

# Package: rxode2parse (via r-universe)

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**Title** Parsing and Code Generation Functions for 'rxode2'

**Version** 2.0.19

**Maintainer** Matthew L. Fidler <matthew.fidler@gmail.com>

**Description** Provides the parsing needed for 'rxode2' (Wang, Hallow and James (2016) <[doi:10.1002/psp4.12052](https://doi.org/10.1002/psp4.12052)>). It also provides the 'stan' based advan linear compartment model solutions with gradients (Carpenter et al (2015), <[doi:10.48550/arXiv.1509.07164](https://doi.org/10.48550/arXiv.1509.07164)>) needed in 'nlmixr2' (Fidler et al (2019) <[doi:10.1002/psp4.12445](https://doi.org/10.1002/psp4.12445)>). This split will reduce computational burden of recompiling 'rxode2'.

**License** GPL (>= 3)

**URL** <https://nlmixr2.github.io/rxode2parse/>,  
<https://github.com/nlmixr2/rxode2parse/>

**BugReports** <https://github.com/nlmixr2/rxode2parse/issues/>

**Depends** R (>= 4.0.0)

**Imports** checkmate, crayon, dparser, knitr, methods, qs, Rcpp (>= 1.0.8), utils, digest, rex, symengine, data.table (>= 1.12.4)

**Suggests** testthat (>= 3.0.0), dplyr, nlmixr2data, devtools, rmarkdown

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**Author** Matthew L. Fidler [aut, cre]  
(<<https://orcid.org/0000-0001-8538-6691>>), Wenping Wang [aut],  
Richard Upton [ctb], Gabriel Staples [ctb], Goro Fuji [ctb],  
Morwenn [ctb], Igor Kushnir [ctb], Kevin Ushey [ctb], David  
Cooley [ctb]

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.toClassicEvid	<i>This converts NONMEM-style EVIDs to classic RxODE events</i>
----------------	---

---

## Description

This converts NONMEM-style EVIDs to classic RxODE events

## Usage

```
.toClassicEvid(cmt = 1L, amt = 0, rate = 0, dur = 0, ii = 0, evid = 0L, ss = 0)
```

## Arguments

cmt	compartment flag
amt	dose amount
rate	dose rate
dur	dose duration
ii	inter-dose interval
evid	event id
ss	steady state

**Value**

classic evids, excluding evids that are added (you need to add them manually) or simply use etTran. This is mostly for testing and really shouldn't be used directly.

**Author(s)**

Matthew L. Fidler

**Examples**

```
.toClassicEvid(cmt=10, amt=3, evid=1)
.toClassicEvid(cmt=10, amt=3, rate=2, evid=1)
.toClassicEvid(cmt=10, amt=3, rate=-1, evid=1)
.toClassicEvid(cmt=10, amt=3, rate=-2, evid=1)
.toClassicEvid(cmt=10, amt=3, dur=2, evid=1)
.toClassicEvid(cmt=304, amt=3, dur=2, evid=1)
.toClassicEvid(cmt=7, amt=0, rate=2, evid=1, ss=1)
.toClassicEvid(cmt=-10, amt=3, evid=1)
.toClassicEvid(cmt=10, amt=3, evid=5)
.toClassicEvid(cmt=6, amt=3, evid=6)
.toClassicEvid(cmt=6, amt=3, evid=7)
.toClassicEvid(evid=2)
.toClassicEvid(evid=4)
```

---

print.rxModelVars      *Print Values*

---

**Description**

print prints its argument and returns it *invisibly* (via `invisible(x)`). It is a generic function which means that new printing methods can be easily added for new `classes`.

**Usage**

```
## S3 method for class 'rxModelVars'
print(x, ...)
```

**Arguments**

x                    an object used to select a method.  
...                   further arguments passed to or from other methods.

## Details

The default method, `print.default` has its own help page. Use `methods("print")` to get all the methods for the `print` generic.

`print.factor` allows some customization and is used for printing `ordered` factors as well.

`print.table` for printing `tables` allows other customization. As of R 3.0.0, it only prints a description in case of a table with 0-extents (this can happen if a classifier has no valid data).

See `noquote` as an example of a class whose main purpose is a specific `print` method.

## Value

This returns invisibly the model variables object

## References

Chambers, J. M. and Hastie, T. J. (1992) *Statistical Models in S*. Wadsworth & Brooks/Cole.

## See Also

The default method `print.default`, and help for the methods above; further `options`, `noquote`.

For more customizable (but cumbersome) printing, see `cat`, `format` or also `write`. For a simple prototypical `print` method, see `.print.via.format` in package `tools`.

## Examples

```
require(stats)

ts(1:20) #-- print is the "Default function" --> print.ts(.) is called
for(i in 1:3) print(1:i)

## Printing of factors
attenu$station ## 117 levels -> 'max.levels' depending on width

## ordered factors: levels "l1 < l2 < .."
esoph$agegp[1:12]
esoph$alcgp[1:12]

## Printing of sparse (contingency) tables
set.seed(521)
t1 <- round(abs(rt(200, df = 1.8)))
t2 <- round(abs(rt(200, df = 1.4)))
table(t1, t2) # simple
print(table(t1, t2), zero.print = ".") # nicer to read

## same for non-integer "table":
T <- table(t2,t1)
T <- T * (1+round(rlnorm(length(T)))/4)
print(T, zero.print = ".") # quite nicer,
print.table(T[,2:8] * 1e9, digits=3, zero.print = ".")
## still slightly inferior to Matrix::Matrix(T) for larger T
```

```
## Corner cases with empty extents:
table(1, NA) # < table of extent 1 x 0 >
```

---

rxDerived	<i>Calculate derived parameters for the 1-, 2-, and 3- compartment linear models.</i>
-----------	---

---

### Description

This calculates the derived parameters based on what is provided in a data frame or arguments

### Usage

```
rxDerived(..., verbose = FALSE, digits = 0)
```

### Arguments

...	The input can be: <ul style="list-style-type: none"> <li>• A data frame with PK parameters in it; This should ideally be a data frame with one pk parameter per row since it will output a data frame with one PK parameter per row.</li> <li>• PK parameters as either a vector or a scalar</li> </ul>
verbose	boolean that when TRUE provides a message about the detected pk parameters and the detected compartmental model. By default this is FALSE.
digits	represents the number of significant digits for the output; If the number is zero or below (default), do not round.

### Value

Return a data.frame of derived PK parameters for a 1-, 2-, or 3-compartment linear model given provided clearances and volumes based on the inferred model type.

The model parameters that will be provided in the data frame are:

- vc: Central Volume (for 1-, 2- and 3- compartment models)
- ke1: First-order elimination rate (for 1-, 2-, and 3-compartment models)
- k12: First-order rate of transfer from central to first peripheral compartment; (for 2- and 3-compartment models)
- k21: First-order rate of transfer from first peripheral to central compartment, (for 2- and 3-compartment models)
- k13: First-order rate of transfer from central to second peripheral compartment; (3-compartment model)
- k31: First-order rate of transfer from second peripheral to central compartment (3-compartment model)
- vp: Peripheral Volume (for 2- and 3- compartment models)

- vp2: Peripheral Volume for 3rd compartment (3- compartment model)
- vss: Volume of distribution at steady state; (1-, 2-, and 3-compartment models)
- t12alpha:  $t_{1/2,\alpha}$ ; (1-, 2-, and 3-compartment models)
- t12beta:  $t_{1/2,\beta}$ ; (2- and 3-compartment models)
- t12gamma:  $t_{1/2,\gamma}$ ; (3-compartment model)
- alpha:  $\alpha$ ; (1-, 2-, and 3-compartment models)
- beta:  $\beta$ ; (2- and 3-compartment models)
- gamma:  $\beta$ ; (3-compartment model)
- A: true A; (1-, 2-, and 3-compartment models)
- B: true B; (2- and 3-compartment models)
- C: true C; (3-compartment model)
- fracA: fractional A; (1-, 2-, and 3-compartment models)
- fracB: fractional B; (2- and 3-compartment models)
- fracC: fractional C; (3-compartment model)

### Author(s)

Matthew Fidler and documentation from Justin Wilkins, <justin.wilkins@occams.com>

### References

Shafer S. L. CONVERT.XLS

Rowland M, Tozer TN. Clinical Pharmacokinetics and Pharmacodynamics: Concepts and Applications (4th). Clipping Williams & Wilkins, Philadelphia, 2010.

### Examples

```
## Note that rxode2 parses the names to figure out the best PK parameter
params <- rxDerived(c1 = 29.4, v = 23.4, vp = 114, vp2 = 4614, q = 270, q2 = 73)

## That is why this gives the same results as the value before
params <- rxDerived(CL = 29.4, V1 = 23.4, V2 = 114, V3 = 4614, Q2 = 270, Q3 = 73)

## You may also use micro-constants alpha/beta etc.
params <- rxDerived(k12 = 0.1, k21 = 0.2, k13 = 0.3, k31 = 0.4, kel = 10, v = 10)

## or you can mix vectors and scalars
params <- rxDerived(CL = 29.4, V = 1:3)

## If you want, you can round to a number of significant digits
## with the `digits` argument:
params <- rxDerived(CL = 29.4, V = 1:3, digits = 2)
```

---

rxode2parse	<i>Internal translation to get model variables list</i>
-------------	---

---

## Description

Internal translation to get model variables list

## Usage

```
rxode2parse(  
  model,  
  linear = FALSE,  
  linCmtSens = c("linCmtA", "linCmtB", "linCmtC"),  
  verbose = FALSE,  
  code = NULL,  
  envir = parent.frame()  
)
```

## Arguments

model	Model (either file name or string)
linear	boolean indicating if linear compartment model should be generated from <code>linCmt()</code> (default FALSE)
linCmtSens	Linear compartment model sensitivity type
verbose	is a boolean indicating the type of model detected with <code>linCmt()</code> parsing
code	is a file name where the c code is written to (for testing purposes mostly, it needs rxode2 to do anything fancy)
envir	is the environment to look for R user functions (defaults to parent environment)

## Value

A `rxModelVars` object that has the model variables of a rxode2 syntax expression

## Examples

```
rxode2parse("a=3")
```

rxode2parseAssignTranslation

*This assigns the c level linkages for a roxde2 model*

---

**Description**

This assigns the c level linkages for a roxde2 model

**Usage**

```
rxode2parseAssignTranslation(df)
```

**Arguments**

df                    data frame containing the character column names rxFun, fun, type, package, packageFun and the integer column names argMin and argMax

**Value**

Nothing called for side effects

**Author(s)**

Matthew L. Fidler

**Examples**

```
rxode2parseAssignTranslation(rxode2parseGetTranslation())
```

---

rxode2parseD

*This gives the derivative table for rxode2*

---

**Description**

This will help allow registration of functions in rxode2

**Usage**

```
rxode2parseD()
```



**Details**

This environment is a derivative table;

For example:

```
Derivative(f(a,b,c), a) = fa() Derivative(f(a,b,c), b) = fb() Derivative(f(a,b,c), c) = fc()
```

Then the derivative table for f would be:

```
assign("f", list(fa(a,b,c), fb(a,b,c), fc(a,b,c)), rxode2parseD())
```

fa translates the arguments to the derivative with respect to a fb translates the arguments to the derivative with respect to b

If any of the list is NULL then rxode2 won't know how to take a derivative with respect to the argument.

If the list is shorter than the length of the arguments then the argument then the derivative of arguments that are not specified cannot be taken.

**Value**

Derivative table environment for rxode2

**Author(s)**

Matthew L. Fidler

---

rxode2parseGetPackagesToLoad

*Control the packages that are loaded when a rxode2 model dll is loaded*

---

**Description**

Control the packages that are loaded when a rxode2 model dll is loaded

**Usage**

```
rxode2parseGetPackagesToLoad()
```

```
rxode2parseAssignPackagesToLoad(pkgs = rxode2parseGetPackagesToLoad())
```

**Arguments**

pkgs                    The packages to make sure are loaded every time you load an rxode2 model.

**Value**

List of packages to load

**Author(s)**

Matthew Fidler

**Examples**

```
rxode2parseGetPackagesToLoad()
```

```
rxode2parseAssignPackagesToLoad(rxode2parseGetPackagesToLoad())
```

---

```
rxode2parseGetPointerAssignment
```

*This function gets the currently assigned function pointer assignments*

---

**Description**

This function gets the currently assigned function pointer assignments

**Usage**

```
rxode2parseGetPointerAssignment()
```

**Value**

The currently assigned pointer assignments

**Author(s)**

Matthew L. Fidler

**Examples**

```
rxode2parseGetTranslation()
```

---

```
rxode2parseGetTranslation
```

*This function gets the currently assigned translations*

---

**Description**

This function gets the currently assigned translations

**Usage**

```
rxode2parseGetTranslation()
```

**Value**

The currently assigned translations

**Author(s)**

Matthew L. Fidler

**Examples**

```
rxode2parseGetTranslation()
```

---

<code>rxode2parseMd5</code>	<i>Get the MD5 hash of the current language revision</i>
-----------------------------	--

---

**Description**

Get the MD5 hash of the current language revision

**Usage**

```
rxode2parseMd5()
```

**Value**

md5 hash of language revision

**Author(s)**

Matthew L. Fidler

**Examples**

```
rxode2parseMd5()
```

rxParseSuppressMsg     *Respect suppress messages*

---

### **Description**

This turns on the silent REprintf in C when suppressMessages() is turned on. This makes the REprintf act like messages in R, they can be suppressed with suppressMessages()

### **Usage**

```
rxParseSuppressMsg()
```

### **Value**

Nothing

### **Author(s)**

Matthew Fidler

### **Examples**

```
# rxParseSuppressMsg() is called with rxode2()

# Note the errors are output to the console

try(rxode2parse("d/dt(matt)=/3"), silent = TRUE)

# When using suppressMessages, the output is suppressed

suppressMessages(try(rxode2parse("d/dt(matt)=/3"), silent = TRUE))

# In rxode2, we use REprintf so that interrupted threads do not crash R
# if there is a user interrupt. This isn't captured by R's messages, but
# This interface allows the `suppressMessages()` to suppress the C printing
# as well

# If you want to suppress messages from rxode2 in other packages, you can use
# this function
```

---

rxSetIni0	<i>Set Initial conditions to time zero instead of the first observed/dosed time</i>
-----------	---

---

**Description**

Set Initial conditions to time zero instead of the first observed/dosed time

**Usage**

```
rxSetIni0(ini0 = TRUE)
```

**Arguments**

ini0	When TRUE (default), set initial conditions to time zero. Otherwise the initial conditions are the first observed time.
------	---

**Value**

the boolean ini0, though this is called for its side effects

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