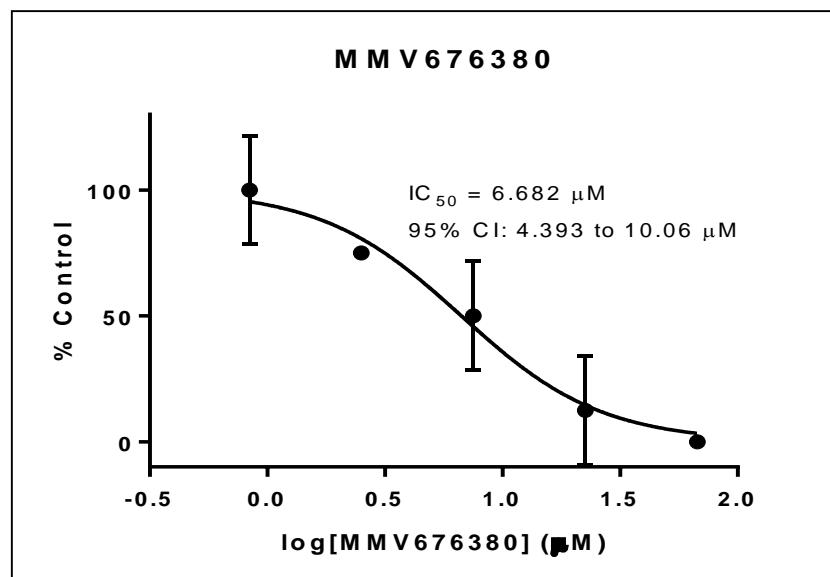
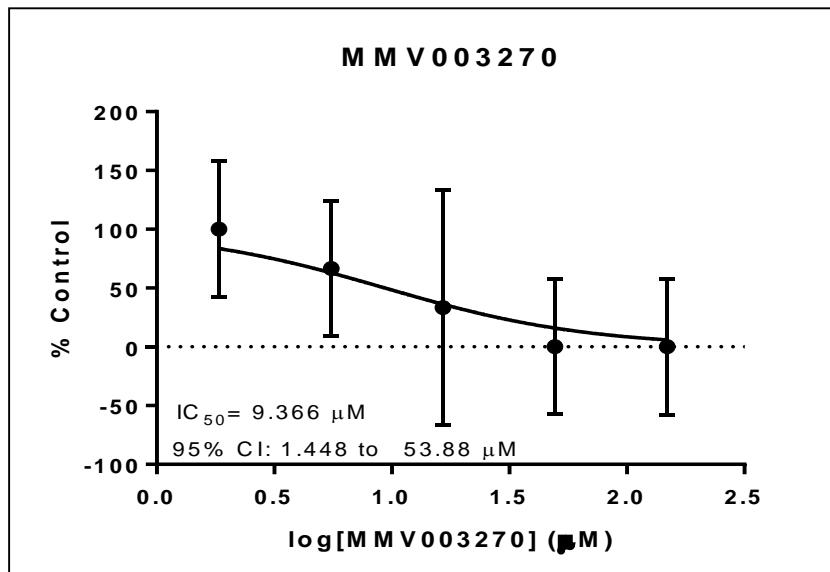


Fig. S1. Dose-response curves (fasciocidal activity) for compounds shown in Table 3.

Data were normalized (% control) as a percentage of the controls (no drug added). Results obtained from triplicated assays.



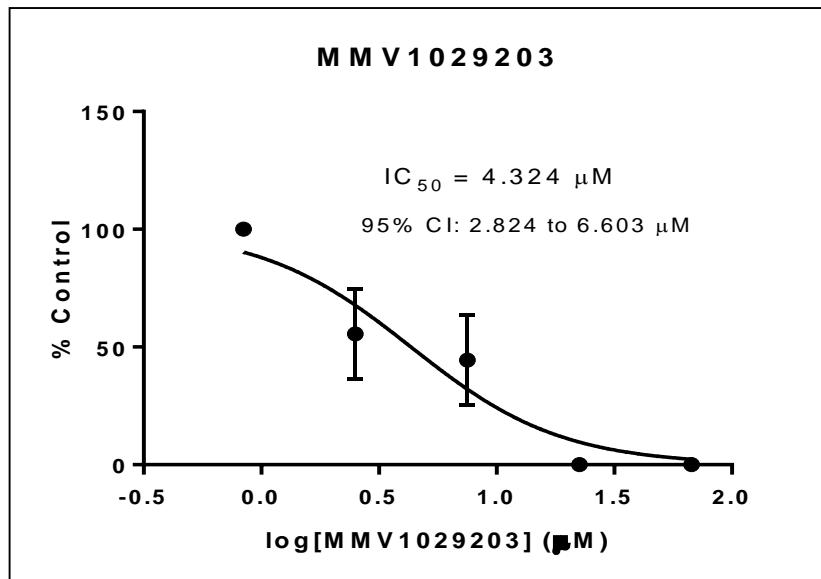
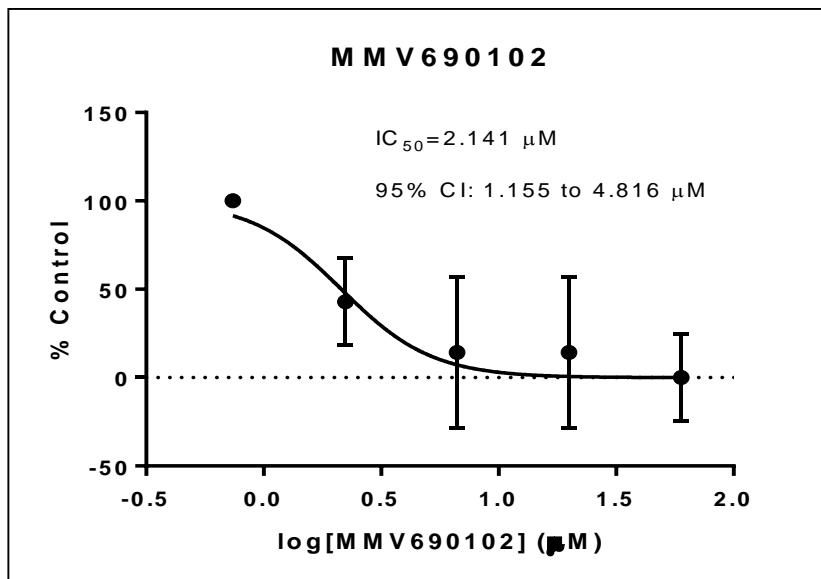


Table S1. Sigmoidal Dose-Response Analysis Data.
The Pathogen Box compounds hit *Fasciola hepatica*.

MMV676380

log(inhibitor) vs. normalized response -- Variable slope

Best-fit values

LogIC50	0.8249
HillSlope	-1.458
IC50	6.682
Std. Error	
LogIC50	0.08178
HillSlope	0.3597
95% CI (profile likelihood)	
LogIC50	0.6428 to 1.003
HillSlope	-2.702 to -0.9179
IC50	4.393 to 10.06
Goodness of Fit	
Degrees of Freedom	13
R square	0.871
Absolute Sum of Squares	3072
Sy.x	15.37

Number of points

# of X values	15
# Y values analyzed	15

MMV1029203

log(inhibitor) vs. normalized response -- Variable slope

Best-fit values

LogIC50	0.6359
HillSlope	-1.36
IC50	4.324
Std. Error	
LogIC50	0.08345
HillSlope	0.3208
95% CI (profile likelihood)	
LogIC50	0.4509 to 0.8198
HillSlope	-2.094 to -0.893
IC50	2.824 to 6.603
Goodness of Fit	
Degrees of Freedom	13
R square	0.8695
Absolute Sum of Squares	2957
Sy.x	15.08

Number of points
of X values 15
Y values analyzed 15

MMV003270

log(inhibitor) vs. normalized response
Best-fit values
LogIC50 0.9716
IC50 9.366
Std. Error
LogIC50 0.3838
95% CI (profile likelihood)
LogIC50 0.1607 to 1.731
IC50 1.448 to 53.88
Goodness of Fit
Degrees of Freedom 14
R square 0.3016
Absolute Sum of Squares 48446
Sy.x 58.83

Number of points
of X values 15
Y values analyzed 15

MMV690102

log(inhibitor) vs. normalized response -- Variable slope

Best-fit values
LogIC50 0.3305
HillSlope -2.25
IC50 2.141
Std. Error
LogIC50 0.1192
HillSlope 1.473
95% CI (profile likelihood)
LogIC50 0.06248 to 0.6827
HillSlope ??? to -0.6474
IC50 1.155 to 4.816
Goodness of Fit
Degrees of Freedom 13
R square 0.6268
Absolute Sum of Squares 10785
Sy.x 28.8

Number of points
of X values 15

Y values analyzed

15