

Package ‘nmw’

April 10, 2018

Version 0.1.4

Title Understanding Nonlinear Mixed Effects Modeling for Population Pharmacokinetics

Description This shows how NONMEM(R) <<http://www.iconplc.com/innovation/nonmem/>> software works. NONMEM's classical estimation methods like 'First Order(FO) approximation', 'First Order Conditional Estimation(FOCE)', and 'Laplacian approximation' are explained.

Depends R (>= 3.0.0), numDeriv

ByteCompile yes

License GPL-3

Copyright 2017-, Kyun-Seop Bae

Author Kyun-Seop Bae

Maintainer Kyun-Seop Bae <k@acr.kr>

URL <https://cran.r-project.org/package=nmw>

NeedsCompilation no

Repository CRAN

Date 2018-04-10 00:01:04 KST

Date/Publication 2018-04-10 12:54:54 UTC

R topics documented:

nmw-package	2
CovStep	3
EstStep	5
InitStep	6
TabStep	8
Index	9

Description

This shows how NONMEM(R) <<http://www.iconplc.com/innovation/nonmem/>> software works.

Details

This package explains 'First Order(FO) approximation' method, 'First Order Conditional Estimation(FOCE)' method, and 'Laplacian(LAPL)' method of NONMEM software.

Author(s)

Kyun-Seop Bae <k@acr.kr>

References

1. NONMEM Users guide
2. Wang Y. Derivation of various NONMEM estimation methods. J Pharmacokinet Pharmacodyn. 2007.
3. Kang D, Bae K, Houp BE, Savic RM, Karlsson MO. Standard Error of Empirical Bayes Estimate in NONMEM(R) VI. K J Physiol Pharmacol. 2012.
4. Kim M, Yim D, Bae K. R-based reproduction of the estimation process hidden behind NONMEM Part 1: First order approximation method. 2015.
5. Bae K, Yim D. R-based reproduction of the estimation process hidden behind NONMEM Part 2: First order conditional estimation. 2016.

Examples

```
DataAll = Theoph
colnames(DataAll) = c("ID", "BWT", "DOSE", "TIME", "DV")
DataAll[, "ID"] = as.numeric(as.character(DataAll[, "ID"]))

nTheta = 3
nEta = 3
nEps = 2

THETAinit = c(2, 50, 0.1)
OMinit = matrix(c(0.2, 0.1, 0.1, 0.1, 0.2, 0.1, 0.1, 0.1, 0.2), nrow=nEta, ncol=nEta)
SGinit = diag(c(0.1, 0.1))

LB = rep(0, nTheta) # Lower bound
UB = rep(1000000, nTheta) # Upper bound

FGD = deriv(~DOSE/(TH2*exp(ETA2))*TH1*exp(ETA1)/(TH1*exp(ETA1) - TH3*exp(ETA3))*  
          (exp(-TH3*exp(ETA3)*TIME)-exp(-TH1*exp(ETA1)*TIME)),
```

```

c("ETA1", "ETA2", "ETA3"),
function.arg=c("TH1", "TH2", "TH3", "ETA1", "ETA2", "ETA3", "DOSE", "TIME"),
func=TRUE, hessian=TRUE)
H = deriv(~F + F*EPS1 + EPS2, c("EPS1", "EPS2"), function.arg=c("F", "EPS1", "EPS2"), func=TRUE)

PRED = function(THETA, ETA, DATAi)
{
  FGDres = FGD(THETA[1], THETA[2], THETA[3], ETA[1], ETA[2], ETA[3], DOSE=320, DATAi[, "TIME"])
  Gres = attr(FGDres, "gradient")
  Hres = attr(H(FGDres, 0, 0), "gradient")

  if (e$METHOD == "LAPL") {
    Dres = attr(FGDres, "hessian")
    Res = cbind(FGDres, Gres, Hres, Dres[,1,1], Dres[,2,1], Dres[,2,2], Dres[,3,])
    colnames(Res) = c("F", "G1", "G2", "G3", "H1", "H2", "D11", "D21", "D22", "D31", "D32", "D33")
  } else {
    Res = cbind(FGDres, Gres, Hres)
    colnames(Res) = c("F", "G1", "G2", "G3", "H1", "H2")
  }
  return(Res)
}

##### First Order Approximation Method
InitStep(DataAll, THETAinit=THETAinit, OMinit=OMinit, SGinit=SGinit, LB=LB, UB=UB,
         Pred=PRED, METHOD="ZERO")
(EstRes = EstStep())           # 4 sec
(CovRes = CovStep())          # 2 sec
PostHocEta() # Using e$FinalPara from EstStep()
#TabStep()   # Commented out for the CRAN CPU time

##### First Order Conditional Estimation with Interaction Method
#InitStep(DataAll, THETAinit=THETAinit, OMinit=OMinit, SGinit=SGinit, LB=LB, UB=UB,
#         Pred=PRED, METHOD="COND")
#(EstRes = EstStep())           # 2 min
#(CovRes = CovStep())          # 1 min
#get("EBC", envir=e)
#TabStep()

##### Laplacian Approximation with Interaction Method
#InitStep(DataAll, THETAinit=THETAinit, OMinit=OMinit, SGinit=SGinit, LB=LB, UB=UB,
#         Pred=PRED, METHOD="LAPL")
#(EstRes = EstStep())           # 4 min
#(CovRes = CovStep())          # 1 min
#get("EBC", envir=e)
#TabStep()

```

Description

It calculates standard errors and various variance matrices with the e\$FinalPara after estimation step.

Usage

`CovStep()`

Details

Because `EstStep` uses nonlinear optimization, covariance step is separated from estimation step. It calculates variance-covariance matrix of estimates in the original scale.

Value

Time	consumed time
Standard Error	standard error of the estimates in the order of theta, omega, and sigma
Covariance Matrix of Estimates	covariance matrix of estimates in the order of theta, omega, and sigma. This is inverse(R) x S x inverse(R) by default.
Correlation Matrix of Estimates	correlation matrix of estimates in the order of theta, omega, and sigma
Inverse Covariance Matrix of Estimates	inverse covariance matrix of estimates in the order of theta, omega, and sigma
Eigen Values	eigen values of covariance matrix
R Matrix	R matrix of NONMEM, the second derivative of log likelihood function with respect to estimation parameters
S Matrix	S matrix of NONMEM, sum of individual cross-product of the first derivative of log likelihood function with respect to estimation parameters

Author(s)

Kyun-Seop Bae <k@acr.kr>

References

NONMEM Users Guide

See Also

[EstStep](#), [InitStep](#)

Examples

```
# Only after InitStep and EstStep
#CovStep()
```

EstStep

Estimation Step

Description

This estimates upon the conditions with `InitStep`.

Usage

`EstStep()`

Details

It does not have arguments. All necessary arguments are stored in the `e` environment. It assumes "INTERACTION" between eta and epsilon for "COND" and "LAPL" options. The output is basically same to NONMEM output.

Value

Initial OFV	initial value of the objective function
Time	time consumed for this step
Optim	the raw output from <code>optim</code> function
Final Estimates	final estimates in the original scale

Author(s)

Kyun-Seop Bae <k@acr.kr>

References

NONMEM Users Guide

See Also

[InitStep](#)

Examples

```
# Only After InitStep  
#EstStep()
```

InitStep*Initialization Step***Description**

It receives parameters for the estimation and stores them into e environment.

Usage

```
InitStep(DataAll, THETAinit, OMinit, SGinit, LB, UB, Pred, METHOD)
```

Arguments

DataAll	Data for all subjects. It should contain columns which Pred function uses.
THETAinit	Theta initial values
OMinit	Omega matrix initial values
SGinit	Sigma matrix initial values
LB	Lower bounds for theta vector
UB	Upper bounds for theta vector
Pred	Prediction function name
METHOD	one of the estimation methods "ZERO", "COND", or "LAPL"

Details

Prediction function should return not only prediction values(F or IPRED) but also G (first derivative with respect to etas) and H (first derivative of Y with respect to epsilon). For the "LAPL", prediction function should return second derivative with respect to eta also. "INTERACTION" is TRUE for "COND" and "LAPL" option, and FALSE for "ZERO". Omega matrix should be full block one. Sigma matrix should be diagonal one.

Value

This does not return values, but stores necessary values into the environment e.

Author(s)

Kyun-Seop Bae <k@acr.kr>

References

NONMEM Users Guide

Examples

```

DataAll = Theoph
colnames(DataAll) = c("ID", "BWT", "DOSE", "TIME", "DV")
DataAll[, "ID"] = as.numeric(as.character(DataAll[, "ID"]))

nTheta = 3
nEta = 3
nEps = 2

THETAinit = c(2, 50, 0.1) # Initial estimate
OMinit = matrix(c(0.2, 0.1, 0.1, 0.1, 0.2, 0.1, 0.1, 0.1, 0.2), nrow=nEta, ncol=nEta)
OMinit
SGinit = diag(c(0.1, 0.1))
SGinit

LB = rep(0, nTheta) # Lower bound
UB = rep(1000000, nTheta) # Upper bound

FGD = deriv(~DOSE/(TH2*exp(ETA2))*TH1*exp(ETA1)/(TH1*exp(ETA1) - TH3*exp(ETA3))*  

            (exp(-TH3*exp(ETA3)*TIME)-exp(-TH1*exp(ETA1)*TIME)),  

            c("ETA1", "ETA2", "ETA3"),  

            function.arg=c("TH1", "TH2", "TH3", "ETA1", "ETA2", "ETA3", "DOSE", "TIME"),  

            func=TRUE, hessian=TRUE)
H = deriv(~F + F*EPS1 + EPS2, c("EPS1", "EPS2"), function.arg=c("F", "EPS1", "EPS2"), func=TRUE)

PRED = function(THETA, ETA, DATAi)  

{
  FGDres = FGD(THETA[1], THETA[2], THETA[3], ETA[1], ETA[2], ETA[3], DOSE=320, DATAi[, "TIME"])
  Gres = attr(FGDres, "gradient")
  Hres = attr(H(FGDres, 0, 0), "gradient")

  if (e$METHOD == "LAPL") {
    Dres = attr(FGDres, "hessian")
    Res = cbind(FGDres, Gres, Hres, Dres[,1,1], Dres[,2,1], Dres[,2,2], Dres[,3,])
    colnames(Res) = c("F", "G1", "G2", "G3", "H1", "H2", "D11", "D21", "D22", "D31", "D32", "D33")
  } else {
    Res = cbind(FGDres, Gres, Hres)
    colnames(Res) = c("F", "G1", "G2", "G3", "H1", "H2")
  }
  return(Res)
}

##### First Order Approximation Method
InitStep(DataAll, THETAinit=THETAinit, OMinit=OMinit, SGinit=SGinit, LB=LB, UB=UB,
         Pred=PRED, METHOD="ZERO")

##### First Order Conditional Estimation with Interaction Method
InitStep(DataAll, THETAinit=THETAinit, OMinit=OMinit, SGinit=SGinit, LB=LB, UB=UB,
         Pred=PRED, METHOD="COND")

##### Laplacian Approximation with Interaction Method
InitStep(DataAll, THETAinit=THETAinit, OMinit=OMinit, SGinit=SGinit, LB=LB, UB=UB,
         Pred=PRED, METHOD="COND")

```

```
Pred=PRED, METHOD="LAPL")
```

TabStep

Table Step

Description

This produces standard table.

Usage

```
TabStep()
```

Details

It does not have arguments. All necessary arguments are stored in the e environment. This is similar to other standard results table.

Value

A table with ID, TIME, DV, PRED, RES, WRES, derivatives of G and H. If the estimation method is other than 'ZERO'(First-order approximation), it includes CWRES, CIPREDI(formerly IPRED), CIRESI(formerly IRES).

Author(s)

Kyun-Seop Bae <k@acr.kr>

References

NONMEM Users Guide

See Also

[EstStep](#)

Examples

```
# Only After EstStep
#TabStep()
```

Index

- *Topic **Covariance Step**
 - CovStep, 3
 - *Topic **Estimation Step**
 - EstStep, 5
 - *Topic **Initialization Step**
 - InitStep, 6
 - *Topic **Nonlinear Mixed Effects Modeling**
 - nmw-package, 2
 - *Topic **Population Pharmacokinetics**
 - nmw-package, 2
 - *Topic **Tabulation Step**
 - TabStep, 8
- CovStep, 3
- EstStep, 4, 5, 8
- InitStep, 4, 5, 6
- nmw (nmw-package), 2
- nmw-package, 2
- TabStep, 8