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```
# CA - scaling 1 vs scaling 2

library (vegan)
data (dune)
CA <- cca (dune)
png ('CA-scaling1-2.png', width = 8, height = 4, units = 'in', res = 300)
par (mfrow = c(1,2))
plot (CA, scaling = 1, main = 'CA on dune data, Scaling 1')
plot (CA, scaling = 2, main = 'CA on dune data, Scaling 2')
dev.off ()
```

From:

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

Permanent link:

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

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