PCNM

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Source: website of Pierre Legendre (http://adn.biol.umontreal.ca/~numericalecology/Rcode/)

Originally part of the `library (PCNM)`, which is no longer supported. Needs to install both `PCNM` function and also supporting `pcoa.all` function. Check how to install instructions, and also the documentation for PCNM and pcoa.all.

Direct import of Legendre's code:

```
source
('https://raw.githubusercontent.com/zdealveindy/anadat-r/master/scripts/PCNM.R')
source
('https://raw.githubusercontent.com/zdealveindy/anadat-
r/master/scripts/pcoa.all.R')
```

From:
https://anadat-r.davidzeleny.net/ - **Analysis of community ecology data in R**

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https://anadat-r.davidzeleny.net/doku.php/en:customized_functions:pcnm

Last update: **2022/04/29 22:45**