Harvard Forest Data Archive HF105-01

Data File:

Name = hf105-01-seedbank-2004.csv
Description = seed bank 2004
Rows = 400 Columns = 46
MD5 checksum = 9e406ce28d596082d27e2b412fa6e2bf

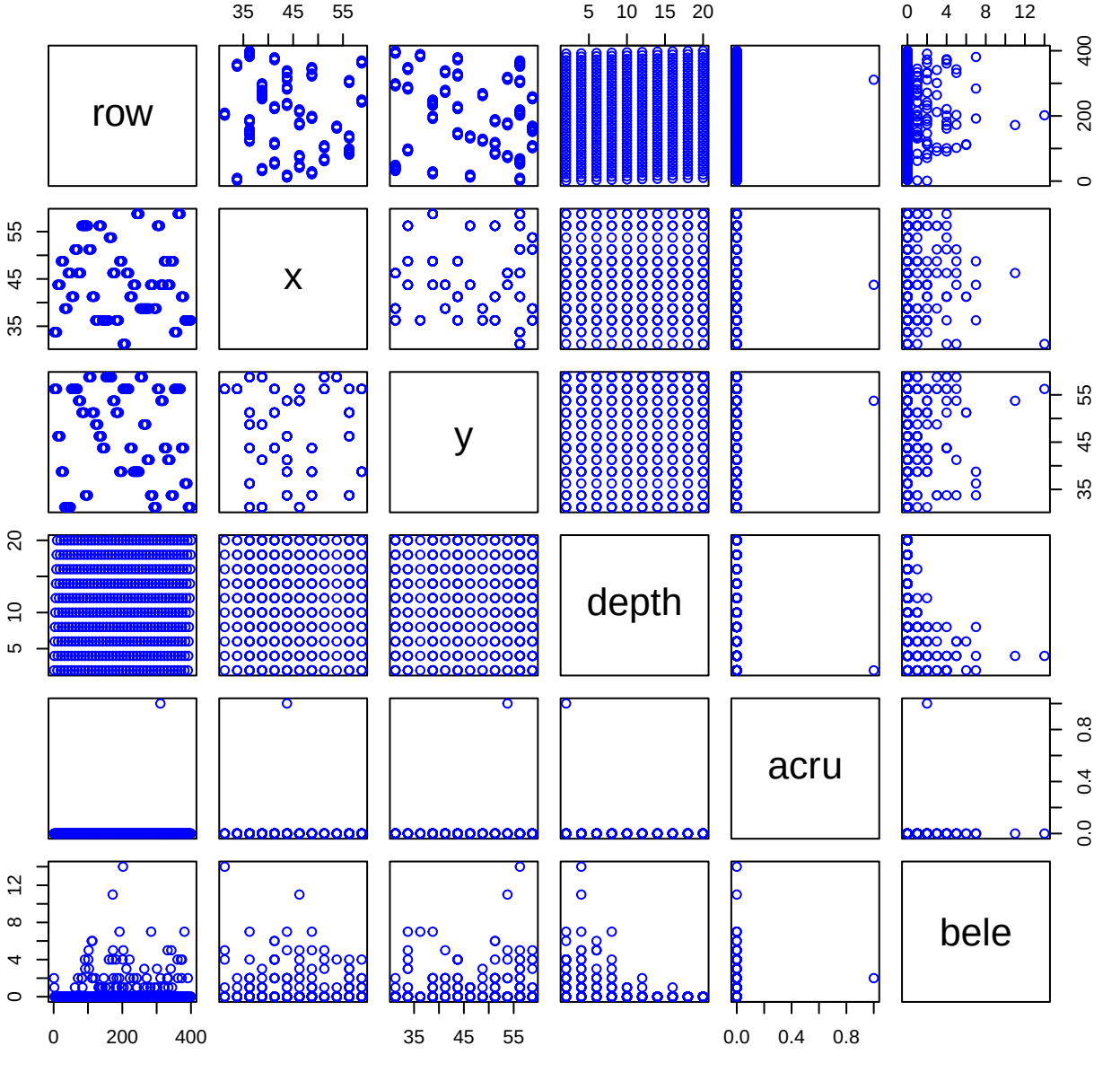
Variables:

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)
depth = depth (in 2 cm increments) of the subsample taken from the
   soil sample for germination trials. Each 20-cm deep sample was divided
   into 10 2-cm thick samples. Value given is the lower depth of the
   subsample. (centimeter)
acru = number of seedlings that emerged in each 2-cm-thick sample
   (number)
bele = number of seedlings that emerged in each 2-cm-thick sample
   (number)
beni = number of seedlings that emerged in each 2-cm-thick sample
   (number)
bepa = number of seedlings that emerged in each 2-cm-thick sample
   (number)
prse = number of seedlings that emerged in each 2-cm-thick sample
   (number)
tsca = number of seedlings that emerged in each 2-cm-thick sample
   (number)
rhty = number of seedlings that emerged in each 2-cm-thick sample
   (number)
rhgl = number of seedlings that emerged in each 2-cm-thick sample
   (number)
rufl = number of seedlings that emerged in each 2-cm-thick sample
   (number)
rusp = number of seedlings that emerged in each 2-cm-thick sample
   (number)
rubus = number of seedlings that emerged in each 2-cm-thick sample
   (number)
sp = number of seedlings that emerged in each 2-cm-thick sample
   (number)
gapr = number of seedlings that emerged in each 2-cm-thick sample
   (number)
mire = number of seedlings that emerged in each 2-cm-thick sample
   (number)
vaan = number of seedlings that emerged in each 2-cm-thick sample
   (number)
amar = number of seedlings that emerged in each 2-cm-thick sample
   (number)

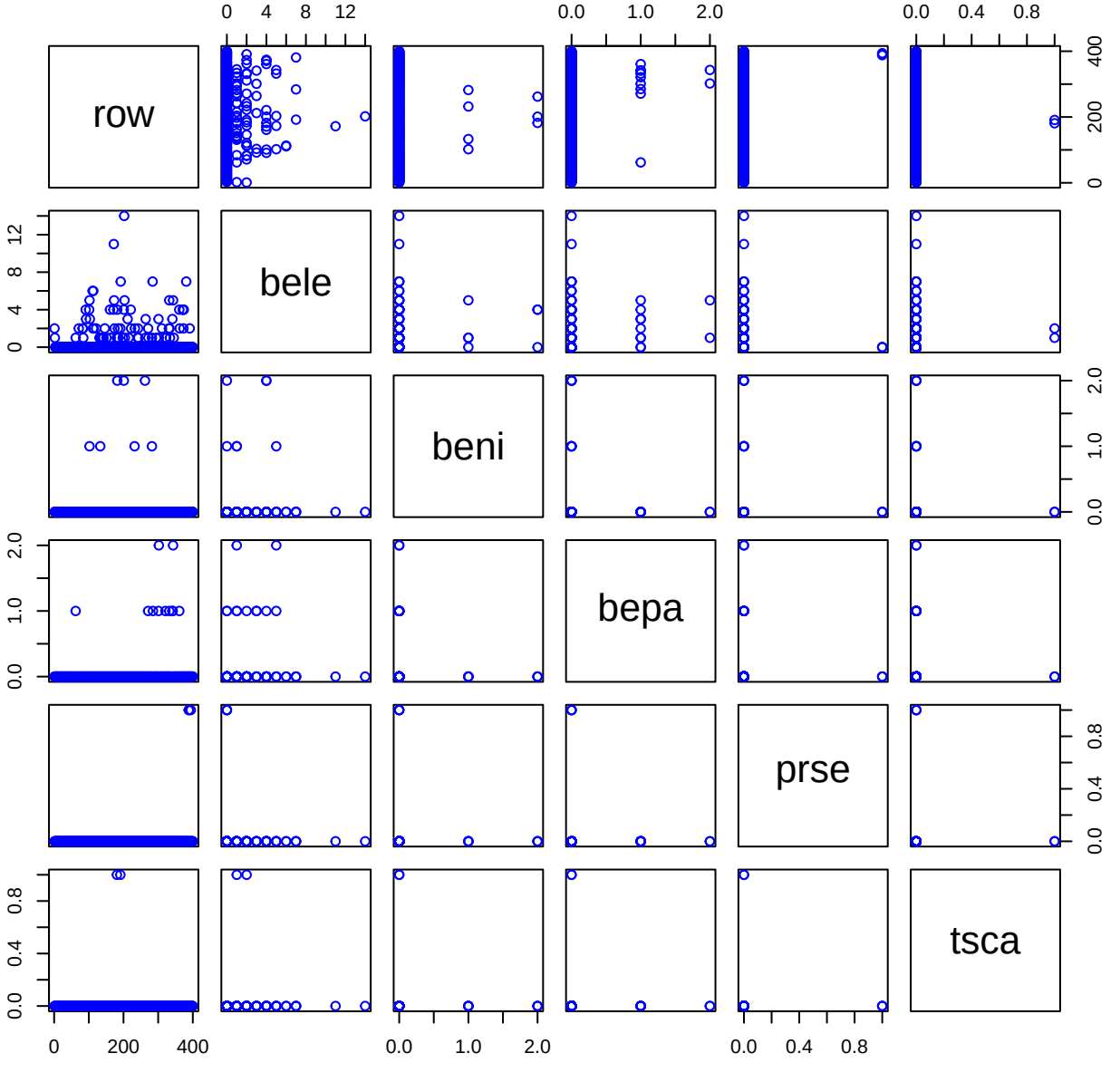
heca = number of seedlings that emerged in each 2−cm−thick sample (number)
hyca = number of seedlings that emerged in each 2−cm−thick sample (number)
hype = number of seedlings that emerged in each 2−cm−thick sample (number)
loin = number of seedlings that emerged in each 2−cm−thick sample (number)
lyci = number of seedlings that emerged in each 2−cm−thick sample (number)
lyqu = number of seedlings that emerged in each 2−cm−thick sample (number)
veth = number of seedlings that emerged in each 2−cm−thick sample (number)
maca = number of seedlings that emerged in each 2−cm−thick sample (number)
move = number of seedlings that emerged in each 2−cm−thick sample (number)
viso = number of seedlings that emerged in each 2−cm−thick sample (number)
caat = number of seedlings that emerged in each 2−cm−thick sample (number)
cade = number of seedlings that emerged in each 2−cm−thick sample (number)
cala = number of seedlings that emerged in each 2−cm−thick sample (number)
cape = number of seedlings that emerged in each 2−cm−thick sample (number)
casp = number of seedlings that emerged in each 2−cm−thick sample (number)
rhal = number of seedlings that emerged in each 2−cm−thick sample (number)
sccy = number of seedlings that emerged in each 2−cm−thick sample (number)
juca = number of seedlings that emerged in each 2−cm−thick sample (number)
jute = number of seedlings that emerged in each 2−cm−thick sample (number)
didi = number of seedlings that emerged in each 2−cm−thick sample (number)
feru = number of seedlings that emerged in each 2−cm−thick sample (number)
pacl = number of seedlings that emerged in each 2−cm−thick sample (number)
pala = number of seedlings that emerged in each 2−cm−thick sample (number)
poan = number of seedlings that emerged in each 2−cm−thick sample (number)
poaceae = number of seedlings that emerged in each 2−cm−thick sample (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
depth	2.000	11.000	11.000	20.000	0
acru	0.000	0.000	0.002	1.000	0
bele	0.000	0.000	0.540	14.000	0
beni	0.000	0.000	0.025	2.000	0
bepa	0.000	0.000	0.032	2.000	0
prse	0.000	0.000	0.007	1.000	0
tsca	0.000	0.000	0.005	1.000	0
rhty	0.000	0.000	0.015	1.000	0
rhgl	0.000	0.000	0.015	2.000	0
rufl	0.000	0.000	0.010	1.000	0
rusp	0.000	0.000	0.180	7.000	0
rubus	0.000	0.000	0.190	7.000	0
sp	0.000	0.000	0.002	1.000	0
gapr	0.000	0.000	0.002	1.000	0
mire	0.000	0.000	0.015	1.000	0
vaan	0.000	0.000	0.002	1.000	0
amar	0.000	0.000	0.010	1.000	0
heca	0.000	0.000	0.015	3.000	0
hyca	0.000	0.000	0.005	1.000	0
hype	0.000	0.000	0.002	1.000	0
loin	0.000	0.000	0.018	2.000	0
lyci	0.000	0.000	0.007	1.000	0
lyqu	0.000	0.000	0.037	1.000	0
veth	0.000	0.000	0.007	1.000	0
maca	0.000	0.000	0.030	3.000	0
move	0.000	0.000	0.007	1.000	0
viso	0.000	0.000	0.007	1.000	0
caat	0.000	0.000	0.007	1.000	0
cade	0.000	0.000	0.002	1.000	0
cala	0.000	0.000	0.037	2.000	0
cape	0.000	0.000	0.185	4.000	0
casp	0.000	0.000	0.012	1.000	0
rhal	0.000	0.000	0.010	2.000	0
sccy	0.000	0.000	0.022	2.000	0
juca	0.000	0.000	0.002	1.000	0
jute	0.000	0.000	0.032	1.000	0
didi	0.000	0.000	0.005	1.000	0
feru	0.000	0.000	0.005	1.000	0
pacl	0.000	0.000	0.002	1.000	0
pala	0.000	0.000	0.030	3.000	0
poan	0.000	0.000	0.090	3.000	0
poaceae	0.000	0.000	0.020	2.000	0

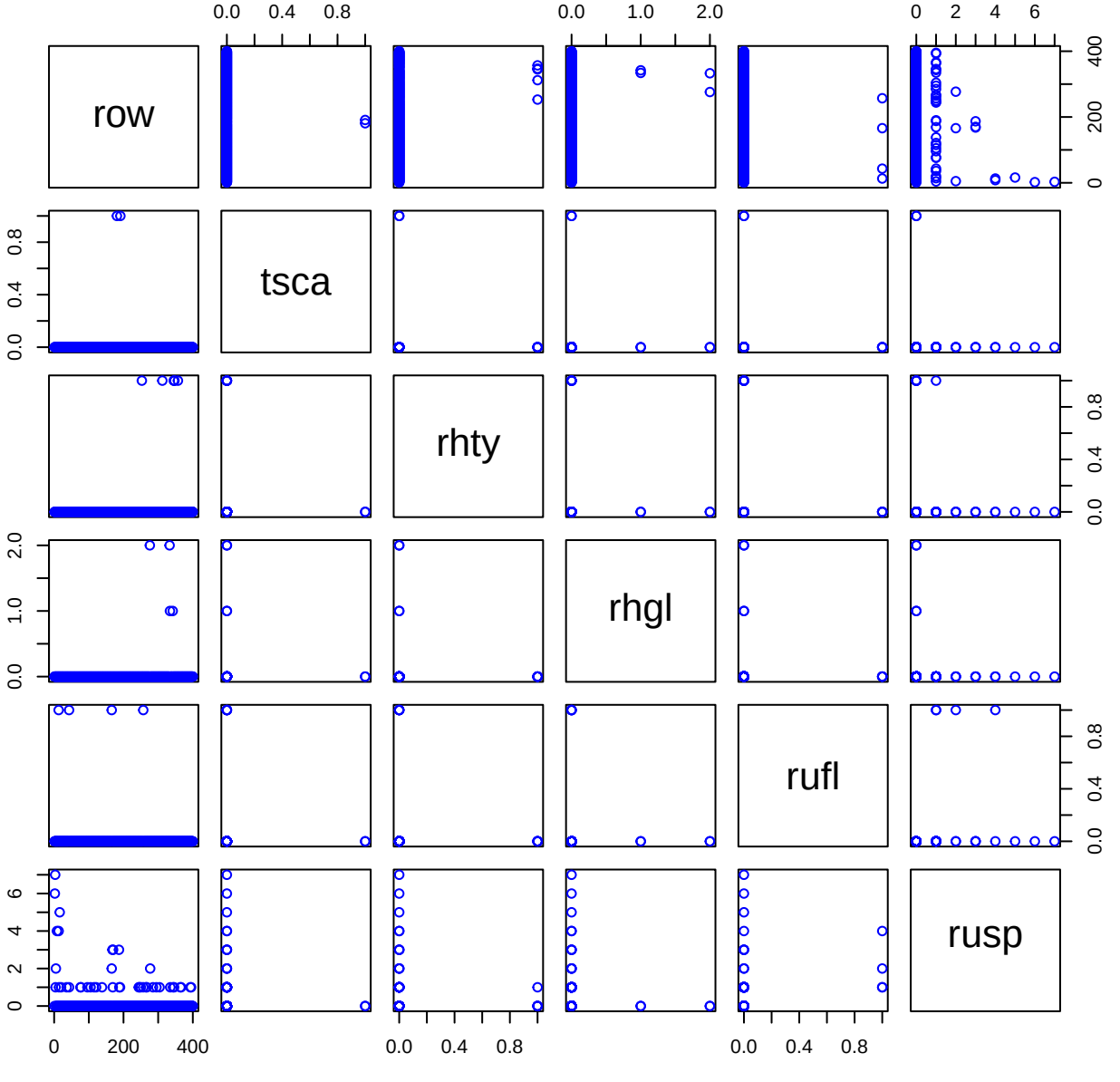
HF105−01 Plot 1



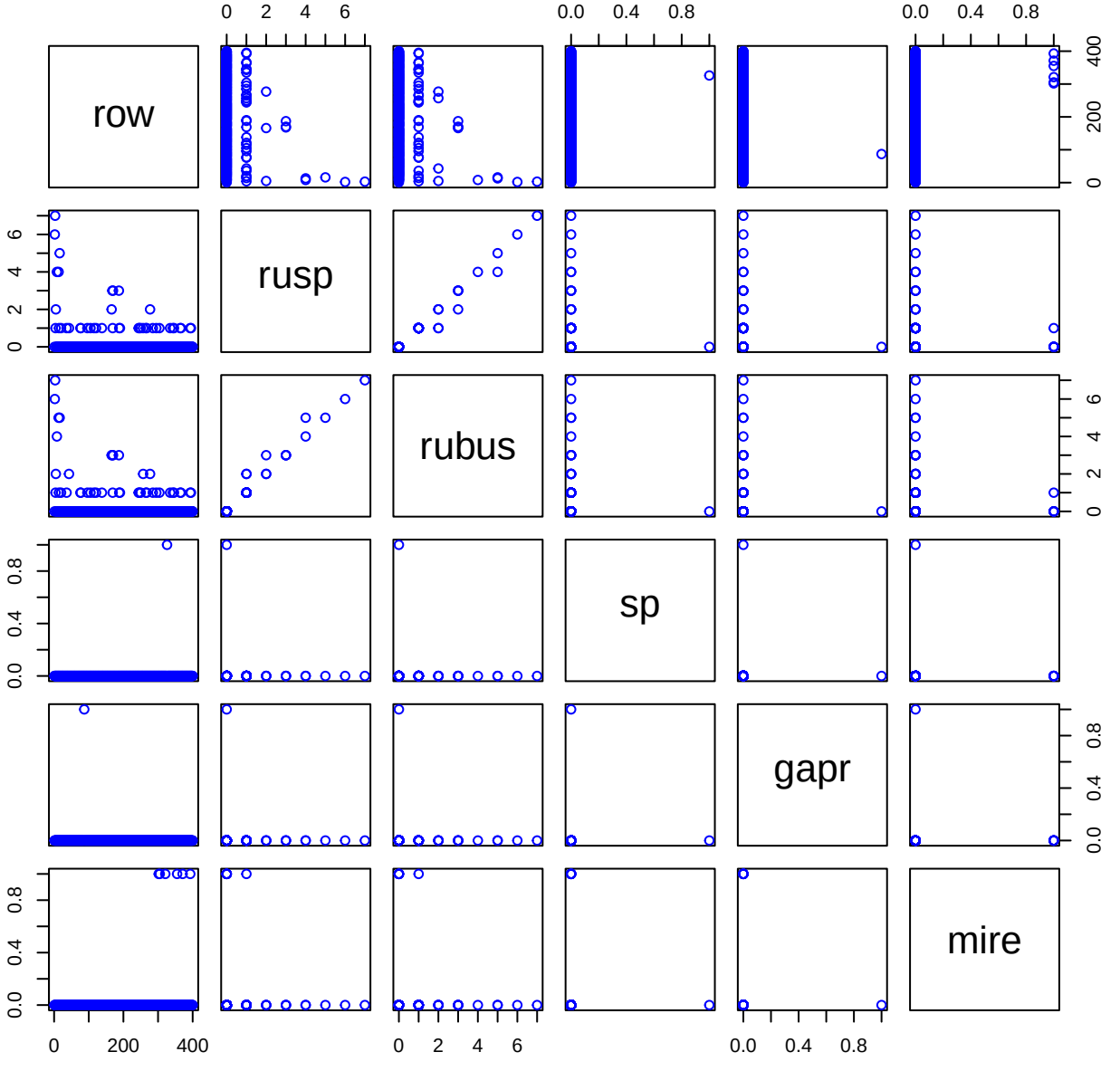
HF105−01 Plot 2



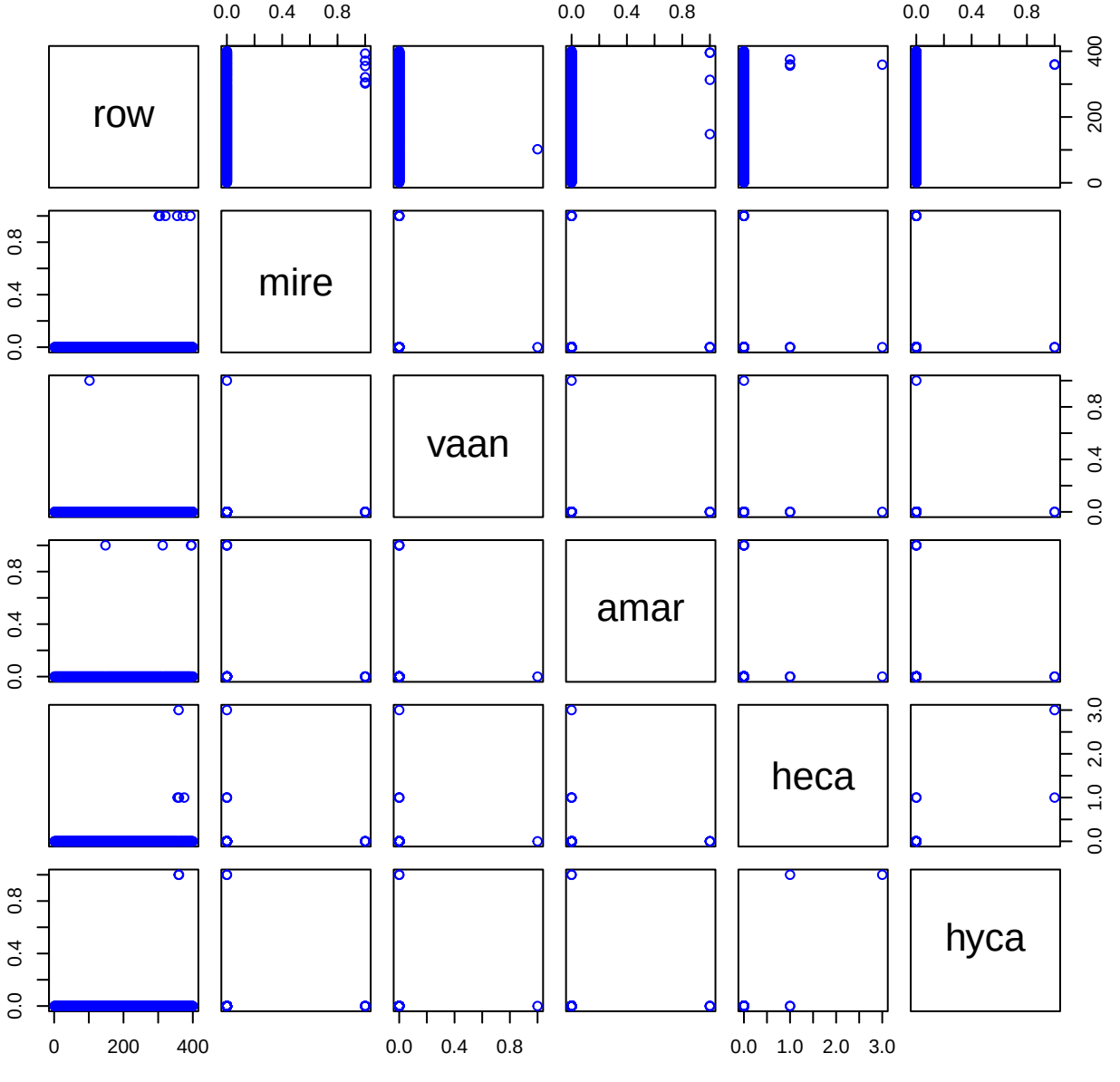
HF105−01 Plot 3



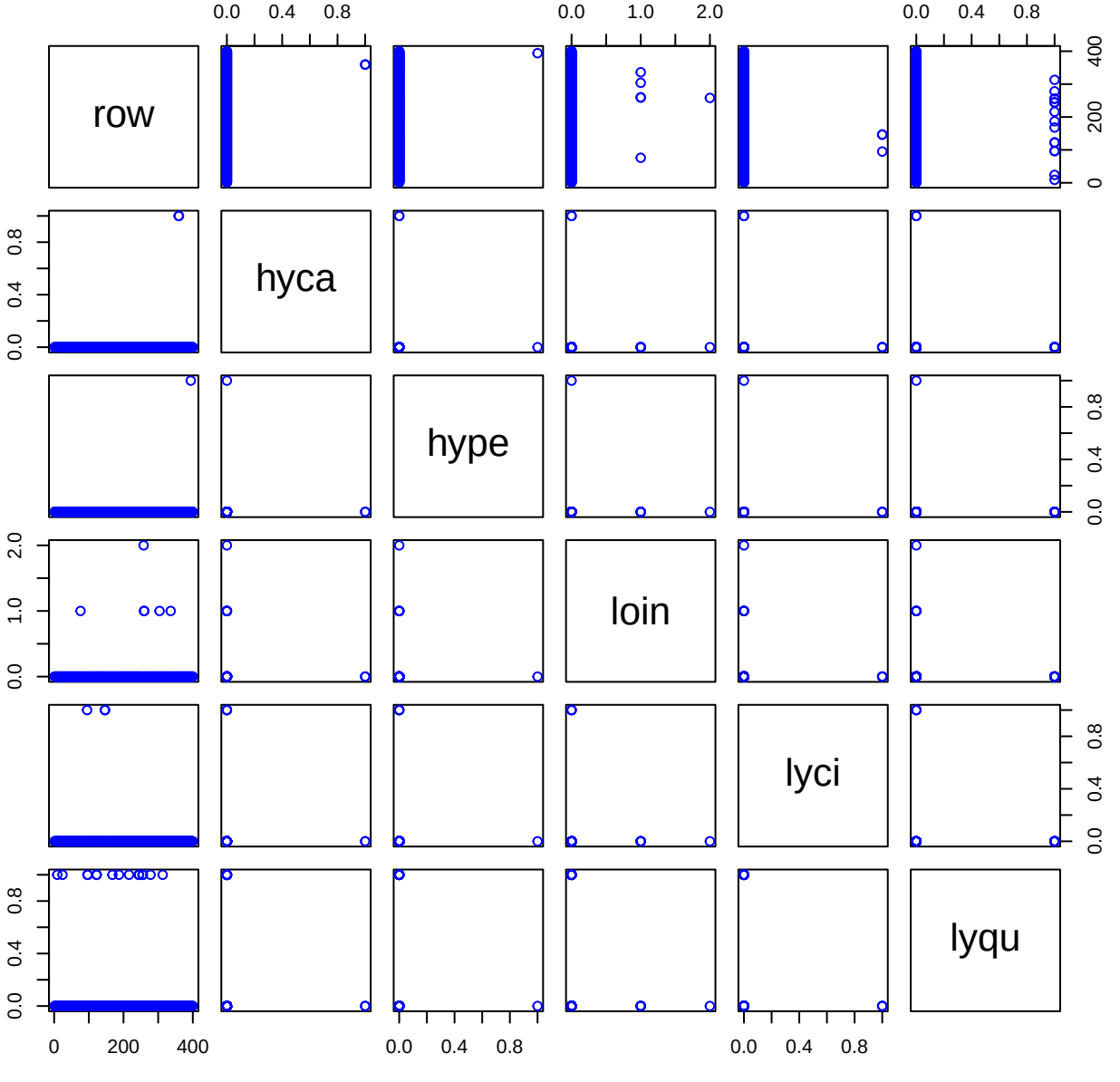
HF105−01 Plot 4



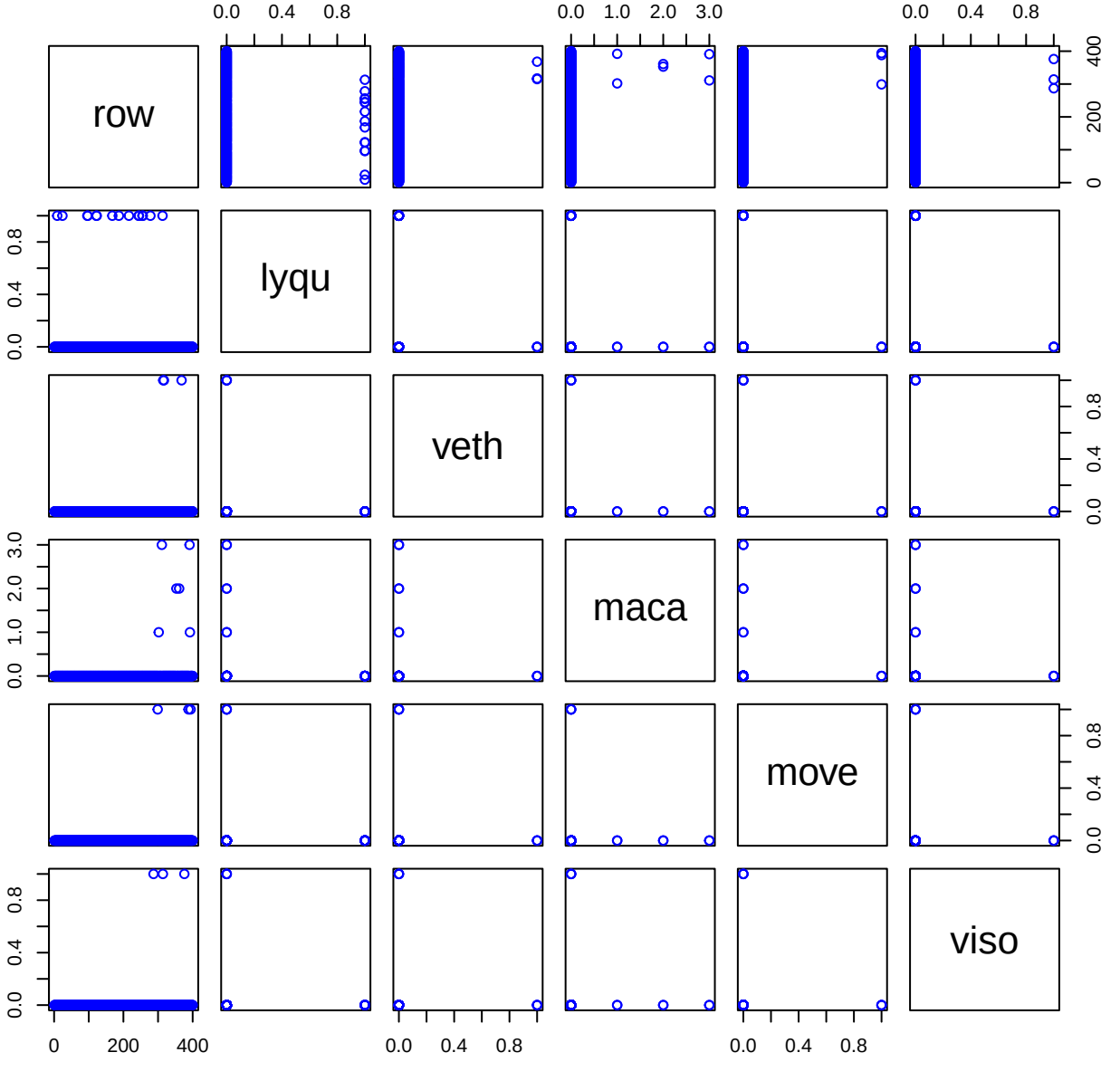
HF105−01 Plot 5



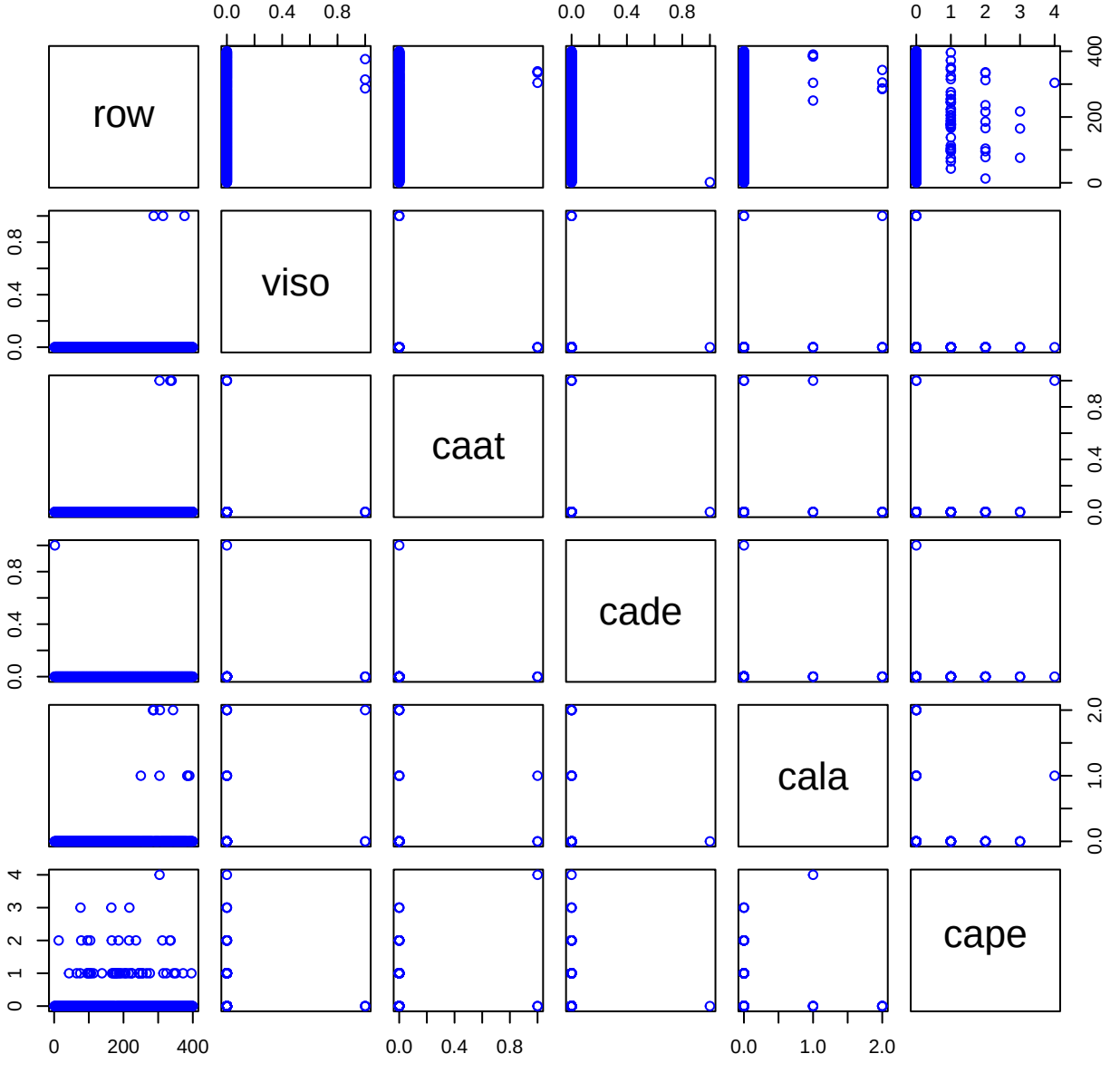
HF105−01 Plot 6



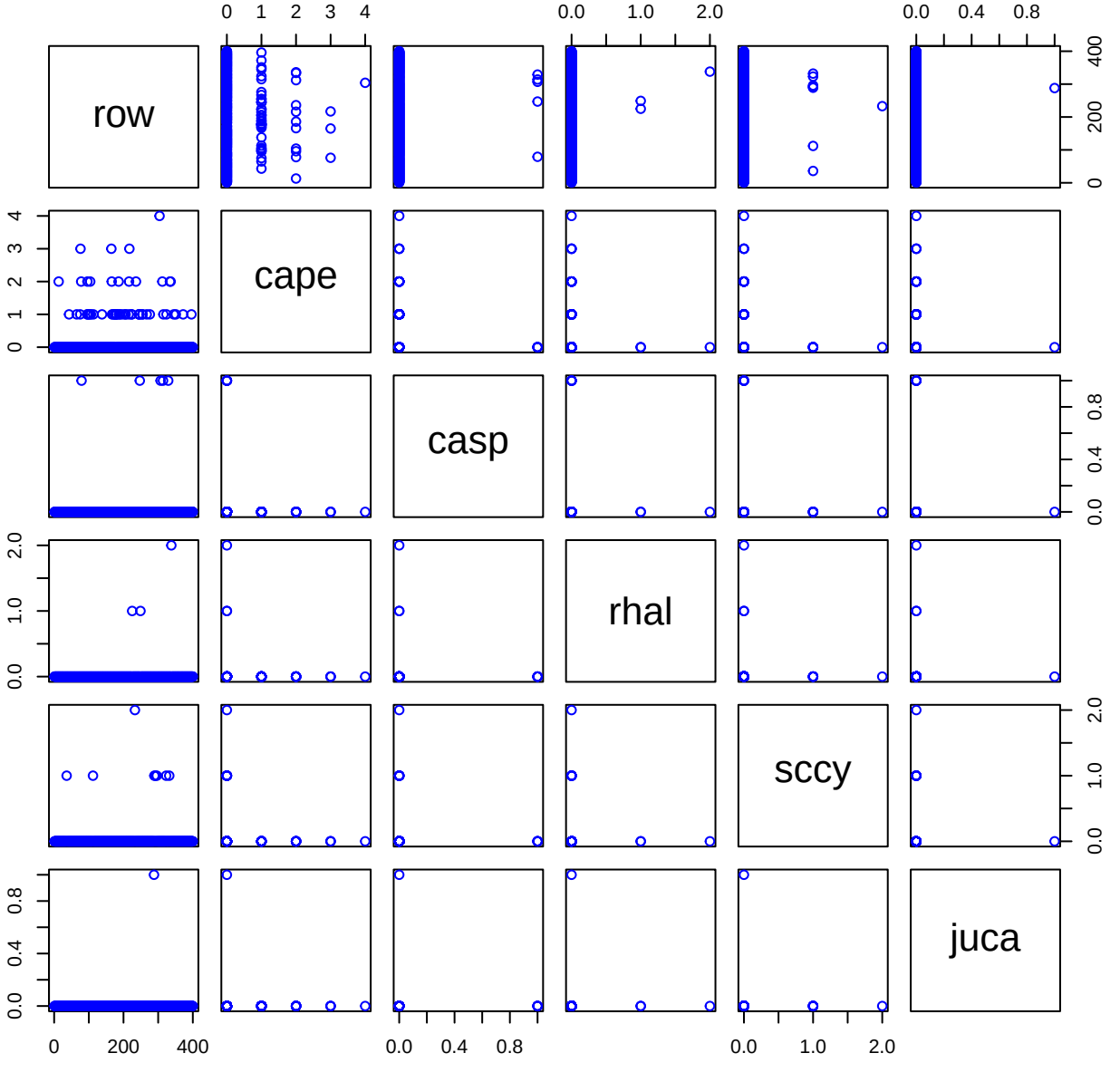
HF105−01 Plot 7



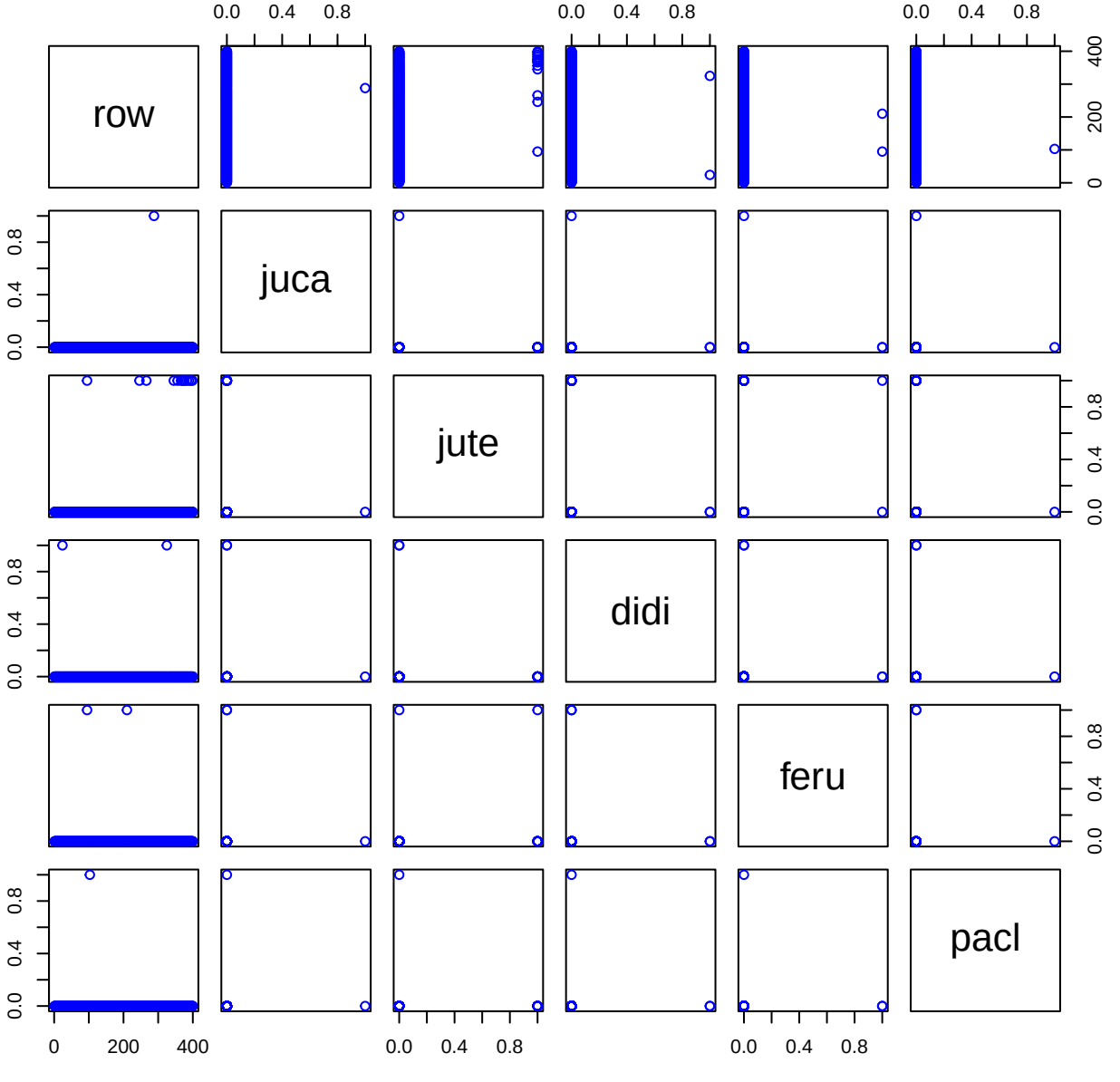
HF105−01 Plot 8

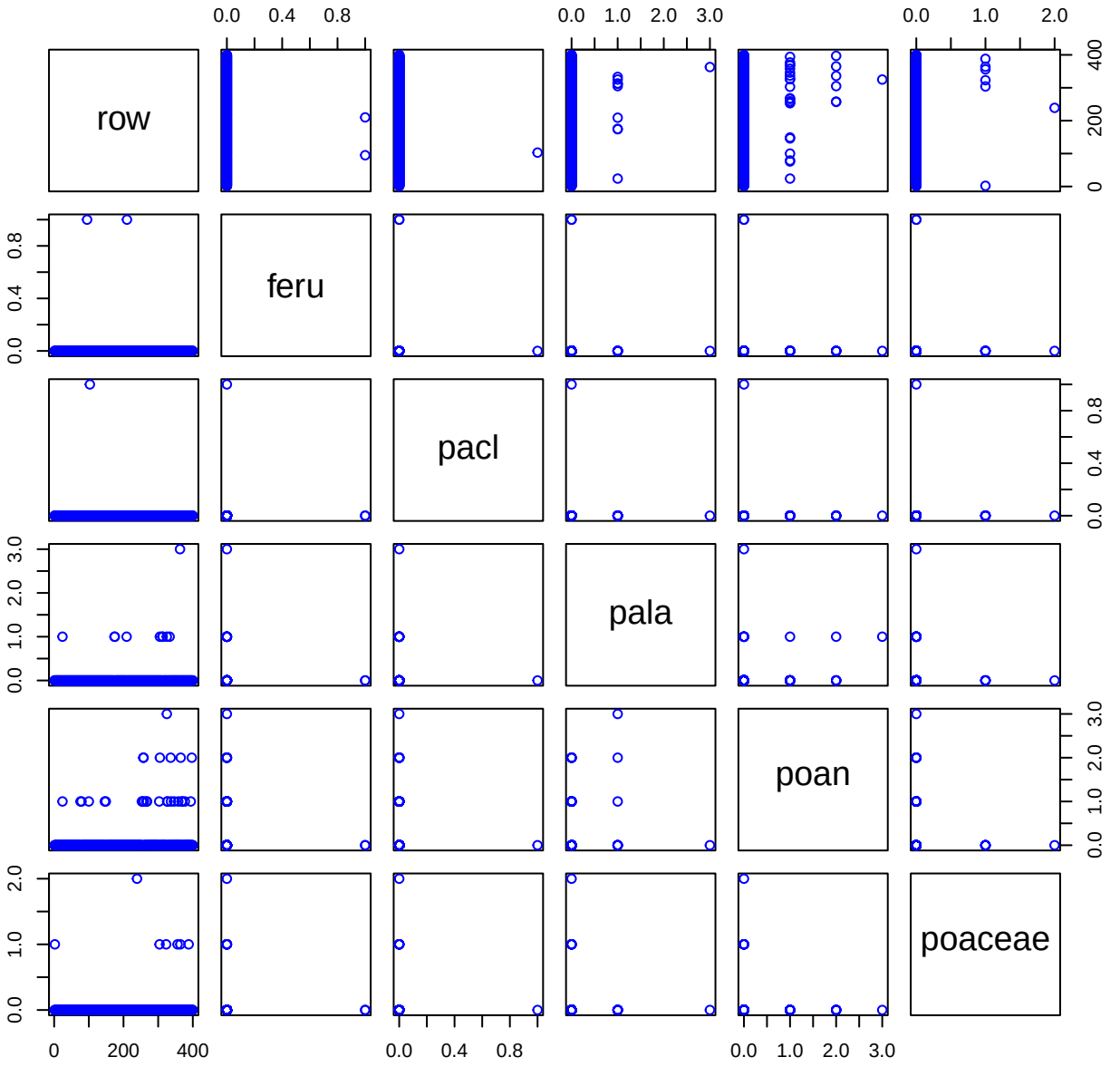


HF105−01 Plot 9



HF105−01 Plot 10




HF105−01 Plot 11
row
feru
pacl
pala
poan
poaceae
0.0
0.4
0.8
0.0
1.0
2.0
3.0
0.0
1.0
2.0
0
200
400
0.0
0.4
0.8
0.0
1.0
2.0
3.0
0
200
400