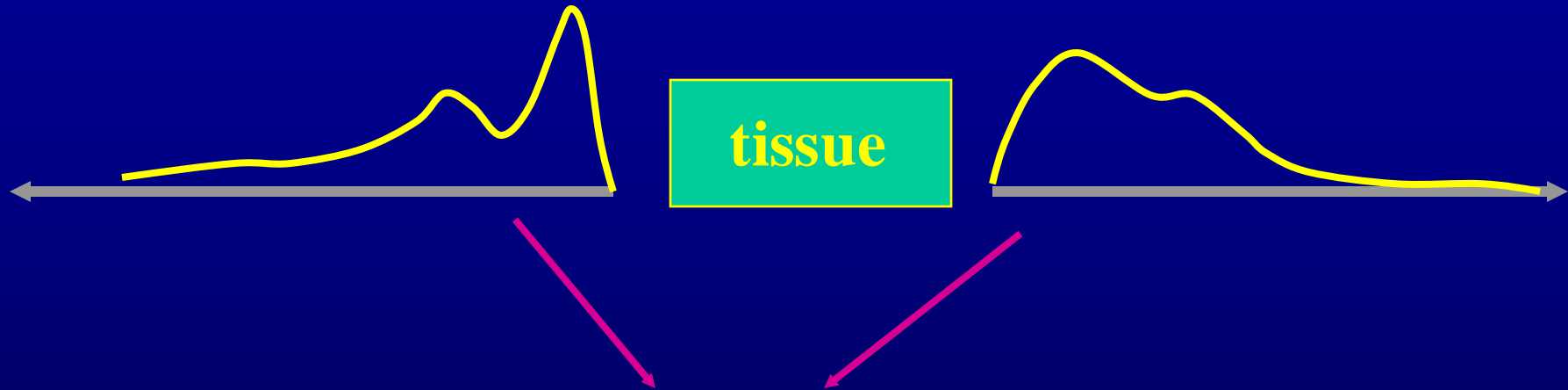


Input :

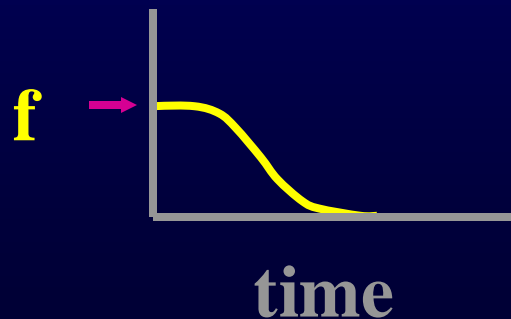
$$C_a(t)$$

Tissue enhancement :

$$C_{tis}(t) = \int f C_a(\tau) RF(t - \tau) d\tau$$



Deconvolution :
find f and $RF(t)$

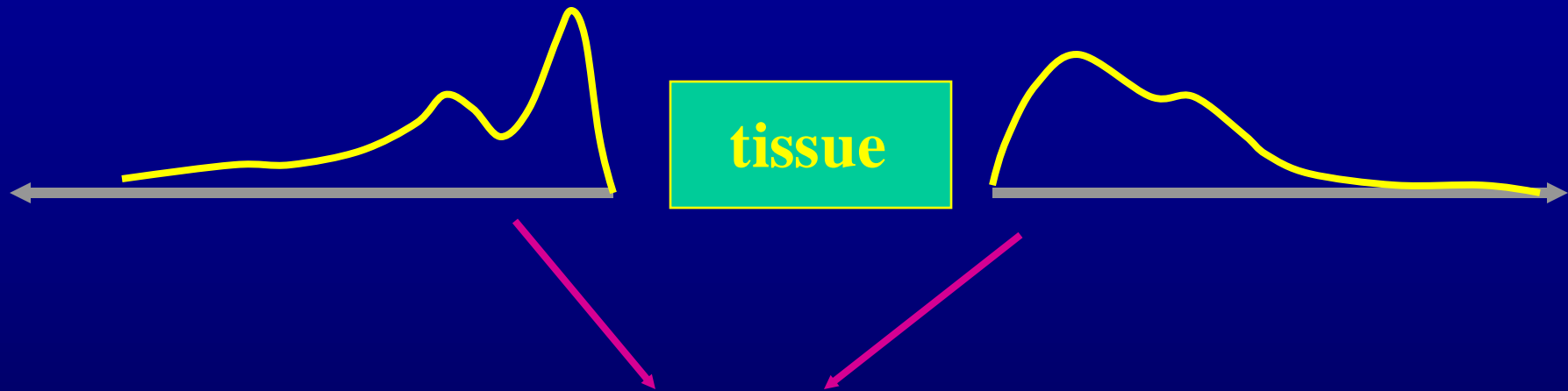


Input :

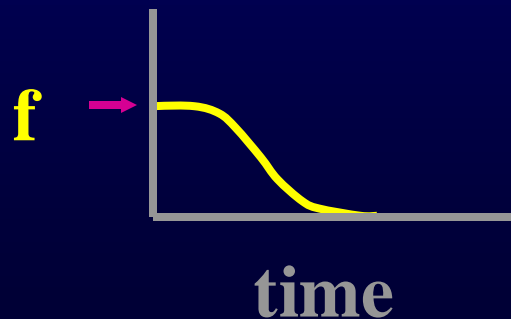
$$C_a(t)$$

Tissue enhancement :

$$C_{tis}(t) = f \int C_a(\tau) RF(t - \tau) d\tau$$



Deconvolution :
find f and $RF(t)$



Conclusion

Measure the
tissue conc

Measure the input conc i.e.
input function

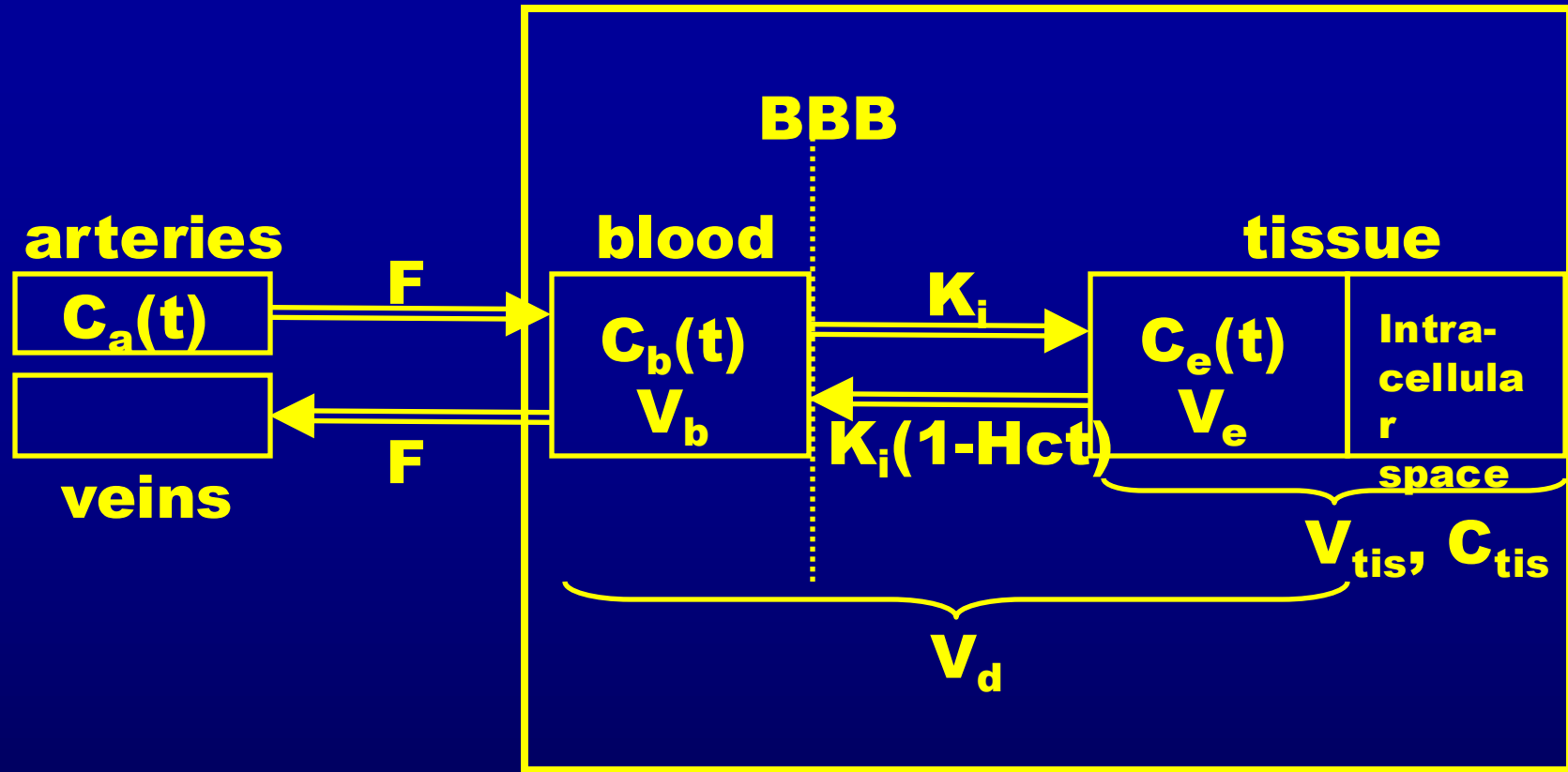
Bolus input : $C_{tis}(t) = f C_a(0) \Delta t RF(t)$

Estimate f and $RF(t)$

Vein injection : $C_{tis}(t) = f \int C_a(\tau) RF(t - \tau) d\tau$

Deconvolution ~ Modelbased

- Use a model e.g.: Monoexponentiel, biexponentiel,
- Optimise the free parameters by least square fit to tissue enhancement curve
- It is robust
- Relative insensitive to noise
- Incorrect if the model is inappropriately chosen



$$V_b \frac{dC_b(t)}{dt} = F C_a(t) - (F + K_i) C_b(t) + K_i(1 - Hct) C_e(t)$$

$$V_e \frac{dC_e(t)}{dt} = K_i C_b(t) - K_i(1 - Hct) C_e(t)$$

$$V_e C_e = V_{\text{tis}} C_{\text{tis}}$$

$$\alpha = \frac{F + K_i}{V_b}$$

$$\beta = \frac{V_{\text{tis}}(1 - \text{Hct})K_i}{V_b V_e}$$

$$\gamma = \frac{K_i}{V_{\text{tis}}}$$

$$\theta = \frac{K_i(1 - \text{Hct})}{V_e}$$

$$(a, b) = \left(\frac{1}{2}[\theta + \alpha + \sqrt{\theta^2 + \alpha^2 - 2\theta\alpha + 4\gamma\beta}], \frac{1}{2}[\theta + \alpha - \sqrt{\theta^2 + \alpha^2 - 2\theta\alpha + 4\gamma\beta}] \right)$$

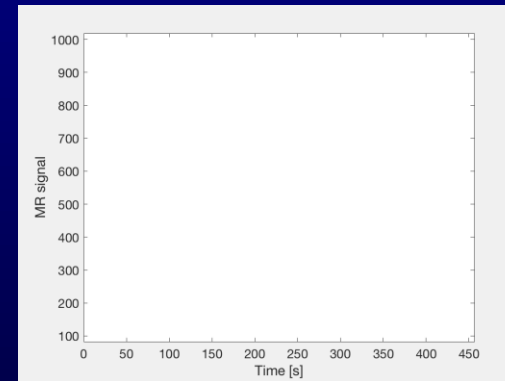
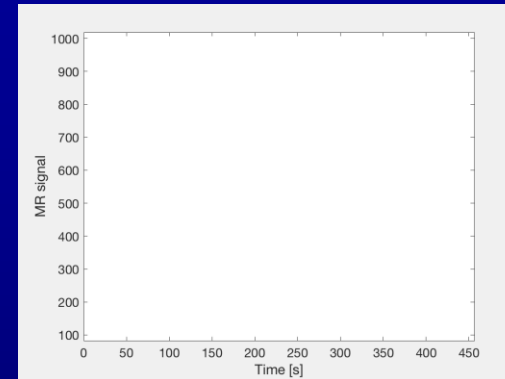
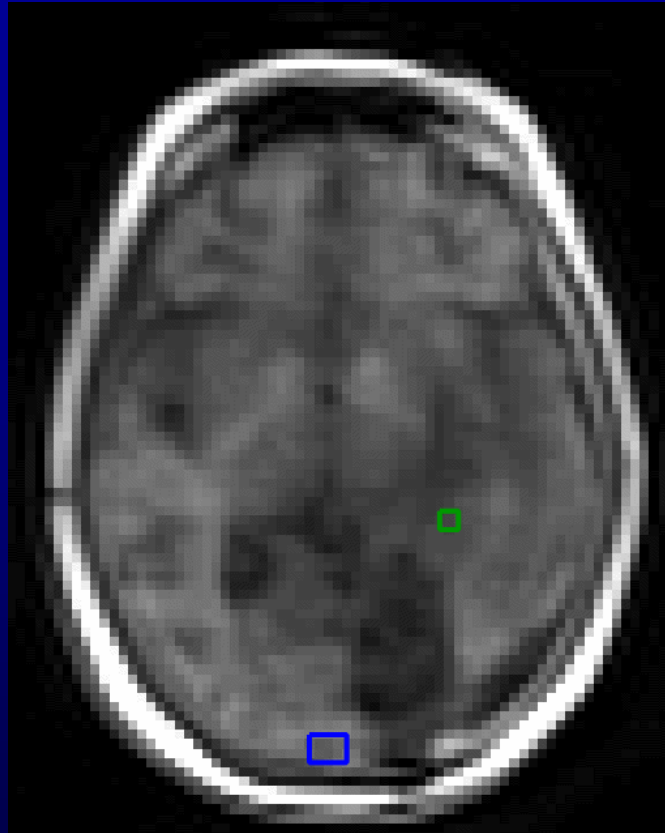
$$C_b(t) = C_a(t) \otimes \frac{F}{V_b} \frac{(a - \theta)e^{-at} - (b - \theta)e^{-bt}}{a - b}$$

$$C_{\text{tis}}(t) = C_a(t) \otimes \frac{F}{V_b} \frac{K_i}{V_{\text{tis}}} \frac{e^{-bt} - e^{-at}}{a - b}$$

$$C_t(t) = V_b C_b(t) + (1 - V_b) C_{\text{tis}}(t) \Leftrightarrow$$

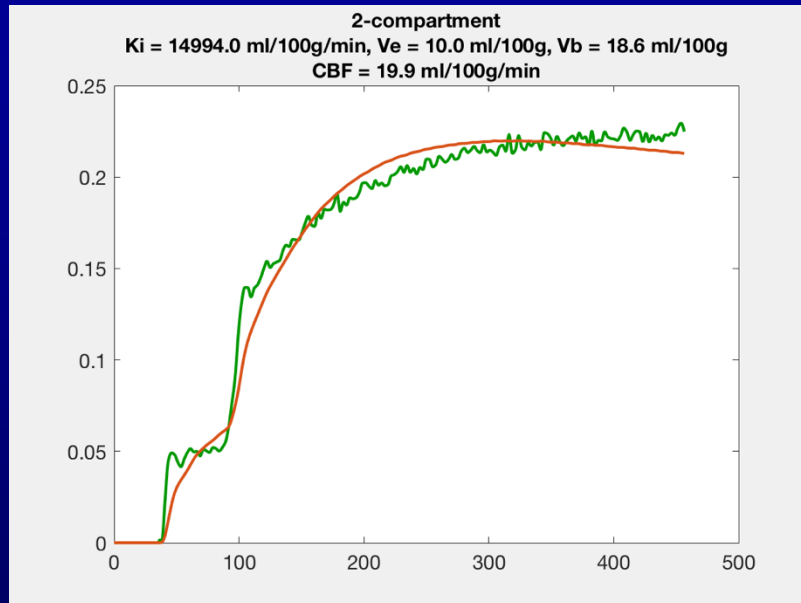
$$C_t(t) = F C_a(t) \otimes \left[\frac{(a - \theta - K_i / V_b) e^{-at} + (-b + \theta + K_i / V_b) e^{-bt}}{a - b} \right]$$

Dynamic Contrast Enhanced

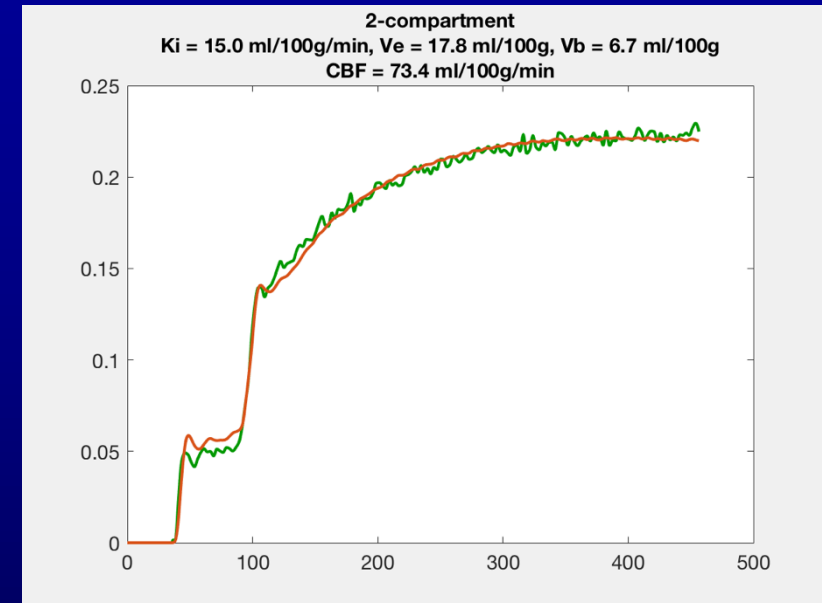


Voxel size: 2.4 x 2.4 x 5 mm
Power injector: 3 mL/s or 1 mL/s
Time resolution: 2.55 sec

No constrain, naïve start guess



Model constrain, model start guess



F: Fixed from Tikhonov (model-free deconvolution)

Ki:

- **LB = 10% of initial guess,**
- **UB**
 - **2000% of initial guess (if from Patlak)**
 - **120% of initial guess (if from CTH)**

Volume: $V_e + V_b = V_d \pm 30\%$ (from Tikhonov)

Deconvolution ~ Modelfree

- No model a priory
- Very flexible: many of free parameters
- A projection
- Very sensitive to noise
- Incorrect if not regularized rigorously
- Fourier transform, SVD, GSVD, Tikhonov, GPD

Convolution written in matrix notation

$$C_t(t) = C_a(t) \otimes f RIF(t) = f \int_0^t C_a(\tau) RIF(t - \tau) d\tau$$

$$\begin{array}{c}
 C_a(t-1) \quad C_a(t-2) \quad \dots \quad C_a(t-(N-1)) \\
 \left[\begin{array}{c} C_t(1) \\ C_t(2) \\ C_t(3) \\ \vdots \\ C_t(N) \end{array} \right] = f\Delta t \left[\begin{array}{cccccc} C_a(1) & 0 & 0 & \dots & 0 \\ C_a(2) & C_a(1) & 0 & \dots & 0 \\ C_a(3) & C_a(2) & C_a(1) & \dots & 0 \\ \vdots & \vdots & \vdots & \ddots & \vdots \\ C_a(N) & C_a(N-1) & C_a(N-2) & \dots & C_a(1) \end{array} \right] \left[\begin{array}{c} RIF(1) \\ RIF(2) \\ RIF(3) \\ \vdots \\ RIF(N) \end{array} \right]
 \end{array}$$

Convolution written in matrix notation

Solution **Successive solving the matrix eq.**

Minimization

$$\min\{\|C_a RIF - C_t\|_2^2\}$$

**Minimization and
regularization**

$$\min\{\|C_a RIF - C_t\|_2^2 + \lambda^2 \|RIF\|_2^2\}$$

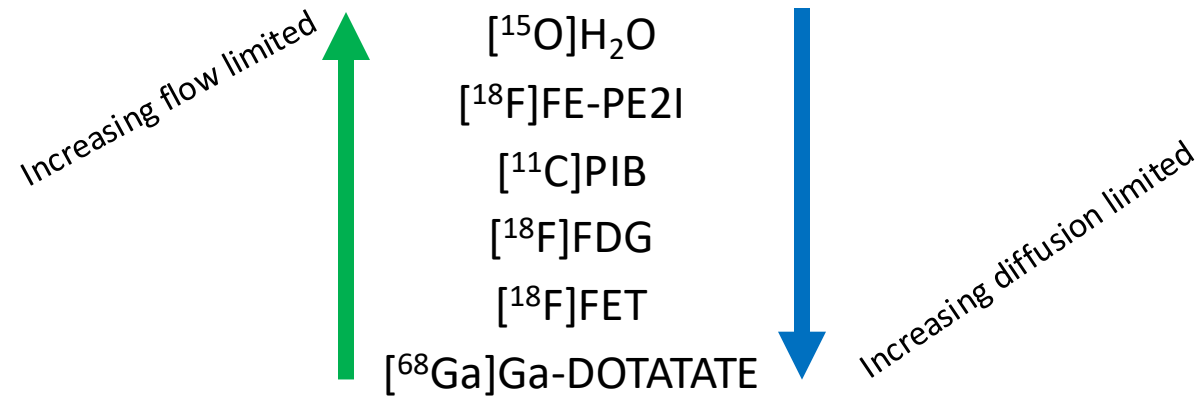
$$\min\{\|C_a RIF - C_t\|_2^2 + \lambda^2 \|L RIF\|_2^2\}$$

$$L = \begin{pmatrix} -1 & 1 & \dots & 0 & 0 \\ \vdots & \vdots & \ddots & \vdots & \vdots \\ 0 & 0 & \dots & -1 & 1 \end{pmatrix}$$

Question

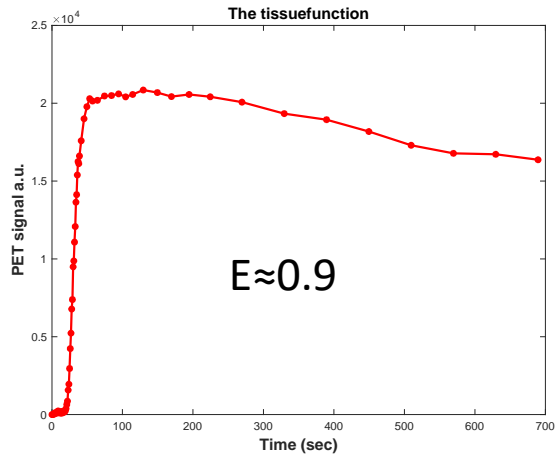
Can we estimate perfusion from any type of
PET tracers

This should work irrespectively
tracer type
flow or diffusion limited

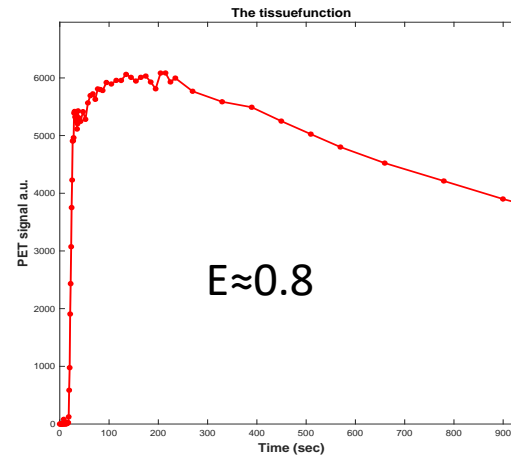


Time resolution : 1 sec

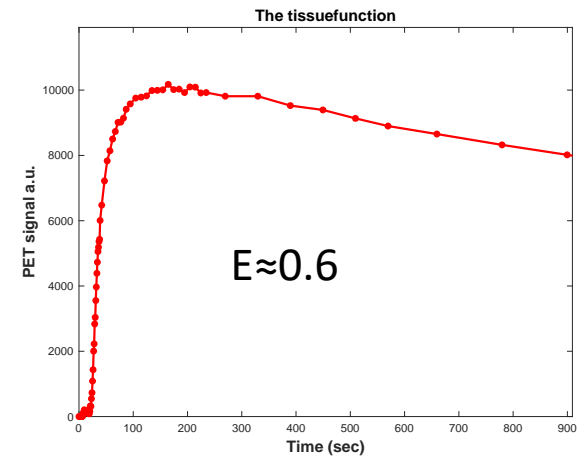
$[^{15}\text{O}]\text{H}_2\text{O}$



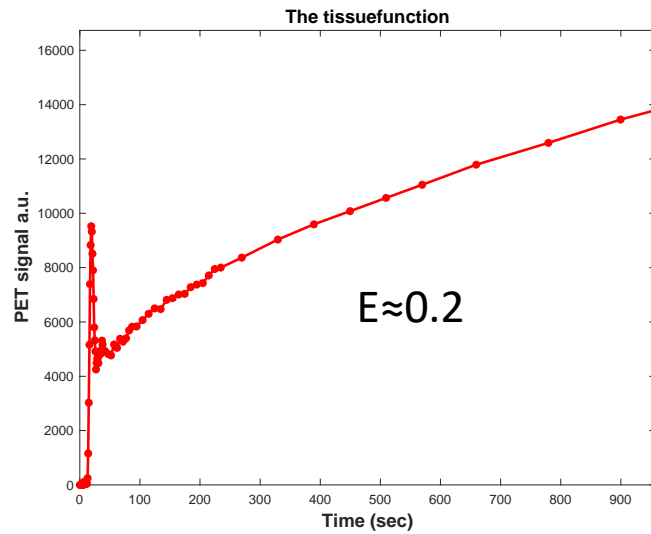
$[^{18}\text{F}]\text{FE-PE2I}$



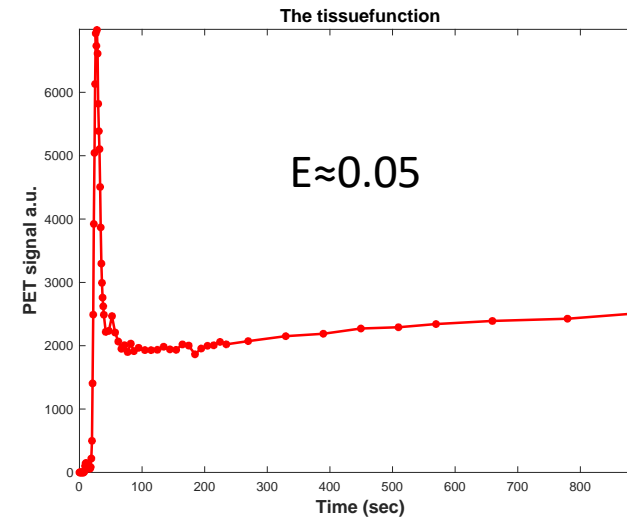
$[^{11}\text{C}]\text{PIB}$



$[^{18}\text{F}]\text{FDG}$

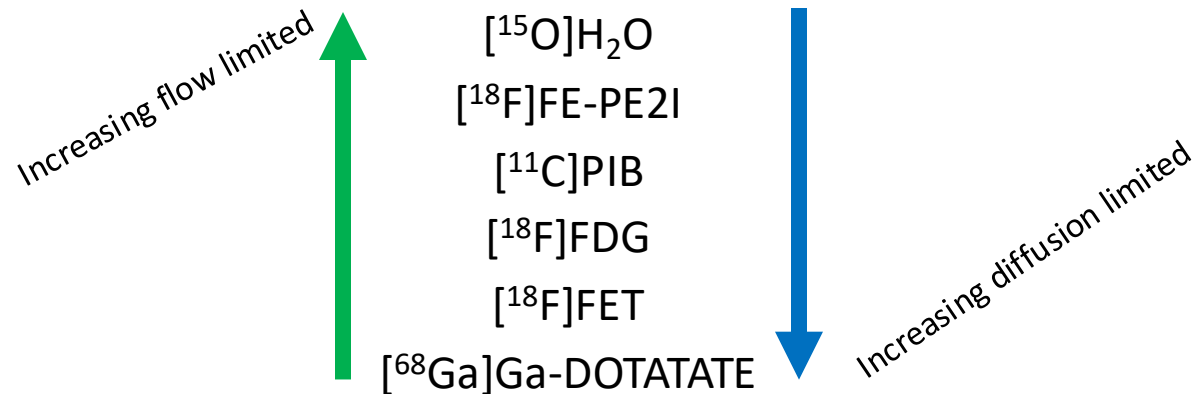


$[^{18}\text{F}]\text{FET}$



What about time resolution ??

This should work irrespectively
tracer type
flow or diffusion limited
if time resolution is high enough

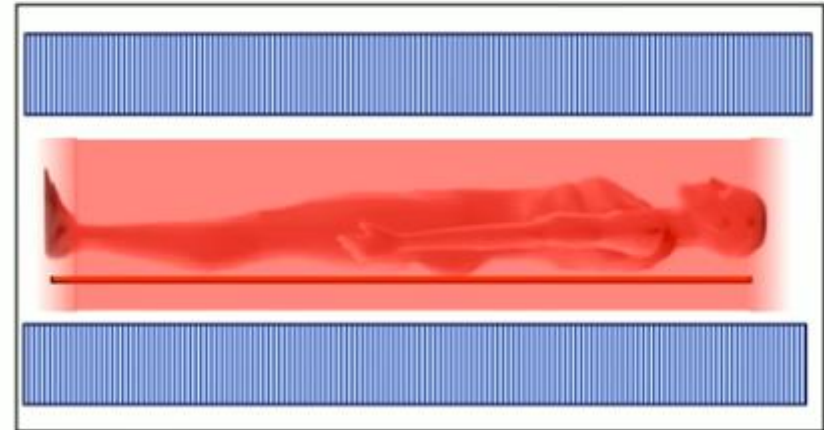


Method

- Siemens Quadra PET/CT scanner
- Reconstruction TrueX PSF 2-3mm Gauss – 1.65x1.65mm²
 - 1.6-3 mm slice thickness
- Framing: 40 x 1 s – 10 x 2 s – 10 x 5 s ... up to 60 min
- Correction for metabolite
- Image derived input function – acquired in descending aorta.
- Standard Injected Activity
- Correction for regional arterial transit time delay between aorta and brain

Long Axial Field of View PET/CT Scanner

- 40x gain in effective sensitivity for total-body imaging!
- 4-5x gain in sensitivity for single organ imaging
- Total-body kinetics
 - All tissues/organs simultaneously
 - Better temporal resolution



CBF

Residue impuls response



SUV - 0-40 seconds $[^{15}\text{O}]\text{H}_2\text{O}$

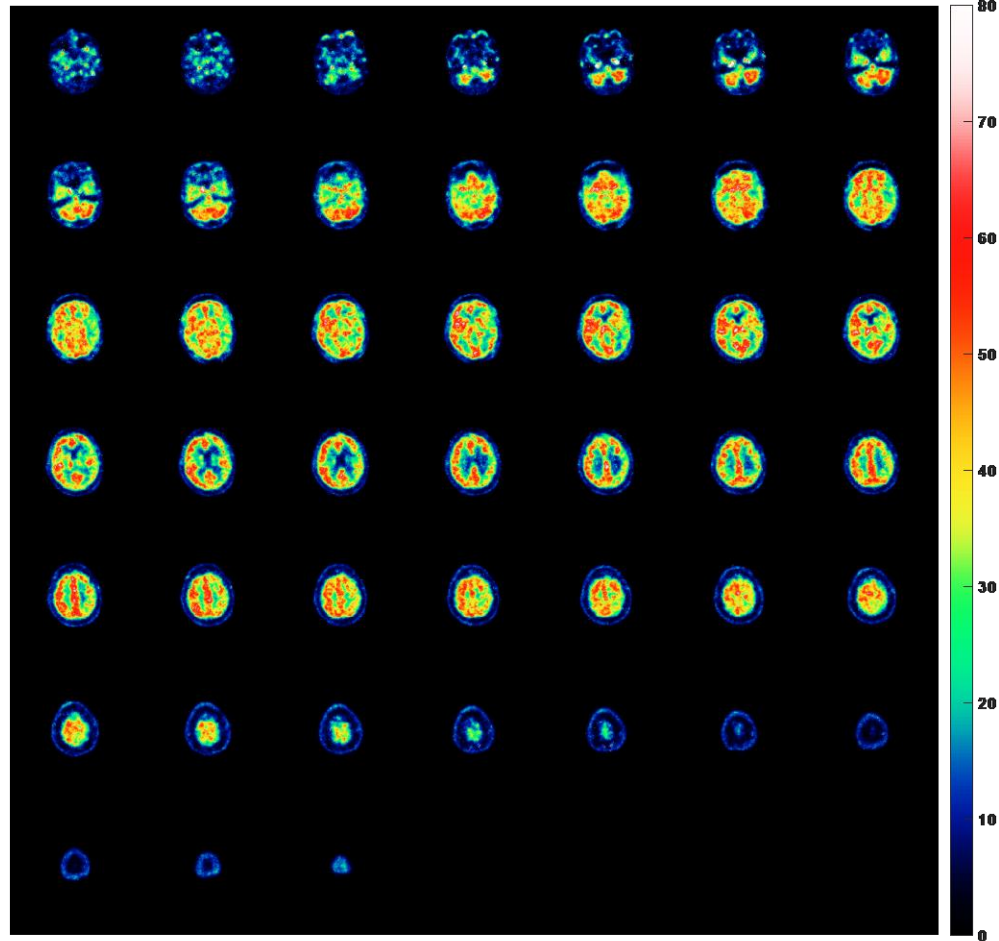
V_d or late images

Extraction fraction

Mean transit time

Arterial transit time delay

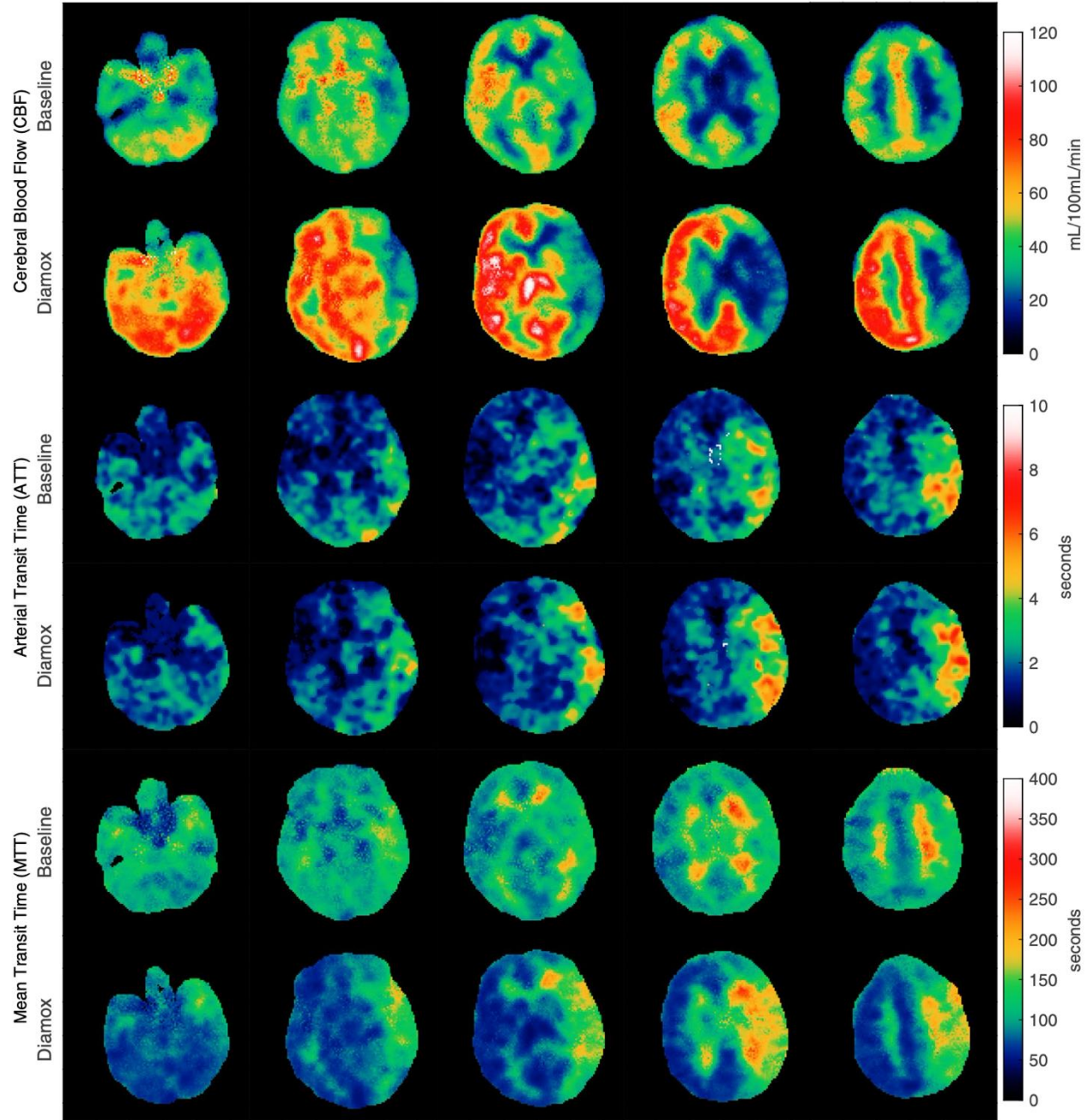
CBF - (ml/100g/min); [¹⁵O]H₂O



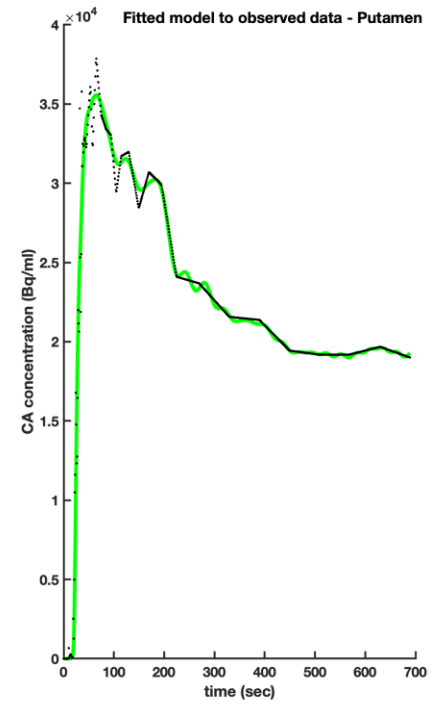
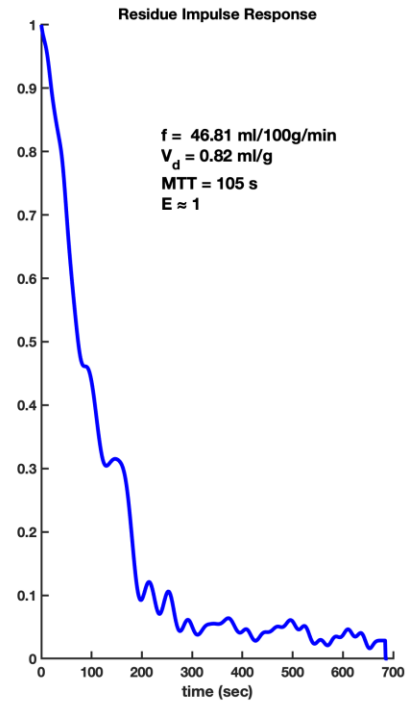
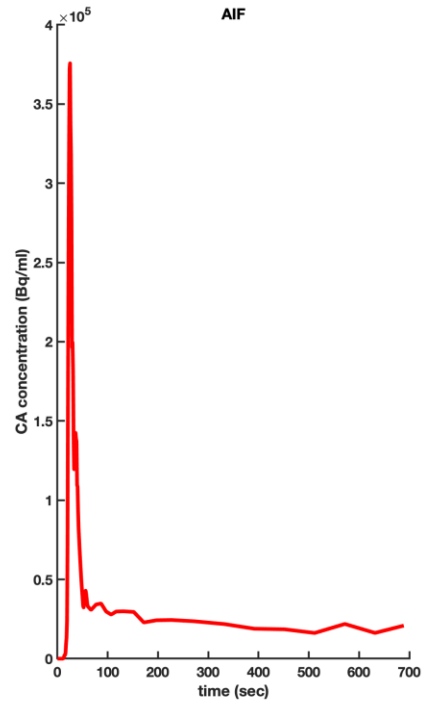
Patient:
Left ICA stenosis

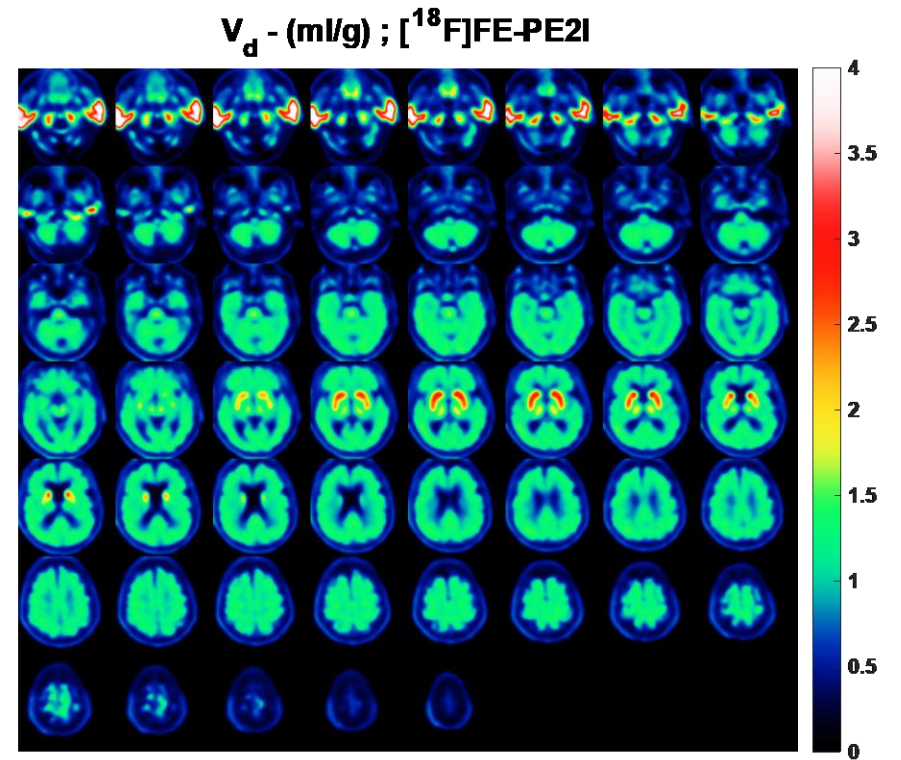
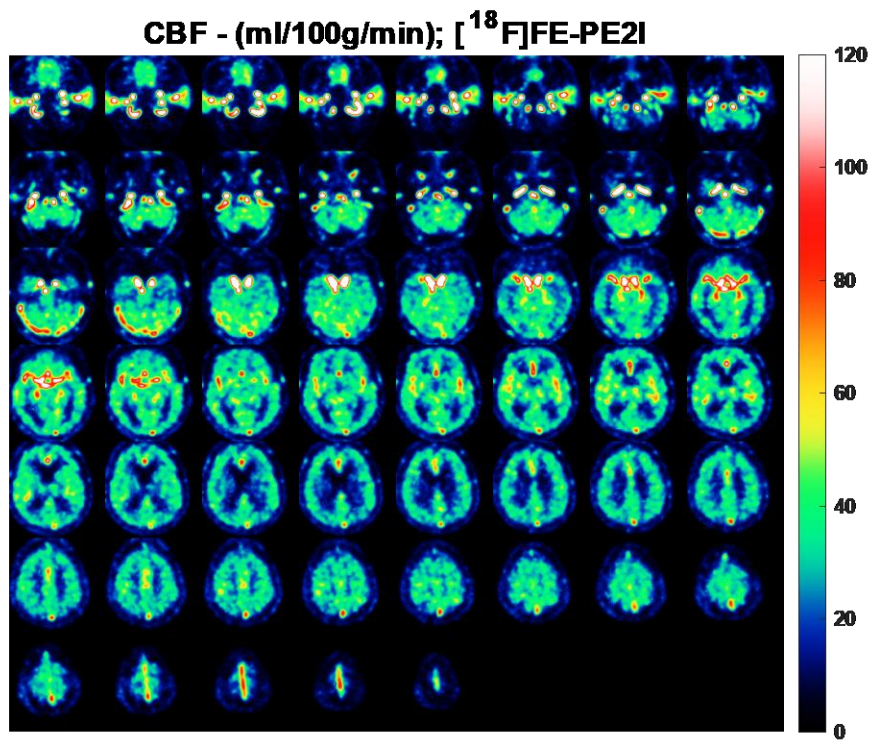
- [¹⁵O]H₂O
- [¹⁸F]FE-PE2I
- [¹¹C]PIB
- [¹⁸F]FDG
- [¹⁸F]FET
- [⁶⁸Ga]Ga-DOTATATE

Increasing diffusion limited



Dynamic [^{15}O]H $_2\text{O}$ Quadra-PET scan

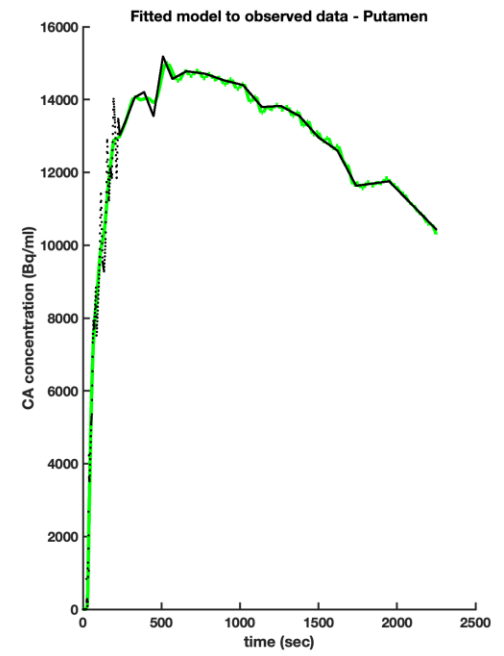
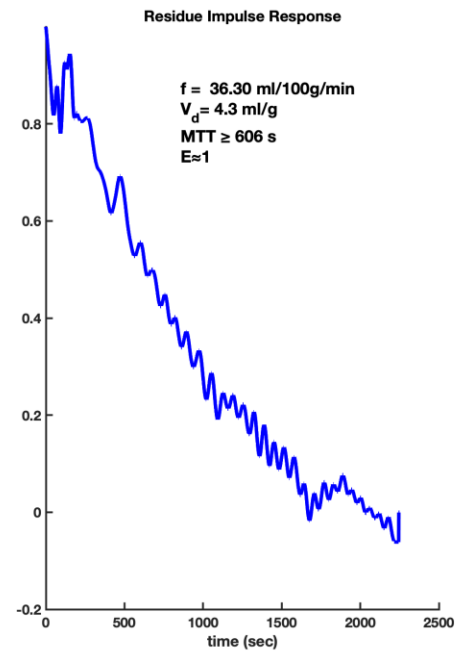
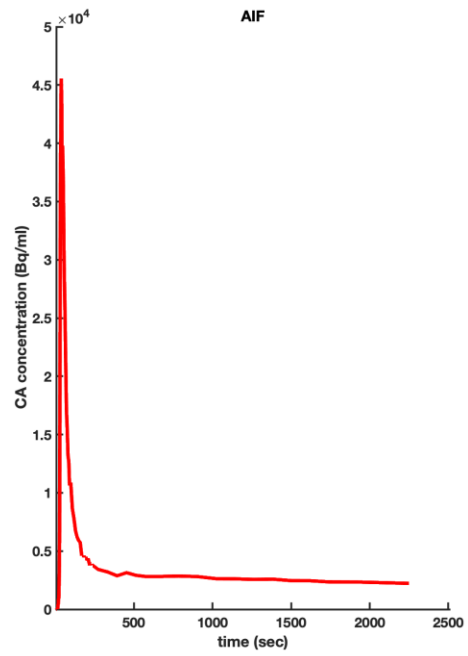


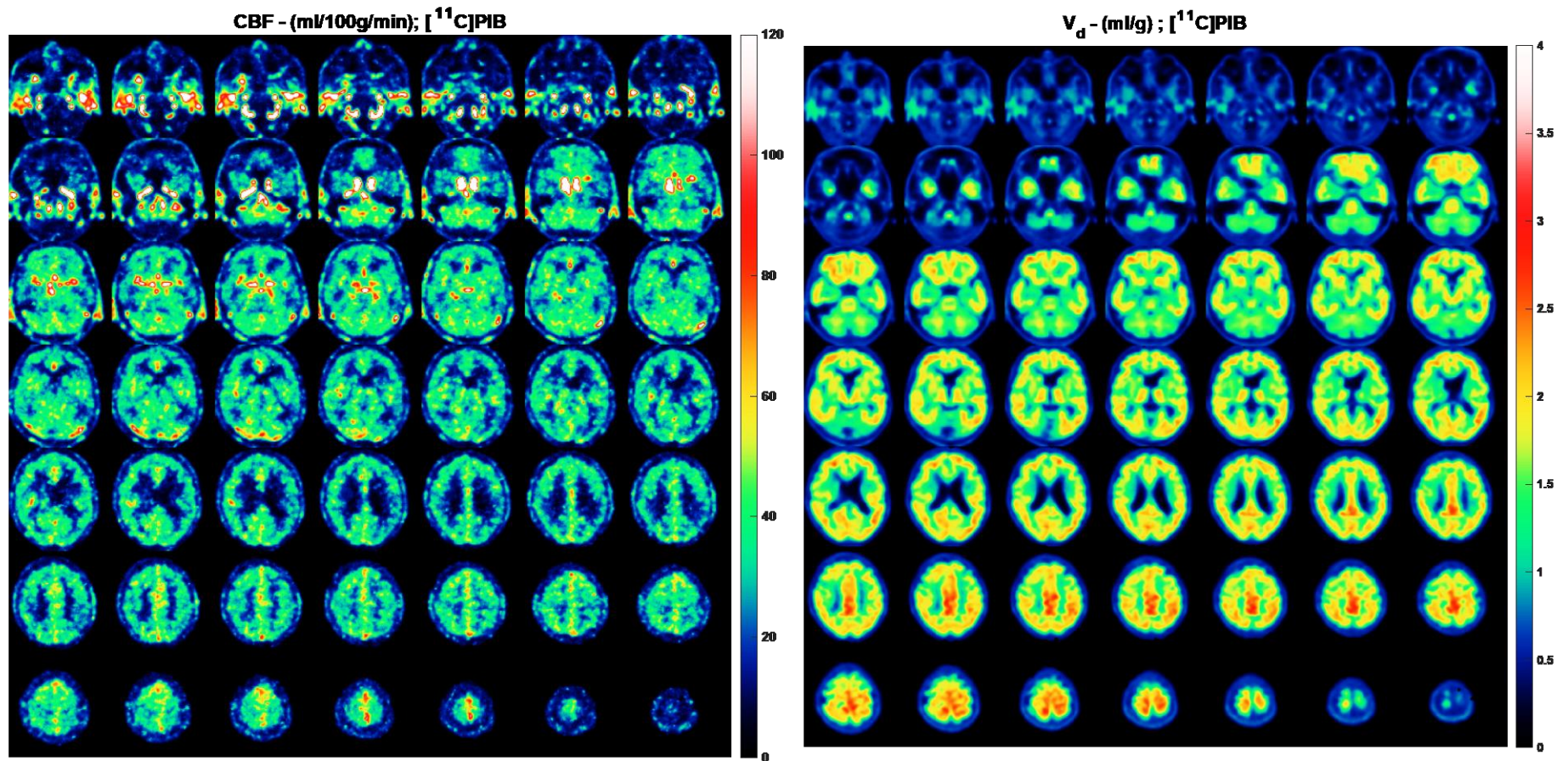


- ↓
- [^{15}O]H₂O
 - [^{18}F]FE-PE2I
 - [^{11}C]PIB
 - [^{18}F]FDG
 - [^{18}F]FET
 - [^{68}Ga]Ga-DOTATATE

Increasing diffusion limited

Dynamic [¹⁸F]FE-PE2I Quadra-PET scan

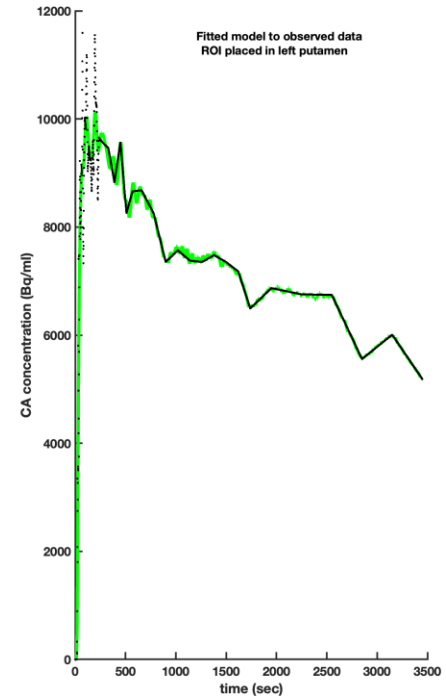
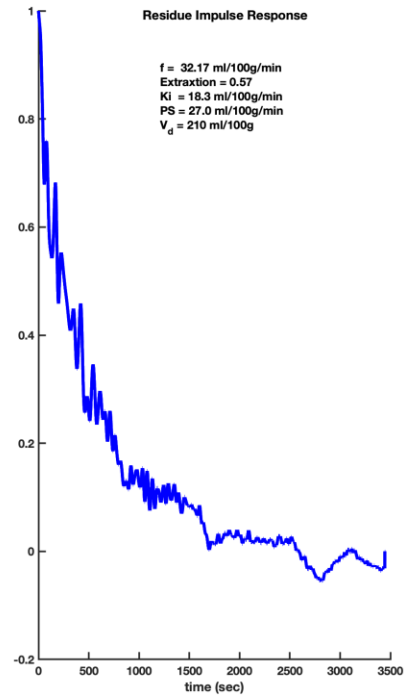
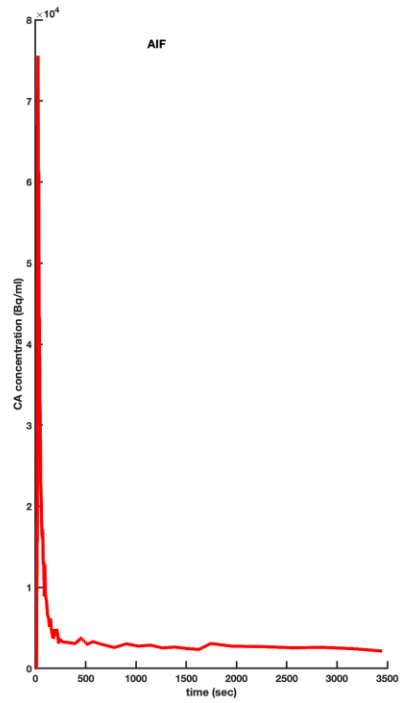


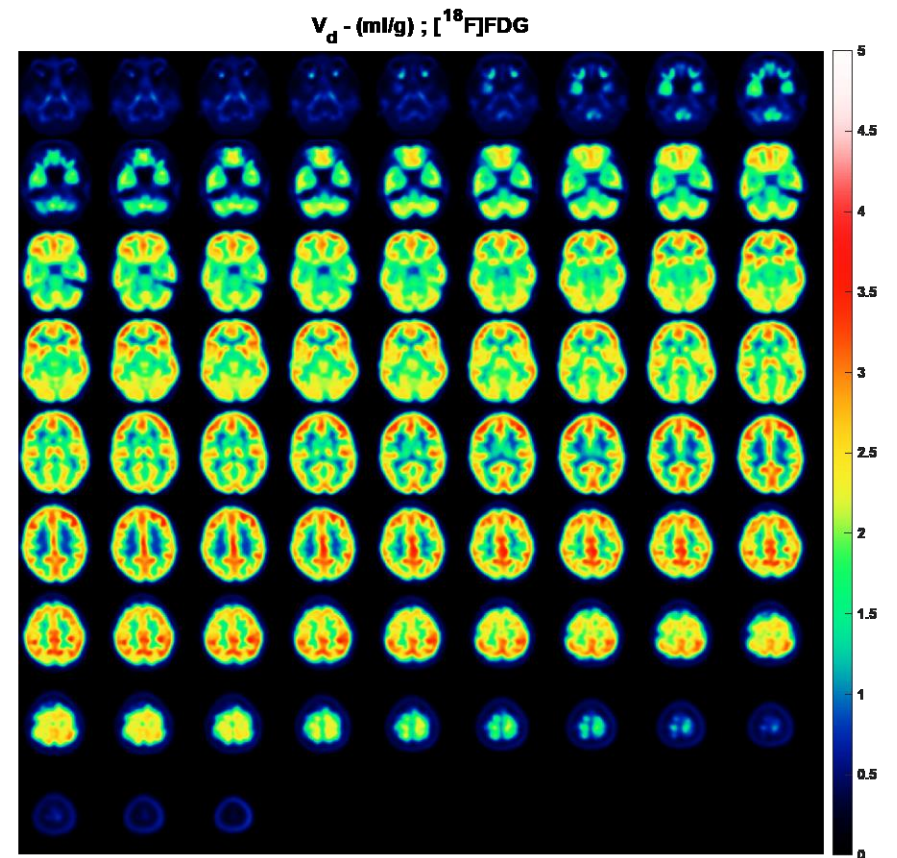
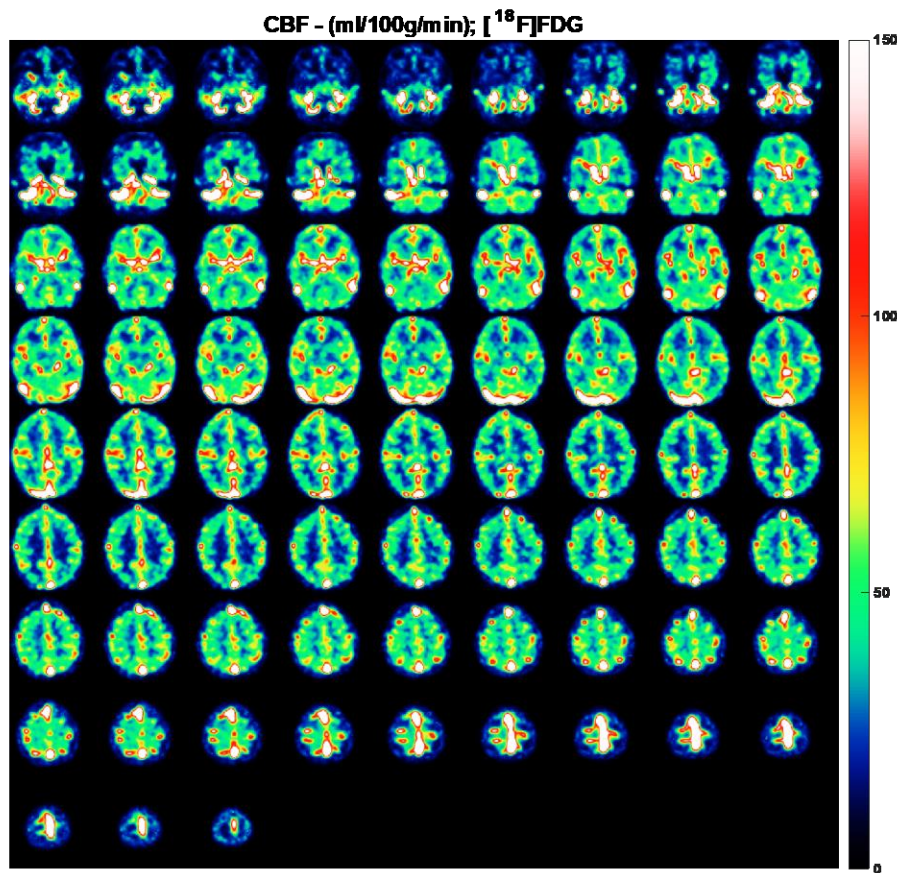


Increasing diffusion limited

- $[^{15}\text{O}]\text{H}_2\text{O}$
- $[^{18}\text{F}]\text{FE-PE2I}$
- $[^{11}\text{C}]\text{PIB}$
- $[^{18}\text{F}]\text{FDG}$
- $[^{18}\text{F}]\text{FET}$
- $[^{68}\text{Ga}]\text{Ga-DOTATATE}$

Dynamic [¹¹C]PIB Quadra-PET scan

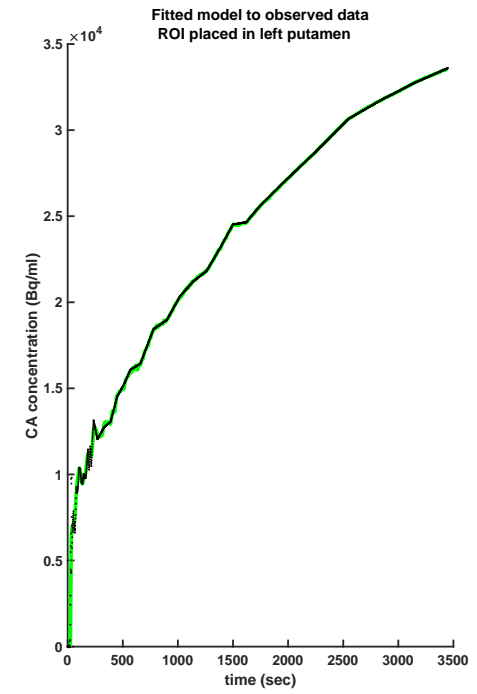
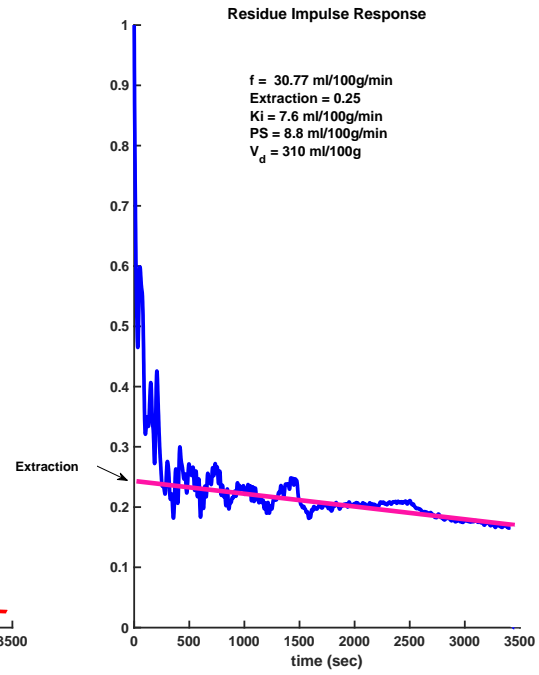
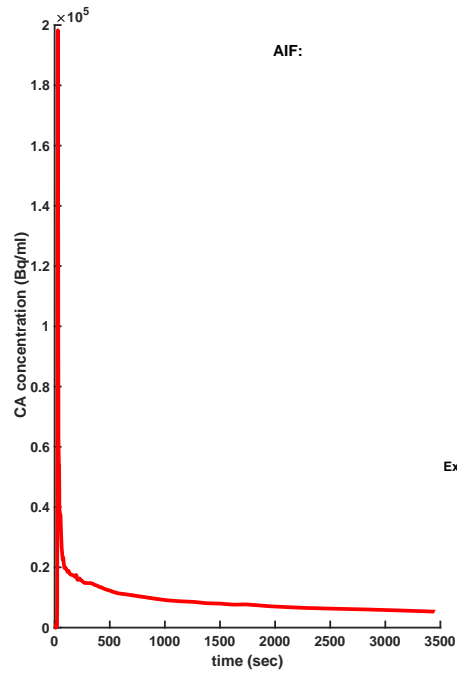




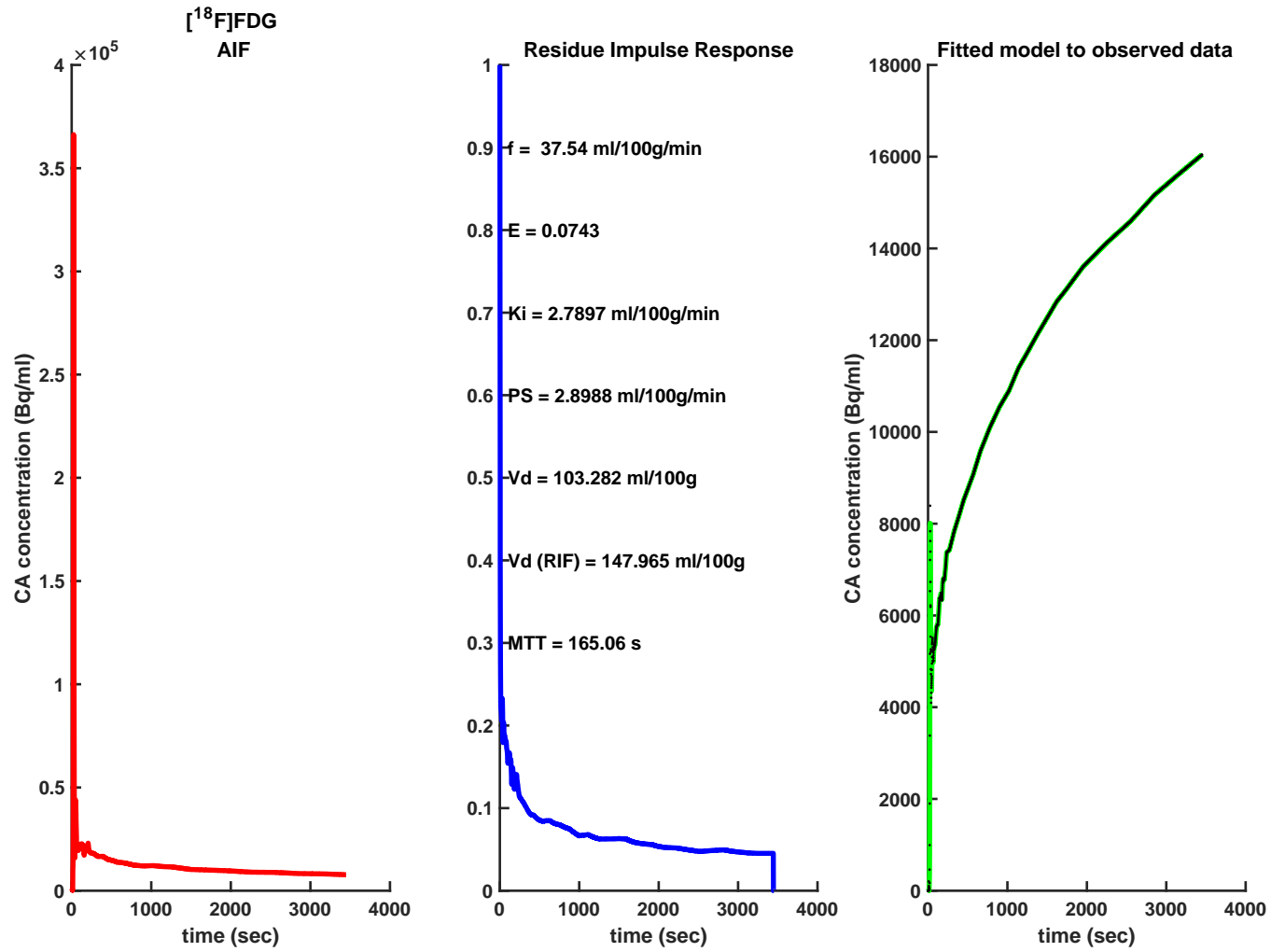
Increasing diffusion limited

- [¹⁵O]H₂O
- [¹⁸F]FE-PE2I
- [¹¹C]PIB
- [¹⁸F]FDG
- [¹⁸F]FET
- [⁶⁸Ga]Ga-DOTATATE

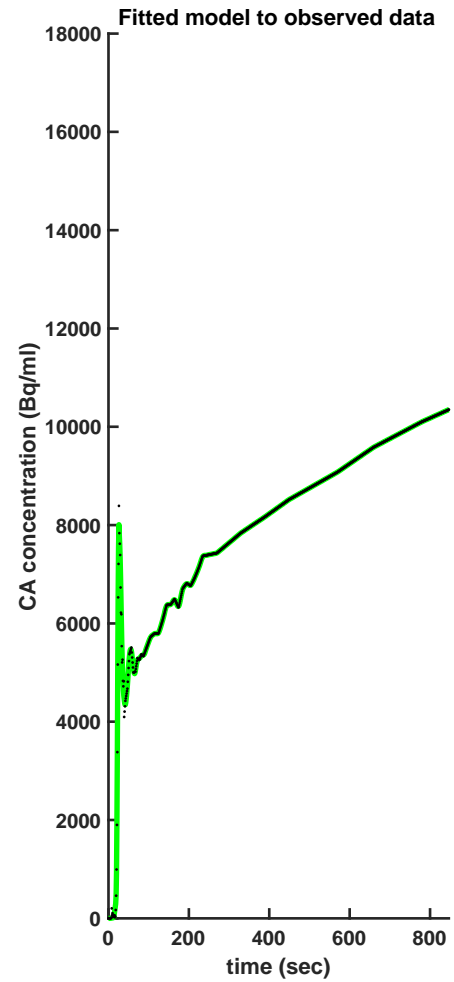
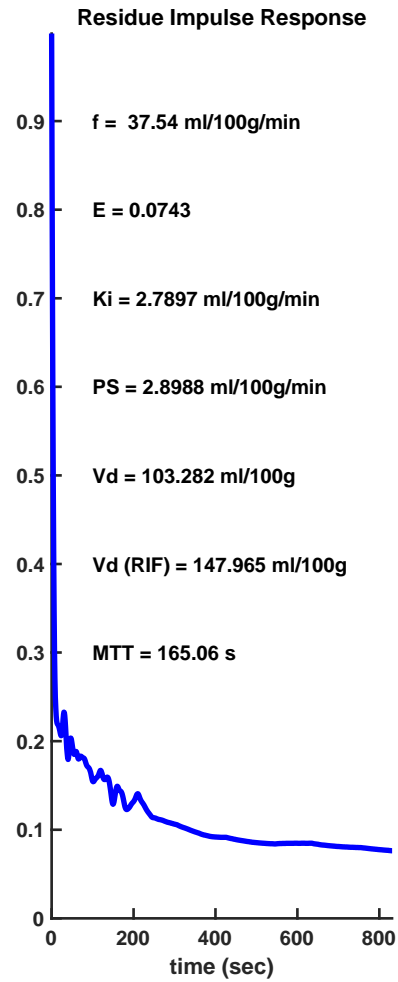
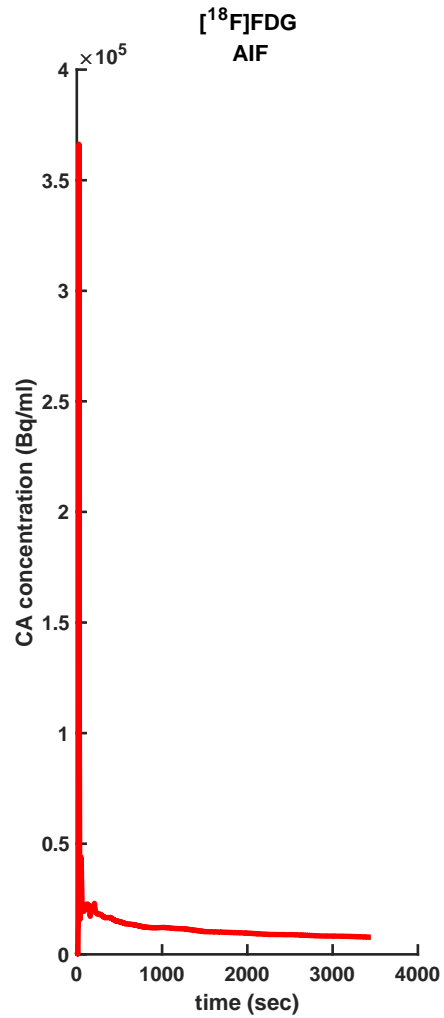
Dynamic [¹⁸F]FDG Quadra-PET scan



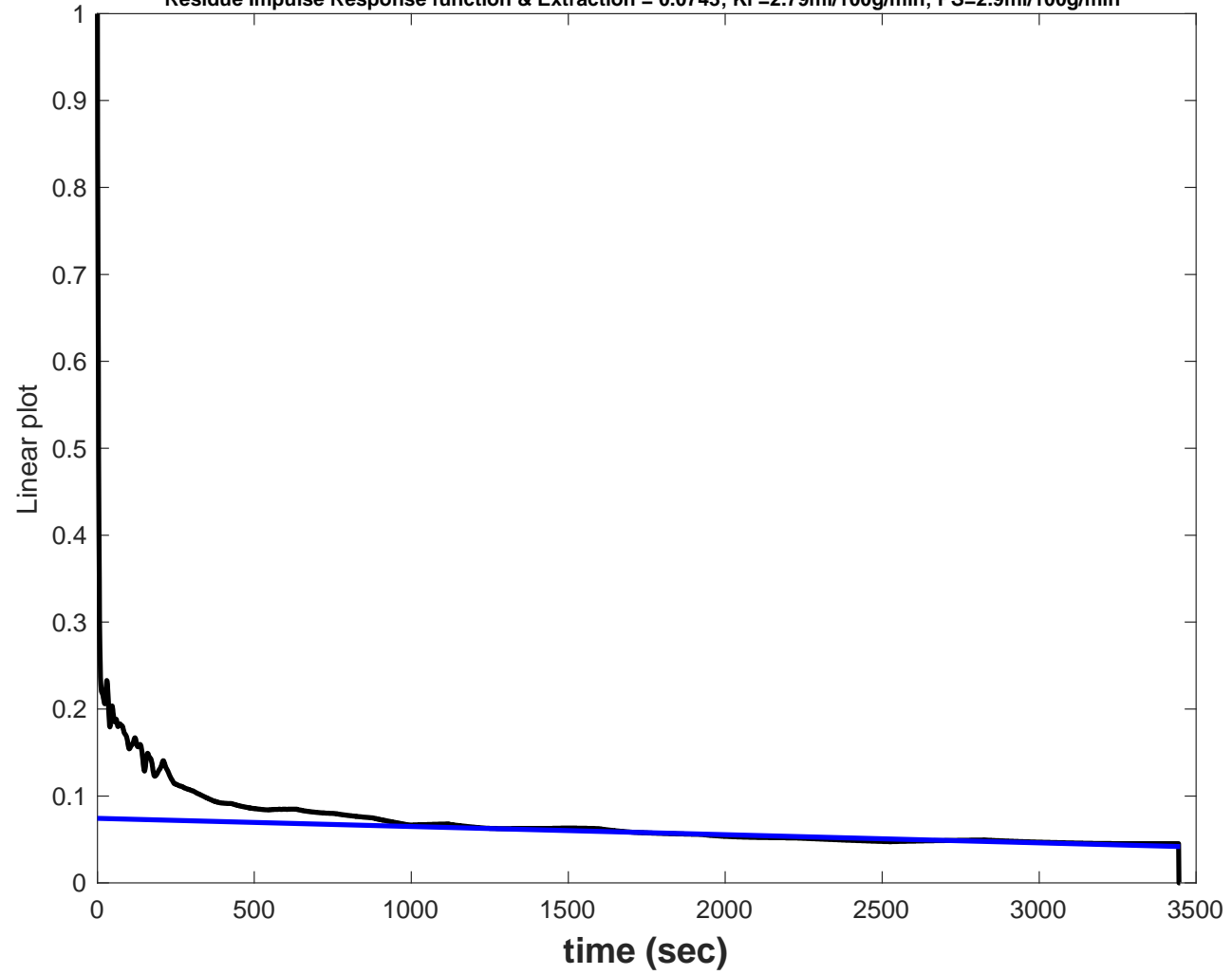
Large ROI – one slice

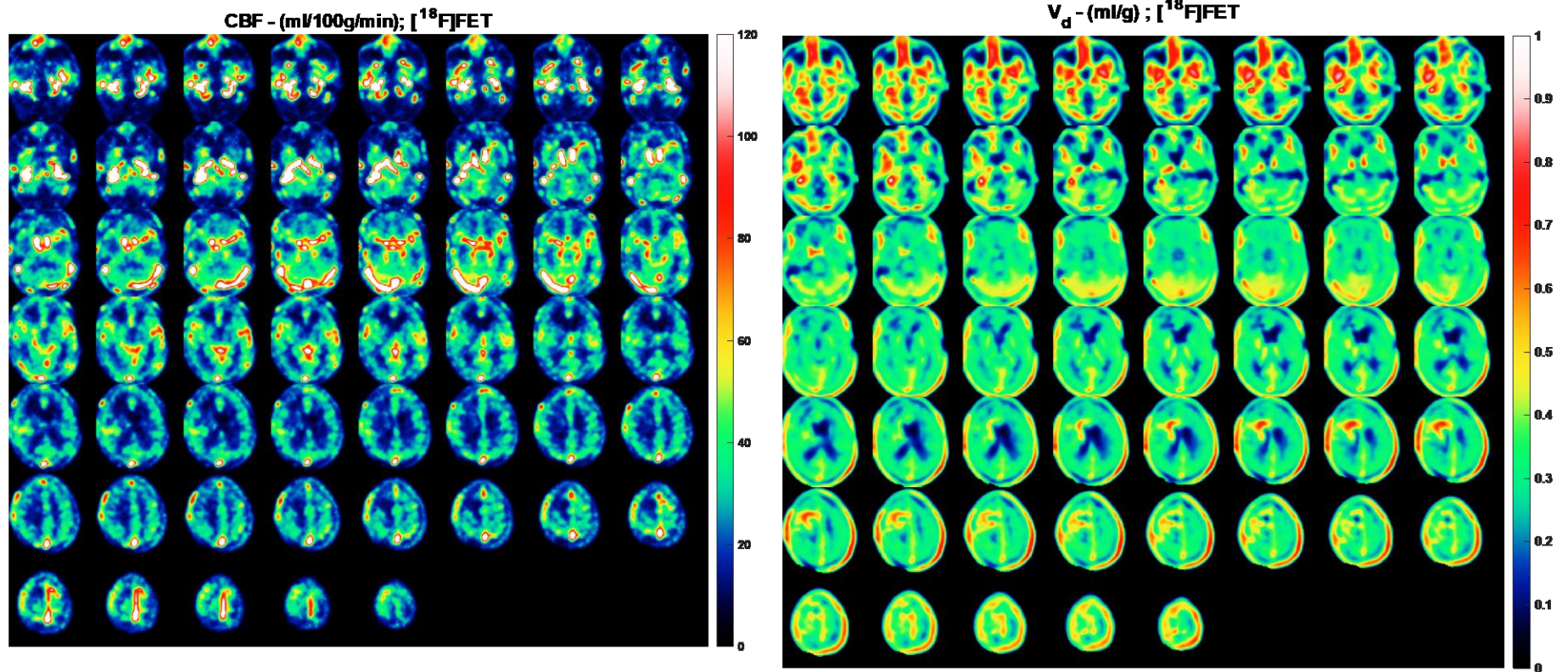


Large ROI – one slice



Residue Impulse Response function & Extraction = 0.0743; $K_i = 2.79\text{ml}/100\text{g}/\text{min}$; $PS = 2.9\text{ml}/100\text{g}/\text{min}$

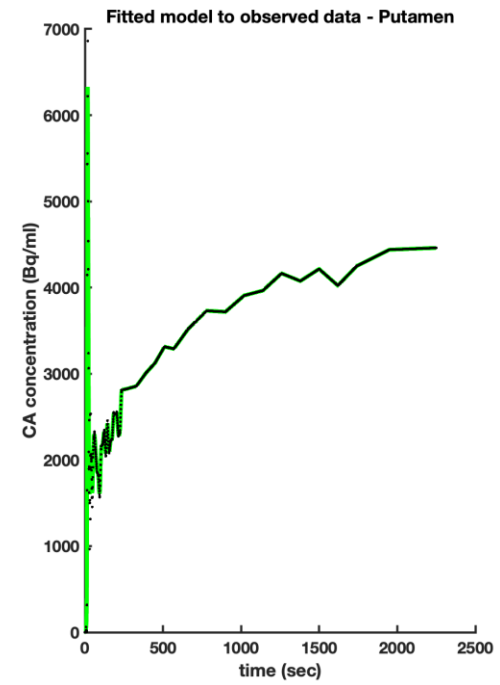
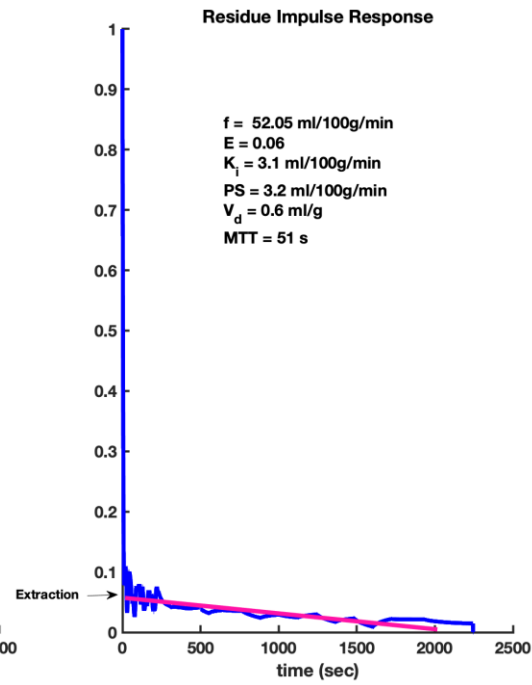
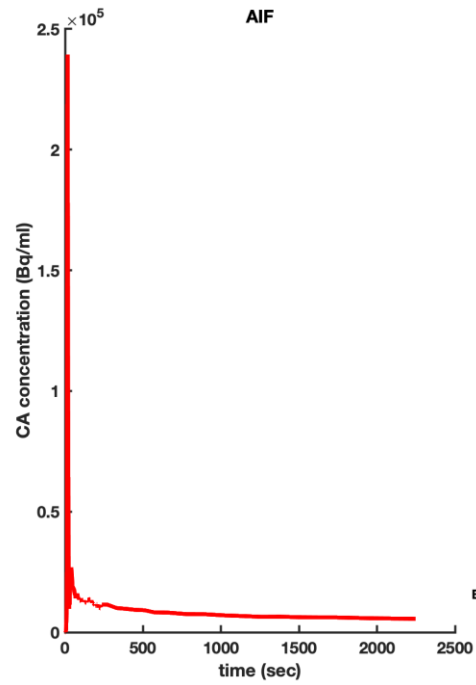




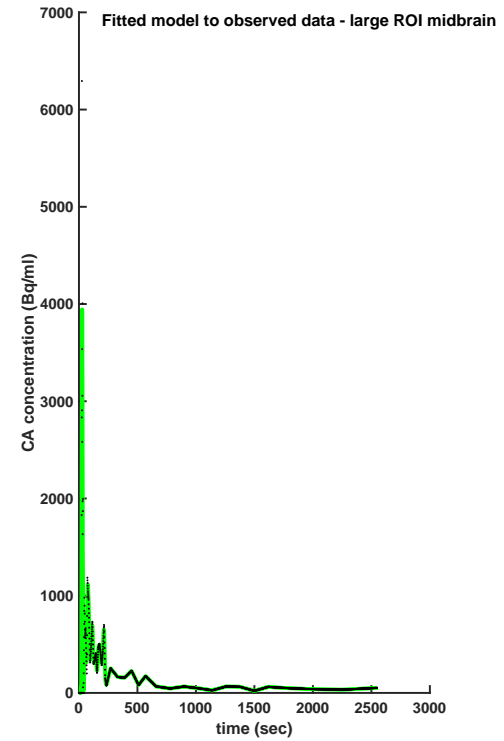
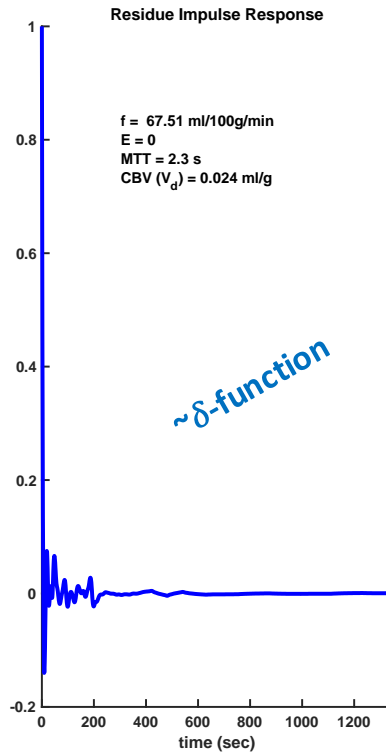
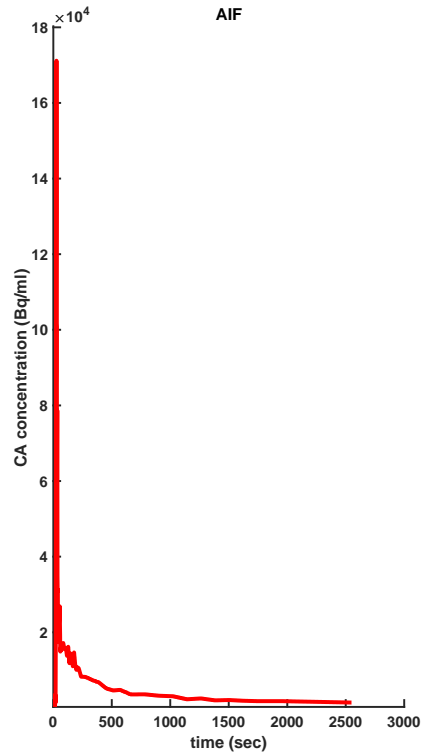
Increasing diffusion limited

- [¹⁵O]H₂O
- [¹⁸F]FE-PE2I
- [¹¹C]PIB
- [¹⁸F]FDG
- [¹⁸F]FET
- [⁶⁸Ga]Dotate

Dynamic [¹⁸F]FET Quadra-PET scan



Dynamic $[^{68}\text{Ga}]\text{Ga-DOTATATE}$ Quadra-PET scan



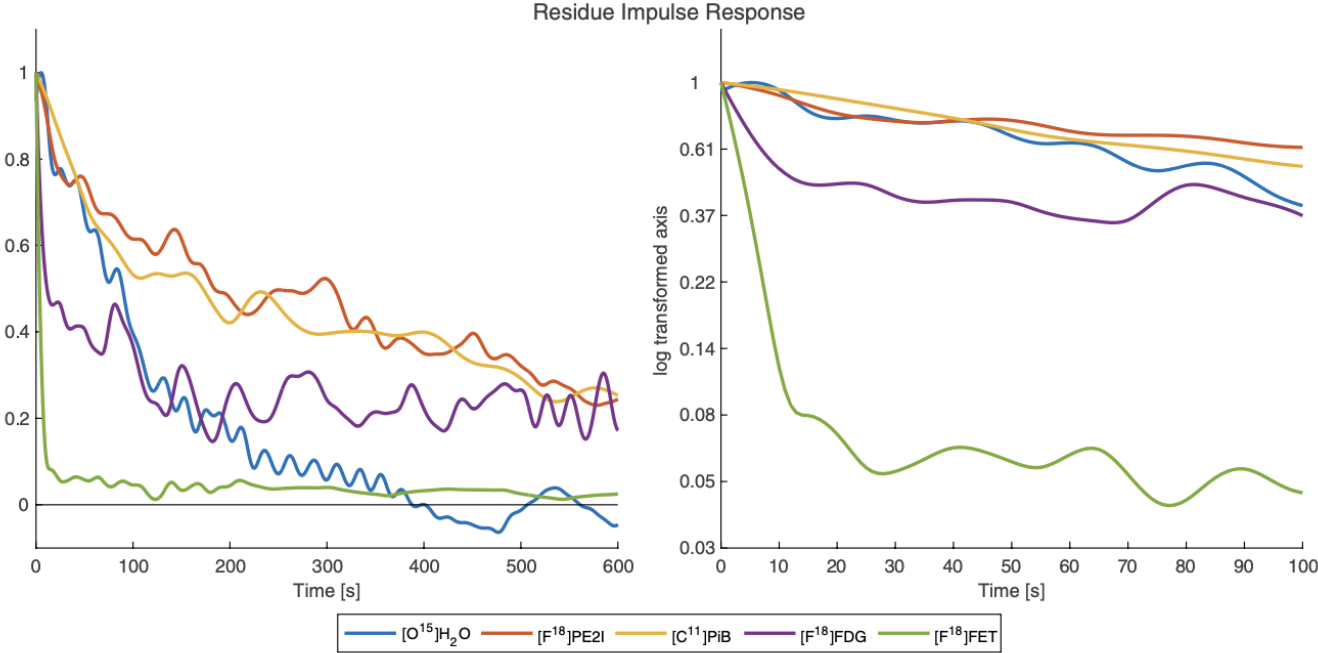


Fig. 2 The residue impulse response functions from Fig. 1 depicted on a smaller time scale for better comparison. Both linear and semilogarithmic plots are shown. The configurations of each tracer correspond to the expected behaviour of the tracers

Fig. 4 Each row shows the perfusion maps and volume of distribution maps for **a** [^{11}C] PIB, **b** [^{18}F]FE-PE2I, **c** 2- ^{18}F]FDG and **d** [^{18}F]FET for four different patients. Patient (a) had Alzheimer's disease and pronounced beta-amyloid accumulation, and the CBF maps show typical parieto-temporal perfusion reduction (left-sided). Patient (b) was eventually diagnosed with major depression, and the CBF and volume of distribution maps were normal. Patient (c) had lung cancer with metastasis, but PET/CT of the brain did not disclose CNS involvement. Patient (d) had previously undergone surgery for brain cancer (glioblastoma), and the CBF maps show CBF reduction/no perfusion corresponding to the resection cavity, and the volume of distribution map shows abnormal frontal subcortical FET uptake, suggesting tumour recurrence. All images are shown in native orientation to avoid interpolation artefacts

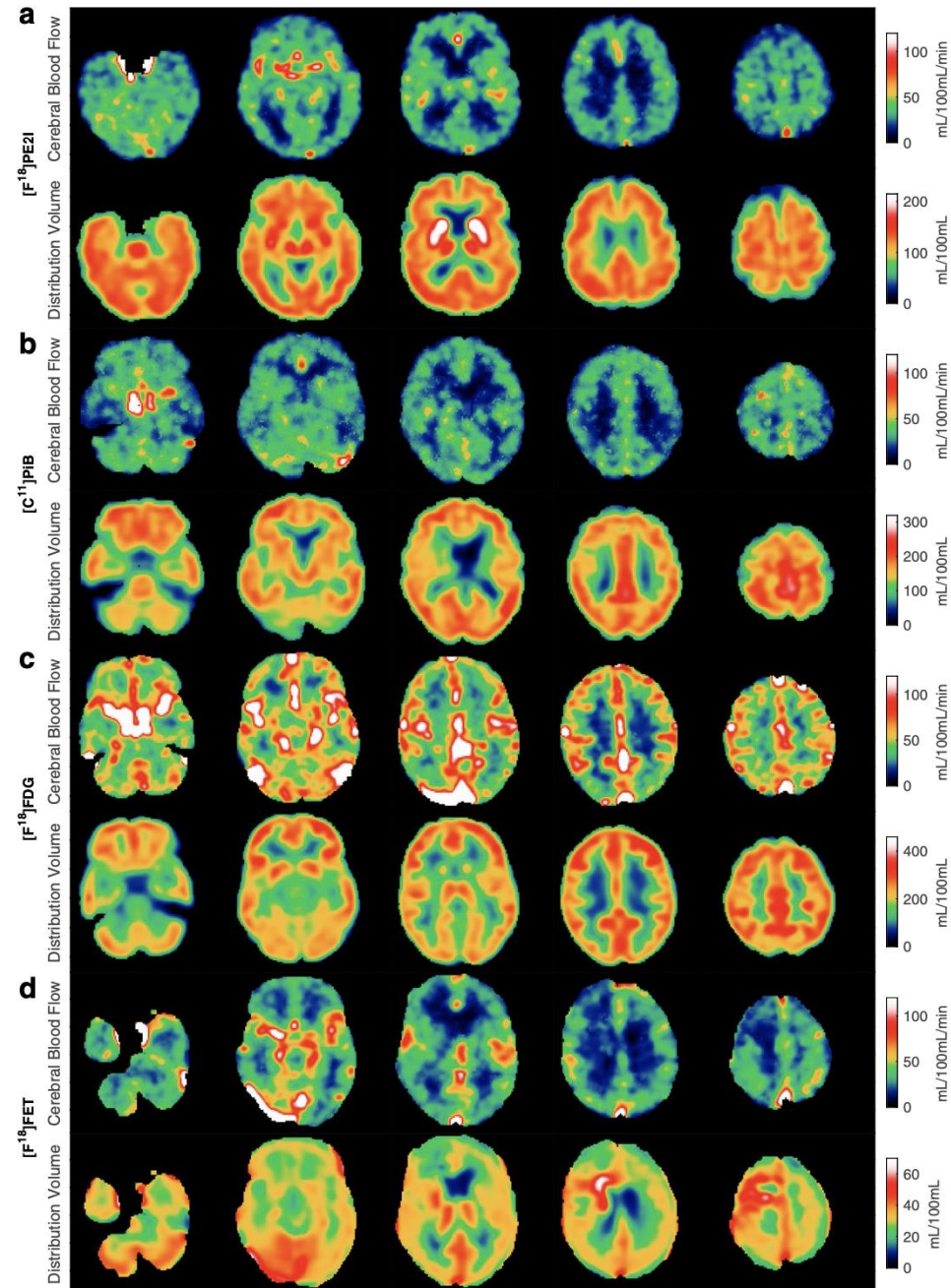


Table 3 Tikhonov method

Radiopharmaceutical	Number of subjects/ROIs	Mean CBF \pm SD (mL/min/100 mL)	Mean $E \pm$ SD	Mean $K_1 \pm$ SD (mL/min/100 mL)	Mean $v_d \pm$ SD (mL/100 mL)
[¹⁵ O]H ₂ O rest	5/5	69 \pm 19	0.94 \pm 0.06	-	90 \pm 5.6
[¹⁵ O]H ₂ O acetazolamide	5/5	115 \pm 31	0.96 \pm 0.04	-	92 \pm 6.9
[¹⁸ F]FE-PE2I	5/5	58 \pm 15	0.78 \pm 0.08	45 \pm 16	383 \pm 98
[¹¹ C]PiB	5/5	50 \pm 7	0.62 \pm 0.08	31 \pm 8	288 \pm 90
[¹⁸ F]FDG	5/5	37 \pm 13	0.19 \pm 0.05	5.5 \pm 1.2	-
[¹⁸ F]FET	5/5	53 \pm 16	0.032 \pm 0.008	1.7 \pm 0.2	48 \pm 9

CBF cerebral blood flow, *E* extraction fraction, *K₁* unidirectional influx constant, *v_d* volume of distribution

European Journal of Nuclear Medicine and Molecular Imaging

Table 4 Conventional compartment model

Radiopharmaceutical	Number of subjects/ROIs	Mean $K_1 \pm$ SD (mL/min/100 mL)	Mean $k_2 \pm$ SD (1/min)	Mean $k_3 \pm$ SD (1/min)	Mean $V_a \pm$ SD (mL/100 mL)
[¹⁵ O]H ₂ O rest	5/5	72.6 \pm 18.3	0.76 \pm 0.17	-	0.88 \pm 0.62
[¹⁵ O]H ₂ O acetazolamide	5/5	115.7 \pm 30.2	1.21 \pm 0.27	-	2.07 \pm 1.16
[¹⁸ F]FE-PE2I	5/5	47.6 \pm 12.3	0.12 \pm 0.03	<0.0004	3.26 \pm 2.02
[¹¹ C]PiB	5/5	41.1 \pm 4.6	0.16 \pm 0.05	0.003 \pm 0.004	4.3 \pm 0.82
[¹⁸ F]FDG	5/5	12.8 \pm 1.2	0.13 \pm 0.03	0.051 \pm 0.008	1.93 \pm 1.79
[¹⁸ F]FET	5/5	2.9 \pm 0.7	0.082 \pm 0.02	0.027 \pm 0.010	2.94 \pm 0.55

K_1 is the unidirectional influx rate constant over the blood–brain barrier, k_2 is a rate constant related to back diffusion of tracer from the reversible compartment to blood, k_3 is a rate constant related to the irreversible binding of tracer in tissue, V_a is the blood volume in tissue

Conclusion

Perfusion can be estimated for (nearly) all types of tracers - high time resolution

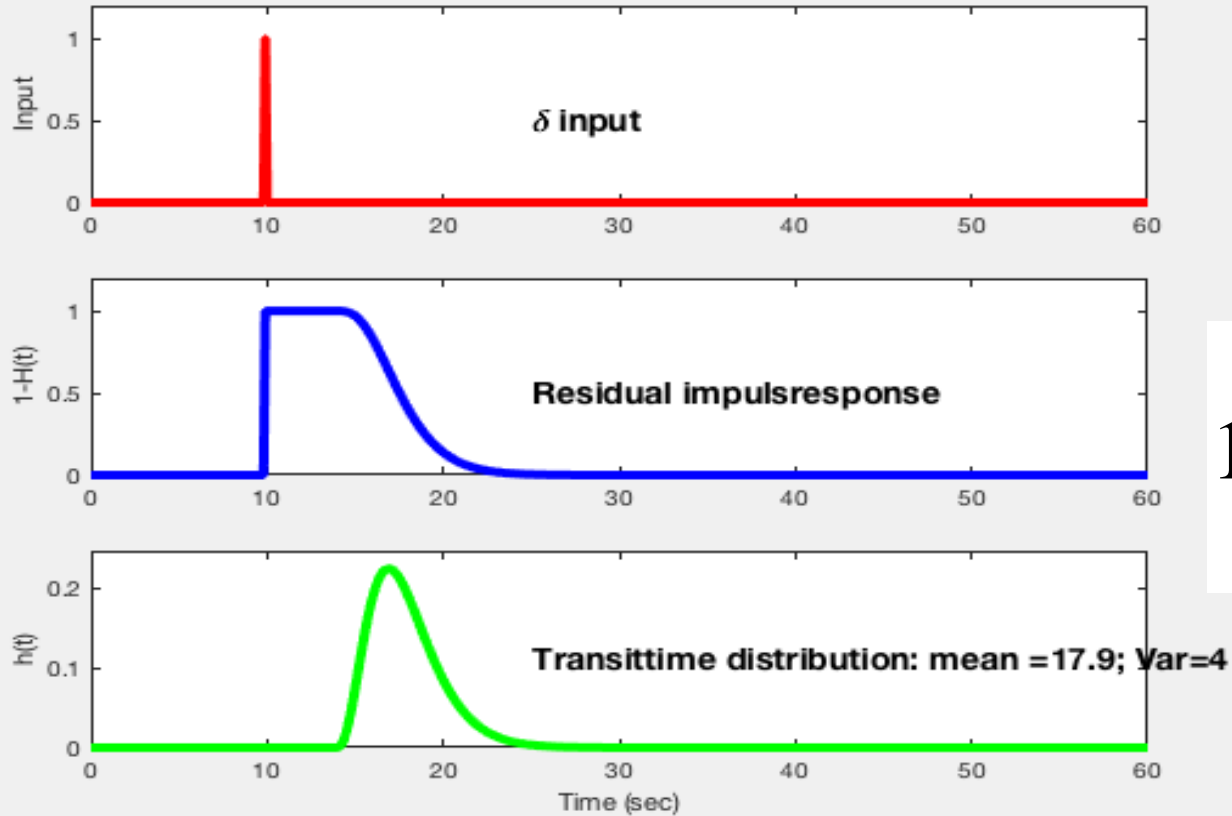
Residue impulse response function can be estimated-

giving K_i and E

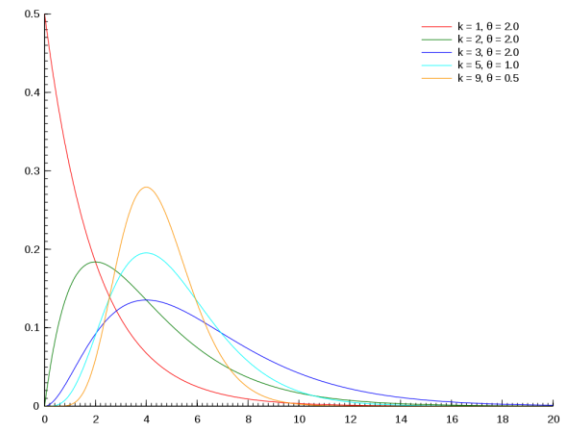
Yes we can !!!!



CTH modeling using gamma distribution



$$1 - H(t) = 1 - \int_0^t h(\tau) d\tau$$



$$f(x; \alpha, \beta) = \frac{x^{\alpha-1} e^{-\beta x} \beta^\alpha}{\Gamma(\alpha)} \quad \text{for } x > 0 \quad \alpha, \beta > 0.$$

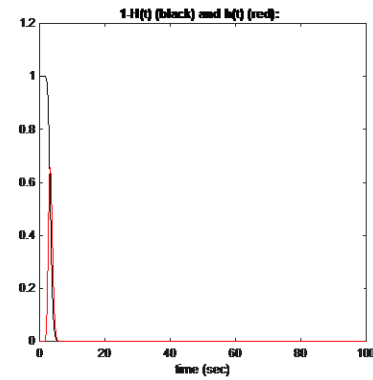
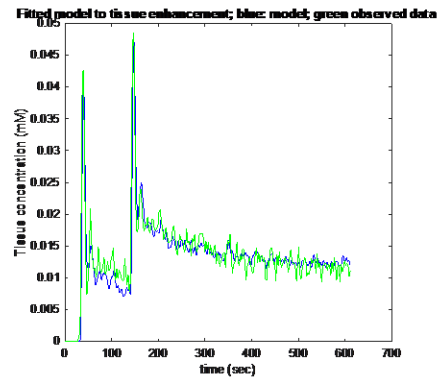
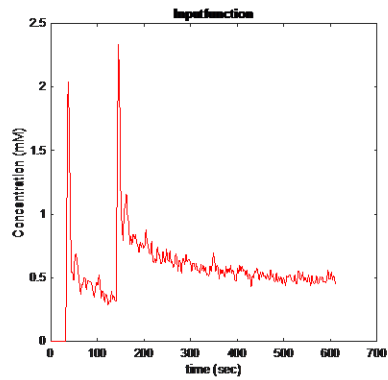
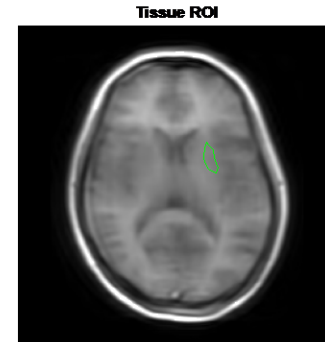
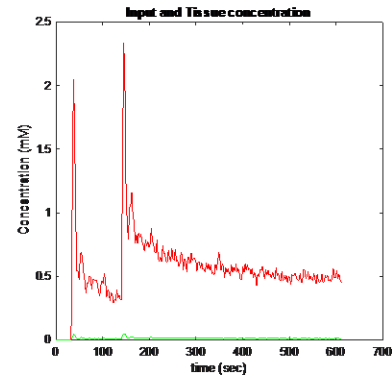
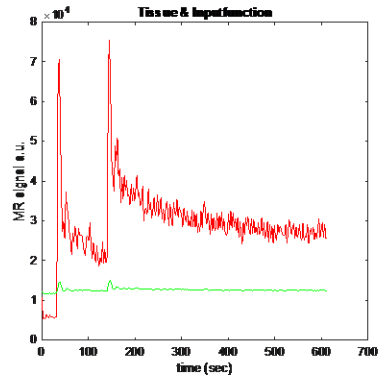
MRI using MR contrast agent

BrainMTCIH model June 2024: IJ+max, mTT, alpha

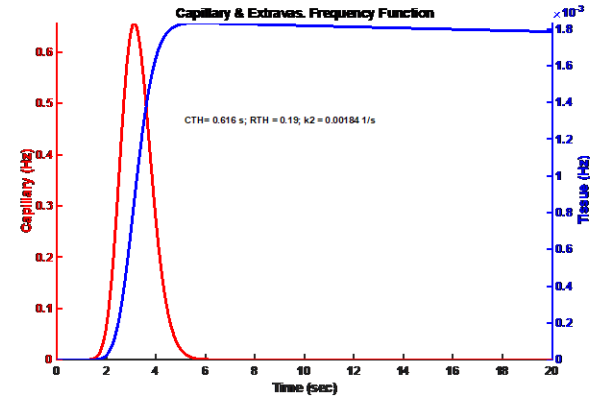
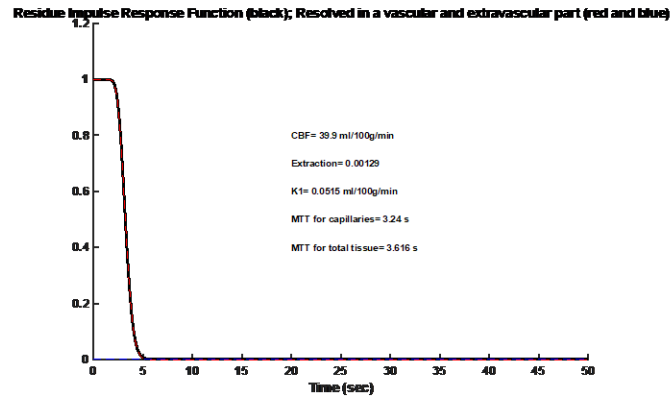
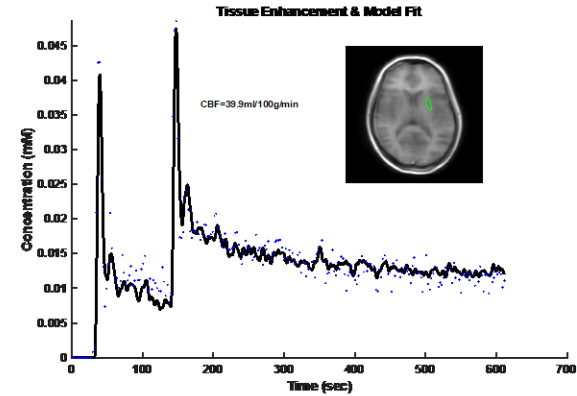
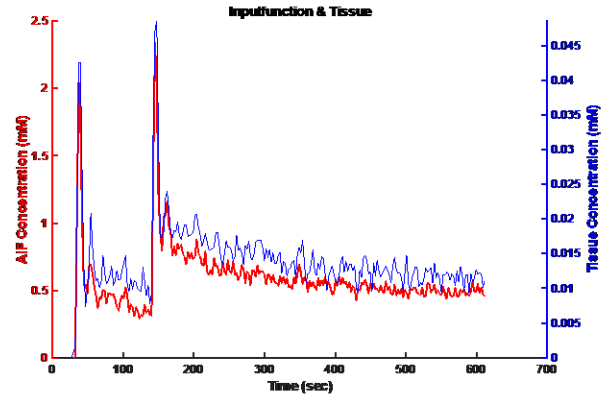
Tissue: $Af_{scs}/hca/c2/D/024032142_job/loc_m_bloodT1_m_sp_pi_fscs_R/Signal_in_loc5_p_stemon_mat_pat$
 Input: $Af_{scs}/hca/c2/D/024032142_job/loc_m_bloodT1_m_sp_pi_fscs_R/Signal_in_loc2_ICA_loc2_calcd.mat$
 Image: $Af_{scs}/hca/c2/D/024032142_job/2024032142_3_PAR$

pixels: tissue = 61; pixels: input = 33

f = 39.9 ml/100g/min; Vd = 2.41 ml/100g; mTT = 0 s; tmax = 3.12s; capil SD (from Mf) = 0.6164s; MTT tissue label = 3.62s; M TT for capillaries = 3.24s; afibcd = 0.853s; r = 0.9241 alpha = 26.6247; ss = 0.052683



MRI using MR contrast agent



PET using $[^{18}\text{F}]\text{-FET}$

Brainfit: Fluoresilytyrosin - CTH model june 2024: f, tmax, mTT, alpha

Tissue: : pat

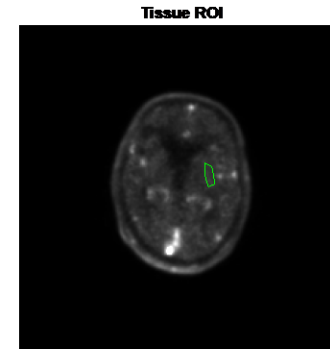
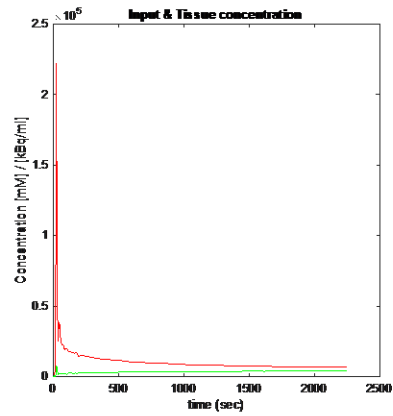
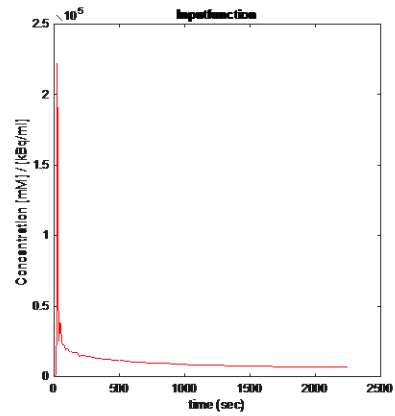
Input:

slice no: 23,

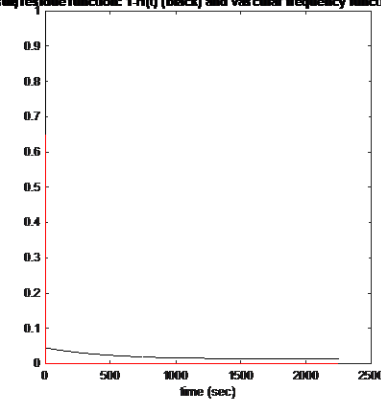
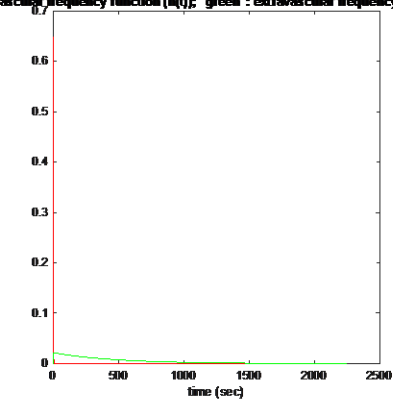
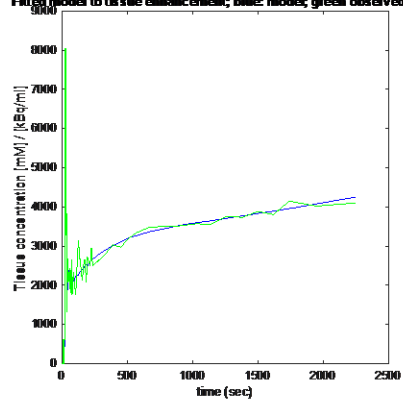
Image

pixels tissue = 75

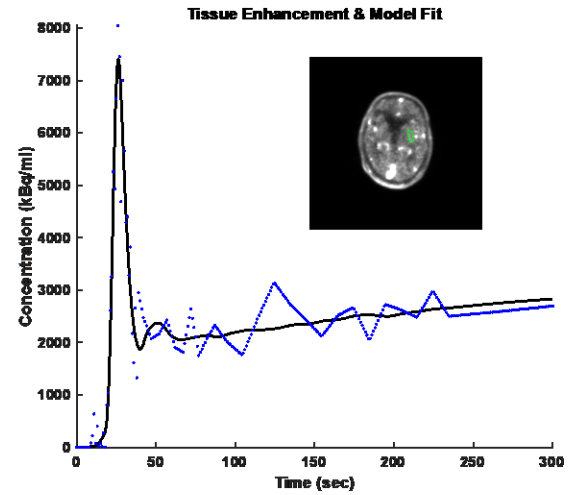
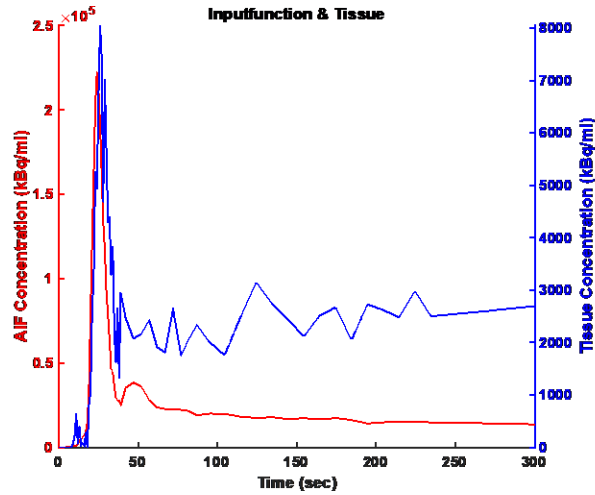
f = 59.9 ml/100g/min; Vd = 33.5ml/100g; mTT = 0 s; tmax = 3.36s; capil SD (from h(t)) = 0.6238s; MTT tissue total = 33.6s; MTT for capillaries = 3.47s; Extraction = 0.0449; k2 = 12.9ml/100g/min; K1 = 2.69ml/100g/min; alpha = 0.067s; r = 0.98031; ss = 144707686.7219



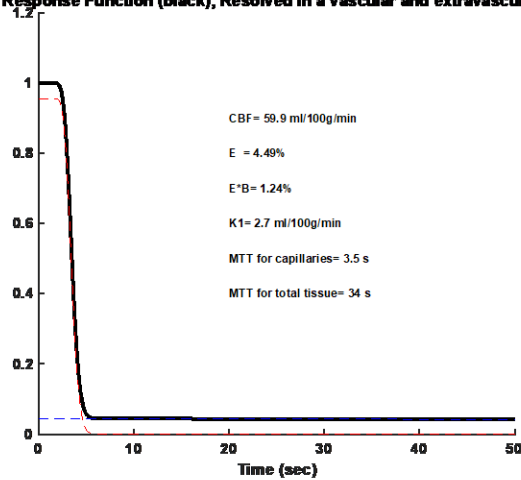
Fitted model to tissue enhancement; blue model; green observed data; vascular frequency function $h(t)$; "green": extravascular frequency $h(t) \cdot (1 - \alpha)$; residue function: $1 - h(t)$ (black) and vascular frequency function: $h(t)$ (red):



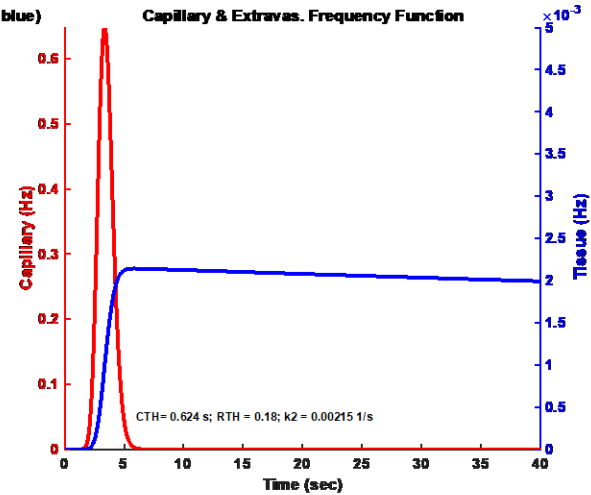
PET using $[^{18}\text{F}]\text{-FET}$



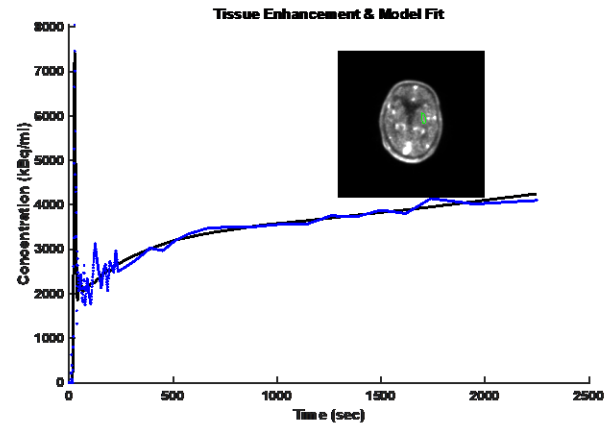
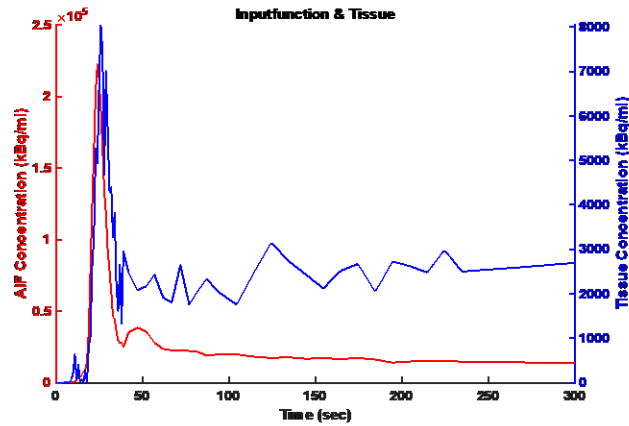
Residue Impulse Response Function (black); Resolved in a vascular and extravascular part (red and blue)



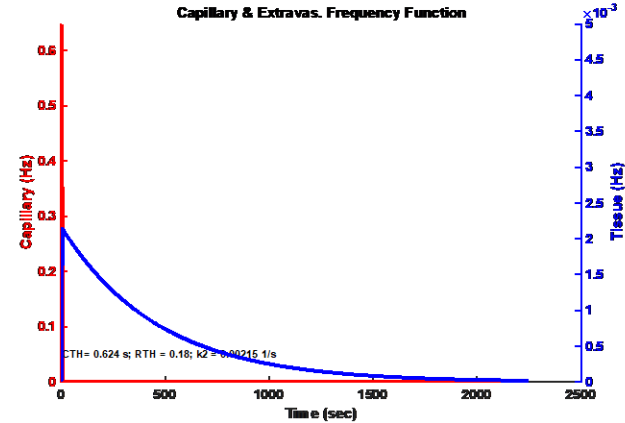
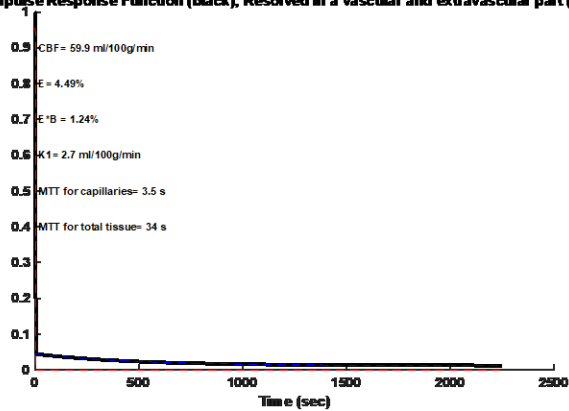
Capillary & Extravas. Frequency Function



PET using $[^{18}\text{F}]\text{-FET}$



Residue Impulse Response Function (black); Resolved in a vascular and extravascular part (red and blue)



Yes we can !!!!



Kety's methods

The inventor of classic tracer kinetic theory

Measurement of local blood flow by the exchange of an inert, diffusible substance

$\text{CF}_3\text{I}^{131}$ and I^{131} -antipyrine

Kety's methods (Residue detection)



$$C_t(t) \equiv \frac{n(t)}{W_{\text{eight}}}$$

or

$$C_t(t) \equiv \frac{n(t)}{V_{\text{olume}}}$$

$$\lambda \equiv \frac{C_t^{\infty}}{C_{\text{blood}}^{\infty}} \approx \frac{C_t(t)}{C_o(t)}$$

Kety's methods (Residue detection)



$$\frac{dn(t)}{dt} = j_i(t) - j_o(t)$$

$$W \frac{dC_t(t)}{dt} = F C_a(t) - F C_o(t)$$

$$W \frac{dC_t(t)}{dt} = F C_a(t) - \frac{F}{\lambda} C_t(t) \Leftrightarrow \frac{dC_t(t)}{dt} = \frac{F}{W} C_a(t) - \frac{F}{W\lambda} C_t(t)$$

Kety's methods (Residue detection)



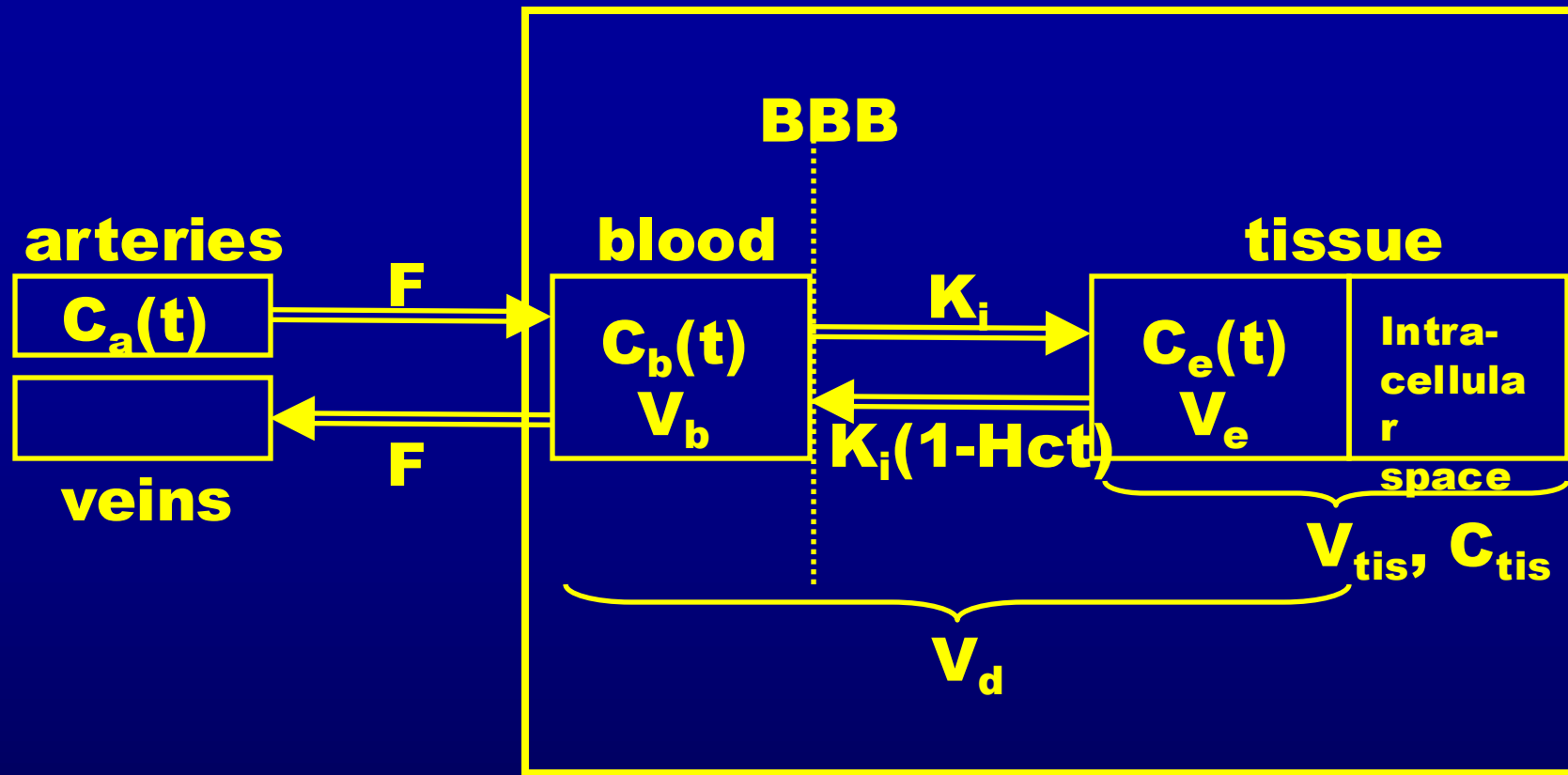
$$\frac{dC_t(t)}{dt} = \frac{F}{W} C_a(t) - \frac{F}{W\lambda} C_t(t)$$

$$\frac{dC_t(t)}{dt} = f C_a(t) - \frac{f}{l} C_t(t)$$

Solution:

$$C_t(t) = f \int_0^t C_a(t) e^{-\frac{f}{l}(t-t)} dt$$

$$\hat{U} C_t(t) = f e^{-\frac{f}{l}t} \int_0^t C_a(t) e^{\frac{f}{l}t} dt$$



$$V_b \frac{dC_b(t)}{dt} = F C_a(t) - (F + K_i) C_b(t) + K_i(1 - Hct) C_e(t)$$

$$V_e \frac{dC_e(t)}{dt} = K_i C_b(t) - K_i(1 - Hct) C_e(t)$$

$$V_e C_e = V_{\text{tis}} C_{\text{tis}}$$

$$\alpha = \frac{F + K_i}{V_b}$$

$$\beta = \frac{V_{\text{tis}}(1 - \text{Hct})K_i}{V_b V_e}$$

$$\gamma = \frac{K_i}{V_{\text{tis}}}$$

$$\theta = \frac{K_i(1 - \text{Hct})}{V_e}$$

$$(a, b) = \left(\frac{1}{2}[\theta + \alpha + \sqrt{\theta^2 + \alpha^2 - 2\theta\alpha + 4\gamma\beta}], \frac{1}{2}[\theta + \alpha - \sqrt{\theta^2 + \alpha^2 - 2\theta\alpha + 4\gamma\beta}] \right)$$

$$C_b(t) = C_a(t) \otimes \frac{F}{V_b} \frac{(a - \theta)e^{-at} - (b - \theta)e^{-bt}}{a - b}$$

$$C_{\text{tis}}(t) = C_a(t) \otimes \frac{F}{V_b} \frac{K_i}{V_{\text{tis}}} \frac{e^{-bt} - e^{-at}}{a - b}$$

$$C_t(t) = V_b C_b(t) + (1 - V_b) C_{\text{tis}}(t) \Leftrightarrow$$

$$C_t(t) = F C_a(t) \otimes \left[\frac{(a - \theta - K_i / V_b) e^{-at} + (-b + \theta + K_i / V_b) e^{-bt}}{a - b} \right]$$

Kety's methods (Residue detection and no outflow)



$$\frac{dn(t)}{dt} = j_i(t)$$

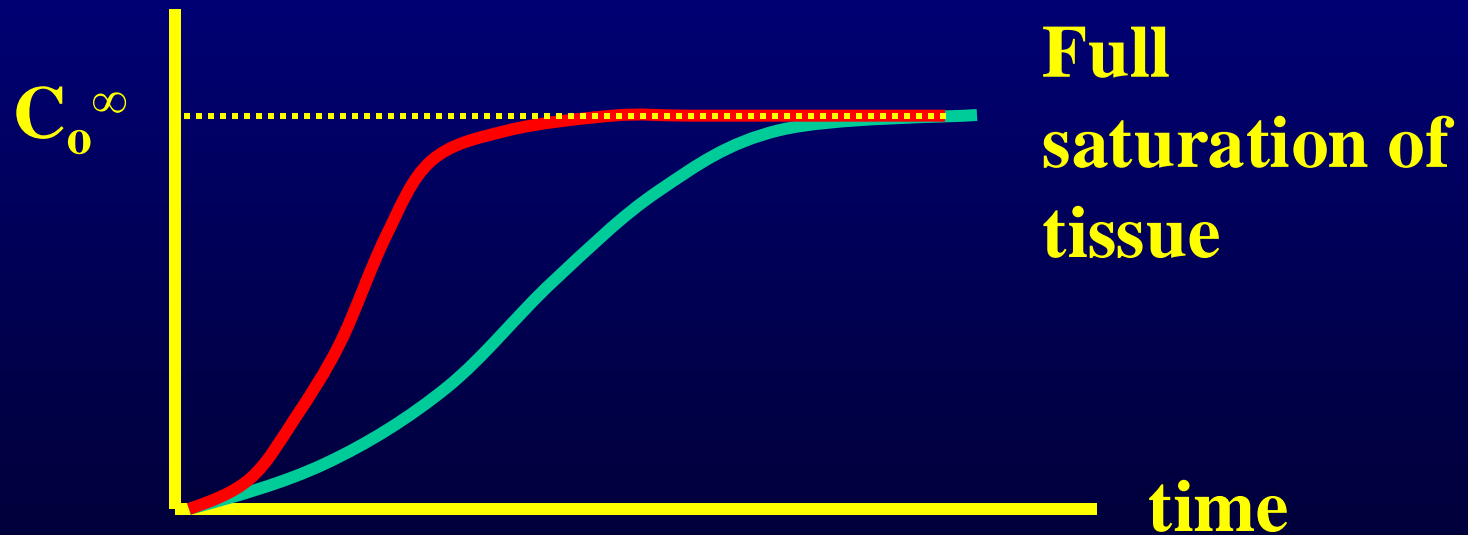
$$\Leftrightarrow W \frac{dC_t(t)}{dt} = F C_a(t)$$

$$\Leftrightarrow dC_t(t) = \frac{F}{W} C_a(t) dt$$

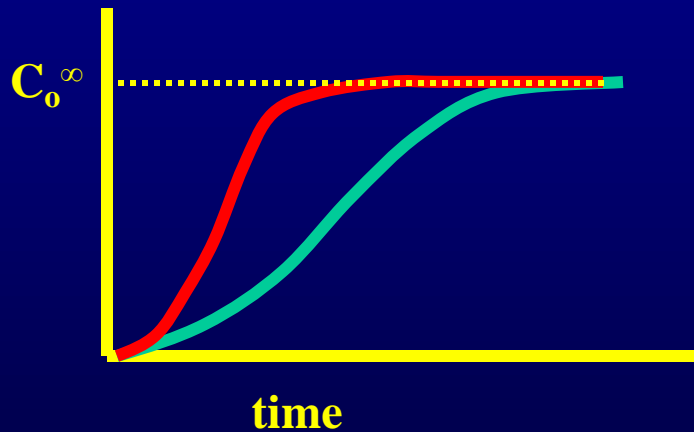
$$\Rightarrow \int_0^T dC_t(t) = \int_0^T \frac{F}{W} C_a(t) dt$$

$$\Leftrightarrow C_t(T) = \frac{F}{W} \int_0^T C_a(t) dt \Leftrightarrow \frac{F}{W} = \frac{C_t(T)}{\int_0^T C_a(t) dt}$$

Kety's methods (Inflow & outflow detection)



Kety's methods (Inflow & outflow detection)

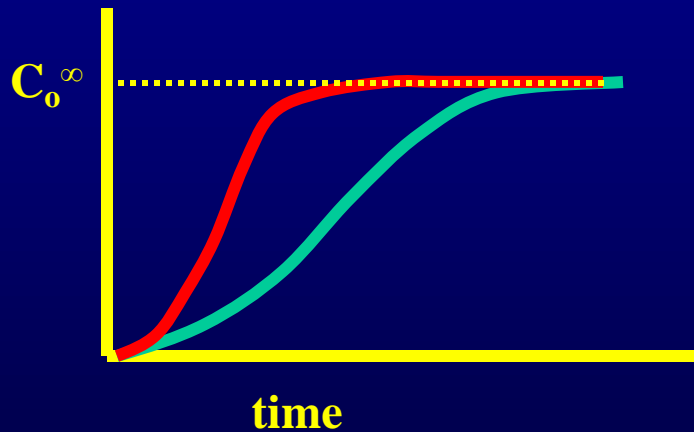


$$\frac{dn(t)}{dt} = j_i(t) - j_o(t)$$

$$\Leftrightarrow W \frac{dC_t(t)}{dt} = F C_a(t) - F C_o(t)$$

$$\Leftrightarrow dC_t(t) = \frac{F}{W} C_a(t) dt - \frac{F}{W} C_o(t) dt$$

Kety's methods (Inflow & outflow detection)



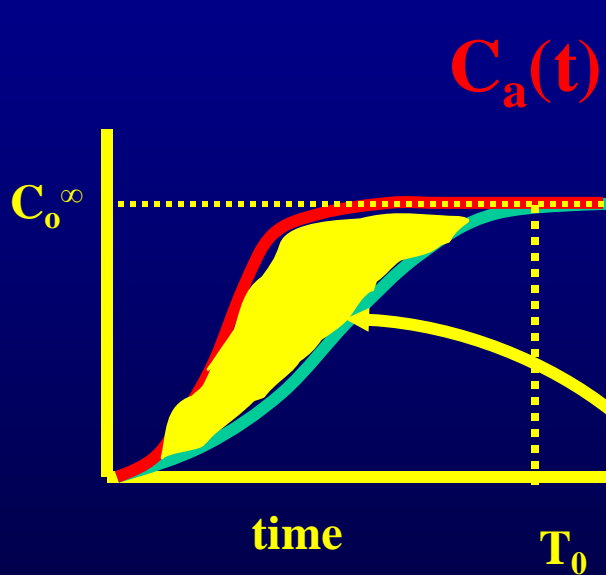
$$\Leftrightarrow dC_t(t) = \frac{F}{W} C_a(t) dt - \frac{F}{W} C_o(t) dt$$

$$\Leftrightarrow dC_t(t) = \frac{F}{W} (C_a(t) - C_o(t)) dt$$

$$\Leftrightarrow \int_0^T dC_t(t) = \frac{F}{W} \int_0^T (C_a(t) - C_o(t)) dt$$

$$\Leftrightarrow C_t(T) = \frac{F}{W} \int_0^T (C_a(t) - C_o(t)) dt$$

Kety's methods (Inflow & outflow detection)



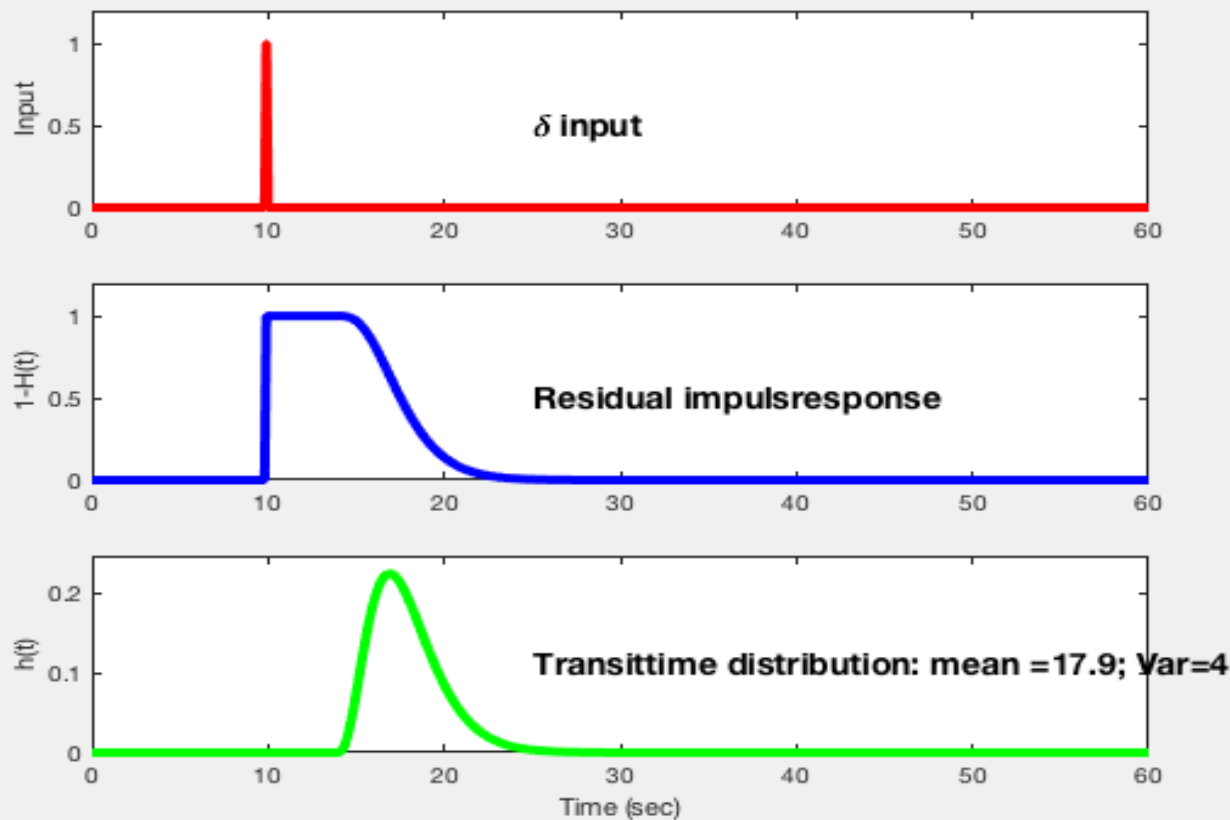
$$C_t(T) = / C_o(T) = / C_o^\infty \quad \text{for } T > T_0$$

$$C_t(T) = \frac{F}{W} \int_0^T (C_a(t) - C_o(t)) dt$$

$$\Rightarrow \frac{F}{W} = \frac{\lambda C_o^\infty}{\int_0^\infty (C_a(t) - C_o(t)) dt}$$

Brain capillary transit time heterogeneity in healthy volunteers measured by dynamic contrast-enhanced T_1 -weighted perfusion MRI.

Larsson HBW^{1,2}, Vestergaard MB¹, Lindberg U¹, Iversen HK^{2,3}, Cramer SP¹.



input



tissue



output

$$C_i(t) = C_a(t) \otimes f RIF(t) = f \int_0^t C_a(\tau) RIF(t - \tau) d\tau \quad [1]$$

$$RIF(t) = 1 - \int_0^t h(\tau) d\tau \quad [2]$$

The mean transit time (MTT) is given as:

$$MTT = \int_0^{\infty} t h(t) dt = \int_0^{\infty} RIF(t) dt \quad [3]$$

CTH can be defined as the standard deviation (SD) of the frequency function, $h(t)$:

$$CTH = \sqrt{Var[h(t)]} = \sqrt{\int (t - MTT)^2 h(t) dt} \quad [4]$$

The frequency function, $h(t)$, can be modelled as a simple gamma-variate function with the parametric form as (15):

$$h(t) = \left[\left(\frac{t - t_0}{t_{\max} - t_0} \right)^{\alpha} \exp \left(\alpha \left(1 - \frac{t - t_0}{t_{\max} - t_0} \right) \right) \right] / A \quad [5]$$

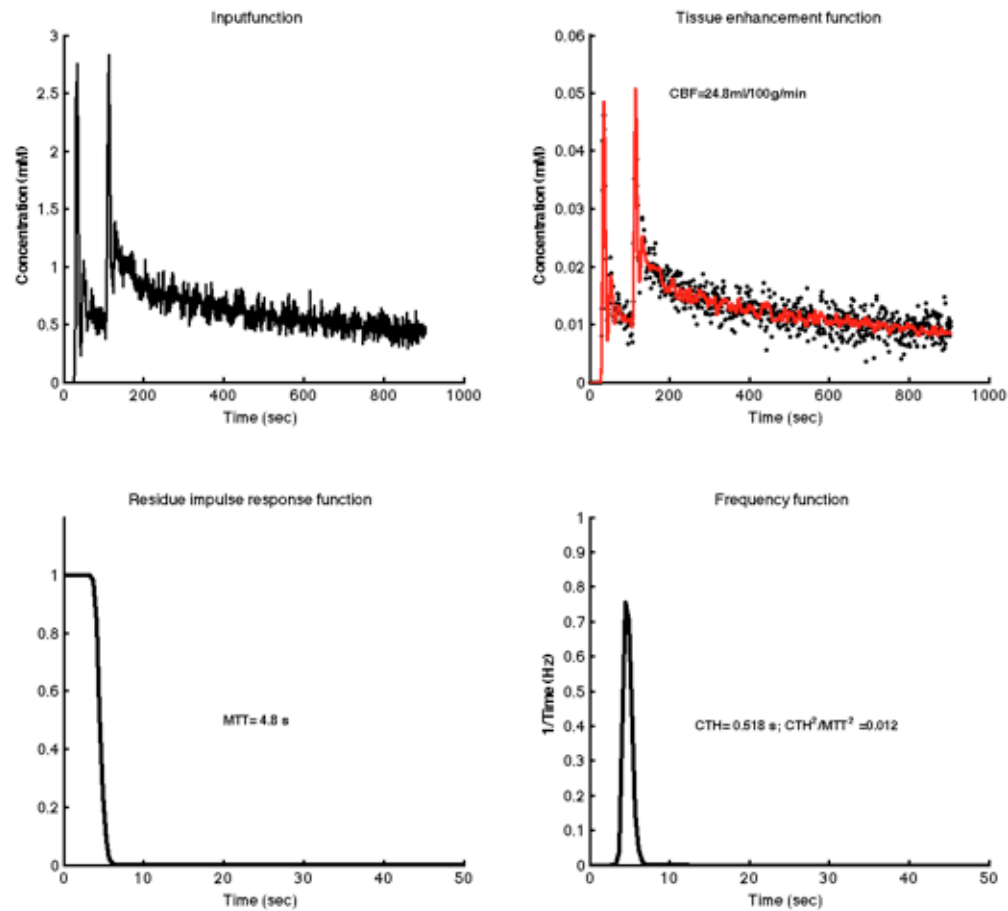


Figure 1. An example of calculation from a ROI placed in thalamus in a young healthy subject. Note the symmetrical shape of the $h(t)$ function. Mean Transit time (MTT), the Capillary Transit time Heterogeneity (CTH) and CTH^2/MTT^2 values are inserted.

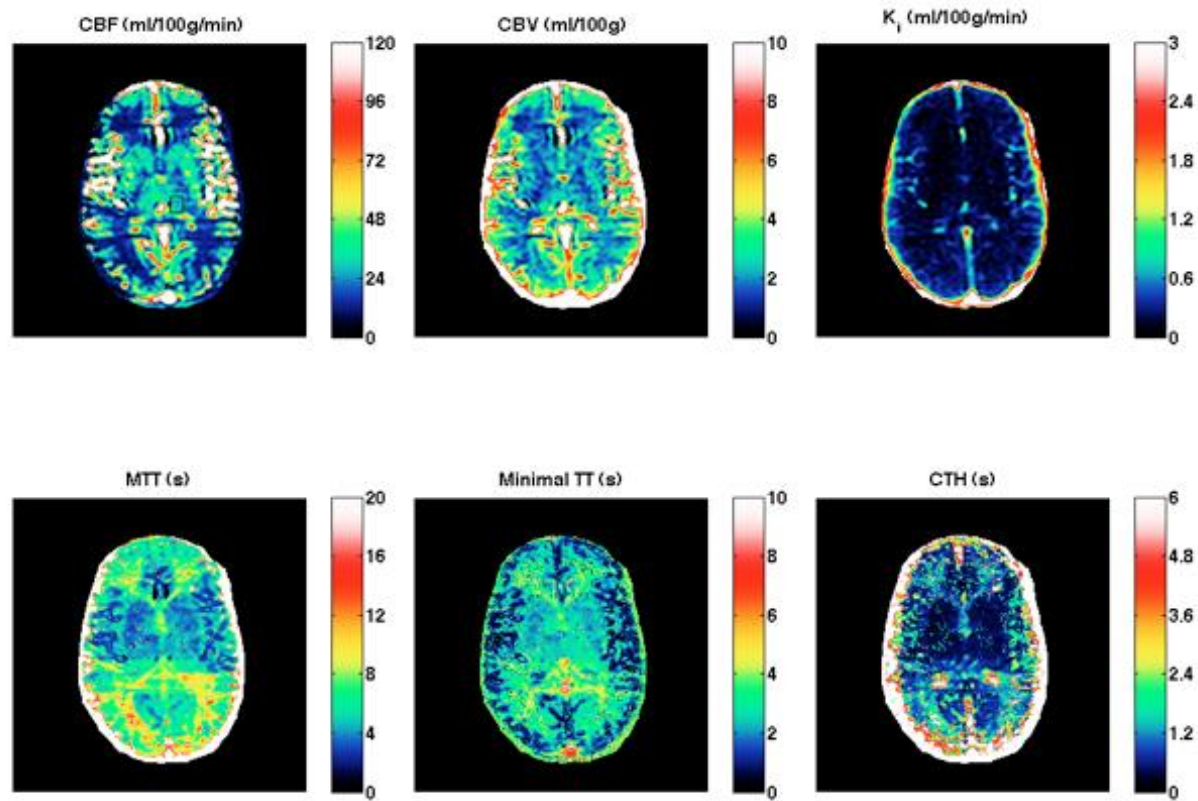


Figure 5. Pixel wise calculated maps of CBF, CBV, permeability K_i , mean transit time (MTT), minimal transit time (Minimal TT), and capillary transit time heterogeneity (CTH), of one healthy subject. The results from the ROI on the CBF map are shown in figure 1.

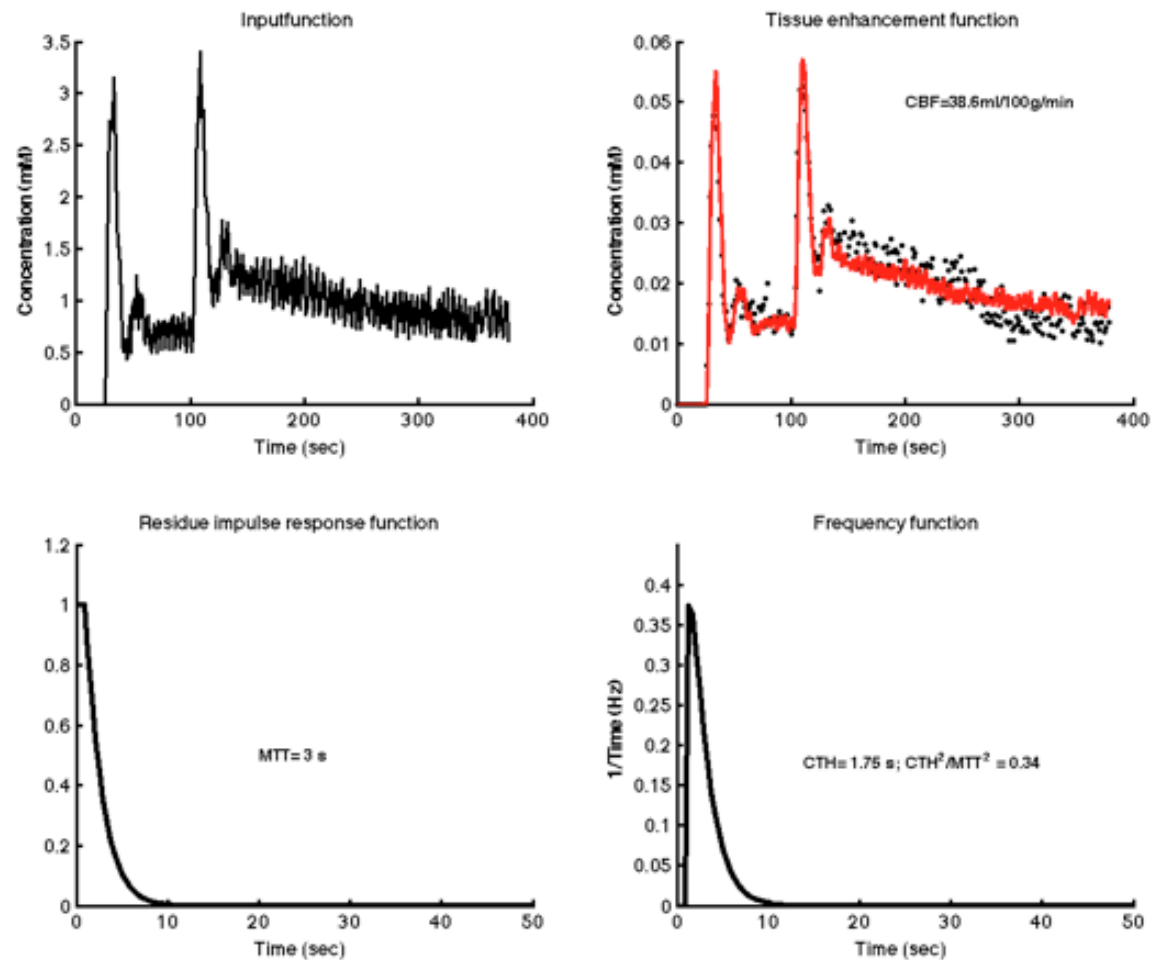


Figure 2. An example of calculation from a ROI placed in frontal WM of a 75-year-old man having internal carotid stenosis contralateral to the ROI placement.

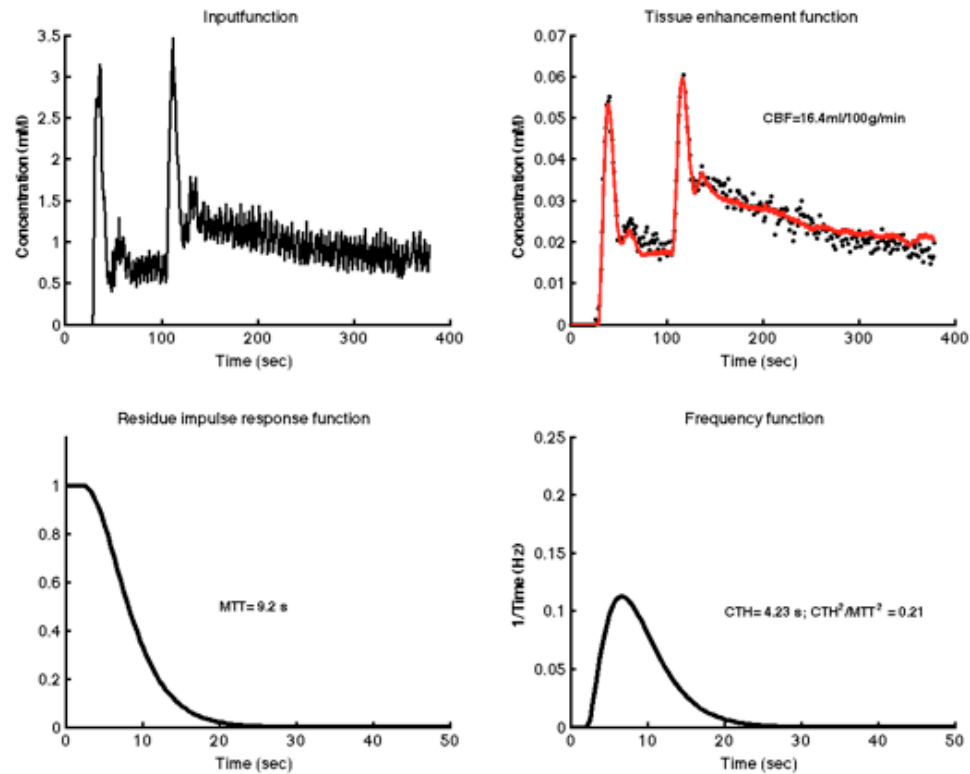


Figure 3. An example of calculation from a ROI placed in parietal WM ipsilateral to an internal carotid stenosis in a 75-year-old man. Mean Transit time (MTT), Capillary Transit time Heterogeneity (CTH) and CTH^2/MTT^2 values are inserted. Note the asymmetry of $h(t)$ signifying a large heterogeneity in capillary transit times.

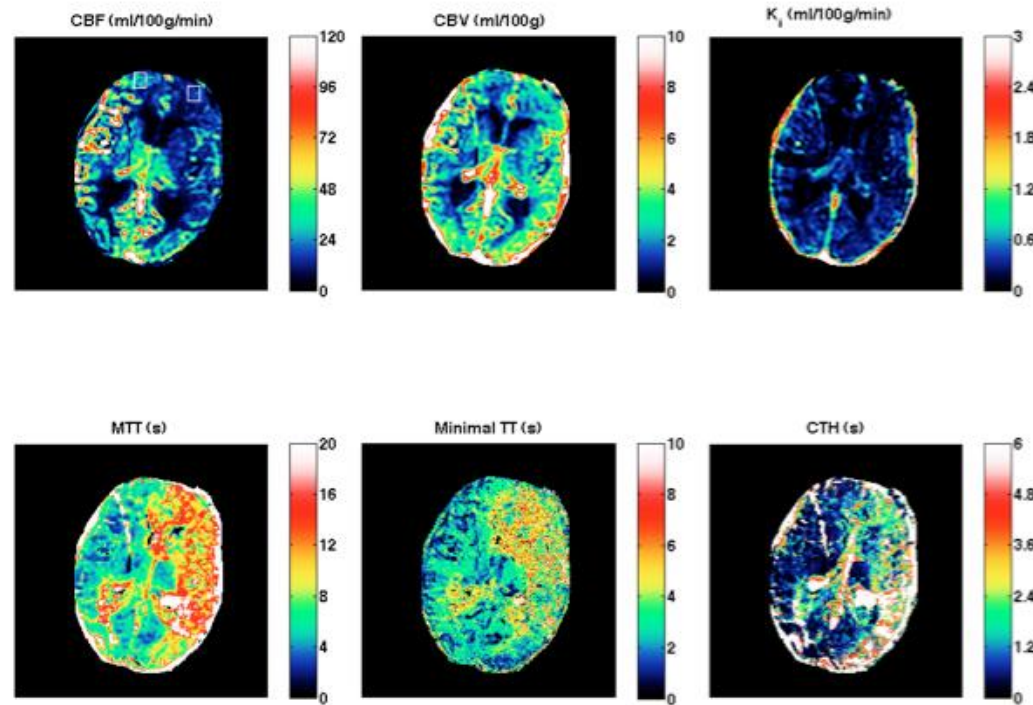


Figure 6. The figure shows results from a patient with left sided internal carotid stenosis, with multiple thrombo-embolic episodes. Perfusion (CBF) is decreased while the cerebral blood volume (CBV) is increased in the fronto-parietal region, but the permeability (K_i) seems relatively normal. The mean transit time (MTT), the minimal transit time (minimal TT) and capillary transit time heterogeneity (CTH) is prolonged in the entire region showing altered perfusion.

