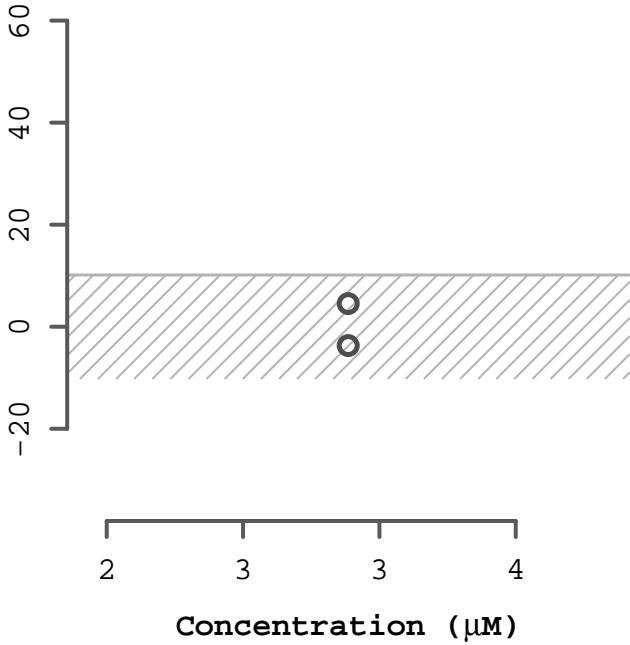


Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sodium benzoate
CHID: 20140 CASRN: 532-32-1
SPID(S): TP0001411G12
M4ID: 30204434

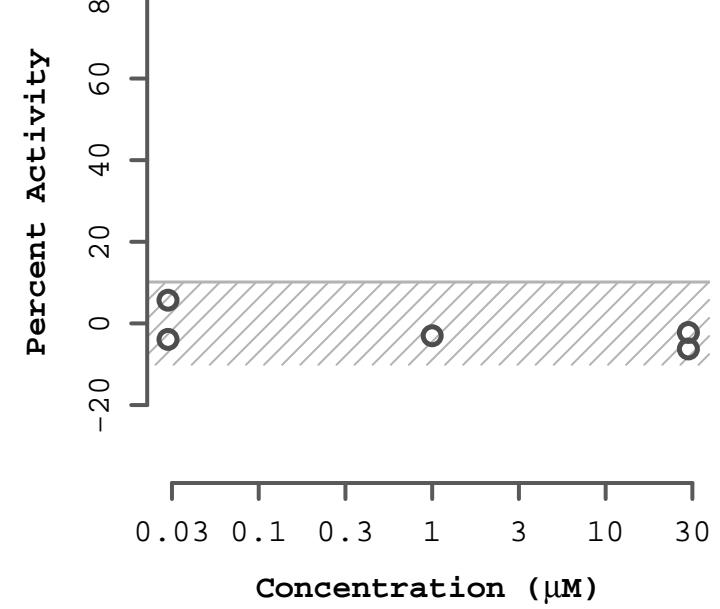
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	NA	NA	NA
PROB:	NA	NA	NA
RMSE:	NA	NA	NA

MAX_MEAN: 0.431 MAX_MED: 0.431 BMAD: 3.38
COFF: 10.1 HIT-CALL: -1 FITC: 2 ACTP: NA

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cadmium chloride
CHID: 20226 CASRN: 10108-64-2
SPID(S): EPAPLT0154E01
M4ID: 30204195

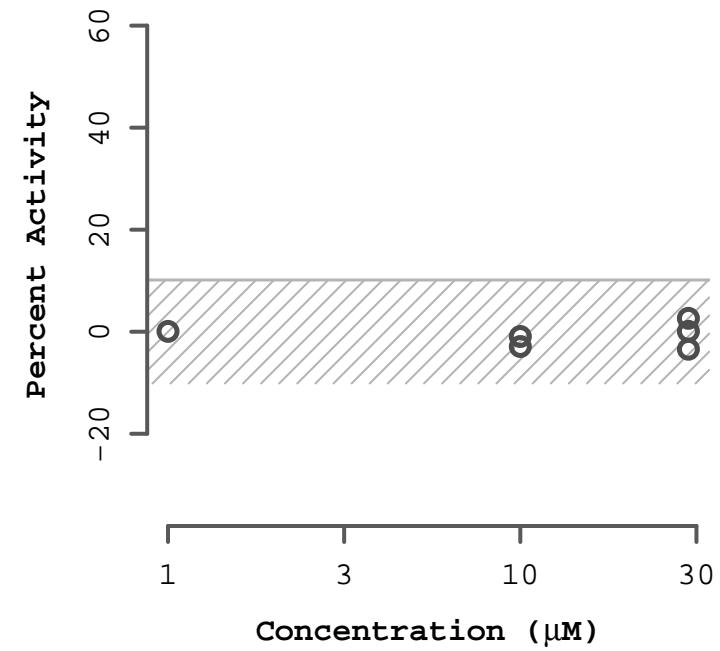
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	NA	NA	NA
PROB:	NA	NA	NA
RMSE:	NA	NA	NA

MAX_MEAN: 27.2 MAX_MED: 0.895 BMAD: 3.38
COFF: 10.1 HIT-CALL: -1 FITC: 2 ACTP: NA

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diphenolic acid
CHID: 22436 CASRN: 126-00-1
SPID(S): EPAPLT0154D01
M4ID: 30204184

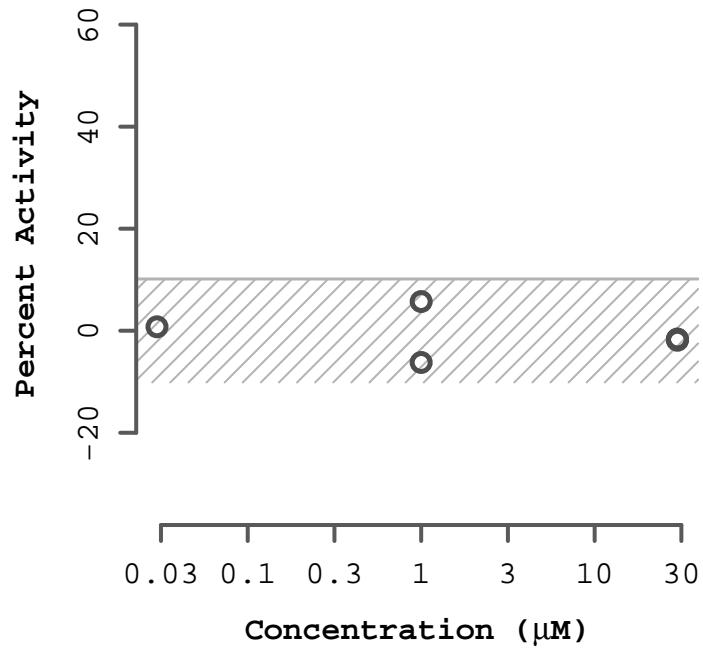
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	NA	NA	NA
PROB:	NA	NA	NA
RMSE:	NA	NA	NA

MAX_MEAN: 0.0716 MAX_MED: 0.0832 BMAD: 3.38
COFF: 10.1 HIT-CALL: -1 FITC: 2 ACTP: NA

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,4-Butanediol
CHID: 24666 CASRN: 110-63-4
SPID(S): EPAPLT0154E02
M4ID: 30204196

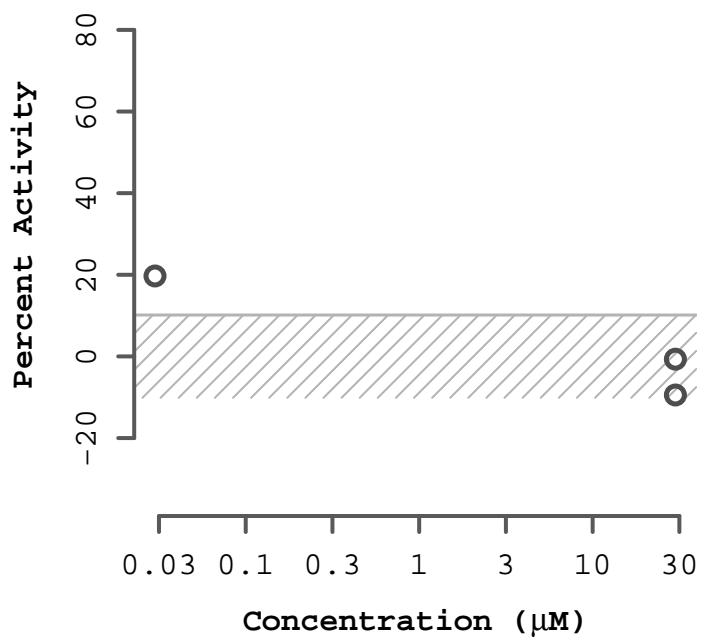
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	NA	NA	NA
PROB:	NA	NA	NA
RMSE:	NA	NA	NA

MAX_MEAN: 0.777 MAX_MED: 0.777 BMAD: 3.38
COFF: 10.1 HIT-CALL: -1 FITC: 2 ACTP: NA

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,2-Diphenoxylethane

CHID: 26706 CASRN: 104-66-5

SPID(S): EPAPLT0154A04

M4ID: 30204156 BRK

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
------	------	------

AIC: NA NA NA

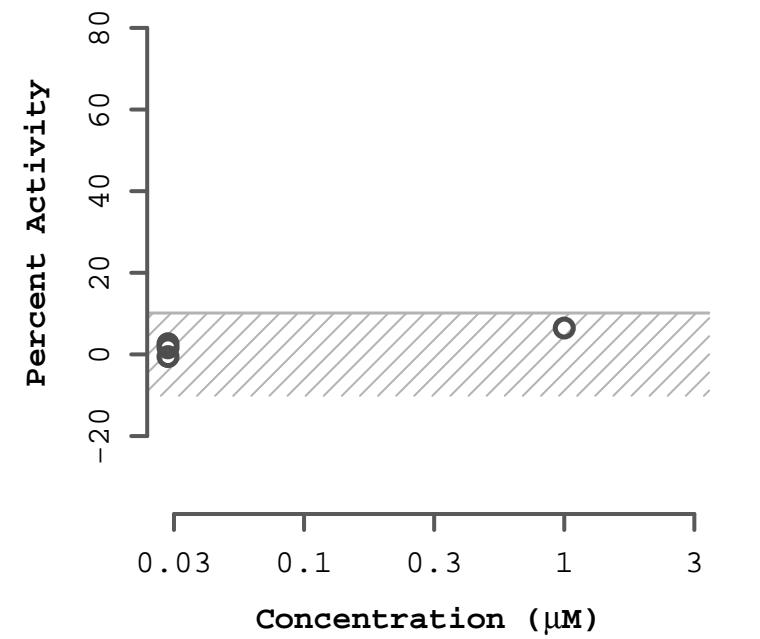
PROB: NA NA NA

RMSE: NA NA NA

MAX_MEAN: 29 MAX_MED: 19.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: -1 FITC: 2 ACTP: NA

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Denatonium saccharide
CHID: 34361 CASRN: 90823-38-4
SPID(S): EPAPLT0154H02
M4ID: 30204225 BRK

HILL MODEL: Not applicable.

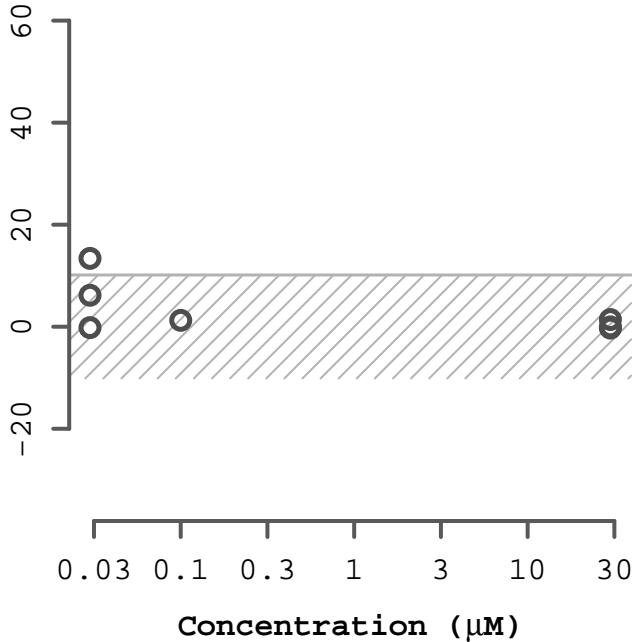
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	NA	NA	NA
PROB:	NA	NA	NA
RMSE:	NA	NA	NA

MAX_MEAN: 98.7 MAX_MED: 98.7 BMAD: 3.38
COFF: 10.1 HIT-CALL: -1 FITC: 2 ACTP: NA

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Trimethoxyphenylsilane
CHID: 40700 CASRN: 2996-92-1
SPID(S): EPAPLT0154H01
M4ID: 30204224

HILL MODEL: Not applicable.

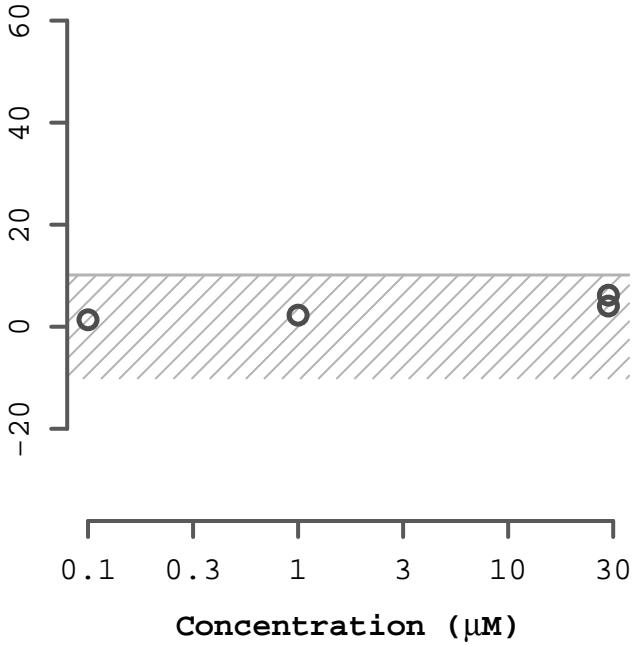
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	NA	NA	NA
PROB:	NA	NA	NA
RMSE:	NA	NA	NA

MAX_MEAN: 6.47 MAX_MED: 6.17 BMAD: 3.38
COFF: 10.1 HIT-CALL: -1 FITC: 2 ACTP: NA

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Methyl phenylacetate
CHID: 44352 CASRN: 101-41-7
SPID(S): EPAPLT0154D07
M4ID: 30204190

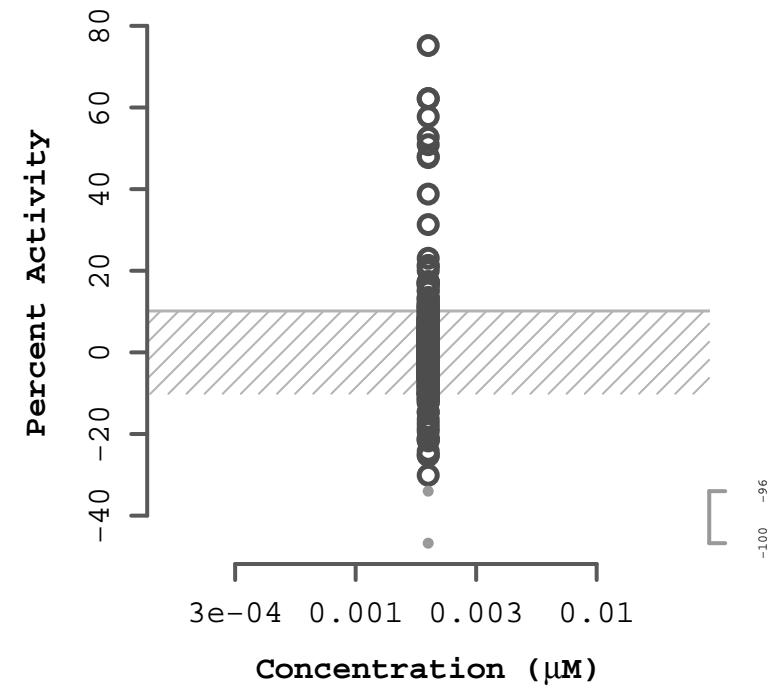
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	NA	NA	NA
PROB:	NA	NA	NA
RMSE:	NA	NA	NA

MAX_MEAN: 5.12 MAX_MED: 5.12 BMAD: 3.38
COFF: 10.1 HIT-CALL: -1 FITC: 2 ACTP: NA

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: DMSO (spid not in DB)

CHID: NA CASRN: NA

SPID(S): DMSO

M4ID: 30204152 BRK

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
------	------	------

AIC: NA	NA	NA
---------	----	----

PROB: NA	NA	NA
----------	----	----

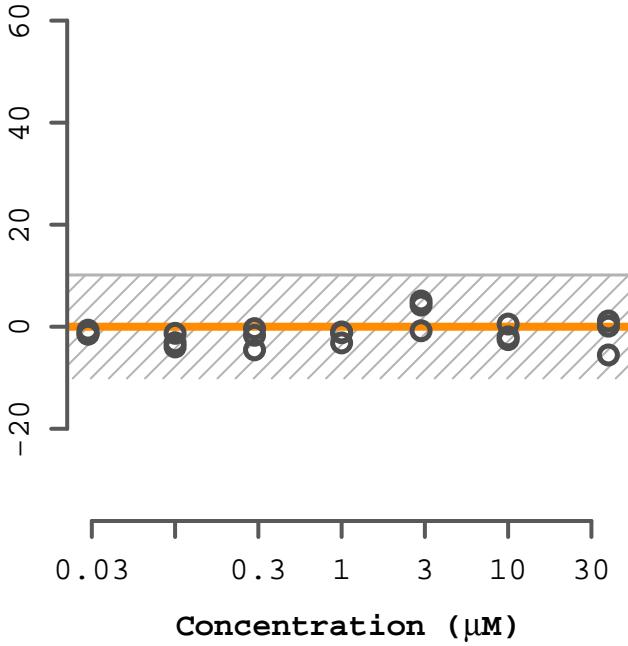
RMSE: NA	NA	NA
----------	----	----

MAX_MEAN: 0.353 MAX_MED: -0.0348 BMAD: 3.38

COFF: 10.1 HIT-CALL: -1 FITC: 2 ACTP: NA

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Acetamide

CHID: 20005 CASRN: 60-35-5

SPID(S): TP0001413E02

M4ID: 30204654

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 104.95

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 2.71

NA

NA

MAX_MEAN: 2.84

MAX_MED: 4.25

BMAD: 3.38

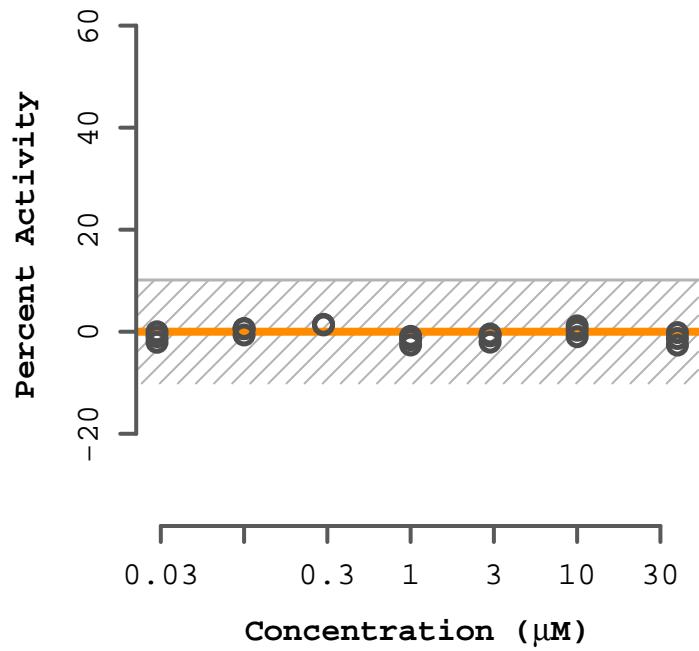
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Acetaminophen

CHID: 20006 CASRN: 103-90-2

SPID(S): TP0001413C09

M4ID: 30204425

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 72.68

HILL

PROB: 1

GNLS

RMSE: 1.34

MAX_MEAN: 1.4

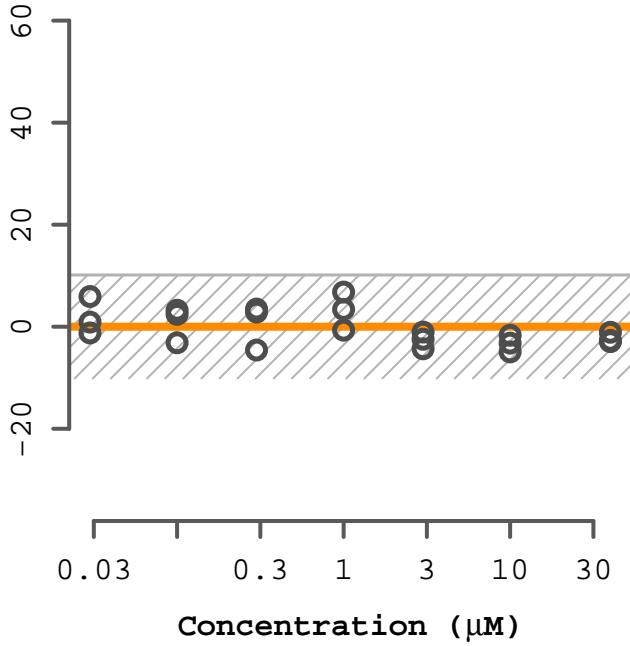
MAX_MED: 1.4

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Acifluorfen

CHID: 20022 CASRN: 50594-66-6

SPID(S): TP0001414F10

M4ID: 30204340

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 115.96

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 3.4

NA

NA

MAX_MEAN: 3.18

MAX_MED: 3.4

BMAD: 3.38

COFF: 10.1

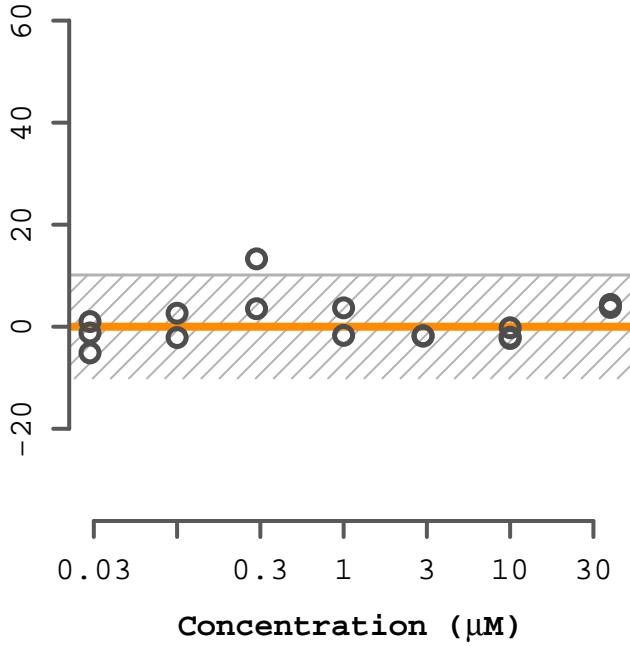
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Acrylamide

CHID: 20027 CASRN: 79-06-1

SPID(S): TP0001413A02

M4ID: 30204648

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 91.4

PROB: 1

RMSE: 4.3

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 8.42

MAX_MED: 8.42

BMAD: 3.38

COFF: 10.1

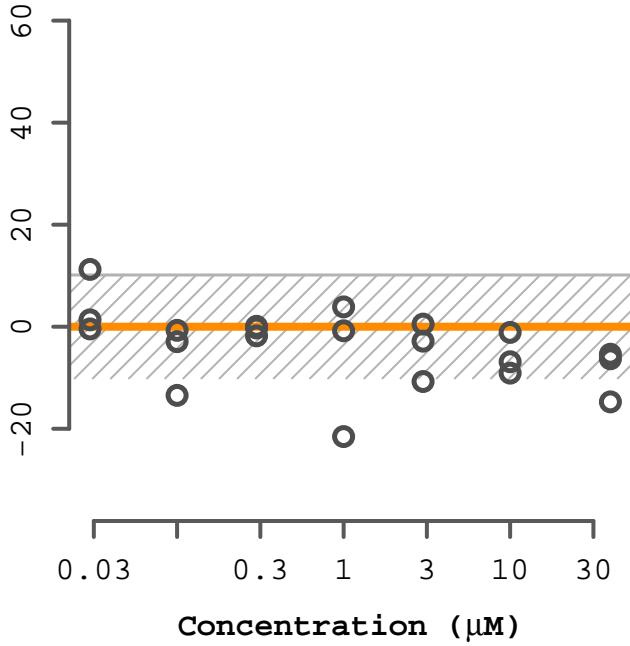
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Aldrin

CHID: 20040 CASRN: 309-00-2

SPID(S): TP0001412G10

M4ID: 30204462

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 147.51

PROB: 1

RMSE: 7.99

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 4.06

MAX_MED: 1.35

BMAD: 3.38

COFF: 10.1

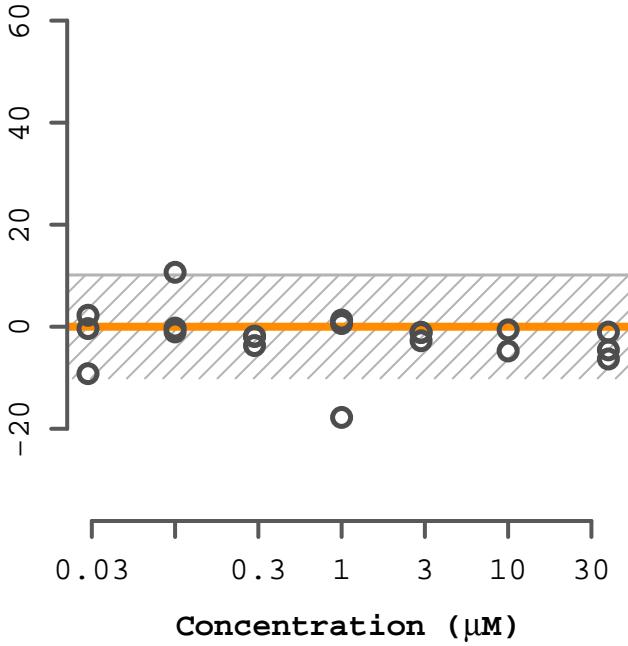
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Aminoanthraquinone
CHID: 20068 CASRN: 117-79-3
SPID(S): TP0001413C04
M4ID: 30204426

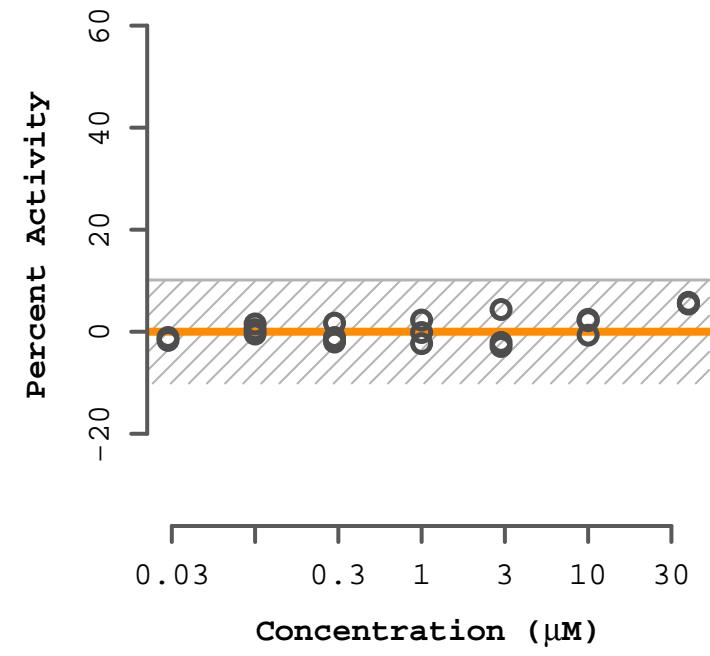
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 122.88	NA	NA
PROB: 1	NA	NA
RMSE: 5.65	NA	NA

MAX_MEAN: 3.14 MAX_MED: 0.631 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: (E)-Anethole

CHID: 20087 CASRN: 4180-23-8

SPID(S): TP0001414D06

M4ID: 30204437

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 101.86

NA

NA

PROB:

1

NA

NA

RMSE: 2.85

NA

NA

MAX_MEAN: 5.59

MAX_MED: 5.55

BMAD: 3.38

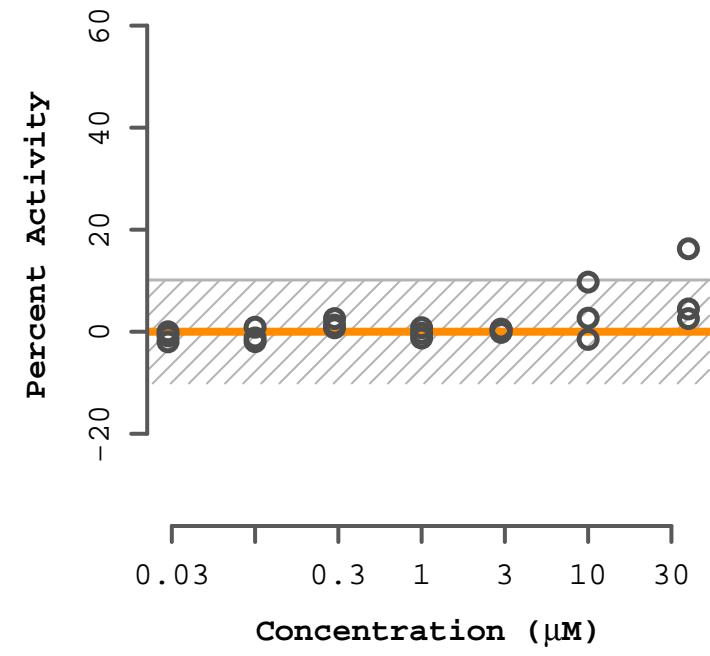
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Anilazine

CHID: 20089 CASRN: 101-05-3

SPID(S): TP0001412F05

M4ID: 30204328

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 110.88

PROB: 1

RMSE: 4.46

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 7.77

MAX_MED: 4.47

BMAD: 3.38

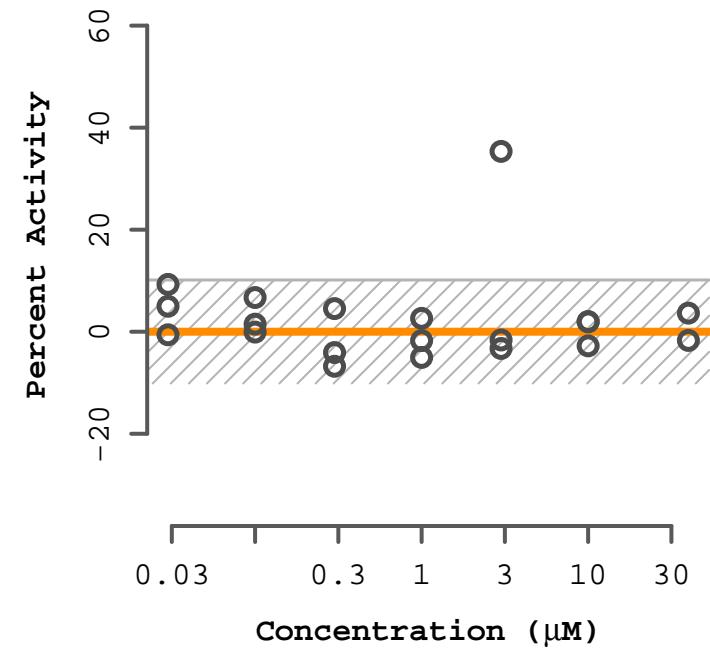
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sodium L-ascorbate
CHID: 20105 CASRN: 134-03-2
SPID(S): TP0001414H01
M4ID: 30204585

HILL MODEL: Not applicable.

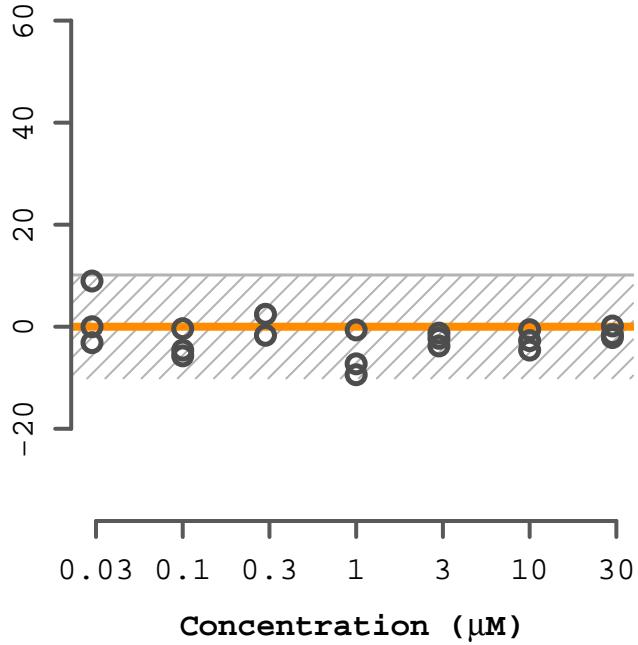
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 131.48	NA	NA
PROB: 1	NA	NA
RMSE: 8.86	NA	NA

MAX_MEAN: 10.2 MAX_MED: 5 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Aspartame

CHID: 20107 CASRN: 22839-47-0

SPID(S): EPAPLT0167D08

M4ID: 30204239

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 116.17

HILL

PROB: 1

NA

RMSE: 4.18

NA

GNLS

MAX_MEAN: 1.95

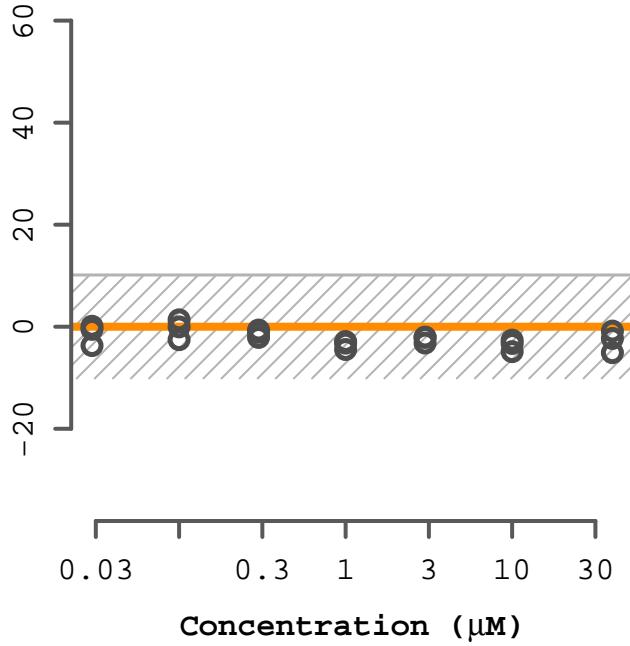
MAX_MED: 0.386

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Aspirin

CHID: 20108 CASRN: 50-78-2

SPID(S): TP0001414H08

M4ID: 30204445

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 106.86

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 2.75

NA

NA

MAX_MEAN: -0.398

MAX_MED: -0.0232

BMAD: 3.38

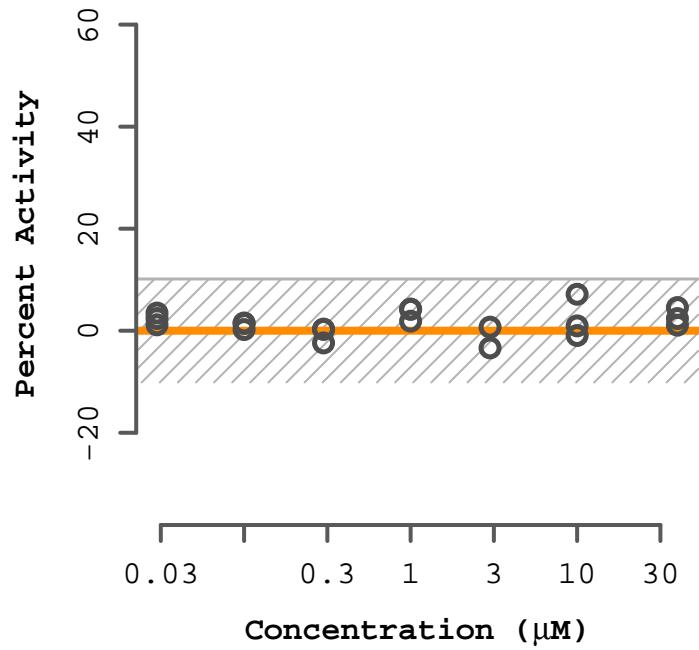
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Azathioprine

CHID: 20119 CASRN: 446-86-6

SPID(S): TP0001412G07

M4ID: 30204481

HILL MODEL: Not applicable.

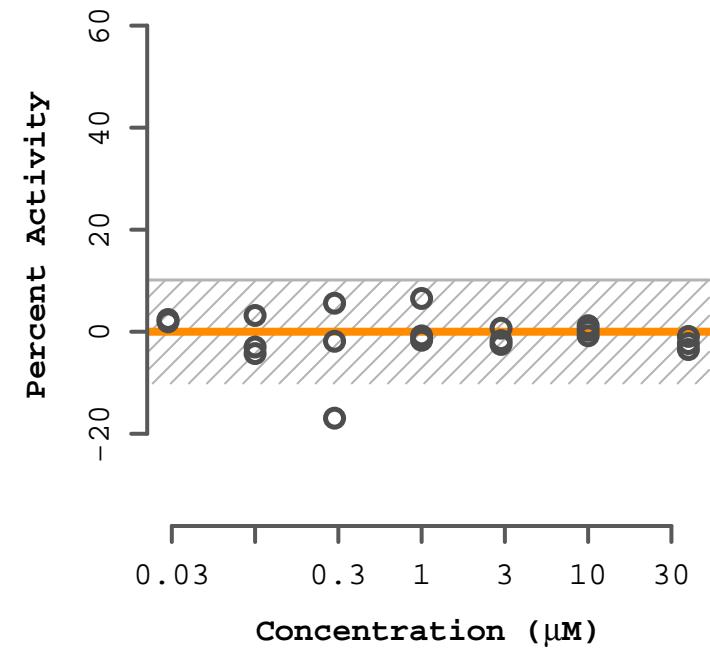
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	101.14	NA	NA
PROB:	1	NA	NA
RMSE:	2.84	NA	NA

MAX_MEAN: 3.45 MAX_MED: 4.22 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sodium azide

CHID: 20121 CASRN: 26628-22-8

SPID(S): TP0001413C02

M4ID: 30204649

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 113.51

NA

NA

PROB:

1

NA

NA

RMSE: 4.72

NA

NA

MAX_MEAN: 2.18

MAX_MED: 2.18

BMAD: 3.38

COFF: 10.1

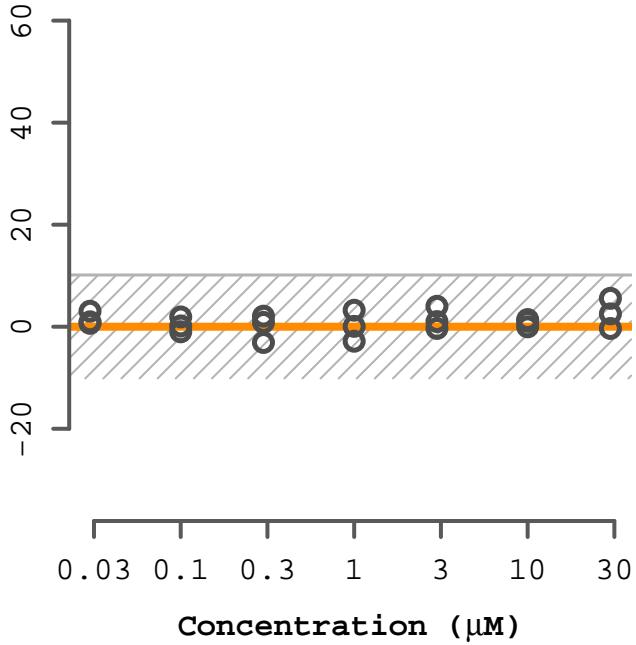
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Azinphos-methyl
CHID: 20122 CASRN: 86-50-0
SPID(S): EPAPLT0167C10
M4ID: 30204240

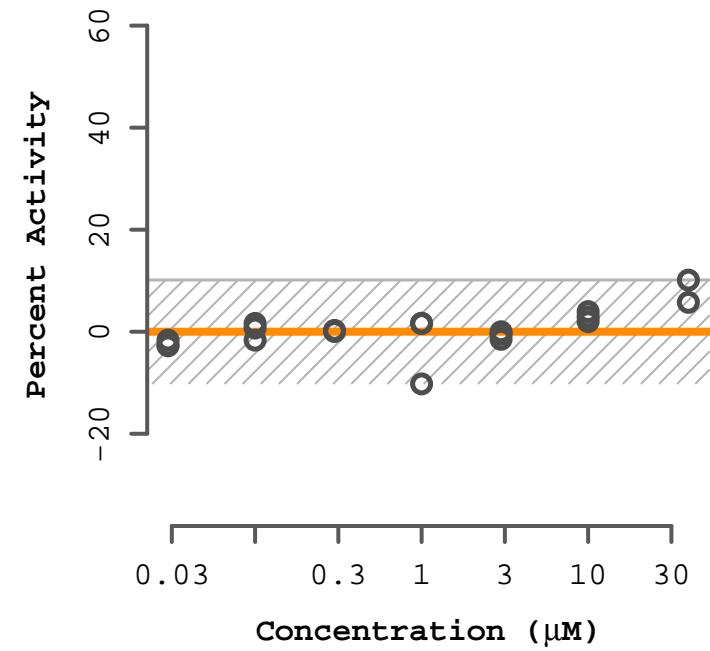
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 96.32	NA	NA
PROB: 1	NA	NA
RMSE: 2.25	NA	NA

MAX_MEAN: 2.58 MAX_MED: 2.53 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Azinphos-methyl
CHID: 20122 CASRN: 86-50-0
SPID(S): TP0001413E09
M4ID: 30204439

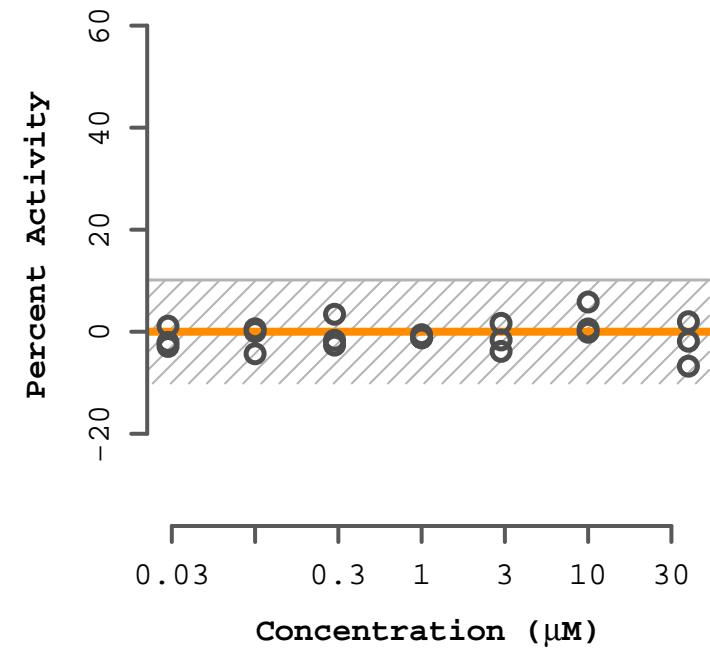
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 104.95	NA	NA
PROB: 1	NA	NA
RMSE: 3.97	NA	NA

MAX_MEAN: 7.95 MAX_MED: 7.95 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 3'-Azido-3'-deoxythymidine

CHID: 20127 CASRN: 30516-87-1

SPID(S): TP0001413H08

M4ID: 30204595

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 105.26

PROB: 1

RMSE: 2.8

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 2.1

MAX_MED: 0.554

BMAD: 3.38

COFF: 10.1

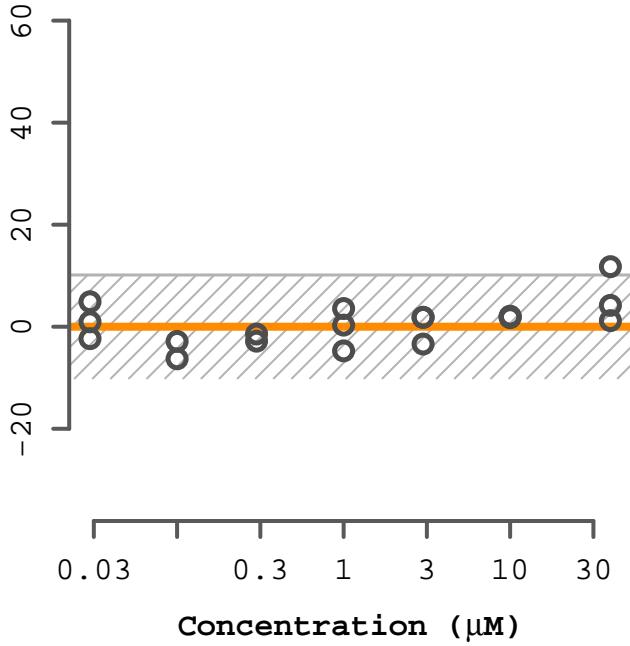
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Benzidine

CHID: 20137 CASRN: 92-87-5

SPID(S): TP0001414E04

M4ID: 30204473

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 108.19

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.04

NA

NA

MAX_MEAN: 5.72

MAX_MED: 4.19

BMAD: 3.38

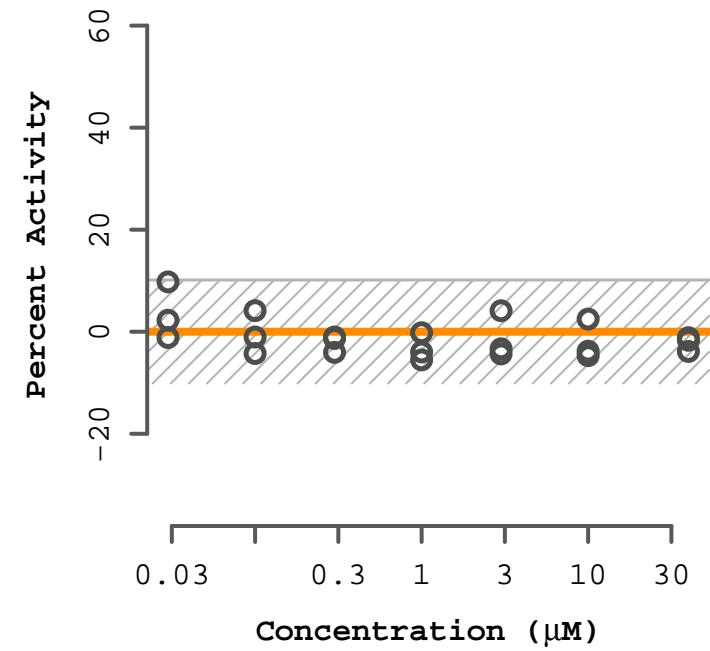
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

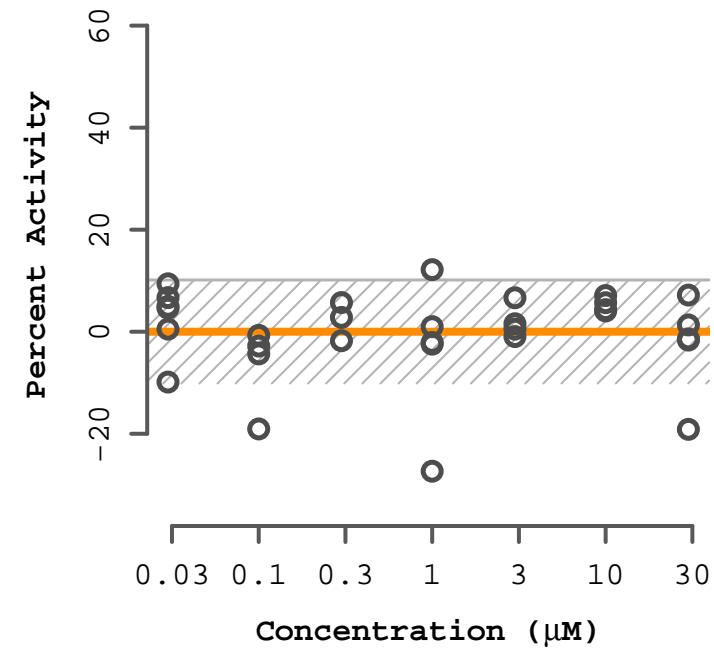
NAME: 2,3-Benzofuran
CHID: 20141 CASRN: 271-89-6
SPID(S): TP0001412B12
M4ID: 30204584

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	120.11	NA	NA
PROB:	1	NA	NA
RMSE:	3.86	NA	NA
MAX_MEAN:	3.64	MAX_MED: 2.26	BMAD: 3.38
COFF:	10.1	HIT-CALL: 0	FITC: 4
ACTP:	0		

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 6-Phenyl-1,3,5-triazine-2,4-diamine

CHID: 20142 CASRN: 91-76-9

SPID(S): EPAPLT0154A11

M4ID: 30204162

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 216.45

PROB: 1

RMSE: 8.44

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 5.3

MAX_MED: 5.02

BMAD: 3.38

COFF: 10.1

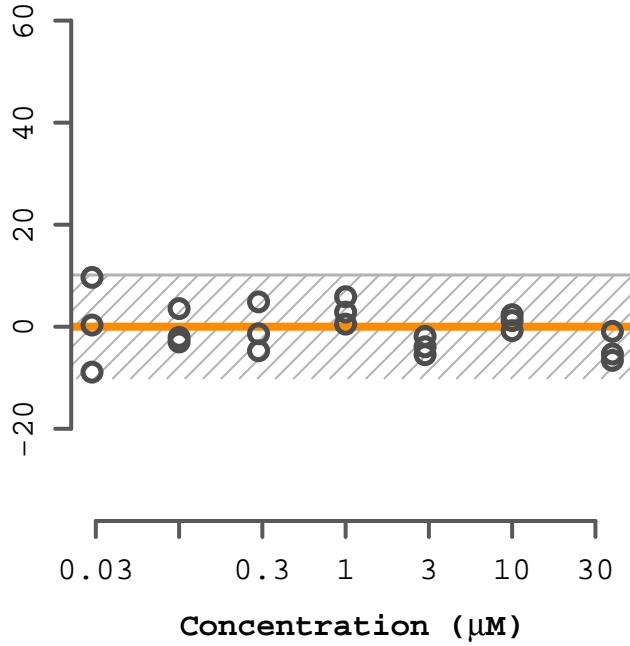
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Benzoic acid

CHID: 20143 CASRN: 65-85-0

SPID(S): TP0001411D07

M4ID: 30204469

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 126.1

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.47

NA

NA

MAX_MEAN: 3.09

MAX_MED: 2.87

BMAD: 3.38

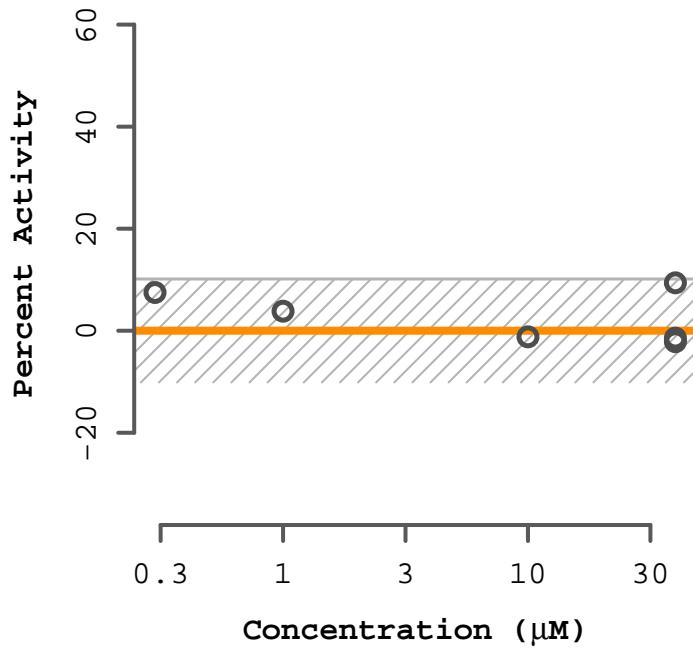
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,2,3-Benzotriazole
CHID: 20147 CASRN: 95-14-7
SPID(S): TP0001411A11
M4ID: 30204395

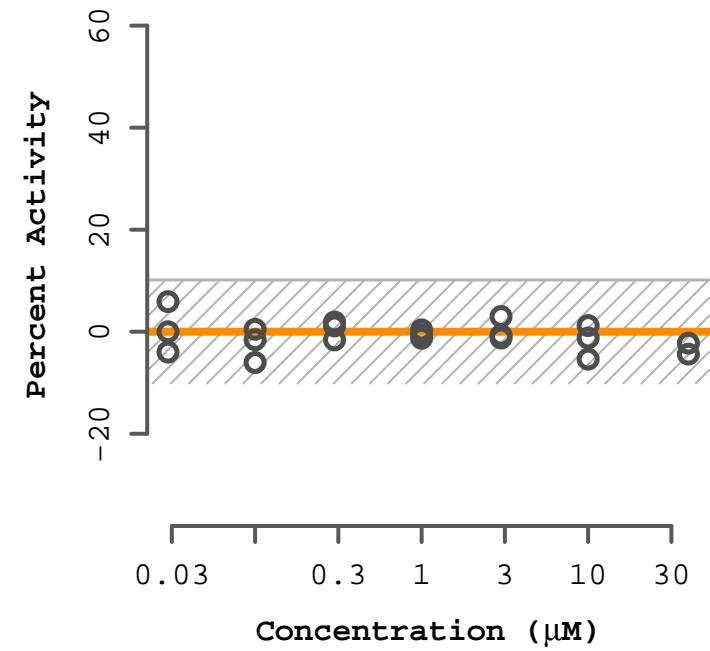
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 39.46	NA	NA
PROB: 1	NA	NA
RMSE: 5.28	NA	NA

MAX_MEAN: 7.5 MAX_MED: 7.5 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Benzotrichloride

CHID: 20148 CASRN: 98-07-7

SPID(S): TP0001413B12

M4ID: 30204470

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 101.42

PROB: 1

RMSE: 2.88

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 0.639

MAX_MED: 1.14

BMAD: 3.38

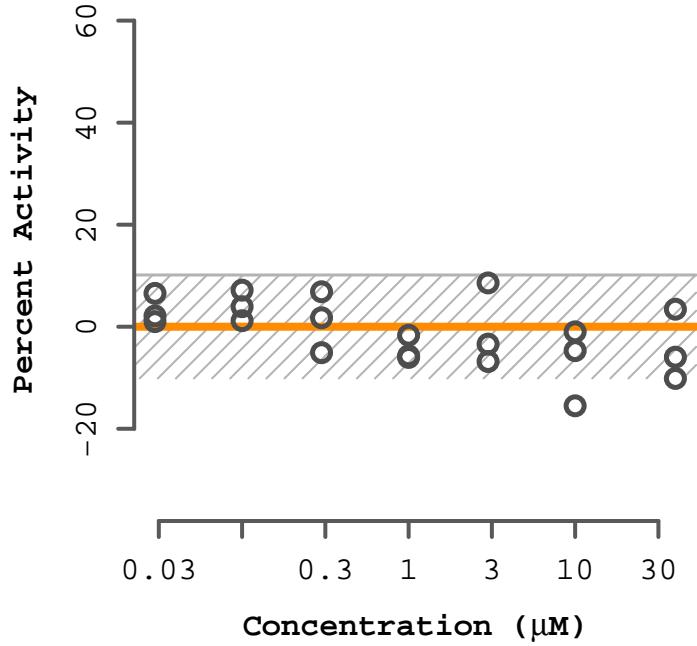
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pentaerythritol dibromide
CHID: 20164 CASRN: 3296-90-0
SPID(S): TP0001414B04
M4ID: 30204430

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

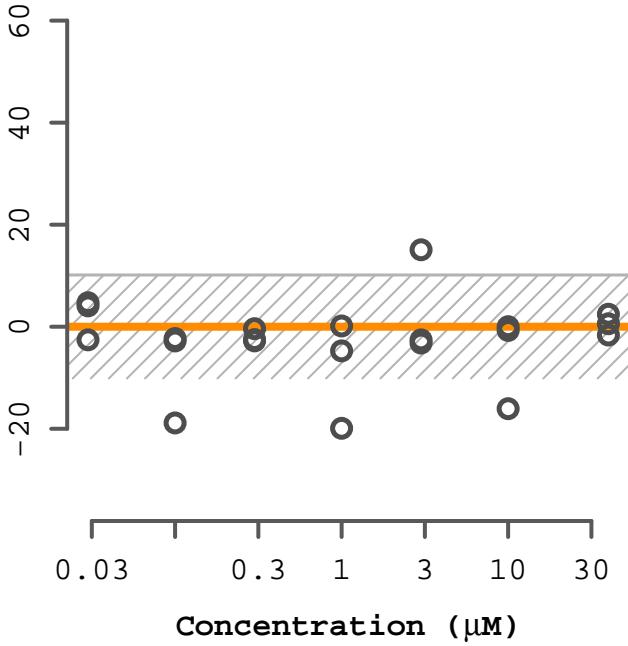
	CNST	HILL	GNLS
AIC:	139.9	NA	NA
PROB:	1	NA	NA
RMSE:	6.21	NA	NA

MAX_MEAN: 4.1 **MAX_MED:** 3.93 **BMAD:** 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS :

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bis(2-chloroethyl) ether
CHID: 20168 CASRN: 111-44-4
SPID(S): TP0001412E02
M4ID: 30204650

HILL MODEL: Not applicable.

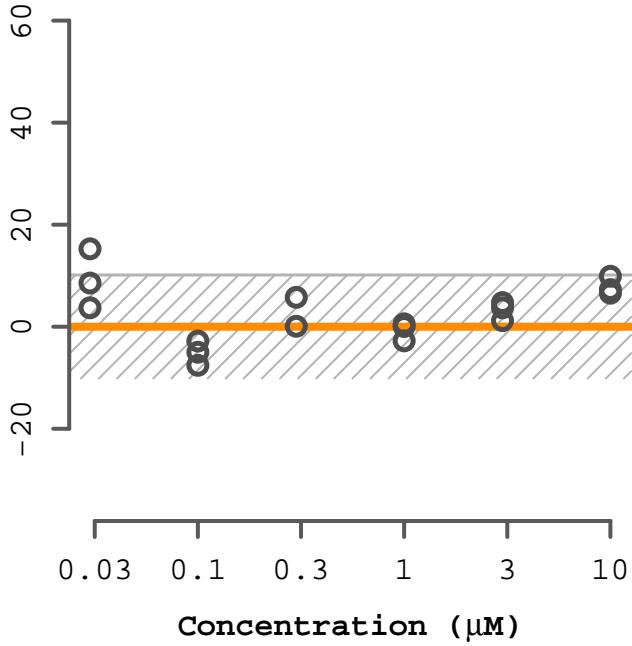
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 144.35	NA	NA
PROB: 1	NA	NA
RMSE: 8.04	NA	NA

MAX_MEAN: 3.11 MAX_MED: 4.09 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bisphenol A

CHID: 20182 CASRN: 80-05-7

SPID(S): TP0001411D06

M4ID: 30204626

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 114.06

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 6.34

NA

NA

MAX_MEAN: 9.19

MAX_MED: 8.57

BMAD: 3.38

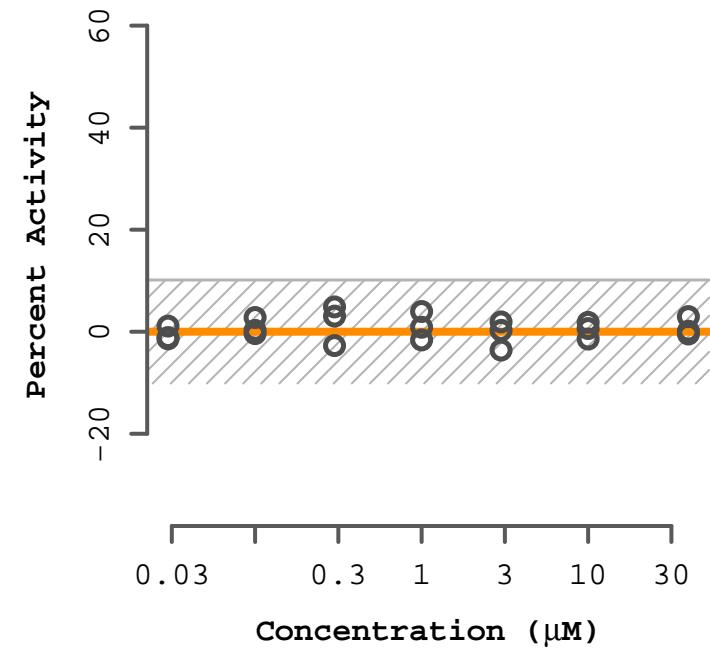
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Boric acid (H₃BO₃)
CHID: 20194 CASRN: 10043-35-3
SPID(S): TP0001413G02
M4ID: 30204446

HILL MODEL: Not applicable.

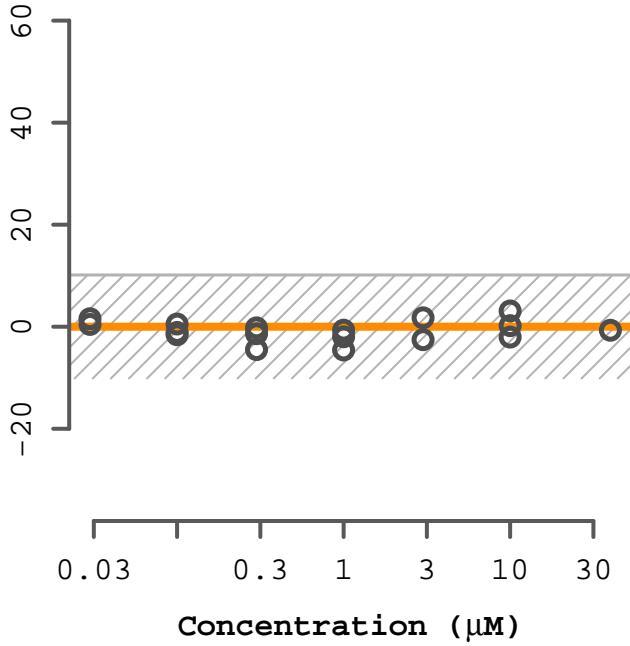
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 96.73	NA	NA
PROB: 1	NA	NA
RMSE: 2.22	NA	NA

MAX_MEAN: 1.74 MAX_MED: 3.07 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Caffeine

CHID: 20232 CASRN: 58-08-2

SPID(S): TP0001411F04

M4ID: 30204356

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 76.73

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 2.13

NA

NA

MAX_MEAN: 1.02

MAX_MED: 1.02

BMAD: 3.38

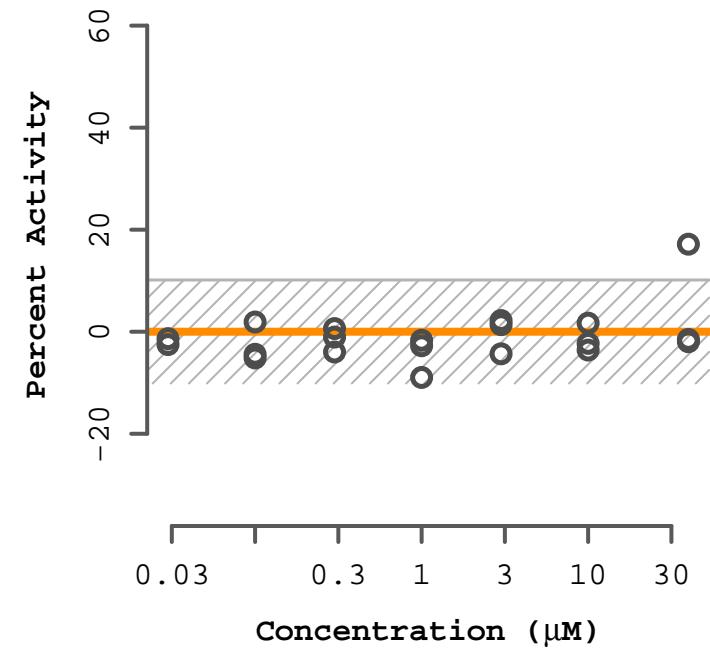
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Captafol

CHID: 20242 CASRN: 2425-06-1

SPID(S): TP0001414D03

M4ID: 30204377

HILL MODEL: Not applicable.

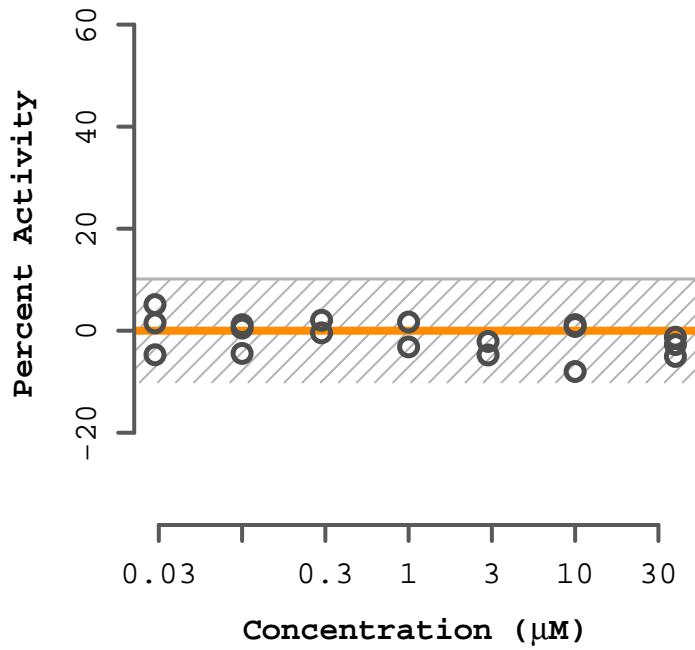
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	118.14	NA	NA
PROB:	1	NA	NA
RMSE:	5.06	NA	NA

MAX_MEAN: 4.57 MAX_MED: 1.39 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Chlorendic acid
CHID: 20268 CASRN: 115-28-6
SPID(S): TP0001414E05
M4ID: 30204479

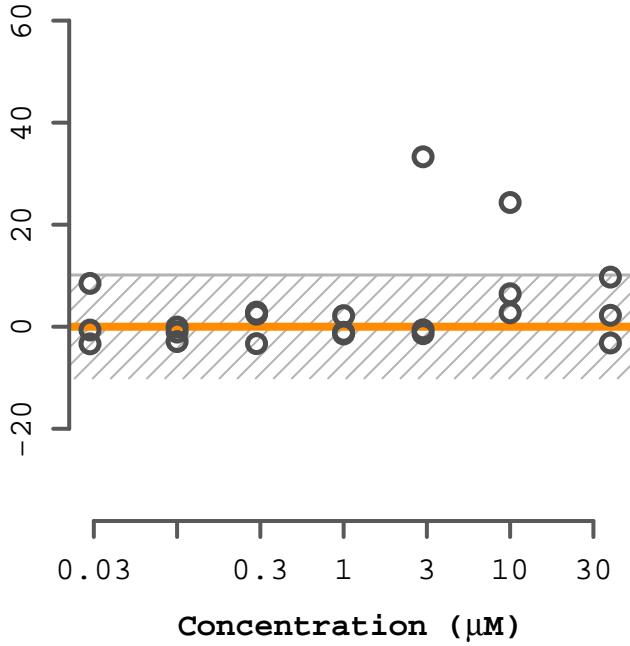
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	99.12	NA	NA
PROB:	1	NA	NA
RMSE:	3.47	NA	NA
MAX_MEAN:	0.827	MAX_MED: 1.54	BMAD: 3.38
COFF:	10.1	HIT-CALL: 0	FITC: 4
ACTP:	0		

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Chlorothalonil

CHID: 20319 CASRN: 1897-45-6

SPID(S): TP0001414A10

M4ID: 30204447

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 144.17

PROB: 1

RMSE: 9.74

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 11.2

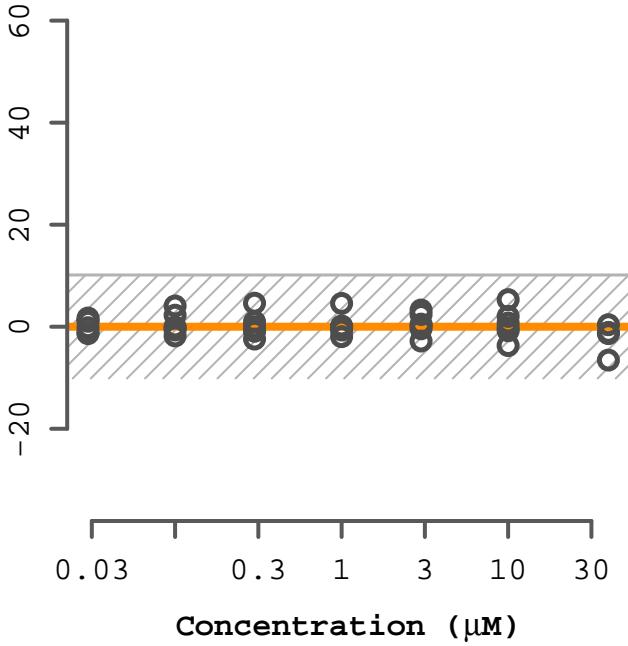
MAX_MED: 6.48

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Daminozide

CHID: 20370 CASRN: 1596-84-5

SPID(S): TP0001413D06

M4ID: 30204323

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 178.4

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 2.27

NA

NA

MAX_MEAN: 0.621

MAX_MED: 0.449

BMAD: 3.38

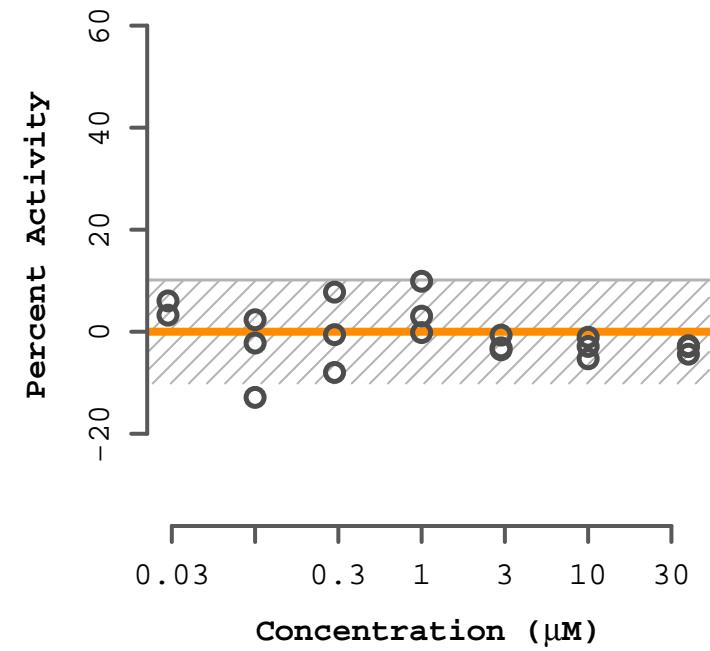
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,4-Diaminotoluene

CHID: 20402 CASRN: 95-80-7

SPID(S): TP0001413D09

M4ID: 30204444

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 125.56

PROB: 1

RMSE: 5.24

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 4.65

MAX_MED: 4.65

BMAD: 3.38

COFF: 10.1

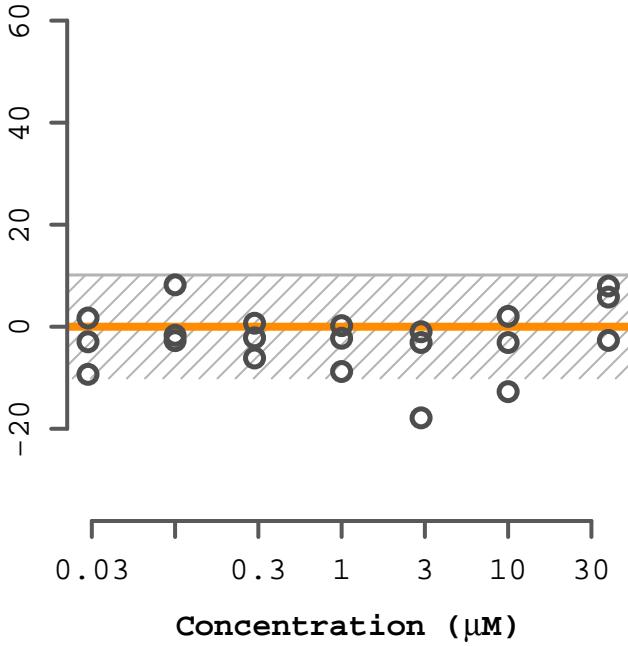
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dimethoate

CHID: 20479 CASRN: 60-51-5

SPID(S): TP0001412D05

M4ID: 30204365

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 139.91

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 6.58

NA

NA

MAX_MEAN: 3.71

MAX_MED: 5.82

BMAD: 3.38

COFF: 10.1

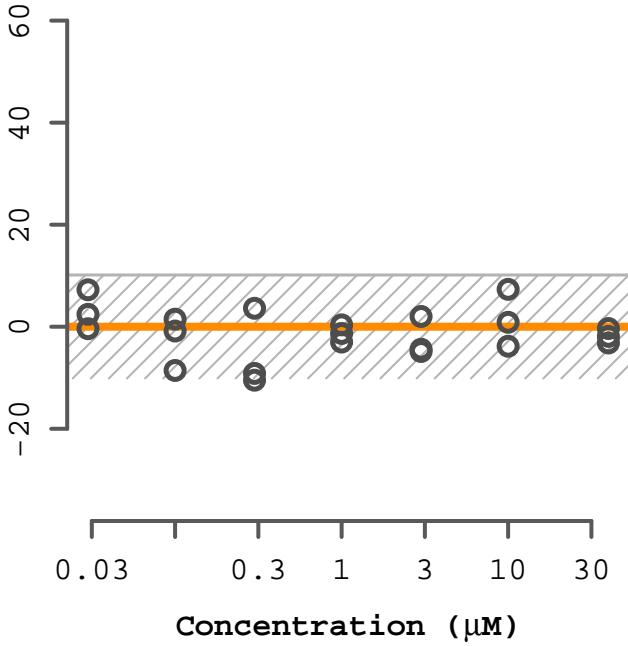
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dimethylaminoethanol
CHID: 20505 CASRN: 108-01-0
SPID(S): TP0001412E01
M4ID: 30204655

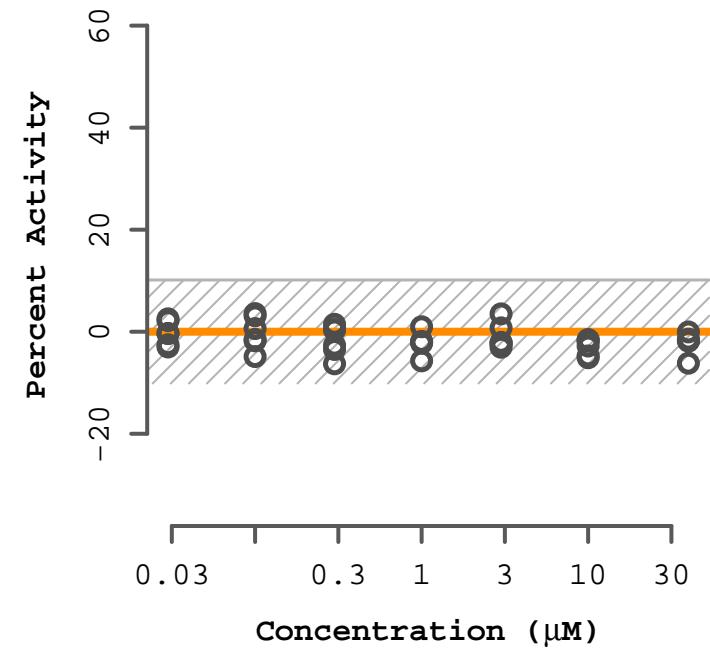
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 128.13	NA	NA
PROB: 1	NA	NA
RMSE: 4.79	NA	NA

MAX_MEAN: 3.1 MAX_MED: 2.43 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: N,N-Dimethylformamide
CHID: 20515 CASRN: 68-12-2
SPID(S): TP0001413D02
M4ID: 30204647

HILL MODEL: Not applicable.

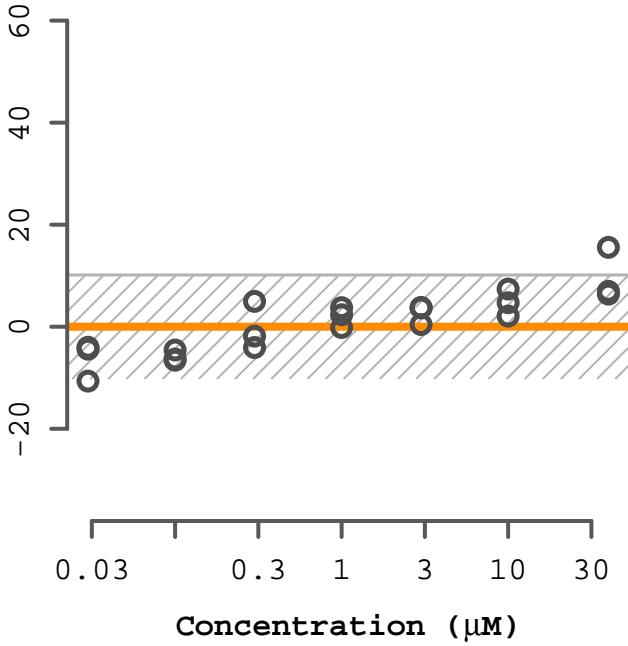
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	204.04	NA	NA
PROB:	1	NA	NA
RMSE:	2.89	NA	NA

MAX_MEAN: -0.137 MAX_MED: -0.346 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Endosulfan

CHID: 20560 CASRN: 115-29-7

SPID(S): TP0001413H09

M4ID: 30204348

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 138.14

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 6.01

NA

NA

MAX_MEAN: 9.64

MAX_MED: 6.94

BMAD: 3.38

COFF: 10.1

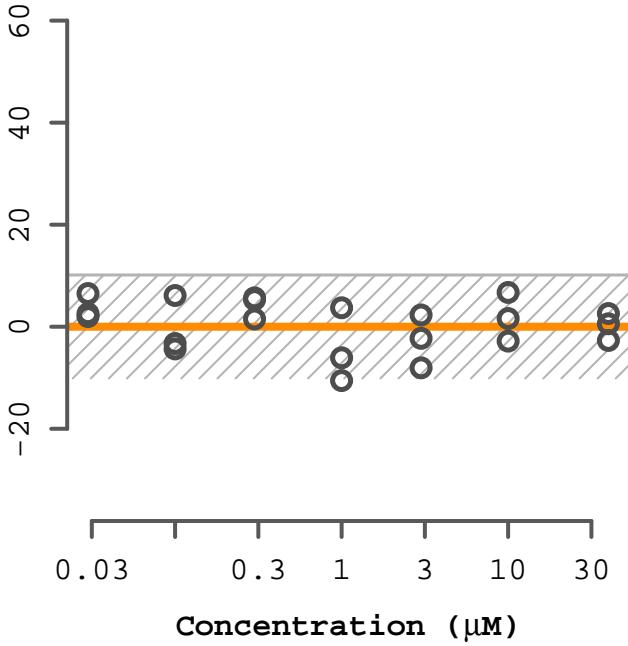
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Furfural

CHID: 20647 CASRN: 98-01-1

SPID(S): TP0001412A07

M4ID: 30204460

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 130.16

NA

NA

PROB:

1

NA

NA

RMSE: 4.83

NA

NA

MAX_MEAN: 4.17

MAX_MED: 5.32

BMAD: 3.38

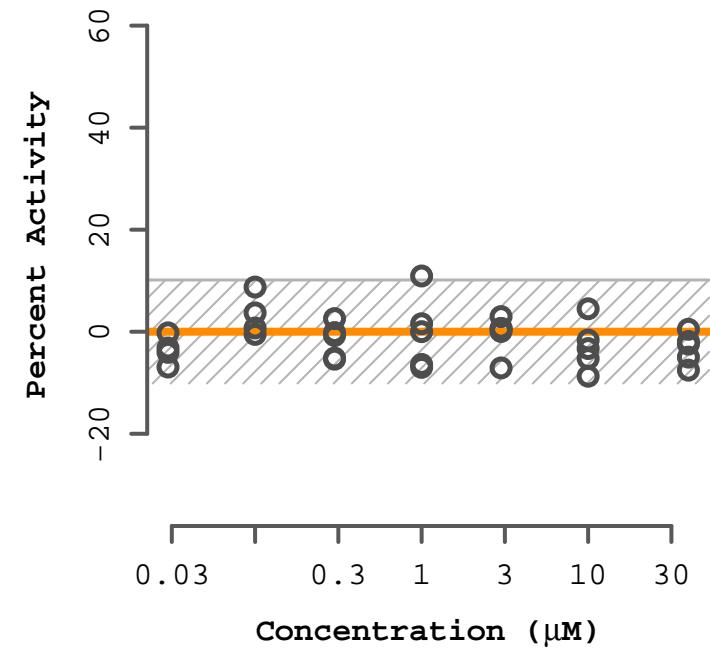
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Hydrochlorothiazide
CHID: 20713 CASRN: 58-93-5
SPID(S): TP0001413B09
M4ID: 30204566

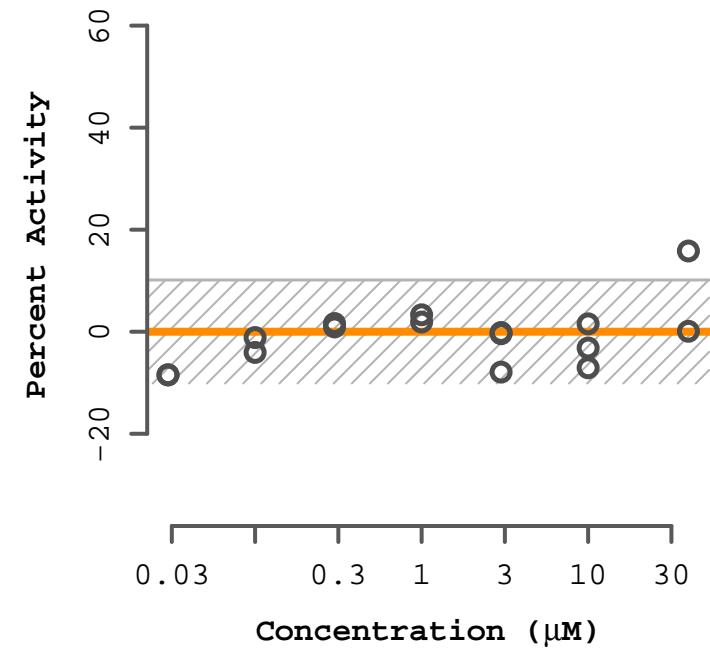
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 199.96	NA	NA
PROB: 1	NA	NA
RMSE: 4.7	NA	NA

MAX_MEAN: 3.17 MAX_MED: 2.22 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Indomethacin

CHID: 20740 CASRN: 53-86-1

SPID(S): TP0001412H12

M4ID: 30204528

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 102.93

PROB: 1

RMSE: 5.89

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 7.96

MAX_MED: 7.96

BMAD: 3.38

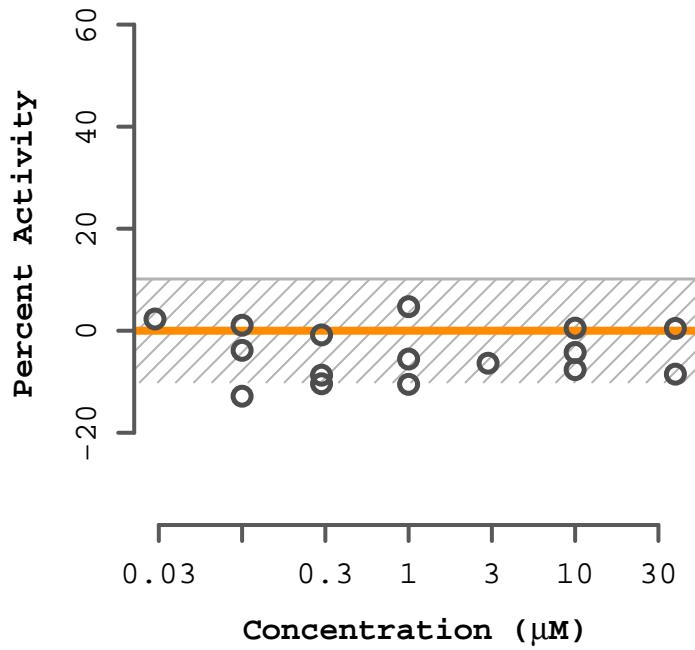
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Malaoxon

CHID: 20790 CASRN: 1634-78-2

SPID(S): TP0001414C10

M4ID: 30204318

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 110.28

PROB: 1

RMSE: 6.72

HILL

NA

NA

NA

GNLS

NA

NA

NA

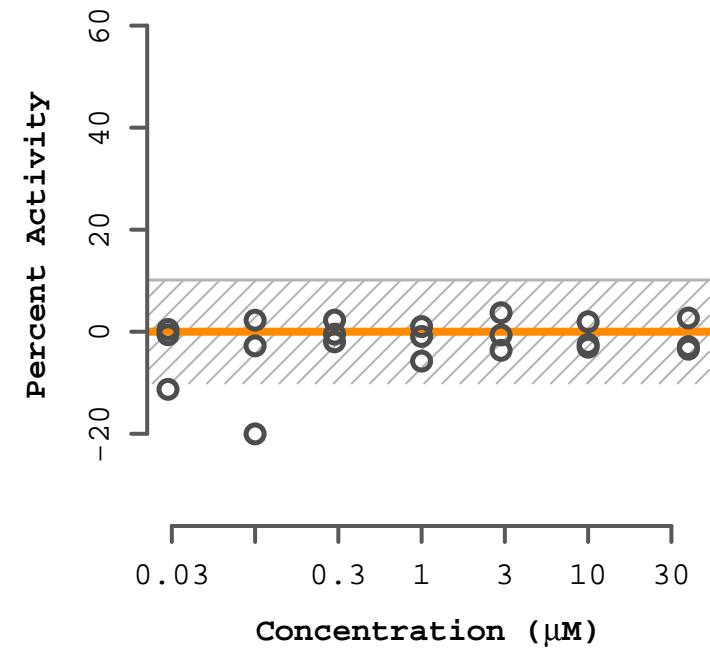
MAX_MEAN: 2.3

MAX_MED: 2.3

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: N-Methyl-2-pyrrolidone
CHID: 20856 CASRN: 872-50-4
SPID(S): TP0001411E04
M4ID: 30204415

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

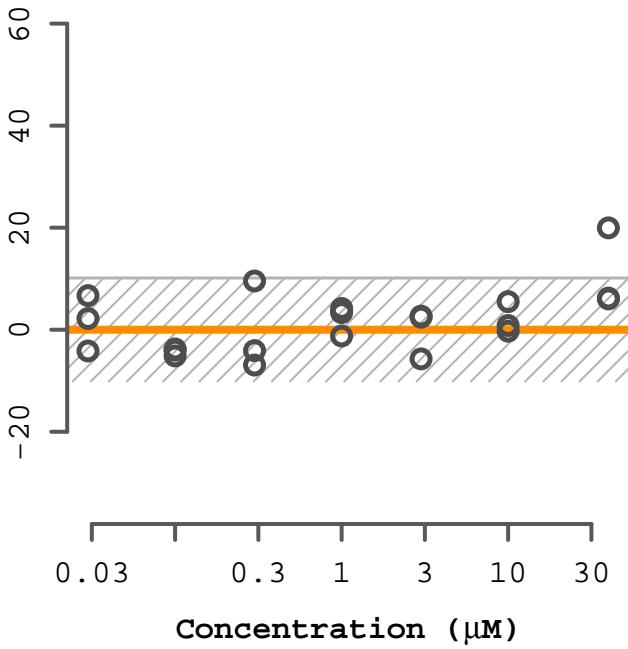
CNST	HILL	GNLS
AIC: 124.86	NA	NA
PROB: 1	NA	NA
RMSE: 5.61	NA	NA

MAX_MEAN: -0.0525 MAX_MED: -0.466 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 6-Methylquinoline
CHID: 20887 CASRN: 91-62-3
SPID(S): TP0001413D12
M4ID: 30204583

HILL MODEL: Not applicable.

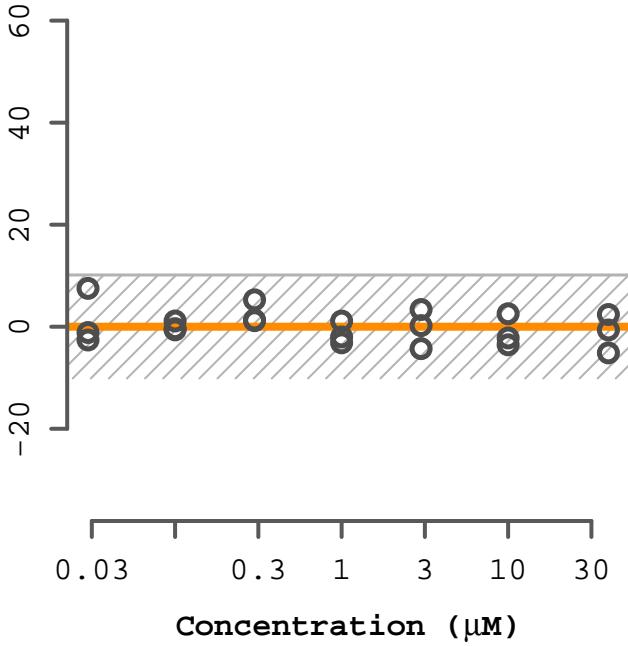
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 138.28	NA	NA
PROB: 1	NA	NA
RMSE: 6.4	NA	NA

MAX_MEAN: 10.7 MAX_MED: 6.21 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Mirex

CHID: 20895 CASRN: 2385-85-5

SPID(S): TP0001413B03

M4ID: 30204600

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 109.81

HILL

PROB: 1

NA

RMSE: 3.08

NA

GNLS

MAX_MEAN: 2.6

MAX_MED: 1.33

BMAD: 3.38

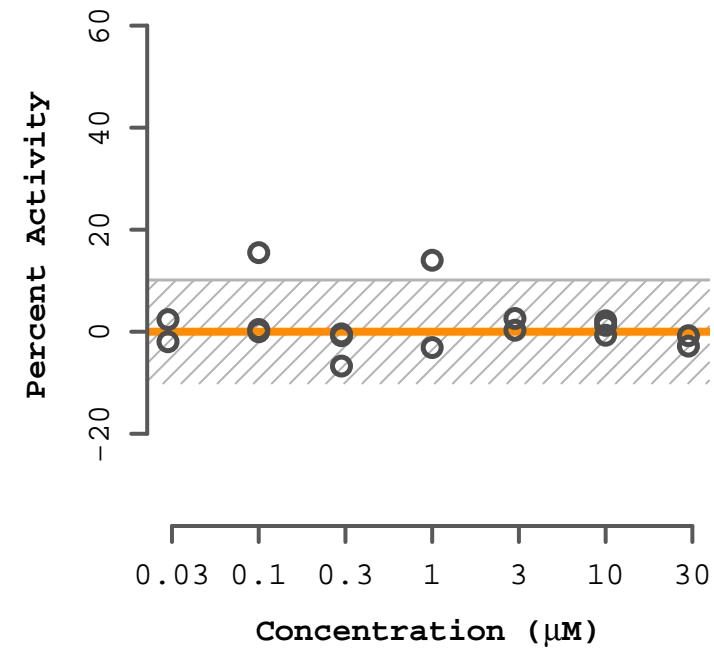
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

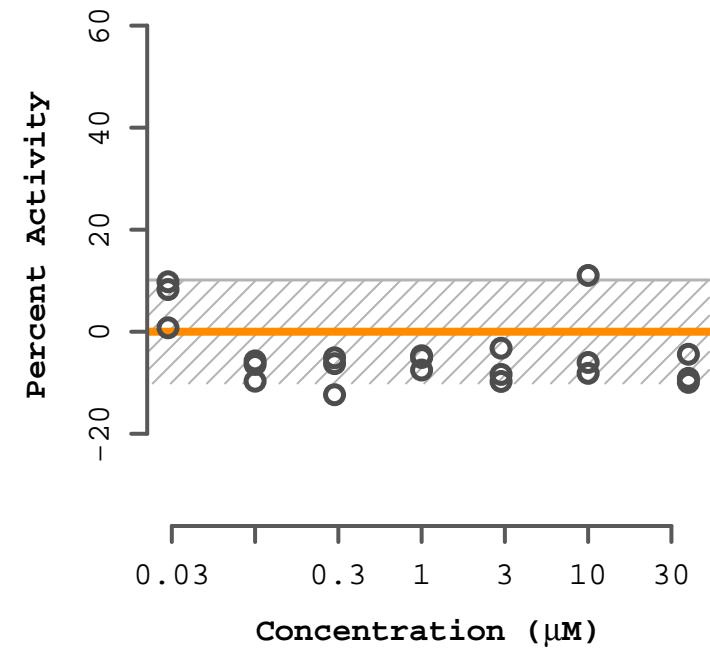
NAME: Nalidixic acid
CHID: 20912 CASRN: 389-08-2
SPID(S): EPAPLT0154E05
M4ID: 30204199

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	101.64	NA	NA
PROB:	1	NA	NA
RMSE:	5.55	NA	NA
MAX_MEAN:	5.46	MAX_MED: 5.46	BMAD: 3.38
COFF:	10.1	HIT-CALL: 0	FITC: 4
ACTP:	0		

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Naphthylamine

CHID: 20921 CASRN: 91-59-8

SPID(S): TP0001412E11

M4ID: 30204392

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 151.59

PROB: 1

RMSE: 7.73

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 6.28

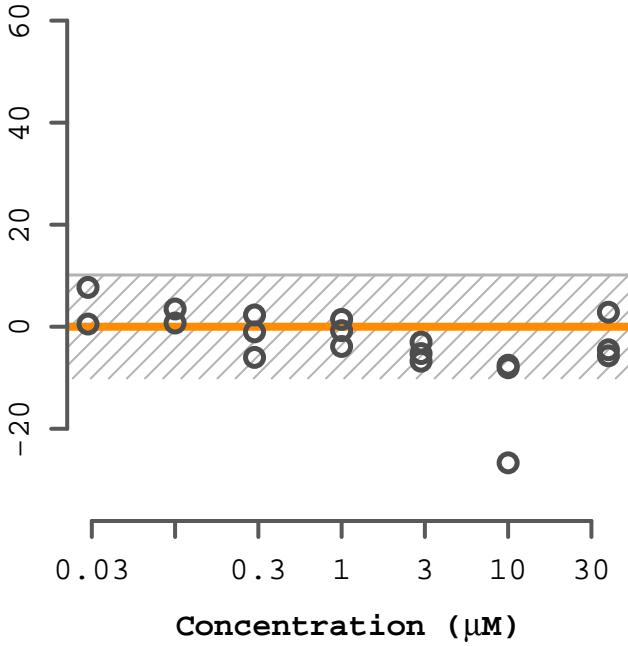
MAX_MED: 8.28

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Nicotine

CHID: 20930 CASRN: 54-11-5

SPID(S): TP0001411C10

M4ID: 30204463

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 127.48

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 7.63

NA

NA

MAX_MEAN: 4.12

MAX_MED: 4.12

BMAD: 3.38

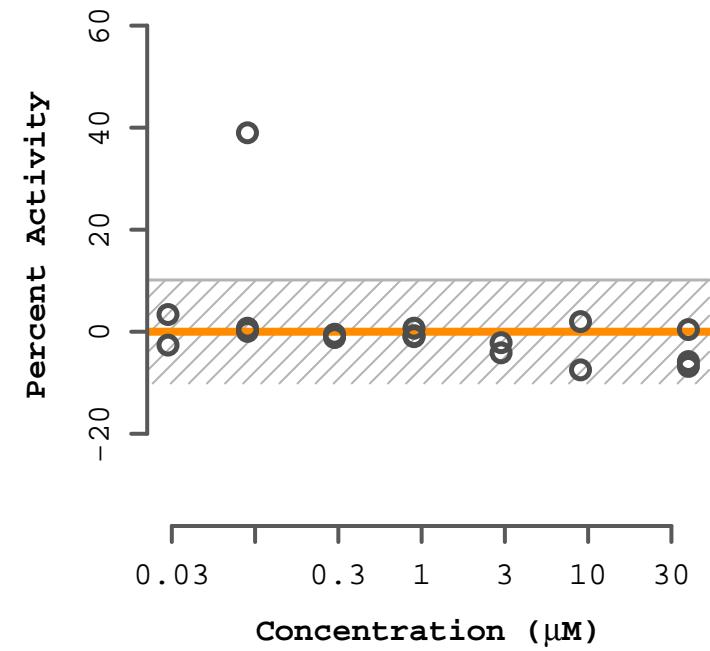
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Nitrilotriacetic acid
CHID: 20939 CASRN: 139-13-9
SPID(S): TP0001414G05
M4ID: 30204452

HILL MODEL: Not applicable.

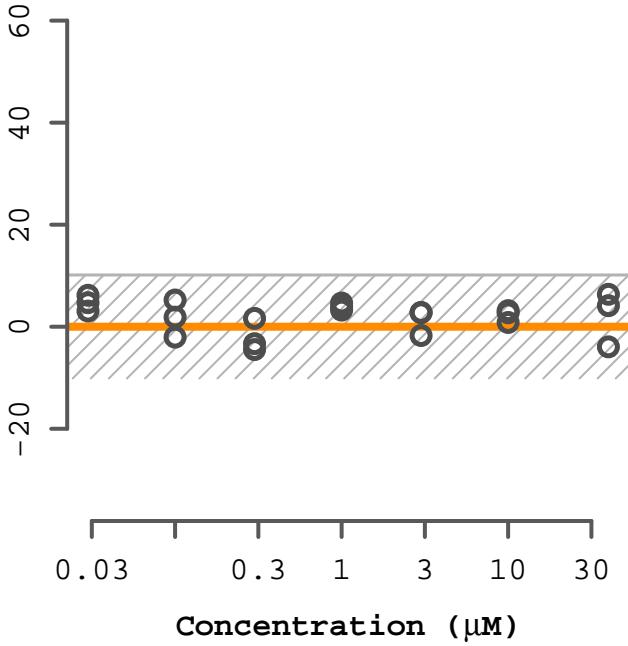
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 109.43	NA	NA
PROB: 1	NA	NA
RMSE: 10.02	NA	NA

MAX_MEAN: 13.3 MAX_MED: 0.701 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: N-Nitrosodiethylamine
CHID: 21028 CASRN: 55-18-5
SPID(S): TP0001412F10
M4ID: 30204646

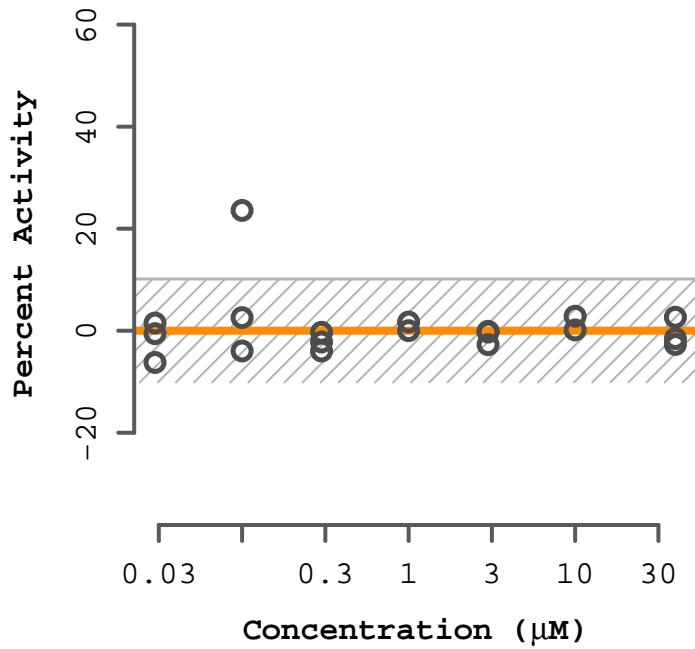
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 120.9	NA	NA
PROB: 1	NA	NA
RMSE: 3.75	NA	NA

MAX_MEAN: 4.66 MAX_MED: 4.72 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: N-Nitrosodipropylamine

CHID: 21032 CASRN: 621-64-7

SPID(S): TP0001411G03

M4ID: 30204454

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 108.17

PROB: 1

RMSE: 5.96

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 7.4

MAX_MED: 2.57

BMAD: 3.38

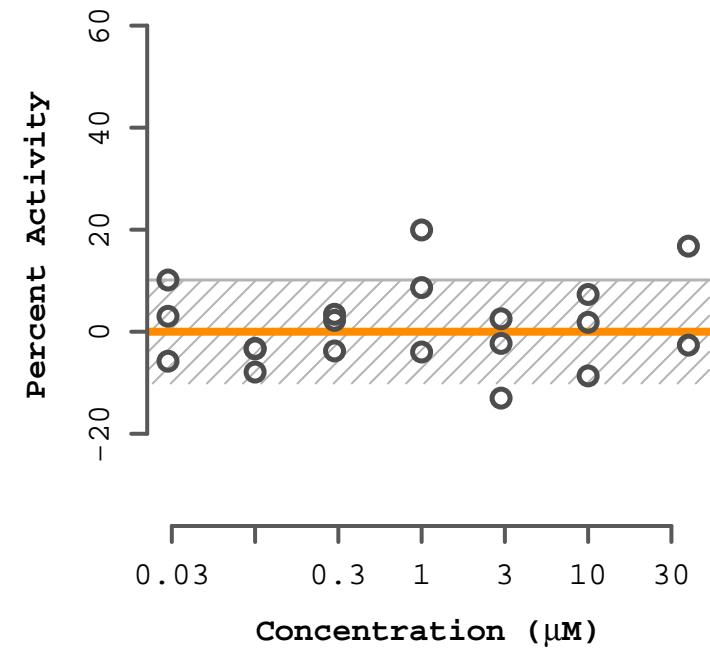
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Oxamyl

CHID: 21086 CASRN: 23135-22-0

SPID(S): TP0001412H07

M4ID: 30204309

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 143.28

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 8.22

NA

NA

MAX_MEAN: 8.21

MAX_MED: 8.66

BMAD: 3.38

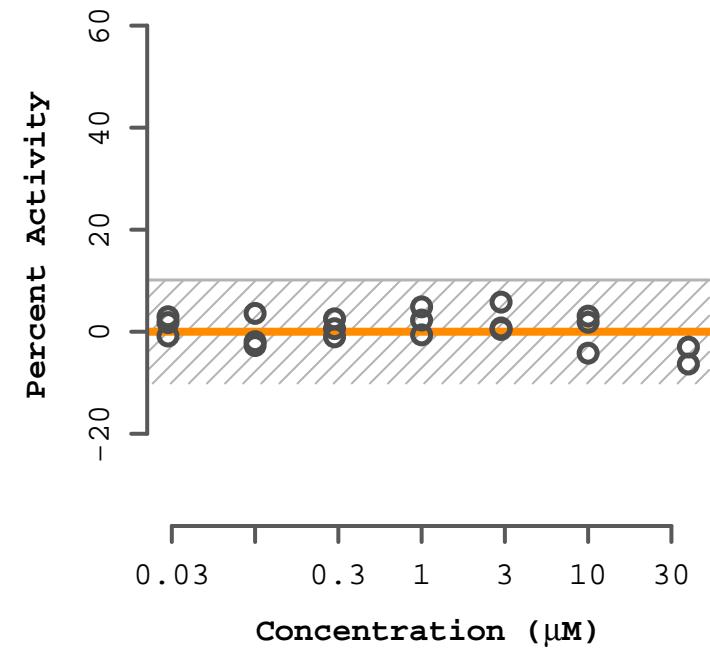
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Oxytetracycline hydrochloride
CHID: 21097 CASRN: 2058-46-0
SPID(S): TP0001414D05
M4ID: 30204604

HILL MODEL: Not applicable.

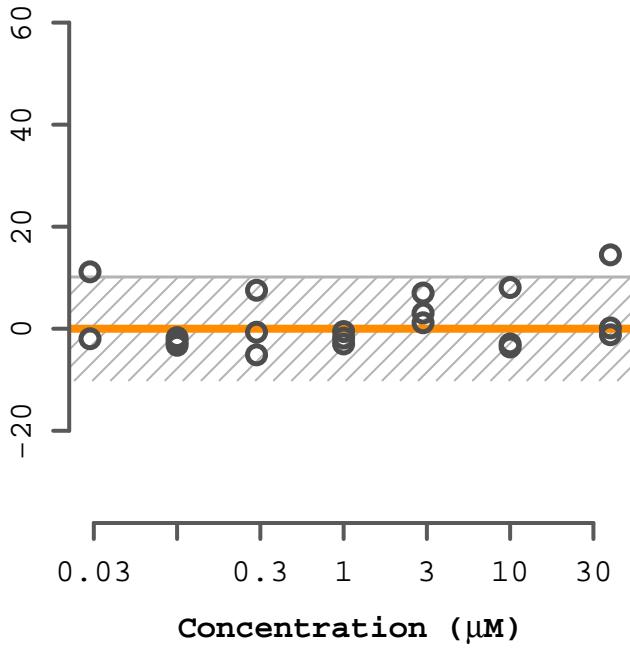
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 105.53	NA	NA
PROB: 1	NA	NA
RMSE: 3.06	NA	NA

MAX_MEAN: 2.33 MAX_MED: 2.3 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Phenobarbital sodium
CHID: 21123 CASRN: 57-30-7
SPID(S): TP0001414H03
M4ID: 30204531

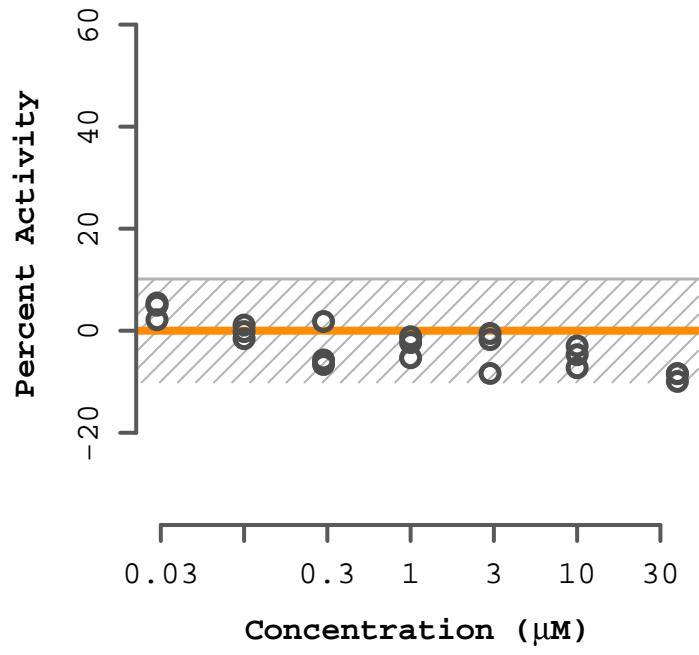
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 125.76	NA	NA
PROB: 1	NA	NA
RMSE: 5.49	NA	NA

MAX_MEAN: 4.6 MAX_MED: 4.6 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Piperazine

CHID: 21164 CASRN: 110-85-0

SPID(S): TP0001412A08

M4ID: 30204652

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 133.5

PROB: 1

RMSE: 5.23

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 4.2

MAX_MED: 5

BMAD: 3.38

COFF: 10.1

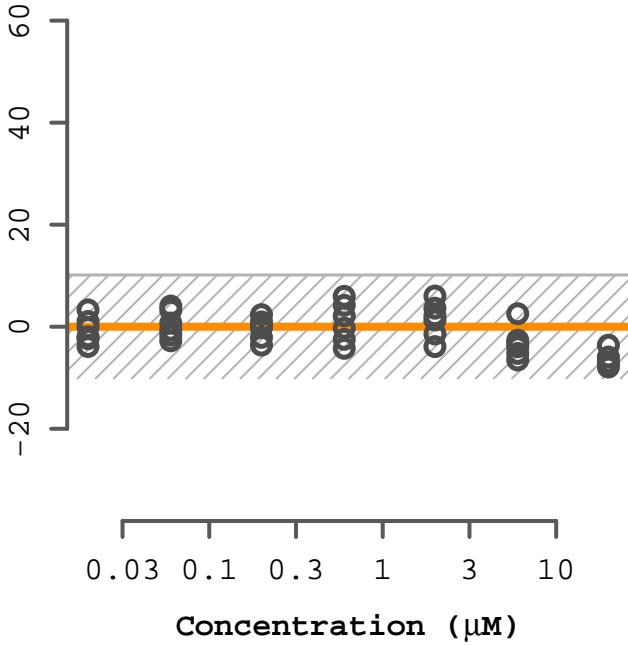
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Rifampicin

CHID: 21244 CASRN: 13292-46-1

SPID(S): TP0001413D05

M4ID: 30204577

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 231.03

HILL

PROB: 1

NA

RMSE: 3.74

NA

GNLS

MAX_MEAN: 1.31

MAX_MED: 1.73

BMAD: 3.38

COFF: 10.1

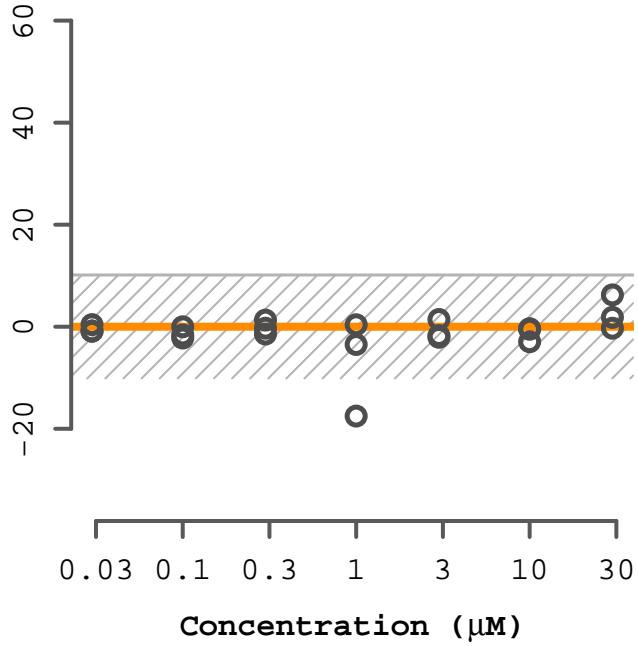
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Saccharin

CHID: 21251 CASRN: 81-07-2

SPID(S): EPAPLT0154G09

M4ID: 30204222

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 105.79

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.32

NA

NA

MAX_MEAN: 2.6

MAX_MED: 1.85

BMAD: 3.38

COFF: 10.1

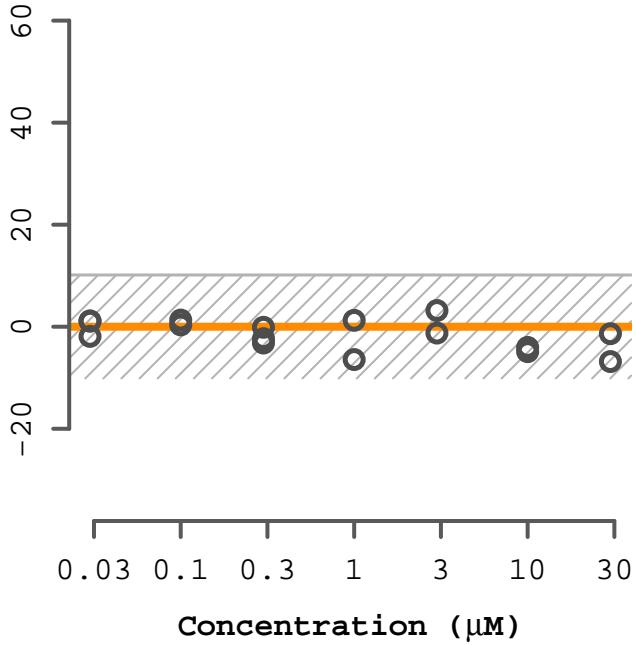
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sulfasalazine

CHID: 21256 CASRN: 599-79-1

SPID(S): EPAPLT0154D03

M4ID: 30204186

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 85.53

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 3.23

NA

NA

MAX_MEAN: 0.975

MAX_MED: 1.12

BMAD: 3.38

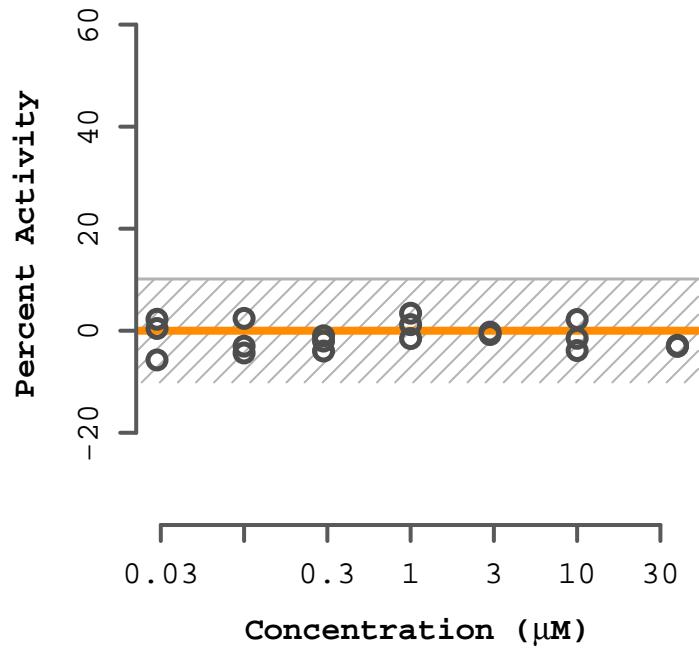
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Theophylline

CHID: 21336 CASRN: 58-55-9

SPID(S): TP0001414H06

M4ID: 30204388

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 102.82

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 2.81

NA

NA

MAX_MEAN: 1.04

MAX_MED: 1.2

BMAD: 3.38

COFF: 10.1

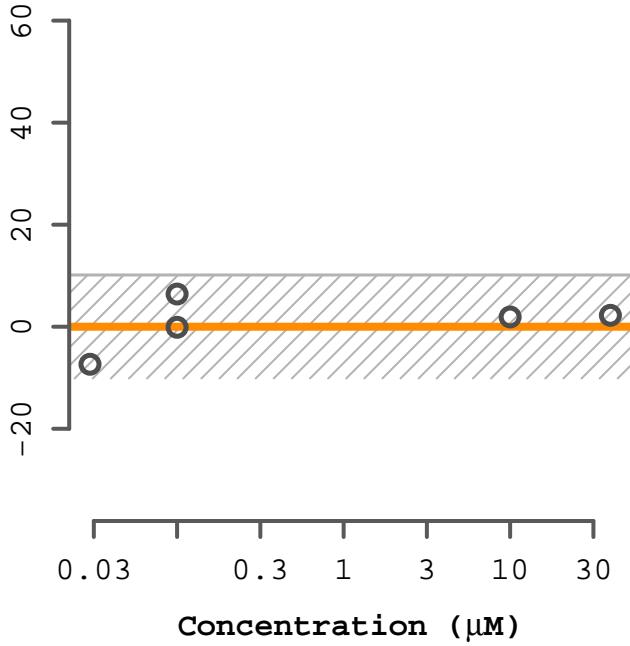
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Trichloroacetic acid
CHID: 21378 CASRN: 76-03-9
SPID(S): TP0001411G10
M4ID: 30204472

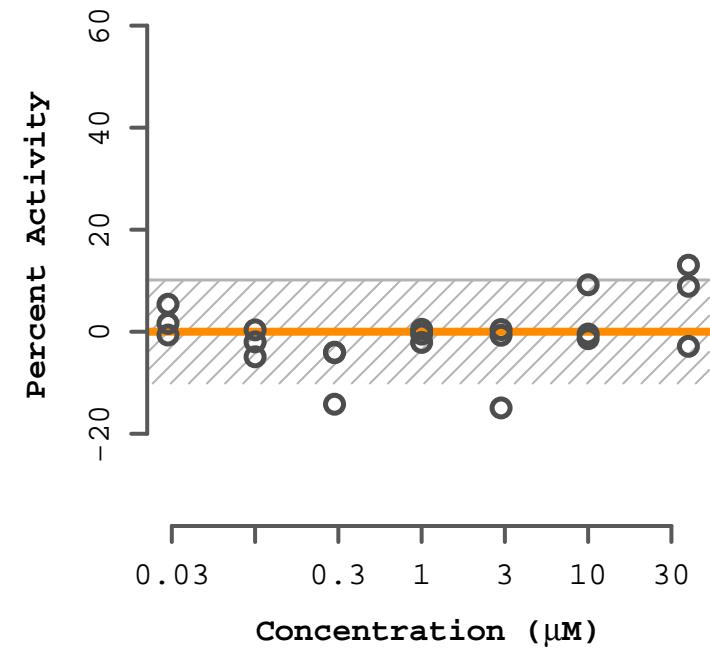
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 31.87	NA	NA
PROB: 1	NA	NA
RMSE: 4.56	NA	NA

MAX_MEAN: 3.15 MAX_MED: 3.15 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Folpet

CHID: 21385 CASRN: 133-07-3

SPID(S): TP0001412E03

M4ID: 30204327

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 137.94

HILL

PROB: 1

NA

RMSE: 6.43

NA

GNLS

NA

NA

MAX_MEAN: 6.37

MAX_MED: 8.91

BMAD: 3.38

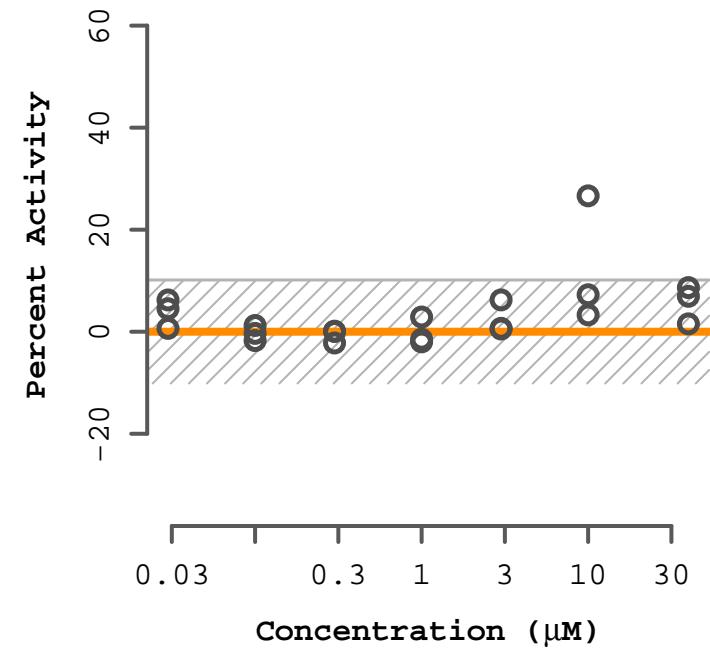
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Trifluralin

CHID: 21395 CASRN: 1582-09-8

SPID(S): TP0001413C10

M4ID: 30204339

HILL MODEL: Not applicable.

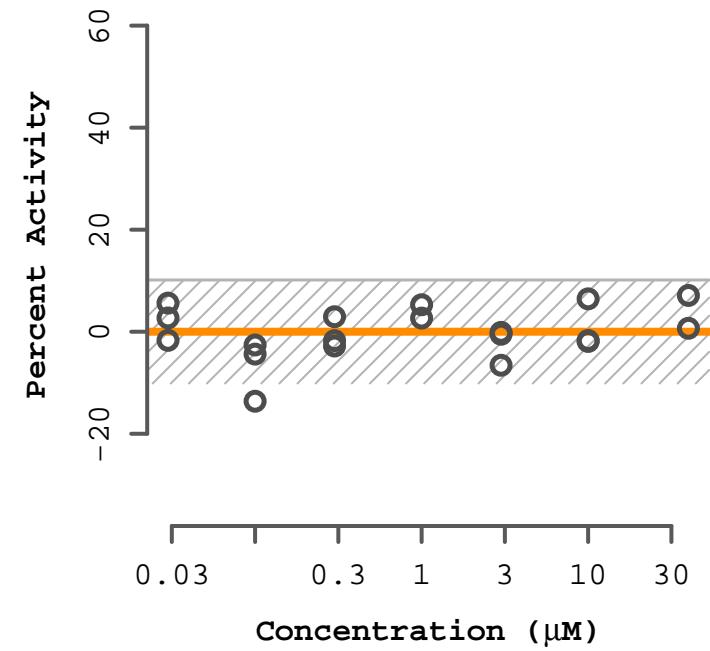
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 132.44	NA	NA
PROB: 1	NA	NA
RMSE: 6.98	NA	NA

MAX_MEAN: 12.4 MAX_MED: 7.26 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tris(2-chloroethyl) phosphate

CHID: 21411 CASRN: 115-96-8

SPID(S): TP0001413A06

M4ID: 30204631

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 115.61

PROB: 1

RMSE: 4.87

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 3.99

MAX_MED: 3.99

BMAD: 3.38

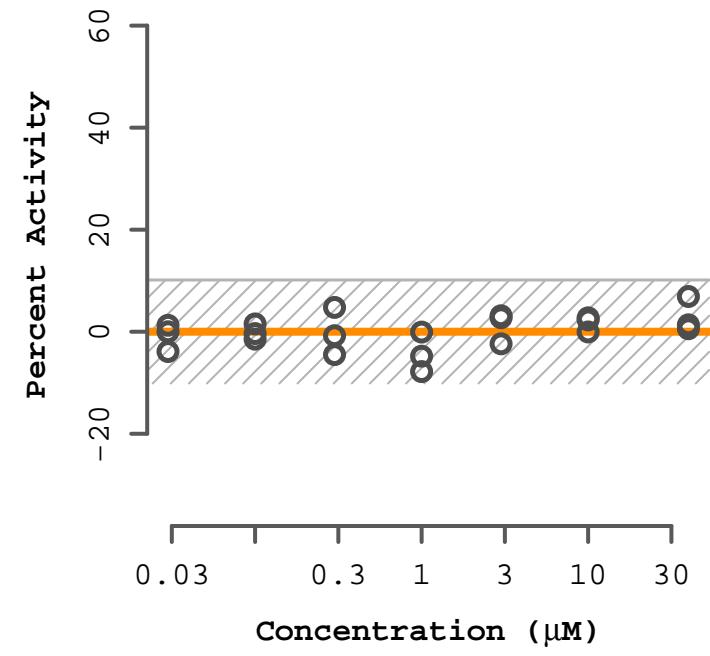
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tris(2-ethylhexyl) phosphate
CHID: 21414 CASRN: 78-42-2
SPID(S): TP0001412D07
M4ID: 30204633

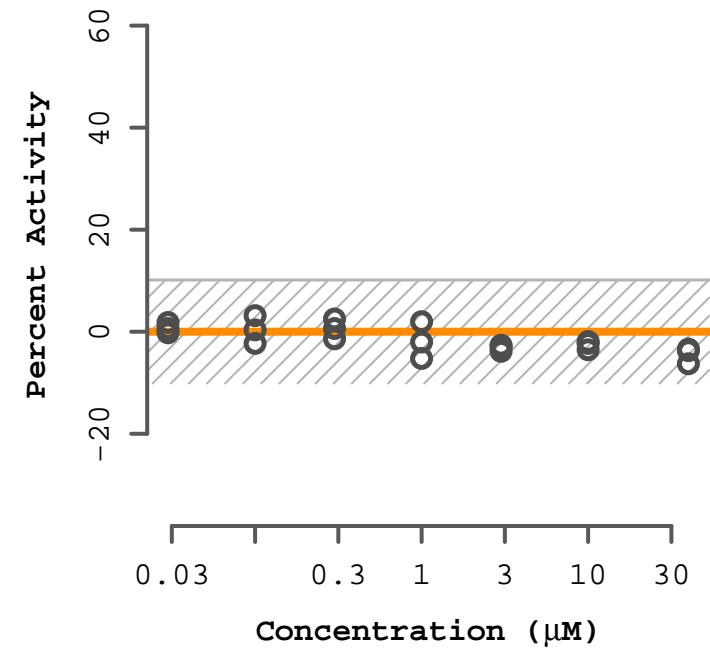
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 112.85	NA	NA
PROB: 1	NA	NA
RMSE: 3.34	NA	NA

MAX_MEAN: 2.96 MAX_MED: 2.75 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: FD&C Yellow 6

CHID: 21456 CASRN: 2783-94-0

SPID(S): TP0001414G03

M4ID: 30204474

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 109.24

PROB: 1

RMSE: 2.92

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 0.749

MAX_MED: 0.641

BMAD: 3.38

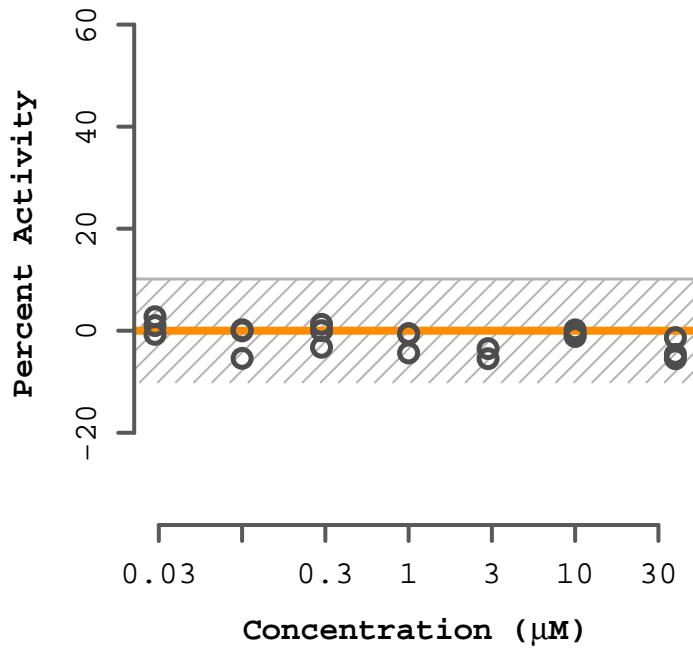
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Butanoic acid

CHID: 21515 CASRN: 107-92-6

SPID(S): TP0001414D07

M4ID: 30204458

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 101.93

HILL

PROB: 1

GNLS

RMSE: 2.89

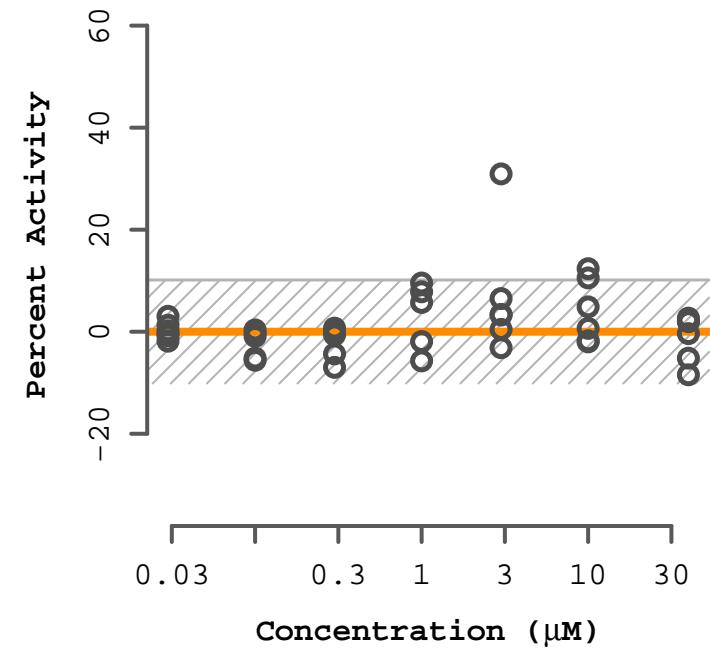
MAX_MEAN: 1.02

MAX_MED: 1.02

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-(2-Butoxyethoxy)ethanol
CHID: 21519 CASRN: 112-34-5
SPID(S): TP0001414C02
M4ID: 30204408

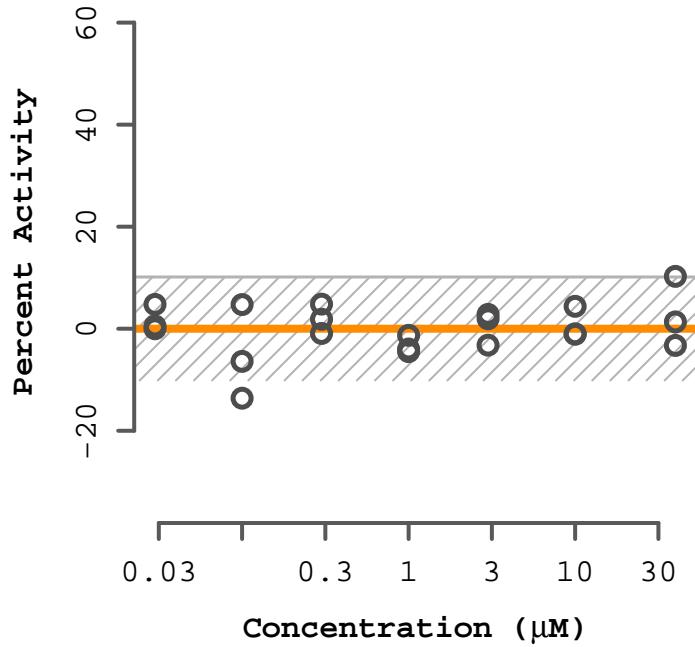
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	246.61	NA	NA
PROB:	1	NA	NA
RMSE:	6.8	NA	NA

MAX_MEAN: 6.91 MAX_MED: 5.75 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Hexanedioic acid
CHID: 21605 CASRN: 124-04-9
SPID(S): TP0001411H11
M4ID: 30204393

HILL MODEL: Not applicable.

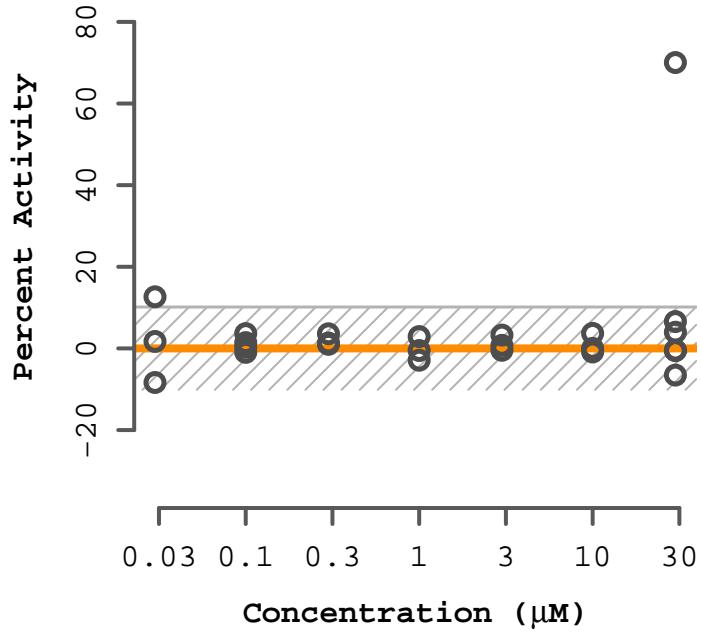
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	126.95	NA	NA
PROB:	1	NA	NA
RMSE:	4.87	NA	NA

MAX MEAN: 2.78 MAX MED: 1.95 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Nonanal

CHID: 21639 CASRN: 124-19-6

SPID(S): EPAPLT0154F04

M4ID: 30204207

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 163.39

PROB: 1

RMSE: 14.58

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 12.2

MAX_MED: 1.79

BMAD: 3.38

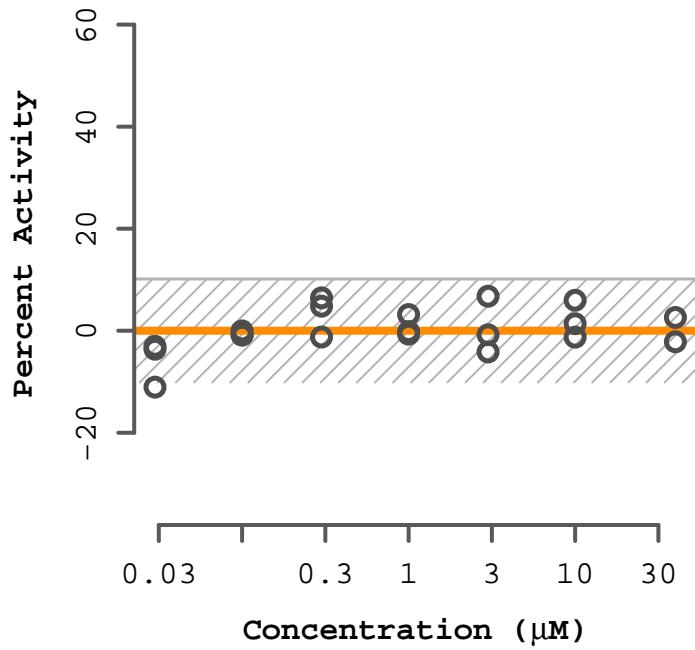
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: N,N-Diethylethanamine
CHID: 21837 CASRN: 100-37-8
SPID(S): TP0001412G09
M4ID: 30204389

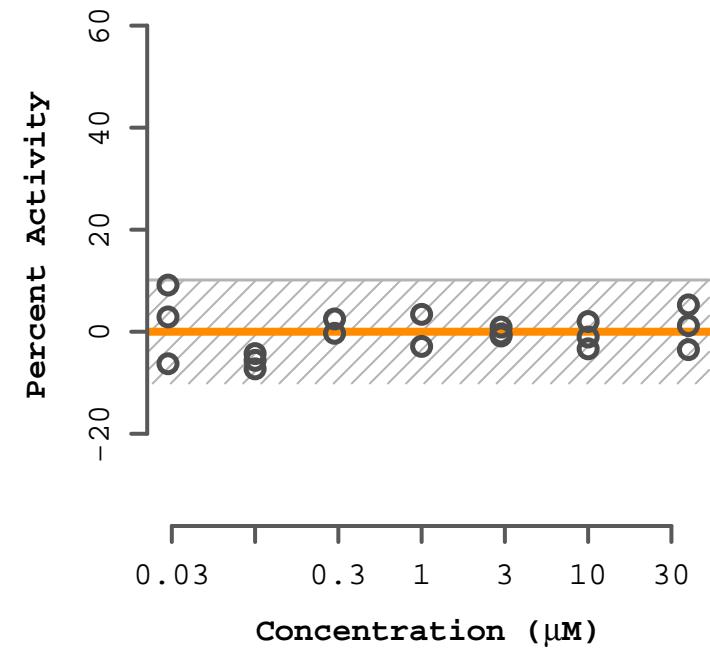
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 114.79	NA	NA
PROB: 1	NA	NA
RMSE: 4.11	NA	NA

MAX_MEAN: 3.35 MAX_MED: 4.82 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Methyl-2,4-pentanediol
CHID: 21885 CASRN: 107-41-5
SPID(S): TP0001411B06
M4ID: 30204433

HILL MODEL: Not applicable.

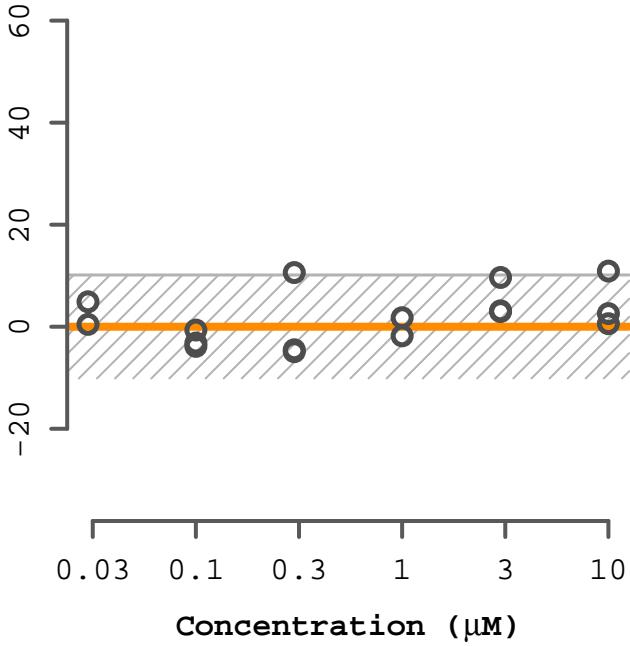
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	110.98	NA	NA
PROB:	1	NA	NA
RMSE:	4.1	NA	NA

MAX_MEAN: 1.93 MAX_MED: 2.9 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tributyl phosphate
CHID: 21986 CASRN: 126-73-8
SPID(S): TP0001412A02
M4ID: 30204632

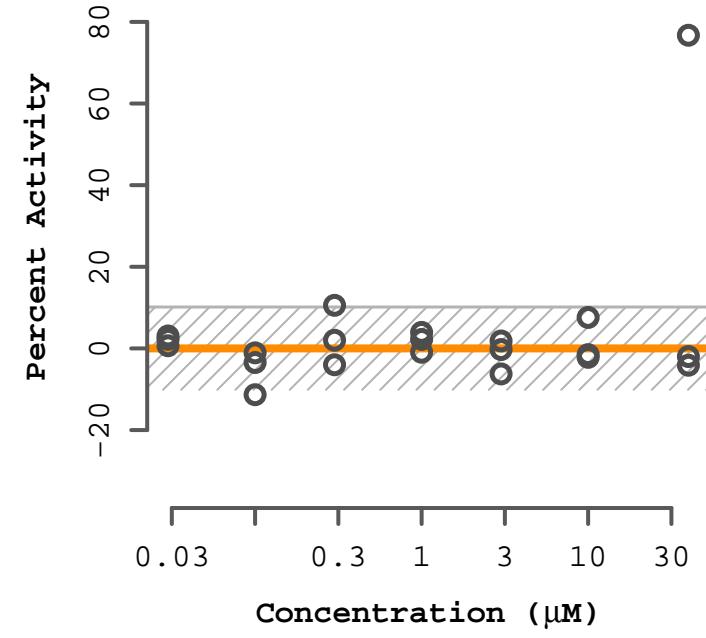
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 101.25	NA	NA
PROB: 1	NA	NA
RMSE: 5.31	NA	NA

MAX_MEAN: 5.25 MAX_MED: 3.12 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: EPN

CHID: 22174 CASRN: 2104-64-5

SPID(S): TP0001413D11

M4ID: 30204489

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 147.36

PROB: 1

RMSE: 17.36

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 23.5

MAX_MED: 2.24

BMAD: 3.38

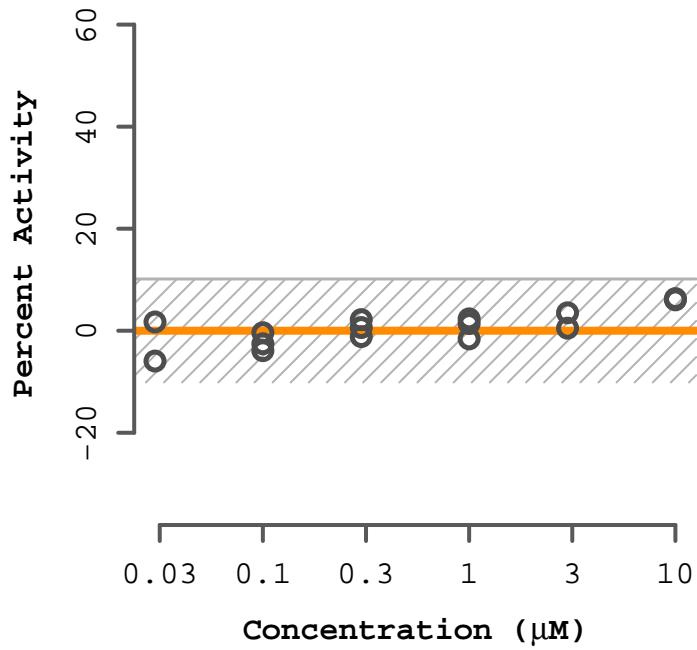
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-(1,1,3,3-Tetramethylbutyl)phenol
CHID: 22360 CASRN: 140-66-9
SPID(S): TP0001414E03
M4ID: 30204410

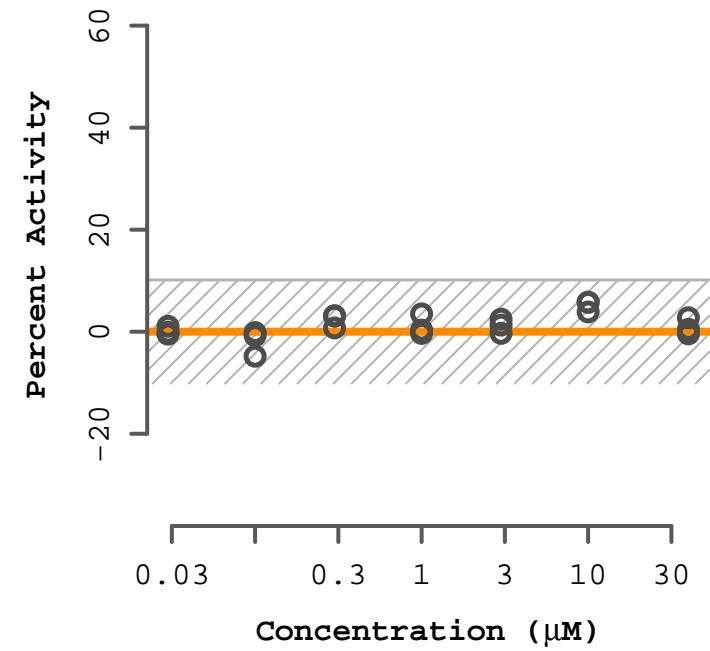
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 81.62	NA	NA
PROB: 1	NA	NA
RMSE: 3.32	NA	NA

MAX_MEAN: 6.18 MAX_MED: 6.18 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 17alpha-Estradiol
CHID: 22377 CASRN: 57-91-0
SPID(S): TP0001413F12
M4ID: 30204487

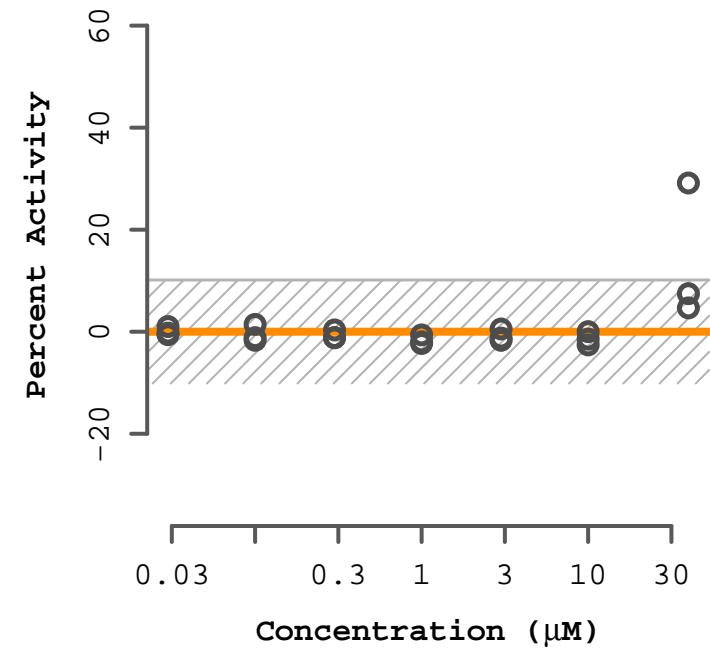
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 103.89	NA	NA
PROB: 1	NA	NA
RMSE: 2.72	NA	NA

MAX_MEAN: 5.15 MAX_MED: 5.79 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: meso-Hexestrol
CHID: 22381 CASRN: 84-16-2
SPID(S): TP0001413F05
M4ID: 30204453

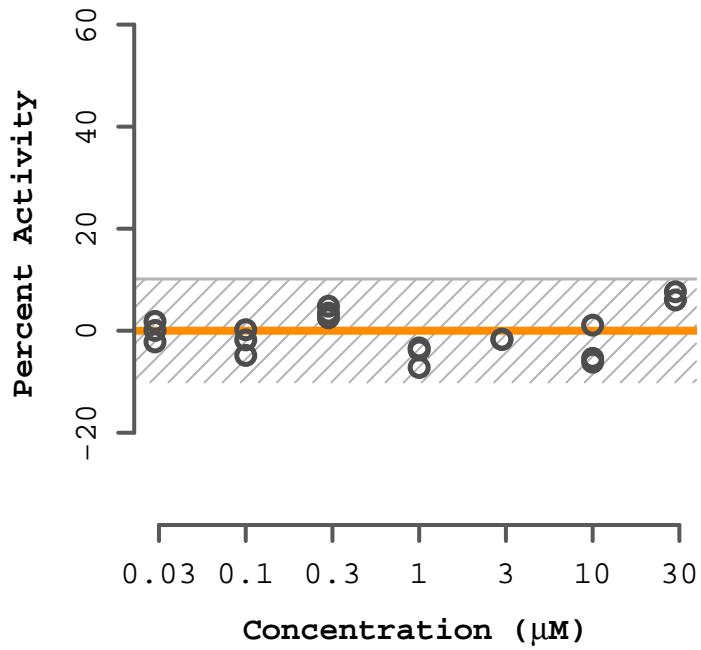
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 113.51	NA	NA
PROB: 1	NA	NA
RMSE: 6.77	NA	NA

MAX_MEAN: 13.8 MAX_MED: 7.47 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bis(4-hydroxyphenyl)methane
CHID: 22445 CASRN: 620-92-8
SPID(S): EPAPLT0154E04
M4ID: 30204198

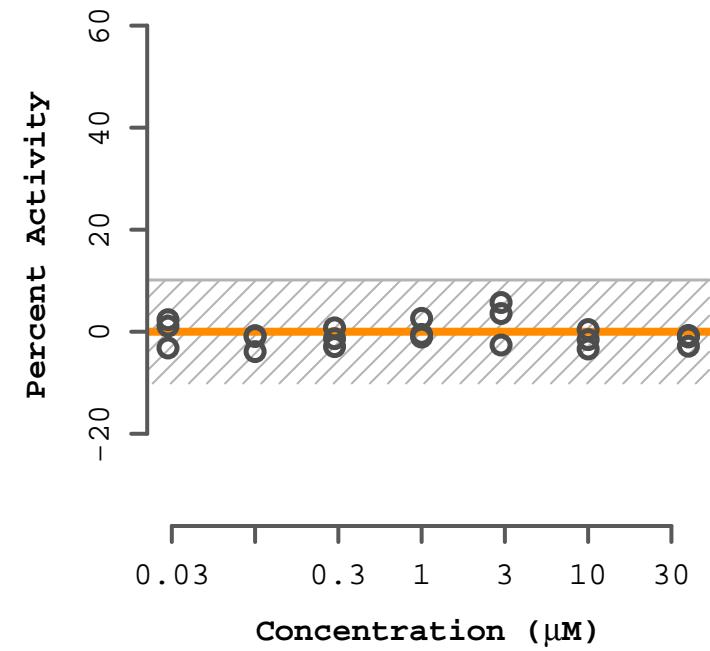
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 112.06	NA	NA
PROB: 1	NA	NA
RMSE: 4.12	NA	NA

MAX_MEAN: 6.84 MAX_MED: 6.84 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Aminofolic acid
CHID: 22588 CASRN: 54-62-6
SPID(S): TP0001413A11
M4ID: 30204617

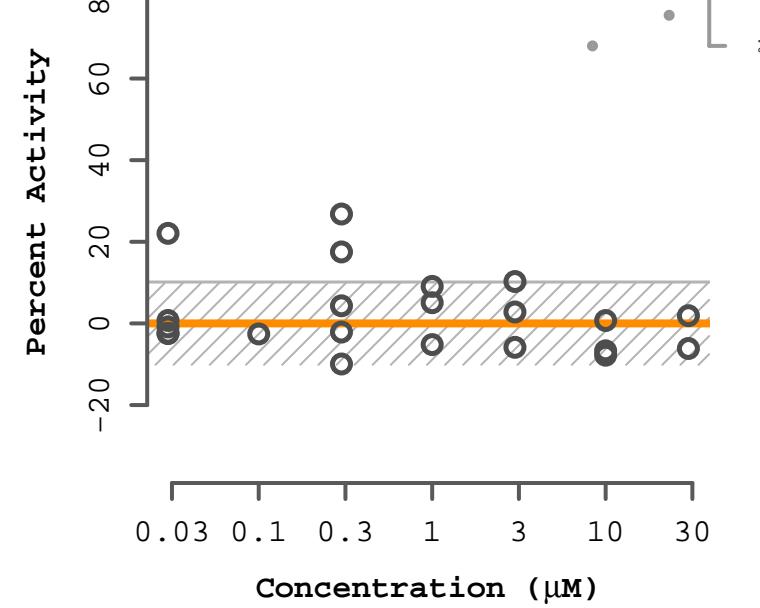
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	101.56	NA	NA
PROB:	1	NA	NA
RMSE:	2.47	NA	NA

MAX_MEAN: 2.24 MAX_MED: 3.57 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Buspirone

CHID: 22707 CASRN: 36505-84-7

SPID(S): EPAPLT0167A09

M4ID: 30204242 BRK

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 231.96

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 34.78

NA

NA

MAX_MEAN: 36.2

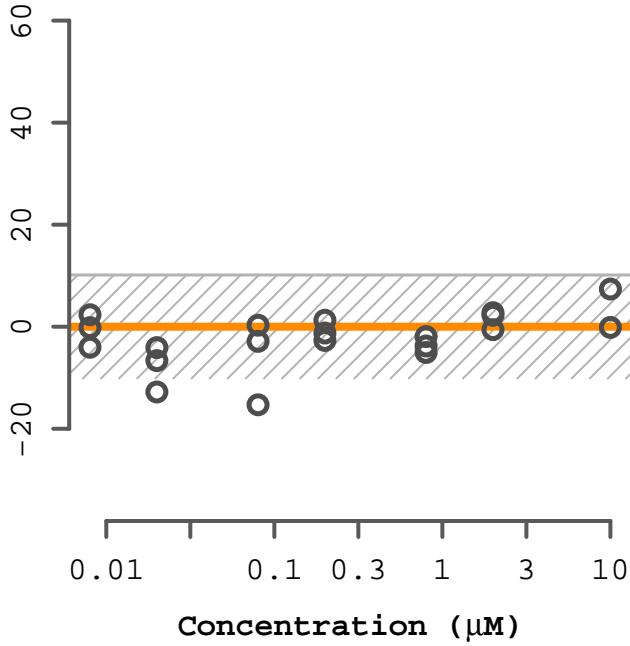
MAX_MED: 5.09

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Erythromycin

CHID: 22991 CASRN: 114-07-8

SPID(S): TP0001412H04

M4ID: 30204559

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 124.4

HILL

PROB: 1

GNLS

RMSE: 5.52

MAX_MEAN: 3.63

MAX_MED: 3.63

BMAD: 3.38

COFF: 10.1

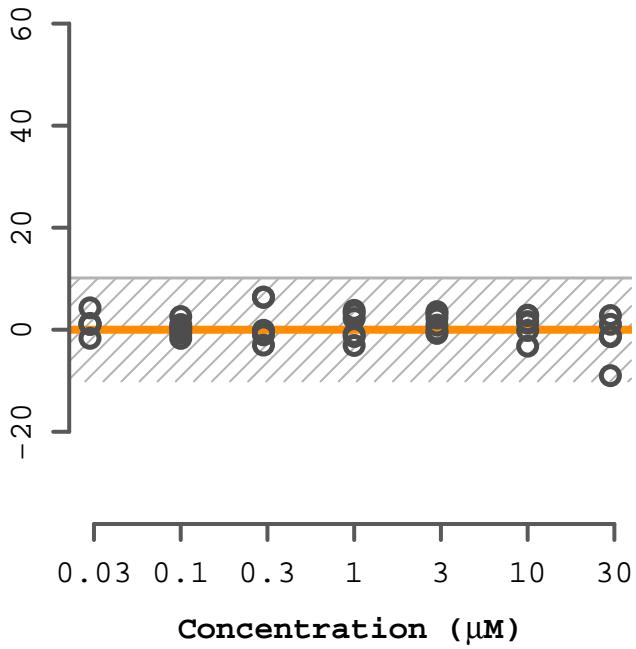
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Famotidine

CHID: 23039 CASRN: 76824-35-6

SPID(S): EPAPLT0167C01

M4ID: 30204253

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 193.71

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 2.88

NA

NA

MAX_MEAN: 1.73

MAX_MED: 1.61

BMAD: 3.38

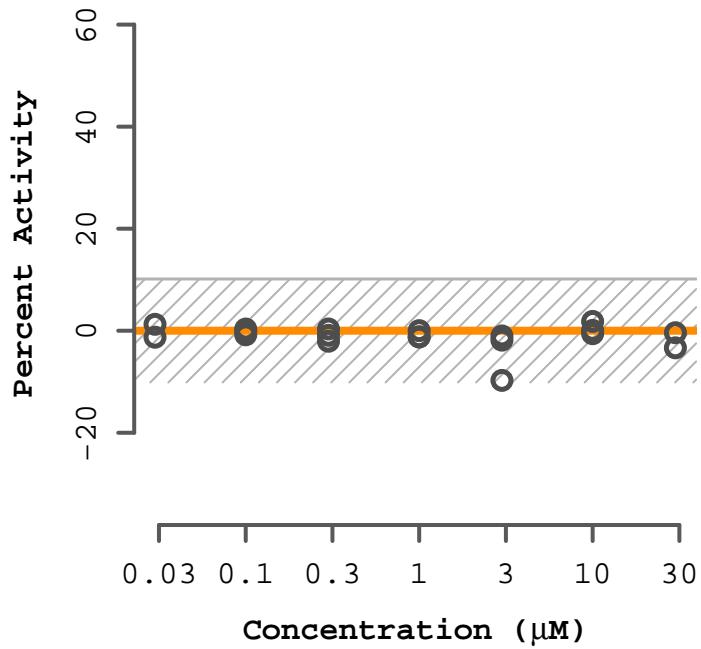
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Methylprednisolone

CHID: 23300 CASRN: 83-43-2

SPID(S): EPAPLT0154H04

M4ID: 30204227

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 80.42

PROB: 1

RMSE: 2.58

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 0.442

MAX_MED: 0.0731

BMAD: 3.38

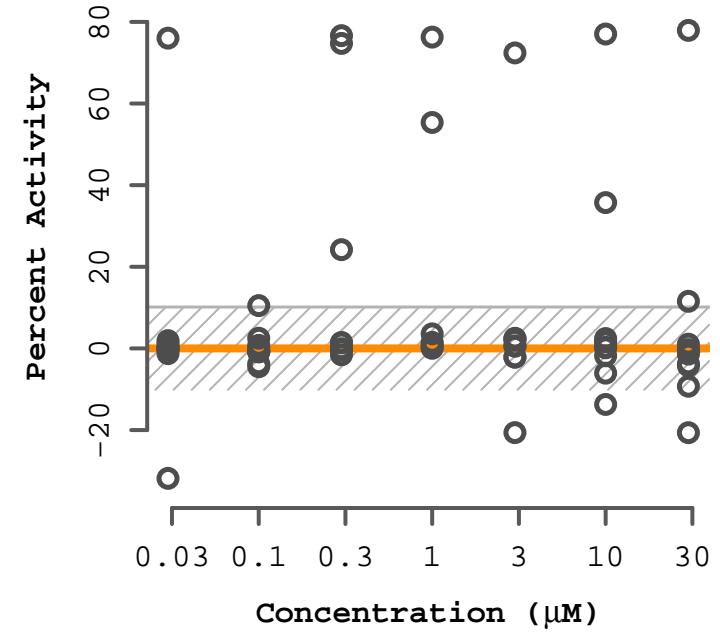
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Metoprolol

CHID: 23309 CASRN: 51384-51-1

SPID(S): EPAPLT0167A04

M4ID: 30204264

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 554.84

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 27.55

NA

NA

MAX_MEAN: 22.1

MAX_MED: 1.49

BMAD: 3.38

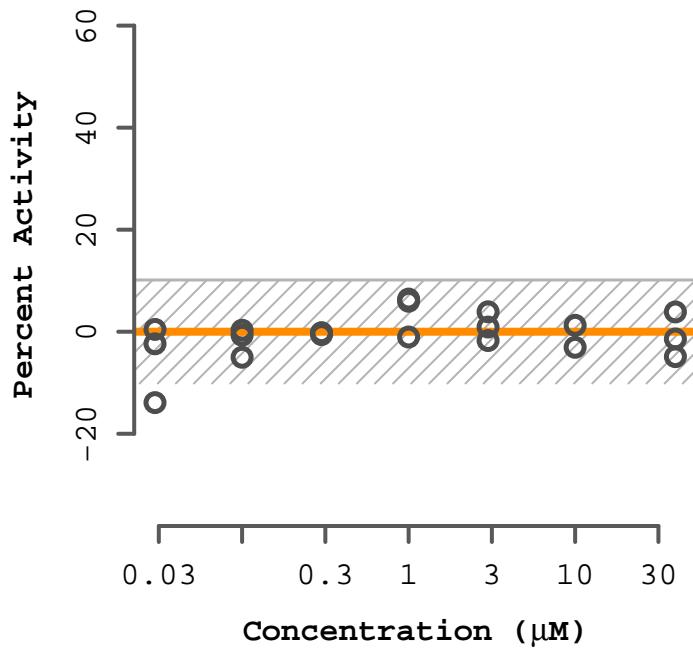
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Simvastatin

CHID: 23581 CASRN: 79902-63-9

SPID(S): TP0001412G08

M4ID: 30204467

HILL MODEL: Not applicable.

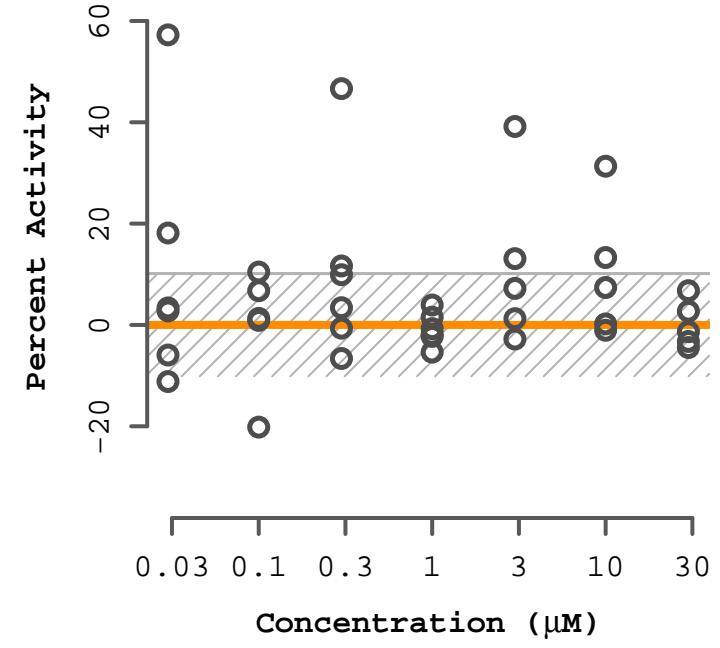
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	113.53	NA	NA
PROB:	1	NA	NA
RMSE:	4.32	NA	NA

MAX_MEAN: 3.79 MAX_MED: 5.99 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: D-Xylose

CHID: 23745 CASRN: 58-86-6

SPID(S): EPAPLT0154D11

M4ID: 30204194

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

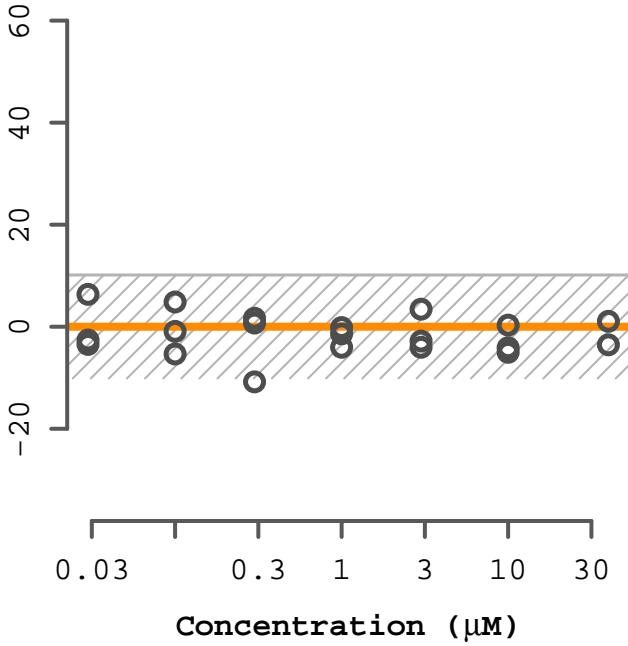
CNST	HILL	GNLS
AIC: 305.57	NA	NA
PROB: 1	NA	NA
RMSE: 16.17	NA	NA

MAX_MEAN: 11.6 MAX_MED: 7.39 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Acephate

CHID: 23846 CASRN: 30560-19-1

SPID(S): TP0001411F12

M4ID: 30204298

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 116.53

NA

NA

PROB:

1

NA

NA

RMSE: 4.13

NA

NA

MAX_MEAN: 0.099

MAX_MED: 0.725

BMAD: 3.38

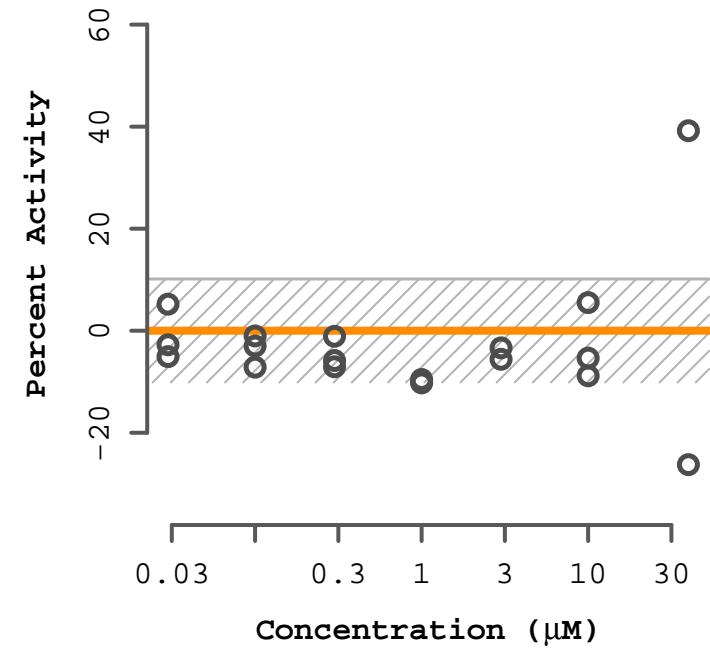
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Amitraz

CHID: 23871 CASRN: 33089-61-1

SPID(S): TP0001414H02

M4ID: 30204324

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 146.41

NA

NA

PROB:

1

NA

NA

RMSE: 12.37

NA

NA

MAX_MEAN: 6.49

MAX_MED: 6.49

BMAD: 3.38

COFF:

10.1

HIT-CALL:

0

FITC:

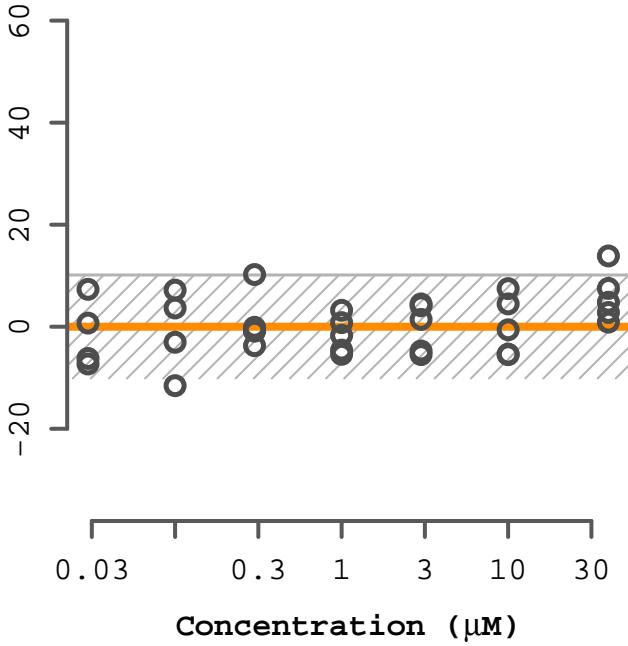
4

ACTP:

0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Clofentezine

CHID: 23881 CASRN: 74115-24-5

SPID(S): TP0001413B06

M4ID: 30204424

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 212.34

HILL

PROB: 1

GNLS

RMSE: 5.63

MAX_MEAN: 5.98

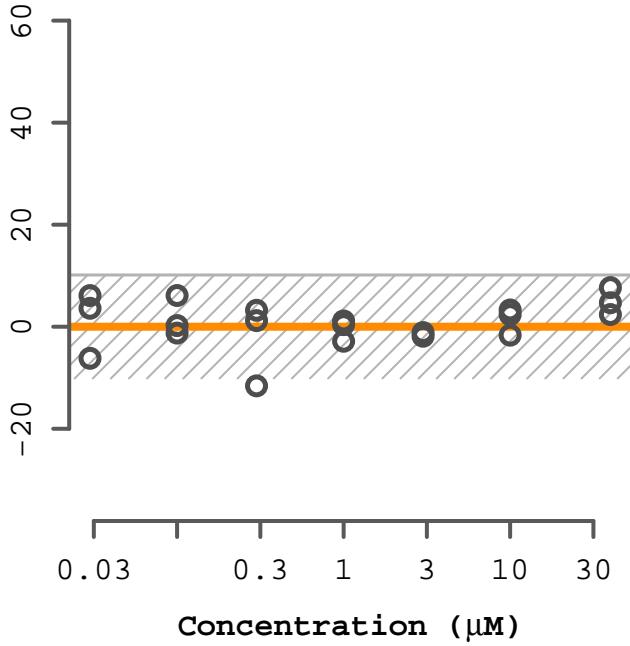
MAX_MED: 4.81

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bentazone

CHID: 23901 CASRN: 25057-89-0

SPID(S): TP0001414B03

M4ID: 30204347

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 118.52

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.44

NA

NA

MAX_MEAN: 4.93

MAX_MED: 4.71

BMAD: 3.38

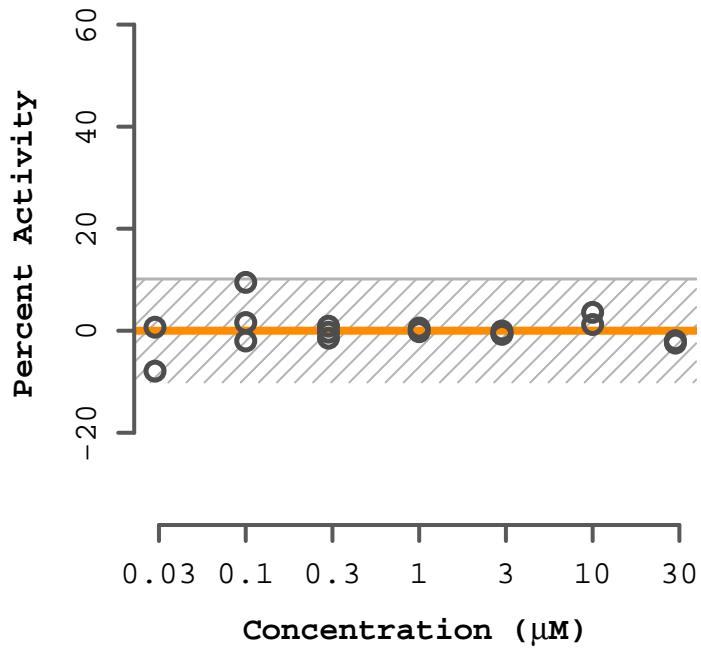
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Butylphthalyl butylglycolate
CHID: 23938 CASRN: 85-70-1
SPID(S): EPAPLT0154E10
M4ID: 30204204

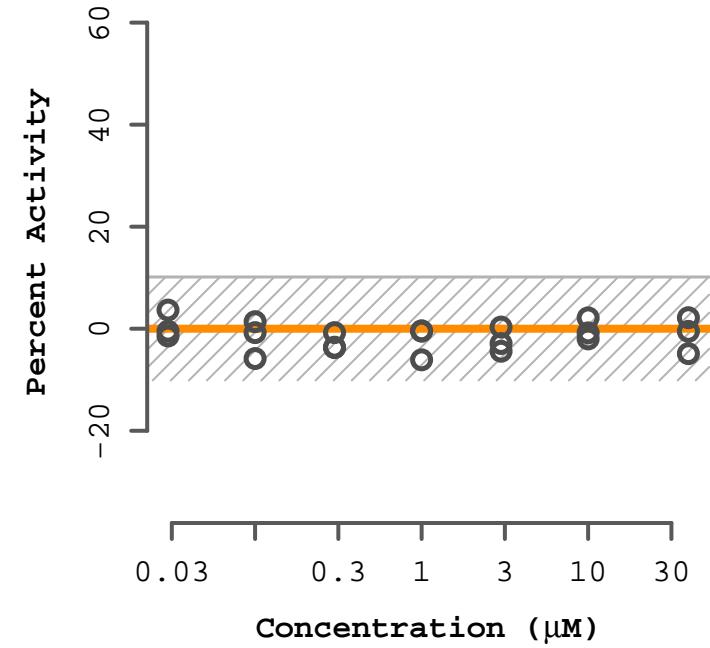
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 85.57	NA	NA
PROB: 1	NA	NA
RMSE: 3.33	NA	NA

MAX_MEAN: 3.03 MAX_MED: 2.41 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dimethipin

CHID: 24052 CASRN: 55290-64-7

SPID(S): TP0001414A09

M4 ID: 30204488

HILL MODEL: Not applicable.

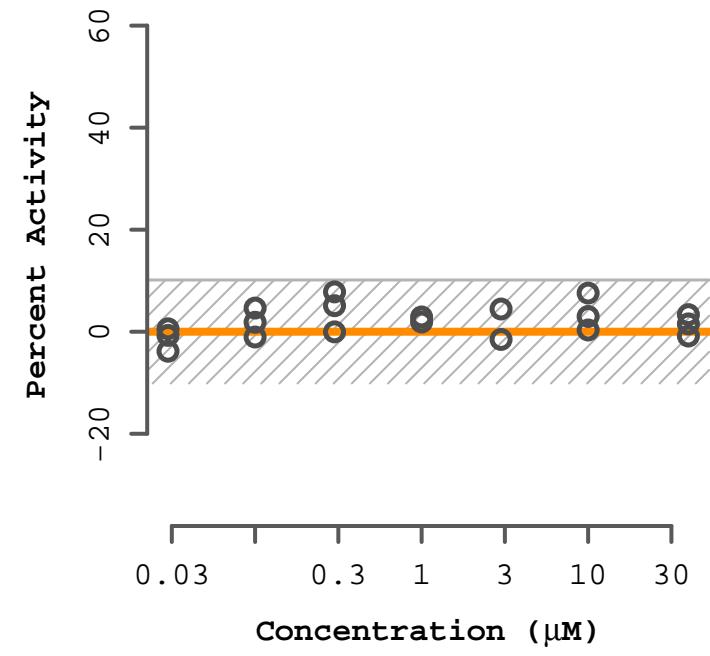
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	108.36	NA	NA
PROB:	1	NA	NA
RMSE:	2.95	NA	NA

MAX MEAN: 0.592 MAX MED: -0.467 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

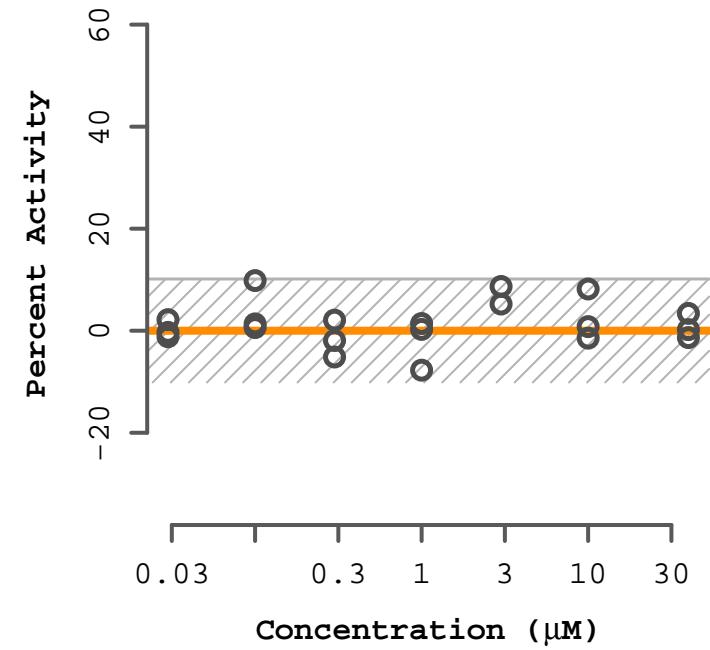
NAME: Dimethyl sulfate
CHID: 24055 CASRN: 77-78-1
SPID(S): TP0001414A06
M4ID: 30204622

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	105.18	NA	NA
PROB:	1	NA	NA
RMSE:	3.58	NA	NA
MAX_MEAN:	4.3	MAX_MED: 5.11	BMAD: 3.38
COFF:	10.1	HIT-CALL: 0	FITC: 4
ACTP:	0		

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,3-Dinitrobenzene

CHID: 24065 CASRN: 99-65-0

SPID(S): TP0001413A09

M4ID: 30204555

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 117.4

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.4

NA

NA

MAX_MEAN: 6.96

MAX_MED: 6.96

BMAD: 3.38

COFF: 10.1

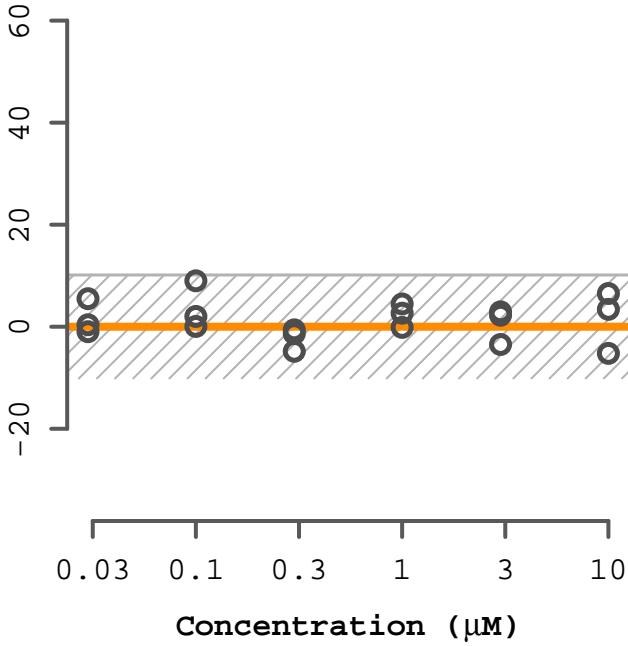
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fenamiphos

CHID: 24102 CASRN: 22224-92-6

SPID(S): TP0001414B10

M4ID: 30204614

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 103.47

HILL

PROB: 1

GNLS

RMSE: 3.92

MAX_MEAN: 3.7

MAX_MED: 3.41

BMAD: 3.38

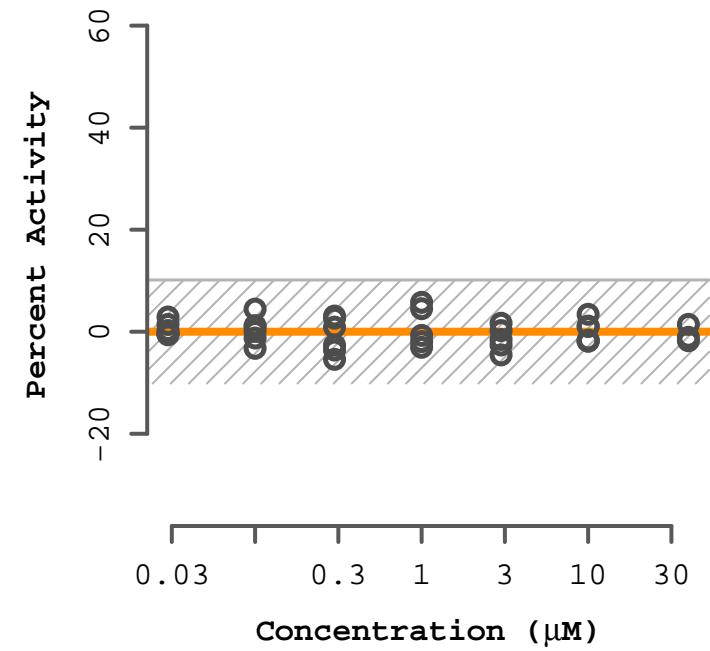
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Mepiquat chloride
CHID: 24170 CASRN: 24307-26-4
SPID(S): TP0001413D03
M4ID: 30204353

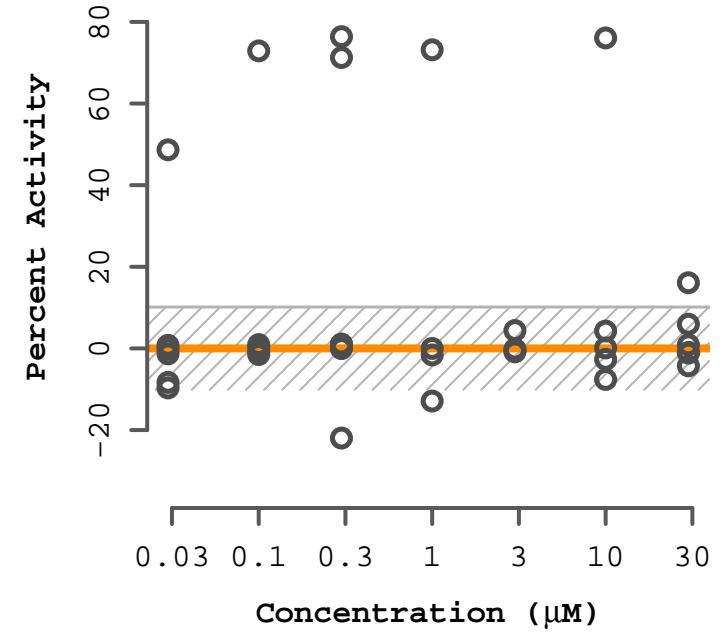
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	179.18	NA	NA
PROB:	1	NA	NA
RMSE:	2.55	NA	NA

MAX_MEAN: 0.922 MAX_MED: 0.531 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Methamidophos

CHID: 24177 CASRN: 10265-92-6

SPID(S): EPAPLT0167A08

M4ID: 30204263

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 327.2

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 29.34

NA

NA

MAX_MEAN: 21.3

MAX_MED: 1.04

BMAD: 3.38

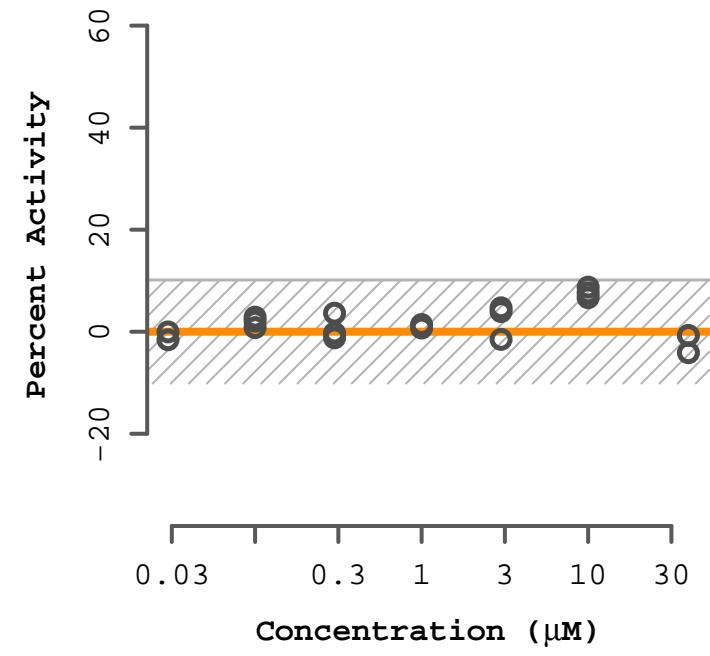
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Oxadiazon

CHID: 24239 CASRN: 19666-30-9

SPID(S): TP0001414G02

M4ID: 30204342

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 102

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 3.87

NA

NA

MAX_MEAN: 7.68

MAX_MED: 7.63

BMAD: 3.38

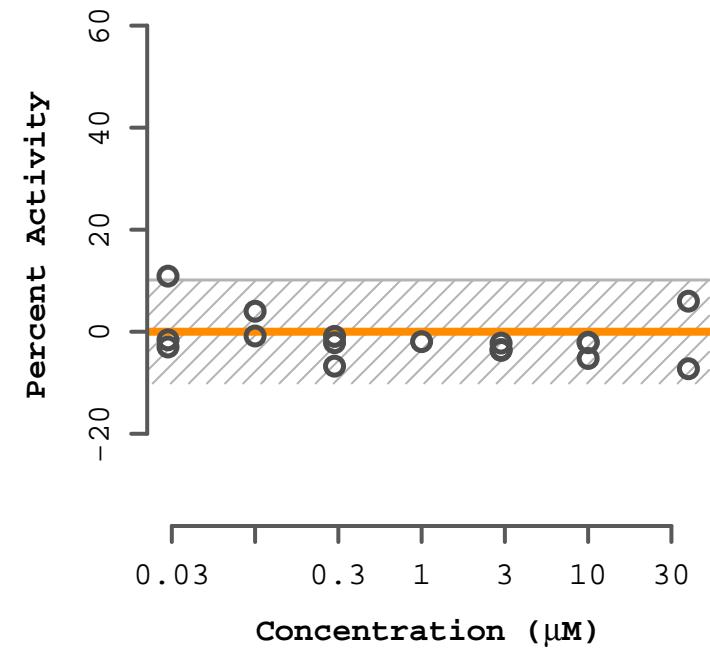
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Paclobutrazol

CHID: 24242 CASRN: 76738-62-0

SPID(S): TP0001411F08

M4ID: 30204301

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 102.95

PROB: 1

RMSE: 4.58

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 2.11

MAX_MED: 1.6

BMAD: 3.38

COFF: 10.1

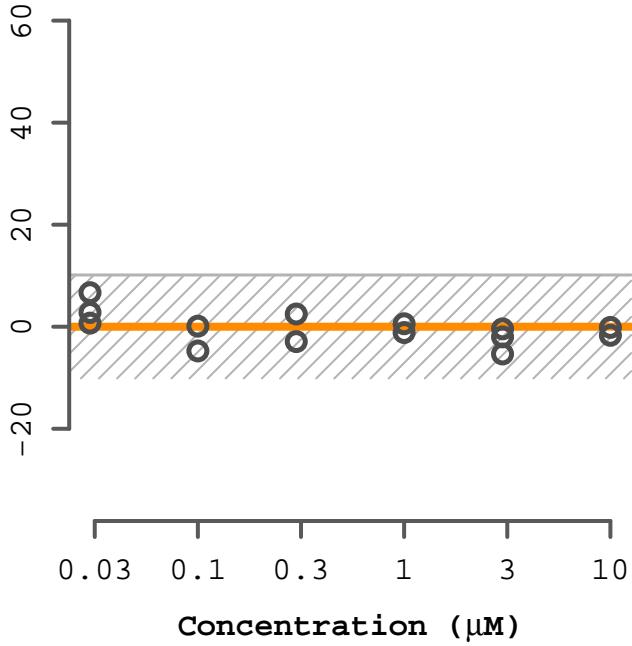
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Imazethapyr

CHID: 24287 CASRN: 81335-77-5

SPID(S): TP0001411F06

M4ID: 30204326

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 72.88

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 3.01

NA

NA

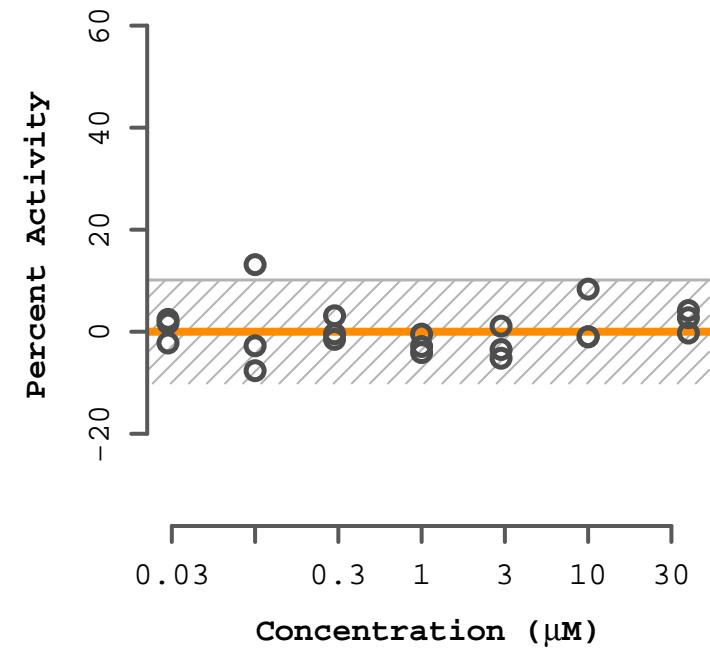
MAX_MEAN: 3.4

MAX_MED: 2.8

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Carbendazim

CHID: 24729 CASRN: 10605-21-7

SPID(S): TP0001412D10

M4ID: 30204536

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 122.96

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.51

NA

NA

MAX_MEAN: 2.22

MAX_MED: 2.75

BMAD: 3.38

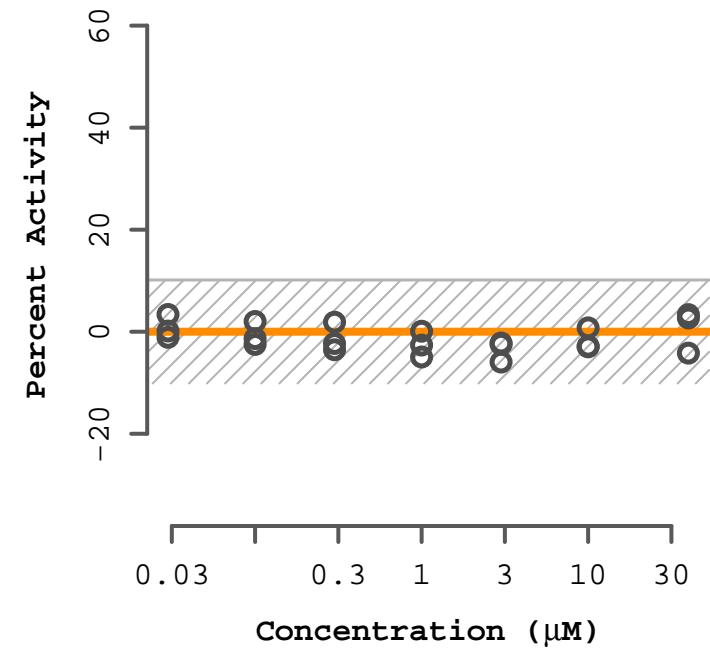
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 3,7-Dimethyl-2,6-octadienal
CHID: 24836 CASRN: 5392-40-5
SPID(S): TP0001414H04
M4ID: 30204414

HILL MODEL: Not applicable.

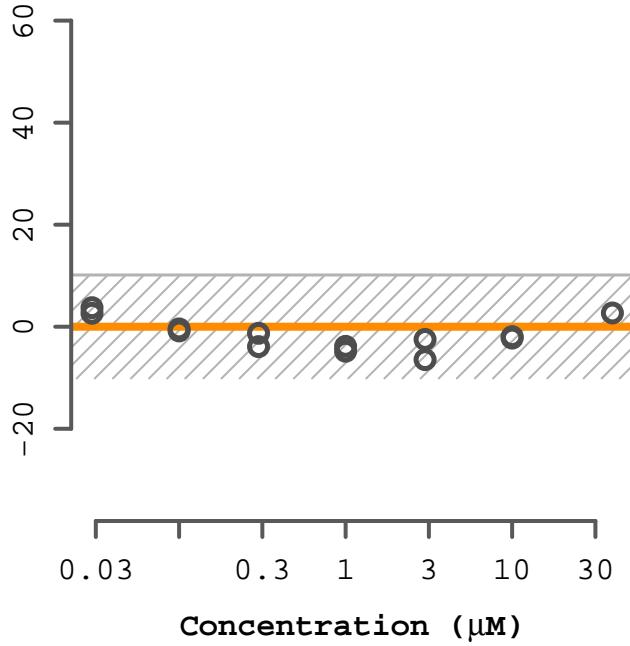
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 109.34	NA	NA
PROB: 1	NA	NA
RMSE: 2.9	NA	NA

MAX_MEAN: 0.814 MAX_MED: 2.69 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cumene hydroperoxide
CHID: 24869 CASRN: 80-15-9
SPID(S): TP0001411D03
M4ID: 30204545

HILL MODEL: Not applicable.

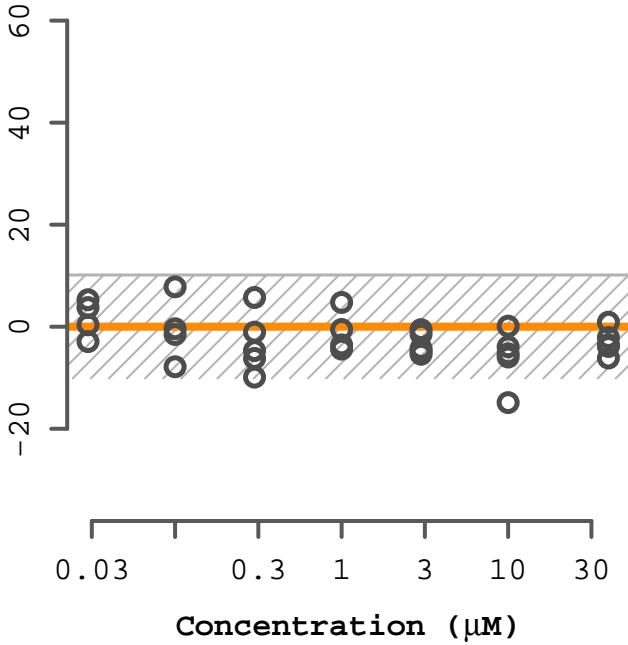
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 71.56	NA	NA
PROB: 1	NA	NA
RMSE: 3.28	NA	NA

MAX_MEAN: 3.19 MAX_MED: 3.19 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dibromoacetonitrile
CHID: 24940 CASRN: 3252-43-5
SPID(S): TP0001413B07
M4ID: 30204574

HILL MODEL: Not applicable.

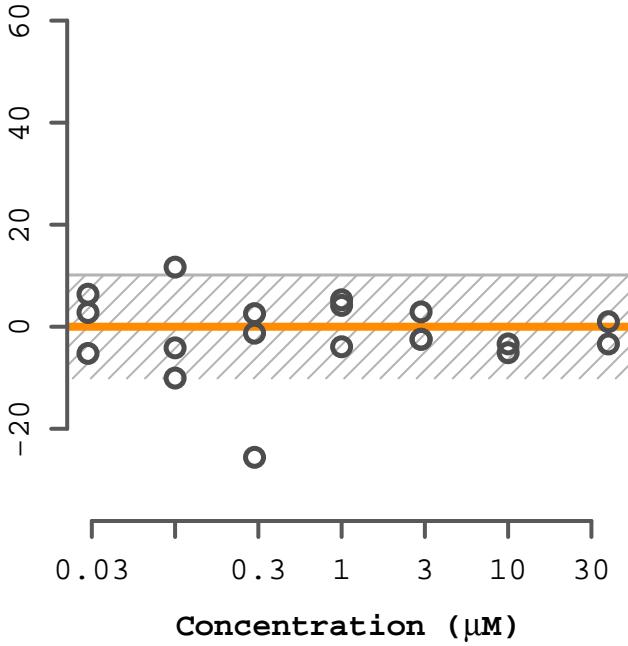
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	199.96	NA	NA
PROB:	1	NA	NA
RMSE:	5.2	NA	NA

MAX_MEAN: 1.64 MAX_MED: 2.08 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diethylene glycol monomethyl ether
CHID: 25049 CASRN: 111-77-3
SPID(S): TP0001414B06
M4ID: 30204406

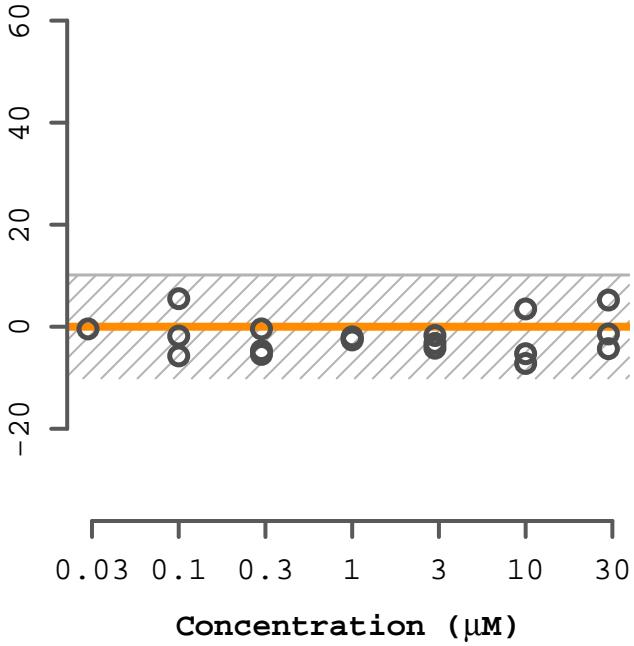
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 128.83	NA	NA
PROB: 1	NA	NA
RMSE: 7.69	NA	NA
MAX_MEAN: 1.85	MAX_MED: 4.19	BMAD: 3.38
COFF: 10.1	HIT-CALL: 0	FITC: 4
		ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diethylenetriamine

CHID: 25050 CASRN: 111-40-0

SPID(S): EPAPLT0154C09

M4ID: 30204181

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 111.69

PROB: 1

RMSE: 4.01

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: -0.153

MAX_MED: -0.426

BMAD: 3.38

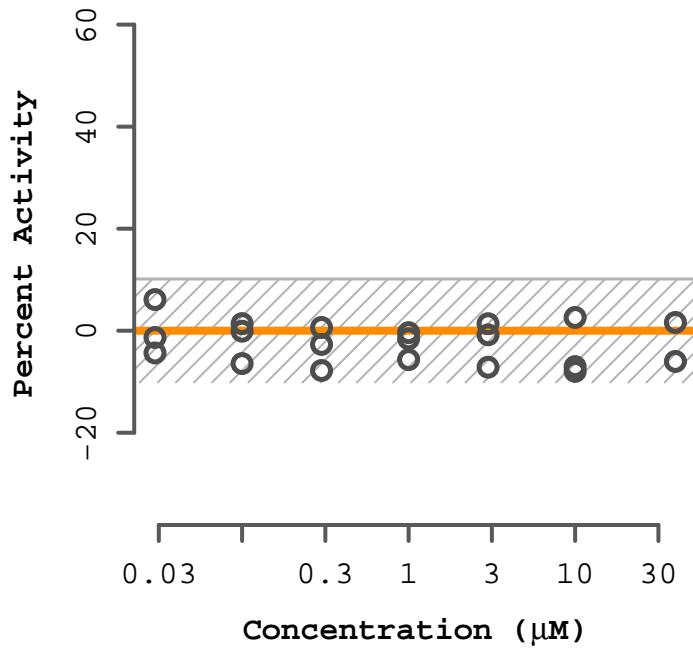
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bis(2-ethylhexyl) decanedioate
CHID: 25055 CASRN: 122-62-3
SPID(S): TP0001412B07
M4ID: 30204611

HILL MODEL: Not applicable.

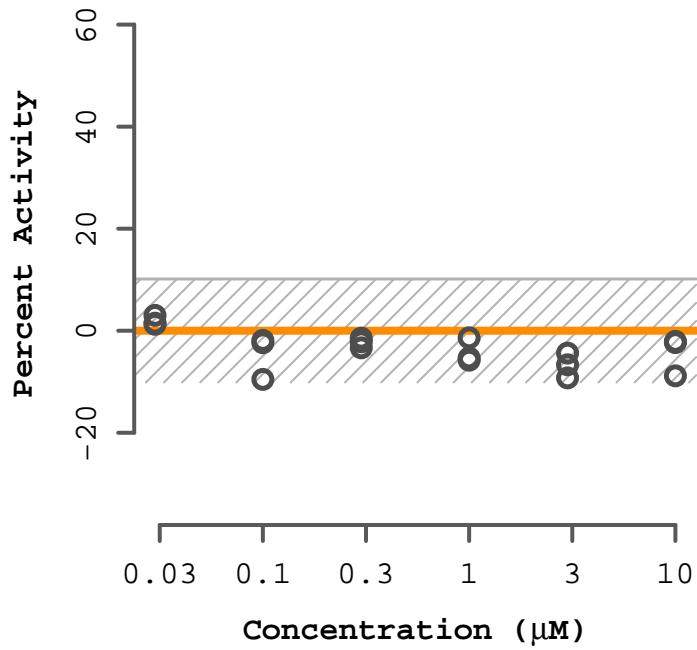
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	121.56	NA	NA
PROB:	1	NA	NA
RMSE:	4.56	NA	NA

MAX_MEAN: 0.151 MAX_MED: -0.0868 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

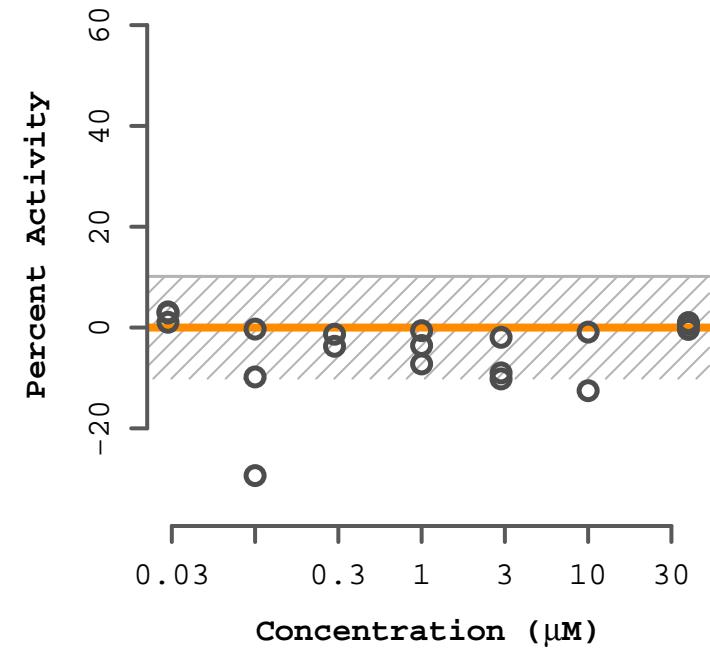
NAME: 1,3-Diphenylguanidine
CHID: 25178 CASRN: 102-06-7
SPID(S): TP0001412D12
M4ID: 30204576

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	111.57	NA	NA
PROB:	1	NA	NA
RMSE:	4.89	NA	NA
MAX_MEAN:	1.92	MAX_MED: 1.49	BMAD: 3.38
COFF:	10.1	HIT-CALL: 0	FITC: 4
ACTP:	0		

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

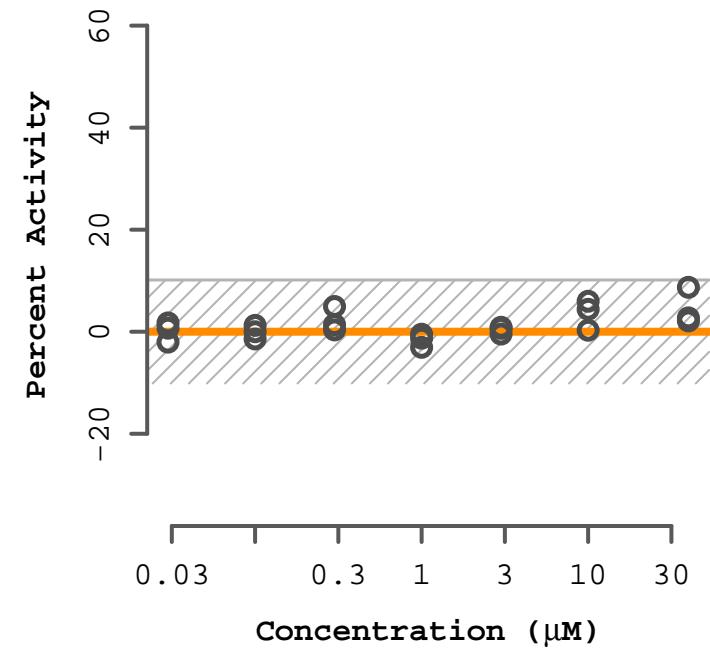
NAME: Isoproterenol hydrochloride
CHID: 25486 CASRN: 51-30-9
SPID(S): TP0001411G01
M4ID: 30204573

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	131.01	NA	NA
PROB:	1	NA	NA
RMSE:	8.6	NA	NA
MAX_MEAN:	2.35	MAX_MED: 2.89	BMAD: 3.38
COFF:	10.1	HIT-CALL: 0	FITC: 4
ACTP:	0		

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,2-Phenylenediamine
CHID: 25881 CASRN: 95-54-5
SPID(S): TP0001413F08
M4ID: 30204442

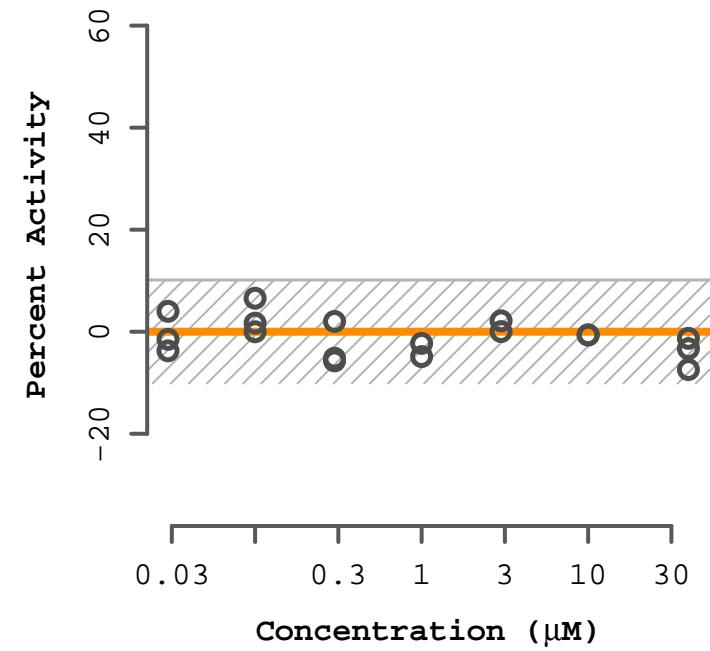
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 105.73	NA	NA
PROB: 1	NA	NA
RMSE: 3.03	NA	NA

MAX_MEAN: 4.52 MAX_MED: 4.4 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sodium dehydroacetate
CHID: 26029 CASRN: 4418-26-2
SPID(S): TP0001414G08
M4ID: 30204403

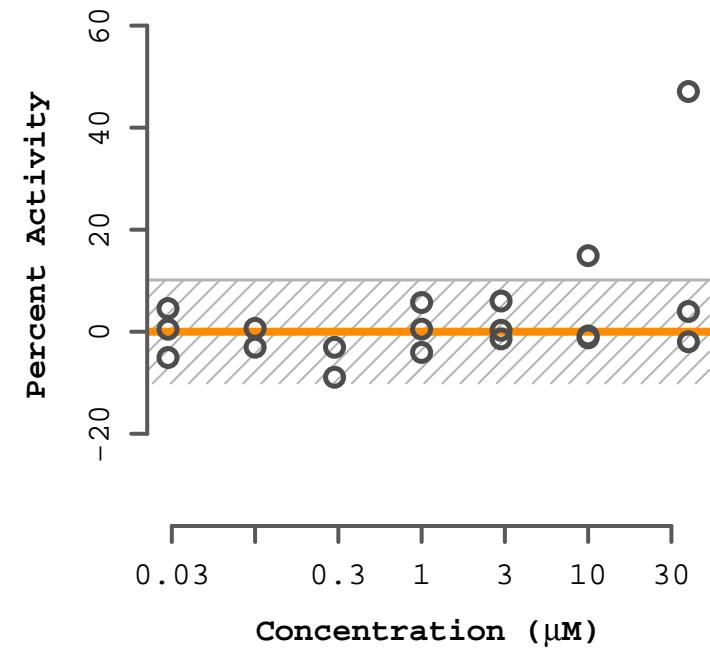
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 101.55	NA	NA
PROB: 1	NA	NA
RMSE: 3.68	NA	NA

MAX_MEAN: 2.75 MAX_MED: 1.69 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

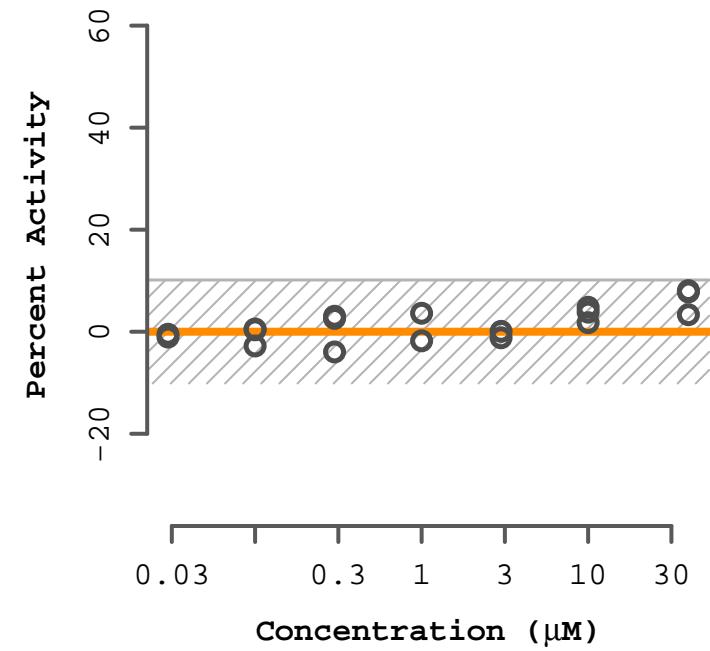
NAME: 1,2,3-Trichlorobenzene
CHID: 26193 CASRN: 87-61-6
SPID(S): TP0001411F02
M4ID: 30204554

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	132.67	NA	NA
PROB:	1	NA	NA
RMSE:	11.91	NA	NA
MAX_MEAN:	16.4	MAX_MED: 3.98	BMAD: 3.38
COFF:	10.1	HIT-CALL: 0	FITC: 4
ACTP:	0		

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tris(1,3-dichloro-2-propyl) phosphate

CHID: 26261 CASRN: 13674-87-8

SPID(S): TP0001413C07

M4ID: 30204630

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 100.14

PROB: 1

RMSE: 3.63

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 6.4

MAX_MED: 7.8

BMAD: 3.38

COFF: 10.1

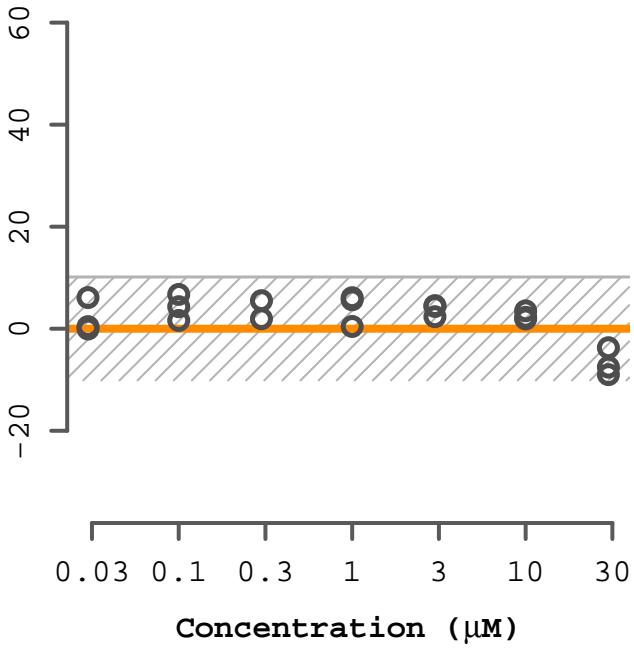
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: p-Xylenol blue
CHID: 26303 CASRN: 125-31-5
SPID(S): EPAPLT0154F10
M4ID: 30204213

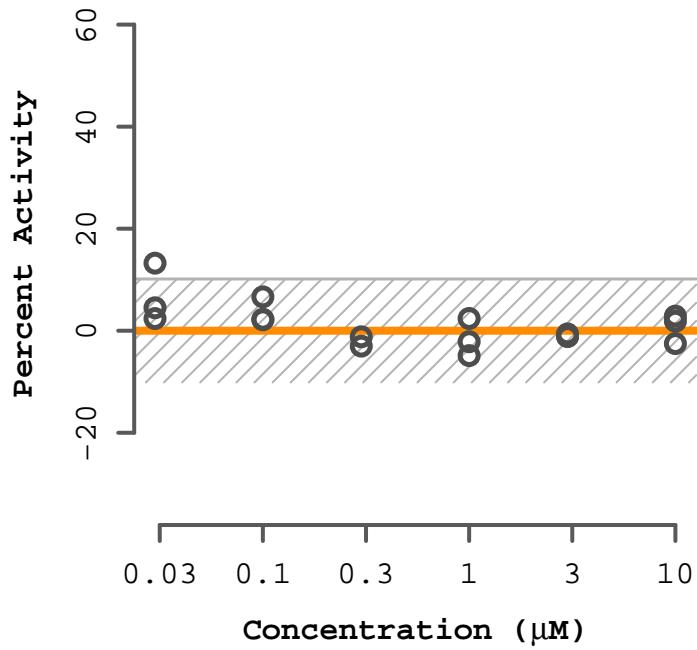
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 122.6	NA	NA
PROB: 1	NA	NA
RMSE: 4.61	NA	NA

MAX_MEAN: 4.23 MAX_MED: 5.68 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Tert-Butyl-5-methylphenol
CHID: 26529 CASRN: 88-60-8
SPID(S): TP0001412C10
M4ID: 30204544

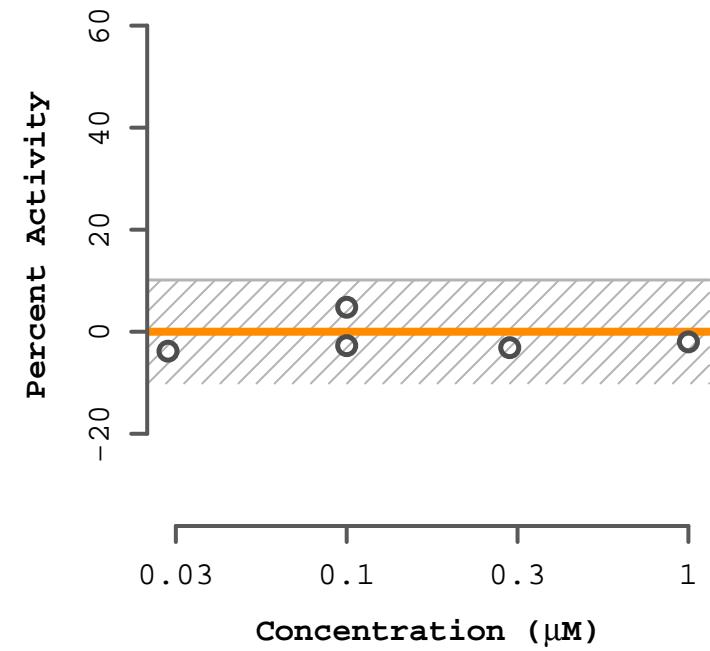
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 97.66	NA	NA
PROB: 1	NA	NA
RMSE: 4.34	NA	NA

MAX_MEAN: 6.72 MAX_MED: 4.5 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Triethylene glycol bis(2-ethylhexanoate)
CHID: 26564 CASRN: 94-28-0
SPID(S): TP0001411H02
M4ID: 30204575

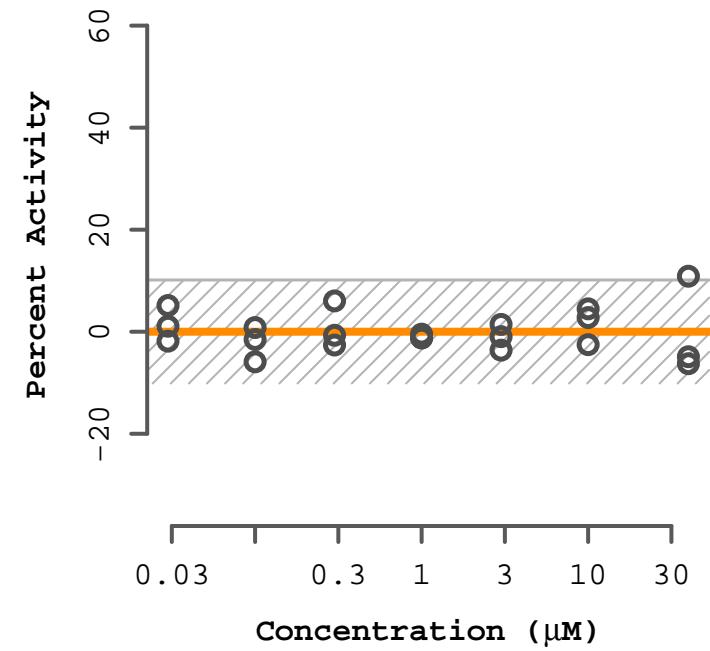
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 29.53	NA	NA
PROB: 1	NA	NA
RMSE: 3.42	NA	NA

MAX_MEAN: 1.04 MAX_MED: 1.04 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 3-Pyridinecarbonitrile
CHID: 26665 CASRN: 100-54-9
SPID(S): TP0001413G07
M4ID: 30204400

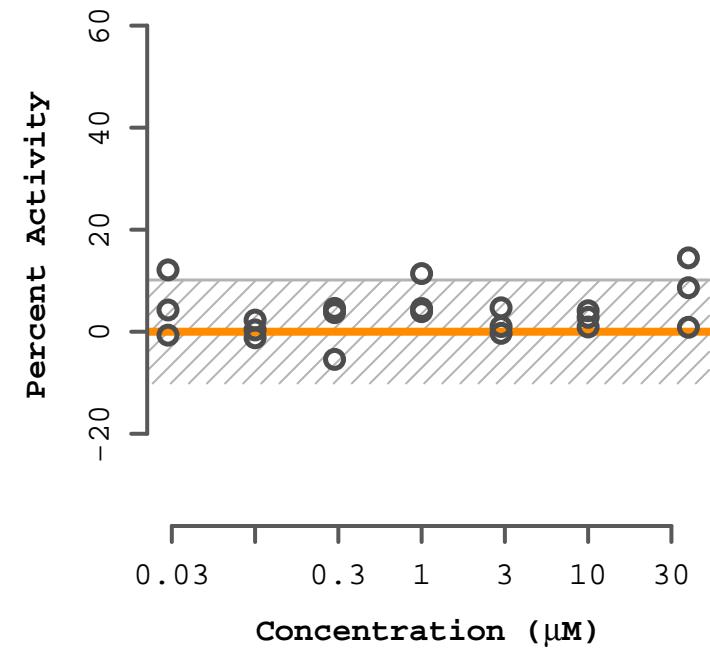
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 120.67	NA	NA
PROB: 1	NA	NA
RMSE: 4.07	NA	NA

MAX_MEAN: 1.58 MAX_MED: 2.79 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

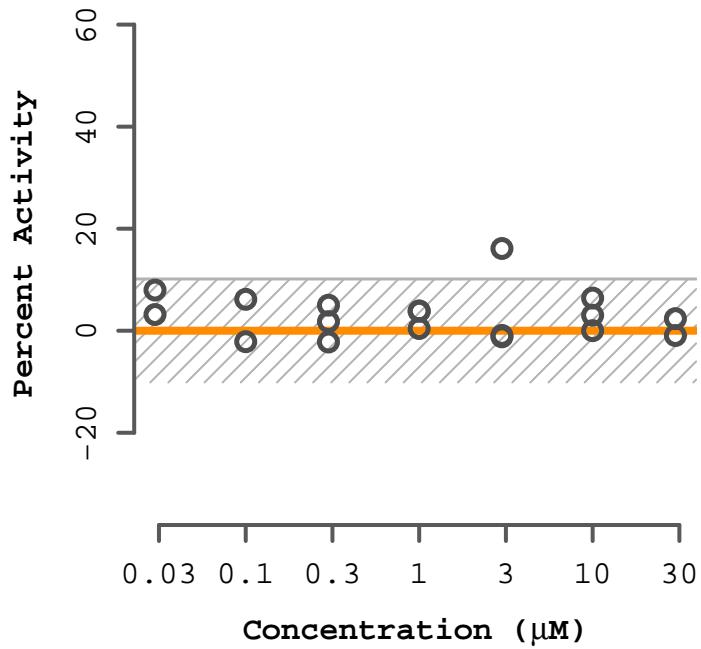
NAME: Dodecylbenzene
CHID: 26994 CASRN: 123-01-3
SPID(S): TP0001412G06
M4ID: 30204569

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 135.74	NA	NA
PROB: 1	NA	NA
RMSE: 5.93	NA	NA
MAX_MEAN: 8	MAX_MED: 8.63	BMAD: 3.38
COFF: 10.1	HIT-CALL: 0	FITC: 4
		ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,2-Dimethylpropane-1,3-diol
CHID: 27036 CASRN: 126-30-7
SPID(S): EPAPLT0154G05
M4ID: 30204218

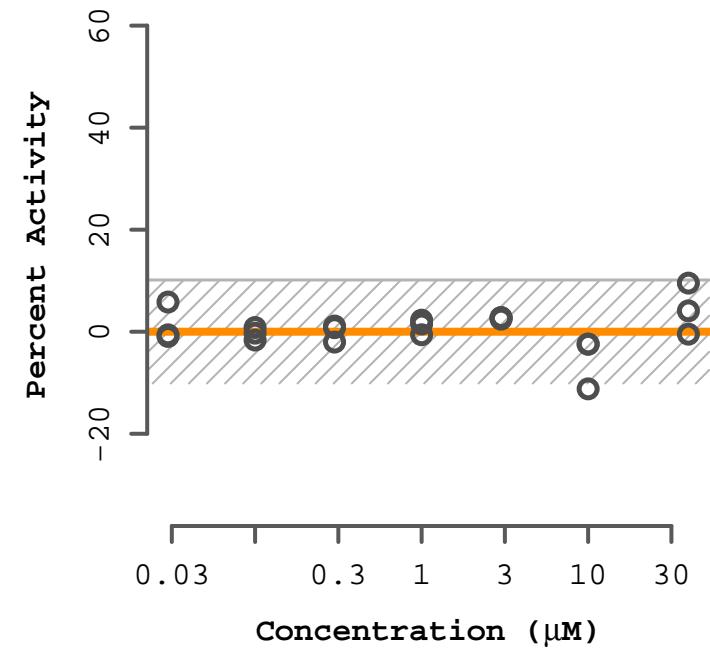
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	110.26	NA	NA
PROB:	1	NA	NA
RMSE:	5.26	NA	NA

MAX_MEAN: 5.58 MAX_MED: 5.58 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sodium dimethyldithiocarbamate
CHID: 27050 CASRN: 128-04-1
SPID(S): TP0001413H12
M4ID: 30204457

HILL MODEL: Not applicable.

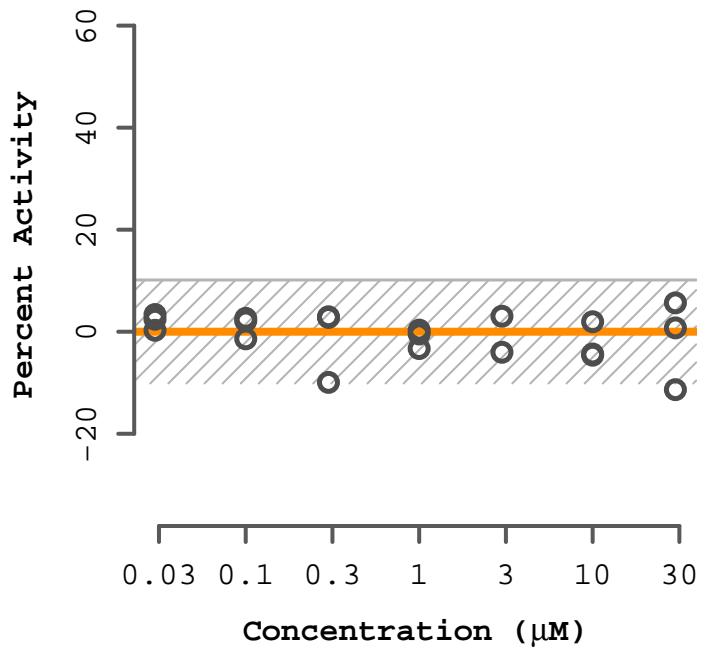
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 114.57	NA	NA
PROB: 1	NA	NA
RMSE: 3.9	NA	NA

MAX_MEAN: 4.39 MAX_MED: 4.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bis(2-ethylhexyl) phosphate

CHID: 27134 CASRN: 298-07-7

SPID(S): EPAPLT0154A06

M4ID: 30204157

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 117.47

PROB: 1

RMSE: 4.4

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 2.04

MAX_MED: 2.81

BMAD: 3.38

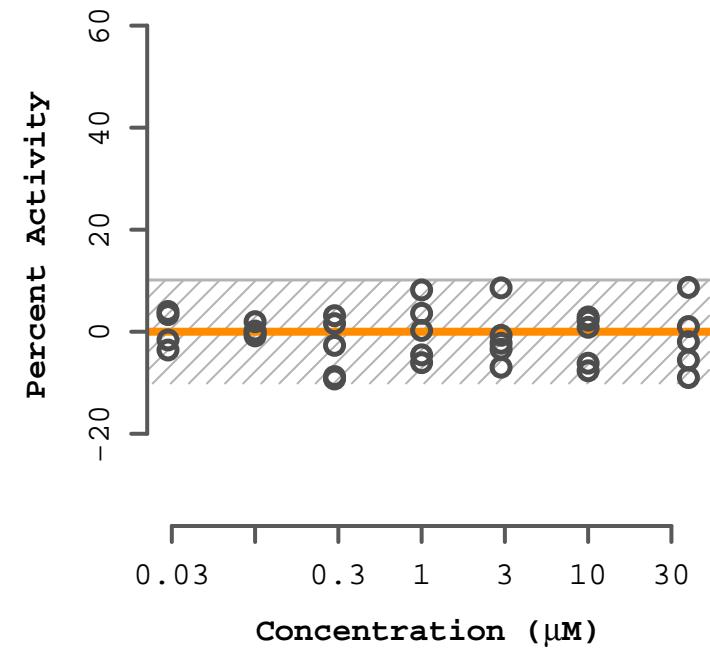
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4,4'-Bipyridine
CHID: 27200 CASRN: 553-26-4
SPID(S): TP0001413B08
M4ID: 30204593

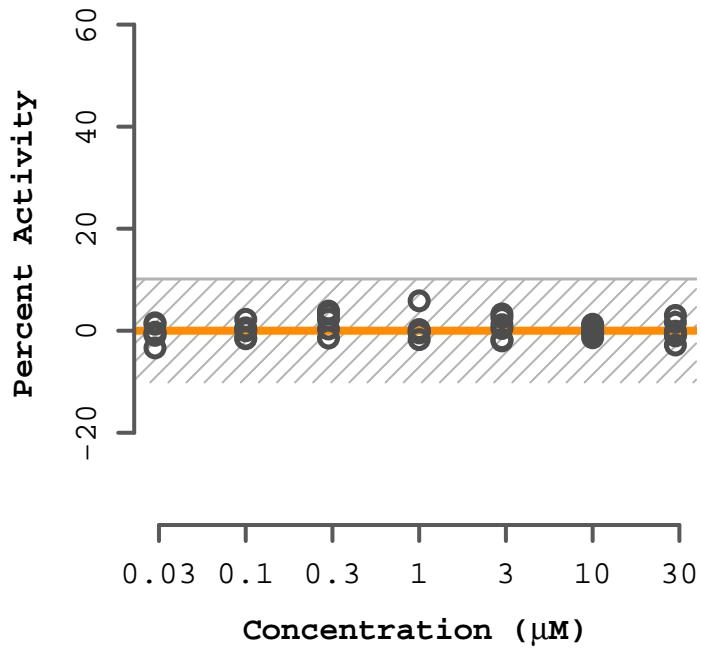
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	204.15	NA	NA
PROB:	1	NA	NA
RMSE:	4.95	NA	NA

MAX_MEAN: 0.544 MAX_MED: 0.885 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Octamethylcyclotetrasiloxane
CHID: 27205 CASRN: 556-67-2
SPID(S): EPAPLT0167H03
M4ID: 30204266

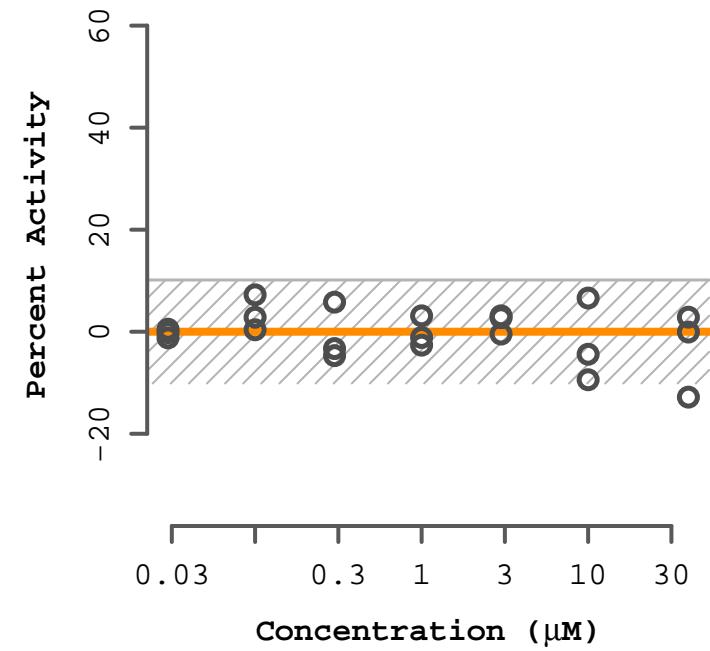
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	175.24	NA	NA
PROB:	1	NA	NA
RMSE:	1.99	NA	NA

MAX_MEAN: 1.99 MAX_MED: 2.74 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sorbitan, mono-(9Z)-9-octadecenoate

CHID: 27397 CASRN: 1338-43-8

SPID(S): TP0001411C09

M4ID: 30204663

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 127.22

PROB: 1

RMSE: 4.82

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 3.49

MAX_MED: 2.84

BMAD: 3.38

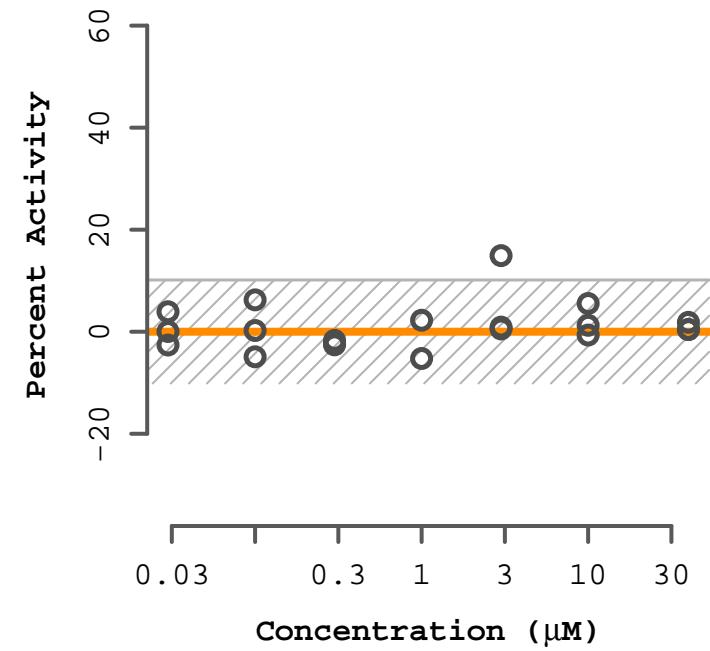
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,2,4-Trimethyl-1,3-pentanediol diisobutyrate

CHID: 27635 CASRN: 6846-50-0

SPID(S): TP0001414G07

M4ID: 30204581

HILL MODEL: Not applicable.

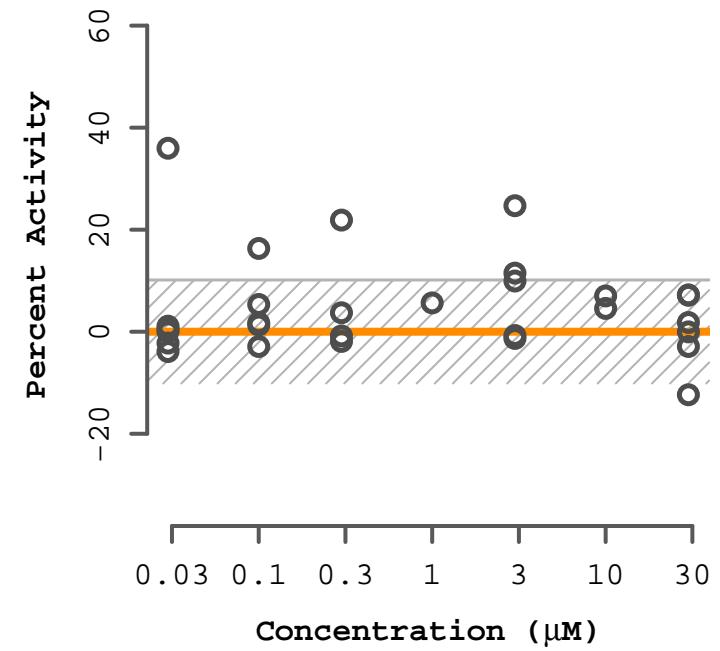
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	113.28	NA	NA
PROB:	1	NA	NA
RMSE:	4.43	NA	NA

MAX_MEAN: 5.46 MAX_MED: 1.27 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ethyltriacetoxy silane
CHID: 27792 CASRN: 17689-77-9
SPID(S): EPAPLT0154F03
M4ID: 30204206

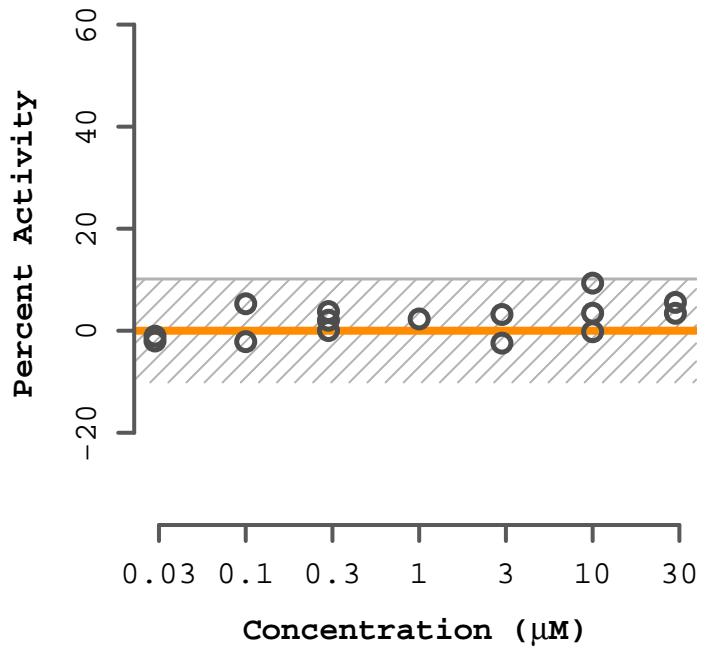
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 213.85	NA	NA
PROB: 1	NA	NA
RMSE: 10.72	NA	NA

MAX_MEAN: 8.82 MAX_MED: 9.95 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Calcium dodecylbenzene sulfonate

CHID: 27891 CASRN: 26264-06-2

SPID(S): EPAPLT0154D02

M4ID: 30204185

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 85.47

HILL

PROB: 1

GNLS

RMSE: 3.82

MAX_MEAN: 4.49

MAX_MED: 4.49

BMAD: 3.38

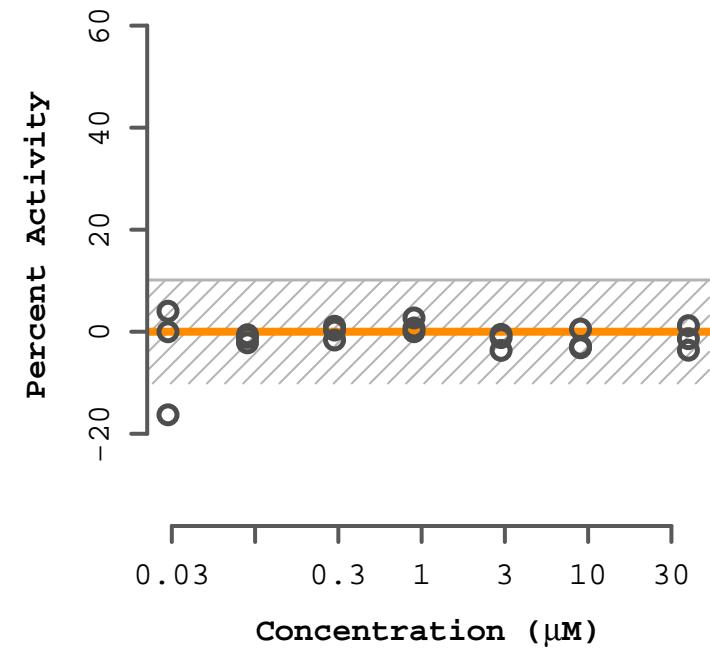
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dodecylbenzene sulfonate triethanolamine(1:1)

CHID: 27932 CASRN: 27323-41-7

SPID(S): TP0001413G01

M4ID: 30204601

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 107.11

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.08

NA

NA

MAX_MEAN: 1.15

MAX_MED: 0.741

BMAD: 3.38

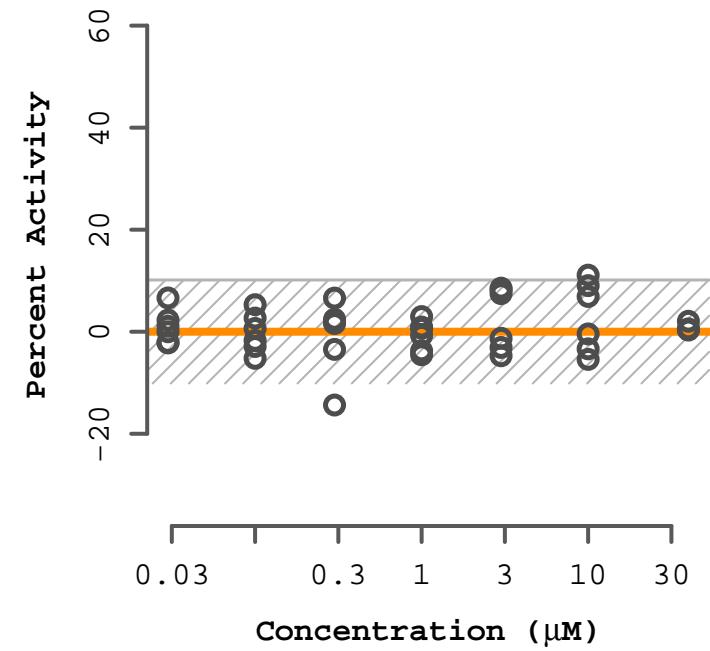
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Hexanedioic acid, diisononyl ester
CHID: 27982 CASRN: 33703-08-1
SPID(S): TP0001414C01
M4ID: 30204565

HILL MODEL: Not applicable.

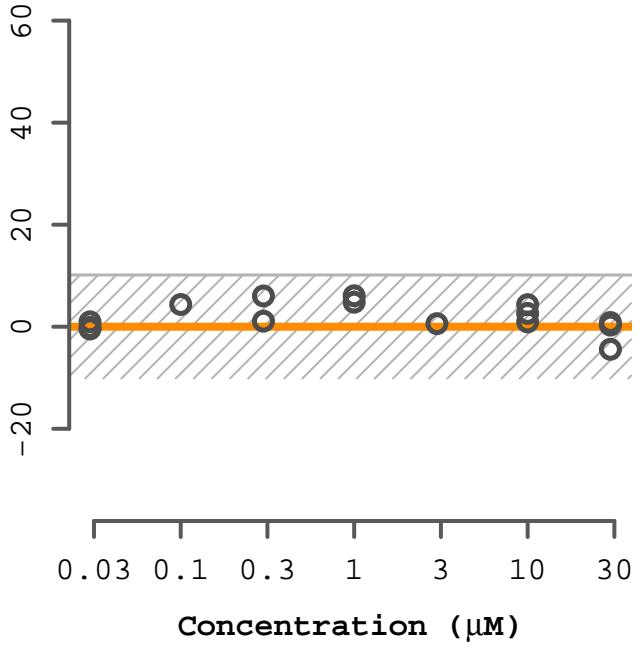
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	243.02	NA	NA
PROB:	1	NA	NA
RMSE:	4.95	NA	NA

MAX_MEAN: 2.98 MAX_MED: 3.25 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Potassium nitrate
CHID: 29692 CASRN: 7757-79-1
SPID(S): EPAPLT0154H07
M4ID: 30204230

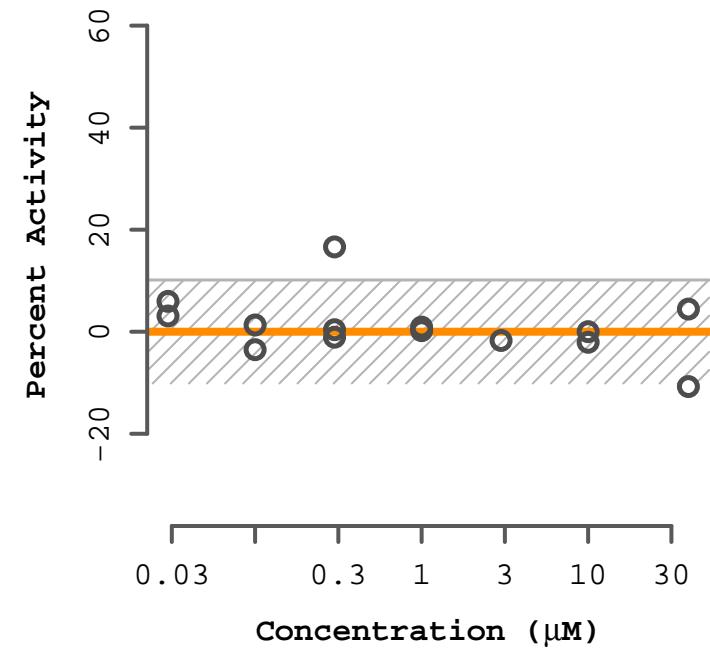
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 81.65	NA	NA
PROB: 1	NA	NA
RMSE: 3.32	NA	NA

MAX_MEAN: 5.45 MAX_MED: 5.45 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 8:2 Fluorotelomer alcohol
CHID: 29904 CASRN: 678-39-7
SPID(S): TP0001411E03
M4ID: 30204466

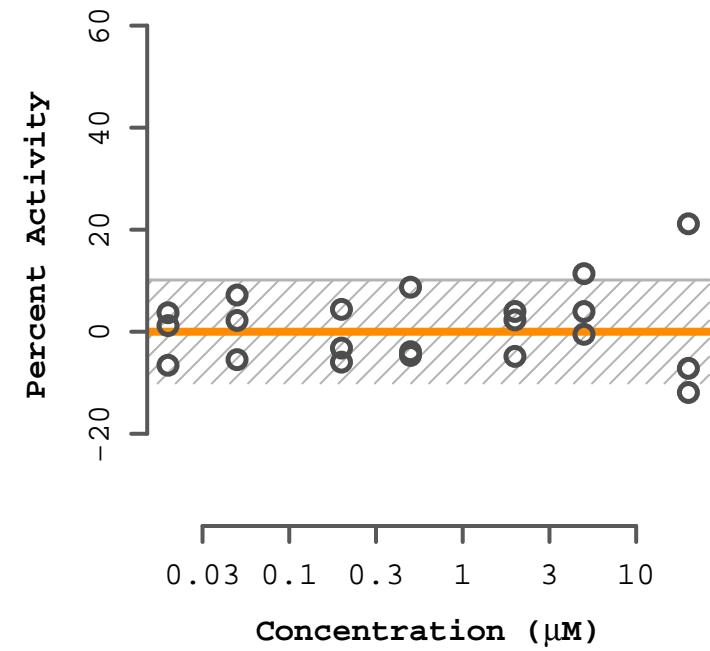
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	87.6	NA	NA
PROB:	1	NA	NA
RMSE:	5.86	NA	NA

MAX_MEAN: 5.32 MAX_MED: 4.52 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diquat dibromide monohydrate
CHID: 31248 CASRN: 6385-62-2
SPID(S): TP0001413E03
M4ID: 30204662

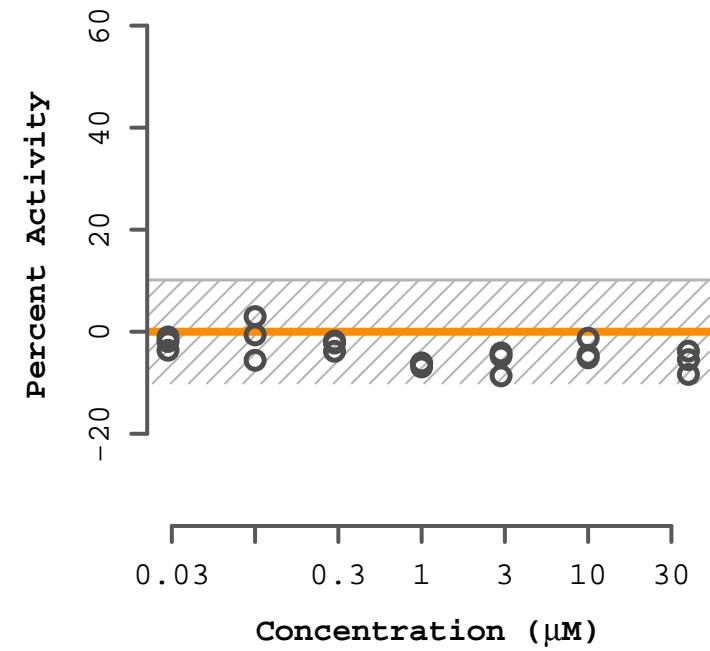
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 145.52	NA	NA
PROB: 1	NA	NA
RMSE: 7.41	NA	NA

MAX_MEAN: 4.95 MAX_MED: 3.96 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Perfluorohexanoic acid
CHID: 31862 CASRN: 307-24-4
SPID(S): TP0001412A09
M4ID: 30204461

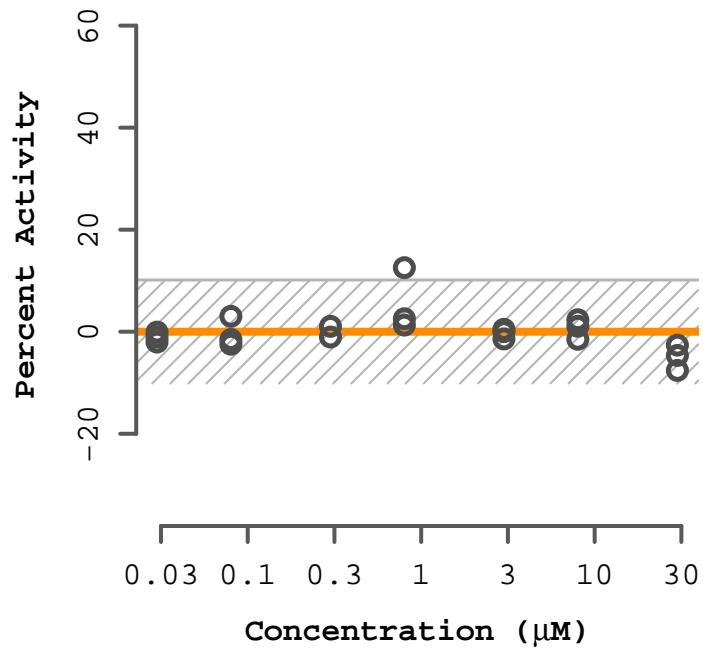
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 130.67	NA	NA
PROB: 1	NA	NA
RMSE: 4.79	NA	NA

MAX_MEAN: -1.08 MAX_MED: -0.614 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Perfluorooctanesulfonic acid
CHID: 31864 CASRN: 1763-23-1
SPID(S): TP0001412H11
M4ID: 30204330

HILL MODEL: Not applicable.

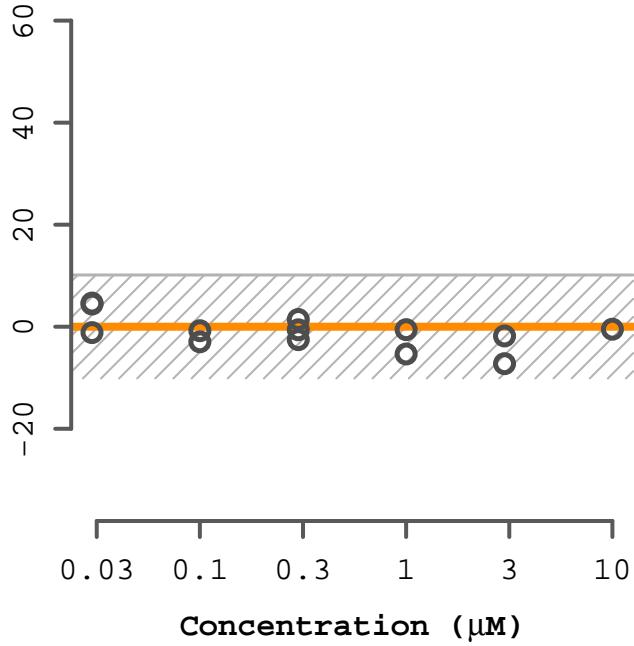
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 106.04	NA	NA
PROB: 1	NA	NA
RMSE: 3.79	NA	NA

MAX_MEAN: 5.47 MAX_MED: 2.55 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tebuconazole

CHID: 32113 CASRN: 107534-96-3

SPID(S): TP0001411F05

M4ID: 30204546

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 70.87

NA

NA

PROB:

1

NA

NA

RMSE: 3.34

NA

NA

MAX_MEAN: 2.65

MAX_MED: 4.45

BMAD: 3.38

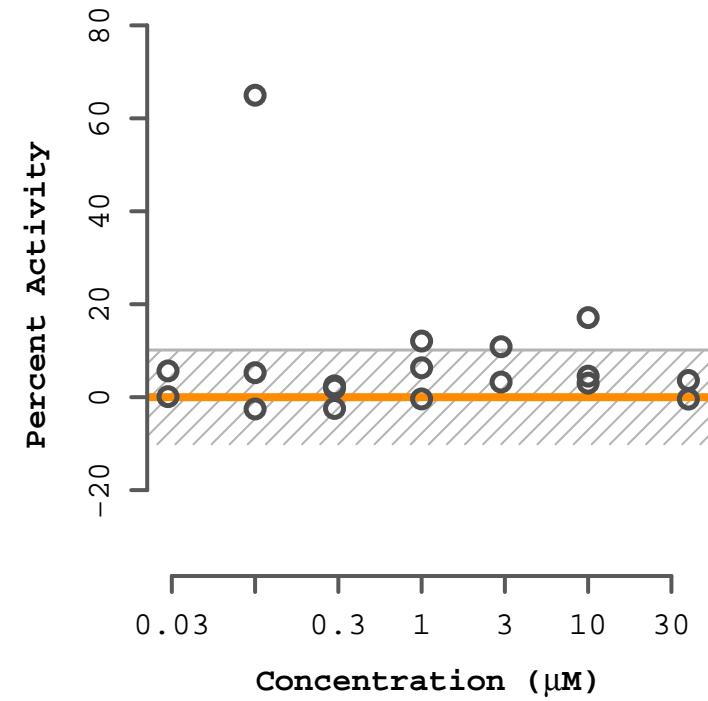
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Butralin

CHID: 32337 CASRN: 33629-47-9

SPID(S): TP0001412B04

M4ID: 30204438

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 136.56

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 16.59

NA

NA

MAX_MEAN: 22.6

MAX_MED: 7.06

BMAD: 3.38

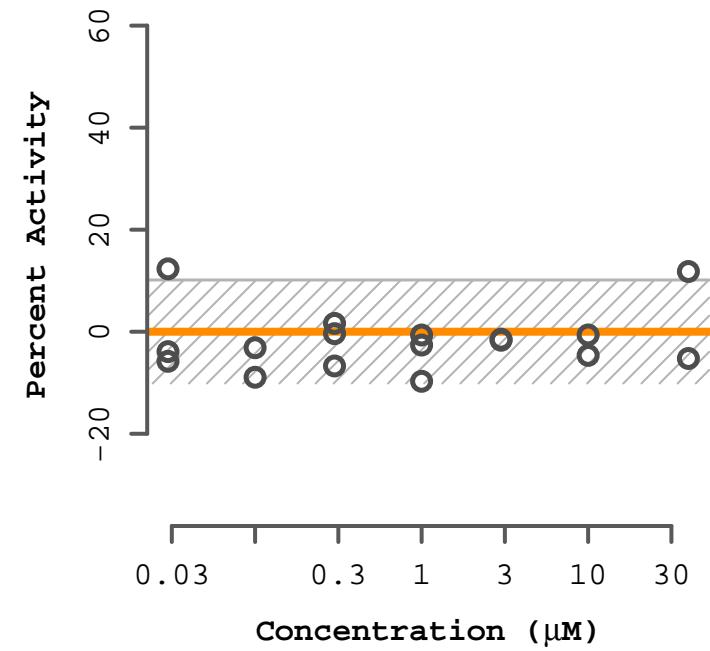
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Clodinafop-propargyl

CHID: 32354 CASRN: 105512-06-9

SPID(S): TP0001412B09

M4ID: 30204346

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 112.72

PROB: 1

RMSE: 6.1

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 3.29

MAX_MED: 3.29

BMAD: 3.38

COFF: 10.1

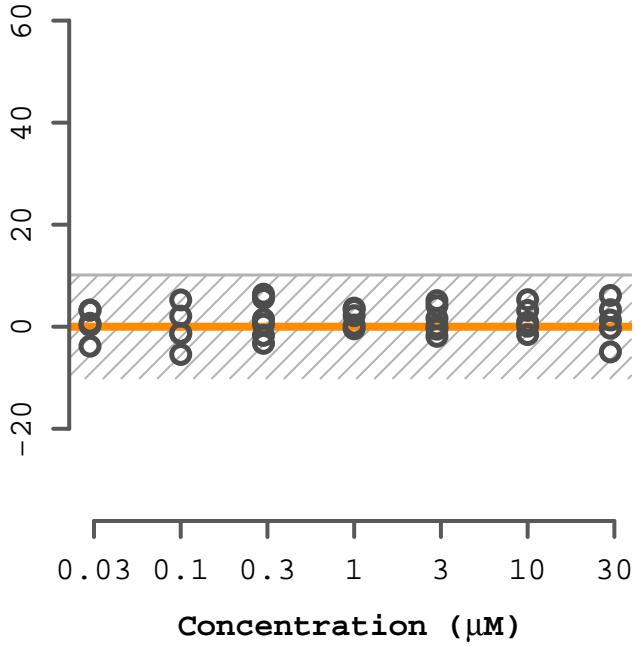
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cymoxanil

CHID: 32358 CASRN: 57966-95-7

SPID(S): EPAPLT0167D07

M4ID: 30204246

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 197.23

HILL

PROB: 1

NA

RMSE: 3.23

NA

GNLS

MAX_MEAN: 1.83

NA

NA

NA

BMAD: 3.38

COFF: 10.1

HIT-CALL: 0

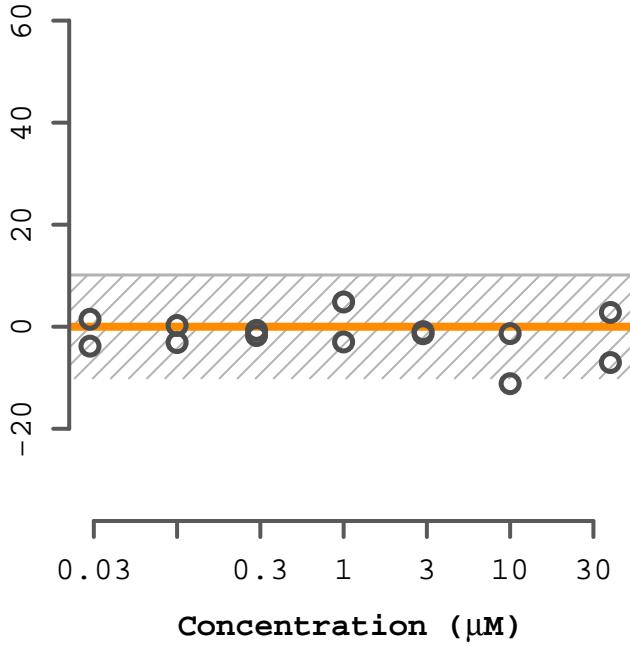
MAX_MED: 2.17

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ethalfluralin

CHID: 32386 CASRN: 55283-68-6

SPID(S): TP0001413G12

M4ID: 30204331

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 80.87

NA

NA

PROB:

1

NA

NA

RMSE: 4.21

NA

NA

MAX_MEAN: 0.938

MAX_MED: 0.938

BMAD: 3.38

COFF:

10.1

HIT-CALL:

0

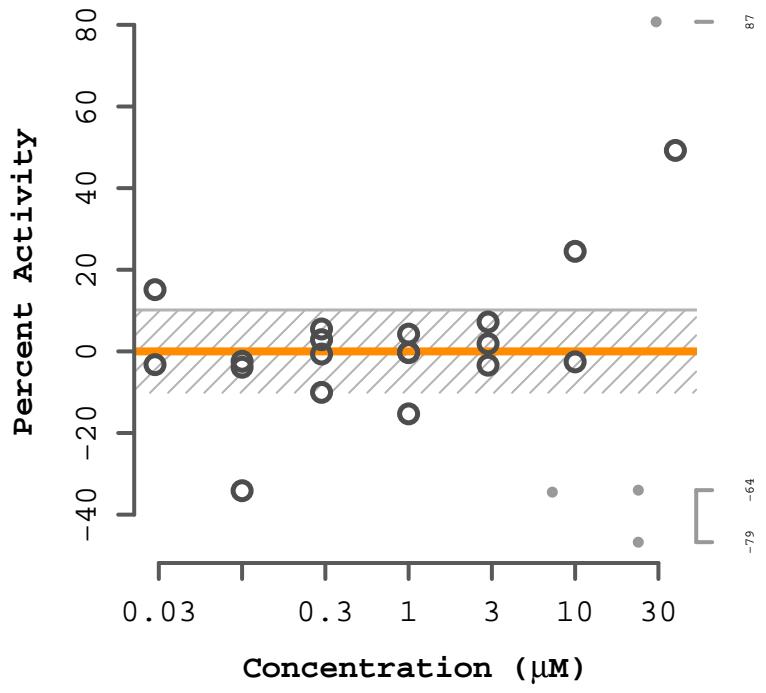
FITC:

4

ACTP:

0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fludioxonil

CHID: 32398 CASRN: 131341-86-1

SPID(S): TP0001411A04

M4ID: 30204362 BRK

HILL MODEL: Not applicable.

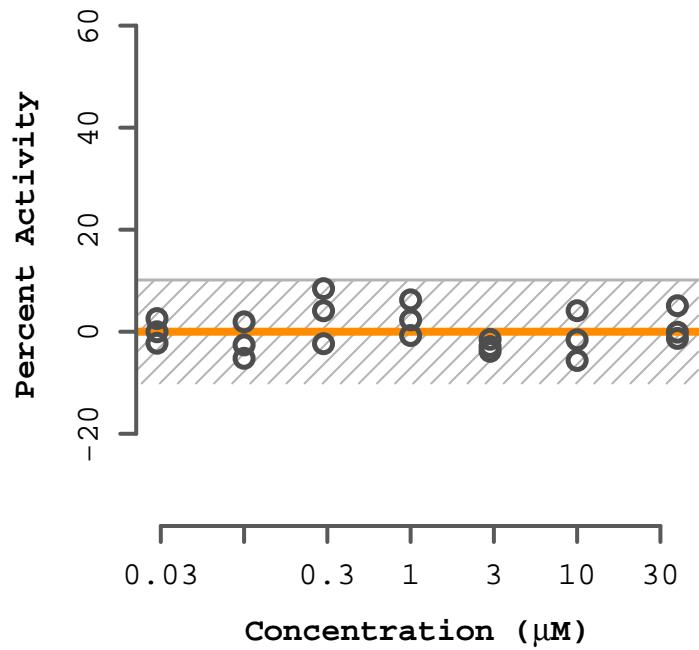
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 225.52	NA	NA
PROB: 1	NA	NA
RMSE: 34.34	NA	NA

MAX_MEAN: 5.94 MAX_MED: 5.94 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Imidacloprid

CHID: 32442 CASRN: 138261-41-3

SPID(S): TP0001413G08

M4ID: 30204306

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 118.87

NA

NA

PROB:

1

NA

NA

RMSE: 3.73

NA

NA

MAX_MEAN: 3.41

MAX_MED: 4.12

BMAD: 3.38

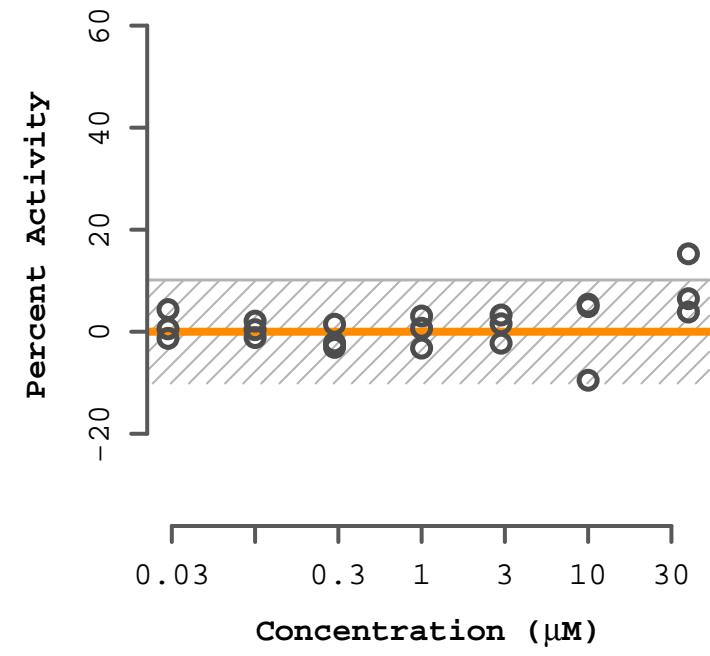
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Hexaflumuron

CHID: 32620 CASRN: 86479-06-3

SPID(S): TP0001413G03

M4ID: 30204478

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 126.15

NA

NA

PROB:

1

NA

NA

RMSE: 4.96

NA

NA

MAX_MEAN: 8.54

MAX_MED: 6.48

BMAD: 3.38

COFF: 10.1

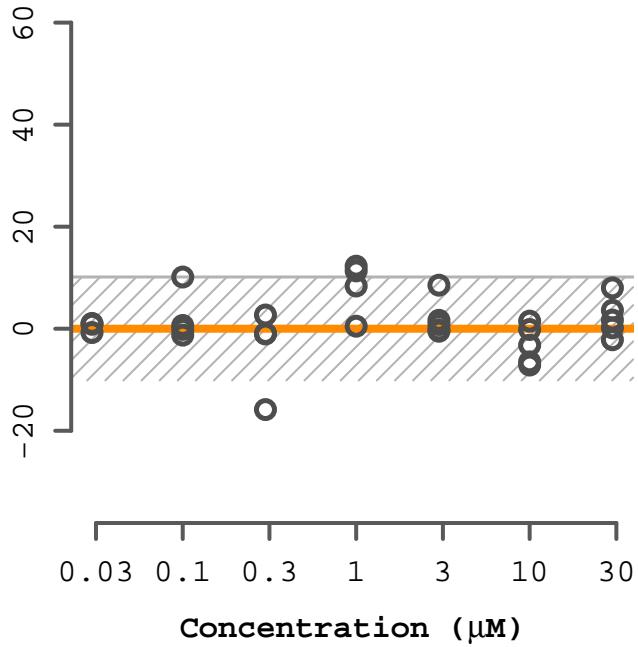
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pymetrozine

CHID: 32637 CASRN: 123312-89-0

SPID(S): EPAPLT0167F07

M4ID: 30204267

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 188.51

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 5.73

NA

NA

MAX_MEAN: 8.12

MAX_MED: 9.86

BMAD: 3.38

COFF: 10.1

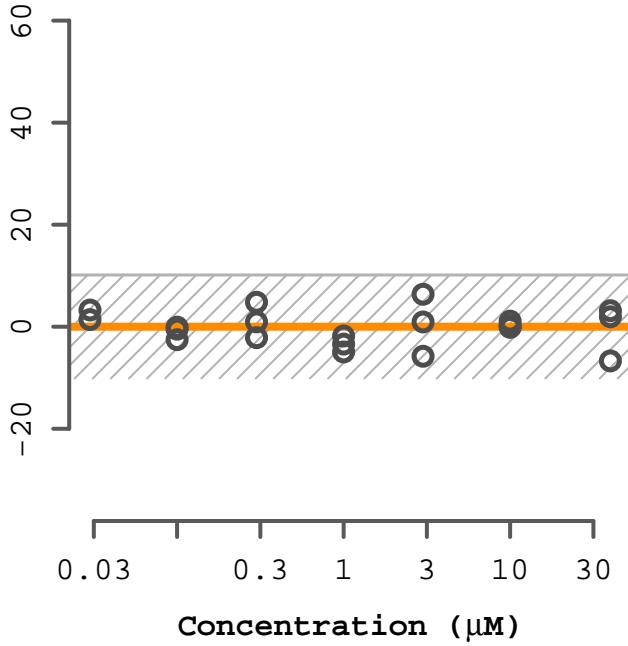
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pymetrozine

CHID: 32637 CASRN: 123312-89-0

SPID(S): TP0001412C02

M4ID: 30204336

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 108.18

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 3.33

NA

NA

MAX_MEAN: 2.34

MAX_MED: 2.34

BMAD: 3.38

COFF: 10.1

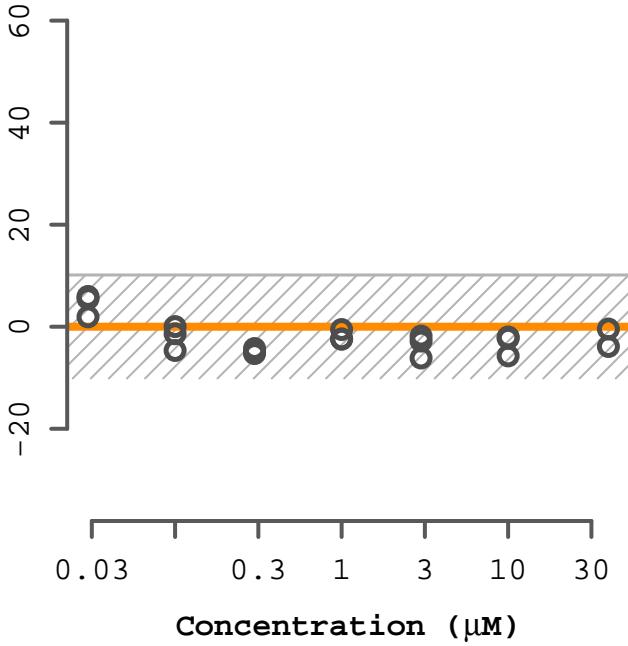
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Triticonazole

CHID: 32655 CASRN: 131983-72-7

SPID(S): TP0001414C07

M4ID: 30204580

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 114.55

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 3.75

NA

NA

MAX_MEAN: 4.47

MAX_MED: 5.52

BMAD: 3.38

COFF: 10.1

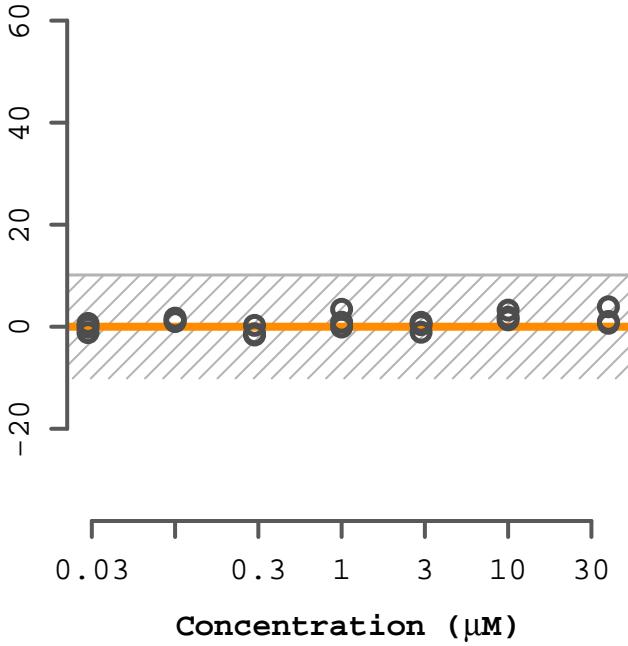
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pyrithiobac-sodium

CHID: 32673 CASRN: 123343-16-8

SPID(S): TP0001413F07

M4ID: 30204369

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 83.82

NA

NA

PROB: 1

NA

NA

RMSE: 1.68

NA

NA

MAX_MEAN: 2.17

MAX_MED: 1.88

BMAD: 3.38

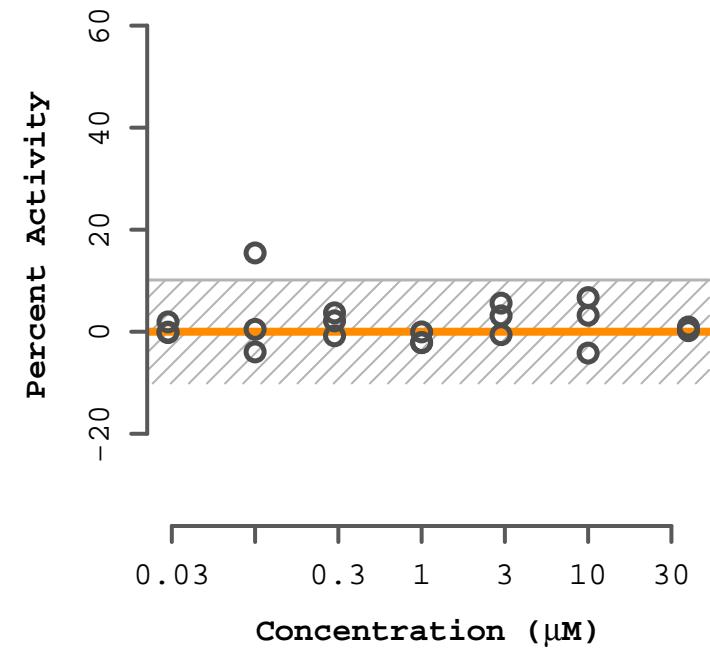
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Prodiamine

CHID: 34210 CASRN: 29091-21-2

SPID(S): TP0001414B09

M4ID: 30204364

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 112.88

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.48

NA

NA

MAX_MEAN: 3.99

MAX_MED: 3.23

BMAD: 3.38

COFF: 10.1

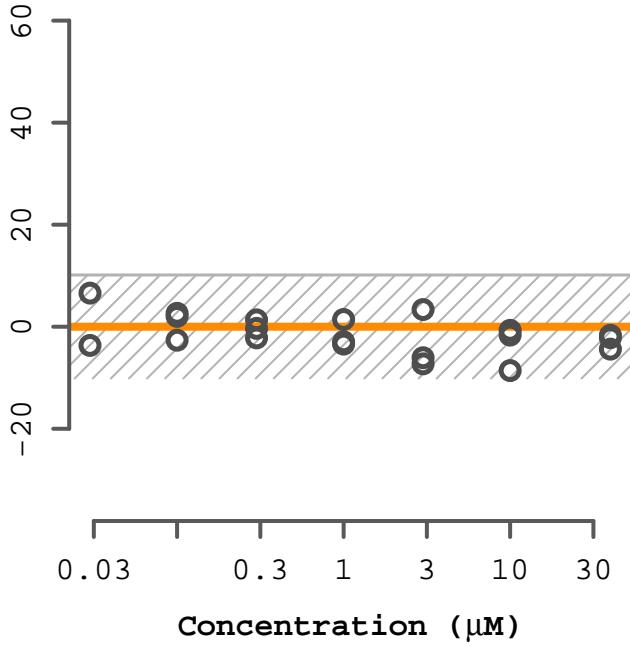
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Spiroxamine

CHID: 34212 CASRN: 118134-30-8

SPID(S): TP0001413G09

M4ID: 30204304

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 114.91

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 3.93

NA

NA

MAX_MEAN: 1.47

MAX_MED: 2.04

BMAD: 3.38

COFF: 10.1

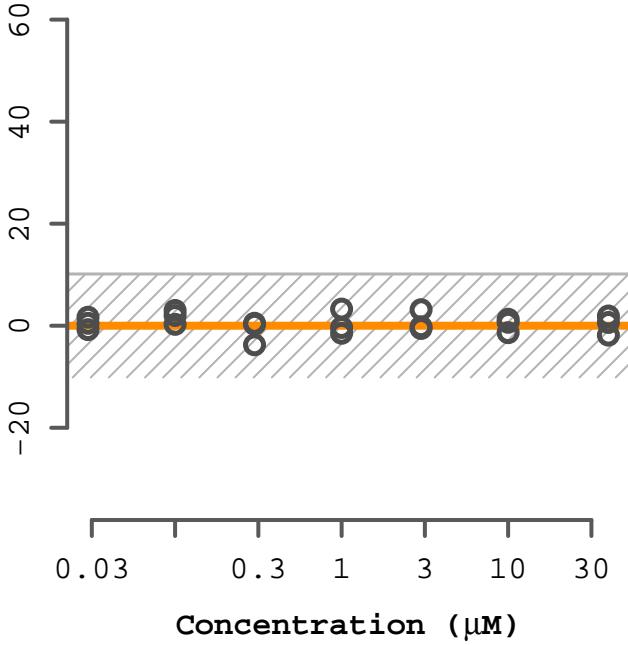
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Clopyralid-olamine

CHID: 34265 CASRN: 57754-85-5

SPID(S): TP0001413F09

M4ID: 30204419

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 86.7

PROB: 1

RMSE: 1.77

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 1.76

MAX_MED: 2.01

BMAD: 3.38

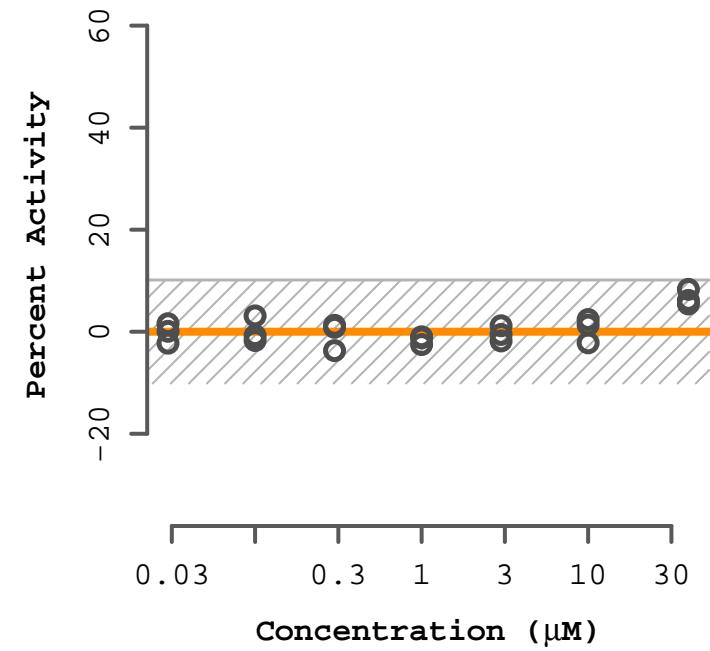
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 5-Heptyldihydro-2(3H)-furanone
CHID: 34287 CASRN: 104-67-6
SPID(S): TP0001413C08
M4ID: 30204396

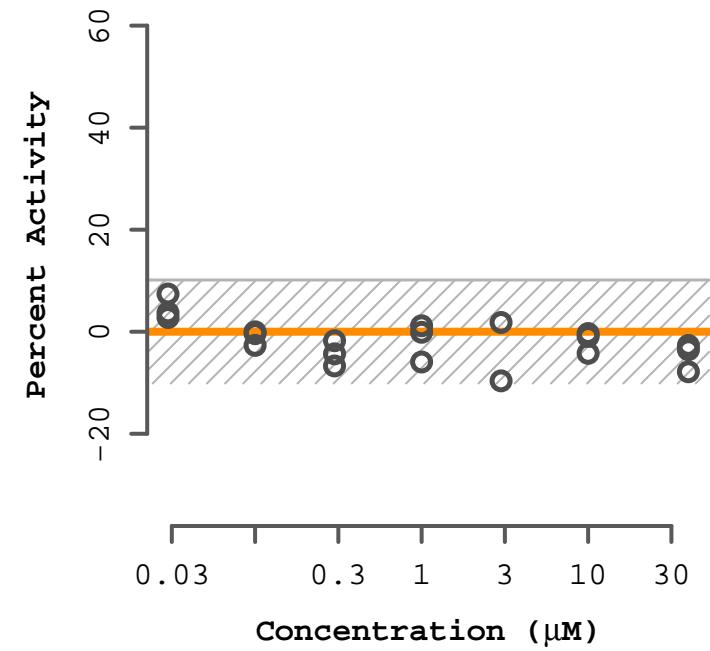
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	107.81	NA	NA
PROB:	1	NA	NA
RMSE:	3.08	NA	NA

MAX_MEAN: 6.64 MAX_MED: 6.15 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Acetamiprid

CHID: 34300 CASRN: 135410-20-7

SPID(S): TP0001411G02

M4ID: 30204319

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 118.92

NA

NA

PROB: 1

NA

NA

RMSE: 4.37

NA

NA

MAX_MEAN: 4.64

MAX_MED: 3.72

BMAD: 3.38

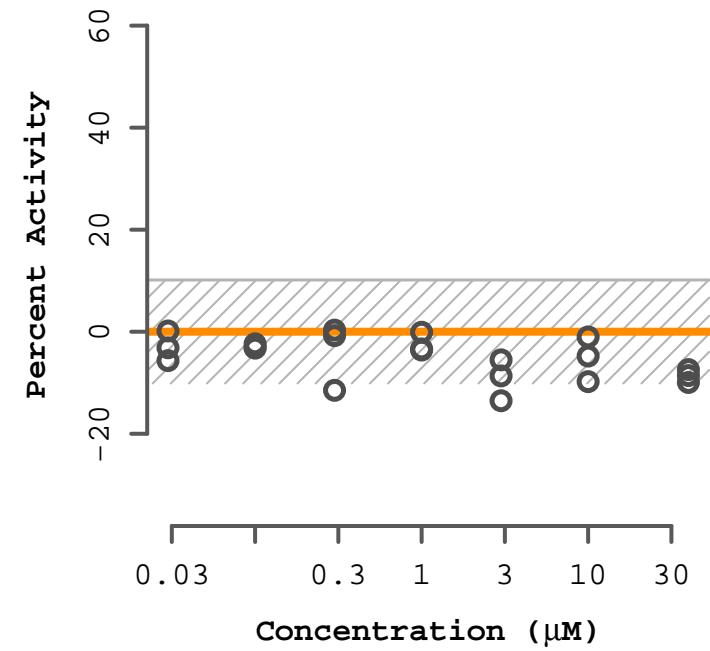
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Clothianidin

CHID: 34465 CASRN: 210880-92-5

SPID(S): TP0001412C05

M4ID: 30204322

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 141.21

PROB: 1

RMSE: 6.39

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: -2.35

MAX_MED: -0.752

BMAD: 3.38

COFF: 10.1

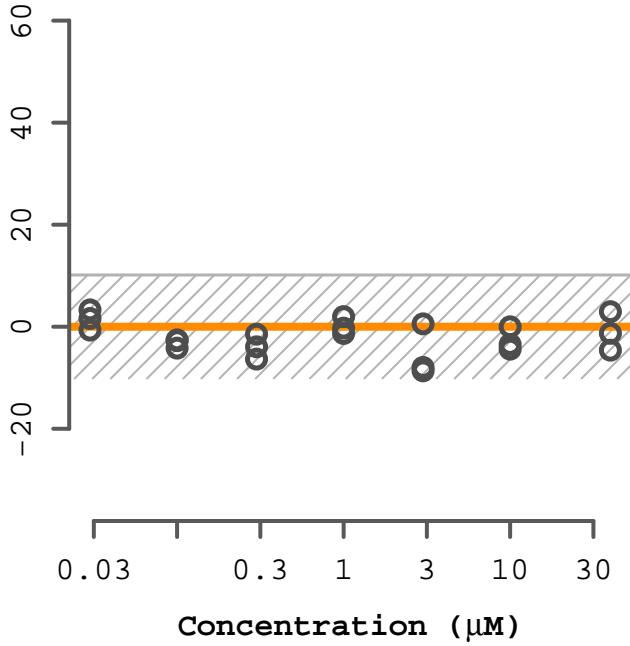
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cyanamide

CHID: 34490 CASRN: 420-04-2

SPID(S): TP0001413C03

M4ID: 30204449

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 119.27

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 3.84

NA

NA

MAX_MEAN: 1.43

MAX_MED: 1.62

BMAD: 3.38

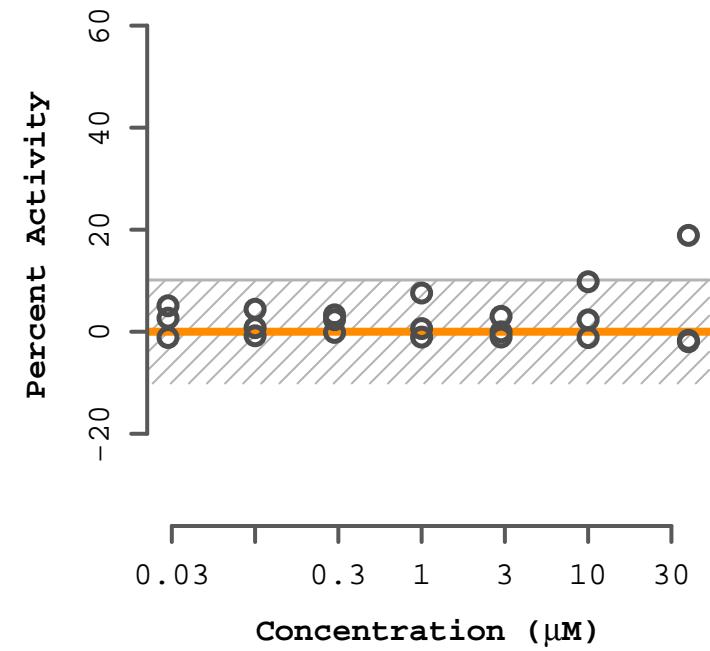
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cyazofamid

CHID: 34492 CASRN: 120116-88-3

SPID(S): TP0001413H10

M4ID: 30204422

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 124

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 5.38

NA

NA

MAX_MEAN: 5.09

MAX_MED: 2.67

BMAD: 3.38

COFF: 10.1

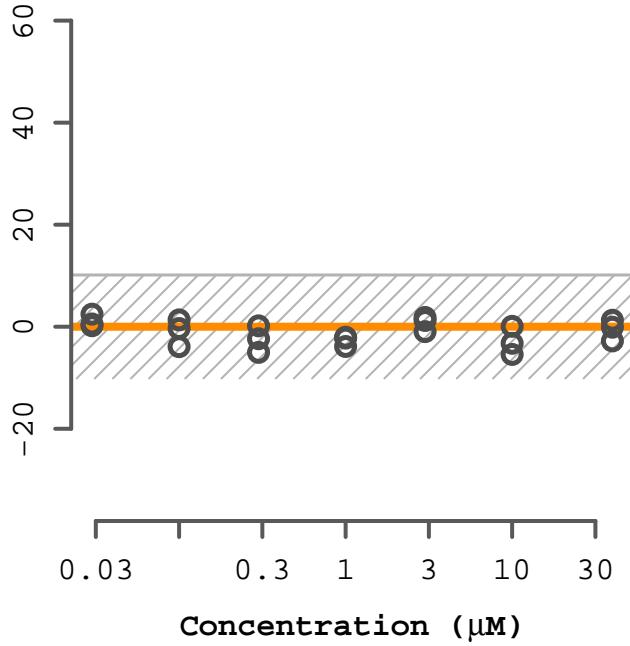
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fluroxypyrr

CHID: 34627 CASRN: 69377-81-7

SPID(S): TP0001414A11

M4ID: 30204307

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 101.79

HILL

PROB: 1

GNLS

RMSE: 2.52

MAX_MEAN: 1.07

MAX_MED: 1.23

BMAD: 3.38

COFF: 10.1

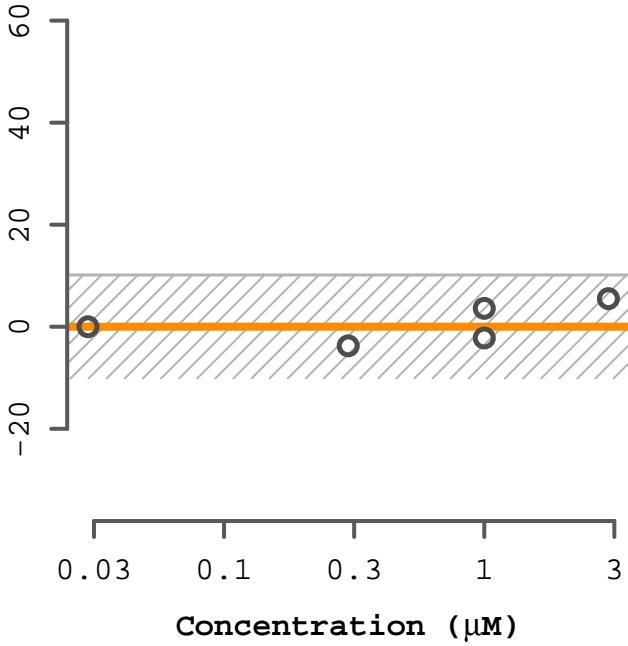
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Hexaconazole

CHID: 34653 CASRN: 79983-71-4

SPID(S): TP0001411H01

M4ID: 30204287

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 29.62

NA

NA

PROB:

1

NA

NA

RMSE: 3.53

NA

NA

MAX_MEAN: 5.51

MAX_MED: 5.51

BMAD: 3.38

COFF: 10.1

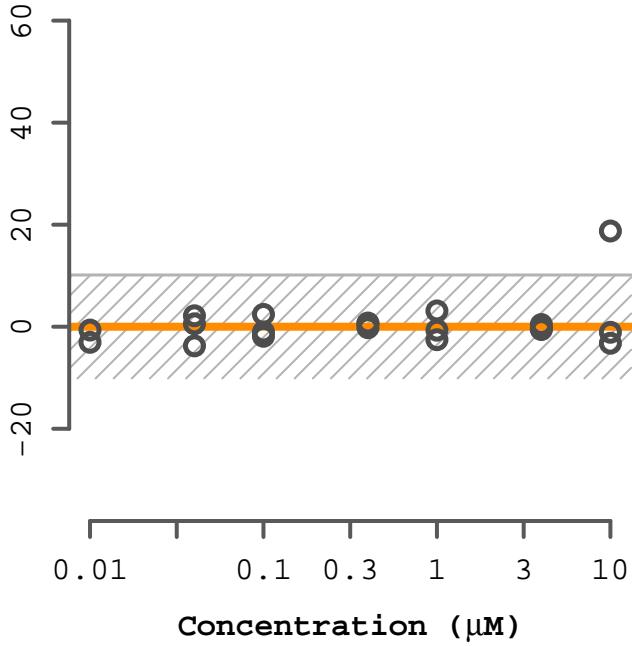
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Topramezone

CHID: 34722 CASRN: 210631-68-8

SPID(S): EPAPLT0167D10

M4ID: 30204274

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 101.92

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.59

NA

NA

MAX_MEAN: 4.82

MAX_MED: 0.592

BMAD: 3.38

COFF: 10.1

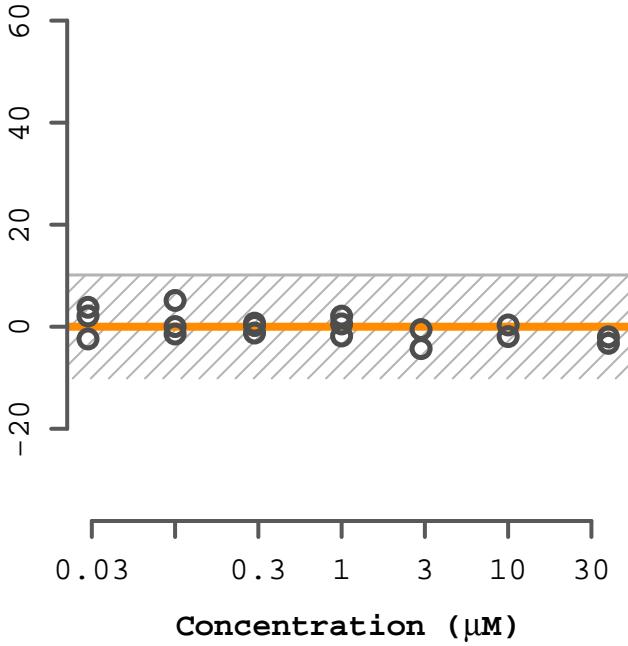
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Monocrotophos

CHID: 34816 CASRN: 6923-22-4

SPID(S): TP0001414H05

M4ID: 30204355

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 85.22

NA

NA

PROB:

1

NA

NA

RMSE: 2.36

NA

NA

MAX_MEAN: 1.26

MAX_MED: 2.13

BMAD: 3.38

COFF: 10.1

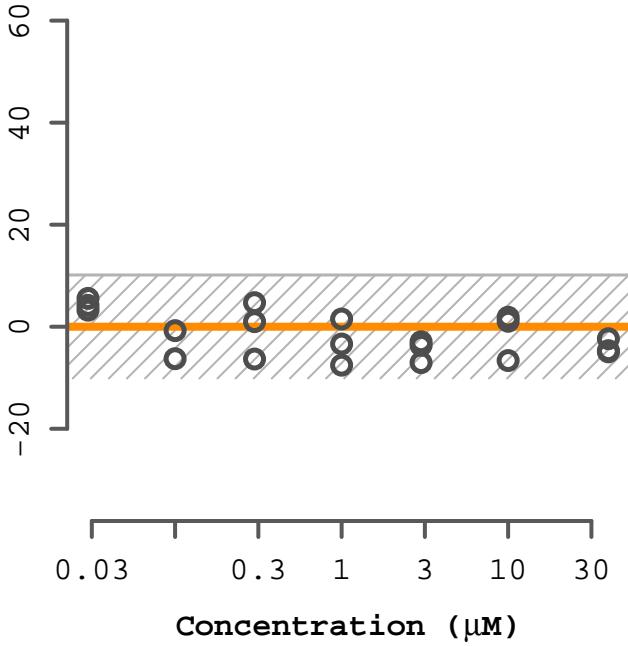
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Quinoxyfen

CHID: 34881 CASRN: 124495-18-7

SPID(S): TP0001412A06

M4ID: 30204283

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 122.08

PROB: 1

RMSE: 4.49

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 4.33

MAX_MED: 4.19

BMAD: 3.38

COFF: 10.1

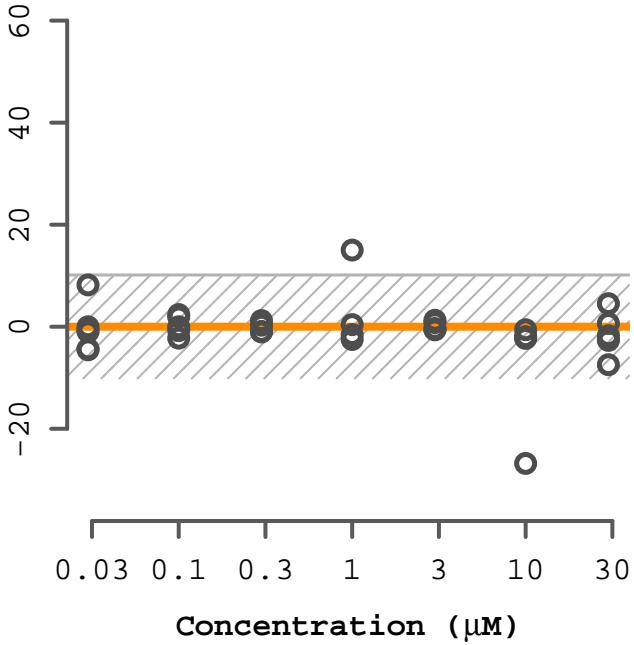
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Spirodiclofen

CHID: 34928 CASRN: 148477-71-8

SPID(S): EPAPLT0167D04

M4ID: 30204270

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 195.67

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 5.82

NA

NA

MAX_MEAN: 2.86

MAX_MED: 0

BMAD: 3.38

COFF: 10.1

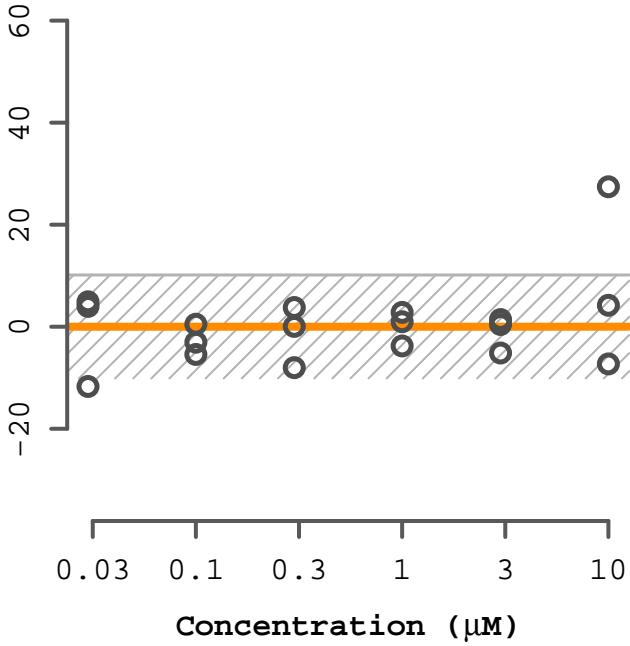
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tetriconazole

CHID: 34956 CASRN: 112281-77-3

SPID(S): TP0001414B07

M4ID: 30204368

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 122.29

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 8.06

NA

NA

MAX_MEAN: 8.12

MAX_MED: 4.21

BMAD: 3.38

COFF: 10.1

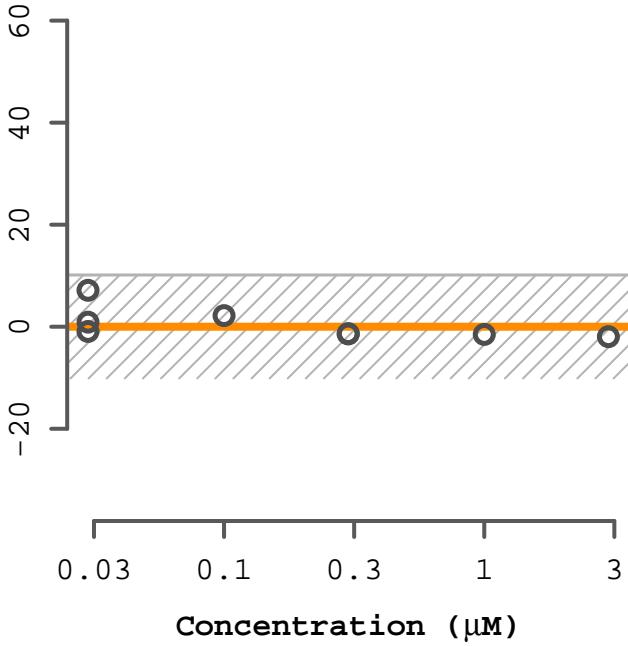
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Thiamethoxam

CHID: 34962 CASRN: 153719-23-4

SPID(S): TP0001411H03

M4ID: 30204303

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 40.59

HILL

PROB: 1

GNLS

RMSE: 2.9

MAX_MEAN: 2.37

MAX_MED: 2.19

BMAD: 3.38

COFF: 10.1

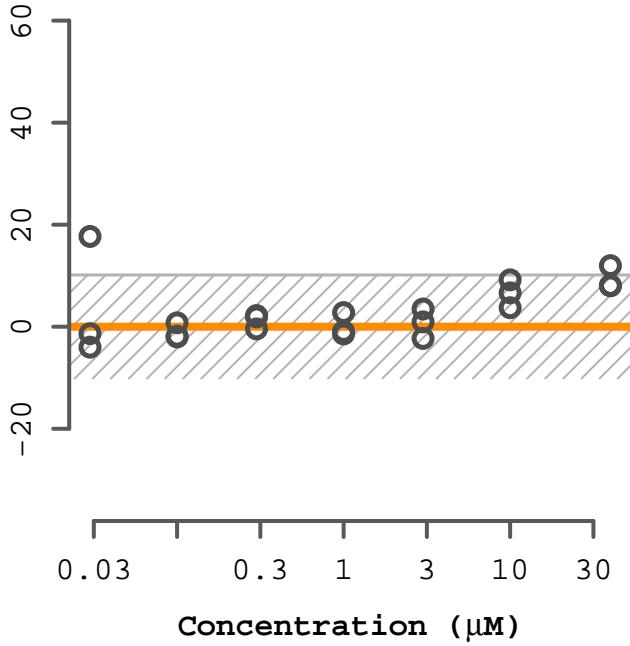
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Thymol

CHID: 34972 CASRN: 89-83-8

SPID(S): TP0001411H10

M4ID: 30204417

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 135.5

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 6.14

NA

NA

MAX_MEAN: 9.37

MAX_MED: 8.09

BMAD: 3.38

COFF: 10.1

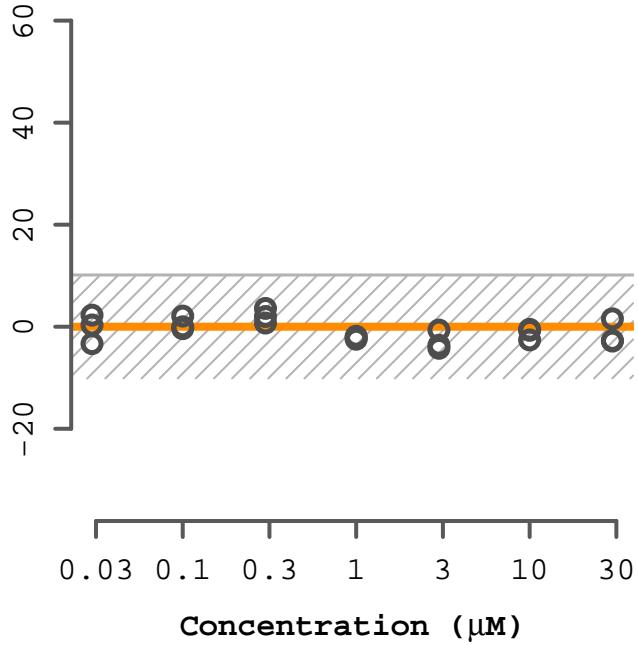
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Amoxicillin

CHID: 37044 CASRN: 26787-78-0

SPID(S): EPAPLT0169F09

M4ID: 30204235

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 99.06

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 2.27

NA

NA

MAX_MEAN: 2.11

MAX_MED: 2.01

BMAD: 3.38

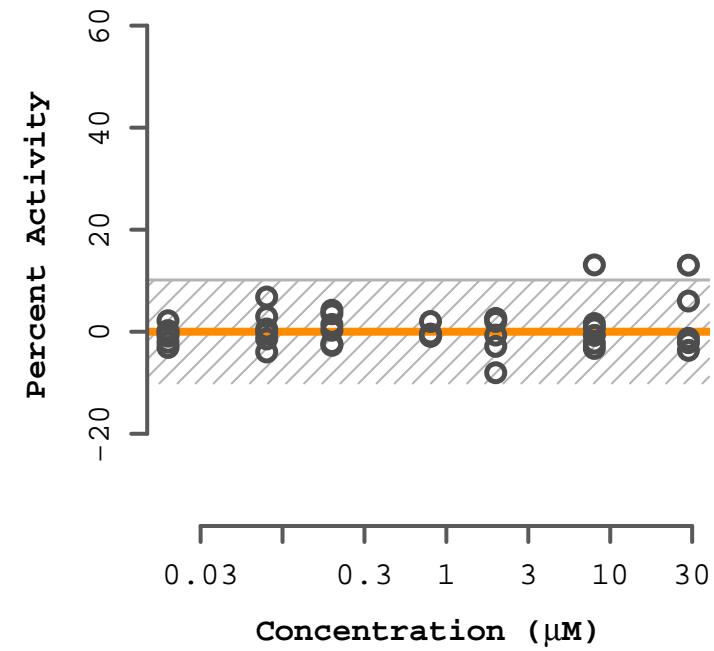
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Amiodarone hydrochloride
CHID: 37185 CASRN: 19774-82-4
SPID(S): TP0001413D07
M4ID: 30204465

HILL MODEL: Not applicable.

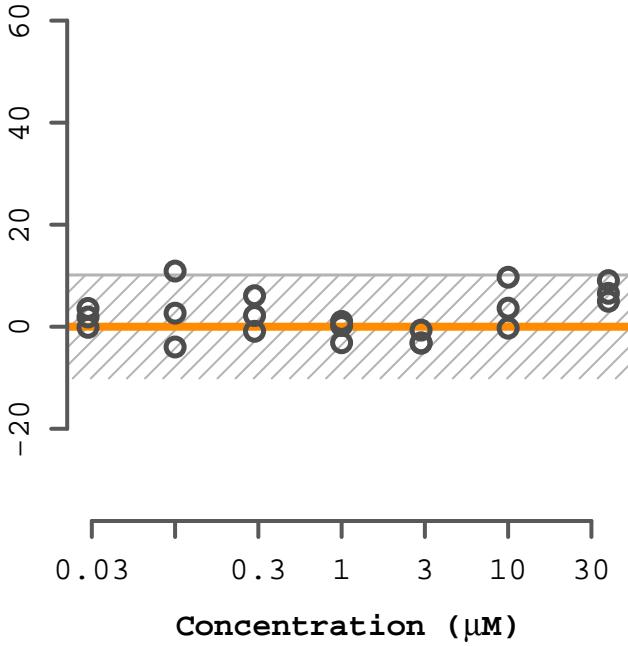
GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	221.83	NA	NA
PROB:	1	NA	NA
RMSE:	3.98	NA	NA

MAX_MEAN: 2.4 MAX_MED: 0.871 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Deisopropylatrazine

CHID: 37495 CASRN: 1007-28-9

SPID(S): TP0001414A05

M4ID: 30204299

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 128.31

PROB: 1

RMSE: 4.83

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 6.87

MAX_MED: 6.55

BMAD: 3.38

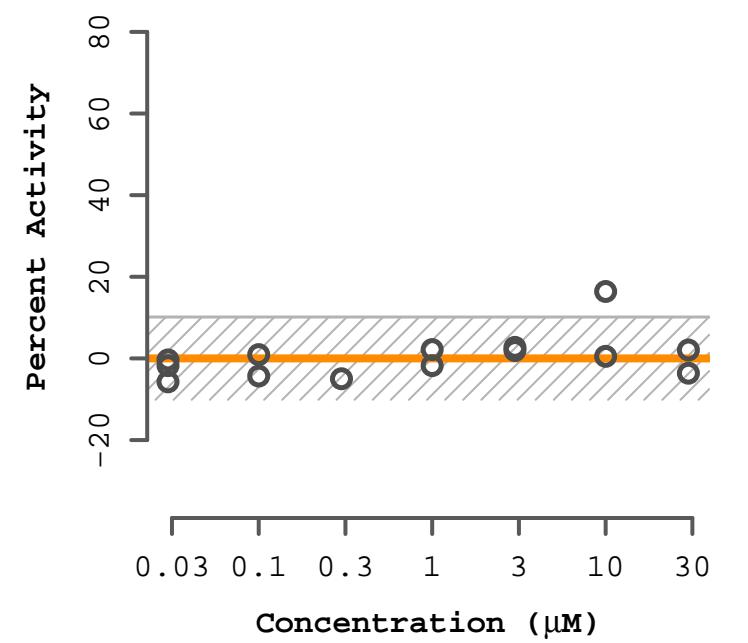
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Neopentyl glycol dibenzoate

CHID: 38822 CASRN: 4196-89-8

SPID(S): EPAPLT0154C10

M4ID: 30204182 BRK

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 113.98

PROB: 1

RMSE: 25.15

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 31.3

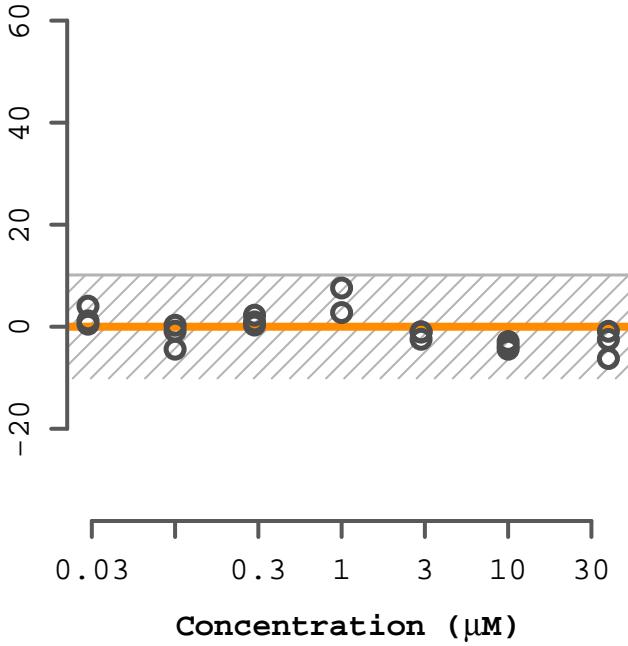
MAX_MED: 8.47

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Aldicarb

CHID: 39223 CASRN: 116-06-3

SPID(S): TP0001411E09

M4ID: 30204291

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 106.08

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 3.2

NA

NA

MAX_MEAN: 5.19

MAX_MED: 5.19

BMAD: 3.38

COFF: 10.1

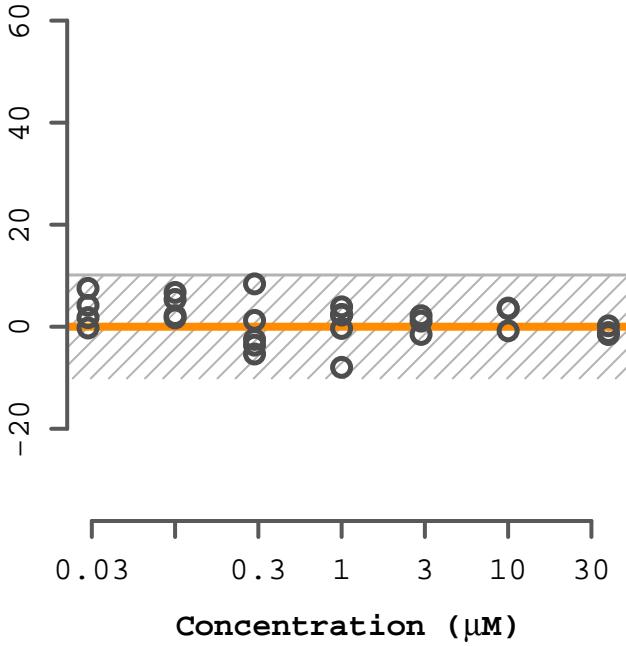
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Monomethyl phthalate
CHID: 40001 CASRN: 4376-18-5
SPID(S): TP0001413B10
M4ID: 30204333

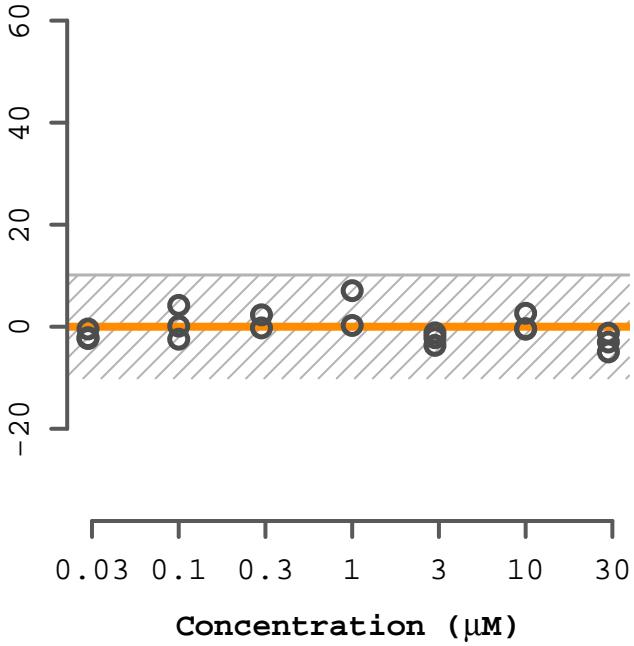
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 153.25	NA	NA
PROB: 1	NA	NA
RMSE: 3.88	NA	NA
MAX_MEAN: 4	MAX_MED: 3.77	BMAD: 3.38
COFF: 10.1	HIT-CALL: 0	FITC: 4
ACTP: 0		

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sucratose

CHID: 40245 CASRN: 56038-13-2

SPID(S): EPAPLT0154F08

M4ID: 30204211

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 95.79

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 2.82

NA

NA

MAX_MEAN: 3.67

MAX_MED: 3.67

BMAD: 3.38

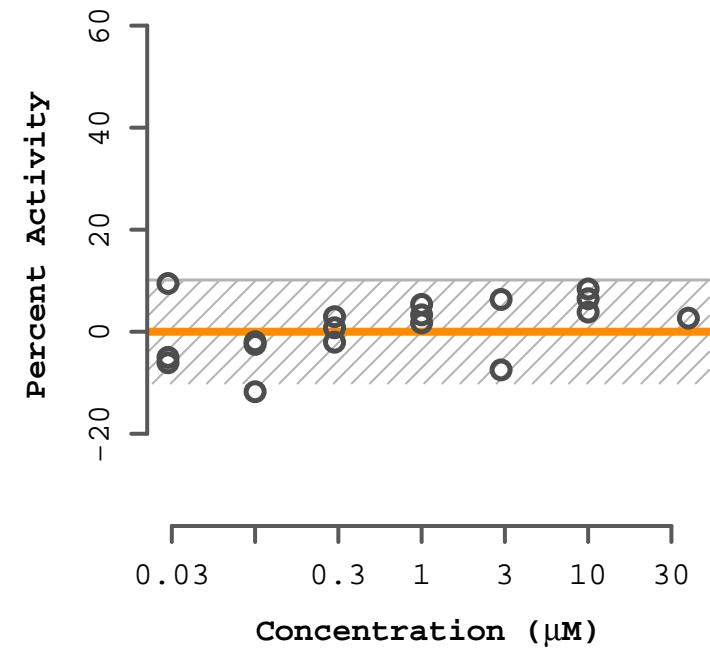
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Cyclohexylcyclohexanol
CHID: 40722 CASRN: 2433-14-9
SPID(S): TP0001413A07
M4ID: 30204476

HILL MODEL: Not applicable.

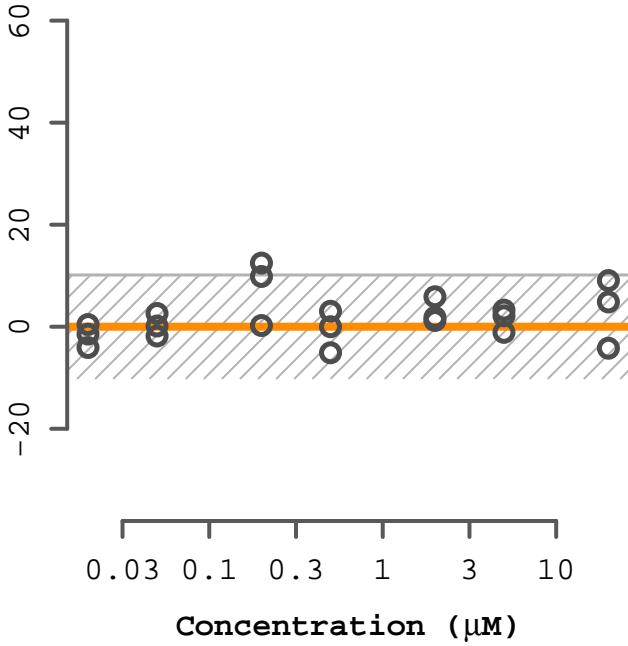
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 117.91	NA	NA
PROB: 1	NA	NA
RMSE: 5.71	NA	NA

MAX_MEAN: 6.24 MAX_MED: 6.48 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Ethylhexyl salicylate
CHID: 40734 CASRN: 118-60-5
SPID(S): EPAPLT0154G07
M4ID: 30204220

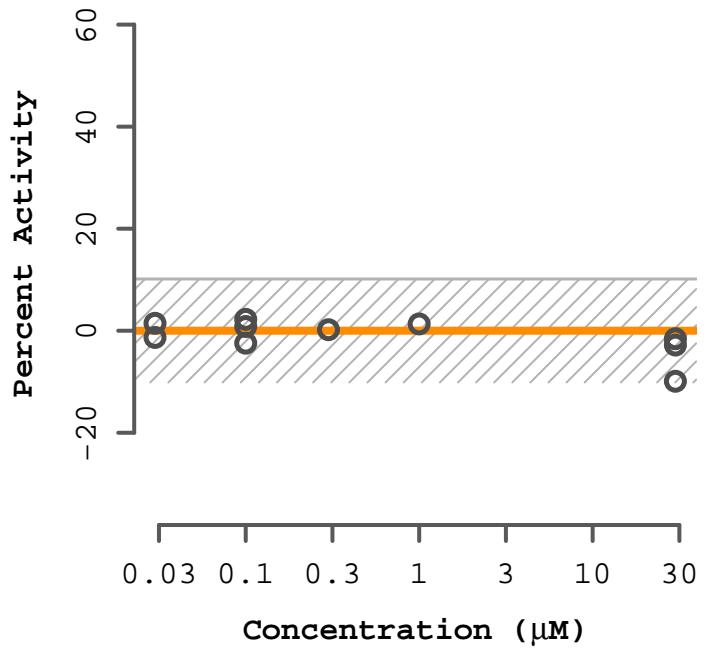
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 127.16	NA	NA
PROB: 1	NA	NA
RMSE: 4.86	NA	NA

MAX_MEAN: 7.53 MAX_MED: 9.85 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: N,N'-Disalicylidene-1,2-diaminopropane
CHID: 41200 CASRN: 94-91-7
SPID(S): EPAPLT0154C05
M4ID: 30204177

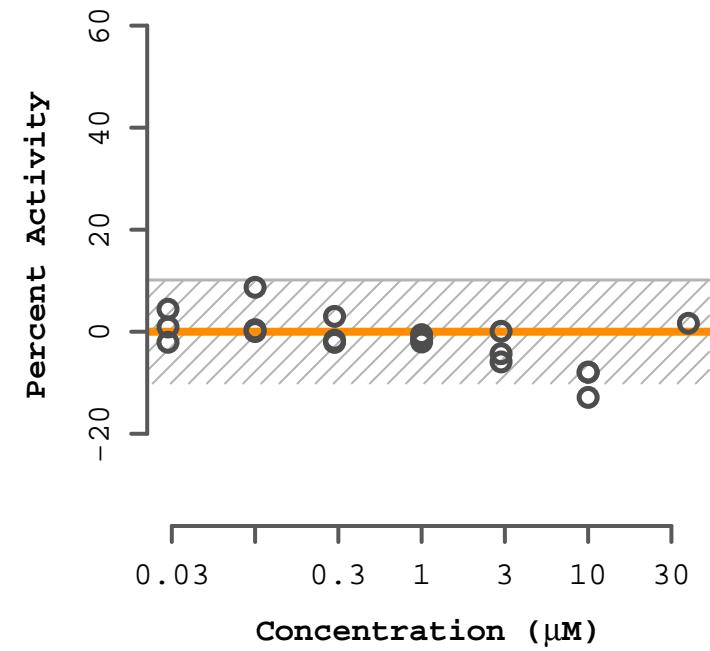
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 52.76	NA	NA
PROB: 1	NA	NA
RMSE: 3.54	NA	NA

MAX_MEAN: 1.34 MAX_MED: 1.34 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,5-Di-tert-butylbenzene-1,4-diol

CHID: 41248 CASRN: 88-58-4

SPID(S): TP0001413D08

M4ID: 30204451

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 116.08

PROB: 1

RMSE: 4.98

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 3.05

MAX_MED: 1.66

BMAD: 3.38

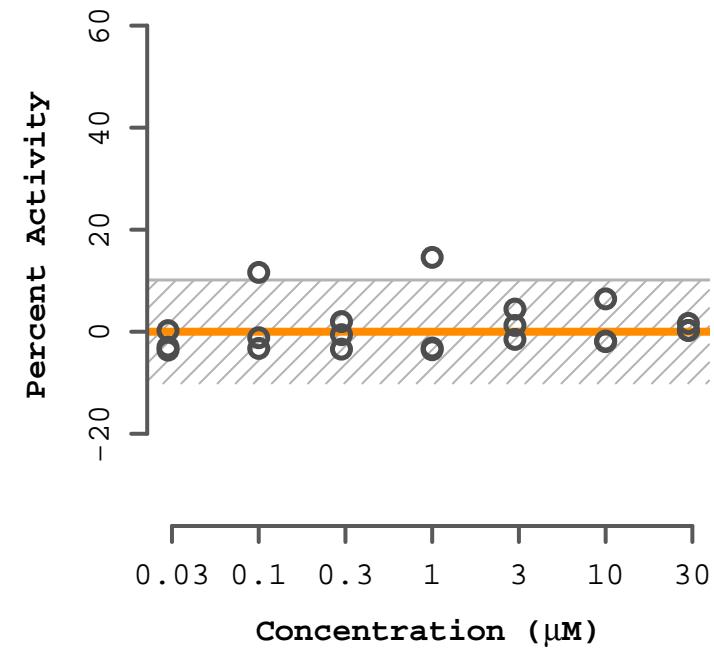
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 3,5,5-Trimethylhexyl acetate
CHID: 41270 CASRN: 58430-94-7
SPID(S): EPAPLT0154G01
M4ID: 30204214

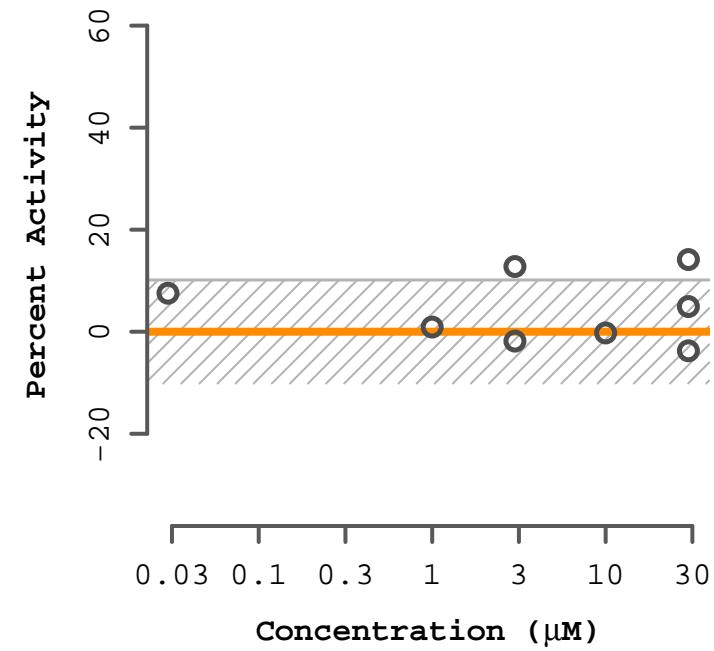
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 123.41	NA	NA
PROB: 1	NA	NA
RMSE: 4.86	NA	NA

MAX_MEAN: 2.62 MAX_MED: 1.24 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Hydroxy-2,2,6,6-tetramethylpiperidin-1-yl)ox:
CHID: 41280 CASRN: 2226-96-2
SPID(S): EPAPLT0154C11
M4ID: 30204183

HILL MODEL: Not applicable.

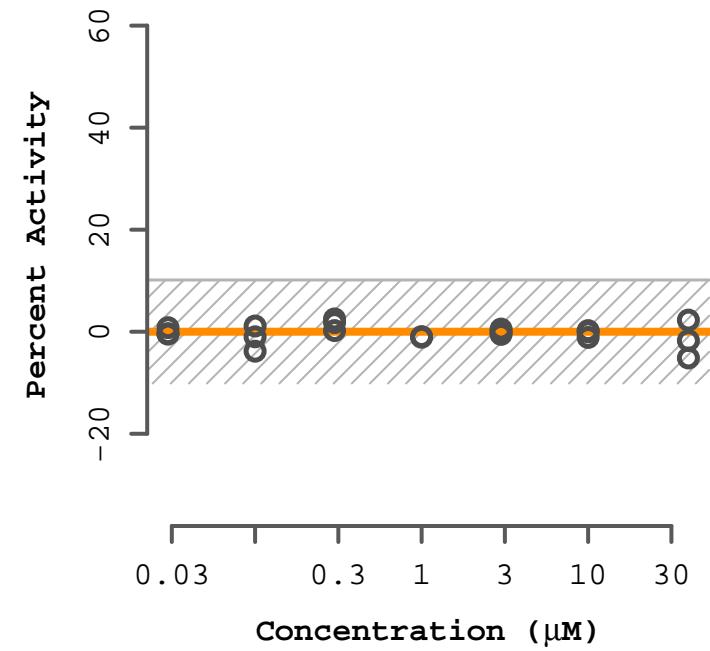
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 57.57	NA	NA
PROB: 1	NA	NA
RMSE: 7.61	NA	NA

MAX_MEAN: 7.56 MAX_MED: 7.56 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,2',4,4'-Tetrahydroxybenzophenone

CHID: 41306 CASRN: 131-55-5

SPID(S): TP0001413G05

M4ID: 30204401

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 80.33

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 1.81

NA

NA

MAX_MEAN: 1.52

MAX_MED: 1.81

BMAD: 3.38

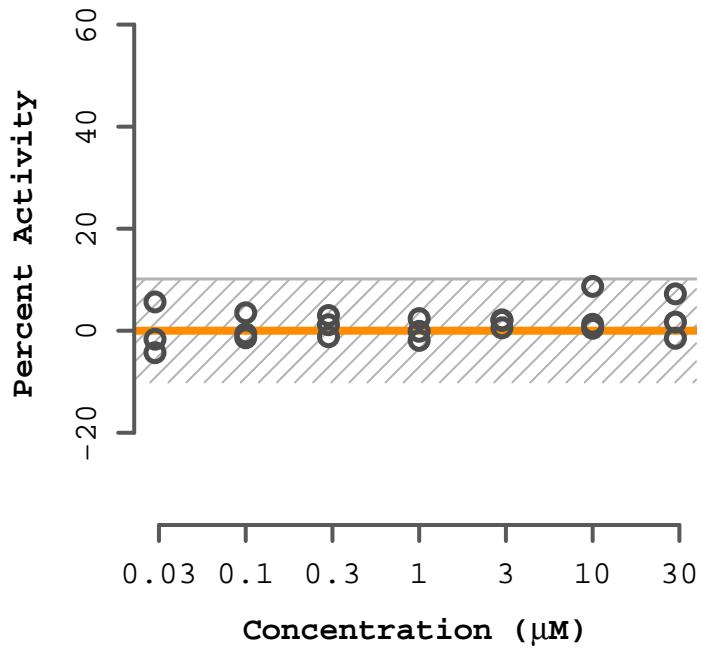
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,6-Dimethyl-2-heptanol
CHID: 41424 CASRN: 13254-34-7
SPID(S): EPAPLT0154C03
M4ID: 30204176

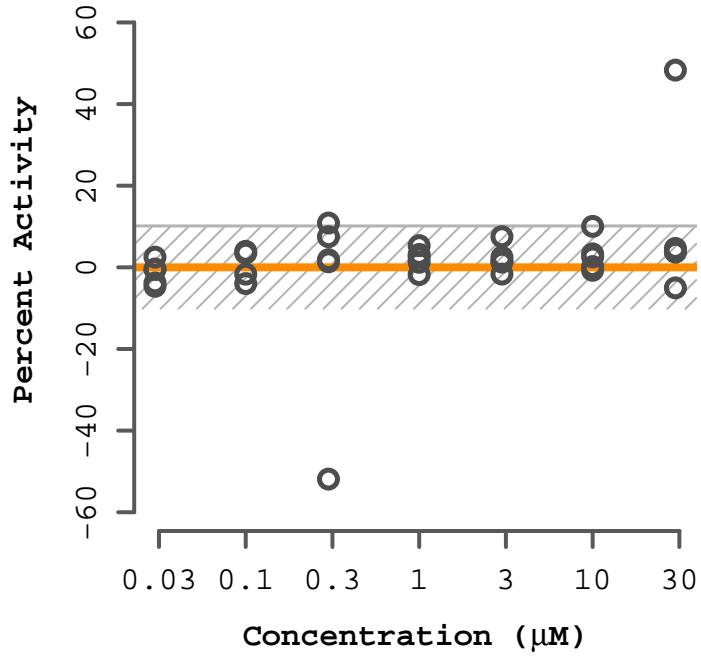
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 110.89	NA	NA
PROB: 1	NA	NA
RMSE: 3.33	NA	NA

MAX_MEAN: 3.48 MAX_MED: 2.03 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: C.I. Direct Red 81 disodium salt

CHID: 41726 CASRN: 2610-11-9

SPID(S): EPAPLT0154B06

M4ID: 30204168

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 233.44

PROB: 1

RMSE: 12.84

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 11.1

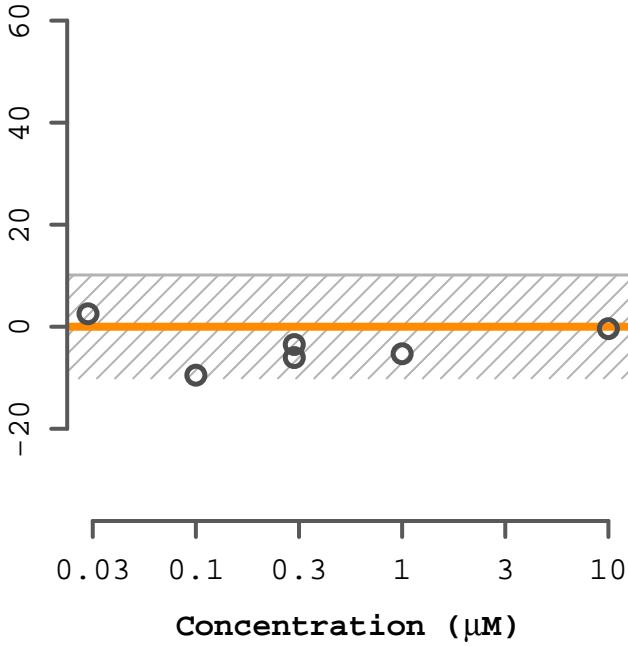
MAX_MED: 3.82

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1-Dodecyl-2-pyrrolidinone

CHID: 42206 CASRN: 2687-96-9

SPID(S): TP0001411G11

M4ID: 30204441

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 39.96

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 5.37

NA

NA

MAX_MEAN: 2.54

MAX_MED: 2.54

BMAD: 3.38

COFF: 10.1

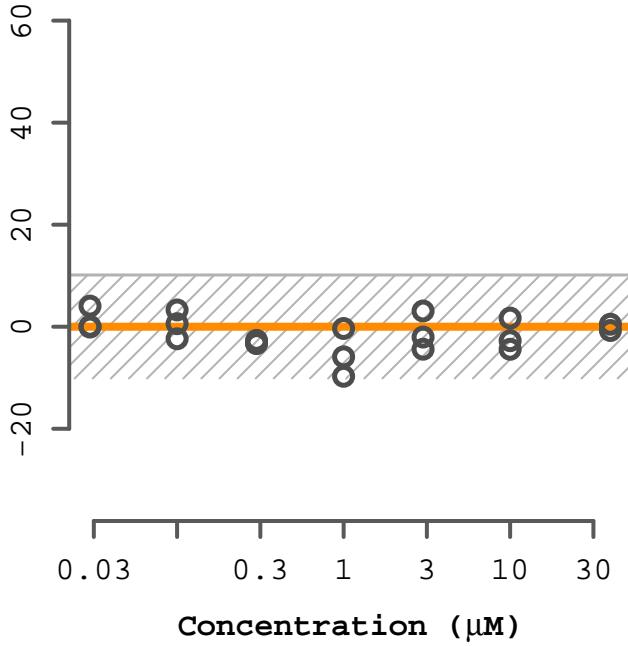
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Propylbenzene

CHID: 42219 CASRN: 103-65-1

SPID(S): TP0001413B04

M4ID: 30204590

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 109.83

HILL

PROB: 1

GNLS

RMSE: 3.56

MAX_MEAN: 1.36

MAX_MED: 0.602

BMAD: 3.38

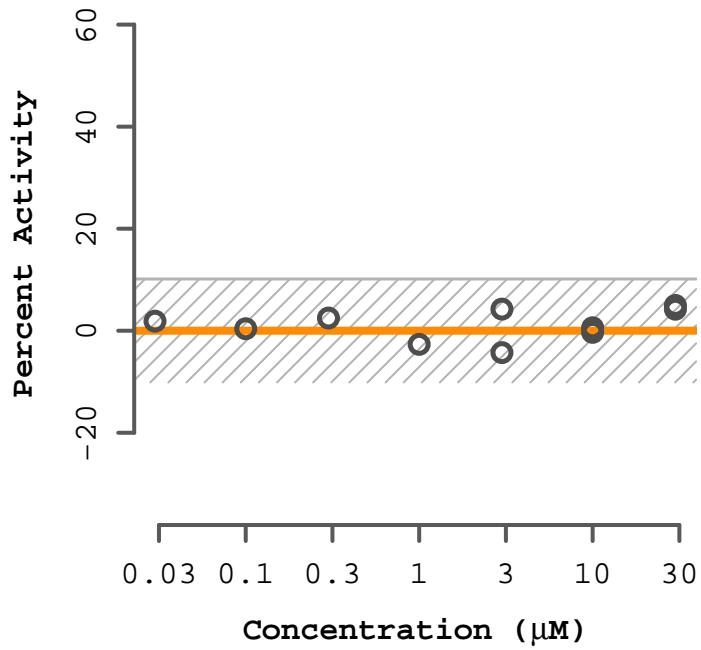
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tetrapropyl orthosilicate
CHID: 42382 CASRN: 682-01-9
SPID(S): EPAPLT0154B03
M4ID: 30204165

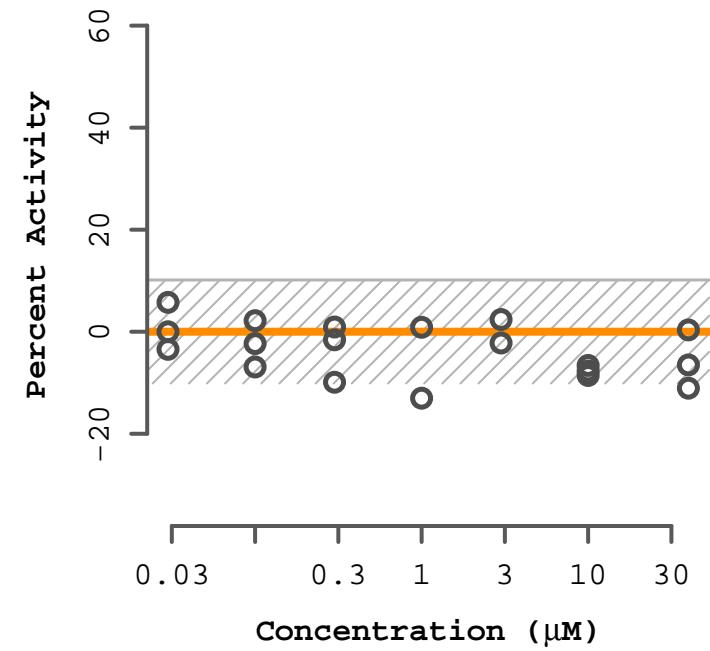
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 54.47	NA	NA
PROB: 1	NA	NA
RMSE: 3.1	NA	NA

MAX_MEAN: 4.59 MAX_MED: 4.59 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sodium (2-pyridylthio)-N-oxide
CHID: 42390 CASRN: 3811-73-2
SPID(S): TP0001411D02
M4ID: 30204436

HILL MODEL: Not applicable.

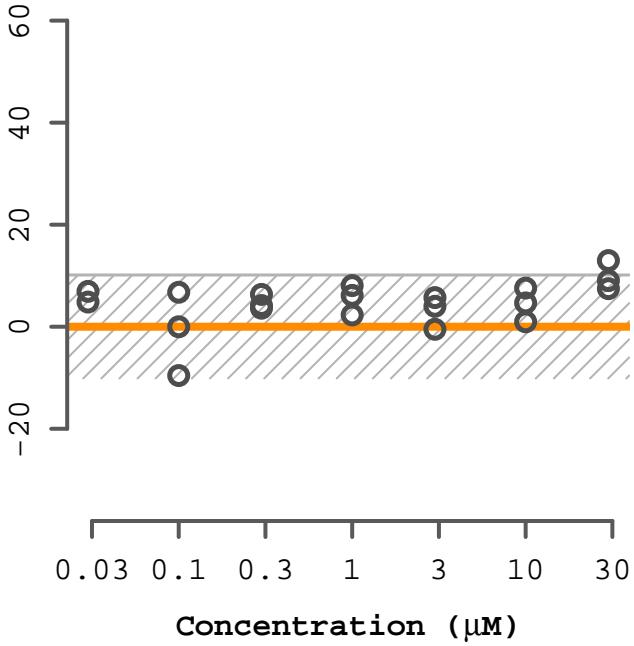
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 126.57	NA	NA
PROB: 1	NA	NA
RMSE: 6.18	NA	NA

MAX_MEAN: 0.773 MAX_MED: 0.0973 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Zoalene

CHID: 42521 CASRN: 148-01-6

SPID(S): EPAPLT0154D09

M4ID: 30204192

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 136.12

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 6.42

NA

NA

MAX_MEAN: 9.83

MAX_MED: 9.04

BMAD: 3.38

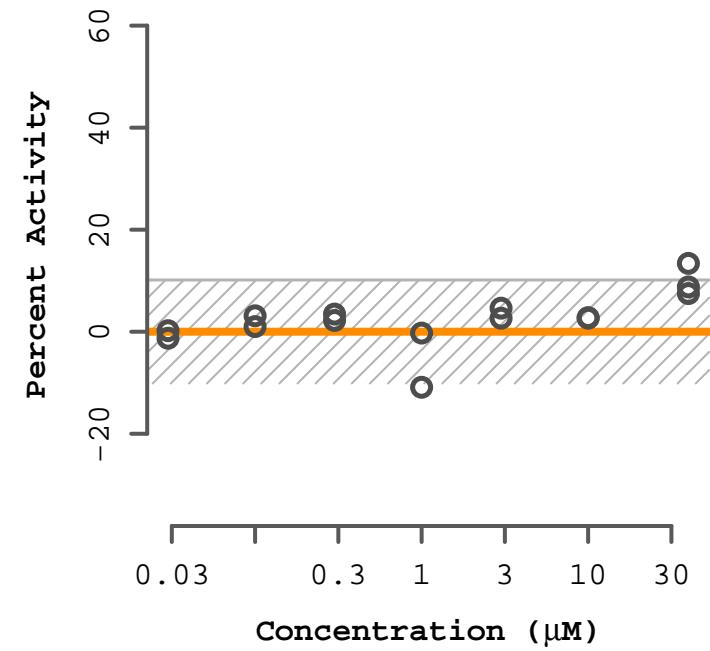
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pioglitazone hydrochloride
CHID: 44203 CASRN: 112529-15-4
SPID(S): TP0001413A01
M4ID: 30204597

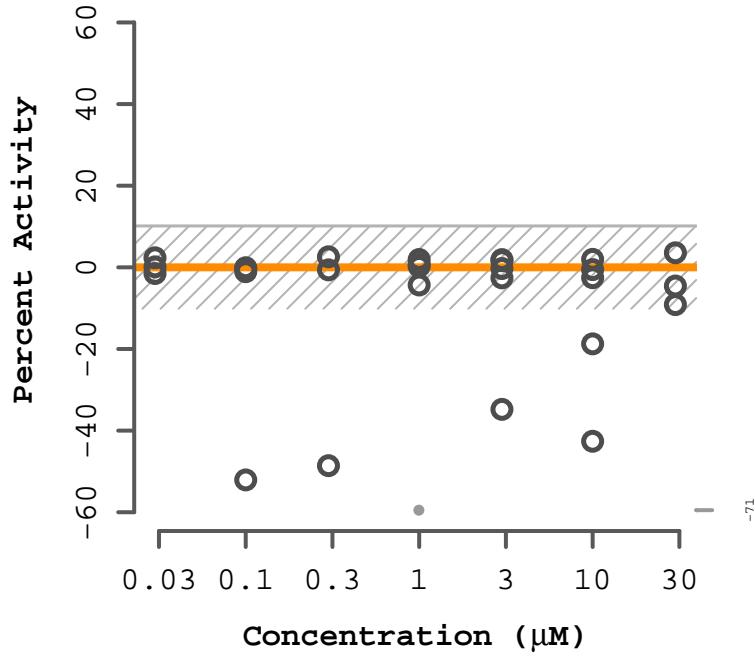
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 97.06	NA	NA
PROB: 1	NA	NA
RMSE: 5.8	NA	NA

MAX_MEAN: 9.89 MAX_MED: 8.79 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: p-Methylacetophenone
CHID: 44374 CASRN: 122-00-9
SPID(S): EPAPLT0154B07
M4ID: 30204169 BRK

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

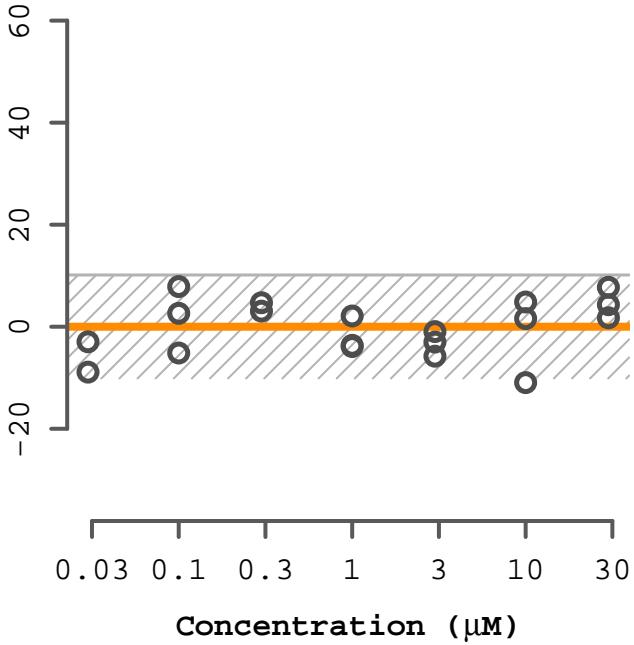
CNST	HILL	GNLS
AIC: 239.07	NA	NA
PROB: 1	NA	NA
RMSE: 22.05	NA	NA

MAX_MEAN: 0.326 MAX_MED: 0.245 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ethenyl(triethoxy)silane
CHID: 44463 CASRN: 78-08-0
SPID(S): EPAPLT0154G08
M4ID: 30204221

HILL MODEL: Not applicable.

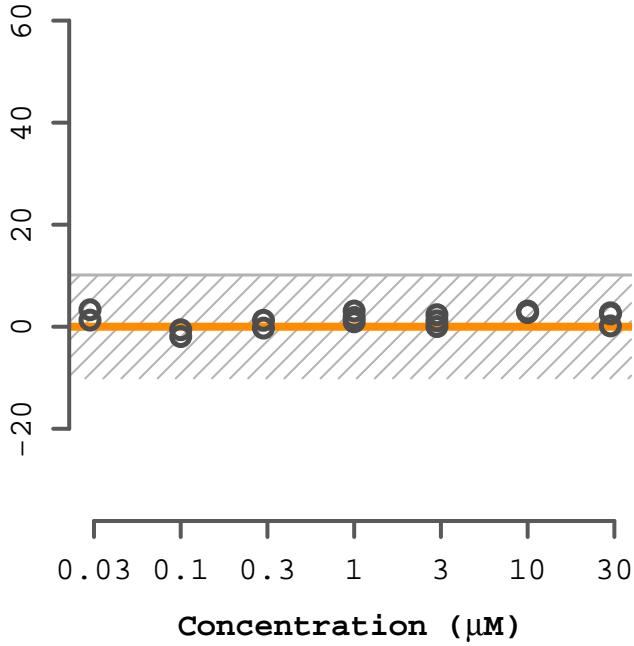
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 120.9	NA	NA
PROB: 1	NA	NA
RMSE: 5.22	NA	NA

MAX_MEAN: 4.61 MAX_MED: 4.33 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Phenylethyl phenylacetate
CHID: 44494 CASRN: 102-20-5
SPID(S): EPAPLT0154H05
M4ID: 30204228

HILL MODEL: Not applicable.

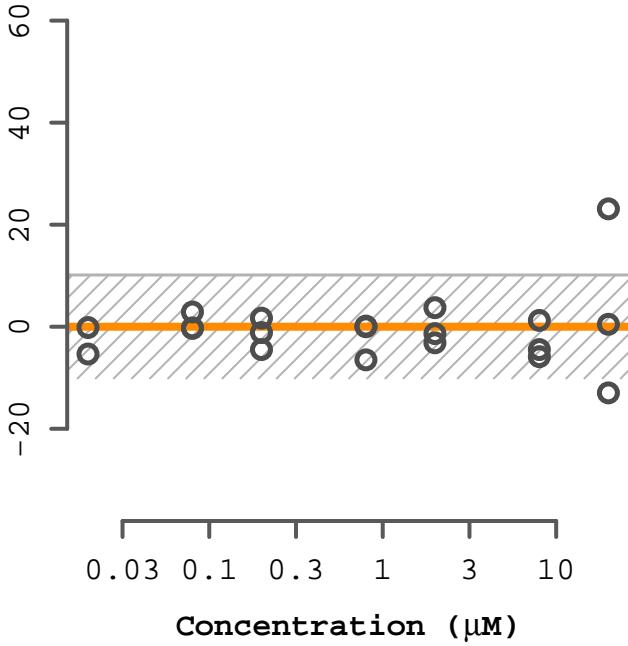
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 76.91	NA	NA
PROB: 1	NA	NA
RMSE: 2.03	NA	NA

MAX_MEAN: 2.93 MAX_MED: 2.93 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tetrabromophenolphthalein ethyl ester
CHID: 44520 CASRN: 1176-74-5
SPID(S): EPAPLT0154G04
M4ID: 30204217

HILL MODEL: Not applicable.

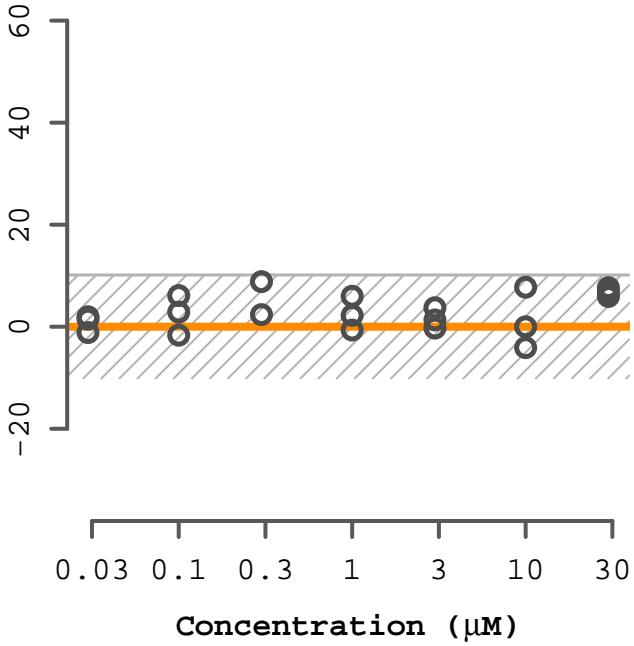
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 126.34	NA	NA
PROB: 1	NA	NA
RMSE: 6.66	NA	NA

MAX_MEAN: 3.55 MAX_MED: 0.485 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: D&C Blue No. 9
CHID: 44532 CASRN: 130-20-1
SPID(S): EPAPLT0154G06
M4ID: 30204219

HILL MODEL: Not applicable.

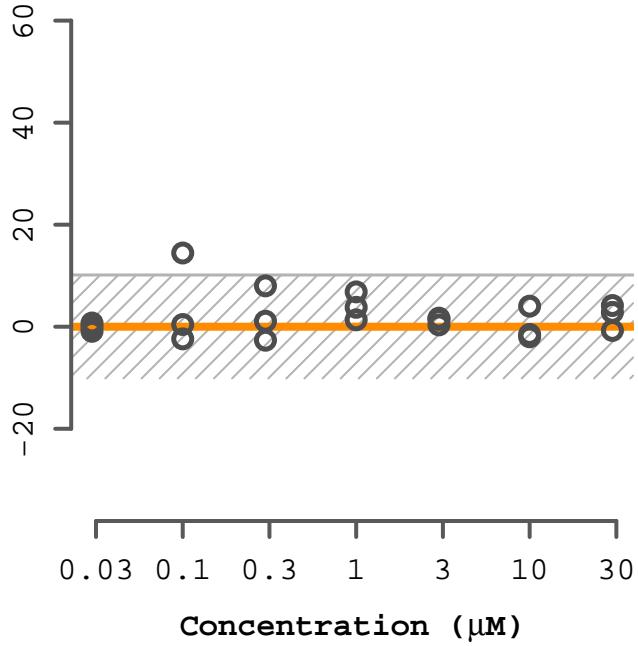
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 121.13	NA	NA
PROB: 1	NA	NA
RMSE: 4.55	NA	NA

MAX_MEAN: 6.79 MAX_MED: 6.75 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Selegiline hydrochloride
CHID: 44584 CASRN: 14611-52-0
SPID(S): EPAPLT0167E07
M4ID: 30204268

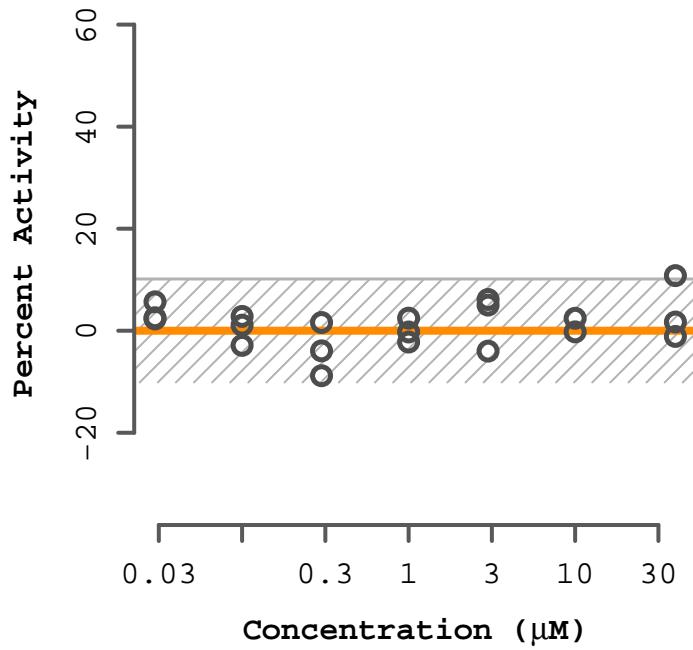
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	118.42	NA	NA
PROB:	1	NA	NA
RMSE:	4.39	NA	NA

MAX_MEAN: 4.18 MAX_MED: 3.81 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Glycidyl trimethylammonium chloride
CHID: 44643 CASRN: 3033-77-0
SPID(S): TP0001412E04
M4ID: 30204651

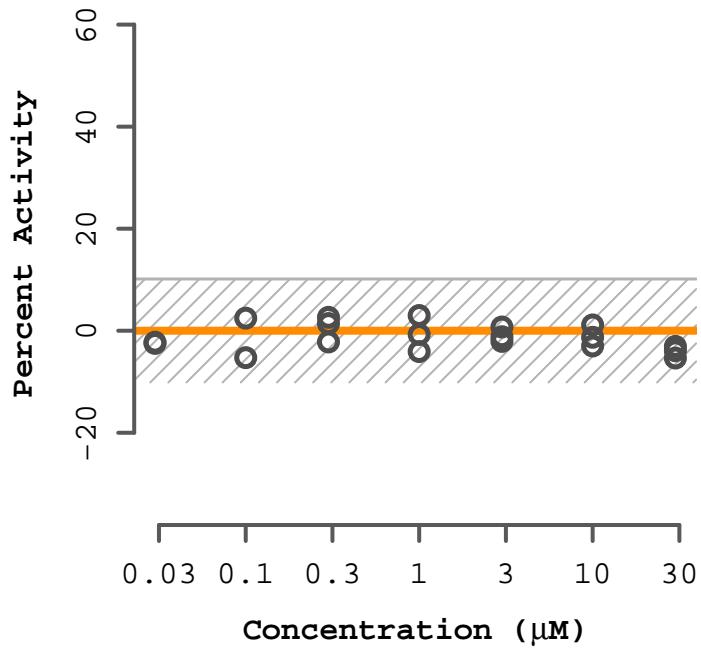
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 117.53	NA	NA
PROB: 1	NA	NA
RMSE: 4.32	NA	NA

MAX_MEAN: 3.8 MAX_MED: 5.03 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Prop-2-en-1-yl 3-cyclohexylpropanoate
CHID: 44755 CASRN: 2705-87-5
SPID(S): EPAPLT0154H06
M4ID: 30204229

HILL MODEL: Not applicable.

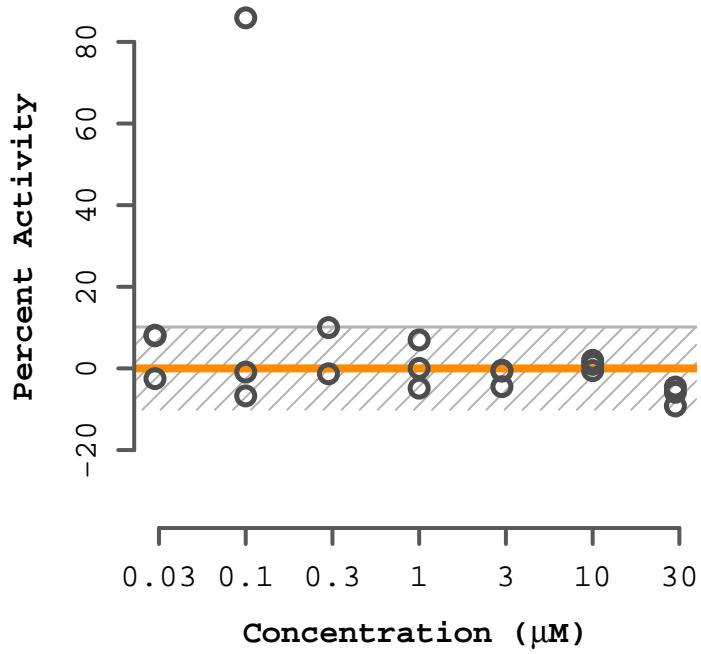
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 98.64	NA	NA
PROB: 1	NA	NA
RMSE: 2.87	NA	NA

MAX_MEAN: 0.616 MAX_MED: 1.41 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Phenylethyl 2-methylpropanoate

CHID: 44764 CASRN: 103-48-0

SPID(S): EPAPLT0154G10

M4ID: 30204223

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 147.56

PROB: 1

RMSE: 19.87

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 26.1

MAX_MED: 7.99

BMAD: 3.38

COFF: 10.1

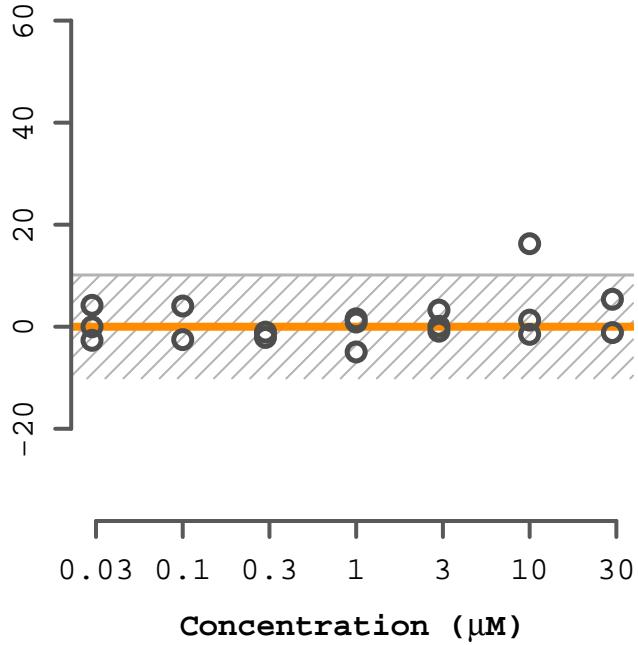
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Benzyl propanoate
CHID: 44791 CASRN: 122-63-4
SPID(S): EPAPLT0154A07
M4ID: 30204158

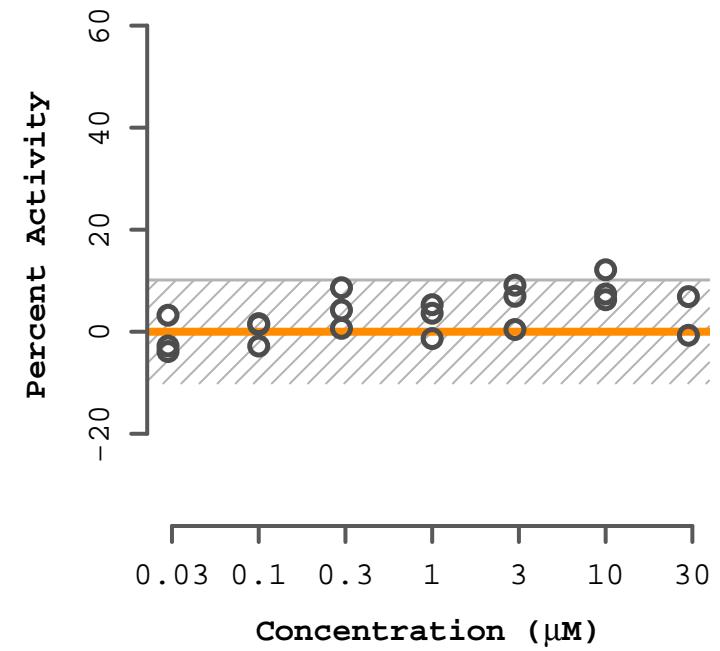
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 101.77	NA	NA
PROB: 1	NA	NA
RMSE: 4.66	NA	NA

MAX_MEAN: 5.36 MAX_MED: 2.12 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ethyl 2-cyano-3,3-diphenylacrylate

CHID: 44821 CASRN: 5232-99-5

SPID(S): EPAPLT0154C08

M4ID: 30204180

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 128.64

PROB: 1

RMSE: 5.49

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 8.61

MAX_MED: 7.4

BMAD: 3.38

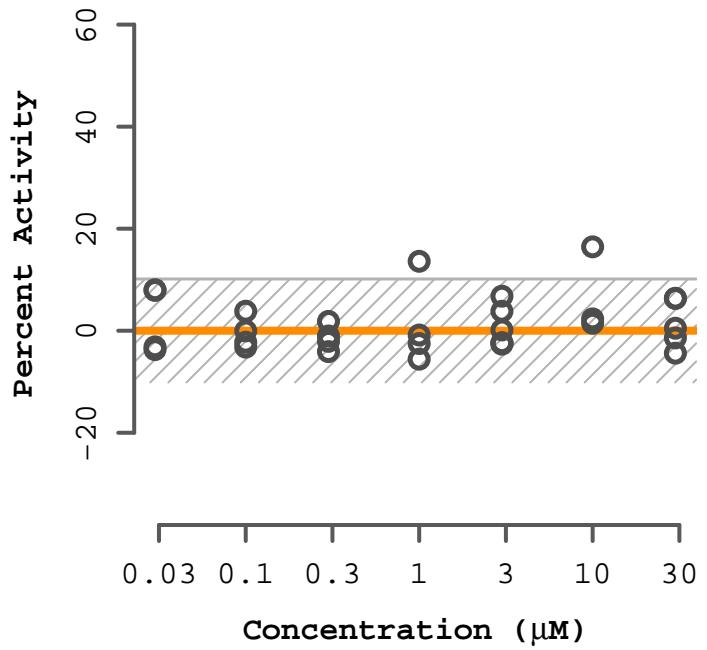
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-[(4-Hydroxyphenyl)methyl]phenol

CHID: 44890 CASRN: 2467-03-0

SPID(S): EPAPLT0154F06

M4ID: 30204209

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 186.2

PROB: 1

RMSE: 5.44

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 6.75

MAX_MED: 2.38

BMAD: 3.38

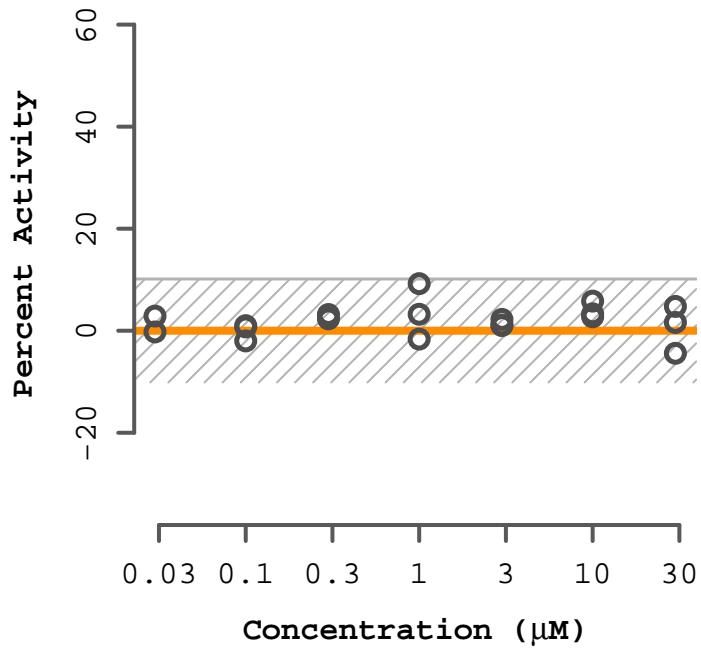
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1-(2-Hydroxyethyl)pyrrolidin-2-one
CHID: 44922 CASRN: 3445-11-2
SPID(S): EPAPLT0154G03
M4ID: 30204216

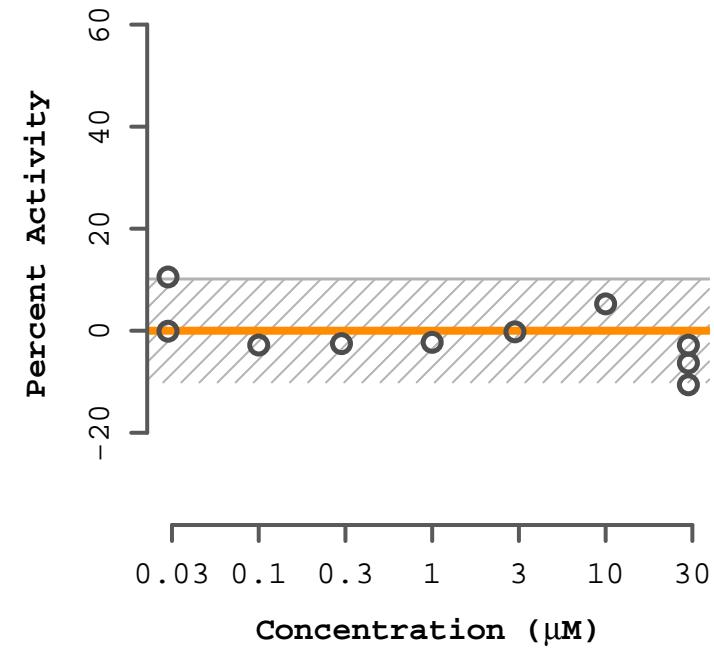
HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 103.93	NA	NA
PROB: 1	NA	NA
RMSE: 3.5	NA	NA

MAX_MEAN: 3.96 MAX_MED: 3.32 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4,5-Dihydroxy-1,3-dimethylimidazolidin-2-one

CHID: 44934 CASRN: 3923-79-3

SPID(S): EPAPLT0154A10

M4ID: 30204161

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 65.45

PROB: 1

RMSE: 5.64

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 5.27

MAX_MED: 5.27

BMAD: 3.38

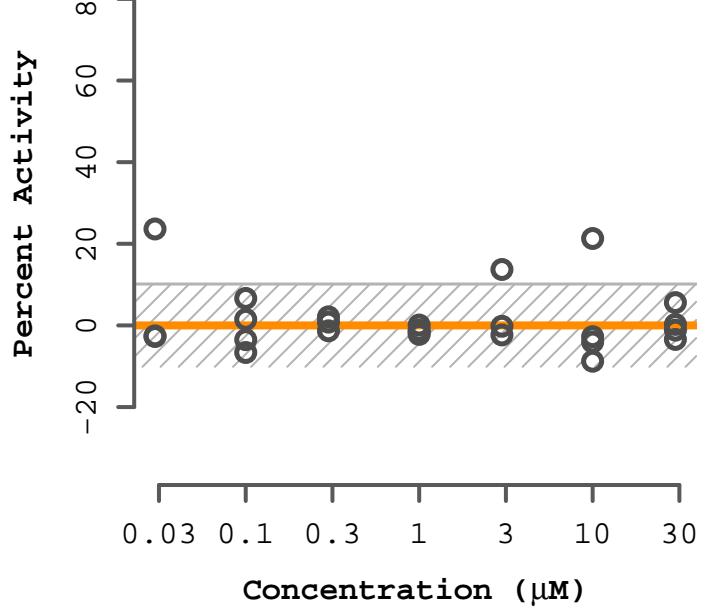
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tembotrione

CHID: 47037 CASRN: 335104-84-2

SPID(S): EPAPLT0167H07

M4ID: 30204272 BRK

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 195

PROB: 1

RMSE: 21.04

HILL

NA

NA

NA

GNLS

NA

NA

NA

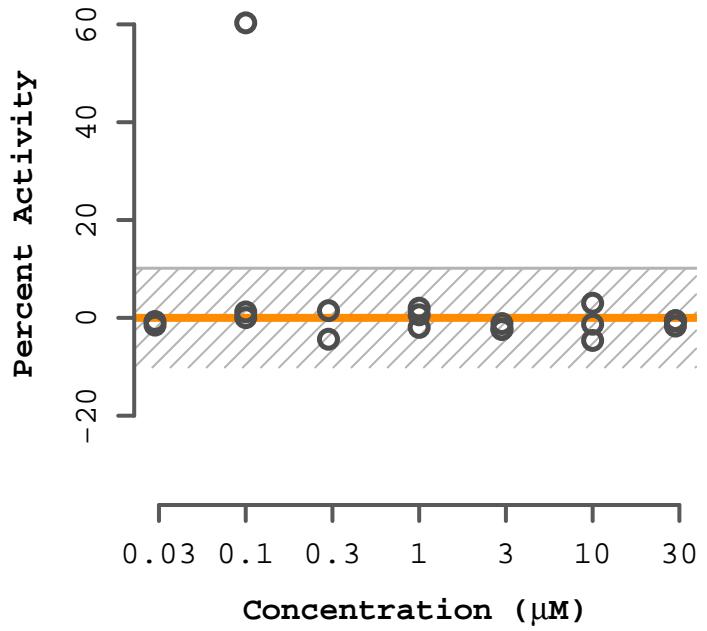
MAX_MEAN: 20.3

MAX_MED: 0.82

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ethyl 2-methylpentanoate
CHID: 47198 CASRN: 39255-32-8
SPID(S): EPAPLT0154H09
M4ID: 30204232

HILL MODEL: Not applicable.

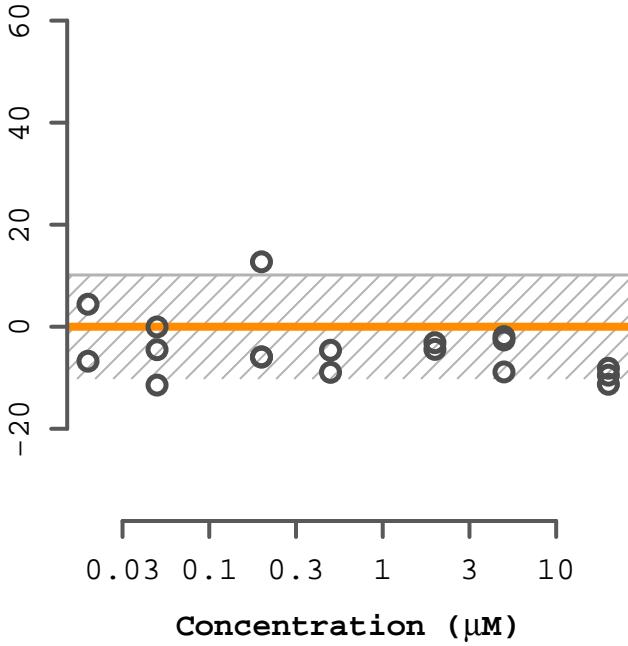
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 112.01	NA	NA
PROB: 1	NA	NA
RMSE: 13.99	NA	NA

MAX_MEAN: 20.5 MAX_MED: 1.22 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

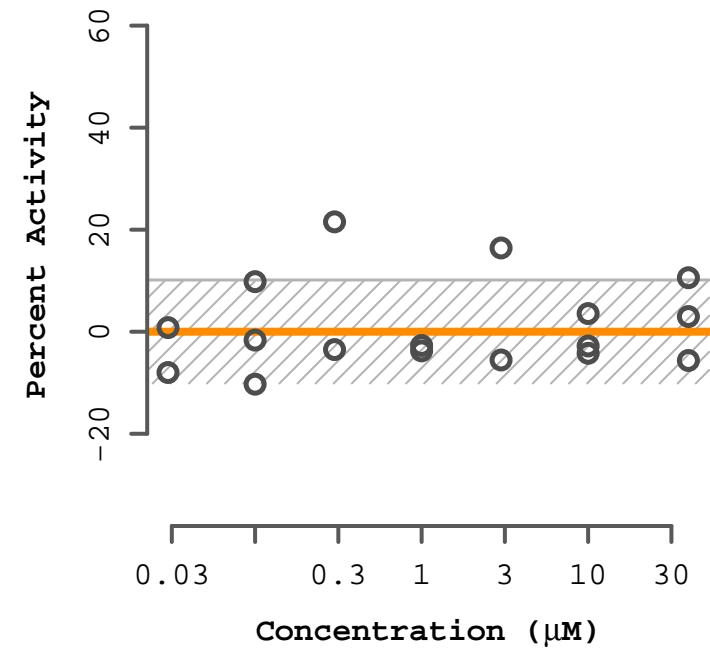
NAME: Fandosentan potassium salt
CHID: 47249 CASRN: 221246-12-4
SPID(S): TP0001414F04
M4ID: 30204491

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

	CNST	HILL	GNLS
AIC:	127.26	NA	NA
PROB:	1	NA	NA
RMSE:	7.27	NA	NA
MAX_MEAN:	0.298	MAX_MED: -1.17	BMAD: 3.38
COFF:	10.1	HIT-CALL: 0	FITC: 4
ACTP:	0		

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: CP-471358

CHID: 47265 CASRN: NOCAS_47265

SPID(S): TP0001412G04

M4ID: 30204496

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 123.27

PROB: 1

RMSE: 8.6

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 9.03

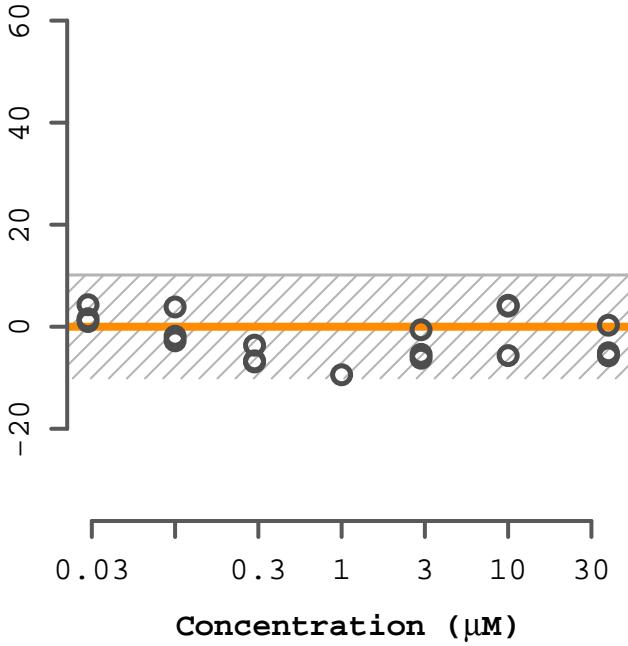
MAX_MED: 9.03

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: CI-959

CHID: 47268 CASRN: 104795-68-8

SPID(S): TP0001411G09

M4ID: 30204497

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 118.24

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 4.78

NA

NA

MAX_MEAN: 2.31

MAX_MED: 4.04

BMAD: 3.38

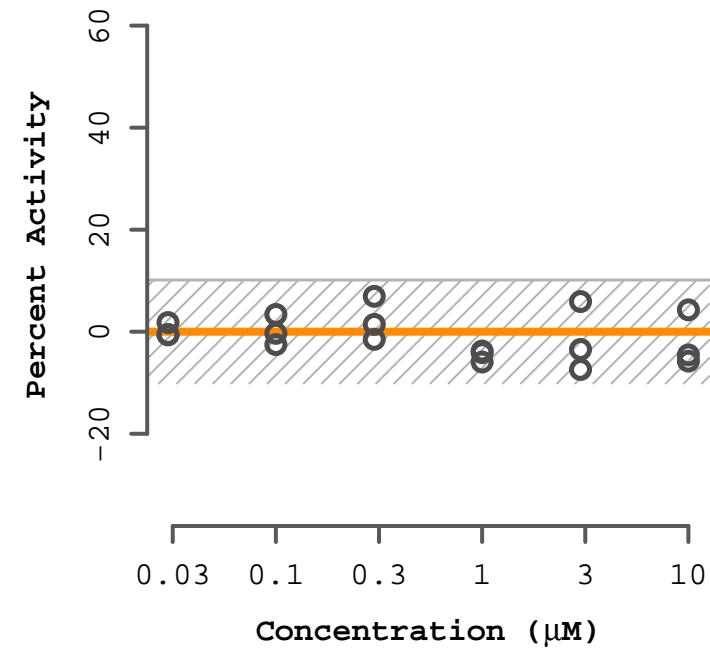
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: CP-728663

CHID: 47283 CASRN: 368832-42-2

SPID(S): TP0001412A11

M4ID: 30204502

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 107.12

PROB: 1

RMSE: 4.18

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 2.3

MAX_MED: 1.44

BMAD: 3.38

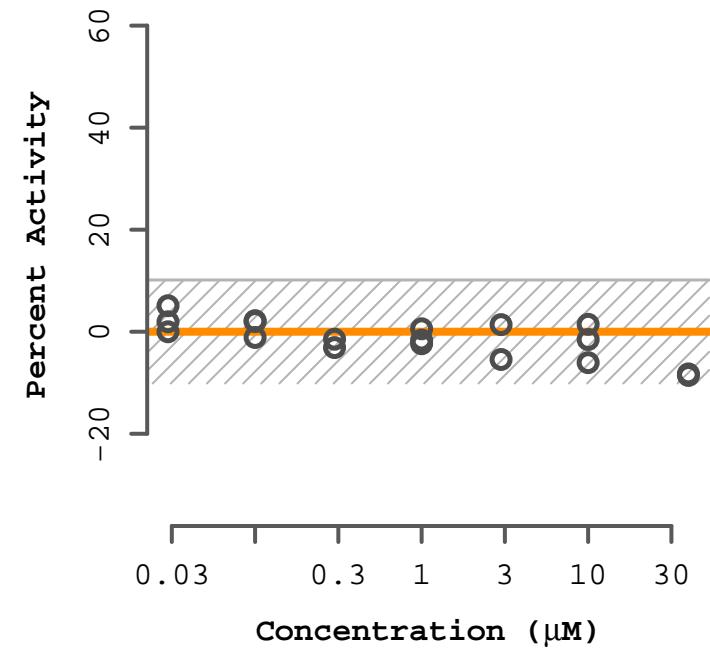
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: PHA-00543613

CHID: 47284 CASRN: 478149-53-0

SPID(S): TP0001414F09

M4ID: 30204503

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 102.22

NA

NA

PROB:

1

NA

NA

RMSE: 3.9

NA

NA

MAX_MEAN: 2.36

MAX_MED: 2.01

BMAD: 3.38

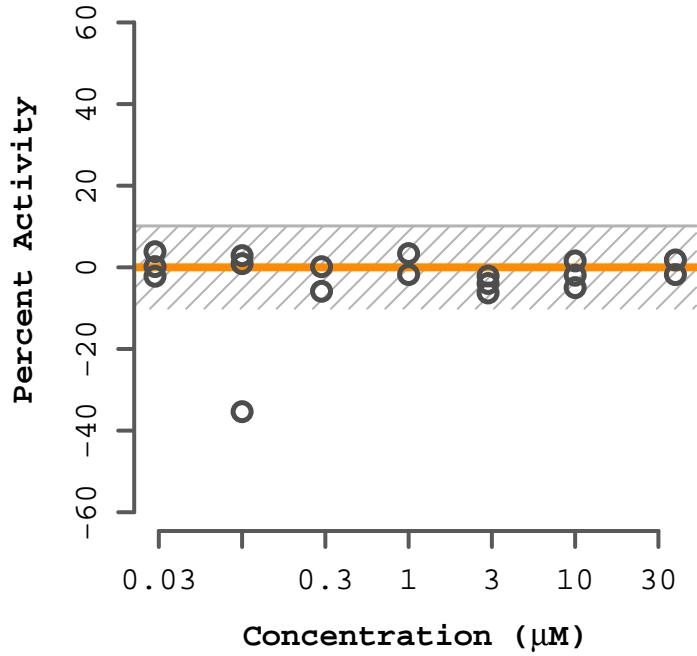
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: PD 0200347

CHID: 47292 CASRN: NOCAS_47292

SPID(S): TP0001411B07

M4ID: 30204506

HILL MODEL: Not applicable.

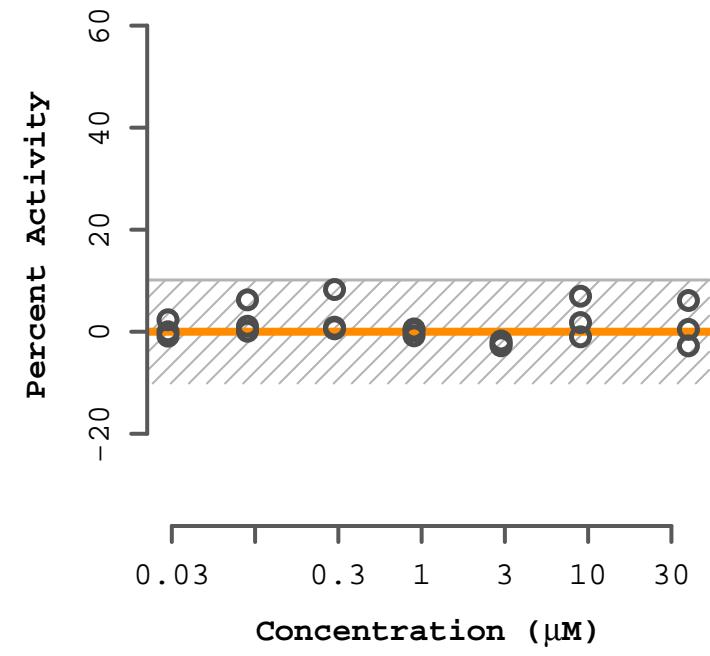
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 113.47	NA	NA
PROB: 1	NA	NA
RMSE: 8.89	NA	NA

MAX_MEAN: 0.811 MAX_MED: 0.916 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SB243213A

CHID: 47322 CASRN: 200940-23-4

SPID(S): TP0001413E01

M4ID: 30204510

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 109.72

PROB: 1

RMSE: 3.32

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 3.25

MAX_MED: 1.8

BMAD: 3.38

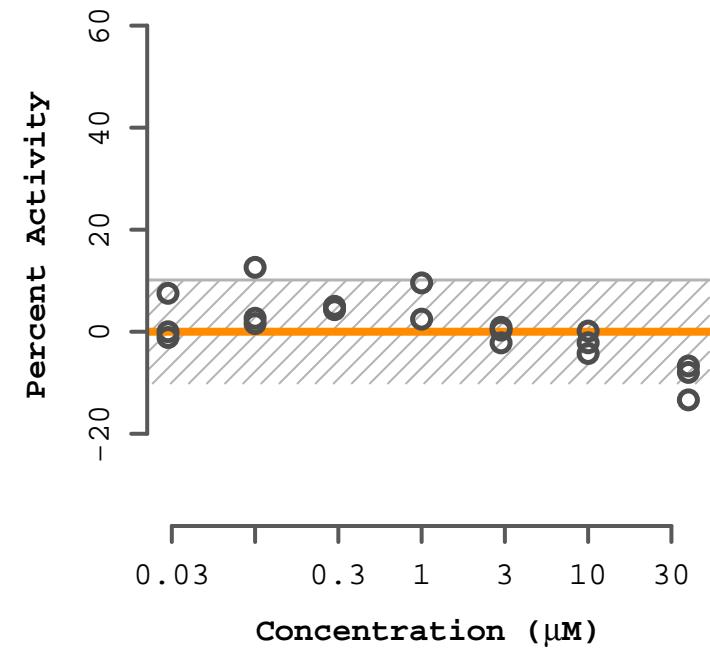
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: PharmaGSID_47333

CHID: 47333 CASRN: NOCAS_47333

SPID(S): TP0001413F04

M4ID: 30204527

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 130.39

PROB: 1

RMSE: 5.94

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 6.02

MAX_MED: 6.02

BMAD: 3.38

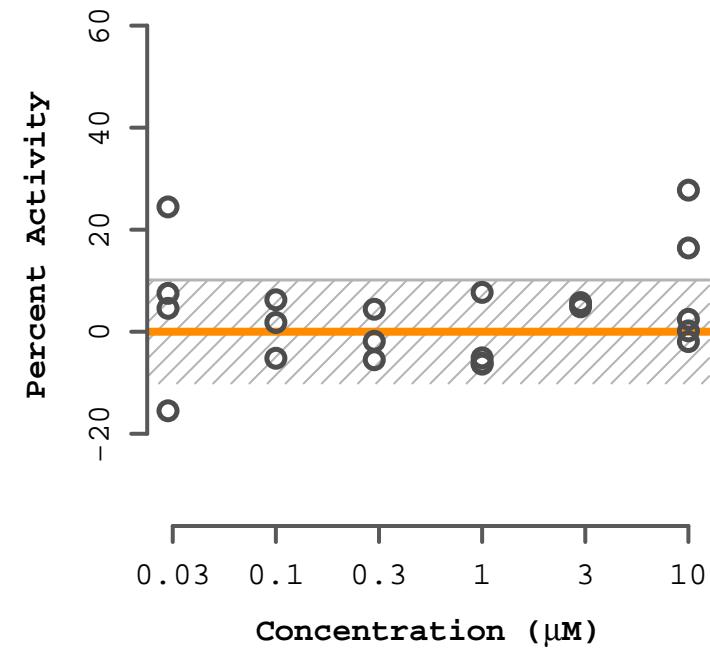
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SR144190

CHID: 47339 CASRN: 181640-09-5

SPID(S): TP0001411A03

M4ID: 30204643

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 158.17

NA

NA

PROB:

1

NA

NA

RMSE: 10.49

NA

NA

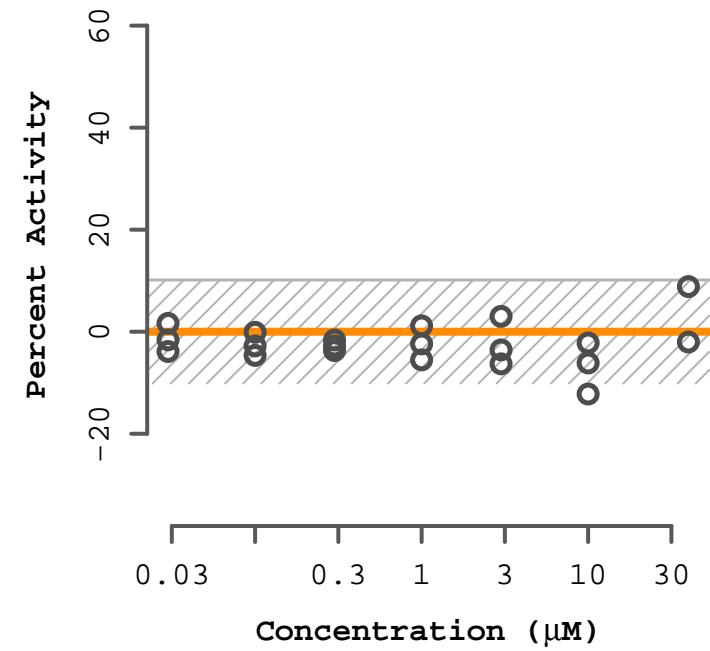
MAX_MEAN: 8.97

MAX_MED: 6.06

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SSR180711

CHID: 47359 CASRN: 298198-52-4

SPID(S): TP0001414G09

M4ID: 30204519

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 121.26

NA

NA

PROB:

1

NA

NA

RMSE: 4.72

NA

NA

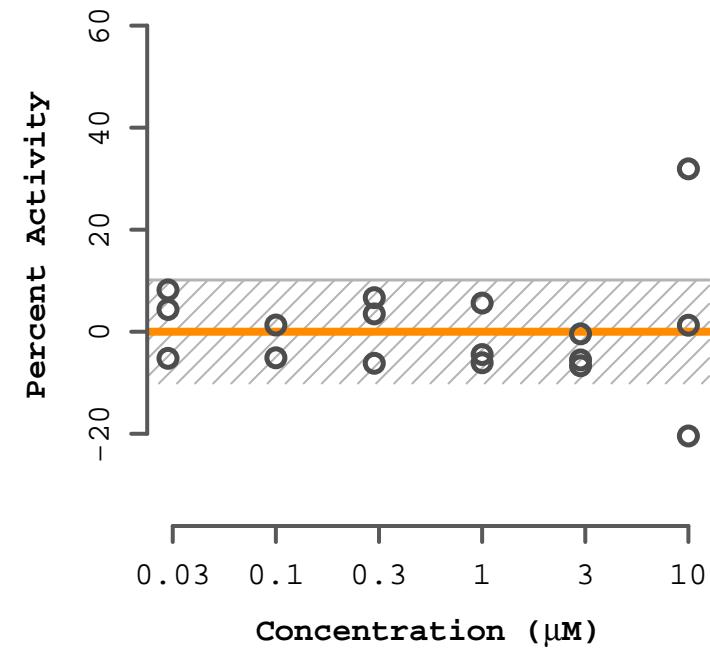
MAX_MEAN: 3.42

MAX_MED: 3.42

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Volinanserin

CHID: 47363 CASRN: 139290-65-6

SPID(S): TP0001414E01

M4ID: 30204520

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

HILL

GNLS

AIC: 125.66

NA

NA

PROB:

1

NA

NA

RMSE: 10.4

NA

NA

MAX_MEAN: 4.26

MAX_MED: 4.36

BMAD: 3.38

COFF:

10.1

HIT-CALL:

0

FITC:

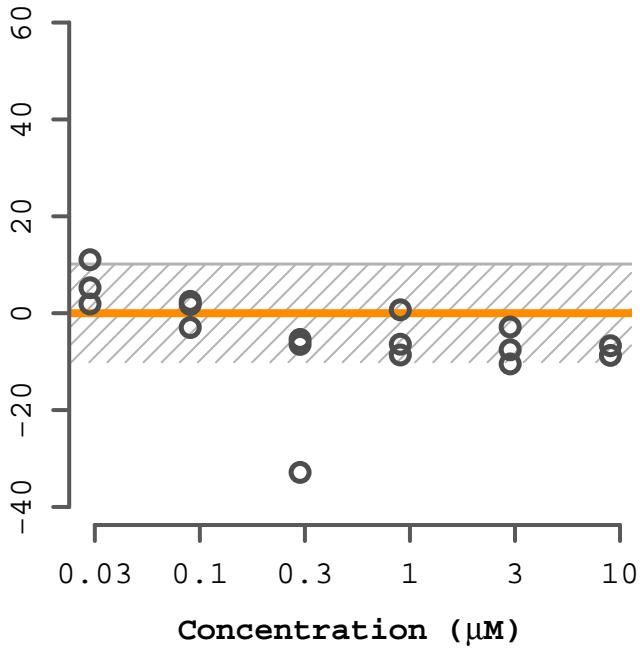
4

ACTP:

0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SSR 103800

CHID: 47364 CASRN: 1075752-90-7

SPID(S): TP0001412A10

M4ID: 30204521

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 125.07

PROB: 1

RMSE: 10.1

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 6.07

MAX_MED: 5.26

BMAD: 3.38

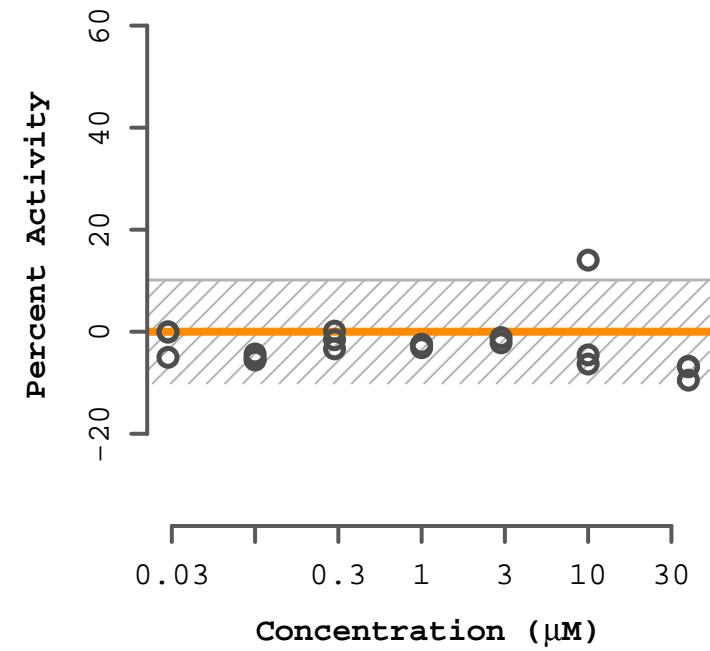
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: AVE9423

CHID: 47375 CASRN: 862243-29-6

SPID(S): TP0001411E10

M4ID: 30204525

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 126.92

HILL

PROB: 1

NA

RMSE: 5.41

NA

GNLS

NA

NA

NA

MAX_MEAN: 1.06

MAX_MED: -0.0791

BMAD: 3.38

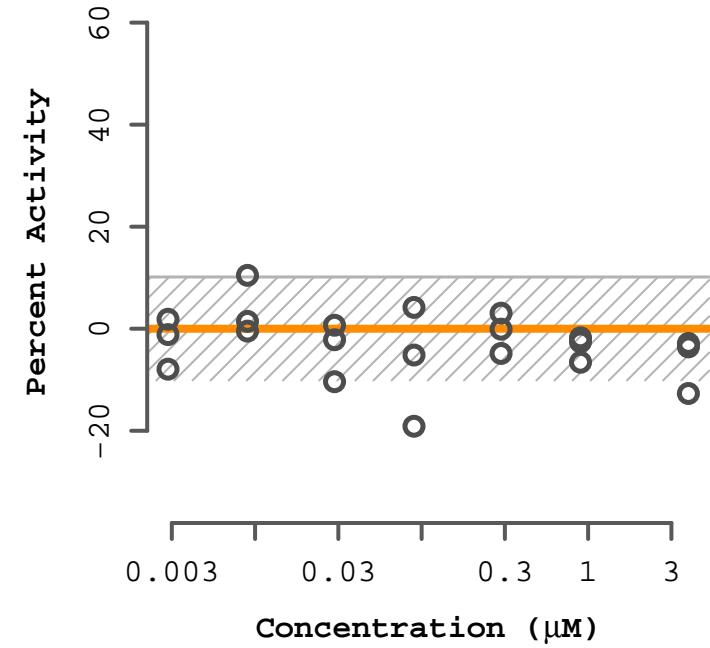
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: AVE8923

CHID: 47381 CASRN: NOCAS 47381

SPID(S): TP0001414C08

M4ID: 30204530

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

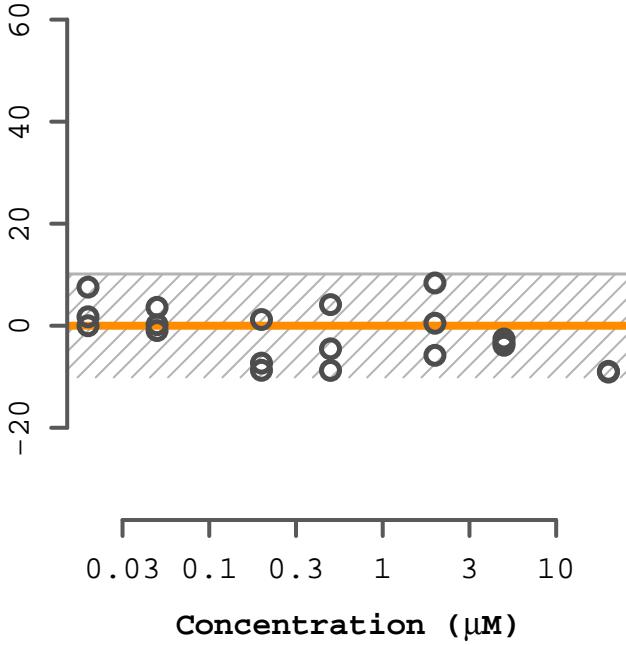
	CNST	HILL	GNLS
AIC:	140.4	NA	NA
PROB:	1	NA	NA
RMSE:	6.79	NA	NA

MAX MEAN: 3.82 MAX MED: 1.45 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pravastatin sodium

CHID: 47525 CASRN: 81131-70-6

SPID(S): TP0001412B05

M4ID: 30204529

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 129.93

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 5.57

NA

NA

MAX_MEAN: 3.11

MAX_MED: 1.76

BMAD: 3.38

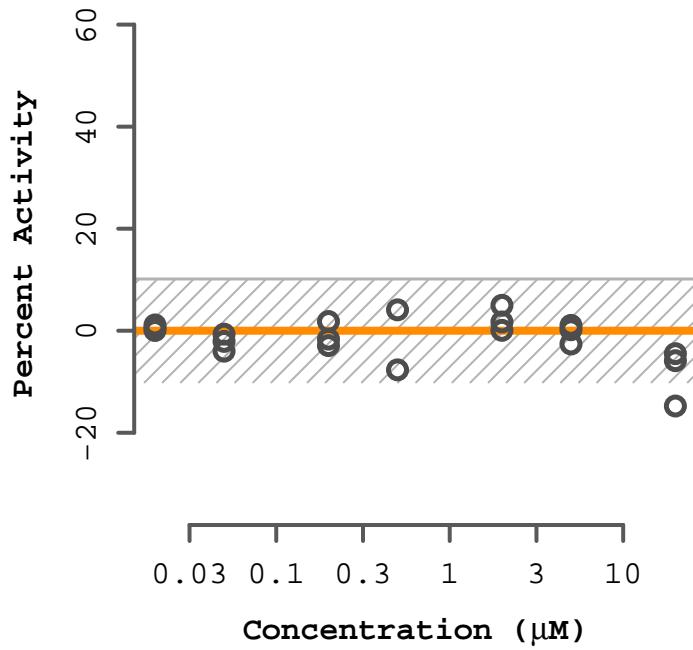
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: C10-21 alkanesulfonic acids phenyl esters

CHID: 47526 CASRN: 91082-17-6

SPID(S): TP0001413B02

M4ID: 30204381

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 115.83

PROB: 1

RMSE: 4.58

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 2.27

MAX_MED: 1.73

BMAD: 3.38

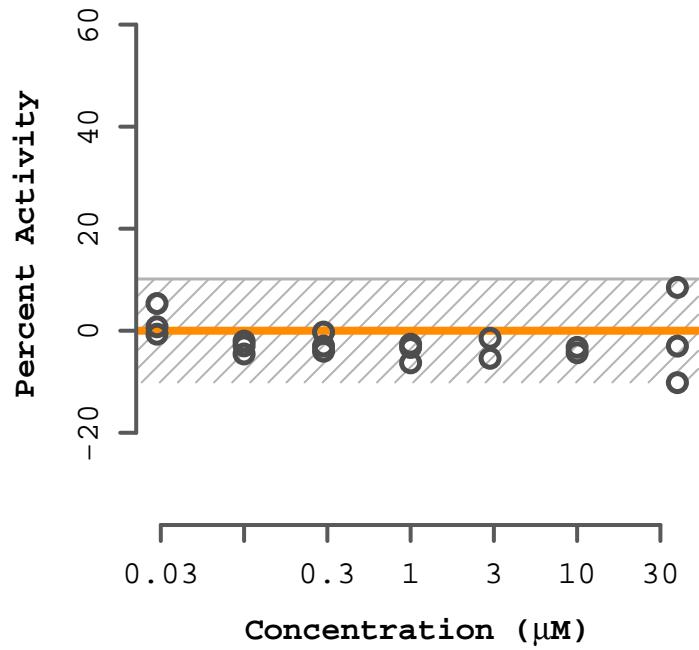
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Perfluoroundecanoic acid
CHID: 47553 CASRN: 2058-94-8
SPID(S): TP0001411E11
M4ID: 30204665

HILL MODEL: Not applicable.

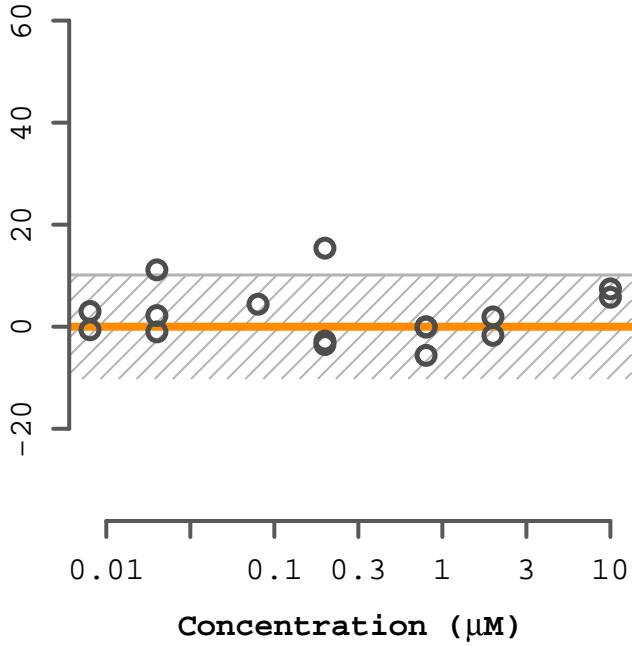
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 115.01	NA	NA
PROB: 1	NA	NA
RMSE: 4.53	NA	NA

MAX_MEAN: 1.81 MAX_MED: 0.772 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1H,1H,2H,2H-Perfluoroctyl iodide

CHID: 47565 CASRN: 2043-57-4

SPID(S): TP0001413A04

M4ID: 30204605

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

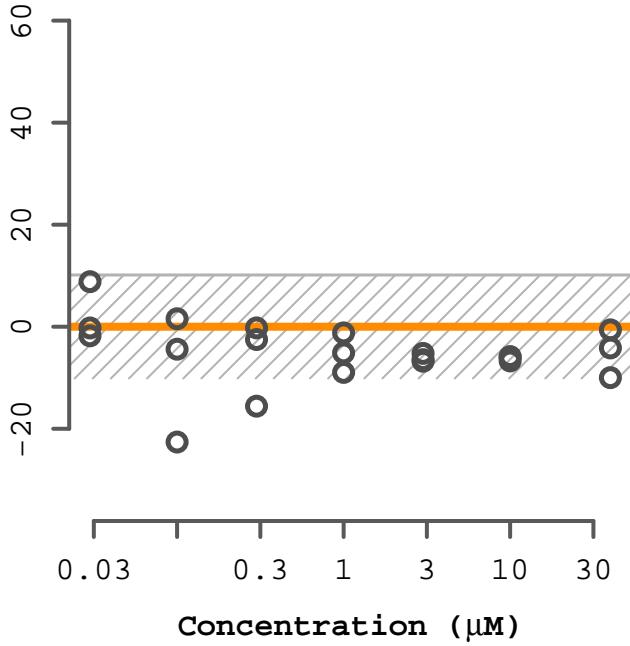
	CNST	HILL	GNLS
AIC:	102.45	NA	NA
PROB:	1	NA	NA
RMSE:	5.83	NA	NA

MAX_MEAN: 6.61 MAX_MED: 6.61 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cotinine

CHID: 47576 CASRN: 486-56-6

SPID(S): TP0001414C05

M4ID: 30204609

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 147.49

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 7.92

NA

NA

MAX_MEAN: 2.28

MAX_MED: -0.236

BMAD: 3.38

COFF: 10.1

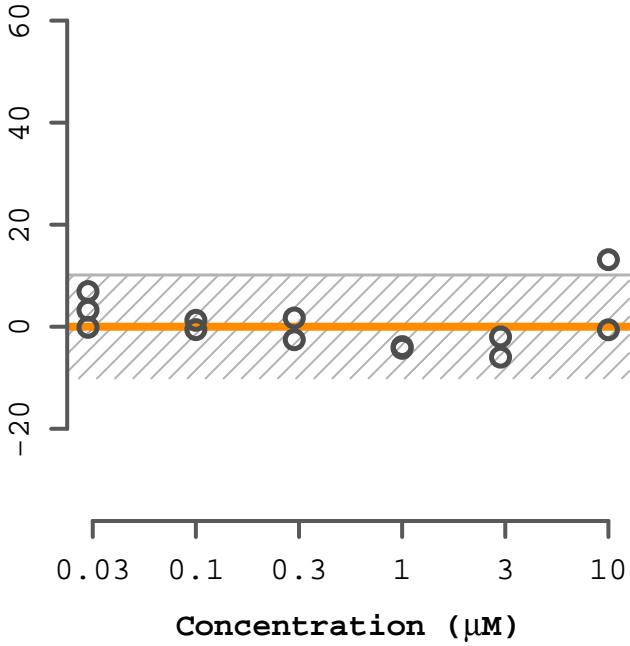
HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 9-Phenanthrol

CHID: 47592 CASRN: 484-17-3

SPID(S): TP0001411F07

M4ID: 30204629

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 79.14

HILL

PROB: 1

GNLS

RMSE: 4.92

MAX_MEAN: 6.28

MAX_MED: 6.28

BMAD: 3.38

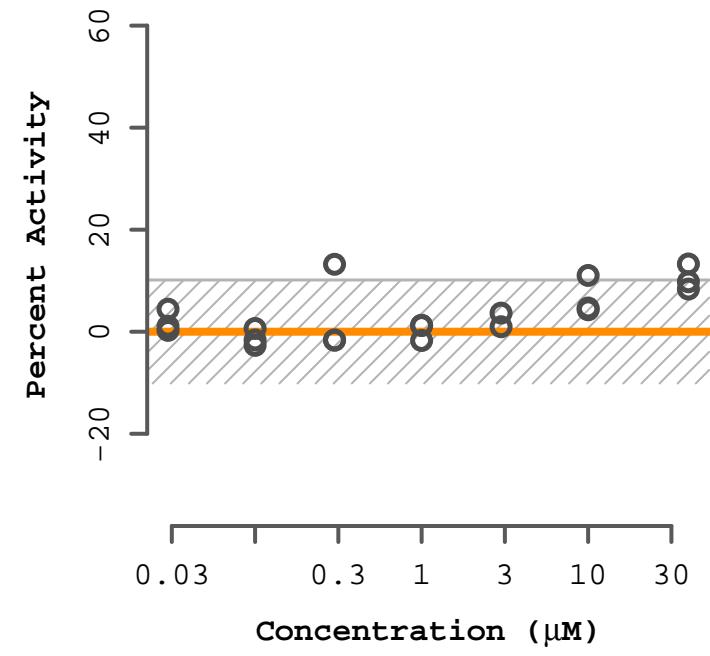
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4,4'-Sulfonylbis[2-(prop-2-en-1-yl)phenol]

CHID: 47598 CASRN: 41481-66-7

SPID(S): TP0001413G04

M4ID: 30204618

HILL MODEL: Not applicable.

GAIN-LOSS MODEL: Not applicable.

CNST

AIC: 130.04

PROB: 1

RMSE: 6.07

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 10.5

MAX_MED: 9.8

BMAD: 3.38

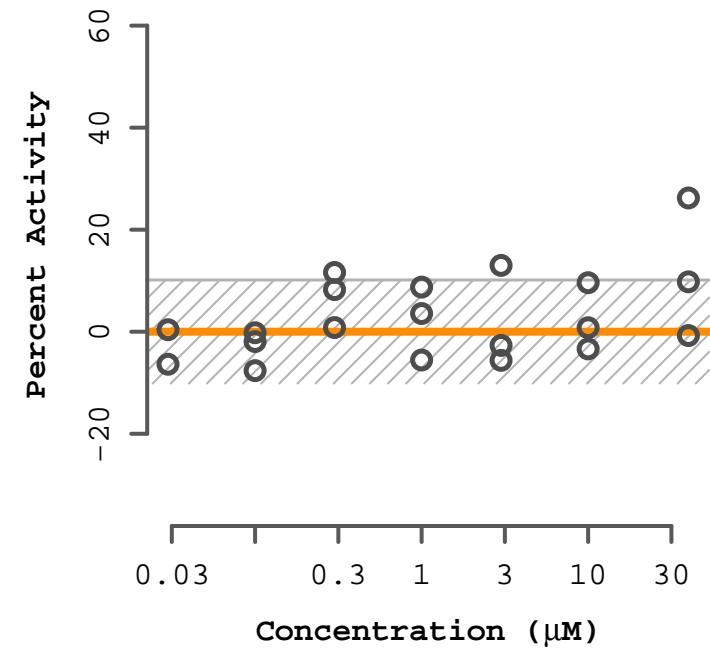
COFF: 10.1

HIT-CALL: 0

FITC: 4

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cyclohexylphenylketone
CHID: 48182 CASRN: 712-50-5
SPID(S): TP0001411H05
M4ID: 30204658

HILL MODEL: Not applicable.

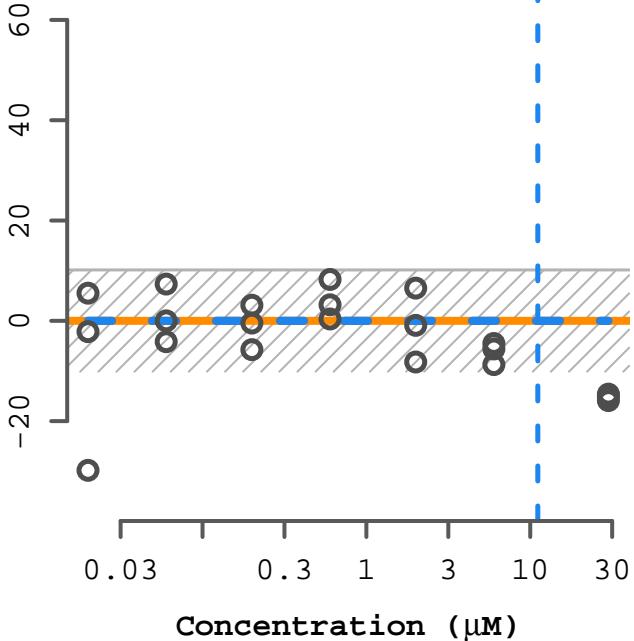
GAIN-LOSS MODEL: Not applicable.

CNST	HILL	GNLS
AIC: 143.92	NA	NA
PROB: 1	NA	NA
RMSE: 8.73	NA	NA

MAX_MEAN: 11.8 MAX_MED: 9.78 BMAD: 3.38
COFF: 10.1 HIT-CALL: 0 FITC: 4 ACTP: 0

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: FD&C Blue No. 1

CHID: 20189 CASRN: 3844-45-9

SPID(S): TP0001413A05

M4ID: 30204560

HILL MODEL (in red):

tp	ga	gw
4.69e-07	1.94	5.21
NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

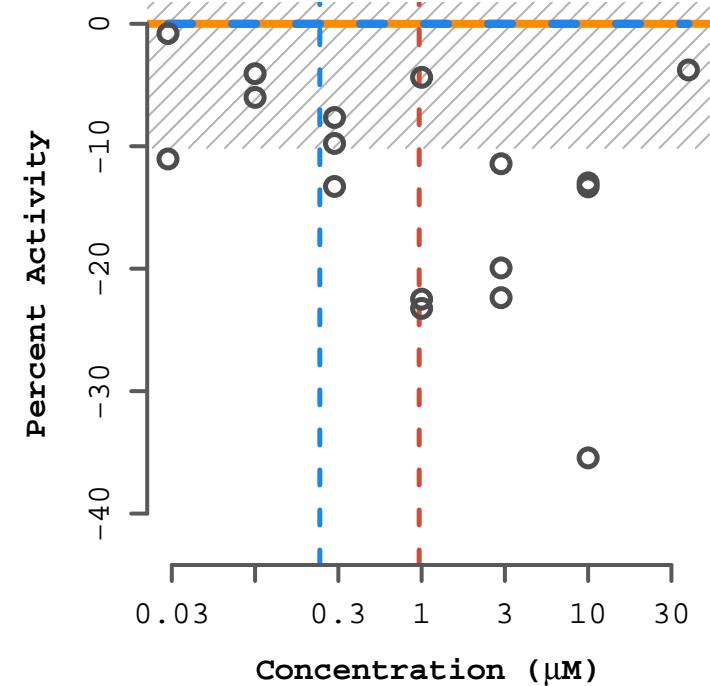
tp	ga	gw	la	lw
3.08e-09	1.05	0.491	1.58	7.03
NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
155.72	161.72	165.72
0.95	0.05	0.01
9.87	9.87	9.87

MAX_MEAN: 3.94 MAX_MED: 3.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Chlordane

CHID: 20267 CASRN: 57-74-9

SPID(S): TP0001411E05

M4ID: 30204598

HILL MODEL (in red):

	tp	ga	gw
val:	6.23e-10	-0.0146	0.885
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

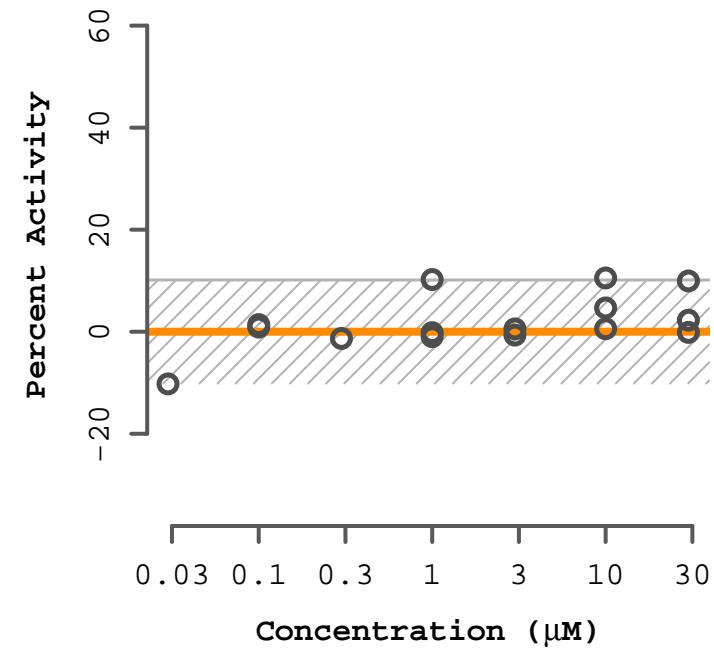
	tp	ga	gw	la	lw
val:	7.65e-10	-0.611	1.85	0.00985	5.64
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	173.94	179.94	183.94
PROB:	0.95	0.05	0.01
RMSE:	14.26	14.26	14.26

MAX_MEAN: -0.205 MAX_MED: 0.686 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Chloro-1,4-diaminobenzene sulfate

CHID: 20284 CASRN: 61702-44-1

SPID(S): EPAPLT0154F07

M4ID: 30204210

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST

AIC: 94.77

PROB: 1

RMSE: 5.5

HILL

NA

NA

NA

GNLS

NA

NA

NA

MAX_MEAN: 5.27

MAX_MED: 4.67

BMAD: 3.38

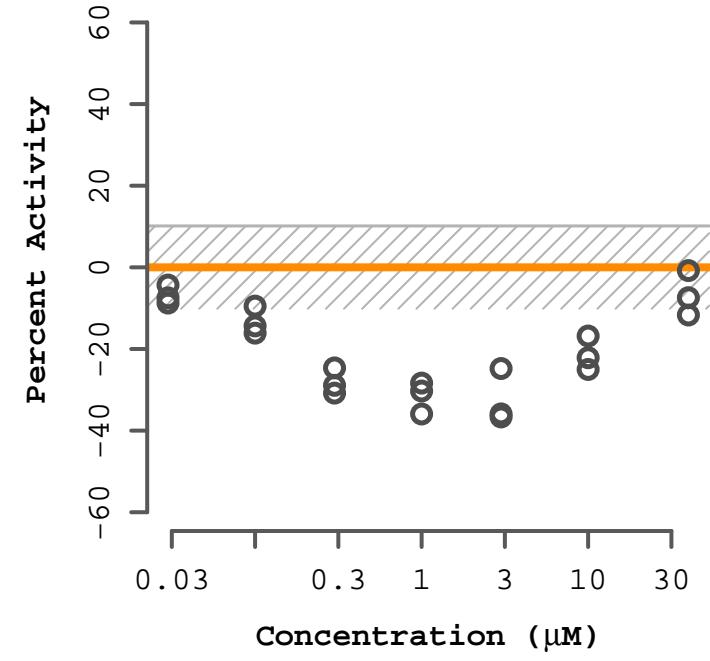
COFF: 10.1

HIT-CALL: 0

FITC: 7

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dieldrin

CHID: 20453 CASRN: 60-57-1

SPID(S): TP0001412F07

M4ID: 30204552

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST

AIC: 196.34

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 22.83

NA

NA

MAX_MEAN: -6.63

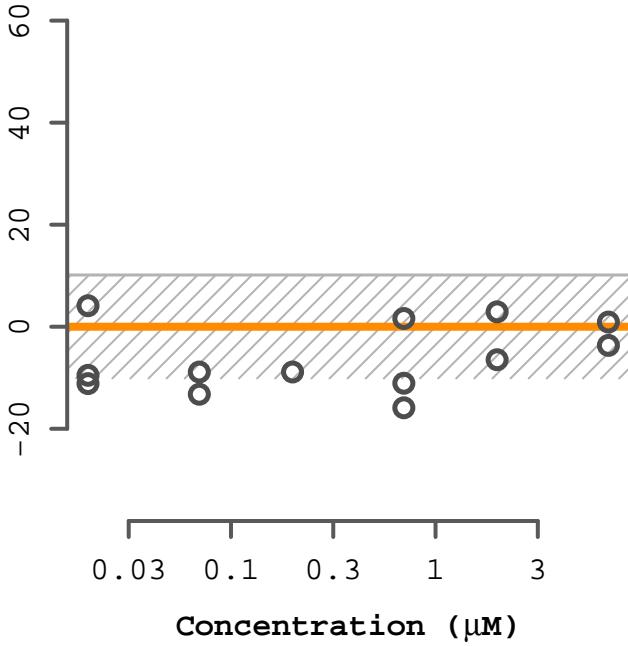
MAX_MED: -7.43

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Methadone hydrochloride
CHID: 20501 CASRN: 1095-90-5
SPID(S): TP0001412E10
M4ID: 30204620

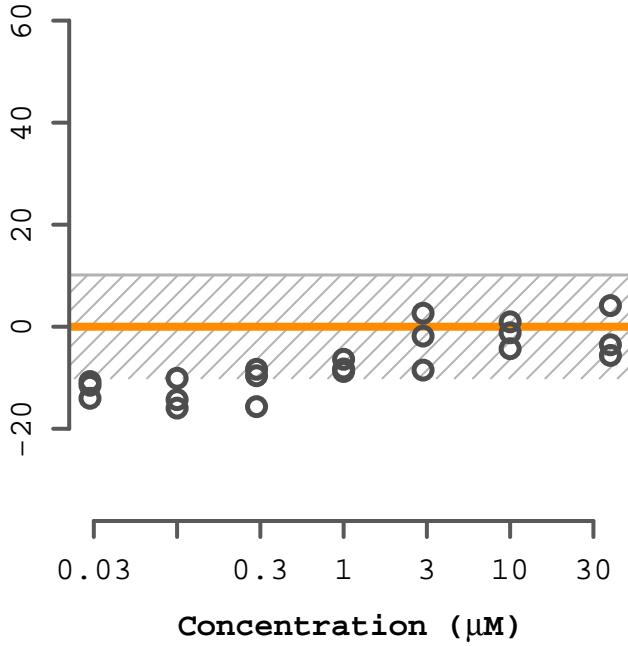
HILL MODEL: Failed to converge.

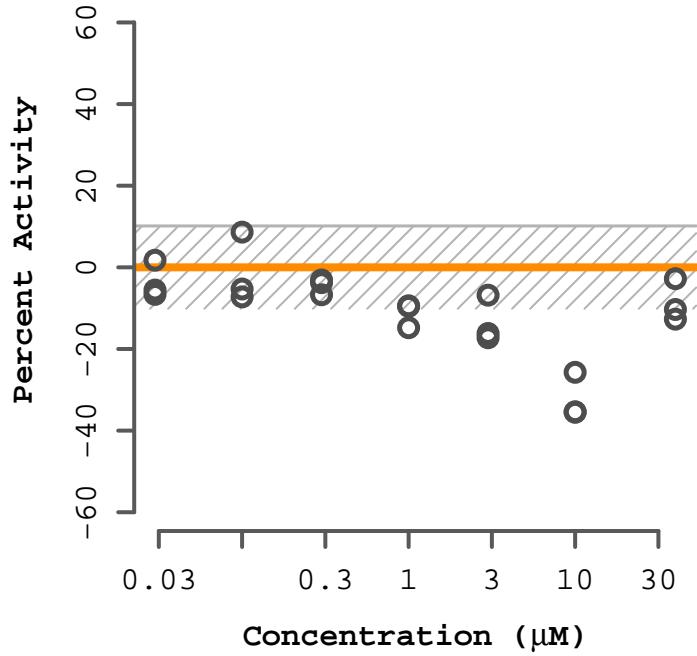
GAIN-LOSS MODEL: Failed to converge.

	CNST	HILL	GNLS
AIC:	97.29	NA	NA
PROB:	1	NA	NA
RMSE:	8.8	NA	NA
MAX_MEAN:	-1.37	MAX_MED: -1.37	BMAD: 3.38
COFF:	10.1	HIT-CALL: 0	FITC: 7
ACTP:	0		

FLAGS:

Percent Activity





ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Heptachlor

CHID: 20679 CASRN: 76-44-8

SPID(S): TP0001411C06

M4ID: 30204564

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST

AIC: 174.99

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 15.04

NA

NA

MAX_MEAN: -1.3

MAX_MED: -3.75

BMAD: 3.38

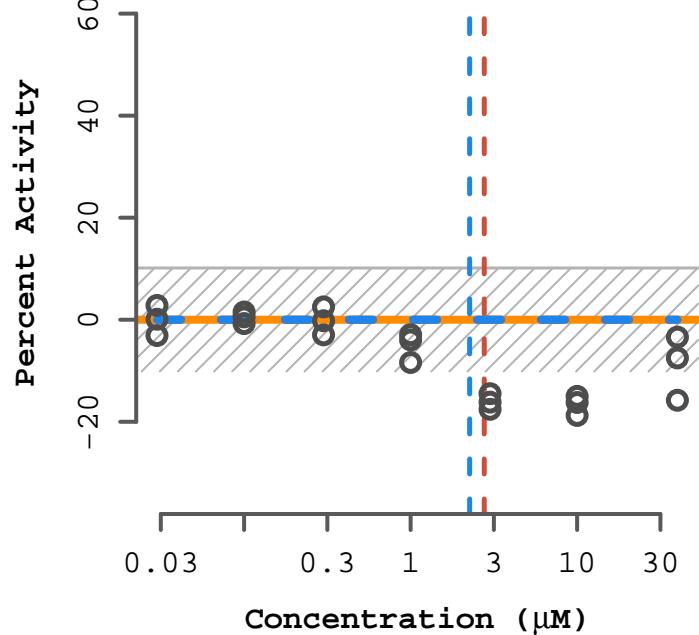
COFF: 10.1

HIT-CALL: 0

FITC: 7

ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Lindane

CHID: 20686 CASRN: 58-89-9

SPID(S): TP0001413B01

M4ID: 30204375

HILL MODEL (in red):

tp	ga	gw
val: 1.14e-09	0.442	1.27
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

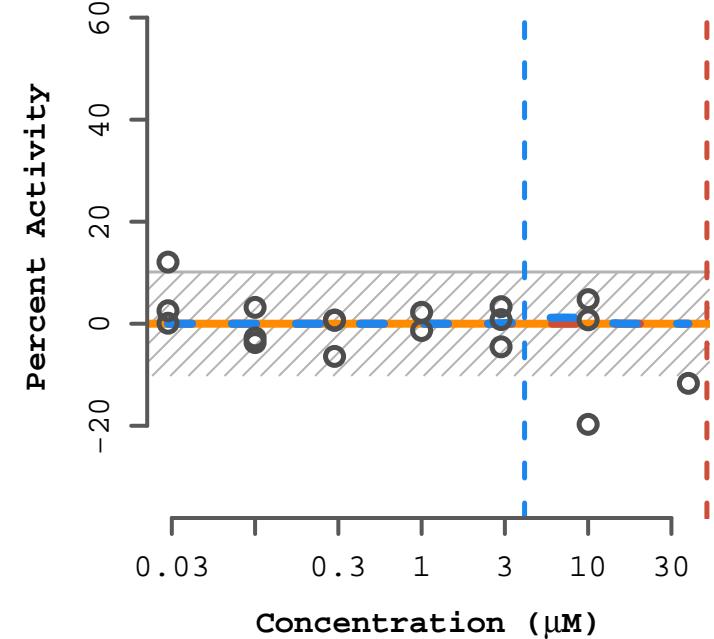
tp	ga	gw	la	lw
val: 4.69e-10	0.354	1.69	1.29	5.11
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 158.96	164.96	168.96
PROB: 0.95	0.05	0.01
RMSE: 9.92	9.92	9.92

MAX_MEAN: 0.446 MAX_MED: 0.561 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Hydroquinone

CHID: 20716 CASRN: 123-31-9

SPID(S): TP0001414G12

M4ID: 30204358

HILL MODEL (in red):

	tp	ga	gw
val:	3.15e-10	1.71	2.5
sd:	6.53e-05	20400	516000

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	1.25	0.618	7.76	1.04	9.97
sd:	73.9	7.92	448	8.09	752

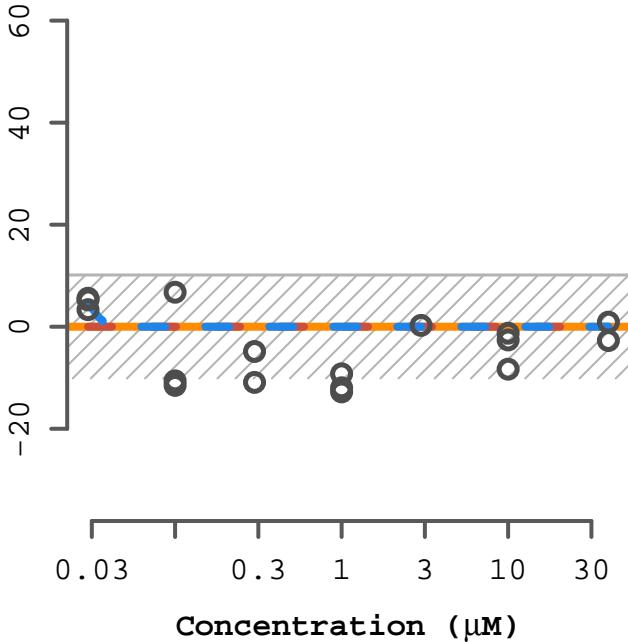
CNST	HILL	GNLS
118.13	124.13	128.06
0.95	0.05	0.01
6.72	6.72	6.84

MAX_MEAN: 4.92 MAX_MED: 2.56 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 6-Propyl-2-thiouracil
CHID: 21209 CASRN: 51-52-5
SPID(S): TP0001412B08
M4ID: 30204542

HILL MODEL (in red):

tp	ga	gw
val: 0.00554	2.07	6.97
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 7.08	-2.22	1.92	-1.49	12.3
sd: NaN	NaN	NaN	NaN	NaN

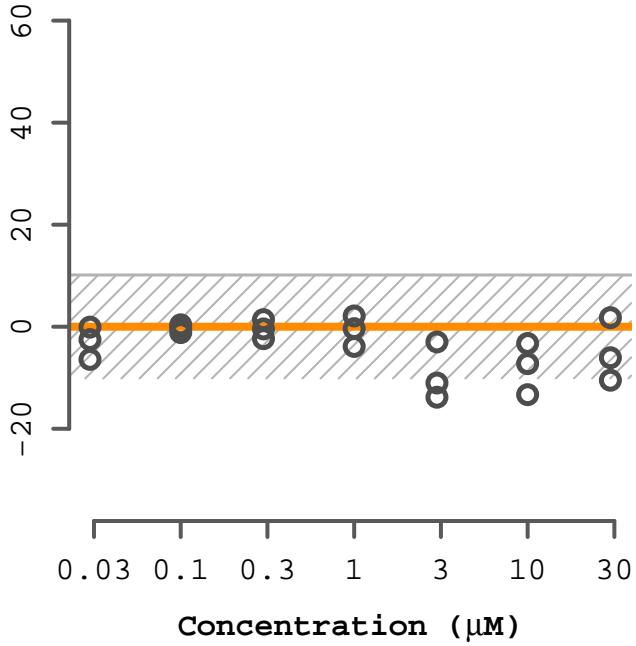
CNST	HILL	GNLS
AIC: 121.49	127.49	129.68
PROB: 0.94	0.05	0.02
RMSE: 7.59	7.59	7.33

MAX_MEAN: 4.74 MAX_MED: 5.26 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.06

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Trichlorfon

CHID: 21389 CASRN: 52-68-6

SPID(S): EPAPLT0170D03

M4ID: 30204278

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST

AIC: 136.86

HILL

PROB: 1

NA

RMSE: 6.12

NA

GNLS

NA

NA

MAX_MEAN: -0.361

MAX_MED: -0.328

BMAD: 3.38

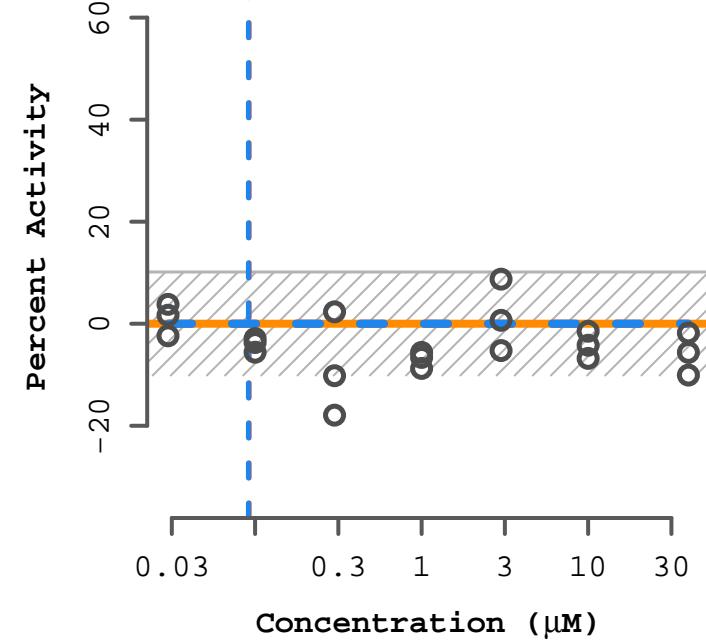
COFF: 10.1

HIT-CALL: 0

FITC: 7

ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Octadecanoic acid
CHID: 21642 CASRN: 57-11-4
SPID(S): TP0001411E07
M4ID: 30204588

HILL MODEL (in red):

	tp	ga	gw
val:	1.61e-09	-1.04	0.684
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

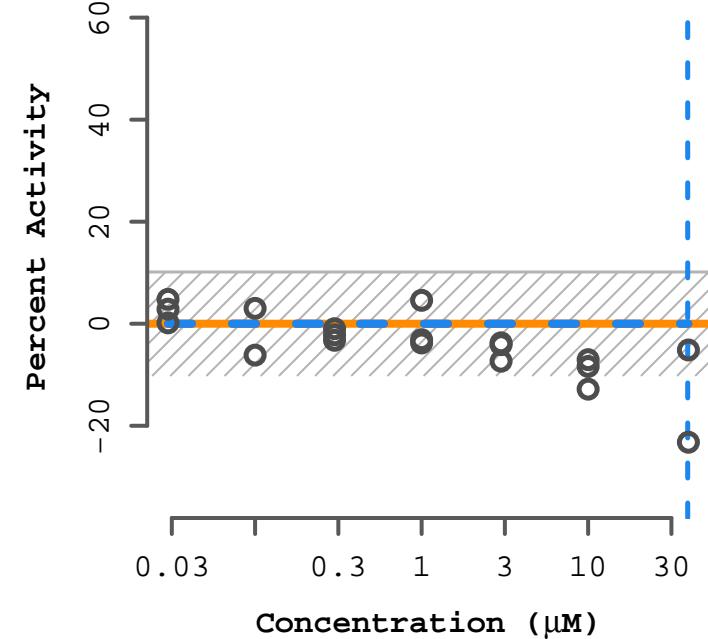
	tp	ga	gw	la	lw
val:	5.97e-10	-1.04	1.7	-0.682	5.84
sd:	NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
142.84	148.84	152.84
0.95	0.05	0.01
6.78	6.78	6.78

MAX_MEAN: 1.39 MAX_MED: 1.67 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Daidzein

CHID: 22310 CASRN: 486-66-8

SPID(S): TP0001414E07

M4ID: 30204594

HILL MODEL (in red):

tp	ga	gw
2.2e-08	2.02	2.89
NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

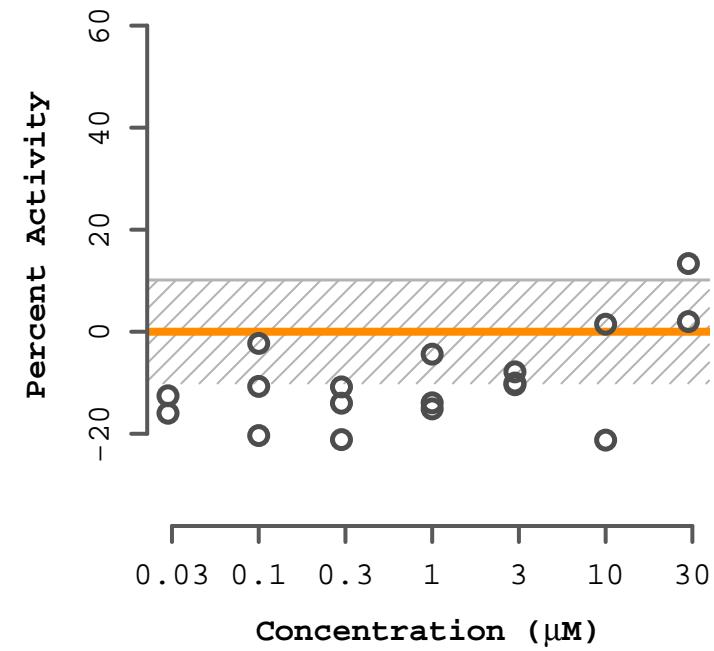
tp	ga	gw	la	lw
5.71e-09	1.6	7.97	1.89	1.7
0.00056	4830	1270000	29700	87100

CNST	HILL	GNLS
130.15	136.15	140.15
0.95	0.05	0.01
7.52	7.52	7.52

MAX_MEAN: 2.61 MAX_MED: 2.86 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

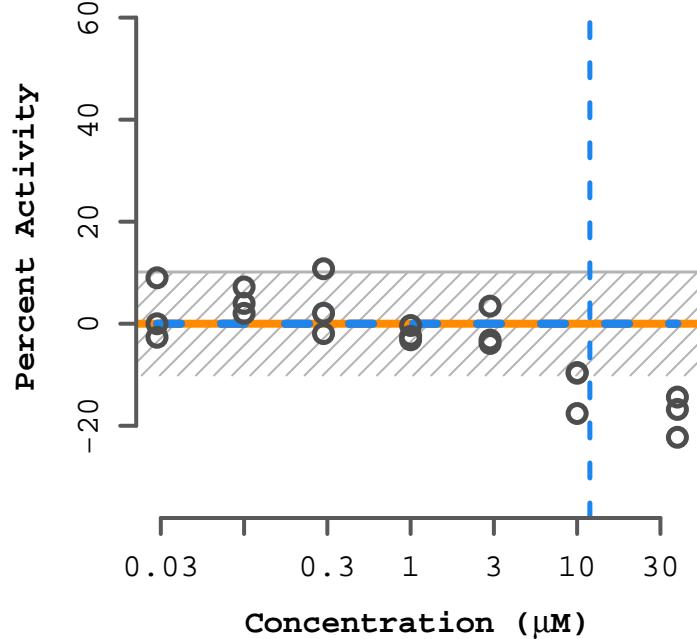
NAME: 2,2'-Dihydroxy-4-methoxybenzophenone
CHID: 22403 CASRN: 131-53-3
SPID(S): EPAPLT0154C02
M4ID: 30204175

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST	HILL	GNLS
AIC: 148.48	NA	NA
PROB: 1	NA	NA
RMSE: 13.03	NA	NA
MAX_MEAN: 7.68	MAX_MED: 7.68	BMAD: 3.38
COFF: 10.1	HIT-CALL: 0	FITC: 7
ACTP: 0		

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Phenolphthalin
CHID: 22439 CASRN: 81-90-3
SPID(S): TP0001411C08
M4ID: 30204539

HILL MODEL (in red):

	tp	ga	gw
val:	0.0809	2.09	7.98
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

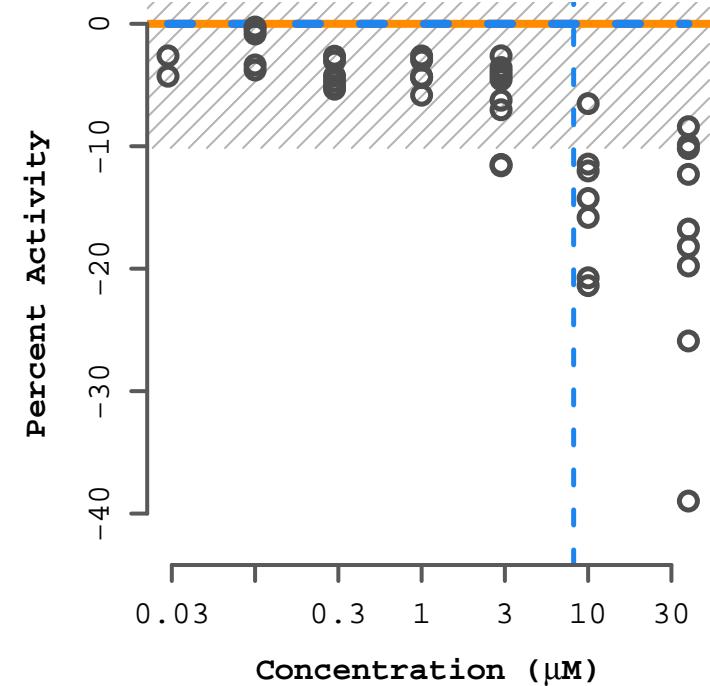
	tp	ga	gw	la	lw
val:	1.06e-08	1.08	0.939	1.35	7.06
sd:	0.000176	6540	29300	5390	363000

	CNST	HILL	GNLS
AIC:	155.28	161.28	165.28
PROB:	0.95	0.05	0.01
RMSE:	9.3	9.3	9.3

MAX_MEAN: 4.41 MAX_MED: 3.99 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 6-Thioguanine

CHID: 23652 CASRN: 154-42-7

SPID(S): TP0001414C03

M4ID: 30204538

HILL MODEL (in red):

	tp	ga	gw
val:	2.91e-09	2	1.74
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

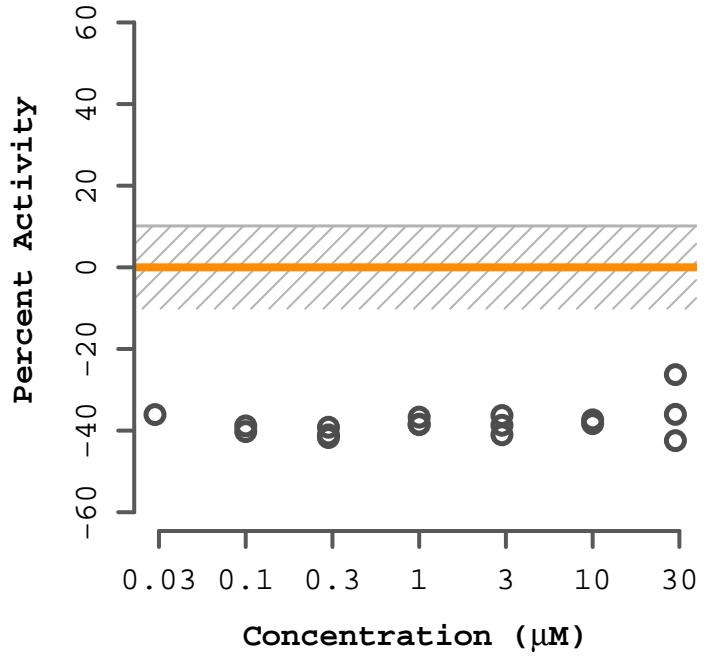
	tp	ga	gw	la	lw
val:	3.06e-10	0.913	4.4	1.23	3.29
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	450.44	456.44	460.44
PROB:	0.95	0.05	0.01
RMSE:	10.2	10.2	10.2

MAX_MEAN: 2.63 MAX_MED: 2.26 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Anthracene

CHID: 23878 CASRN: 120-12-7

SPID(S): EPAPLT0167C05

M4ID: 30204238

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST

AIC: 191.34

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 38.22

NA

NA

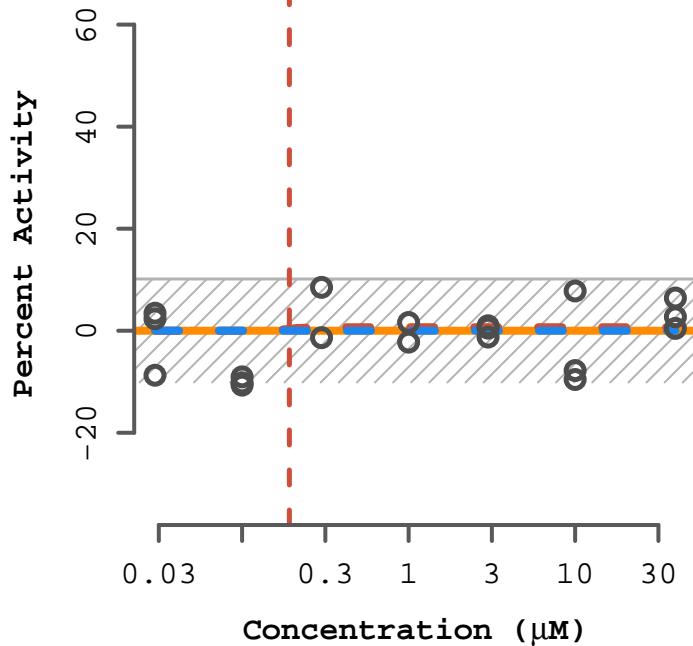
MAX_MEAN: -34.9

MAX_MED: -36

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diisopropyl methylphosphonate
CHID: 24051 CASRN: 1445-75-6
SPID(S): TP0001411D12
M4ID: 30204533

HILL MODEL (in red):

	tp	ga	gw
val:	0.689	-0.716	8
sd:	1.68	0.793	26.4

GAIN-LOSS MODEL (in blue):

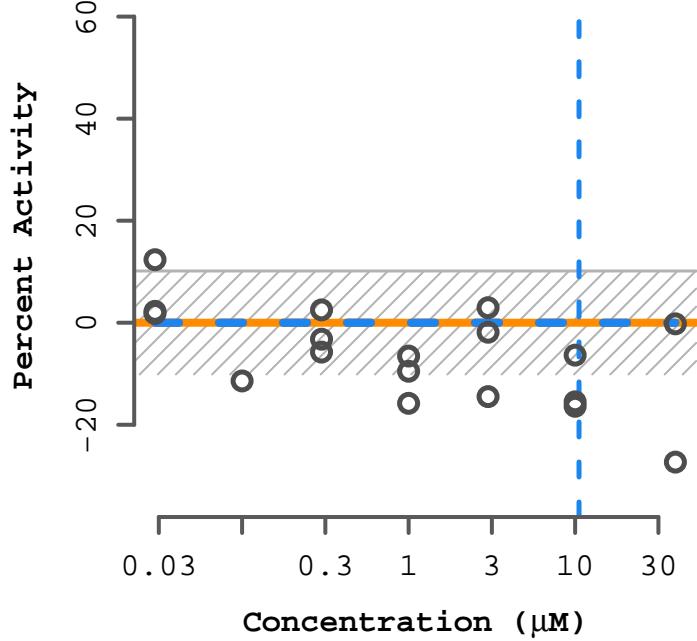
	tp	ga	gw	la	lw
val:	4.82	-2.52	2.1	-1.78	8.23
sd:	3880	2640	5720	51.9	838

CNST	HILL	GNLS
AIC: 127.67	133.5	137.67
PROB: 0.94	0.05	0.01
RMSE: 6.24	6.23	6.24

MAX_MEAN: 3.58 MAX_MED: 3.58 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.06

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,2-Dinitrobenzene
CHID: 24066 CASRN: 528-29-0
SPID(S): TP0001411B09
M4ID: 30204485

HILL MODEL (in red):

tp	ga	gw
7.23e-09	1.83	1.75
NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

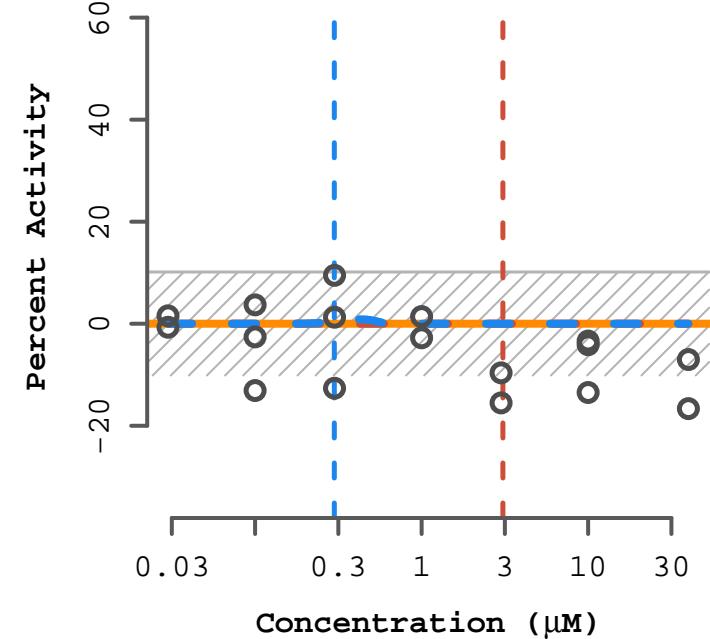
tp	ga	gw	la	lw
3.06e-08	1.02	5.1	1.28	3.88
0.000559	532	118000	2330	86900

CNST	HILL	GNLS
140.62	146.62	150.62
0.95	0.05	0.01
11.15	11.15	11.15

MAX_MEAN: 5.47 MAX_MED: 2.23 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Isoxaben

CHID: 24159 CASRN: 82558-50-7

SPID(S): TP0001413E12

M4ID: 30204285

HILL MODEL (in red):

	tp	ga	gw
val:	3.22e-10	0.488	1.31
sd:	0.000156	82700	509000

GAIN-LOSS MODEL (in blue):

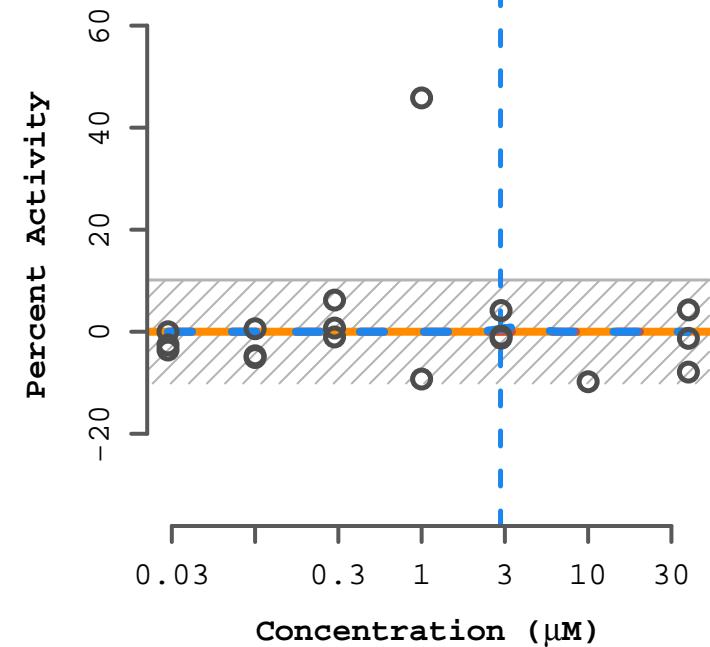
	tp	ga	gw	la	lw
val:	0.949	-0.524	7.89	-0.274	14.8
sd:	106	12.4	214	40.9	2210

CNST	HILL	GNLS
125.72	131.72	135.72
0.95	0.05	0.01
8.83	8.83	8.83

MAX_MEAN: 0.44 MAX_MED: 1.28 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Metalaxyl

CHID: 24175 CASRN: 57837-19-1

SPID(S): TP0001411B08

M4ID: 30204366

HILL MODEL (in red):

	tp	ga	gw
val:	6.53	2.1	7.95
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

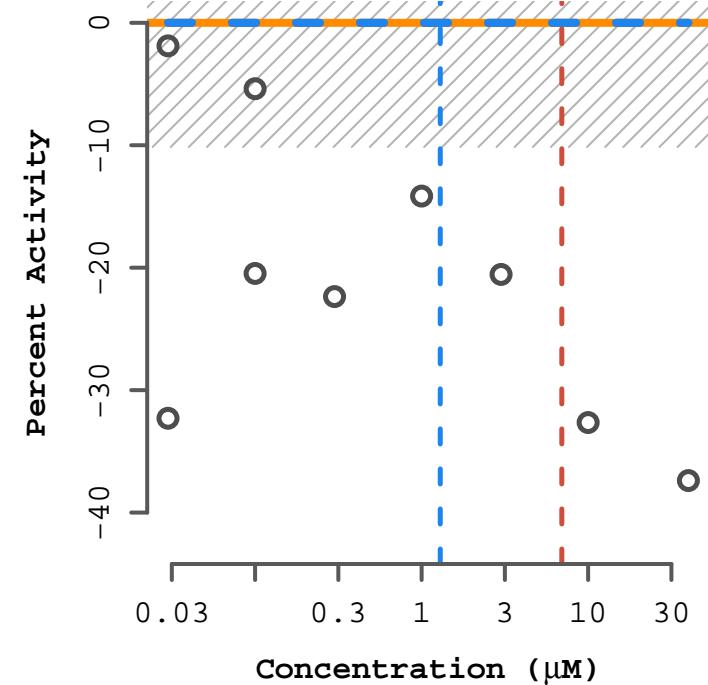
	tp	ga	gw	la	lw
val:	1.09	0.474	8	0.724	17.6
sd:	150	15.3	214	30.4	1940

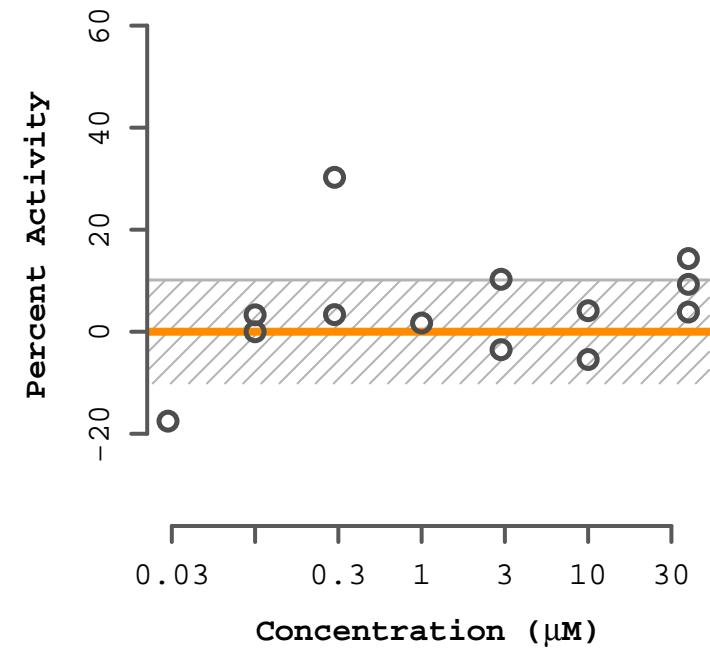
	CNST	HILL	GNLS
AIC:	126.21	132.21	136.17
PROB:	0.95	0.05	0.01
RMSE:	11.77	11.77	11.76

MAX_MEAN: 18.3 MAX_MED: 18.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS: 10





ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,2,4,5-Tetrachlorobenzene

CHID: 24320 CASRN: 95-94-3

SPID(S): TP0001411B03

M4ID: 30204557

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST

AIC: 101.11

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 11.5

NA

NA

MAX_MEAN: 16.8

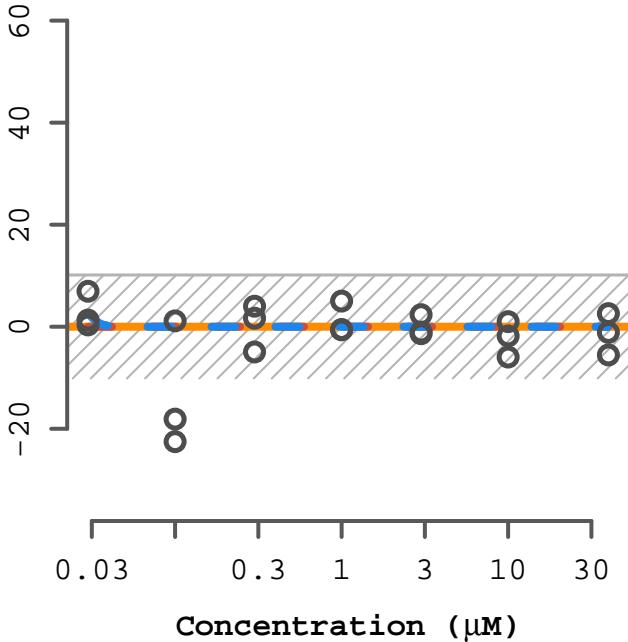
MAX_MED: 16.8

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,3-Dichloro-2-propanol
CHID: 25010 CASRN: 96-23-1
SPID(S): TP0001412G12
M4ID: 30204548

HILL MODEL (in red):

tp	ga	gw
4.75e-10	-1.81	1.81
3.24e-05	33800	275000

GAIN-LOSS MODEL (in blue):

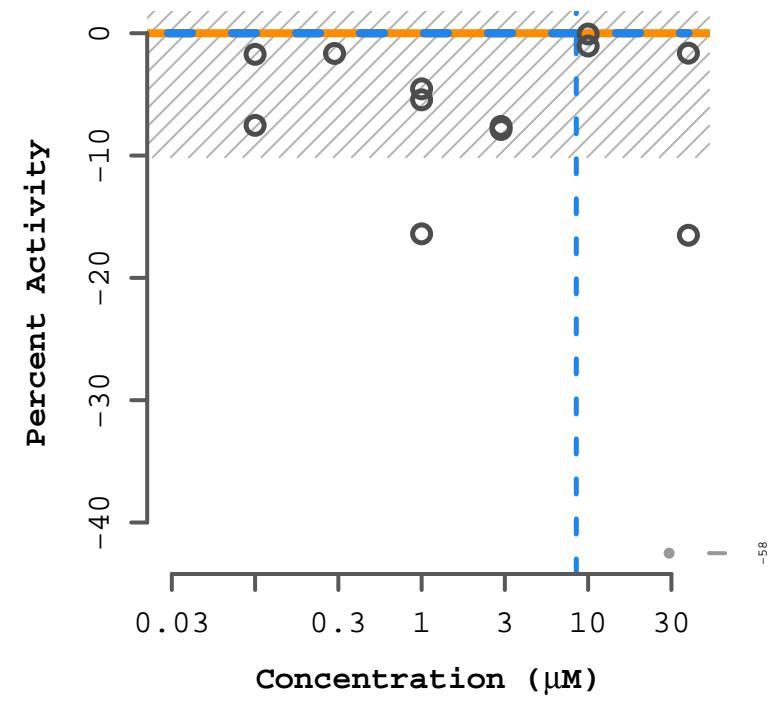
tp	ga	gw	la	lw
4.42	-2.27	2.02	-1.51	13
NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
134.82	140.82	143.57
0.94	0.05	0.01
7.05	7.05	6.96

MAX_MEAN: 2.88 MAX_MED: 1.68 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.06

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Chlorobenzotrifluoride

CHID: 27593 CASRN: 5216-25-1

SPID(S): TP0001414F06

M4ID: 30204586 BRK

HILL MODEL (in red):

tp	ga	gw
val: 0.0342	2.09	7.89
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 2.89e-08	0.929	4.5	1.27	5.98
sd: NaN	NaN	NaN	NaN	NaN

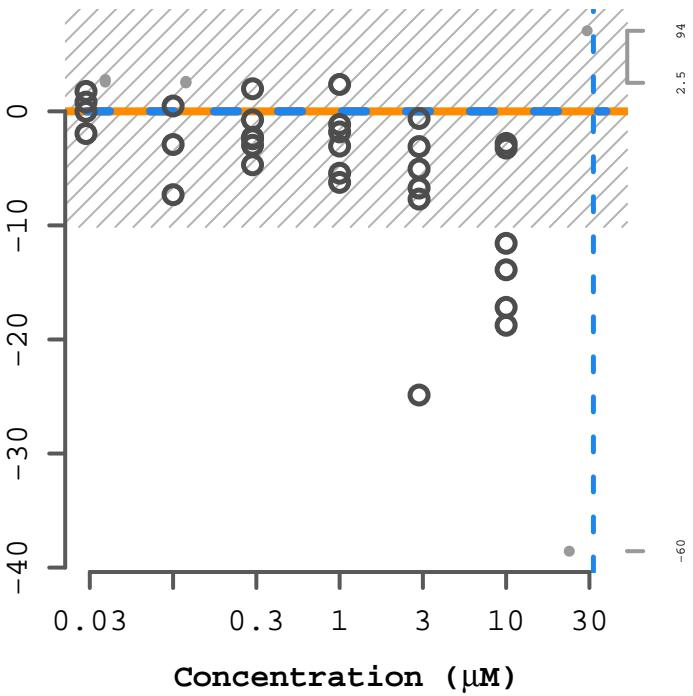
CNST	HILL	GNLS
AIC: 137.94	143.94	147.94
PROB: 0.95	0.05	0.01
RMSE: 15.31	15.31	15.31

MAX_MEAN: 3.88 MAX_MED: 1.79 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ketoconazole

CHID: 29879 CASRN: 65277-42-1

SPID(S): TP0001413H04

M4ID: 30204635 BRK

HILL MODEL (in red):

tp	ga	gw
val: 2.38	2.1	7.83
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

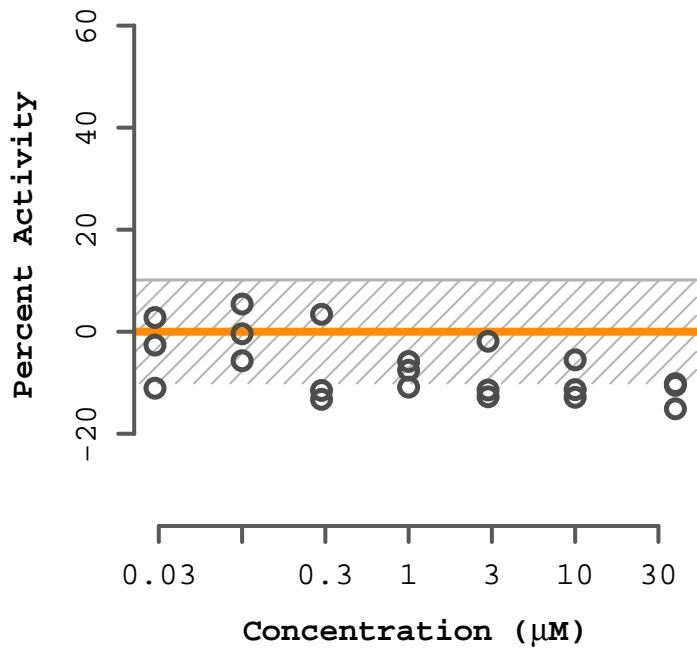
tp	ga	gw	la	lw
val: 6.43e-08	1.52	7.71	2.58	0.61
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 292.4	298.4	302.4
PROB: 0.95	0.05	0.01
RMSE: 19.52	19.52	19.52

MAX_MEAN: 16.7 MAX_MED: 16.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Chlorpyrifos-methyl
CHID: 32352 CASRN: 5598-13-0
SPID(S): TP0001412E12
M4ID: 30204300

HILL MODEL: Failed to converge.

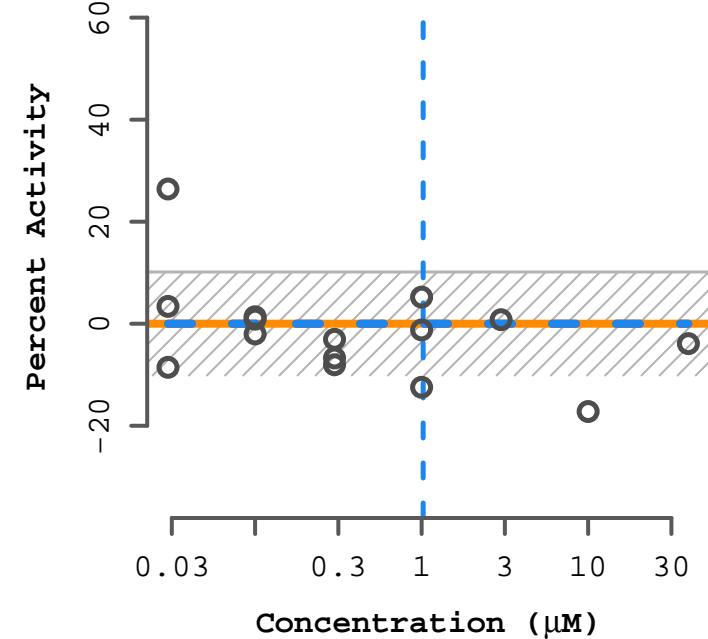
GAIN-LOSS MODEL: Failed to converge.

CNST	HILL	GNLS
AIC: 158.53	NA	NA
PROB: 1	NA	NA
RMSE: 9.23	NA	NA

MAX_MEAN: -0.214 MAX_MED: -0.392 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fenbuconazole

CHID: 32548 CASRN: 114369-43-6

SPID(S): TP0001411A05

M4ID: 30204352

HILL MODEL (in red):

tp	ga	gw
val: 8.39e-08	2.08	2.48
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 2.33e-08	0.00937	4.09	0.375	2.03
sd: NaN	NaN	NaN	NaN	NaN

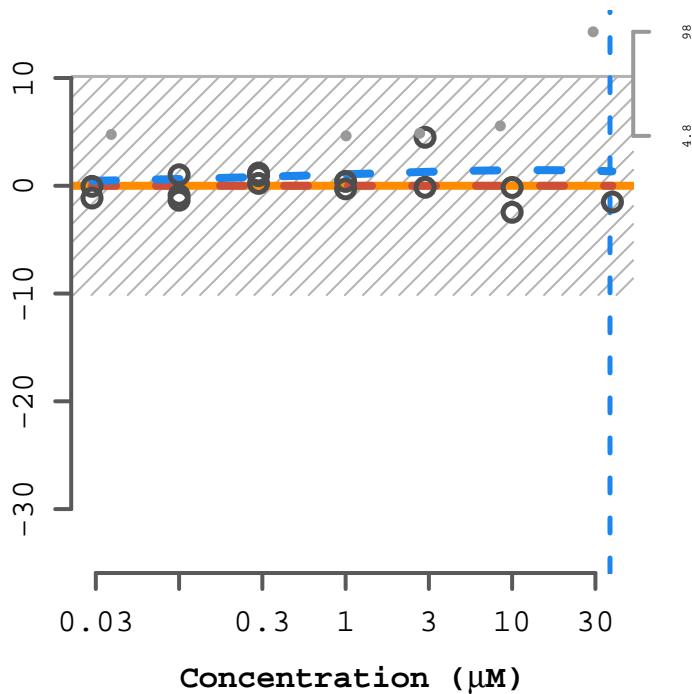
CNST	HILL	GNLS
AIC: 110.68	116.68	120.68
PROB: 0.95	0.05	0.01
RMSE: 9.67	9.67	9.67

MAX_MEAN: 7.09 MAX_MED: 3.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Flufenacet

CHID: 32552 CASRN: 142459-58-3

SPID(S): TP0001414A07

M4ID: 30204311 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	2.55	2.1	7.95
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

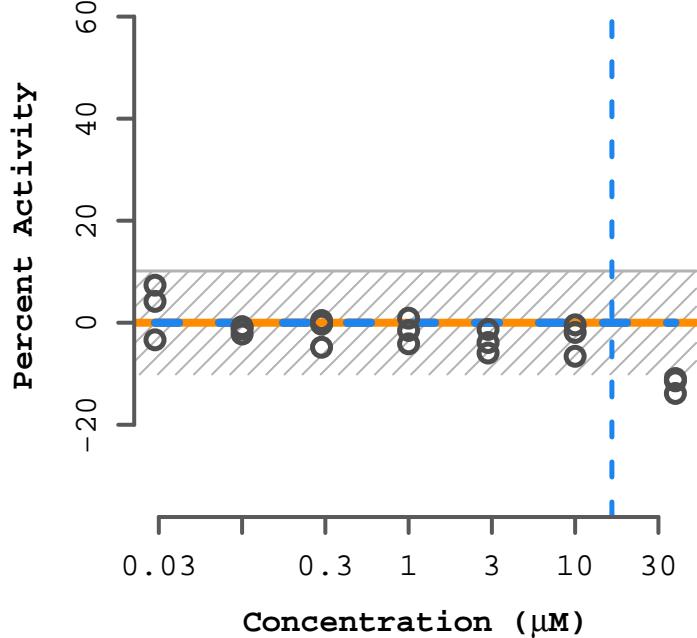
	tp	ga	gw	la	lw
val:	4.41	1.59	0.3	1.84	0.78
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	136.91	142.91	145.45
PROB:	0.94	0.05	0.01
RMSE:	22.35	22.35	21.99

MAX_MEAN: 48.4 MAX_MED: 48.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.06

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Flumetsulam

CHID: 32615 CASRN: 98967-40-9

SPID(S): TP0001412C07

M4ID: 30204379

HILL MODEL (in red):

tp	ga	gw
1.52e-06	2.1	5.66
NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

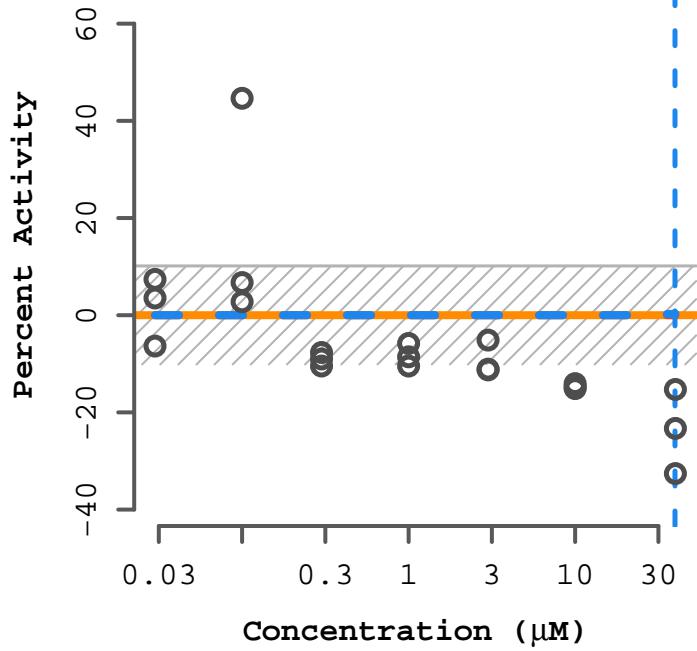
tp	ga	gw	la	lw
4.52e-08	1.22	7.97	1.47	18
NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
133.84	139.84	143.84
0.95	0.05	0.01
5.66	5.66	5.66

MAX_MEAN: 2.74 MAX_MED: 4.19 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Spirodiclofen

CHID: 34928 CASRN: 148477-71-8

SPID(S): TP0001412F02

M4ID: 30204316

HILL MODEL (in red):

tp	ga	gw
val: 0.219	2.1	7.98
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

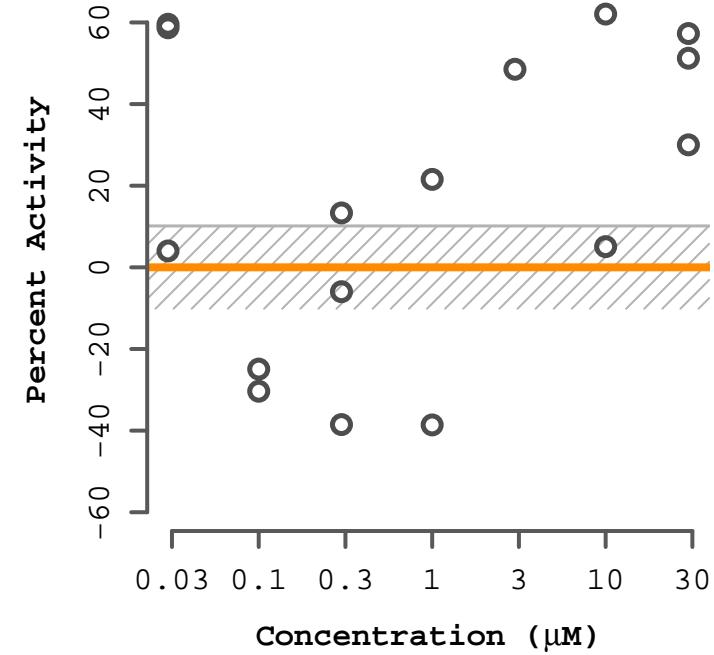
tp	ga	gw	la	lw
val: 0.815	1.6	8	1.85	2.85
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 177.28	183.28	187.48
PROB: 0.95	0.05	0.01
RMSE: 16	16	16.07

MAX_MEAN: 18 MAX_MED: 6.73 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Allethrin

CHID: 35180 CASRN: 584-79-2

SPID(S): EPAPLT0167B05

M4ID: 30204269

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST

HILL

GNLS

AIC: 167.74

NA

NA

PROB:

1

NA

NA

RMSE: 39.72

NA

NA

MAX_MEAN: 48.5

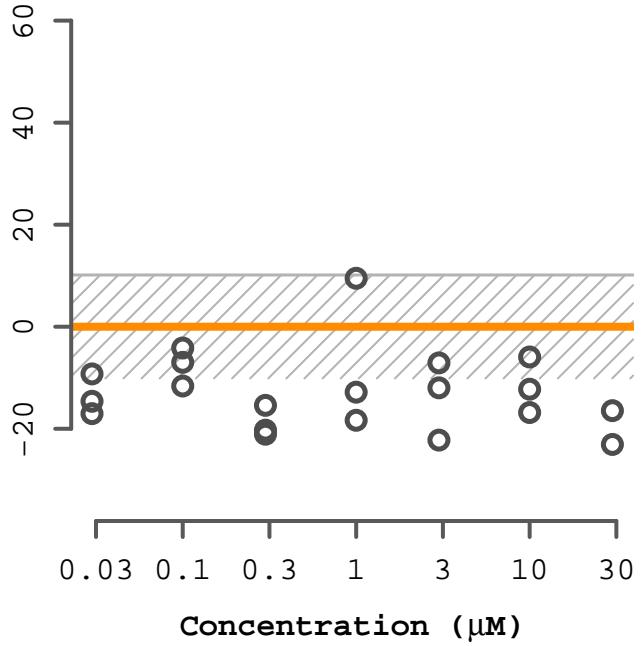
MAX_MED: 58.7

BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0

FLAGS: 8; 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pregnenolone

CHID: 36541 CASRN: 145-13-1

SPID(S): EPAPLT0154B08

M4ID: 30204170

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST

AIC: 170.54

HILL

NA

GNLS

NA

PROB: 1

NA

NA

RMSE: 14.88

NA

NA

MAX_MEAN: -7.22

MAX_MED: -6.98

BMAD: 3.38

COFF: 10.1

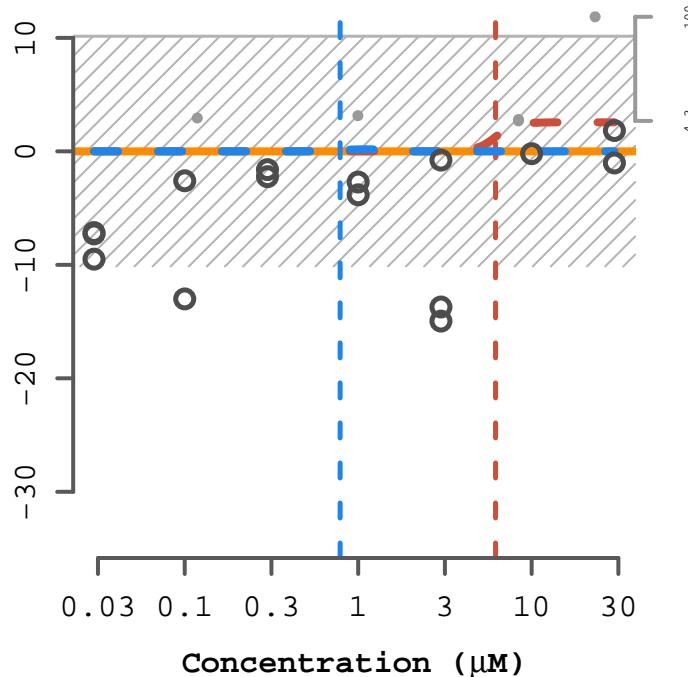
HIT-CALL: 0

FITC: 7

ACTP: 0

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: (3Z)-Hex-3-en-1-yl salicylate

CHID: 37750 CASRN: 65405-77-8

SPID(S): EPAPLT0154D05

M4ID: 30204188 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	2.56	0.792	8
sd:	3.06	0.734	21.9

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	0.199	-0.104	7.98	0.146	14.8
sd:	18.8	31.5	548	56.2	2560

CNST

AIC: 158.55

PROB: 0.93

RMSE: 23.62

HILL

AIC: 163.84

PROB: 0.07

RMSE: 23.05

GNLS

AIC: 168.55

PROB: 0.01

RMSE: 23.62

MAX_MEAN: 33.9

MAX_MED: 4.33

BMAD: 3.38

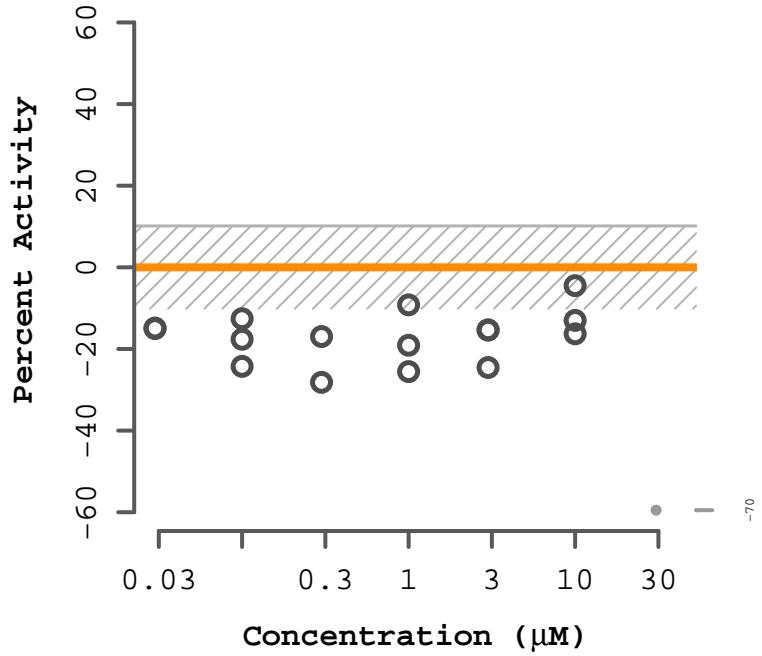
COFF: 10.1

HIT-CALL: 0

FITC: 7

ACTP: 0.07

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

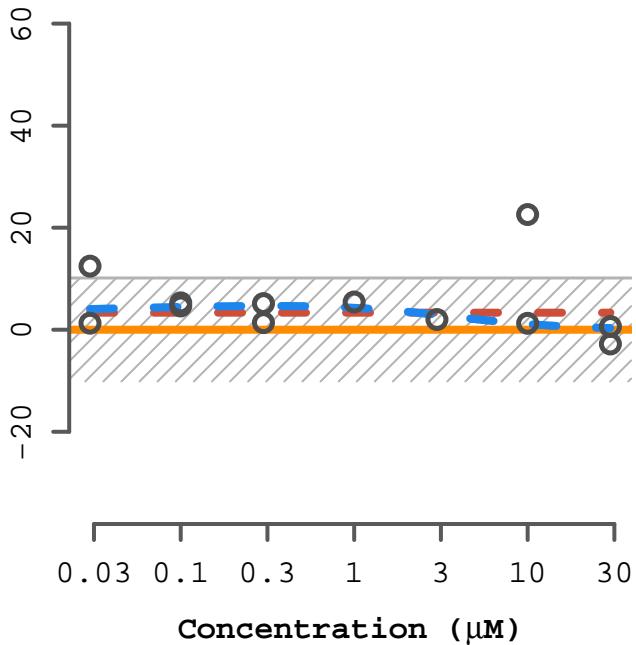
NAME: Chlorpyrifos oxon
CHID: 38666 CASRN: 5598-15-2
SPID(S): TP0001411D01
M4ID: 30204361 BRK

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST	HILL	GNLS
AIC: 158.57	NA	NA
PROB: 1	NA	NA
RMSE: 24.41	NA	NA
MAX_MEAN: -11.2	MAX_MED: -13	BMAD: 3.38
COFF: 10.1	HIT-CALL: 0	FITC: 7
ACTP: 0		
FLAGS: 10		

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Hexyl salicylate

CHID: 38924 CASRN: 6259-76-3

SPID(S): EPAPLT0154H08

M4ID: 30204231

HILL MODEL (in red):

tp	ga	gw
3.34	-2.52	6.28
1.27	357	2240

GAIN-LOSS MODEL (in blue):

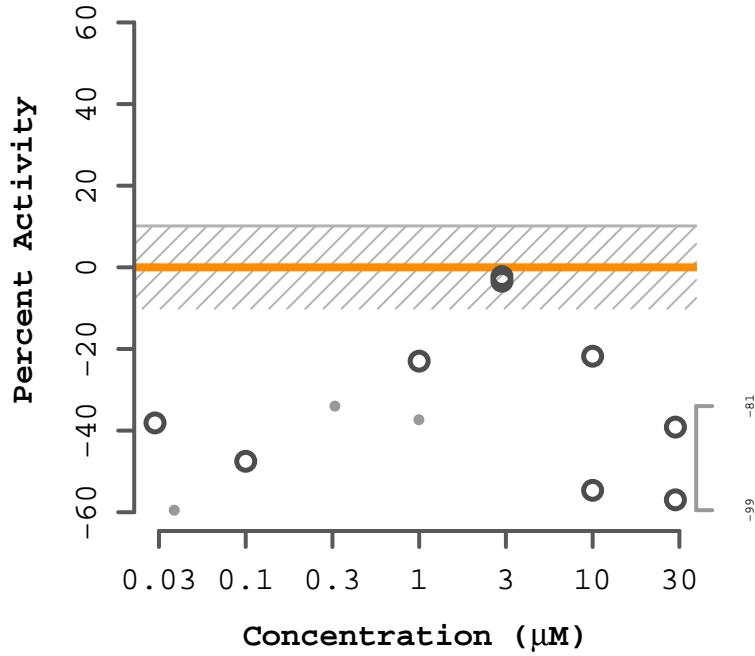
tp	ga	gw	la	lw
6.09	-2.52	0.3	0.531	1.29
NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
83.42	83.44	84.33
0.38	0.38	0.24
8.11	6.62	6.85

MAX_MEAN: 11.9 MAX_MED: 11.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.62

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

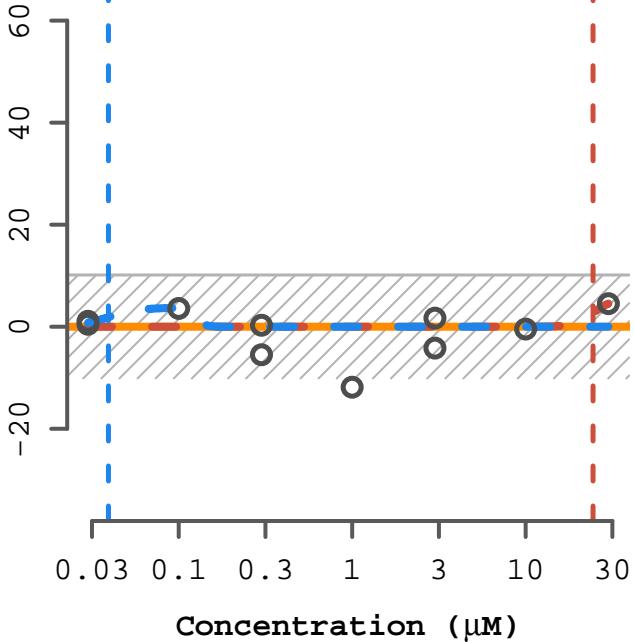
NAME: 2,6-Diaminopyridine
CHID: 40127 CASRN: 141-86-6
SPID(S): EPAPLT0154B02
M4ID: 30204164 BRK

HILL MODEL: Failed to converge.

GAIN-LOSS MODEL: Failed to converge.

CNST	HILL	GNLS
AIC: 133.58	NA	NA
PROB: 1	NA	NA
RMSE: 54.55	NA	NA
MAX_MEAN: -2.82	MAX_MED: -2.82	BMAD: 3.38
COFF: 10.1	HIT-CALL: 0	FITC: 7
ACTP: 0		
FLAGS: 10		

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4,4'-(Oxydiethylene)bis(morpholine)

CHID: 42170 CASRN: 6425-39-4

SPID(S): EPAPLT0154B04

M4ID: 30204166

HILL MODEL (in red):

	tp	ga	gw
val:	5.42	1.39	8
sd:	54.4	2.89	71

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	3.74	-1.41	4.9	-0.907	16.1
sd:	NaN	NaN	NaN	NaN	NaN

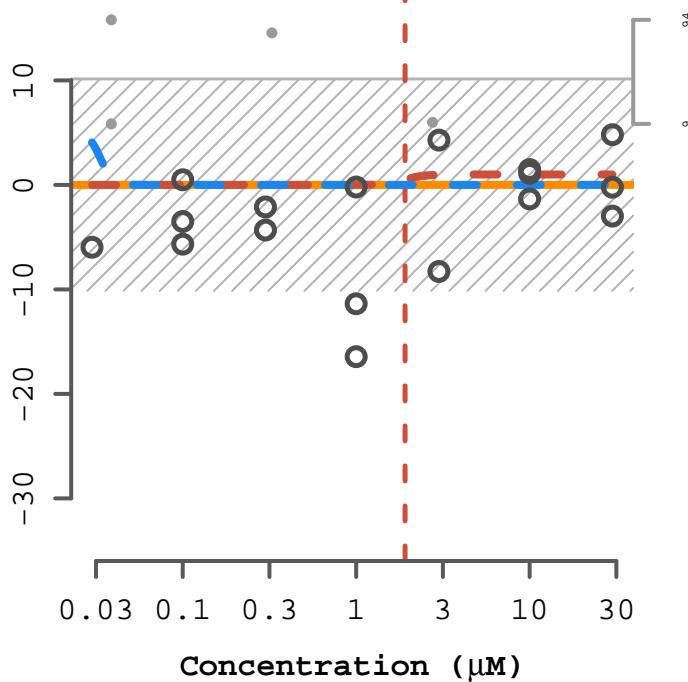
	CNST	HILL	GNLS
AIC:	60.72	64.69	69.25
PROB:	0.87	0.12	0.01
RMSE:	4.75	4.53	4.6

MAX_MEAN: 4.53 MAX_MED: 4.53 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.13

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Triethoxymethylsilane
CHID: 42490 CASRN: 2031-67-6
SPID(S): EPAPLT0154E09
M4ID: 30204203 BRK

HILL MODEL (in red):

tp	ga	gw
val: 0.995	0.282	8
sd: 2.62	0.947	31.4

GAIN-LOSS MODEL (in blue):

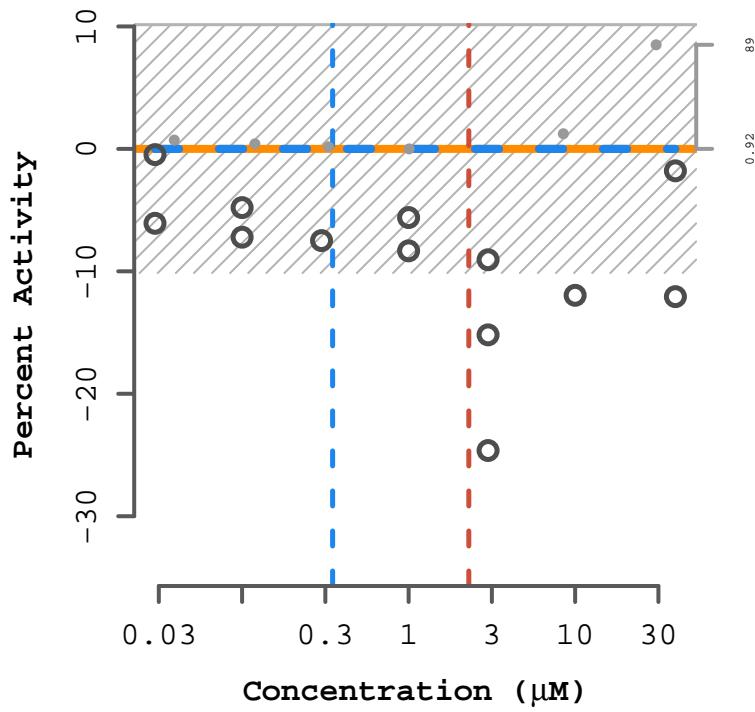
tp	ga	gw	la	lw
val: 5.56	-2.07	2	-1.48	12.4
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 178	183.85	187.66
PROB: 0.94	0.05	0.01
RMSE: 28.16	28.15	27.53

MAX_MEAN: 32.4 MAX_MED: 9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.06

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Zamifenacin

CHID: 47257 CASRN: 127308-82-1

SPID(S): TP0001414E10

M4ID: 30204493 BRK

HILL MODEL (in red):

tp	ga	gw
val: 4.29e-09	0.361	1.22
sd: 0.00016	1770	75900

GAIN-LOSS MODEL (in blue):

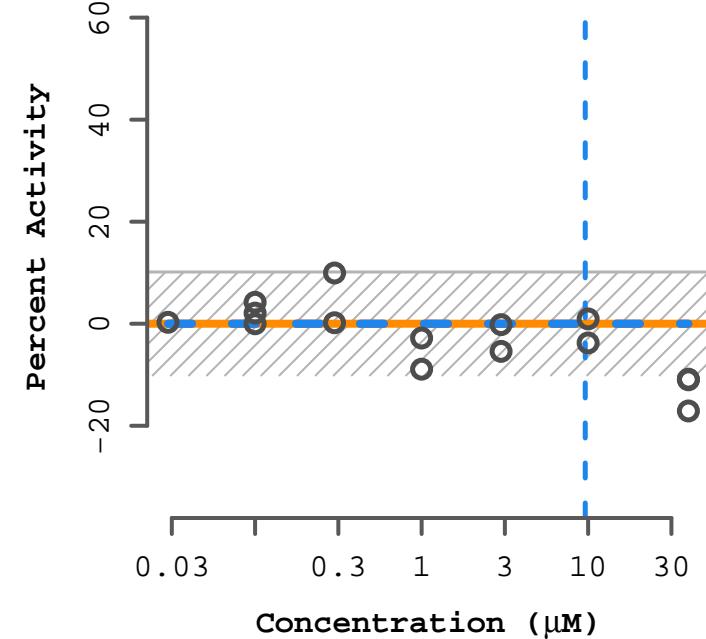
tp	ga	gw	la	lw
val: 4.58e-09	-0.456	2.27	-0.0832	5.77
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 159.3	165.3	169.3
PROB: 0.95	0.05	0.01
RMSE: 22.69	22.69	22.69

MAX_MEAN: 25.2 MAX_MED: 0.927 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: CI-1029

CHID: 47262 CASRN: 207736-05-8

SPID(S): TP0001414E09

M4ID: 30204495

HILL MODEL (in red):

tp	ga	gw
val: 1.62e-06	2.08	4.81
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

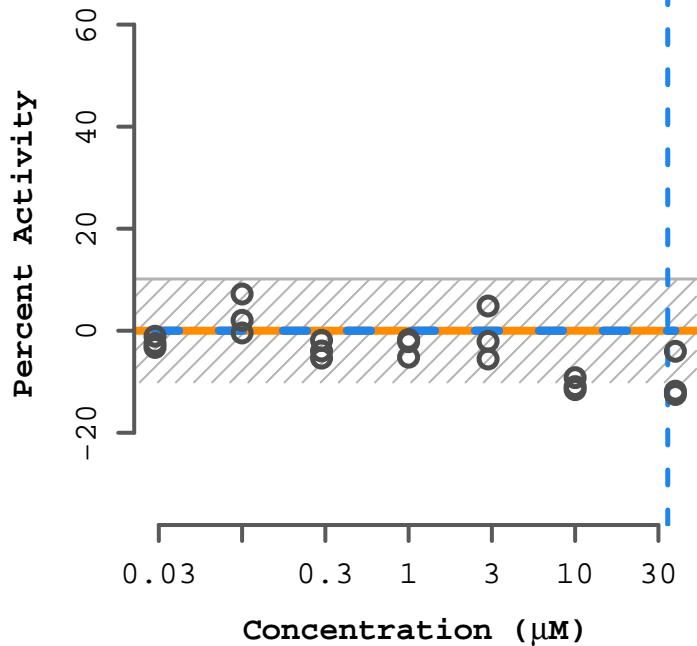
tp	ga	gw	la	lw
val: 1.89e-09	0.983	6.22	1.44	7.48
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 103.92	109.92	113.92
PROB: 0.95	0.05	0.01
RMSE: 7.23	7.23	7.23

MAX_MEAN: 5.07 MAX_MED: 5.07 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SB281832

CHID: 47324 CASRN: 219790-72-4

SPID(S): TP0001414D09

M4ID: 30204512

HILL MODEL (in red):

tp	ga	gw
val: 1.73e-08	2.08	2.11
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

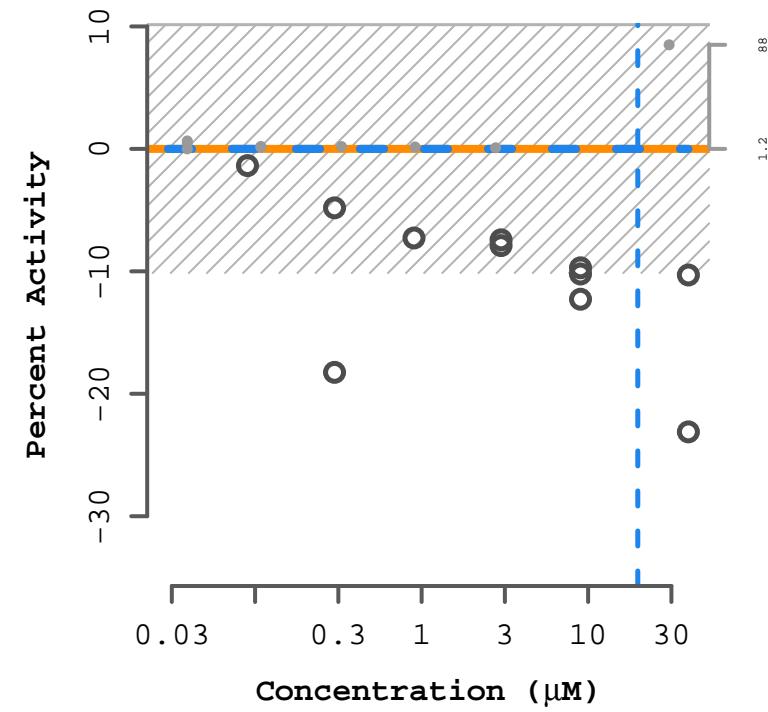
tp	ga	gw	la	lw
val: 1.17e-08	1.56	5.28	1.81	0.352
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 141.21	147.21	151.21
PROB: 0.95	0.05	0.01
RMSE: 6.42	6.42	6.42

MAX_MEAN: 2.94 MAX_MED: 2.05 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SR125047

CHID: 47342 CASRN: NOCAS_47342

SPID(S): TP0001412H10

M4ID: 30204517 BRK

HILL MODEL (in red):

tp	ga	gw
3.6e-09	1.98	3.05
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
4.84e-08	1.3	4.53	1.55	3.1
sd: NaN	NaN	NaN	NaN	NaN

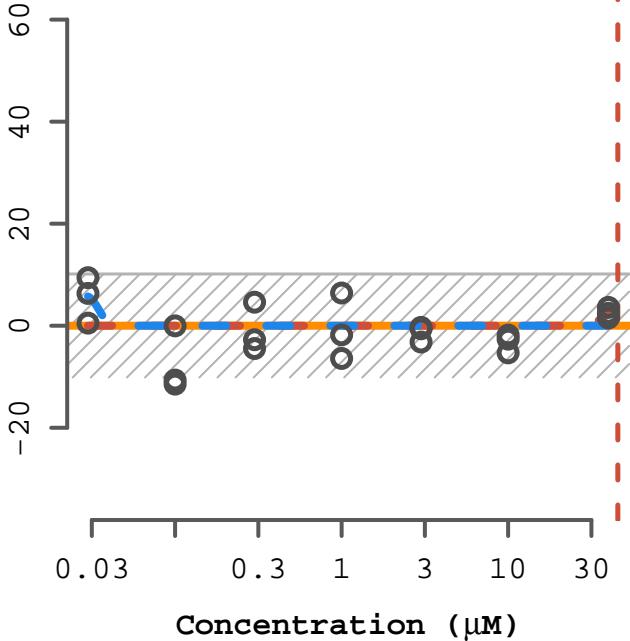
CNST	HILL	GNLS
164.47	170.47	174.47
PROB: 0.95	0.05	0.01
RMSE: 21.75	21.75	21.75

MAX_MEAN: 18.3 MAX_MED: 6.23 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 7 ACTP: 0.05

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Benzyl alcohol

CHID: 20152 CASRN: 100-51-6

SPID(S): TP0001411C02

M4ID: 30204390

HILL MODEL (in red):

tp	ga	gw
9.55	1.66	7.85
NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
6.75	-2.4	1.4	-1.46	15
NaN	NaN	NaN	NaN	NaN

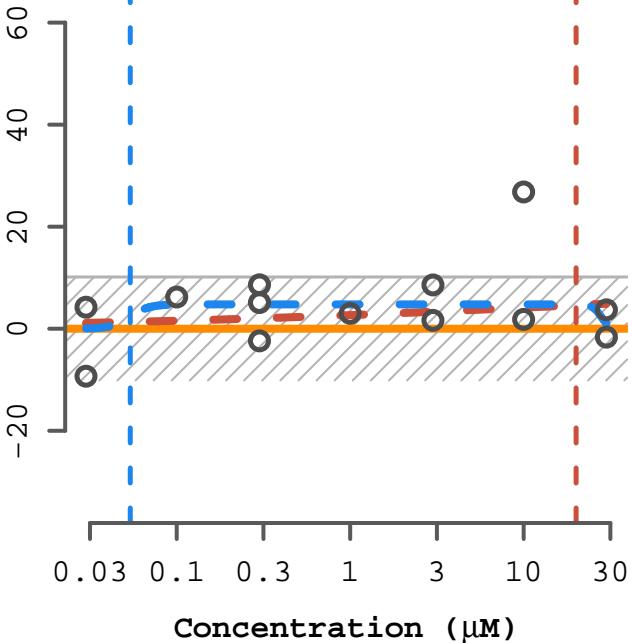
CNST	HILL	GNLS
132.21	136.96	138.3
0.88	0.08	0.04
5.25	5.16	4.83

MAX_MEAN: 5.43 MAX_MED: 6.34 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 9 ACTP: 0.12

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: N-Butyldiethanolamine
CHID: 44354 CASRN: 102-79-4
SPID(S): EPAPLT0154D06
M4ID: 30204189

HILL MODEL (in red):

tp	ga	gw
val: 9.23	1.3	0.3
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

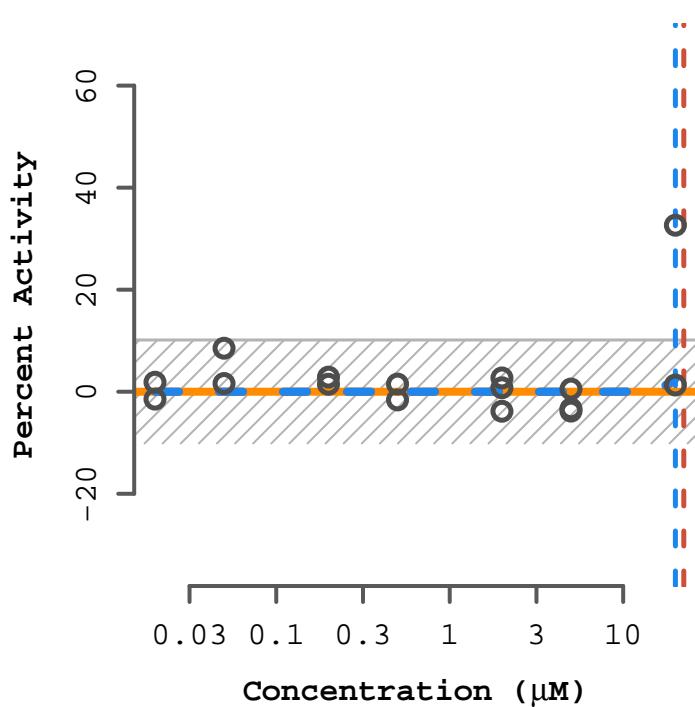
tp	ga	gw	la	lw
val: 4.77	-1.27	8	1.45	17.9
sd: 1.77	0.716	21.9	0.969	561

CNST	HILL	GNLS
AIC: 93.41	96.12	97.32
PROB: 0.71	0.18	0.1
RMSE: 9.09	7.88	7.39

MAX_MEAN: 14.3 MAX_MED: 14.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 9 ACTP: 0.29

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Darbufelone mesylate
CHID: 47246 CASRN: 139340-56-0
SPID(S): TP0001414D02
M4ID: 30204490

HILL MODEL (in red):

	tp	ga	gw
val:	8.81	1.35	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

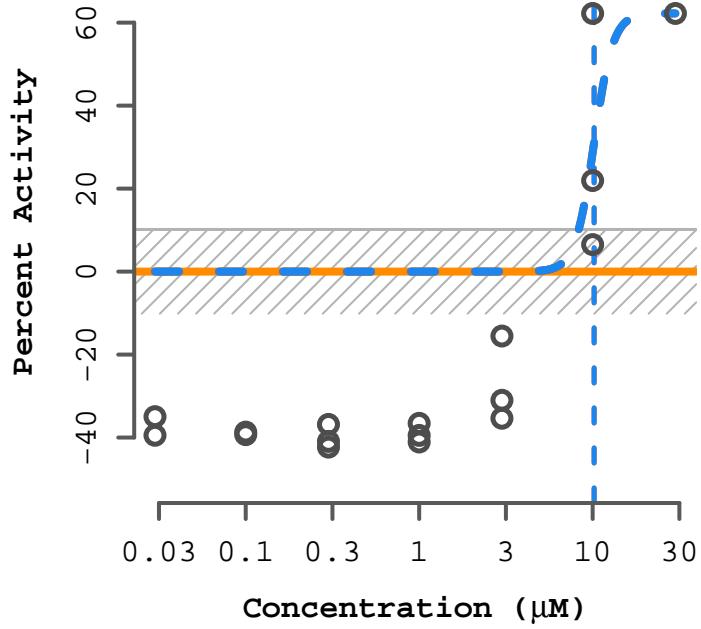
	tp	ga	gw	la	lw
val:	5.57	1.3	8	3.23	0.546
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	100.26	105.44	109.44
PROB:	0.92	0.07	0.01
RMSE:	8.7	8.11	8.11

MAX_MEAN: 17 MAX_MED: 17 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 9 ACTP: 0.08

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 5-Azacytidine

CHID: 20116 CASRN: 320-67-2

SPID(S): EPAPLT0167A06

M4ID: 30204247

HILL MODEL (in red):

tp	ga	gw
val: 62.2	1.01	8
sd: 29.3	0.102	50.1

GAIN-LOSS MODEL (in blue):

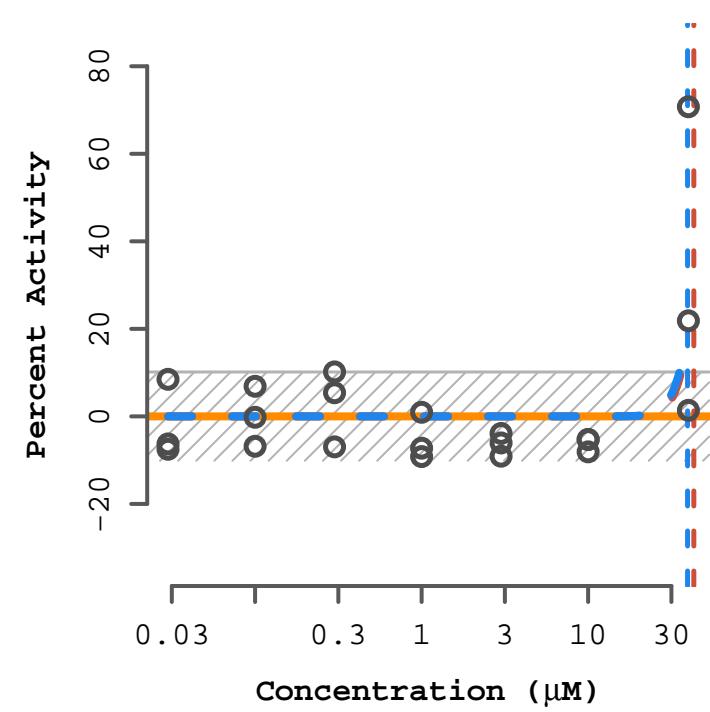
tp	ga	gw	la	lw
val: 62.3	1.01	8	2.82	2.32
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 178.22	179.45	183.45
PROB: 0.62	0.34	0.05
RMSE: 39.06	33.72	33.72

MAX_MEAN: 62.2 MAX_MED: 62.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.38

FLAGS: 8; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Biphenyl

CHID: 20161 CASRN: 92-52-4

SPID(S): TP0001412F04

M4ID: 30204459

HILL MODEL (in red):

	tp	ga	gw
val:	51.2	1.63	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

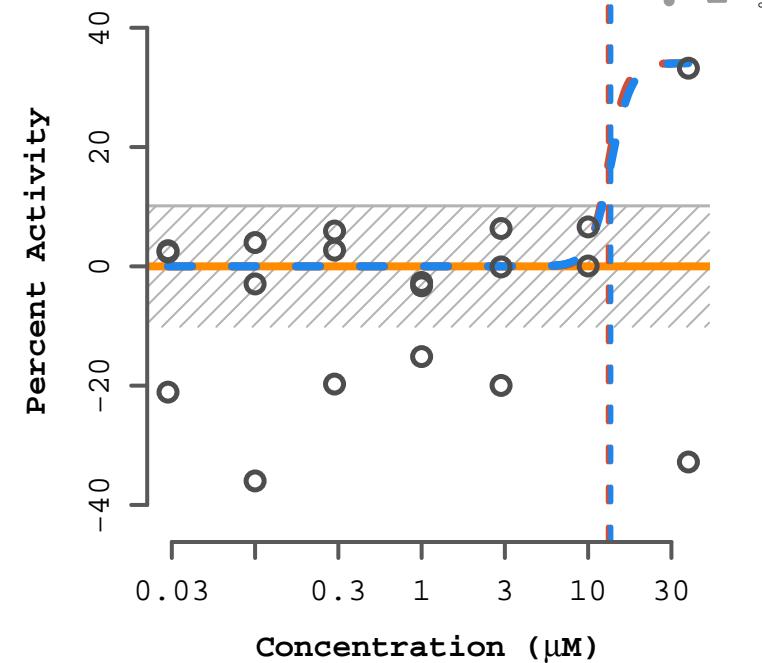
	tp	ga	gw	la	lw
val:	34.6	1.6	7.92	3.4	10.9
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	165.58	167.47	171.47
PROB:	0.69	0.27	0.04
RMSE:	17.35	13.63	13.63

MAX_MEAN: 31.3 MAX_MED: 21.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.31

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dinoseb

CHID: 20207 CASRN: 88-85-7

SPID(S): TP0001412C11

M4ID: 30204486 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	34.1	1.12	7.98
sd:	13.1	3.48	229

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	34.1	1.14	7.07	3.6	1.66
sd:	NaN	NaN	NaN	NaN	NaN

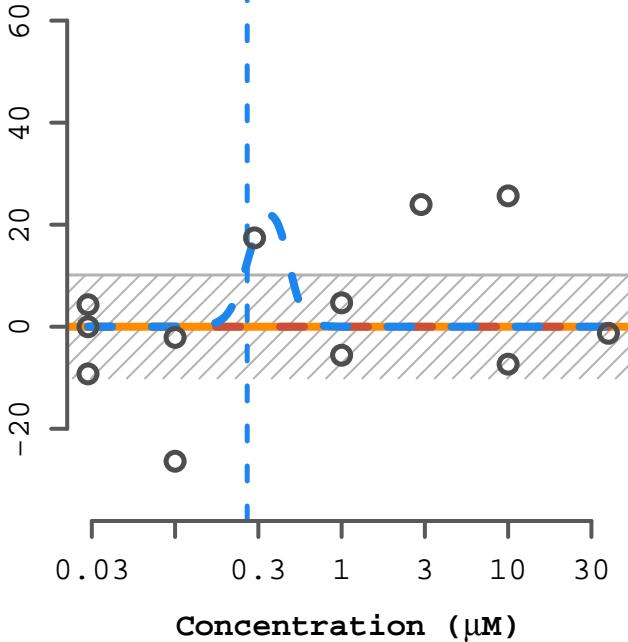
	CNST	HILL	GNLS
AIC:	181.49	184.12	188.12
PROB:	0.77	0.21	0.03
RMSE:	26.21	23.28	23.28

MAX_MEAN: 31.1 MAX_MED: 33.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.23

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Clofibrate

CHID: 20336 CASRN: 637-07-0

SPID(S): TP0001411A06

M4ID: 30204541

HILL MODEL (in red):

	tp	ga	gw
val:	21.4	2.1	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

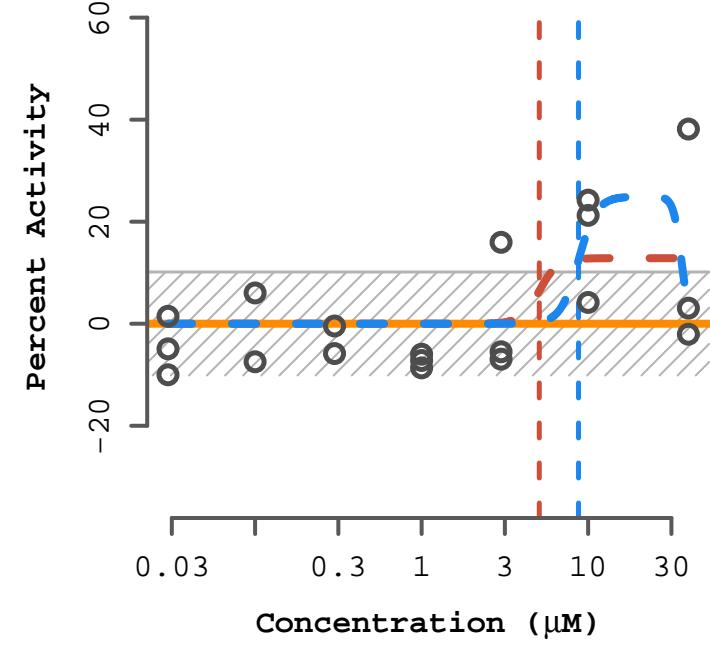
	tp	ga	gw	la	lw
val:	25.4	-0.567	8	-0.317	10
sd:	249	1.62	31.9	4.23	139

	CNST	HILL	GNLS
AIC:	100.28	106.28	107.68
PROB:	0.93	0.05	0.02
RMSE:	14.28	14.28	13.37

MAX_MEAN: 24 MAX_MED: 24 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.07

FLAGS: 10; 8



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dichlorprop

CHID: 20440 CASRN: 120-36-5

SPID(S): TP0001412G03

M4ID: 30204371

HILL MODEL (in red):

	tp	ga	gw
val:	12.8	0.707	8
sd:	6.14	0.442	13.5

GAIN-LOSS MODEL (in blue):

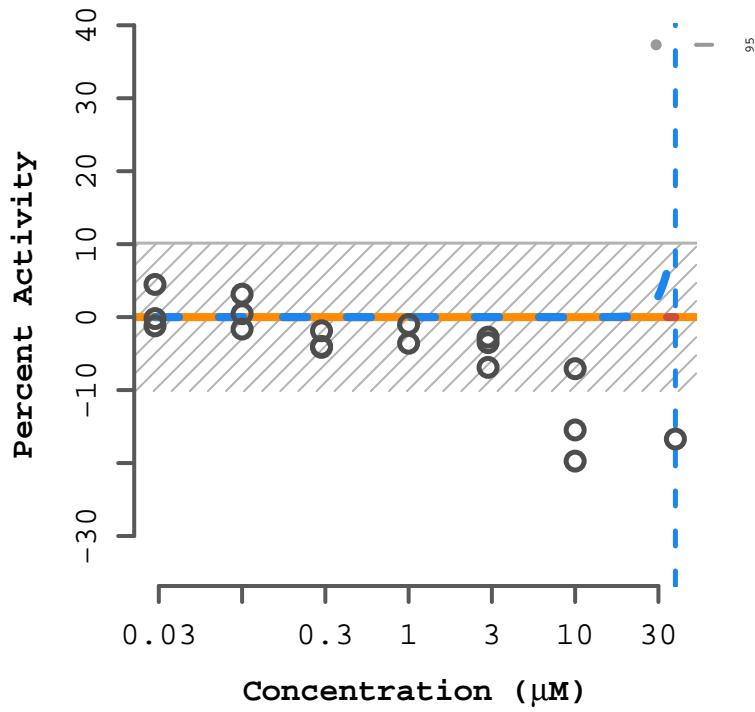
	tp	ga	gw	la	lw
val:	24.9	0.942	8	1.56	18
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	151.18	151.75	153.34
PROB:	0.48	0.36	0.16
RMSE:	13.13	10.21	10.8

MAX_MEAN: 16.6 MAX_MED: 21.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.52

FLAGS: 12; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Reserpine

CHID: 21237 CASRN: 50-55-5

SPID(S): TP0001413C12

M4ID: 30204666 BRK

HILL MODEL (in red):

tp ga gw

val: 2.08 2.1 7.97

sd: NaN NaN NaN

GAIN-LOSS MODEL (in blue):

tp ga gw la lw

val: 22 1.6 7.99 3.6 18

sd: NA NA NA NA NA

CNST

AIC: 151.11

PROB: 0.95

RMSE: 23.06

HILL

157.11

0.05

23.06

GNLS

163.45

0

21.31

MAX_MEAN: 39.1

MAX_MED: 39.1

BMAD: 3.38

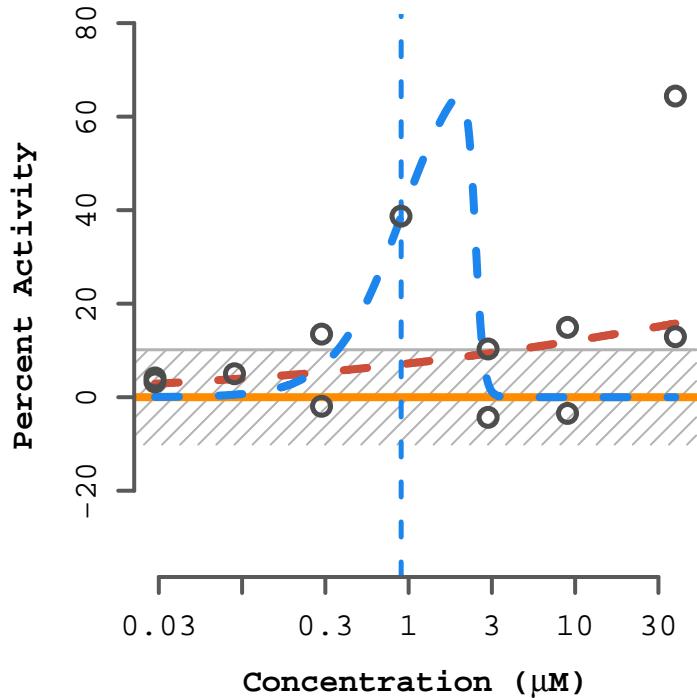
COFF: 10.1

HIT-CALL: 0

FITC: 10

ACTP: 0.05

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tamoxifen citrate

CHID: 21301 CASRN: 54965-24-1

SPID(S): TP0001411B01

M4ID: 30204537

HILL MODEL (in red):

	tp	ga	gw
val:	37.9	2.1	0.3
sd:	219	12.9	0.375

GAIN-LOSS MODEL (in blue):

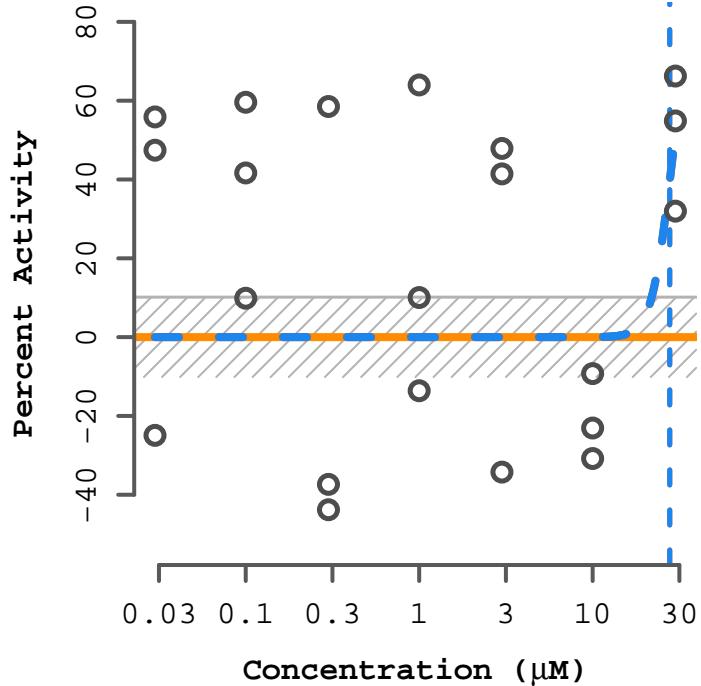
	tp	ga	gw	la	lw
val:	77.3	-0.0449	2.16	0.401	17.8
sd:	287	1.5	3.79	9.91	2300

CNST	HILL	GNLS
AIC: 107.85	108.56	110.21
PROB: 0.5	0.35	0.15
RMSE: 23.12	18.1	20.05

MAX_MEAN: 38.7 MAX_MED: 38.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.5

FLAGS: 12; 10; 8



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tetramethylthiuram monosulfide
 CHID: 21333 CASRN: 97-74-5
 SPID(S): EPAPLT0154C01
 M4ID: 30204174

HILL MODEL (in red):

	tp	ga	gw
val:	79.5	1.44	8
sd:	784	1.43	31.8

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	79.3	1.44	7.98	3.46	9.62
sd:	NA	NA	NA	NA	NA

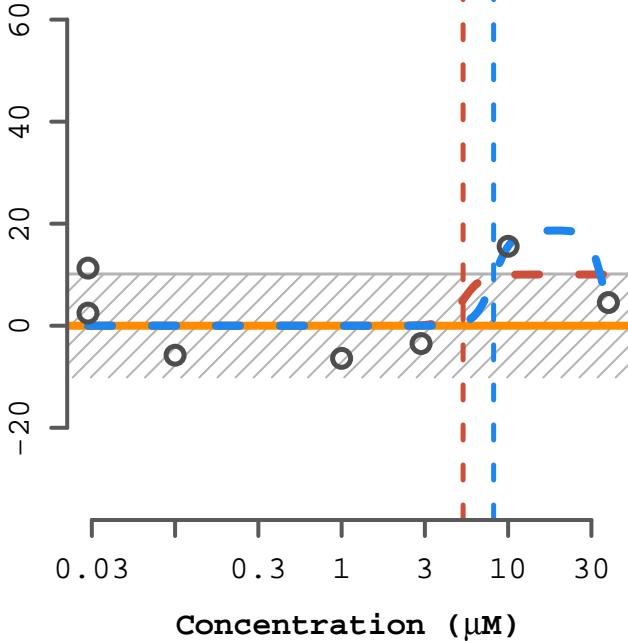
	CNST	HILL	GNLS
AIC:	222.7	223.27	227.27
PROB:	0.54	0.41	0.05
RMSE:	42.36	37.71	37.71

MAX_MEAN: 51 MAX_MED: 54.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.46

FLAGS: 8; 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dalapon

CHID: 21575 CASRN: 75-99-0

SPID(S): TP0001411A07

M4ID: 30204429

HILL MODEL (in red):

	tp	ga	gw
val:	10	0.729	8
sd:	5.17	0.363	11.1

GAIN-LOSS MODEL (in blue):

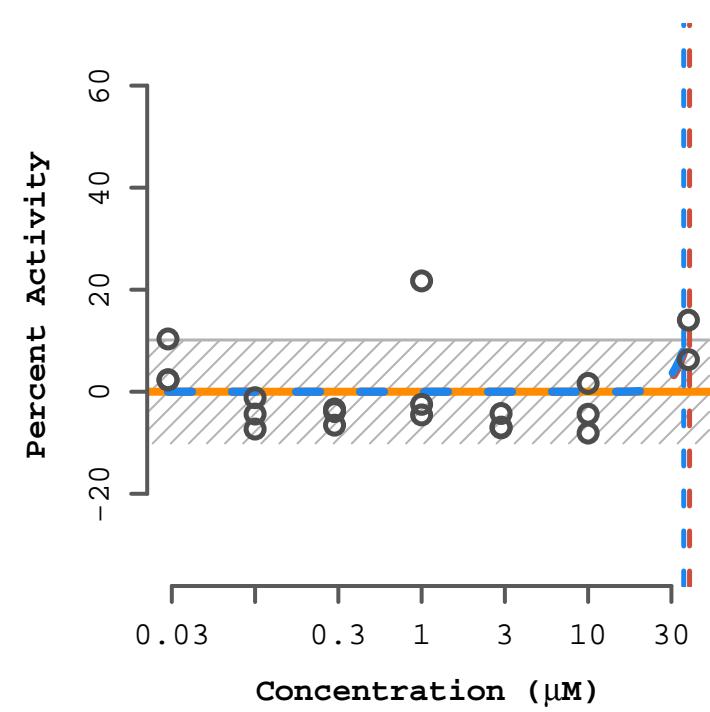
	tp	ga	gw	la	lw
val:	18.7	0.913	8	1.56	11.3
sd:	127	1.89	38.6	0.455	70

CNST	HILL	GNLS
AIC: 52.19	54.85	56.35
PROB: 0.72	0.19	0.09
RMSE: 8.31	6.35	5.61

MAX_MEAN: 15.6 MAX_MED: 15.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.28

FLAGS:



Percent Activity

Concentration (μM)

ASSAY:

AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME:

Pentanal

CHID:

21653 CASRN: 110-62-3

SPID(S):

TP0001412H05

M4ID:

30204653

HILL MODEL (in red):

tp

ga

gw

val:

21.7

1.61

8

sd:

NaN

NaN

NaN

GAIN-LOSS MODEL (in blue):

tp

ga

gw

la

lw

val:

16.7

1.57

8

3.58

0.774

sd:

NaN

NaN

NaN

NaN

NaN

CNST

HILL

GNLS

AIC:

140.28

141.46

145.46

PROB:

0.61

0.34

0.05

RMSE:

7.75

7.05

7.05

MAX_MEAN:

10.2

MAX_MED:

10.2

BMAD: 3.38

COFF:

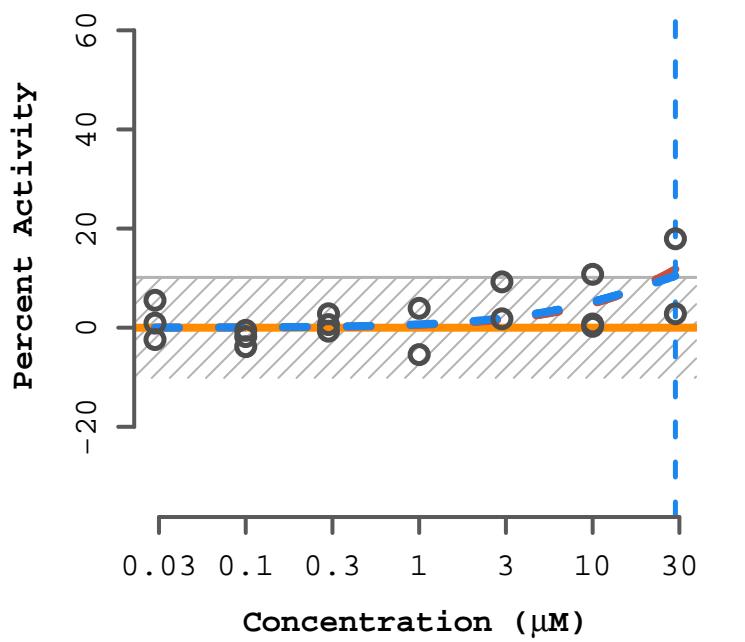
10.1

HIT-CALL: 0

FITC: 10

ACTP: 0.39

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Disulfoton

CHID: 22018 CASRN: 298-04-4

SPID(S): EPAPLT0167C04

M4ID: 30204249 BRK

HILL MODEL (in red):

tp	ga	gw
val: 47.6	1.98	0.965
sd: 212	2.62	1.01

GAIN-LOSS MODEL (in blue):

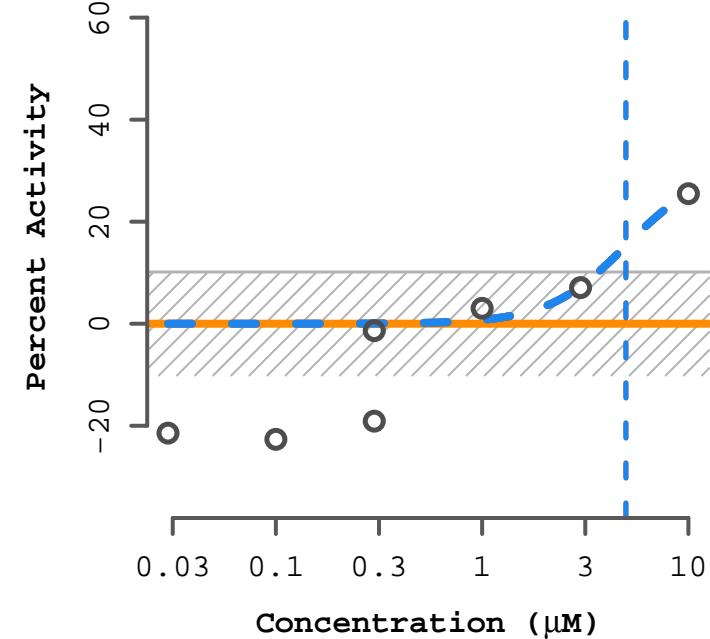
tp	ga	gw	la	lw
val: 20.9	1.48	1	2.28	17.8
sd: 90	2.51	1.28	272000	2050000

CNST	HILL	GNLS
AIC: 143.08	143.17	147.3
PROB: 0.48	0.46	0.06
RMSE: 23.48	20.51	20.8

MAX_MEAN: 40 MAX_MED: 17.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.52

FLAGS: 12; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bisphenol B

CHID: 22442 CASRN: 77-40-7

SPID(S): TP0001411B04

M4ID: 30204483

HILL MODEL (in red):

	tp	ga	gw
val:	30.6	0.697	2.29
sd:	52.7	1.1	6.12

GAIN-LOSS MODEL (in blue):

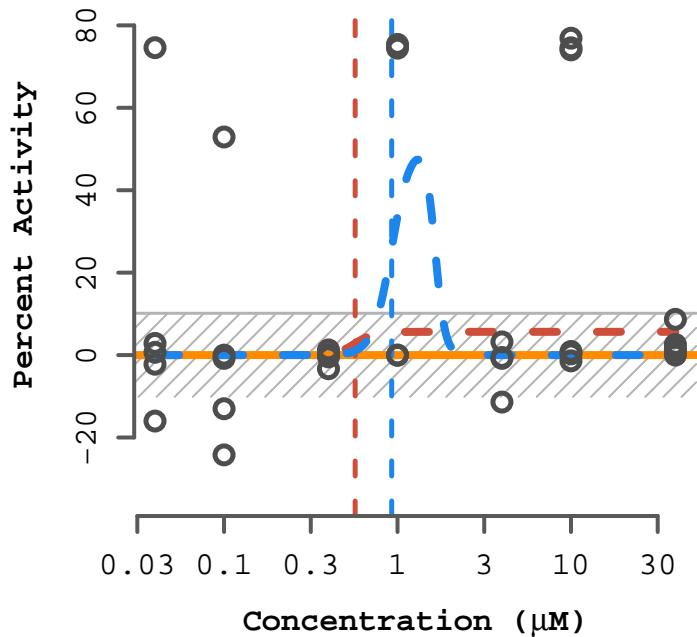
	tp	ga	gw	la	lw
val:	30.6	0.697	2.29	2.05	5.66
sd:	51.8	1.08	6	1820	9790

	CNST	HILL	GNLS
AIC:	62.7	65.14	69.14
PROB:	0.75	0.22	0.03
RMSE:	17.11	13.87	13.87

MAX_MEAN: 25.5 MAX_MED: 25.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.25

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Metformin

CHID: 23270 CASRN: 657-24-9

SPID(S): EPAPLT0167A02

M4ID: 30204262

HILL MODEL (in red):

tp	ga	gw
val: 5.64	-0.244	8
sd: 4.97	0.561	22.6

GAIN-LOSS MODEL (in blue):

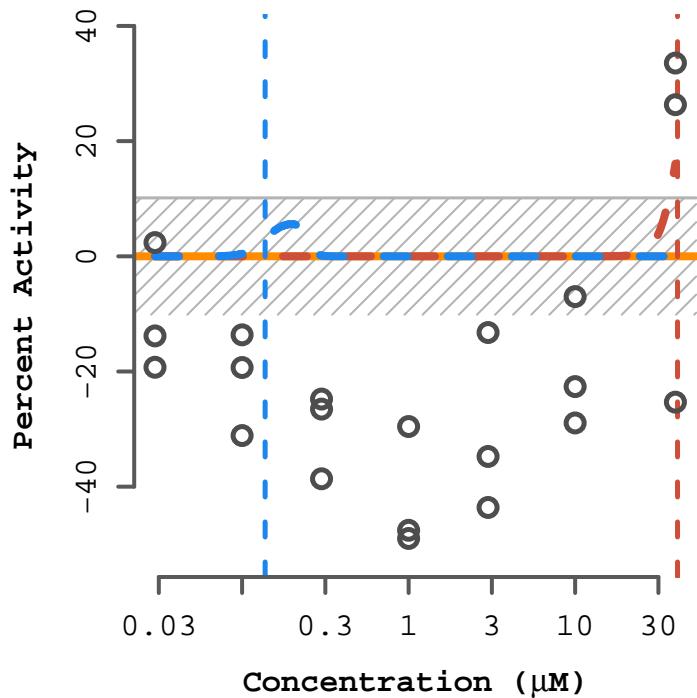
tp	ga	gw	la	lw
val: 51.9	-0.0335	8	0.217	15.3
sd: 813	2.27	53.6	20.7	918

CNST	HILL	GNLS
AIC: 317.75	322.17	325.75
PROB: 0.89	0.1	0.02
RMSE: 33.85	32.18	31.27

MAX_MEAN: 37.5 MAX_MED: 37.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.11

FLAGS: 8; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Heptachlor epoxide B
CHID: 24126 CASRN: 1024-57-3
SPID(S): TP0001412F11
M4ID: 30204615

HILL MODEL (in red):

	tp	ga	gw
val:	36.5	1.61	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

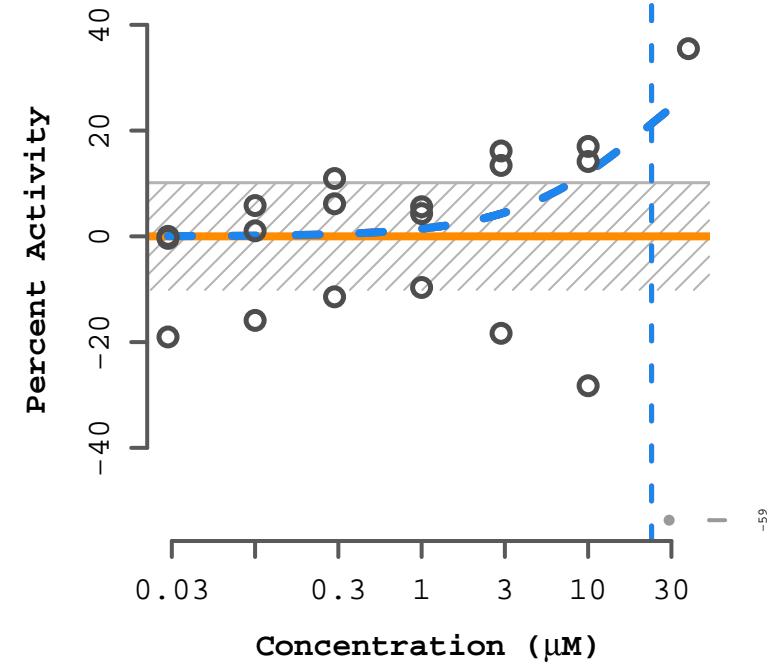
	tp	ga	gw	la	lw
val:	6.03	-0.862	8	-0.612	18
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	206.55	211.83	216.67
PROB:	0.93	0.07	0.01
RMSE:	28.93	28.66	29

MAX_MEAN: 11.5 MAX_MED: 26.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.07

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Lactofen

CHID: 24160 CASRN: 77501-63-4

SPID(S): TP0001412D02

M4ID: 30204337 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	42.6	1.38	1.05
sd:	60.9	1.2	1.09

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	42.6	1.38	1.05	2.76	8.14
sd:	60.9	1.2	1.09	15700	110000

CNST

AIC: 176.61

PROB: 0.85

RMSE: 19.89

HILL

AIC: 180.29

PROB: 0.13

RMSE: 23.41

GNLS

AIC: 184.29

PROB: 0.02

RMSE: 23.41

MAX_MEAN: 3.75

COFF: 10.1

FLAGS: 8; 10

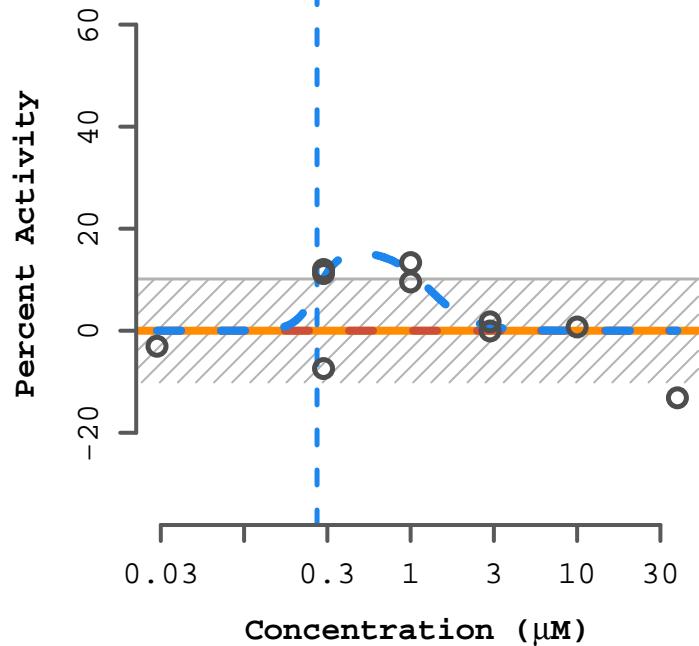
MAX_MED: 14.2

HIT-CALL: 0

FITC: 10

BMAD: 3.38

ACTP: 0.15



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Colchicine

CHID: 24845 CASRN: 64-86-8

SPID(S): TP0001411B02

M4ID: 30204596

HILL MODEL (in red):

	tp	ga	gw
val:	0.0203	2.1	7.66
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	16	-0.562	6.6	0.116	3.44
sd:	217	2.94	306	4.95	30.7

CNST

AIC: 75.41

PROB: 0.6

RMSE: 8.83

HILL

AIC: 81.41

PROB: 0.03

RMSE: 8.83

GNLS

AIC: 76.38

PROB: 0.37

RMSE: 7.12

MAX_MEAN: 11.5

MAX_MED: 11.5

BMAD: 3.38

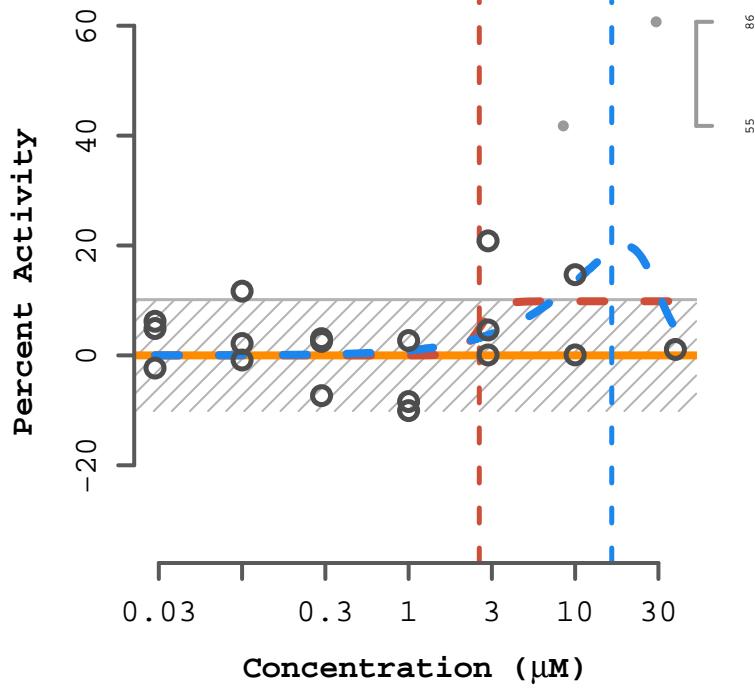
COFF: 10.1

HIT-CALL: 0

FITC: 10

ACTP: 0.4

FLAGS: 8



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Flutamide

CHID: 32004 CASRN: 13311-84-7

SPID(S): TP0001412F03

M4ID: 30204567 BRK

HILL MODEL (in red):

tp	ga	gw
val: 9.85	0.424	8
sd: 6.63	0.288	33.1

GAIN-LOSS MODEL (in blue):

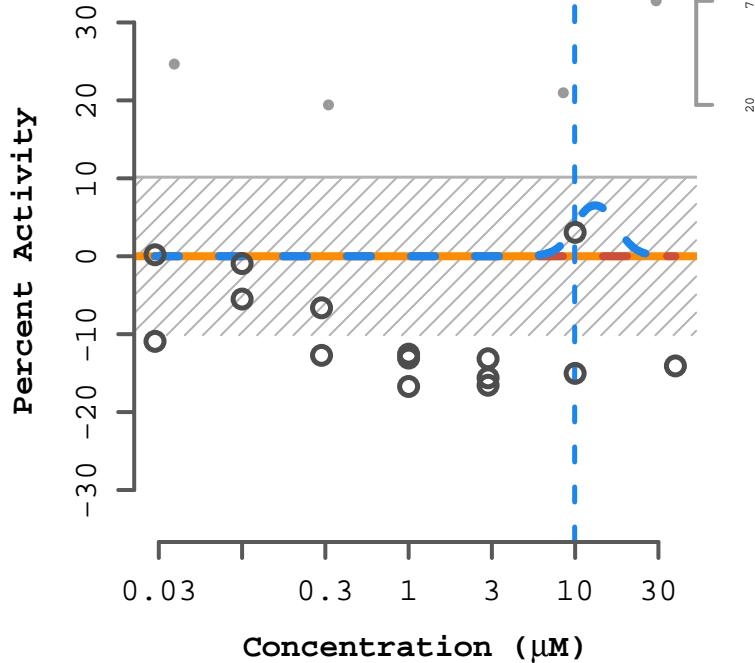
tp	ga	gw	la	lw
val: 39.8	1.22	1.33	1.47	5.42
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 170.01	171.74	176.04
PROB: 0.68	0.29	0.03
RMSE: 23.96	20.83	21.5

MAX_MEAN: 43.5 MAX_MED: 43.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.32

FLAGS: 10; 8



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Haloperidol

CHID: 34150 CASRN: 52-86-8

SPID(S): TP0001412F01

M4ID: 30204535 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	18.5	2.1	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

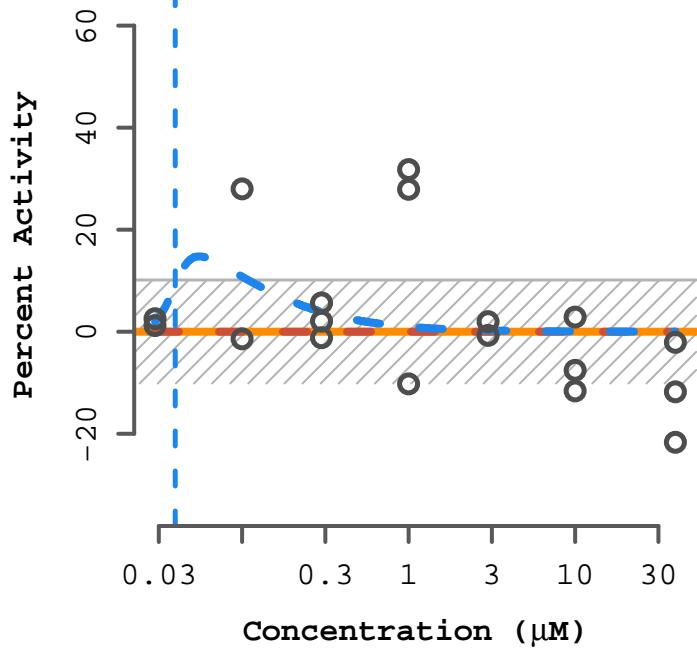
	tp	ga	gw	la	lw
val:	7.99	0.997	8	1.25	7.62
sd:	556	7.71	102	19	412

	CNST	HILL	GNLS
AIC:	172.74	178.74	182.61
PROB:	0.95	0.05	0.01
RMSE:	23.94	23.94	23.86

MAX_MEAN: 31.5 MAX_MED: 31.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.05

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diclosulam

CHID: 34528 CASRN: 145701-21-9

SPID(S): TP0001412E08

M4ID: 30204372

HILL MODEL (in red):

	tp	ga	gw
val:	0.0582	2.1	7.92
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

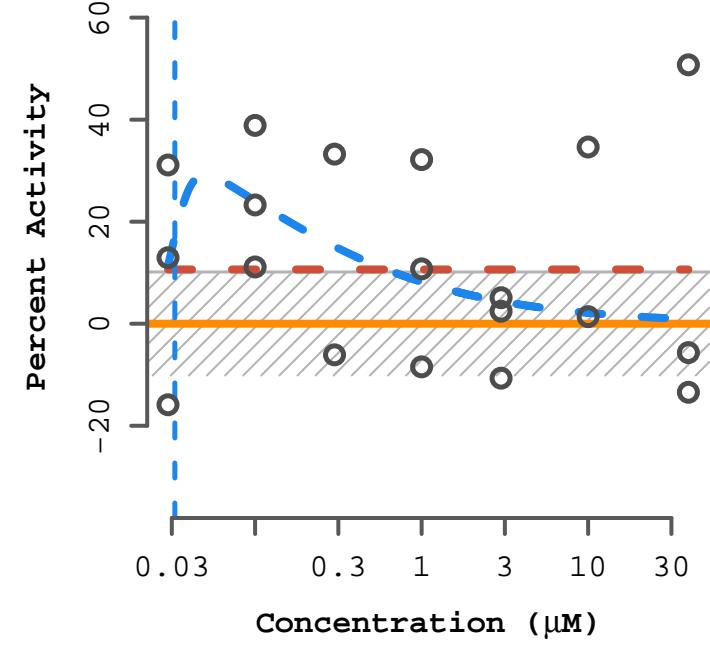
	tp	ga	gw	la	lw
val:	27.5	-1.4	8	-1.15	1.24
sd:	134	0.797	51.5	2.36	2.84

CNST	HILL	GNLS
AIC: 146.77	152.77	155.28
PROB: 0.94	0.05	0.01
RMSE: 14.03	14.03	13.12

MAX_MEAN: 16.5 MAX_MED: 27.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.06

FLAGS: 8; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Imazapyr

CHID: 34665 CASRN: 81334-34-1

SPID(S): TP0001412E06

M4ID: 30204302

HILL MODEL (in red):

	tp	ga	gw
val:	10.6	-2.46	5.6
sd:	5.09	481	2780

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	57.9	-1.48	7.59	-1.23	0.635
sd:	390	0.958	175	8.33	1.51

CNST

AIC: 176.3

PROB: 0.6

RMSE: 23.02

HILL

AIC: 177.93

PROB: 0.26

RMSE: 19.7

GNLS

AIC: 179.23

PROB: 0.14

RMSE: 19.44

MAX_MEAN: 24.4

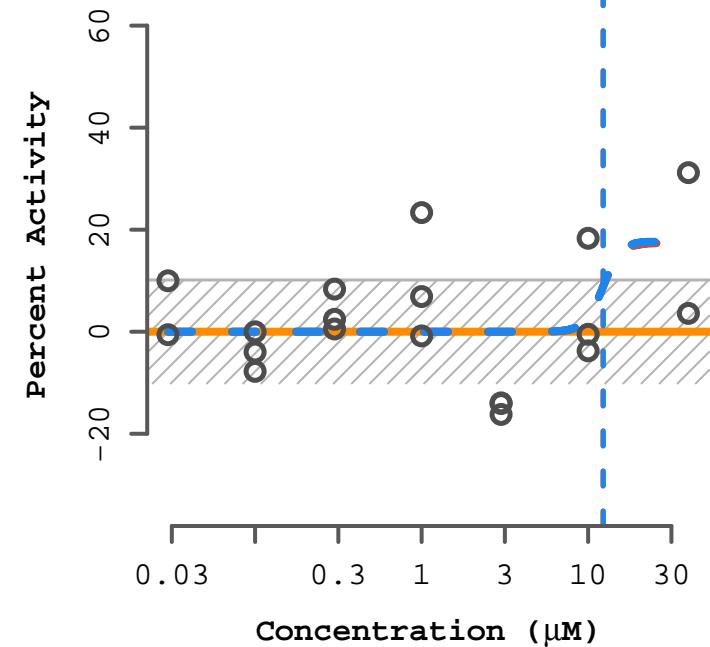
MAX_MED: 23.3

BMAD: 3.38

COFF:

10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.4

FLAGS: 8; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diisobutyl adipate
CHID: 36690 CASRN: 141-04-8
SPID(S): TP0001412H08
M4ID: 30204397

HILL MODEL (in red):

	tp	ga	gw
val:	17.4	1.09	8
sd:	13.7	0.976	86.7

GAIN-LOSS MODEL (in blue):

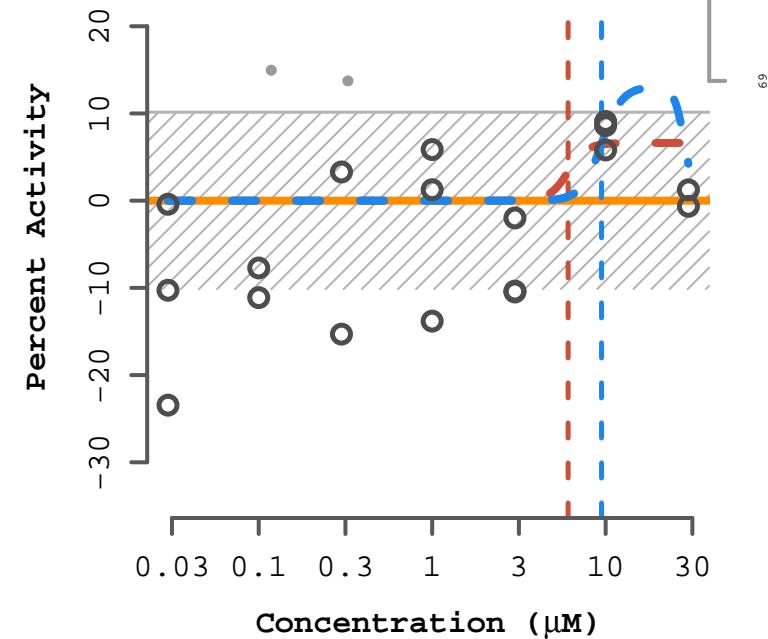
	tp	ga	gw	la	lw
val:	18.5	1.09	7.97	3.19	0.755
sd:	400	1.27	85.8	239	119

	CNST	HILL	GNLS
AIC:	150.49	153.79	157.79
PROB:	0.82	0.16	0.02
RMSE:	12.22	10.7	10.7

MAX_MEAN: 17.4 MAX_MED: 17.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.18

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Calcium nitrate

CHID: 39719 CASRN: 10124-37-5

SPID(S): EPAPLT0154A02

M4ID: 30204154 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	6.61	0.784	8
sd:	6.1	0.714	21.4

GAIN-LOSS MODEL (in blue):

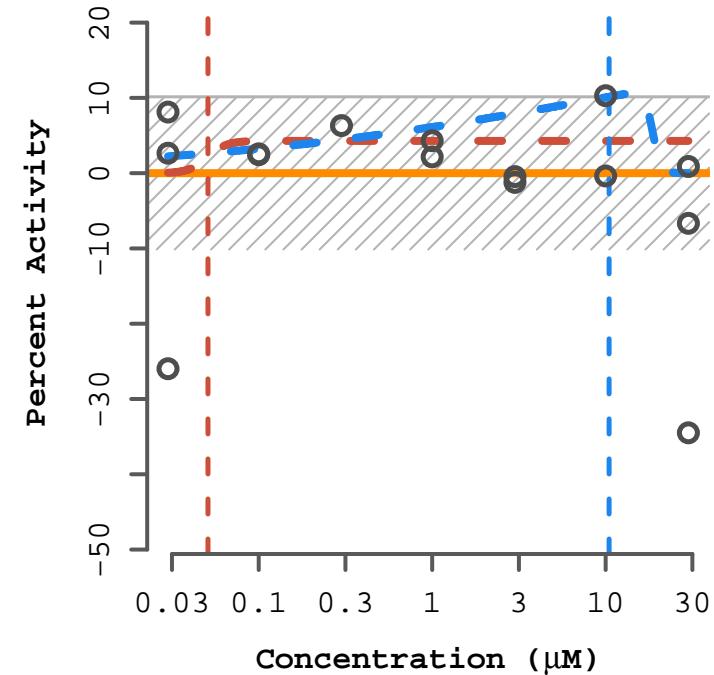
	tp	ga	gw	la	lw
val:	13.1	0.976	7.4	1.46	14.7
sd:	NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 196.06	200.84	204.74
PROB: 0.91	0.08	0.01
RMSE: 31.32	30.31	30.6

MAX_MEAN: 31.9 MAX_MED: 8.61 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.09

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Morpholinecarboxaldehyde
CHID: 44427 CASRN: 4394-85-8
SPID(S): EPAPLT0154E03
M4ID: 30204197

HILL MODEL (in red):

tp	ga	gw
val:	4.28	-1.29
sd:	3.74	1.07
		32.6

GAIN-LOSS MODEL (in blue):

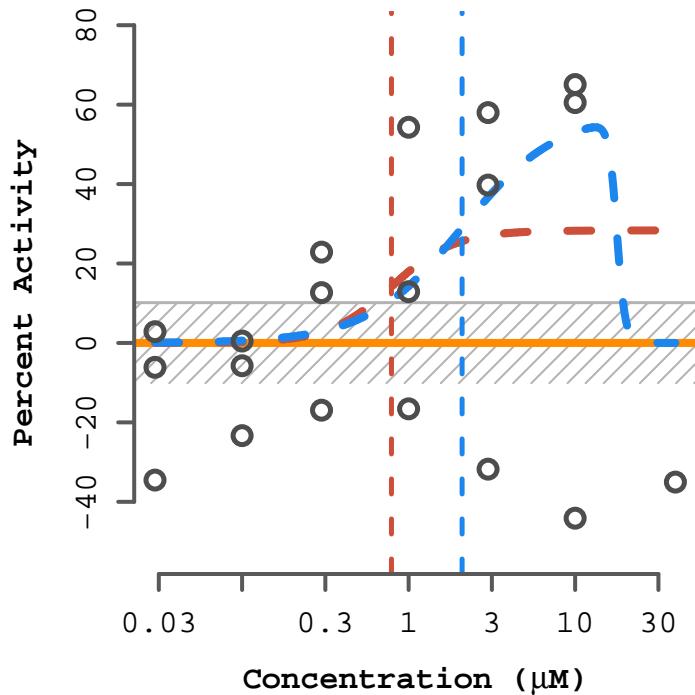
tp	ga	gw	la	lw
val:	20.3	1.02	0.357	1.27
sd:	101	10.6	0.67	2.58
				223

CNST	HILL	GNLS
AIC: 157.56	162.18	164.45
PROB: 0.88	0.09	0.03
RMSE: 19.09	18.22	17.18

MAX_MEAN: 15.1 MAX_MED: 10.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.12

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-tert-Butyl-4-ethylphenol
 CHID: 44581 CASRN: 96-70-8
 SPID(S): TP0001412D01
 M4ID: 30204456

HILL MODEL (in red):

tp	ga	gw
val: 28.3	-0.103	2.3
sd: 23.8	0.785	3.55

GAIN-LOSS MODEL (in blue):

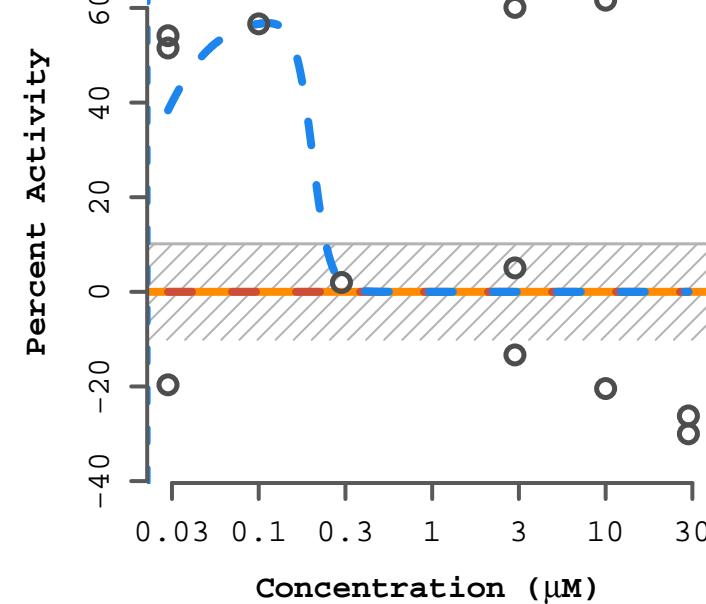
tp	ga	gw	la	lw
val: 58	0.321	1.51	1.25	17.5
sd: 31	0.53	1.52	12.3	709

CNST	HILL	GNLS
AIC: 193	196.77	196.74
PROB: 0.77	0.12	0.12
RMSE: 34.95	33.63	33.49

MAX_MEAN: 27.2 MAX_MED: 60.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.23

FLAGS: 8; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,4,4'-Trihydroxybenzophenone
CHID: 44836 CASRN: 1470-79-7
SPID(S): EPAPLT0154B11
M4ID: 30204173

HILL MODEL (in red):

	tp	ga	gw
val:	3.7	1.97	7.98
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

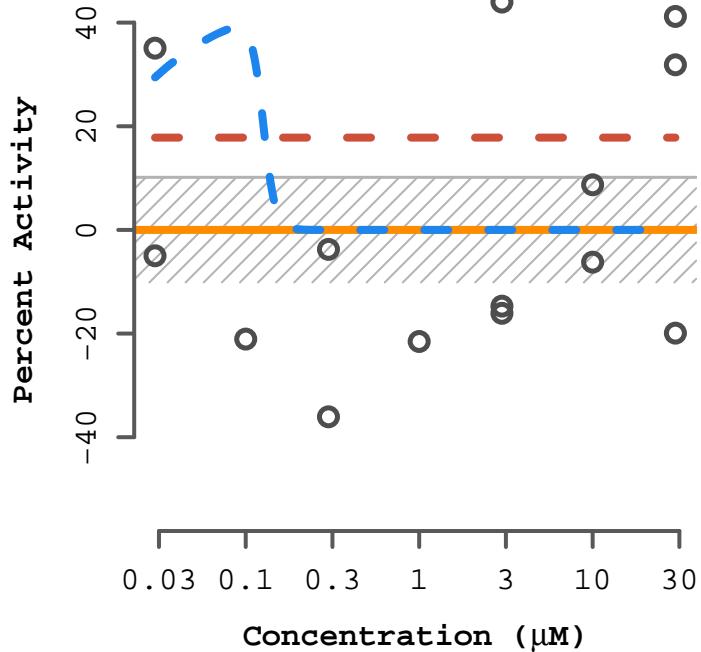
	tp	ga	gw	la	lw
val:	58.2	-1.64	2.48	-0.69	8.69
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	126.02	132.02	130.79
PROB:	0.88	0.04	0.08
RMSE:	39.59	39.59	33.43

MAX_MEAN: 56.6 MAX_MED: 56.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.12

FLAGS: 8; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ethenylsilanetriyl triacetate

CHID: 44938 CASRN: 4130-08-9

SPID(S): EPAPLT0154B10

M4ID: 30204172

HILL MODEL (in red):

	tp	ga	gw
val:	17.8	-2.52	5.99
sd:	8.87	269	1610

GAIN-LOSS MODEL (in blue):

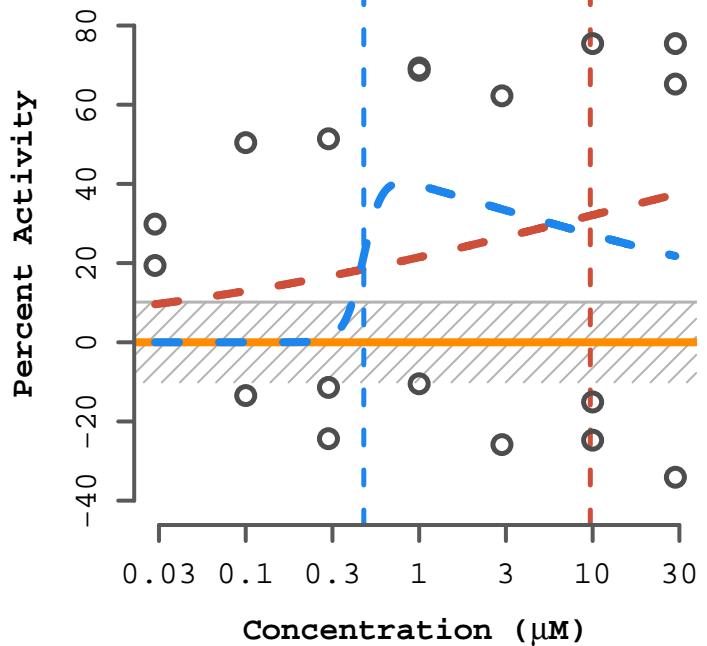
	tp	ga	gw	la	lw
val:	44.1	-1.75	1.33	-0.899	12.8
sd:	2900	75	326	24.9	1980

CNST	HILL	GNLS
AIC: 206.06	208.16	211.77
PROB: 0.71	0.25	0.04
RMSE: 36.91	32.62	33.68

MAX_MEAN: 29.1 MAX_MED: 51.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.29

FLAGS: 10; 8



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: (-)-Ambroxide

CHID: 47113 CASRN: 6790-58-5

SPID(S): EPAPLT0154B09

M4ID: 30204171

HILL MODEL (in red):

tp	ga	gw
val: 63.9	0.987	0.3
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 74.2	-0.319	8	0.206	0.3
sd: 803	1.1	31.8	28.9	1.05

CNST

AIC: 193.96

PROB: 0.75

RMSE: 46.65

HILL

196.41

0.22

41.17

GNLS

200.52

0.03

41.24

MAX_MEAN: 42.5

MAX_MED: 68.7

BMAD: 3.38

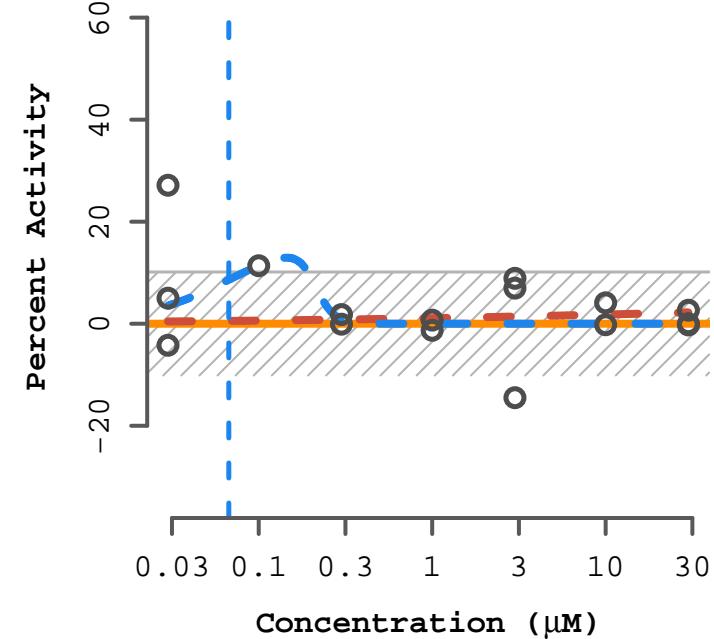
COFF: 10.1

HIT-CALL: 0

FITC: 10

ACTP: 0.25

FLAGS: 10; 8



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Vanillin isobutyrate

CHID: 47201 CASRN: 20665-85-4

SPID(S): EPAPLT0154A08

M4ID: 30204159

HILL MODEL (in red):

	tp	ga	gw
val:	5.16	1.9	0.3
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

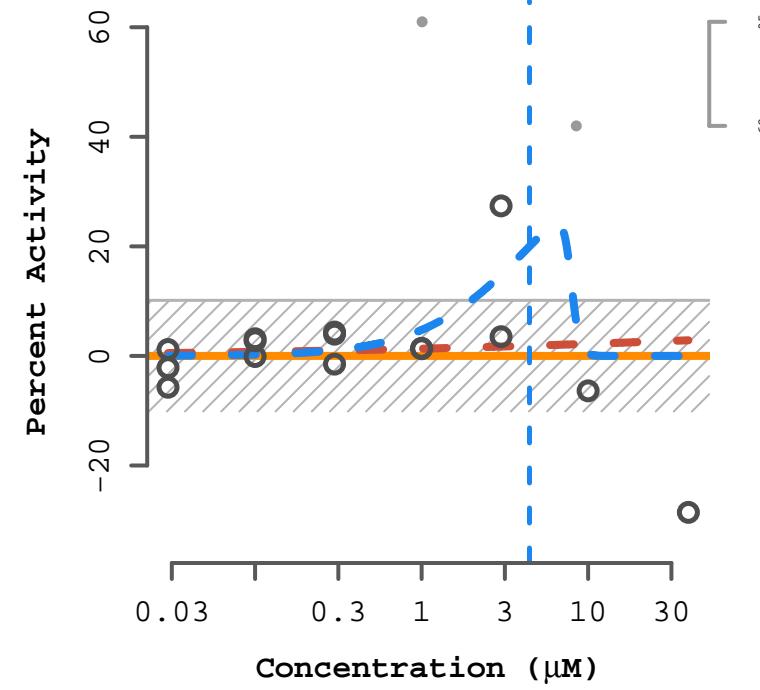
	tp	ga	gw	la	lw
val:	17.2	-1.17	1.68	-0.675	8.2
sd:	1160	41.6	94.7	9.27	487

	CNST	HILL	GNLS
AIC:	113.4	118.35	118.52
PROB:	0.86	0.07	0.07
RMSE:	8.93	8.8	7.85

MAX_MEAN: 11.4 MAX_MED: 11.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.14

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: CP-122721

CHID: 47251 CASRN: 145742-28-5

SPID(S): TP0001414E08

M4ID: 30204492 BRK

HILL MODEL (in red):

tp	ga	gw
val: 6.62	2	0.3
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

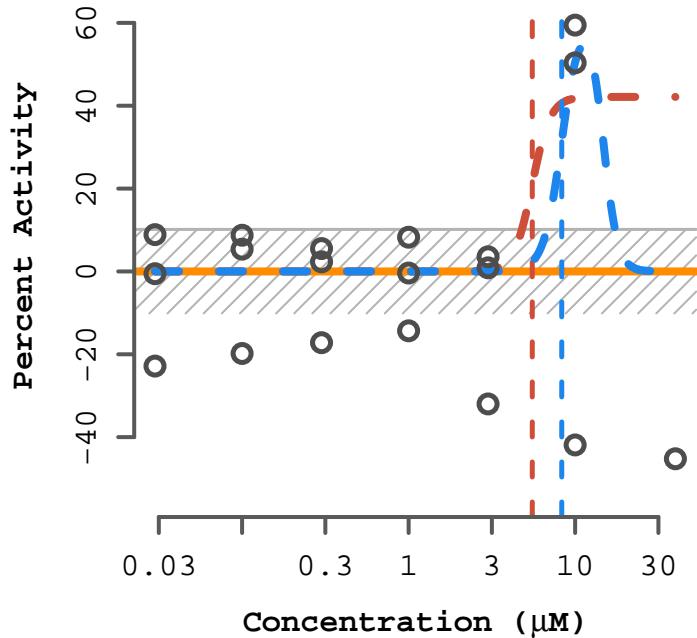
tp	ga	gw	la	lw
val: 40	0.648	1.34	0.898	17.9
sd: 208	2.85	2.49	1.05	184

CNST	HILL	GNLS
AIC: 153.87	159.73	161.13
PROB: 0.93	0.05	0.02
RMSE: 29.29	28.85	27.85

MAX_MEAN: 32.5 MAX_MED: 28 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.07

FLAGS: 10; 8



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: CP-607366

CHID: 47280 CASRN: 289716-94-5

SPID(S): TP0001412C12

M4ID: 30204501

HILL MODEL (in red):

tp	ga	gw
val: 42.1	0.743	8
sd: 16.8	0.282	8.56

GAIN-LOSS MODEL (in blue):

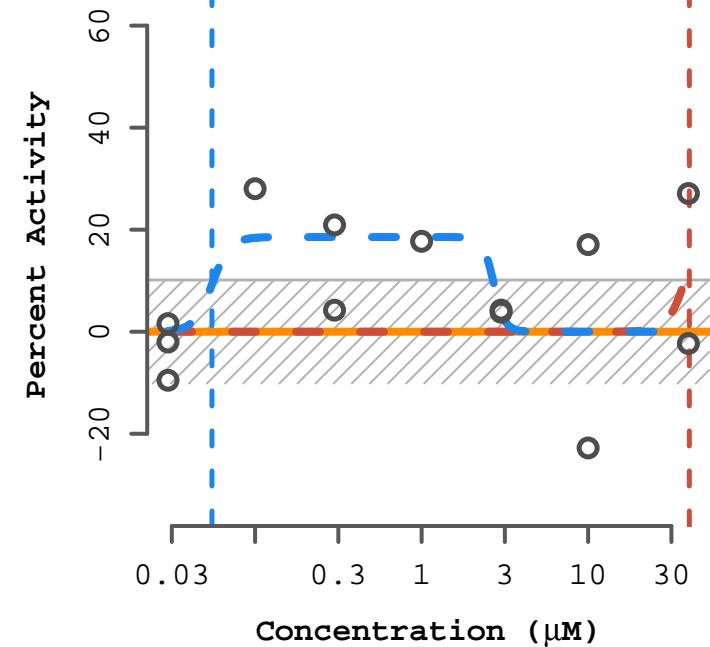
tp	ga	gw	la	lw
val: 63.1	0.92	8	1.17	10.2
sd: 304	1.22	30.6	2.52	76.4

CNST	HILL	GNLS
AIC: 178.84	184.07	182.2
PROB: 0.79	0.06	0.15
RMSE: 25.76	30.58	26.55

MAX_MEAN: 22.7 MAX_MED: 50.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.21

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Nelivaptan

CHID: 47358 CASRN: 439687-69-1

SPID(S): TP0001411A08

M4ID: 30204641

HILL MODEL (in red):

tp	ga	gw
val: 26	1.61	7.3
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

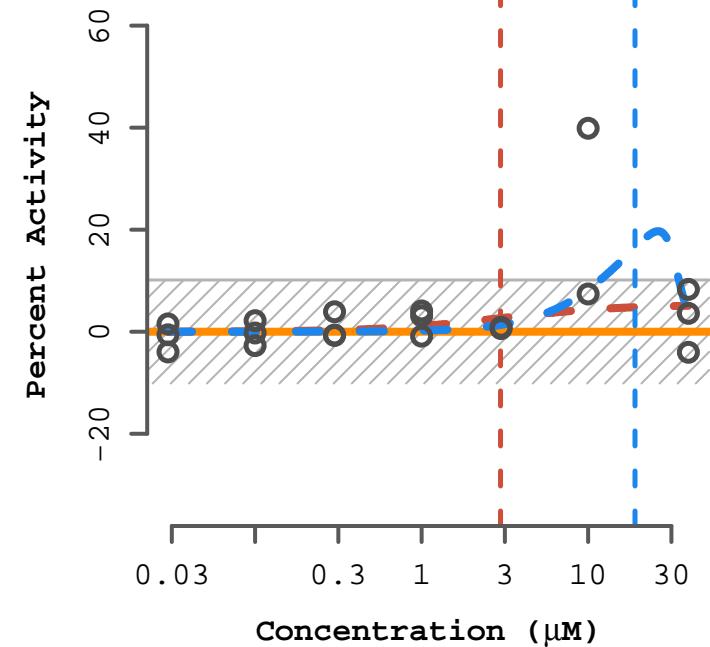
tp	ga	gw	la	lw
val: 18.6	-1.26	8	0.435	13
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 111.84	116.91	113.79
PROB: 0.69	0.05	0.26
RMSE: 15.78	15.01	12.26

MAX_MEAN: 28 MAX_MED: 28 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.31

FLAGS: 10; 8



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SAR115740

CHID: 47366 CASRN: NOCAS_47366

SPID(S): TP0001414G01

M4ID: 30204522

HILL MODEL (in red):

tp	ga	gw
val: 5.3	0.474	1.22
sd: 2.62	0.594	1.57

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 33.1	1.28	1.7	1.53	10.9
sd: NaN	NAN	NAN	NAN	NAN

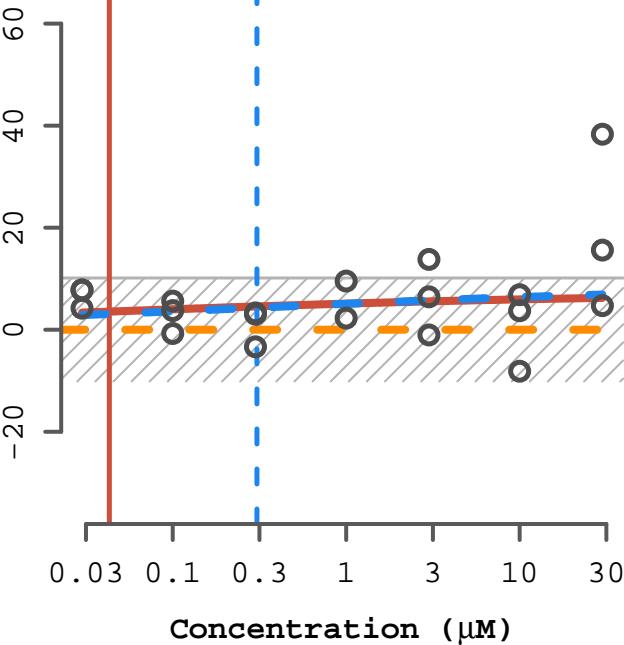
CNST	HILL	GNLS
AIC: 122.03	122.36	123.68
PROB: 0.44	0.37	0.19
RMSE: 9.79	8.71	7.79

MAX_MEAN: 23.6 MAX_MED: 23.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 10 ACTP: 0.56

FLAGS: 12

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bis{2-[2-(propan-2-yl)-1,3-oxazolidin-3-yl]ethyl}
CHID: 44961 CASRN: 59719-67-4
SPID(S): EPAPLT0154D04
M4ID: 30204187

HILL MODEL (in red):

tp	ga	gw
val: 7.11	-1.37	0.3
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 8.64	-0.515	0.3	3.4	3.87
sd: NaN	NaN	NaN	NaN	NaN

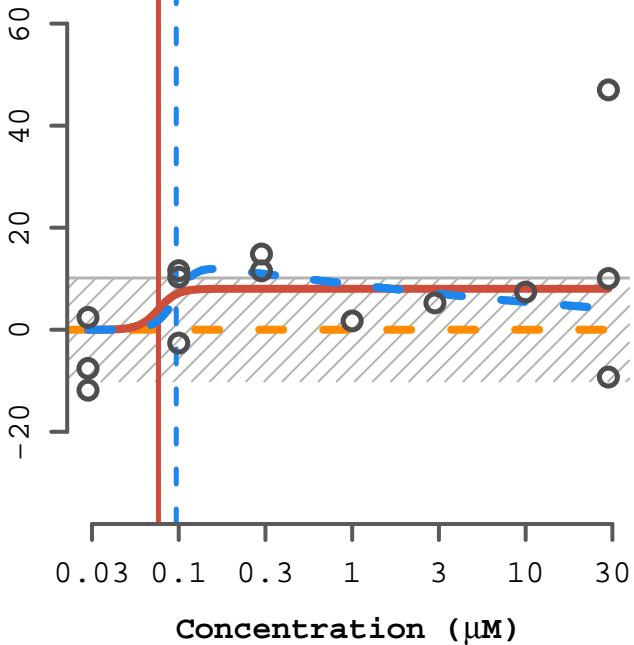
CNST	HILL	GNLS
AIC: 148.67	144.05	148.07
PROB: 0.08	0.81	0.11
RMSE: 10.97	8.93	8.83

MAX_MEAN: 19.6 MAX_MED: 15.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 15 ACTP: 0.92

FLAGS: 12

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Ethylhexyl glycidyl ether
CHID: 25301 CASRN: 2461-15-6
SPID(S): EPAPLT0154A09
M4ID: 30204160

HILL MODEL (in red):

tp	ga	gw
val: 8.03	-1.12	8
sd: 3.16	0.487	26

GAIN-LOSS MODEL (in blue):

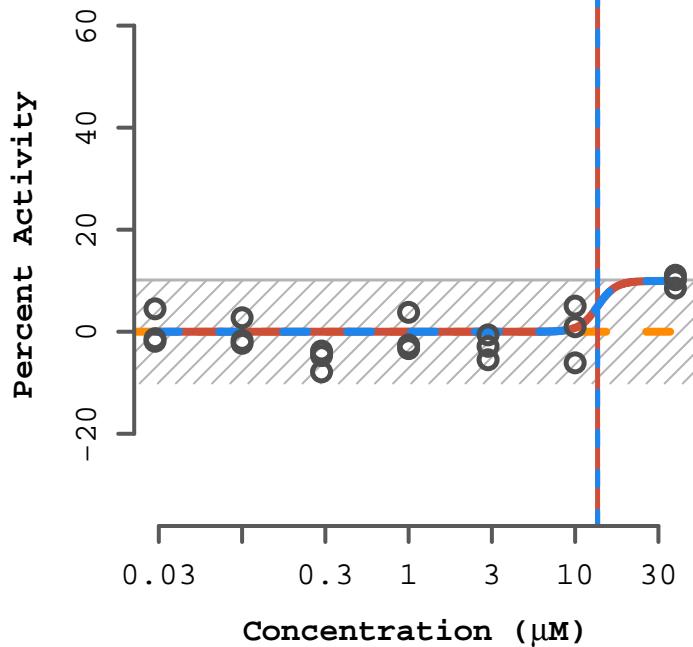
tp	ga	gw	la	lw
val: 24.1	-1.01	7.78	-0.764	0.303
sd: 87.2	0.164	37.2	8.91	0.468

CNST	HILL	GNLS
AIC: 115	114.71	118.21
PROB: 0.42	0.49	0.09
RMSE: 15.32	12.7	13.24

MAX_MEAN: 15.9 MAX_MED: 13.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 16 ACTP: 0.58

FLAGS: 12; 10; 8



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,4-Dichlorophenoxybutyric acid
CHID: 24035 CASRN: 94-82-6
SPID(S): TP0001412B01
M4ID: 30204613

HILL MODEL (in red):

	tp	ga	gw
val:	9.95	1.13	8
sd:	1.73	2.15	127

GAIN-LOSS MODEL (in blue):

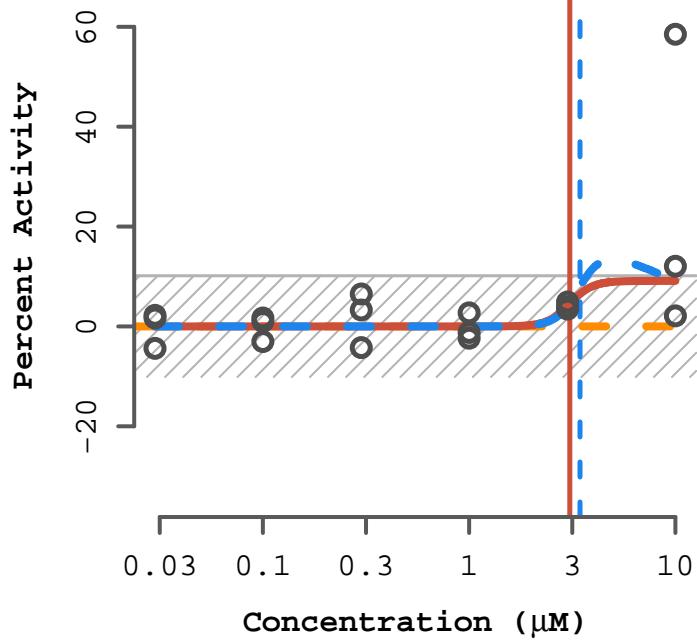
	tp	ga	gw	la	lw
val:	10	1.14	8	3.6	1.13
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	132.77	123.9	127.9
PROB:	0.01	0.87	0.12
RMSE:	5.22	3.64	3.64

MAX_MEAN: 9.94 MAX_MED: 10.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 20 ACTP: 0.99

FLAGS: 12



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Celecoxib

CHID: 22777 CASRN: 169590-42-5

SPID(S): TP0001413G06

M4ID: 30204603

HILL MODEL (in red):

tp	ga	gw
val: 9.11	0.488	8
sd: 4.14	0.133	85.5

GAIN-LOSS MODEL (in blue):

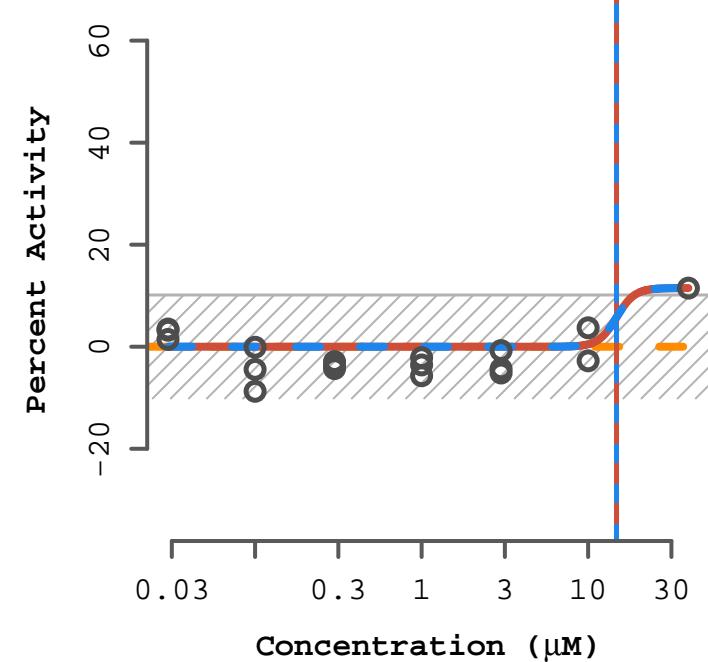
tp	ga	gw	la	lw
val: 18.9	0.538	7.95	0.981	1.65
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 125.43	123.14	127.14
PROB: 0.22	0.69	0.09
RMSE: 14.44	12.08	12.08

MAX_MEAN: 24.2 MAX_MED: 12.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 0 FITC: 21 ACTP: 0.78

FLAGS: 12; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Azobenzene

CHID: 20123 CASRN: 103-33-3

SPID(S): TP0001411G08

M4ID: 30204551

HILL MODEL (in red):

	tp	ga	gw
val:	11.5	1.17	7.99
sd:	3.29	3.65	170

GAIN-LOSS MODEL (in blue):

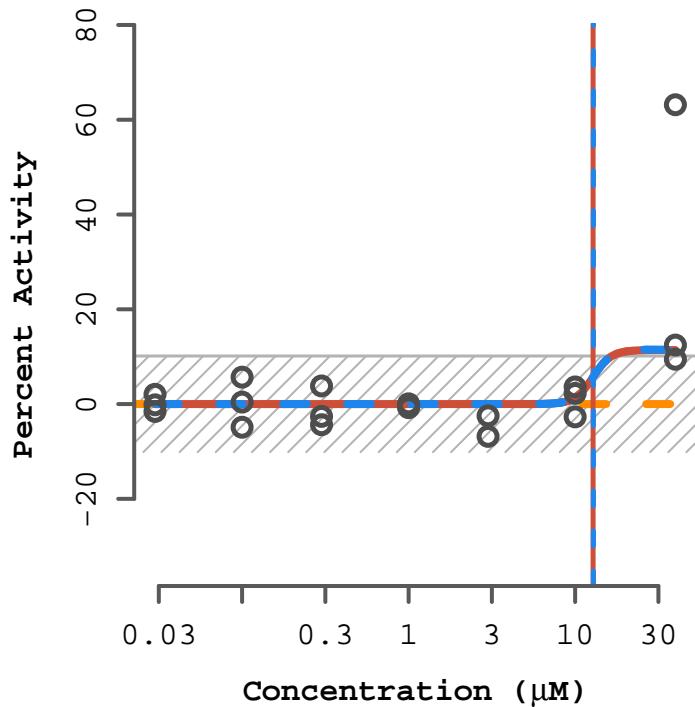
	tp	ga	gw	la	lw
val:	11.5	1.17	7.99	3.09	9.16
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	110.61	110.61	114.61
PROB:	0.47	0.47	0.06
RMSE:	4.77	3.93	3.93

MAX_MEAN: 11.5 MAX_MED: 11.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 37 ACTP: 0.53

FLAGS: 17; 16; 11; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Acibenzolar-S-methyl
CHID: 32519 CASRN: 135158-54-2
SPID(S): TP0001412C06
M4ID: 30204421

HILL MODEL (in red):

tp	ga	gw
val: 11.4	1.11	8
sd: 2.17	1.33	98.6

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 11.5	1.11	7.93	1.95	5.38
sd: 68.4	1.38	95.9	103	1620

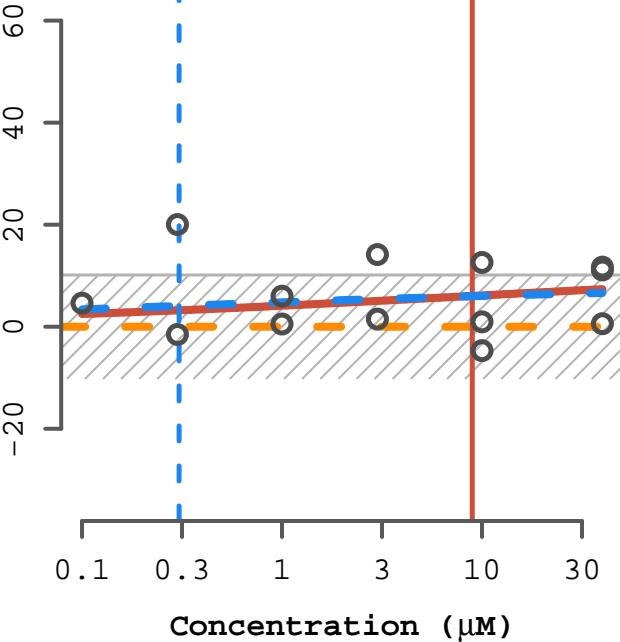
CNST	HILL	GNLS
AIC: 139.86	132.43	136.43
PROB: 0.02	0.86	0.12
RMSE: 14.86	11.97	11.97

MAX_MEAN: 28.4 MAX_MED: 12.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 37 ACTP: 0.98

FLAGS: 17; 11; 10; 7

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Thiacloprid

CHID: 34961 CASRN: 111988-49-9

SPID(S): TP0001411A12

M4ID: 30204289

HILL MODEL (in red):

	tp	ga	gw
val:	12.1	0.951	0.3
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

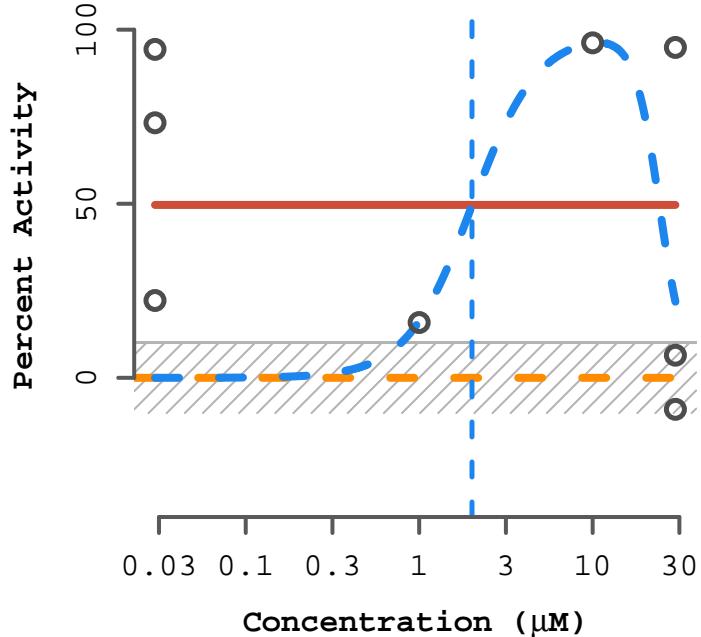
	tp	ga	gw	la	lw
val:	8.15	-0.515	0.3	3.6	11.9
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	97.23	97.2	101.15
PROB:	0.46	0.47	0.07
RMSE:	9.23	7.27	7.15

MAX_MEAN: 9.26 MAX_MED: 11.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 38 ACTP: 0.54

FLAGS: 17; 16; 11; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Linalool

CHID: 25502 CASRN: 78-70-6

SPID(S): EPAPLT0154A03

M4ID: 30204155

HILL MODEL (in red):

tp	ga	gw
val: 49.7	-2.52	5.88
sd: 18.3	158	927

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 99	0.304	2.32	1.39	5.97
sd: NaN	NaN	NaN	NaN	NaN

CNST

HILL

GNLS

AIC: 92.37

92.14

98.35

PROB: 0.46

0.52

0.02

RMSE: 64.62

41.79

51.58

MAX_MEAN: 96.2

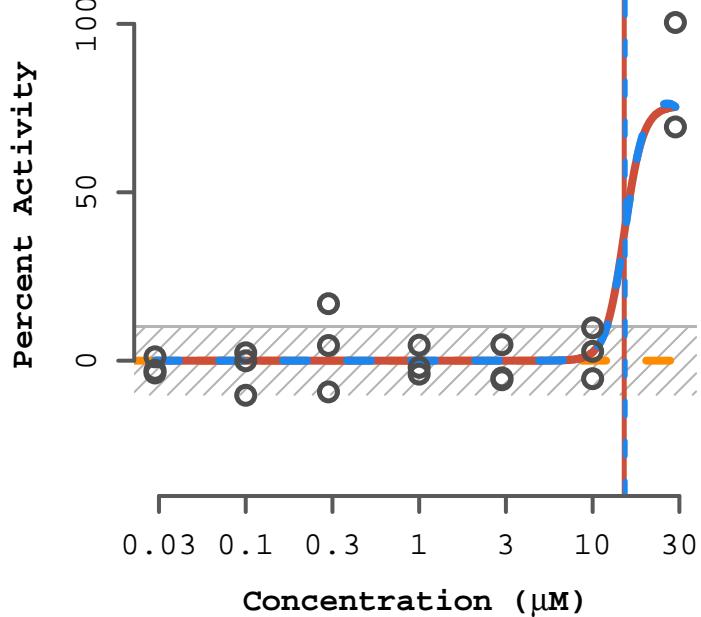
MAX_MED: 96.2

BMAD: 3.38

COFF:

10.1 HIT-CALL: 1 FITC: 40 ACTP: 0.54

FLAGS: 18; 17; 16; 11; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Amino-4-nitrophenol
CHID: 20062 CASRN: 99-57-0
SPID(S): EPAPLT0154D08
M4ID: 30204191

HILL MODEL (in red):

tp	ga	gw
val: 75.7	1.18	8
sd: 45.3	2.81	122

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 77.9	1.19	7.83	1.6	12.1
sd: NaN	NAN	NAN	NAN	NAN

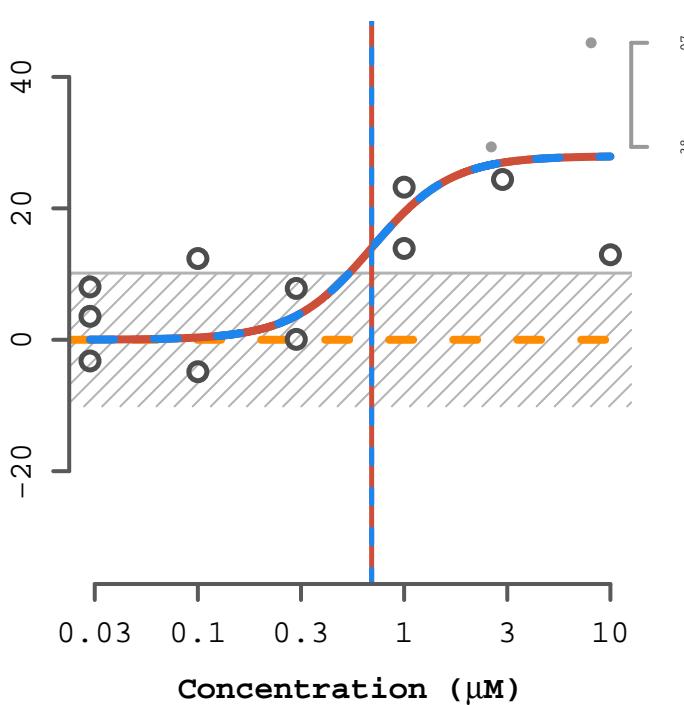
CNST	HILL	GNLS
AIC: 170.22	147.82	151.82
PROB: 0	0.88	0.12
RMSE: 28	8.45	8.45

MAX_MEAN: 84.9 MAX_MED: 84.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 7

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Clorophene

CHID: 20154 CASRN: 120-32-1

SPID(S): TP0001414E06

M4ID: 30204579 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	28	-0.157	2.25
sd:	8.42	0.244	1.83

GAIN-LOSS MODEL (in blue):

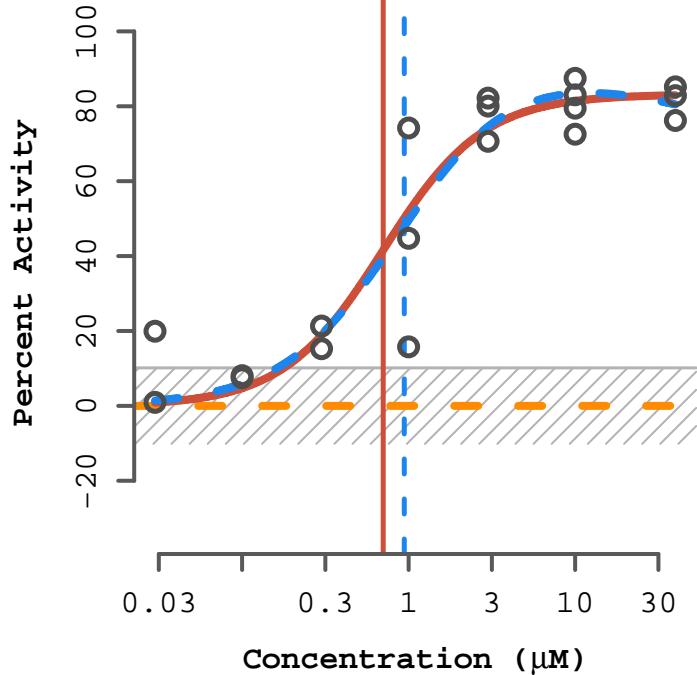
	tp	ga	gw	la	lw
val:	28	-0.157	2.25	2.67	18
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	123.12	113.42	117.42
PROB:	0.01	0.87	0.12
RMSE:	31.22	20.47	20.47

MAX_MEAN: 54.9 MAX_MED: 54.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.99

FLAGS: 17; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bifenthrin

CHID: 20160 CASRN: 82657-04-3

SPID(S): TP0001411B11

M4ID: 30204350

HILL MODEL (in red):

tp	ga	gw
val: 83.2	-0.153	1.44
sd: 3.1	0.0868	0.282

GAIN-LOSS MODEL (in blue):

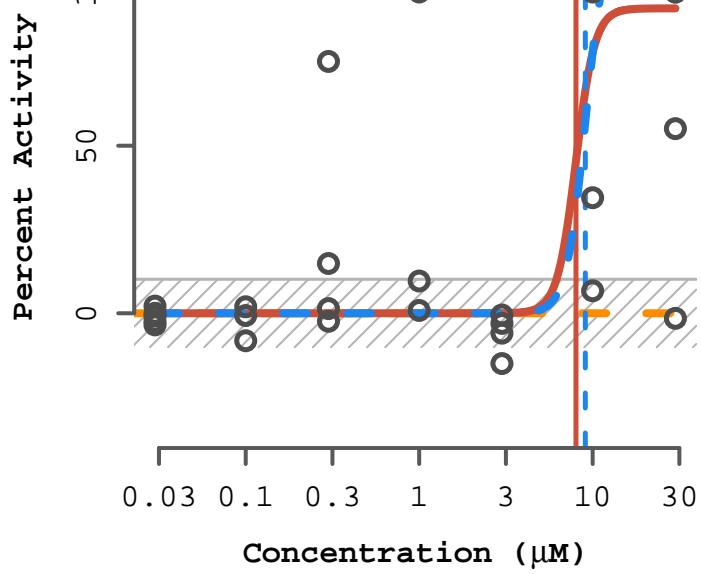
tp	ga	gw	la	lw
val: 105	-0.0261	1.24	3.41	0.3
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 215.98	149.54	152.8
PROB: 0	0.84	0.16
RMSE: 62.06	11.57	11.34

MAX_MEAN: 81.4 MAX_MED: 82.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Butylated hydroxyanisole
CHID: 20215 CASRN: 25013-16-5
SPID(S): EPAPLT0167B11
M4ID: 30204243

HILL MODEL (in red):

tp	ga	gw
val: 91	0.904	8
sd: 8.88	0.2	14.6

GAIN-LOSS MODEL (in blue):

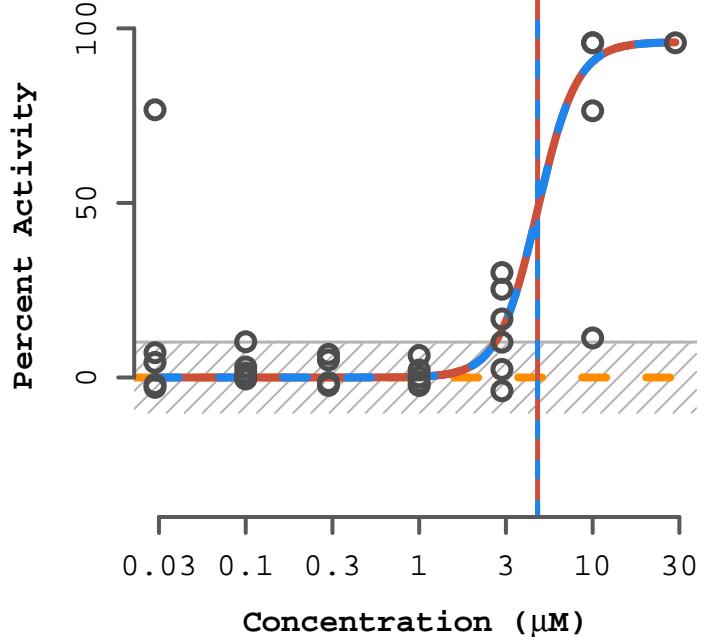
tp	ga	gw	la	lw
val: 114	0.958	7.84	1.59	5.08
sd: 1100	1.56	24.7	1.17	149

CNST	HILL	GNLS
AIC: 311.18	283.64	287.64
PROB: 0	0.88	0.12
RMSE: 51.87	33.94	33.95

MAX_MEAN: 74.6 MAX_MED: 95.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Capsaicin

CHID: 20241 CASRN: 404-86-4

SPID(S): EPAPLT0154B01

M4ID: 30204163

HILL MODEL (in red):

tp	ga	gw
val: 96.1	0.683	3.81
sd: 6.32	0.0795	1.32

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 96.1	0.683	3.81	2.5	8.24
sd: NaN	NaN	NaN	NaN	NaN

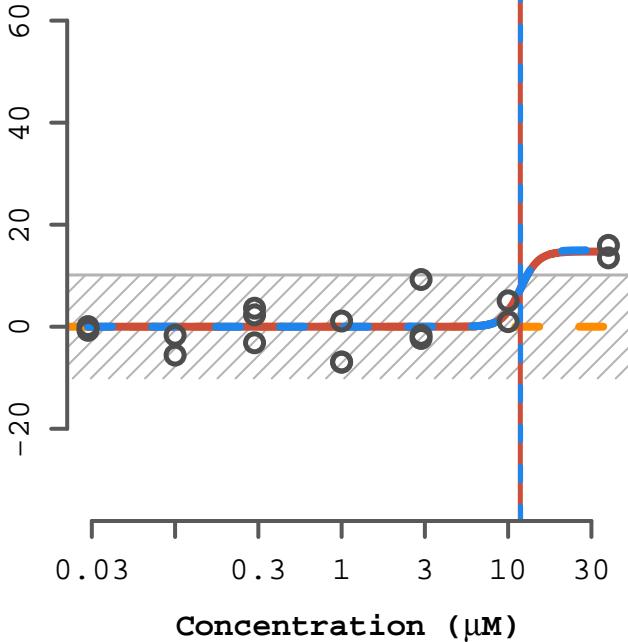
CNST	HILL	GNLS
AIC: 297.73	250.99	254.99
PROB: 0	0.88	0.12
RMSE: 36.68	20.94	20.94

MAX_MEAN: 95.9 MAX_MED: 95.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Carbaryl

CHID: 20247 CASRN: 63-25-2

SPID(S): TP0001414F07

M4ID: 30204332

HILL MODEL (in red):

	tp	ga	gw
val:	14.7	1.07	7.99
sd:	2.1	2.21	241

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	15	1.07	7.95	1.75	11.9
sd:	NaN	NaN	NaN	NaN	NaN

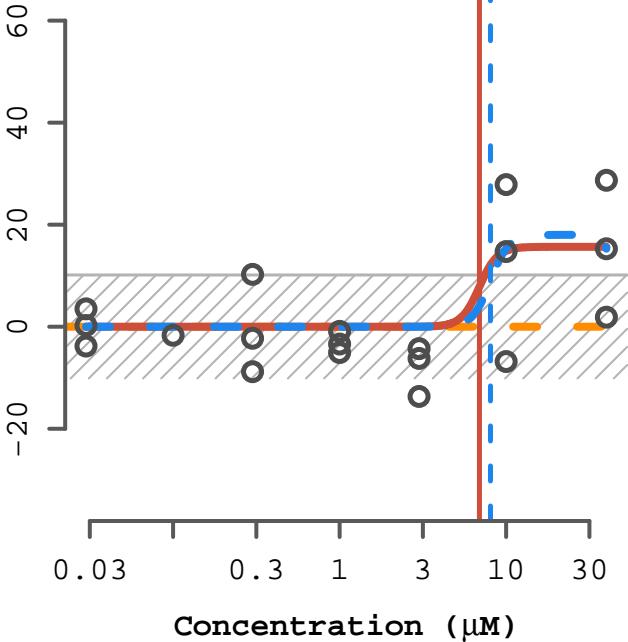
	CNST	HILL	GNLS
AIC:	113.11	102.99	106.99
PROB:	0.01	0.88	0.12
RMSE:	6.5	3.95	3.95

MAX_MEAN: 14.7 MAX_MED: 14.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.99

FLAGS: 17; 7

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Clomiphene citrate (1:1)
CHID: 20337 CASRN: 50-41-9
SPID(S): TP0001411H07
M4ID: 30204645

HILL MODEL (in red):

	tp	ga	gw
val:	15.7	0.84	8
sd:	5.36	0.366	11.5

GAIN-LOSS MODEL (in blue):

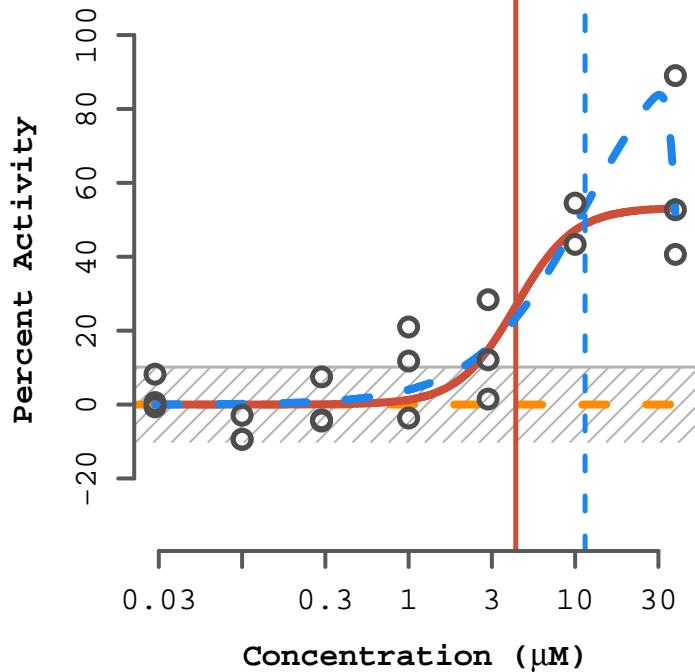
	tp	ga	gw	la	lw
val:	18	0.906	8	1.64	18
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	147.82	145.26	149.24
PROB:	0.2	0.71	0.1
RMSE:	11.7	8.89	8.91

MAX_MEAN: 15.3 MAX_MED: 15.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.8

FLAGS: 17; 11



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: p,p'-DDD

CHID: 20373 CASRN: 72-54-8

SPID(S): TP0001412B02

M4ID: 30204480

HILL MODEL (in red):

	tp	ga	gw
val:	53.3	0.643	2.49
sd:	7.84	0.122	1.81

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	107	1.06	1.33	1.61	17.4
sd:	NaN	NaN	NaN	NaN	NaN

CNST

AIC: 192.91

PROB: 0

RMSE: 30.91

HILL

161.22

0.83

11.73

GNLS

164.38

0.17

11.7

MAX_MEAN: 60.8

MAX_MED: 52.7

BMAD: 3.38

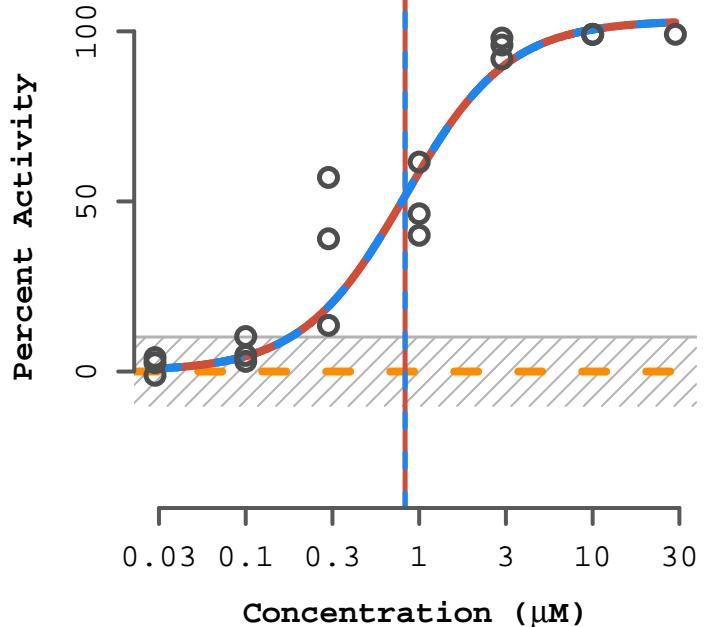
COFF: 10.1

HIT-CALL: 1

FITC: 41

ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Deltamethrin

CHID: 20381 CASRN: 52918-63-5

SPID(S): EPAPLT0169B07

M4ID: 30204248

HILL MODEL (in red):

tp	ga	gw
val: 103	-0.0812	1.46
sd: 3.77	0.0728	0.304

GAIN-LOSS MODEL (in blue):

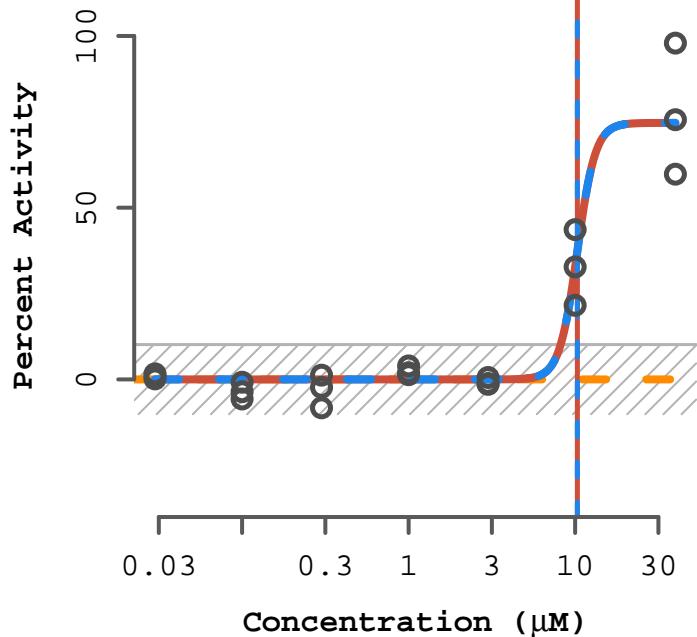
tp	ga	gw	la	lw
val: 103	-0.0812	1.46	2.72	8.03
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 216.55	148.62	152.62
PROB: 0	0.88	0.12
RMSE: 64.59	11.62	11.62

MAX_MEAN: 99.1 MAX_MED: 99.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diethylstilbestrol
 CHID: 20465 CASRN: 56-53-1
 SPID(S): TP0001414B02
 M4ID: 30204563

HILL MODEL (in red):

tp	ga	gw
val: 74.7	1.01	8
sd: 4.51	0.0285	16.6

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 74.7	1.01	8	3.5	10.9
sd: NA	NA	NA	NA	NA

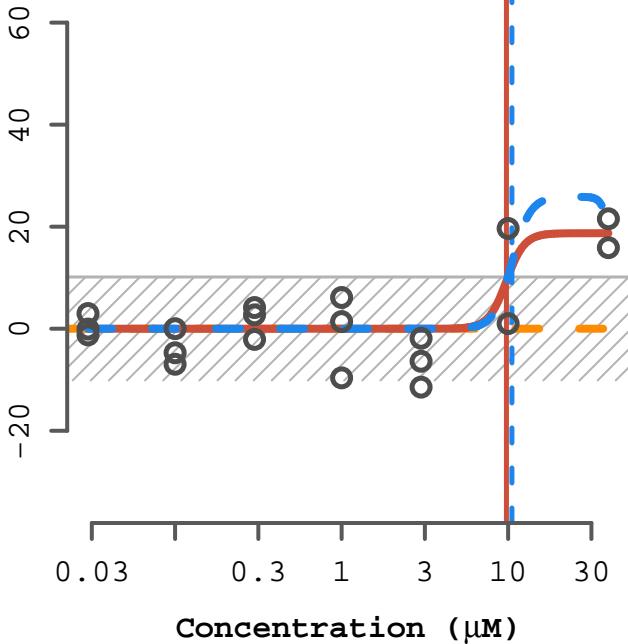
CNST	HILL	GNLS
AIC: 199.69	145.96	149.96
PROB: 0	0.88	0.12
RMSE: 32.71	7.42	7.42

MAX_MEAN: 77.8 MAX_MED: 75.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Eugenol

CHID: 20617 CASRN: 97-53-0

SPID(S): TP0001412H09

M4ID: 30204427

HILL MODEL (in red):

	tp	ga	gw
val:	18.7	0.99	8
sd:	3.45	0.296	28.3

GAIN-LOSS MODEL (in blue):

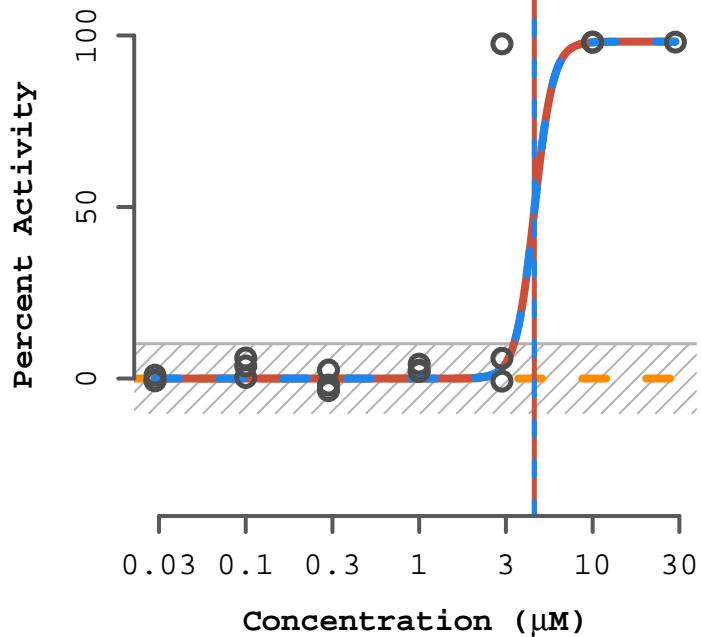
	tp	ga	gw	la	lw
val:	25.9	1.02	8	1.63	18
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	137.85	129.29	133.28
PROB:	0.01	0.87	0.12
RMSE:	8.95	5.65	5.65

MAX_MEAN: 18.7 MAX_MED: 18.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.99

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fluoxetine hydrochloride
CHID: 20635 CASRN: 56296-78-7
SPID(S): EPAPLT0169B09
M4ID: 30204259

HILL MODEL (in red):

tp	ga	gw
val: 98.2	0.664	8
sd: 2.07	0.217	9.21

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 98.2	0.664	8	2.87	7.74
sd: NaN	NaN	NaN	NaN	NaN

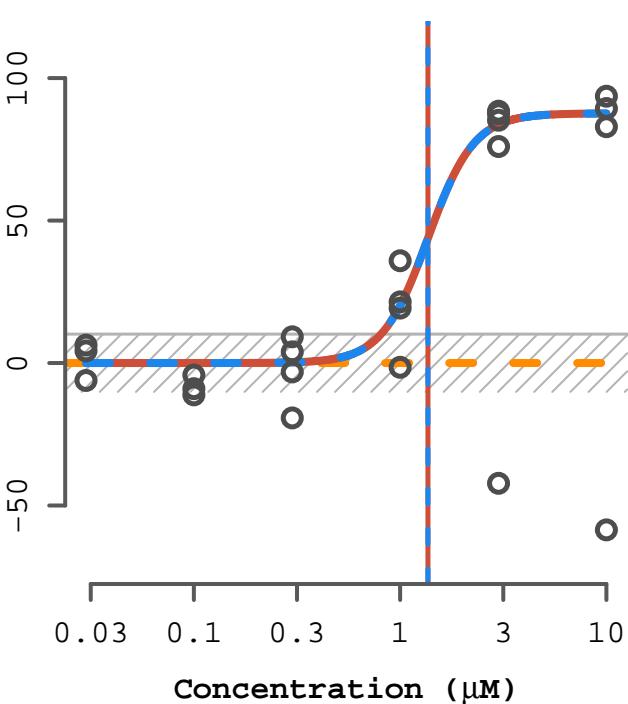
CNST	HILL	GNLS
AIC: 184.45	124.43	128.43
PROB: 0	0.88	0.12
RMSE: 46.26	22.43	22.43

MAX_MEAN: 98.1 MAX_MED: 98.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Gentian Violet
CHID: 20653 CASRN: 548-62-9
SPID(S): TP0001411A02
M4ID: 30204591

HILL MODEL (in red):

tp	ga	gw	
val:	87.6	0.135	3.88
sd:	5.73	0.0751	1.76

GAIN-LOSS MODEL (in blue):

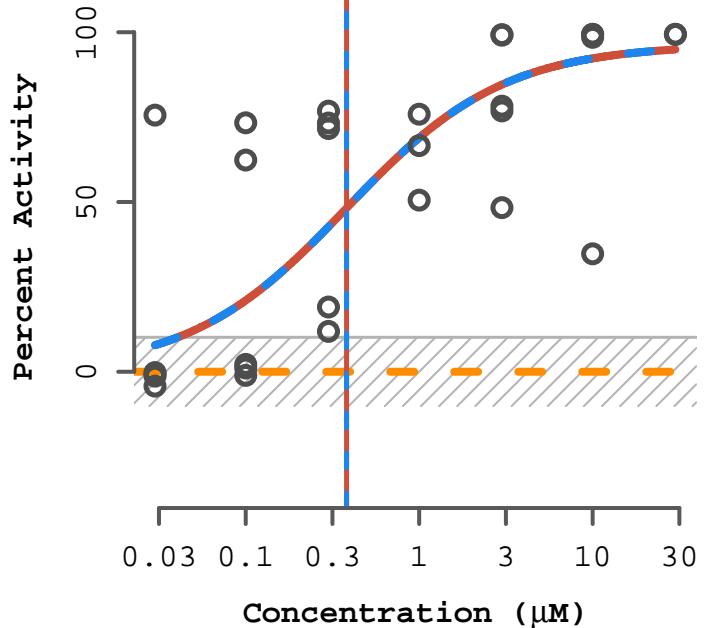
tp	ga	gw	la	lw	
val:	87.6	0.135	3.88	1.61	9.8
sd:	5.74	0.0751	1.76	262	4210

CNST	HILL	GNLS
AIC: 249.27	213.96	217.96
PROB: 0	0.88	0.12
RMSE: 51.19	41.12	41.12

MAX_MEAN: 59 MAX_MED: 86.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Hexachlorophene

CHID: 20690 CASRN: 70-30-4

SPID(S): EPAPLT0169C05

M4ID: 30204260

HILL MODEL (in red):

	tp	ga	gw
val:	96.4	-0.419	0.954
sd:	13.1	0.279	0.389

GAIN-LOSS MODEL (in blue):

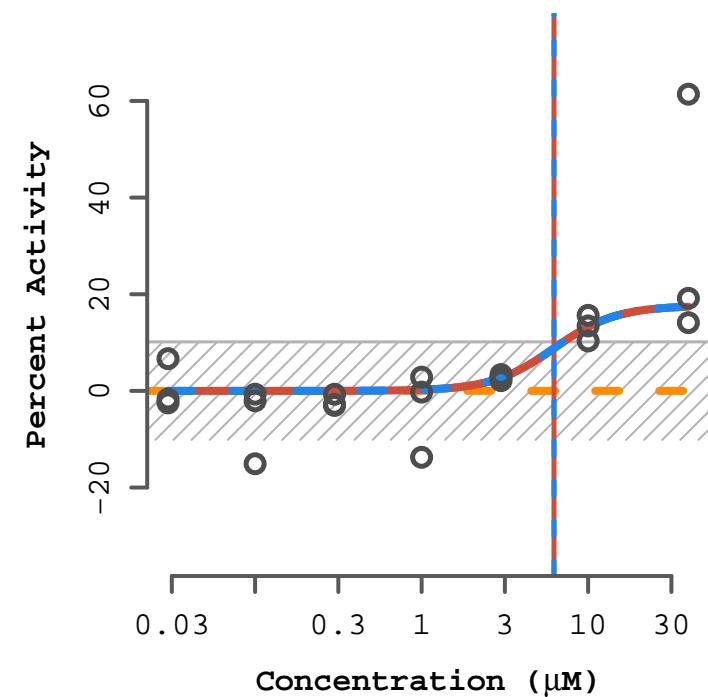
	tp	ga	gw	la	lw
val:	96.4	-0.419	0.954	3.31	17.9
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	332.52	281.24	285.24
PROB:	0	0.88	0.12
RMSE:	66.73	26.79	26.79

MAX_MEAN: 99.4 MAX_MED: 99.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Hexylresorcinol
CHID: 20699 CASRN: 136-77-6
SPID(S): TP0001412G05
M4ID: 30204402

HILL MODEL (in red):

	tp	ga	gw
val:	17.6	0.795	2.34
sd:	3.11	0.137	0.936

GAIN-LOSS MODEL (in blue):

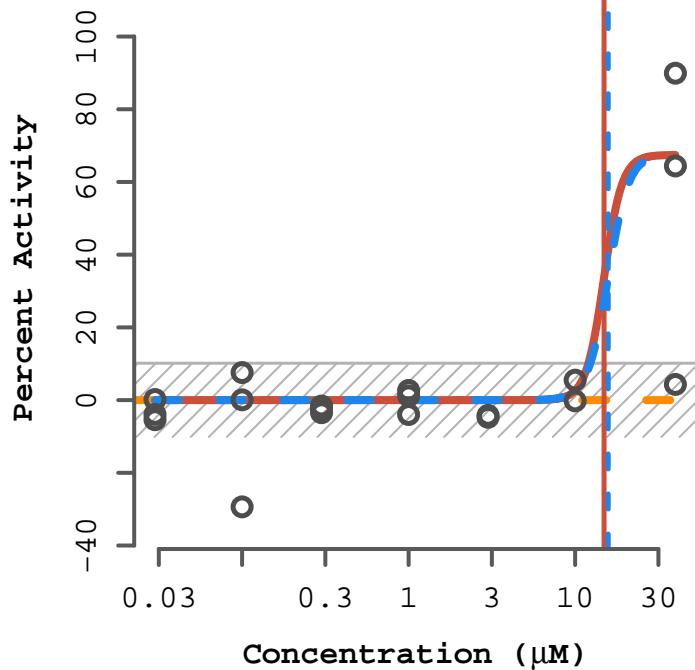
	tp	ga	gw	la	lw
val:	17.6	0.795	2.34	2.34	8.62
sd:	3.11	0.137	0.936	624	7270

CNST	HILL	GNLS
AIC: 167.31	146.61	150.61
PROB: 0	0.88	0.12
RMSE: 16.03	10.84	10.84

MAX_MEAN: 31.6 MAX_MED: 19.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,2-Diphenylhydrazine
CHID: 20710 CASRN: 122-66-7
SPID(S): TP0001414F02
M4ID: 30204550

HILL MODEL (in red):

tp	ga	gw
val: 67.4	1.17	8
sd: 7.35	1.79	82.1

GAIN-LOSS MODEL (in blue):

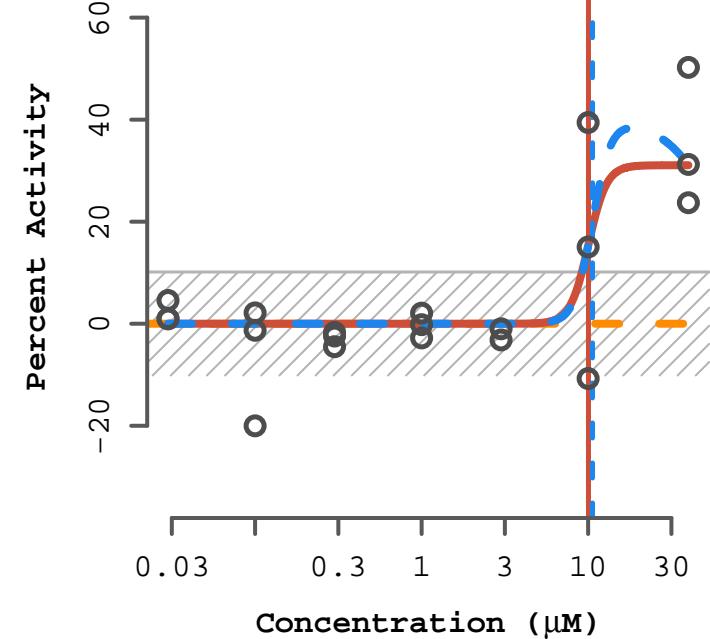
tp	ga	gw	la	lw
val: 68	1.2	7.01	3.6	1.08
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 161.46	154.13	158.13
PROB: 0.02	0.86	0.12
RMSE: 26.49	17.11	17.11

MAX_MEAN: 52.9 MAX_MED: 64.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.98

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Kepone

CHID: 20770 CASRN: 143-50-0

SPID(S): TP0001413C06

M4ID: 30204416

HILL MODEL (in red):

	tp	ga	gw
val:	31.1	1	8
sd:	4.84	0.0457	48.4

GAIN-LOSS MODEL (in blue):

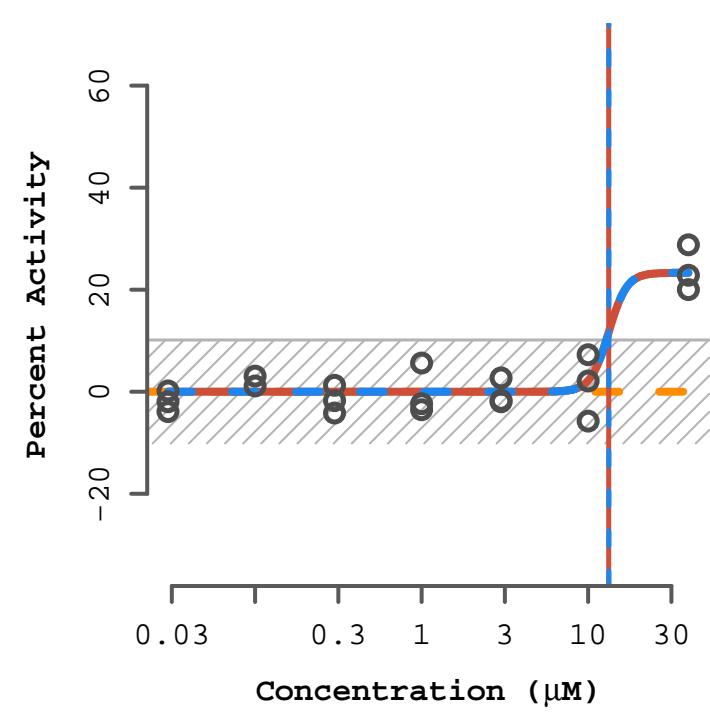
	tp	ga	gw	la	lw
val:	40.2	1.03	7.99	1.8	2.72
sd:	1100	1.78	111	7.65	291

	CNST	HILL	GNLS
AIC:	169.67	152.95	156.95
PROB:	0	0.88	0.12
RMSE:	17.95	10.4	10.4

MAX_MEAN: 35.1 MAX_MED: 31.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Methoxychlor

CHID: 20827 CASRN: 72-43-5

SPID(S): TP0001413G10

M4ID: 30204612

HILL MODEL (in red):

	tp	ga	gw
val:	23.3	1.12	7.99
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

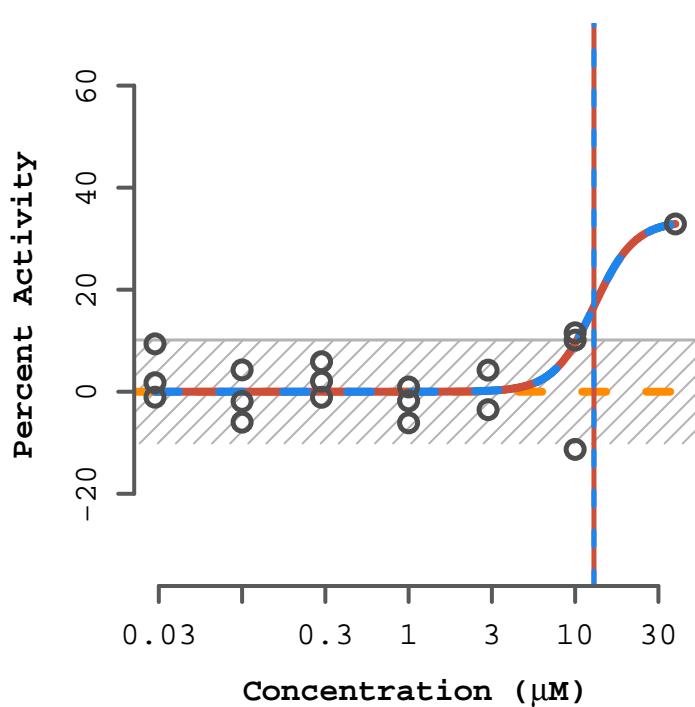
	tp	ga	gw	la	lw
val:	23.4	1.12	7.96	3.58	1.13
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	143.02	116.67	120.67
PROB:	0	0.88	0.12
RMSE:	9.89	3.5	3.5

MAX_MEAN: 23.9 MAX_MED: 22.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4,4'-Diamino-3,3'-dimethyldiphenylmethane
CHID: 20867 CASRN: 838-88-0
SPID(S): TP0001411C07
M4ID: 30204484

HILL MODEL (in red):

	tp	ga	gw
val:	33.5	1.11	3.55
sd:	10.6	0.426	11.7

GAIN-LOSS MODEL (in blue):

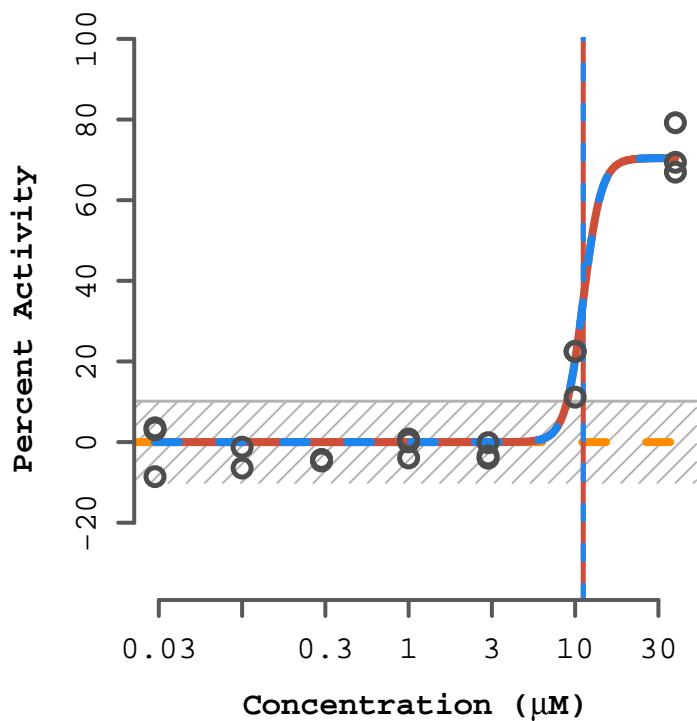
	tp	ga	gw	la	lw
val:	33.5	1.11	3.55	2.18	6.6
sd:	21.5	0.438	11.7	347	3860

CNST	HILL	GNLS
AIC: 129.57	119.47	123.47
PROB: 0.01	0.88	0.12
RMSE: 9.74	6.23	6.23

MAX_MEAN: 32.9 MAX_MED: 32.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.99

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Nitrofen

CHID: 20970 CASRN: 1836-75-5

SPID(S): TP0001412C09

M4ID: 30204540

HILL MODEL (in red):

	tp	ga	gw
val:	70.4	1.05	8
sd:	2.95	0.0383	6.36

GAIN-LOSS MODEL (in blue):

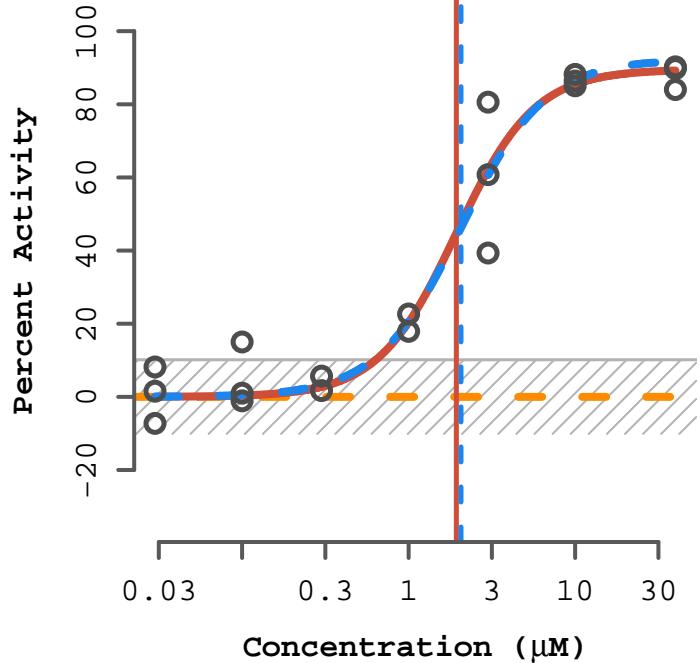
	tp	ga	gw	la	lw
val:	70.4	1.05	8	3.14	6.8
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	184.29	127.62	131.62
PROB:	0	0.88	0.12
RMSE:	29.11	4.62	4.62

MAX_MEAN: 71.9 MAX_MED: 69.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Phenylmercuric acetate
 CHID: 21150 CASRN: 62-38-4
 SPID(S): TP0001412G11
 M4ID: 30204599

HILL MODEL (in red):

tp	ga	gw
val: 89.6	0.287	1.85
sd: 2.43	0.0445	0.261

GAIN-LOSS MODEL (in blue):

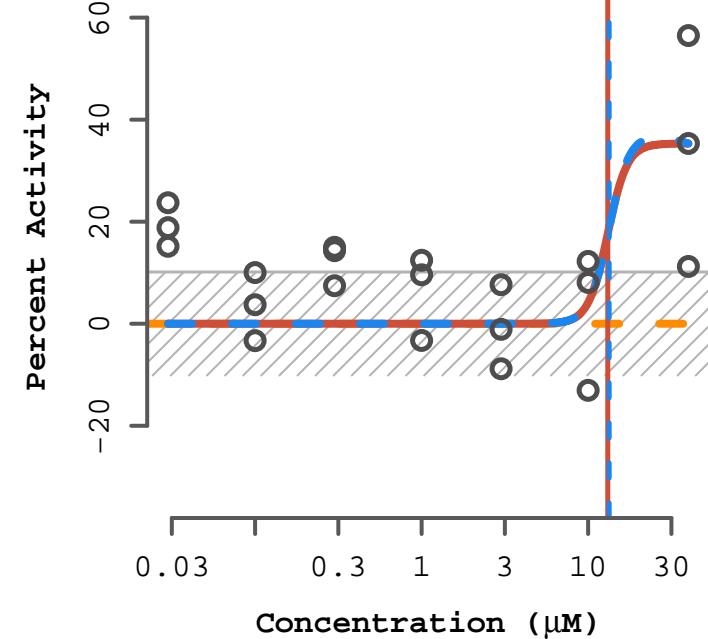
tp	ga	gw	la	lw
val: 92.4	0.315	1.71	1.7	13.7
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 210.22	136.31	139.83
PROB: 0	0.85	0.15
RMSE: 55.58	8.14	8.06

MAX_MEAN: 88 MAX_MED: 89.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Rotenone

CHID: 21248 CASRN: 83-79-4

SPID(S): TP0001412B10

M4ID: 30204343

HILL MODEL (in red):

	tp	ga	gw
val:	35.3	1.12	7.43
sd:	10.6	3.76	236

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	37.7	1.13	7.18	2.12	2.26
sd:	430	3.36	187	39.8	175

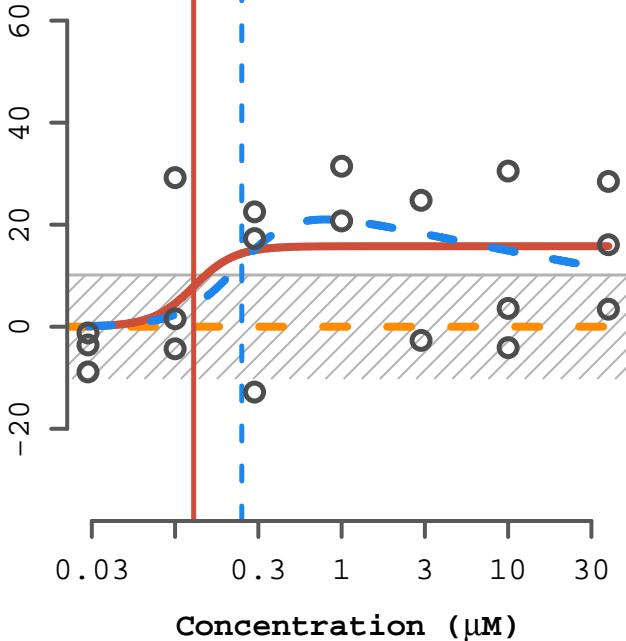
	CNST	HILL	GNLS
AIC:	181.64	177.91	181.91
PROB:	0.12	0.77	0.1
RMSE:	18.4	13.02	13.02

MAX_MEAN: 34.4 MAX_MED: 35.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.88

FLAGS: 17; 11; 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,4,6-Tris(tert-butyl)phenol
CHID: 21311 CASRN: 732-26-3
SPID(S): TP0001412E09
M4ID: 30204432

HILL MODEL (in red):

	tp	ga	gw
val:	15.8	-0.888	3.49
sd:	4.99	0.555	8.73

GAIN-LOSS MODEL (in blue):

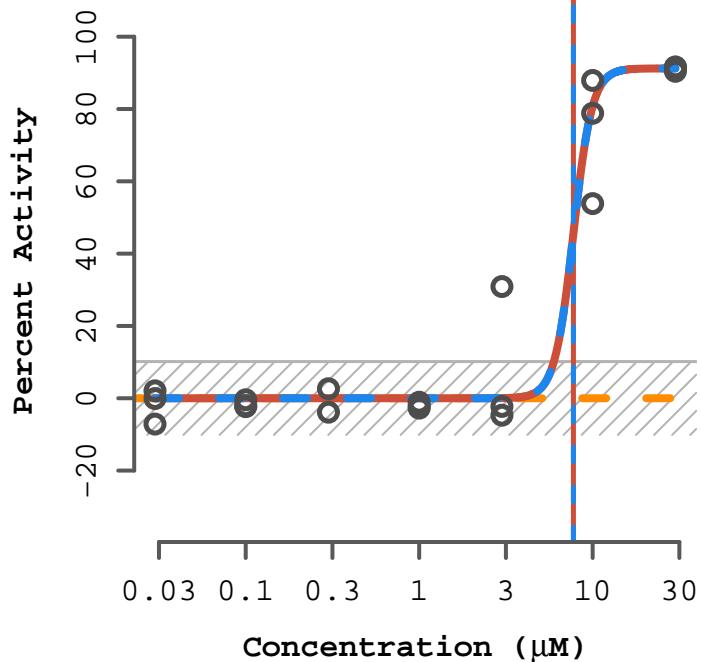
	tp	ga	gw	la	lw
val:	37.7	-0.599	2.56	0.397	0.305
sd:	62	0.363	2.48	4.53	0.424

	CNST	HILL	GNLS
AIC:	167.19	162.27	165.61
PROB:	0.07	0.79	0.15
RMSE:	17.8	13.36	13.39

MAX_MEAN: 26.1 MAX_MED: 26.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.93

FLAGS: 17; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bithionol

CHID: 21342 CASRN: 97-18-7

SPID(S): EPAPLT0154D10

M4ID: 30204193

HILL MODEL (in red):

tp	ga	gw
val: 91.2	0.889	8
sd: 1.97	0.116	8.23

GAIN-LOSS MODEL (in blue):

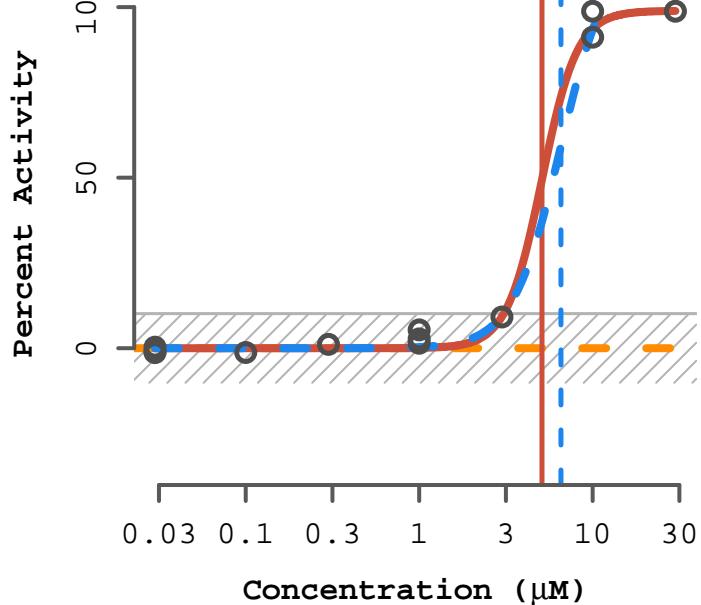
tp	ga	gw	la	lw
val: 91.2	0.889	8	1.86	10.1
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 210.28	140.72	144.72
PROB: 0	0.88	0.12
RMSE: 46.31	9.64	9.64

MAX_MEAN: 91.2 MAX_MED: 91.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tris(methylphenyl) phosphate
CHID: 21391 CASRN: 1330-78-5
SPID(S): EPAPLT0169C01
M4ID: 30204282

HILL MODEL (in red):

	tp	ga	gw
val:	98.9	0.708	4.27
sd:	1.73	0.044	0.807

GAIN-LOSS MODEL (in blue):

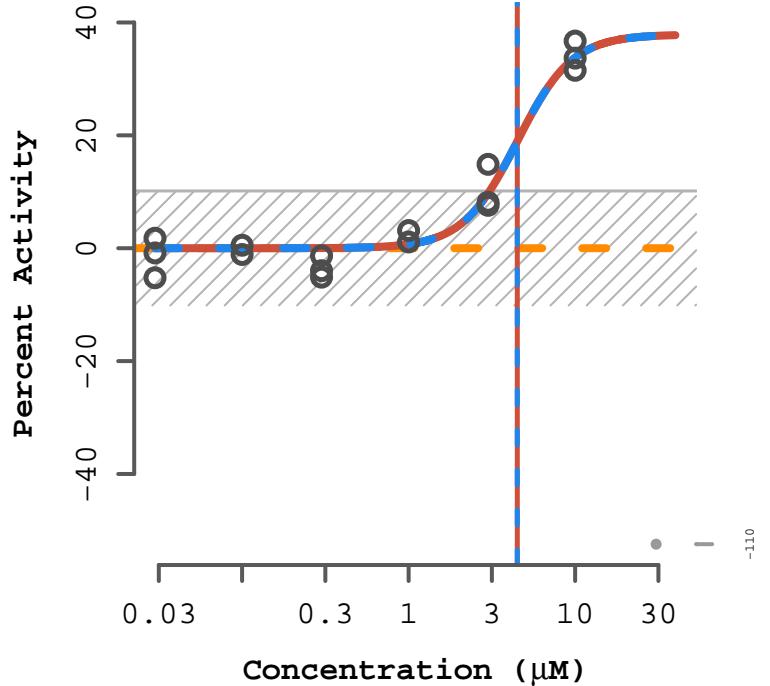
	tp	ga	gw	la	lw
val:	119	0.817	3.12	1.54	12.2
sd:	8.26	0.0391	0.324	0.00604	3.03

	CNST	HILL	GNLS
AIC:	126.27	63.71	66.94
PROB:	0	0.83	0.17
RMSE:	48.28	2.5	2.45

MAX_MEAN: 98.8 MAX_MED: 98.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,2-Bis(4-hydroxyphenyl)-1,1,1-trichloroethane
 CHID: 22325 CASRN: 2971-36-0
 SPID(S): TP0001411C04
 M4ID: 30204335 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	37.9	0.653	2.63
sd:	10.8	0.188	1.86

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	37.9	0.653	2.63	3.57	11.7
sd:	NA	NA	NA	NA	NA

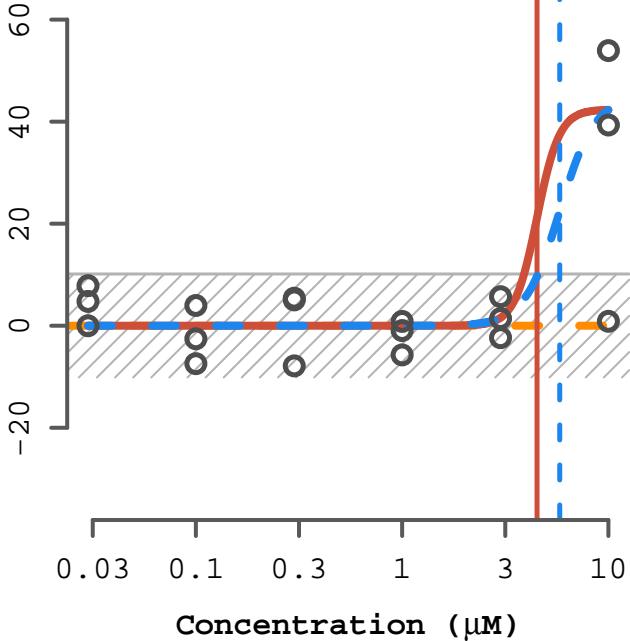
	CNST	HILL	GNLS
AIC:	155.06	125.3	129.3
PROB:	0	0.88	0.12
RMSE:	30.56	35.8	35.8

MAX_MEAN: 34 MAX_MED: 33.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 10; 7

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-(Butan-2-yl)phenol
CHID: 22331 CASRN: 89-72-5
SPID(S): TP0001414C06
M4ID: 30204391

HILL MODEL (in red):

tp	ga	gw
val:	42.4	0.654
sd:	13.1	2.97
		133

GAIN-LOSS MODEL (in blue):

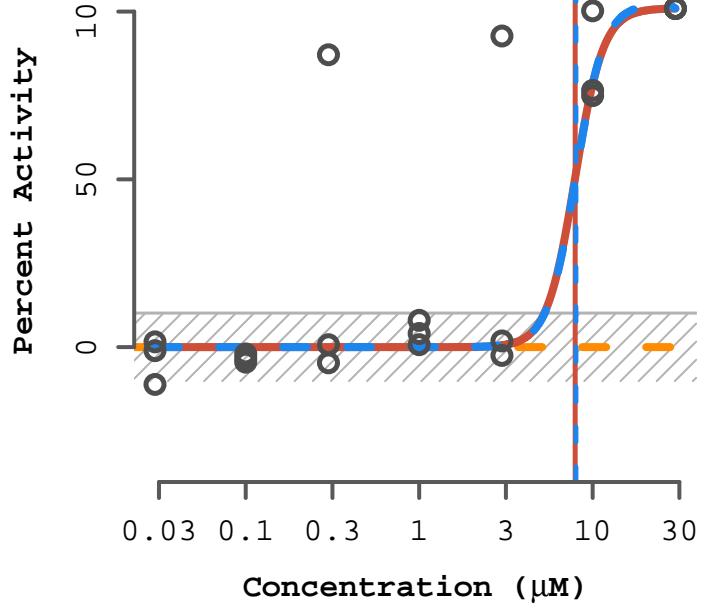
tp	ga	gw	la	lw
val:	45	0.764	5.1	3
sd:	128	2.38	37.3	101
				73

CNST	HILL	GNLS
AIC:	140.14	134.68
PROB:	0.05	0.83
RMSE:	16.35	11.07

MAX_MEAN: 31.4 MAX_MED: 39.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.95

FLAGS: 17; 10; 6



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,2'-Bisphenol F
CHID: 22446 CASRN: 2467-02-9
SPID(S): EPAPLT0154E06
M4ID: 30204200

HILL MODEL (in red):

	tp	ga	gw
val:	101	0.899	5.36
sd:	3.13	0.123	6.54

GAIN-LOSS MODEL (in blue):

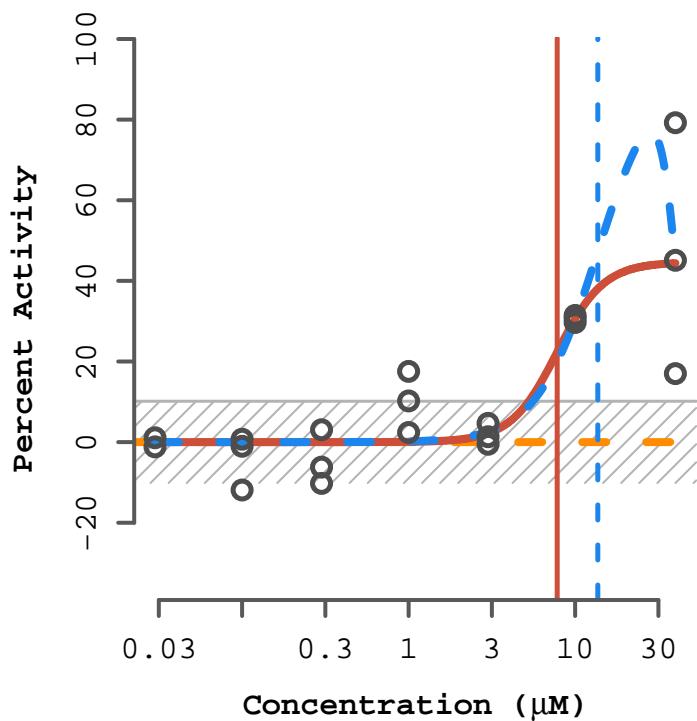
	tp	ga	gw	la	lw
val:	102	0.904	5.33	1.6	15.6
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	231.75	180.55	184.55
PROB:	0	0.88	0.12
RMSE:	57.13	28.34	28.34

MAX_MEAN: 101 MAX_MED: 101 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Metolachlor

CHID: 22448 CASRN: 51218-45-2

SPID(S): TP0001411E06

M4ID: 30204317

HILL MODEL (in red):

tp	ga	gw
44.7	0.892	3.11
6.75	0.0826	1.61

GAIN-LOSS MODEL (in blue):

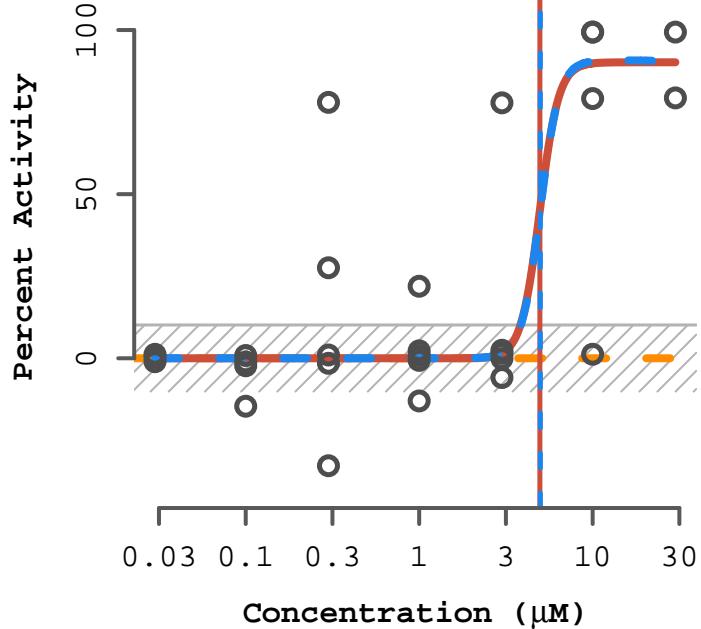
tp	ga	gw	la	lw
94.6	1.14	2.36	1.6	9.09
NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
181.68	156.11	160.05
0	0.88	0.12
24.65	11.6	11.59

MAX_MEAN: 47.1 MAX_MED: 45.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Mifepristone

CHID: 23322 CASRN: 84371-65-3

SPID(S): EPAPLT0167D09

M4ID: 30204265

HILL MODEL (in red):

tp	ga	gw
val: 90.2	0.696	8
sd: 7.3	0.415	14.9

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 90.8	0.699	8	1.71	7.84
sd: NaN	NaN	NaN	NaN	NaN

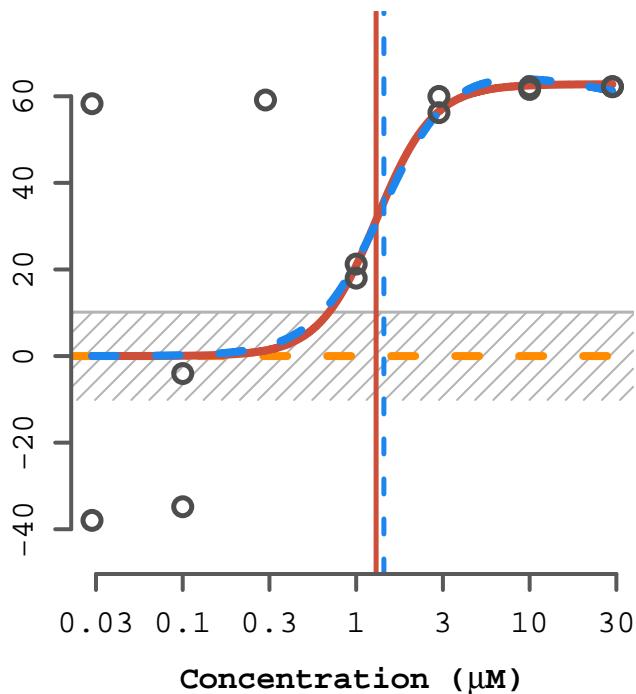
CNST	HILL	GNLS
AIC: 290.6	259.73	263.72
PROB: 0	0.88	0.12
RMSE: 45.15	28.67	28.74

MAX_MEAN: 89.4 MAX_MED: 89.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Acephate

CHID: 23846 CASRN: 30560-19-1

SPID(S): EPAPLT0167A01

M4ID: 30204257

HILL MODEL (in red):

	tp	ga	gw
val:	62.8	0.115	2.63
sd:	7.9	0.134	2.47

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	74.6	0.16	2.18	3.48	0.332
sd:	NaN	NaN	NaN	NaN	NaN

CNST

AIC: 143.43

PROB: 0

RMSE: 50

HILL

128.62

0.88

26.95

GNLS

132.59

0.12

26.83

MAX_MEAN: 62.2

MAX_MED: 62.2

BMAD: 3.38

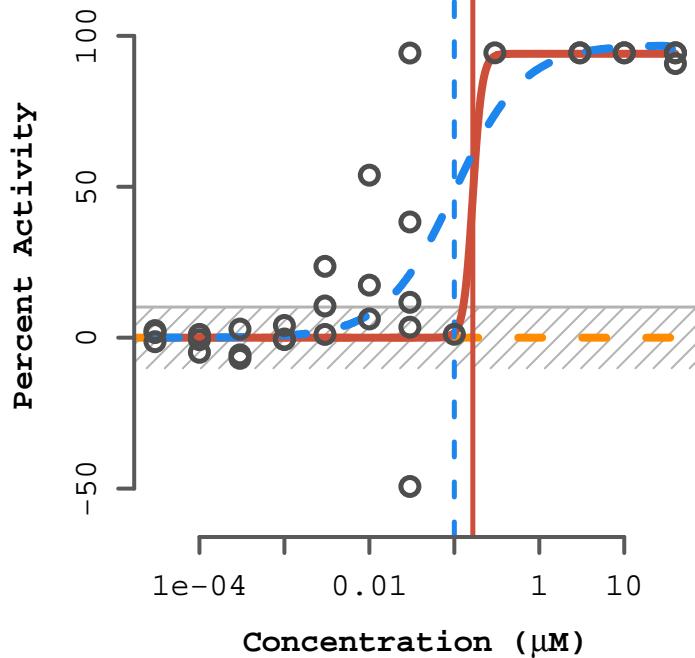
COFF: 10.1

HIT-CALL: 1

FITC: 41

ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Abamectin

CHID: 23892 CASRN: 71751-41-2

SPID(S): TP0001414B08

M4ID: 30204634

HILL MODEL (in red):

	tp	ga	gw
val:	94.1	-0.782	8
sd:	2.52	0.181	6.28

GAIN-LOSS MODEL (in blue):

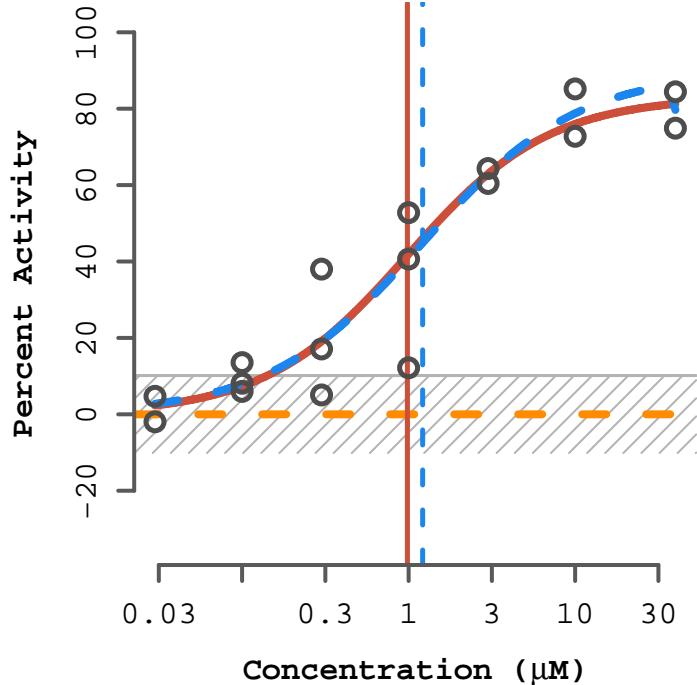
	tp	ga	gw	la	lw
val:	97	-1	1.06	1.71	13.4
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	345.35	272.99	284.9
PROB:	0	1	0
RMSE:	52.33	23.06	22.76

MAX_MEAN: 94.4 MAX_MED: 94.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fenpropathrin

CHID: 24002 CASRN: 39515-41-8

SPID(S): TP0001413F11

M4ID: 30204294

HILL MODEL (in red):

tp	ga	gw
val: 83.1	-0.00732	1.02
sd: 5.5	0.11	0.182

GAIN-LOSS MODEL (in blue):

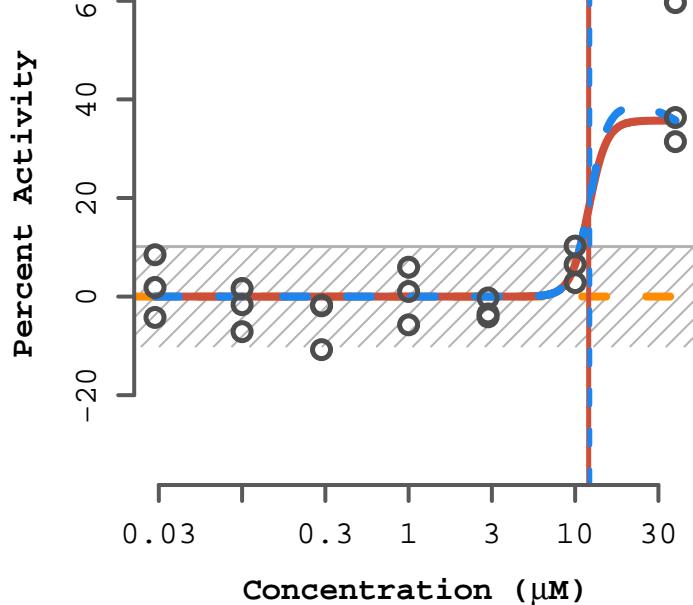
tp	ga	gw	la	lw
val: 89.7	0.0843	0.935	1.7	11.4
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 183.77	132.17	135.78
PROB: 0	0.86	0.14
RMSE: 48.36	10.24	10.04

MAX_MEAN: 79.7 MAX_MED: 79.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ethion

CHID: 24086 CASRN: 563-12-2

SPID(S): TP0001411G07

M4ID: 30204464

HILL MODEL (in red):

	tp	ga	gw
val:	35.7	1.08	8
sd:	3.38	0.832	81.8

GAIN-LOSS MODEL (in blue):

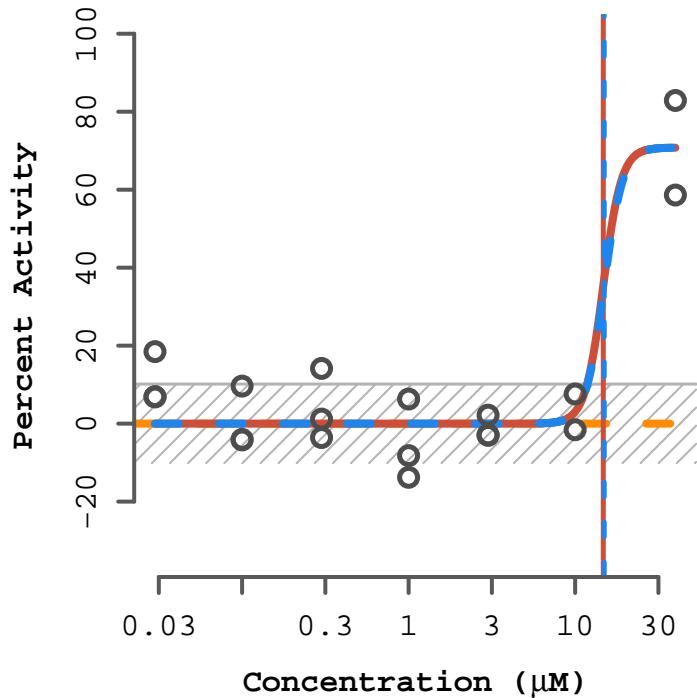
	tp	ga	gw	la	lw
val:	39.4	1.09	7.98	1.94	2.94
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	170.14	142.61	146.61
PROB:	0	0.88	0.12
RMSE:	17.44	6.86	6.86

MAX_MEAN: 42.5 MAX_MED: 36.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Phosalone

CHID: 24259 CASRN: 2310-17-0

SPID(S): TP0001412H06

M4ID: 30204349

HILL MODEL (in red):

	tp	ga	gw
val:	70.8	1.17	7.99
sd:	14	2.73	129

GAIN-LOSS MODEL (in blue):

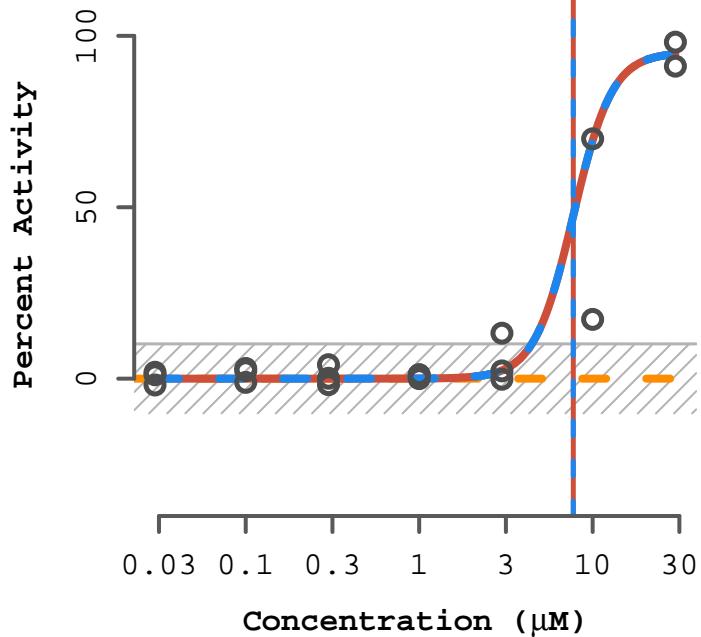
	tp	ga	gw	la	lw
val:	70.8	1.17	7.73	1.88	12
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	164.61	145.9	149.9
PROB:	0	0.88	0.12
RMSE:	24.57	8.68	8.68

MAX_MEAN: 70.8 MAX_MED: 70.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tri-allate

CHID: 24344 CASRN: 2303-17-5

SPID(S): EPAPLT0167F05

M4ID: 30204275

HILL MODEL (in red):

	tp	ga	gw
val:	95.2	0.889	3.86
sd:	3.35	0.0259	0.767

GAIN-LOSS MODEL (in blue):

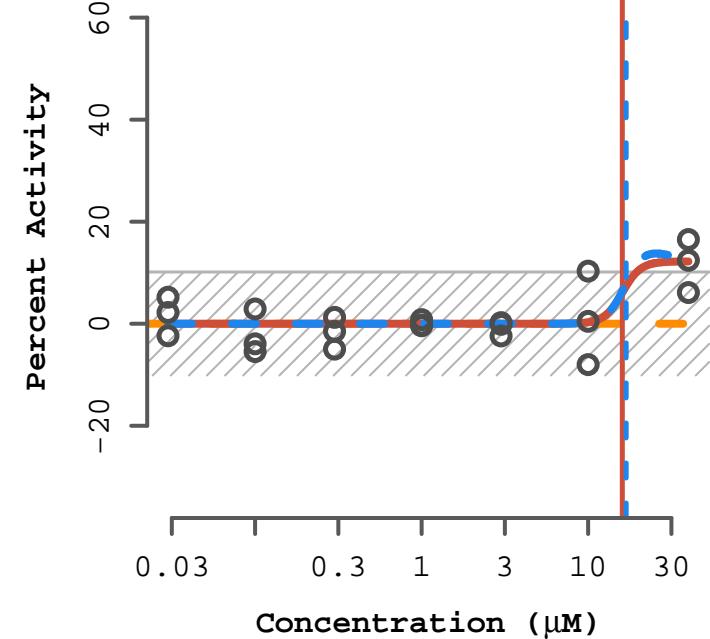
	tp	ga	gw	la	lw
val:	95.2	0.889	3.86	2.43	12.1
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	176.41	125.3	129.3
PROB:	0	0.88	0.12
RMSE:	35.07	12.38	12.38

MAX_MEAN: 94.7 MAX_MED: 94.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dicumyl peroxide
CHID: 25017 CASRN: 80-43-3
SPID(S): TP0001411D08
M4ID: 30204399

HILL MODEL (in red):

	tp	ga	gw
val:	12.2	1.21	7.92
sd:	5	9.17	351

GAIN-LOSS MODEL (in blue):

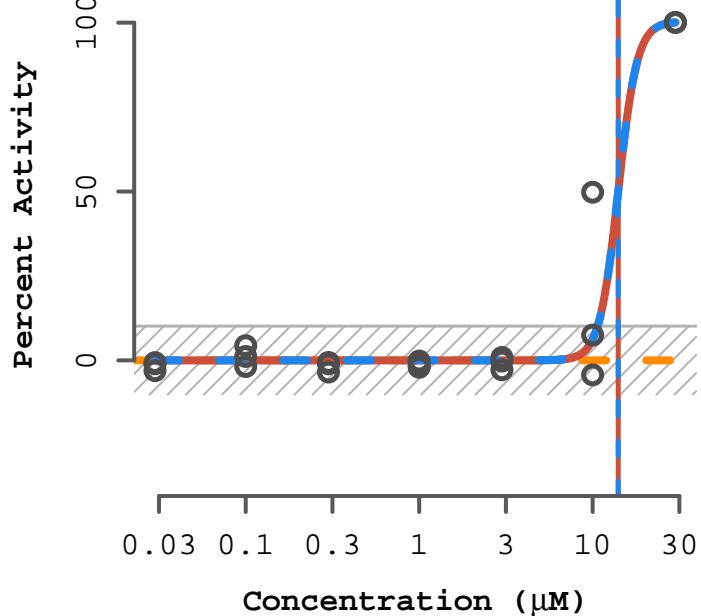
	tp	ga	gw	la	lw
val:	19.4	1.23	7.8	1.8	1.13
sd:	2350	9.7	330	65.2	362

	CNST	HILL	GNLS
AIC:	135.02	126.86	130.86
PROB:	0.01	0.87	0.12
RMSE:	6.02	4.09	4.09

MAX_MEAN: 11.7 MAX_MED: 12.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.99

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Ethylhexyl diphenyl phosphate

CHID: 25300 CASRN: 1241-94-7

SPID(S): EPAPLT0169D05

M4ID: 30204234

HILL MODEL (in red):

	tp	ga	gw
val:	100	1.15	7.96
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

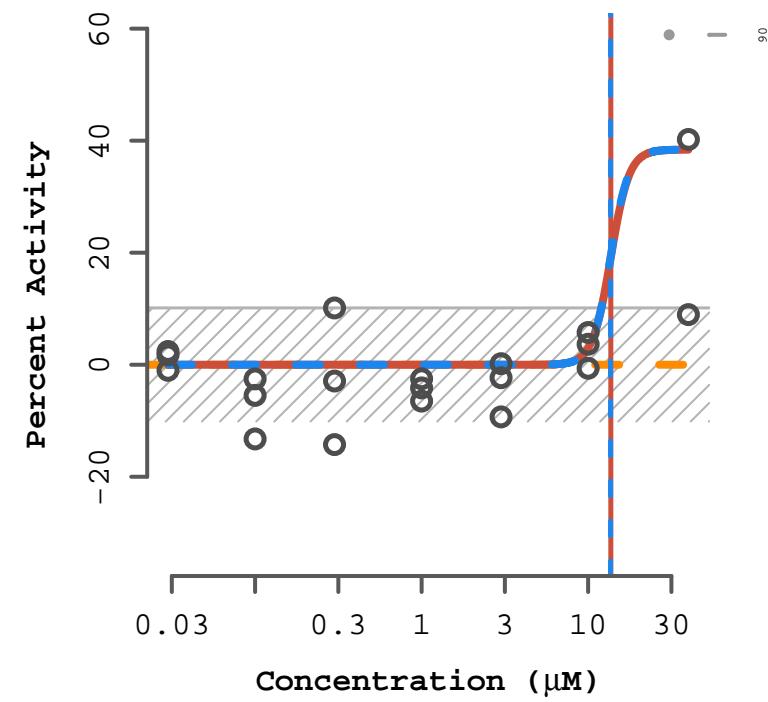
	tp	ga	gw	la	lw
val:	100	1.15	7.96	2.94	11.5
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	174.05	124.54	128.54
PROB:	0	0.88	0.12
RMSE:	33.65	10.17	10.17

MAX_MEAN: 100 MAX_MED: 100 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Methylene bis(thiocyanate)

CHID: 25599 CASRN: 6317-18-6

SPID(S): TP0001411F11

M4ID: 30204637 BRK

HILL MODEL (in red):

tp	ga	gw
38.4	1.14	7.9
sd: 6.98	1.89	110

GAIN-LOSS MODEL (in blue):

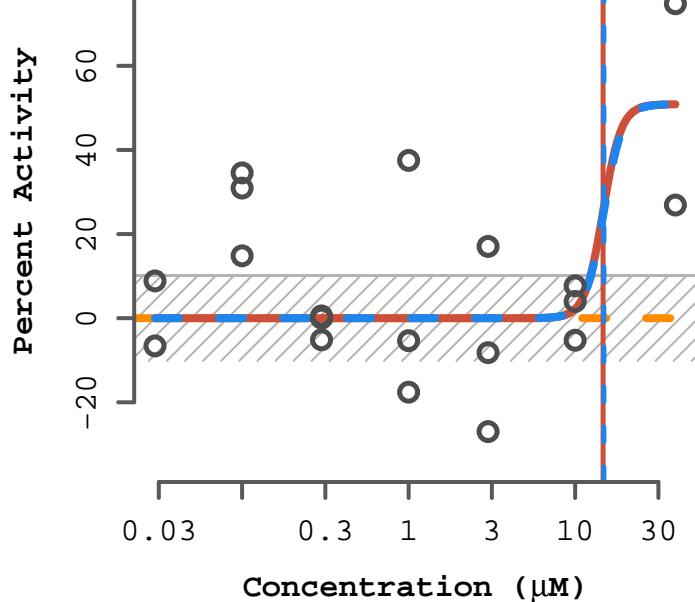
tp	ga	gw	la	lw
38.4	1.13	8	3.18	7.73
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 171.11	167.03	171.03
PROB: 0.1	0.79	0.11
RMSE: 22.37	14.19	14.19

MAX_MEAN: 46.3 MAX_MED: 40.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.9

FLAGS: 17; 11; 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 3,3',5,5'-Tetrabromobisphenol A

CHID: 26081 CASRN: 79-94-7

SPID(S): TP0001412E07

M4ID: 30204587

HILL MODEL (in red):

	tp	ga	gw
val:	50.9	1.17	7.97
sd:	25.2	4.91	233

GAIN-LOSS MODEL (in blue):

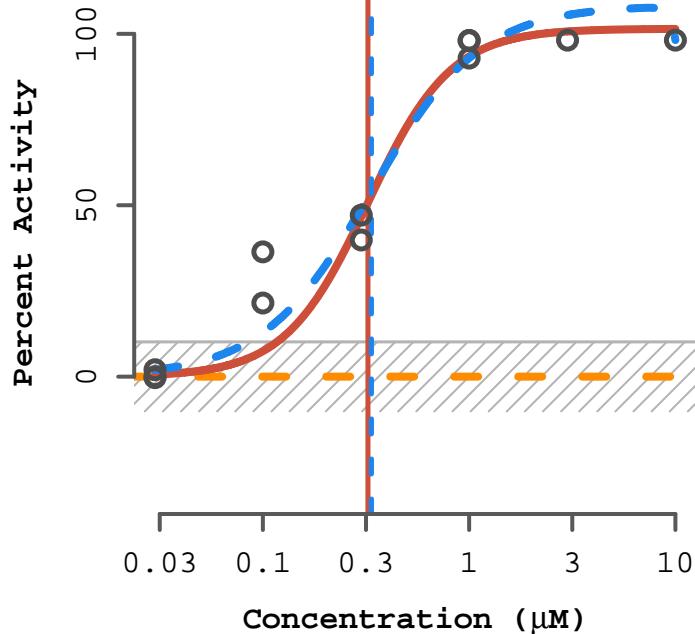
	tp	ga	gw	la	lw
val:	50.9	1.17	7.75	3.57	9.76
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	175.78	174.35	178.35
PROB:	0.3	0.62	0.08
RMSE:	24.89	18.62	18.62

MAX_MEAN: 50.8 MAX_MED: 50.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.7

FLAGS: 16; 11; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tributyltin chloride
CHID: 27403 CASRN: 1461-22-9
SPID(S): EPAPLT0167G09
M4ID: 30204277

HILL MODEL (in red):

tp	ga	gw
val: 101	-0.491	2.16
sd: 5.01	0.0383	1

GAIN-LOSS MODEL (in blue):

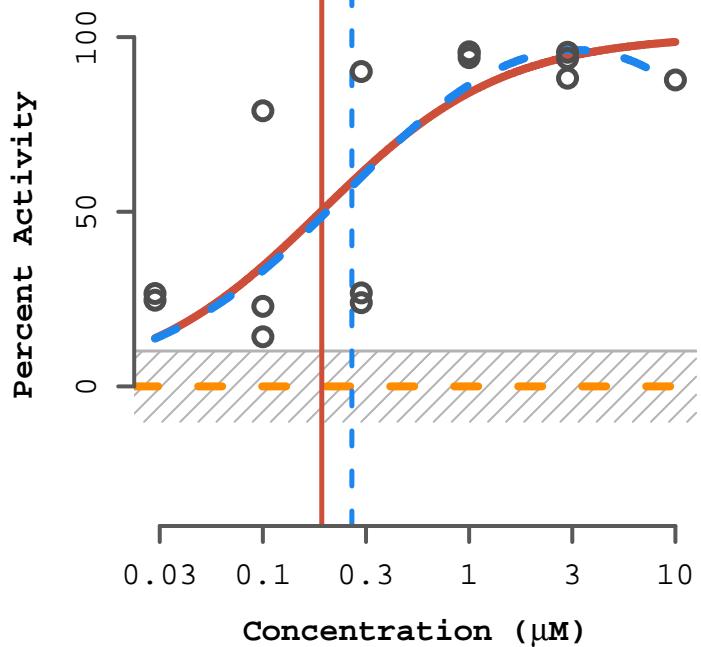
tp	ga	gw	la	lw
val: 108	-0.474	1.65	1.06	18
sd: 9.28	0.0658	0.482	0.0603	23.1

CNST	HILL	GNLS
AIC: 138.22	91.69	94.39
PROB: 0	0.79	0.21
RMSE: 66.3	9.78	8.32

MAX_MEAN: 98.2 MAX_MED: 98.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tributyltin chloride
 CHID: 27403 CASRN: 1461-22-9
 SPID(S): TP0001414G06
 M4ID: 30204571

HILL MODEL (in red):

tp	ga	gw
val: 101	-0.714	0.988
sd: 12.3	0.256	0.464

GAIN-LOSS MODEL (in blue):

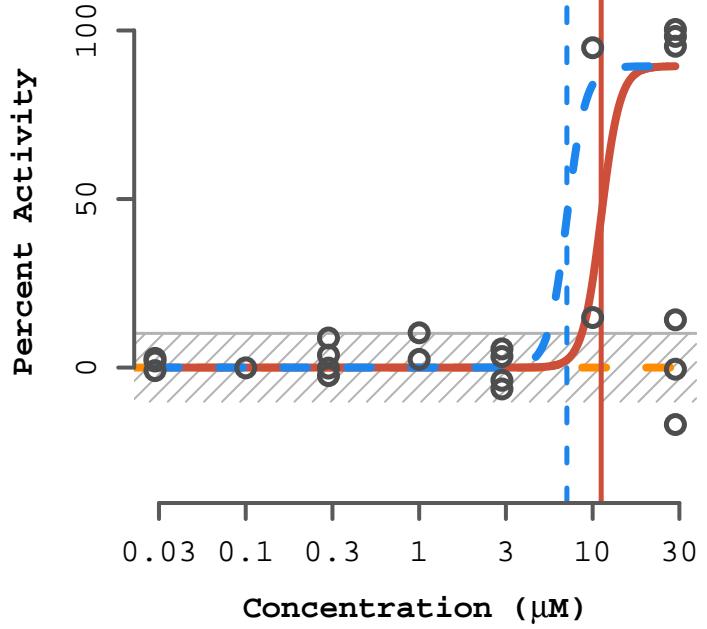
tp	ga	gw	la	lw
val: 115	-0.568	0.91	1.49	1.12
sd: 35.9	0.371	0.42	0.679	1.59

CNST	HILL	GNLS
AIC: 175.6	142.95	145.81
PROB: 0	0.81	0.19
RMSE: 72.25	21.17	20.75

MAX_MEAN: 95.1 MAX_MED: 95.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Triphenyl phosphates isopropylated

CHID: 28880 CASRN: 68937-41-7

SPID(S): EPAPLT0169A03

M4ID: 30204280

HILL MODEL (in red):

	tp	ga	gw
val:	89.4	1.05	7.99
sd:	14.1	0.978	158

GAIN-LOSS MODEL (in blue):

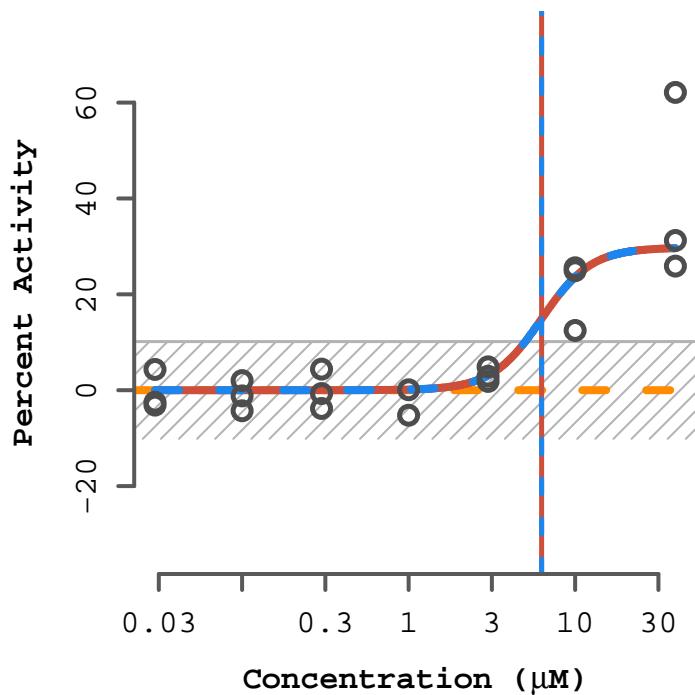
	tp	ga	gw	la	lw
val:	89.4	0.852	8	3.15	8.26
sd:	12.8	0.494	25.7	330000	868000

	CNST	HILL	GNLS
AIC:	218.24	217.45	221.45
PROB:	0.37	0.55	0.07
RMSE:	42.01	37.22	37.21

MAX_MEAN: 54.9 MAX_MED: 54.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.63

FLAGS: 16; 11; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,4-Bis(1-methyl-1-phenylethyl)phenol

CHID: 29241 CASRN: 2772-45-4

SPID(S): TP0001411B05

M4ID: 30204582

HILL MODEL (in red):

	tp	ga	gw
val:	29.8	0.799	2.92
sd:	2.95	0.082	0.771

GAIN-LOSS MODEL (in blue):

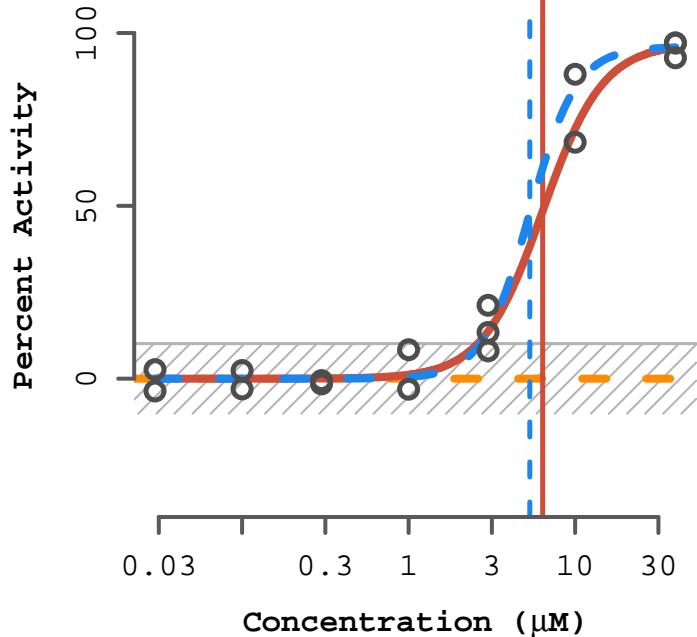
	tp	ga	gw	la	lw
val:	29.8	0.799	2.92	2.32	8.52
sd:	2.95	0.082	0.771	394	4660

CNST	HILL	GNLS
AIC: 169.31	132.33	136.33
PROB: 0	0.88	0.12
RMSE: 18.83	8.15	8.15

MAX_MEAN: 39.8 MAX_MED: 31.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Clotrimazole

CHID: 29871 CASRN: 23593-75-1

SPID(S): TP0001411E01

M4ID: 30204482

HILL MODEL (in red):

	tp	ga	gw
val:	97	0.805	2.39
sd:	2.68	0.0495	0.381

GAIN-LOSS MODEL (in blue):

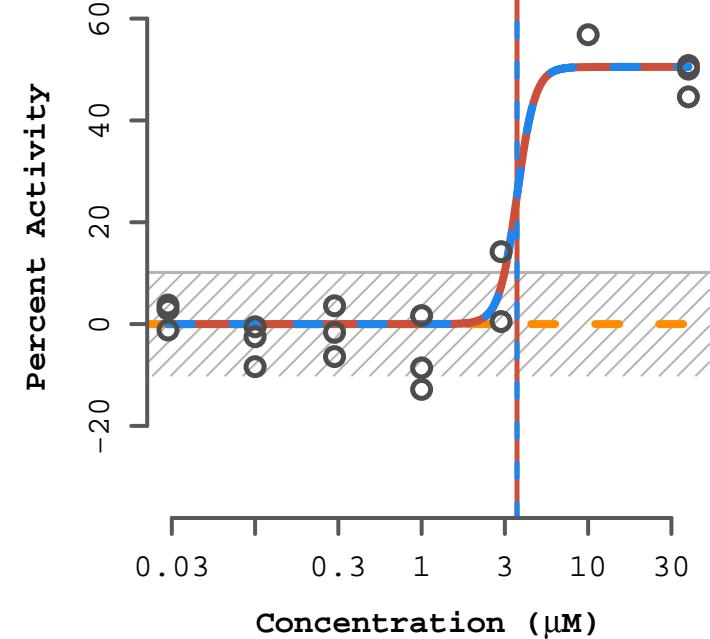
	tp	ga	gw	la	lw
val:	96	0.728	3.11	3.15	12.8
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	171.87	105.77	109.84
PROB:	0	0.88	0.12
RMSE:	50.46	5.43	5.44

MAX_MEAN: 95.7 MAX_MED: 97.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cycloate

CHID: 32356 CASRN: 1134-23-2

SPID(S): TP0001412B06

M4ID: 30204296

HILL MODEL (in red):

	tp	ga	gw
val:	50.5	0.573	8
sd:	2.65	0.146	11.6

GAIN-LOSS MODEL (in blue):

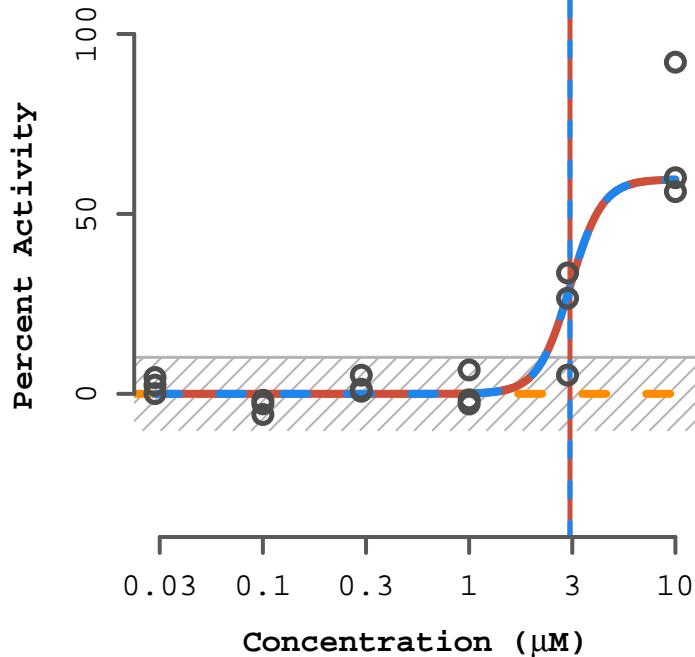
	tp	ga	gw	la	lw
val:	50.5	0.573	8	3.03	6.84
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	164.78	122.55	126.55
PROB:	0	0.88	0.12
RMSE:	24.62	5.61	5.61

MAX_MEAN: 56.8 MAX_MED: 56.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fenarimol

CHID: 32390 CASRN: 60168-88-9

SPID(S): TP0001412C01

M4ID: 30204385

HILL MODEL (in red):

tp	ga	gw
val: 59.6	0.489	5.54
sd: 4.29	0.0645	24.9

GAIN-LOSS MODEL (in blue):

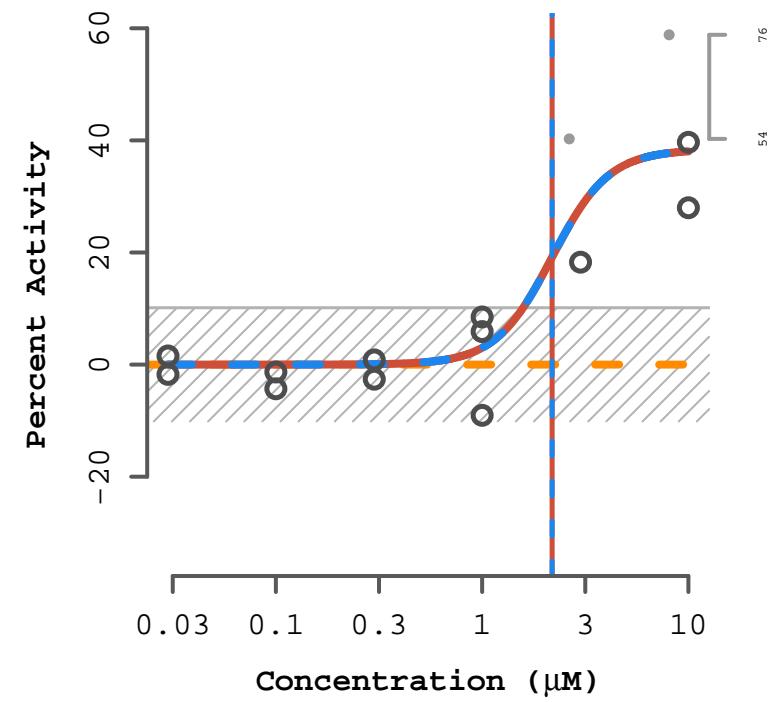
tp	ga	gw	la	lw
val: 59.6	0.489	5.54	1.86	7.16
sd: 4.83	0.0647	24.9	8910	73900

CNST	HILL	GNLS
AIC: 170.06	131.33	135.33
PROB: 0	0.88	0.12
RMSE: 30.99	9.91	9.91

MAX_MEAN: 69.5 MAX_MED: 60.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Triclosan

CHID: 32498 CASRN: 3380-34-5

SPID(S): TP0001413F01

M4ID: 30204657 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	38.3	0.339	3.16
sd:	7.15	0.277	3.08

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	38.4	0.339	3.16	1.54	8.06
sd:	7.64	0.278	3.09	198	2940

CNST

AIC: 133.05

PROB: 0

RMSE: 28.77

HILL

116.4

0.88

13.42

GNLS

120.4

0.12

13.42

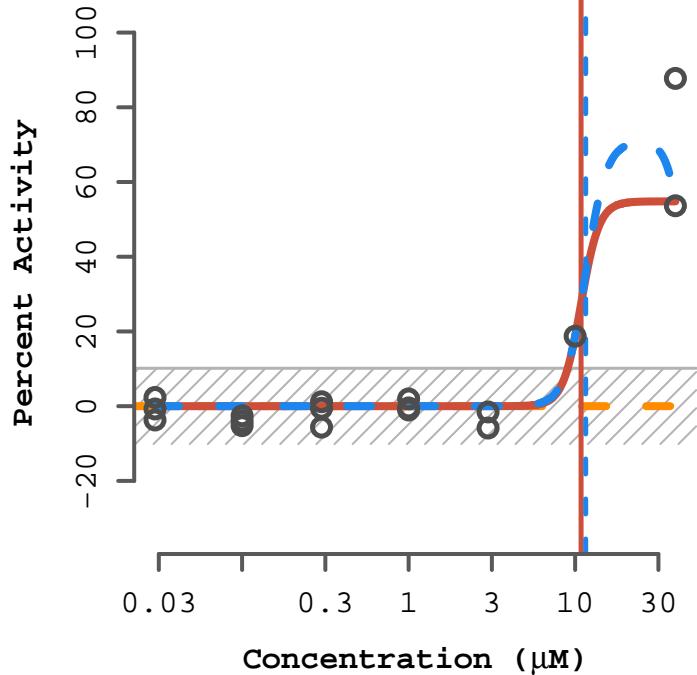
MAX_MEAN: 48

MAX_MED: 39.7

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Azoxystrobin

CHID: 32520 CASRN: 131860-33-8

SPID(S): TP0001413C01

M4ID: 30204321

HILL MODEL (in red):

	tp	ga	gw
val:	54.8	1.04	8
sd:	3.03	0.0308	6.83

GAIN-LOSS MODEL (in blue):

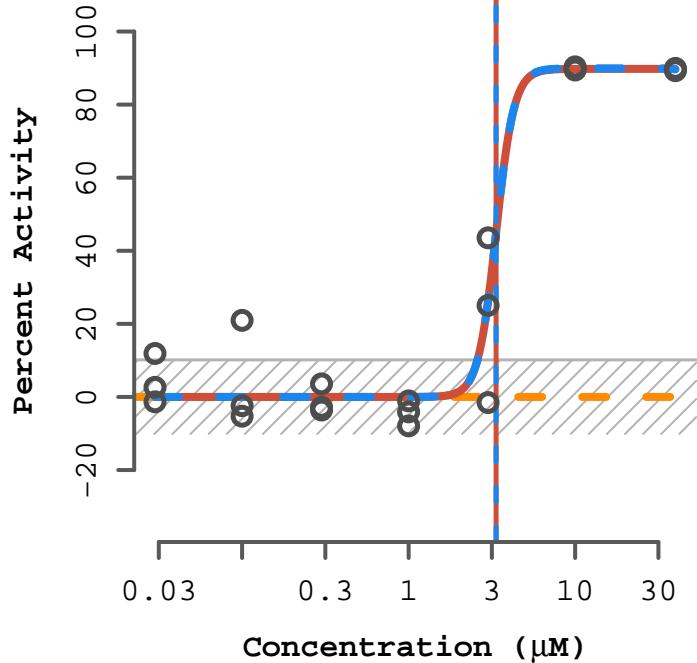
	tp	ga	gw	la	lw
val:	71.1	1.06	7.04	1.66	8.89
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	142.25	111.62	115.62
PROB:	0	0.88	0.12
RMSE:	25.52	8.5	8.51

MAX_MEAN: 70.7 MAX_MED: 70.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fenpyroximate (Z,E)

CHID: 32550 CASRN: 111812-58-9

SPID(S): TP0001412F08

M4ID: 30204338

HILL MODEL (in red):

tp	ga	gw
val: 89.8	0.525	8
sd: 1.86	0.0872	14.3

GAIN-LOSS MODEL (in blue):

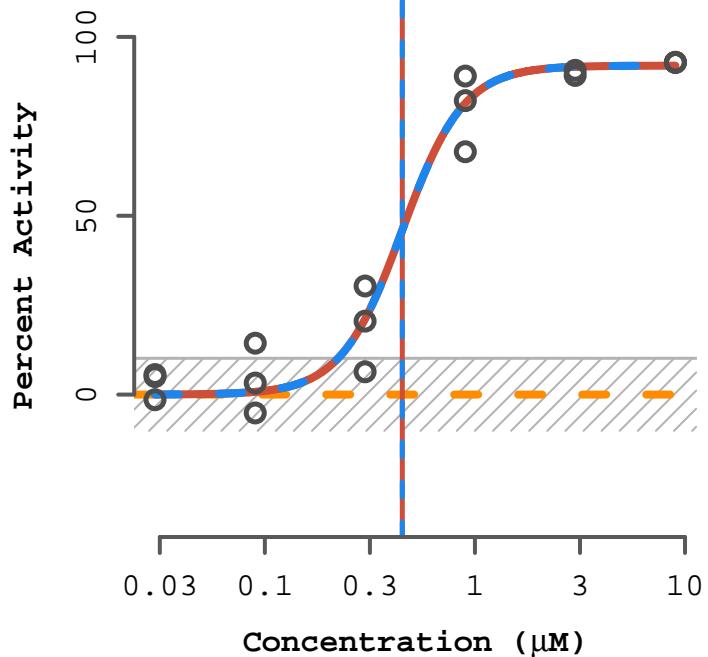
tp	ga	gw	la	lw
val: 89.9	0.525	8	1.77	14.4
sd: 2.65	0.0921	15	1.06	97.5

CNST	HILL	GNLS
AIC: 224.66	154.25	158.25
PROB: 0	0.88	0.12
RMSE: 49.58	9.28	9.28

MAX_MEAN: 89.9 MAX_MED: 89.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fluazinam

CHID: 32551 CASRN: 79622-59-6

SPID(S): TP0001412A04

M4ID: 30204313

HILL MODEL (in red):

tp	ga	gw
92	-0.346	2.96
2.28	0.0417	0.583

GAIN-LOSS MODEL (in blue):

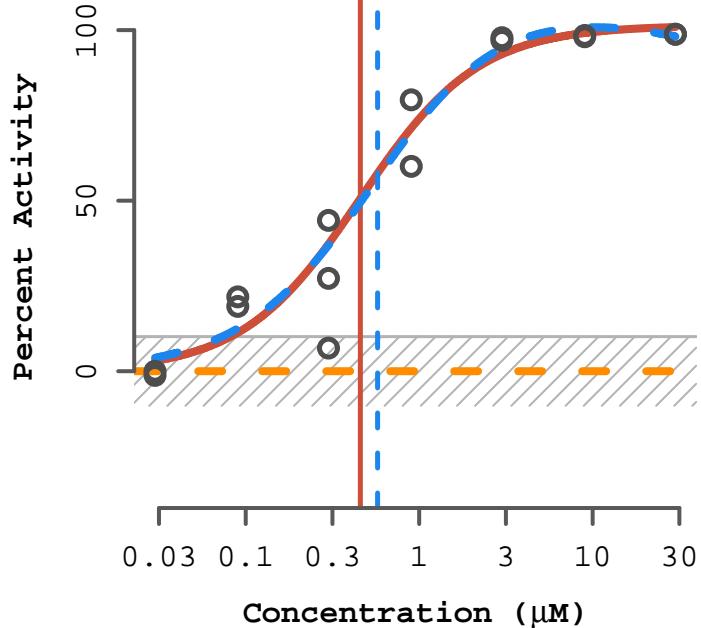
tp	ga	gw	la	lw
92	-0.346	2.96	1.81	14.7
2.28	0.0417	0.583	116000	2020000

CNST	HILL	GNLS
AIC: 191.78	122.43	126.43
PROB: 0	0.88	0.12
RMSE: 60.91	7.03	7.03

MAX_MEAN: 92.9 MAX_MED: 92.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: ..-Cyhalothrin
 CHID: 32559 CASRN: 91465-08-6
 SPID(S): EPAPLT0167H06
 M4ID: 30204236

HILL MODEL (in red):

tp	ga	gw
val: 101	-0.339	1.28
sd: 3.53	0.0959	0.265

GAIN-LOSS MODEL (in blue):

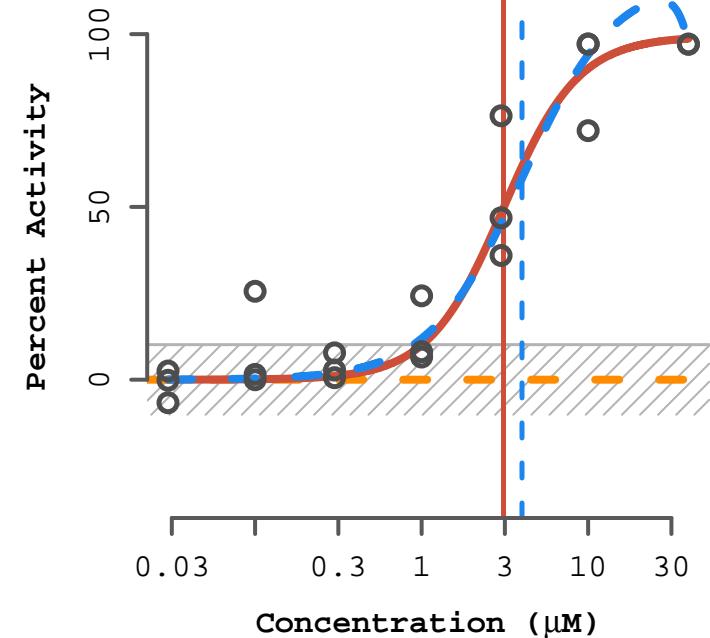
tp	ga	gw	la	lw
val: 119	-0.24	1.14	3.46	0.359
sd: 43.5	0.179	0.252	2.62	0.776

CNST	HILL	GNLS
AIC: 173.73	117.19	120.04
PROB: 0	0.81	0.19
RMSE: 68.96	10.08	9.81

MAX_MEAN: 98.8 MAX_MED: 98.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pyridaben

CHID: 32573 CASRN: 96489-71-3

SPID(S): TP0001414H07

M4ID: 30204380

HILL MODEL (in red):

tp	ga	gw
99.4	0.492	1.93
4.65	0.0593	0.458

GAIN-LOSS MODEL (in blue):

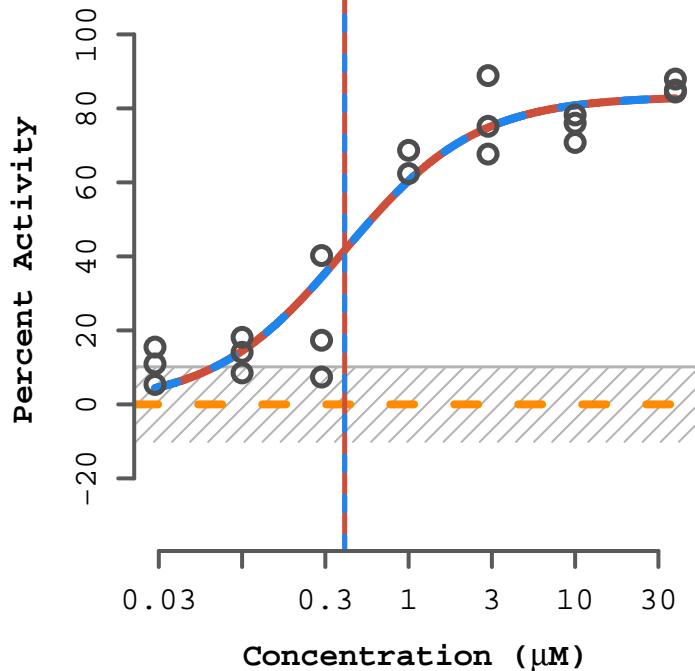
tp	ga	gw	la	lw
116	0.603	1.57	1.71	7.13
28.6	0.194	0.558	0.081	9.15

CNST	HILL	GNLS
AIC: 216.62	156.92	159.69
PROB: 0	0.8	0.2
RMSE: 51.83	10.83	11.13

MAX_MEAN: 97.1 MAX_MED: 97.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tefluthrin

CHID: 32577 CASRN: 79538-32-2

SPID(S): TP0001413D10

M4ID: 30204284

HILL MODEL (in red):

tp	ga	gw
val: 83.3	-0.384	1.1
sd: 3.53	0.0958	0.222

GAIN-LOSS MODEL (in blue):

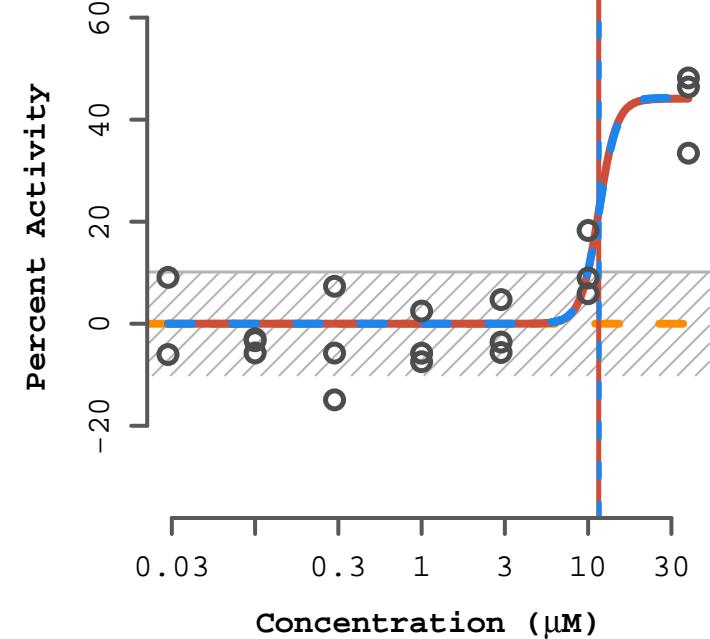
tp	ga	gw	la	lw
val: 83.3	-0.384	1.1	3.16	7.27
sd: 3.53	0.0958	0.222	22400	105000

CNST	HILL	GNLS
AIC: 224.61	152.68	156.68
PROB: 0	0.88	0.12
RMSE: 58.61	9.37	9.37

MAX_MEAN: 85.8 MAX_MED: 85 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pyriproxyfen

CHID: 32640 CASRN: 95737-68-1

SPID(S): TP0001411C01

M4ID: 30204354

HILL MODEL (in red):

	tp	ga	gw
val:	44.1	1.06	8
sd:	4.12	0.392	49.2

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	44.5	1.07	7.71	2.66	1.98
sd:	95	0.409	46.5	93.4	172

CNST

AIC: 170.17

PROB: 0

RMSE: 18.28

HILL

142.45

0.88

6.57

GNLS

146.45

0.12

6.57

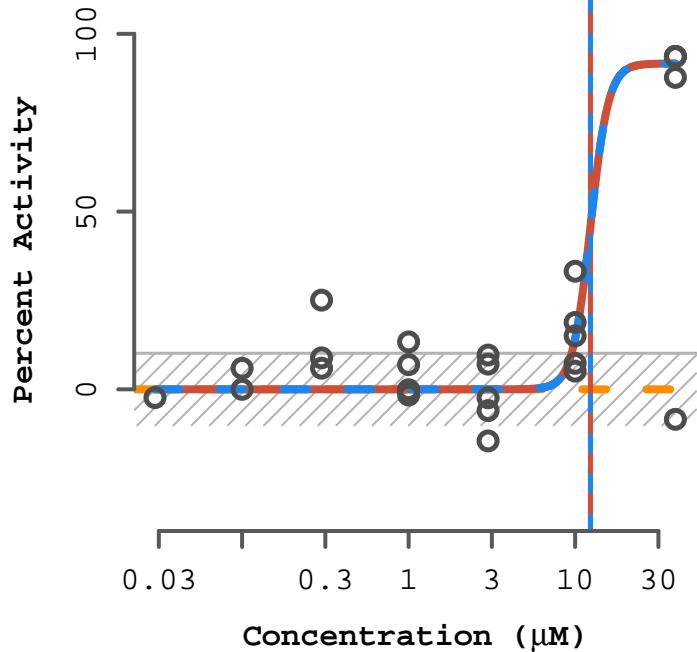
MAX_MEAN: 42.7

MAX_MED: 46.4

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tetramethrin

CHID: 32649 CASRN: 7696-12-0

SPID(S): TP0001412A12

M4ID: 30204315

HILL MODEL (in red):

	tp	ga	gw
val:	91.6	1.09	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

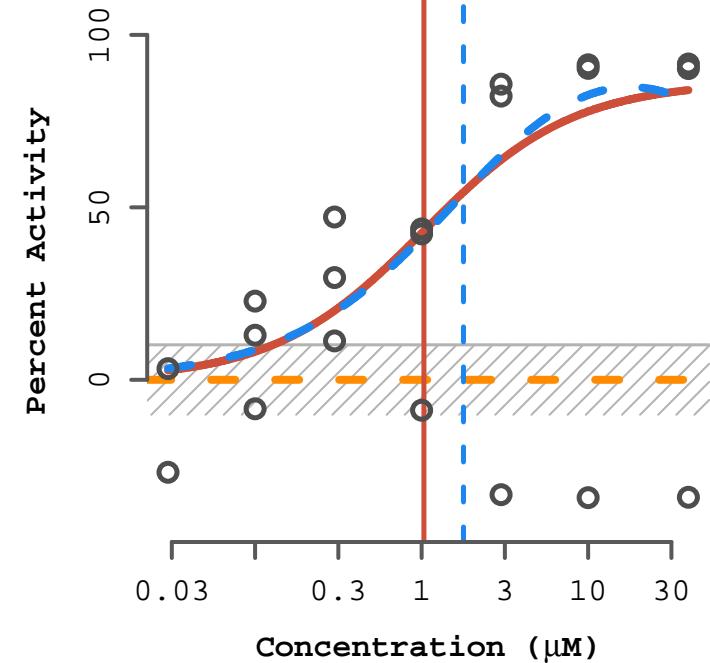
	tp	ga	gw	la	lw
val:	91.6	1.09	8	2.82	11.2
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	254.24	211.83	215.83
PROB:	0	0.88	0.12
RMSE:	37.85	21.45	21.45

MAX_MEAN: 72.1 MAX_MED: 93.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Esfenvalerate

CHID: 32667 CASRN: 66230-04-4

SPID(S): TP0001412D04

M4ID: 30204295

HILL MODEL (in red):

	tp	ga	gw
val:	86.4	0.0138	0.969
sd:	19	0.351	0.518

GAIN-LOSS MODEL (in blue):

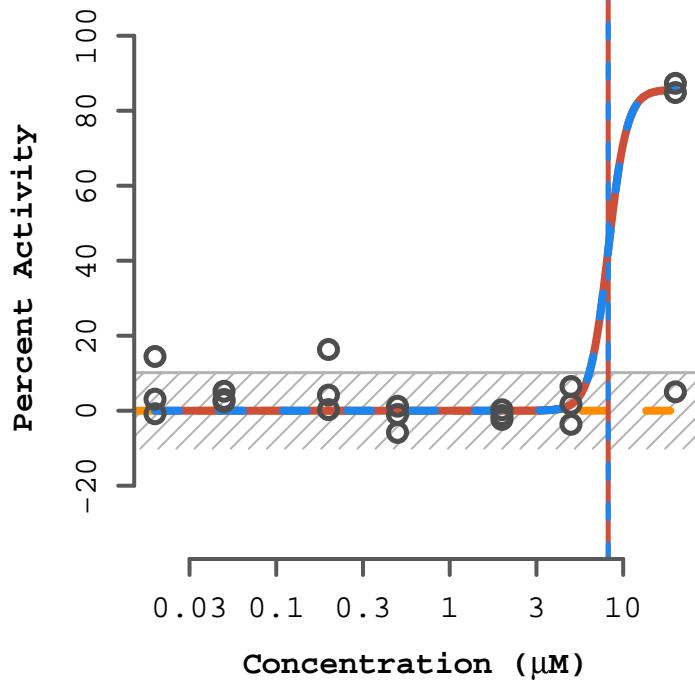
	tp	ga	gw	la	lw
val:	110	0.251	0.851	2.24	0.879
sd:	183	1.36	0.592	2.31	5.98

	CNST	HILL	GNLS
AIC:	220.57	212.22	216.07
PROB:	0.01	0.86	0.13
RMSE:	54.35	45.87	46.03

MAX_MEAN: 49.3 MAX_MED: 90.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.99

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tamoxifen

CHID: 34187 CASRN: 10540-29-1

SPID(S): TP0001412F09

M4ID: 30204568

HILL MODEL (in red):

	tp	ga	gw
val:	85.7	0.914	8
sd:	7.6	2	74.3

GAIN-LOSS MODEL (in blue):

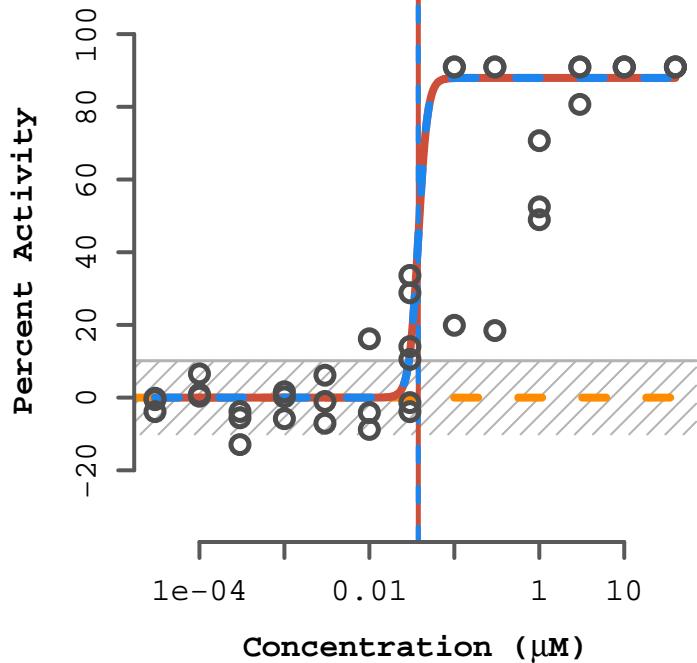
	tp	ga	gw	la	lw
val:	85.7	0.914	8	3.27	14.2
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	174.01	157.14	161.14
PROB:	0	0.88	0.12
RMSE:	27.17	18.43	18.43

MAX_MEAN: 59.1 MAX_MED: 84.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Emamectin benzoate
 CHID: 34566 CASRN: 155569-91-8
 SPID(S): TP0001413H03
 M4ID: 30204382

HILL MODEL (in red):

tp	ga	gw
val: 88	-1.42	8
sd: 2.38	0.24	19.4

GAIN-LOSS MODEL (in blue):

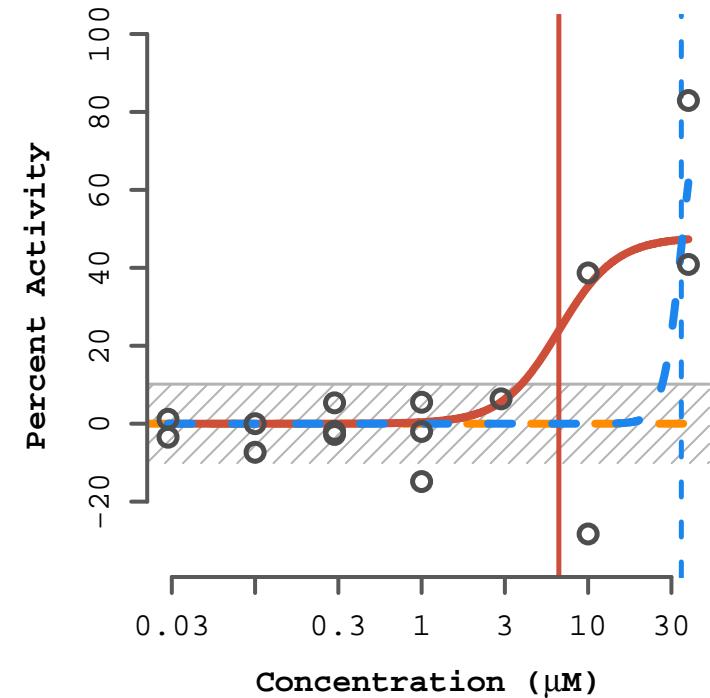
tp	ga	gw	la	lw
val: 88	-1.42	8	3.59	15.9
sd: NA	NA	NA	NA	NA

CNST	HILL	GNLS
AIC: 456.54	347.08	351.08
PROB: 0	0.88	0.12
RMSE: 53.4	18.66	18.66

MAX_MEAN: 91 MAX_MED: 91 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Etoxazole

CHID: 34586 CASRN: 153233-91-1

SPID(S): TP0001411E02

M4ID: 30204660

HILL MODEL (in red):

	tp	ga	gw
val:	47.8	0.824	2.54
sd:	12.8	0.207	1.4

GAIN-LOSS MODEL (in blue):

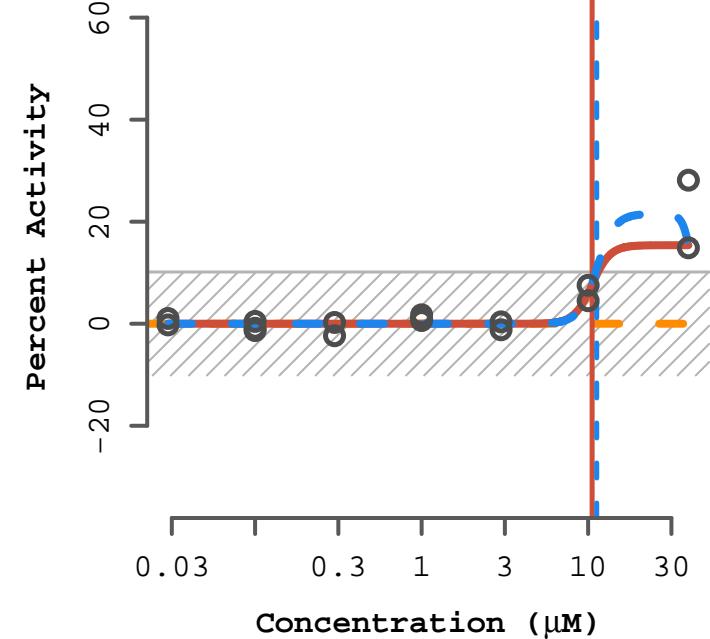
	tp	ga	gw	la	lw
val:	90.6	1.56	7.87	3.07	2.98
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	138.87	129.58	135.47
PROB:	0.01	0.94	0.05
RMSE:	27.39	19.54	15.46

MAX_MEAN: 61.9 MAX_MED: 61.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 0.99

FLAGS: 17; 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fenamidone

CHID: 34590 CASRN: 161326-34-7

SPID(S): TP0001411F09

M4ID: 30204320

HILL MODEL (in red):

	tp	ga	gw
val:	15.4	1.02	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

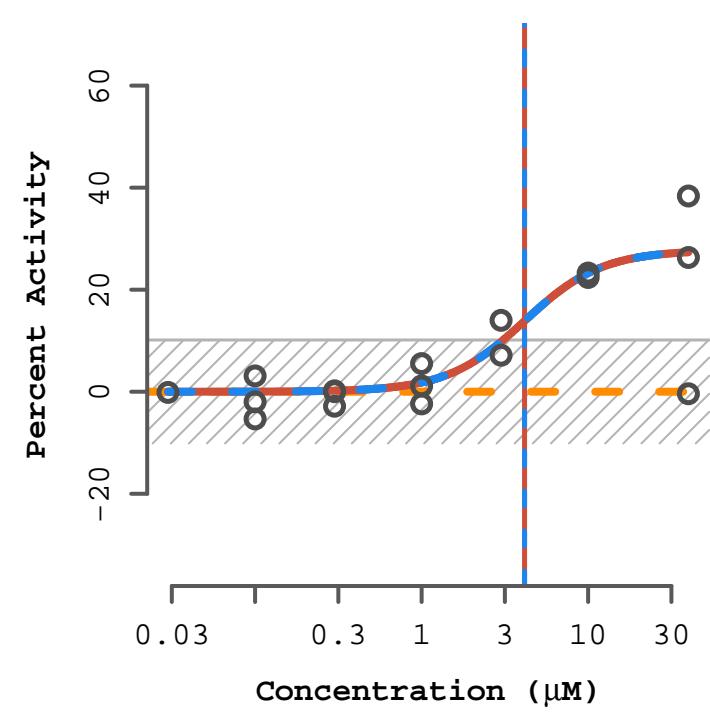
	tp	ga	gw	la	lw
val:	21.6	1.05	7.98	1.63	16.6
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	103.11	76.56	80.56
PROB:	0	0.88	0.12
RMSE:	8.32	3.38	3.38

MAX_MEAN: 21.5 MAX_MED: 21.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Allethrin

CHID: 35180 CASRN: 584-79-2

SPID(S): TP0001414A08

M4ID: 30204636

HILL MODEL (in red):

	tp	ga	gw
val:	27.7	0.618	1.87
sd:	4.18	0.147	0.672

GAIN-LOSS MODEL (in blue):

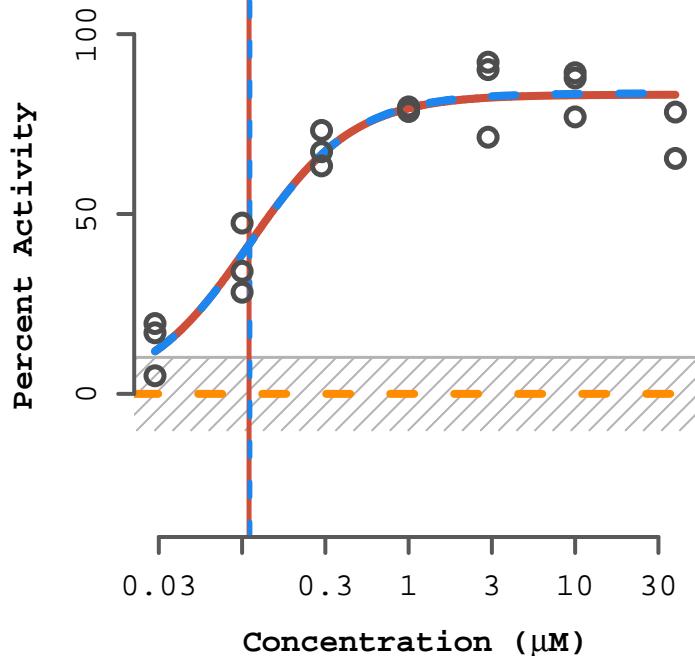
	tp	ga	gw	la	lw
val:	27.7	0.618	1.87	2.75	6.58
sd:	4.18	0.147	0.672	969	5570

	CNST	HILL	GNLS
AIC:	138.31	113.89	117.89
PROB:	0	0.88	0.12
RMSE:	14.44	7.67	7.67

MAX_MEAN: 22.8 MAX_MED: 26.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tributyltin methacrylate
 CHID: 35204 CASRN: 2155-70-6
 SPID(S): TP0001414D04
 M4ID: 30204475

HILL MODEL (in red):

tp	ga	gw
val: 83.2	-0.958	1.38
sd: 2.97	0.0675	0.244

GAIN-LOSS MODEL (in blue):

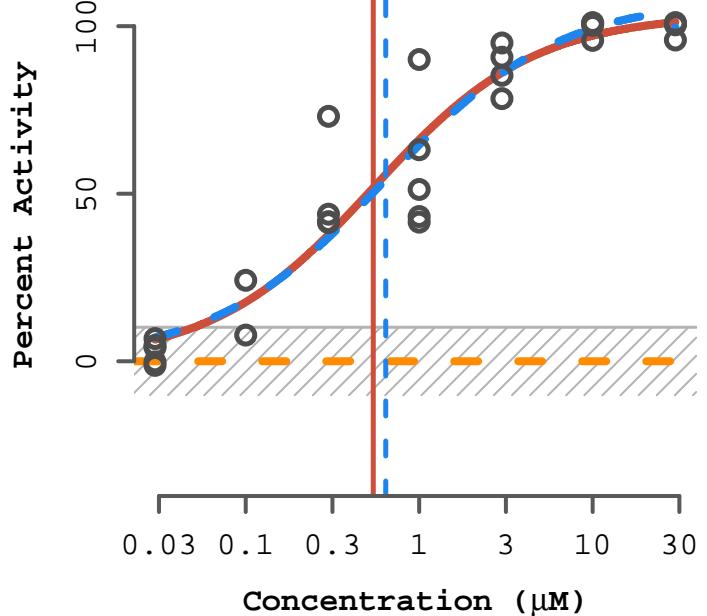
tp	ga	gw	la	lw
val: 83.6	-0.953	1.37	3.6	18
sd: NA	NA	NA	NA	NA

CNST	HILL	GNLS
AIC: 231.28	147.11	151.13
PROB: 0	0.88	0.12
RMSE: 67.6	7.46	7.49

MAX_MEAN: 84.7 MAX_MED: 90.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cyfluthrin

CHID: 35957 CASRN: 68359-37-5

SPID(S): EPAPLT0167F10

M4ID: 30204245

HILL MODEL (in red):

tp	ga	gw
val: 104	-0.265	0.934
sd: 3.87	0.0905	0.118

GAIN-LOSS MODEL (in blue):

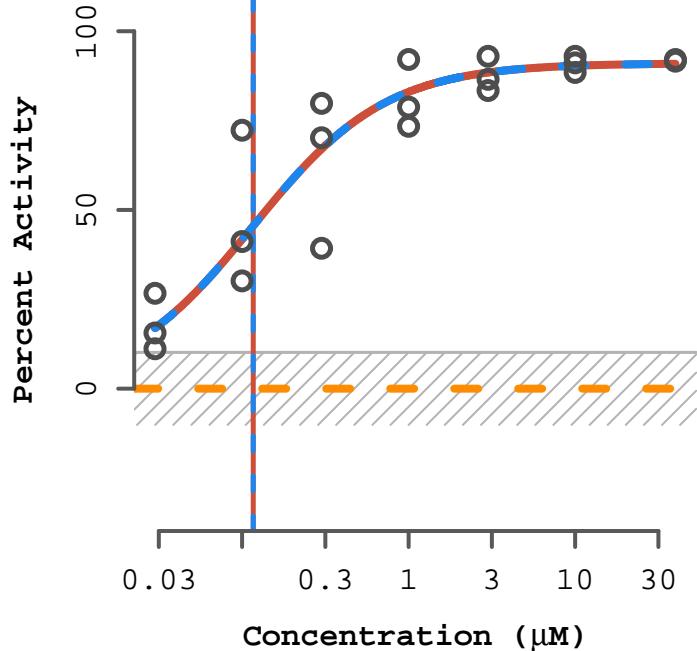
tp	ga	gw	la	lw
val: 108	-0.193	0.87	1.63	8.63
sd: NaN	NAN	NAN	NAN	NAN

CNST	HILL	GNLS
AIC: 324.46	218.59	221.88
PROB: 0	0.84	0.16
RMSE: 70.64	11.54	11.48

MAX_MEAN: 99.5 MAX_MED: 101 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cyfluthrin

CHID: 35957 CASRN: 68359-37-5

SPID(S): TP0001411E12

M4ID: 30204292

HILL MODEL (in red):

tp	ga	gw
val: 91	-0.933	1.09
sd: 2.83	0.0858	0.217

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 91	-0.933	1.09	3.46	4.45
sd: 2.83	0.0858	0.217	1950	4660

CNST

AIC: 234.39

PROB: 0

RMSE: 73.01

HILL

156.17

0.88

10.95

GNLS

160.17

0.12

10.95

MAX_MEAN: 91.9

MAX_MED: 91.9

BMAD: 3.38

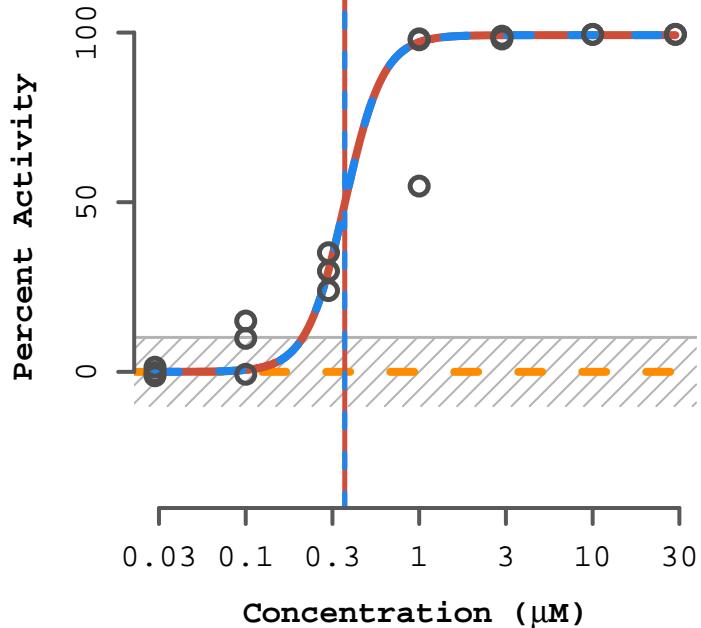
COFF: 10.1

HIT-CALL: 1

FITC: 41

ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: tau-Fluvalinate

CHID: 37555 CASRN: 102851-06-9

SPID(S): EPAPLT0167G07

M4ID: 30204271

HILL MODEL (in red):

	tp	ga	gw
val:	99.2	-0.429	3.9
sd:	0.977	0.024	0.894

GAIN-LOSS MODEL (in blue):

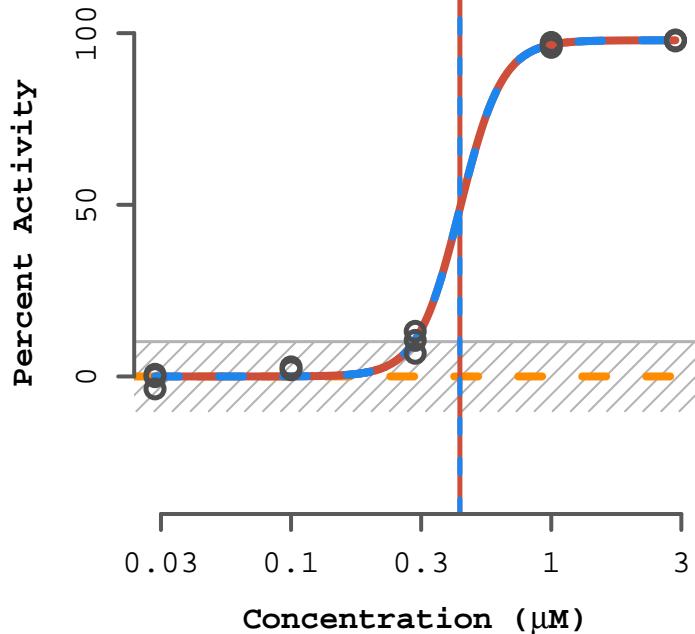
	tp	ga	gw	la	lw
val:	99.2	-0.429	3.9	3.23	15.8
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	220.09	129.67	133.67
PROB:	0	0.88	0.12
RMSE:	70.33	10.67	10.67

MAX_MEAN: 99.5 MAX_MED: 99.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Basic Blue 7

CHID: 38888 CASRN: 2390-60-5

SPID(S): EPAPLT0154G02

M4ID: 30204215

HILL MODEL (in red):

tp	ga	gw
val: 97.9	-0.351	5.33
sd: 0.696	0.0222	0.65

GAIN-LOSS MODEL (in blue):

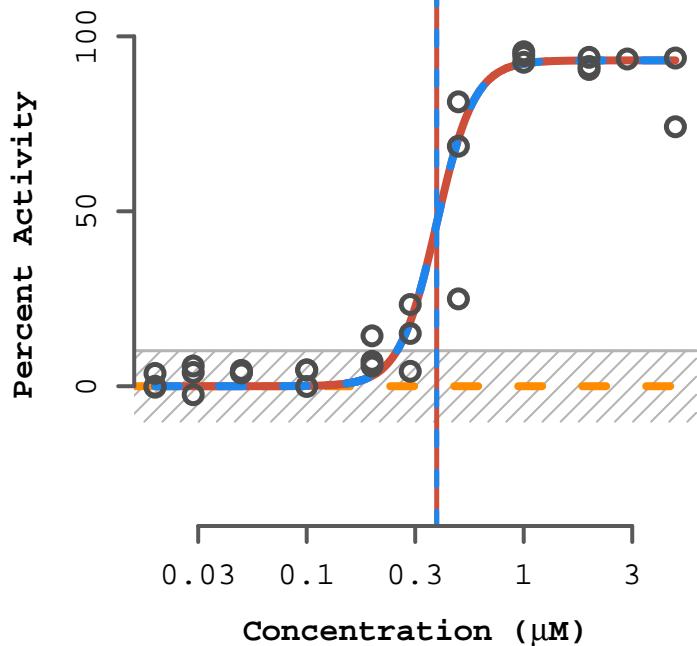
tp	ga	gw	la	lw
val: 97.9	-0.351	5.33	1.43	11.8
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 159.11	64.42	68.42
PROB: 0	0.88	0.12
RMSE: 63.88	1.84	1.84

MAX_MEAN: 97.9 MAX_MED: 98 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Niclosamide

CHID: 40362 CASRN: 50-65-7

SPID(S): TP0001413D04

M4ID: 30204627

HILL MODEL (in red):

tp	ga	gw
val: 93.1	-0.401	4.99
sd: 1.66	0.0175	0.832

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 93.1	-0.401	4.99	1.75	10.8
sd: NaN	NaN	NaN	NaN	NaN

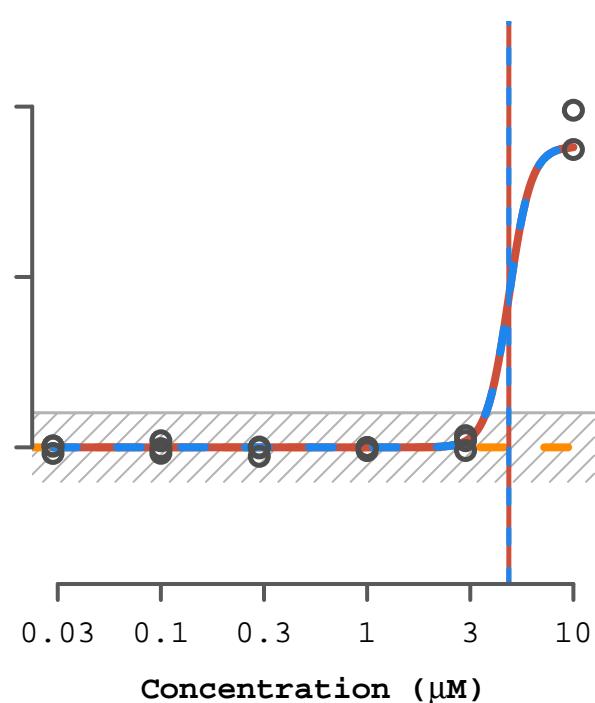
CNST	HILL	GNLS
AIC: 327.26	211.51	215.51
PROB: 0	0.88	0.12
RMSE: 54.24	10.24	10.24

MAX_MEAN: 94.2 MAX_MED: 94.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Octyl gallate

CHID: 40713 CASRN: 1034-01-1

SPID(S): TP0001413F06

M4ID: 30204534

HILL MODEL (in red):

tp	ga	gw
val: 88.4	0.687	8
sd: 7.67	0.607	23

GAIN-LOSS MODEL (in blue):

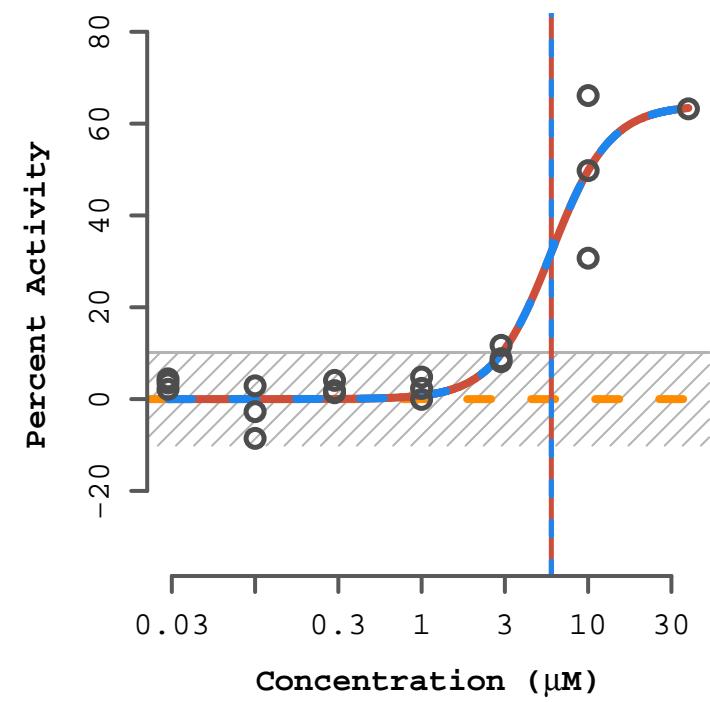
tp	ga	gw	la	lw
val: 88.4	0.687	8	2.03	17
sd: NA	NA	NA	NA	NA

CNST	HILL	GNLS
AIC: 130.98	81.08	85.08
PROB: 0	0.88	0.12
RMSE: 32.05	2.94	2.94

MAX_MEAN: 93.2 MAX_MED: 93.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 6



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Mepanipyrim

CHID: 42121 CASRN: 110235-47-7

SPID(S): TP0001411D05

M4ID: 30204621

HILL MODEL (in red):

tp	ga	gw
64	0.779	2.46
3.63	0.0511	0.404

GAIN-LOSS MODEL (in blue):

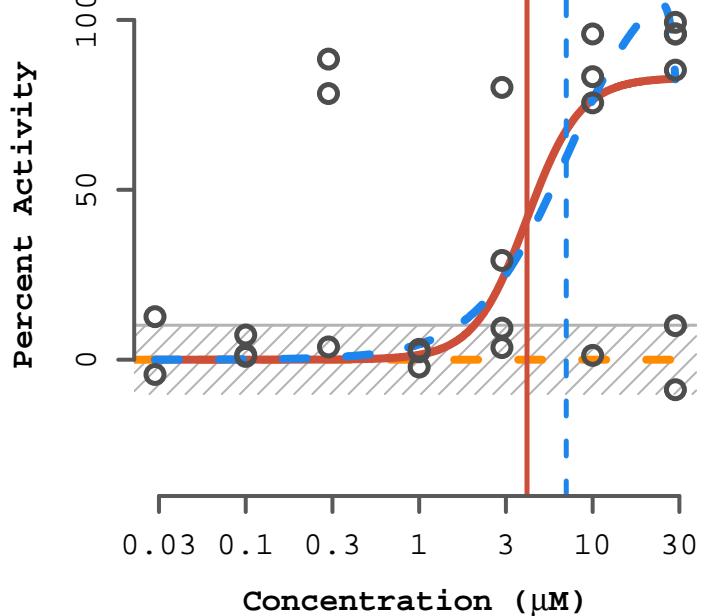
tp	ga	gw	la	lw
64	0.779	2.46	3.17	8.09
NA	NA	NA	NA	NA

CNST	HILL	GNLS
AIC: 171.71	126.56	130.56
PROB: 0	0.88	0.12
RMSE: 25.39	6.51	6.51

MAX_MEAN: 63.2 MAX_MED: 63.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Butyloctan-1-ol
 CHID: 44818 CASRN: 3913-02-8
 SPID(S): EPAPLT0154C06
 M4ID: 30204178

HILL MODEL (in red):

tp	ga	gw
val: 83	0.622	2.82
sd: 14.4	0.186	2.68

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 119	0.848	1.65	1.5	17.7
sd: NaN	NaN	NaN	NaN	NaN

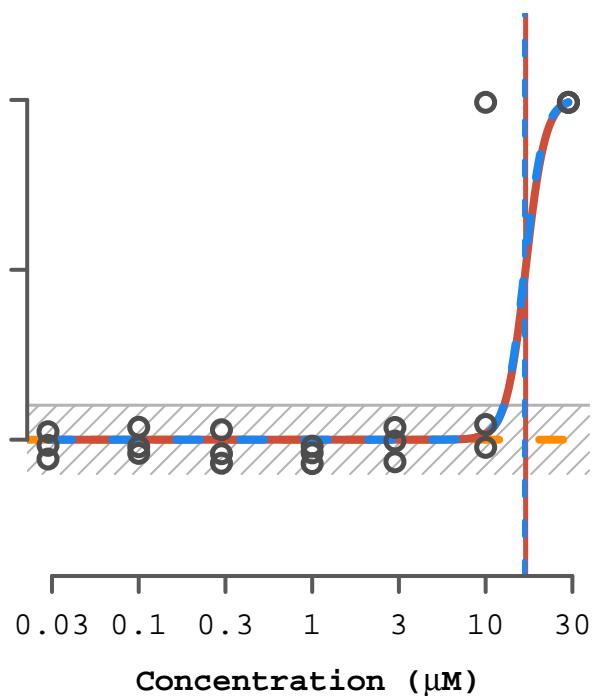
CNST	HILL	GNLS
AIC: 275.66	258.03	261.99
PROB: 0	0.88	0.12
RMSE: 56.27	39.21	39.09

MAX_MEAN: 64 MAX_MED: 90.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,6-Di-tert-butyl-4-[(dimethylamino)methyl]phenol
CHID: 44997 CASRN: 88-27-7
SPID(S): EPAPLT0154C07
M4ID: 30204179

HILL MODEL (in red):

	tp	ga	gw
val:	101	1.23	7.69
sd:	74.4	1.63	52.8

GAIN-LOSS MODEL (in blue):

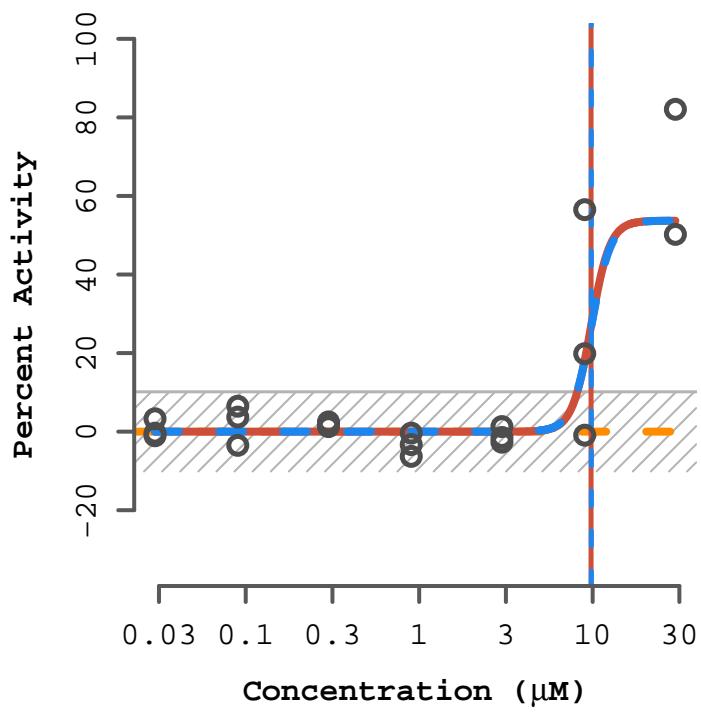
	tp	ga	gw	la	lw
val:	100	1.22	7.91	2.28	18
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	208	151.48	155.48
PROB:	0	0.88	0.12
RMSE:	43.51	21.63	21.63

MAX_MEAN: 99.3 MAX_MED: 99.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SB236057A

CHID: 47320 CASRN: 180084-01-9

SPID(S): TP0001412C04

M4ID: 30204509

HILL MODEL (in red):

	tp	ga	gw
val:	53.7	0.991	8
sd:	6.09	0.1	21.6

GAIN-LOSS MODEL (in blue):

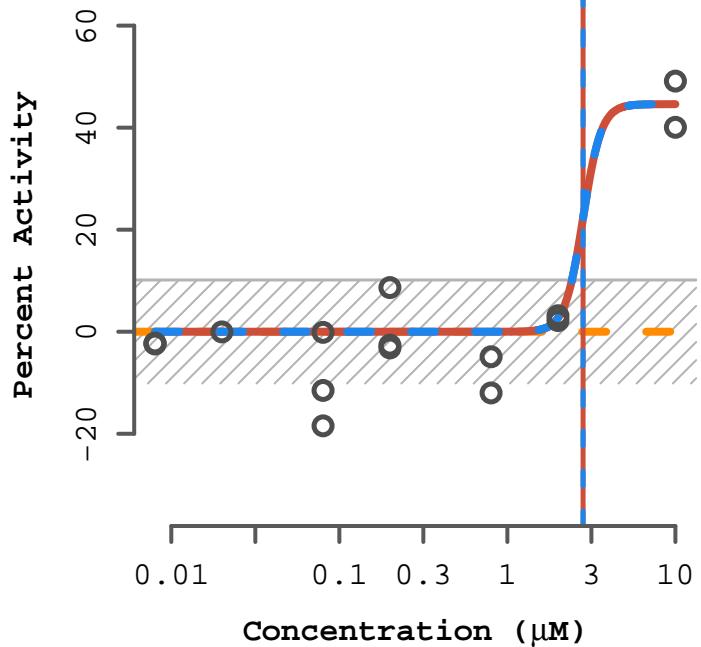
	tp	ga	gw	la	lw
val:	53.7	0.992	7.68	3.48	11.3
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	173.39	151.59	155.59
PROB:	0	0.88	0.12
RMSE:	25.49	11.85	11.85

MAX_MEAN: 66.1 MAX_MED: 66.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SSR69071

CHID: 47368 CASRN: 344930-95-6

SPID(S): TP0001414E11

M4ID: 30204523

HILL MODEL (in red):

tp	ga	gw
val: 44.6	0.451	8
sd: 4.28	0.724	38.4

GAIN-LOSS MODEL (in blue):

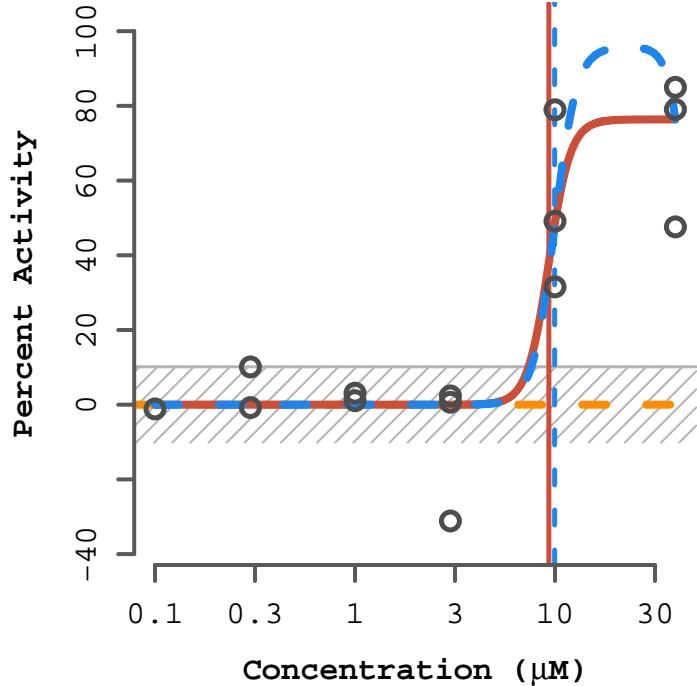
tp	ga	gw	la	lw
val: 44.6	0.451	8	1.76	6.55
sd: 13.1	0.72	38.2	3440	29800

CNST	HILL	GNLS
AIC: 132.55	114.31	118.31
PROB: 0	0.88	0.12
RMSE: 17.29	7.02	7.02

MAX_MEAN: 44.6 MAX_MED: 44.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 17; 6



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SAR102779

CHID: 47387 CASRN: NOCAS_47387

SPID(S): TP0001411C05

M4ID: 30204642

HILL MODEL (in red):

tp	ga	gw
val: 76.4	0.969	8
sd: 8.1	0.12	30.4

GAIN-LOSS MODEL (in blue):

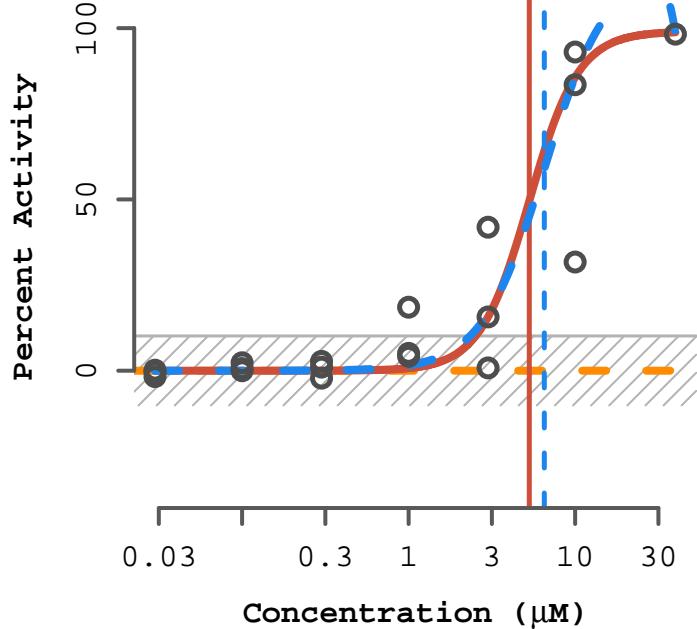
tp	ga	gw	la	lw
val: 95.8	0.998	7.94	1.66	10.7
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 147.6	122.68	126.68
PROB: 0	0.88	0.12
RMSE: 43.49	15.14	15.14

MAX_MEAN: 70.6 MAX_MED: 79.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Picoxystrobin

CHID: 47542 CASRN: 117428-22-5

SPID(S): TP0001414B01

M4ID: 30204616

HILL MODEL (in red):

tp	ga	gw
val: 99	0.725	2.93
sd: 5.43	0.0544	0.659

GAIN-LOSS MODEL (in blue):

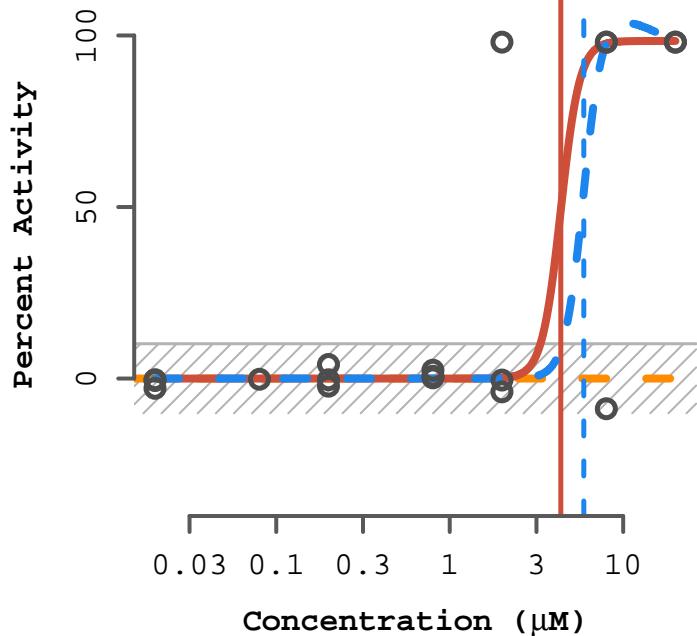
tp	ga	gw	la	lw
val: 118	0.815	2.3	1.68	9.79
sd: 68.6	0.293	1.43	0.0314	21.4

CNST	HILL	GNLS
AIC: 188.58	151.38	154.8
PROB: 0	0.85	0.15
RMSE: 38.86	14.95	14.86

MAX_MEAN: 98.3 MAX_MED: 98.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Flubendiamide

CHID: 47672 CASRN: 272451-65-7

SPID(S): EPAPLT0167F09

M4ID: 30204256

HILL MODEL (in red):

tp	ga	gw
val: 98.4	0.64	8
sd: 1.32	0.0918	2.58

GAIN-LOSS MODEL (in blue):

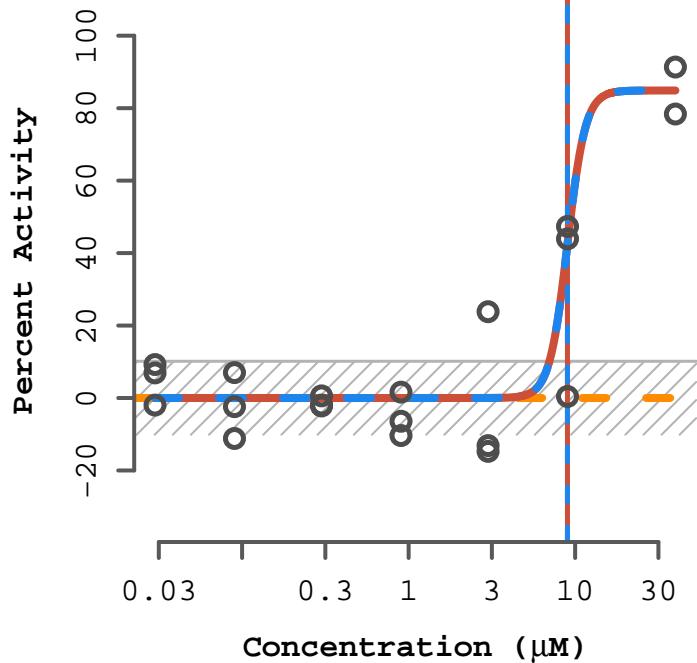
tp	ga	gw	la	lw
val: 118	0.773	8	2.24	0.749
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 192.49	145.51	149.2
PROB: 0	0.86	0.14
RMSE: 51.76	34.12	34.21

MAX_MEAN: 98.1 MAX_MED: 98.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fabezetron hydrochloride
 CHID: 48168 CASRN: 129299-90-7
 SPID(S): TP0001412D09
 M4ID: 30204619

HILL MODEL (in red):

tp	ga	gw
val: 84.9	0.954	8
sd: 6.61	0.0175	20.5

GAIN-LOSS MODEL (in blue):

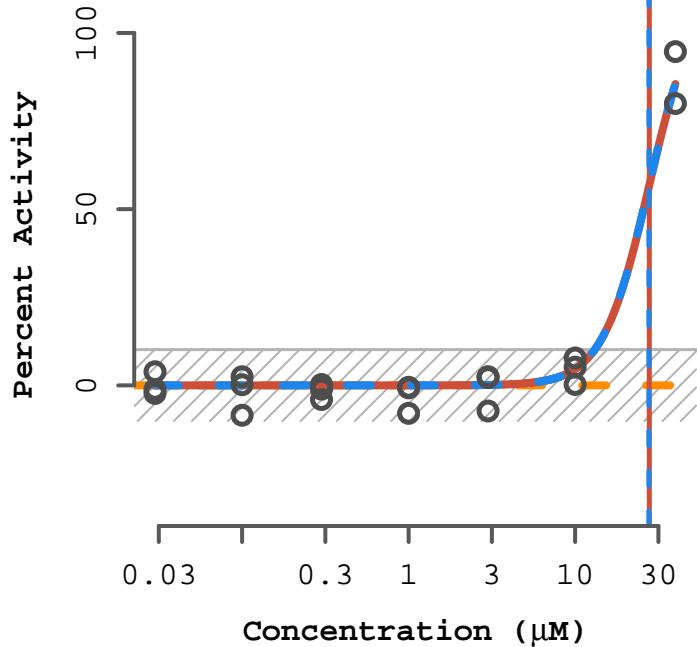
tp	ga	gw	la	lw
val: 84.9	0.954	8	3.48	2.63
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 190.16	162.26	166.26
PROB: 0	0.88	0.12
RMSE: 31.7	12.88	12.88

MAX_MEAN: 84.9 MAX_MED: 84.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 41 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-Amino-5-azotoluene
CHID: 20069 CASRN: 97-56-3
SPID(S): TP0001413E04
M4ID: 30204570

HILL MODEL (in red):

tp	ga	gw
val: 114	1.45	3.08
sd: 125	0.51	2.41

GAIN-LOSS MODEL (in blue):

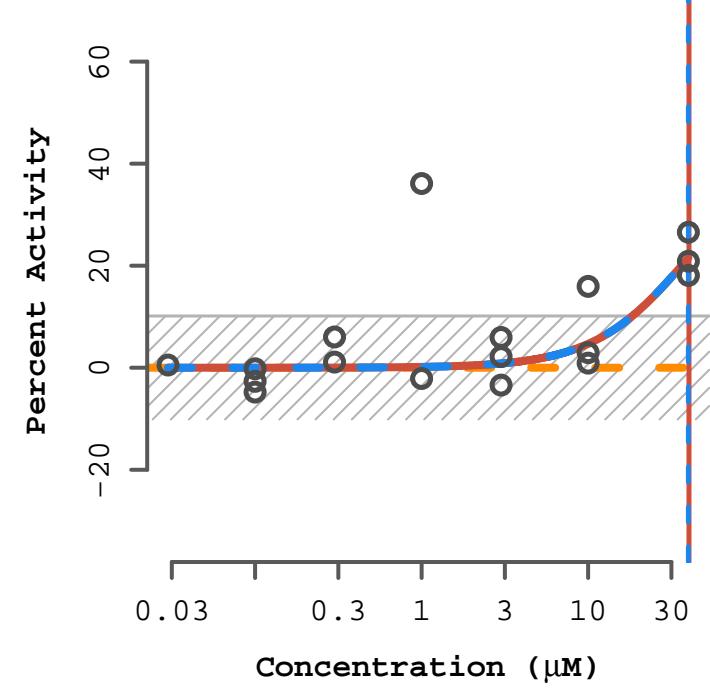
tp	ga	gw	la	lw
val: 113	1.44	3.09	1.96	17.8
sd: 120	0.493	2.37	533	25300

CNST	HILL	GNLS
AIC: 162.44	124.88	128.88
PROB: 0	0.88	0.12
RMSE: 28.02	4.41	4.41

MAX_MEAN: 87.4 MAX_MED: 87.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Atrazine

CHID: 20112 CASRN: 1912-24-9

SPID(S): TP0001411B12

M4ID: 30204374

HILL MODEL (in red):

	tp	ga	gw
val:	43.3	1.61	1.51
sd:	95.9	1.26	1.52

GAIN-LOSS MODEL (in blue):

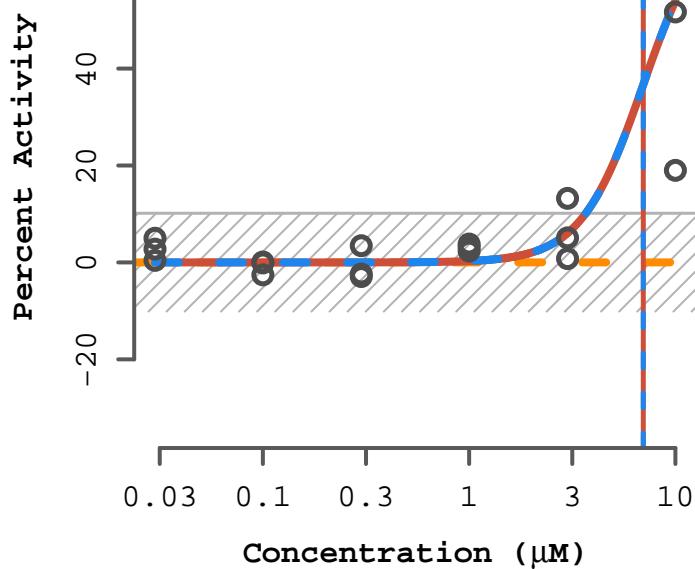
	tp	ga	gw	la	lw
val:	43.1	1.6	1.52	1.98	17.8
sd:	123	1.62	1.87	390	18300

	CNST	HILL	GNLS
AIC:	136.54	120.78	124.78
PROB:	0	0.88	0.12
RMSE:	13.64	9.68	9.68

MAX_MEAN: 21.8 MAX_MED: 20.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Benzyl butyl phthalate
CHID: 20205 CASRN: 85-68-7
SPID(S): TP0001413E05
M4ID: 30204638

HILL MODEL (in red):

tp	ga	gw
val: 73.4	0.844	2.85
sd: 73	0.442	2.47

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 73.4	0.844	2.85	1.78	17.1
sd: NA	NA	NA	NA	NA

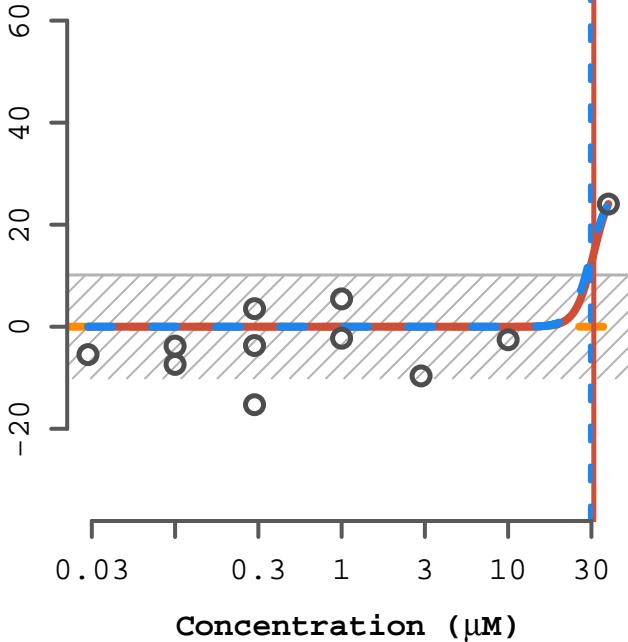
CNST	HILL	GNLS
AIC: 147.18	122.78	126.78
PROB: 0	0.88	0.12
RMSE: 19.82	8.97	8.97

MAX_MEAN: 43.9 MAX_MED: 51.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 6

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Butylparaben

CHID: 20209 CASRN: 94-26-8

SPID(S): TP0001411A10

M4ID: 30204387

HILL MODEL (in red):

	tp	ga	gw
val:	28.9	1.51	7.97
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	27.3	1.49	8	3.6	3.72
sd:	105	1.62	44.7	17300	32200

CNST

AIC: 83.01

PROB: 0.28

RMSE: 9.87

HILL

81.41

0.63

6.69

GNLS

85.41

0.09

6.69

MAX_MEAN: 24.1

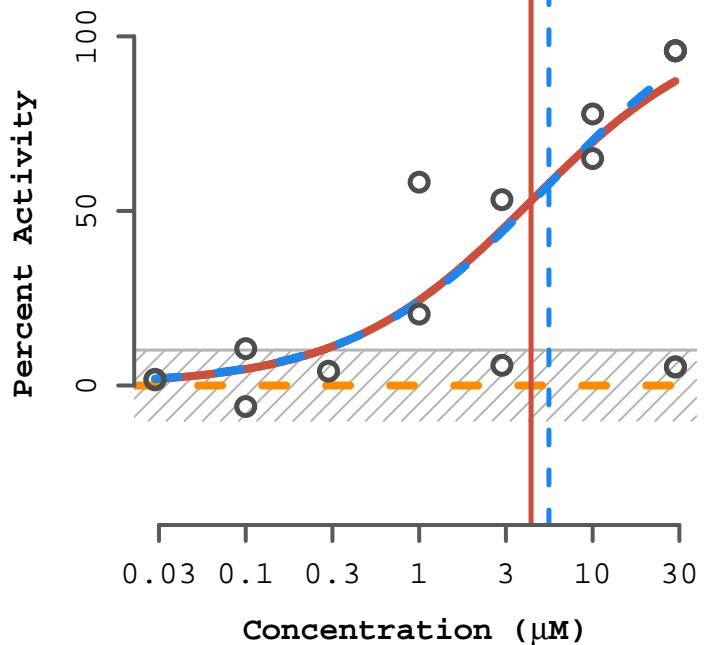
MAX_MED: 24.1

BMAD: 3.38

COFF: 10.1

HIT-CALL: 1 FITC: 42 ACTP: 0.72

FLAGS: 17; 16; 11; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Butylated hydroxyanisole
CHID: 20215 CASRN: 25013-16-5
SPID(S): EPAPLT0154B05
M4ID: 30204167

HILL MODEL (in red):

tp	ga	gw
val: 106	0.645	0.809
sd: 41.3	0.582	0.427

GAIN-LOSS MODEL (in blue):

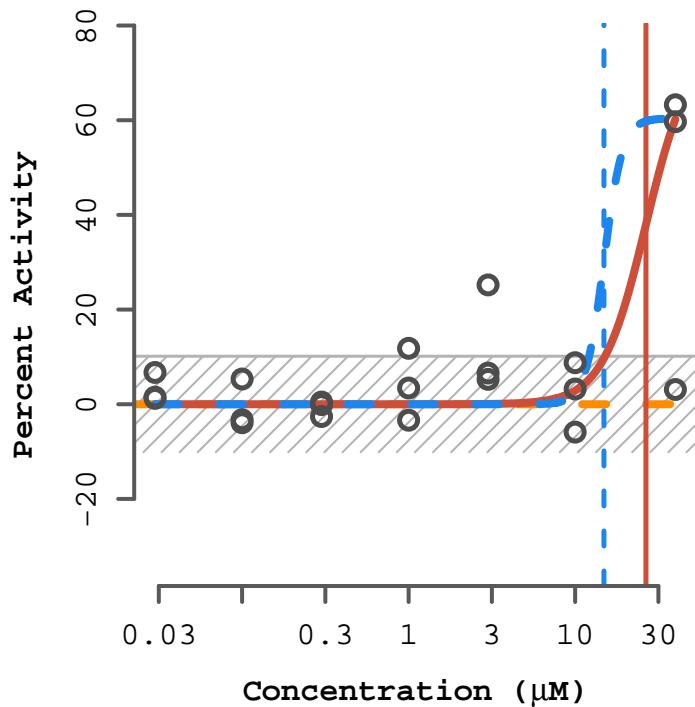
tp	ga	gw	la	lw
val: 115	0.748	0.773	1.56	17.7
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 175.07	159.33	163.32
PROB: 0	0.88	0.12
RMSE: 52.89	32.26	32.16

MAX_MEAN: 71.4 MAX_MED: 95.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: tert-Butylhydroquinone
CHID: 20220 CASRN: 1948-33-0
SPID(S): TP0001413A10
M4ID: 30204405

HILL MODEL (in red):

tp	ga	gw
val: 75.9	1.43	3.3
sd: 185	1.16	7.53

GAIN-LOSS MODEL (in blue):

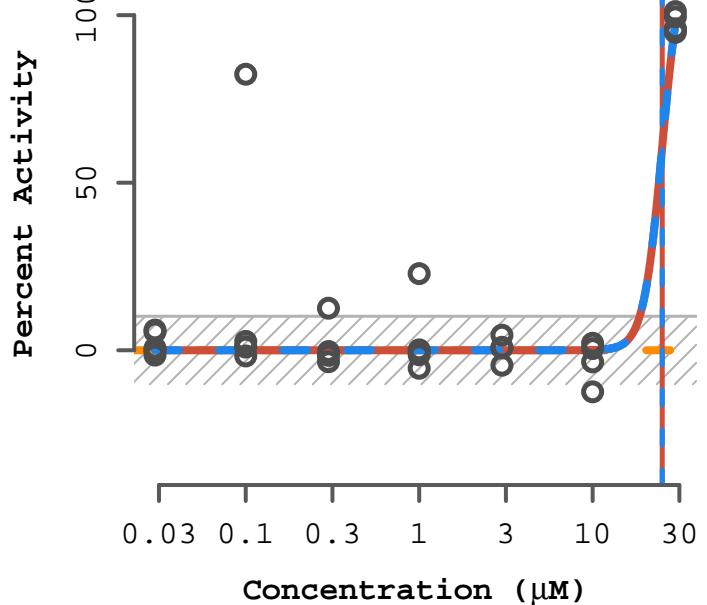
tp	ga	gw	la	lw
val: 60.4	1.17	7.84	2.06	5.69
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 172.62	162.03	166.08
PROB: 0	0.88	0.12
RMSE: 20.33	14.41	14.42

MAX_MEAN: 42 MAX_MED: 59.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Chlorpheniramine maleate
CHID: 20321 CASRN: 113-92-8
SPID(S): EPAPLT0167C07
M4ID: 30204244

HILL MODEL (in red):

tp	ga	gw
val: 121	1.4	8
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

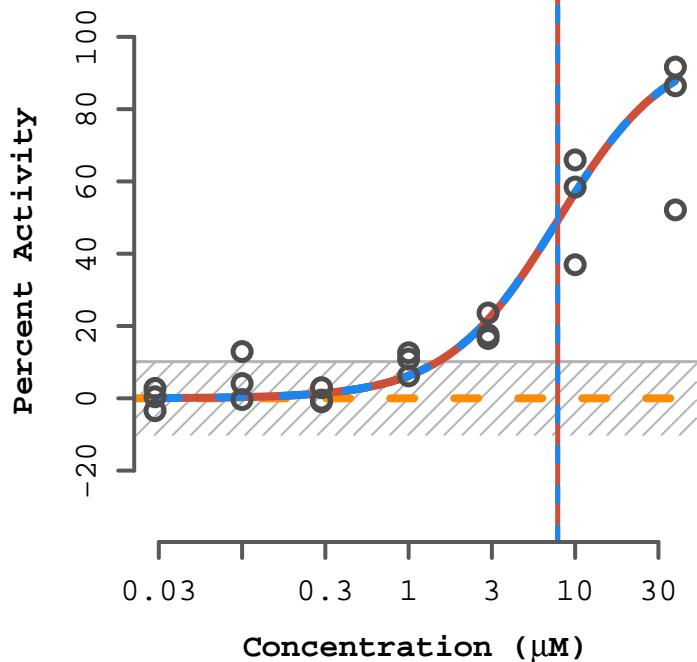
tp	ga	gw	la	lw
val: 121	1.4	8	3.48	18
sd: NA	NA	NA	NA	NA

CNST	HILL	GNLS
AIC: 322.07	225	229
PROB: 0	0.88	0.12
RMSE: 40.94	15.41	15.41

MAX_MEAN: 97.4 MAX_MED: 95.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Coumaphos

CHID: 20347 CASRN: 56-72-4

SPID(S): TP0001414D12

M4ID: 30204384

HILL MODEL (in red):

	tp	ga	gw
val:	98.4	0.895	1.31
sd:	9.59	0.11	0.269

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	98.4	0.895	1.31	2.86	10.3
sd:	NaN	NaN	NaN	NaN	NaN

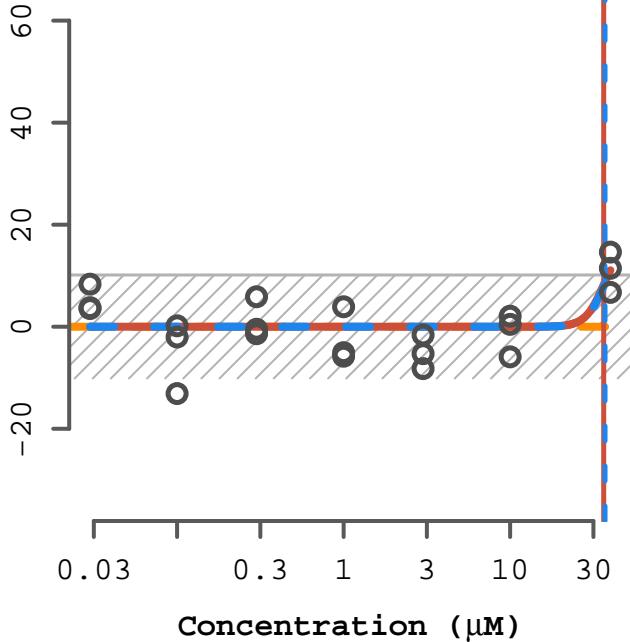
	CNST	HILL	GNLS
AIC:	211.02	153.6	157.6
PROB:	0	0.88	0.12
RMSE:	37.41	9.98	9.98

MAX_MEAN: 76.7 MAX_MED: 86.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: p,p'-DDE

CHID: 20374 CASRN: 72-55-9

SPID(S): TP0001411H04

M4ID: 30204428

HILL MODEL (in red):

	tp	ga	gw
val:	16.4	1.56	7.78
sd:	388	3.01	210

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	17.2	1.57	7.98	2.67	3.34
sd:	NaN	NaN	NaN	NaN	NaN

CNST

AIC: 141.98

PROB: 0.08

RMSE: 6.59

HILL

137.22

0.81

5.13

GNLS

141.22

0.11

5.13

MAX_MEAN: 10.9

MAX_MED: 11.5

BMAD: 3.38

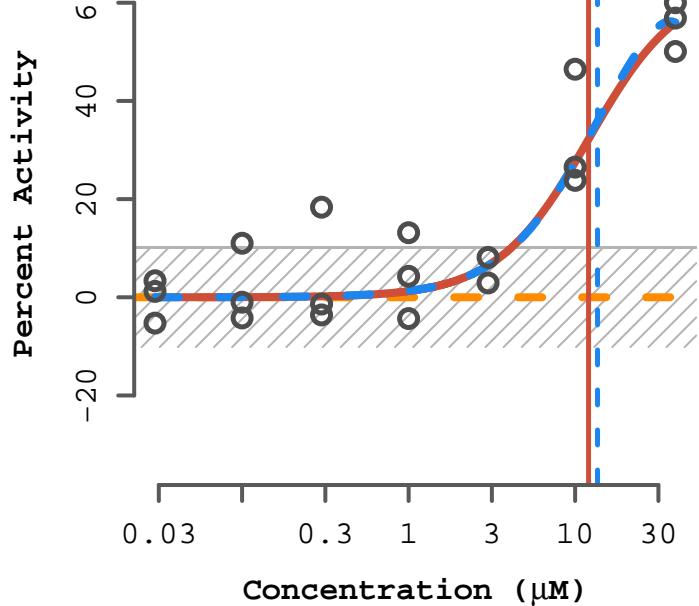
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 0.92

FLAGS: 17; 11; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dichlorodiphenyltrichloroethane

CHID: 20375 CASRN: 50-29-3

SPID(S): TP0001412C08

M4ID: 30204398

HILL MODEL (in red):

	tp	ga	gw
val:	64.4	1.08	1.58
sd:	13.2	0.177	0.691

GAIN-LOSS MODEL (in blue):

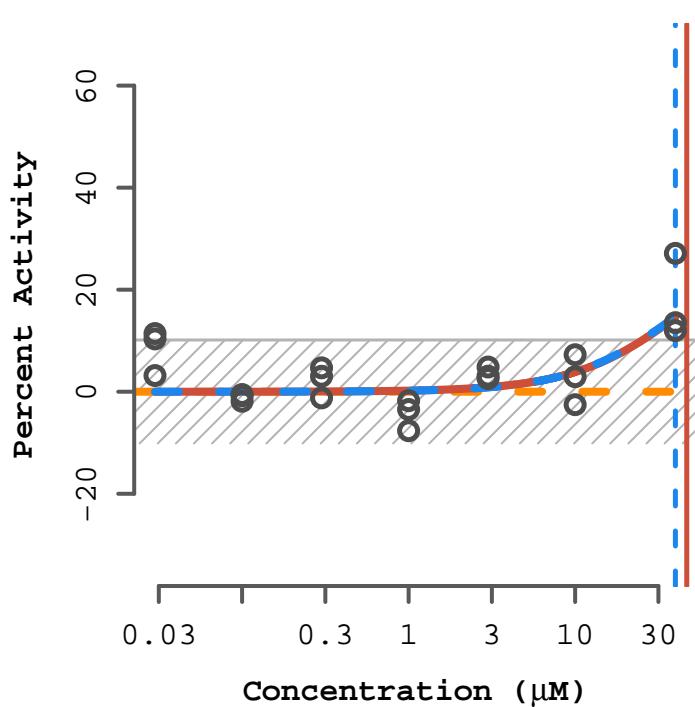
	tp	ga	gw	la	lw
val:	71.9	1.13	1.53	1.8	5.52
sd:	164	1.14	1.18	2.74	101

	CNST	HILL	GNLS
AIC:	186.96	143.77	147.75
PROB:	0	0.88	0.12
RMSE:	26.07	7.58	7.57

MAX_MEAN: 55.7 MAX_MED: 56.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diallyl phthalate
CHID: 20392 CASRN: 131-17-9
SPID(S): TP0001411D09
M4ID: 30204440

HILL MODEL (in red):

	tp	ga	gw
val:	32.5	1.67	1.3
sd:	57.5	1.11	0.998

GAIN-LOSS MODEL (in blue):

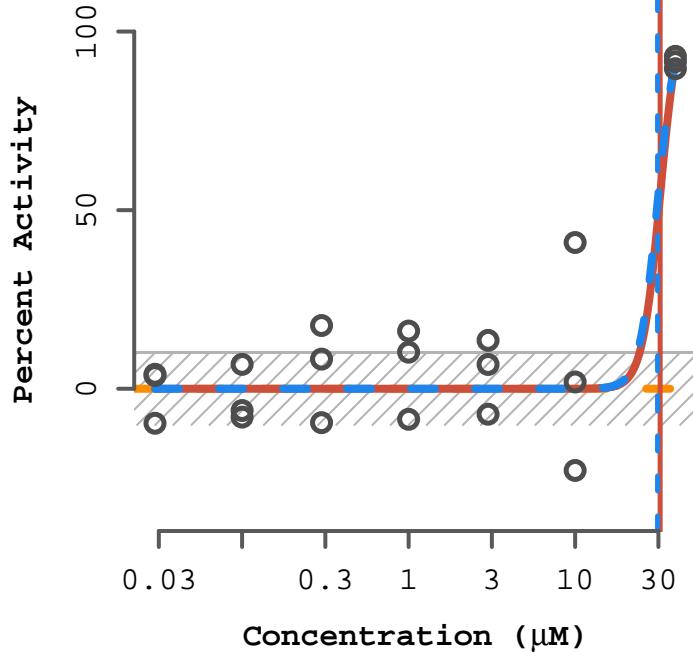
	tp	ga	gw	la	lw
val:	29.3	1.6	1.36	2.94	5.18
sd:	53.4	1.15	1.16	1400	5430

	CNST	HILL	GNLS
AIC:	148.19	137.63	141.65
PROB:	0	0.88	0.12
RMSE:	8.53	5.36	5.36

MAX_MEAN: 17.5 MAX_MED: 13.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dicloran

CHID: 20426 CASRN: 99-30-9

SPID(S): TP0001412F12

M4ID: 30204448

HILL MODEL (in red):

	tp	ga	gw
val:	108	1.51	7.99
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

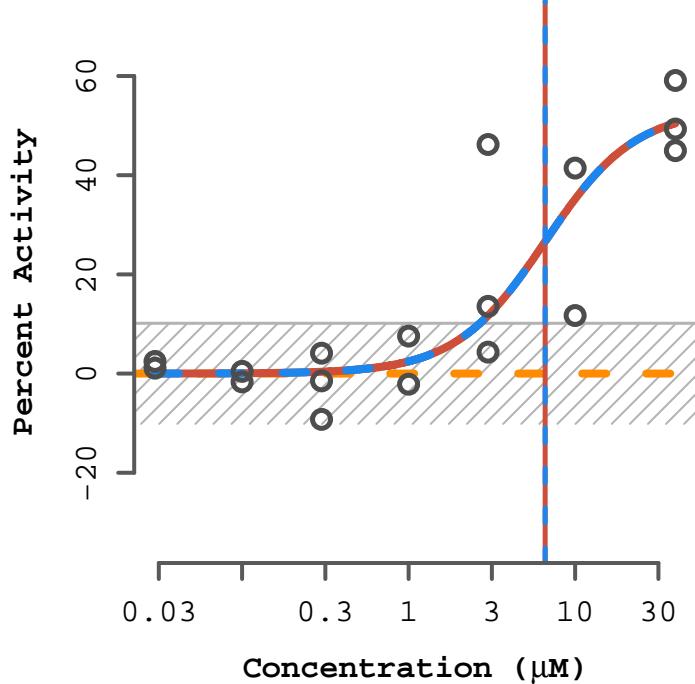
	tp	ga	gw	la	lw
val:	104	1.49	7.98	3.56	11.9
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	205.78	173.81	177.82
PROB:	0	0.88	0.12
RMSE:	37	13.22	13.22

MAX_MEAN: 91.4 MAX_MED: 91.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dicofol

CHID: 20450 CASRN: 115-32-2

SPID(S): TP0001414C12

M4ID: 30204363

HILL MODEL (in red):

tp	ga	gw
val: 53.3	0.82	1.61
sd: 8.35	0.226	0.817

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 53.3	0.82	1.61	2.2	11.7
sd: 8.36	0.226	0.817	818	16100

CNST

AIC: 161.09

PROB: 0

RMSE: 26.91

HILL

AIC: 132.16

PROB: 0.88

RMSE: 11.15

GNLS

AIC: 136.16

PROB: 0.12

RMSE: 11.15

MAX_MEAN: 51.1

MAX_MED: 49.3

BMAD: 3.38

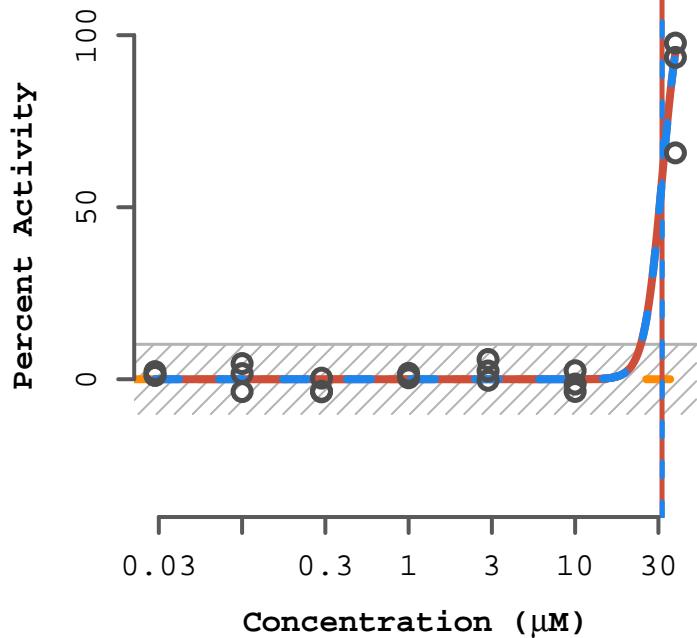
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 1

FLAGS: 17; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 7,12-Dimethylbenz(a)anthracene

CHID: 20510 CASRN: 57-97-6

SPID(S): TP0001414A01

M4ID: 30204553

HILL MODEL (in red):

	tp	ga	gw
val:	117	1.52	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

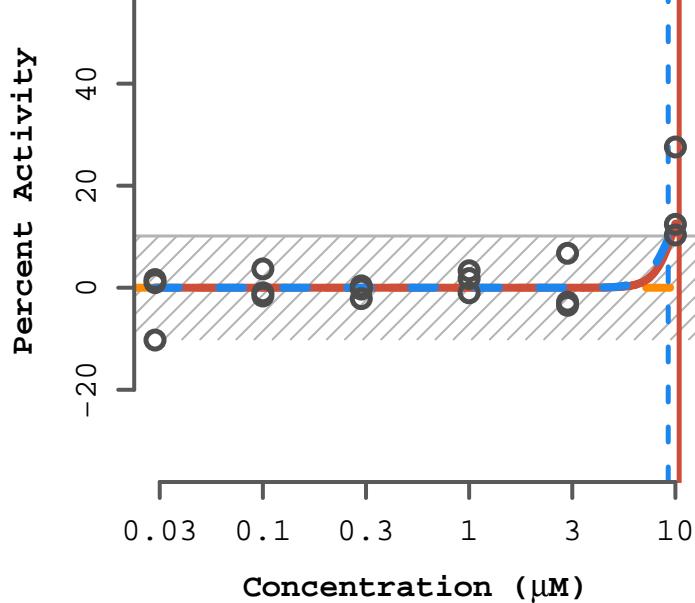
	tp	ga	gw	la	lw
val:	117	1.52	7.99	3.51	1.43
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	181.17	127.75	131.75
PROB:	0	0.88	0.12
RMSE:	32.93	6.86	6.86

MAX_MEAN: 85.7 MAX_MED: 93.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diphenhydramine hydrochloride
CHID: 20537 CASRN: 147-24-0
SPID(S): TP0001413E08
M4ID: 30204606

HILL MODEL (in red):

	tp	ga	gw
val:	30.2	1.02	8
sd:	55.4	0.212	53.1

GAIN-LOSS MODEL (in blue):

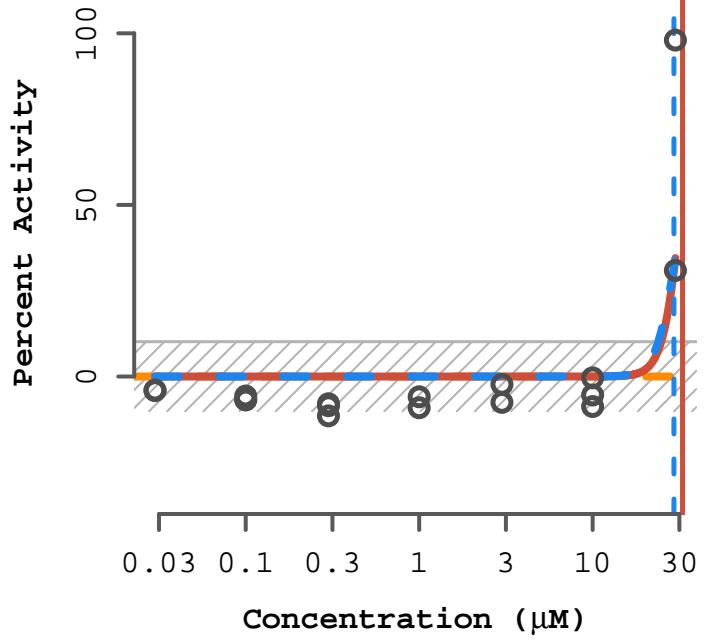
	tp	ga	gw	la	lw
val:	19.2	0.965	8	1.29	10.7
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	122.63	111.87	115.87
PROB:	0	0.88	0.12
RMSE:	8.28	4.95	4.95

MAX_MEAN: 16.8 MAX_MED: 12.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 6



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Endosulfan

CHID: 20560 CASRN: 115-29-7

SPID(S): EPAPLT0167G10

M4ID: 30204251

HILL MODEL (in red):

	tp	ga	gw
val:	108	1.52	8
sd:	142	0.0543	23.1

GAIN-LOSS MODEL (in blue):

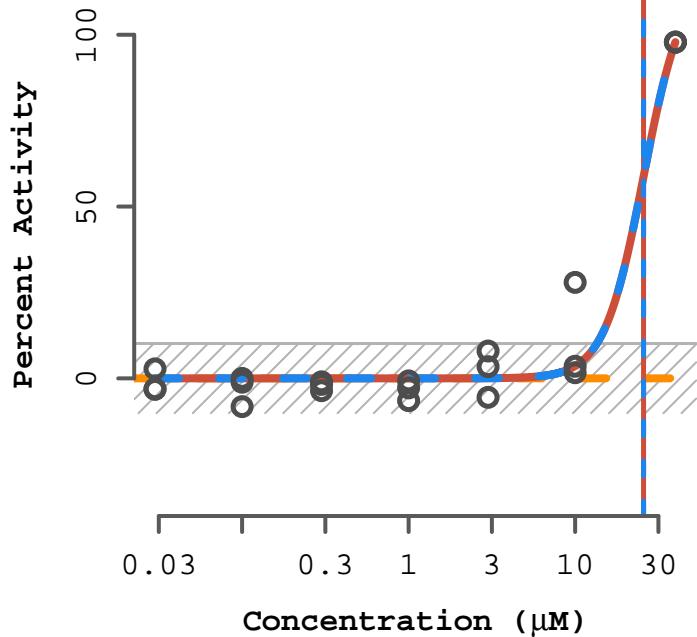
	tp	ga	gw	la	lw
val:	64.1	1.47	8	2.7	2.3
sd:	272	0.444	49.3	92.4	174

	CNST	HILL	GNLS
AIC:	143.92	138.58	142.58
PROB:	0.06	0.83	0.11
RMSE:	25.75	16.73	16.73

MAX_MEAN: 64.5 MAX_MED: 64.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.94

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fenthion

CHID: 20620 CASRN: 55-38-9

SPID(S): TP0001414A03

M4ID: 30204344

HILL MODEL (in red):

tp	ga	gw
val: 117	1.41	3.65
sd: 99.2	0.428	3.42

GAIN-LOSS MODEL (in blue):

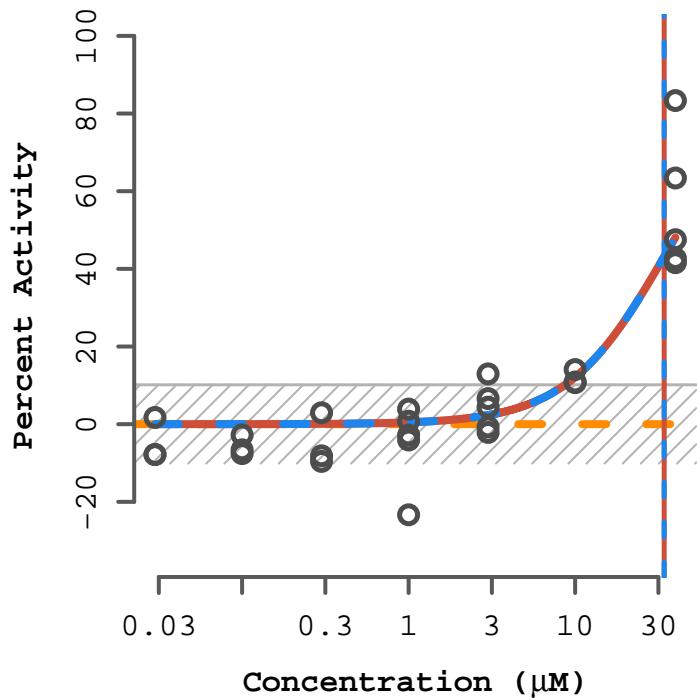
tp	ga	gw	la	lw
val: 117	1.41	3.66	3.6	3.08
sd: 113	0.489	3.9	1830	2830

CNST	HILL	GNLS
AIC: 196.93	135.05	139.05
PROB: 0	0.88	0.12
RMSE: 37.67	6.42	6.42

MAX_MEAN: 97.9 MAX_MED: 97.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Chlorpropham

CHID: 20764 CASRN: 101-21-3

SPID(S): TP0001411A01

M4ID: 30204376

HILL MODEL (in red):

	tp	ga	gw
val:	86.3	1.53	1.47
sd:	108	0.783	0.901

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	86.3	1.53	1.47	2.13	9.93
sd:	108	0.784	0.901	186	3520

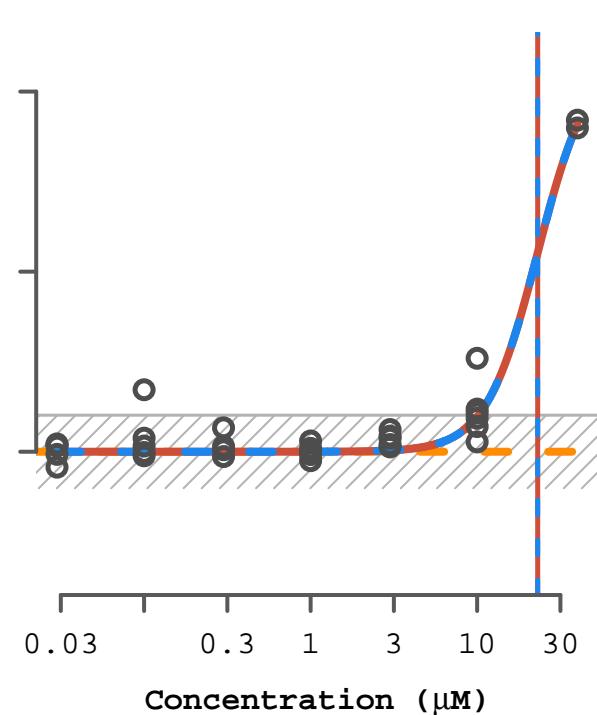
	CNST	HILL	GNLS
AIC:	245.82	201.85	205.85
PROB:	0	0.88	0.12
RMSE:	26.13	10.07	10.07

MAX_MEAN: 55.7 MAX_MED: 47.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Mercuric chloride
CHID: 20811 CASRN: 7487-94-7
SPID(S): TP0001413F02
M4ID: 30204549

HILL MODEL (in red):

	tp	ga	gw
val:	110	1.36	2.79
sd:	46.5	0.249	1.42

GAIN-LOSS MODEL (in blue):

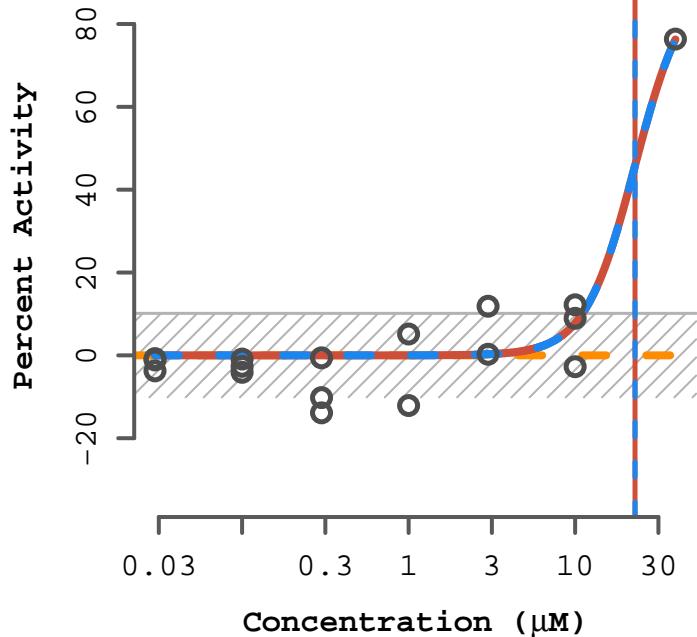
	tp	ga	gw	la	lw
val:	110	1.36	2.79	2.96	12
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	250.88	188.35	192.35
PROB:	0	0.88	0.12
RMSE:	23.8	5.01	5.01

MAX_MEAN: 91 MAX_MED: 91 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Phenolphthalein
CHID: 21125 CASRN: 77-09-8
SPID(S): TP0001413H05
M4ID: 30204543

HILL MODEL (in red):

tp	ga	gw
val: 91.6	1.36	2.87
sd: 102	0.665	4.03

GAIN-LOSS MODEL (in blue):

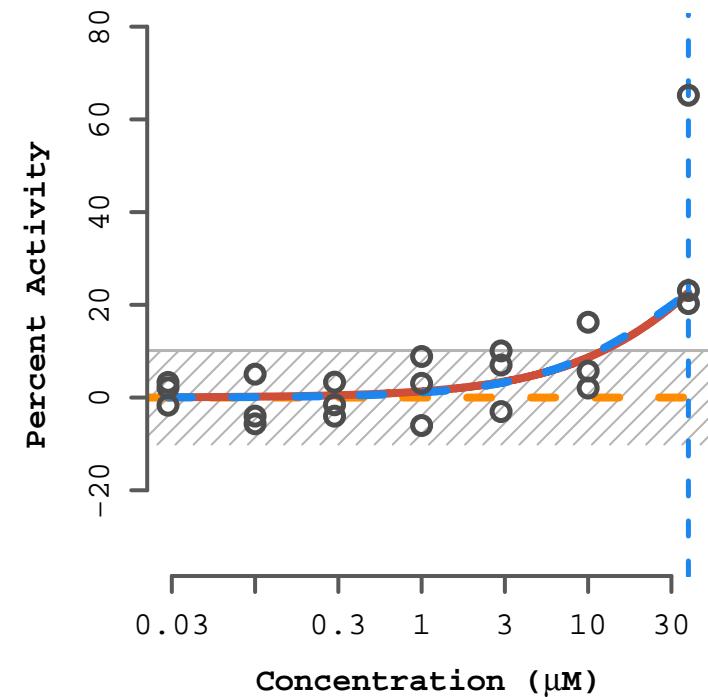
tp	ga	gw	la	lw
val: 91.6	1.36	2.87	3.16	5.58
sd: 102	0.666	4.04	9670	34600

CNST	HILL	GNLS
AIC: 135.9	121.49	125.49
PROB: 0	0.88	0.12
RMSE: 19.88	6.76	6.76

MAX_MEAN: 76.4 MAX_MED: 76.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Phenothiazine

CHID: 21126 CASRN: 92-84-2

SPID(S): TP0001411E08

M4ID: 30204409

HILL MODEL (in red):

tp	ga	gw
val: 78.2	2.06	0.856
sd: 180	1.83	0.465

GAIN-LOSS MODEL (in blue):

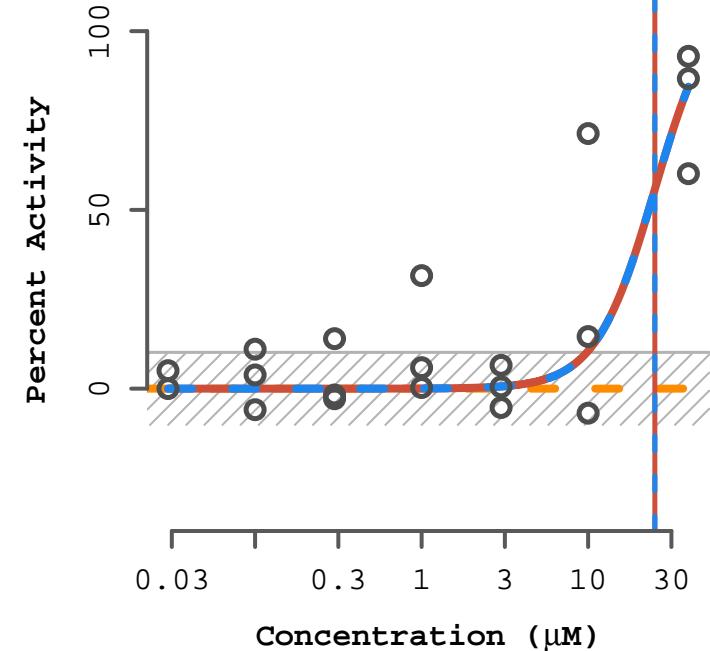
tp	ga	gw	la	lw
val: 45.2	1.6	1	3.48	3.72
sd: 56.2	1.06	0.607	1310	2600

CNST	HILL	GNLS
AIC: 166.71	150.78	154.93
PROB: 0	0.89	0.11
RMSE: 16.77	10.39	10.41

MAX_MEAN: 36.2 MAX_MED: 23.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ziram

CHID: 21464 CASRN: 137-30-4

SPID(S): TP0001412G02

M4ID: 30204556

HILL MODEL (in red):

	tp	ga	gw
val:	112	1.4	2.45
sd:	452	2.22	8.94

GAIN-LOSS MODEL (in blue):

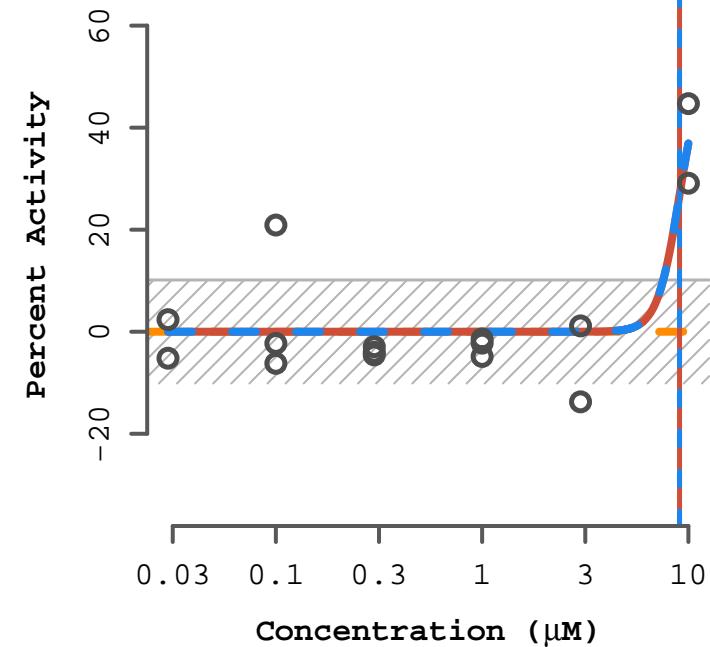
	tp	ga	gw	la	lw
val:	112	1.4	2.45	3.59	4.71
sd:	381	1.88	7.56	57700	136000

	CNST	HILL	GNLS
AIC:	196.84	171.26	175.26
PROB:	0	0.88	0.12
RMSE:	36.5	17.6	17.6

MAX_MEAN: 80 MAX_MED: 86.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-(2-Methylbutan-2-yl)phenol

CHID: 21771 CASRN: 80-46-6

SPID(S): TP0001414F01

M4ID: 30204412

HILL MODEL (in red):

tp	ga	gw
val: 53.6	0.957	8
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

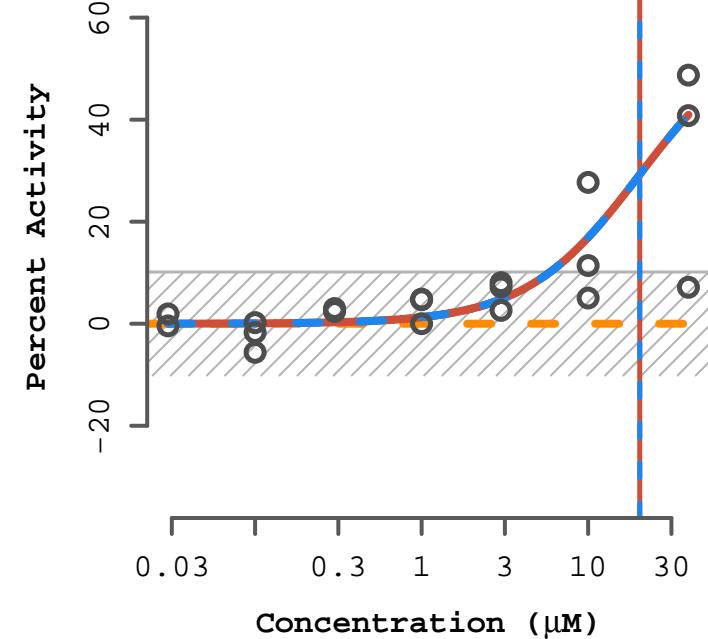
tp	ga	gw	la	lw
val: 53.6	0.957	8	3	9.84
sd: NA	NA	NA	NA	NA

CNST	HILL	GNLS
AIC: 122.47	110.76	114.76
PROB: 0	0.88	0.12
RMSE: 15.55	7.75	7.75

MAX_MEAN: 36.9 MAX_MED: 36.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 6



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1-Decanol

CHID: 21946 CASRN: 112-30-1

SPID(S): TP0001414F03

M4ID: 30204404

HILL MODEL (in red):

	tp	ga	gw
val:	58.5	1.31	1.26
sd:	18.2	0.273	0.401

GAIN-LOSS MODEL (in blue):

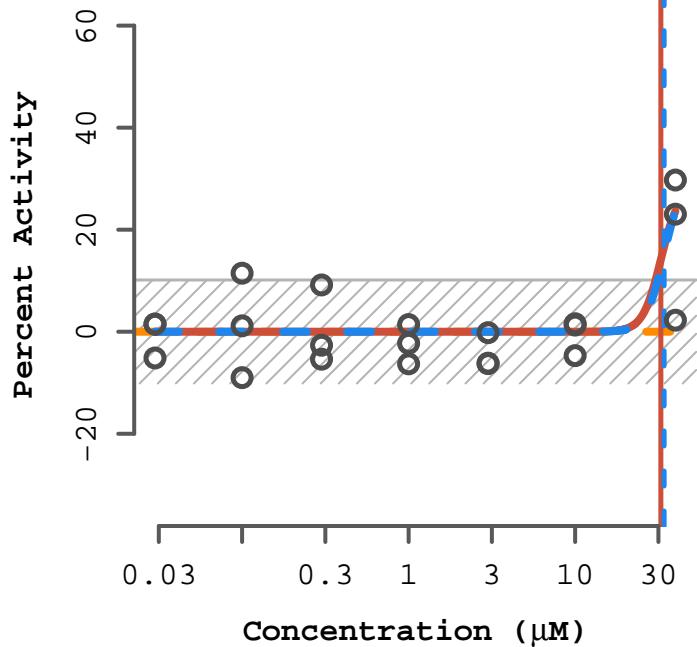
	tp	ga	gw	la	lw
val:	58.5	1.31	1.26	3.46	10.9
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	160.66	140.74	144.74
PROB:	0	0.88	0.12
RMSE:	16.19	8.93	8.93

MAX_MEAN: 32.2 MAX_MED: 40.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,3,6-Trimethylphenol
CHID: 22187 CASRN: 2416-94-6
SPID(S): TP0001412D08
M4ID: 30204394

HILL MODEL (in red):

	tp	ga	gw
val:	28.4	1.51	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

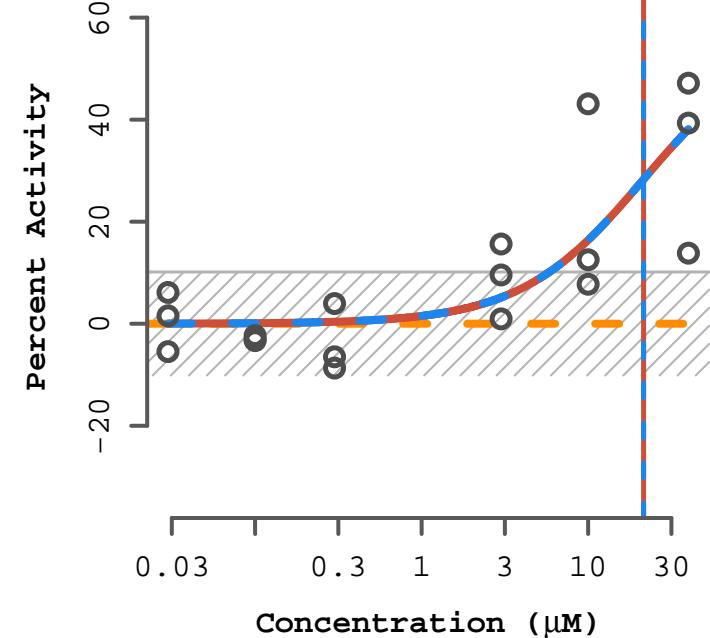
	tp	ga	gw	la	lw
val:	30.5	1.53	8	1.99	6.44
sd:	107	0.0693	86.7	115	2350

CNST	HILL	GNLS
AIC: 150.82	146.72	150.72
PROB: 0.1	0.79	0.11
RMSE: 9.6	6.93	6.93

MAX_MEAN: 18.4 MAX_MED: 23 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.9

FLAGS: 17; 11; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Resmethrin

CHID: 22253 CASRN: 10453-86-8

SPID(S): TP0001413E07

M4ID: 30204477

HILL MODEL (in red):

	tp	ga	gw
val:	56.6	1.33	1.17
sd:	33.8	0.523	0.567

GAIN-LOSS MODEL (in blue):

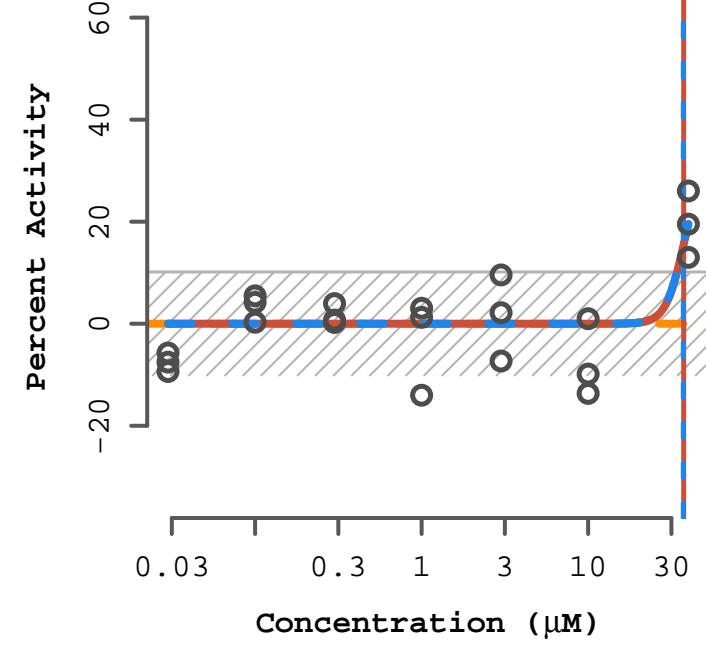
	tp	ga	gw	la	lw
val:	56.6	1.33	1.17	3.05	9.96
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	155.96	140.33	144.33
PROB:	0	0.88	0.12
RMSE:	19.15	10.17	10.17

MAX_MEAN: 33.4 MAX_MED: 39.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Permethrin

CHID: 22292 CASRN: 52645-53-1

SPID(S): TP0001414F11

M4ID: 30204345

HILL MODEL (in red):

	tp	ga	gw
val:	31.1	1.57	7.97
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

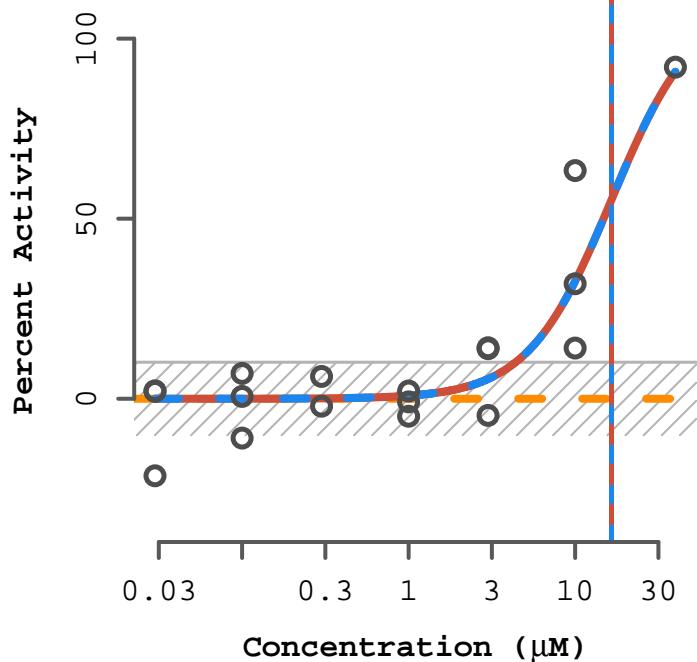
	tp	ga	gw	la	lw
val:	31.1	1.57	8	2.23	2.9
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	158.35	149.74	153.74
PROB:	0.01	0.87	0.12
RMSE:	10.02	6.79	6.79

MAX_MEAN: 19.5 MAX_MED: 19.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.99

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Octylphenol

CHID: 22312 CASRN: 1806-26-4

SPID(S): TP0001411C11

M4ID: 30204443

HILL MODEL (in red):

	tp	ga	gw
val:	111	1.22	1.73
sd:	29.2	0.203	0.74

GAIN-LOSS MODEL (in blue):

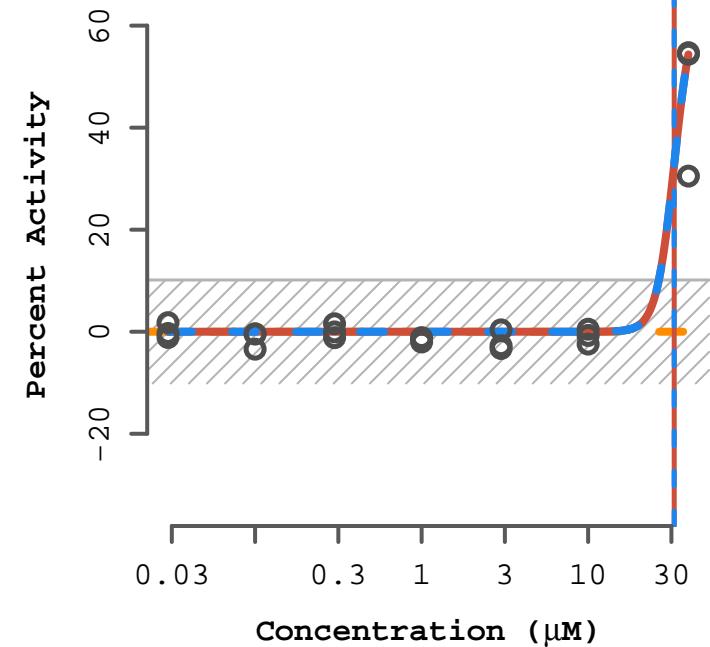
	tp	ga	gw	la	lw
val:	111	1.22	1.73	3.44	16.9
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	165.22	143.95	147.95
PROB:	0	0.88	0.12
RMSE:	28.71	11.19	11.19

MAX_MEAN: 92.1 MAX_MED: 92.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-(Butan-2-yl)phenol
CHID: 22332 CASRN: 99-71-8
SPID(S): TP0001413F10
M4ID: 30204411

HILL MODEL (in red):

tp	ga	gw
val: 65.7	1.52	8
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

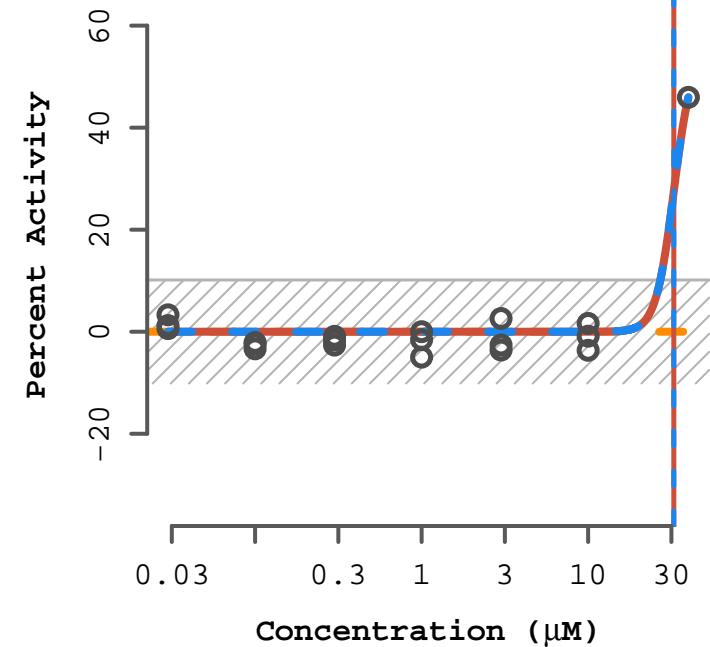
tp	ga	gw	la	lw
val: 65.7	1.52	8	3.45	3.01
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 156.74	109.01	113.01
PROB: 0	0.88	0.12
RMSE: 18.17	5.44	5.44

MAX_MEAN: 46.5 MAX_MED: 54.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 5alpha-Dihydrotestosterone
CHID: 22364 CASRN: 521-18-6
SPID(S): TP0001413E06
M4ID: 30204640

HILL MODEL (in red):

	tp	ga	gw
val:	55.1	1.51	7.98
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	55.1	1.51	8	3.56	13.6
sd:	NA	NA	NA	NA	NA

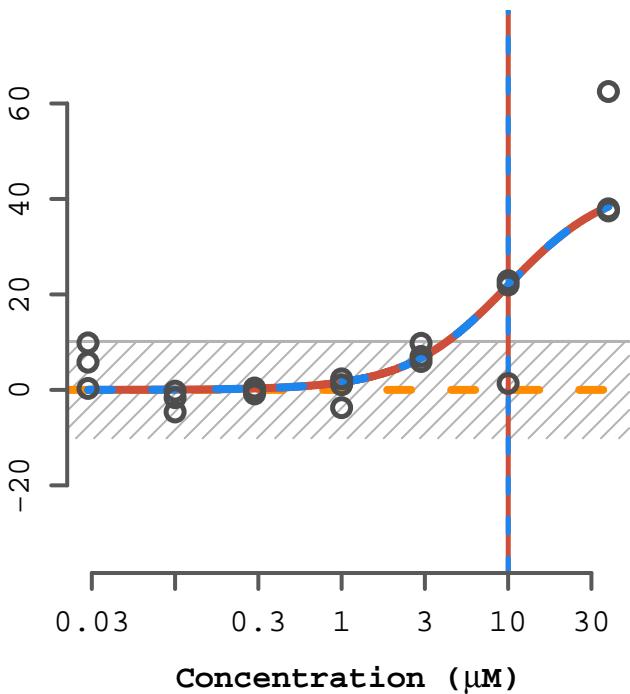
	CNST	HILL	GNLS
AIC:	115.06	98.58	102.58
PROB:	0	0.88	0.12
RMSE:	10.82	2.46	2.46

MAX_MEAN: 45.9 MAX_MED: 45.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Diisobutyl phthalate
CHID: 22522 CASRN: 84-69-5
SPID(S): TP0001412B11
M4ID: 30204607

HILL MODEL (in red):

	tp	ga	gw
val:	43.8	1	1.42
sd:	5.33	0.117	0.301

GAIN-LOSS MODEL (in blue):

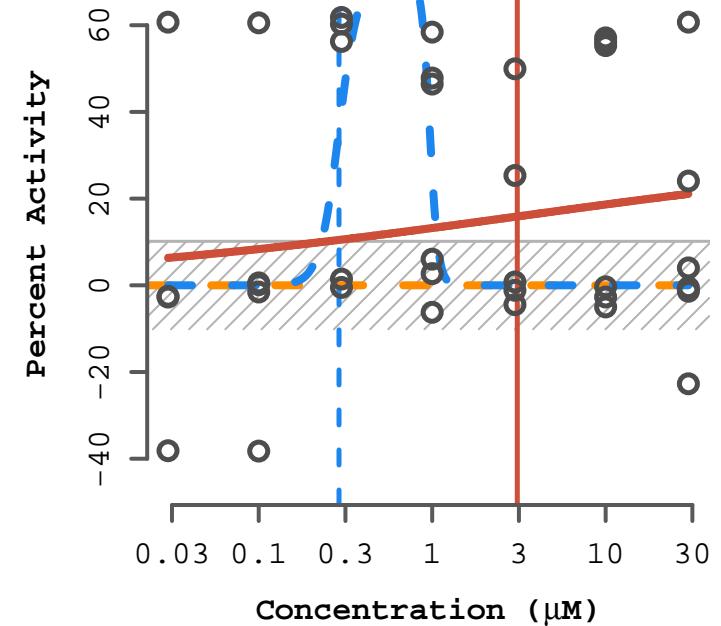
	tp	ga	gw	la	lw
val:	43.8	1	1.42	2.75	6.22
sd:	5.33	0.117	0.301	880	4760

	CNST	HILL	GNLS
AIC:	179.22	139.62	143.62
PROB:	0	0.88	0.12
RMSE:	19.66	7.56	7.56

MAX_MEAN: 46 MAX_MED: 37.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Ampicillin

CHID: 22602 CASRN: 69-53-4

SPID(S): EPAPLT0167C02

M4ID: 30204237

HILL MODEL (in red):

tp	ga	gw
val: 31.8	0.491	0.3
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

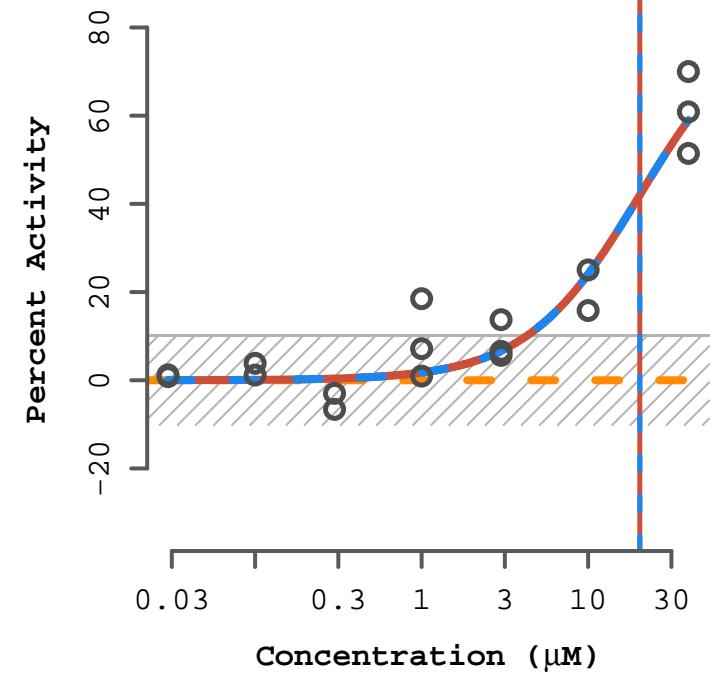
tp	ga	gw	la	lw
val: 71.8	-0.537	7.83	-0.014	17.9
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 373.13	371.61	373.76
PROB: 0.26	0.55	0.19
RMSE: 35.38	30.77	31.18

MAX_MEAN: 35.9 MAX_MED: 56.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.74

FLAGS: 17; 11; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Mifepristone

CHID: 23322 CASRN: 84371-65-3

SPID(S): TP0001414D01

M4ID: 30204656

HILL MODEL (in red):

	tp	ga	gw
val:	84	1.31	1.28
sd:	21.4	0.211	0.319

GAIN-LOSS MODEL (in blue):

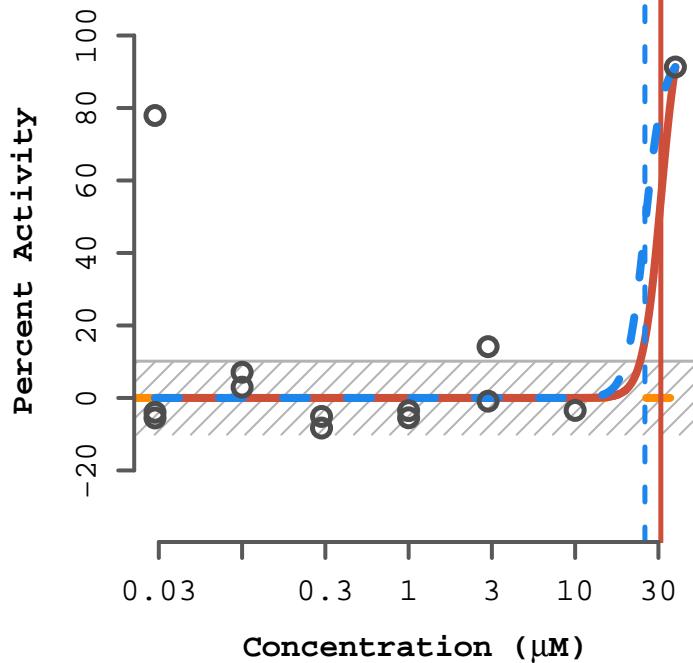
	tp	ga	gw	la	lw
val:	84	1.31	1.28	3.21	14
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	159.13	118.51	122.51
PROB:	0	0.88	0.12
RMSE:	27.5	6.37	6.37

MAX_MEAN: 60.8 MAX_MED: 60.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Troglitazone

CHID: 23719 CASRN: 97322-87-7

SPID(S): TP0001414E12

M4ID: 30204511

HILL MODEL (in red):

	tp	ga	gw
val:	110	1.51	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

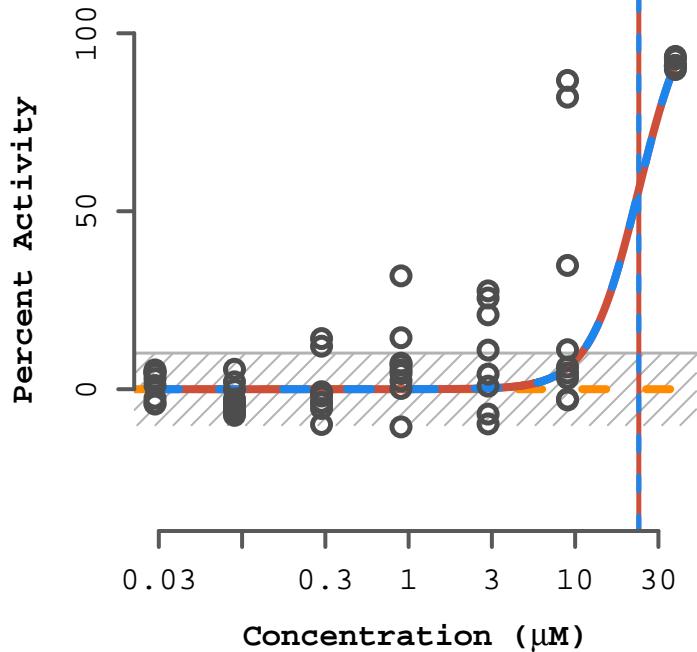
	tp	ga	gw	la	lw
val:	94.4	1.42	8	3.6	4.53
sd:	52.6	0.691	16.7	32500	73800

	CNST	HILL	GNLS
AIC:	121.64	110.86	114.86
PROB:	0	0.88	0.12
RMSE:	33.82	22.41	22.41

MAX_MEAN: 91.3 MAX_MED: 91.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pentamidine isethionate
CHID: 23796 CASRN: 140-64-7
SPID(S): TP0001414C04
M4ID: 30204578

HILL MODEL (in red):

	tp	ga	gw
val:	112	1.38	2.91
sd:	112	0.569	3.05

GAIN-LOSS MODEL (in blue):

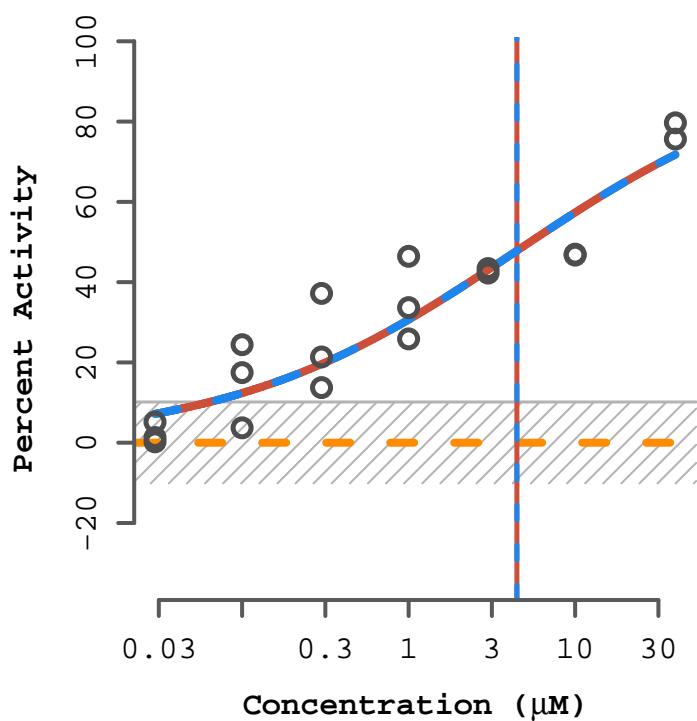
	tp	ga	gw	la	lw
val:	112	1.38	2.91	2.7	10.3
sd:	112	0.572	3.06	66200	614000

	CNST	HILL	GNLS
AIC:	573.48	470.4	474.4
PROB:	0	0.88	0.12
RMSE:	36.19	17.22	17.22

MAX_MEAN: 91.4 MAX_MED: 91 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Cypermethrin

CHID: 23998 CASRN: 52315-07-8

SPID(S): TP0001413H11

M4ID: 30204308

HILL MODEL (in red):

	tp	ga	gw
val:	95.6	0.65	0.502
sd:	23.2	0.486	0.109

GAIN-LOSS MODEL (in blue):

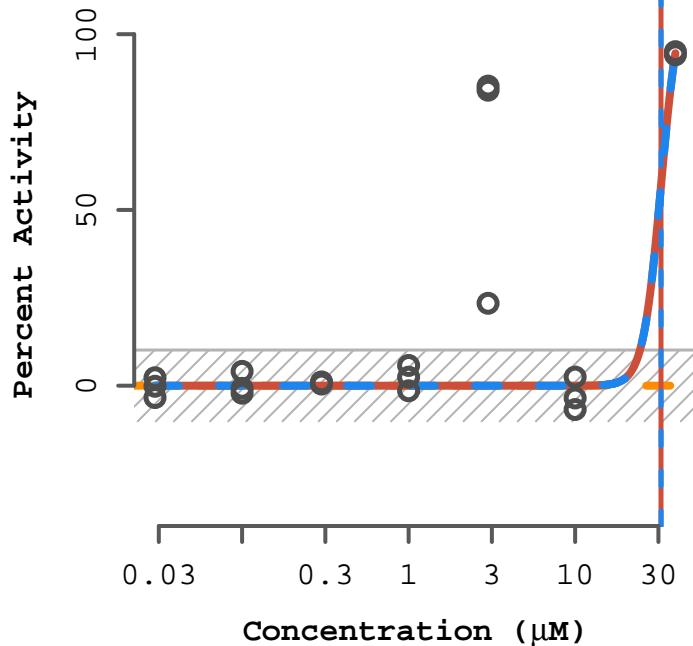
	tp	ga	gw	la	lw
val:	95.6	0.65	0.502	3.31	17.8
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	186.47	137.13	141.13
PROB:	0	0.88	0.12
RMSE:	38.78	8.37	8.37

MAX_MEAN: 77.7 MAX_MED: 77.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Difenozoquat metilsulfate
CHID: 24048 CASRN: 43222-48-6
SPID(S): TP0001413A12
M4ID: 30204420

HILL MODEL (in red):

	tp	ga	gw
val:	114	1.52	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

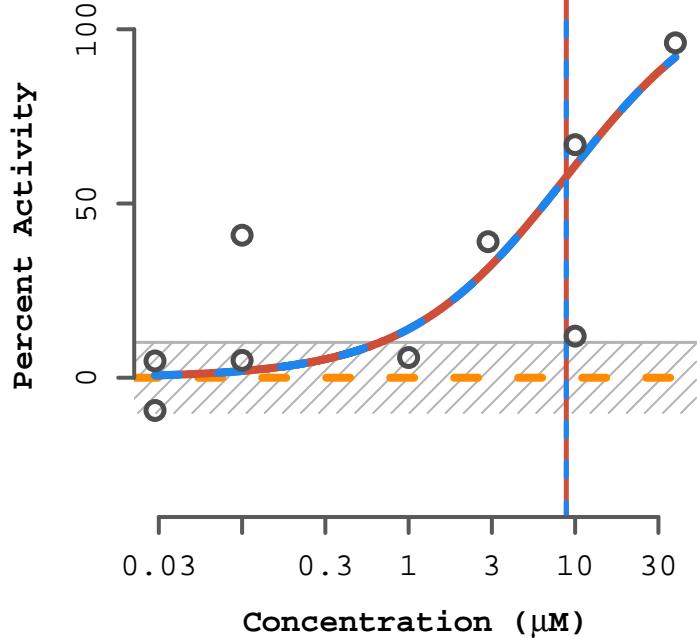
	tp	ga	gw	la	lw
val:	114	1.52	7.99	3.6	3.6
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	198.54	170.36	174.36
PROB:	0	0.88	0.12
RMSE:	40.58	27.42	27.42

MAX_MEAN: 94.6 MAX_MED: 94.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Imazalil

CHID: 24151 CASRN: 35554-44-0

SPID(S): TP0001411A09

M4ID: 30204286

HILL MODEL (in red):

	tp	ga	gw
val:	115	0.947	0.909
sd:	45.3	0.494	0.503

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	115	0.947	0.909	2.32	10
sd:	45.3	0.494	0.503	296	4150

CNST

AIC: 95.3

PROB: 0.02

RMSE: 43.75

HILL

87.42

0.87

21.65

GNLS

91.42

0.12

21.65

MAX_MEAN: 96.1

MAX_MED: 96.1

BMAD: 3.38

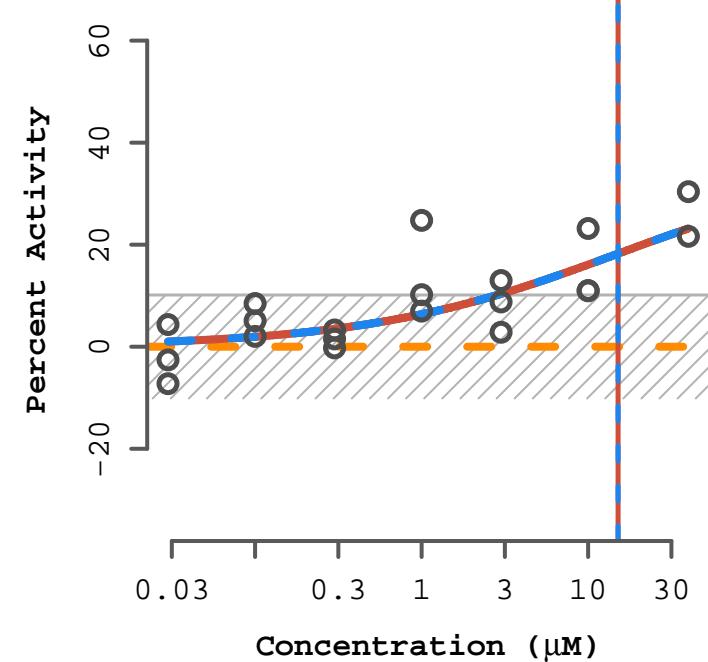
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 0.98

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Linuron

CHID: 24163 CASRN: 330-55-2

SPID(S): TP0001414F08

M4ID: 30204351

HILL MODEL (in red):

tp	ga	gw
36.5	1.18	0.571
18.8	0.809	0.194

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
36.5	1.18	0.571	2.88	5.69
18.7	0.809	0.194	481	2140

CNST

AIC: 161.49

PROB: 0

RMSE: 13.01

HILL

135.63

0.88

6.08

GNLS

139.63

0.12

6.08

MAX_MEAN: 26

MAX_MED: 26

BMAD: 3.38

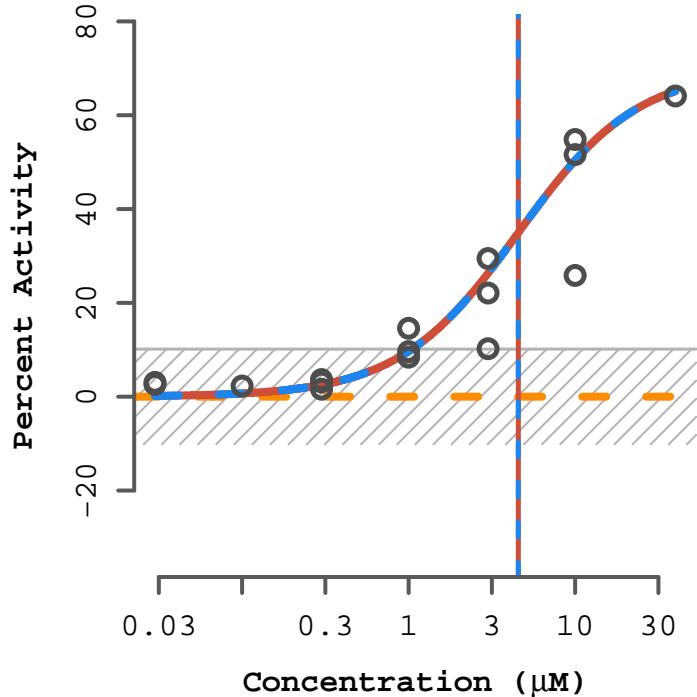
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 1

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Flusilazole

CHID: 24235 CASRN: 85509-19-9

SPID(S): TP0001414H11

M4ID: 30204288

HILL MODEL (in red):

	tp	ga	gw
val:	69.9	0.659	1.21
sd:	5.4	0.0952	0.183

GAIN-LOSS MODEL (in blue):

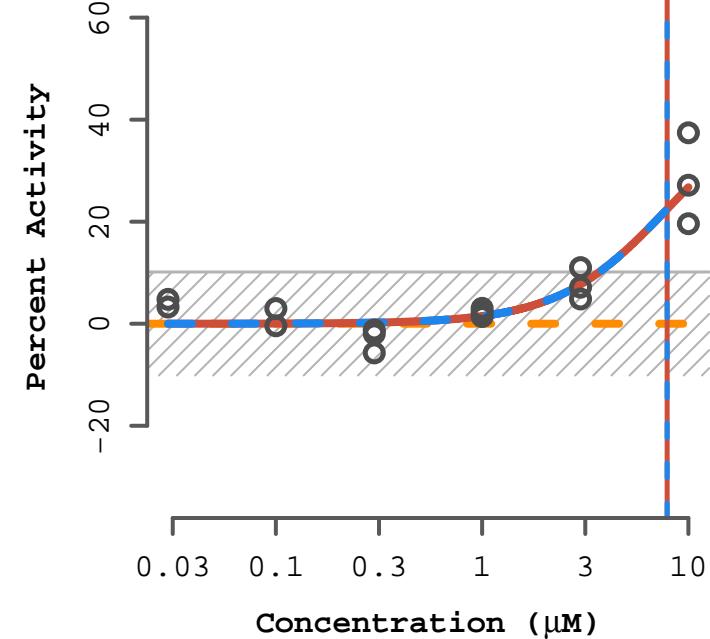
	tp	ga	gw	la	lw
val:	69.9	0.659	1.21	2.88	13.5
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	152.63	110.06	114.06
PROB:	0	0.88	0.12
RMSE:	27.77	7.73	7.73

MAX_MEAN: 64.1 MAX_MED: 64.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Oryzalin

CHID: 24238 CASRN: 19044-88-3

SPID(S): TP0001411C03

M4ID: 30204610

HILL MODEL (in red):

tp	ga	gw
val: 44.9	0.897	1.63
sd: 46.4	0.607	0.969

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 44.9	0.897	1.63	1.47	17.8
sd: 46.1	0.605	0.964	3320	126000

CNST

AIC: 125.73

PROB: 0

RMSE: 13.27

HILL

98.03

0.88

4.12

GNLS

102.03

0.12

4.12

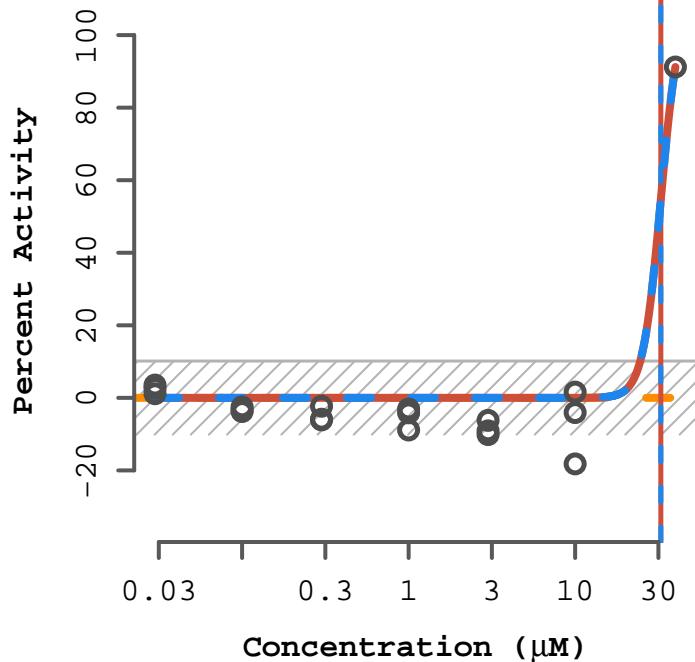
MAX_MEAN: 28.1

MAX_MED: 27.2

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 6



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Propargite

CHID: 24276 CASRN: 2312-35-8

SPID(S): TP0001412C03

M4ID: 30204378

HILL MODEL (in red):

tp	ga	gw
val: 109	1.51	8
sd: 488	0.353	103

GAIN-LOSS MODEL (in blue):

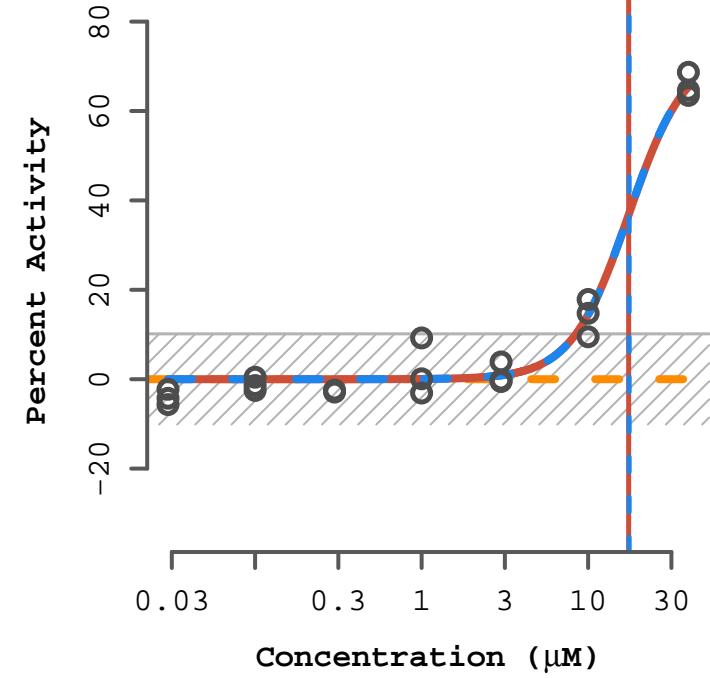
tp	ga	gw	la	lw
val: 109	1.51	8	2.26	5.3
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 147.55	132.69	136.69
PROB: 0	0.88	0.12
RMSE: 21.92	6.52	6.52

MAX_MEAN: 91.2 MAX_MED: 91.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Propiconazole

CHID: 24280 CASRN: 60207-90-1

SPID(S): TP0001413E10

M4ID: 30204341

HILL MODEL (in red):

	tp	ga	gw
val:	73.6	1.24	2.51
sd:	21.9	0.227	1.63

GAIN-LOSS MODEL (in blue):

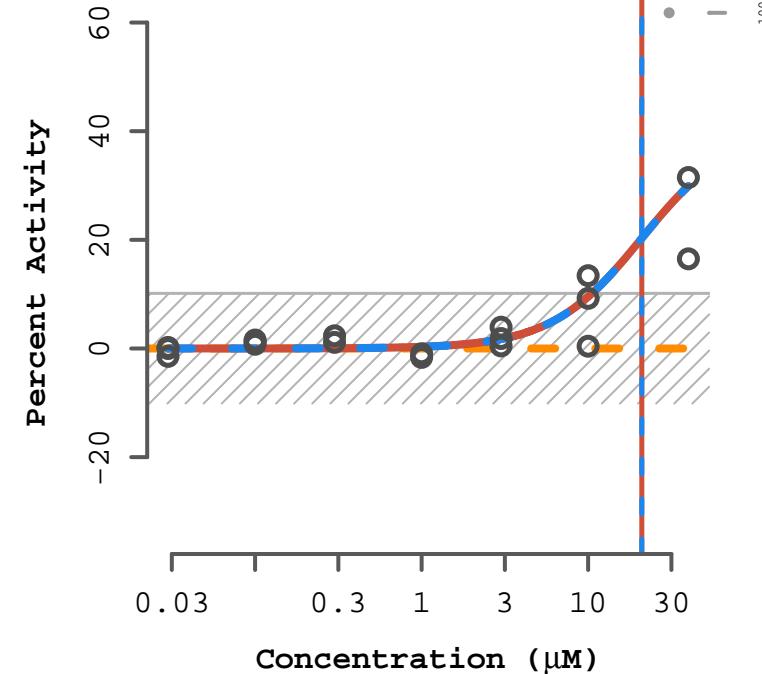
	tp	ga	gw	la	lw
val:	74.6	1.25	2.51	3.02	1.37
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	178.59	114.47	118.47
PROB:	0	0.88	0.12
RMSE:	26.21	3.43	3.43

MAX_MEAN: 65.6 MAX_MED: 64.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Hexythiazox

CHID: 24299 CASRN: 78587-05-0

SPID(S): TP0001413H07

M4ID: 30204661 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	40.6	1.32	1.6
sd:	22.6	0.411	0.85

GAIN-LOSS MODEL (in blue):

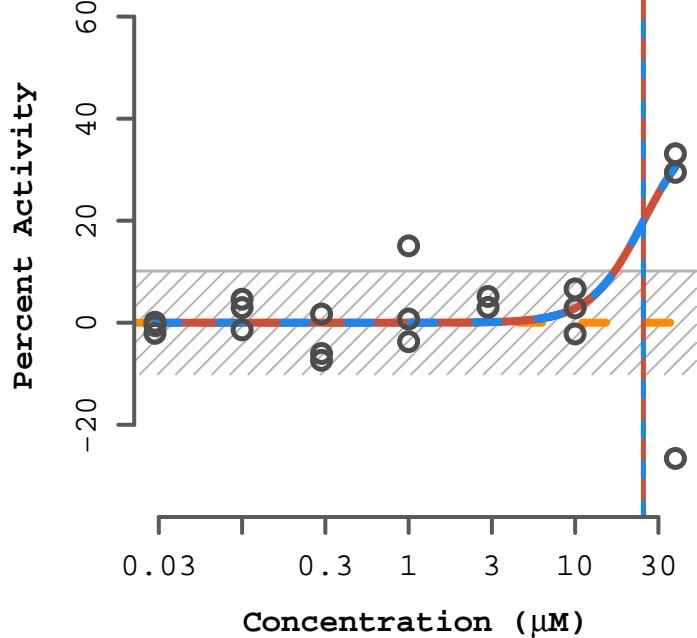
	tp	ga	gw	la	lw
val:	40.6	1.32	1.6	2.33	15.7
sd:	22.6	0.411	0.85	93600	1700000

	CNST	HILL	GNLS
AIC:	151.82	131.79	135.79
PROB:	0	0.88	0.12
RMSE:	24.62	16.52	16.52

MAX_MEAN: 49.2 MAX_MED: 31.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sethoxydim

CHID: 24304 CASRN: 74051-80-2

SPID(S): TP0001411D11

M4ID: 30204360

HILL MODEL (in red):

	tp	ga	gw
val:	39.8	1.41	2.71
sd:	72.3	0.931	4.71

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	39.8	1.41	2.71	3.55	17.5
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	154.06	144.62	148.62
PROB:	0.01	0.87	0.12
RMSE:	12.53	13.63	13.63

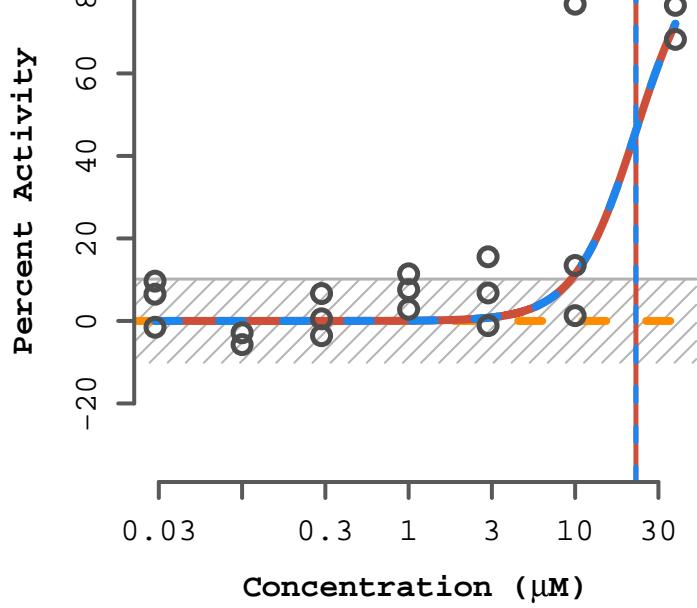
MAX_MEAN: 12

MAX_MED: 29.5

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.99

FLAGS: 17; 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tri-allate

CHID: 24344 CASRN: 2303-17-5

SPID(S): TP0001412B03

M4ID: 30204423

HILL MODEL (in red):

	tp	ga	gw
val:	92.1	1.36	2.34
sd:	79.9	0.491	2.54

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	92.1	1.36	2.34	3.6	6.12
sd:	80.3	0.494	2.55	163000	454000

CNST

AIC: 175.88

PROB: 0

RMSE: 30.19

HILL

151.87

0.88

16.4

GNLS

155.87

0.12

16.4

MAX_MEAN: 72.4

MAX_MED: 72.4

BMAD: 3.38

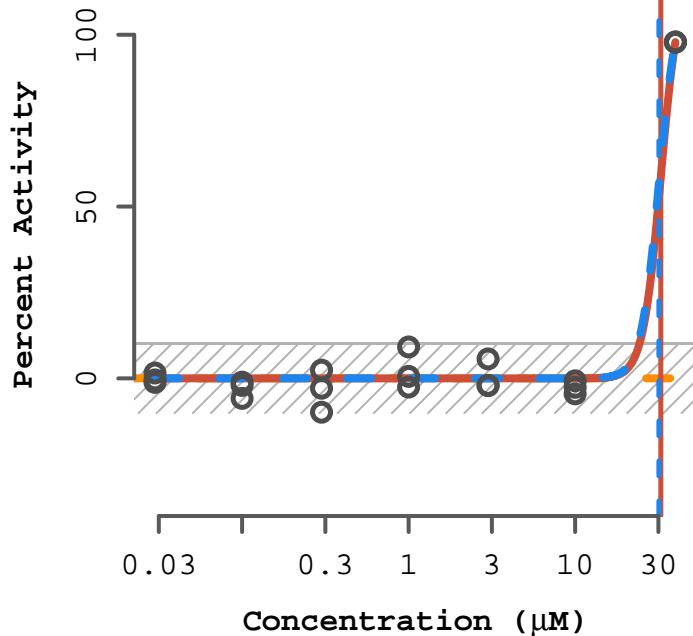
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,4,5-Trichlorophenol
CHID: 24359 CASRN: 95-95-4
SPID(S): TP0001414A02
M4ID: 30204471

HILL MODEL (in red):

	tp	ga	gw
val:	117	1.51	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

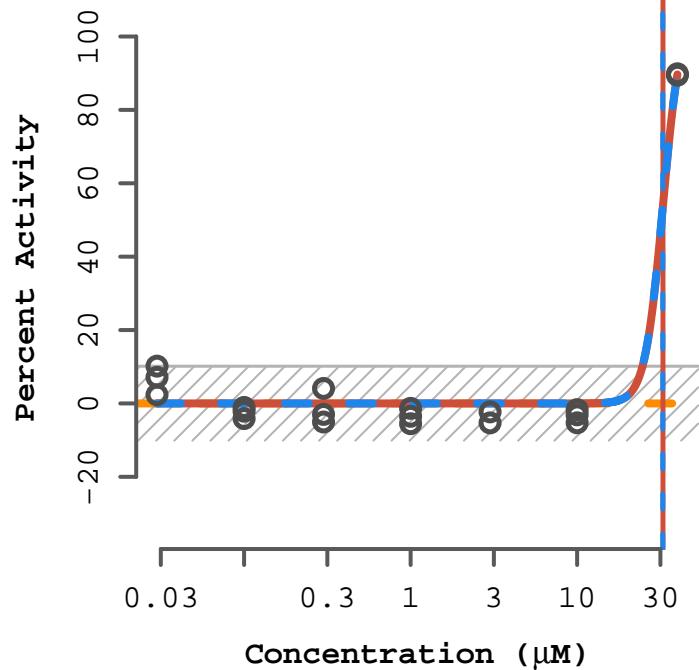
	tp	ga	gw	la	lw
val:	114	1.5	8	1.8	18
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	190.55	122.43	126.43
PROB:	0	0.88	0.12
RMSE:	37.19	3.86	3.86

MAX_MEAN: 97.9 MAX_MED: 97.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bronopol

CHID: 24652 CASRN: 52-51-7

SPID(S): TP0001411G06

M4ID: 30204435

HILL MODEL (in red):

tp	ga	gw
val: 108	1.52	8
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

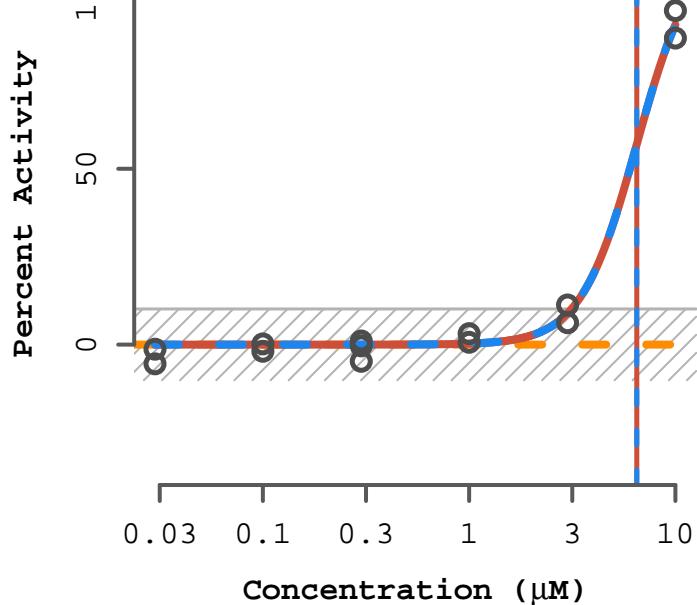
tp	ga	gw	la	lw
val: 107	1.51	8	1.85	17.9
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 158.38	118.94	122.94
PROB: 0	0.88	0.12
RMSE: 29.4	4.3	4.3

MAX_MEAN: 89.7 MAX_MED: 89.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: tert-Butylphenyl diphenyl phosphate

CHID: 24701 CASRN: 56803-37-3

SPID(S): EPAPLT0169B03

M4ID: 30204273

HILL MODEL (in red):

	tp	ga	gw
val:	114	0.813	3.19
sd:	84.8	0.36	2.44

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	114	0.813	3.19	2.8	3.85
sd:	88.1	0.374	2.54	1210	2590

CNST

AIC: 120.65

PROB: 0

RMSE: 36.01

HILL

74.17

0.88

2.92

GNLS

78.17

0.12

2.91

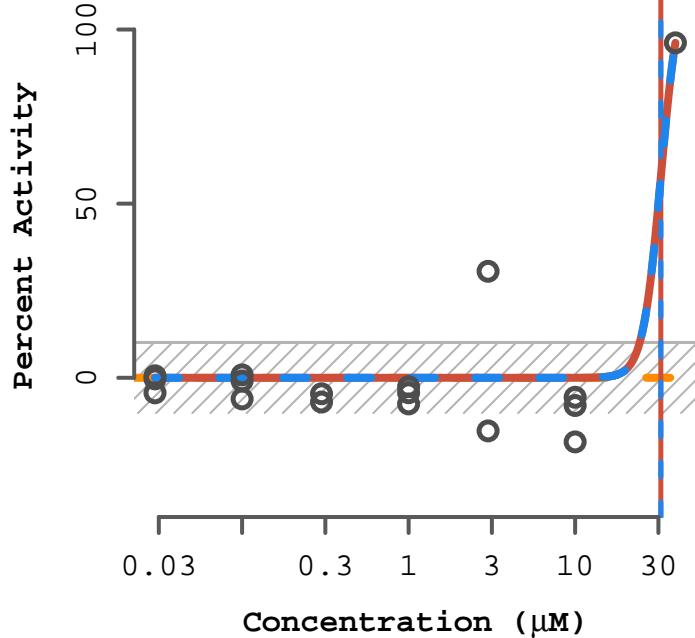
MAX_MEAN: 91.1

MAX_MED: 91.1

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 6



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Chlorpromazine hydrochloride
CHID: 24827 CASRN: 69-09-0
SPID(S): TP0001414F05
M4ID: 30204561

HILL MODEL (in red):

tp	ga	gw
val: 115	1.51	8
sd: 87.5	0.117	15.1

GAIN-LOSS MODEL (in blue):

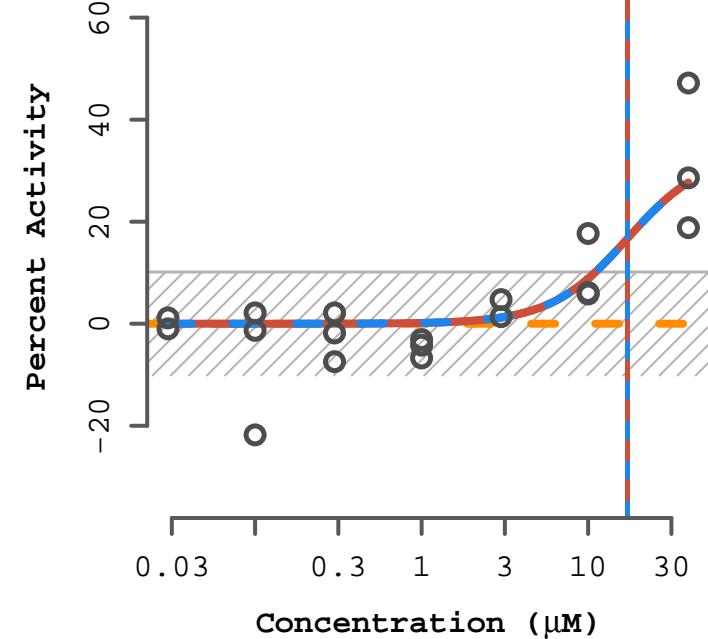
tp	ga	gw	la	lw
val: 115	1.51	8	3.6	3.57
sd: 87.6	0.117	15.1	8300	14900

CNST	HILL	GNLS
AIC: 144.43	132.16	136.16
PROB: 0	0.88	0.12
RMSE: 25.51	10.31	10.31

MAX_MEAN: 96.2 MAX_MED: 96.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 3,4-Diaminotoluene
CHID: 24930 CASRN: 496-72-0
SPID(S): TP0001411C12
M4ID: 30204558

HILL MODEL (in red):

tp	ga	gw
val: 33.4	1.24	1.87
sd: 18.9	0.414	1.56

GAIN-LOSS MODEL (in blue):

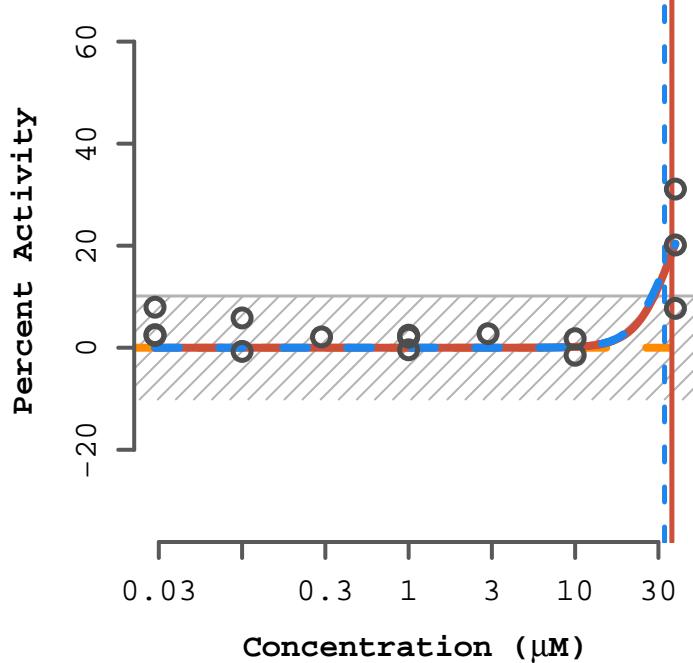
tp	ga	gw	la	lw
val: 33.4	1.24	1.87	2.06	11.7
sd: 18.9	0.414	1.56	236	5990

CNST	HILL	GNLS
AIC: 154.47	136.53	140.53
PROB: 0	0.88	0.12
RMSE: 15.26	7.91	7.91

MAX_MEAN: 31.5 MAX_MED: 28.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Sodium dodecylbenzenesulfonate

CHID: 25219 CASRN: 25155-30-0

SPID(S): TP0001413A03

M4ID: 30204664

HILL MODEL (in red):

	tp	ga	gw
val:	37.2	1.58	4.01
sd:	108	0.671	12.6

GAIN-LOSS MODEL (in blue):

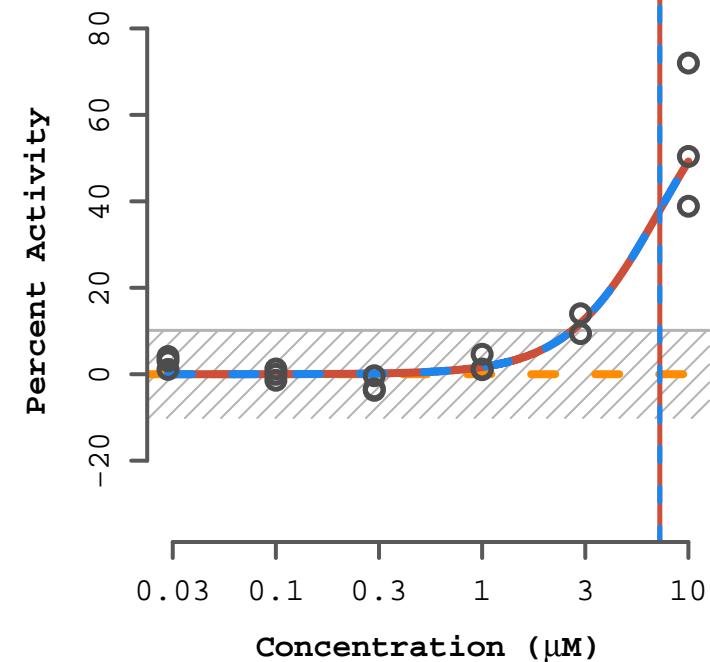
	tp	ga	gw	la	lw
val:	31.2	1.54	4.21	3.57	2.9
sd:	233	1.99	18.8	12700	18900

	CNST	HILL	GNLS
AIC:	107.87	98.98	102.98
PROB:	0.01	0.87	0.12
RMSE:	10.23	5.25	5.25

MAX_MEAN: 19.7 MAX_MED: 20.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.99

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Octhilinone

CHID: 25805 CASRN: 26530-20-1

SPID(S): TP0001411D10

M4ID: 30204589

HILL MODEL (in red):

	tp	ga	gw
val:	76	0.861	1.91
sd:	54.4	0.392	0.912

GAIN-LOSS MODEL (in blue):

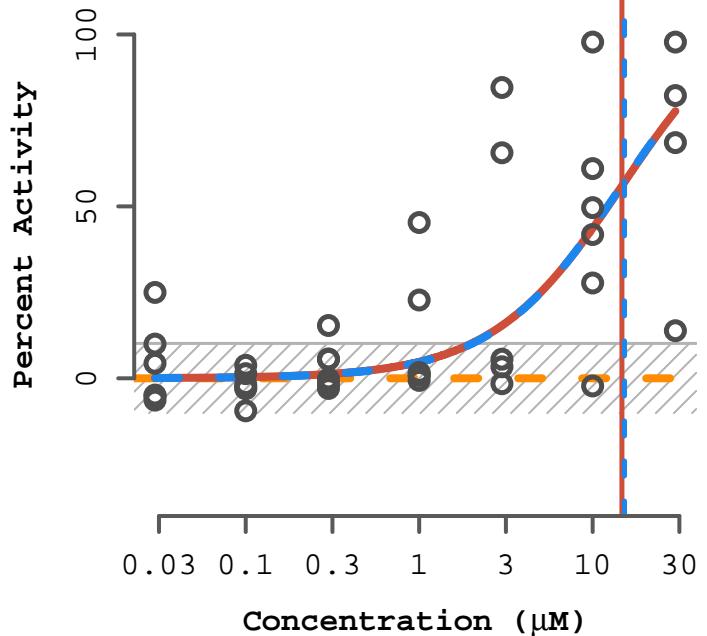
	tp	ga	gw	la	lw
val:	76	0.861	1.91	1.52	9.58
sd:	54.4	0.392	0.912	174	3190

	CNST	HILL	GNLS
AIC:	148.56	109.02	113.02
PROB:	0	0.88	0.12
RMSE:	23.76	6.44	6.44

MAX_MEAN: 53.8 MAX_MED: 50.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: N-Phenyl-1-naphthylamine

CHID: 25892 CASRN: 90-30-2

SPID(S): EPAPLT0154F05

M4ID: 30204208

HILL MODEL (in red):

	tp	ga	gw
val:	112	1.17	1.16
sd:	74.9	0.587	0.743

GAIN-LOSS MODEL (in blue):

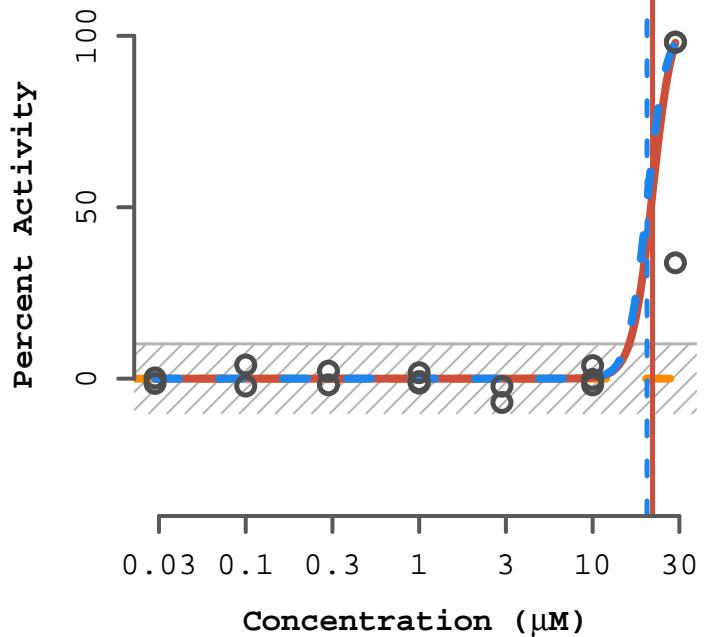
	tp	ga	gw	la	lw
val:	117	1.18	1.17	3.43	0.72
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	387.43	354.1	358.1
PROB:	0	0.88	0.12
RMSE:	37.5	23.38	23.39

MAX_MEAN: 65.6 MAX_MED: 75.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tris(1,3-dichloro-2-propyl) phosphate

CHID: 26261 CASRN: 13674-87-8

SPID(S): EPAPLT0167H02

M4ID: 30204281

HILL MODEL (in red):

	tp	ga	gw
val:	109	1.35	7.33
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

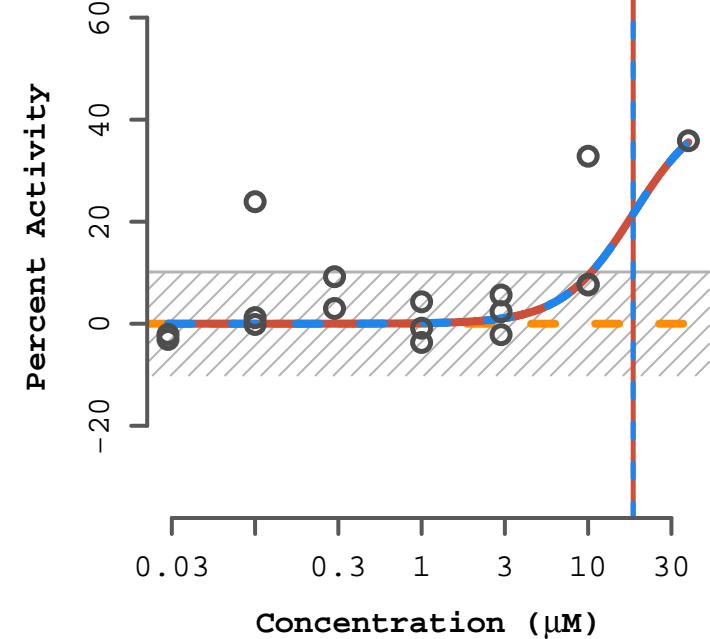
	tp	ga	gw	la	lw
val:	103	1.31	7.99	2.55	3.58
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	152.78	113.41	117.41
PROB:	0	0.88	0.12
RMSE:	34.75	15.76	15.76

MAX_MEAN: 76.7 MAX_MED: 98.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-tert-Butylphenol
CHID: 26525 CASRN: 88-18-6
SPID(S): TP0001413H02
M4ID: 30204357

HILL MODEL (in red):

	tp	ga	gw
val:	43.1	1.27	2.04
sd:	23.2	0.375	1.74

GAIN-LOSS MODEL (in blue):

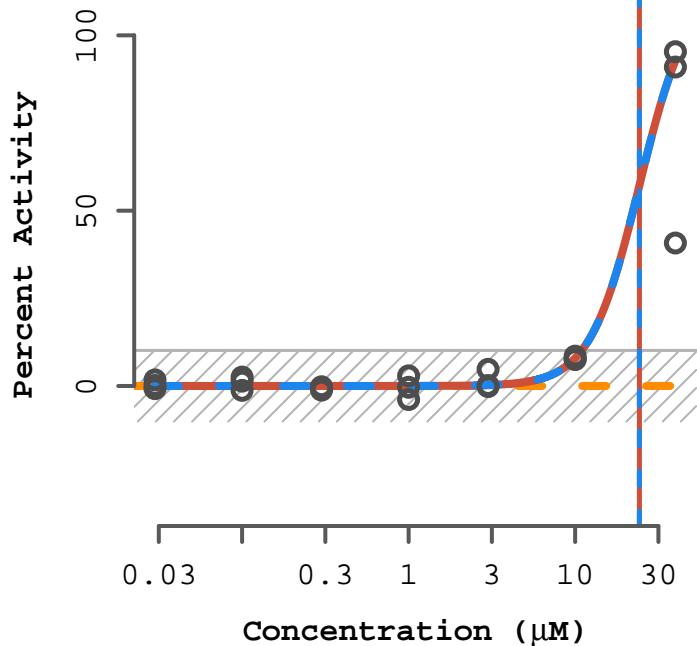
	tp	ga	gw	la	lw
val:	43.1	1.27	2.04	3.22	4.9
sd:	23.2	0.375	1.74	2980	9000

	CNST	HILL	GNLS
AIC:	134.52	122.26	126.26
PROB:	0	0.88	0.12
RMSE:	13.81	8.75	8.75

MAX_MEAN: 35.9 MAX_MED: 35.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1-Dodecanol

CHID: 26918 CASRN: 112-53-8

SPID(S): TP0001412A03

M4ID: 30204431

HILL MODEL (in red):

	tp	ga	gw
val:	114	1.39	2.92
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

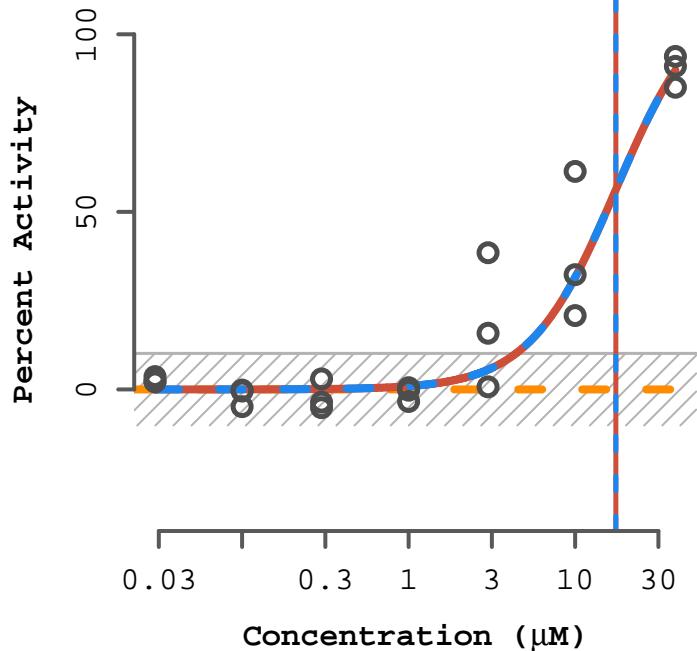
	tp	ga	gw	la	lw
val:	114	1.39	2.92	3.6	3.99
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	179.7	120.28	124.28
PROB:	0	0.88	0.12
RMSE:	30.3	11.49	11.49

MAX_MEAN: 75.7 MAX_MED: 91 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,4-Di-tert-pentylphenol
CHID: 26974 CASRN: 120-95-6
SPID(S): TP0001413A08
M4ID: 30204413

HILL MODEL (in red):

tp	ga	gw
val: 113	1.24	1.67
sd: 26.8	0.183	0.6

GAIN-LOSS MODEL (in blue):

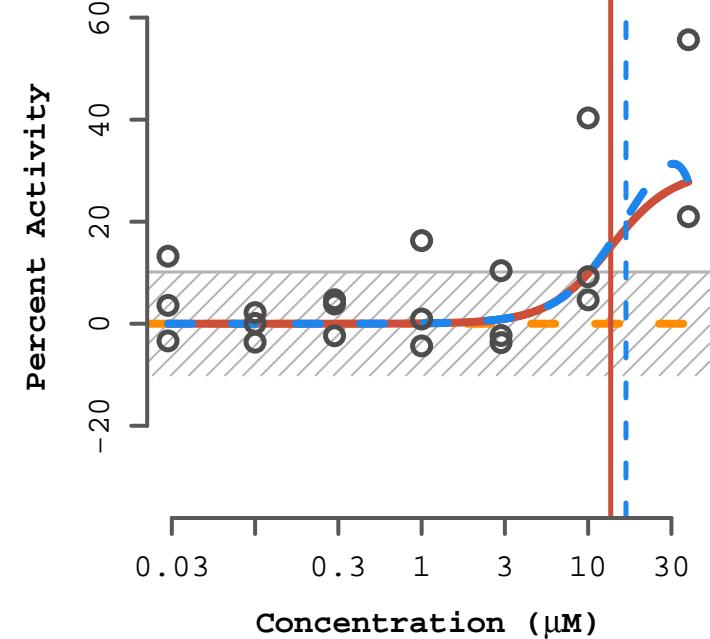
tp	ga	gw	la	lw
val: 113	1.24	1.67	2.97	7.58
sd: 26.8	0.183	0.599	20400	113000

CNST	HILL	GNLS
AIC: 209.98	156.24	160.24
PROB: 0	0.88	0.12
RMSE: 38.67	10.64	10.64

MAX_MEAN: 90 MAX_MED: 91 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: alpha-Isomethylionone
CHID: 27047 CASRN: 127-51-5
SPID(S): TP0001411F03
M4ID: 30204450

HILL MODEL (in red):

	tp	ga	gw
val:	30.1	1.14	2.32
sd:	29.5	0.688	6.78

GAIN-LOSS MODEL (in blue):

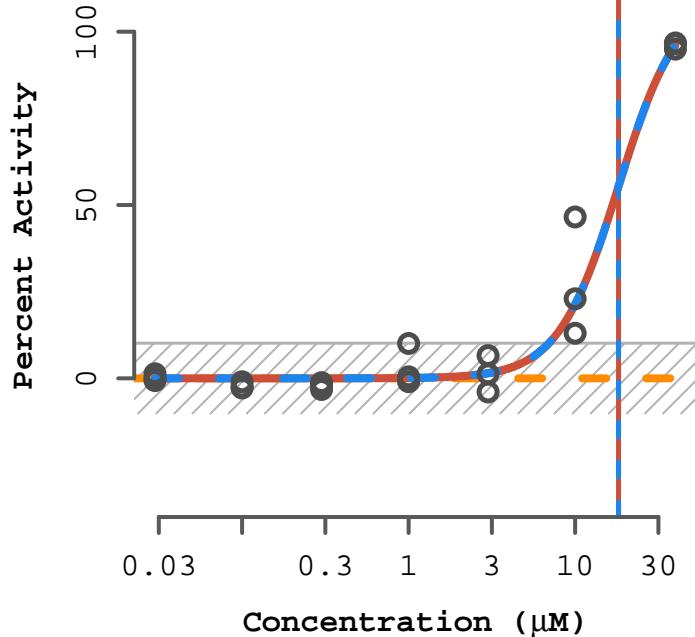
	tp	ga	gw	la	lw
val:	41.1	1.23	2.21	1.68	7.44
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	164.14	156.22	160.22
PROB:	0.02	0.87	0.12
RMSE:	17.25	11.06	11.06

MAX_MEAN: 38.3 MAX_MED: 38.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.98

FLAGS: 17; 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 3-Iodo-2-propynyl-N-butylcarbamate

CHID: 28038 CASRN: 55406-53-6

SPID(S): TP0001413G11

M4ID: 30204334

HILL MODEL (in red):

	tp	ga	gw
val:	111	1.26	2.39
sd:	32.3	0.215	1.3

GAIN-LOSS MODEL (in blue):

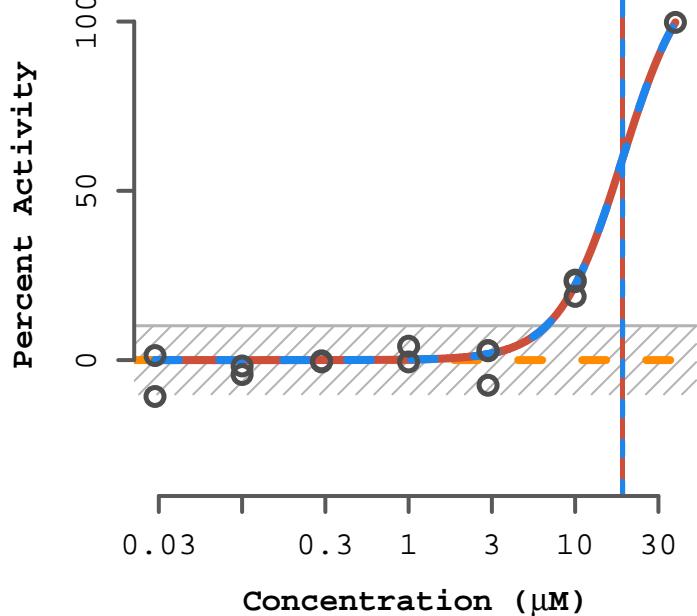
	tp	ga	gw	la	lw
val:	111	1.26	2.39	2.83	8.59
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	203.6	132.61	136.61
PROB:	0	0.88	0.12
RMSE:	38.27	6.53	6.53

MAX_MEAN: 96.1 MAX_MED: 96.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Nonylphenol, branched
CHID: 29055 CASRN: 84852-15-3
SPID(S): TP0001413H06
M4ID: 30204592

HILL MODEL (in red):

	tp	ga	gw
val:	119	1.28	2.25
sd:	39.2	0.228	1.17

GAIN-LOSS MODEL (in blue):

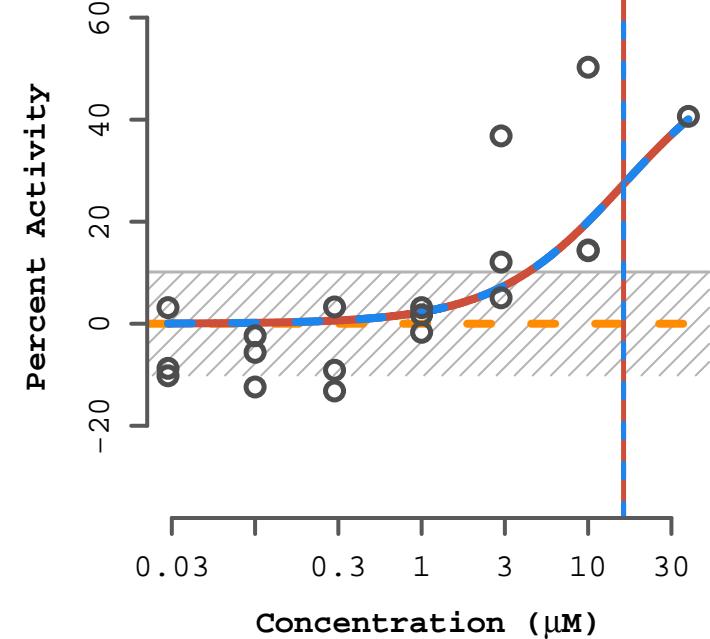
	tp	ga	gw	la	lw
val:	120	1.29	2.25	3.52	1.21
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	131.16	90.02	94.02
PROB:	0	0.88	0.12
RMSE:	27.84	4.13	4.13

MAX_MEAN: 99.8 MAX_MED: 99.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Dipentyl phthalate
CHID: 31131 CASRN: 131-18-0
SPID(S): TP0001411B10
M4ID: 30204572

HILL MODEL (in red):

tp	ga	gw
val: 54.7	1.21	1.12
sd: 34	0.608	0.525

GAIN-LOSS MODEL (in blue):

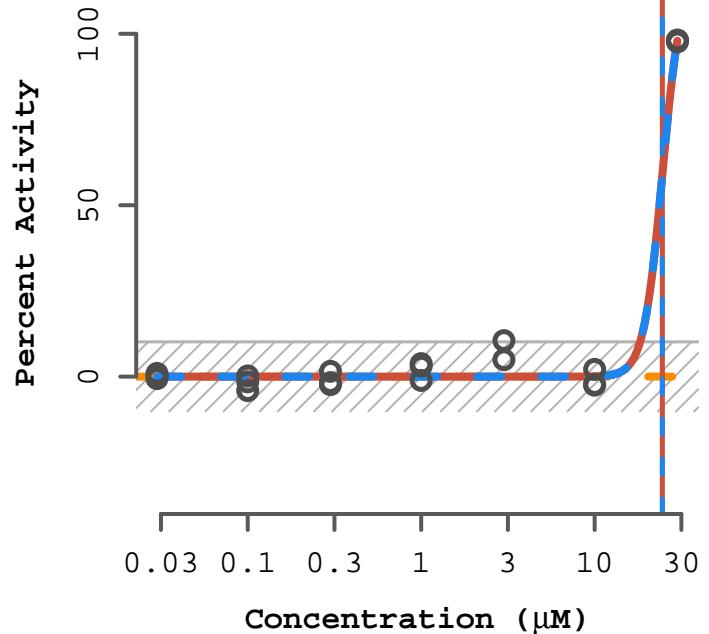
tp	ga	gw	la	lw
val: 54.7	1.21	1.12	2.62	6.54
sd: 34	0.609	0.525	572	3670

CNST	HILL	GNLS
AIC: 164.56	152.23	156.23
PROB: 0	0.88	0.12
RMSE: 18.9	11.67	11.67

MAX_MEAN: 40.7 MAX_MED: 40.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tri-o-cresyl phosphate
CHID: 32192 CASRN: 78-30-8
SPID(S): EPAPLT0167B02
M4ID: 30204279

HILL MODEL (in red):

	tp	ga	gw
val:	118	1.39	8
sd:	13.3	0.01	2.92

GAIN-LOSS MODEL (in blue):

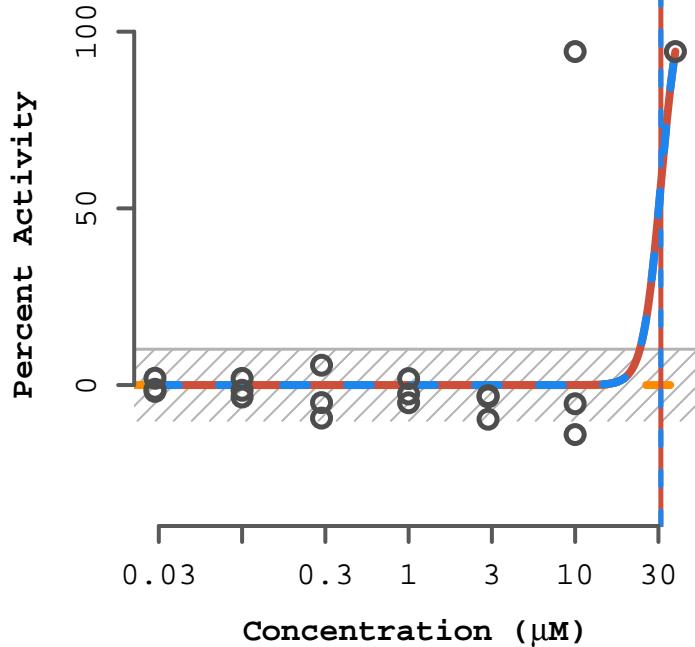
	tp	ga	gw	la	lw
val:	118	1.39	8	2.73	17.9
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	182.17	107.71	111.71
PROB:	0	0.88	0.12
RMSE:	38.08	3.22	3.22

MAX_MEAN: 98 MAX_MED: 98.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bensulide

CHID: 32329 CASRN: 741-58-2

SPID(S): TP0001414B05

M4ID: 30204297

HILL MODEL (in red):

	tp	ga	gw
val:	113	1.51	8
sd:	172	0.13	34.8

GAIN-LOSS MODEL (in blue):

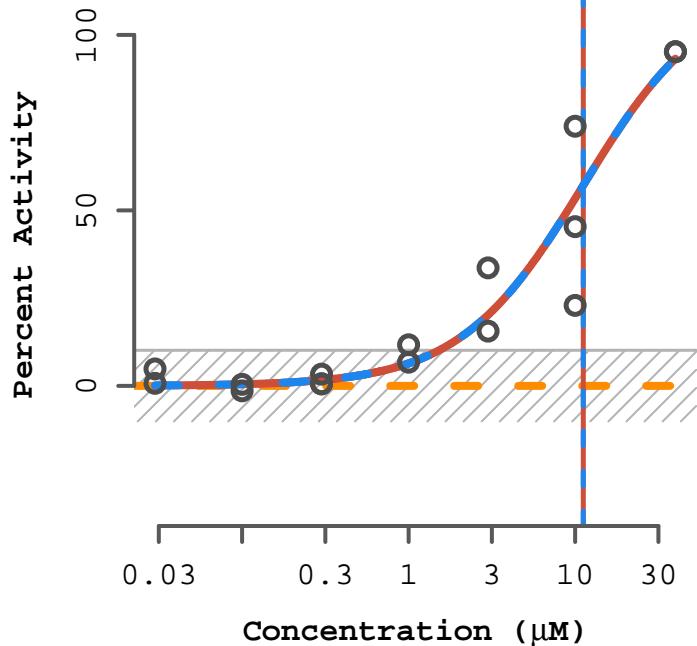
	tp	ga	gw	la	lw
val:	113	1.51	8	3.33	8.2
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	162.84	148.25	152.25
PROB:	0	0.88	0.12
RMSE:	31.09	22.3	22.3

MAX_MEAN: 94.4 MAX_MED: 94.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Difenoconazole

CHID: 32372 CASRN: 119446-68-3

SPID(S): TP0001414H12

M4ID: 30204314

HILL MODEL (in red):

	tp	ga	gw
val:	114	1.05	1.17
sd:	19.1	0.235	0.228

GAIN-LOSS MODEL (in blue):

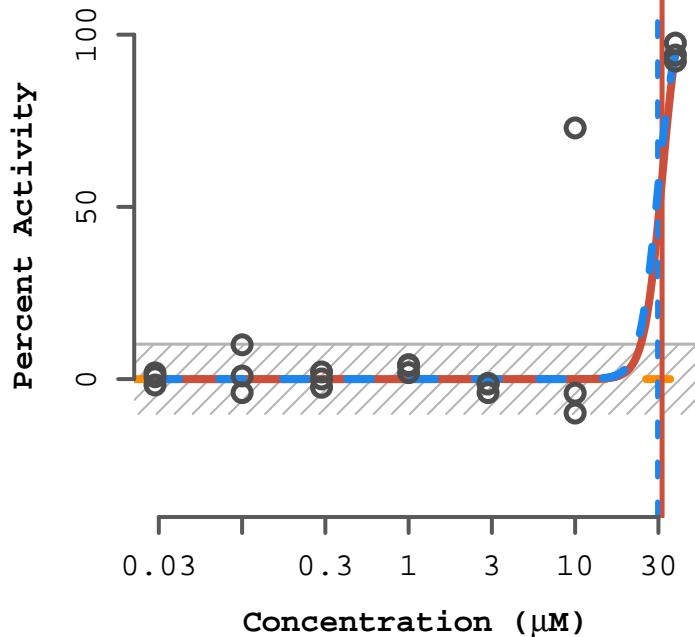
	tp	ga	gw	la	lw
val:	114	1.05	1.17	2.99	11.9
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	164.18	122.34	126.34
PROB:	0	0.88	0.12
RMSE:	41.68	10.26	10.26

MAX_MEAN: 95.2 MAX_MED: 95.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tebupirimfos

CHID: 32482 CASRN: 96182-53-5

SPID(S): TP0001414A04

M4ID: 30204386

HILL MODEL (in red):

	tp	ga	gw
val:	116	1.52	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

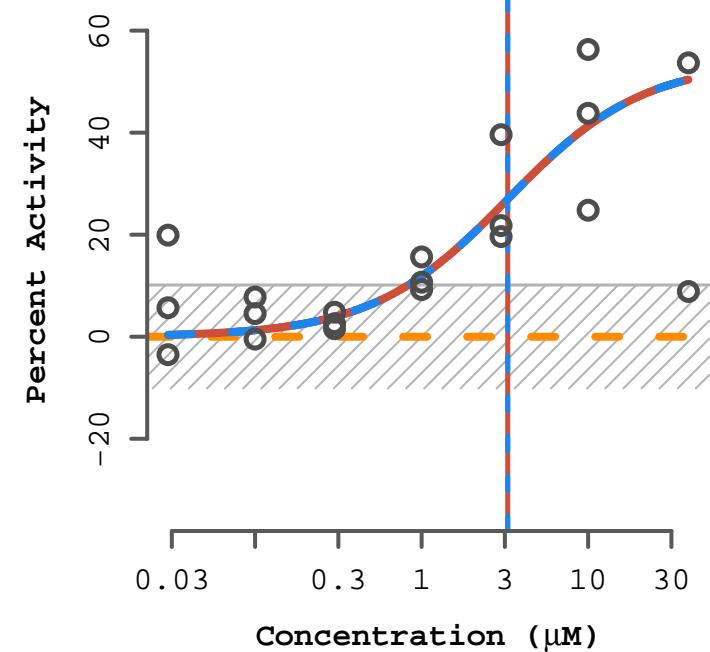
	tp	ga	gw	la	lw
val:	108	1.5	8	3.57	1.34
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	203.76	147.17	151.17
PROB:	0	0.88	0.12
RMSE:	39.35	16.38	16.38

MAX_MEAN: 94.7 MAX_MED: 94.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Triflumizole

CHID: 32500 CASRN: 68694-11-1

SPID(S): TP0001414G11

M4ID: 30204370

HILL MODEL (in red):

	tp	ga	gw
val:	53.9	0.517	1.07
sd:	11.4	0.259	0.41

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	53.9	0.517	1.07	2.61	9.36
sd:	NaN	NaN	NaN	NaN	NaN

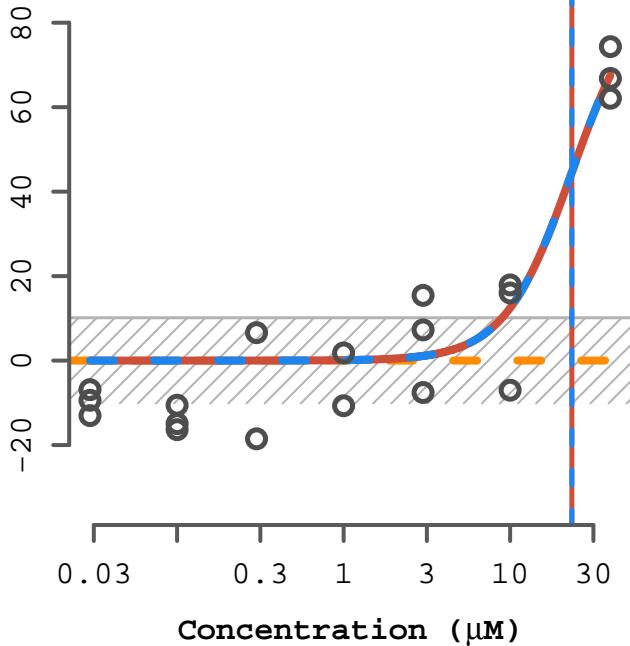
	CNST	HILL	GNLS
AIC:	186.04	158.28	162.28
PROB:	0	0.88	0.12
RMSE:	24.59	12.24	12.24

MAX_MEAN: 41.6 MAX_MED: 43.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 1,2-Benzisothiazolin-3-one
CHID: 32523 CASRN: 2634-33-5
SPID(S): TP0001412E05
M4ID: 30204547

HILL MODEL (in red):

	tp	ga	gw
val:	89.2	1.37	2.13
sd:	92.9	0.626	2.33

GAIN-LOSS MODEL (in blue):

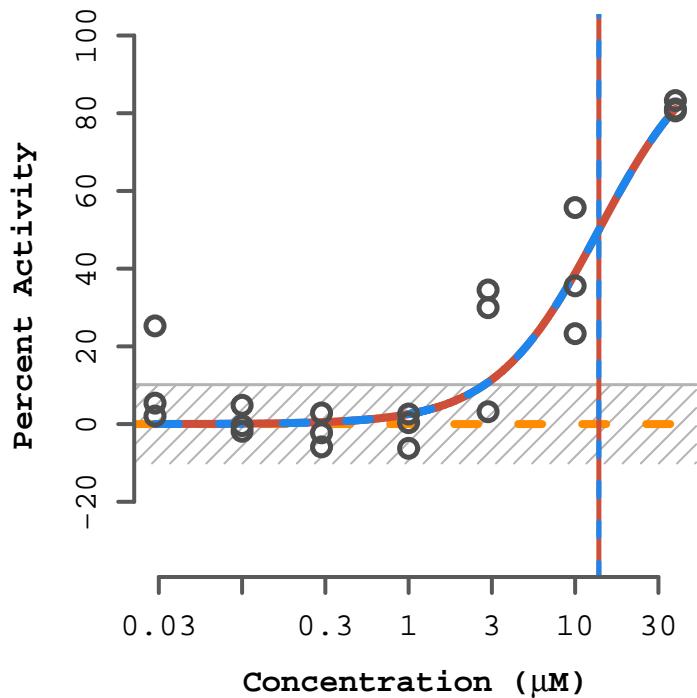
	tp	ga	gw	la	lw
val:	89.2	1.37	2.13	3.24	13.7
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	180.25	155.15	159.15
PROB:	0	0.88	0.12
RMSE:	29.23	10.83	10.83

MAX_MEAN: 67.8 MAX_MED: 66.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Didecyldimethylammonium chloride

CHID: 32537 CASRN: 7173-51-5

SPID(S): TP0001411H09

M4ID: 30204602

HILL MODEL (in red):

	tp	ga	gw
val:	99.9	1.14	1.38
sd:	20.5	0.201	0.47

GAIN-LOSS MODEL (in blue):

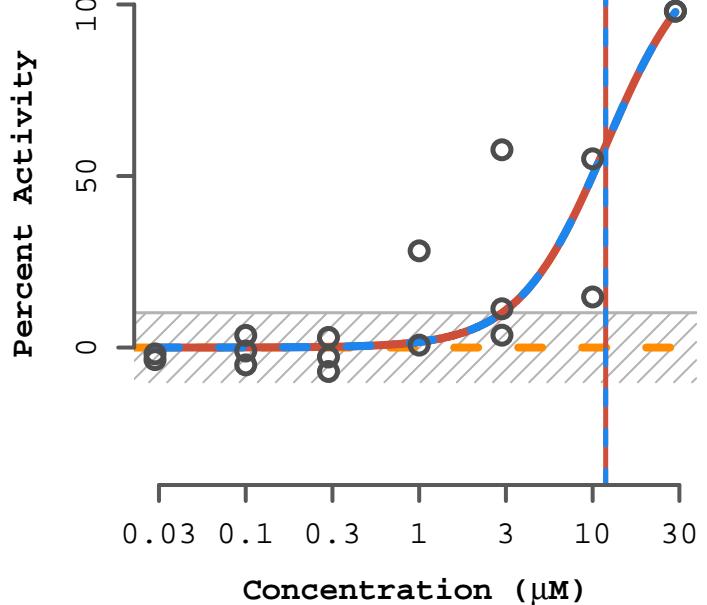
	tp	ga	gw	la	lw
val:	99.9	1.14	1.38	3.6	7.26
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	209.88	164.86	168.86
PROB:	0	0.88	0.12
RMSE:	36.4	10.62	10.62

MAX_MEAN: 81.7 MAX_MED: 81.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Flufenacet

CHID: 32552 CASRN: 142459-58-3

SPID(S): EPAPLT0167E04

M4ID: 30204258

HILL MODEL (in red):

	tp	ga	gw
val:	118	1.08	1.72
sd:	24.9	0.173	0.61

GAIN-LOSS MODEL (in blue):

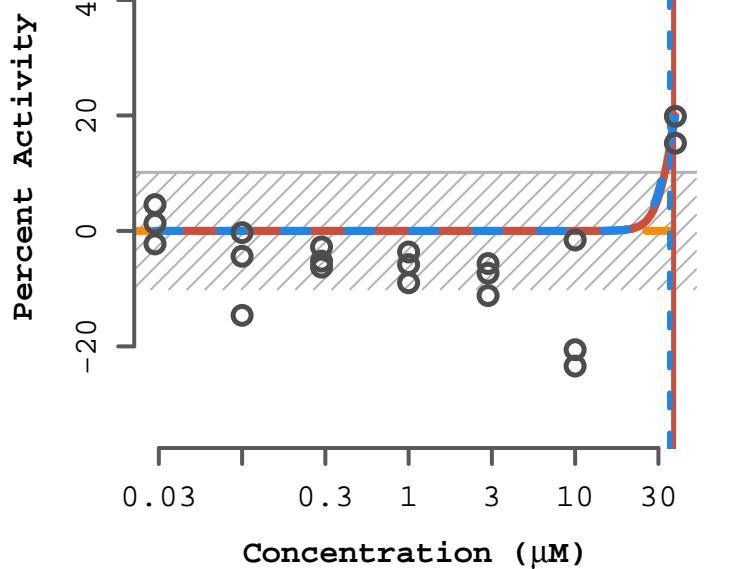
	tp	ga	gw	la	lw
val:	118	1.08	1.72	3.46	5.38
sd:	24.9	0.173	0.61	53900	146000

	CNST	HILL	GNLS
AIC:	170.93	142.31	146.31
PROB:	0	0.88	0.12
RMSE:	39.75	16.11	16.11

MAX_MEAN: 98.1 MAX_MED: 98.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Octylbicycloheptenedicarboximide

CHID: 32562 CASRN: 113-48-4

SPID(S): TP0001412A05

M4ID: 30204418 BRK

HILL MODEL (in red):

	tp	ga	gw
val:	36.3	1.59	7.88
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

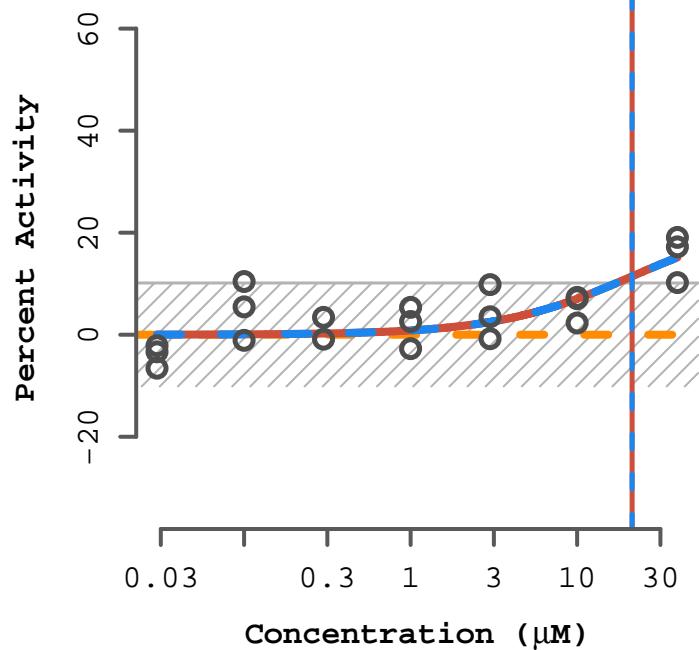
	tp	ga	gw	la	lw
val:	30.3	1.57	8	3.57	2.08
sd:	537	2.41	97.5	1670	1770

	CNST	HILL	GNLS
AIC:	174.42	171.18	175.18
PROB:	0.15	0.75	0.1
RMSE:	19.12	14.71	14.71

MAX_MEAN: 36.2 MAX_MED: 19.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.85

FLAGS: 17; 11; 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pirimicarb

CHID: 32569 CASRN: 23103-98-2

SPID(S): TP0001412F06

M4ID: 30204305

HILL MODEL (in red):

	tp	ga	gw
val:	22.9	1.33	1.07
sd:	12.3	0.482	0.644

GAIN-LOSS MODEL (in blue):

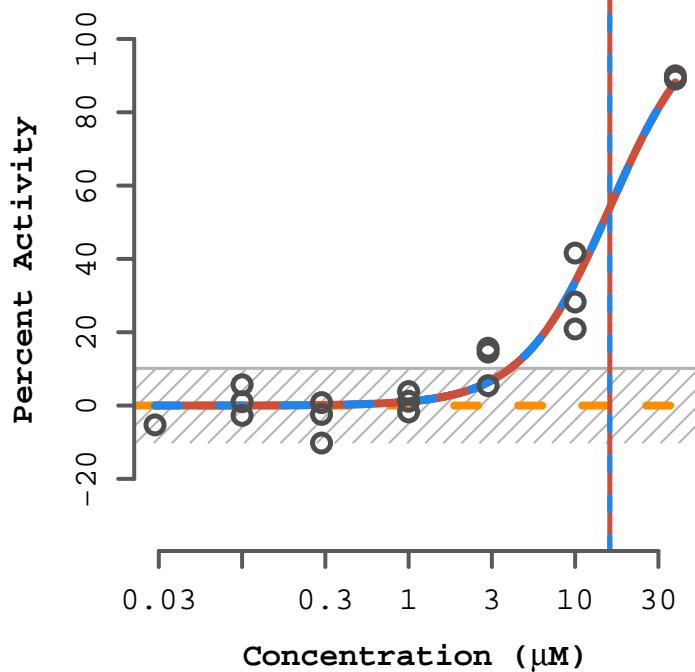
	tp	ga	gw	la	lw
val:	22.9	1.33	1.07	3.59	4.04
sd:	12.3	0.482	0.644	2350	4780

	CNST	HILL	GNLS
AIC:	141.21	124.96	128.96
PROB:	0	0.88	0.12
RMSE:	7.87	4.34	4.34

MAX_MEAN: 15.5 MAX_MED: 17.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Prallethrin

CHID: 32572 CASRN: 23031-36-9

SPID(S): TP0001414A12

M4ID: 30204367

HILL MODEL (in red):

	tp	ga	gw
val:	108	1.21	1.65
sd:	14.2	0.111	0.388

GAIN-LOSS MODEL (in blue):

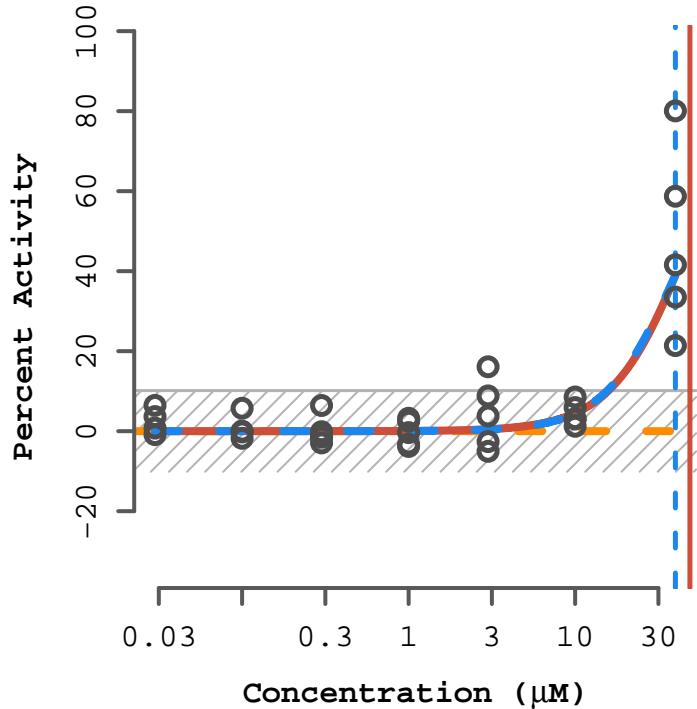
	tp	ga	gw	la	lw
val:	108	1.21	1.65	3.49	7.06
sd:	14.2	0.111	0.388	383000	1400000

	CNST	HILL	GNLS
AIC:	171.58	123.5	127.5
PROB:	0	0.88	0.12
RMSE:	33.06	5.84	5.84

MAX_MEAN: 89.5 MAX_MED: 89.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Thiodicarb

CHID: 32578 CASRN: 59669-26-0

SPID(S): TP0001413B05

M4ID: 30204373

HILL MODEL (in red):

	tp	ga	gw
val:	96.1	1.69	1.9
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

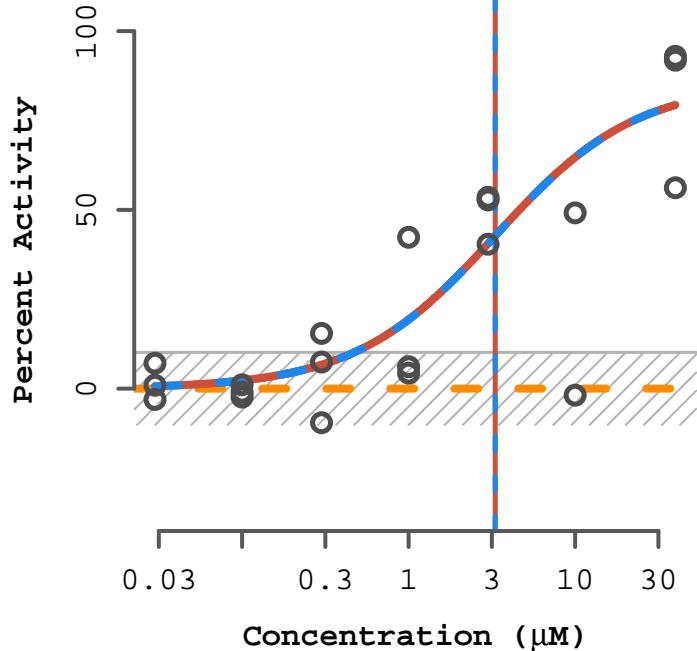
	tp	ga	gw	la	lw
val:	78.1	1.6	2.02	2.44	12.9
sd:	306	1.69	2.72	107000	1670000

	CNST	HILL	GNLS
AIC:	268.74	230.82	234.85
PROB:	0	0.88	0.12
RMSE:	20.5	9.55	9.55

MAX_MEAN: 47.1 MAX_MED: 41.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Trifloxystrobin

CHID: 32580 CASRN: 141517-21-7

SPID(S): TP0001414D11

M4ID: 30204310

HILL MODEL (in red):

	tp	ga	gw
val:	85.5	0.52	1.03
sd:	28	0.454	0.626

GAIN-LOSS MODEL (in blue):

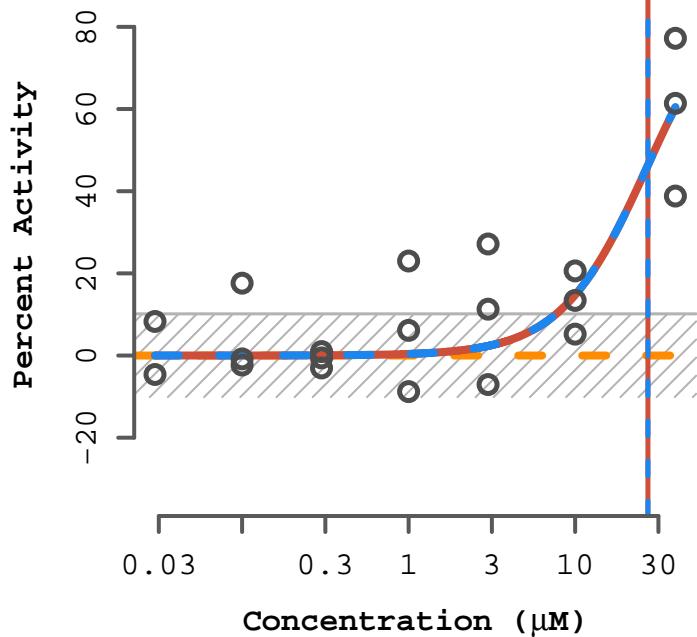
	tp	ga	gw	la	lw
val:	85.5	0.52	1.03	2.32	14.5
sd:	28	0.454	0.626	5390	110000

	CNST	HILL	GNLS
AIC:	205.54	176.98	180.98
PROB:	0	0.88	0.12
RMSE:	40.16	19	19

MAX_MEAN: 80.3 MAX_MED: 91.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Methyl (RS)-2-[4-(2,4-dichlorophenoxy)phenoxy]benzoic acid
 CHID: 32605 CASRN: 51338-27-3
 SPID(S): TP0001412G01
 M4ID: 30204293

HILL MODEL (in red):

	tp	ga	gw
val:	92.7	1.44	1.65
sd:	81.1	0.548	1.24

GAIN-LOSS MODEL (in blue):

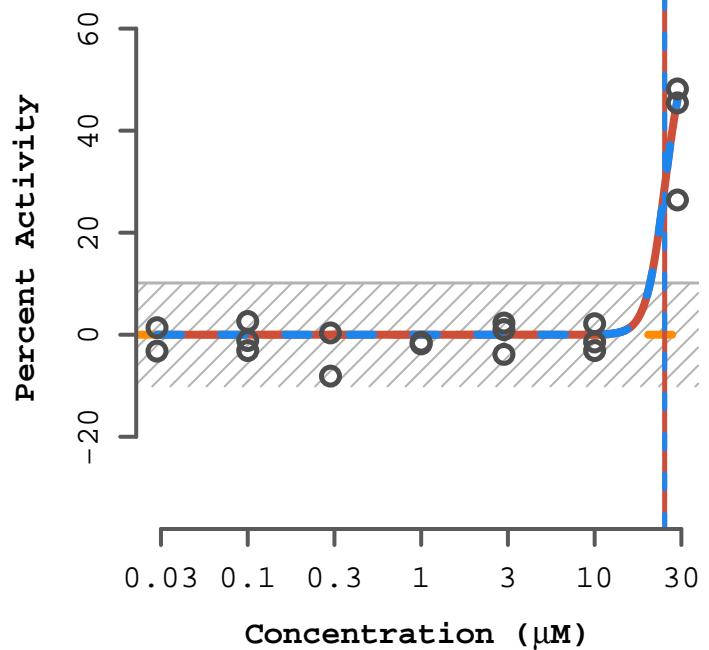
	tp	ga	gw	la	lw
val:	92.7	1.44	1.65	2.84	6.79
sd:	81	0.547	1.23	3280	18100

	CNST	HILL	GNLS
AIC:	184.65	163.38	167.38
PROB:	0	0.88	0.12
RMSE:	26.31	11.63	11.63

MAX_MEAN: 59.2 MAX_MED: 61.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Etofenprox

CHID: 32610 CASRN: 80844-07-1

SPID(S): EPAPLT0167B09

M4ID: 30204252

HILL MODEL (in red):

	tp	ga	gw
val:	57.8	1.4	8
sd:	50.8	0.079	18.1

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	57.8	1.4	8	2.81	4.71
sd:	44.2	0.0707	15.8	1590	5620

CNST

AIC: 142.8

PROB: 0

RMSE: 17.04

HILL

109.99

0.88

5.4

GNLS

113.99

0.12

5.4

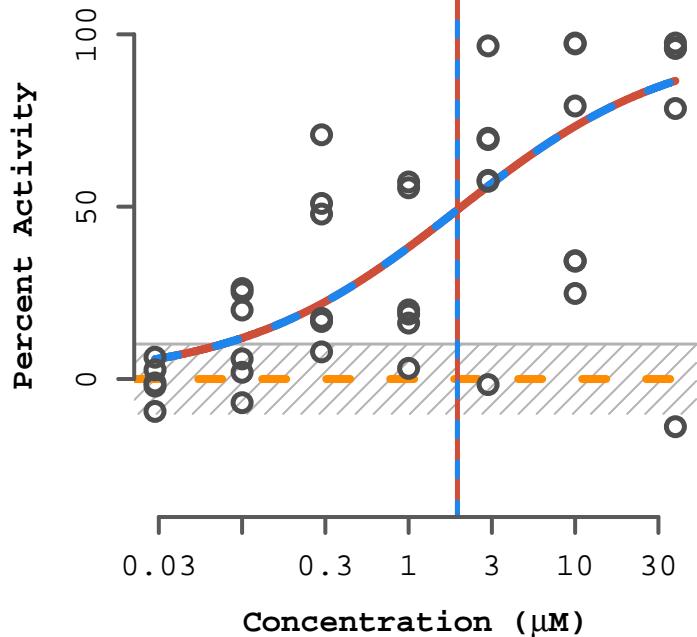
MAX_MEAN: 40

MAX_MED: 45.5

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Pyraclostrobin

CHID: 32638 CASRN: 175013-18-0

SPID(S): TP0001414B12

M4ID: 30204329

HILL MODEL (in red):

	tp	ga	gw
val:	98.1	0.293	0.666
sd:	23.1	0.443	0.242

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	98.1	0.293	0.666	2.14	13.6
sd:	23.1	0.443	0.242	301	7660

CNST

AIC: 434.17

PROB: 0

RMSE: 51.98

HILL

382.87

0.88

27.8

GNLS

386.87

0.12

27.8

MAX_MEAN: 70.8

MAX_MED: 95.9

BMAD: 3.38

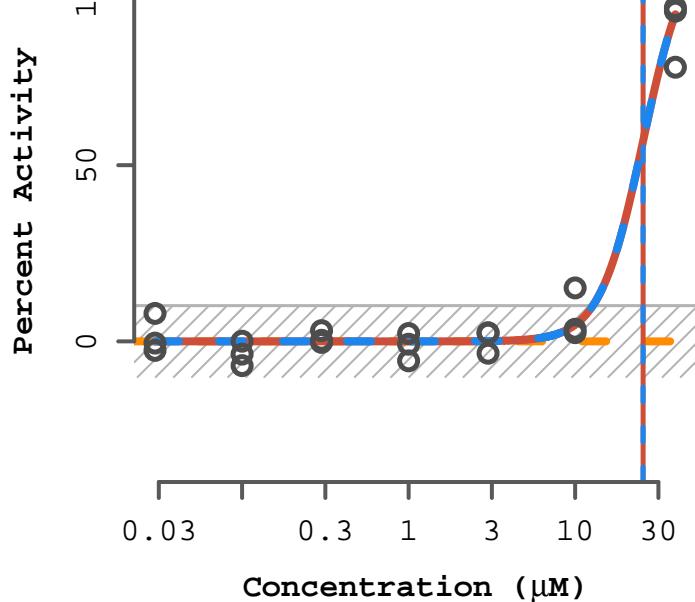
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-(Thiocyanomethylthio)benzothiazole

CHID: 32647 CASRN: 21564-17-0

SPID(S): TP0001414G10

M4ID: 30204532

HILL MODEL (in red):

	tp	ga	gw
val:	114	1.41	3.34
sd:	128	0.586	3.77

GAIN-LOSS MODEL (in blue):

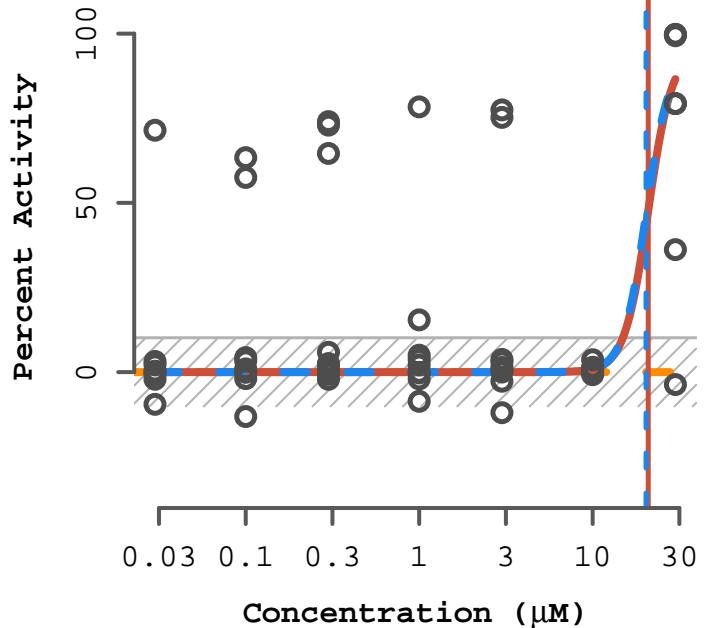
	tp	ga	gw	la	lw
val:	114	1.41	3.34	2.34	14.4
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	189.46	132.7	136.7
PROB:	0	0.88	0.12
RMSE:	33.99	5.09	5.09

MAX_MEAN: 88.8 MAX_MED: 93.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Indoxacarb

CHID: 32690 CASRN: 173584-44-6

SPID(S): EPAPLT0167E10

M4ID: 30204261

HILL MODEL (in red):

	tp	ga	gw
val:	95.4	1.32	6.32
sd:	967	5.26	90.5

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	93.3	1.31	6.55	2.03	7.76
sd:	753	4.96	94.6	279	3920

CNST

AIC: 569.95

PROB: 0

RMSE: 39.7

HILL

538.49

0.88

31.74

GNLS

542.49

0.12

31.74

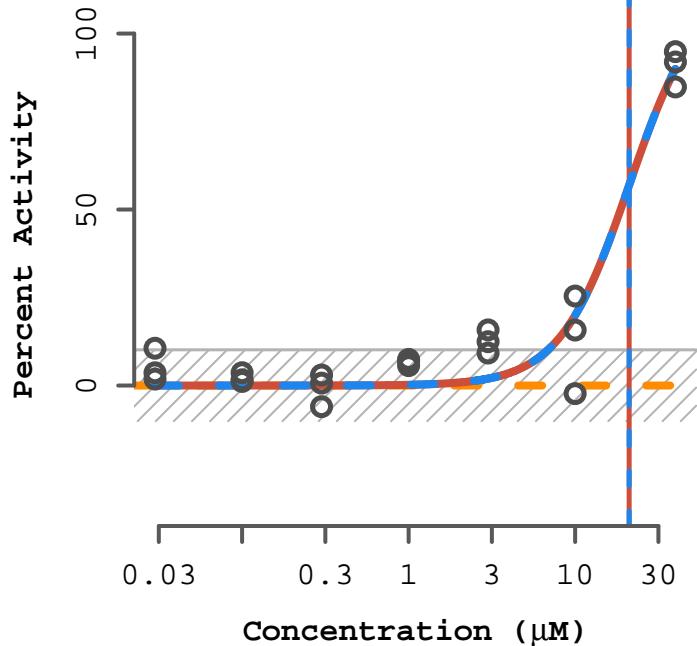
MAX_MEAN: 70

MAX_MED: 79.3

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Nonylphenol

CHID: 33836 CASRN: 104-40-5

SPID(S): TP0001413E11

M4ID: 30204562

HILL MODEL (in red):

	tp	ga	gw
val:	114	1.32	2.07
sd:	56.2	0.282	1.66

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	114	1.32	2.07	3.11	9.37
sd:	NaN	NaN	NaN	NaN	NaN

CNST

AIC: 199.78

PROB: 0

RMSE: 35.43

HILL

153.08

0.88

7.78

GNLS

157.08

0.12

7.78

MAX_MEAN: 90.5

MAX_MED: 91.9

BMAD: 3.38

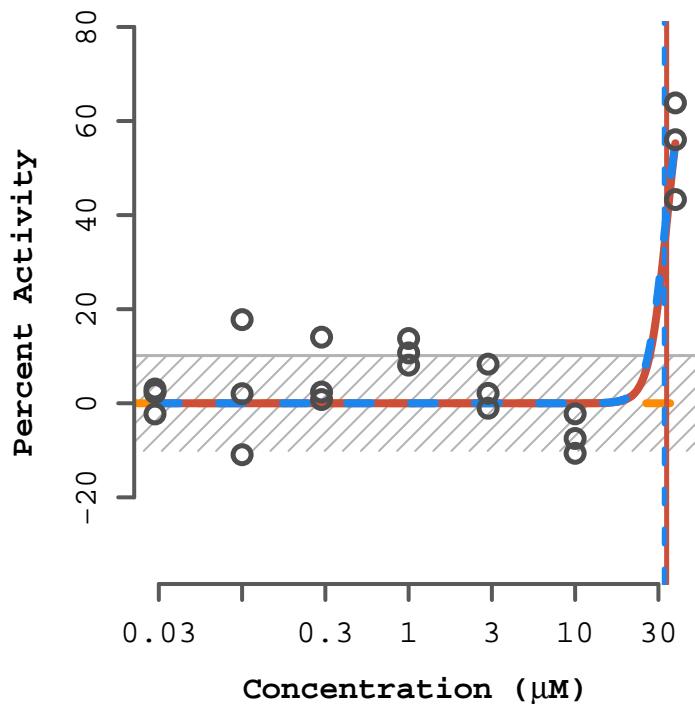
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Raloxifene hydrochloride
CHID: 34181 CASRN: 82640-04-8
SPID(S): TP0001412H01
M4ID: 30204639

HILL MODEL (in red):

	tp	ga	gw
val:	76.3	1.55	8
sd:	77.8	0.148	23.2

GAIN-LOSS MODEL (in blue):

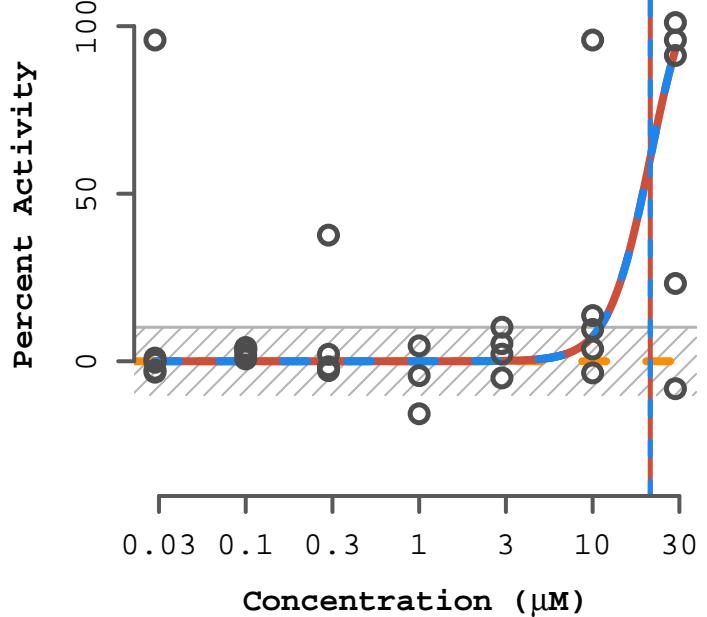
	tp	ga	gw	la	lw
val:	71.4	1.54	8	1.91	13
sd:	79.8	0.188	20.9	29.7	1170

	CNST	HILL	GNLS
AIC:	184.33	159.13	163.13
PROB:	0	0.88	0.12
RMSE:	22.23	8.45	8.45

MAX_MEAN: 54.4 MAX_MED: 56 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Boscalid

CHID: 34392 CASRN: 188425-85-6

SPID(S): EPAPLT0167D05

M4ID: 30204241

HILL MODEL (in red):

	tp	ga	gw
val:	121	1.33	3.57
sd:	493	1.63	12.1

GAIN-LOSS MODEL (in blue):

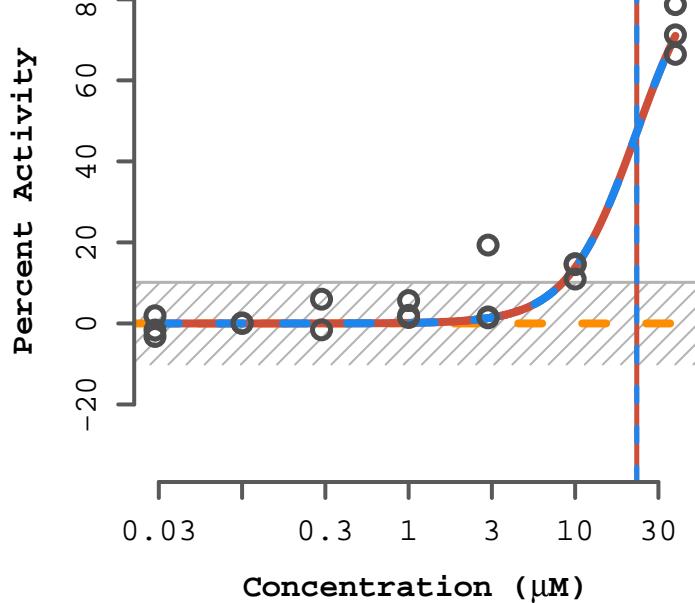
	tp	ga	gw	la	lw
val:	121	1.33	3.57	2.73	5.03
sd:	375	1.24	9.25	1240	5000

	CNST	HILL	GNLS
AIC:	310.25	294.4	298.4
PROB:	0	0.88	0.12
RMSE:	39.12	32.75	32.75

MAX_MEAN: 60.6 MAX_MED: 91.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Buprofezin

CHID: 34401 CASRN: 69327-76-0

SPID(S): TP0001412H02

M4ID: 30204312

HILL MODEL (in red):

	tp	ga	gw
val:	94.4	1.37	2.07
sd:	40.1	0.26	0.89

GAIN-LOSS MODEL (in blue):

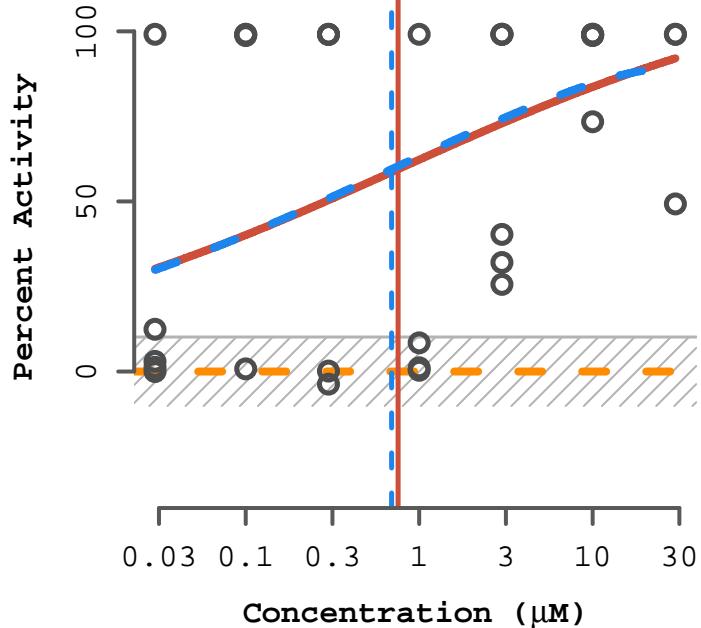
	tp	ga	gw	la	lw
val:	94.4	1.37	2.07	3.02	5.51
sd:	40	0.259	0.888	1280	4990

	CNST	HILL	GNLS
AIC:	175.62	116.56	120.56
PROB:	0	0.88	0.12
RMSE:	29.65	5.16	5.16

MAX_MEAN: 72.1 MAX_MED: 71.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Emamectin benzoate

CHID: 34566 CASRN: 155569-91-8

SPID(S): EPAPLT0167H10

M4ID: 30204250

HILL MODEL (in red):

tp	ga	gw
val: 119	-0.121	0.334
sd: 96.1	2.5	0.326

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 119	-0.159	0.348	2.13	1.92
sd: NaN	NaN	NaN	NaN	NaN

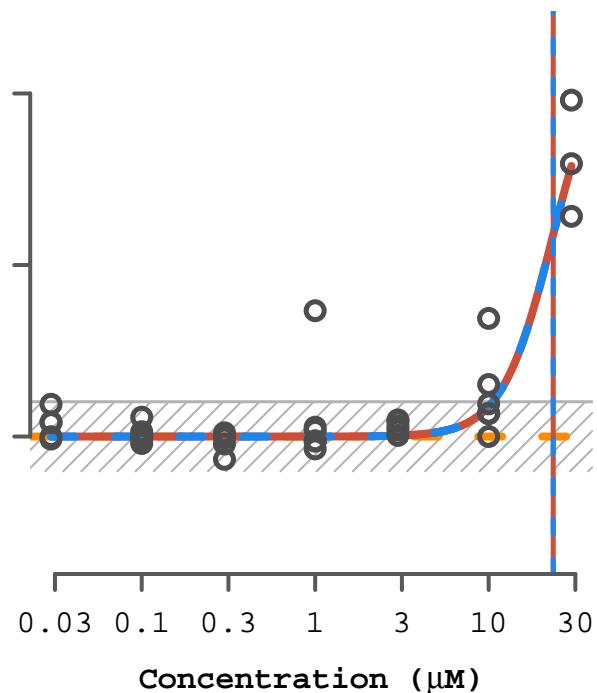
CNST	HILL	GNLS
AIC: 720.19	351.48	355.42
PROB: 0	0.88	0.12
RMSE: 75.5	40.51	40.53

MAX_MEAN: 94.7 MAX_MED: 99.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fenamidone

CHID: 34590 CASRN: 161326-34-7

SPID(S): EPAPLT0167E02

M4ID: 30204254

HILL MODEL (in red):

tp	ga	gw
val: 118	1.37	2.94
sd: 56.6	0.185	0.956

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 118	1.37	2.94	3.46	5.6
sd: 56.8	0.186	0.96	81800	226000

CNST

AIC: 300.1

PROB: 0

RMSE: 25.45

HILL

247.76

0.88

9.13

GNLS

251.76

0.12

9.13

MAX_MEAN: 80.6

MAX_MED: 79.6

BMAD: 3.38

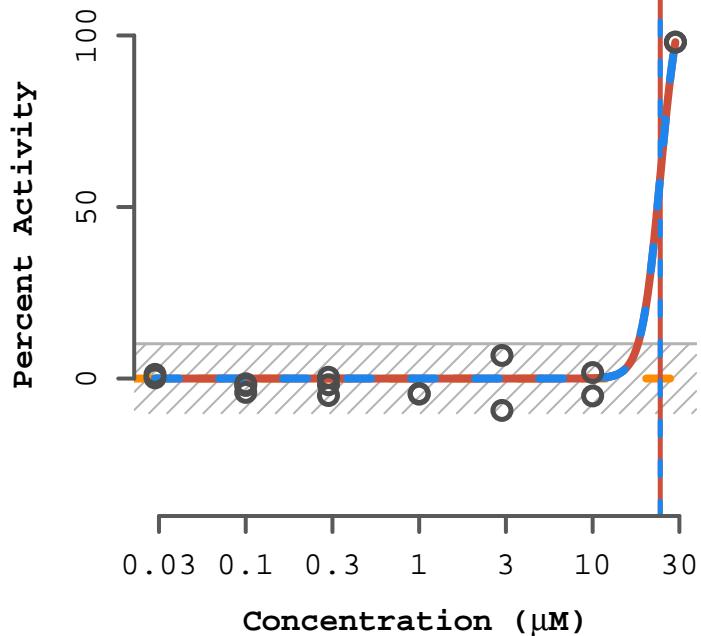
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fipronil

CHID: 34609 CASRN: 120068-37-3

SPID(S): EPAPLT0167B08

M4ID: 30204255

HILL MODEL (in red):

	tp	ga	gw
val:	118	1.39	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

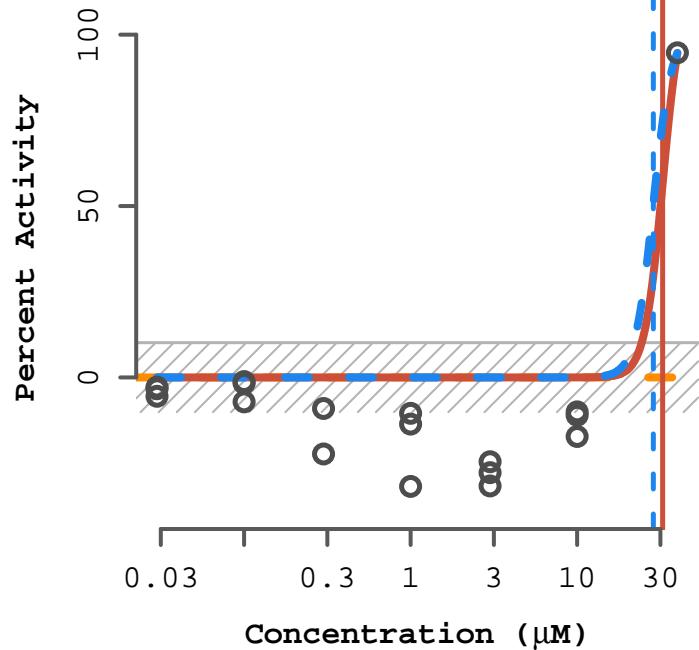
	tp	ga	gw	la	lw
val:	118	1.39	8	3.47	9.42
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	140.49	98.09	102.09
PROB:	0	0.88	0.12
RMSE:	34.9	3.97	3.97

MAX_MEAN: 98.1 MAX_MED: 98.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fipronil

CHID: 34609 CASRN: 120068-37-3

SPID(S): TP0001412H03

M4ID: 30204359

HILL MODEL (in red):

	tp	ga	gw
val:	113	1.51	8
sd:	338	0.928	18.9

GAIN-LOSS MODEL (in blue):

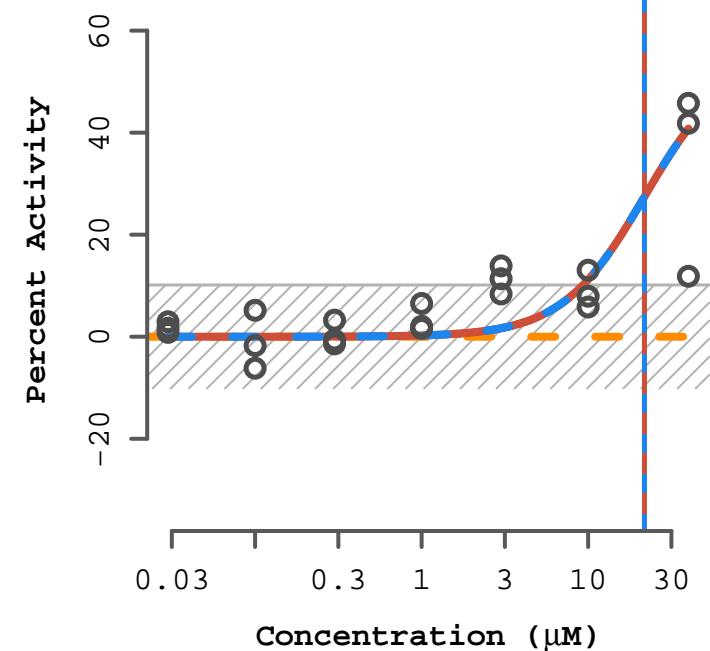
	tp	ga	gw	la	lw
val:	101	1.46	8	3.45	2.36
sd:	100	0.662	14.7	240	307

	CNST	HILL	GNLS
AIC:	166.91	161.09	165.1
PROB:	0.05	0.84	0.11
RMSE:	27.75	16.48	16.48

MAX_MEAN: 94.7 MAX_MED: 94.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.95

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Fluoxastrobin

CHID: 34625 CASRN: 361377-29-9

SPID(S): TP0001411F10

M4ID: 30204290

HILL MODEL (in red):

	tp	ga	gw
val:	54.9	1.34	1.75
sd:	24.6	0.275	1.29

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	54.9	1.34	1.75	3.28	6.72
sd:	24.5	0.274	1.29	42800	171000

CNST

AIC: 167.06

PROB: 0

RMSE: 15.1

HILL

146.83

0.88

7.97

GNLS

150.83

0.12

7.97

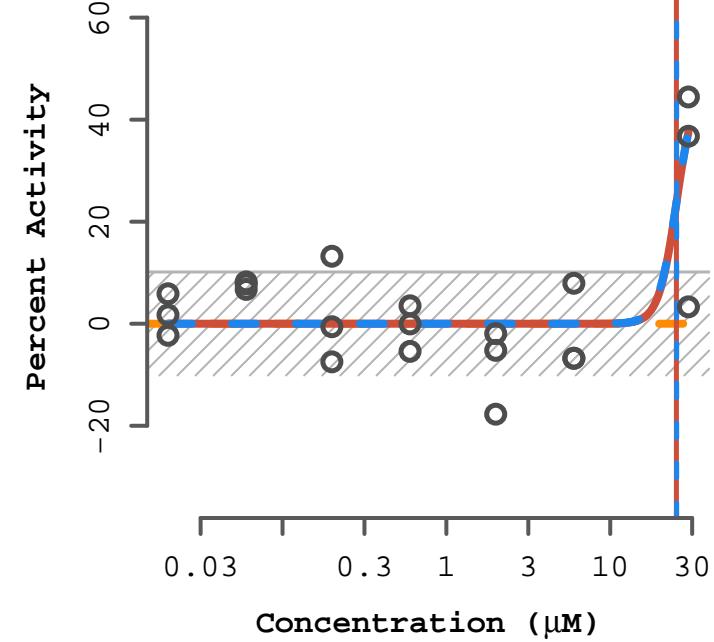
MAX_MEAN: 33.1

MAX_MED: 41.8

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4-Hydroxytamoxifen

CHID: 37094 CASRN: 68392-35-8

SPID(S): TP0001411H06

M4ID: 30204623

HILL MODEL (in red):

	tp	ga	gw
val:	47.3	1.4	7.94
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	48.2	1.41	7.67	3.46	1.3
sd:	NaN	NaN	NaN	NaN	NaN

CNST

AIC: 166.26

PROB: 0.06

RMSE: 14.35

HILL

AIC: 161.1

PROB: 0.83

RMSE: 10.24

GNLS

AIC: 165.1

PROB: 0.11

RMSE: 10.24

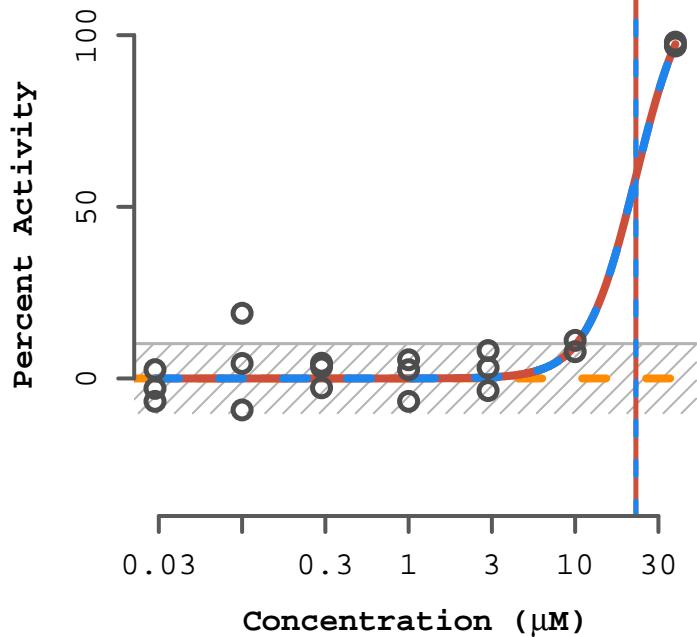
MAX_MEAN: 28.2

MAX_MED: 36.8

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.94

FLAGS: 17; 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4,4',4''-Ethane-1,1,1-triyltriphenol
CHID: 37712 CASRN: 27955-94-8
SPID(S): TP0001411H12
M4ID: 30204468

HILL MODEL (in red):

	tp	ga	gw
val:	118	1.36	2.88
sd:	127	0.643	3.76

GAIN-LOSS MODEL (in blue):

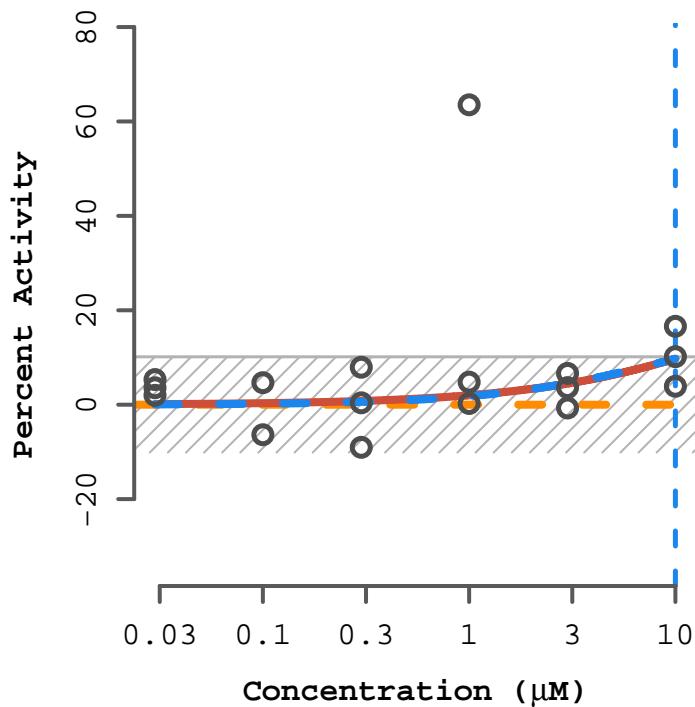
	tp	ga	gw	la	lw
val:	118	1.36	2.88	3.37	13.3
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	168.42	129.39	133.39
PROB:	0	0.88	0.12
RMSE:	32.36	6.21	6.21

MAX_MEAN: 97.4 MAX_MED: 97.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Bisphenol AF

CHID: 37717 CASRN: 1478-61-1

SPID(S): TP0001412A01

M4ID: 30204455

HILL MODEL (in red):

	tp	ga	gw
val:	34.5	1.5	0.808
sd:	136	3.36	0.995

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	19.4	1	0.988	2.99	9.08
sd:	NA	NA	NA	NA	NA

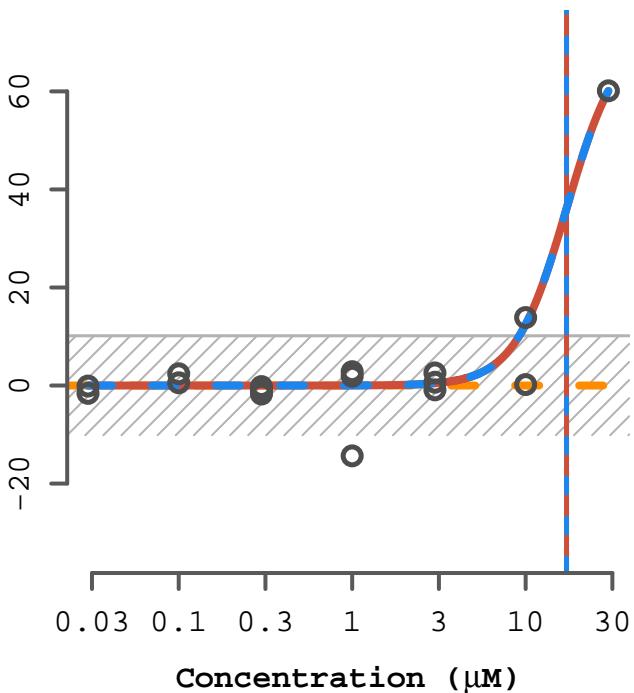
	CNST	HILL	GNLS
AIC:	131.36	129.8	133.9
PROB:	0.29	0.63	0.08
RMSE:	16.74	15.65	15.7

MAX_MEAN: 22.9 MAX_MED: 10.2 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.71

FLAGS: 17; 16; 11; 10; 6

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 4,4'-(9H-Fluorene-9,9-diyl)diphenol

CHID: 37731 CASRN: 3236-71-3

SPID(S): EPAPLT0154H10

M4ID: 30204233

HILL MODEL (in red):

	tp	ga	gw
val:	72.2	1.24	2.89
sd:	89	0.661	5.38

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	72.2	1.24	2.89	1.96	17.9
sd:	87.8	0.652	5.31	1540	57000

CNST

AIC: 114.37

PROB: 0

RMSE: 15.9

HILL

95.22

0.88

4.93

GNLS

99.22

0.12

4.93

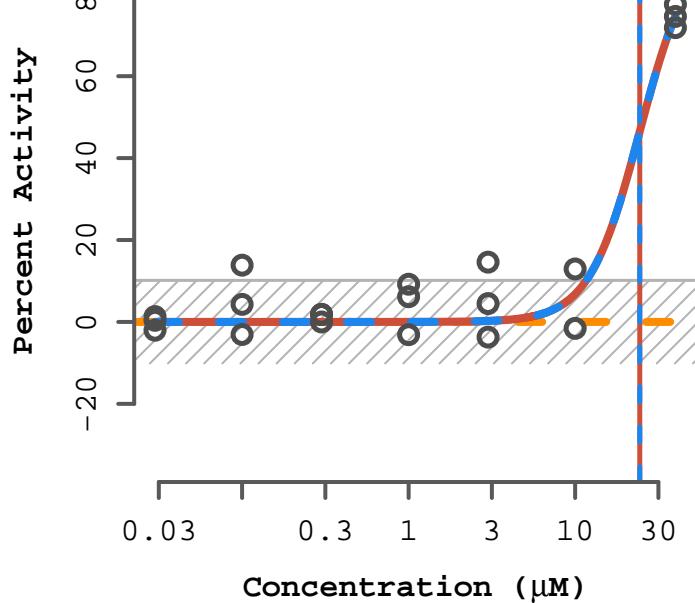
MAX_MEAN: 60.1

MAX_MED: 60.1

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: S-Bioallethrin

CHID: 39336 CASRN: 28434-00-6

SPID(S): TP0001414C11

M4ID: 30204383

HILL MODEL (in red):

	tp	ga	gw
val:	93	1.39	2.84
sd:	164	0.92	6

GAIN-LOSS MODEL (in blue):

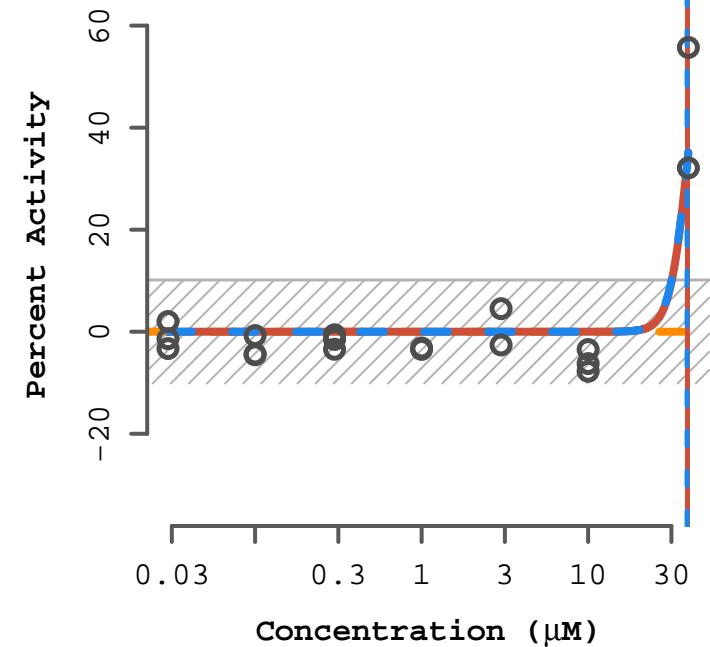
	tp	ga	gw	la	lw
val:	93	1.39	2.84	3.48	4.46
sd:	165	0.926	6.03	8450	20100

	CNST	HILL	GNLS
AIC:	182.1	136.06	140.06
PROB:	0	0.88	0.12
RMSE:	29.6	6.04	6.04

MAX_MEAN: 74.7 MAX_MED: 74.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2-tert-Butyl-4-methoxyphenol
CHID: 40788 CASRN: 121-00-6
SPID(S): TP0001414G04
M4ID: 30204407

HILL MODEL (in red):

	tp	ga	gw
val:	66.8	1.6	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

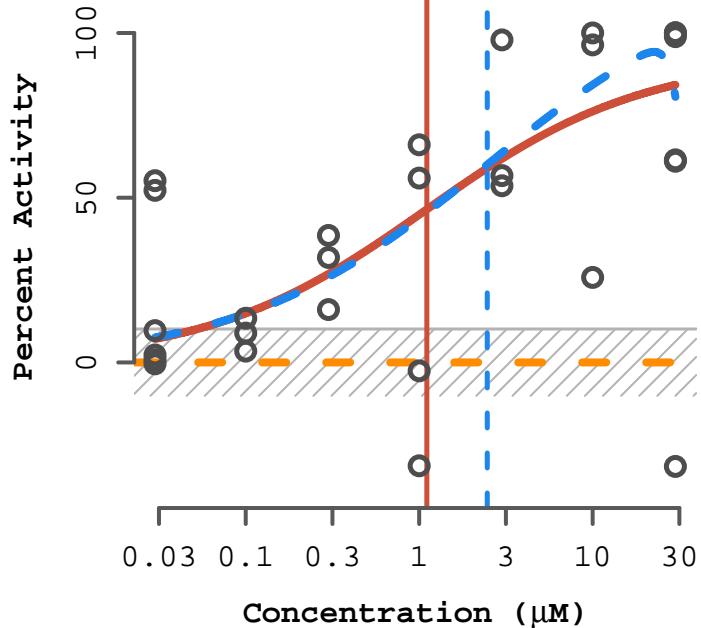
	tp	ga	gw	la	lw
val:	65.7	1.59	7.91	3.28	2.19
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	134.89	119.07	123.07
PROB:	0	0.88	0.12
RMSE:	15.58	6.07	6.07

MAX_MEAN: 43.9 MAX_MED: 43.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: alpha-Cypermethrin
 CHID: 41201 CASRN: 67375-30-8
 SPID(S): EPAPLT0167F02
 M4ID: 30204276

HILL MODEL (in red):

	tp	ga	gw
val:	92.9	0.0442	0.689
sd:	23.6	0.497	0.368

GAIN-LOSS MODEL (in blue):

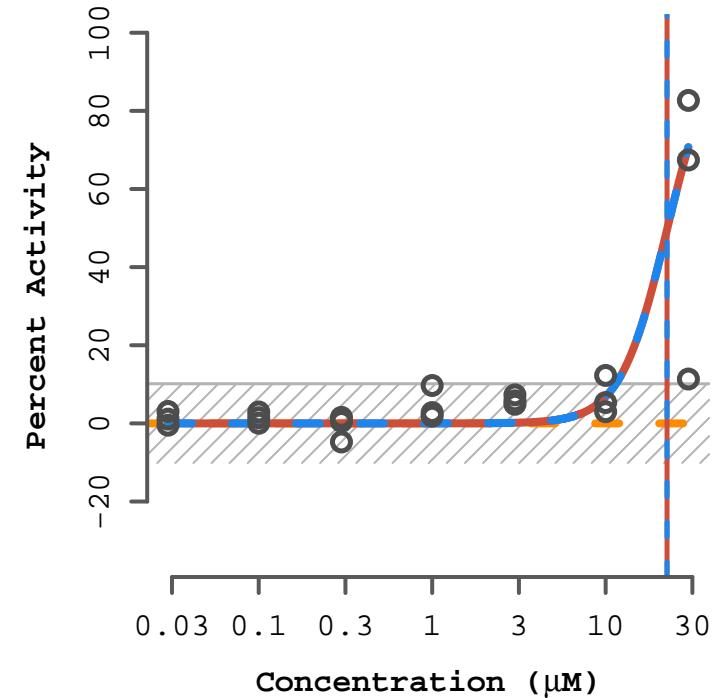
	tp	ga	gw	la	lw
val:	120	0.393	0.612	1.54	10.1
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	310.77	280.78	284.21
PROB:	0	0.85	0.15
RMSE:	57.12	34.66	34.26

MAX_MEAN: 74.1 MAX_MED: 96.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: N-Dodecanoyl-N-methylglycine
CHID: 42011 CASRN: 97-78-9
SPID(S): EPAPLT0154F09
M4ID: 30204212

HILL MODEL (in red):

	tp	ga	gw
val:	99.2	1.35	3.2
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

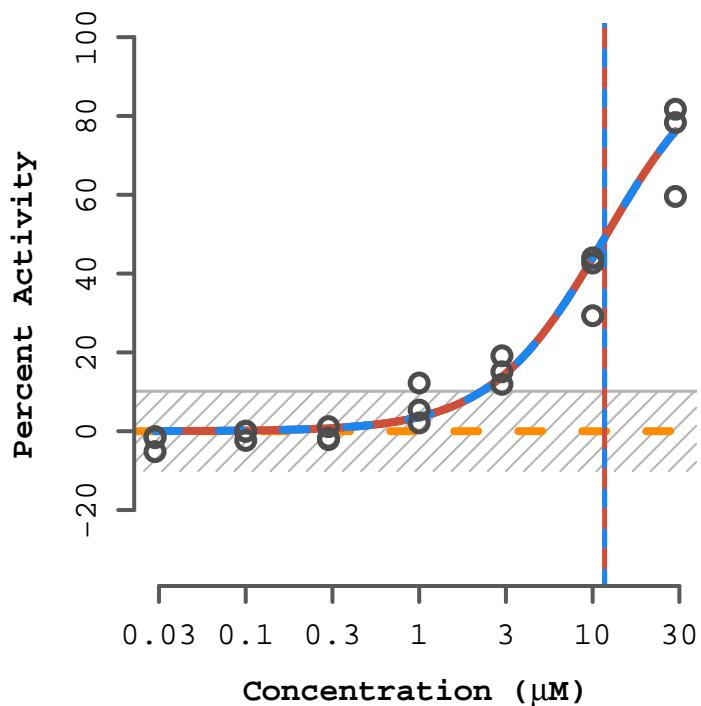
	tp	ga	gw	la	lw
val:	99.2	1.35	3.2	3.27	8.15
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	170.95	152.09	156.09
PROB:	0	0.88	0.12
RMSE:	23.87	13.77	13.77

MAX_MEAN: 53.8 MAX_MED: 67.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tetrabutylammonium bromide
 CHID: 44400 CASRN: 1643-19-2
 SPID(S): EPAPLT0154A01
 M4ID: 30204153

HILL MODEL (in red):

tp	ga	gw
val: 98.1	1.07	1.33
sd: 12	0.0915	0.16

GAIN-LOSS MODEL (in blue):

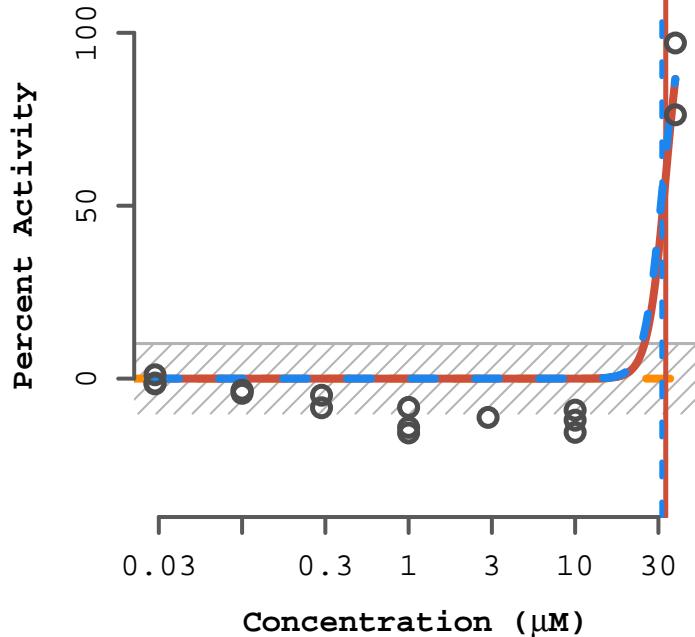
tp	ga	gw	la	lw
val: 98.1	1.07	1.33	2.16	18
sd: 12	0.0915	0.16	11200	295000

CNST	HILL	GNLS
AIC: 203.69	135.34	139.34
PROB: 0	0.88	0.12
RMSE: 32.33	5.74	5.74

MAX_MEAN: 73.2 MAX_MED: 78.3 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Enadoline

CHID: 47258 CASRN: 124378-77-4

SPID(S): TP0001413C11

M4ID: 30204494

HILL MODEL (in red):

	tp	ga	gw
val:	116	1.54	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

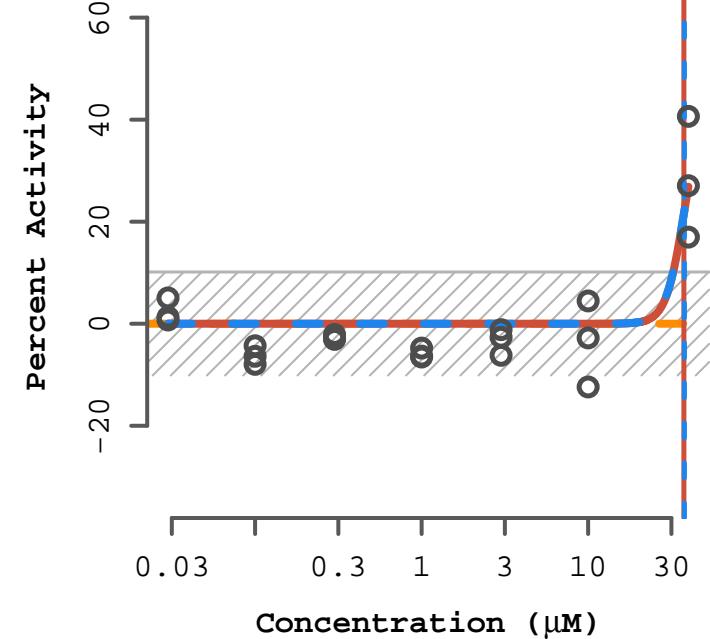
	tp	ga	gw	la	lw
val:	106	1.52	8	3.58	18
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	155.64	134.44	138.45
PROB:	0	0.88	0.12
RMSE:	31.15	9.28	9.28

MAX_MEAN: 86.7 MAX_MED: 86.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: PD 0343701

CHID: 47271 CASRN: 676116-04-4

SPID(S): TP0001412D11

M4ID: 30204498

HILL MODEL (in red):

	tp	ga	gw
val:	43.2	1.57	8
sd:	80.1	0.134	55.8

GAIN-LOSS MODEL (in blue):

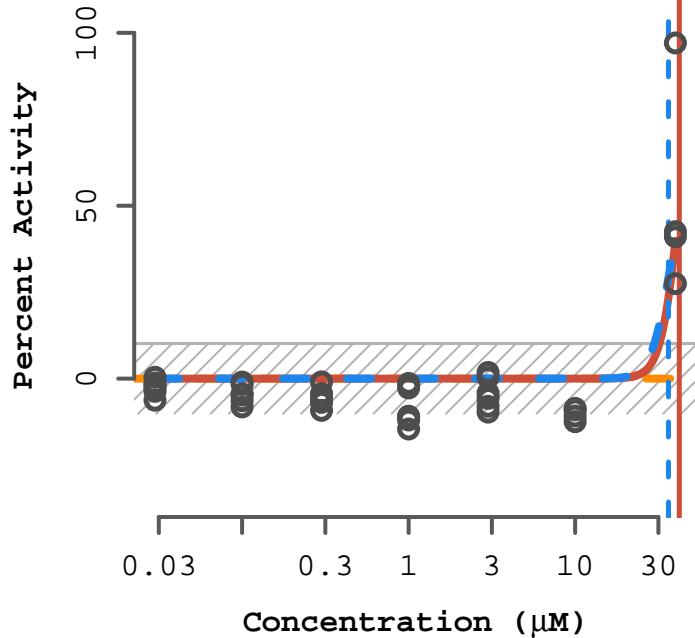
	tp	ga	gw	la	lw
val:	44.3	1.58	7.79	2.82	6.64
sd:	68.1	0.127	48	110000	613000

	CNST	HILL	GNLS
AIC:	159.59	145.26	149.26
PROB:	0	0.88	0.12
RMSE:	12.3	6.15	6.15

MAX_MEAN: 28.2 MAX_MED: 27.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Elzasonan

CHID: 47277 CASRN: 361343-19-3

SPID(S): TP0001414B11

M4ID: 30204499

HILL MODEL (in red):

	tp	ga	gw
val:	101	1.63	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	58.5	1.56	7.96	3.44	2.11
sd:	59.7	0.113	20.5	536	620

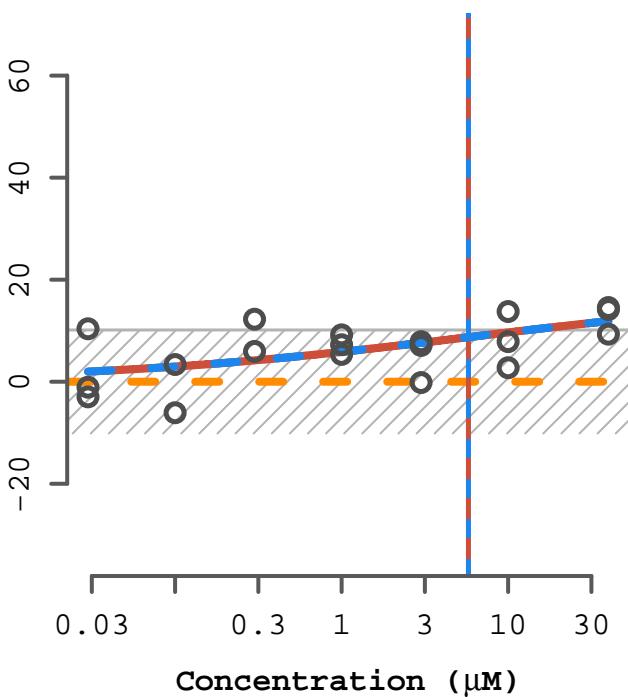
	CNST	HILL	GNLS
AIC:	307.68	282.59	286.6
PROB:	0	0.88	0.12
RMSE:	20.1	11.61	11.61

MAX_MEAN: 52 MAX_MED: 41.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 10; 7

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: CP-532623

CHID: 47279 CASRN: 261947-38-0

SPID(S): TP0001411D04

M4ID: 30204500

HILL MODEL (in red):

	tp	ga	gw
val:	17.5	0.762	0.391
sd:	17.7	2.48	0.328

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	17.5	0.762	0.391	3.59	3.77
sd:	17.7	2.48	0.328	1300	2480

CNST

AIC: 140.32

PROB: 0

RMSE: 8.55

HILL

121.75

0.88

4.68

GNLS

125.75

0.12

4.68

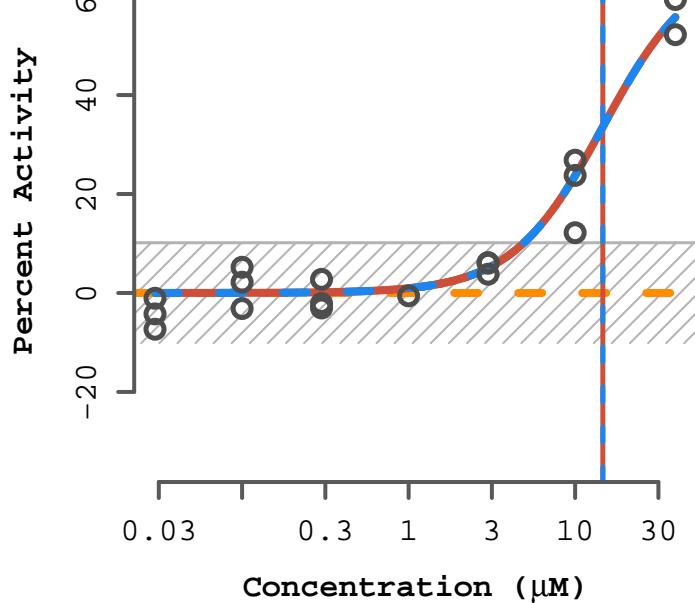
MAX_MEAN: 12.7

MAX_MED: 14.2

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: UK-333747

CHID: 47287 CASRN: 197077-55-7

SPID(S): TP0001411G04

M4ID: 30204504

HILL MODEL (in red):

	tp	ga	gw
val:	67.2	1.17	1.58
sd:	10.9	0.132	0.363

GAIN-LOSS MODEL (in blue):

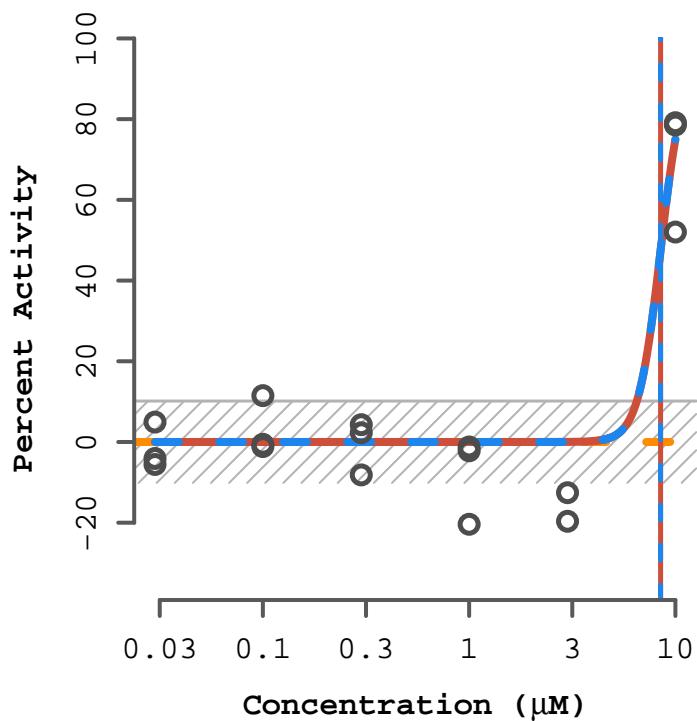
	tp	ga	gw	la	lw
val:	67.2	1.17	1.58	2.6	6.57
sd:	10.9	0.132	0.363	925	6120

	CNST	HILL	GNLS
AIC:	155.66	109.64	113.64
PROB:	0	0.88	0.12
RMSE:	20.95	4.15	4.15

MAX_MEAN: 55.8 MAX_MED: 55.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: UK-337312

CHID: 47288 CASRN: 203942-49-8

SPID(S): TP0001414D10

M4ID: 30204659

HILL MODEL (in red):

tp	ga	gw
val: 94.9	0.928	8
sd: 149	0.344	12.1

GAIN-LOSS MODEL (in blue):

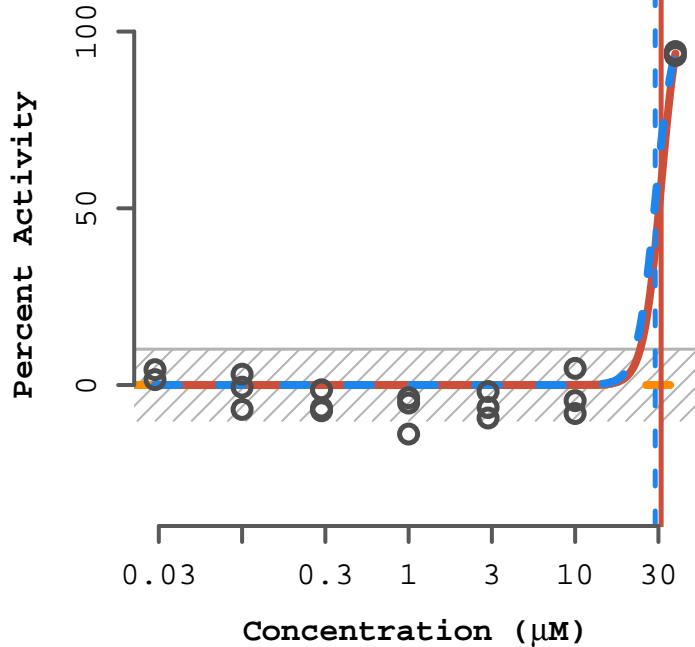
tp	ga	gw	la	lw
val: 94.9	0.928	8	1.88	17.8
sd: NA	NA	NA	NA	NA

CNST	HILL	GNLS
AIC: 161.08	135.2	139.2
PROB: 0	0.88	0.12
RMSE: 31.06	10.35	10.35

MAX_MEAN: 69.9 MAX_MED: 78.6 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 6



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: UK-416244

CHID: 47290 CASRN: 402910-27-4

SPID(S): TP0001414C09

M4ID: 30204505

HILL MODEL (in red):

	tp	ga	gw
val:	113	1.52	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	104	1.48	8	3.6	13.2
sd:	NA	NA	NA	NA	NA

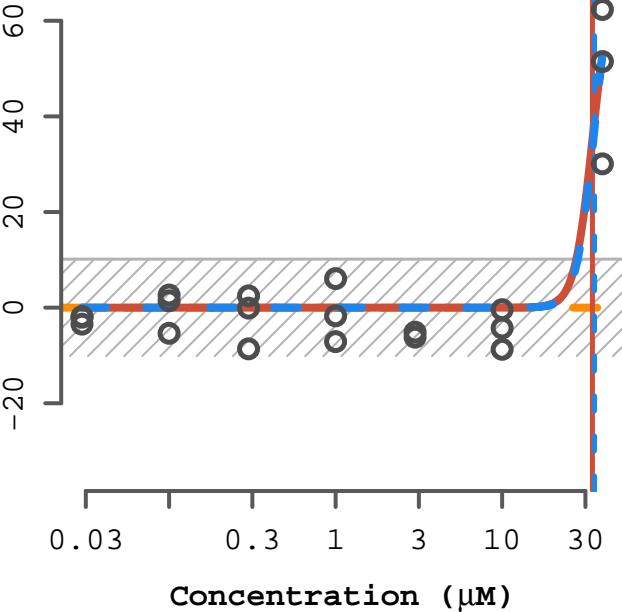
	CNST	HILL	GNLS
AIC:	164.82	130.33	134.33
PROB:	0	0.88	0.12
RMSE:	30.98	5.82	5.82

MAX_MEAN: 93.8 MAX_MED: 93.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: UK-373911

CHID: 47295 CASRN: 291305-06-1

SPID(S): TP0001411H08

M4ID: 30204507

HILL MODEL (in red):

tp	ga	gw
val: 70.3	1.54	8
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

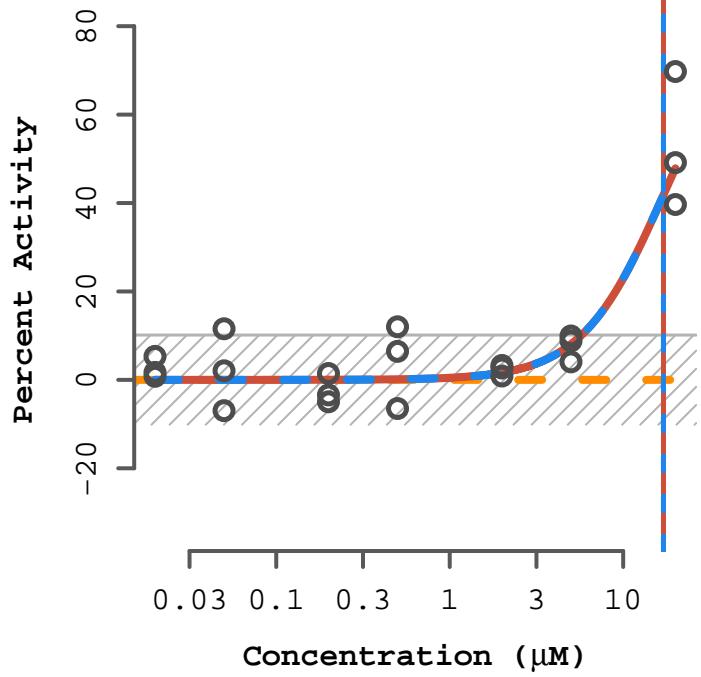
tp	ga	gw	la	lw
val: 74.8	1.55	8	2.78	5.3
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 164.94	139.94	143.94
PROB: 0	0.88	0.12
RMSE: 19.79	7.06	7.06

MAX_MEAN: 48 MAX_MED: 51.5 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: CJ-013790

CHID: 47303 CASRN: 179465-71-5

SPID(S): TP0001412D06

M4ID: 30204508

HILL MODEL (in red):

	tp	ga	gw
val:	83.7	1.23	1.82
sd:	106	0.659	1.13

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	83.7	1.23	1.82	3.08	3.99
sd:	108	0.675	1.15	1940	4360

CNST

AIC: 178.36

PROB: 0

RMSE: 21.33

HILL

146.88

0.88

7.09

GNLS

150.88

0.12

7.09

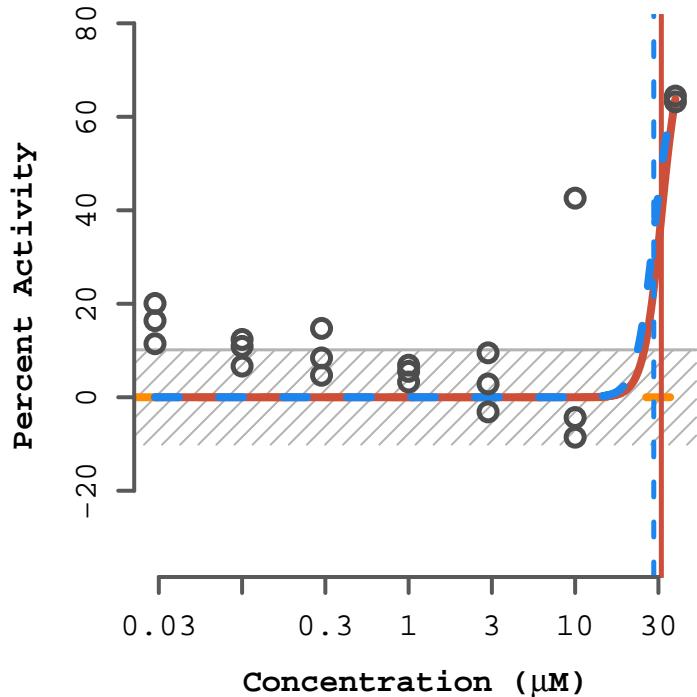
MAX_MEAN: 52.9

MAX_MED: 49.2

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: PharmaGSID_47330
CHID: 47330 CASRN: NOCAS_47330
SPID(S): TP0001413D01
M4ID: 30204514

HILL MODEL (in red):

	tp	ga	gw
val:	77.3	1.52	7.99
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

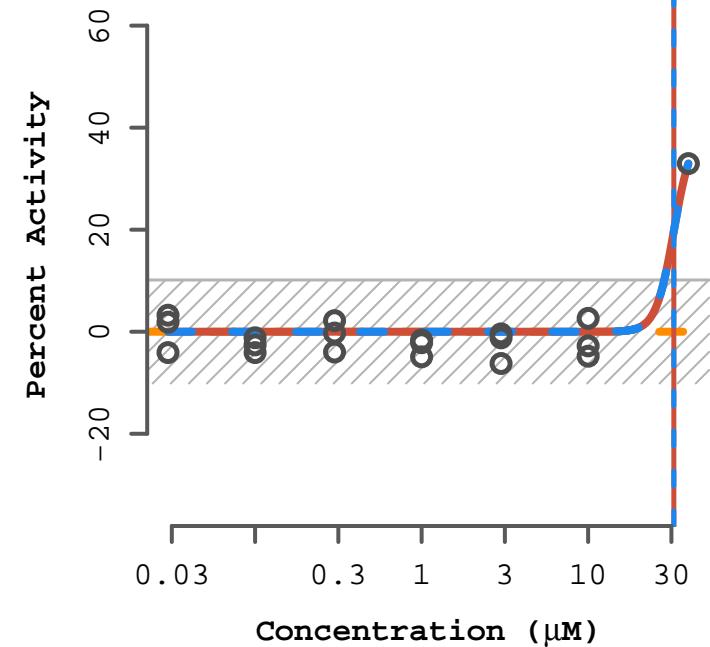
	tp	ga	gw	la	lw
val:	69.7	1.47	8	1.79	18
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	180.77	165.39	169.39
PROB:	0	0.88	0.12
RMSE:	24.17	13.29	13.29

MAX_MEAN: 63.8 MAX_MED: 63.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: HMR1426

CHID: 47338 CASRN: 262376-75-0

SPID(S): TP0001411G05

M4ID: 30204515

HILL MODEL (in red):

	tp	ga	gw
val:	39.6	1.51	8
sd:	56.8	0.221	27.6

GAIN-LOSS MODEL (in blue):

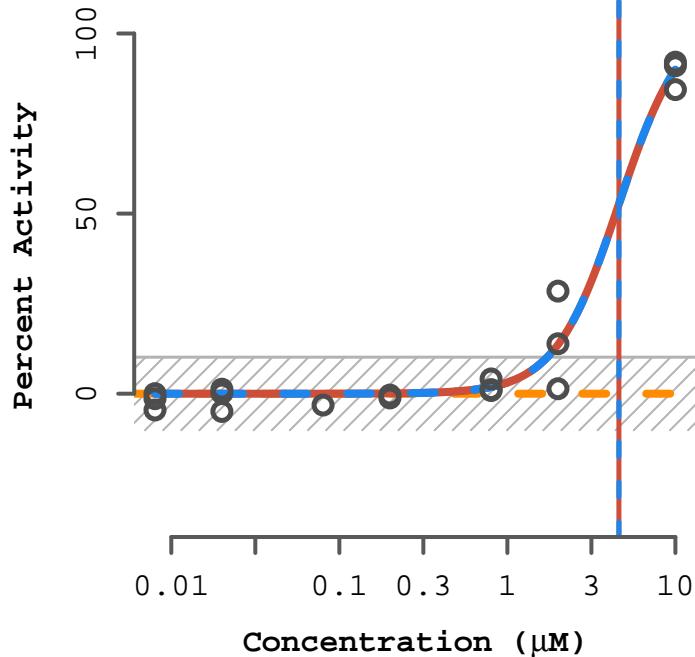
	tp	ga	gw	la	lw
val:	39.6	1.51	8	3.6	11.8
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	118.56	107.61	111.61
PROB:	0	0.88	0.12
RMSE:	8.18	3.12	3.12

MAX_MEAN: 33 MAX_MED: 33 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SSR146977

CHID: 47341 CASRN: 264618-44-2

SPID(S): TP0001414D08

M4ID: 30204516

HILL MODEL (in red):

	tp	ga	gw
val:	105	0.664	2.28
sd:	32.5	0.251	1.04

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	105	0.664	2.28	3	2.86
sd:	32.6	0.251	1.04	4190	6210

CNST

AIC: 173.61

PROB: 0

RMSE: 37.25

HILL

112.31

0.88

5.13

GNLS

116.31

0.12

5.13

MAX_MEAN: 89.2

MAX_MED: 91.1

BMAD: 3.38

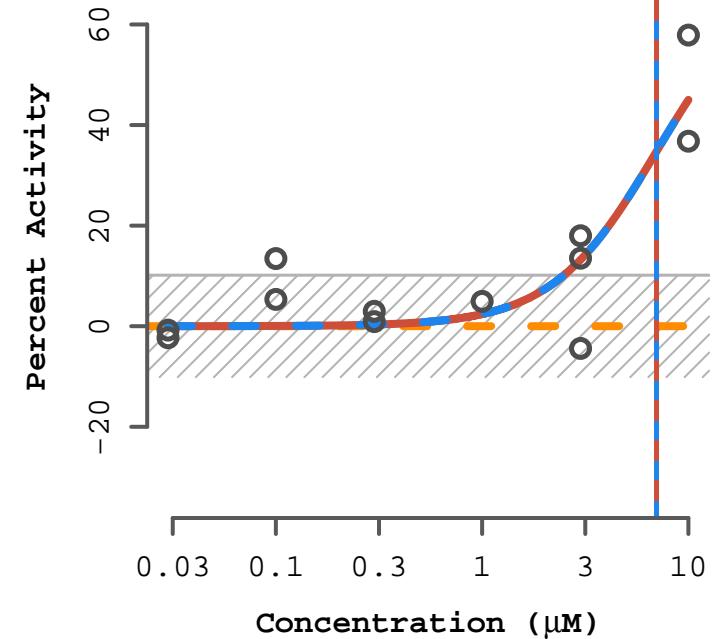
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SSR504734

CHID: 47348 CASRN: 742693-38-5

SPID(S): TP0001413H01

M4ID: 30204518

HILL MODEL (in red):

	tp	ga	gw
val:	69.5	0.845	1.71
sd:	68.7	0.601	1.61

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	69.5	0.845	1.71	2.1	8.73
sd:	68.7	0.6	1.61	11600	92000

CNST

AIC: 106.5

PROB: 0

RMSE: 21.37

HILL

92.37

0.88

8.14

GNLS

96.37

0.12

8.14

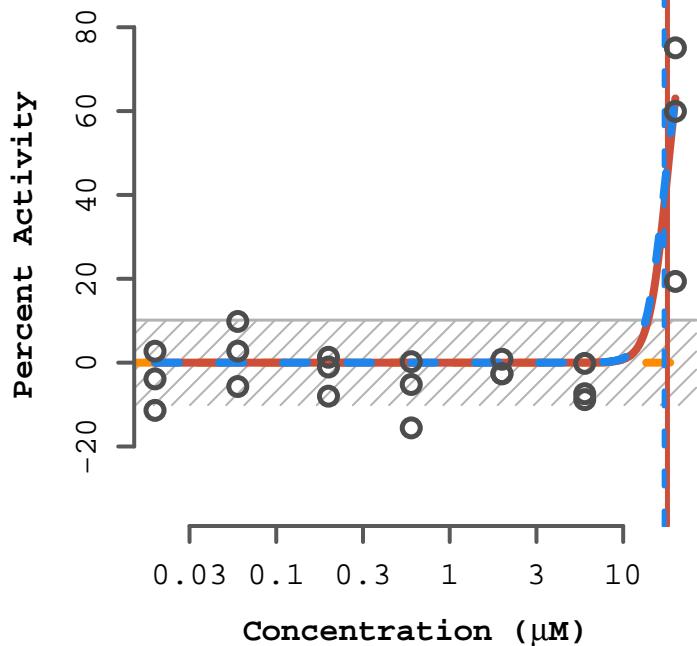
MAX_MEAN: 47.3

MAX_MED: 47.3

BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 17



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SSR 241586 HCl

CHID: 47353 CASRN: NOCAS_47353

SPID(S): TP0001411F01

M4ID: 30204644

HILL MODEL (in red):

tp	ga	gw
val: 90.1	1.26	8
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

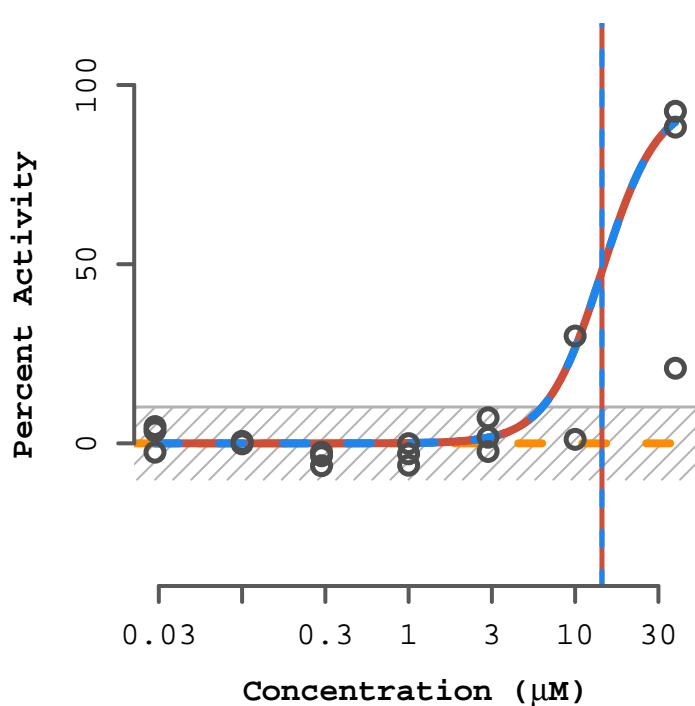
tp	ga	gw	la	lw
val: 82.1	1.24	8	3.3	3.24
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 176.48	161.17	165.17
PROB: 0	0.88	0.12
RMSE: 22.23	11.61	11.61

MAX_MEAN: 51.5 MAX_MED: 59.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: AVE5638

CHID: 47371 CASRN: 725228-45-5

SPID(S): TP0001413B11

M4ID: 30204524

HILL MODEL (in red):

	tp	ga	gw
val:	96.2	1.16	2.59
sd:	15.2	0.143	1.44

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	96.2	1.16	2.59	2.63	5.8
sd:	15.2	0.143	1.44	1480	8340

CNST

AIC: 169.36

PROB: 0

RMSE: 30.71

HILL

145.33

0.88

17.16

GNLS

149.33

0.12

17.16

MAX_MEAN: 67.3

MAX_MED: 88.3

BMAD: 3.38

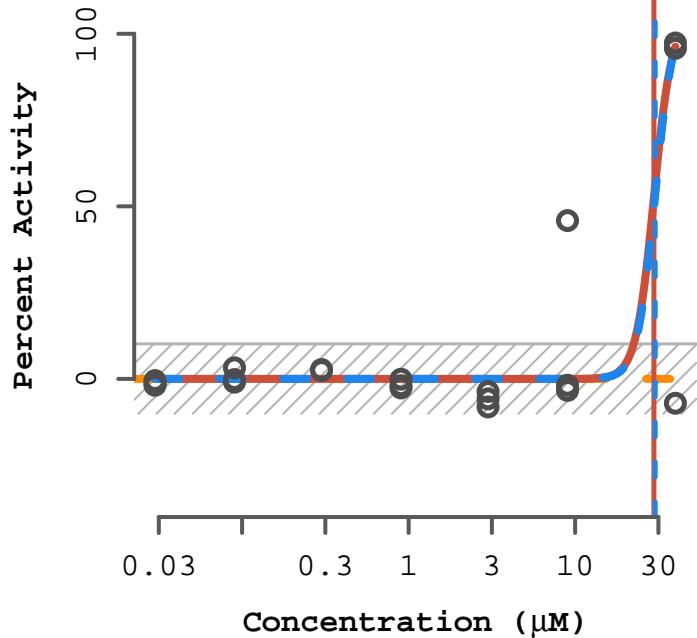
COFF: 10.1

HIT-CALL: 1

FITC: 42

ACTP: 1

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: SAR 150640

CHID: 47389 CASRN: 433212-21-6

SPID(S): TP0001414F12

M4ID: 30204526

HILL MODEL (in red):

tp	ga	gw
val: 105	1.47	8
sd: 73.2	0.243	15.9

GAIN-LOSS MODEL (in blue):

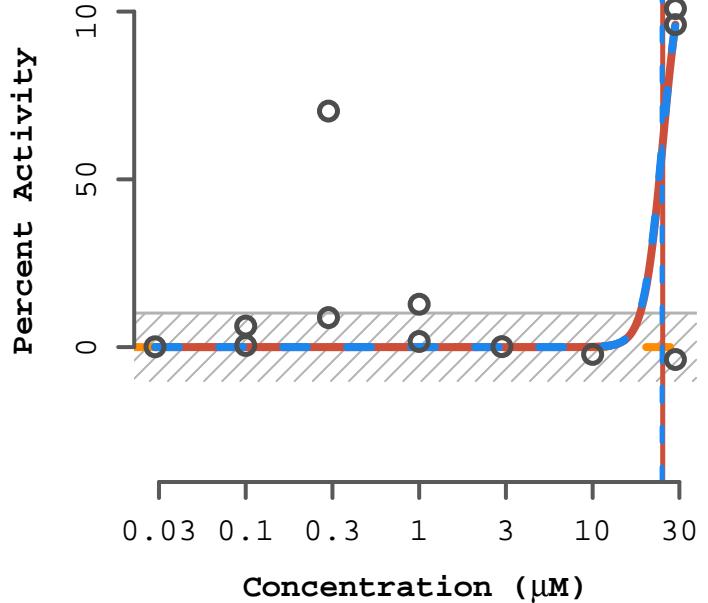
tp	ga	gw	la	lw
val: 107	1.48	8	3.57	2.39
sd: NaN	NaN	NaN	NaN	NaN

CNST	HILL	GNLS
AIC: 175.2	156.62	160.62
PROB: 0	0.88	0.12
RMSE: 32.38	25.45	25.45

MAX_MEAN: 62 MAX_MED: 95.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 10; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 2,2,2-Trichloro-1-phenylethyl acetate

CHID: 47725 CASRN: 90-17-5

SPID(S): EPAPLT0154E07

M4ID: 30204201

HILL MODEL (in red):

	tp	ga	gw
val:	121	1.4	8
sd:	288	0.279	41.8

GAIN-LOSS MODEL (in blue):

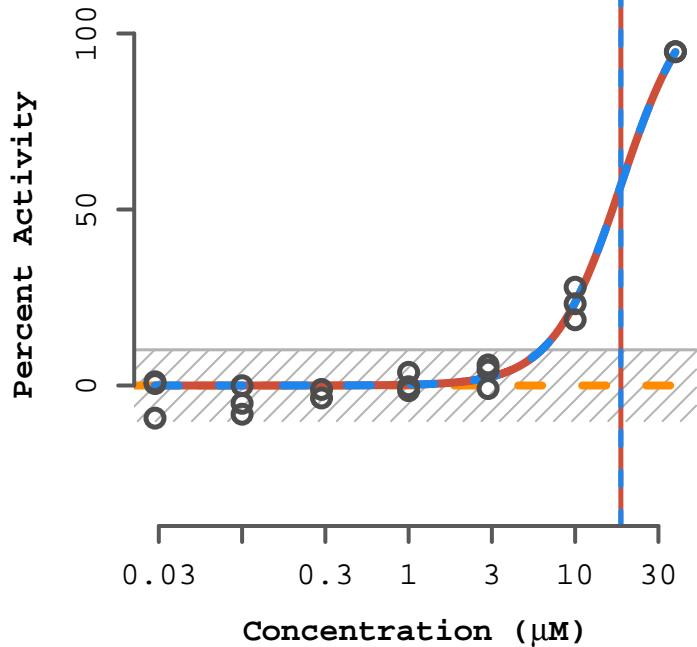
	tp	ga	gw	la	lw
val:	119	1.4	8	3.4	8.12
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	124.69	119.45	123.45
PROB:	0.06	0.83	0.11
RMSE:	45.35	35.62	35.61

MAX_MEAN: 64.5 MAX_MED: 96.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 0.94

FLAGS: 10



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: PharmaGSID_48509

CHID: 48509 CASRN: NOCAS_48509

SPID(S): TP0001414H10

M4ID: 30204624

HILL MODEL (in red):

	tp	ga	gw
val:	114	1.27	2.12
sd:	18.8	0.116	0.555

GAIN-LOSS MODEL (in blue):

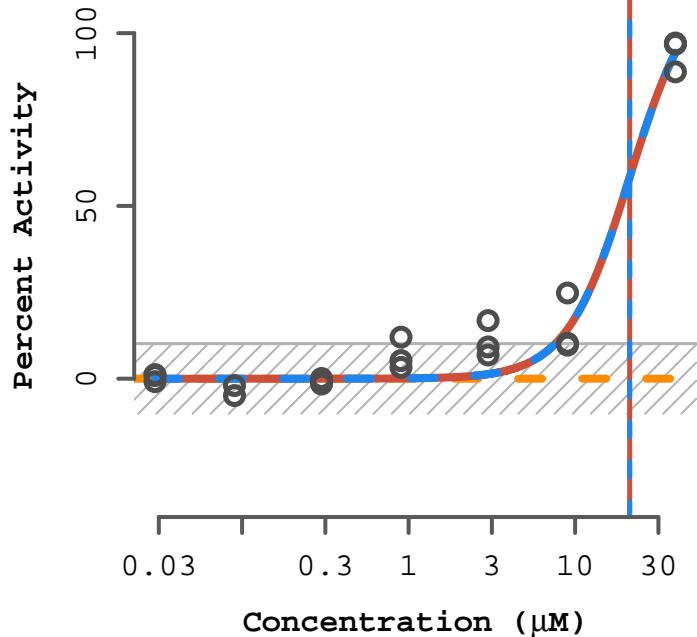
	tp	ga	gw	la	lw
val:	114	1.27	2.12	2.94	8.82
sd:	18.8	0.116	0.555	79800	527000

	CNST	HILL	GNLS
AIC:	173.93	112.81	116.81
PROB:	0	0.88	0.12
RMSE:	32.38	3.83	3.83

MAX_MEAN: 94.8 MAX_MED: 94.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: PharmaGSID_48511

CHID: 48511 CASRN: 1062243-51-9

SPID(S): TP0001414H09

M4ID: 30204625

HILL MODEL (in red):

	tp	ga	gw
val:	117	1.33	2.28
sd:	44	0.233	1.24

GAIN-LOSS MODEL (in blue):

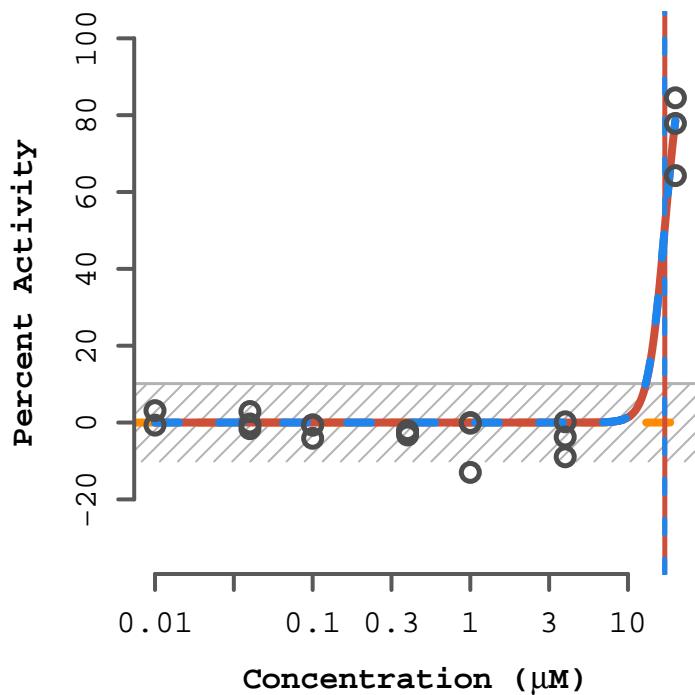
	tp	ga	gw	la	lw
val:	117	1.33	2.28	3.57	8.53
sd:	NA	NA	NA	NA	NA

	CNST	HILL	GNLS
AIC:	192.19	136.54	140.54
PROB:	0	0.88	0.12
RMSE:	37.51	6.08	6.08

MAX_MEAN: 94.3 MAX_MED: 96.8 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS:



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: HMR1171 trifluoroacetate (1:1)

CHID: 48522 CASRN: NOCAS_48522

SPID(S): TP0001413F03

M4ID: 30204628

HILL MODEL (in red):

tp	ga	gw
val: 101	1.23	8
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 101	1.23	7.92	3.25	8.75
sd: NA	NA	NA	NA	NA

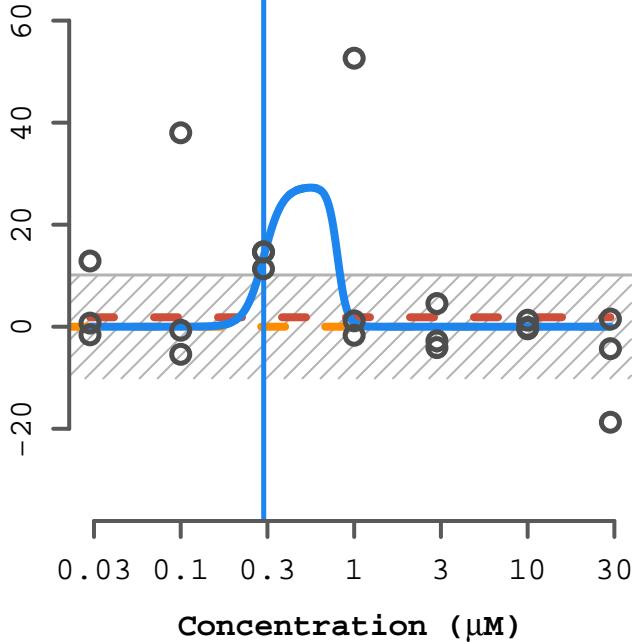
CNST	HILL	GNLS
AIC: 176.6	127.35	131.35
PROB: 0	0.88	0.12
RMSE: 29.72	5.3	5.3

MAX_MEAN: 75.6 MAX_MED: 77.9 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 42 ACTP: 1

FLAGS: 7

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: Tripropylene glycol
CHID: 27837 CASRN: 24800-44-0
SPID(S): EPAPLT0154H03
M4ID: 30204226

HILL MODEL (in red):

tp	ga	gw
val: 1.85	-2.52	5.43
sd: 2.14	322	1750

GAIN-LOSS MODEL (in blue):

tp	ga	gw	la	lw
val: 27.5	-0.521	8	-0.0959	15.9
sd: 939	3.66	51.2	12.2	2040

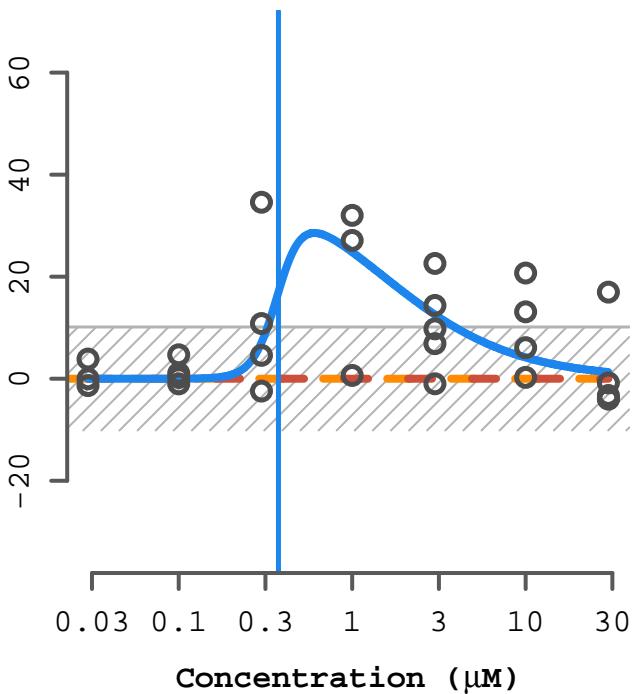
CNST	HILL	GNLS
AIC: 169.24	174.45	168.77
PROB: 0.43	0.03	0.54
RMSE: 16.03	15.5	15.06

MAX_MEAN: 17.4 MAX_MED: 14.7 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 50 ACTP: 0.57

FLAGS: 17; 16; 11; 10; 7

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: 3-(Dimethylphosphono)-N-methylolpropionamide
CHID: 40215 CASRN: 20120-33-6
SPID(S): EPAPLT0154F02
M4ID: 30204205

HILL MODEL (in red):

tp	ga	gw
val: 18.2	1.98	8
sd: NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

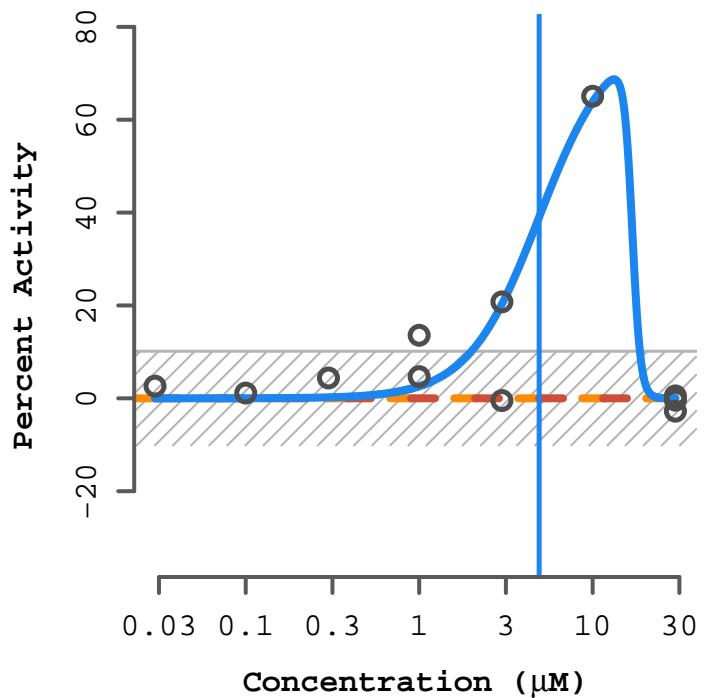
tp	ga	gw	la	lw
val: 41.5	-0.425	6.15	0.155	1.13
sd: 31.3	0.269	16.3	0.583	0.569

CNST	HILL	GNLS
AIC: 223.23	229.24	214.3
PROB: 0.01	0	0.99
RMSE: 13.33	13.33	9.59

MAX_MEAN: 19.9 MAX_MED: 27.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 50 ACTP: 0.99

FLAGS: 17; 7



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: (4-Methoxyphenyl)methanol
CHID: 44357 CASRN: 105-13-5
SPID(S): EPAPLT0154E08
M4ID: 30204202

HILL MODEL (in red):

	tp	ga	gw
val:	51.1	1.98	8
sd:	NaN	NaN	NaN

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	78.1	0.692	2.12	1.23	18
sd:	21.2	0.154	1.06	0.735	55.3

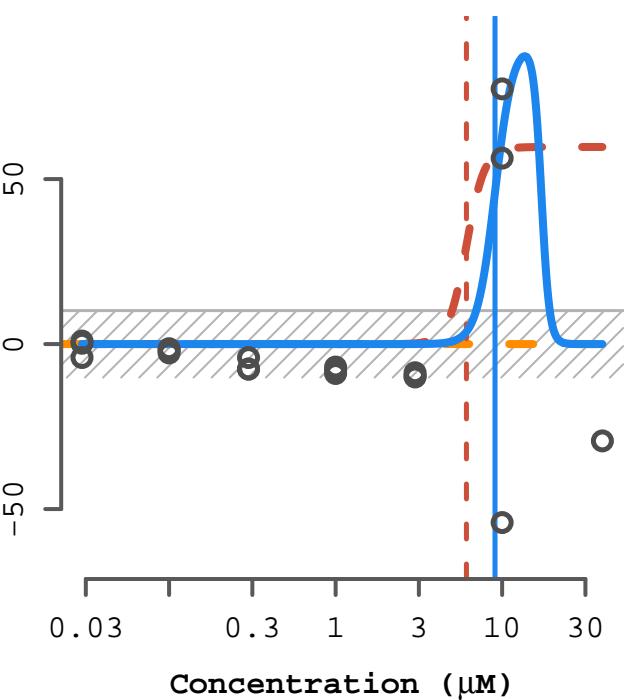
	CNST	HILL	GNLS
AIC:	92.39	98.39	82.43
PROB:	0.01	0	0.99
RMSE:	21.12	21.12	7.34

MAX_MEAN: 65.1 MAX_MED: 65.1 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 50 ACTP: 0.99

FLAGS: 16

Percent Activity



ASSAY: AEID2455 (NHEERL_MEA_acute_synchrony_index_dn)

NAME: MK-274

CHID: 47328 CASRN: NOCAS_47328

SPID(S): TP0001413C05

M4ID: 30204513

HILL MODEL (in red):

	tp	ga	gw
val:	59.7	0.785	8
sd:	10.4	0.159	4.72

GAIN-LOSS MODEL (in blue):

	tp	ga	gw	la	lw
val:	92	0.957	8	1.24	17.9
sd:	NaN	NaN	NaN	NaN	NaN

	CNST	HILL	GNLS
AIC:	173.66	175.82	169.81
PROB:	0.12	0.04	0.84
RMSE:	26.71	33.73	28.55

MAX_MEAN: 26.5 MAX_MED: 56.4 BMAD: 3.38

COFF: 10.1 HIT-CALL: 1 FITC: 50 ACTP: 0.88

FLAGS: 16; 11; 10; 7