

# Package ‘sysid’

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**Description** The sysid package provides functions for constructing mathematical models of dynamic systems from measured input-output data. The package contains functions for data visualization, data preprocessing, parametric and non-parametric model estimation, and model predictions and validation.

**License** GPL-3

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armax	<i>Estimate ARMAX Models</i>
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## Description

Fit an ARMAX model of the specified order given the input-output data

## Usage

```
armax(x, order = c(0, 1, 1, 0), options = optimOptions())
```

## Arguments

x	an object of class idframe
options	Estimation Options, setup using <a href="#">optimOptions</a>
order:	Specification of the orders: the four integer components (na,nb,nc,nk) are the order of polynomial A, order of polynomial B + 1, order of the polynomial C, and the input-output delay respectively

## Details

SISO ARMAX models are of the form

$$y[k] + a_1 y[k-1] + \dots + a_n y[k-na] = b_{nk} u[k-nk] + \dots + b_{nk+nb} u[k-nk-nb] + c_1 e[k-1] + \dots + c_{nc} e[k-nc] + e[k]$$

The function estimates the coefficients using non-linear least squares (Levenberg-Marquardt Algorithm) \ The data is expected to have no offsets or trends. They can be removed using the [detrend](#) function.

## Value

An object of class estpoly containing the following elements:

sys	an idpoly object containing the fitted ARMAX coefficients
fitted.values	the predicted response
residuals	the residuals
input	the input data used
call	the matched call

stats	A list containing the following fields: vcov - the covariance matrix of the fitted coefficients sigma - the standard deviation of the innovations
options	Option set used for estimation. If no custom options were configured, this is a set of default options
termination	Termination conditions for the iterative search used for prediction error minimization: WhyStop - Reason for termination iter - Number of Iterations iter - Number of Function Evaluations

## References

Arun K. Tangirala (2015), *Principles of System Identification: Theory and Practice*, CRC Press, Boca Raton. Sections 14.4.1, 21.6.2

## Examples

```
data(armaxsim)
z <- dataSlice(data,end=1533) # training set
mod_armax <- armax(z,c(1,2,1,2))
mod_armax
```

---

arx

---

Estimate ARX Models

---

## Description

Fit an ARX model of the specified order given the input-output data

## Usage

```
arx(x, order = c(0, 1, 0))
```

## Arguments

x	an object of class idframe
order:	Specification of the orders: the three integer components (na,nb,nk) are the order of polynolnomial A, (order of polynomial B + 1) and the input-output delay

## Details

SISO ARX models are of the form

$$y[k] + a_1y[k-1] + \dots + a_nay[k-na] = b_nku[k-nk] + \dots + b_{nk+nb}u[k-nk-nb] + e[k]$$

The function estimates the coefficients using linear least squares (with no regularization). Future versions may include regularization parameters as well \ The data is expected to have no offsets or trends. They can be removed using the [detrnd](#) function.

**Value**

An object of class `estpoly` containing the following elements:

<code>sys</code>	an <code>idpoly</code> object containing the fitted ARX coefficients
<code>fitted.values</code>	the predicted response
<code>residuals</code>	the residuals
<code>input</code>	the input data used
<code>call</code>	the matched call
<code>stats</code>	A list containing the following fields: <code>vcov</code> - the covariance matrix of the fitted coefficients <code>sigma</code> - the standard deviation of the innovations <code>df</code> - the residual degrees of freedom

**References**

Arun K. Tangirala (2015), *Principles of System Identification: Theory and Practice*, CRC Press, Boca Raton. Section 21.6.1

Lennart Ljung (1999), *System Identification: Theory for the User*, 2nd Edition, Prentice Hall, New York. Section 10.1

**Examples**

```
data(arxsim)
model <- arx(data,c(2,1,1))
model
plot(model) # plot the predicted and actual responses
```

---

`compare`

*Compare the measured output and the predicted output(s)*

---

**Description**

Plots the output predictions of model(s) superimposed over validation data, `data`, for comparison.

**Usage**

```
compare(data, nahead = 1, ...)
```

**Arguments**

<code>data</code>	validation data in the form of an <code>idframe</code> object
<code>nahead</code>	number of steps ahead at which to predict (Default:1). For infinite- step ahead predictions, supply <code>Inf</code> .
<code>...</code>	models whose predictions are to be compared

**See Also**

[predict.estpoly](#) for obtaining model predictions

**Examples**

```
data(arxsim)
compare(data,nahead=1,mod1,mod2,mod3)
```

dataSlice

*Subset or Resample idframe data***Description**

dataSlice is a subsetting method for objects of class idframe. It extracts the subset of the object data observed between indices start and end. If a frequency is specified, the series is then re-sampled at the new frequency.

**Usage**

```
dataSlice(data, start = NULL, end = NULL, freq = NULL)
```

**Arguments**

data	an object of class idframe
start	the start index
end	the end index
freq	fraction of the original frequency at which the series to be sampled.

**Details**

The dataSlice function extends the [window](#) function for idframe objects

**Value**

an idframe object

**See Also**

[window](#)

**Examples**

```
data(cstr)
cstrsub <- dataSlice(cstr,start=200,end=400) # extract between indices 200 and 400
cstrTrain <- dataSlice(cstr,end=4500) # extract upto index 4500
cstrTest <- dataSlice(cstr,start=6501) # extract from index 6501 till the end
cstr_new <- dataSlice(cstr,freq=0.5) # resample data at half the original frequency
```

---

detrend	<i>Remove offsets and linear trends</i>
---------	---

---

### Description

Removes the offsets or linear trends in each of the input and output matrices.

### Usage

```
detrend(x, type = c("constant", "linear")[1])
```

### Arguments

x	an object of class idframe
type	trend type - "constant" or "linear". (Default: "constant")

### Value

A list containing the following elements

fitted.values	idframe object with detrended variables
output_trend	list containing trend fits for each output variable
input_trend	list containing trend fits for each input variable

### See Also

[predict.detrend, lm](#)

### Examples

```
data(cstr)
fit <- detrend(cstr,type="linear") # remove linear trends
Zdetrend <- predict(fit) # get the detrended data

demean <- detrend(cstr) # remove offsets
Zcent <- predict(demean) # get the centered data
```

---

etfe	<i>Estimate empirical transfer function</i>
------	---

---

### Description

Estimates the emperical transfer function from the data by taking the ratio of the fourier transforms of the output and the input variables

### Usage

```
etfe(data)
```

**Arguments**

data                      an object of class idframe

**Value**

an idfrd object containing the estimated frequency response

**References**

Arun K. Tangirala (2015), *Principles of System Identification: Theory and Practice*, CRC Press, Boca Raton. Sections 5.3 and 20.4.2

**See Also**

[fft](#)

**Examples**

```
data(frf)
frf <- etfe(data)
```

---

fitch	<i>Fit Characteristics</i>
-------	----------------------------

---

**Description**

Returns quantitative assessment of the estimated model as a list

**Usage**

```
fitch(x)
```

**Arguments**

x                      the estimated model

**Value**

A list containing the following elements

MSE	Mean Square Error measure of how well the response of the model fits the estimation data
FPE	Final Prediction Error
FitPer	Normalized root mean squared error (NRMSE) measure of how well the response of the model fits the estimation data, expressed as a percentage.
AIC	Raw Akaike Information Criteria (AIC) measure of model quality
AICc	Small sample-size corrected AIC
nAIC	Normalized AIC
BIC	Bayesian Information Criteria (BIC)

---

getcov	<i>Parameter covariance of the identified model</i>
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---

### Description

Obtain the parameter covariance matrix of the linear, identified parametric model

### Usage

```
getcov(sys)
```

### Arguments

sys	a linear, identified parametric model
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---

idframe	<i>S3 class for storing input-output data.</i>
---------	--

---

### Description

idframe is an S3 class for storing and manipulating input-output data. It supports discrete time and frequency domain data.

### Usage

```
idframe(output = NULL, input = NULL, Ts = 1, start = 0, end = NULL,
        unit = c("seconds", "minutes", "hours", "days")[1])
```

### Arguments

output	dataframe/matrix/vector containing the outputs
input	dataframe/matrix/vector containing the inputs
Ts	sampling interval (Default: 1)
start	Time of the first observation
end	Time of the last observation Optional Argument
unit	Time Unit (Default: "seconds")

### Value

an idframe object

### See Also

[plot.idframe](#), the plot method for idframe objects, [summary.idframe](#), the summary method for idframe objects

### Examples

```
dataMatrix <- matrix(rnorm(1000),ncol=5)
data <- idframe(output=dataMatrix[,3:5],input=dataMatrix[,1:2],Ts=1)
```



---

idfrd	<i>S3 class for storing frequency response data</i>
-------	---

---

**Description**

S3 class for storing frequency response data

**Usage**

```
idfrd(response, freq, Ts)
```

**Arguments**

response	complex vector/matrix containing the response
freq	the frequencies at which the response is observed/estimated
Ts	sampling time of data

**Value**

an idfrd object

**Note**

The class can currently store only SISO Responses. Future versions will have support for multivariate data

**See Also**

[plot.idfrd](#) for generating bode plots; [spa](#) and [etfe](#) for estimating the frequency response given input/output data

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idinput	<i>function to generate input singals (rgs/rbs/prbs/sine)</i>
---------	---

---

**Description**

idinput is a function for generating input signals (rgs/rbs/prbs/sine) for identification purposes

**Usage**

```
idinput(n, type = "rgs", band = c(0, 1), levels = c(-1, 1))
```

**Arguments**

n	integer length of the input signal to be generated
type	the type of input signal to be generated. 'rgs' - generates random gaussian signal 'rbs' - generates random binary signal 'prbs' - generates pseudorandom binary signal 'sine' - generates a signal that is a sum of sinusoids Default value is type='rgs'
band	determines the frequency content of the signal. For type='rbs'/'sine', band = [wlow,whigh] which specifies the lower and the upper bound of the passband frequencies(expressed as fractions of Nyquist frequency). Default is c(0,1) For type='prbs', band=[0,B] where B is such that the signal is constant over 1/B (clock period). Default is c(0,1)
levels	row vector defining the input level. It is of the form levels=c(minu, maxu) For 'rbs','prbs', 'sine', the generated signal always between minu and maxu. For 'rgs', minu=mean value of signal minus one standard deviation and maxu=mean value of signal plus one standard deviation Default value is levels=c(-1,1)

idpoly

*Polynomial model with identifiable parameters***Description**

Creates a polynomial model with identifiable coefficients

**Usage**

```
idpoly(A = 1, B = 1, C = 1, D = 1, F1 = 1, ioDelay = 0, Ts = 1)
```

**Arguments**

A	Autoregressive coefficients
B, F1	Coefficients of the numerator and denominator respectively of the deterministic model between the input and output
C, D	Coefficients of the numerator and denominator respectively of the stochastic model
ioDelay	the delay in the input-output channel
Ts	sampling interval

**Details**

Discrete-time polynomials are of the form

$$A(q^{-1})y[k] = \frac{B(q^{-1})}{F1(q^{-1})}u[k] + \frac{C(q^{-1})}{D(q^{-1})}e[k]$$

**Examples**

```
# define output-error model
mod_oe <- idpoly(B=c(0.6,-0.2),F1=c(1,-0.5),ioDelay = 2,Ts=0.1)

# define box-jenkins model
B <- c(0.6,-0.2)
C <- c(1,-0.3)
D <- c(1,1.5,0.7)
F1 <- c(1,-0.5)
mod_bj <- idpoly(1,B,C,D,F1,ioDelay=1)
```

---

impulseest	<i>Estimate Impulse Response Coefficients</i>
------------	---

---

**Description**

impulseest is used to estimate impulse response coefficients from the data

**Usage**

```
impulseest(x, M = 30, K = NULL, regul = F, lambda = 1)
```

**Arguments**

x	an object of class idframe
M	Order of the FIR Model (Default:30)
K	Transport delay in the estimated impulse response (Default:NULL)
regul	Parameter indicating whether regularization should be used. (Default:FALSE)
lambda	The value of the regularization parameter. Valid only if regul=TRUE. (Default:1)

**Details**

The IR Coefficients are estimated using linear least squares. Future Versions will provide support for multivariate data.

**References**

Arun K. Tangirala (2015), *Principles of System Identification: Theory and Practice*, CRC Press, Boca Raton. Sections 17.4.11 and 20.2

**See Also**

[step](#)

**Examples**

```
uk <- rnorm(1000,1)
yk <- filter (uk,c(0.9,-0.4),method="recursive") + rnorm(1000,1)
data <- idframe(output=data.frame(yk),input=data.frame(uk))
fit <- impulseest(data)
plot(fit)
```

---

misdata	<i>Replace Missing Data by Interpolation</i>
---------	--

---

**Description**

Function for replacing missing values with interpolated ones. This is an extension of the `na.approx` function from the `zoo` package. The missing data is indicated using the value `NA`.

**Usage**

```
misdata(data)
```

**Arguments**

data	an object of class <code>idframe</code>
------	---

**Value**

data (an `idframe` object) with missing data replaced.

**See Also**

[na.approx](#)

**Examples**

```
data(cstr_mis)
summary(cstr_mis) # finding out the number of NAs
cstr <- misdata(cstr_mis)
```

---

oe	<i>Estimate Output-Error Models</i>
----	-------------------------------------

---

**Description**

Fit an output-error model of the specified order given the input-output data

**Usage**

```
oe(x, order = c(1, 1, 0), options = optimOptions())
```

**Arguments**

x	an object of class <code>idframe</code>
order	Specification of the orders: the four integer components (nb,nf,nk) are order of polynomial B + 1, order of the polynomial F, and the input-output delay respectively
options	Estimation Options, setup using <a href="#">optimOptions</a>

## Details

SISO OE models are of the form

$$y[k] + f_1 y[k-1] + \dots + f_n y[k-n] = b_n u[k-n] + \dots + b_{n+b} u[k-n-b] + f_1 e[k-1] + \dots + f_n e[k-n] + e[k]$$

The function estimates the coefficients using non-linear least squares (Levenberg-Marquardt Algorithm) \ The data is expected to have no offsets or trends. They can be removed using the [detrrend](#) function.

## Value

An object of class `estpoly` containing the following elements:

<code>sys</code>	an <code>idpoly</code> object containing the fitted OE coefficients
<code>fitted.values</code>	the predicted response
<code>residuals</code>	the residuals
<code>input</code>	the input data used
<code>call</code>	the matched call
<code>stats</code>	A list containing the following fields: <code>vcov</code> - the covariance matrix of the fitted coefficients <code>sigma</code> - the standard deviation of the innovations
<code>options</code>	Option set used for estimation. If no custom options were configured, this is a set of default options
<code>termination</code>	Termination conditions for the iterative search used for prediction error minimization: <code>WhyStop</code> - Reason for termination <code>iter</code> - Number of Iterations <code>iter</code> - Number of Function Evaluations

## References

Arun K. Tangirala (2015), *Principles of System Identification: Theory and Practice*, CRC Press, Boca Raton. Sections 14.4.1, 17.5.2, 21.6.3

## Examples

```
data(oesim)
z <- dataSlice(data,end=1533) # training set
mod_oe <- oe(z,c(2,1,2))
mod_oe
plot(mod_oe) # plot the predicted and actual responses
```

---

optimOptions

Create optimization options

---

## Description

Specify optimization options that are to be passed to the numerical estimation routines

**Usage**

```
optimOptions(tol = 1e-05, maxIter = 20, LMinit = 100, LMstep = 8)
```

**Arguments**

tol	Minimum ratio of the improvement to the current loss function. Iterations stop if this ratio goes below the tolerance limit (Default: 1e-5)
maxIter	Maximum number of iterations to be performed
LMinit	Starting value of search-direction length in the Levenberg-Marquardt method.
LMstep	Size of the Levenberg-Marquardt step

---

plot.idframe	<i>Plotting idframe objects</i>
--------------	---------------------------------

---

**Description**

Plotting method for objects inheriting from class idframe

**Usage**

```
## S3 method for class 'idframe'
plot(x, par = list(mar = c(3, 4, 2, 2)),
     col = "steelblue", ...)
```

**Arguments**

x	an object of class idframe
par	a list of arguments passed to par() before plotting.
col	line color, to be passed to plot.(Default="steelblue")
...	additional arguments to be passed to the tfplot function

**See Also**

[tfplot](#)

**Examples**

```
data(cstr)
plot(cstr,col="blue")
```

---

plot.idfrd	<i>Plotting idfrd objects</i>
------------	-------------------------------

---

**Description**

Generates the bode plot of the given frequency response data. It uses the ggplot2 plotting engine

**Usage**

```
## S3 method for class 'idfrd'  
plot(x)
```

**Arguments**

x	An object of class idframe
---	----------------------------

**See Also**

[ggplot](#)

**Examples**

```
data(frf)  
frf <- spa(data) # Estimates the frequency response from data  
plot(frf)
```

---

plot.impulseest	<i>Impulse Response Plots</i>
-----------------	-------------------------------

---

**Description**

Plots the estimated IR coefficients along with the significance limits at each lag.

**Usage**

```
## S3 method for class 'impulseest'  
plot(model, sig = 0.975)
```

**Arguments**

model	an object of class impulseest
sig	Significance Limits (Default: 0.975)

**See Also**

[impulseest,step](#)

---

predict.detrend	<i>Detrend data based on linear trend fits</i>
-----------------	--

---

**Description**

Returns detrended idframe object based on linear trend fit

**Usage**

```
## S3 method for class 'detrend'
predict(model, newdata = NULL, ...)
```

**Arguments**

model	an object of class detrend
newdata	An optional idframe object in which to look for variables with which to predict. If omitted, the original detrended idframe object is used

**Value**

an idframe object

**Examples**

```
data(cstr)
train <- dataSlice(cstr,end=5000)
test <- dataSlice(cstr,start=6001)
fit <- detrend(train)
Ztrain <- predict(fit)
Ztest <- predict(fit,test)
```

---

predict.estpoly	<i>Predictions of identified model</i>
-----------------	--

---

**Description**

Predicts the output of an identified model (estpoly) object K steps ahead.

**Usage**

```
## S3 method for class 'estpoly'
predict(x, newdata = NULL, nahead = 1)
```

**Arguments**

x	estpoly object containing the identified model
newdata	optional dataset to be used for predictions. If not supplied, predictions are made on the training set.
nahead	number of steps ahead at which to predict (Default:1). For infinite- step ahead predictions or pure simulation, supply Inf.



**Value**

Time-series containing the predictions

**References**

Arun K. Tangirala (2015), *Principles of System Identification: Theory and Practice*, CRC Press, Boca Raton. Chapter 18

**Examples**

```
data(arxsim)
Yhat <- predict(mod1,data) # 1-step ahead predictions
Yhat_2 <- predict(mod1,data,nahead=2) # 2-step ahead predictions
Yhat_inf <- predict(mod1,data,nahead=Inf) # Infinite-step ahead predictions
```

---

read.idframe	<i>Data input into a idframe object</i>
--------------	---

---

**Description**

Read the contents of a data.frame/matrix into a idframe object.

**Usage**

```
read.idframe(data, ninputs = NULL, Ts = 1, unit = c("seconds", "minutes",
"hours", "days")[1])
```

**Arguments**

data	a data.frame object
ninputs	the number of input columns. (Default: 0)
Ts	sampling interval (Default: 1)
unit	Time Unit (Default: "seconds")

**Value**

an idframe object

**Examples**

```
data(cstr)
data <- read.idframe(cstrData,ninputs=1,Ts= 1,unit="minutes")
```

---

read.odf.idframe	<i>Reading from .odf documents</i>
------------------	------------------------------------

---

### Description

Read the contents of an a .odf document into a idframe object.

### Usage

```
read.odf.idframe(file, sheetName, header = TRUE, ninputs = 0, Ts = 1,
  unit = c("seconds", "minutes", "hours", "days")[1], ...)
```

### Arguments

file	the path to the file to read
sheetName	a character string with the sheet name
header	a logical value indicating whether the first row corresponding to the first element of the rowIndex vector contains the names of the variables.
ninputs	the number of input columns. (Default: 0)
Ts	sampling interval (Default: 1)
unit	Time Unit (Default: "seconds")
...	additional arguments to be passed to the <a href="#">read.xlsx2</a> function

### Details

The read.odf.idframe function uses the [read.gnumeric.sheet](#) function, provided by the **xlsx** package, to read data from a .odf file and then calls the [read.idframe](#) function to read the data into a idframe object

### Value

an idframe object

### See Also

[read.xlsx2](#)

---

read.table.idframe	<i>Read the contents of a table-formatted file</i>
--------------------	--

---

### Description

Read the contents of an file in table format into a idframe object.

### Usage

```
read.table.idframe(file, header = TRUE, sep = ",", ninputs = 0, Ts = 1,
  unit = c("seconds", "minutes", "hours", "days")[1], ...)
```

**Arguments**

file	the path to the file to read
header	a logical value indicating whether the first row corresponding to the first element of the rowIndex vector contains the names of the variables. (Default: TRUE)
sep	the field separator character. Values on each line of the file are separated by this character. (Default: ",")
ninputs	the number of input columns. (Default: 0)
Ts	sampling interval (Default: 1)
unit	Time Unit (Default: "seconds")
...	additional arguments to be passed to the <a href="#">read.table</a> function

**Details**

The `read.table.idframe` function uses the [read.table](#) function, provided by the **utils** package, to read data from a table-formatted file and then calls the [read.idframe](#) function to read the data into a `idframe` object

**Value**

an `idframe` object

**See Also**

[read.table](#)

**Examples**

```
dataMatrix <- data.frame(matrix(rnorm(1000),ncol=5))
colnames(dataMatrix) <- c("u1","u2","y1","y2","y3")
write.csv(dataMatrix,file="test.csv",row.names=FALSE)

data <- read.table.idframe("test.csv",ninputs=2,unit="minutes")
```

---

read.xls.idframe

*Read the contents of a worksheet into a idframe object*


---

**Description**

Read the contents of an excel worksheet into a `idframe` object.

**Usage**

```
read.xls.idframe(file, sheetName, header = TRUE, ninputs = 0, Ts = 1,
  unit = c("seconds", "minutes", "hours", "days")[1], ...)
```

**Arguments**

file	the path to the file to read
sheetName	a character string with the sheet name
header	a logical value indicating whether the first row corresponding to the first element of the rowIndex vector contains the names of the variables.
ninputs	the number of input columns. (Default: 0)
Ts	sampling interval (Default: 1)
unit	Time Unit (Default: "seconds")
...	additional arguments to be passed to the <a href="#">read.xlsx2</a> function

**Details**

The `read.xlsx.idframe` function uses the [read.xlsx2](#) function, provided by the **xlsx** package, to read data from an excel file and then calls the [read.idframe](#) function to read the data into a `idframe` object

The function requires the java runtime to be installed on the system (Requirement of the **xlsx** package).

**Value**

an `idframe` object

**See Also**

[read.xlsx2](#)

**Examples**

```
library(xlsx)
dataMatrix <- data.frame(matrix(rnorm(1000),ncol=5))
colnames(dataMatrix) <- c("u1","u2","y1","y2","y3")
write.xlsx2(dataMatrix,file="test.xlsx",row.names=FALSE)

data <- read.xls.idframe("test.xlsx","Sheet1",ninputs=2,unit="minutes")
```

---

sim

*Simulate dynamic system*


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**Description**

Simulate the response of a system given the input

**Usage**

```
sim(model, input, sigma = 0, seed = NULL)
```

**Arguments**

model	the system model to simulate
input	a vector/matrix containing the input
sigma	standard deviation of the innovations (Default= 0)
seed	integer indicating the seed value of the random number generator

**Details**

The routine is currently built only for SISO systems. Future Versions will include support for MIMO systems. Current support

**Value**

a vector containing the output

**See Also**

[sim.idpoly](#) for simulating polynomial models

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sim.idpoly

*Simulate from a Polynomial Model*


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**Description**

Simulate the response of a system governed by a polynomial model , given the input

**Usage**

```
## S3 method for class 'idpoly'
sim(model, input, sigma = 0, seed = NULL)
```

**Arguments**

model	an object of class idpoly containing the coefficients
input	a vector/matrix containing the input
sigma	standard deviation of the innovations (Default= 0)
seed	integer indicating the seed value of the random number generator

**Details**

The routine is currently built only for SISO systems. Future Versions will include support for MIMO systems

**Value**

a vector containing the output

**See Also**

[idpoly](#) for defining polynomial models

## Examples

```
# ARX Model
u <- rnorm(200,sd=1)
model <- idpoly(A=c(1,-1.5,0.7),B=c(0.8,-0.25),ioDelay=1)
y <- sim(model,u,sigma=0.1)
```

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spa

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*Estimate frequency response*


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## Description

Estimates Frequency Response with fixed frequency resolution using spectral analysis

## Usage

```
spa(data, npad = 255)
```

## Arguments

data	an idframe object
npad	an integer representing the total length of each time series to analyze after padding with zeros. This argument allows the user to control the spectral resolution of the SDF estimates: the normalized frequency interval is $\text{deltaf}=1/\text{npad}$ . (Default: 255)

## Details

The function calls the SDF function in the sapa package to compute the cross-spectral densities. The method used is **Welch's Overlapped Segment Averaging** with a normalized **Hanning** window. The overlap used is 50

## Value

an idfrd object containing the estimated frequency response

## References

Arun K. Tangirala (2015), *Principles of System Identification: Theory and Practice*, CRC Press, Boca Raton. Sections 16.5 and 20.4

## See Also

[SDF](#)

## Examples

```
data(frf)
frf <- spa(data)
```

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`step`*Step Response Plots*

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**Description**

Plots the step response of a system, given the IR model

**Usage**

```
step(model)
```

**Arguments**

`model`                      an object of class `impulseeest`

**See Also**

[impulseeest](#)

**Examples**

```
uk <- rnorm(1000,1)
yk <- filter (uk,c(0.9,-0.4),method="recursive") + rnorm(1000,1)
data <- idframe(output=data.frame(yk),input=data.frame(uk))
fit <- impulseeest(data)
step(fit)
```

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