

Harvard Forest Data Archive HF105-04

Data File:

Name = hf105-04-understory-2010.csv
Description = understory vegetation 2010
Rows = 40 Columns = 50
MD5 checksum = bc8cb2cf8ea494385e5da2509e3b0c48

Variables:

date = data from both the May and August field sampling were combined; date shown is 2011-08-15
depu = number of *Dennstaedtia punctilobula* (Michx.) T. Moore seedlings identified in plot surrounding the core (number)
mire = number of *Mitchella repens* L. seedlings identified in plot surrounding the core (number)
dede = number of *Dendrolycopodium dendroideum* (Michx.) A. Haines seedlings identified in plot surrounding the core (number)
maca = number of *Maianthemum canadense* Desf. seedlings identified in plot surrounding the core (number)
acru = number of *Acer rubrum* L. seedlings identified in plot surrounding the core (number)
pist = number of *Pinus strobus* L. seedlings identified in plot surrounding the core (number)
arnu = number of *Aralia nudicaulis* L. seedlings identified in plot surrounding the core (number)
fagr = number of *Fagus grandifolia* Ehrh. seedlings identified in plot surrounding the core (number)
trbo = number of *Trientalis borealis* Raf. ssp. *borealis* seedlings identified in plot surrounding the core (number)
vaan = number of *Vaccinium angustifolium* Ait. seedlings identified in plot surrounding the core (number)
prse = number of *Prunus serotina* Ehrh. var. *serotina* seedlings identified in plot surrounding the core (number)
uvse = number of *Uvularia sessilifolia* L. seedlings identified in plot surrounding the core (number)
moun = number of *Monotropa uniflora* L. seedlings identified in plot surrounding the core (number)
tsca = number of *Tsuga canadensis* (L.) Carr. seedlings identified in plot surrounding the core (number)
hulu = number of *Huperzia lucidula* (Michx.) Trevisan seedlings identified in plot surrounding the core (number)
trer = number of *Trillium erectum* L. seedlings identified in plot surrounding the core (number)
bele = number of *Betula lenta* L. seedlings identified in plot surrounding the core (number)
coco = number of *Corylus cornuta* Marsh. ssp. *cornuta* seedlings identified in plot surrounding the core (number)
cape = number of *Carex pensylvanica* Lam. seedlings identified in plot surrounding the core (number)

guru = number of *Quercus rubra* L. seedlings identified in plot
surrounding the core (number)

vaco = number of *Vaccinium corymbosum* L. seedlings identified in
plot surrounding the core (number)

pham = number of *Phytolacca americana* L. var. *americana* seedlings
identified in plot surrounding the core (number)

rual = number of *Rubus allegheniensis* Porter seedlings identified in
plot surrounding the core (number)

arhi = number of *Aralia hispida* Vent. seedlings identified in plot
surrounding the core (number)

ruhi = number of *Rubus hispidus* L. seedlings identified in plot
surrounding the core (number)

lyqu = number of *Lysimachia quadrifolia* L. seedlings identified in
plot surrounding the core (number)

ruoc = number of *Rubus occidentalis* L. seedlings identified in plot
surrounding the core (number)

beal = number of *Betula alleghaniensis* Brett. seedlings identified
in plot surrounding the core (number)

beth = number of *Berberis thunbergii* DC. seedlings identified in
plot surrounding the core (number)

bepa = number of *Betula papyrifera* Marsh. seedlings identified in
plot surrounding the core (number)

ampsp = number of *Amelanchier* Medik. sp. seedlings identified in plot
surrounding the core (number)

drin = number of *Dryopteris intermedia* (Muhl. ex Willd.) Gray
seedlings identified in plot surrounding the core (number)

gapr = number of *Gaultheria procumbens* L. seedlings identified in
plot surrounding the core (number)

daco = number of *Danthonia compressa* Austin ex Peck seedlings
identified in plot surrounding the core (number)

pano = number of *Parathelypteris noveboracensis* (L.) Ching seedlings
identified in plot surrounding the core (number)

qual = number of *Quercus alba* L. seedlings identified in plot
surrounding the core (number)

drca = number of *Dryopteris carthusiana* (Vill.) H. P. Fuchs
seedlings identified in plot surrounding the core (number)

crsp = number of *Crataegus* L. sp. seedlings identified in plot
surrounding the core (number)

cotr = number of *Coptis trifolia* (L.) Salisb. seedlings identified
in plot surrounding the core (number)

acsa = number of *Acer saccharum* Marsh. seedlings identified in plot
surrounding the core (number)

lycl = number of *Lycopodium clavatum* L. seedlings identified in plot
surrounding the core (number)

mevi = number of *Medeola virginiana* L. seedlings identified in plot
surrounding the core (number)

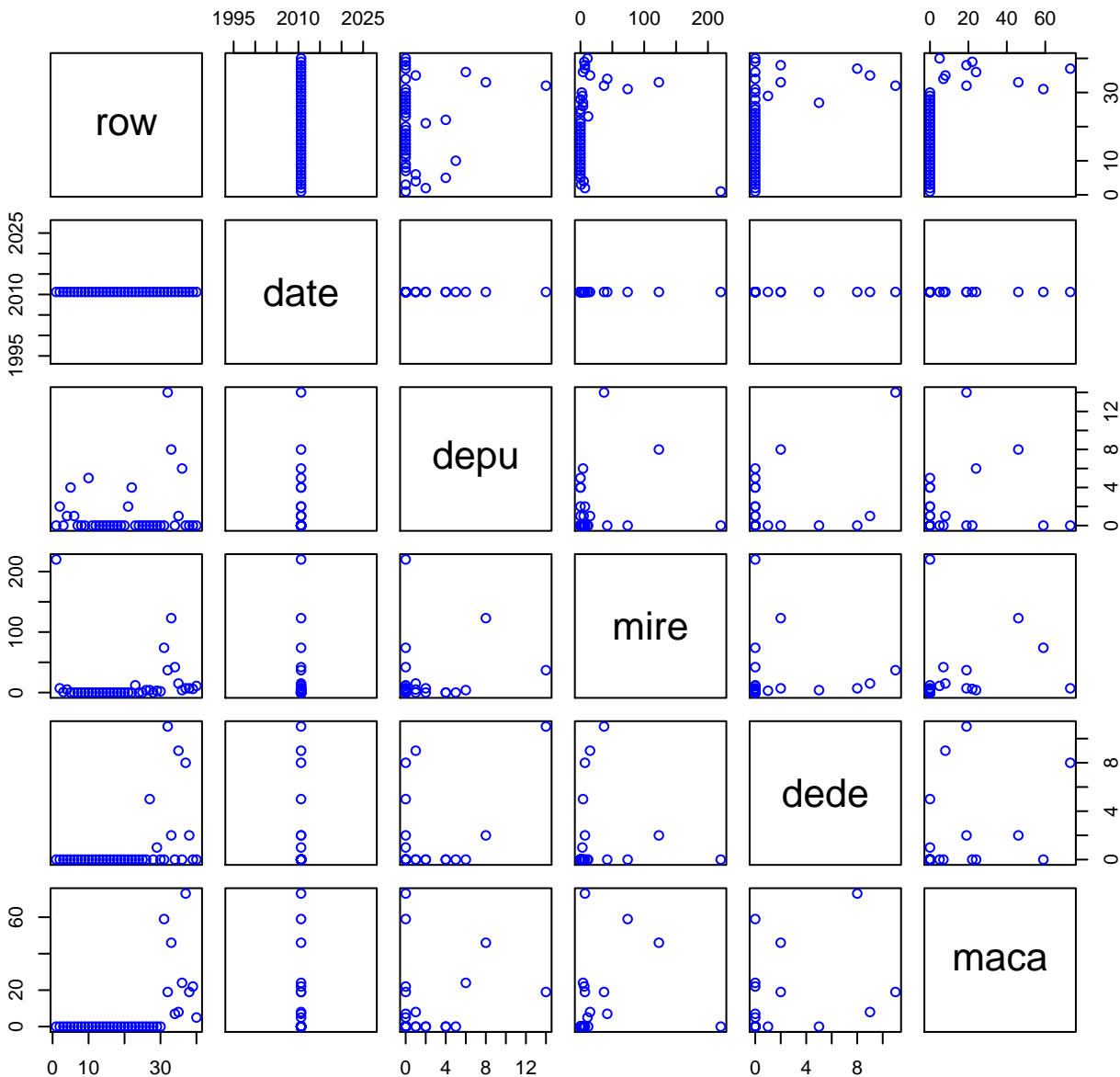
oscl = number of *Osmunda claytoniana* L. seedlings identified in plot
surrounding the core (number)

rufl = number of *Rubus flagellaris* Willd. seedlings identified in
plot surrounding the core (number)

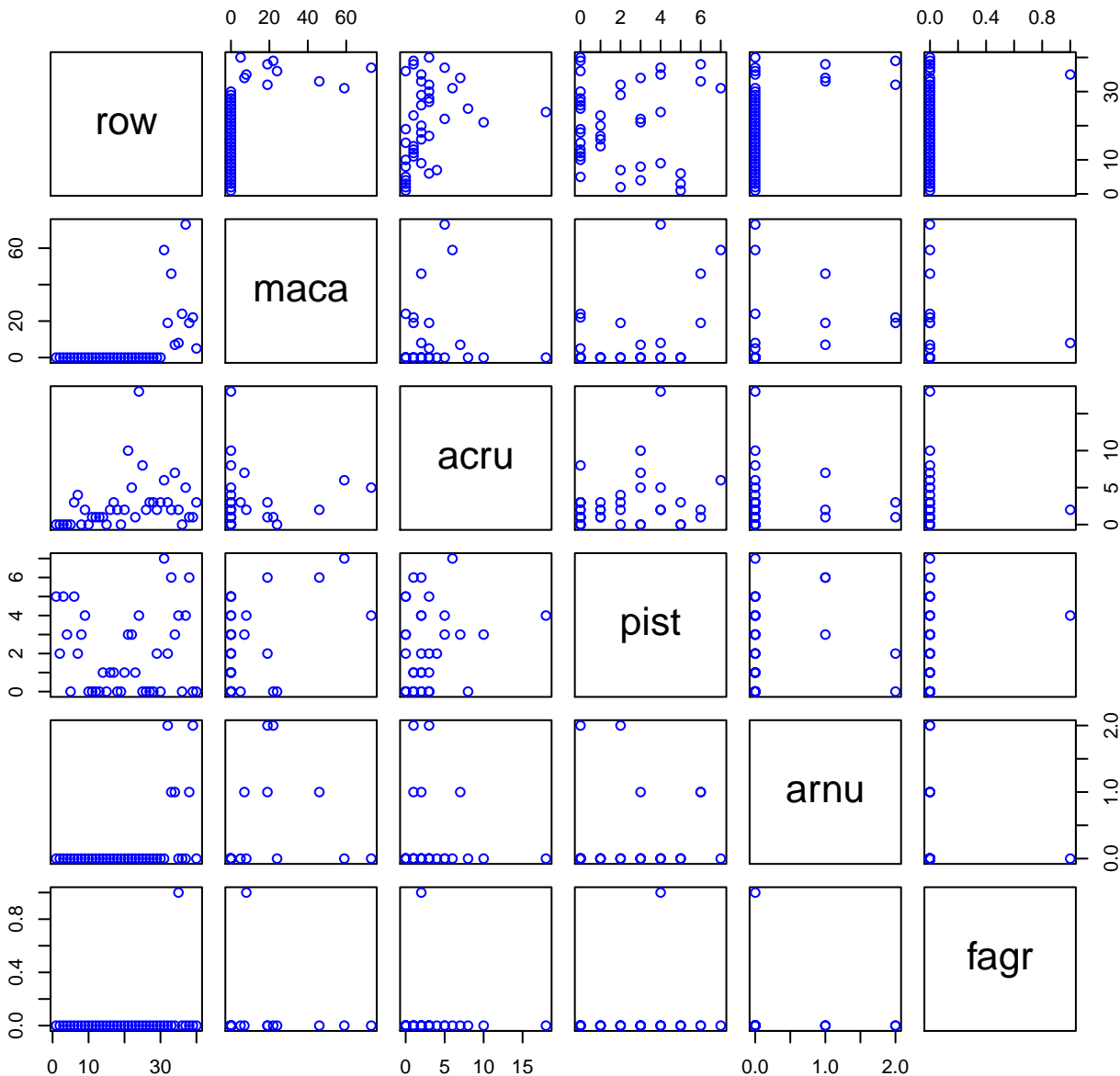
cope = number of *Comptonia peregrina* (L.) Coult. seedlings
identified in plot surrounding the core (number)

Variable	Min	Median	Mean	Max	NAs
date	2010-08-15	2010-08-15	2010-08-15	2010-08-15	0
depu	0.000	0.000	1.200	14.000	0
mire	0.000	0.000	14.600	220.000	0
dede	0.000	0.000	0.950	11.000	0
maca	0.000	0.000	7.050	73.000	0
acru	0.000	2.000	2.675	18.000	0
pist	0.000	1.000	1.950	7.000	0
arnu	0.000	0.000	0.175	2.000	0
fagr	0.000	0.000	0.025	1.000	0
trbo	0.000	0.000	1.100	6.000	0
vaan	0.000	0.000	0.525	5.000	0
prse	0.000	0.000	0.500	7.000	0
uvse	0.000	0.000	0.225	6.000	0
moun	0.000	0.000	0.025	1.000	0
tsca	0.000	0.000	1.025	6.000	0
hulu	0.000	0.000	0.100	3.000	0
trrer	0.000	0.000	0.025	1.000	0
bele	0.000	1.500	3.000	16.000	0
coco	0.000	0.000	0.125	4.000	0
cape	0.000	0.000	2.275	26.000	0
quru	0.000	0.000	0.450	7.000	0
vaco	0.000	0.000	0.625	7.000	0
pham	0.000	0.000	0.025	1.000	0
rual	0.000	0.000	0.700	6.000	0
arhi	0.000	0.000	1.525	12.000	0
ruhi	0.000	0.000	2.875	50.000	0
lyqu	0.000	0.000	0.550	6.000	0
ruoc	0.000	0.000	0.225	5.000	0
beal	0.000	0.000	0.125	2.000	0
beth	0.000	0.000	0.050	1.000	0
bepa	0.000	0.000	0.200	7.000	0
amsp	0.000	0.000	0.025	1.000	0
drin	0.000	0.000	0.125	2.000	0
gapr	0.000	0.000	0.150	5.000	0
daco	0.000	0.000	0.025	1.000	0
pano	0.000	0.000	0.100	4.000	0
qual	0.000	0.000	0.150	6.000	0
drca	0.000	0.000	0.100	4.000	0
crsp	0.000	0.000	0.375	6.000	0
cotr	0.000	0.000	0.150	6.000	0
acsa	0.000	0.000	0.175	7.000	0
lycl	0.000	0.000	0.150	6.000	0
mevi	0.000	0.000	0.300	8.000	0
oscl	0.000	0.000	0.075	3.000	0
ruf1	0.000	0.000	0.825	6.000	0
cope	0.000	0.000	0.100	4.000	0

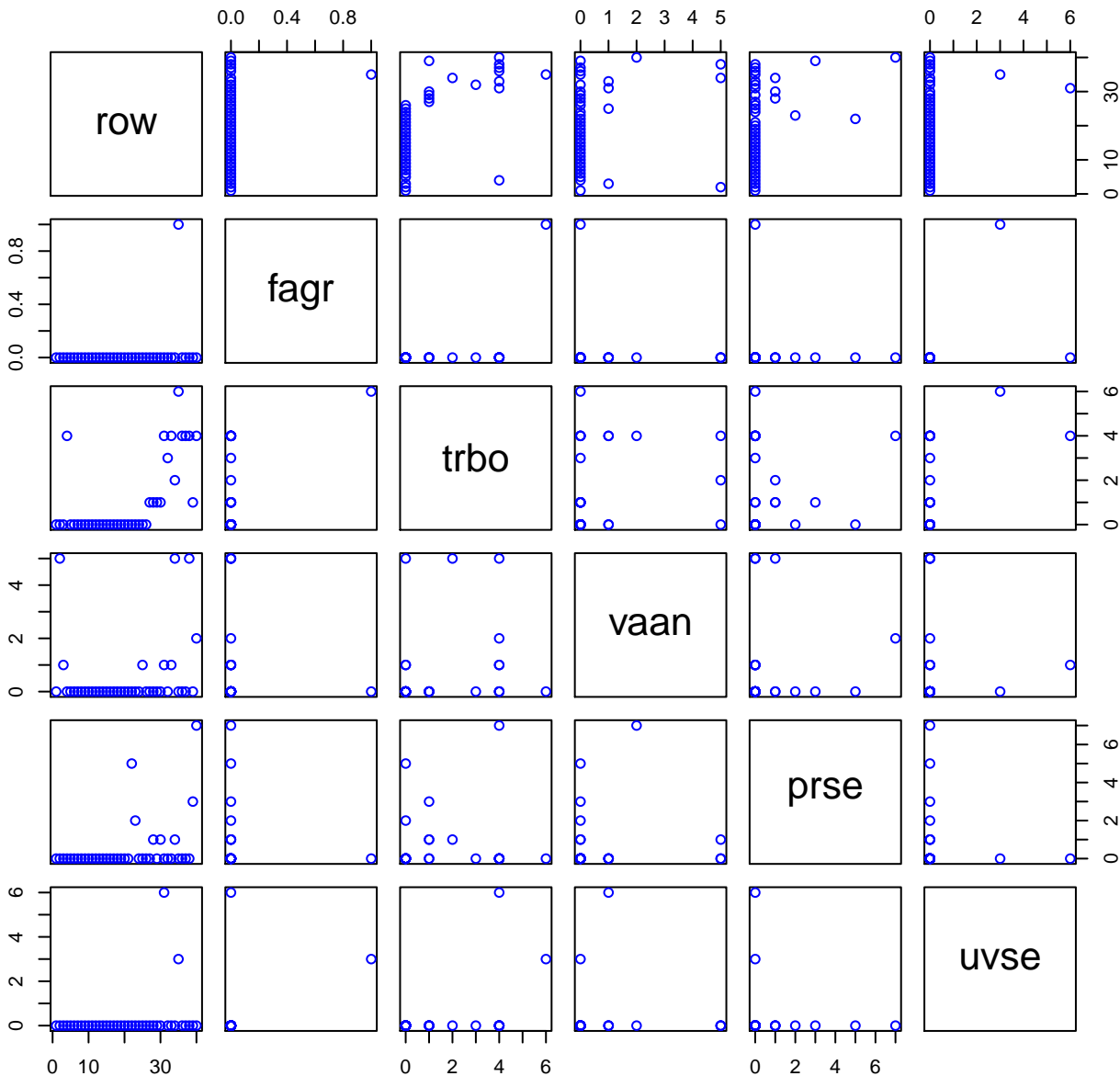
HF105-04 Plot 1



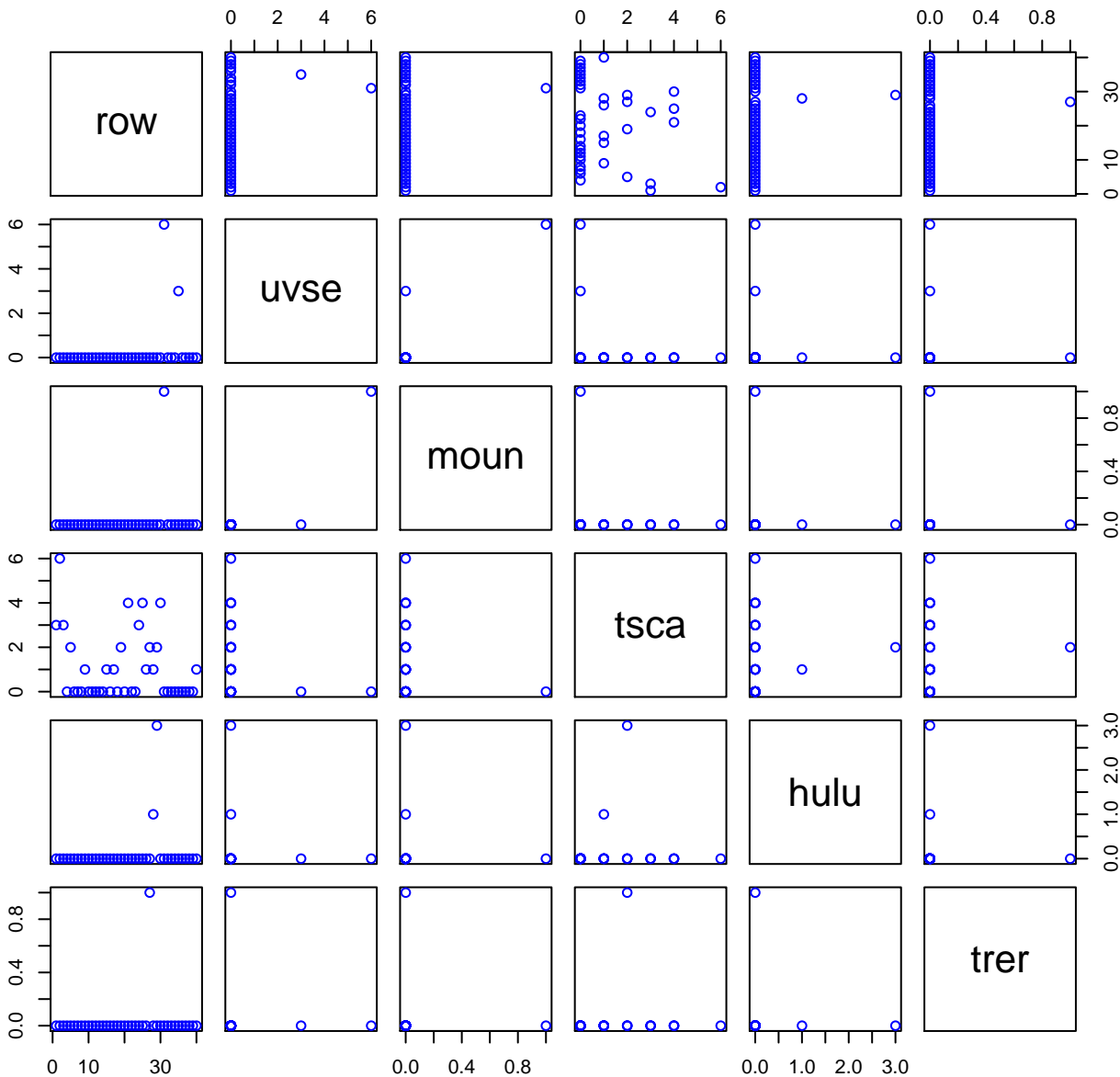
HF105-04 Plot 2



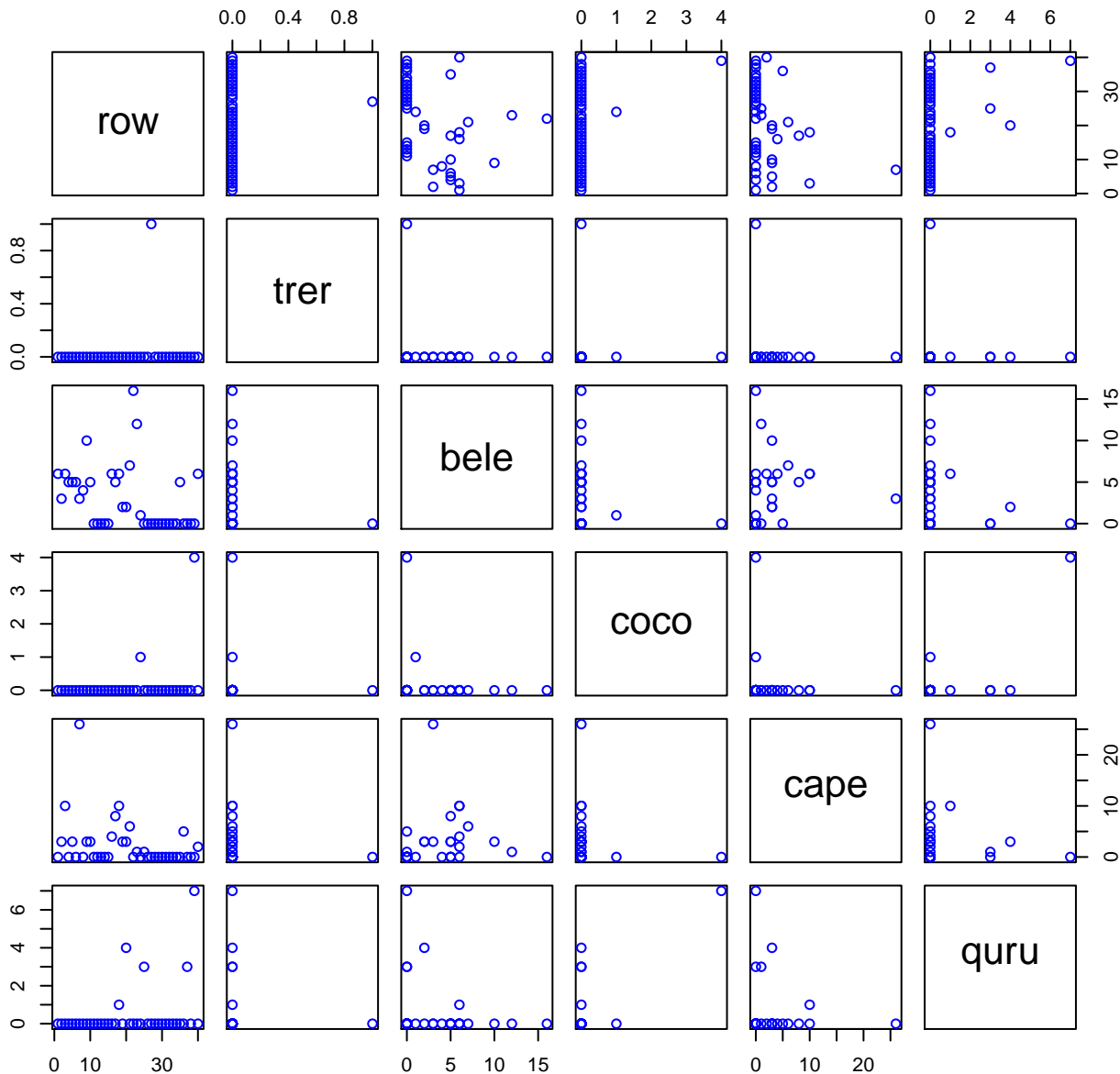
HF105-04 Plot 3



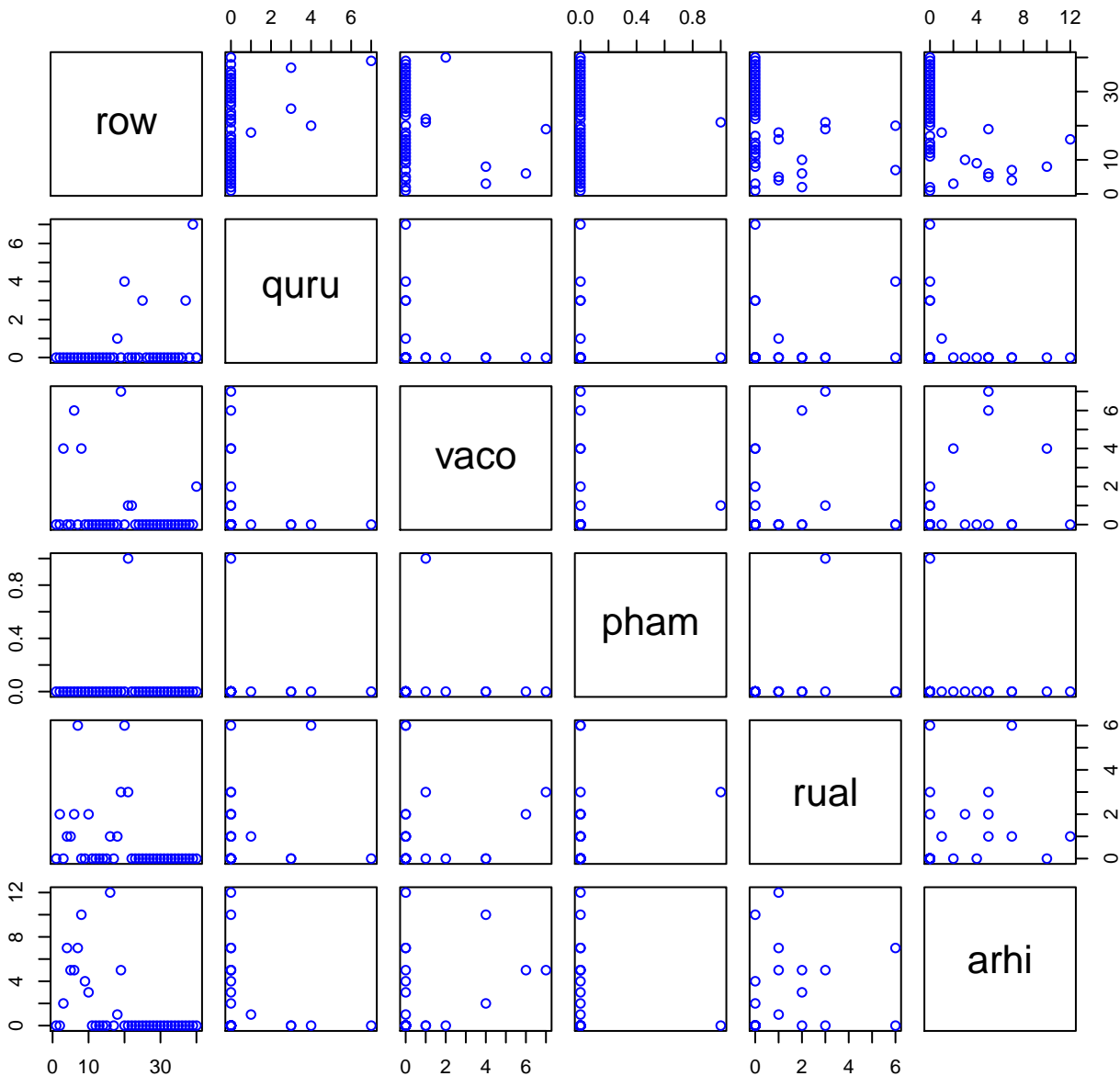
HF105-04 Plot 4



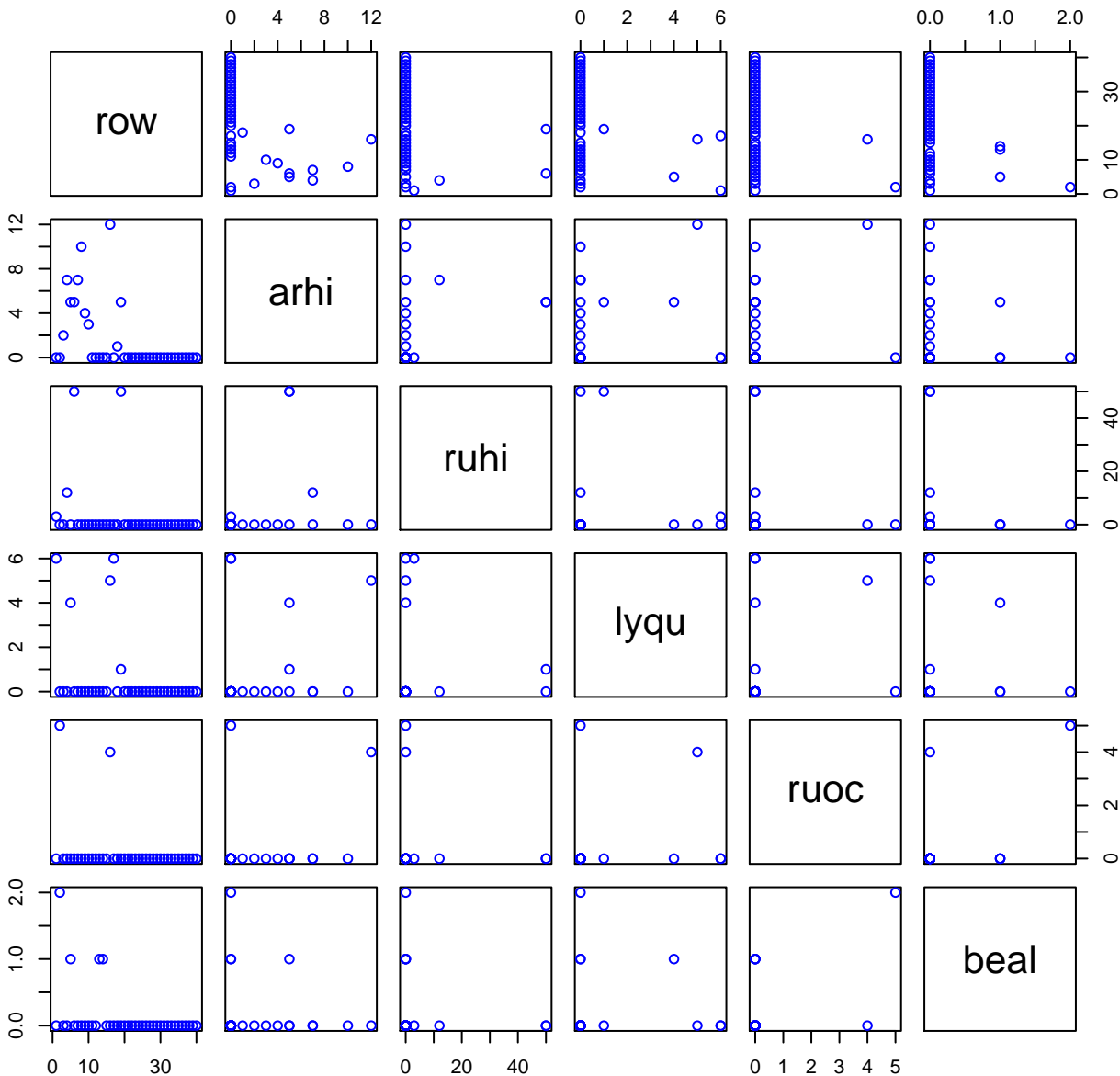
HF105-04 Plot 5



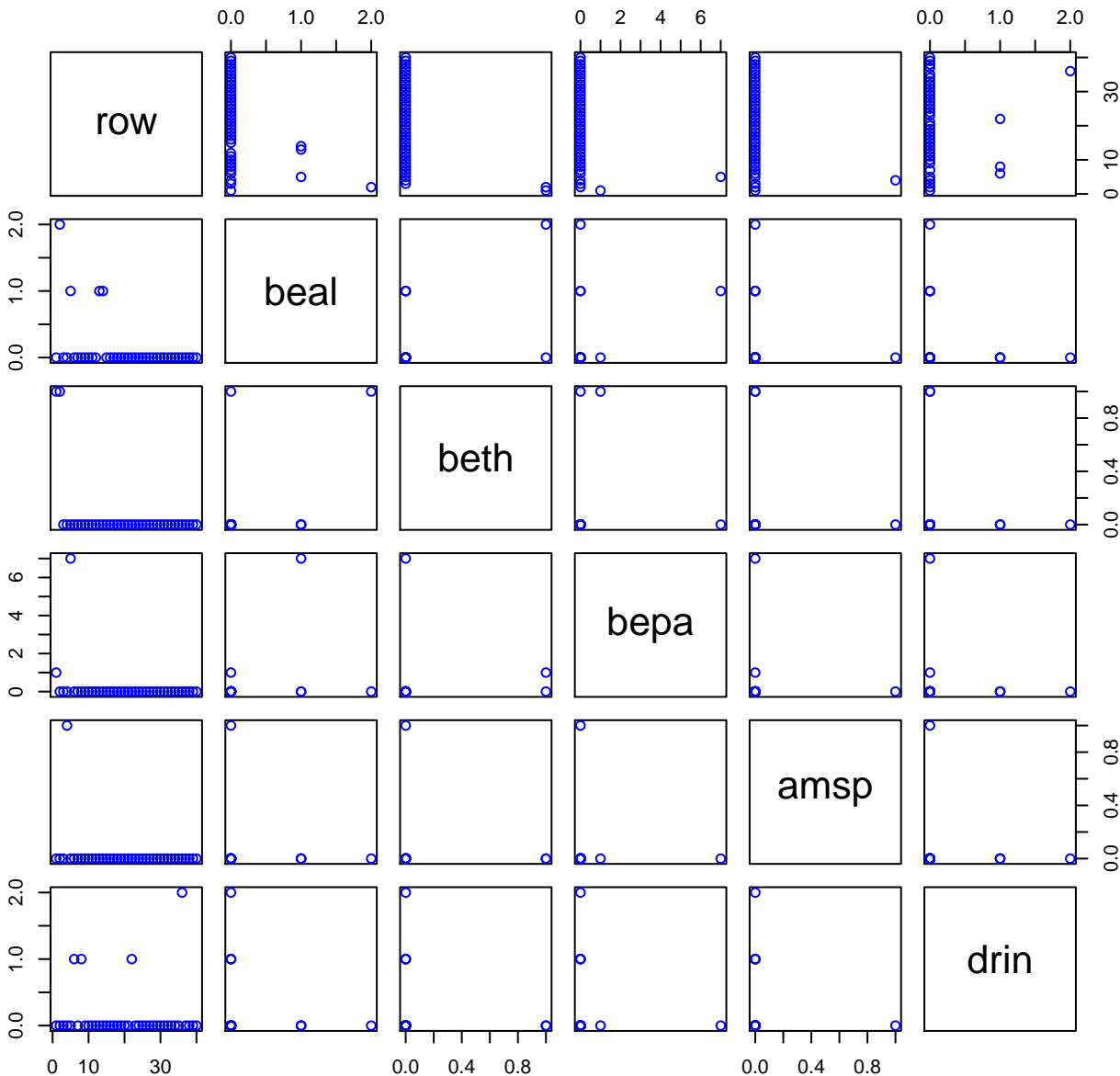
HF105-04 Plot 6



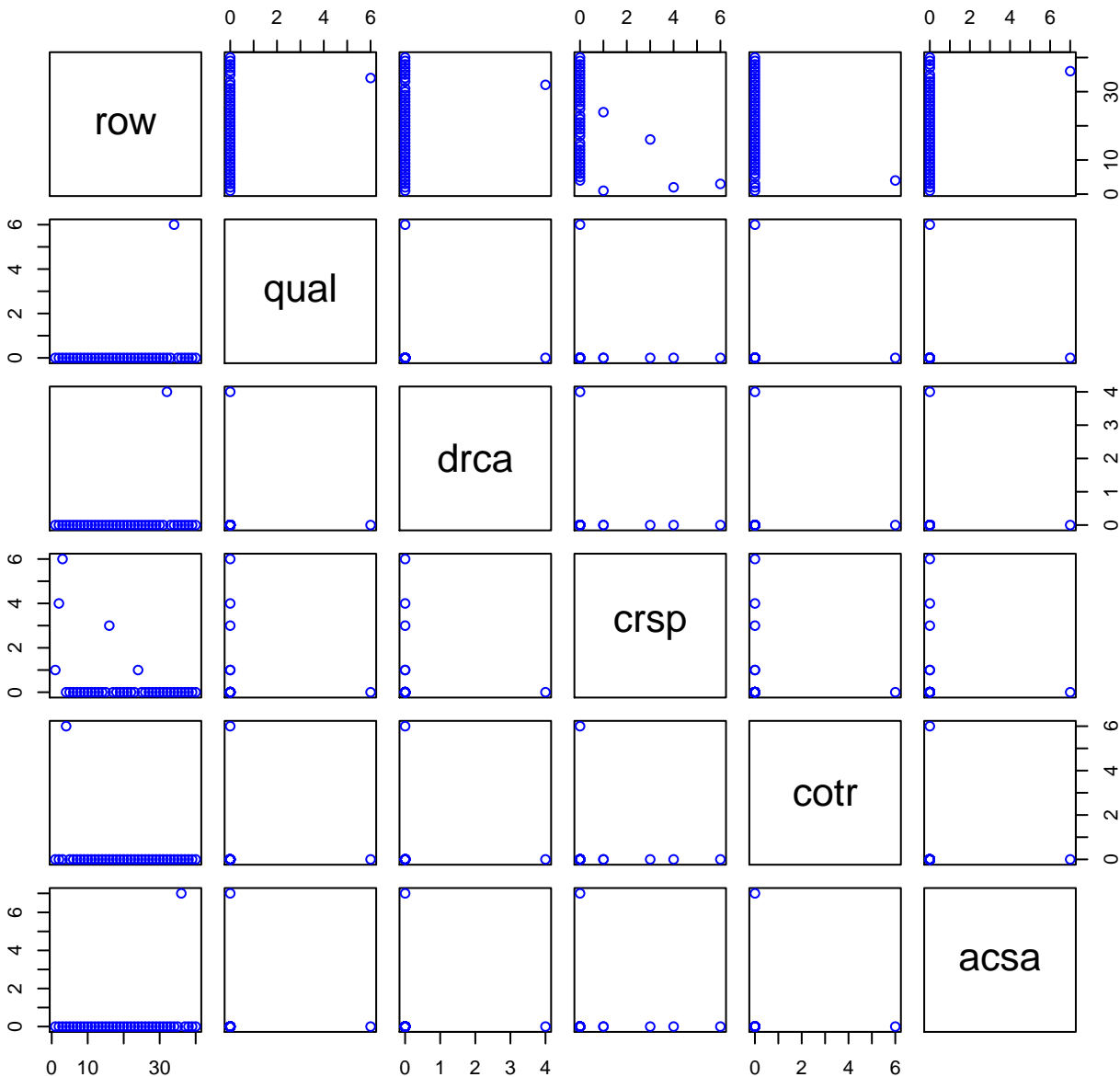
HF105-04 Plot 7



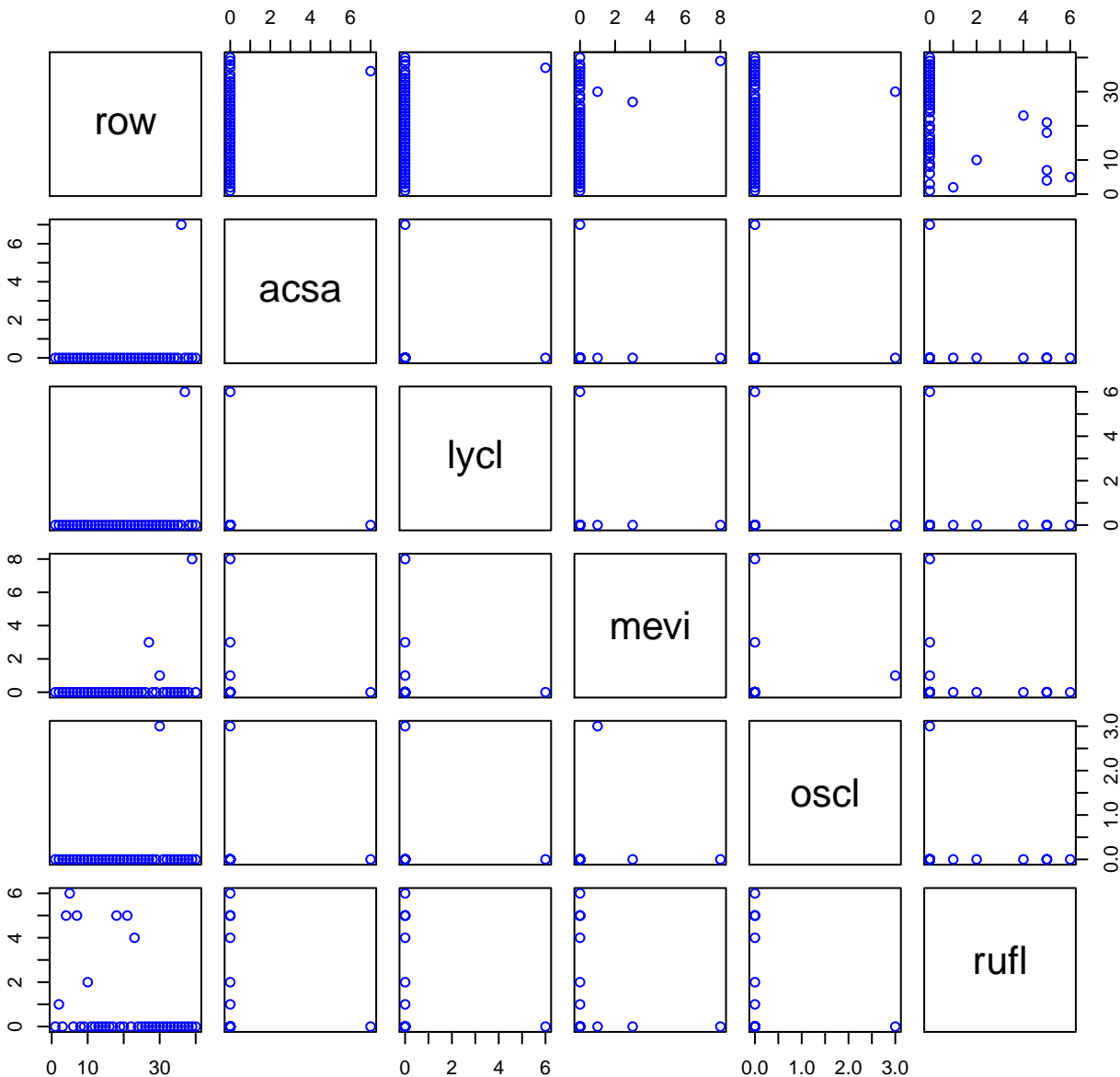
HF105-04 Plot 8



HF105-04 Plot 10



HF105-04 Plot 11



HF105-04 Plot 12

