

# Package ‘DynNom’

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**Type** Package

**Title** Visualising Statistical Models using Dynamic Nomograms

**Version** 5.1

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**Description** Demonstrate the results of a statistical model object as a dynamic nomogram in an RStudio panel or web browser. The package provides two generics functions: DynNom, which display statistical model objects as a dynamic nomogram; DNbuilder, which builds required scripts to publish a dynamic nomogram on a web server such as the <<https://www.shinyapps.io/>>. Current version of 'DynNom' supports stats::lm, stats::glm, survival::coxph, rms::ols, rms::Glm, rms::lrm, rms::cph, and mgcv::gam model objects.

**License** GPL-2

**Imports** magrittr, shiny, ggplot2 (> 2.1.0), plotly, stargazer, dplyr, compare, BBmisc, broom, rms, survival

**Suggests** mgcv

**NeedsCompilation** no

**Repository** CRAN

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## Contents

DNbuilder . . . . .	2
DynNom . . . . .	4
getclass.DN . . . . .	7
getdata.DN . . . . .	8
getpred.DN . . . . .	9
<b>Index</b>	<b>10</b>

**Description**

DNbuilder is a generic function which builds required scripts to publish a dynamic nomogram on a web server such as the <https://www.shinyapps.io/>. This application can then access through a URL and be used independent of R. DNbuilder supports a large number of model objects from a variety of packages.

**Usage**

```
DNbuilder(model, data = NULL, clevel = 0.95, m.summary = c("raw", "formatted"),
  covariate = c("slider", "numeric"), ptype = c("st", "1-st"),
  DNtitle = NULL, DNxlab = NULL, DNYlab = NULL, DNlimits = NULL,
  KMtitle = NULL, KMxlab = NULL, KMYlab = NULL)
```

```
DNbuilder.core(model, data, clevel, m.summary, covariate,
  DNtitle, DNxlab, DNYlab, DNlimits)
```

```
DNbuilder.surv(model, data, clevel, m.summary, covariate,
  ptype, DNtitle, DNxlab, DNYlab, KMtitle, KMxlab, KMYlab)
```

**Arguments**

model	an lm, glm, coxph, ols, Glm, lrm, cph or mgcv : : gam model object.
data	a dataframe of the accompanying dataset for the model (if required).
clevel	a confidence level for constructing the confidence interval. If not specified, a 95% level will be used.
m.summary	an option to choose the type of the model output represented in the 'Summary Model' tab. "raw" (the default) returns an unformatted summary of the model; "formatted" returns a formatted table of the model summary using stargazer package.
covariate	an option to choose the type of input control widgets used for numeric values. "slider" (the default) picks out sliderInput; "numeric" picks out numericInput.
ptype	an option for coxph or cph model objects to choose the type of plot which displays in "Survival plot" tab. "st" (the default) returns plot of estimated survivor probability $S(t)$ . "1-st" returns plot of estimated failure probability $1-S(t)$ .
DNtitle	a character vector used as the app's title. If not specified, "Dynamic Nomogram" will be used.
DNxlab	a character vector used as the title for the x-axis in "Graphical Summary" tab. If not specified, "Probability" will be used for logistic model and Cox proportional model objects; or "Response variable" for other model objects.
DNYlab	a character vector used as the title for the y-axis in "Graphical Summary" tab (default is NULL).

DNlimits	a vector of 2 numeric values used to set x-axis limits in "Graphical Summary" tab. Note: This also removes the 'Set x-axis ranges' widget in the sidebar panel.
KMtitle	a character vector used as KM plot's title in "Survival plot" tab. If not specified, "Estimated Survival Probability" for ptype = "st" and "Estimated Probability" for ptype = "1-st" will be used.
KMxlab	a character vector used as the title for the x-axis in "Survival plot" tab. If not specified, "Follow Up Time" will be used.
KMylab	a character vector used as the title for the y-axis in "Survival plot" tab. If not specified, "S(t)" for ptype = "st" and "F(t)" for ptype = "1-st" will be used.

### Value

A new folder called 'DynNomapp' will be created in the current working directory which contains all the required scripts to deploy this dynamic nomogram on a host server such as the <https://www.shinyapps.io/>. This folder includes ui.R, server.R, global.R and data.RData which needs to publish the app. A user guide text file (README.txt) will be also added to explain how to deploy the app using these files.

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### References

Banks, J. 2006. Nomograms. Encyclopedia of Statistical Sciences. 8.  
 Easy web applications in R. <https://www.rstudio.com/products/shiny/>  
 Frank E Harrell Jr (2017). rms: Regression Modeling Strategies. R package version 4.5-0. <https://CRAN.R-project.org/package=rms/>

### See Also

[DynNom](#), [getpred.DN](#)

### Examples

```
## Not run:
# Simple linear regression models
fit1 <- lm(uptake ~ Plant + conc + Plant * conc, data = C02)
DNbuilder(fit1)

t.data <- datadist(swiss)
options(datadist = 't.data')
ols(Fertility ~ Agriculture + Education + rcs(Catholic, 4), data = swiss) %>%
```

```

DNbuilder(clevel = 0.9, m.summary="formatted")

# Generalized regression models
fit2 <- glm(Survived ~ Age + Class + Sex,
  data = as.data.frame(Titanic), weights = Freq, binomial("probit"))
DNbuilder(fit2, DNtitle="Titanic", DNxlab = "Probability of survival")

counts <- c(18, 17, 15, 20, 10, 20, 25, 13, 12)
outcome <- gl(3, 1, 9)
treatment <- gl(3, 3)
d <- datadist(treatment, outcome)
options(datadist = "d")
Glm((2 * counts) ~ outcome + treatment, family = poisson(),
  data = data.frame(counts, outcome, treatment)) %>%
  DNbuilder()

# Proportional hazard models
coxph(Surv(time, status) ~ age + strata(sex) + ph.ecog, data = lung) %>%
  DNbuilder()

data.kidney <- kidney
data.kidney$sex <- as.factor(data.kidney$sex)
levels(data.kidney$sex) <- c("male", "female")
coxph(Surv(time, status) ~ age + strata(sex) + disease, data.kidney) %>%
  DNbuilder(ptype = "1-st")

d <- datadist(veteran)
options(datadist = "d")
fit3 <- cph((Surv(time/30, status)) ~ rcs(age, 4) * strat(trt) + diagtime +
  strat(prior) + lsp(karno, 60), veteran)
DNbuilder(fit3, DNxlab = "Survival probability",
  KMtitle="Kaplan-Meier plot", KMxlab = "Time (Days)", KMyLab = "Survival probability")

# Generalized additive models
mgcv::gam(Fertility ~ s(Agriculture) + Education + s(Catholic), data=swiss) %>%
  DNbuilder(DNlimits = c(0, 110), m.summary="formatted")

## End(Not run)
if (interactive()) {
  data(rock)
  lm(area~I(log(peri)), data = rock) %>%
    DNbuilder()
}

```

**Description**

DynNom is a generic function to display the results of statistical model objects as a dynamic nomogram in an 'RStudio' panel or web browser. DynNom supports a large number of model objects from

a variety of packages.

### Usage

```
DynNom(model, data = NULL, clevel = 0.95, m.summary = c("raw", "formatted"),
        covariate = c("slider", "numeric"), ptype = c("st", "1-st"),
        DNtitle = NULL, DNxlab = NULL, DNYlab = NULL, DNlimits = NULL,
        KMtitle = NULL, KMxlab = NULL, KMylab = NULL)
```

```
DynNom.core(model, data, clevel, m.summary, covariate, DNtitle, DNxlab, DNYlab, DNlimits)
```

```
DynNom.surv(model, data, clevel, m.summary, covariate,
            ptype, DNtitle, DNxlab, DNYlab, KMtitle, KMxlab, KMylab)
```

### Arguments

model	an lm, glm, coxph, ols, Glm, lrm, cph or mgcv::gam model object.
data	a dataframe of the accompanying dataset for the model (if required).
clevel	a confidence level for constructing the confidence interval. If not specified, a 95% level will be used.
m.summary	an option to choose the type of the model output represented in the 'Summary Model' tab. "raw" (the default) returns an unformatted summary of the model; "formatted" returns a formatted table of the model summary using stargazer package.
covariate	an option to choose the type of input control widgets used for numeric values. "slider" (the default) picks out sliderInput; "numeric" picks out numericInput.
ptype	an option for coxph or cph model objects to choose the type of plot which displays in "Survival plot" tab. "st" (the default) returns plot of estimated survivor probability (S(t)). "1-st" returns plot of estimated failure probability (1-S(t)).
DNtitle	a character vector used as the app's title. If not specified, "Dynamic Nomogram" will be used.
DNxlab	a character vector used as the title for the x-axis in "Graphical Summary" tab. If not specified, "Probability" will be used for logistic model and Cox proportional model objects; or "Response variable" for other model objects.
DNYlab	a character vector used as the title for the y-axis in "Graphical Summary" tab (default is NULL).
DNlimits	a vector of 2 numeric values used to set x-axis limits in "Graphical Summary" tab. Note: This also removes the 'Set x-axis ranges' widget in the sidebar panel.
KMtitle	a character vector used as KM plot's title in "Survival plot" tab. If not specified, "Estimated Survival Probability" for ptype = "st" and "Estimated Probability" for ptype = "1-st" will be used.
KMxlab	a character vector used as the title for the x-axis in "Survival plot" tab. If not specified, "Follow Up Time" will be used.
KMylab	a character vector used as the title for the y-axis in "Survival plot" tab. If not specified, "S(t)" for ptype = "st" and "F(t)" for ptype = "1-st" will be used.

**Value**

A dynamic nomogram in a shiny application providing individual predictions which can be used as a model visualisation or decision-making tools.

The individual predictions with a relative confidence interval are calculated using the `predict` function, displaying either graphically as an interactive plot in the Graphical Summary tab or a table in the Numerical Summary tab. A table of model output is also available in the Model Summary tab. In the case of the Cox proportional hazards model, an estimated survivor/failure function will be additionally displayed in a new tab.

**Please cite as:**

Jalali A, Alvarez-Iglesias A, Roshan D, Newell J (2019) Visualising statistical models using dynamic nomograms. PLOS ONE 14(11): e0225253. <https://doi.org/10.1371/journal.pone.0225253>

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**References**

Banks, J. 2006. Nomograms. Encyclopedia of Statistical Sciences. 8.

Easy web applications in R. <https://www.rstudio.com/products/shiny/>

Frank E Harrell Jr (2017). rms: Regression Modeling Strategies. R package version 4.5-0. <https://CRAN.R-project.org/package=rms/>

**See Also**

[DNbuilder](#), [getpred.DN](#)

**Examples**

```
## Not run:
# Simple linear regression models
fit1 <- lm(uptake ~ Plant + conc + Plant * conc, data = CO2)
DynNom(fit1)

t.data <- datadist(swiss)
options(datadist = 't.data')
ols(Fertility ~ Agriculture + Education + rcs(Catholic, 4), data = swiss) %>%
  DynNom(clevel = 0.9, m.summary="formatted")

# Generalized regression models
fit2 <- glm(Survived ~ Age + Class + Sex,
  data = as.data.frame(Titanic), weights = Freq, family = binomial("probit"))
DynNom(fit2, DNtitle="Titanic", DNxlab = "Probability of survival")

counts <- c(18, 17, 15, 20, 10, 20, 25, 13, 12)
outcome <- gl(3, 1, 9)
treatment <- gl(3, 3)
```

```

d <- datadist(treatment, outcome)
options(datadist = "d")
Glm((2 * counts) ~ outcome + treatment, family = poisson(),
  data = data.frame(counts, outcome, treatment)) %>%
  DynNom()

# Proportional hazard models
coxph(Surv(time, status) ~ age + strata(sex) + ph.ecog, data = lung) %>%
  DynNom()

data.kidney <- kidney
data.kidney$sex <- as.factor(data.kidney$sex)
levels(data.kidney$sex) <- c("male", "female")
coxph(Surv(time, status) ~ age + strata(sex) + disease, data.kidney) %>%
  DynNom(ptype = "1-st")

d <- datadist(veteran)
options(datadist = "d")
fit3 <- cph((Surv(time/30, status)) ~ rcs(age, 4) * strat(trt) + diagtime +
  strat(prior) + lsp(karno, 60), veteran)
DynNom(fit3, DNxlab = "Survival probability",
  KMtitle="Kaplan-Meier plot", KMxlab = "Time (Days)", KMylab = "Survival probability")

# Generalized additive models
mgcv::gam(Fertility ~ s(Agriculture) + Education + s(Catholic), data=swiss) %>%
  DynNom(DNlimits = c(0, 110), m.summary="formatted")

## End(Not run)
if (interactive()) {
  data(rock)
  lm(area~I(log(peri)), data = rock) %>%
    DynNom()
}

```

---

getclass.DN

*Extract class and family of a model object*


---

## Description

getclass.DN extracts class and family of a model object (supported in DynNom).

## Usage

```
getclass.DN(model)
```

## Arguments

model            an lm, glm, coxph, ols, Glm, lrm, cph or mgcv::gam model objects.

**Value**

A list including the model class and the family name of the model (if relevant).

**See Also**

[DynNom](#), [DNbuilder](#)

**Examples**

```
fit1 <- glm(Survived ~ Age + Class + Sex, data = as.data.frame(Titanic),
  weights = Freq, family = binomial("probit"))
getclass.DN(fit1)

library(survival)
fit2 <- coxph(Surv(time, status) ~ age + strata(sex) + ph.ecog, data = lung)
getclass.DN(fit2)
```

---

getdata.DN

*Extract dataset from a model object*

---

**Description**

getdata.DN extracts dataset that was used to produce the model object (supported in DynNom).

**Usage**

```
getdata.DN(model)
```

**Arguments**

model                    an lm, glm, coxph, ols, Glm, lrm, cph or mgcv: :gam model objects.

**Value**

A data.frame containing the dataset used in the fitted model object.

**See Also**

[DynNom](#), [DNbuilder](#)

**Examples**

```
fit1 <- glm(Survived ~ Age + Class + Sex, data = as.data.frame(Titanic),
  weights = Freq, family = binomial("probit"))
getdata.DN(fit1)

library(survival)
fit2 <- coxph(Surv(time, status) ~ age + strata(sex) + ph.ecog, data = lung)
getdata.DN(fit2)
```



---

`getpred.DN`*Extract predictions from a Model Object*

---

**Description**

`getpred.DN` extracts class, family and inverse of link function from a model object (supported in `DynNom`).

**Usage**

```
getpred.DN(model, newd, set.rms=F)
```

**Arguments**

<code>model</code>	an <code>lm</code> , <code>glm</code> , <code>coxph</code> , <code>ols</code> , <code>Glm</code> , <code>lrm</code> , <code>cph</code> or <code>mgcv</code> : :gam model objects.
<code>newd</code>	a data frame of predictors for prediction
<code>set.rms</code>	a logical value indicating if data should be updated in the model object (required for <code>rms</code> model objects in <code>DNbuilder</code> ).

**Value**

A list including the prediction (`pred`) and the standard error of prediction (`SEpred`).

**See Also**

[DynNom](#), [DNbuilder](#)

**Examples**

```
fit1 <- glm(Survived ~ Age + Class + Sex, data = as.data.frame(Titanic),
  weights = Freq, family = binomial("probit"))
getpred.DN(fit1, newd = data.frame(Class="1st", Sex="Male", Age="Child"))
```

# Index

\* **dynamic nomogram**

DNbuilder, [2](#)

DynNom, [4](#)

\* **model visualisation**

DNbuilder, [2](#)

DynNom, [4](#)

\* **shiny**

DNbuilder, [2](#)

DynNom, [4](#)

DNbuilder, [2](#), [6](#), [8](#), [9](#)

DynNom, [3](#), [4](#), [8](#), [9](#)

getclass.DN, [7](#)

getdata.DN, [8](#)

getpred.DN, [3](#), [6](#), [9](#)