

Additional functions for transforming soil particlesize distributions

Wei Shangguan

August 31, 2010

1 Load the soiltexture package

The soiltexture package can be installed from CRAN with the following commands:

```
install.packages("soiltexture")
```

And loaded with the following commands:

```
require("soiltexture")
```

2 Transforming soil texture data using many Particle-Size Distribution models (from 3 or more particle size classes)

`TT.text.transf.Xm()` is used to transform soil texture data from 3 or more particle size classes using various Particle-Size Distribution (PSD) models. The `drc` package and its associate packages(`lattice`,`magic`,`nlme`, `plotrix`) are required in the PSD model fitting. Compared to `TT.text.transf()`, the following check is not needed (and not done) :

- When the 1st value of input `tri.data` and output particle size classes limits is 0, The 2nd value of the output particle size classes limits must be higher or equal to the 2nd value of the input particle size classes limits.”

We need first to create a dummy dataset with more than 3 particle size classes:

```
my.text4 <- data.frame(  
  "CLAY" = c(05,60,15,05,25,05,25,45,65,75,13,47),  
  "FSILT" = c(02,04,10,15,25,40,35,20,10,05,10,20),  
  "CSILT" = c(03,04,05,10,30,45,30,25,05,10,07,23),  
  "SAND" = c(90,32,70,70,20,10,10,10,20,10,70,10)  
) #
```

Transform this data frame from 4 particle size classes to 3 particle size classes:

```

res <- TT.text.transf.Xm(
  tri.data      = my.text4,
  base.ps.lim   = c(0,1,50,2000),
  dat.ps.lim    = c(0,2,30,60,2000),
  psdmodel      = "AD"
) #
#
round( res[,1:6], 3 )

      0-1    1-50 50-2000 f0:(Intercept) b:(Intercept)
[1,]  4.341  4.651  91.007           0.584           0.364
[2,] 59.657  6.931  33.412           0.807           0.148
[3,] 13.657 14.860  71.483           0.763           0.477
[4,]  3.408 23.472  73.119           0.571           0.412
[5,] 24.116 49.480  26.403           0.619           0.265
[6,]  4.432 81.454  14.123           0.520           0.318
[7,] 24.363 62.045  13.592           0.620           0.255
[8,] 44.532 41.597  13.889           0.722           0.189
[9,] 63.849 14.739  21.412           0.833           0.171
[10,] 74.778 11.982  13.239           0.874           0.087
[11,] 11.934 15.827  72.239           0.611           0.361
[12,] 46.489 39.836  13.679           0.731           0.183
      c:(Intercept)
[1,]           4.276
[2,]           3.211
[3,]           1.314
[4,]           1.630
[5,]           4.298
[6,]           9.168
[7,]           6.745
[8,]           5.913
[9,]           1.102
[10,]          4.801
[11,]           1.989
[12,]          5.531

#
round( res[,7:ncol(res)], 3 )

      r0:(Intercept)    dev
[1,]           0.613 0.783
[2,]           0.138 0.000
[3,]           0.773 0.003
[4,]           0.179 0.000
[5,]           0.039 0.000
[6,]           0.032 0.000
[7,]           0.031 0.000
[8,]           0.035 0.002
[9,]           0.090 0.000
[10,]          0.052 0.000

```

```
[11,]          0.231 0.000
[12,]          0.035 0.000
```

The first 3 columns are the predicted values with a sum not equal to 100% (can be normalised by `TT.normalise.sum.X()`). The following 4 columns are the fitted PSD model parameters. And the last column is the Residual Sum of Squares (deviance). Note that the transforming results may be slightly different even with the same function parameters. This is caused by the nature of `drc` package in fitting dose-response models.

Sometimes, the fitting will fail for the iteration is not converged or some errors and warnings happened. These can be ignored, as you can get the transforming results.

The following PSD models are implemented: Anderson (AD), Fredlund4P (F4P), Fredlund3P (F3P), modified logistic growth (ML), Offset-Nonrenormalized Lognormal (ONL), Offset-Renormalized Lognormal (ORL), Skaggs (S), van Genuchten type(VG), van Genuchten modified, Weibull (W), Logarithm(L), Logistic growth (LG), Simple Lognormal (SL), Shiozawa and Campbell (SC). The performance of PSD models is influenced by many aspects like soil texture class, number and position (or closeness) of observation points, clay content etc. The latter four PSD models perform worse than the former ten. The AD, F4P, S, and W model is recommended for most of texture classes. And it will be even better to compare several different PSD models and using the results of the model with the minimum residual sum of squares. Except S and W models, all the PSD models could be used to predict the content below the minimum input limit. The "psdmodel" option could be changed to any other of the above models:

```
res <- TT.text.transf.Xm(
  tri.data      = my.text4,
  base.ps.lim   = c(0,1,50,2000),
  dat.ps.lim    = c(0,2,30,60,2000),
  psdmodel      = "ML"
) #
#
round( res[,1:6], 3 )

      0-1    1-50 50-2000 a:(Intercept) b:(Intercept)
[1,]  4.942  3.946 91.112          19.472          13.364
[2,] 59.849  6.849 33.302           0.675           5.739
[3,] 14.721 13.805 70.984           6.473           4.910
[4,]  4.413 22.511 72.511          53.861           7.420
[5,] 24.466 46.833 28.700           3.162          62.767
[6,]  4.377 76.265 19.359          26.139          57.107
[7,] 24.185 58.851 16.964           3.259          72.583
[8,] 44.788 38.541 16.671           1.238         180.300
[9,] 64.027 14.560 21.321           0.615           4.399
[10,] 74.978 11.682 13.340           0.334         249.053
[11,] 12.405 15.396 71.925           8.171           5.553
```

```

[12,] 46.747 37.139 16.114          1.146          144.537
      c:(Intercept)
[1,]          1.014
[2,]          0.983
[3,]          0.549
[4,]          0.304
[5,]          1.140
[6,]          0.834
[7,]          1.090
[8,]          1.534
[9,]          0.563
[10,]         1.927
[11,]         0.527
[12,]         1.467

#
round( res[,7:ncol(res)], 3 )

[1] 0.000 0.000 1.669 1.071 0.000 0.000 0.000 0.000 0.011 0.000
[11] 0.205 0.000

```

Because the current PSD model fitting is quite time-consuming and some models are not always successful for all soils, you can change the PSD model, or optimization method potentially at the cost of some accuracy. The default "omethod" option (i.e. "all") is to run all methods and choose the best results with minimum residual sum of squares. The optional methods are "Nelder-Mead", "BFGS", "CG", "L-BFGS-B", "SANN" (see `optim()` for details.)

```

res <- TT.text.transf.Xm(
  tri.data      = my.text4,
  base.ps.lim   = c(0,1,50,2000),
  dat.ps.lim    = c(0,2,30,60,2000),
  psdmodel      = "ML",
  omethod       = "SANN"
) #
#
round( res[,1:6], 3 )

      0-1    1-50 50-2000 a:(Intercept) b:(Intercept)
[1,]  4.941  3.946 91.113          19.473          13.367
[2,] 59.848  6.849 33.302           0.675           5.738
[3,] 14.721 13.805 70.984           6.473           4.911
[4,]  4.413 22.512 72.511          53.852           7.420
[5,] 24.467 46.832 28.702           3.162          62.730
[6,]  4.375 76.264 19.361          26.155          57.066
[7,] 24.186 58.850 16.964           3.259          72.587
[8,] 44.788 38.541 16.671           1.238         180.309
[9,] 64.027 14.560 21.322           0.614           4.403
[10,] 74.978 11.682 13.340           0.334         248.948
[11,] 12.406 15.396 71.926           8.170           5.555
[12,] 46.748 37.137 16.115           1.146          144.549

```

```

      c:(Intercept)
[1,]      1.014
[2,]      0.983
[3,]      0.549
[4,]      0.304
[5,]      1.140
[6,]      0.834
[7,]      1.090
[8,]      1.534
[9,]      0.564
[10,]     1.927
[11,]     0.527
[12,]     1.467

#
round( res[,7:ncol(res)], 3 )

[1] 0.000 0.000 1.669 1.071 0.000 0.000 0.000 0.000 0.011 0.000
[11] 0.205 0.000

```

3 Normalizing soil texture data (sum of X texture classes)

TT.normalise.sum.X() is similar to TT.normalise.sum(). But it normalize the sum of the X (X>1) texture classes instead of 3. The option tri.data should be a data.frame with only soil texture data (no additional extra columns should be present).

```

my.text5 <- data.frame(
  "CLAY" = c(05,60,15,04.9,25,05,25,45,65,75,13,47),
  "FSILT" = c(02,04.3,10,15,25,40,35,20,10,05,10,20),
  "CSILT" = c(03,04,05,10,30,45,30,25,05,10,07.2,23.3),
  "SAND" = c(90.5,32,70,70,20.3,10.9,9.3,9.4,20,10,70,10)
) #
#
res <- TT.normalise.sum.X(
  tri.data = my.text5,
  residuals = TRUE
) #

[1] 100.5 100.3 100.0 99.9 100.3 100.9 99.3 99.4 100.0 100.0
[11] 100.2 100.3

#
res

      CLAY      FSILT      CSILT      SAND residuals
[1,] 4.975124 1.990050 2.985075 90.049751      0.5
[2,] 59.820538 4.287139 3.988036 31.904287      0.3
[3,] 15.000000 10.000000 5.000000 70.000000      0.0

```

[4,]	4.904905	15.015015	10.010010	70.070070	-0.1
[5,]	24.925224	24.925224	29.910269	20.239282	0.3
[6,]	4.955401	39.643211	44.598612	10.802775	0.9
[7,]	25.176234	35.246727	30.211480	9.365559	-0.7
[8,]	45.271630	20.120724	25.150905	9.456740	-0.6
[9,]	65.000000	10.000000	5.000000	20.000000	0.0
[10,]	75.000000	5.000000	10.000000	10.000000	0.0
[11,]	12.974052	9.980040	7.185629	69.860279	0.2
[12,]	46.859422	19.940179	23.230309	9.970090	0.3