

20211122_spinrelaxation_plotting_R.R

kelse

2022-06-01

```
# 20211122 plotting of spin relaxation data collected on 500 MHz and 600 MHz  
# spectrometers. Plots of R1, R2, NOE, and calculated order parameters by residue.
```

```
library(tidyverse)
```

```
## -- Attaching packages ----- tidyverse 1.3.0 --
```

```
## v ggplot2 3.3.3      v purrr  0.3.4  
## v tibble  3.1.0      v dplyr  1.0.5  
## v tidyr   1.1.3      v stringr 1.4.0  
## v readr   1.4.0      v forcats 0.5.1
```

```
## Warning: package 'stringr' was built under R version 4.0.5
```

```
## -- Conflicts ----- tidyverse_conflicts() --  
## x dplyr::filter() masks stats::filter()  
## x dplyr::lag()     masks stats::lag()
```

```
library(readxl)
```

```
#load file - edited in excel  
#raw data included in folder
```

```
order_param <- read_excel("20211122_spinrelaxation_plotting.xlsx", sheet = 1)
```

```
R1 <- read_excel("20211122_spinrelaxation_plotting.xlsx", sheet = 2)
```

```
R2 <- read_excel("20211122_spinrelaxation_plotting.xlsx", sheet = 3)
```

```
NOE <- read_excel("20211122_spinrelaxation_plotting.xlsx", sheet = 4)
```

```
#Combined data - R1, R2, and HetNOE
```

```
#R1
```

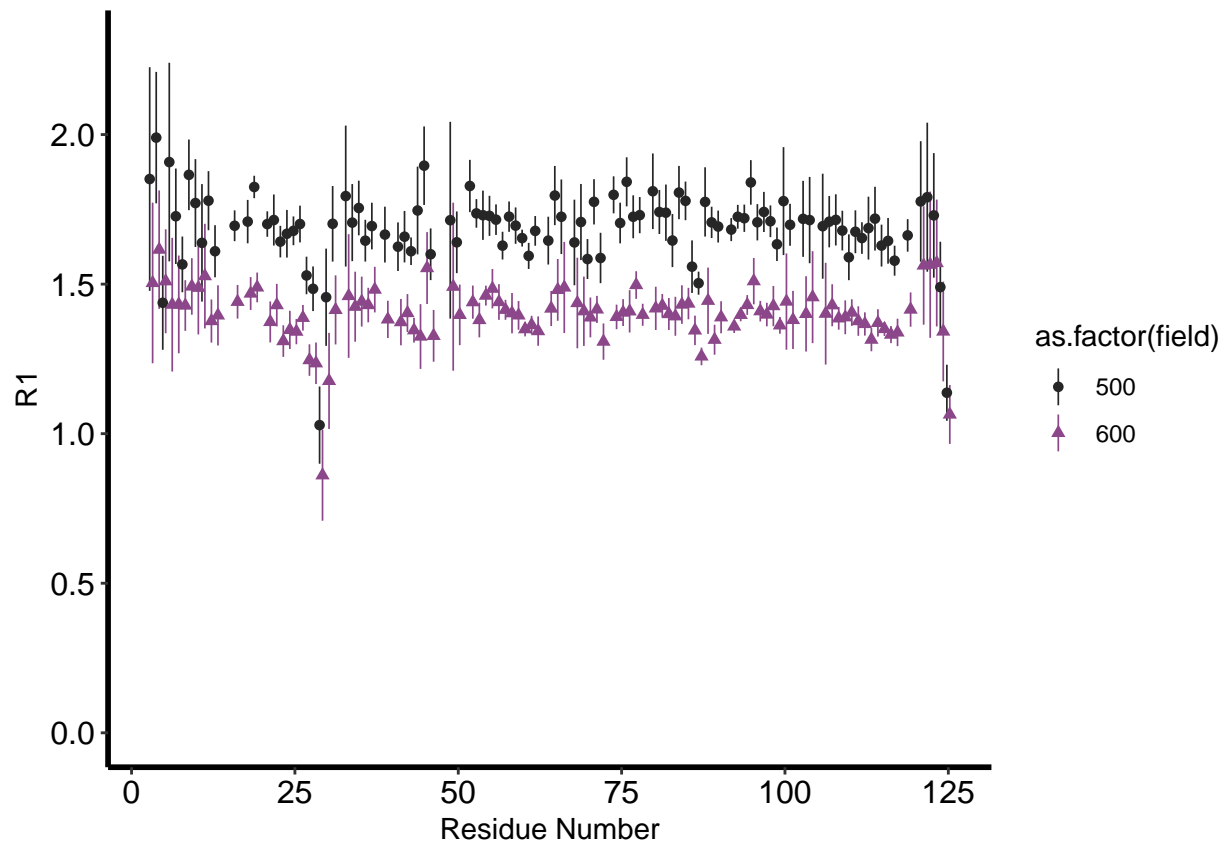
```
R1 %>%
```

```
  ggplot(., aes(x = residue, y = R1, color = as.factor(field), shape=as.factor(field))) +  
  geom_pointrange(aes(ymin=R1-R1err, ymax=R1+R1err), position=position_dodge(0.9),  
                  size=0.3) +  
  scale_color_manual(values=c("gray15", "orchid4")) +
```

```

labs(x = "Residue Number", y = "R1") +
theme_bw() +
scale_y_continuous(limits = c(0, 2.3), breaks = c(0, 0.5, 1.0, 1.5, 2.0)) +
theme(axis.text = element_text(color = "black", size = 12),
      panel.grid.major = element_blank(),
      panel.grid.minor = element_blank(),
      panel.border = element_blank(),
      panel.background = element_blank(),
      axis.line.x = element_line(color = "black", size = 1, linetype = 1),
      axis.line.y = element_line(color = "black", size = 1, linetype = 1)
)

```



```

ggsave("20211122_Fis1_R1.png",
       width = 16, height = 10, units = "cm")

ggsave("20211122_Fis1_R1.pdf",
       width = 16, height = 10, units = "cm")

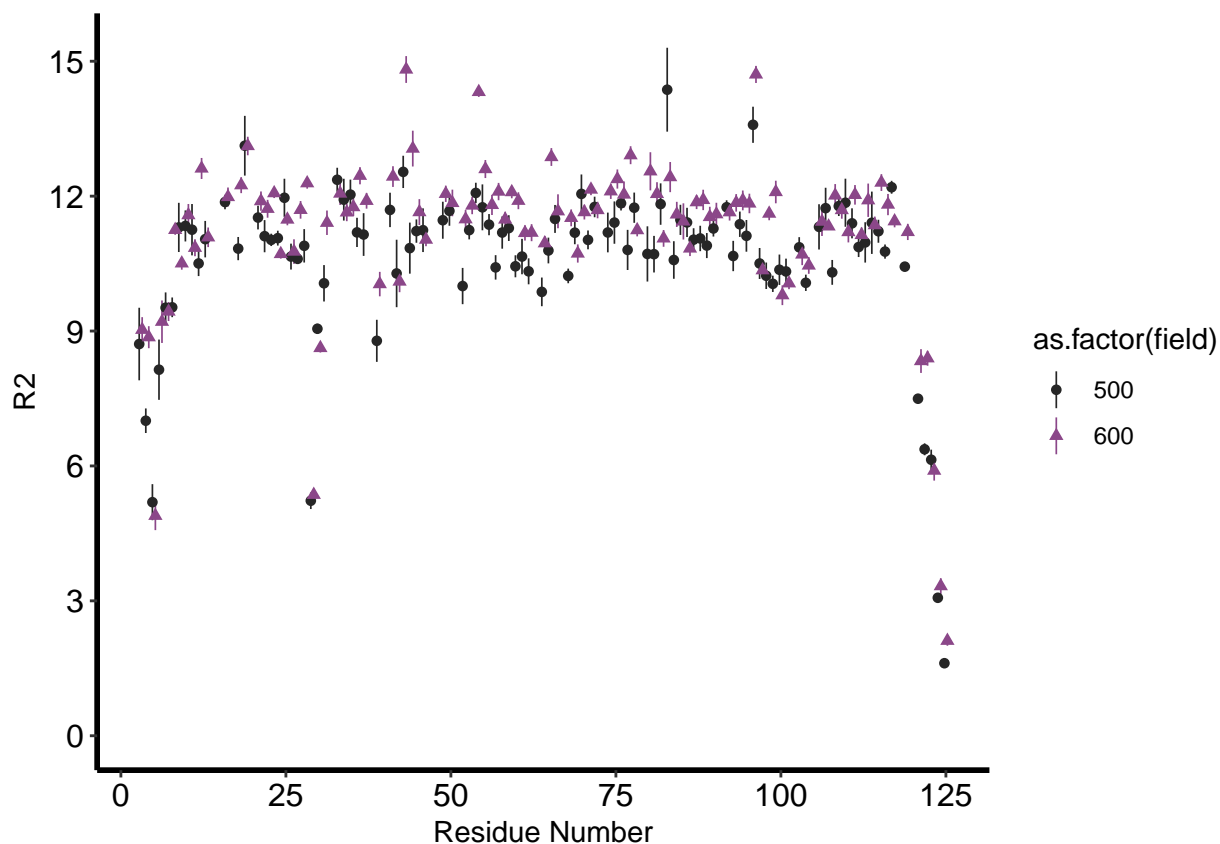
#R2
R2 %>%
  ggplot(., aes(x = residue, y = R2, color = as.factor(field), shape=as.factor(field))) +
  geom_pointrange(aes(ymin=R2-R2err, ymax=R2+R2err), position=position_dodge(0.9),
                 size=0.3) +
  scale_color_manual(values=c("gray15", "orchid4")) +
  labs(x = "Residue Number", y = "R2") +

```

```

theme_bw() +
scale_y_continuous(limits = c(0, 15.3), breaks = c(0, 3, 6, 9, 12, 15)) +
theme(axis.text = element_text(color = "black", size = 12),
      panel.grid.major = element_blank(),
      panel.grid.minor = element_blank(),
      panel.border = element_blank(),
      panel.background = element_blank(),
      axis.line.x = element_line(color = "black", size = 1, linetype = 1),
      axis.line.y = element_line(color = "black", size = 1, linetype = 1)
)

```



```

ggsave("20211122_Fis1_R2.png",
       width = 16, height = 10, units = "cm")

ggsave("20211122_Fis1_R2.pdf",
       width = 16, height = 10, units = "cm")

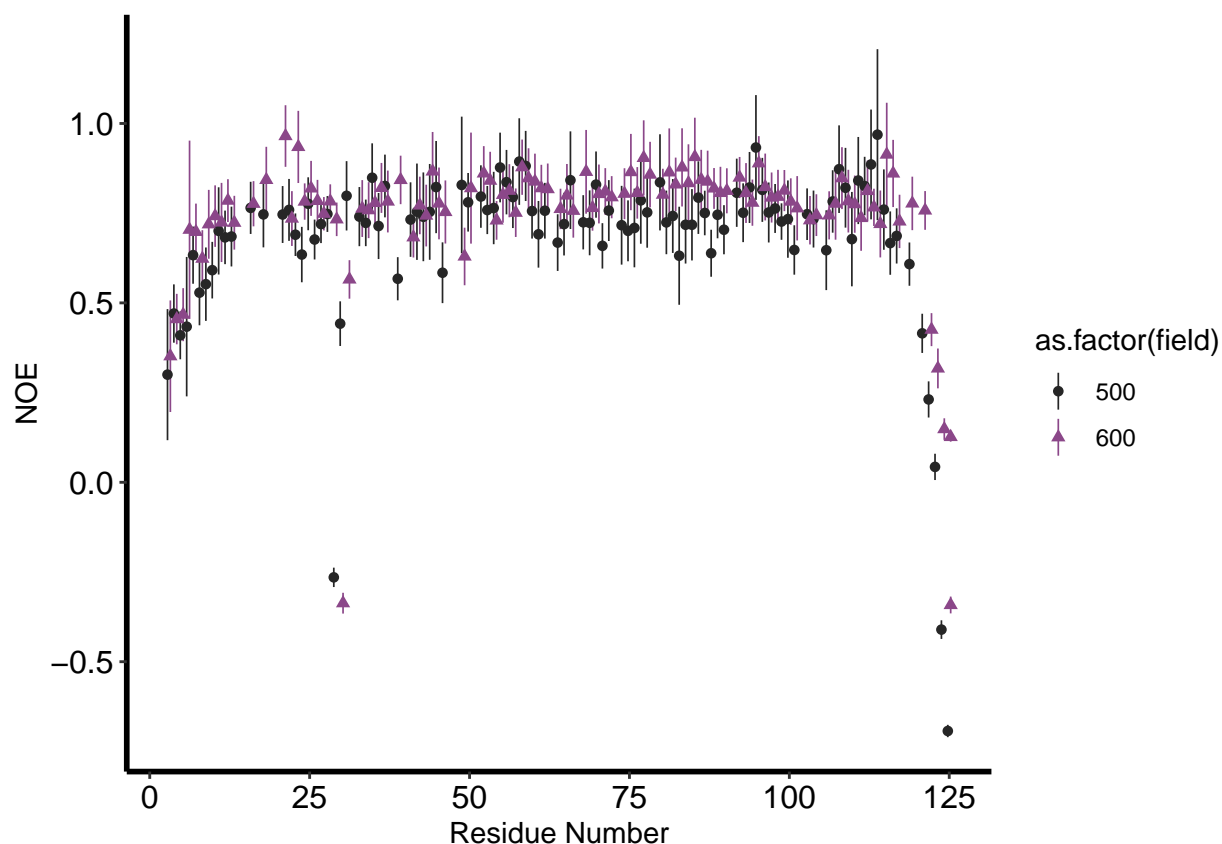
#NOE
NOE %>%
  ggplot(., aes(x = residue, y = NOE, color = as.factor(field), shape=as.factor(field))) +
  geom_pointrange(aes(ymin=NOE-NOEerr, ymax=NOE+NOEerr), position=position_dodge(0.9),
                 size=0.3) +
  scale_color_manual(values=c("gray15", "orchid4")) +
  labs(x = "Residue Number", y = "NOE") +
  theme_bw() +

```

```

theme(axis.text = element_text(color = "black", size = 12),
      panel.grid.major = element_blank(),
      panel.grid.minor = element_blank(),
      panel.border = element_blank(),
      panel.background = element_blank(),
      axis.line.x = element_line(color = "black", size = 1, linetype = 1),
      axis.line.y = element_line(color = "black", size = 1, linetype = 1)
)

```



```

ggsave("20211122_Fis1_NOE.png",
       width = 16, height = 10, units = "cm")

ggsave("20211122_Fis1_NOE.pdf",
       width = 16, height = 10, units = "cm")

#Plot order parameter figure
order_param %>%
  filter(., residue!= 5) %>%
  ggplot(., aes(x = residue, y = S2, color = secondary)) +
  # geom_point(stat = "identity", color = "gray40", size = 1.0) +
  geom_pointrange(aes(ymin=S2-S2err, ymax=S2+S2err), size = 0.3) +
  scale_y_continuous(limits = c(0, 1.1), breaks = c(0, 0.2, 0.4, 0.6, 0.8, 1.0)) +
  scale_color_manual(breaks = c("arm", "helix", "loop"),
                    labels = c("N-arm", "Helix", "Loop"),

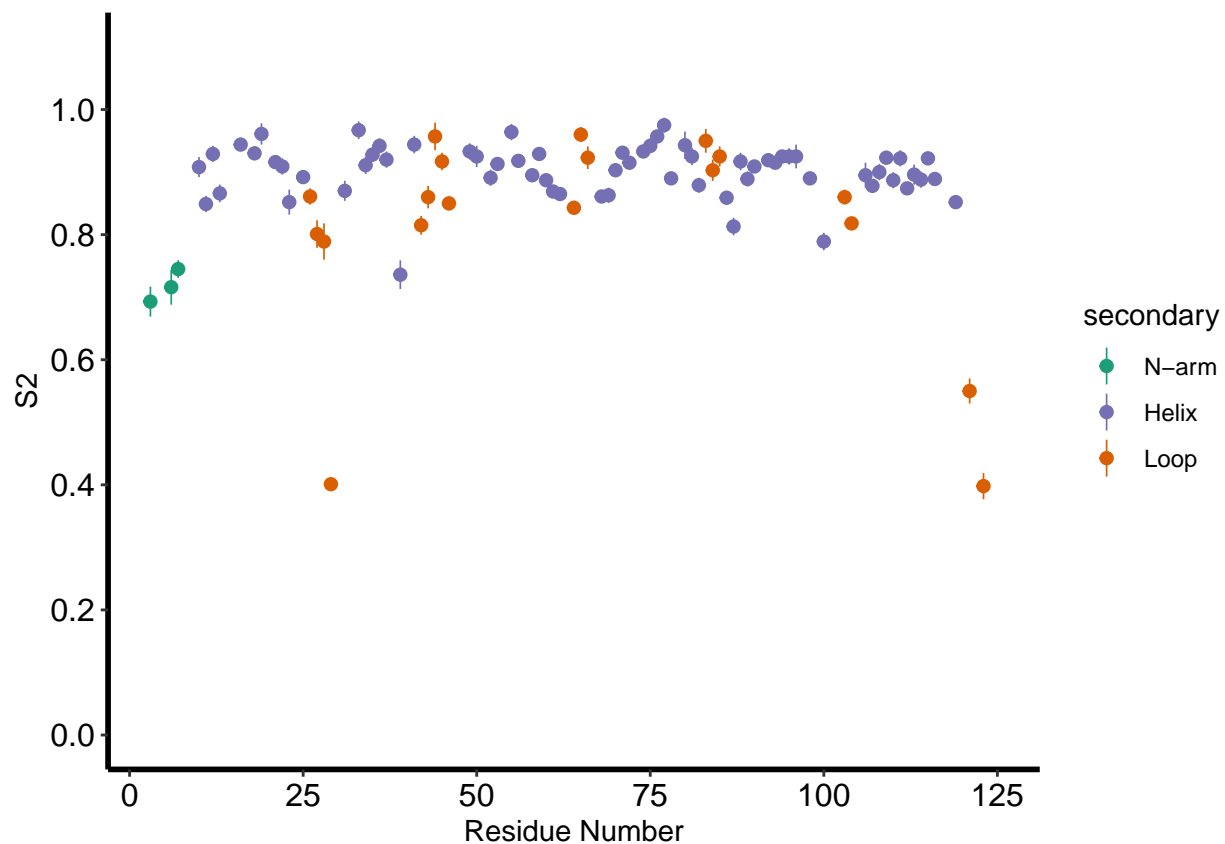
```

```

        values = c("#1b9e77", "#7570b3", "#d95f02")) +
labs(x = "Residue Number", y = "S2") +
theme_bw() +
theme(axis.text = element_text(color = "black", size = 12),
      panel.grid.major = element_blank(),
      panel.grid.minor = element_blank(),
      panel.border = element_blank(),
      panel.background = element_blank(),
      axis.line.x = element_line(color = "black", size = 1, linetype = 1),
      axis.line.y = element_line(color = "black", size = 1, linetype = 1)
)

```

Warning: Removed 14 rows containing missing values (geom_pointrange).



```

ggsave("20211122_Fis1_order_parameters_color.png",
       width = 16, height = 10, units = "cm")

```

Warning: Removed 14 rows containing missing values (geom_pointrange).

```

ggsave("20211122_Fis1_order_parameters_color.pdf",
       width = 16, height = 10, units = "cm")

```

Warning: Removed 14 rows containing missing values (geom_pointrange).