Serodynamics Analysis for Multiple Subjects

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Introduction

This document demonstrates the full workflow using the **serodynamics** package. In this analysis, we run JAGS models for three different subjects:

- **sees_npl_128 (HlyE_IgA)**
- **sees_npl_131 (HlyE_IgA)**
- **sees_npl_133 (HlyE_IgG)**

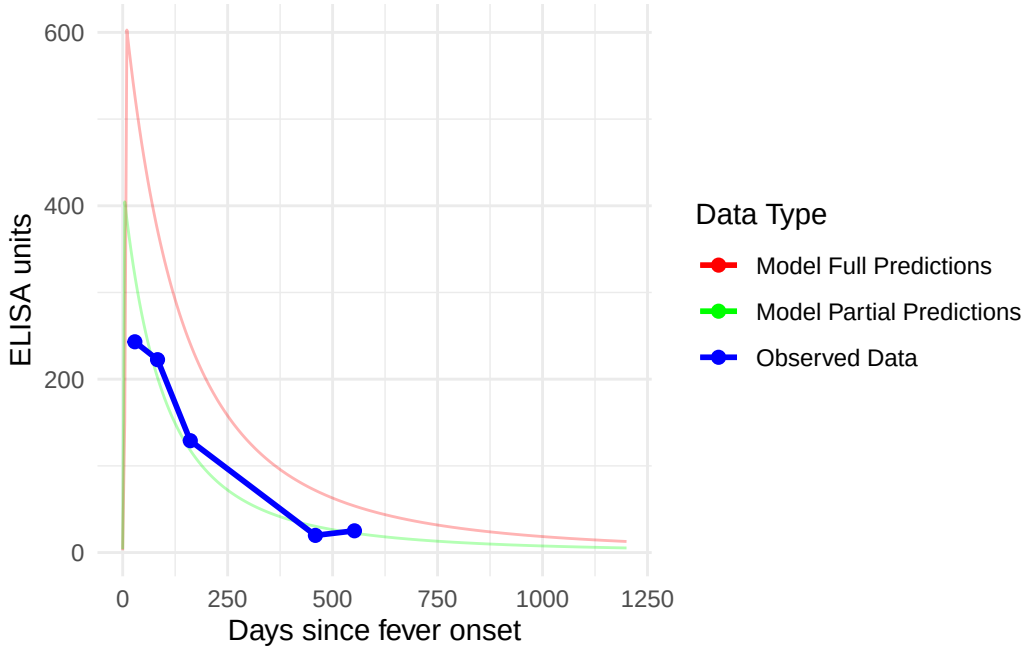
For each subject, we process the JAGS output to extract median parameter estimates and generate predicted antibody response curve plots.

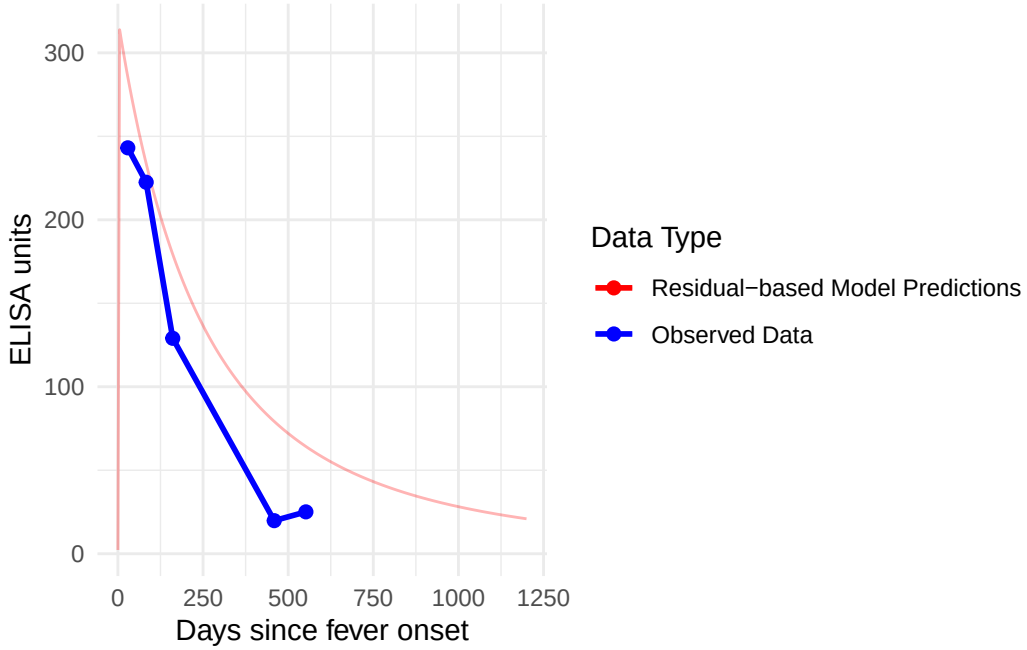
Load Required Libraries

Run Models for Each Subject

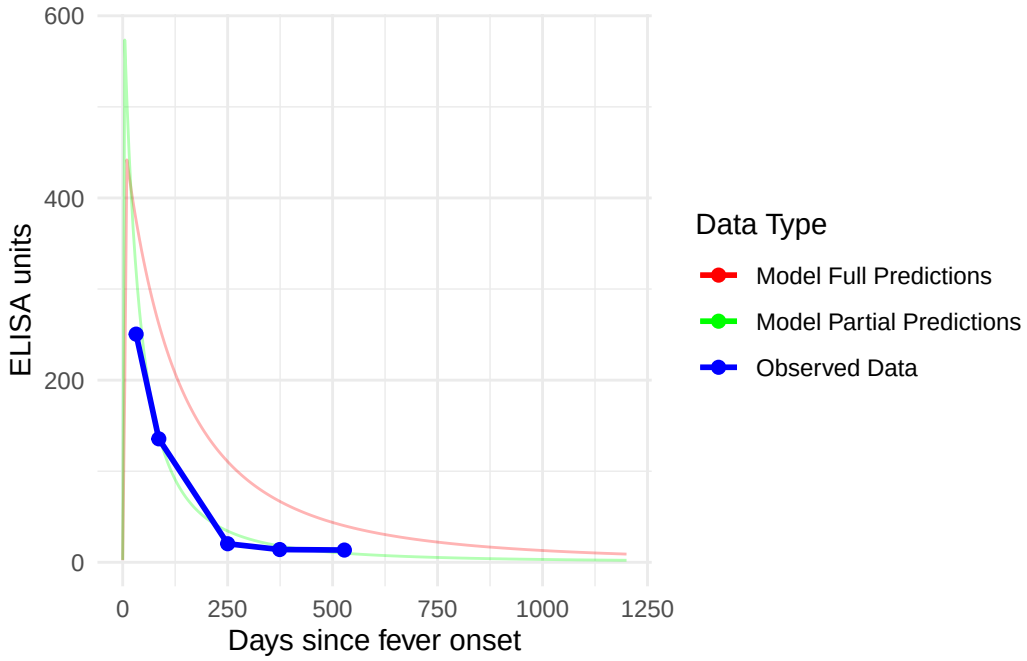
```
# A tibble: 3 x 4
  Country id           antigen_iso visit_num
  <chr>   <chr>        <chr>           <int>
1 Nepal   sees_npl_128 HlyE_IgA            5
2 Nepal   sees_npl_131 HlyE_IgA            5
3 Nepal   sees_npl_133 HlyE_IgG            5
```

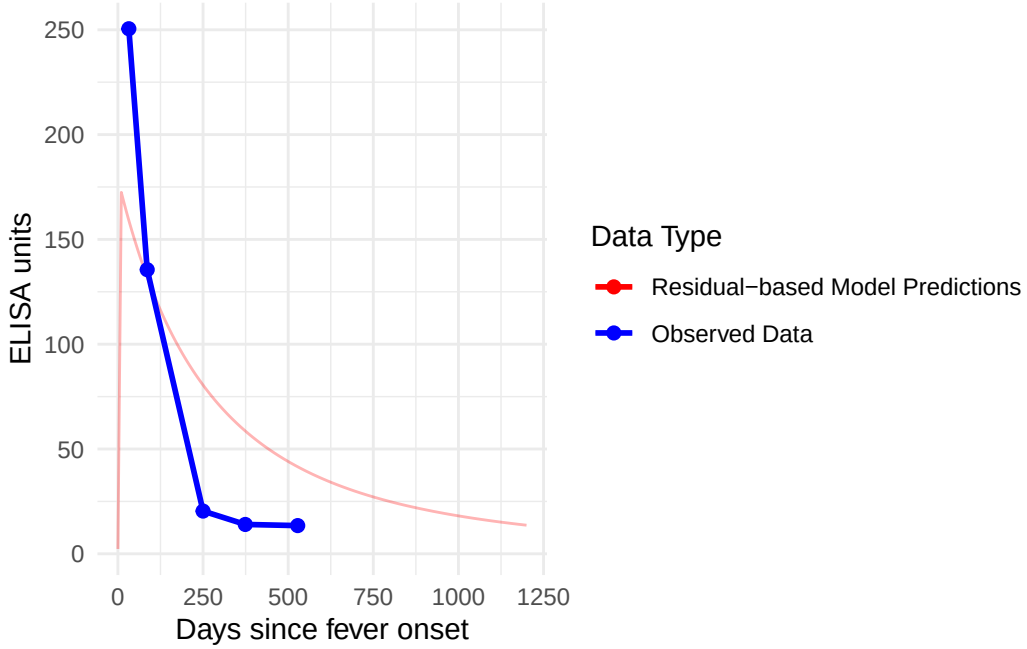
Subject: sees_npl_128 (HlyE_IgA)



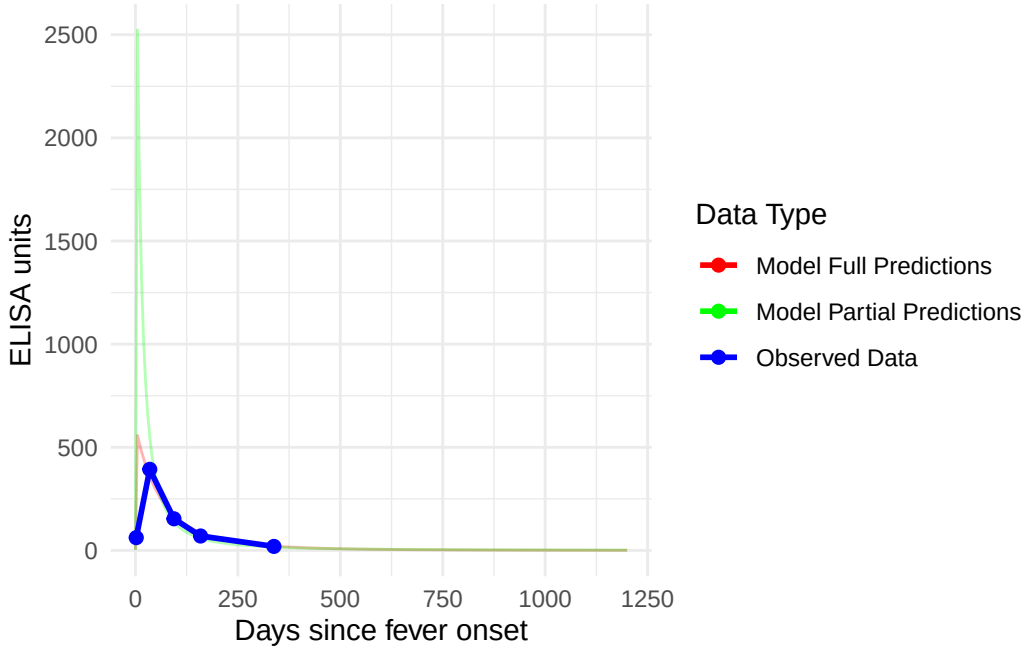


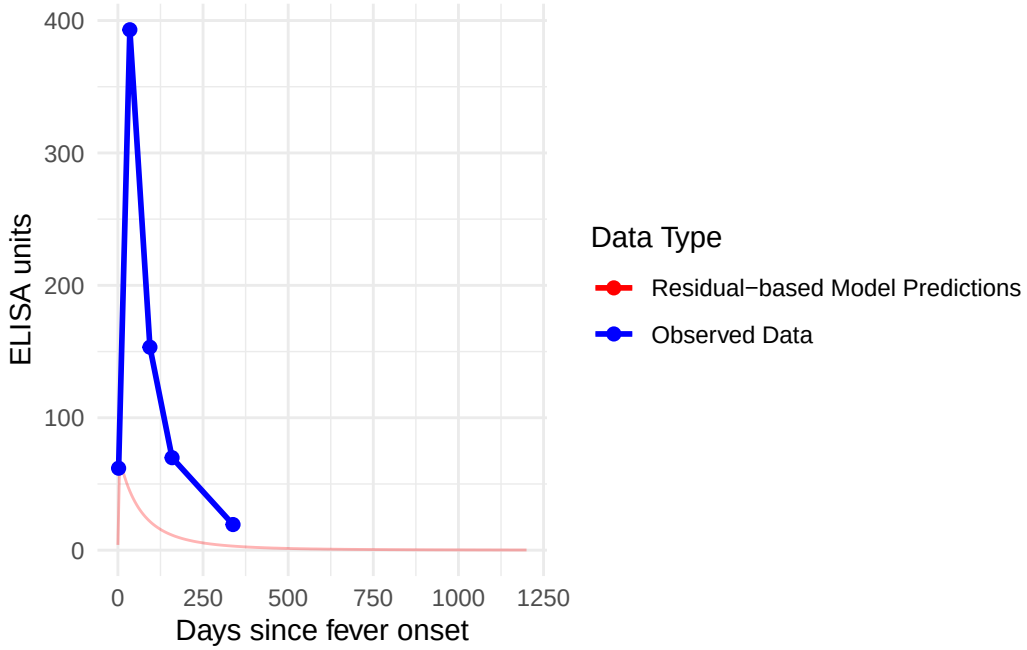
Subject: sees_npl_131 (HlyE_IgA)





Subject: sees_npl_133 (HlyE_IgG)





Conclusion

This document generated JAGS models and corresponding predicted antibody response curves for three subjects:

- sees_npl_128 (HlyE_IgA)
- sees_npl_131 (HlyE_IgA)
- sees_npl_133 (HlyE_IgG)