

# Package ‘Rquefts’

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**Type** Package

**Title** Quantitative Evaluation of the Native Fertility of Tropical Soils

**Version** 1.0-7

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**LinkingTo** Rcpp

**SystemRequirements** C++11

**Imports** meteor, methods, Rcpp (>= 0.12.4)

**Description** An implementation of the QUEFTS (Quantitative Evaluation of the Native Fertility of Tropical Soils) model. The model (1) estimates native nutrient (N, P, K) supply of soils from a few soil chemical properties; and (2) computes crop yield given that supply, fertilizer application and crop parameters. See Janssen et al. (1990) <doi:10.1016/0016-7061(90)90021-Z> for the technical details and Sartari et al. (2014) <doi:10.1016/j.fcr.2013.12.005> for a recent evaluation and improvements.

**License** GPL (>= 3)

**BugReports** <https://github.com/cropmodels/Rquefts/issues>

**URL** <https://github.com/cropmodels/Rquefts/>

**NeedsCompilation** yes

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Rquefts-package	<i>Quantitative Evaluation of the Native Fertility of Tropical Soils</i>
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### Description

This package provides a R interface to a C++ implementation of the QUEFTS model. QUEFTS (Quantitative Evaluation of the Native Fertility of Tropical Soils) model (1) estimates native nutrient (N, P, K) supply of soils from a few soil chemical properties; and (2) computes crop yield given that supply, fertilizer application and crop parameters. See Janssen et al. (1990) <doi:10.1016/0016-7061(90)90021-Z> for the technical details and Sattari et al. (2014) <doi:10.1016/j.fcr.2013.12.005> for a recent evaluation and improvements.

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Fertilizers	<i>Helper functions to go from fertilizers to nutrients</i>
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### Description

Computes the amount of nutrients given a rate of fertilizer.

### Usage

```
fertilizers()
nutrientRates(supply, treatment)
```

### Arguments

supply	data.frame with columns "N", "P", "K" expressed as percentage of the product (row)
treatment	amounts applied

### Examples

```
# fertilizer product list
fert <- fertilizers()
# shortening some of the names for display
fert[,2] = substr(fert[,2], 1, 20)
# contents are expressed as a percentage.
fert

myferts <- fert[c(8,15), ]
nutrientRates(myferts, c(100,50))
```

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 nutSupply

*Soil nutrients supply for QUEFTS model*


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### Description

nutSupply1 computes the base (unfertilized) soil supply of N, P and K according to Janssen et al. (1990), Table 2. For use with the QUEFTS model.

nutSupply2 is a modified version following Sattari et al. (2014). It has an additional variable "temperature", and P-total is required. Sattari et al suggest that, for soils that have not been fertilized with P, you can estimate P-total as  $95 * P\text{-Olsen}$ . Emperically, using soilGrids, I found  $55 * P\text{-Olsen}$ .

### Usage

```
nutSupply1(pH, SOC, Kex, Polsen, Ptotal=NA)
nutSupply2(temp, pH, SOC, Kex, Polsen, Ptotal)
```

### Arguments

temp	average growing season temperature (C)
pH	soil pH (H <sub>2</sub> O)
SOC	soil organic carbon (g/kg)
Kex	exchangeble K in the soil (mmol/kg)
Polisen	soil P measured with the P-Olsen method (mg/kg)
Ptotal	total soil P (mg/kg)

### Value

Matrix with three columns: Nsup, Psup and Ksup. These are the potential supply of N, P and K of the unfertilized soil (kg/ha).

### References

Janssen B.H., F.C.T. Guiking, D. van der Eijk, E.M.A. Smaling, J. Wolf and H. van Reuler, 1990. A system for the quantitative evaluation of the fertility of tropical soils (QUEFTS). Geoderma 46: 299-318

Sattari, S.Z., M.K. van Ittersum, A.F. Bouwman, A.L. Smit, and B.H. Janssen, 2014. Crop yield response to soil fertility and N, P, K inputs in different environments: Testing and improving the QUEFTS model. Field Crops Research 157: 35-46

### Examples

```
s1 <- nutSupply1(6, c(23, 11, 35), 15, c(1.6, 2.6, 2.4))
s1
s2 <- nutSupply2(20, 6, c(23, 11, 35), 15, c(1.6, 2.6, 2.4), 225)
s2
```

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 quefts

*QUEFTS model*


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### Description

Create a QUEFTS model, set parameters, and run it to compute nutrient requirements and nutrient limited yield.

A number of default crop parameter sets are provided, as well as one example soil. You need to provide attainable crop production (in this context that is the maximum production in the absence of nutrient limitation), or target dry-matter biomass for leaves, stems and the storage organ (e.g. grain, root or tuber). Some crops are grown for the stems/leaves, in which case there is no relevant storage organ (e.g. sugarcane, jute). production yield estimates can be obtained with a crop growth model.

For a cereal crop you can assume that 50

### Usage

```
quefts(soil, crop, fert, biom)
quefts_soil()
quefts_fert()
quefts_crop(name="")
quefts_biom()
crop(x) <- value
soil(x) <- value
fert(x) <- value
biom(x) <- value
run(x, ...)
```

### Arguments

soil	list with named soil parameters. See Details. An example is returned by <code>quefts_soil()</code>
crop	list with named crop parameters. See Details. An example is returned by <code>quefts_crop()</code>
fert	list with named fertilizer parameters (N, P and K). An example is returned by <code>quefts_fert()</code>
biom	list with named biomass and growing season length parameters. An example is returned by <code>quefts_biom()</code>
name	character. crop name
x	QueftsModel object
value	list with soil, crop, fertilizer, or biomass parameters as above
...	additional arguments. None implemented

### Details

**Input Parameters****Soil**

N\_base\_supply, P\_base\_supply, K\_base\_supply

N\_recovery, P\_recovery, K\_recovery

UptakeAdjust

**Crop**

\_minVeg, \_maxVeg, \_minStore, \_maxStore

Yzero

Nfix

**Management**

N, P, K

**Crop yield**

leaf\_att, stem\_att, store\_att

SeasonLength

.

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**Output Variables**

N\_actual\_supply, P\_actual\_supply, K\_actual\_supply

leaf\_lim, stem\_lim, store\_lim

N\_gap, P\_gap, K\_gap

**Explanation**

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Potential supply (kg/ha) of N, P and K of the (unfertilized) soil

Fertilizer recovery, that is, the fraction of applied fertilizer that is recovered

Two-column matrix to compute the fraction uptake from soil

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minimum and maximum concentration of "\_" (N, P, or K) in soil

the maximum biomass of vegetative organs at zero yield of soil

the fraction of a crop's nitrogen uptake supplied by biological N fixation

.

N, P, and K fertilizer applied.

.

Attainable (in the absence of nutrient limitation), or target crop yield

Length of the growing season (days)

**Explanation**

nutrient uptake from soil (not fertilizer) (kg/ha)

nutrient limited biomass of leaves, stems, and storage organs (kg/ha)

fertilizer required to reach the specified biomass (kg/ha)

**Value**

vector with output variables as described in the Details

**References**

Janssen B.H., F.C.T. Guiking, D. van der Eijk, E.M.A. Smaling, J. Wolf and H. van Reuler, 1990. A system for the quantitative evaluation of the fertility of tropical soils (QUEFTS). *Geoderma* 46: 299-318

Sattari, S.Z., M.K. van Ittersum, A.F. Bouwman, A.L. Smit, and B.H. Janssen, 2014. Crop yield response to soil fertility and N, P, K inputs in different environments: Testing and improving the QUEFTS model. *Field Crops Research* 157: 35-46

**Examples**

```
# create a QUEFTS model
# 1. get parameters
soiltype <- quefts_soil()
barley <- quefts_crop("Barley")
fertilizer <- list(N=0, P=0, K=0)
att_yield <- list(leaf_att=2200, stem_att=2700, store_att=4800, SeasonLength=110)

# 2. create a model
q <- quefts(soiltype, barley, fertilizer, att_yield)
```

```

# 3. run the model
run(q)

# change some parameters
q$SeasonLength <- 162
q$leaf_att <- 2651
q$stem_att <- 5053
q$store_att <- 8208

q$N <- 100
q$P <- 50
q$K <- 50

run(q)

## note that Rquefts uses C++ reference classes.
## This means that if you copy a quefts model, you do not create a
## new instance of the model, but you point to the same one!
q <- quefts()
q["N"]
k <- q
k["N"] <- 150
k["N"]
# the value of q has also changed!
q["N"]

## different ways of subsetting / replacement
q <- quefts()
q$N
q$N <- 30
q["N"]
q["N"] <- 90
q["model", "N"]
q["model", "N"] <- 60
q$N

q$soil$N_recovery
q["soil$N_recovery"]
q["soil$N_recovery"] <- .6
q["soil", "N_recovery"]
q["soil", "N_recovery"] <- .4
q$soil$N_recovery

```

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QueftsModel-class

*QueftsModel class*


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### Description

These are the classes!

*QueftsModel-class*

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**Objects from the Class**

QueftsModel\$new()

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