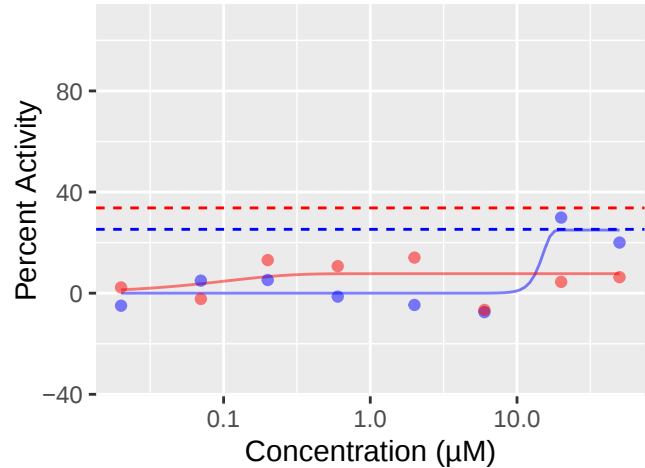


DTXSID2045232 Tiratricol
SPID:TP0000904H05

A: AEID:623 AENM:NVS_GPCR_h5HT5A
M4ID:1833668

B: AEID:679 AENM:NVS_GPCR_rOpiate_NonSelective
M4ID:1839401



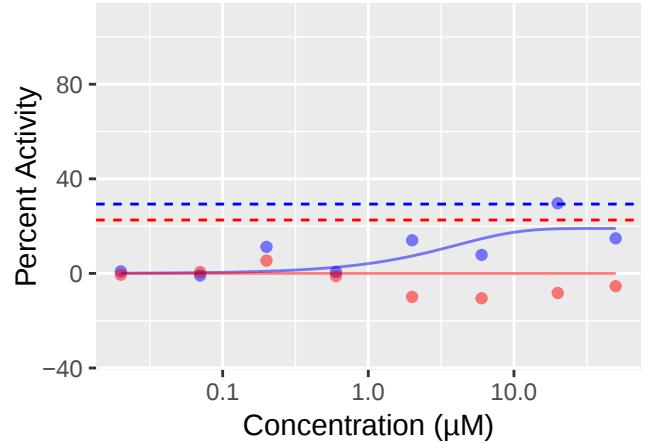
Flags:
A(3): Borderline;
Average number of replicates per conc is less than 2;
Multiple points above baseline, inactive

B(1): Average number of replicates per conc is less than 2

DTXSID2045232 Tiratricol
SPID:TP0000904H05

A: AEID:626 AENM:NVS_GPCR_hAdoRA1
M4ID:1834094

B: AEID:704 AENM:NVS_LGIC_rGlyRStrySens
M4ID:1842443



Flags:
A(3): Average number of replicates per conc is less than 2;
Multiple points above baseline, inactive;
Bmd > ac50, indication of high baseline variability

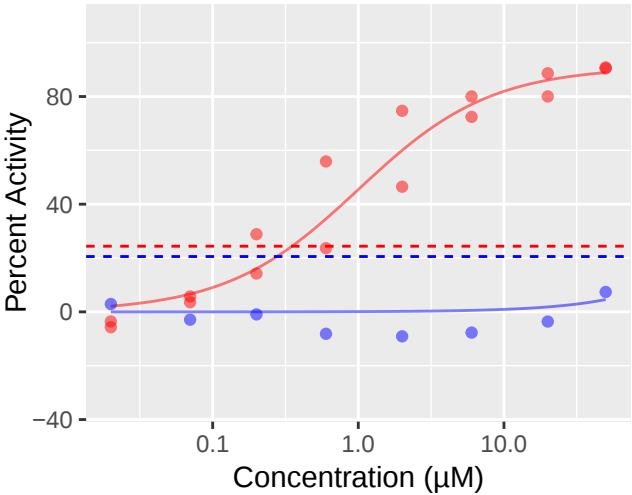
B(2): Bmd > ac50, indication of high baseline variability;
Average number of replicates per conc is less than 2

	Hitcall	BMD	AC50
A	0.388	13.392	14.428
B	3.7e-15	NA	0.073

DTXSID2045232 Tiratricol
SPID:TP0000904H05

A: AEID:645 AENM:NVS_GPCR_hM4
M4ID:1836233

B: AEID:714 AENM:NVS_NR_hER
M4ID:1847540



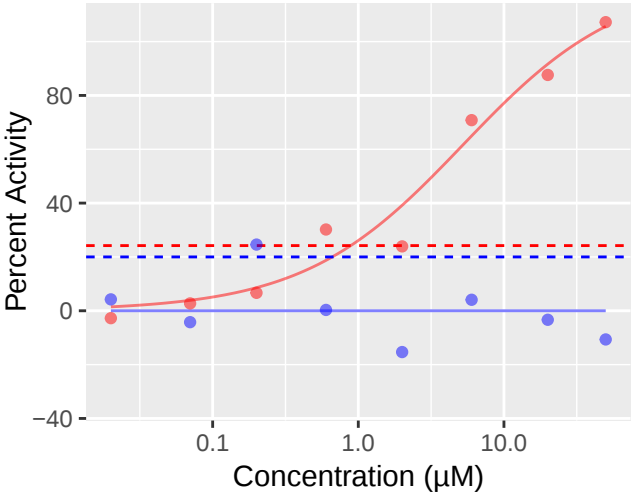
Flags:
A(2): Average number of replicates per conc is less than 2;
Bmd > ac50, indication of high baseline variability

B(1): Bmd > ac50, indication of high baseline variability

DTXSID2045232 Tiratricol
SPID:TP0000904H05

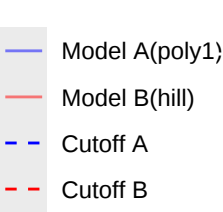
A: AEID:647 AENM:NVS_GPCR_hNK2
M4ID:1836494

B: AEID:719 AENM:NVS_NR_hPPARg
M4ID:1850045



Flags:
A(2): Average number of replicates per conc is less than 2;
Bmd > ac50, indication of high baseline variability

B(1): Average number of replicates per conc is less than 2



	Hitcall	BMD	AC50
A	4.5e-05	144.046	25.000
B	1.000	1.166	1.003