

Harvard Forest Data Archive HF105-02

Data File:

Name = hf105-02-understory-2004.csv

Description = understory vegetation 2004

Rows = 40 Columns = 28

MD5 checksum = f21fa6e5d0d5d2977b821540ffallb94

Variables:

x = x-coordinate location, in meters, of the sample. Each plot of  
the Simes Tract is 90 x 90 m. (meter)  
y = y-coordinate location, in meters, of the sample. Each plot at  
the Simes Tract is 90 x 90 m. (meter)  
acru = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
acsa = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
be = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
coam = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
osvi = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
pist = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
prse = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
quru = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
tsca = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
rufl = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
vaan = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
arnu = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
fram = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
maca = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
mevi = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
mire = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
smra = number of seedings that emerged in each 2-cm-thick subsample  
(number)  
cape = number of seedings that emerged in each 2-cm-thick subsample  
(number)

jute = number of seedlings that emerged in each 2-cm-thick subsamp  
(number)

dasp = number of seedlings that emerged in each 2-cm-thick subsamp  
(number)

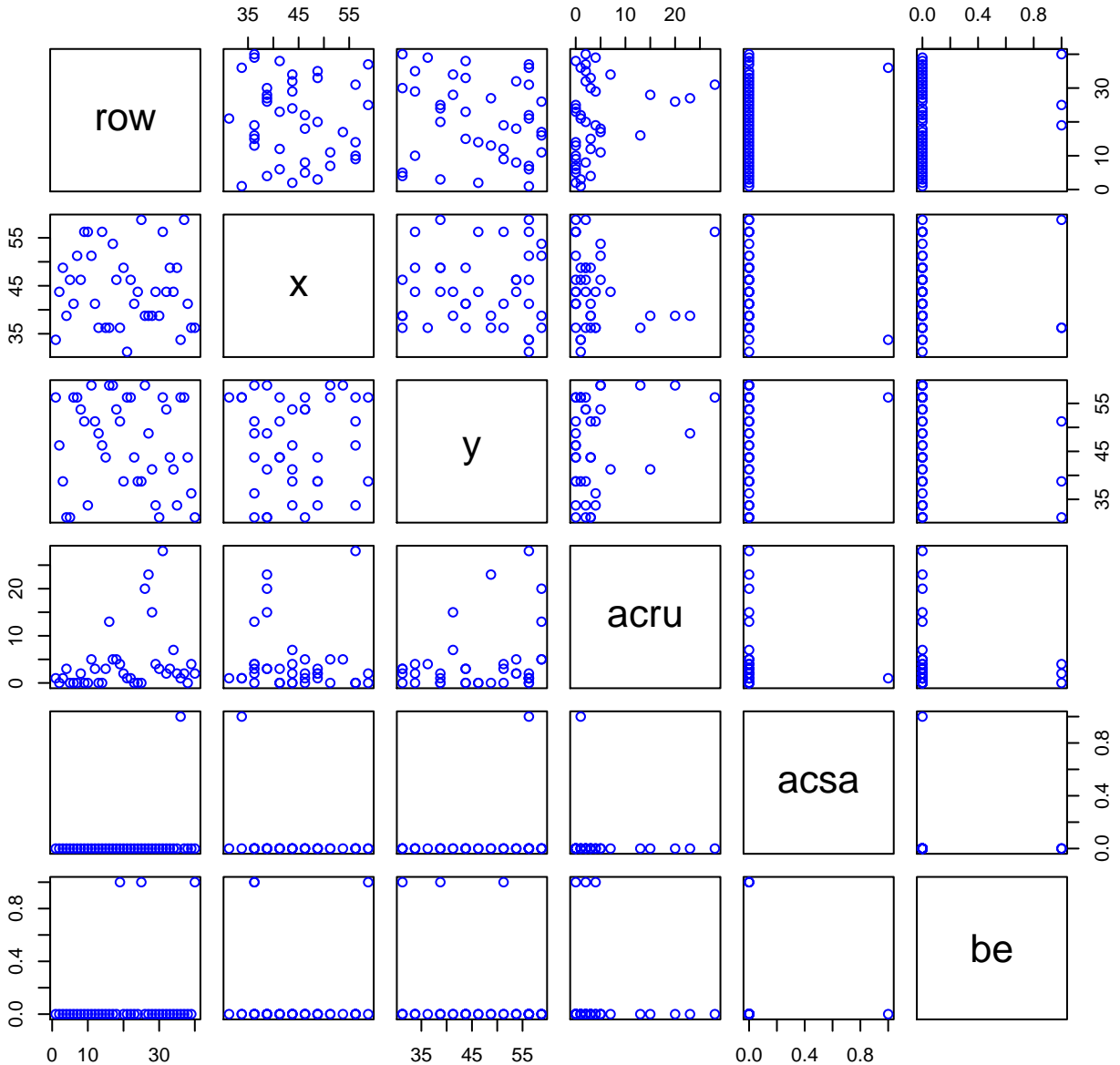
depu = number of seedlings that emerged in each 2-cm-thick subsamp  
(number)

drsp = number of seedlings that emerged in each 2-cm-thick subsamp  
(number)

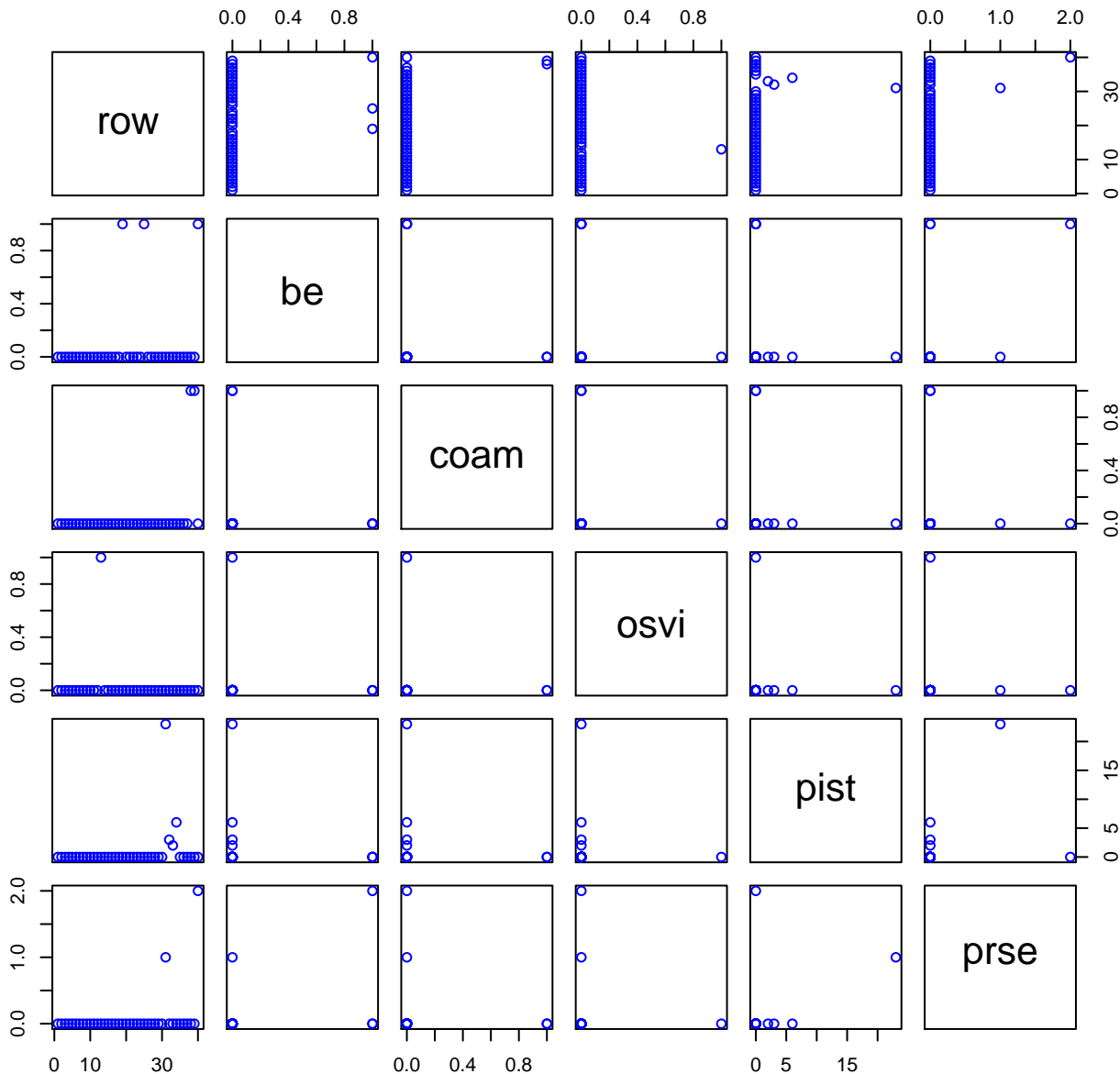
deob = number of seedlings that emerged in each 2-cm-thick subsamp  
(number)

| Variable | Min    | Median | Mean   | Max     | NAs |
|----------|--------|--------|--------|---------|-----|
| x        | 31.250 | 43.750 | 44.312 | 58.750  | 0   |
| y        | 31.250 | 47.500 | 46.625 | 58.750  | 0   |
| acru     | 0.000  | 2.000  | 4.125  | 28.000  | 0   |
| acsa     | 0.000  | 0.000  | 0.025  | 1.000   | 0   |
| be       | 0.000  | 0.000  | 0.075  | 1.000   | 0   |
| coam     | 0.000  | 0.000  | 0.050  | 1.000   | 0   |
| osvi     | 0.000  | 0.000  | 0.025  | 1.000   | 0   |
| pist     | 0.000  | 0.000  | 0.850  | 23.000  | 0   |
| prse     | 0.000  | 0.000  | 0.075  | 2.000   | 0   |
| guru     | 0.000  | 0.000  | 0.150  | 1.000   | 0   |
| tsca     | 0.000  | 0.000  | 0.700  | 7.000   | 0   |
| ruf1     | 0.000  | 0.000  | 0.025  | 1.000   | 0   |
| vaan     | 0.000  | 0.000  | 0.850  | 15.000  | 0   |
| arnu     | 0.000  | 0.000  | 0.050  | 1.000   | 0   |
| fram     | 0.000  | 0.000  | 0.025  | 1.000   | 0   |
| maca     | 0.000  | 0.000  | 21.675 | 140.000 | 0   |
| mevi     | 0.000  | 0.000  | 1.150  | 14.000  | 0   |
| mire     | 0.000  | 0.000  | 8.675  | 133.000 | 0   |
| smra     | 0.000  | 0.000  | 0.475  | 10.000  | 0   |
| cape     | 0.000  | 0.000  | 1.625  | 27.000  | 0   |
| jute     | 0.000  | 0.000  | 0.075  | 3.000   | 0   |
| dasp     | 0.000  | 0.000  | 0.225  | 3.000   | 0   |
| depu     | 0.000  | 0.000  | 0.850  | 19.000  | 0   |
| drsp     | 0.000  | 0.000  | 0.025  | 1.000   | 0   |
| deob     | 0.000  | 0.000  | 0.400  | 8.000   | 0   |

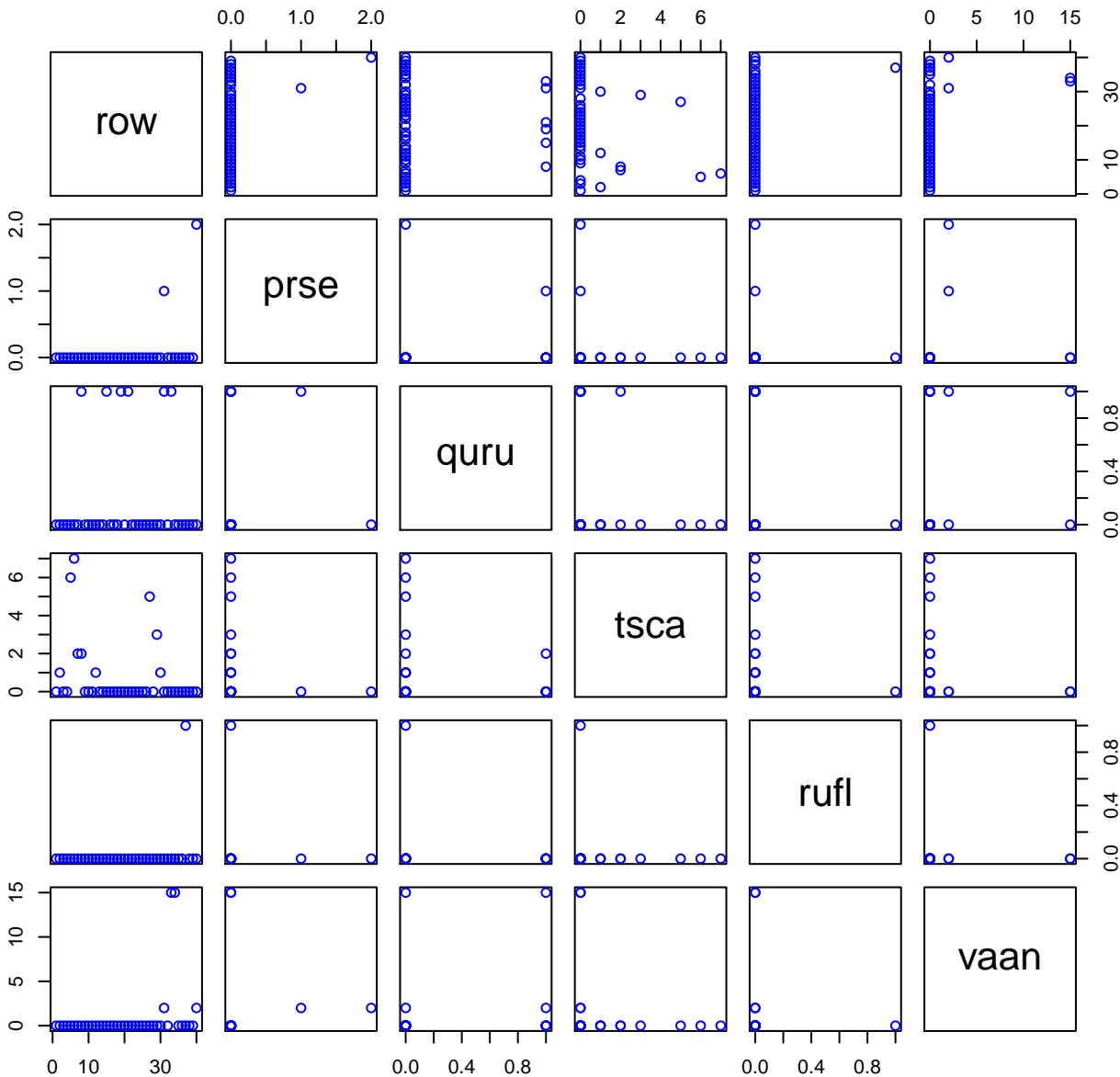
# HF105-02 Plot 1



# HF105-02 Plot 2

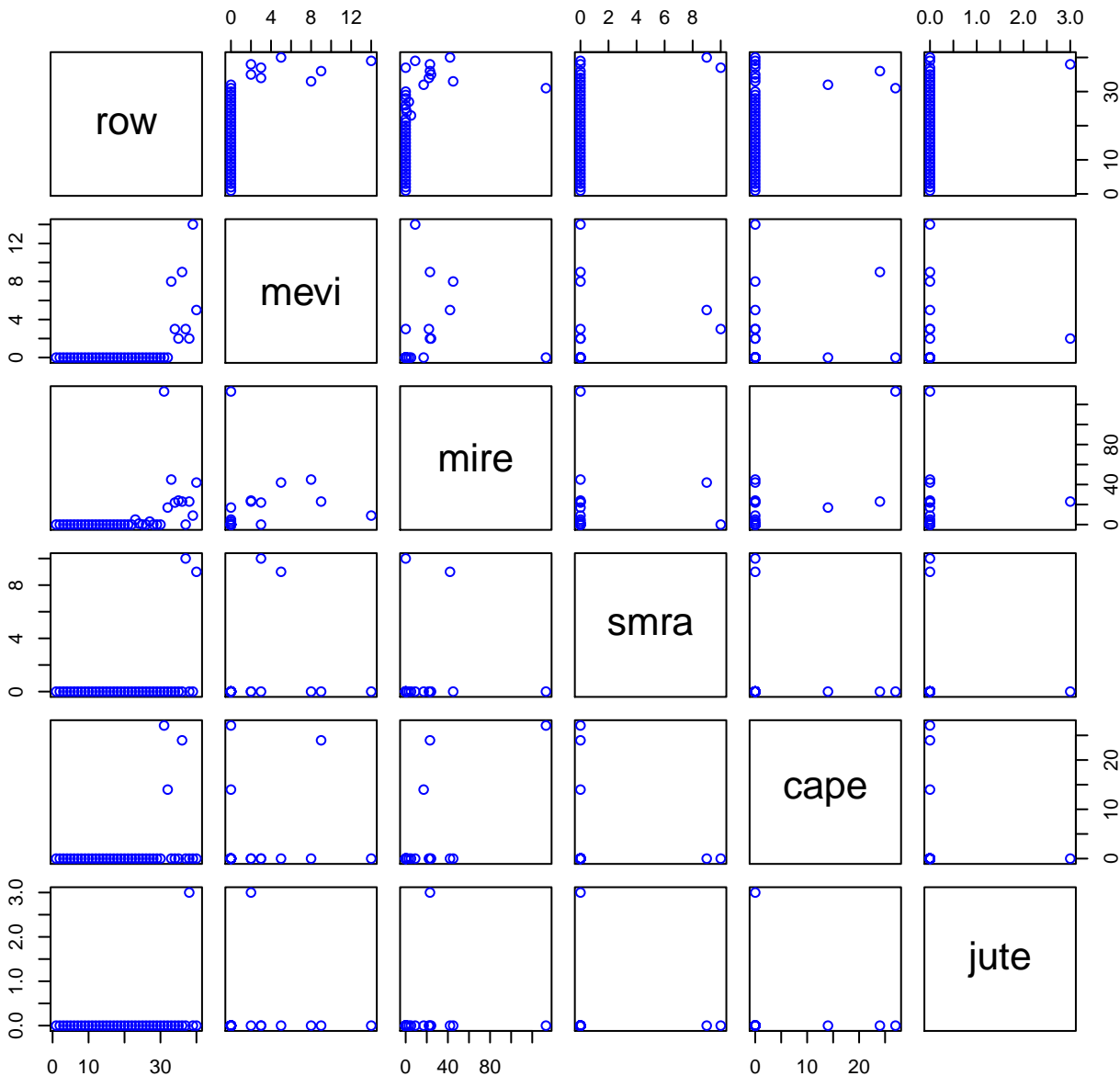


# HF105-02 Plot 3





# HF105-02 Plot 5





# HF105-02 Plot 6

