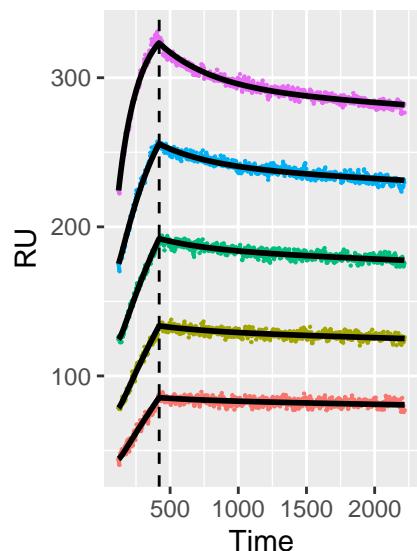


CH505

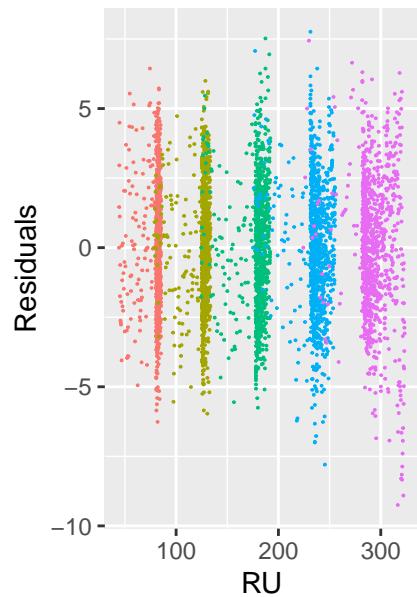
Bivalent Analyte Model–1 with Extended Length of Dissociation



Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

Residuals

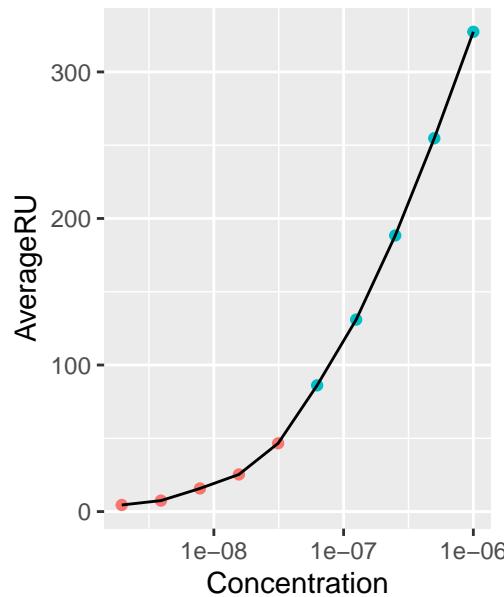


Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

 $ka1$ 1.58e+03 1.77e+01 $ka2$ 7.78e-05 1.46e-06 $kd1$ 2.56e-03 5.82e-05 $kd2$ 1.05e-05 3.84e-07 $Rmax$ 1 8.84e+02 8.19e+00 $Rmax$ 2 7.01e+02 1.42e+01 $Rmax$ 3 6.48e+02 3.74e+00 $Rmax$ 4 6.34e+02 2.41e+00 $Rmax$ 5 6.73e+02 1.49e+00 p 1 1.00e+00 NA p 2 6.81e-01 1.28e-01 p 3 1.00e+00 NA p 4 1.00e+00 NA

CH505

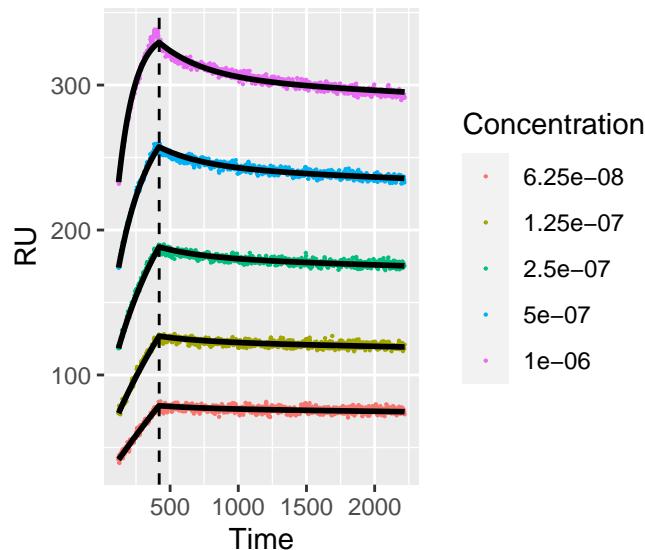


Included

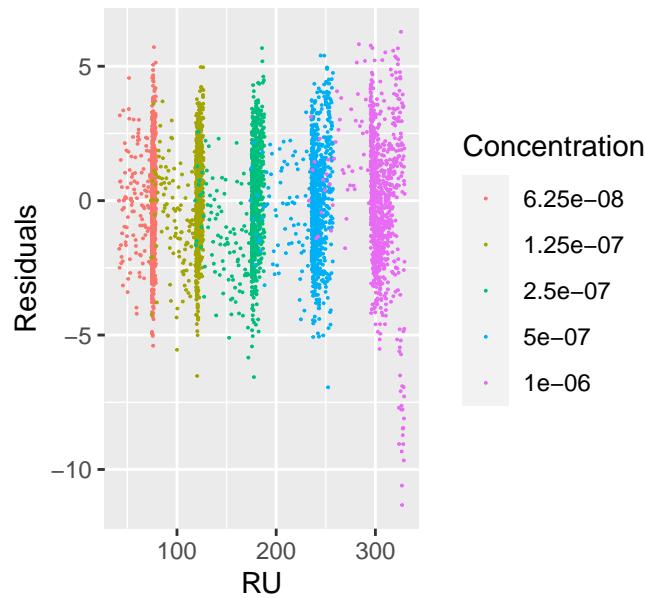
- No
- Yes

CH505

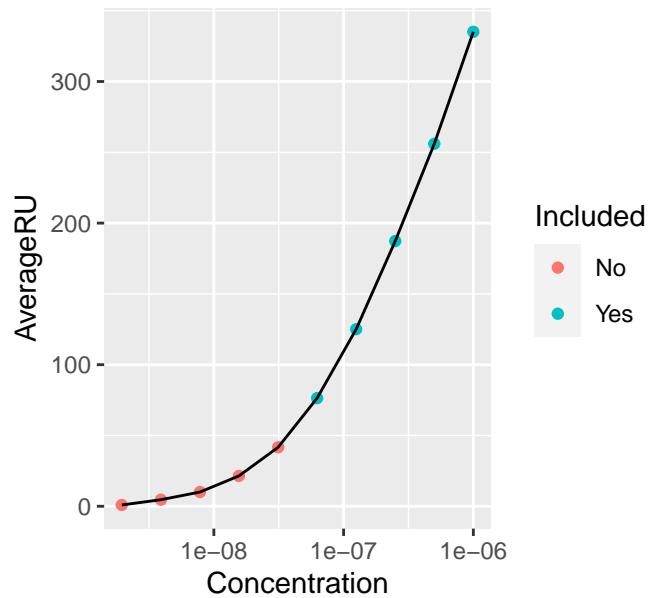
Bivalent Analyte Model–1 with Extended Length of Dissociation



Residuals

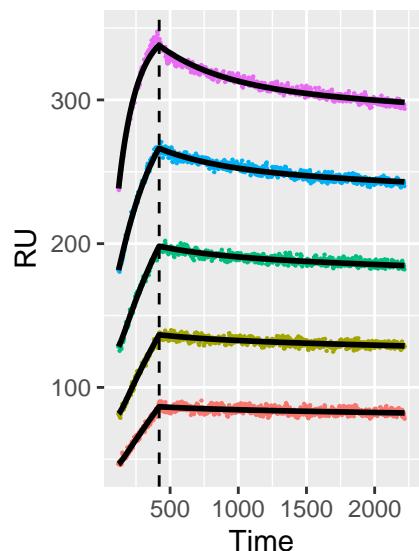
 k_{a1} 2.14e+03 4.59e+01 k_{a2} 1.07e-04 2.42e-06 k_{d1} 2.91e-03 1.02e-04 k_{d2} 8.06e-06 4.76e-07 R_{max} 1 5.82e+02 1.64e+01 R_{max} 2 4.83e+02 1.19e+01 R_{max} 3 4.29e+02 9.66e+00 R_{max} 4 4.11e+02 2.16e+01 R_{max} 5 4.36e+02 1.28e+01 p 1 1.99e-01 1.94e-01 p 2 0.00e+00 9.91e-02 p 3 0.00e+00 7.14e-02 p 4 0.00e+00 1.28e-01

CH505



CH505

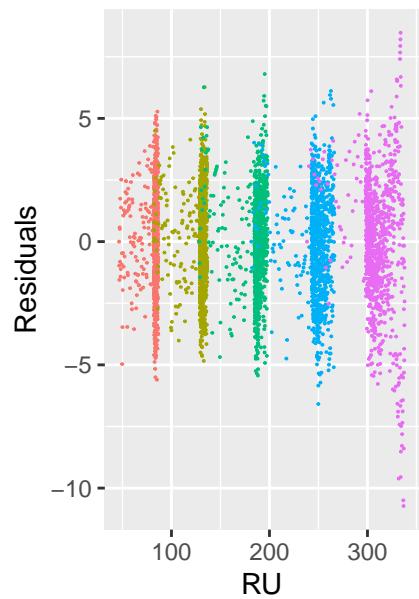
Bivalent Analyte Model–1 with Extended Length of Dissociation



Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

Residuals

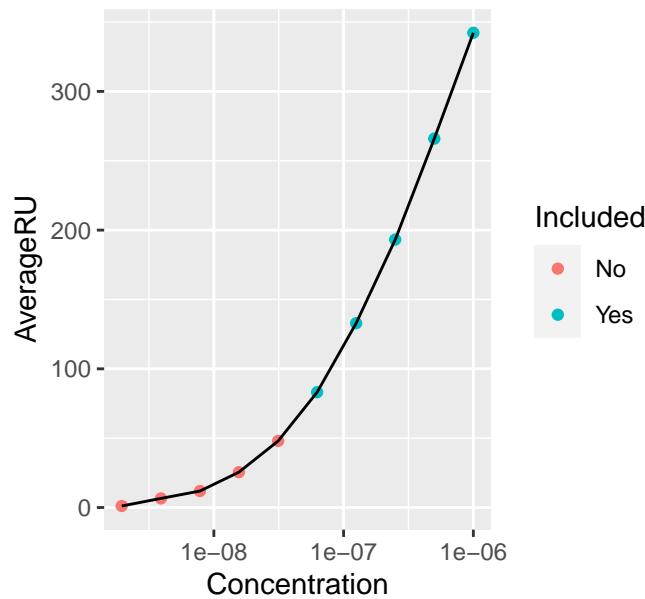


Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

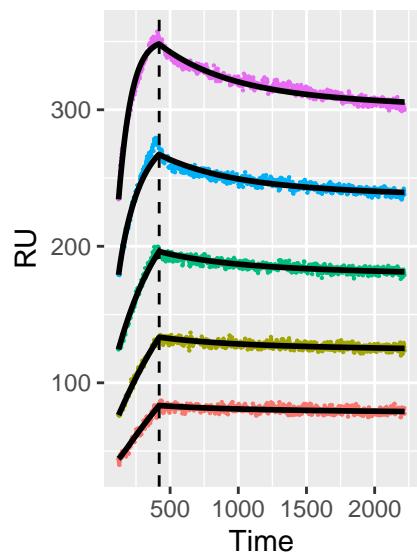
 ka_1 1.72e+03 2.11e+01 ka_2 8.41e-05 1.66e-06 kd_1 2.09e-03 5.18e-05 kd_2 8.52e-06 4.43e-07 R_{max} 1 8.01e+02 8.29e+00 R_{max} 2 6.87e+02 5.90e+00 R_{max} 3 6.31e+02 3.97e+00 R_{max} 4 6.35e+02 2.55e+00 R_{max} 5 5.64e+02 1.47e+01 p 1 1.00e+00 NA p 2 1.00e+00 NA p 3 1.00e+00 NA p 4 1.00e+00 NA

CH505

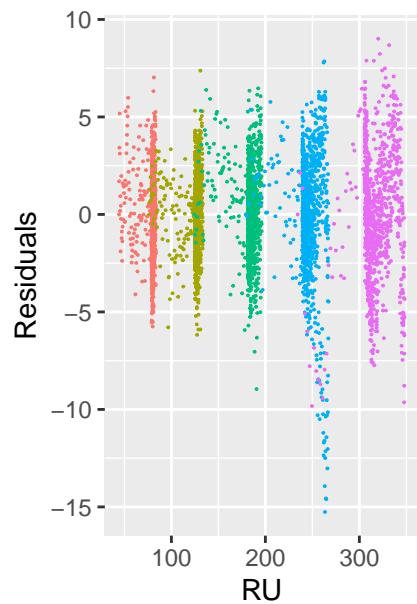


CH505

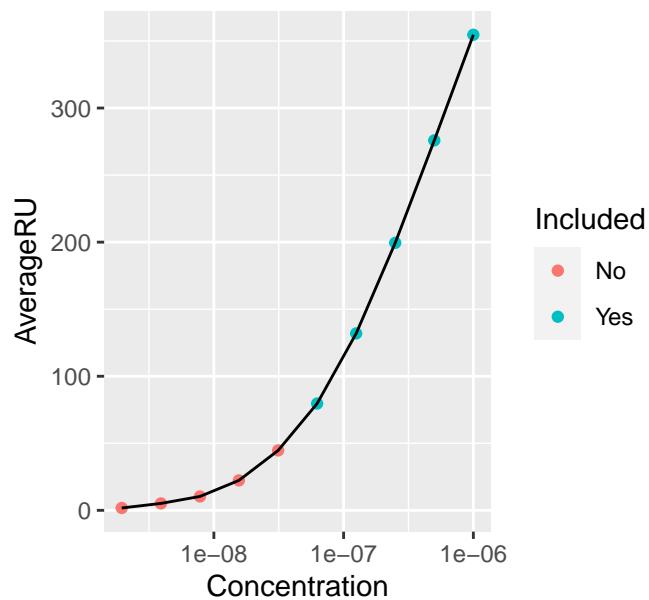
Bivalent Analyte Model–1 with Extended Length of Dissociation



Residuals

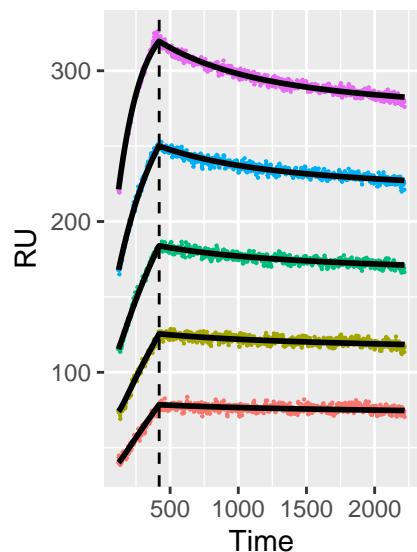
 ka_1 2.88e+03 3.27e+01 ka_2 9.80e-05 2.13e-06 kd_1 1.95e-03 5.49e-05 kd_2 4.81e-06 6.71e-07 R_{max} 1 5.22e+02 4.75e+00 R_{max} 2 4.35e+02 1.80e+01 R_{max} 3 4.03e+02 3.69e+01 R_{max} 4 5.62e+02 1.57e+00 R_{max} 5 4.43e+02 4.25e+00 p 1 1.00e+00 NA p 2 2.34e-01 2.06e-01 p 3 1.98e-01 2.88e-01 p 4 1.00e+00 NA

CH505



CH505

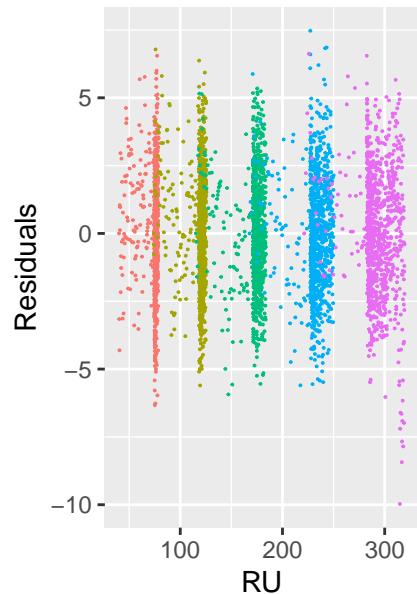
Bivalent Analyte Model–1 with Extended Length of Dissociation



Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

Residuals

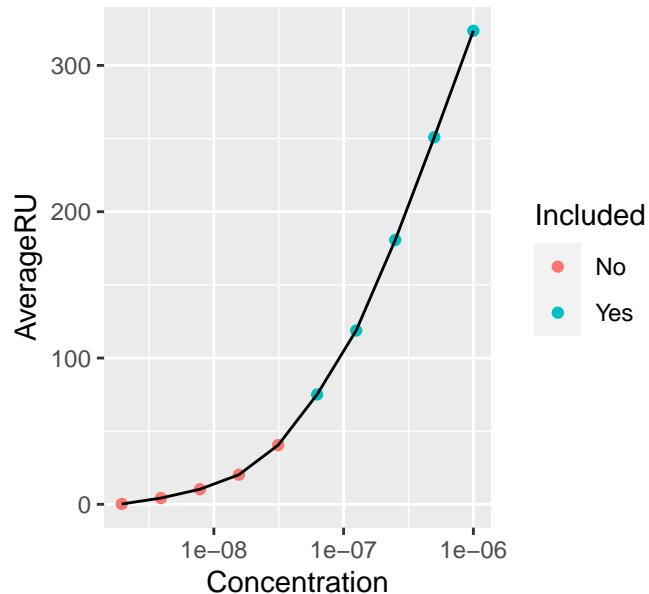


Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

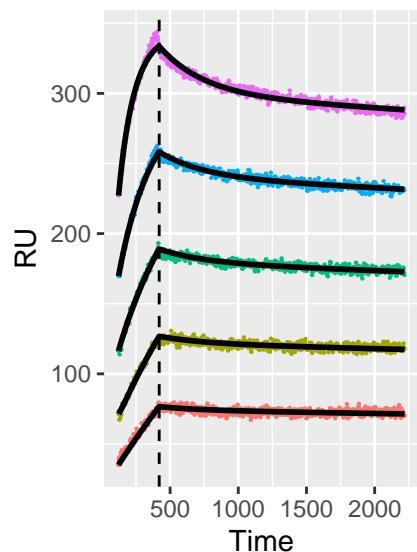
| | | |
|-------------|----------|----------|
| k_a1 | 1.57e+03 | 2.84e+01 |
| k_a2 | 7.10e-05 | 2.25e-06 |
| k_d1 | 1.62e-03 | 5.77e-05 |
| k_d2 | 7.12e-06 | 7.19e-07 |
| $R_{max} 1$ | 8.13e+02 | 1.28e+01 |
| $R_{max} 2$ | 6.75e+02 | 9.01e+00 |
| $R_{max} 3$ | 5.60e+02 | 2.00e+01 |
| $R_{max} 4$ | 5.16e+02 | 6.77e+01 |
| $R_{max} 5$ | 4.39e+02 | 8.29e+00 |
| p_1 | 1.00e+00 | NA |
| p_2 | 1.00e+00 | NA |
| p_3 | 5.32e-01 | 1.68e-01 |
| p_4 | 4.49e-01 | 3.89e-01 |

CH505



CH505

Bivalent Analyte Model–1 with Extended Length of Dissociation

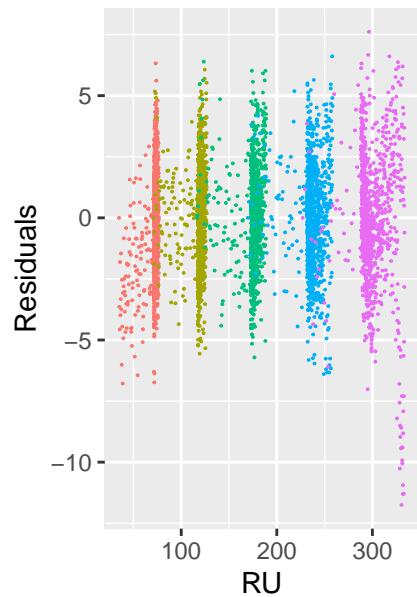


Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

| | | |
|-------------|----------|----------|
| k_{a1} | 2.00e+03 | 3.62e+01 |
| k_{a2} | 8.00e-05 | 1.38e-06 |
| k_{d1} | 3.05e-03 | 9.02e-05 |
| k_{d2} | 1.03e-05 | 5.21e-07 |
| $R_{max} 1$ | 6.67e+02 | 1.71e+01 |
| $R_{max} 2$ | 5.68e+02 | 1.20e+01 |
| $R_{max} 3$ | 4.61e+02 | 9.43e+00 |
| $R_{max} 4$ | 4.35e+02 | 1.55e+01 |
| $R_{max} 5$ | 6.22e+02 | 4.60e+01 |
| p_1 | 0.00e+00 | 2.21e-01 |
| p_2 | 4.09e-01 | 1.04e-01 |
| p_3 | 0.00e+00 | 6.48e-02 |
| p_4 | 0.00e+00 | 8.99e-02 |

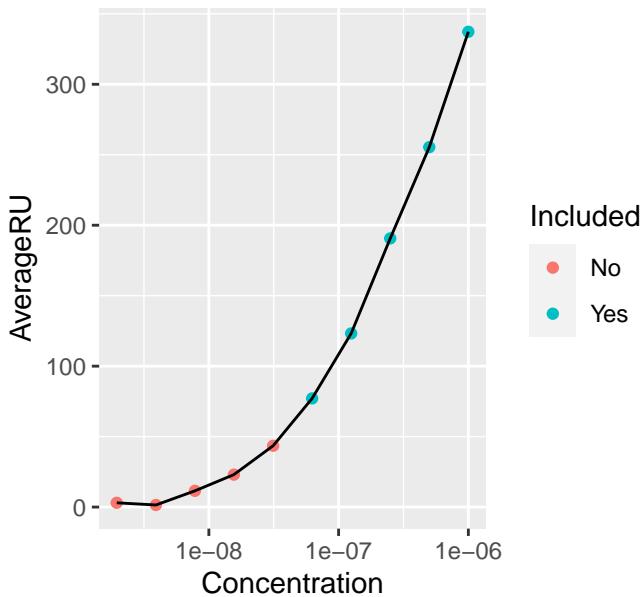
Residuals



Concentration

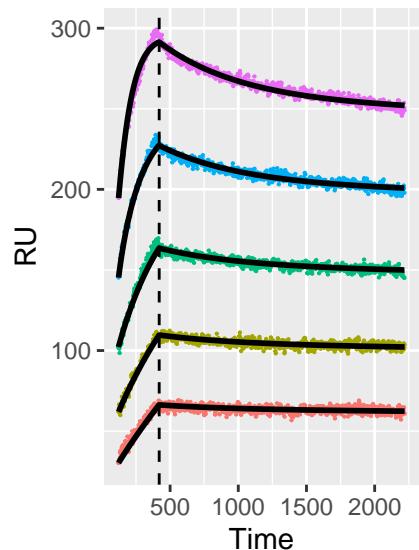
- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

CH505



CH505

Bivalent Analyte Model–1 with Extended Length of Dissociation

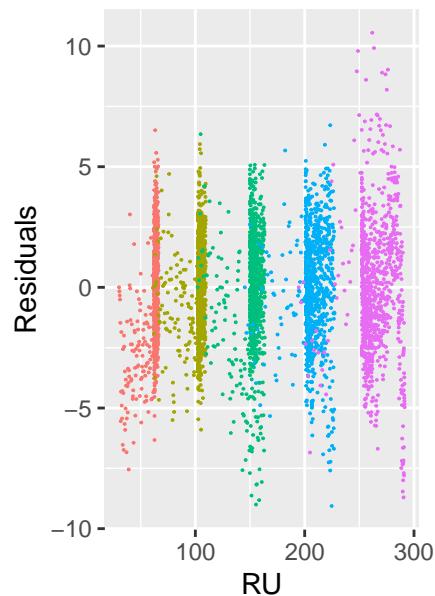


Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

| | | |
|-------------|----------|----------|
| k_{a1} | 2.78e+03 | 6.98e+01 |
| k_{a2} | 9.70e-05 | 2.56e-06 |
| k_{d1} | 1.65e-03 | 7.37e-05 |
| k_{d2} | 4.26e-06 | 1.24e-06 |
| $R_{max} 1$ | 4.28e+02 | 1.94e+01 |
| $R_{max} 2$ | 3.52e+02 | 1.76e+01 |
| $R_{max} 3$ | 3.22e+02 | 4.32e+01 |
| $R_{max} 4$ | 4.67e+02 | 3.99e+01 |
| $R_{max} 5$ | 3.68e+02 | 6.01e+00 |
| p_1 | 0.00e+00 | 4.44e-01 |
| p_2 | 8.48e-04 | 2.47e-01 |
| p_3 | 0.00e+00 | 4.11e-01 |
| p_4 | 9.14e-01 | 2.58e-01 |

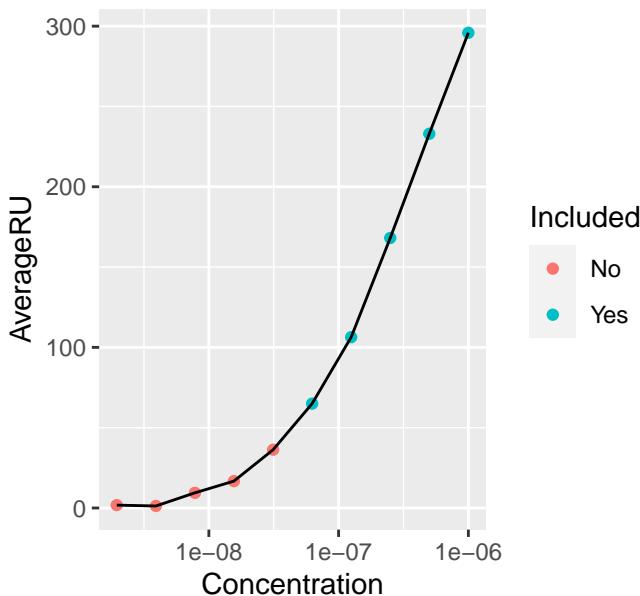
Residuals



Concentration

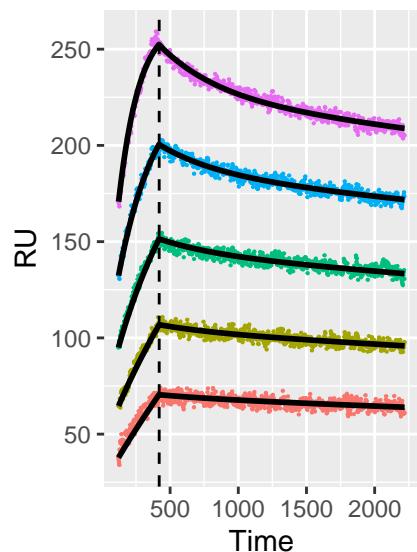
- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

CH505



CH505

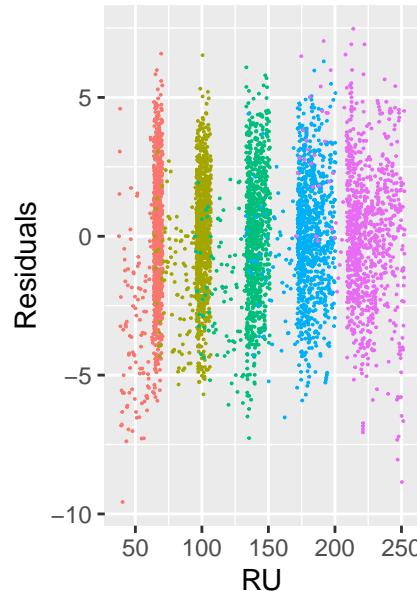
Bivalent Analyte Model–1 with Extended Length of Dissociation



Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

Residuals



Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

ka_1 1.73e+03 4.48e+01

ka_2 7.46e-05 2.65e-06

kd_1 2.29e-03 9.16e-05

kd_2 2.09e-05 8.34e-07

R_{max} 1 6.05e+02 2.17e+01

R_{max} 2 4.58e+02 1.50e+01

R_{max} 3 3.90e+02 1.25e+01

R_{max} 4 3.50e+02 1.88e+01

R_{max} 5 3.48e+02 7.68e+00

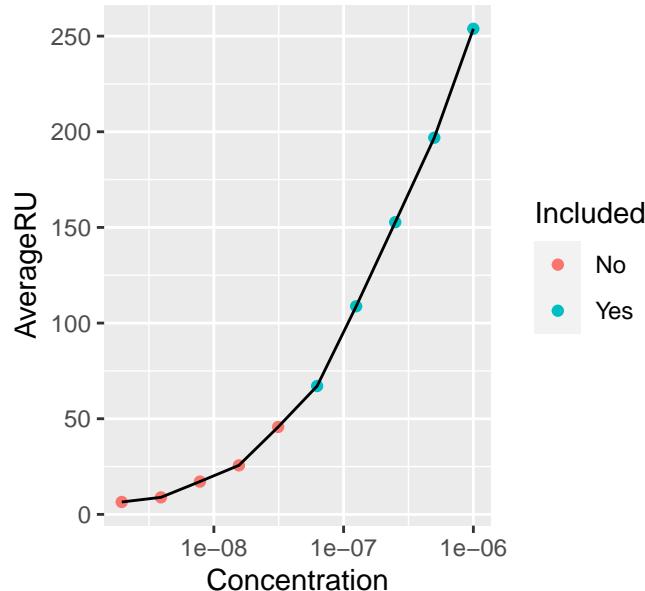
p 1 0.00e+00 2.44e-01

p 2 0.00e+00 1.25e-01

p 3 0.00e+00 1.01e-01

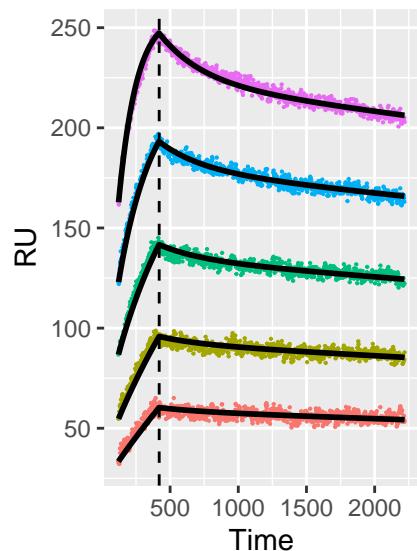
p 4 0.00e+00 1.27e-01

CH505

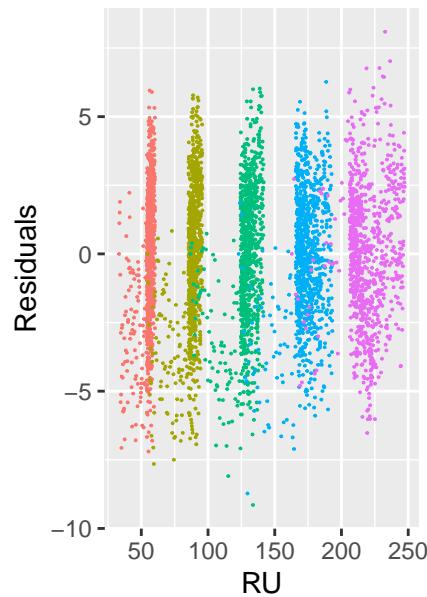


CH505

Bivalent Analyte Model–1 with Extended Length of Dissociation



Residuals



ka_1 $1.93e+03$ $5.19e+01$

ka_2 $9.24e-05$ $3.04e-06$

kd_1 $3.19e-03$ $1.39e-04$

kd_2 $2.16e-05$ $7.55e-07$

R_{max} 1 $4.64e+02$ $1.78e+01$

R_{max} 2 $4.09e+02$ $1.33e+01$

R_{max} 3 $3.57e+02$ $1.02e+01$

R_{max} 4 $3.39e+02$ $1.39e+01$

R_{max} 5 $3.46e+02$ $1.27e+01$

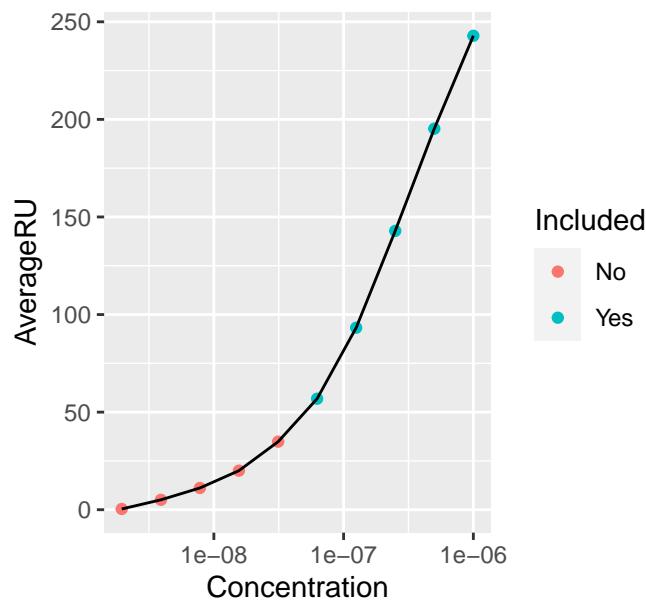
p 1 $0.00e+00$ $2.00e-01$

p 2 $0.00e+00$ $1.18e-01$

p 3 $0.00e+00$ $8.17e-02$

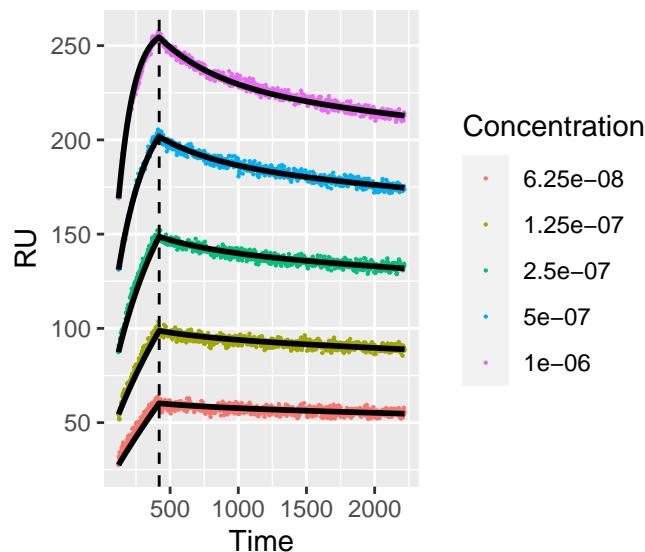
p 4 $0.00e+00$ $1.10e-01$

CH505

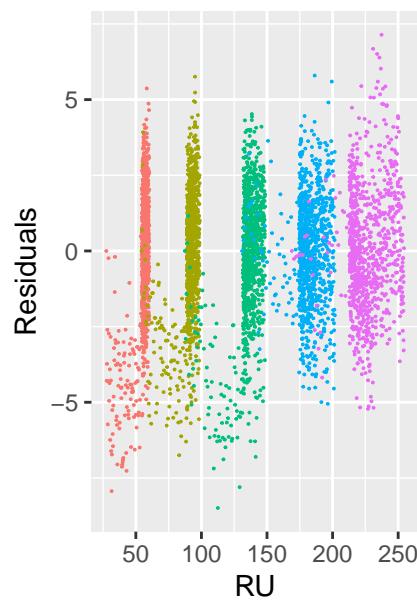


CH505

Bivalent Analyte Model–1 with Extended Length of Dissociation

 $ka1$ $2.36e+03$ $6.34e+01$ $ka2$ $1.02e-04$ $2.72e-06$ $kd1$ $2.24e-03$ $8.97e-05$ $kd2$ $1.82e-05$ $8.57e-07$ $Rmax$ 1 $4.52e+02$ $1.65e+01$ $Rmax$ 2 $3.68e+02$ $1.22e+01$ $Rmax$ 3 $3.39e+02$ $1.40e+01$ $Rmax$ 4 $3.18e+02$ $2.59e+01$ $Rmax$ 5 $3.35e+02$ $9.11e+00$ p 1 $0.00e+00$ $3.24e-01$ p 2 $0.00e+00$ $1.61e-01$ p 3 $0.00e+00$ $1.60e-01$ p 4 $5.19e-04$ $1.78e-01$

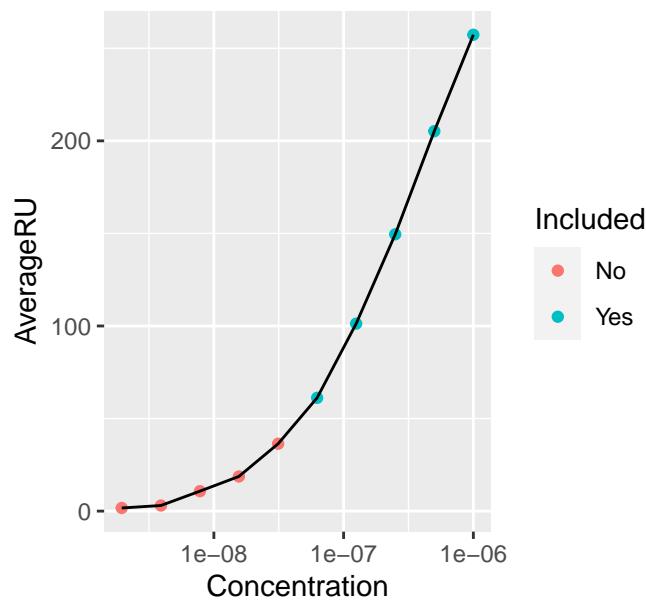
Residuals



Concentration

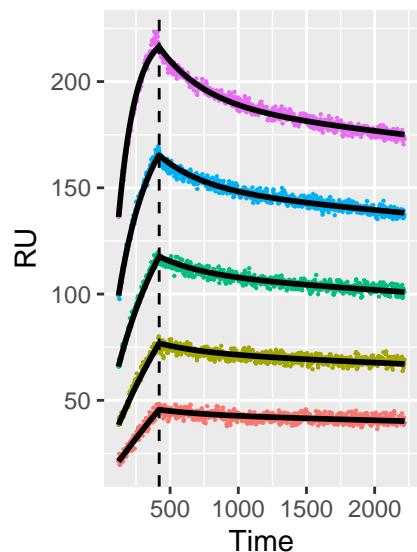
- $6.25e-08$
- $1.25e-07$
- $2.5e-07$
- $5e-07$
- $1e-06$

CH505



CH505

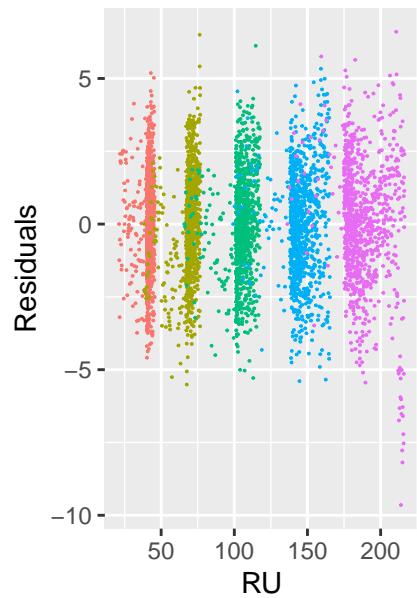
Bivalent Analyte Model–1 with Extended Length of Dissociation



Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

Residuals

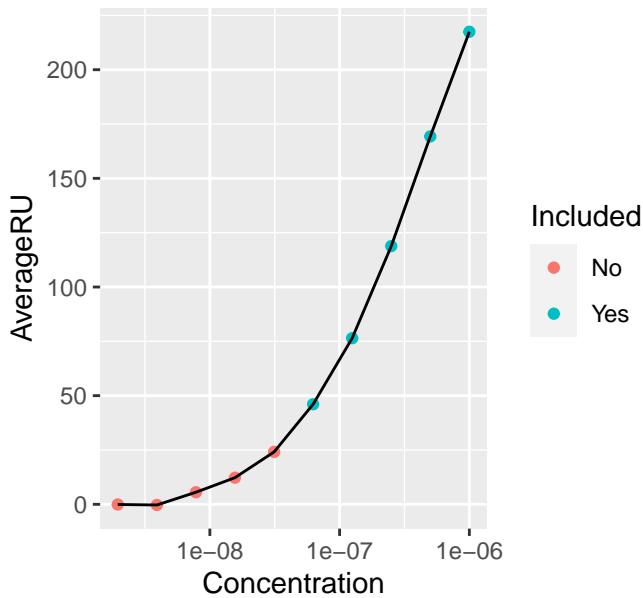


Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

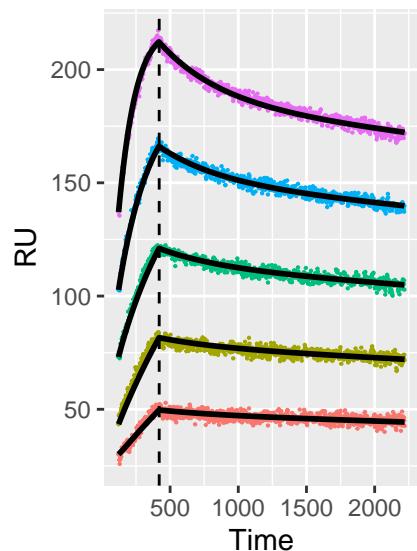
| | | |
|-------------|----------|----------|
| ka_1 | 2.13e+03 | 4.01e+01 |
| ka_2 | 8.63e-05 | 2.15e-06 |
| kd_1 | 2.96e-03 | 9.44e-05 |
| kd_2 | 2.09e-05 | 7.99e-07 |
| R_{max} 1 | 3.90e+02 | 1.17e+01 |
| R_{max} 2 | 3.44e+02 | 9.03e+00 |
| R_{max} 3 | 3.03e+02 | 7.78e+00 |
| R_{max} 4 | 2.90e+02 | 1.19e+01 |
| R_{max} 5 | 3.01e+02 | 5.42e+00 |
| p 1 | 2.62e-01 | 2.22e-01 |
| p 2 | 0.00e+00 | 1.22e-01 |
| p 3 | 0.00e+00 | 8.66e-02 |
| p 4 | 0.00e+00 | 1.09e-01 |

CH505

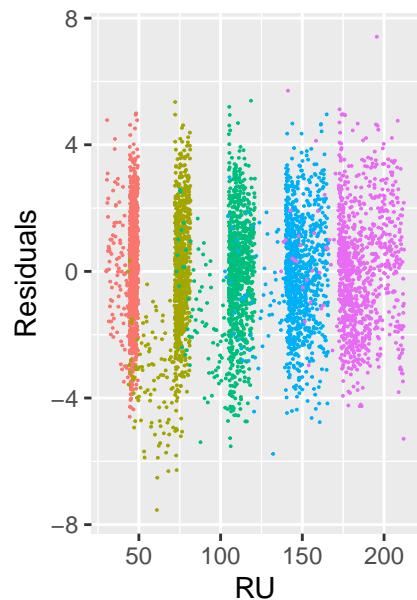


CH505

Bivalent Analyte Model–1 with Extended Length of Dissociation

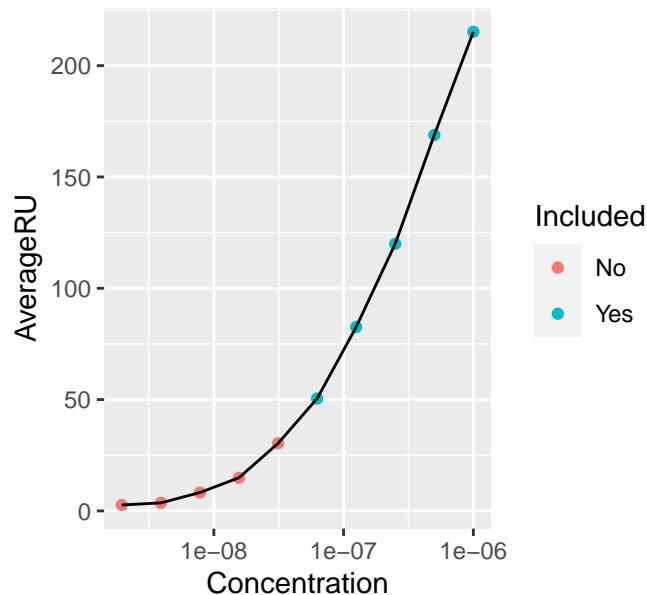


Residuals



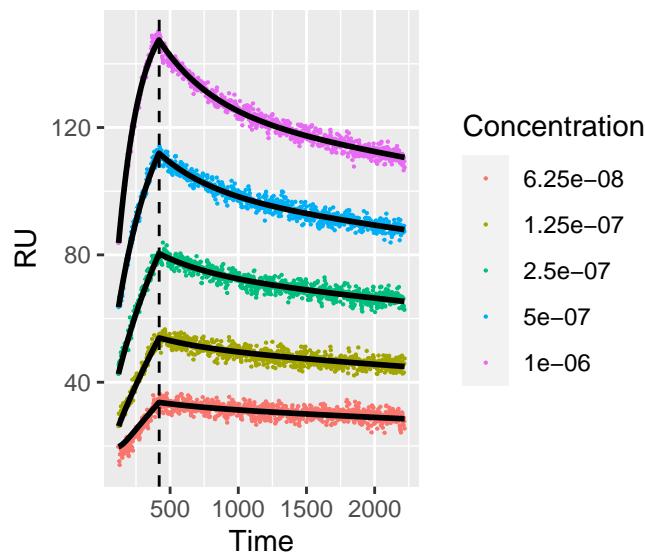
| | | |
|-------------|------------|------------|
| ka_1 | $2.15e+03$ | $4.93e+01$ |
| ka_2 | $9.44e-05$ | $2.69e-06$ |
| kd_1 | $2.28e-03$ | $8.58e-05$ |
| kd_2 | $2.15e-05$ | $9.53e-07$ |
| $R_{max} 1$ | $3.32e+02$ | $1.22e+01$ |
| $R_{max} 2$ | $3.39e+02$ | $1.04e+01$ |
| $R_{max} 3$ | $2.86e+02$ | $1.03e+01$ |
| $R_{max} 4$ | $2.80e+02$ | $1.66e+01$ |
| $R_{max} 5$ | $2.87e+02$ | $6.19e+00$ |
| p_1 | $3.39e-01$ | $1.94e-01$ |
| p_2 | $0.00e+00$ | $1.53e-01$ |
| p_3 | $0.00e+00$ | $1.23e-01$ |
| p_4 | $5.71e-04$ | $1.42e-01$ |

CH505

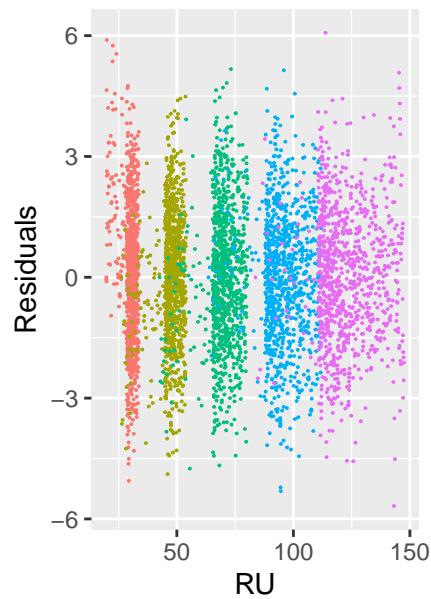


CH505

Bivalent Analyte Model–1 with Extended Length of Dissociation



Residuals

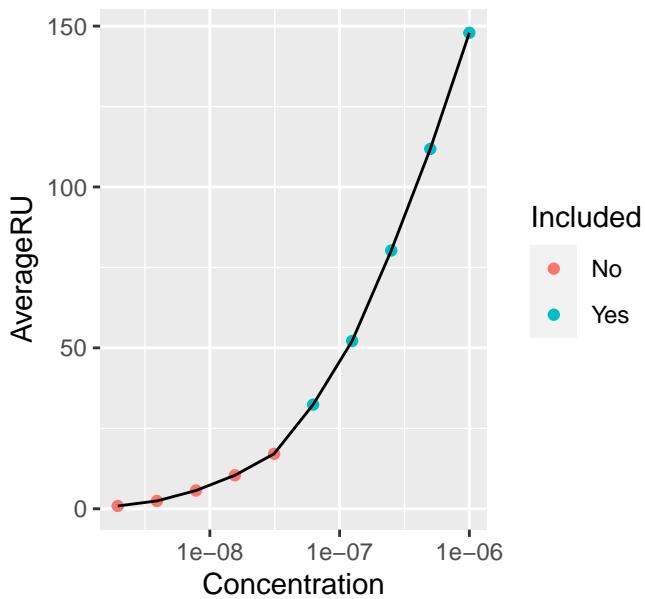


Concentration

- 6.25e-08
- 1.25e-07
- 2.5e-07
- 5e-07
- 1e-06

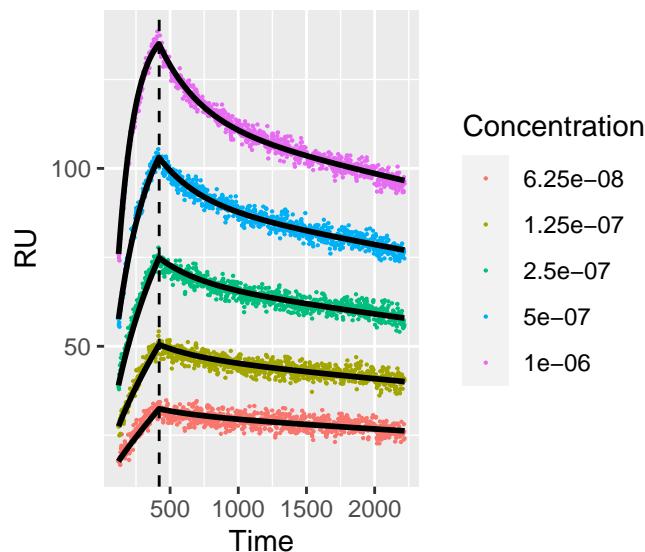
| | | |
|----------|----------|----------|
| $ka1$ | 1.64e+03 | 4.18e+01 |
| $ka2$ | 8.23e-05 | 2.88e-06 |
| $kd1$ | 2.52e-03 | 9.68e-05 |
| $kd2$ | 3.39e-05 | 1.24e-06 |
| $Rmax$ 1 | 3.32e+02 | 8.30e+00 |
| $Rmax$ 2 | 3.03e+02 | 1.02e+01 |
| $Rmax$ 3 | 2.54e+02 | 8.24e+00 |
| $Rmax$ 4 | 2.27e+02 | 9.32e+00 |
| $Rmax$ 5 | 2.27e+02 | 5.60e+00 |
| p 1 | 1.00e+00 | NA |
| p 2 | 0.00e+00 | 1.69e-01 |
| p 3 | 0.00e+00 | 1.23e-01 |
| p 4 | 1.02e-02 | 1.17e-01 |

CH505



CH505

Bivalent Analyte Model–1 with Extended Length of Dissociation

 k_{a1} 2.08e+03 5.08e+01 k_{a2} 9.81e-05 3.26e-06 k_{d1} 3.41e-03 1.21e-04 k_{d2} 4.53e-05 1.08e-06 R_{max} 1 2.50e+02 1.04e+01 R_{max} 2 2.22e+02 7.83e+00 R_{max} 3 2.12e+02 6.61e+00 R_{max} 4 1.95e+02 7.66e+00 R_{max} 5 2.01e+02 4.93e+00 p 1 0.00e+00 1.76e-01 p 2 0.00e+00 1.16e-01 p 3 0.00e+00 1.03e-01 p 4 0.00e+00 1.09e-01

CH505

