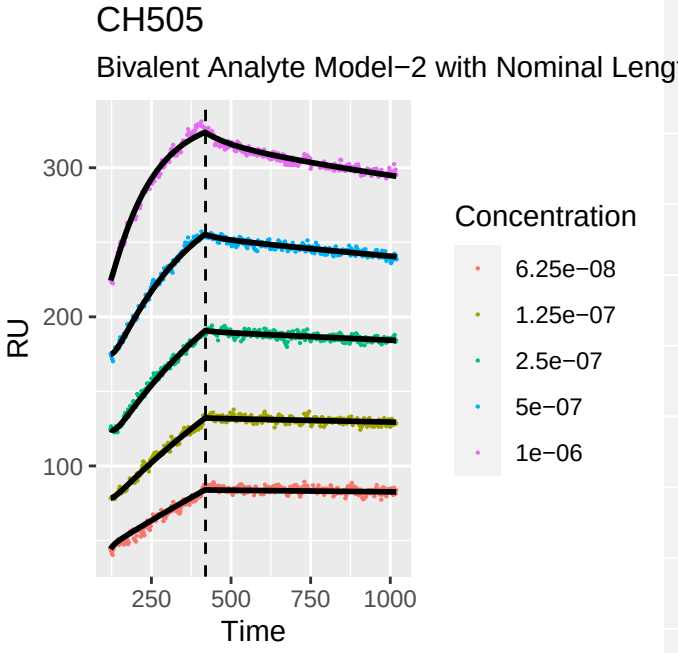
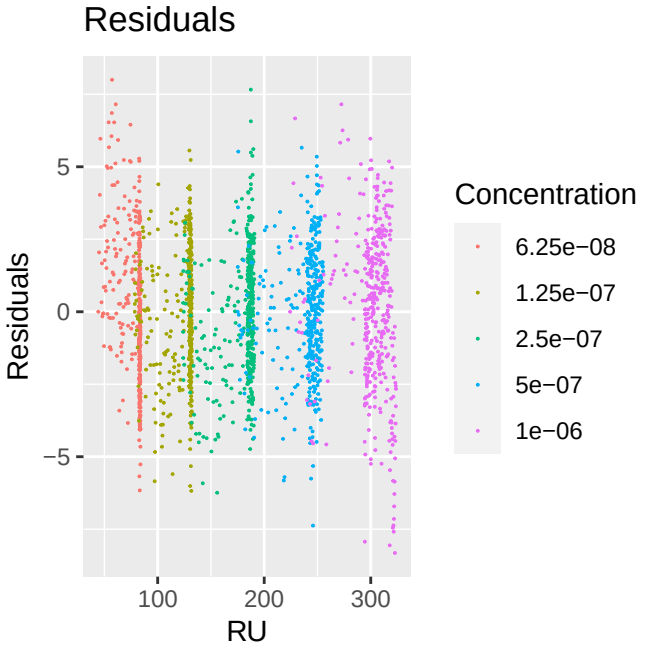
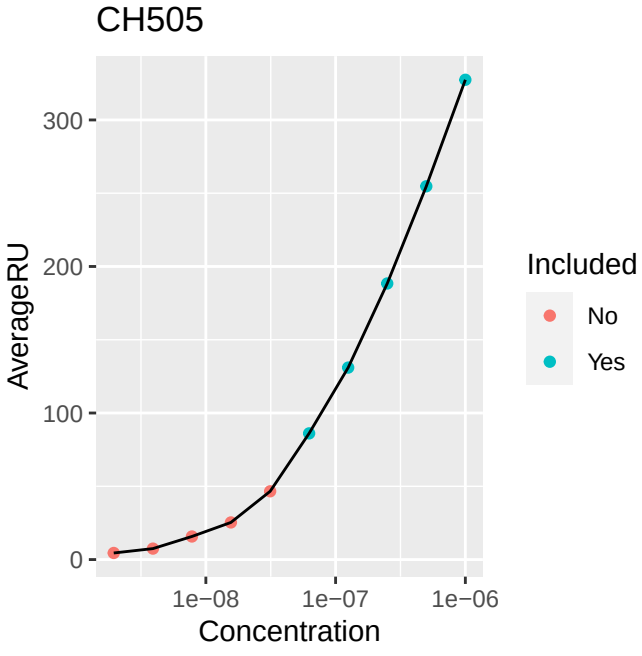


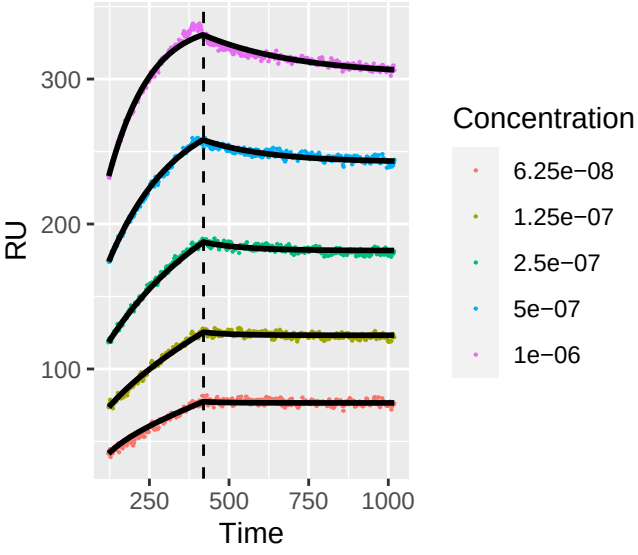
	Estimate	Std. Error
ka1	1.95e+03	35.3400501
ka2	1.47e−04	0.0000137
kd1	5.34e−03	0.0004421
kd2	1.96e−04	0.0000127
Rmax	6.02e+02	8.6025689
AL20 1	8.35e+01	3.7312787
AL20 2	3.38e+01	5.3043565
AL20 3	0.00e+00	7.0342862
AL20 4	1.50e+01	8.1949459
AL20 5	7.77e+01	9.1715230





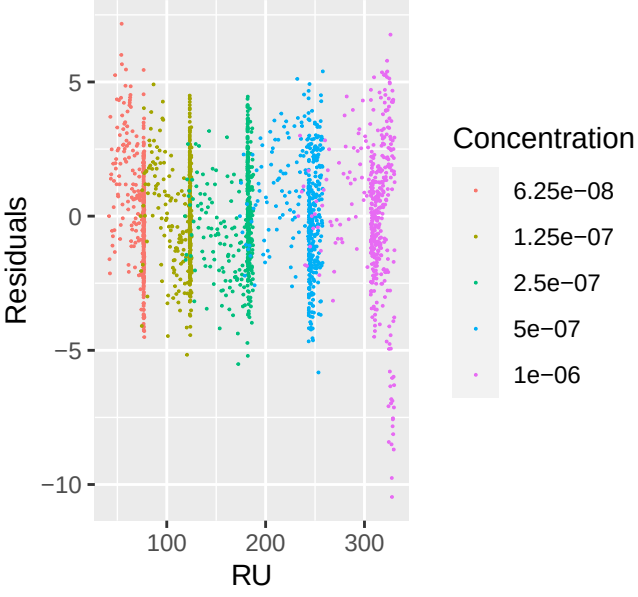
CH505

Bivalent Analyte Model−2 with Nominal Length of Dissociation

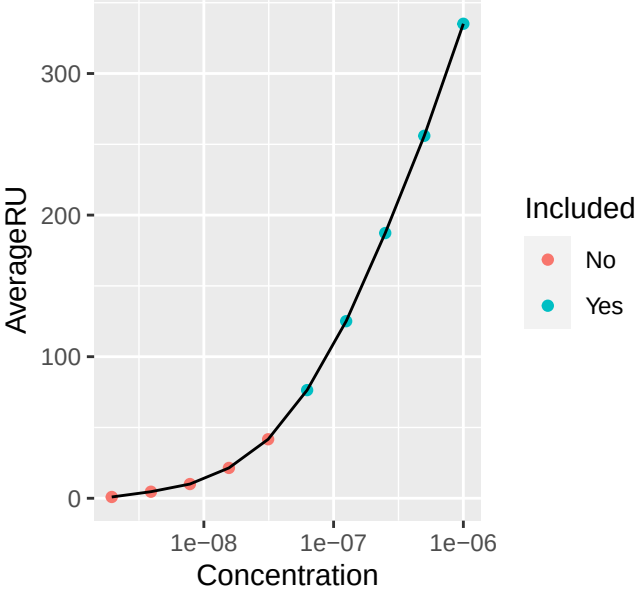


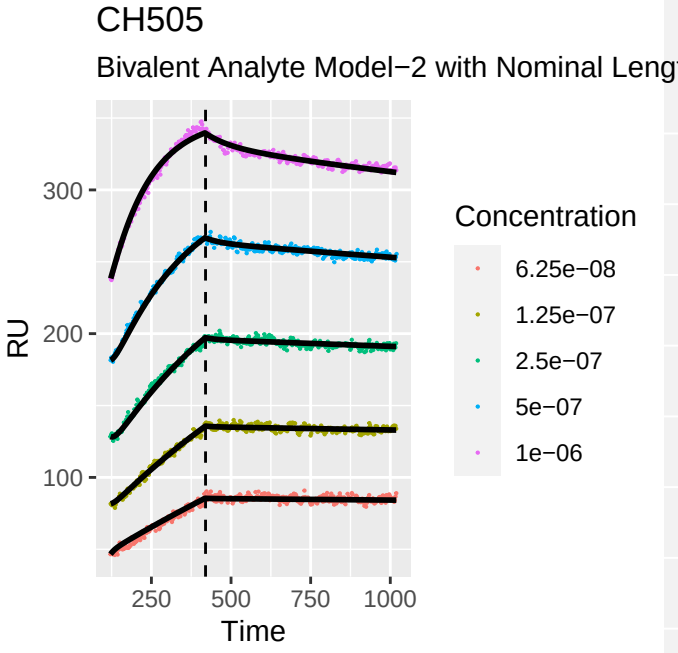
	Estimate	Std. Error
ka1	2.53e+03	4.51e+01
ka2	3.95e−05	3.23e−06
kd1	1.90e−03	9.08e−05
kd2	0.00e+00	1.01e−05
Rmax	3.81e+02	4.22e+00
AL20 1	9.46e+01	1.87e+00
AL20 2	1.04e+02	2.42e+00
AL20 3	1.25e+02	3.05e+00
AL20 4	1.75e+02	4.09e+00
AL20 5	2.64e+02	5.35e+00

Residuals

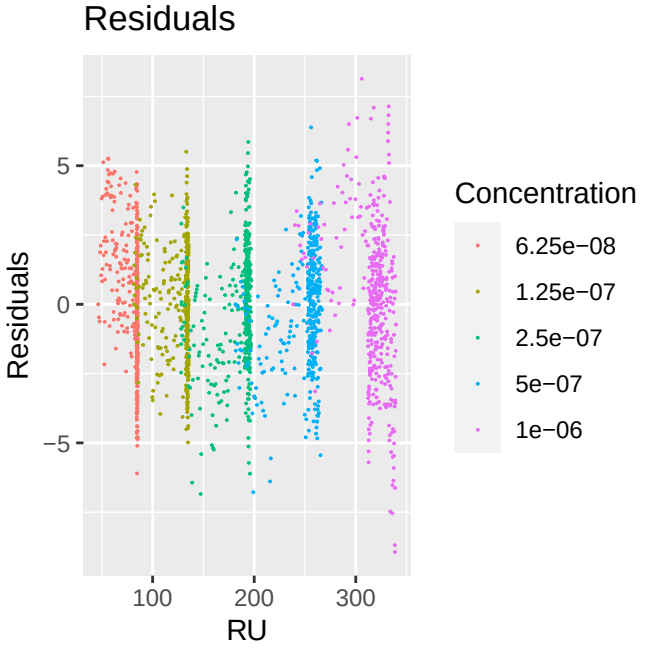


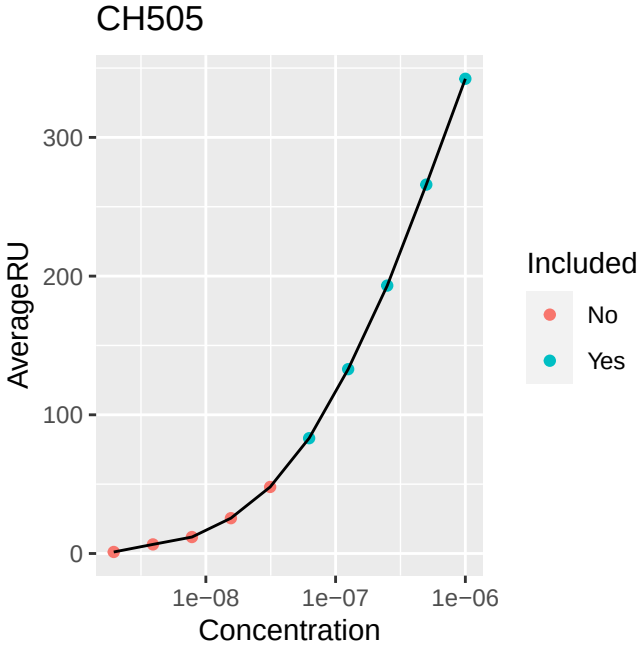
CH505





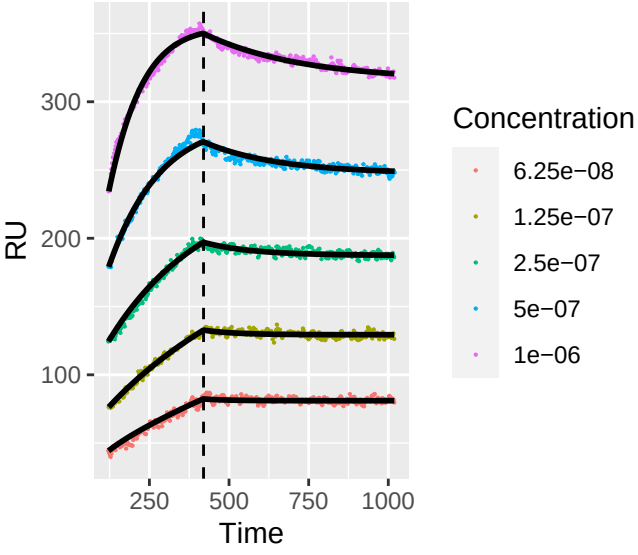
	Estimate	Std. Error
ka1	2.10e+03	3.38e+01
ka2	1.26e−04	9.33e−06
kd1	6.13e−03	4.89e−04
kd2	1.22e−04	7.17e−06
Rmax	5.76e+02	5.94e+00
AL20 1	7.50e+01	2.65e+00
AL20 2	4.82e+01	3.68e+00
AL20 3	3.38e+01	5.02e+00
AL20 4	6.28e+01	5.91e+00
AL20 5	1.34e+02	6.69e+00





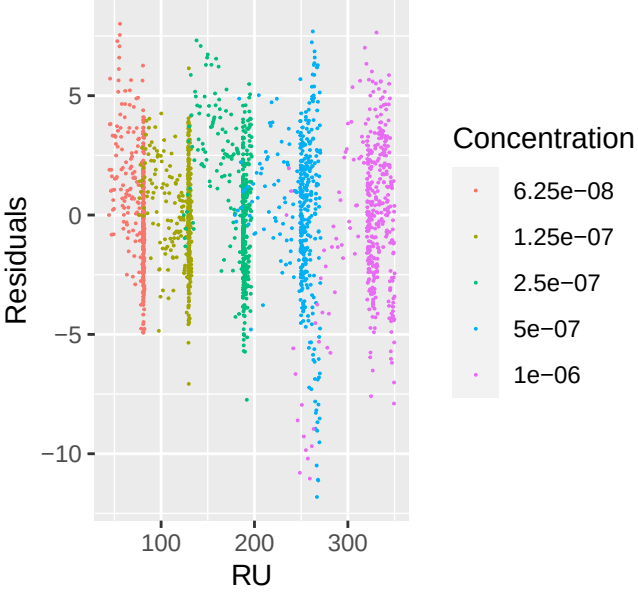
CH505

Bivalent Analyte Model−2 with Nominal Length of Dissociation

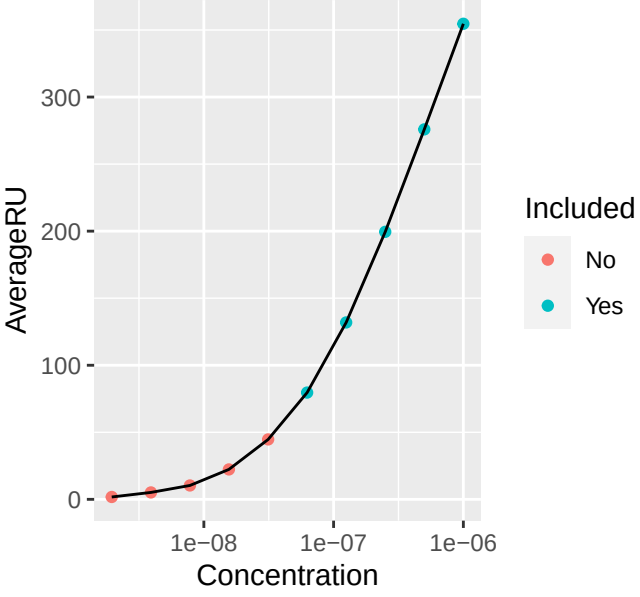


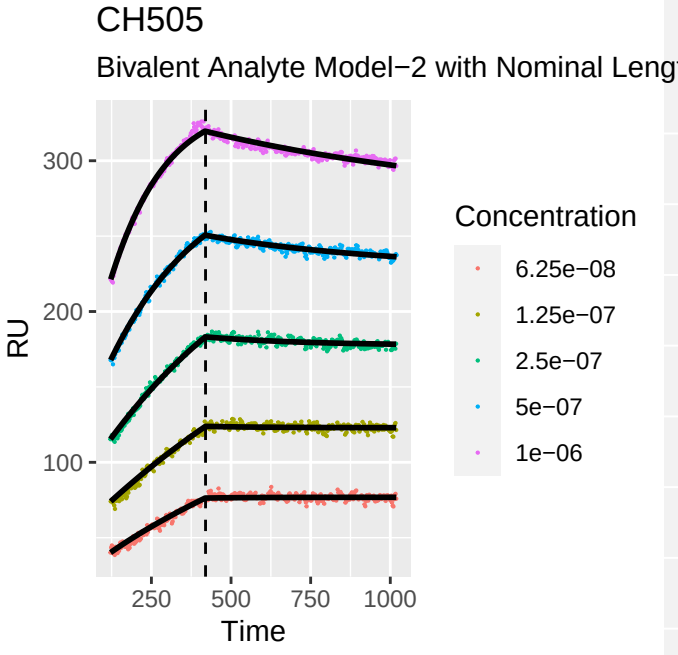
	Estimate	Std. Error
ka1	3.31e+03	6.18e+01
ka2	3.88e−05	3.22e−06
kd1	2.01e−03	1.41e−04
kd2	0.00e+00	1.26e−05
Rmax	3.90e+02	5.61e+00
AL20 1	6.22e+01	2.64e+00
AL20 2	7.42e+01	3.97e+00
AL20 3	9.80e+01	5.57e+00
AL20 4	1.63e+02	7.29e+00
AL20 5	2.81e+02	8.46e+00

Residuals

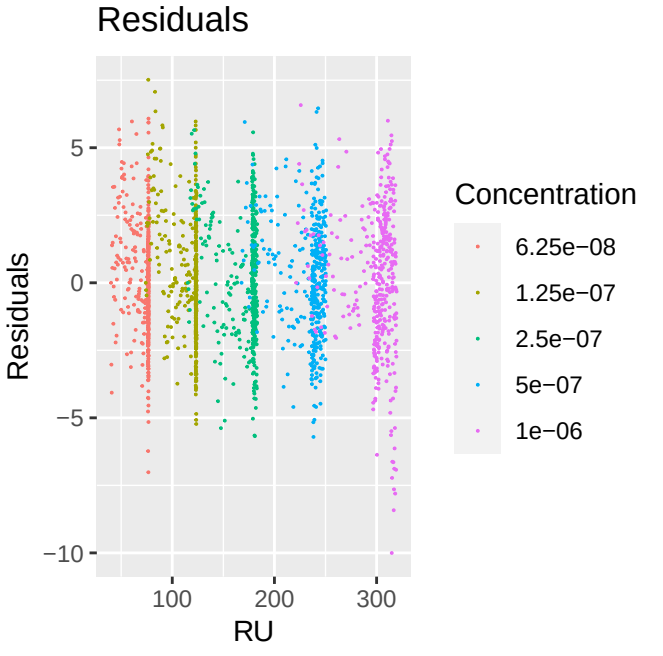


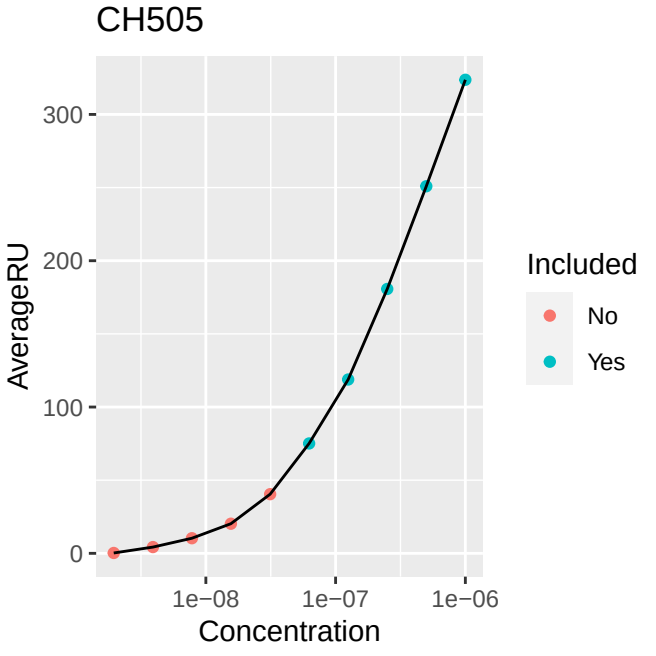
CH505





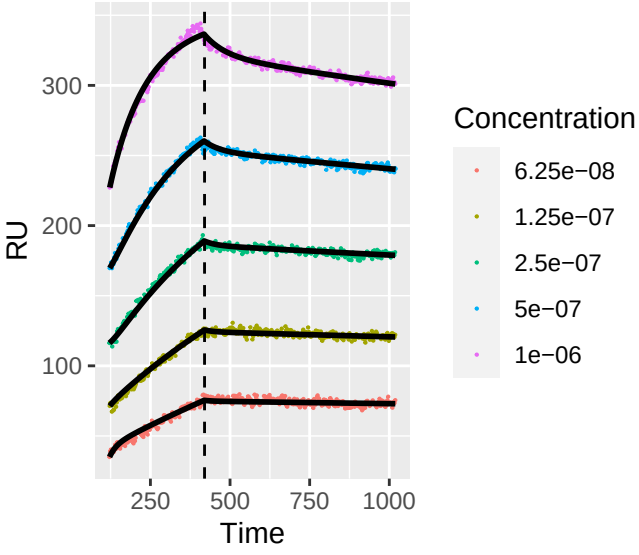
	Estimate	Std. Error
ka1	2.00e+03	3.30e+01
ka2	1.45e−05	9.47e−07
kd1	4.63e−04	4.38e−05
kd2	0.00e+00	1.97e−05
Rmax	3.96e+02	3.50e+00
AL20 1	1.66e+02	6.71e+00
AL20 2	1.40e+02	2.82e+00
AL20 3	1.25e+02	4.62e+00
AL20 4	1.22e+02	7.91e+00
AL20 5	1.75e+02	1.04e+01





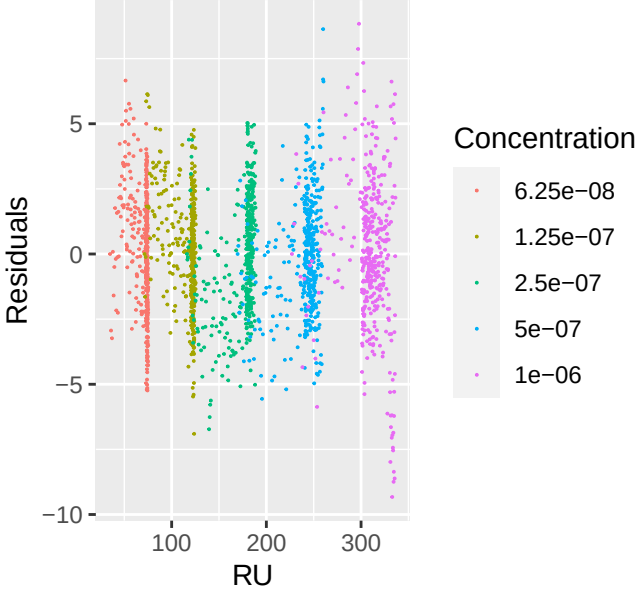
CH505

Bivalent Analyte Model−2 with Nominal Length of Dissociation



	Estimate	Std. Error
ka1	2.26e+03	3.15e+01
ka2	7.88e−05	3.60e−06
kd1	9.04e−03	6.14e−04
kd2	1.06e−04	4.31e−06
Rmax	5.32e+02	4.38e+00
AL20 1	7.44e+01	2.11e+00
AL20 2	7.38e+01	1.59e+00
AL20 3	8.59e+01	1.73e+00
AL20 4	1.20e+02	2.06e+00
AL20 5	1.91e+02	2.68e+00

Residuals



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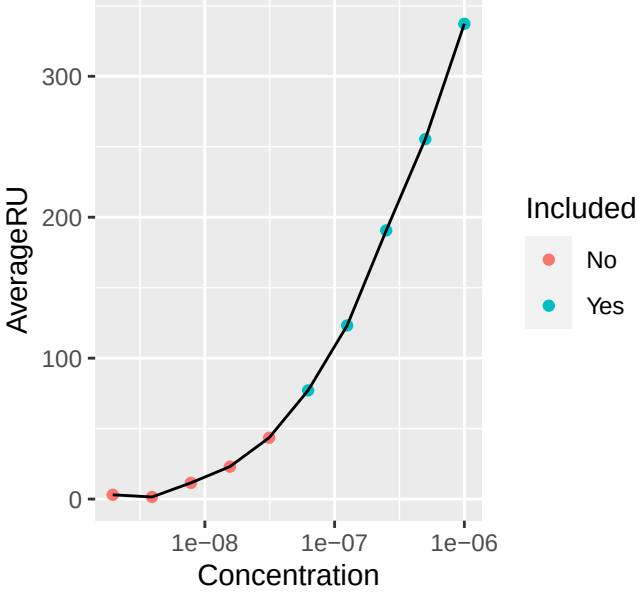
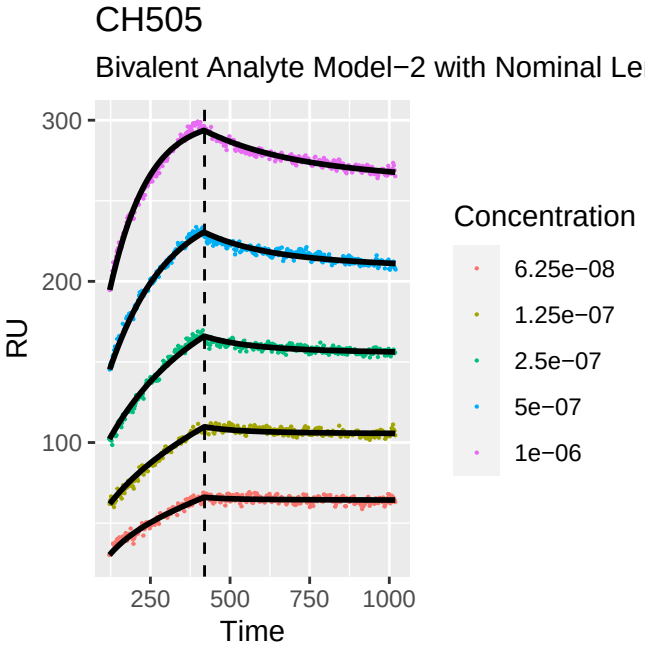
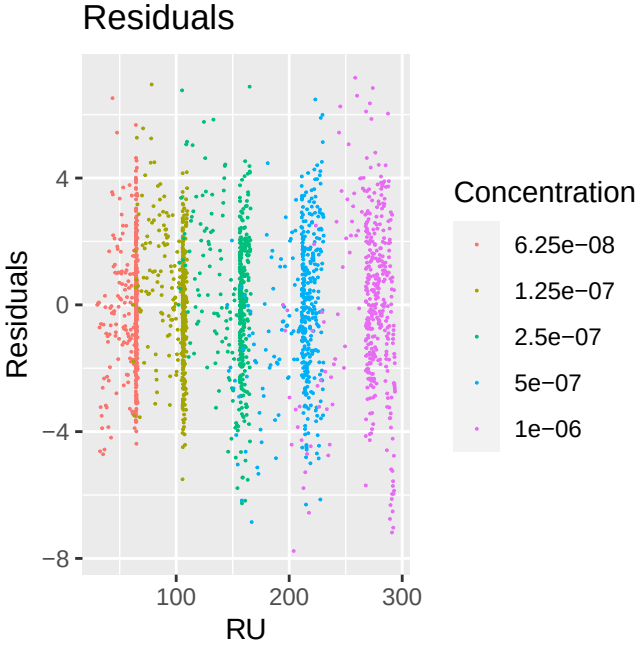
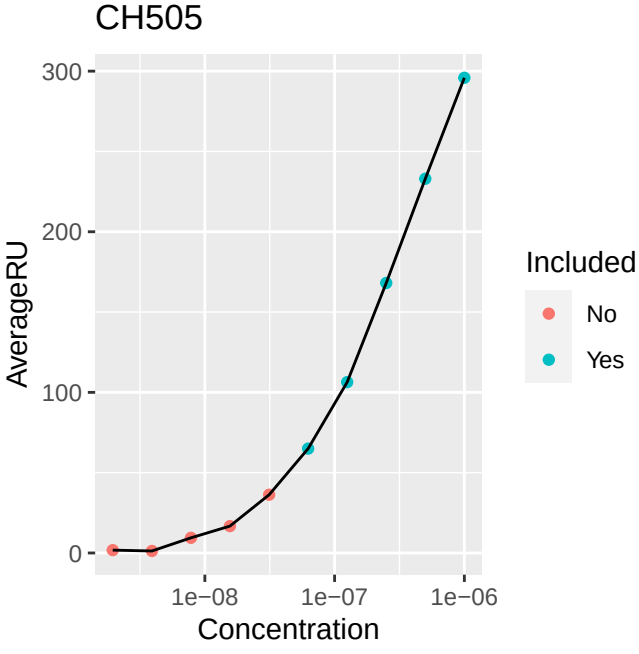
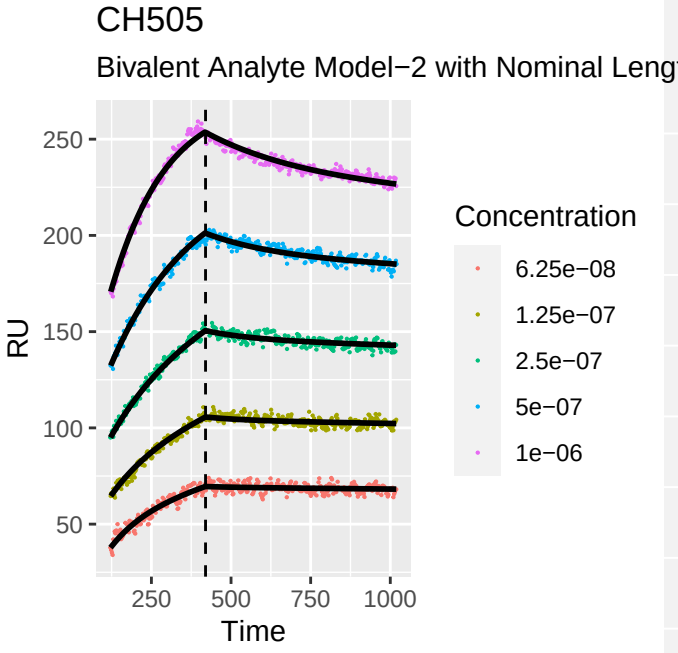




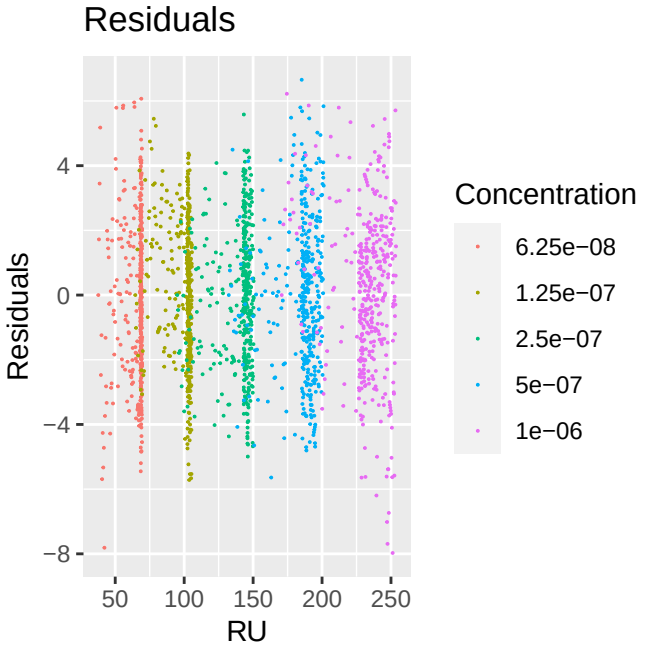
	Estimate	Std. Error
ka1	2.91e+03	5.93e+01
ka2	3.77e−05	2.76e−06
kd1	2.52e−03	8.31e−05
kd2	2.66e−05	8.36e−06
Rmax	3.18e+02	3.72e+00
AL20 1	8.01e+01	1.55e+00
AL20 2	9.01e+01	2.06e+00
AL20 3	1.19e+02	2.48e+00
AL20 4	1.77e+02	3.09e+00
AL20 5	2.59e+02	4.11e+00

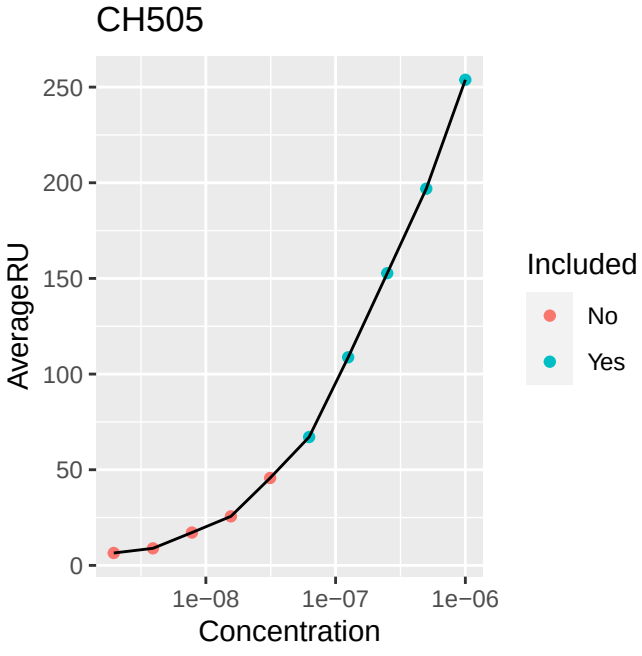


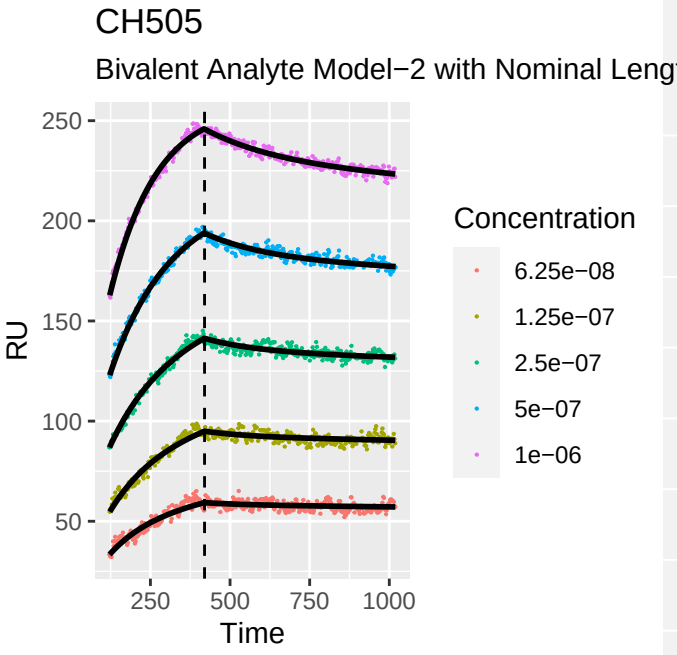




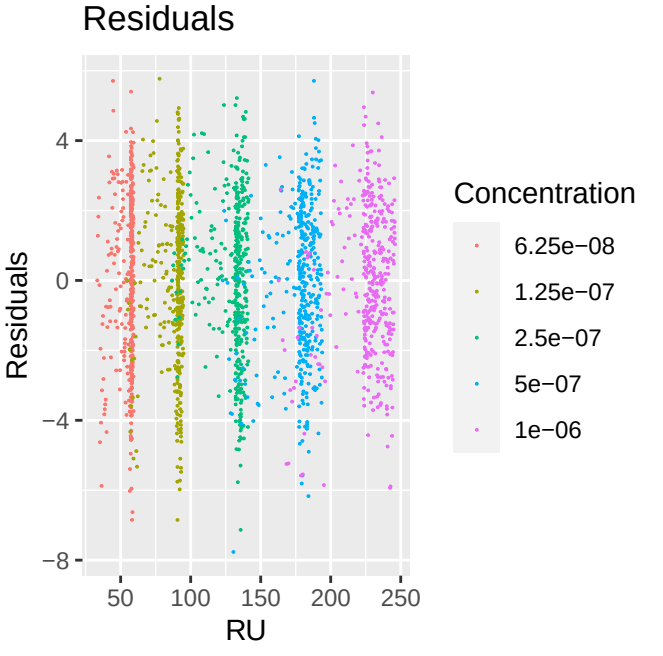
	Estimate	Std. Error
ka1	1.81e+03	7.54e+01
ka2	2.52e−05	2.84e−06
kd1	2.02e−03	7.49e−05
kd2	5.47e−05	1.07e−05
Rmax	3.08e+02	4.90e+00
AL20 1	1.14e+02	3.97e+00
AL20 2	1.26e+02	1.95e+00
AL20 3	1.46e+02	2.29e+00
AL20 4	1.68e+02	2.68e+00
AL20 5	2.08e+02	2.74e+00

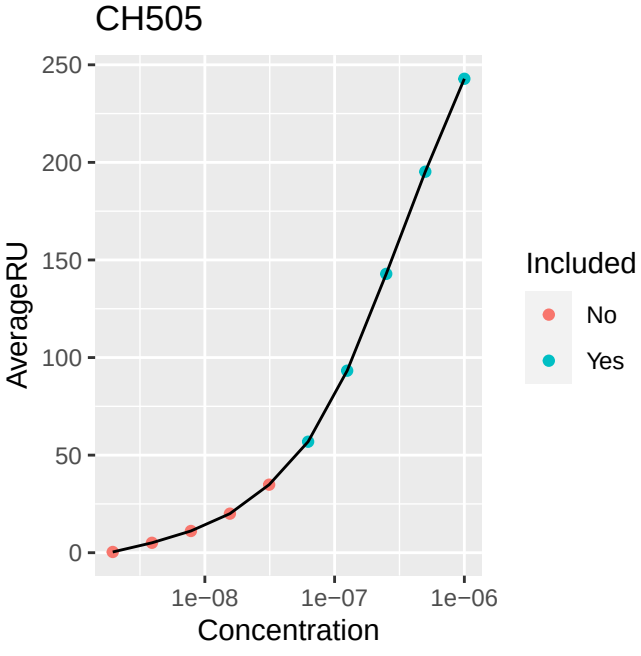






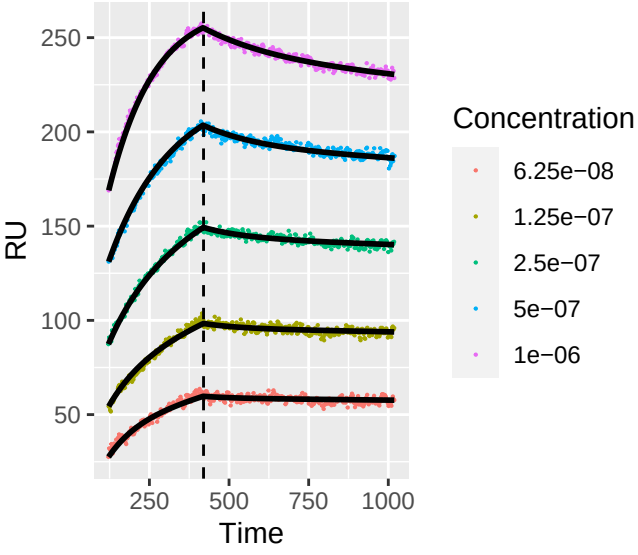
	Estimate	Std. Error
ka1	1.75e+03	1.27e+02
ka2	1.67e−05	4.09e−06
kd1	3.37e−03	1.46e−04
kd2	3.38e−05	6.75e−06
Rmax	2.86e+02	6.34e+00
AL20 1	6.98e+01	2.16e+00
AL20 2	1.04e+02	1.91e+00
AL20 3	1.39e+02	1.41e+00
AL20 4	1.81e+02	1.64e+00
AL20 5	2.29e+02	1.80e+00





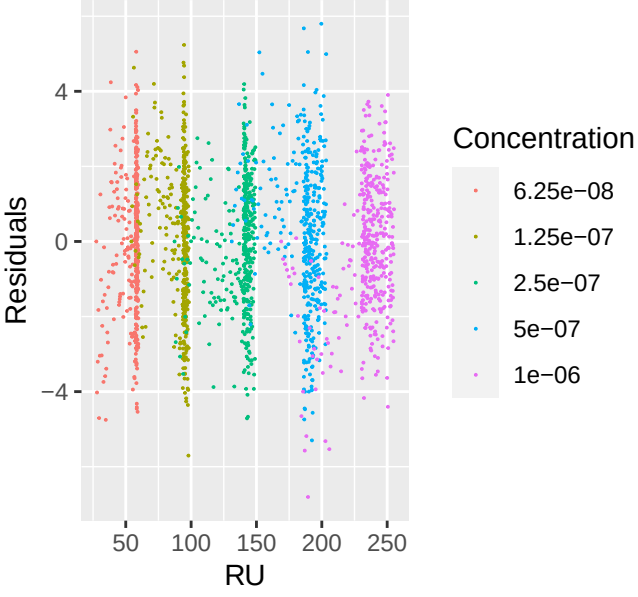
CH505

Bivalent Analyte Model−2 with Nominal Length of Dissociation

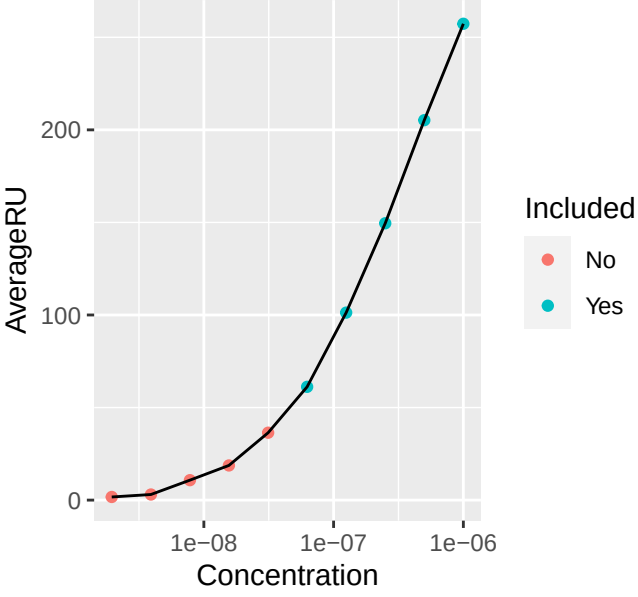


	Estimate	Std. Error
ka1	2.09e+03	7.35e+01
ka2	2.96e−05	3.09e−06
kd1	2.97e−03	5.57e−05
kd2	5.42e−05	7.27e−06
Rmax	2.84e+02	4.29e+00
AL20 1	8.84e+01	2.21e+00
AL20 2	1.14e+02	1.46e+00
AL20 3	1.51e+02	1.46e+00
AL20 4	1.86e+02	1.47e+00
AL20 5	2.38e+02	2.45e+00

Residuals

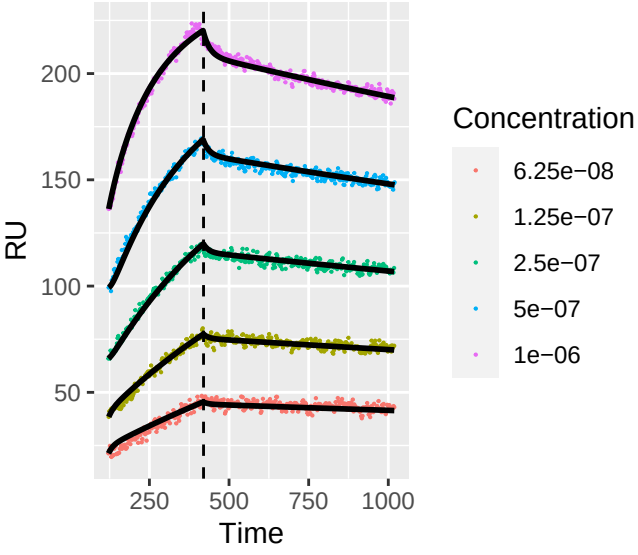


CH505



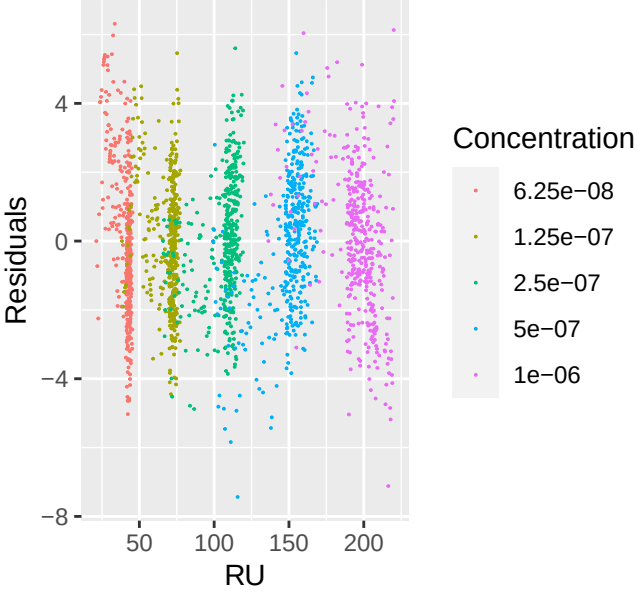
CH505

Bivalent Analyte Model−2 with Nominal Length of Dissociation

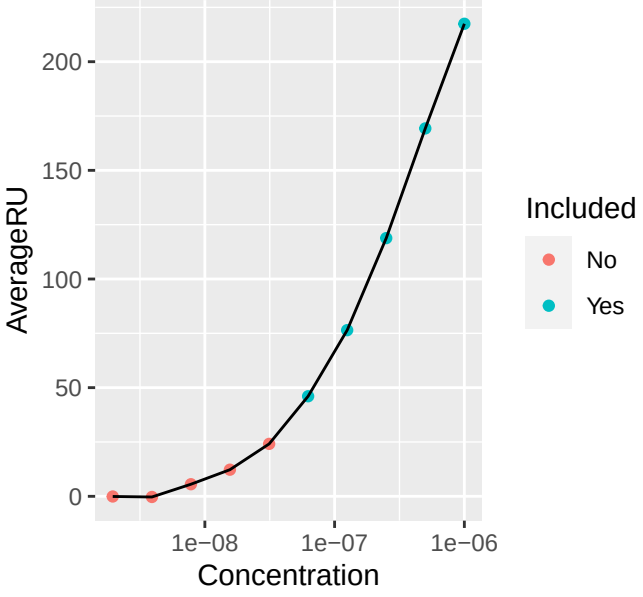


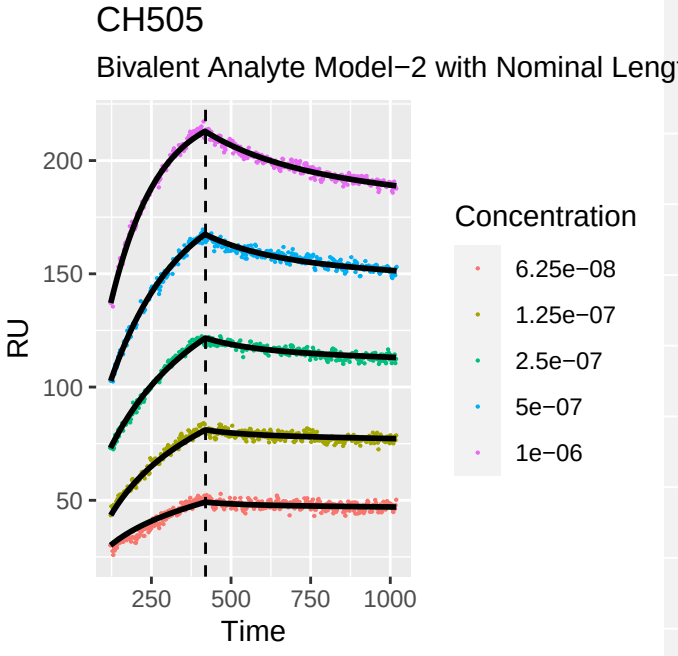
	Estimate	Std. Error
ka1	3.14e+03	1.30e+02
ka2	9.40e−05	3.50e−06
kd1	3.77e−02	3.09e−03
kd2	1.13e−04	3.00e−06
Rmax	4.28e+02	3.20e+00
AL20 1	2.60e+01	5.20e−01
AL20 2	4.00e+01	5.97e−01
AL20 3	5.41e+01	6.13e−01
AL20 4	7.43e+01	8.17e−01
AL20 5	1.08e+02	1.22e+00

Residuals

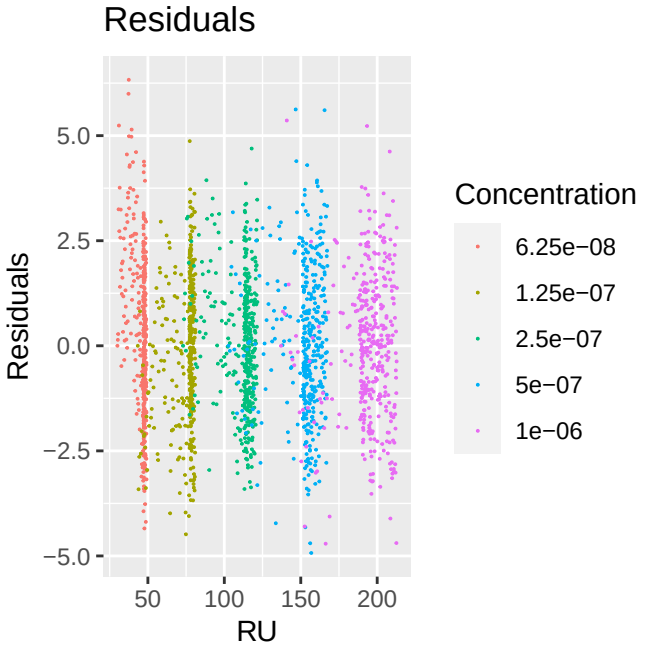


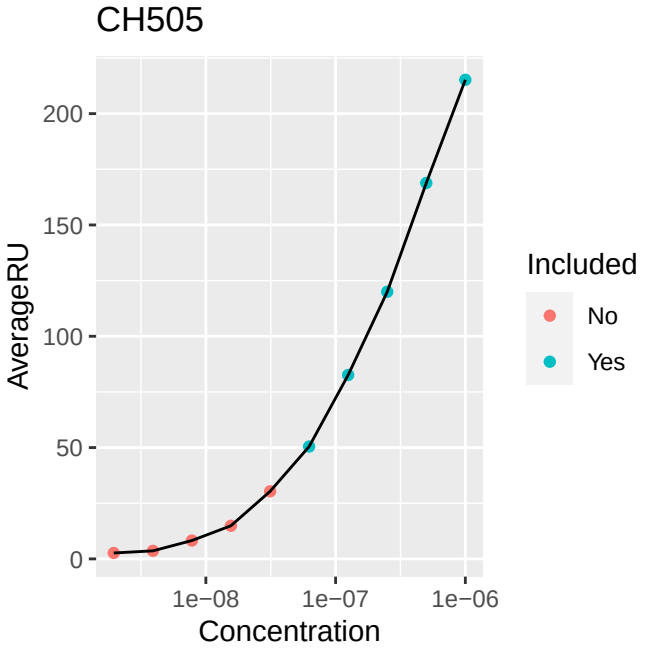
CH505





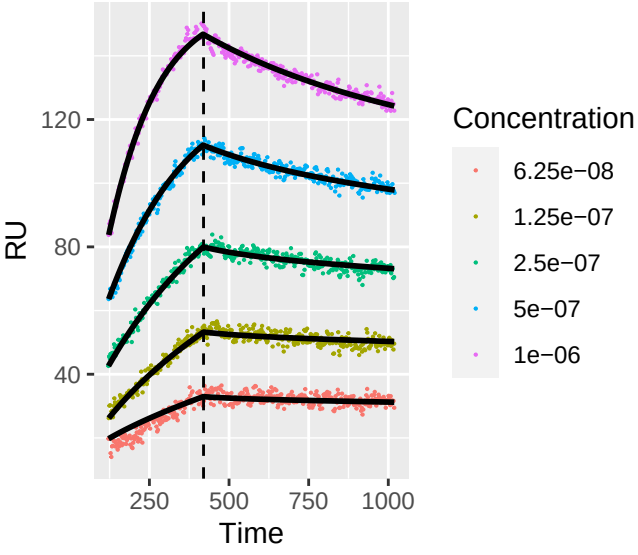
	Estimate	Std. Error
ka1	2.09e+03	7.29e+01
ka2	2.95e−05	3.63e−06
kd1	2.64e−03	6.41e−05
kd2	5.19e−05	1.07e−05
Rmax	2.47e+02	3.79e+00
AL20 1	5.19e+01	8.76e−01
AL20 2	9.12e+01	1.51e+00
AL20 3	1.11e+02	1.47e+00
AL20 4	1.43e+02	1.59e+00
AL20 5	1.85e+02	2.30e+00





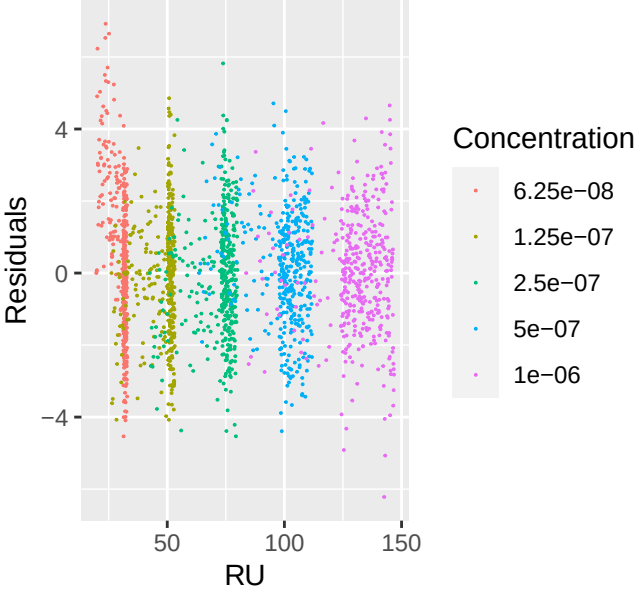
CH505

Bivalent Analyte Model−2 with Nominal Length of Dissociation

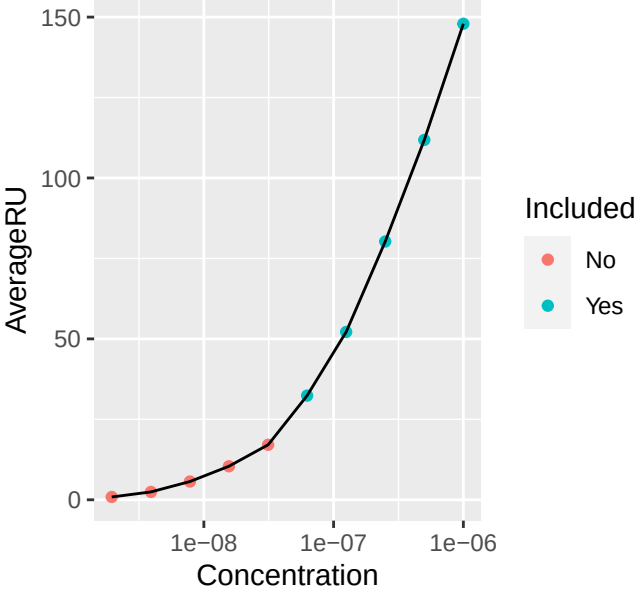


	Estimate	Std. Error
ka1	2.39e+03	83.9518885
ka2	4.42e−05	0.0000086
kd1	1.19e−03	0.0000895
kd2	1.98e−04	0.0000568
Rmax	1.77e+02	4.3148043
AL20 1	2.83e+01	1.5204061
AL20 2	5.43e+01	2.0567353
AL20 3	5.44e+01	2.9070057
AL20 4	6.43e+01	4.3120607
AL20 5	9.47e+01	5.8813628

Residuals

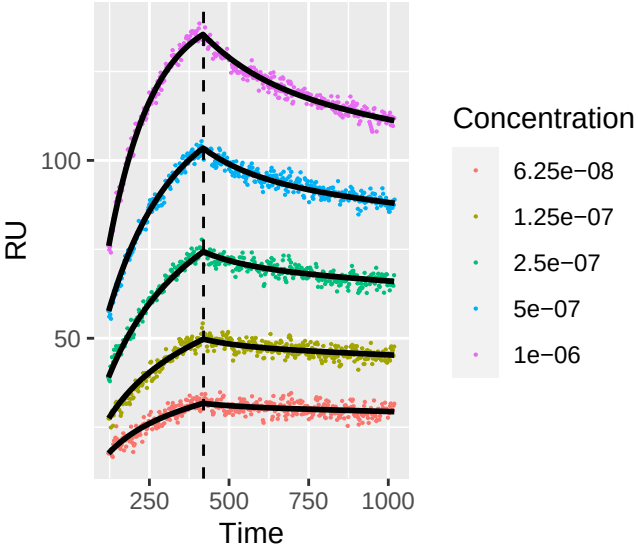


CH505



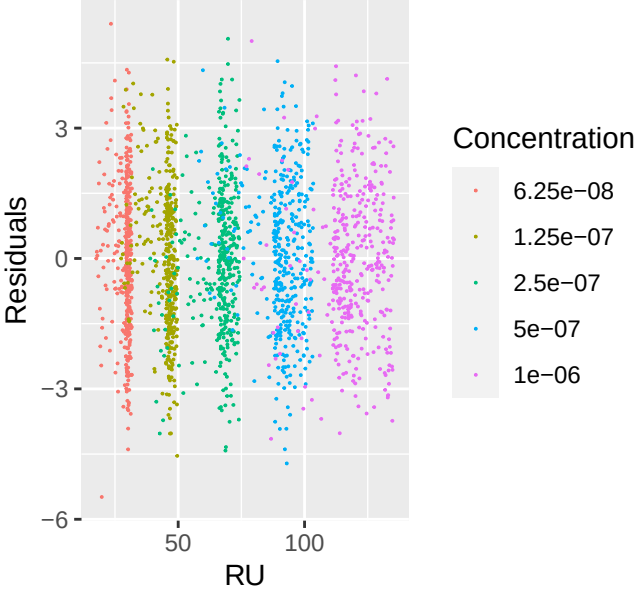
CH505

Bivalent Analyte Model−2 with Nominal Length of Dissociation



	Estimate	Std. Error
ka1	2.03e+03	9.09e+01
ka2	3.81e−05	5.23e−06
kd1	3.13e−03	9.03e−05
kd2	1.13e−04	1.81e−05
Rmax	1.69e+02	3.73e+00
AL20 1	3.56e+01	8.63e−01
AL20 2	4.94e+01	9.90e−01
AL20 3	6.76e+01	1.25e+00
AL20 4	8.27e+01	1.39e+00
AL20 5	1.09e+02	1.84e+00

Residuals



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