

2022 Bayer Hooded Sprayer

Maxwel C Oliveira

```
library(tidyverse)

library(drc)

readxl::read_excel("dataset.xlsx") %>%

  janitor::clean_names() %>%

  mutate(rating_dat = as_factor(rating_dat)) %>%

  mutate_if(is_character, as_factor) -> dat

dat %>%

# filter(rating_dat == 14) %>%

  ggplot(aes(x = distance_m, y = rating_avg3plants, color = boom)) +

  facet_grid(rating_dat ~ location) +

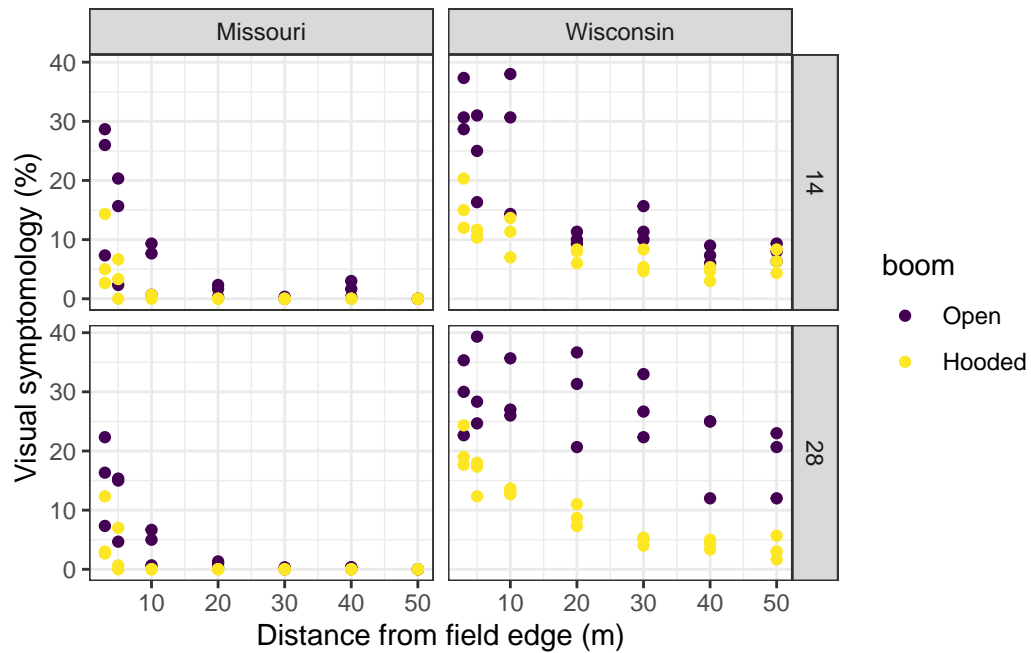
  geom_point() +

  scale_color_viridis_d() +

  theme_bw() +

  labs(y = "Visual symptomology (%)",

       x = "Distance from field edge (m)")
```



Wisconsin

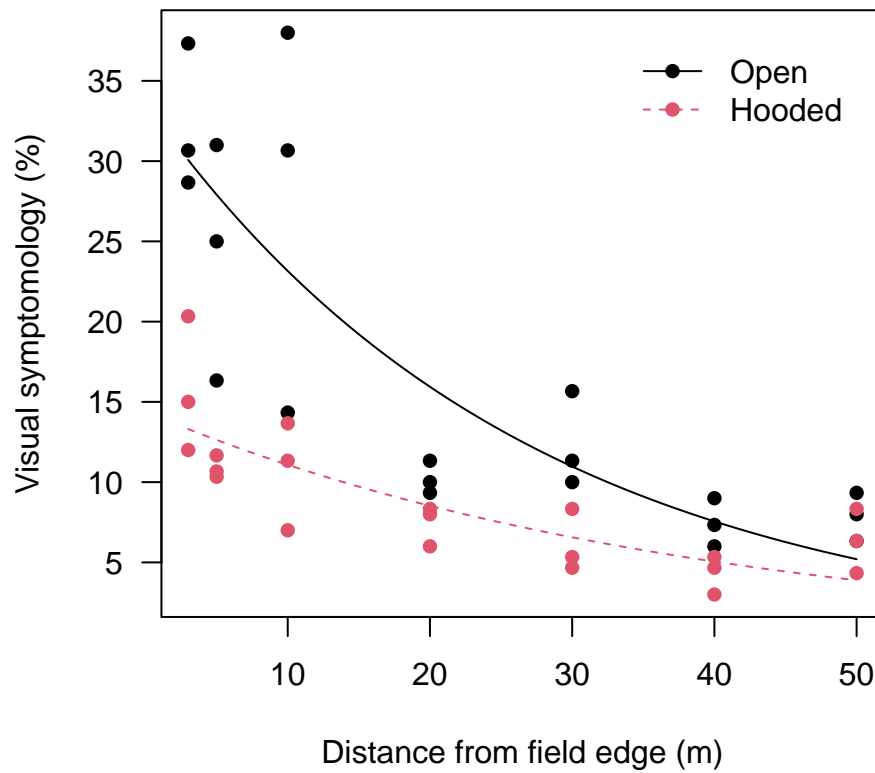
14 days after application

```
wisc14 <- dat %>%
  filter(rating_dat == "14" & location == "Wisconsin")

model_14_wi <- drm(rating_avg3plants ~ distance_m, boom,
  fct = EXD.2(),
  data = wisc14)

plot(model_14_wi, type = "all",
  col = c(1,2),
  pch = 16, log = "",
```

```
ylab = "Visual symptomology (%)",
xlab = "Distance from field edge (m)")
```



```
summary(model_14_wi)
```

Model fitted: Exponential decay with lower limit at 0 (2 parms)

Parameter estimates:

	Estimate	Std. Error	t-value	p-value
d:Open	33.6379	2.5369	13.2594	8.364e-16 ***
d:Hooded	14.3956	2.3099	6.2322	2.735e-07 ***
e:Open	26.7826	4.2638	6.2814	2.342e-07 ***
e:Hooded	38.1935	14.8301	2.5754	0.01403 *

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error:

4.725031 (38 degrees of freedom)

```
ED(model_14_wi, c(5, 10, 50, 90))
```

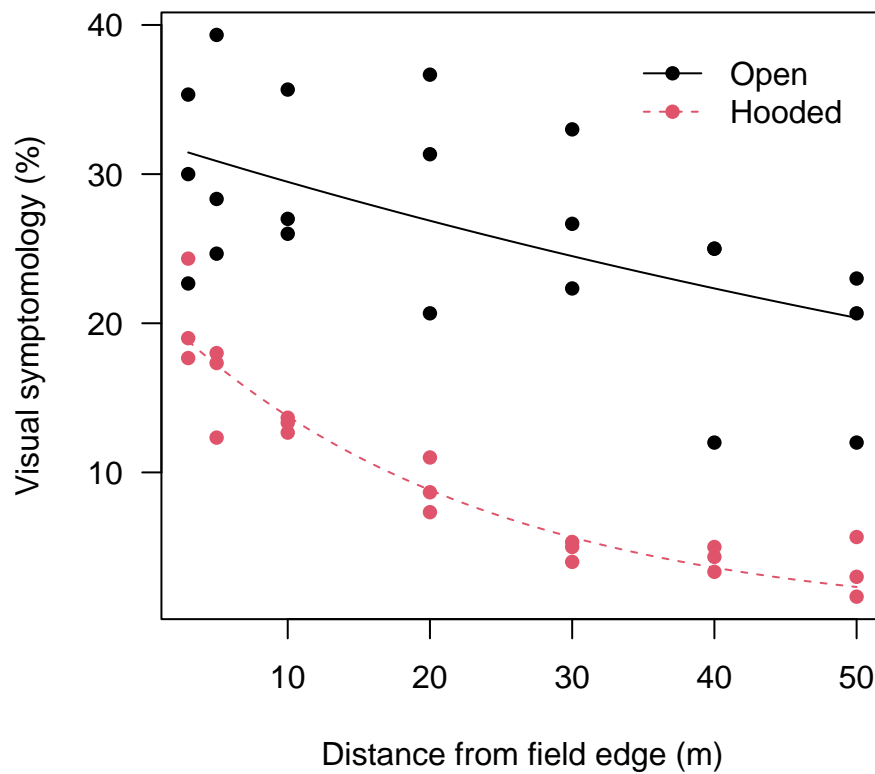
Estimated effective doses

	Estimate	Std. Error
e:Hooded:5	1.95907	0.76068
e:Hooded:10	4.02409	1.56250
e:Hooded:50	26.47375	10.27941
e:Hooded:90	87.94388	34.14745
e:Open:5	1.37377	0.21870
e:Open:10	2.82182	0.44923
e:Open:50	18.56426	2.95541

e:Open:90 61.66913 9.81768

28 days after application

```
wisc28 <- dat %>%  
  filter(rating_dat == "28" & location == "Wisconsin")  
  
model_28_wi <- drm(rating_avg3plants ~ distance_m, boom, fct = EXD.2(), data = wisc28)  
  
plot(model_28_wi, type = "all",  
      col = c(1,2),  
      pch = 16, log = "",  
      ylab = "Visual symptomology (%)",  
      xlab = "Distance from field edge (m)")
```



```
summary(model_28_wi)
```

Model fitted: Exponential decay with lower limit at 0 (2 parms)

Parameter estimates:

	Estimate	Std. Error	t-value	p-value
d:Open	32.3366	1.7967	17.9979	< 2.2e-16 ***

```

d:Hooded  21.4973      2.6228  8.1962 6.308e-10 ***
e:Open    108.1046     27.3537  3.9521 0.0003254 ***
e:Hooded  22.4464      5.5530  4.0422 0.0002491 ***

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error:

4.541804 (38 degrees of freedom)

```
ED(model_28_wi, c(5, 10, 50, 90))
```

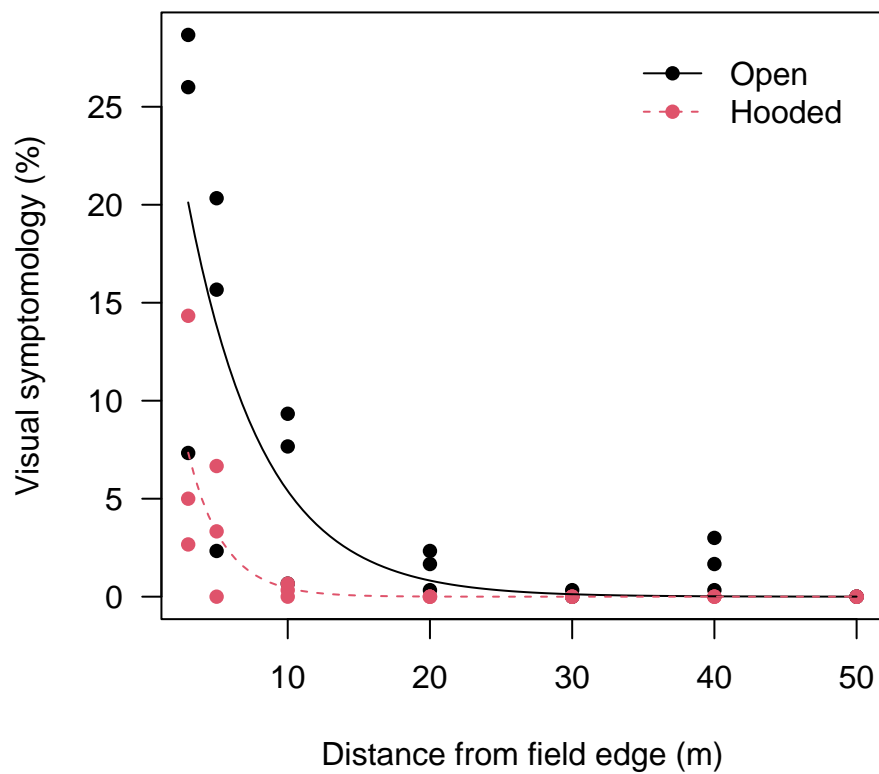
Estimated effective doses

	Estimate	Std. Error
e:Hooded:5	1.15135	0.28483
e:Hooded:10	2.36496	0.58507
e:Hooded:50	15.55863	3.84907
e:Hooded:90	51.68466	12.78634
e:Open:5	5.54504	1.40306
e:Open:10	11.38996	2.88200
e:Open:50	74.93241	18.96012
e:Open:90	248.92009	62.98416

Missouri

14 days after application

```
miss14 <- dat %>%  
  filter(rating_dat == "14" & location == "Missouri") %>%  
  mutate(rating_avg3plants = case_when(  
    rating_avg3plants == 0.000000 ~ 0.000001,  
    TRUE ~ rating_avg3plants  
  ))  
  
model_14_mo <- drm(rating_avg3plants ~ distance_m, boom,  
  fct = EXD.2(),  
  data = miss14)  
  
plot(model_14_mo, type = "all",  
  col = c(1,2),  
  pch = 16, log = "",  
  ylab = "Visual symptomology (%)",  
  xlab = "Distance from field edge (m)")
```

```
summary(model_14_mo)
```

Model fitted: Exponential decay with lower limit at 0 (2 parms)

Parameter estimates:

	Estimate	Std. Error	t-value	p-value
d:Open	35.3675	8.7849	4.0259	0.0002615 ***

```

d:Hooded 24.5941 28.8420 0.8527 0.3991577
e:Open 5.3172 1.6489 3.2247 0.0025928 **
e:Hooded 2.4835 2.0922 1.1870 0.2425937

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error:

3.992089 (38 degrees of freedom)

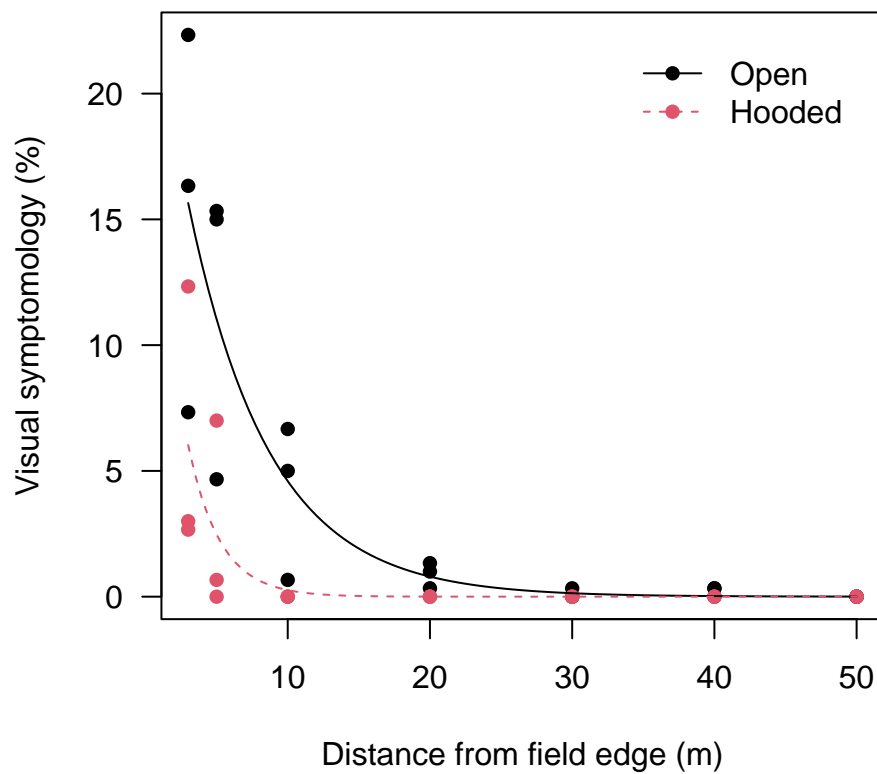
```
ED(model_14_mo, c(5, 10, 50, 90))
```

Estimated effective doses

	Estimate	Std. Error
e:Hooded:5	0.127386	0.107317
e:Hooded:10	0.261661	0.220438
e:Hooded:50	1.721418	1.450221
e:Hooded:90	5.718427	4.817530
e:Open:5	0.272737	0.084578
e:Open:10	0.560224	0.173730
e:Open:50	3.685611	1.142934
e:Open:90	12.243335	3.796746

28 days after application

```
miss28 <- dat %>%  
  filter(rating_dat == "28" & location == "Missouri") %>%  
  mutate(rating_avg3plants = case_when(  
    rating_avg3plants == 0.000000 ~ 0.000001,  
    TRUE ~ rating_avg3plants  
  ))  
  
model_28_mo <- drm(rating_avg3plants ~ distance_m, boom,  
  fct = EXD.2(),  
  data = miss28)  
  
plot(model_28_mo, type = "all",  
  col = c(1,2),  
  pch = 16, log = "",  
  ylab = "Visual symptomology (%)",  
  xlab = "Distance from field edge (m)")
```



```
summary(model_28_mo)
```

Model fitted: Exponential decay with lower limit at 0 (2 parms)

Parameter estimates:

	Estimate	Std. Error	t-value	p-value
d:Open	26.4652	5.1850	5.1042	9.584e-06 ***

```

d:Hooded  23.1460    24.8630  0.9309 0.3577615
e:Open    5.7097     1.4360  3.9760 0.0003032 ***
e:Hooded  2.2299     1.5764  1.4146 0.1653400

```

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error:

2.810194 (38 degrees of freedom)

```
summary(model_28_mo)
```

Model fitted: Exponential decay with lower limit at 0 (2 parms)

Parameter estimates:

	Estimate	Std. Error	t-value	p-value
d:Open	26.4652	5.1850	5.1042	9.584e-06 ***
d:Hooded	23.1460	24.8630	0.9309	0.3577615
e:Open	5.7097	1.4360	3.9760	0.0003032 ***
e:Hooded	2.2299	1.5764	1.4146	0.1653400

Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error:

2.810194 (38 degrees of freedom)

```
ED(model_28_mo, c(5, 10, 50, 90))
```

Estimated effective doses

	Estimate	Std. Error
e:Hooded:5	0.114378	0.080858
e:Hooded:10	0.234942	0.166088
e:Hooded:50	1.545637	1.092663
e:Hooded:90	5.134493	3.629749
e:Open:5	0.292871	0.073659
e:Open:10	0.601581	0.151302
e:Open:50	3.957690	0.995389
e:Open:90	13.147163	3.306611