



Modelización de la distribución del ^{18}F -FDG en el organismo.

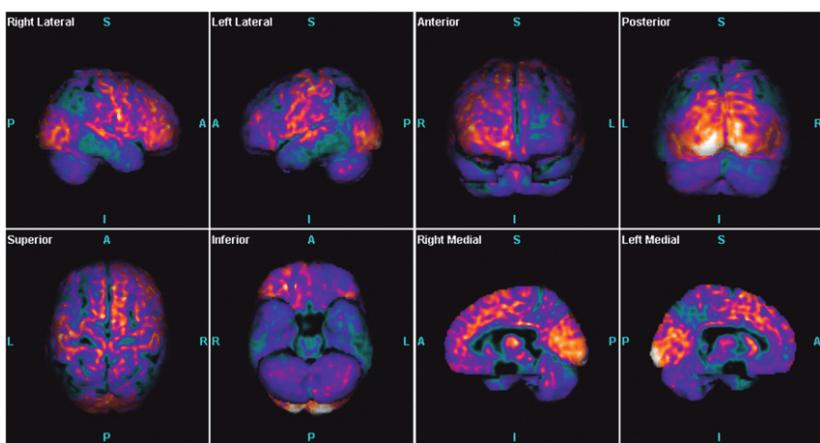
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Objetivo

En la medicina nuclear diagnóstica ocupa un papel importante la tomografía por emisión de positrones (PET). En este tipo de exploraciones el marcador más utilizado es la ^{18}F -fludesoxiglucosa (FDG), un radiofármaco que contiene al isótopo radiactivo flúor-18, con el que se generan imágenes que permiten el estudio del metabolismo celular de la glucosa.



Un problema fundamental es calcular las desintegraciones en los distintos órganos como consecuencia de la administración de FDG. Para ello se utiliza modelización compartimental que nos permite conocer la cantidad $x_i(t)$ retenida en un compartimento i . A partir de ello se calcula la dosis.

Para evaluar la dosis en los distintos órganos del paciente se emplea las tablas de la ICRP 128, que se basa en el modelo compartimental: Hays y Segall A mathematical model for the distribution of fluorodeoxyglucose in humans. J. Nucl. Med. 40, 1358-1356.(1999). EURADOS ha detectado discrepancias significativas con los valores de la ICRP 128.

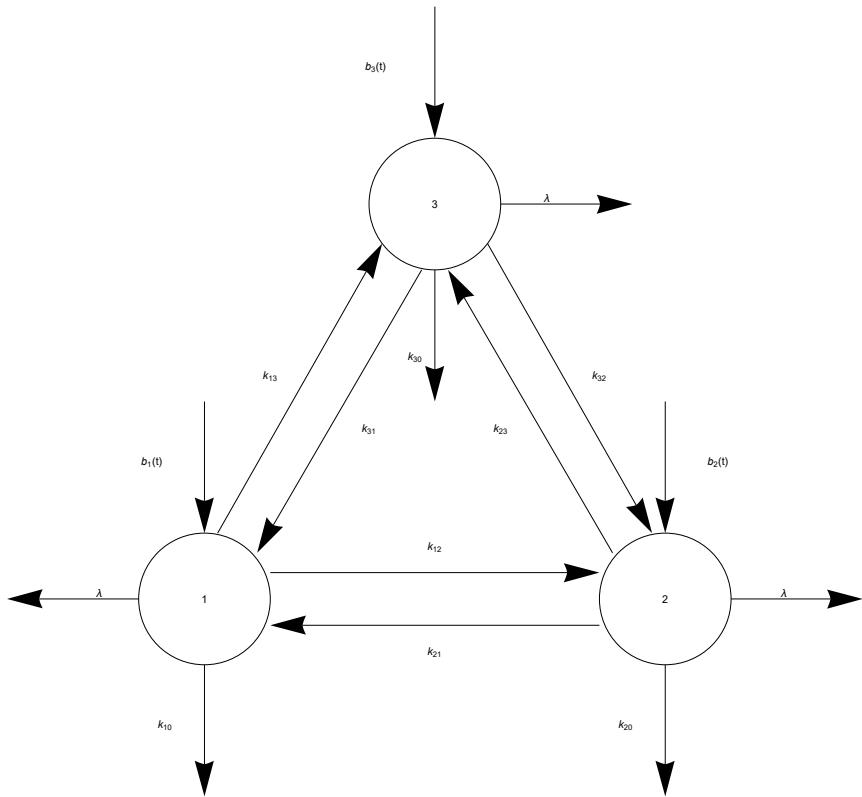
Nosotros hemos modelado independientemente la metabolización del FDG coincidiendo con los resultados de EURADOS. Nuestra aportación más importante es que hemos obtenido las soluciones analíticas del modelo, lo que facilita su aplicación para distintos fines. Además, el modelo desarrollado lo hemos integrado como parte de la aplicación web BIOKMOD, permitiendo que los usuarios modifiquen distintos parámetros.

Puede ejecutarse en:

<http://oed.usal.es/webMathematica/Biokmod/biokmod13fluor.jsp> o

<http://oed.usal.es/webMathematica/Biokmod/index.html>.

El modelo general compartmental



Si el caso anterior lo extendemos a n compartimentos, la ecuación que da el contenido en cualquier compartimento i esta dada por

$$\dot{\mathbf{x}}(t) = \mathbf{A} \mathbf{x} + \mathbf{b}(t), \quad t \geq 0$$

$$\mathbf{x}(0) = \mathbf{x}_0$$

donde:

$\mathbf{x}(t) = \{x_1(t), x_2(t), \dots, x_n(t)\}^T$ $x_i(t)$ denota la cantidad o contenido de un sustancia en el compartimento i en el momento t .

A: $n \times n$ es la matriz compartmental o matriz del sistema

b(t) = {b₁(t), b₂(t), ..., b_n(t)}^{T}} {b_i(t)} es la tasa de entrada hacia el compartimento i desde fuera del sistema.

x(0) = {x₁(0), x₂(0), ..., x_n(0)}^T son las condiciones iniciales que corresponden al contenido de un sustancia en el compartimento i en el momento $t = 0$.

Time integrated activity coefficients (TIAC)

The time - integrated activity is calculated as the area under the curve (AUC *) that describes the activity as a function of time in the source region after the administration of the radiopharmaceutical . The time integrated activity coefficient (TIAC) is defined as the time - integrated activity divided by the administered activity and is given in hours .

(*) The area under the curve or AUC, in the field of pharmacokinetics, is the definite integral of a curve that describes the variation of a drug concentration in blood plasma as a function of time

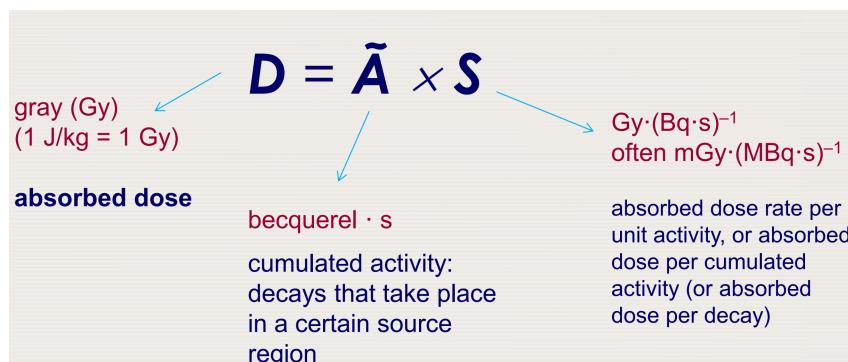
Symbol	Quantity	Unit
$\tilde{A}(r_S, T_D)$	Time-integrated activity	Bq · s
$\tilde{a}(r_S, T_D)$	Time-integrated activity coefficient	s
$D(r_T)$	Absorbed dose to the target region r_T	Gy
\dot{D}	Absorbed dose rate	Gy/s

18.1.1. Basic concepts

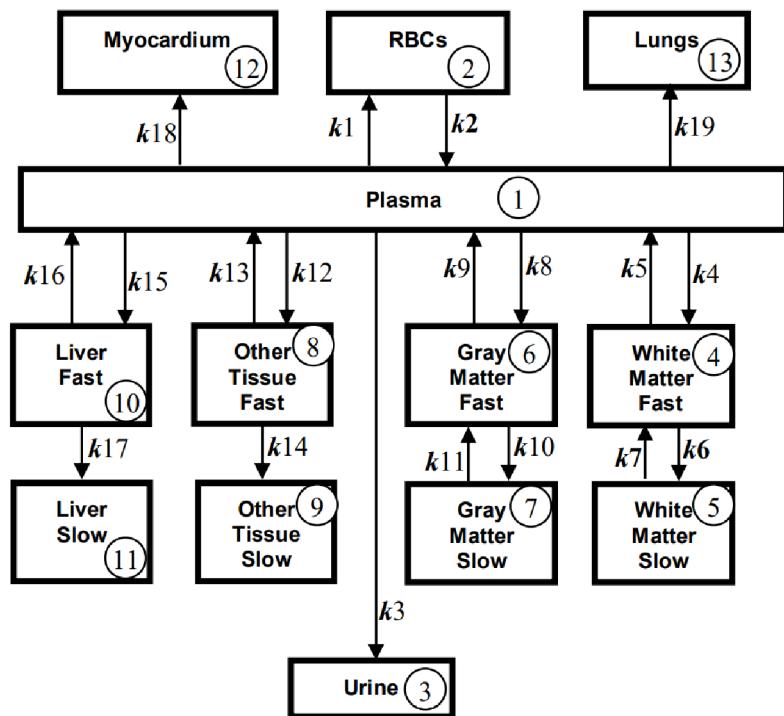
The time-integration period is commonly chosen from the time of administration of the radiopharmaceutical until infinite time. However, the integration period should be matched to the biological endpoint studied in combination with the time period in which the relevant absorbed dose is delivered (T_D).

$$\tilde{A}(r_S) = \int A(r_S, t) dt = \int_0^{T_D} A(r_S, t) dt = \tilde{a}(r_S) \cdot A_0$$

$\tilde{a}(r_S) = \frac{\tilde{A}(r_S)}{A_0}$ Is defined as the **time-integrated activity coefficient**, being A_0 the administered activity; it has the unit of time (e.g. s, or h). In the MIRD Primer it was named '**residence time**'



Modelo biocinético del F-18

Table 1: Transfer rates* of the compartmental model for ^{18}F -FDG by Hays and Segall.

From	Organ To	Parameter	Transfer rate (min^{-1}) Arithmetic mean	Transfer rate (min^{-1}) Geometric mean
Plasma	RBCs	k_1	4.8	4.07
RBCs	plasma	k_2	8.07	7.35
Plasma	Urine bladder	k_3	0.0088	0.0085
Plasma	WM Fast	k_4	0.054	0.052
WM Fast	Plasma	k_5	0.109	0.10
WM Fast	WM Slow	k_6	0.045	0.042
WM Slow	WM Fast	k_7	0.0058	0.0055
Plasma	GM Fast	k_8	0.102	0.099
GM Fast	GM Slow	k_9	0.13	0.115
GM Slow	GM Fast	k_{10}	0.062	0.059
Plasma	Other Fast	k_{11}	0.0068	0.0066
Plasma	Other Fast	k_{12}	0.371	0.348
Other Fast	Plasma	k_{13}	0.102	0.097
Other Fast	Other Slow	k_{14}	0.0167	0.015
Plasma	Liver Fast	k_{15}	0.068	0.038
Liver Fast	Plasma	k_{16}	0.219	0.186
Liver Fast	Liver Slow	k_{17}	0.018	0.006
Plasma	Myocardium	k_{18}	0.0053	0.003
Plasma	Lungs	k_{19}	0.0017	0.0016

The half-life of ^{18}F was taken from ICRP 107 (ICRP, 2008); $T_{1/2} = 109.77$ min, i.e. the physical decay constant is $\lambda = 0.0063 \text{ min}^{-1}$.

La resolución del modelo compartimental nos dará la función respuesta (contenido de F presente en un compartimento i, tras la inyección en $t=0$, en el compartimento 1 de una cantidad $q(0) = 1$; $r_t(t)$). Lo hemos calculado utilizando ambos tipos de tasas de trasferencia

El paquete

El modelo lo hemos incorporado en un paquete de Mathematica

```
In[1]:= Needs["Biokmod`OIRModels`"]
SysModel, version 2.0.b7 2020-11-30
OIRModels 1.1b2 2022-03-24

In[2]:= CompartmentalnumbersF
Out[2]= {{Plasma, 1}, {RBCs, 2}, {Urine, 3}, {WhiteMatterFast, 4}, {WhiteMatterSlow, 5},
{GrayMatterFast, 6}, {GrayMatterSlow, 7}, {OtherTissueFast, 8},
{OtherTissueSlow, 9}, {LiverFast, 10}, {LiverSlow, 11}, {Myocardium, 12}, {Lungs, 13}}
```

Características isotópicas

```
In[3]:= IsotopeDecayModes["Fluorine18"]
"BetaPlusDecay"
"\u03b2"+"
1. ` 3.
633.023` 6. keV
BetaPlusDecay
\beta^+
1.00
633.023 keV

In[4]:= fluorine-18 ISOTOPE ["Properties"]
Out[4]= {atomic mass, atomic number, atomic symbol, binding energy per nucleon, biological half-life,
biological lifetime, branching ratios, critical diameter, critical mass, critical organs,
daughter nuclides, decay constant, Q-value, decay modes, decay modes, decay products,
diagnostic applications, diagnostic applications to diseases, effective half-life, effective lifetime,
entity classes, entity type list, E_level, half-life \u03c4\u2081/\u2082, excited state lifetimes, excited state parities,
excited state spins, width \u0394, external exposure advisory, final decay products, full symbol,
half-life, half-value layer, isotope abundance, mean lifetime, magnetic moment, mass defect,
mass excess, mass number, diagnostic applications, memberships, molar mass, molar radioactivity,
name, neutron number, parity, electric quadrupole moment, type of particle, radioactivity,
specific activity, spin, spin parity J^\u03c0, stable, brief symbol, tenth-value layer, width}
```

```
In[5]:= fluorine-18 ISOTOPE ["HalfLife"]
```

```
Out[5]= 109.73 min
```

8 Funciones respuesta:

```
In[=]* qF18Injection[t] // Chop // Short
Out[=]//Short=
{x1[t] → 0.395683 e-13.1132 t + 0.430092 e-0.51053 t +
 <<4>> + 0.000262063 e-0.0105465 t + 0.0161431 e-0.00868373 t,
 x2[t] → <<1>>, <<9>>, <<1>>, x13[t] → <<15>> + <<21>> e<<1>>}
```



```
In[=]* qF18InjectionG[t] // Chop // Short
Out[=]//Short=
{x1[t] → 0.379079 e-11.6304 t + 0.454068 e-0.465429 t +
 <<4>> + 0.000254002 e-0.010303 t + 0.0183932 e-0.00835775 t,
 x2[t] → <<1>>, <<10>>, x13[t] → <<15>> + <<21>> e<<1>>}
```

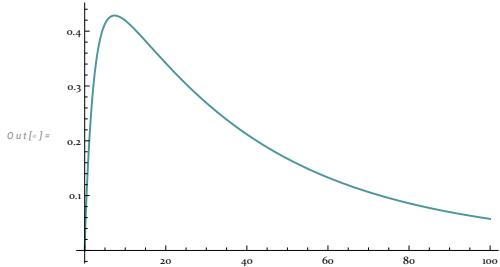
Incoporaciones de distinto tipo

A partir de la función respuesta pueden realizarse multitud calculos. Por ejemplo; El total, como suma en todos los compartimentos, o realizarse representaciones gráficas

```
In[1]:= Total[Table[xi[t], {i, 1, 13}] /. qF18Injection[t]] // Chop
Out[1]= 5.55112×10-17 e-13.1132 t + 3.21965×10-15 e-0.51053 t + 2.08167×10-16 e-0.230225 t +
1.38778×10-16 e-0.186303 t + 5.55112×10-17 e-0.156732 t + 1.43219×10-14 e-0.0294228 t +
2.62984×10-15 e-0.0105465 t + 6.24223×10-14 e-0.00868373 t + 1. e-0.00631449 t

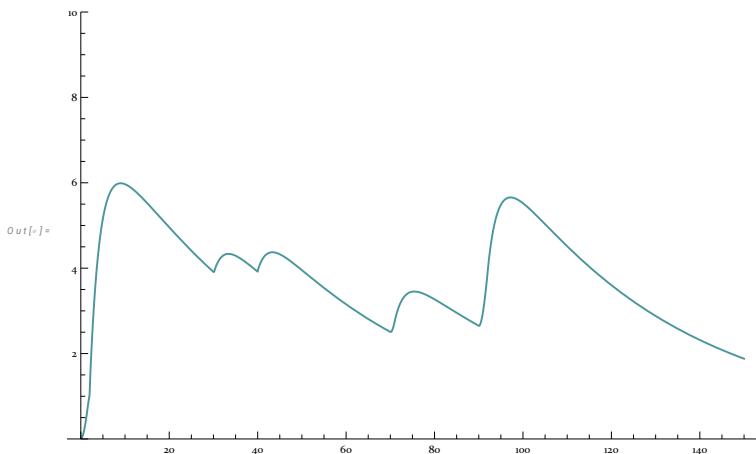
In[2]:= u[t_] = x8[t] /. qF18InjectionG[t] // ExpandAll // Chop
Out[2]= -0.0114592 e-11.6304 t - 0.455226 e-0.465429 t - 0.00851962 e-0.194332 t - 0.0589342 e-0.170587 t -
0.0411823 e-0.14548 t + 0.516291 e-0.0278065 t + 0.000818364 e-0.010303 t + 0.0582123 e-0.00835775 t
```

```
In[3]:= Plot[u[t], {t, 0, 100}]
```



Inputs multiples

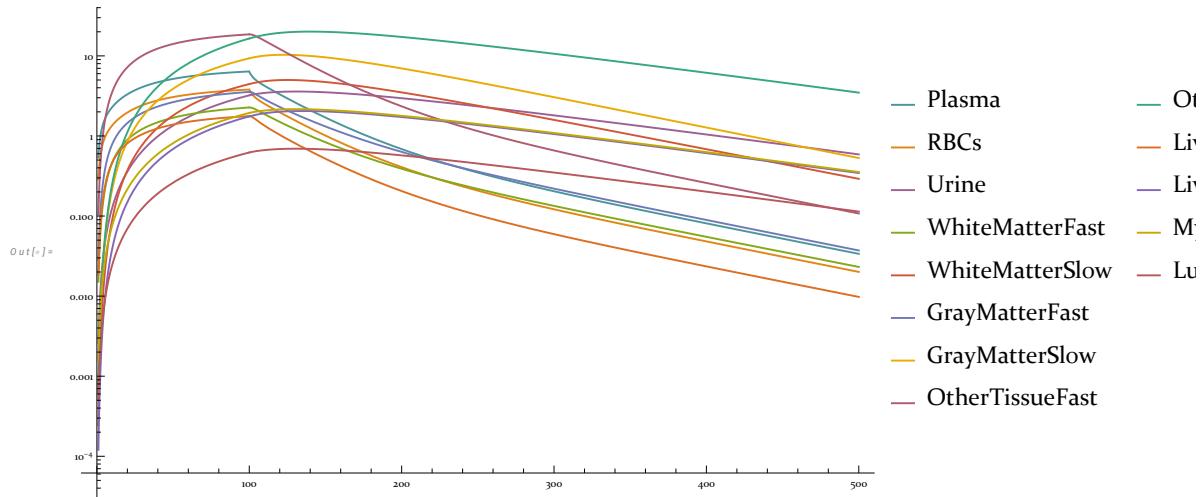
```
In[4]:= inputs = {{5, 0, 1.5}, {9, 2}, {2, 30, 0.3}, {2, 40}, {3, 70, 1}, {8, 90, 2}};
Plot[qMultiple[inputs, {u[t], t}, t1], {t1, 0, 150},
PlotRange → {0, 10}, ExclusionsStyle → {Blue, Blue}, ImageSize → Large]
```



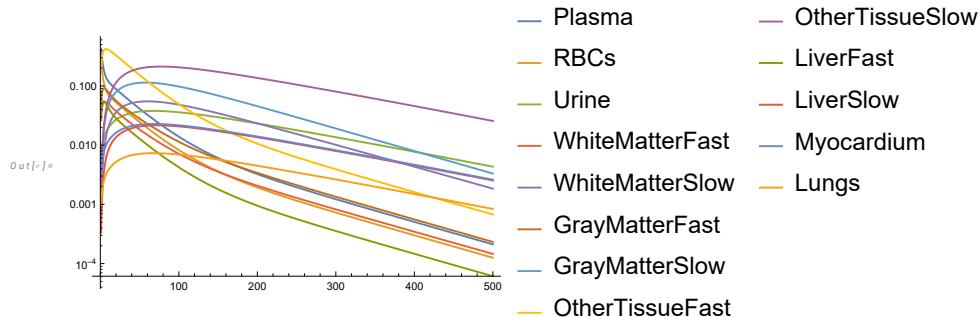
Incorporación constante.

```
In[+]:= compf = {"Plasma", "RBCs", "Urine", "WhiteMatterFast",
  "WhiteMatterSlow", "GrayMatterFast", "GrayMatterSlow",
  "OtherTissueFast", "OtherTissueSlow",
  "LiverFast", "LiverSlow", "Myocardium", "Lungs"};
```

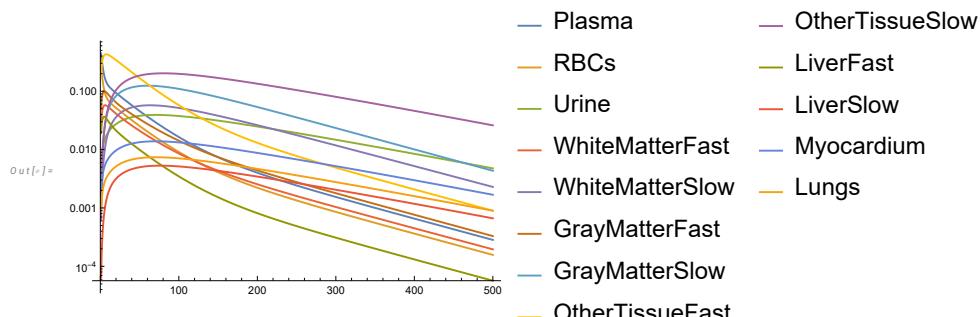
```
In[+]:= LogPlot[Evaluate[qF18InjectionCte[t, 100]],
 {t, 1, 500}, PlotRange -> All, PlotLegends -> compf]
```



```
In[+]:= ResponseReport["Fluorine", "Injection",
  "Acute", "ArithmeticMean", "GraphicsReport", {1, 500}]
```



```
In[+]:= ResponseReport["Fluorine", "Injection",
  "Acute", "GeometricMean", "GraphicsReport", {1, 500}]
```



Time integrated activity coefficients (TIAC)

```
In[+]:= ResponseReport["Fluorine", "Injection",
  "Acute", "ArithmeticMean", "ResponseFunction", t] // Chop // Short
```

$$\text{Out}[+]\text{//Short} = \left\{ x_1[t] \rightarrow 0.395683 e^{-13.1132 t} + 0.430092 e^{-0.51053 t} + \dots + 0.0161431 e^{-0.00868373 t}, \dots, x_{13}[t] \rightarrow \dots + \dots e^{\dots} \right\}$$

```
In[+]:= ResponseReport["Fluorine", "Injection",
  "Acute", "GeometricMean", "ResponseFunction", t] // Chop // Short
```

$$\text{Out}[+]\text{//Short} = \{ \dots \}$$

```
In[+]:= ResponseReport["Fluorine", "Injection",
  "Acute", "ArithmeticandGeometricMean", "TIAC", 1000]
```

Plasma	RBCs	Urine	WhiteMatterFast	WhiteMatterSlow	GrayMatterFast	GrayMatterSlow	OtherTi
0.123683	0.0735088	0.17182	0.0481276	0.178739	0.0759195	0.358863	0.36
0.123683	0.0735088	0.17182	0.0481276	0.178739	0.0759195	0.358863	0.36

Se han integrado en BIOKMOD

<http://oed.usal.es/webMathematica/Biokmod/biokmod13fluor.jsp>

O

[http://oed.usal.es/webMathematica/Biokmod/index.html.](http://oed.usal.es/webMathematica/Biokmod/index.html)

BIOKMOD.
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Sánchez G, López-Fidalgo J "Mathematical Techniques for Solving Analytically Large Compartmental Systems" Health Physics.. 85 (2): 2003. ISSN/ISBN: 0017-9078
López-Fidalgo J; Sánchez G; Statistical Criteria to Establish Bioassay Programs. Health Physics. 89 (4). 2005. ISSN/ISBN: 0017-9078."/>


Referencias

<https://diarium.usal.es/guillermo/biokmod/>