Figure 1

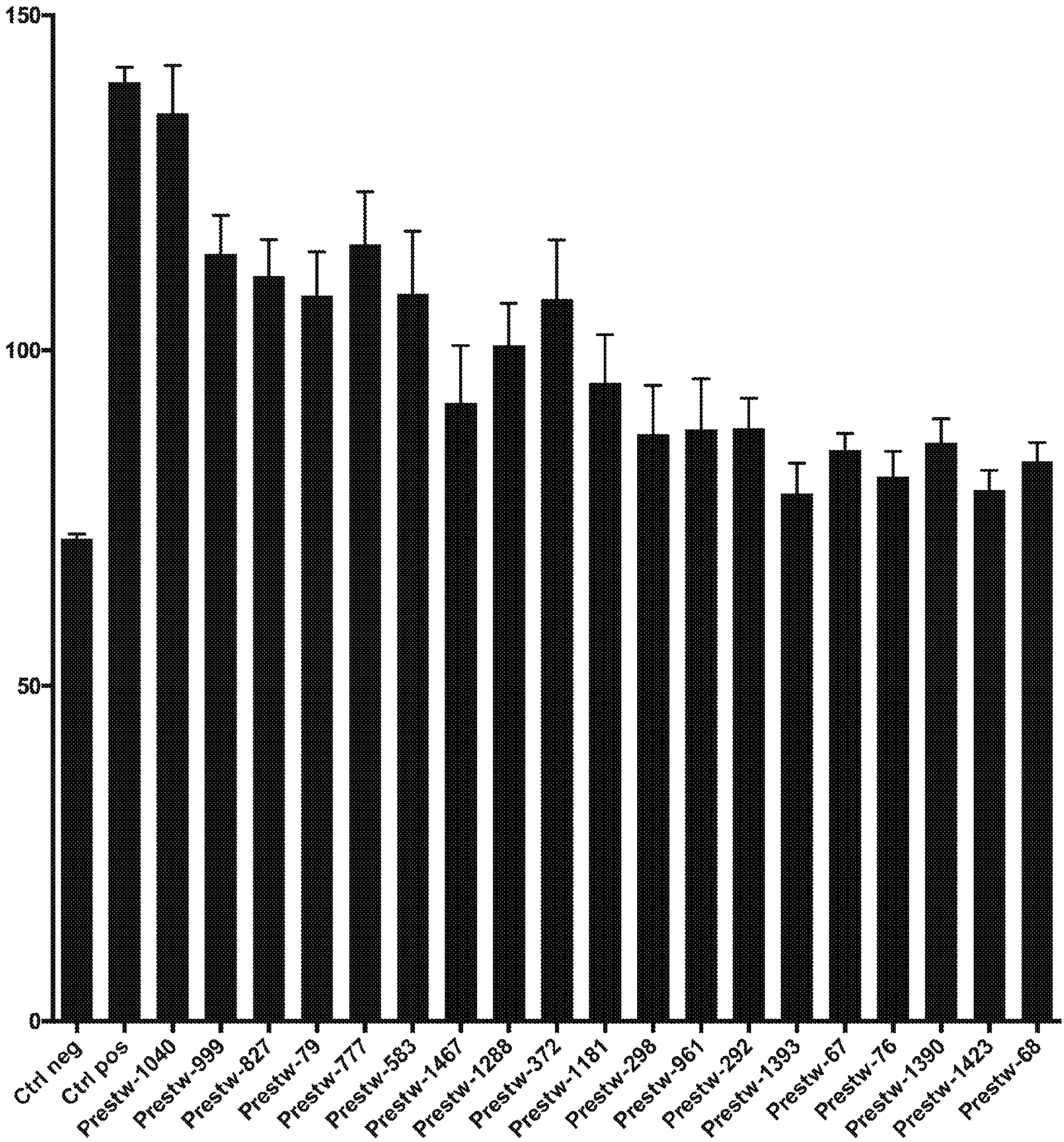

150
100
50
0
Lactate (uM)
Ctrl neg
Ctrl pos
Prestw-1040
Prestw-999
Prestw-827
Prestw-79
Prestw-777
Prestw-583
Prestw-1467
Prestw-1288
Prestw-372
Prestw-1181
Prestw-298
Prestw-961
Prestw-292
Prestw-1393
Prestw-67
Prestw-76
Prestw-1390
Prestw-1423
Prestw-68

Figure 2

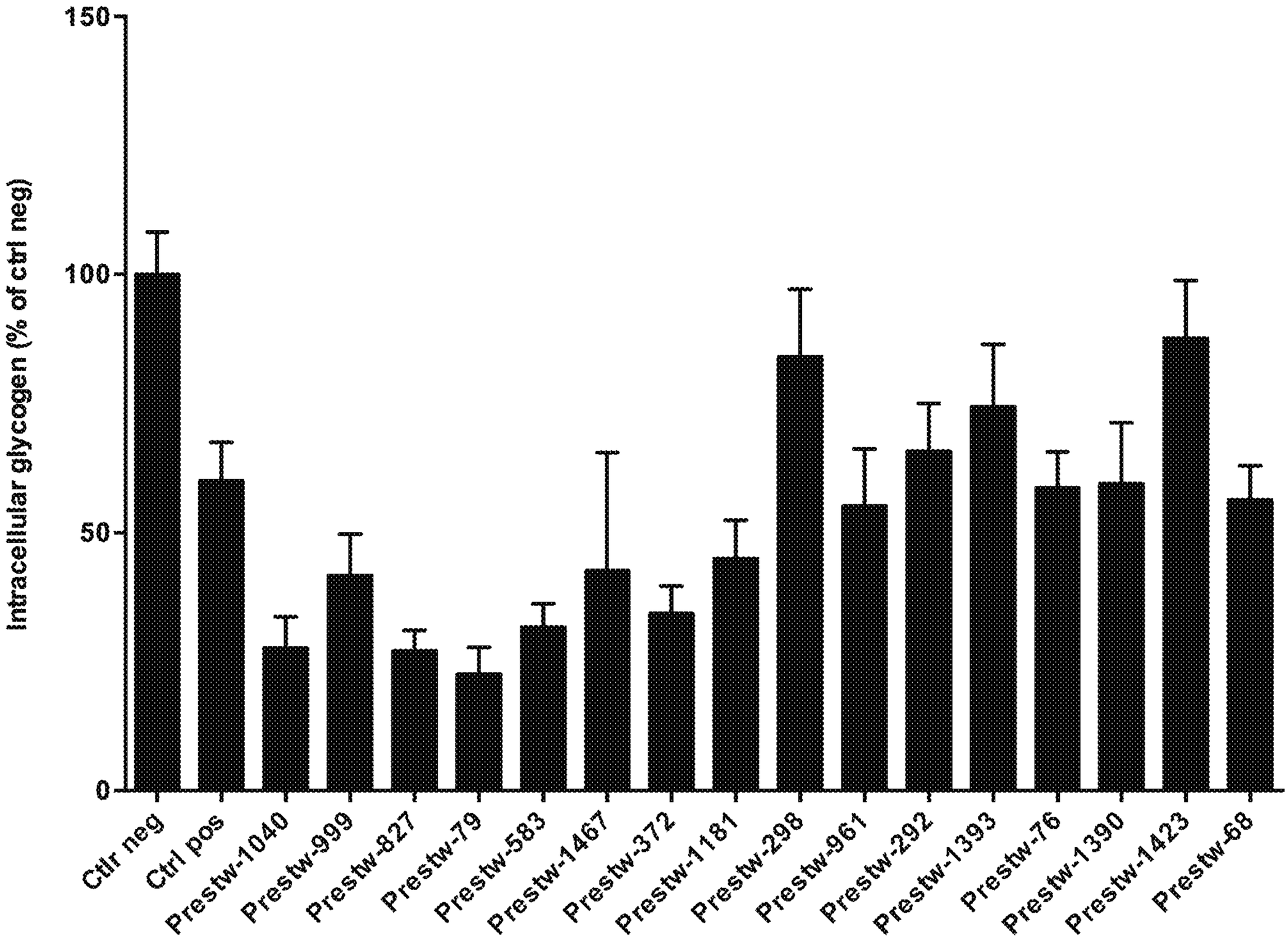

Intracellular glycogen (% of ctrl neg)
150
100
50
0
Ctlr neg
Ctrl pos
Prestw-1040
Prestw-999
Prestw-827
Prestw-79
Prestw-583
Prestw-1467
Prestw-372
Prestw-1181
Prestw-298
Prestw-961
Prestw-292
Prestw-1393
Prestw-76
Prestw-1390
Prestw-1423
Prestw-68

Figure 3

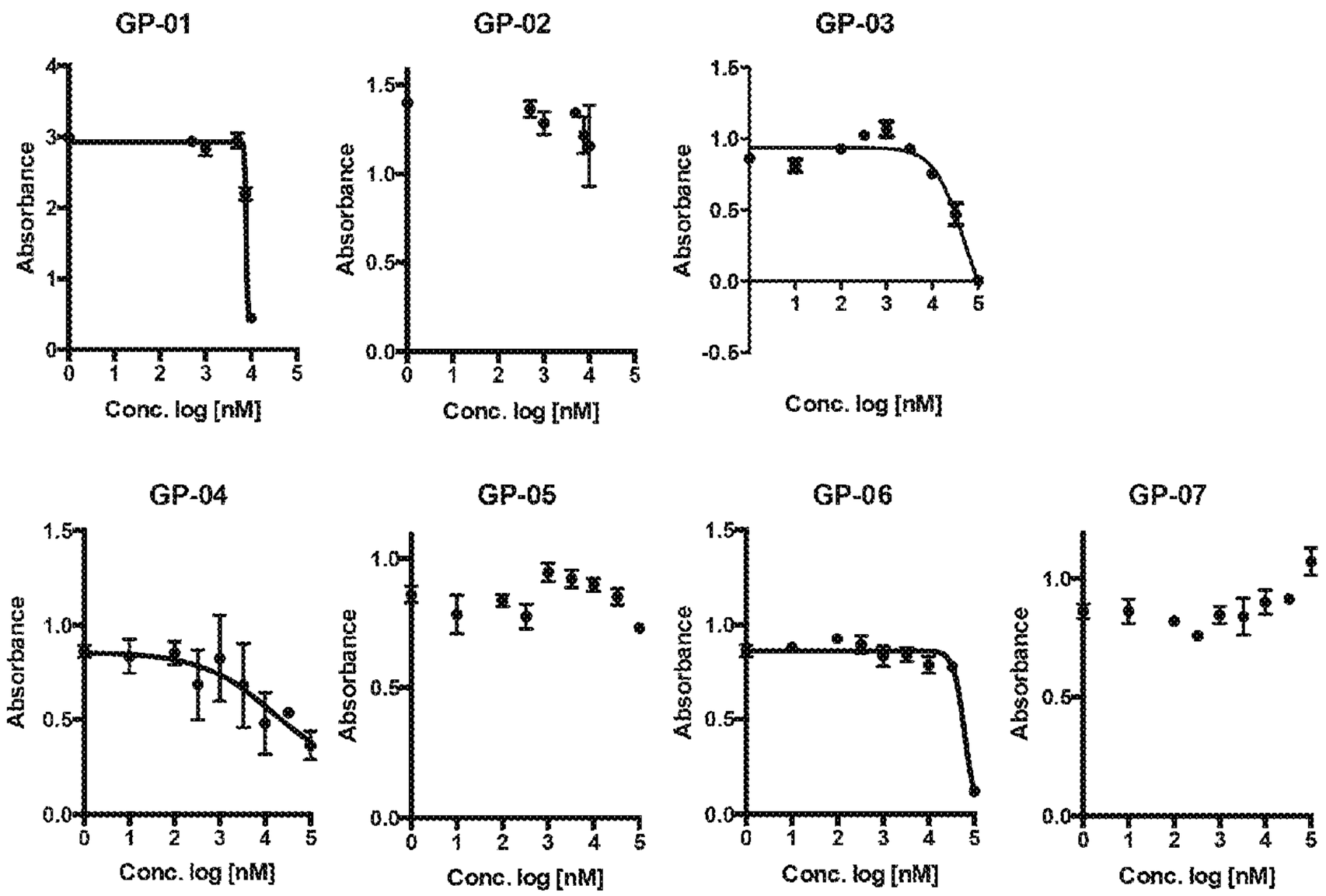


Figure 4

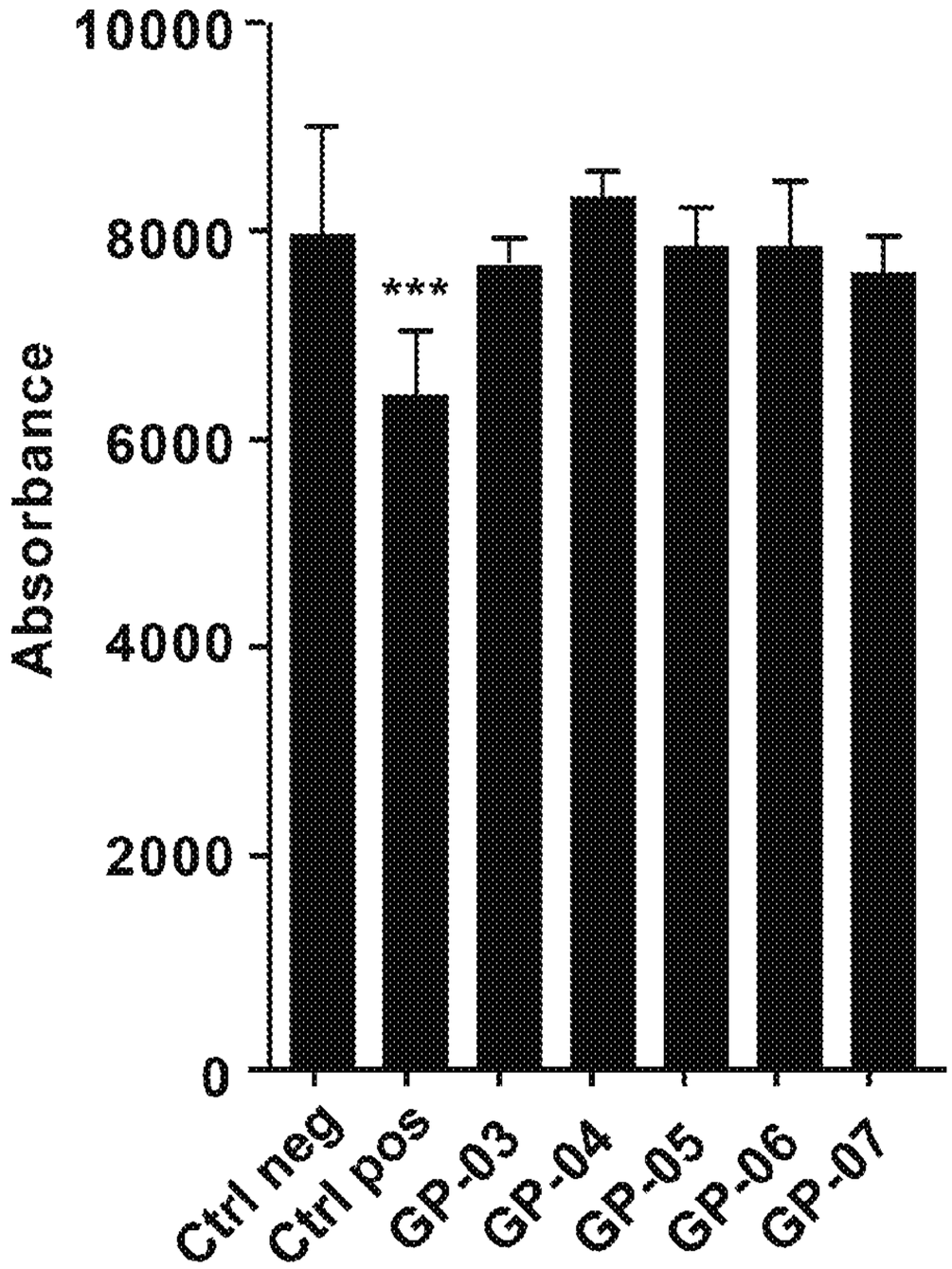

10000
8000
6000
4000
2000
0
Absorbance
***
Ctrl neg
Ctrl pos
GP-03
GP-04
GP-05
GP-06
GP-07

Figure 5A

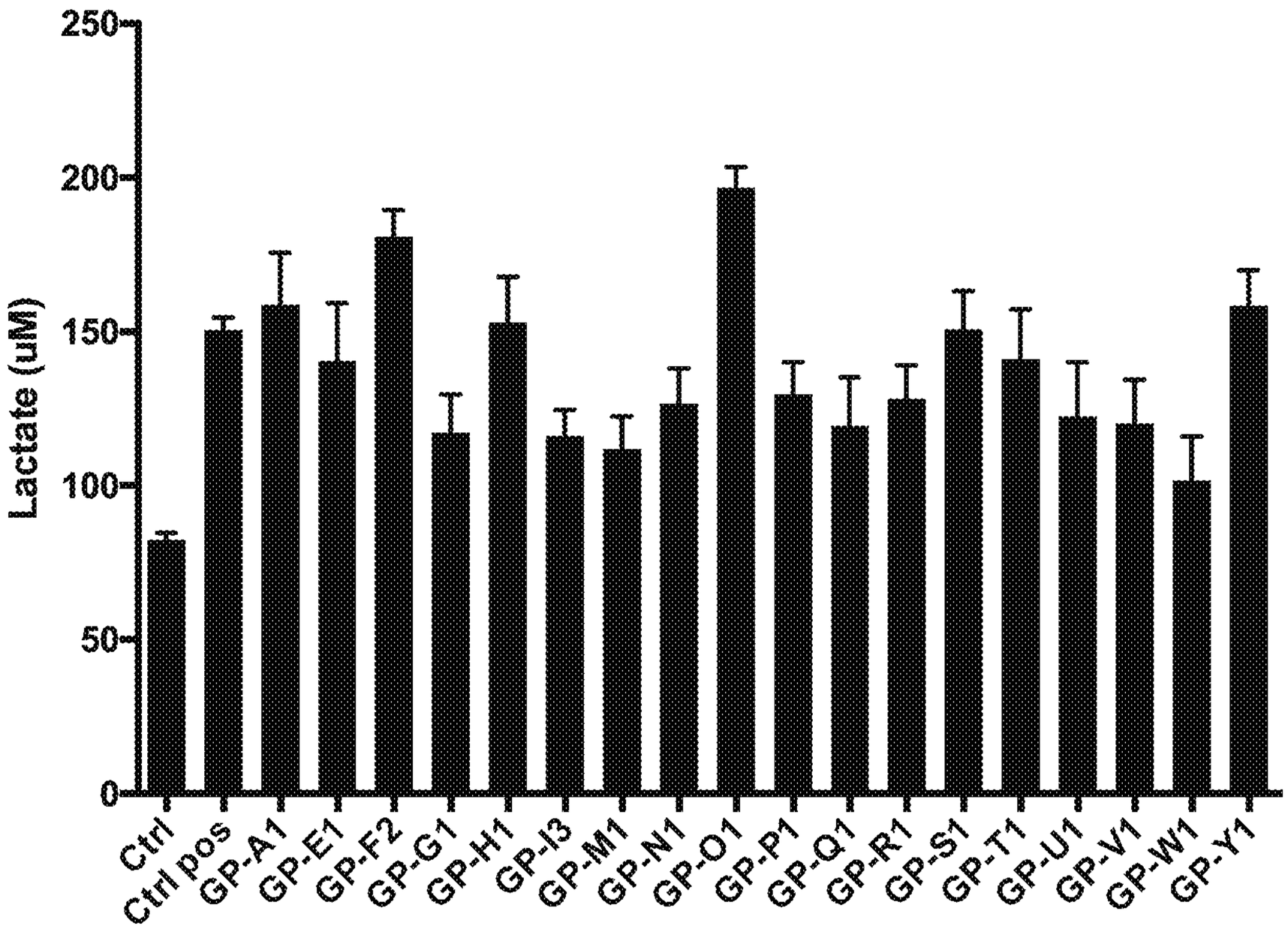

Lactate (uM)
250
200
150
100
50
0
Ctrl
Ctrl pos
GP-A1
GP-E1
GP-F2
GP-G1
GP-H1
GP-I3
GP-M1
GP-N1
GP-O1
GP-P1
GP-Q1
GP-R1
GP-S1
GP-T1
GP-U1
GP-V1
GP-W1
GP-Y1

Figure 5B

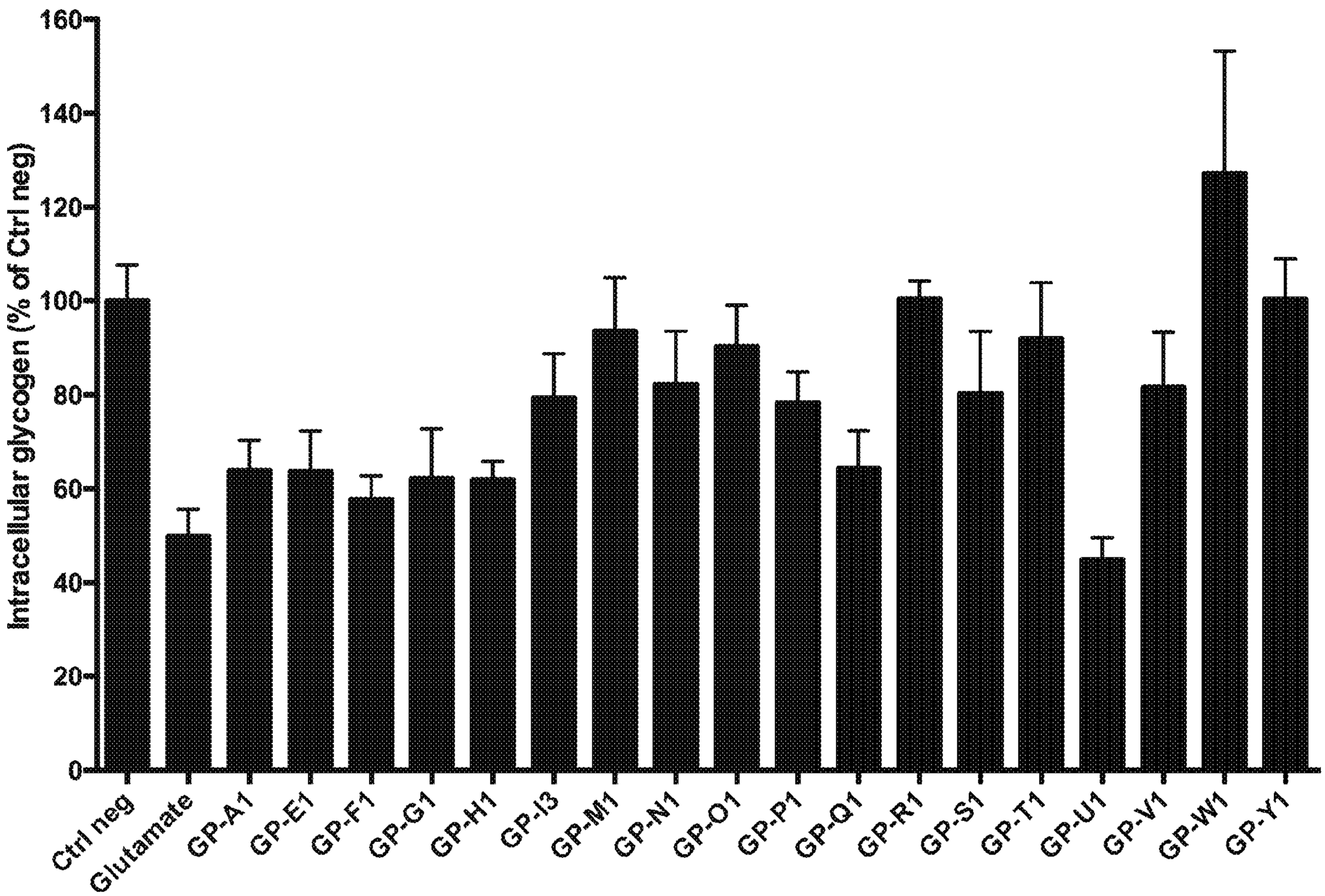

Intracellular glycogen (% of Ctrl neg)
160
140
120
100
80
60
40
20
0
Ctrl neg
Glutamate
GP-A1
GP-E1
GP-F1
GP-G1
GP-H1
GP-I3
GP-M1
GP-N1
GP-O1
GP-P1
GP-Q1
GP-R1
GP-S1
GP-T1
GP-U1
GP-V1
GP-W1
GP-Y1

Figure 6

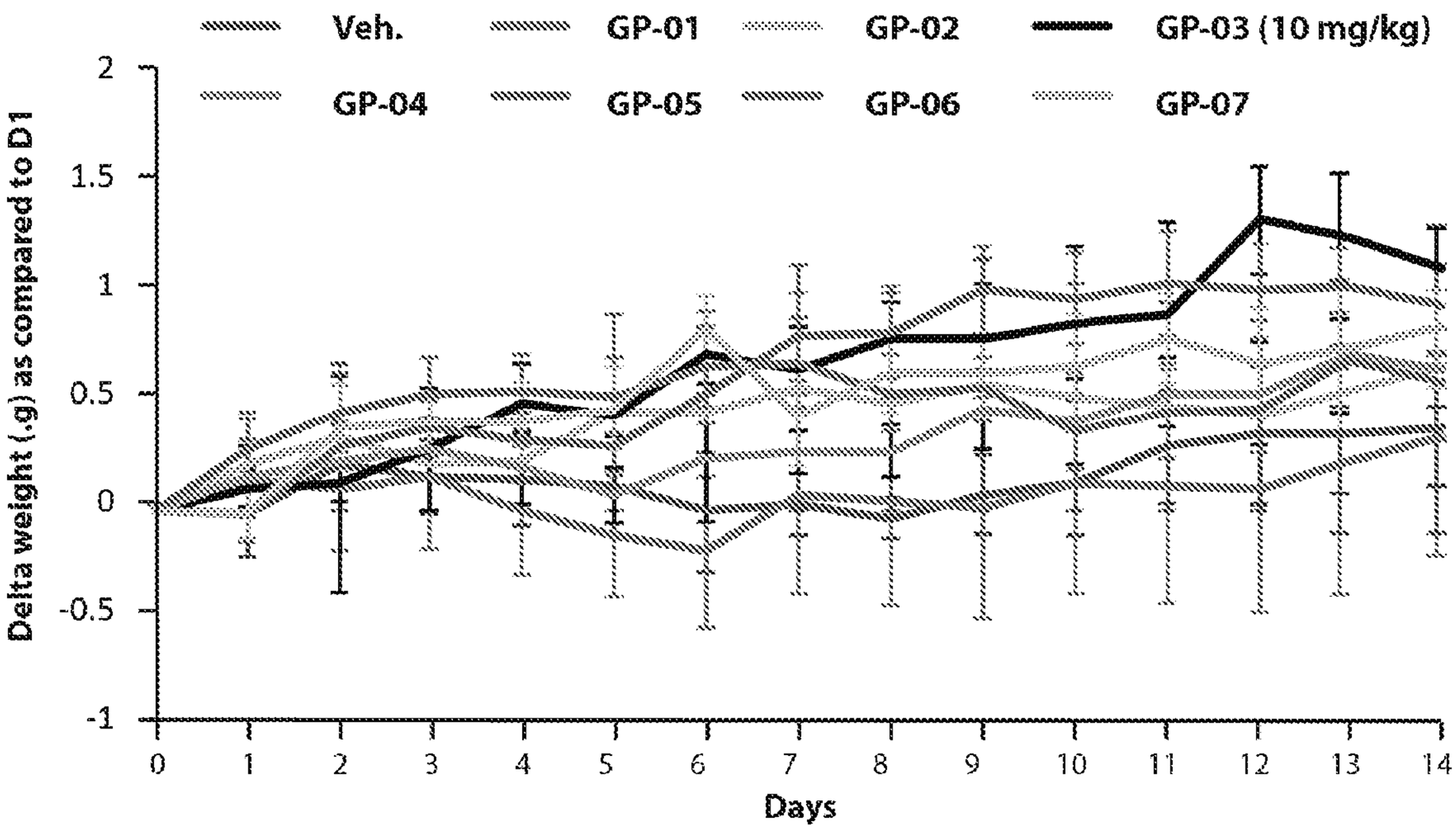

Veh.
GP-01
GP-02
GP-03 (10 mg/kg)
GP-04
GP-05
GP-06
GP-07
Delta weight (.g) as compared to D1
2
1.5
1
0.5
0
-0.5
-1
0
1
2
3
4
5
6
7
8
9
10
11
12
13
14
Days

Figure 7

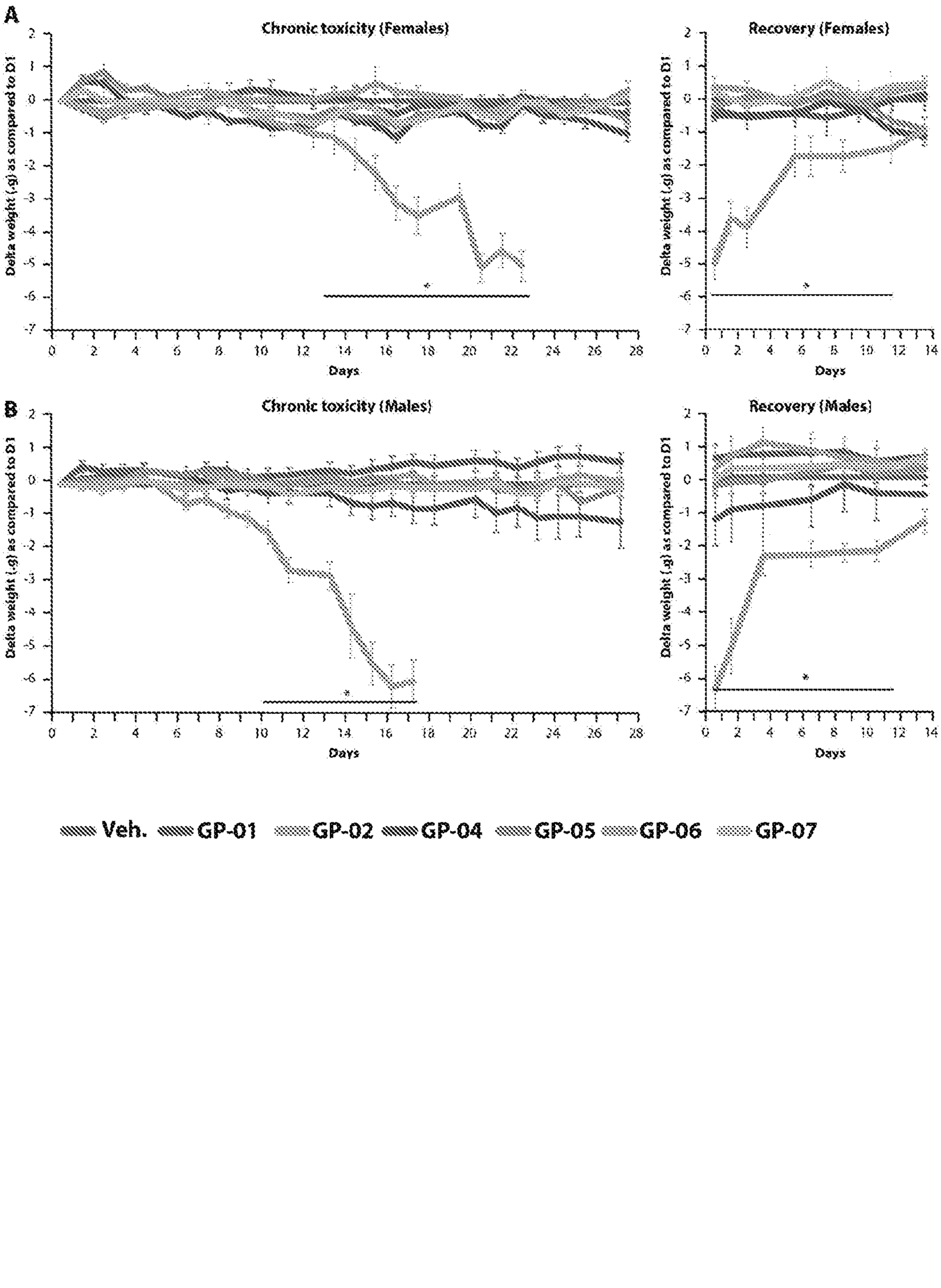

A
Chronic toxicity (Females)
Recovery (Females)
Delta weight (.g) as compared to D1
Days
B
Chronic toxicity (Males)
Recovery (Males)
Delta weight (.g) as compared to D1
Days
Veh.
GP-01
GP-02
GP-04
GP-05
GP-06
GP-07

Figure 8

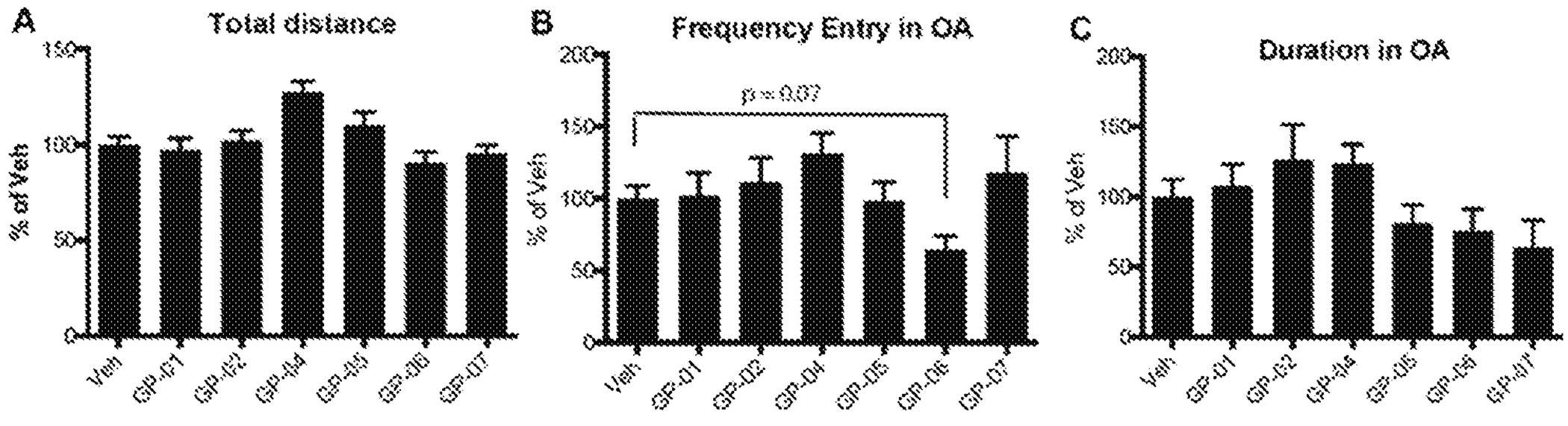

A
Total distance
% of Veh
150
100
50
0
Veh
GP-01
GP-02
GP-04
GP-05
GP-06
GP-07
B
Frequency Entry in OA
p = 0.07
% of Veh
200
150
100
50
0
Veh
GP-01
GP-02
GP-04
GP-05
GP-06
GP-07
C
Duration in OA
% of Veh
200
150
100
50
0
Veh
GP-01
GP-02
GP-04
GP-05
GP-06
GP-07

Figure 9

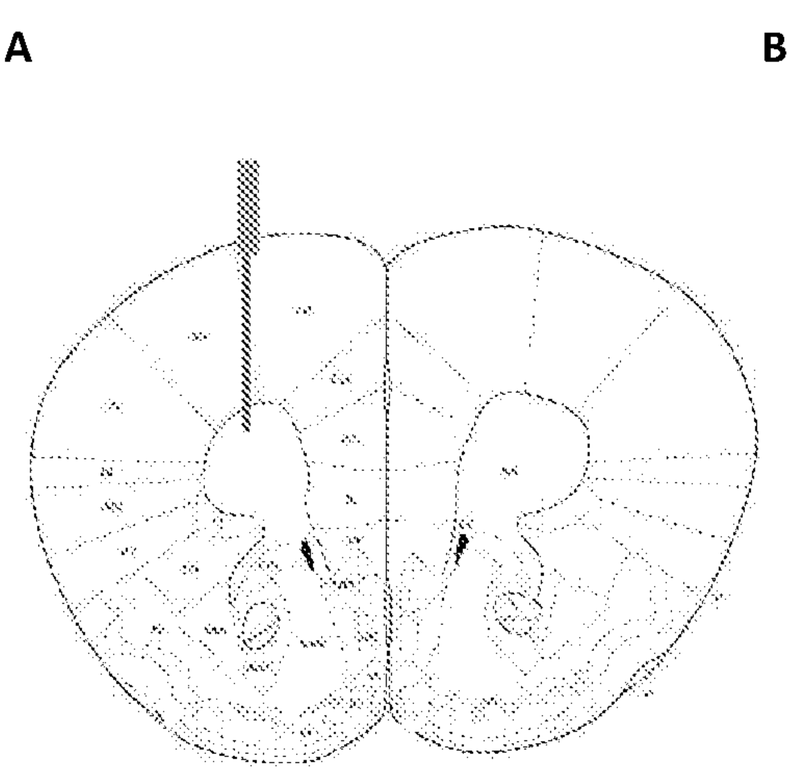

A
B
Veh.
GP-07
250
200
150
100
50
0
Lactate (μM)
0
60
120
180
240
300
360 (min.)
Veh. AUC
GP-07 AUC
30
60
90
120
150
180 (min.)
C
D
Area Under Curve Ratio (Treatment / Veh)
2
1.5
1
0.5
0
**
***
*
***
Veh.
GP-01
GP-02
GP-03
GP-04
GP-05
GP-06
GP-07
10 mg/kg
2.5
*
*
GP-A
GP-I
GP-P
GP-Q
GP-R
GP-V
100 mg/kg

Figure 10

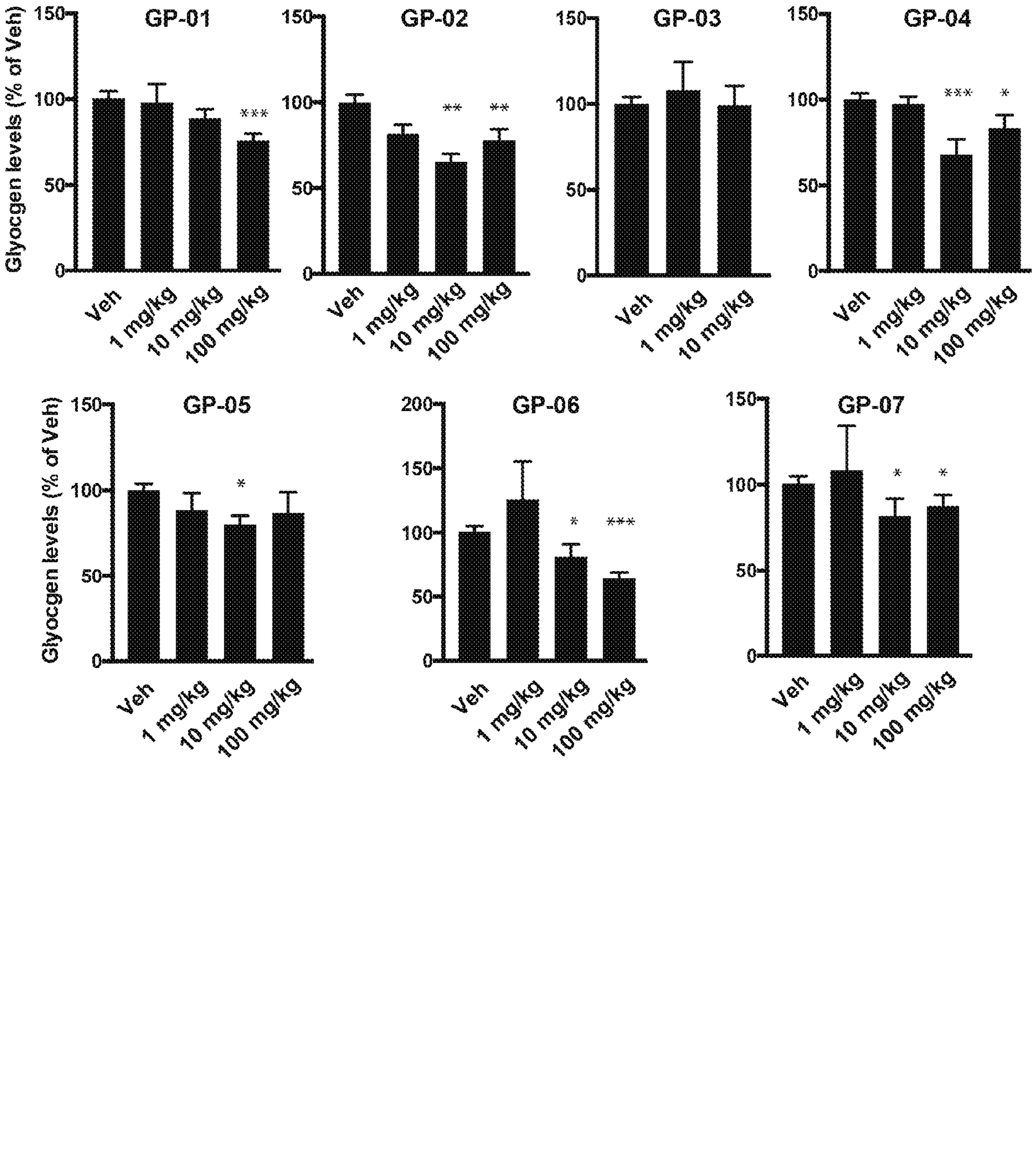

GP-01
GP-02
GP-03
GP-04
GP-05
GP-06
GP-07
Glyocgen levels (% of Veh)
Veh
1 mg/kg
10 mg/kg
100 mg/kg

Figure 11A

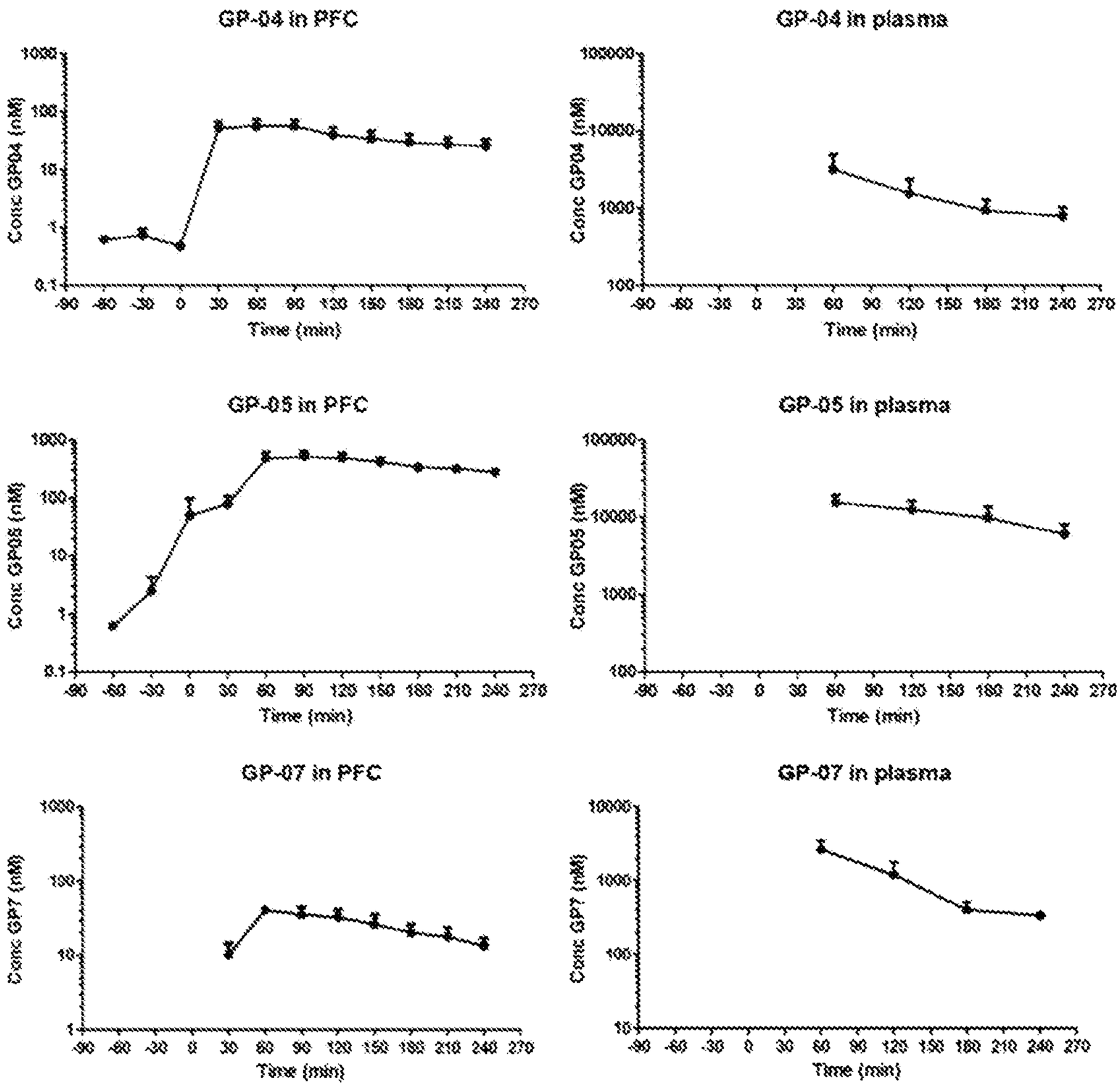

GP-04 in PFC
Conc GP04 (nM)
Time (min)
GP-04 in plasma
Conc GP04 (nM)
Time (min)
GP-05 in PFC
Conc GP05 (nM)
Time (min)
GP-05 in plasma
Conc GP05 (nM)
Time (min)
GP-07 in PFC
Conc GP7 (nM)
Time (min)
GP-07 in plasma
Conc GP7 (nM)
Time (min)

Figure 11B

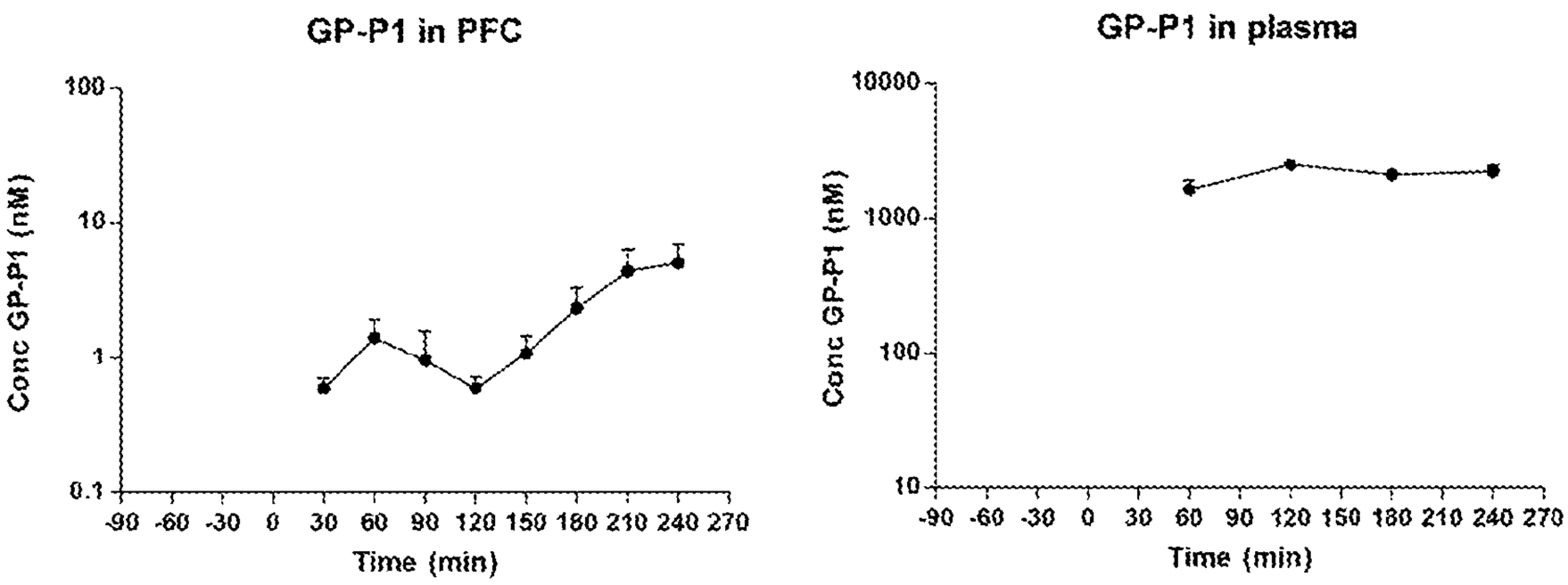

GP-P1 in PFC
Conc GP-P1 (nM)
100
10
1
0.1
-90 -60 -30 0 30 60 90 120 150 180 210 240 270
Time (min)
GP-P1 in plasma
Conc GP-P1 (nM)
10000
1000
100
10
-90 -60 -30 0 30 60 90 120 150 180 210 240 270
Time (min)
GP-R1 in PFC
Conc GP-R1 (nM)
1000
100
10
1
-90 -60 -30 0 30 60 90 120 150 180 210 240 270
Time (min)
GP-R1 in plasma
Conc GP-R1 (nM)
10000
1000
100
10
-90 -60 -30 0 30 60 90 120 150 180 210 240 270
Time (min)

Figure 12

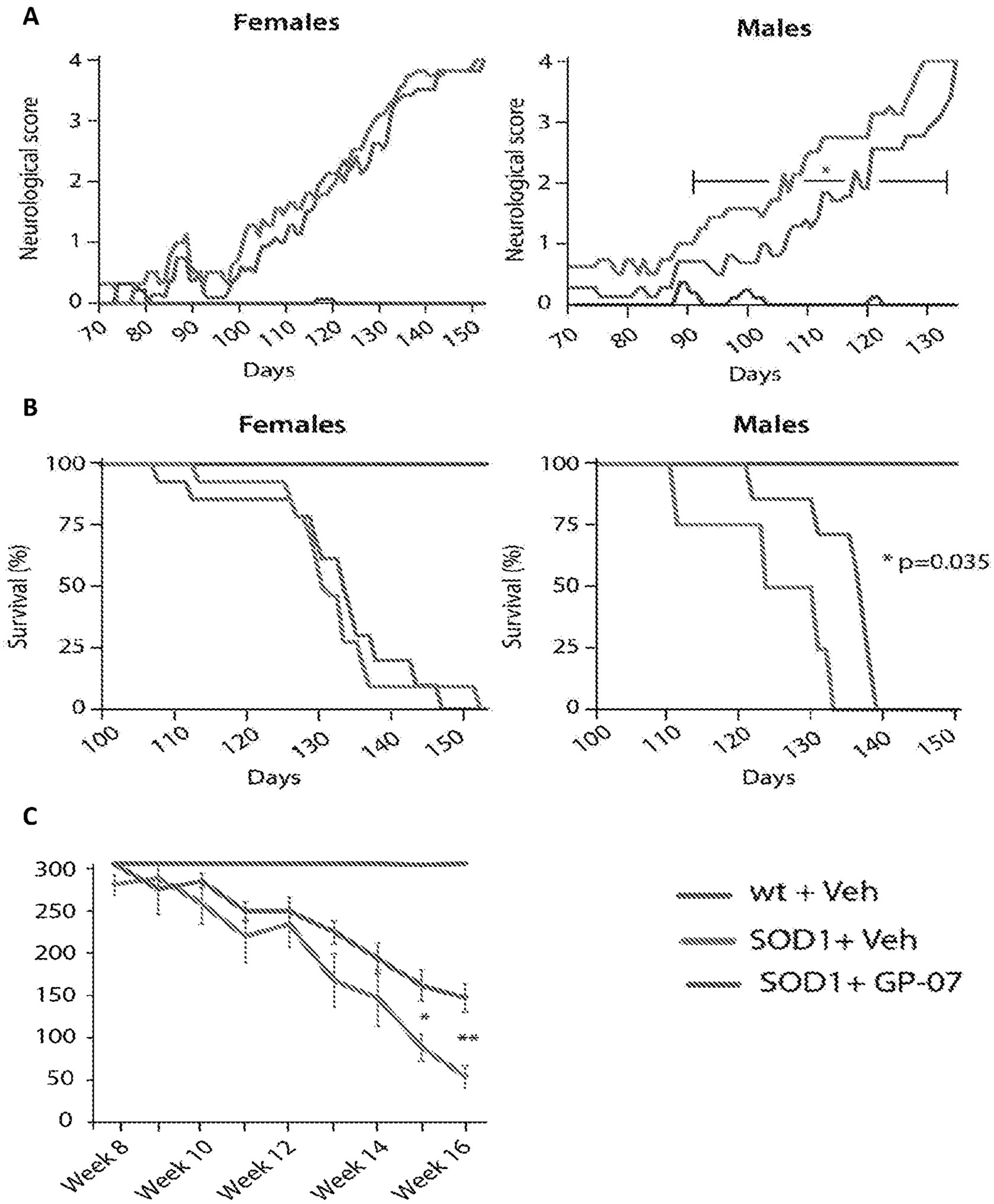

A
Females
Males
Neurological score
Days
B
Females
Males
Survival (%)
* p=0.035
Days
C
Week 8
Week 10
Week 12
Week 14
Week 16
wt + Veh
SOD1+ Veh
SOD1+ GP-07

Figure 13

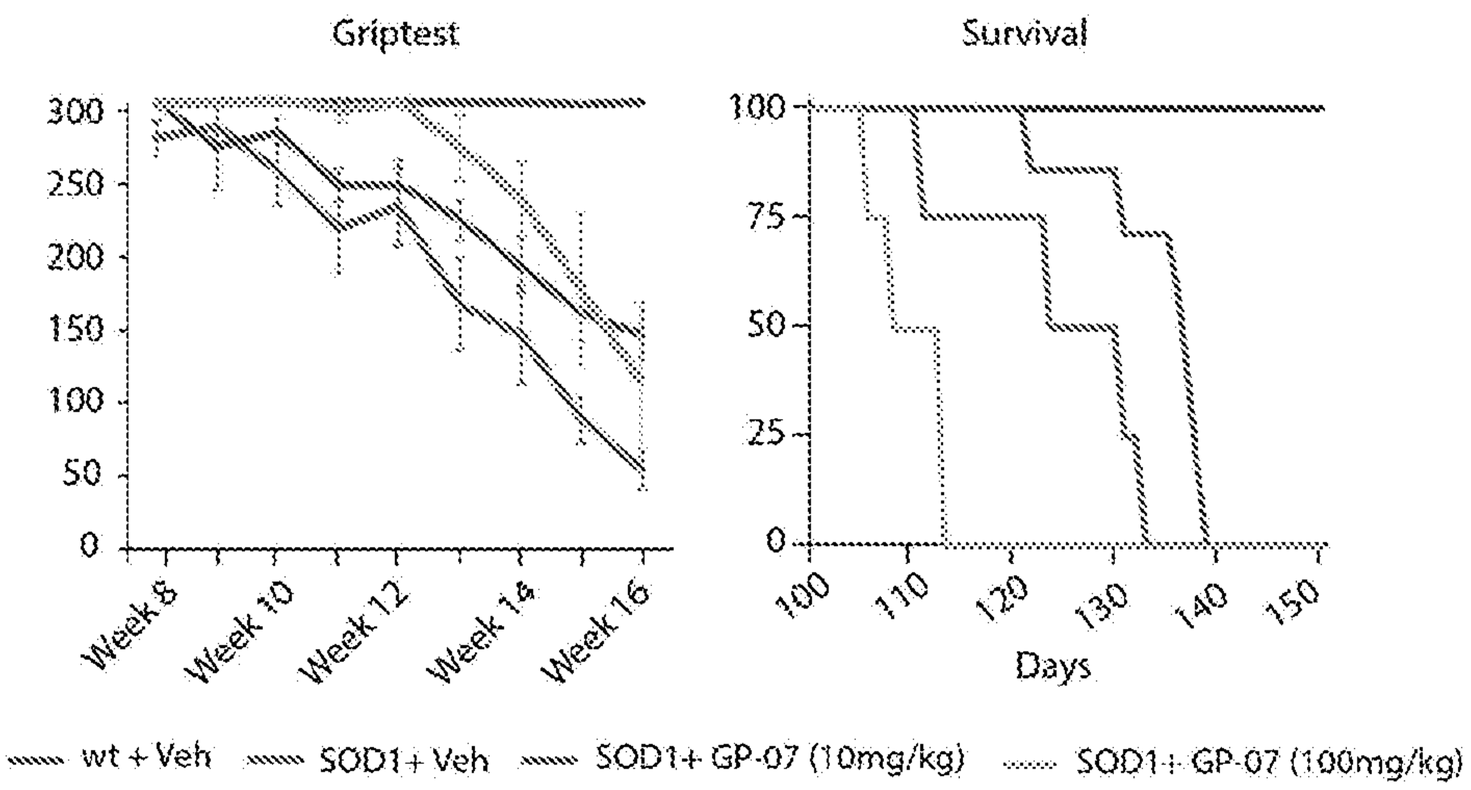
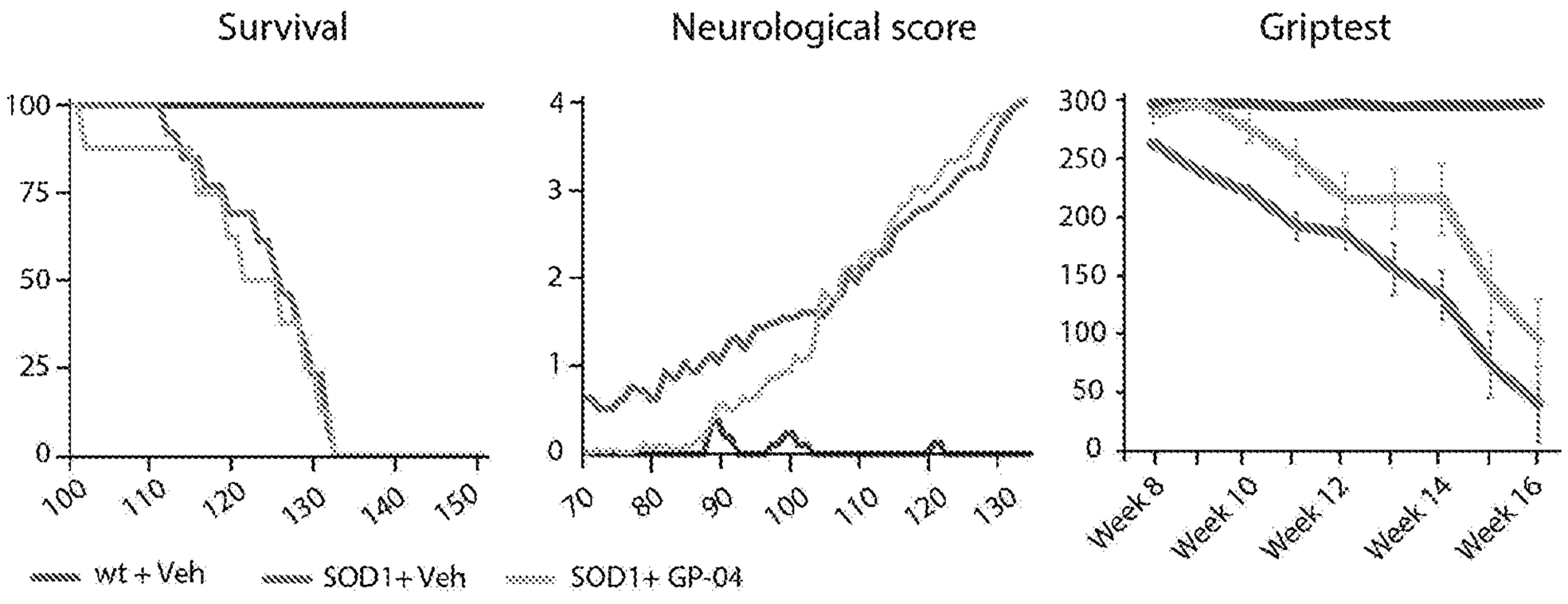

Griptest
300
250
200
150
100
50
0
Week 8
Week 10
Week 12
Week 14
Week 16
Survival
100
75
50
25
0
100
110
120
130
140
150
Days
wt + Veh
SOD1+ Veh
SOD1+ GP-07 (10mg/kg)
SOD1+ GP-07 (100mg/kg)

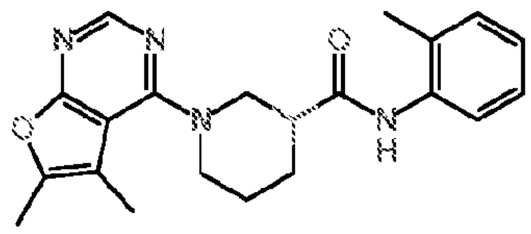
Figure 14


Survival
Neurological score
Griptest
100
75
50
25
0
100
110
120
130
140
150
4
3
2
1
0
70
80
90
100
110
120
130
300
250
200
150
100
50
0
Week 8
Week 10
Week 12
Week 14
Week 16
wt + Veh
SOD1+ Veh
SOD1+ GP-04

Appendix I

Family A

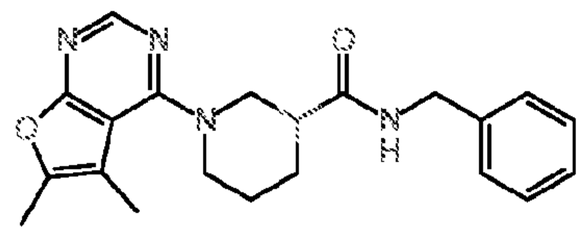
A1



unknown chirality

CatalogID	F228-0422
Score	1.057299178434214
Score_SNARF	0.7701853152834461
Score_LACTATE	0.6363638379469401

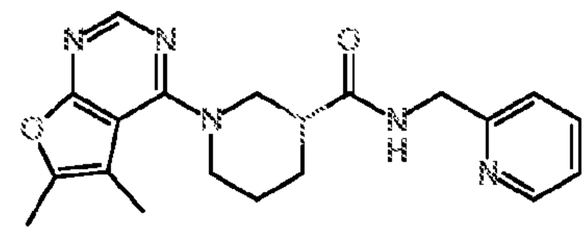
Toxic (MTT)



unknown chirality

CatalogID	F228-0365
Score	1.0347684901720595
Score_SNARF	0.942866402185284
Score_LACTATE	0.9000644924199342

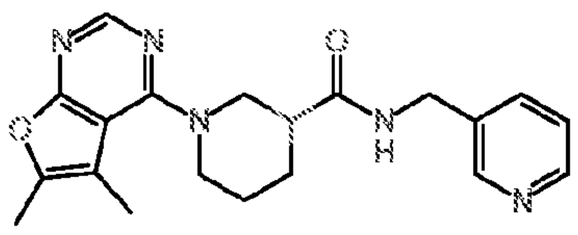
A2



unknown chirality

CatalogID	F228-0350
Score	0.9575255500229646
Score_SNARF	1.0109377915014888
Score_LACTATE	1.022932010836847

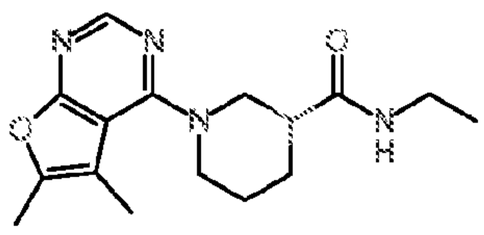
Toxic (MTT)



unknown chirality

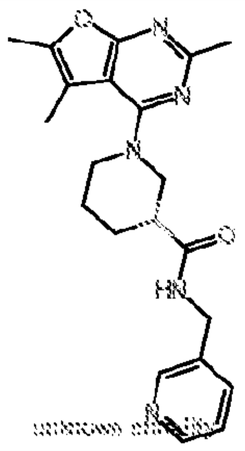
CatalogID	F228-0351
Score	0.5920192604665382
Score_SNARF	0.819975594094603
Score_LACTATE	0.898536677505049

A3



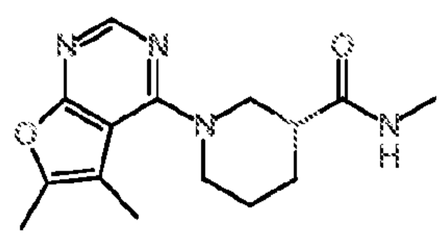
unknown chirality

CatalogID	F228-0534
Score	0.35368992473276867
Score_SNARF	0.5203267581394291
Score_LACTATE	0.3387117841389162



unknown chirality

CatalogID	F228-0856
Score	0.18647369720387383
Score_SNARF	0
Score_LACTATE	0

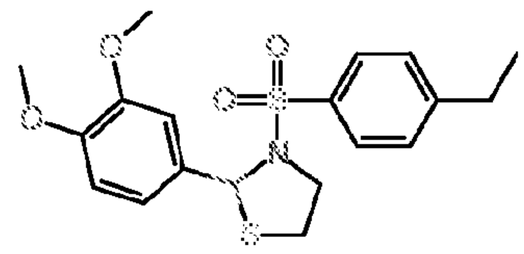


unknown chirality

CatalogID	F228-0541
Score	0.15277599688164334
Score_SNARF	0.36093980523381536
Score_LACTATE	0

Family C

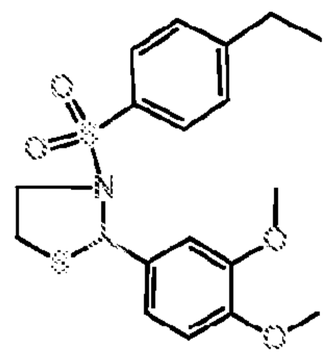
C1



unknown chirality

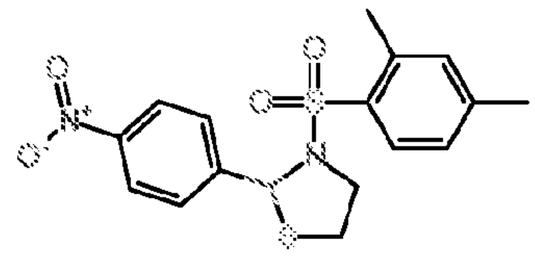
CatalogID	T5463586
Score	0.7706610463405941
Score_SNARF	0.5553240531901028
Score_LACTATE	0.3935782163943059

C2



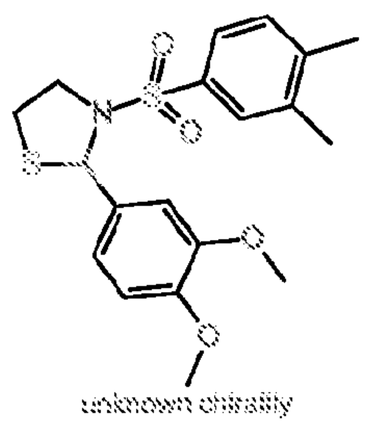
unknown chirality

CatalogID	4052-4304
Score	0.4940426943326044
Score_SNARF	0.4491394402072394
Score_LACTATE	0.433527881002696



unknown chirality

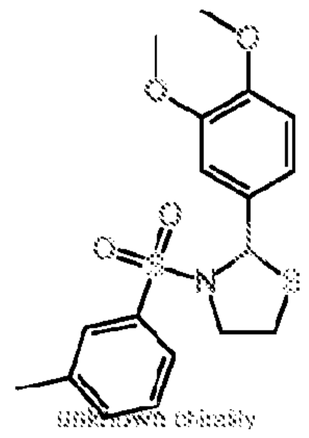
CatalogID	T5464782
Score	0.27666864801157476
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



unknown chirality

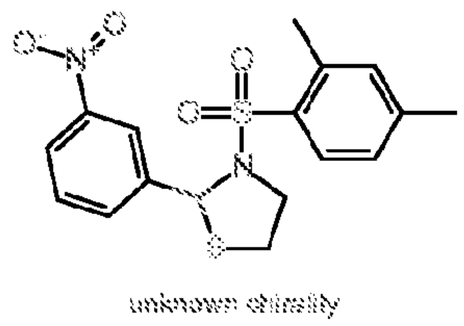
CatalogID	F1462-0491
Score	0.27239461193483955
Score_SNARF	0
Score_LACTATE	0

C3



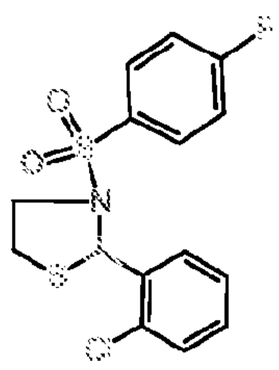
unknown chirality

CatalogID	T5463658
Score	0.2721822922258924
Score_SNARF	0.4683062878625885
Score_LACTATE	0.37725037983836096



unknown chirality

CatalogID	T5463709
Score	0.24395452401314383
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

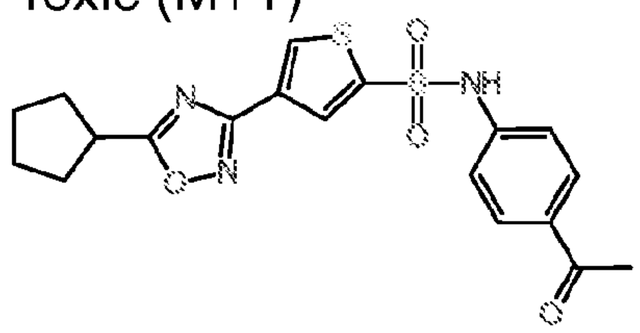


unknown chirality

CatalogID	4052-4279
Score	0.15619544912441768
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

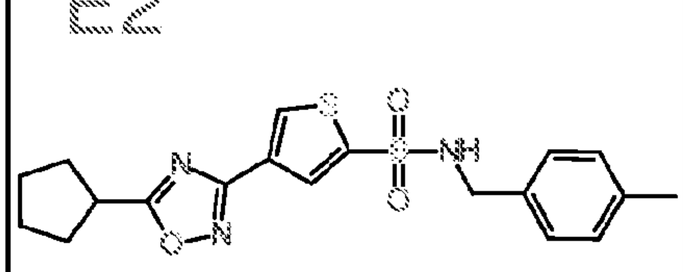
Family E

Toxic (MTT)



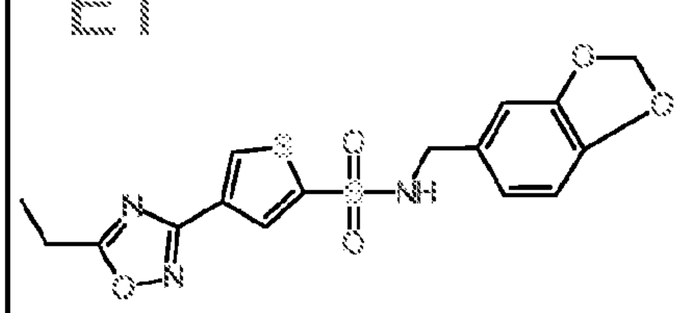
CatalogID	L287-1577
Score	1.1941965522070632
Score_SNARF	1.059135232742917
Score_LACTATE	1.2154260492545208

E2



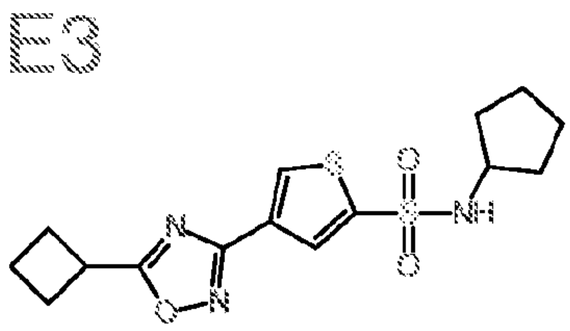
CatalogID	L287-1641
Score	0.7717661404373028
Score_SNARF	0.6259648254476997
Score_LACTATE	0.8035139544488019

E1



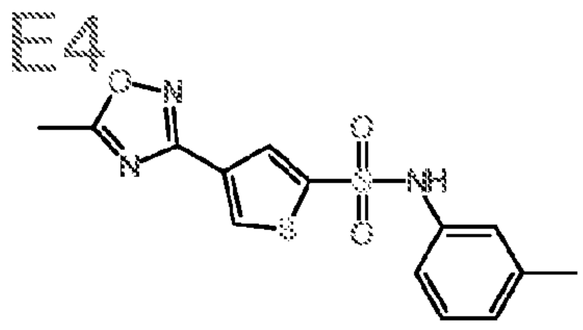
CatalogID	L287-0468
Score	0.597113045524908
Score_SNARF	0.6592317135195043
Score_LACTATE	0.6179394415979561

E3

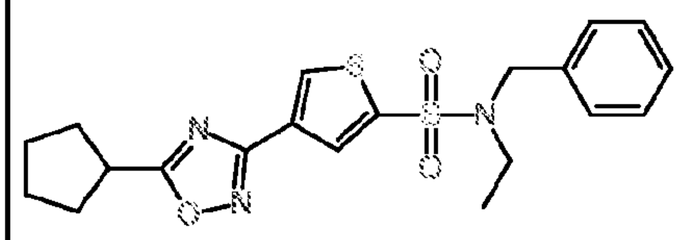


CatalogID	L287-1221
Score	0.5155681934872474
Score_SNARF	0.5994953570205764
Score_LACTATE	0.6601796155644594

E4



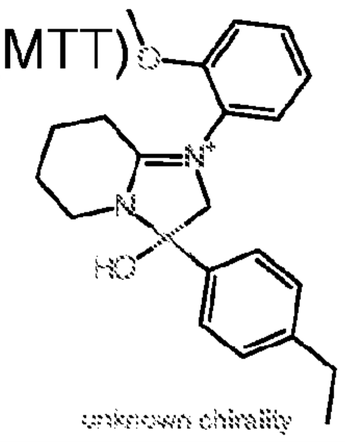
CatalogID	L287-0220
Score	0.35245757621656926
Score_SNARF	0.5828763746073296
Score_LACTATE	1.5779605457379224



CatalogID	L287-1758
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

Family F (1/3)

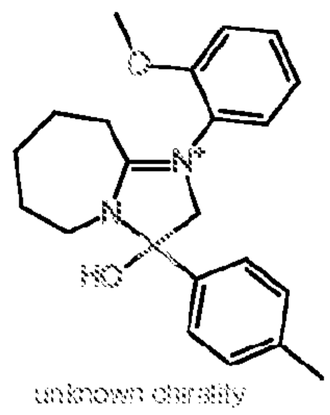
Toxic (MTT)



unknown chirality

CatalogID	K404-0672
Score	1.245294619968429
Score_SNARF	1.0137401503371428
Score_LACTATE	1.3564152530836169

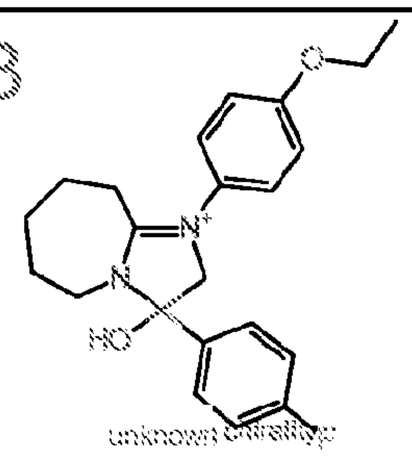
F1



unknown chirality

CatalogID	K404-0834
Score	1.1228184799775751
Score_SNARF	1.022137061560947
Score_LACTATE	1.6126572223635596

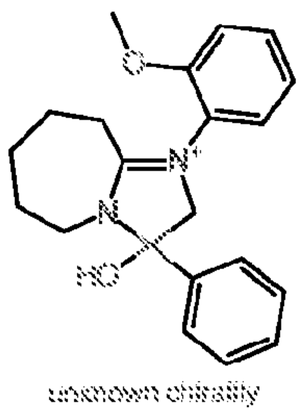
F3



unknown chirality

CatalogID	K404-0885
Score	1.0632020143035092
Score_SNARF	0.9078708580769379
Score_LACTATE	0.9014739788461116

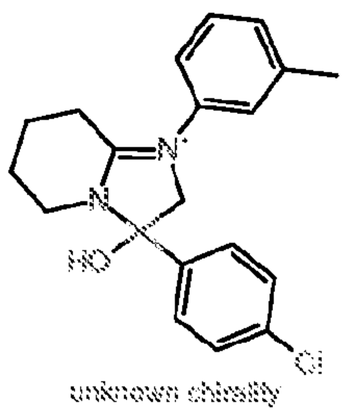
F2



unknown chirality

CatalogID	K404-0838
Score	1.030964579539889
Score_SNARF	0.8328339622087815
Score_LACTATE	1.0509377415150052

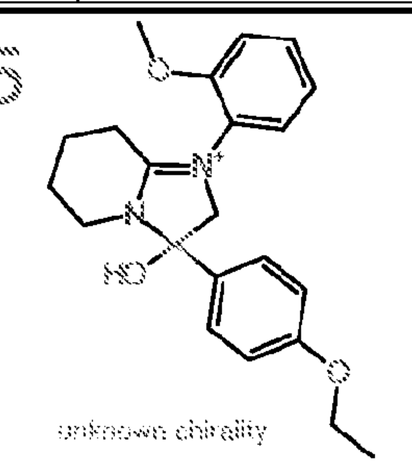
F4



unknown chirality

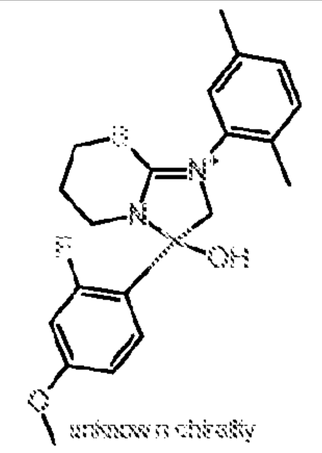
CatalogID	K404-0800
Score	0.9729596471903071
Score_SNARF	0.7385730422972991
Score_LACTATE	0.8385058046355538

F5



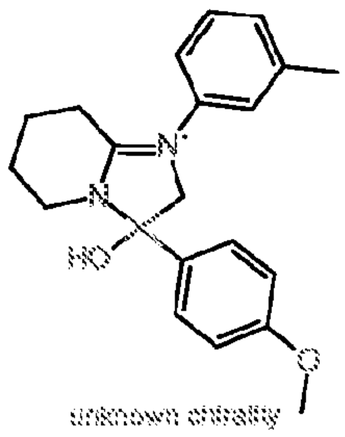
unknown chirality

CatalogID	K404-0673
Score	0.9305928701278892
Score_SNARF	0.975128679661051
Score_LACTATE	2.0243685786220746



unknown chirality

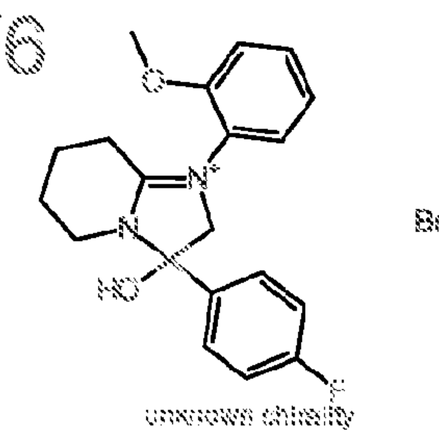
CatalogID	K404-0183
Score	0.895586999693623
Score_SNARF	0
Score_LACTATE	0



unknown chirality

CatalogID	K404-0796
Score	0.8606816213733115
Score_SNARF	0.6402071193972154
Score_LACTATE	0

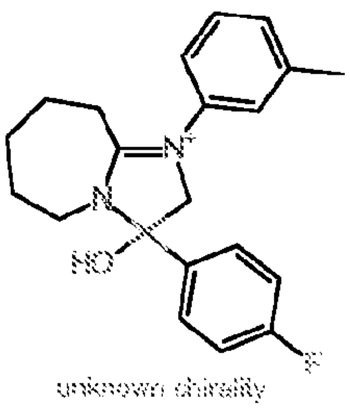
F6



unknown chirality

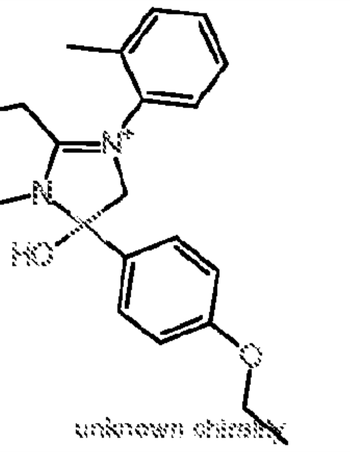
CatalogID	F0524-0338
Score	0.7360202432760756
Score_SNARF	0.6339637469346097
Score_LACTATE	1.584755646134777

F7



unknown chirality

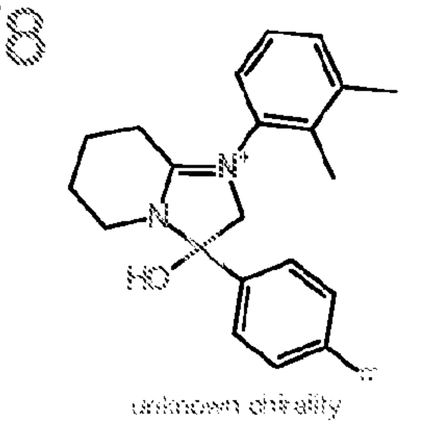
CatalogID	K404-0910
Score	0.7255494073967776
Score_SNARF	0.587837672209965
Score_LACTATE	0.6219124056361738



unknown chirality

CatalogID	F0524-0511
Score	0.7004940551078669
Score_SNARF	0.5555152736849276
Score_LACTATE	0

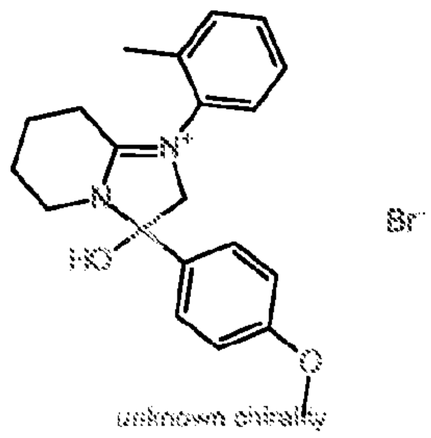
F8



unknown chirality

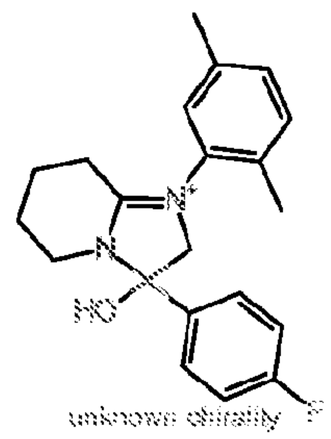
CatalogID	K404-0685
Score	0.6788640765960137
Score_SNARF	0.6561775100211843
Score_LACTATE	0.6821438644674629

Family F (23)



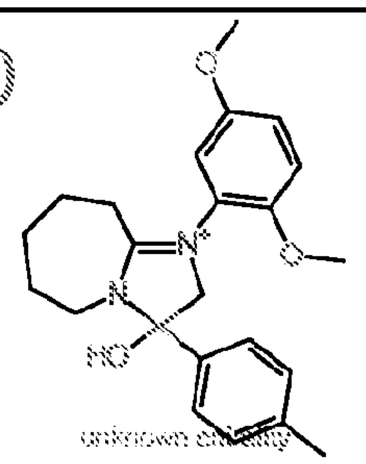
CatalogID	F0524-0507
Score	0.6120319085702075
Score_SNARF	0.4490842880759446
Score_LACTATE	0

F9



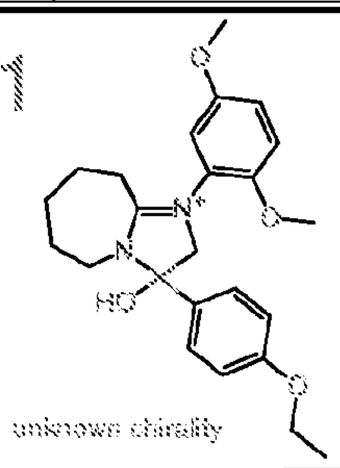
CatalogID	K404-0697
Score	0.5955558493686706
Score_SNARF	0.5122850686232165
Score_LACTATE	0.6526243981885684

F10

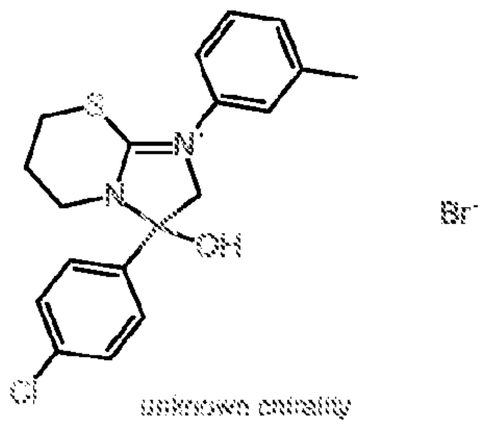


CatalogID	K404-0855
Score	0.5849297608174214
Score_SNARF	0.49441470822619804
Score_LACTATE	0.6276573393012033

F11

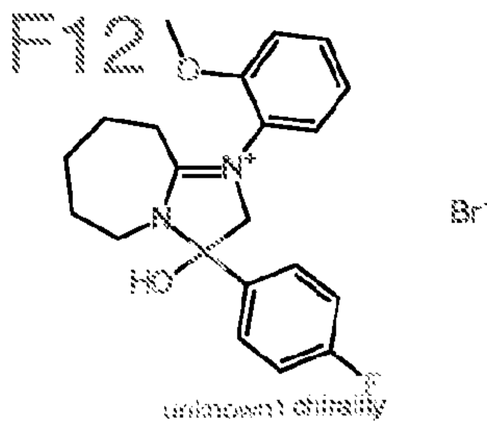


CatalogID	K404-0860
Score	0.5789542930848154
Score_SNARF	0.6137065353631177
Score_LACTATE	0.4765859547890614

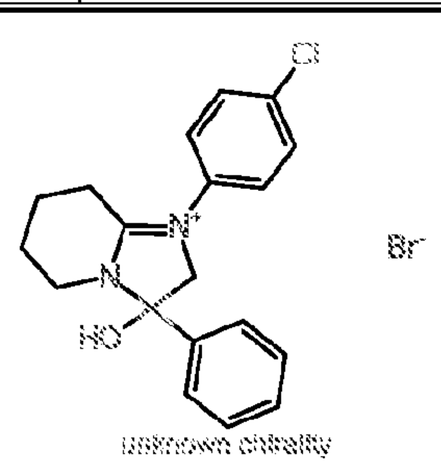


CatalogID	F0522-0533
Score	0.5572373039563334
Score_SNARF	0.47052230558046815
Score_LACTATE	0

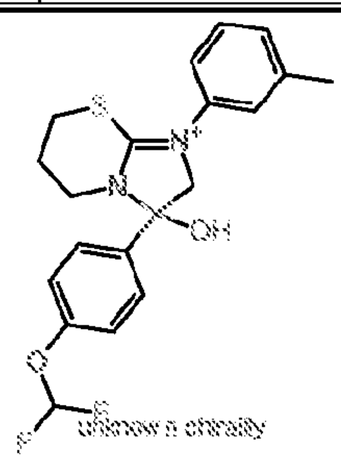
F12



CatalogID	F0524-0611
Score	0.5339163037048325
Score_SNARF	0.5953907345776432
Score_LACTATE	1.5467390894846487

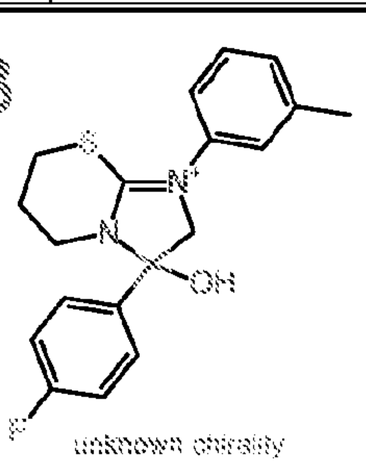


CatalogID	F0524-0488
Score	0.5117725163099112
Score_SNARF	0.3488588317771778
Score_LACTATE	0

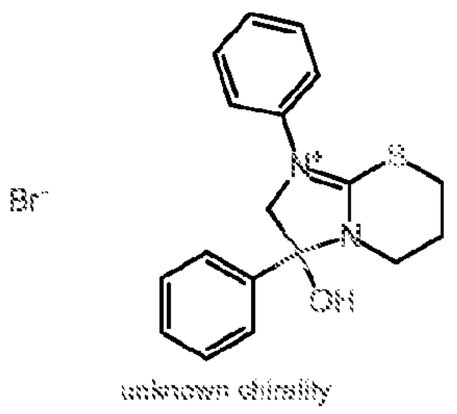


CatalogID	K404-0400
Score	0.3634628173863156
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

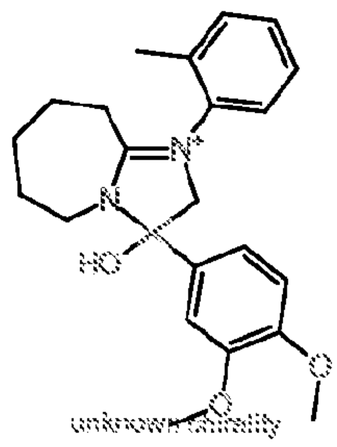
F13



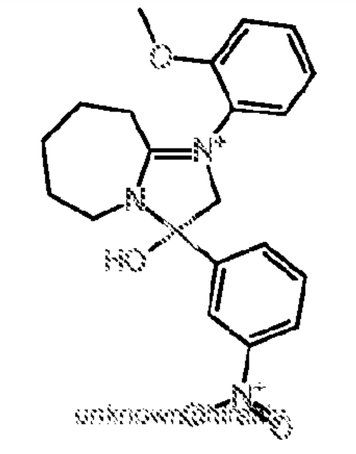
CatalogID	K404-0394
Score	0.3467068112662994
Score_SNARF	0.3308952459455954
Score_LACTATE	0.4428535839996976



CatalogID	T0507-8442
Score	0.26840498230024956
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

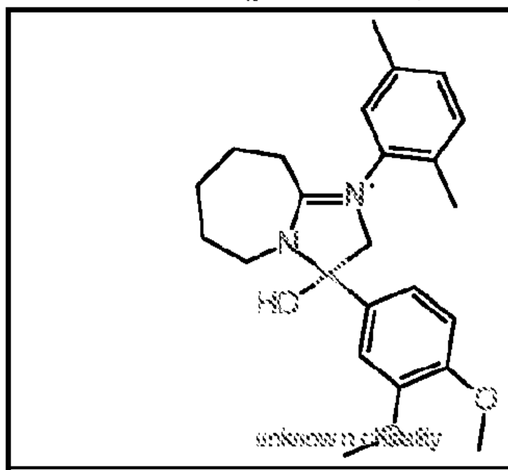


CatalogID	K404-0906
Score	0.1913767711510388
Score_SNARF	0
Score_LACTATE	0

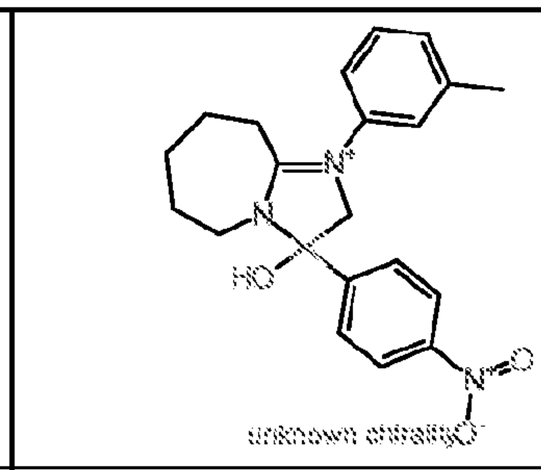


CatalogID	K404-0842
Score	0.169031494172405
Score_SNARF	0.2021699692940943
Score_LACTATE	0

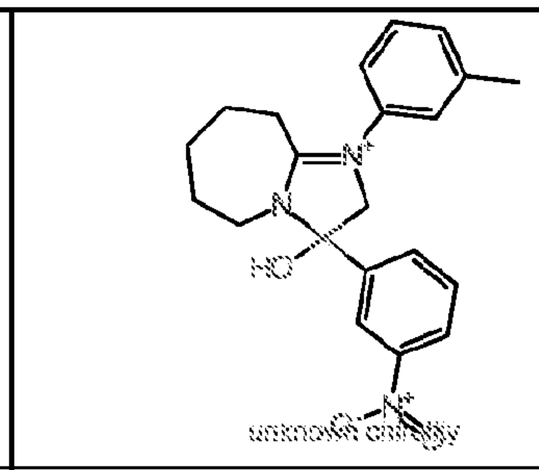
Family F (3/3)



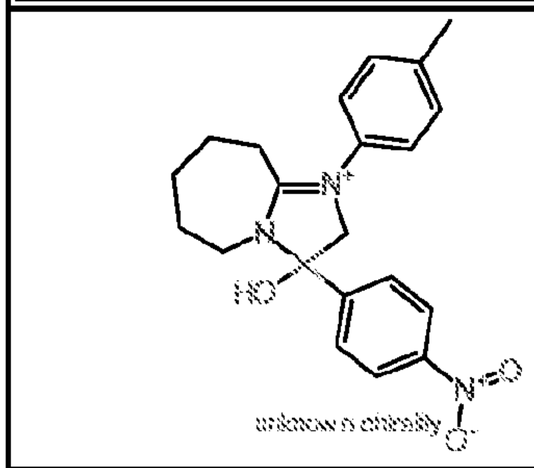
CatalogID	K404-0852
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



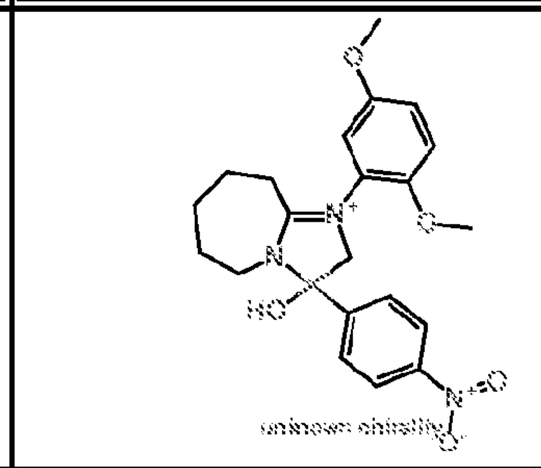
CatalogID	K404-0914
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



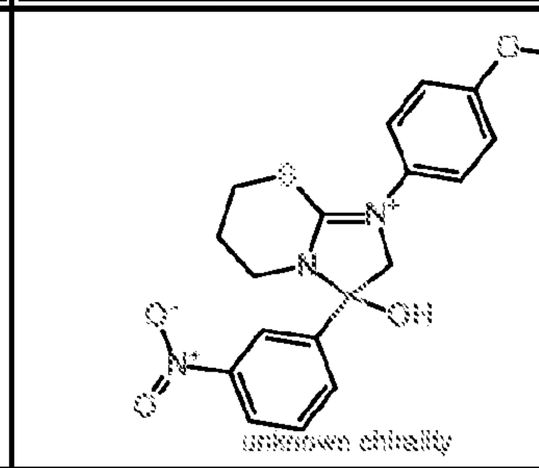
CatalogID	K404-0915
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



CatalogID	K404-0828
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



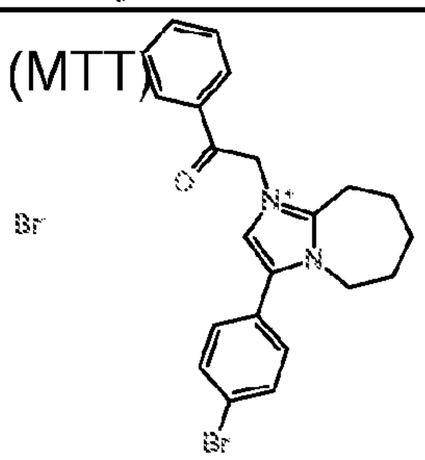
CatalogID	K404-0863
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

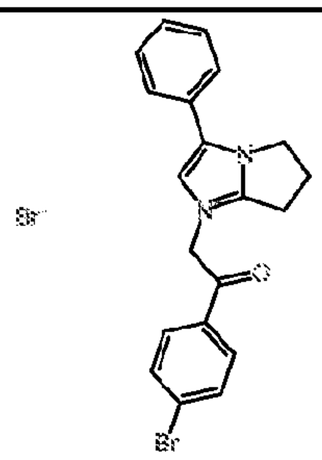


CatalogID	K404-0277
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

Family F2

Toxic (MTT)



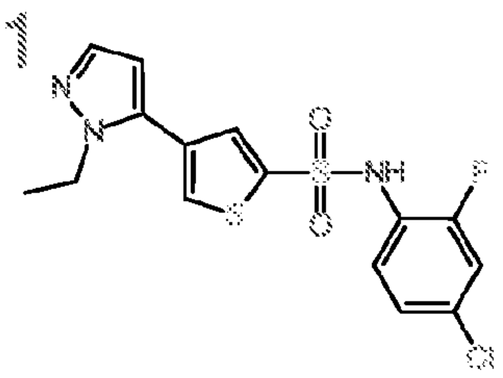


CatalogID	T0500-4648
Score	1.0945456348515448
Score_SNARF	1.0336425682807298
Score_LACTATE	1.1162396727491835

CatalogID	T0502-5560
Score	1.0186741404078639
Score_SNARF	0.8504454491757103
Score_LACTATE	0.8523597142194843

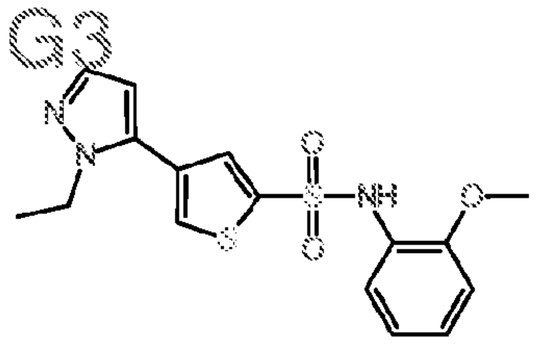
Family G

G1



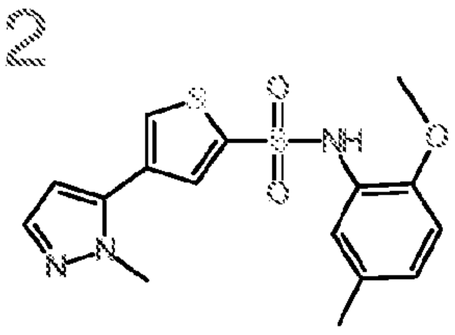
CatalogID	L924-1031
Score	0.9285626587383733
Score_SNARF	0.8713278858649929
Score_LACTATE	1.6324528424335565

G3



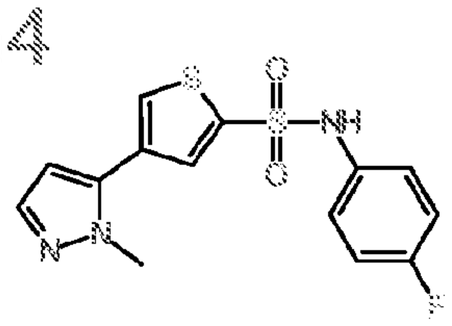
CatalogID	L924-1088
Score	0.7850885946772328
Score_SNARF	0.8834624954983981
Score_LACTATE	0.9105778321422937

G2



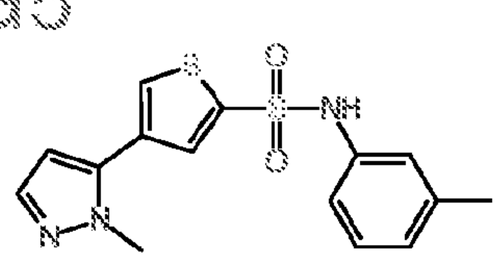
CatalogID	L924-0830
Score	0.7820290835785534
Score_SNARF	0.7153389708932881
Score_LACTATE	0.9168484167457057

G4



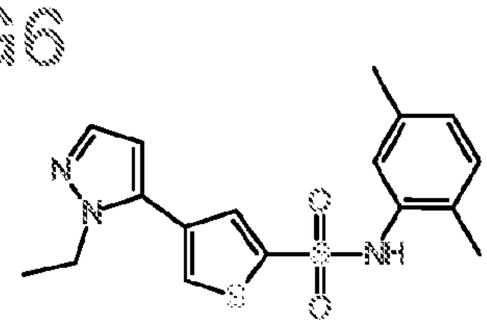
CatalogID	L924-0760
Score	0.6913326730640292
Score_SNARF	0.7695123014835685
Score_LACTATE	0.7898333850310828

G5



CatalogID	L924-0884
Score	0.6560790709954556
Score_SNARF	0.7507266569597015
Score_LACTATE	0.7362417013630416

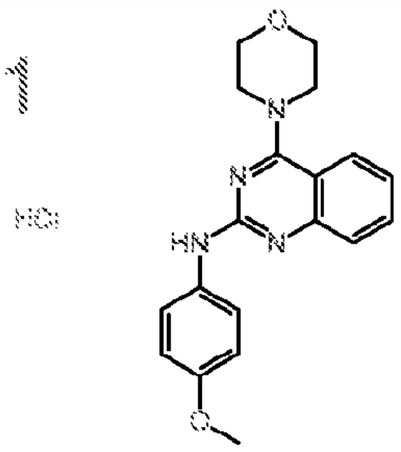
G6

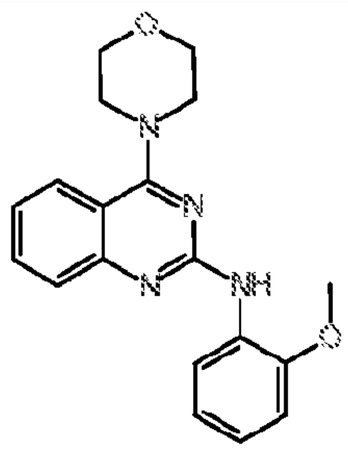


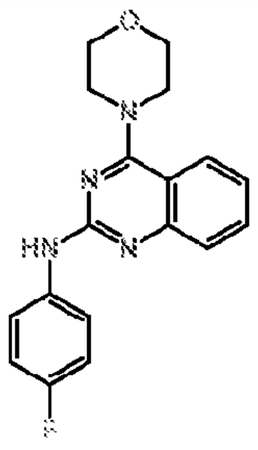
CatalogID	L924-0988
Score	0.5266337896361102
Score_SNARF	0.6651843383172913
Score_LACTATE	0.6138869900753379

Family H

H1







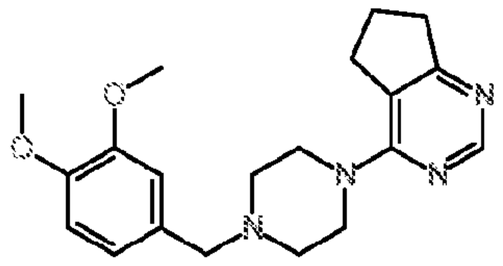
CatalogID	T0508-5190
Score	0.4587604202347154
Score_SNARF	0.4029122435484707
Score_LACTATE	0.2892472208157204

CatalogID	F3007-0009
Score	0.19840697203688884
Score_SNARF	0
Score_LACTATE	0

CatalogID	T0508-5193
Score	0.19512447407576727
Score_SNARF	0.30521182534409874
Score_LACTATE	0

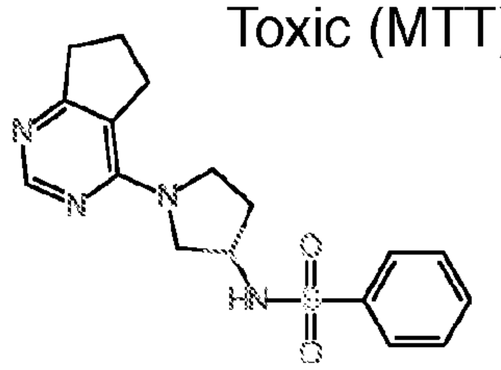
Family 1 (1/2)

Toxic (MTT)



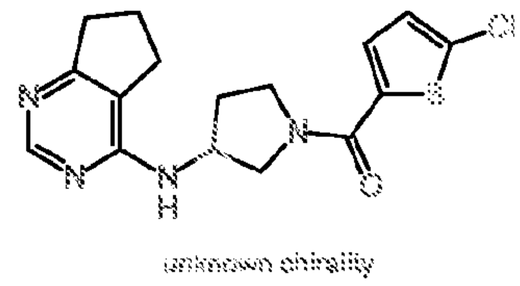
CatalogID	T636-2007
Score	1.0151791427998305
Score_SNARF	0.8595986424585913
Score_LACTATE	0.9841180645962743

Toxic (MTT)



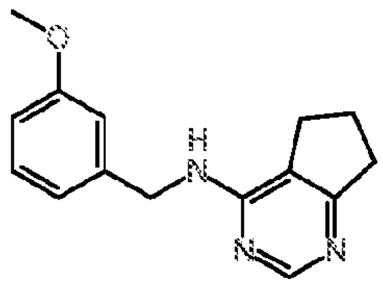
CatalogID	T636-1250
Score	0.9644615645907111
Score_SNARF	0.8587018300764222
Score_LACTATE	0.8982224440823987

Toxic (MTT)



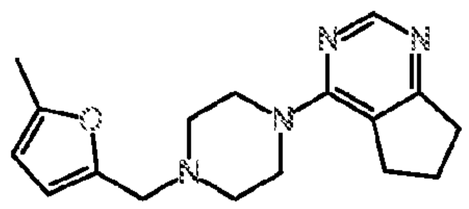
CatalogID	T636-2391
Score	0.932345345636482
Score_SNARF	0.8743350166506818
Score_LACTATE	0.8854443852559006

14



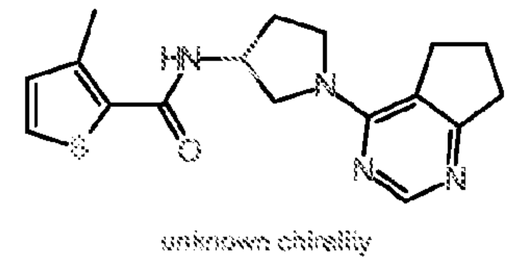
CatalogID	T636-0134
Score	0.7513378916885404
Score_SNARF	0
Score_LACTATE	0.3283315307869653

11

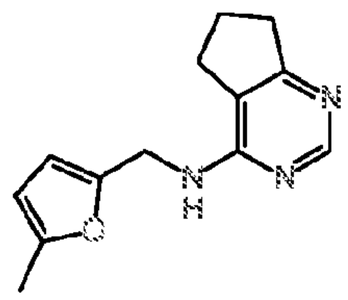


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Score_LACTATE	0.8044660300257542

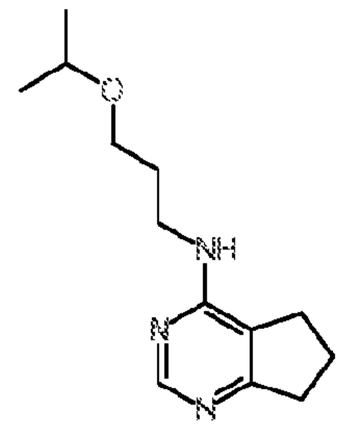
12



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Score_LACTATE	0.547728388970824

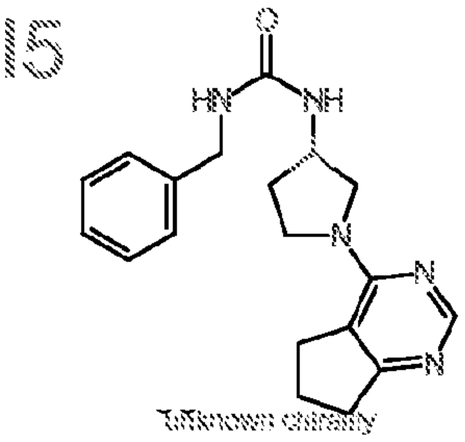


CatalogID	T636-0054
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Score_LACTATE	0

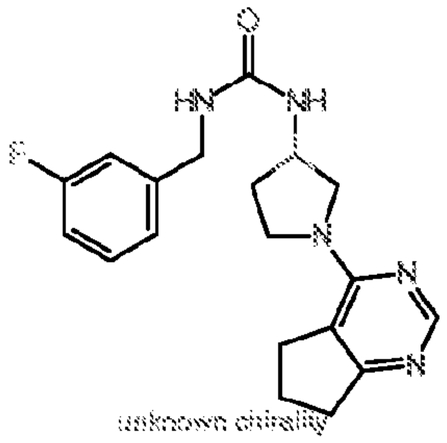


CatalogID	T636-0027
Score	0.5361875692026127
Score_SNARF	0
Score_LACTATE	0

15

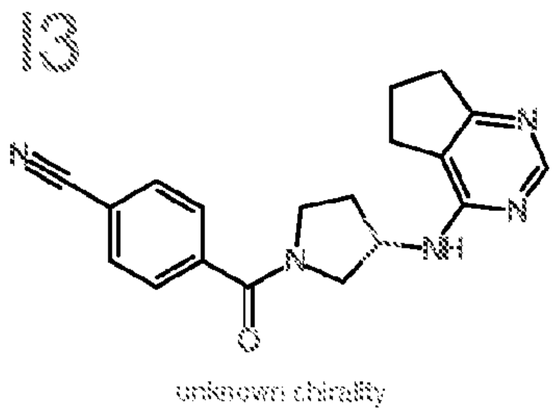


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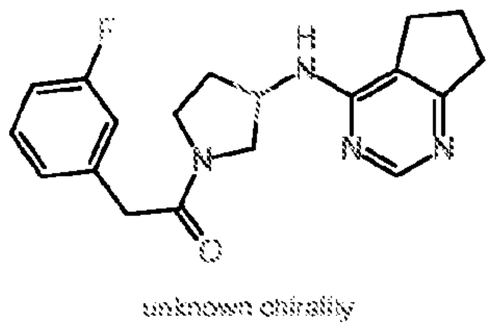


CatalogID	T636-1243
Score	0.45504840873980434
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Score_LACTATE	0

13

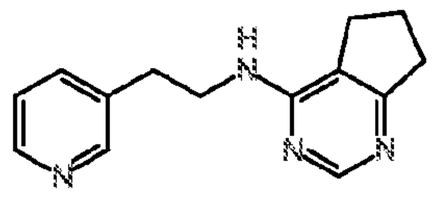


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Score_SNARF	0.4808881204258762
Score_LACTATE	0.3005490442971268

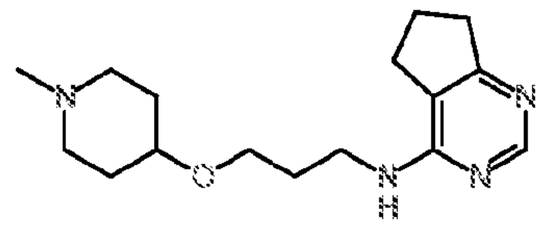


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Score_LACTATE	0.41303393247609077

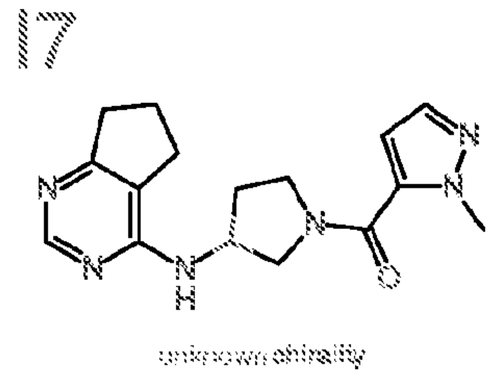
Family 1 (2/2)



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Score_LACTATE	NOT_TESTED

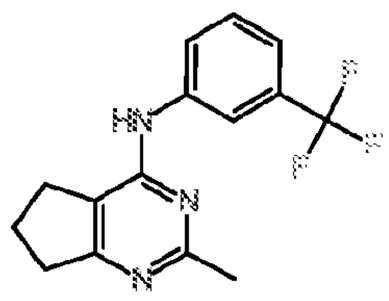


CatalogID	T636-0181
Score	0.22771221504714034
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Score_LACTATE	NOT_TESTED

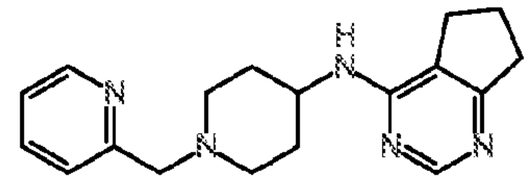


unknown chirality

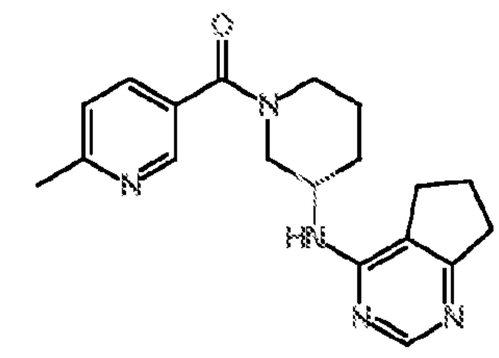
CatalogID	T636-2425
Score	0.21958972496943524
Score_SNARF	0.27247555941127144
Score_LACTATE	0.30269809289243466



CatalogID	D278-0514
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

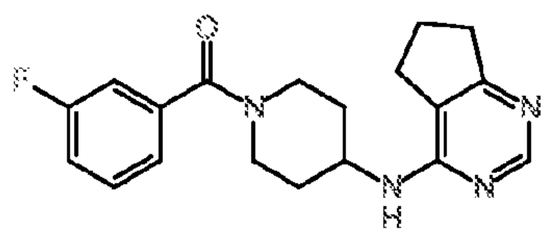


CatalogID	T636-1715
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Score_LACTATE	NOT_TESTED

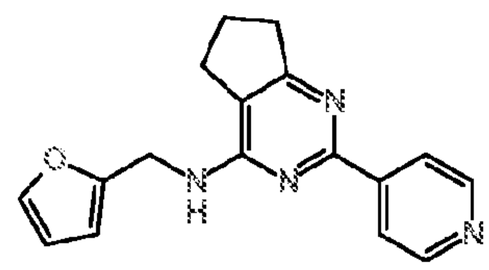


unknown chirality

CatalogID	T636-2144
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Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



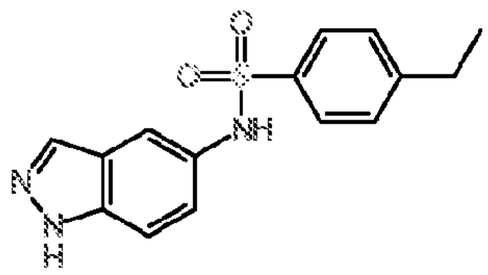
CatalogID	T636-1601
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



CatalogID	T636-0973
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

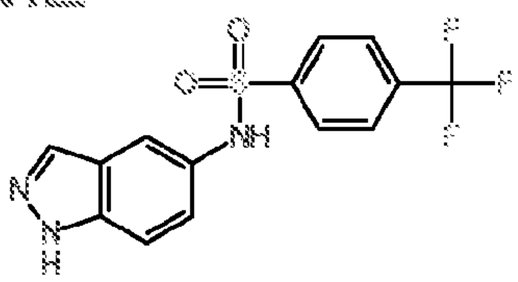
Family M

M1

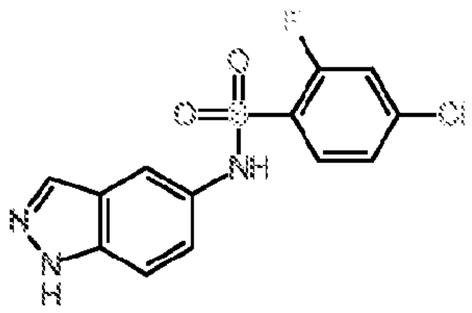


CatalogID	T5599014
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Score_SNARF	0.5778347174521993
Score_LACTATE	0.4418517946231513

M2



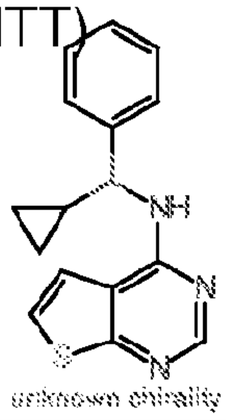
CatalogID	T5653029
Score	0.36218450009714054
Score_SNARF	0.5641600427647684
Score_LACTATE	1.3535979987810212



CatalogID	T5436375
Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

Family N

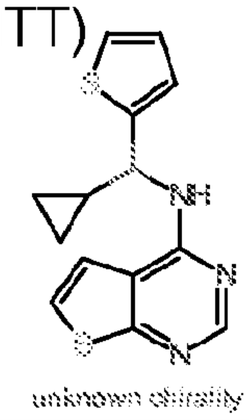
Toxic (MTT)



unknown chirality

CatalogID	T6016487
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Score_SNARF	1.081418939812701
Score_LACTATE	1.8565016432738268

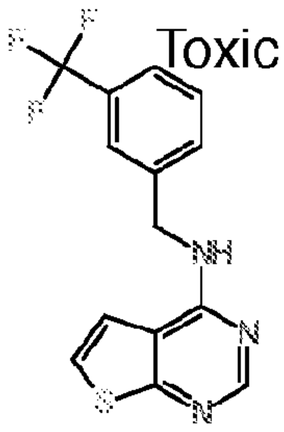
Toxic (MTT)



unknown chirality

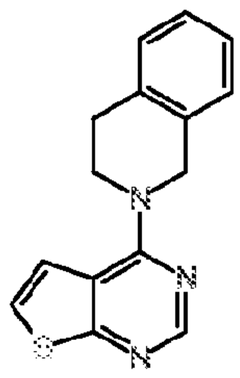
CatalogID	T5713637
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Score_SNARF	1.1838450634294277
Score_LACTATE	1.287213975075479

Toxic (MTT)



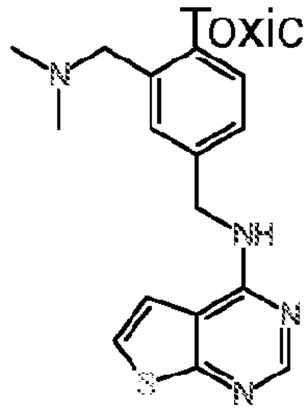
CatalogID	T5346988
Score	0.9960475361592487
Score_SNARF	0.9569371376034257
Score_LACTATE	0.9326889609464317

N1



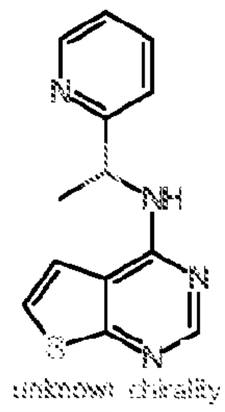
CatalogID	T0517-8250
Score	0.9574785194936402
Score_SNARF	0.9123967008705844
Score_LACTATE	0.9269003205038194

Toxic (MTT)



CatalogID	T6783542
Score	0.8569906585878225
Score_SNARF	0.9183439213520057
Score_LACTATE	1.0325028610945488

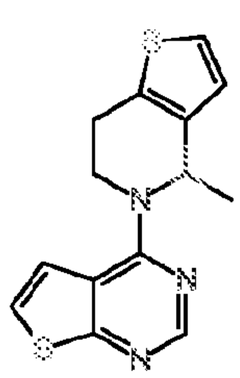
N2



unknown chirality

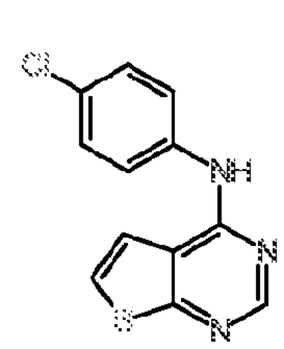
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Score_SNARF	0.7508314360875484
Score_LACTATE	0.5882588561328497

N3

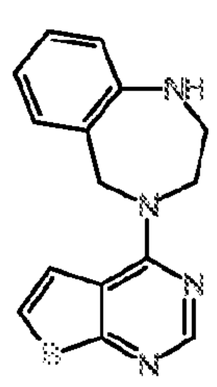


unknown chirality

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Score_SNARF	0.48158494627623916
Score_LACTATE	0.6999954284930354



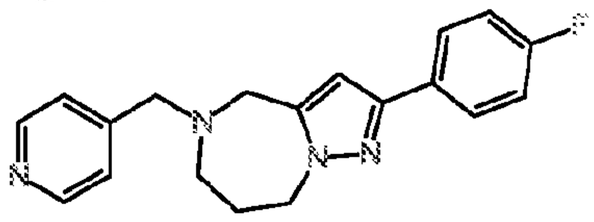
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Score	0.3693019544327658
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



CatalogID	T6768300
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Score_SNARF	0.3741744712882059
Score_LACTATE	0

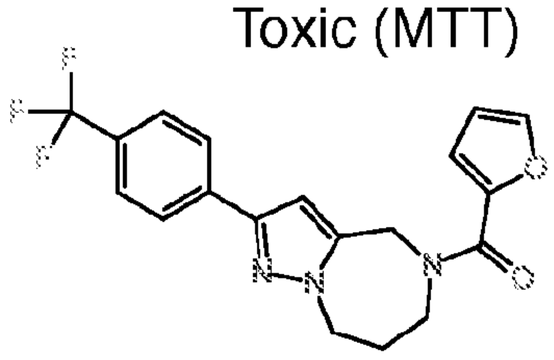
Family O

O1

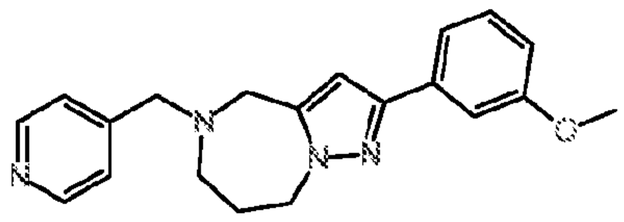


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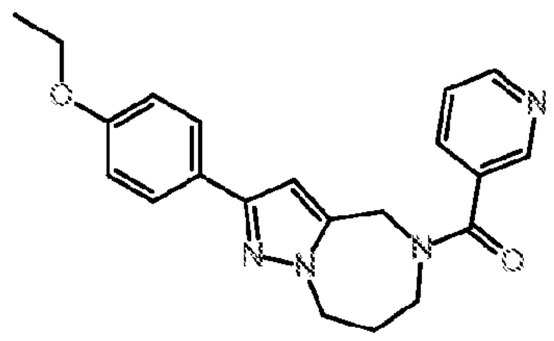
Toxic (MTT)



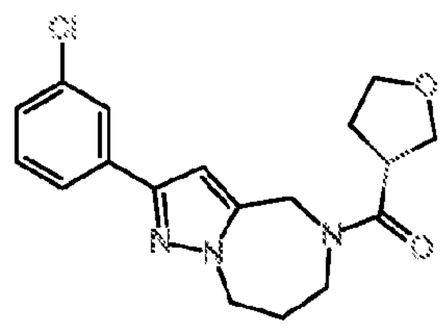
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Score	0.5618497687392737
Score_SNARF	0.38039779657169775
Score_LACTATE	0.2896301510809021



CatalogID	T202-0973
Score	0.34101692441028175
Score_SNARF	0.2834303332326549
Score_LACTATE	0



CatalogID	T200-0707
Score	0.16740550045889108
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

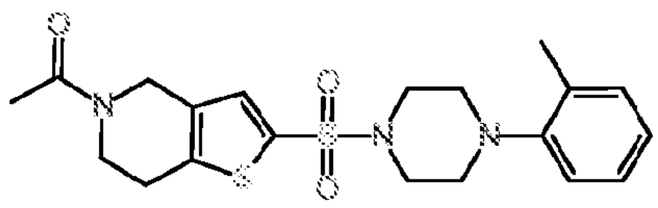


unknown chirality

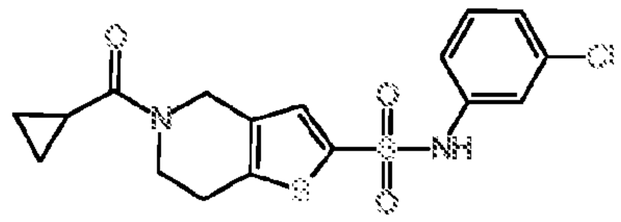
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Score	0
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

Family P

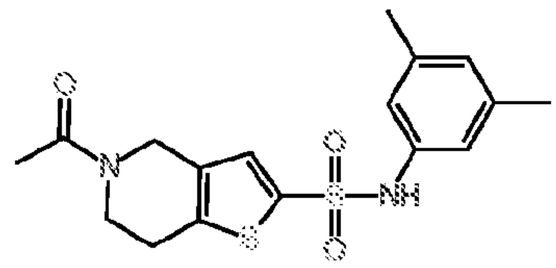
P1



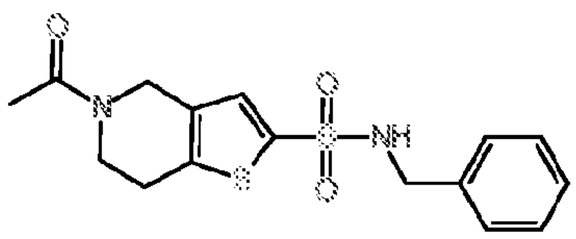
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Score_SNARF	0.8006744658424207
Score_LACTATE	0.9231665502291709



CatalogID	P025-0462
Score	0.3872119888600468
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



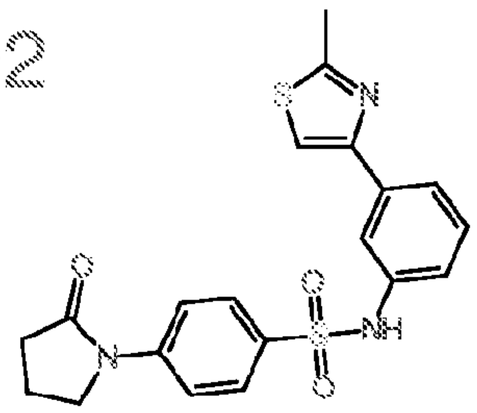
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Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



CatalogID	P025-0168
Score	0.14492617861885423
Score_SNARF	NOT_TESTED
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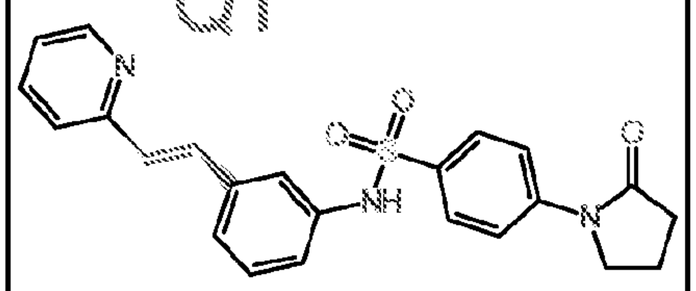
Family Q

Q2



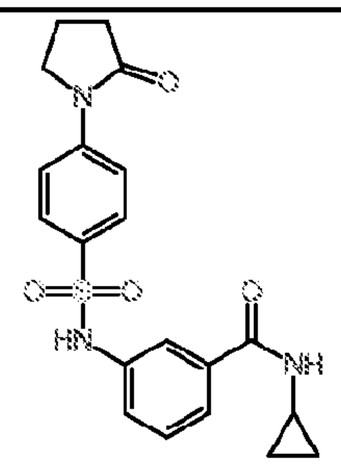
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Score	0.855937067832691
Score_SNARF	0.8856396558188464
Score_LACTATE	0.7495521570358064

Q1



CatalogID	T5644989
Score	0.6796130774342963
Score_SNARF	0.8385482212594828
Score_LACTATE	0.7858882841885826

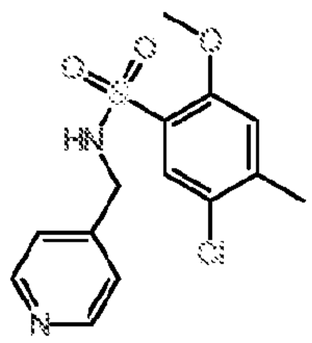
Q3



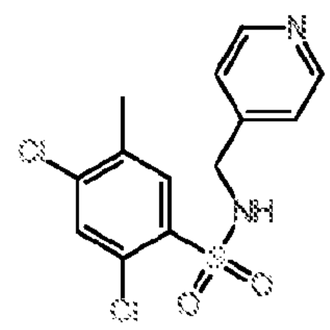
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Score	0.41563542866343284
Score_SNARF	0.5352874527303951
Score_LACTATE	0.3601184130676996

Family R

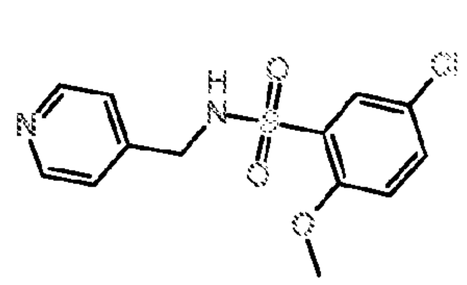
R1



CatalogID	T5580243
Score	0.8528253990291386
Score_SNARF	0.8564967067233966
Score_LACTATE	0.7229025303296649



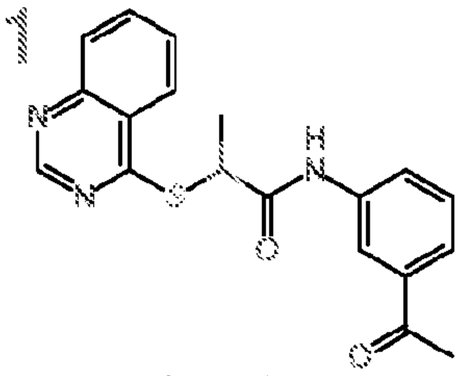
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Score_SNARF	0.5050374641230523
Score_LACTATE	0



CatalogID	F0376-0203
Score	0.20288513101448935
Score_SNARF	0
Score_LACTATE	0

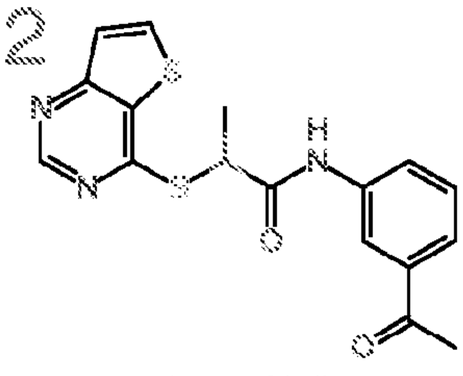
Family S

S1



unknown chirality

S2



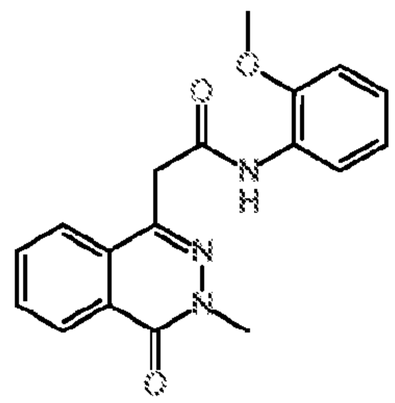
unknown chirality

CatalogID	T0511-9200
Score	0.8441832214748136
Score_SNARF	0.48656473560837266
Score_LACTATE	0.7517887234187022

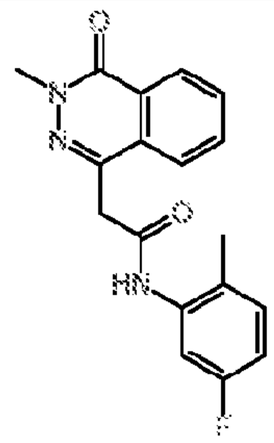
CatalogID	T5627721
Score	0.7237159041769383
Score_SNARF	0.7384840988075632
Score_LACTATE	0.9486502359872533

Family T

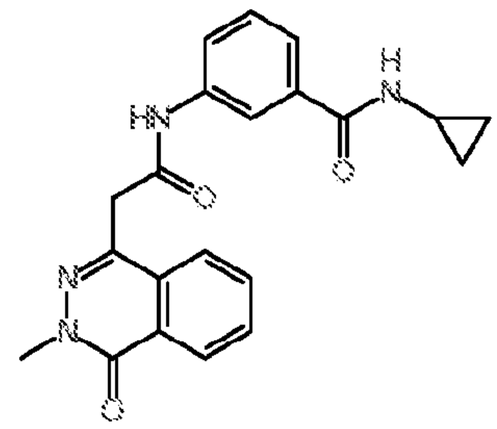
T1



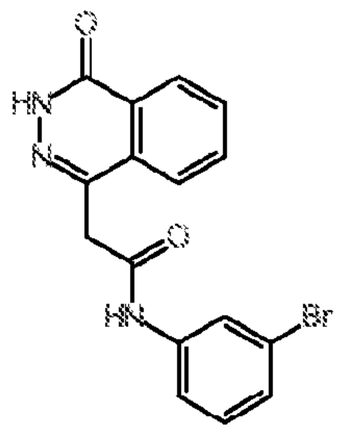
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Score	0.722280122898277
Score_SNARF	0.7568885849430411
Score_LACTATE	0.9280741657657858



CatalogID	T6302989
Score	0.4051145241844445
Score_SNARF	0.407229804897944
Score_LACTATE	0

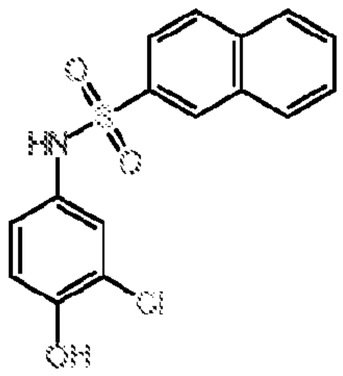


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Score_SNARF	NOT_TESTED
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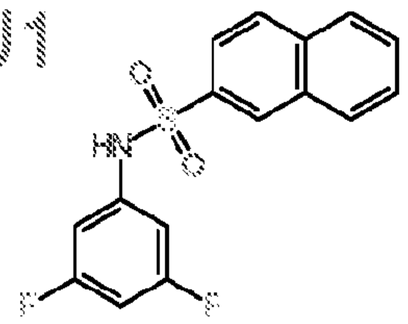
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Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

Family U



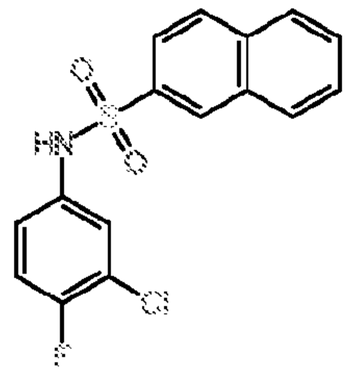
CatalogID	T0517-8939
Score	0.7679530086775413
Score_SNARF	0.6381895919510732
Score_LACTATE	0

U1



CatalogID	T5884038
Score	0.7207565908025609
Score_SNARF	0.8379087709023765
Score_LACTATE	0.9262350987060564

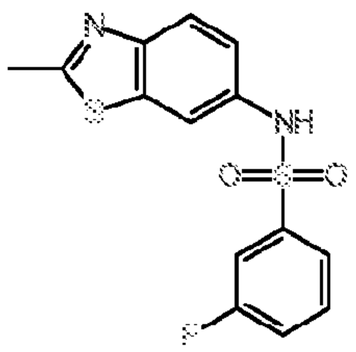
U2



CatalogID	T5231424
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Score_SNARF	0.3706614739849921
Score_LACTATE	0.5258096899997711

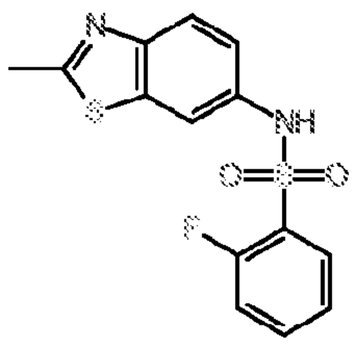
Family V

V1

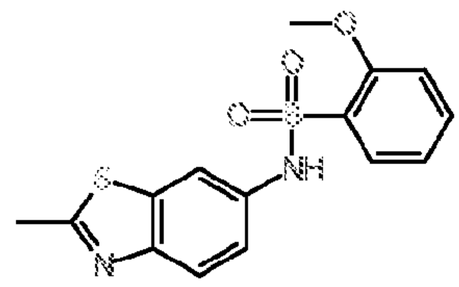


CatalogID	T6937001
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Score_SNARF	0.855659897786533
Score_LACTATE	0.9087796548938434

V2



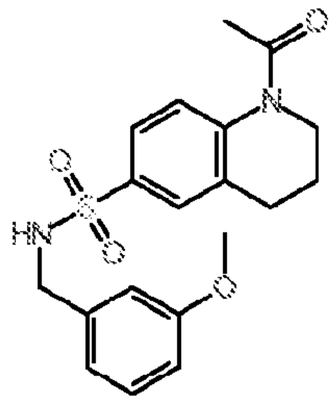
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Score_SNARF	0.5544595058134321
Score_LACTATE	0.3824503515573746



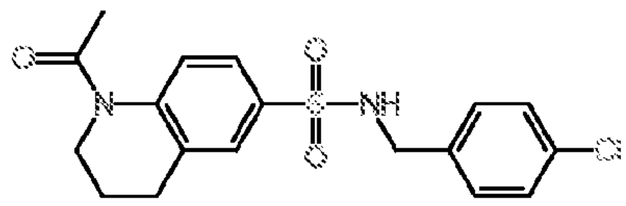
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Score	0.2408425006175835
Score_SNARF	0.24545703685650536
Score_LACTATE	0

Family W

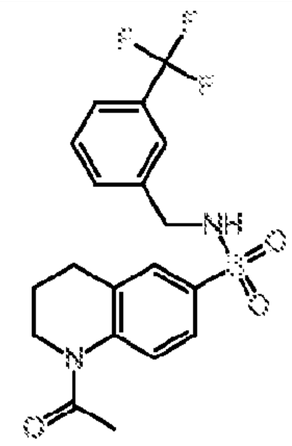
W1



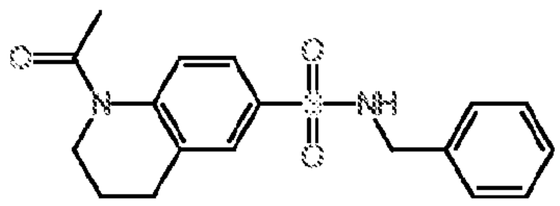
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Score_SNARF	0.2788913341775572
Score_LACTATE	0.45168048543941974



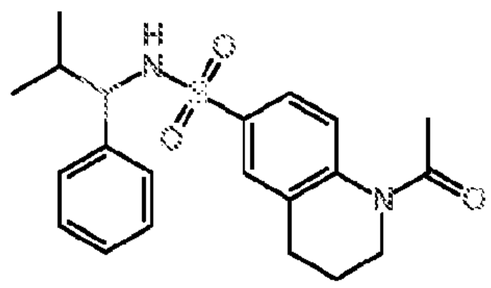
CatalogID	G855-0033
Score	0.34741470465778507
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED



CatalogID	T6099519
Score	0.2755618623360358
Score_SNARF	0.3943862632059517 7
Score_LACTATE	0



CatalogID	G855-0210
Score	0.20620606338380704
Score_SNARF	0.31231048770176095
Score_LACTATE	0

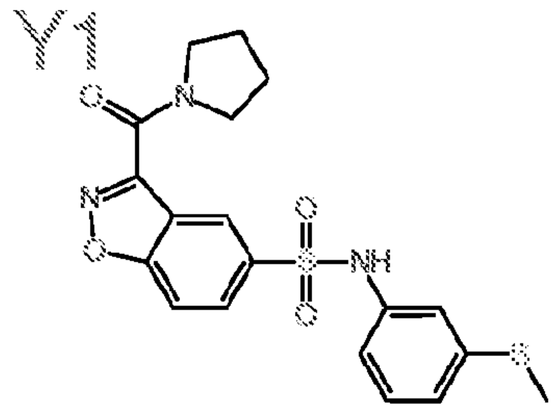


unknown chirality

CatalogID	T6067366
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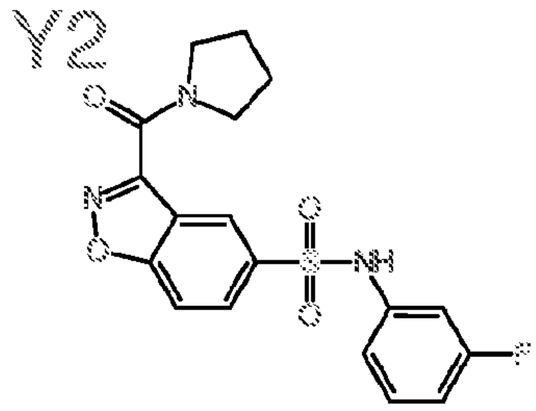
Family Y

Y1

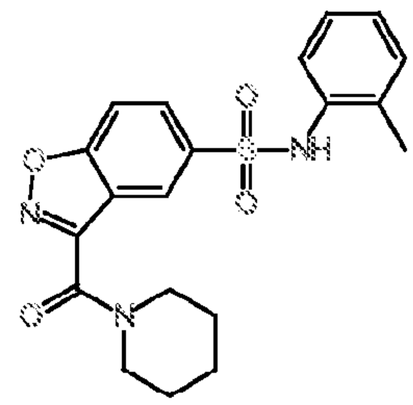


CatalogID	L995-0125
Score	0.8544527003879917
Score_SNARF	0.8016477664825784
Score_LACTATE	0.9122179865041484

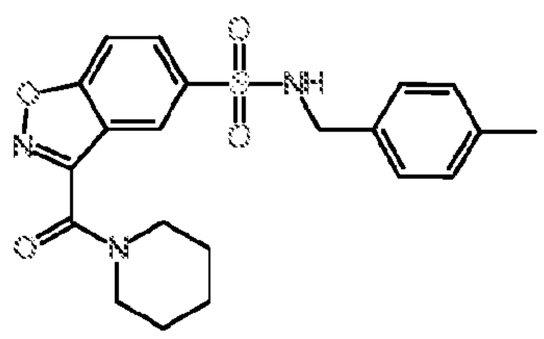
Y2



CatalogID	L995-0058
Score	0.42257261140984437
Score_SNARF	0.4197252825194825
Score_LACTATE	0.4240067586981341

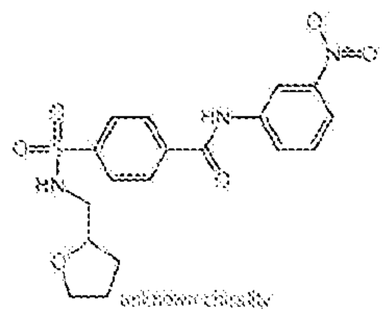


CatalogID	L995-0405
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Score_LACTATE	0

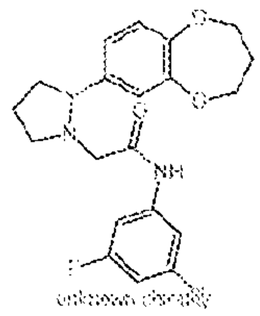


CatalogID	L995-0386
Score	0.18817082101410132
Score_SNARF	NOT_TESTED
Score_LACTATE	NOT_TESTED

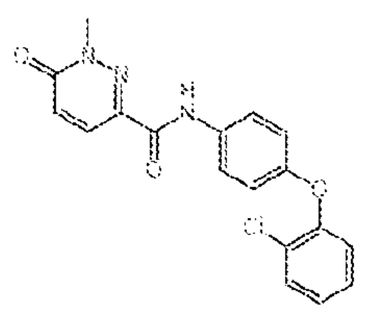
Appendix II



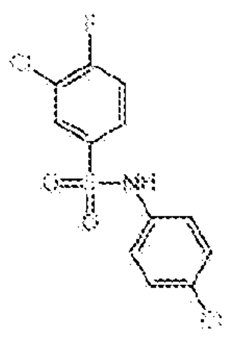
Catalog ID	T6010789
Score Lact.	1.31
EC50	5.82



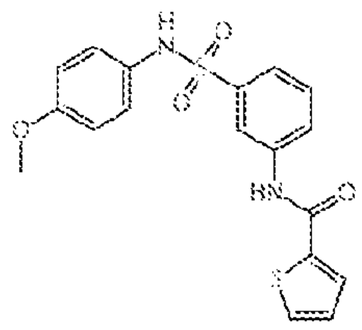
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EC50	8.81



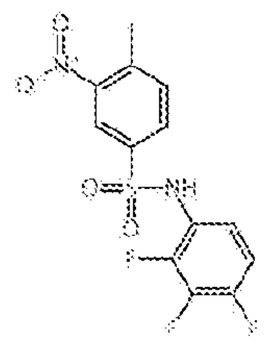
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EC50	4.10



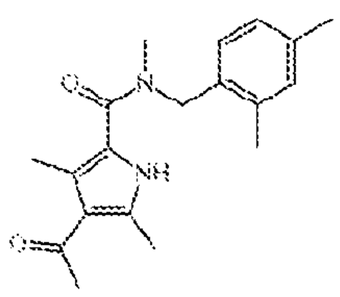
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EC50	5.50



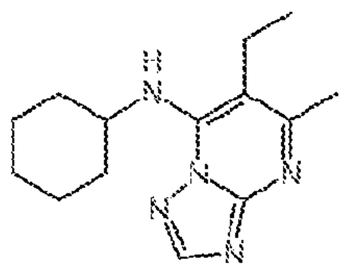
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EC50	5.43



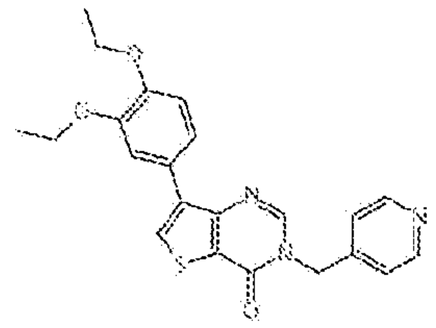
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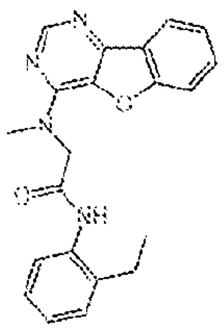
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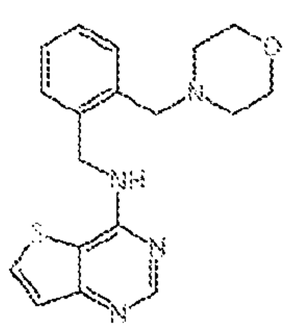
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Score Lact.	0.83
EC50	4.11



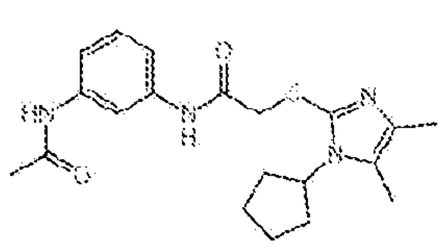
Catalog ID	L115-0403
Score Lact.	0.81
EC50	3.28



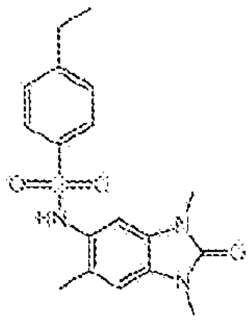
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Score Lact.	0.8
EC50	1.14



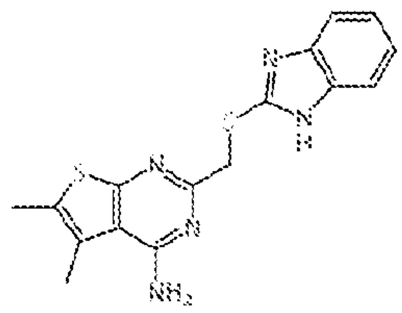
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Score Lact.	0.79
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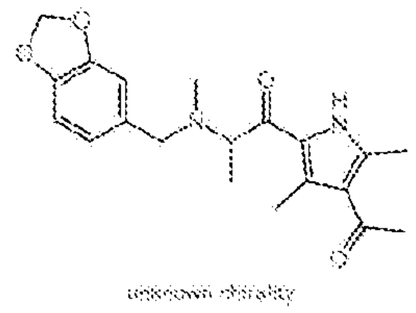
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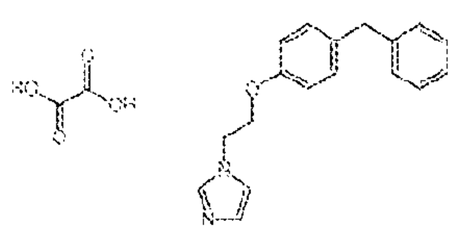
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Score Lact.	0.74
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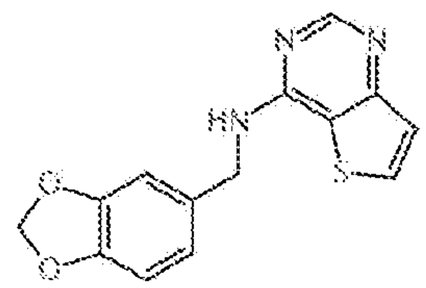
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Score Lact.	0.74
EC50	2.85



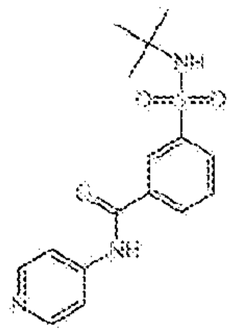
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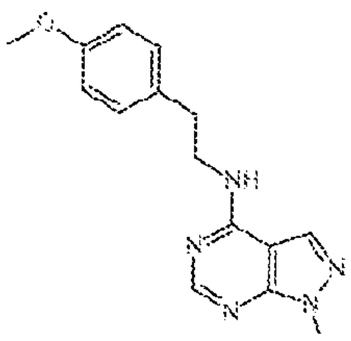
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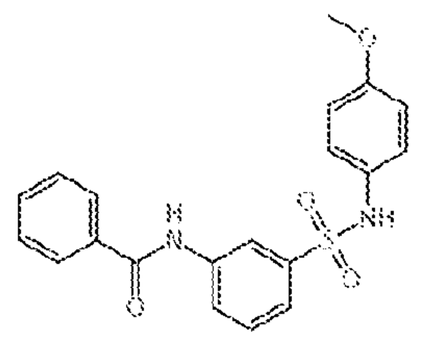
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Score Lact.	0.71
EC50	3.89



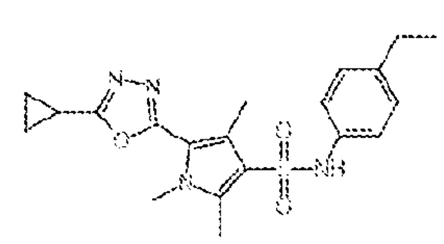
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EC50	5.02



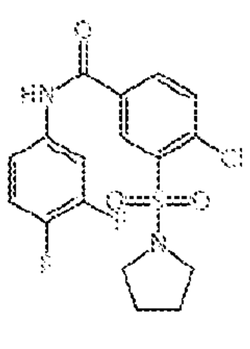
Catalog ID	K405-2595
Score Lact.	0.70
EC50	3.51



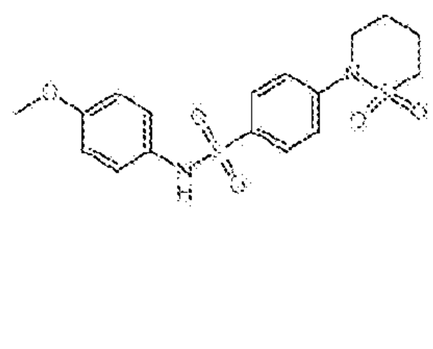
Catalog ID	T5274959
Score Lact.	0.68
EC50	4.12



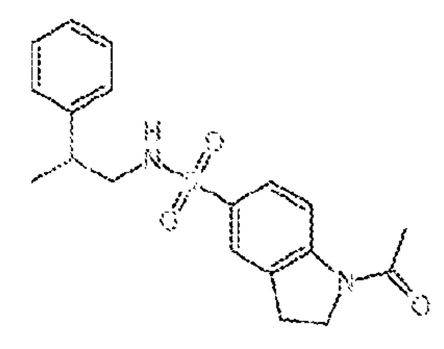
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Score Lact.	0.67
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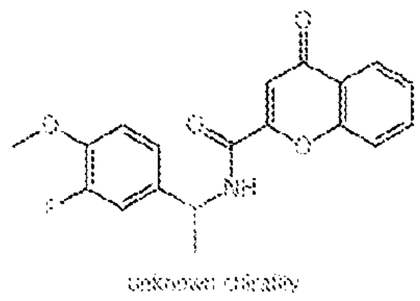
Catalog ID	T5450239
Score Lact.	0.67
EC50	4.42



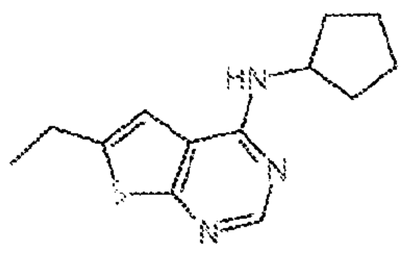
Catalog ID	G508-0015
Score Lact.	0.67
EC50	4.83



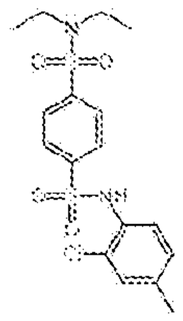
Catalog ID	T5707230
Score Lact.	0.67
EC50	4.88



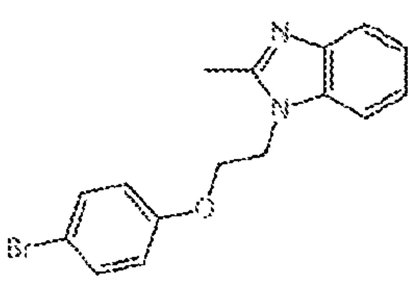
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Score Lact.	0.67
EC50	6.31



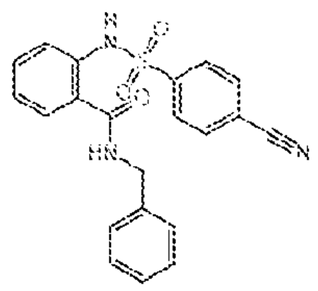
Catalog ID	887-0183
Score Lact.	0.66
EC50	3.94



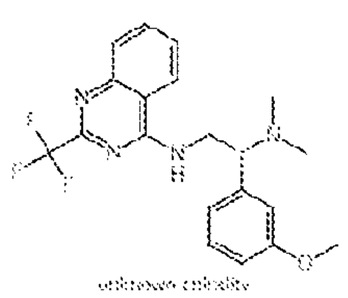
Catalog ID	T5453923
Score Lact.	0.66
EC50	5.36



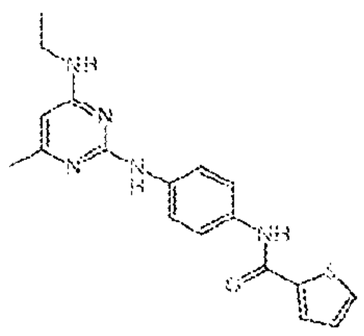
Catalog ID	T0505-4087
Score Lact.	0.64
EC50	7.07



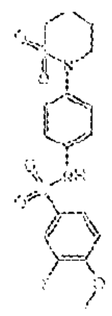
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Score Lact.	0.63
EC50	7.42



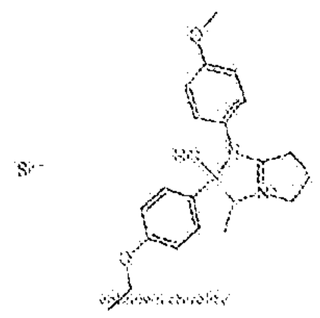
Catalog ID	T5800607
Score Lact.	063
EC50	5.87



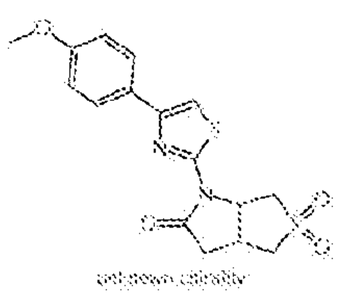
Catalog ID	G869-0071
Score Lact.	0.63
EC50	4.9



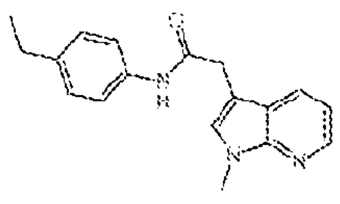
Catalog ID	F2794-0128
Score Lact.	0.62
EC50	9.85



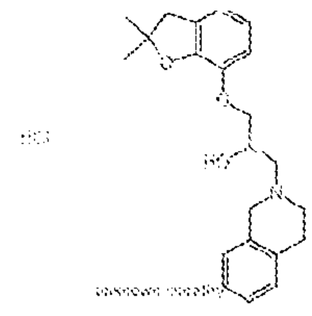
Catalog ID	T0500-6629
Score Lact.	0.61
EC50	2.61



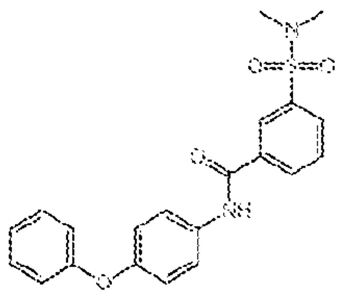
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Score Lact.	0.60
EC50	4.02



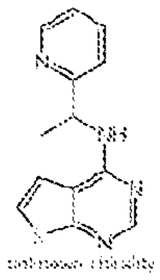
Catalog ID	M508-0370
Score Lact.	0.60
EC50	4.58



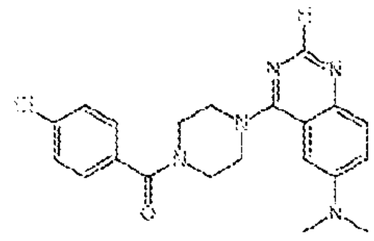
Catalog ID	T0515-1783
Score Lact.	0.59
EC50	5.39



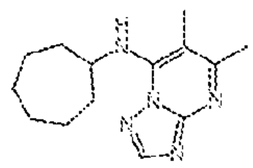
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Score Lact.	0.59
EC50	3.43



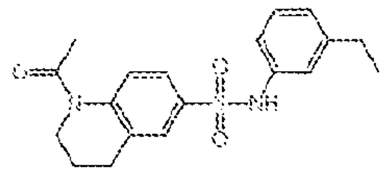
Catalog ID	T5672380
Score Lact.	0.58
EC50	3.91



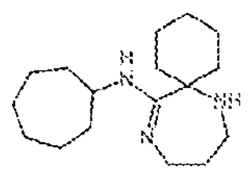
Catalog ID	M381-0730
Score Lact.	0.57
EC50	3.19



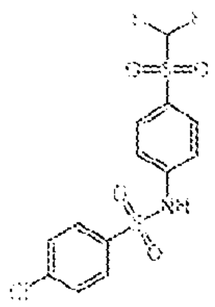
Catalog ID	Z606-8287
Score Lact.	0.57
EC50	4.29



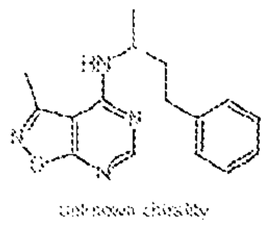
Catalog ID	G855-0143
Score Lact.	0.56
EC50	4.72



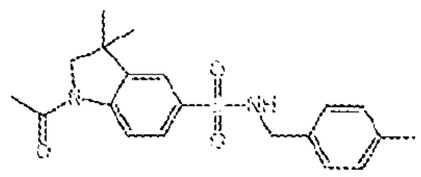
Catalog ID	Z076-0028
Score Lact.	0.56
EC50	8.01



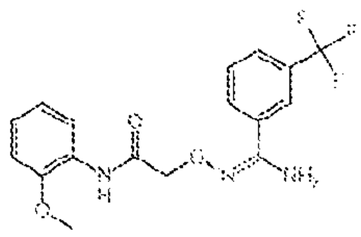
Catalog ID	T5311200
Score Lact.	0.56
EC50	6.02



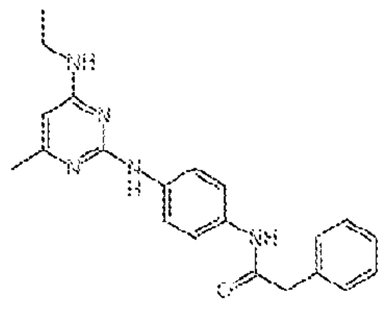
Catalog ID	E944-0182
Score Lact.	0.55
EC50	3.99



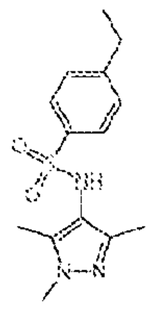
Catalog ID	L302-0069
Score Lact.	0.55
EC50	6.27



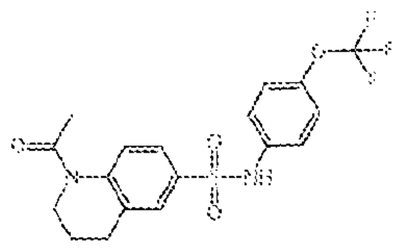
Catalog ID	T5770640
Score Lact.	0.55
EC50	4.86



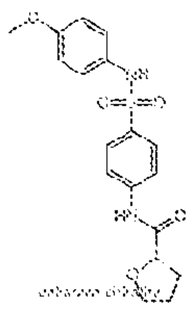
Catalog ID	G869-0064
Score Lact.	0.54
EC50	4.26



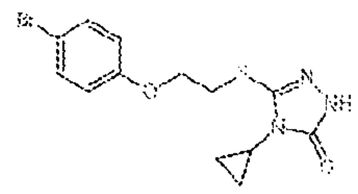
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Score Lact.	0.54
EC50	6.41



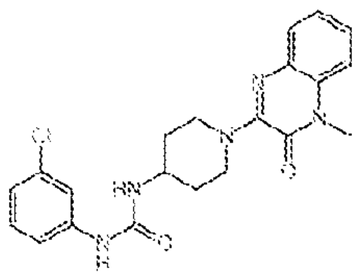
Catalog ID	G855-0183
Score Lact.	0.54
EC50	6.0



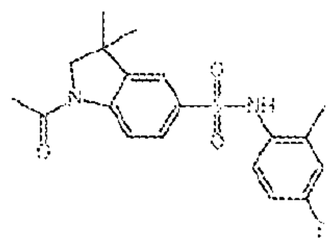
Catalog ID	T5329723
Score Lact.	0.53
EC50	4.61



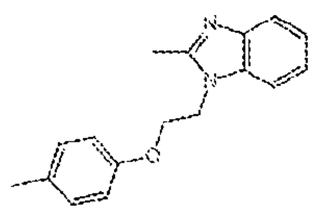
Catalog ID	T533260
Score Lact.	0.53
EC50	6.84



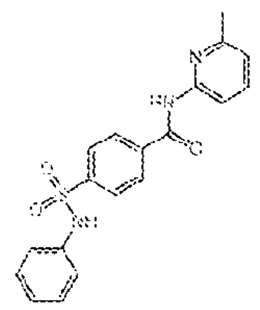
Catalog ID	L932-0267
Score Lact.	0.53
EC50	4.64



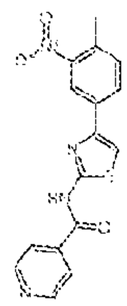
Catalog ID	L302-0181
Score Lact.	0.52
EC50	5.05



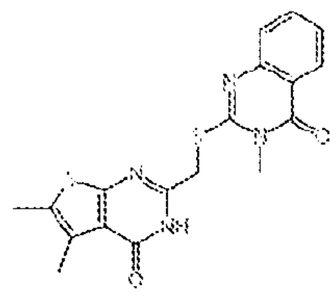
Catalog ID	T5444083
Score Lact.	0.51
EC50	4.59



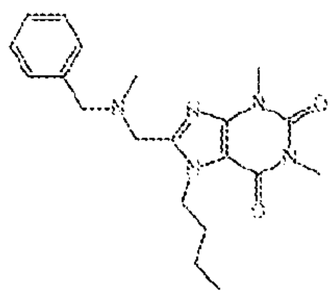
Catalog ID	T6125251
Score Lact.	0.51
EC50	4.56



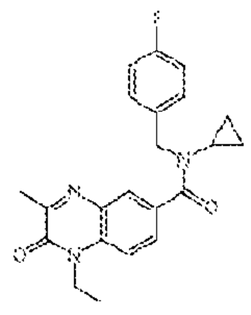
Catalog ID	T5694329
Score Lact.	0.47
EC50	1.59



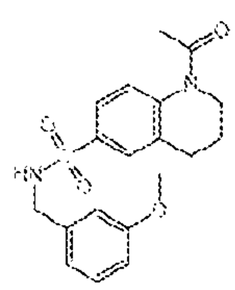
Catalog ID	T0517-2783
Score Lact.	0.46
EC50	3.84



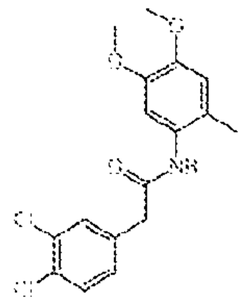
Catalog ID	T5788545
Score Lact.	0.45
EC50	3.59



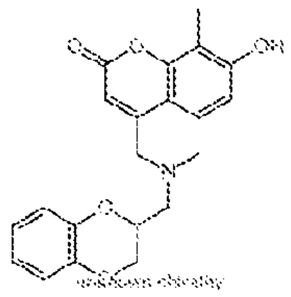
Catalog ID	T5586091
Score Lact.	0.45
EC50	5.36



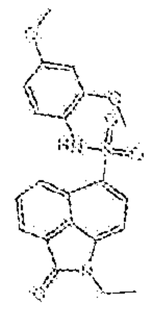
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Score Lact.	0.45
EC50	9.67



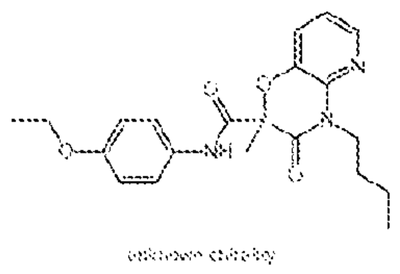
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Score Lact.	0.44
EC50	6.11



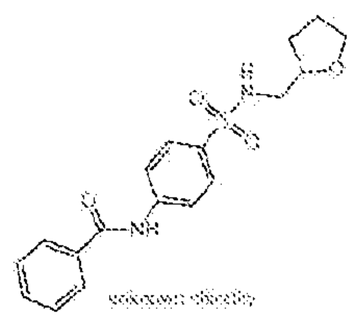
Catalog ID	T5494352
Score Lact.	0.44
EC50	5024



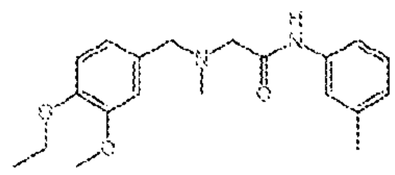
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Score Lact.	0.44
EC50	5.10



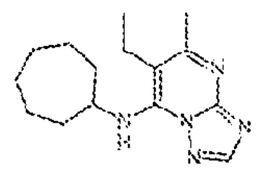
Catalog ID	P621-1364
Score Lact.	0.43
EC50	5.86



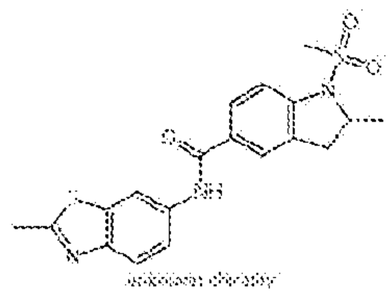
Catalog ID	Y031-0361
Score Lact.	0.43
EC50	8.23



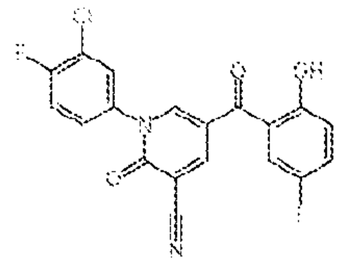
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Score Lact.	0.42
EC50	5.45



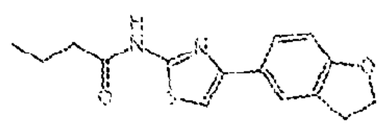
Catalog ID	Z606-8351
Score Lact.	0.42
EC50	5.0



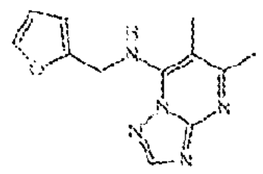
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0.41	0.38
EC50	4.61



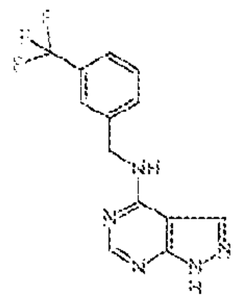
Catalog ID	T0516-6894
Score Lact.	0.38
EC50	8.21



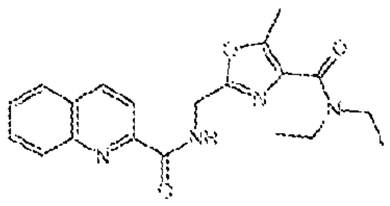
Catalog ID	T5691896
Score Lact.	0.38
EC50	5.21



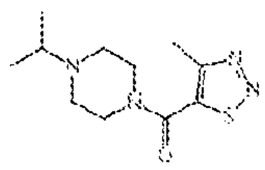
Catalog ID	Z606-8298
Score Lact.	0.38
EC50	4.89



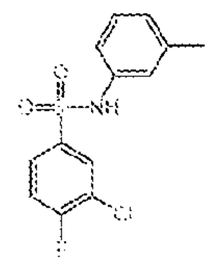
Catalog ID	F5285-0069
Score Lact.	0.36
EC50	6.29



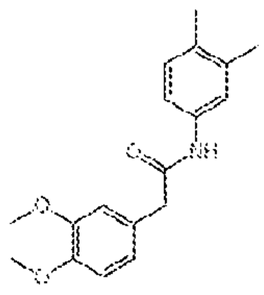
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Score Lact.	0.36
EC50	7.99



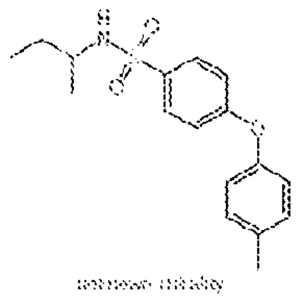
Catalog ID	Z606-5341
Score Lact.	0.36
EC50	ND



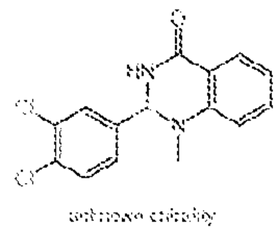
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Score Lact.	0.33
EC50	5.87



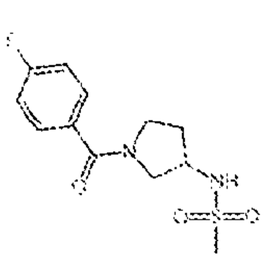
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Score Lact.	0.34
EC50	ND



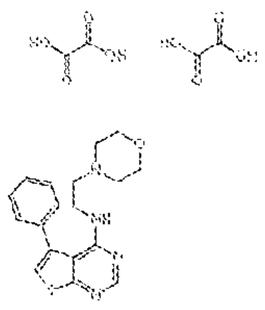
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Score Lact.	0.33
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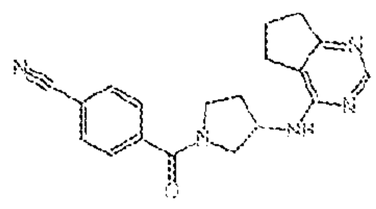
Catalog ID	T5389517
Score Lact.	0.33
EC50	5.74



Catalog ID	Z603-8037
Score Lact.	0.32
EC50	ND



Catalog ID	T0513-0213
Score Lact.	0.30
EC50	3.99



Catalog ID	T636-2387
Score Lact.	0.30
EC50	6.83