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UNITED STATES ENVIRONMENTAL PROTECTION AGENCY WASHINGTON, D.C. 20460

OPP OFFICIAL RECORD HEALTH EFFECTS DIVISION SCIENTIFIC DATA REVIEWS EPA SERIES 361

OFFICE OF PREVENTION, PESTICIDES AND TOXIC SUBSTANCES

June 9, 2006

MEMORANDUM

SUBJECT:

Benchmark dose analysis of cholinesterase inhibition data in neonatal and adult

rats (MRID no. 46688914) following exposure to DDVP

PC Code: 084001. DP Baroode DP 328793. TXR No. 0054223

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The purpose of the current memo is to provide BMD estimates calculated by the Health Effects Division for the acute and repeated dosing studies in adult and juvenile rats in DDVP (MRID nos. 45805703, 45842301, 46433201, 46153304). The results of this memo are intended to inform the FQPA 10X factor for dichlorvos (also called DDVP) in the single chemical aggregate and cumulative risk assessments.

Summary

This memo describes a benchmark dose analysis for red blood cell (RBC) and brain cholinesterase (ChE) inhibition data in adult and juvenile rats exposed orally to DDVP (MRID nos. 45805703, 45842301, 464433201, 46153304). This analysis was developed using the exponential model provided in EPA's OPCum Risk software. The estimated dose to result in 10% inhibition (BMD₁₀) and the lower 95% confidence limit on the BMD₁₀ (BMDL₁₀) were calculated. For acute exposures, the calculated BMDs₁₀ range from approximately 1.3 mg/kg to 2.0 mg/kg for each compartment, sex and age group (young adult and postnatal day 8 [PND8] rats). Regarding repeated exposures, the Agency has used both concurrent and historical controls to estimate RBC and brain BMD₁₀. For most groups, the calculated the BMDs₁₀ ranged from approximately 0.5 mg/kg to 1.2 mg/kg when using the historical or concurrent controls. Overall, for the repeated exposure, the BMDs are similar between compartments, sexes and age groups.

Background

The Agency's draft BMD technical guidance indicates that the BMD approach is a preferable alternative to the NOAEL/LOAEL approach (USEPA, 2000). The Office of Pesticide Programs is increasing its use of BMD techniques in its hazard assessments and risk characterizations. BMDs are preferred over the NOAEL/LOAEL as NOAELs/LOAELs are highly dependent on dose selection in that they are limited to the doses included in a study. Moreover, the NOAEL/LOAEL approach does not account for the uncertainty in the estimate of the dose-response. Benchmark dose analysis attempts to model the doseresponse relationship with a dose-response curve that can be described by a mathematical function. The dose-response curve that is estimated based on the experimental observations is used to estimate the magnitude of the response for any dose within the experimental dose range. Various mathematical models can be used to model this dose-response curve. Once a BMD model has been selected, the dose associated with a specified response (i.e. benchmark response, BMR) is determined. For cholinesterase data, the BMR is typically expressed as a certain percent change in the control group response (i.e., background). The dose resulting in the specified BMR and estimated based on the modeled dose-response curve is termed the "benchmark dose." Generally, the dose resulting in a BMR of X% change in background is referred to as the BMD_X. The corresponding 95% lower confidence limit on the BMD_x is referred to as the BMDL_x. In the case of ChE inhibition for DDVP, EPA has used a BMR of 10% for ChE inhibition. This BMR was selected based on both statistical and toxicological considerations. The reader is referred to the Revised Organophosphate (OP) Cumulative Assessment which provides a detailed power analysis of over 1000 measurements of brain ChE inhibition which indicates that 10% is a level that can be reliably measured in the majority of rat toxicity studies. In the case of the ChE inhibiting pesticides, it is important to consider the degree to which the response level for inhibition of ChE is protective for other toxicities which may result from pesticide exposure. The Agency believes that at a BMR of 10% for brain ChE inhibition is, indeed, protective of other toxicities whereas higher BMR values may not be.

Methods

Selected Model

The BMD analysis of ChE data from the DDVP oral gavage acute and repeated comparative sensitivity toxicity studies was performed using EPA's OPCumRisk program. The exponential function used for modeling the effect of DDVP on ChE activity was:

$$y = B + (A - B) e^{-m \times dose}$$

Where **y** is ChE activity, **dose** is the dose in mg/kg, **m** is the dose scale factor, **A** is background ChE activity, and **B** is the limiting high-dose ChE activity. Both y (ChE activity) and dose were extracted from the above referenced toxicity study. The equation for the exponential model reflects the observation that

ChE activity decreases to a limiting value (B) as dose increases. The model has three parameters to be estimated: m (dose scale factor), A (background), and B (limiting high-dose ChE activity).

The OPCumRisk program utilizes the same dose-response model (i.e., decreasing exponential model) as utilized in the Preliminary OP Cumulative Risk Assessment (CRA; USEPA, 2001). The OPCumRisk program can be obtained at www.epa.gov/pesticides/cumulative/EPA approach methods.htm. These dose-response models and the respective computer code are publicly available for download, review, and use. This method has been previously evaluated by the FIFRA SAP (www.epa.gov/scipoly/sap/2001/index.htm). For the revised OP CRA (USEPA, 2002), the exponential model was expanded to include a "low dose shoulder." The low dose shoulder corresponds to the portion of the dose-response curve where the slope at the low end of the dose-response curve is more flat compared to the slope at higher doses. This low dose shoulder is not modeled in the OPCumRisk program. For DDVP brain and RBC ChE data in the comparative ChE study, there is little evidence that a low dose shoulder exists and as such, the current BMD analysis is considered adequate and supportable. It is important to note that the exponential model is also being used by EPA in its cumulative risk assessment for the OP common mechanism group (of which DDVP is a member; USEPA, 2002).

The calculated BMD values represent the dose at which a 10% reduction in background ChE activity levels is expected (i.e. BMD₁₀ values were calculated). For each ChE dataset, parameters were initially estimated including all dose groups. The OPCumRisk program utilizes a decision algorithm for selecting from among various options for the exponential model. Generally the model is fitted until an adequate p-value for the $\chi 2$ goodness-of-fit (GoF) statistic is obtained. The decision algorithm is provided below.

- 1. If the p-value for the GoF statistic is greater than 0.05, then the model's fit was considered adequate and the initial parameter estimates were used.
- 2. Otherwise (that is, if the p-value was less than 0.05, or no estimates resulted because the model did not converge), the horizontal asymptote was set to zero and the model was refit to the data.
- 3. If the p-value was still less than 0.05, or there was no model fit at all, then the highest dose was dropped and the model was refit with the horizontal asymptote set to zero until either the p-value exceeded 0.05, or there are only three doses remaining.

Although the user can specify options that are not consistent with the default decision algorithm utilized by OPCumRisk, all BMD values provided in these analyses are based on the default decision algorithm. The decision algorithm and technical details of the "basic" exponential model used in this BMD analysis can be obtained at www.epa.gov/scipoly/sap/2001/september/rpfappendix1.pdf.

Results

BMD values (both BMD₁₀ and BMDL₁₀) were calculated for the brain and RBC ChE data from the oral gavage acute comparative sensitivity toxicity study. These BMD values and the goodness of fit information from the OPCumRisk model runs are provided in Tables 1 and 2 for acute and repeated exposures, respectively. Details of this analysis can be found below in Appendix 1 and 2.

Based on statistical and/or visual fit, the exponential model adequately fit the RBC and brain ChE data from the acute exposures to adult and PND8 rats. Moreover, for acute exposures, all of calculated the BMD₁₀ range from approximately 1.3 mg/kg to 2.0 mg/kg. The BMDs are similar between compartments, sexes and age groups. It is also notable that there is a narrow margin between the BMD₁₀ and BMDL₁₀ indicating quality data and quality model fit.

Table 1. BMD Summary of OPP's Analysis of ChE Data from the DDVP Oral Acute Gavage Comparative Sensitivity Toxicity Study (MRID no. 45805703, 45842301) Based on the Exponential Model

Brain	Male	Adult	1.46	0.76	0.53
		Neonate	1.50	1.40	0.46
	Female	Adult	1.29	0.65	0.51
		Neonate	2.02	1.80	0.95
RBC	Male	Adult	1.66	1.31	0.20
		Neonate	1.63	1.41	0.95
	Female	Adult	1.55	1.26	0.65
		Neonate	1.47	0.83	0.15

Based on statistical and/or visual fit, the exponential model adequately fit most of the sex-age-compartment scenarios for the RBC and brain ChE data from the repeated exposures to DDVP (Table 2). As described in the Data Evaluation Record (DER), the ChE activity measurements in some control groups are unusually high for the laboratory which conducted the repeated exposure study. The registrant, AMVAC, provided historical control values for brain and RBC ChE activity. BMD estimates were developed using both the concurrent and pooled historical control values. For most groups, the calculated the BMDs₁₀ ranged from approximately 0.5 mg/kg to 1.2 mg/kg when using the historical or concurrent controls. For the female brain ChE from juvenile rats, when using the concurrent controls, the statistical fit was not adequate (p < 0.05); however, when using the historical controls, the fit was improved. For the female adult RBC ChE data, when using the concurrent controls, the BMDs were substantially lower than values calculated for the other groups; when the historical control values were considered, the BMD for this group increased to be consistent with the remaining groups. Overall, for the repeated exposure, the BMDs are similar between compartments, sexes and age groups. It is also notable that the BMD₁₀ and BMDL₁₀ are close together in magnitude indicating quality data for most treatment groups and quality model fit.

Table 2. BMD Summary of OPP's Analysis of ChE Data from the DDVP Oral Multi-dosing Gavage Comparative Sensitivity Toxicity Study (MRID no. 46433201, 46153304) Based on the Exponential Model

Brain (concurrent controls)	Male	Adult	0.72	0.55	0.71
		Neonate	0.88	0.75	0.0081
	Female	Adult	0.88	0.71	0.84
		Neonate	0.95	0.80	0.022
Brain (pooled historical controls)	Male	Adult	0.77	0.62	0.63
		Neonate	0.84	0.65	0.37
	Female	Adult	0.92	0.75	0.75
		Neonate	0.85	0.62	0.39
RBC (concurrent controls)	Male	Adult	0.71	0.45	0.11
		Neonate	0.50	0.35	0.41
	Female	Adult	0.09	0.054	0.66
		Neonate	0.57	0.43	0.69
RBC (pooled historical controls)	Male	Adult	0.75	0.50	0.30
		Neonate	0.55	0.38	0.094
	Female	Adult	1.17	1.00	0.90
		Neonate	0.61	0.46	0.50

Conclusions

The BMD analysis described here provides a robust consideration of the brain and RBC ChE data from the acute and repeated exposure studies in young adult and juvenile rats exposed to DDVP orally. This analysis indicates that for acute and repeated exposures that young adult and juvenile rats respond similarly to oral dosing to DDVP.

References

USEPA, 2000. "Benchmark Dose Technical Guidance Document" Draft report. Risk Assessment Forum, Office of Research and Development, U.S. Environmental Protection Agency. Washington, DC. EPA/630/R-00/001

USEPA (2001). Preliminary Organophosphorus Pesticide Cumulative Risk Assessment. Office of Pesticide Programs, U.S. Environmental Protection Agency. Washington, DC.http://www.epa.gov/pesticides/cumulative/pra_op/.

USEPA, 2002. Revised Organophosphorous Pesticide Cumulative Risk Assessment; June 10, 2002. Office of Pesticide Programs, U.S. Environmental Protection Agency. Washington, D.C. Available: http://www.epa.gov/pesticides/cumulative/rra-op/

Appendix 1: Benchmark dose runs for the acute dosing (MRID no. 45805703, 45842301)

DDVP:1-D:BRAIN:F:MAIN Tue Apr 25 16:46:22 2006

MRID: ADULT Guideline: 81-1

Continuous Exponential Model (Decreasing) Formula: $chei = B + (A-B) * exp(-(m*dose)^g)$

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 85.75863 89.74156 -38.87931

Coefficients:

Value Std.Error

A 8.7164482 0.9536187

B 2.0810054 2.6396297

m 0.1093959 0.1071854

Correlation:

В

A 1.0000000 0.4940059 0.5941367

в 0.4940059 1.0000000 0.9731630

m 0.5941367 0.9731630 1.0000000

Approximate 95% confidence intervals

Coefficients:

est. upper lower

A 6.91981135 8.7164482 10.9795579 B 0.14322729 2.0810054 30.2357431

m 0.01384314 0.1093959 0.8645054

Residual standard error:

lower est. upper

2.063742 2.750235 4.122999

Degrees of freedom: 20 total; 17 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.4272 with 1 degrees of freedom. P = 0.513

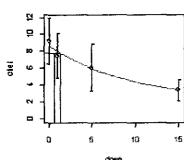
dose n chei Expected sd Exp.SD X2 Resid.

1 0 5 9.17 8.716448 2.19 2.637271 0.38455377

1 5 7.48 8.028853 2.19 2.409555 -0.50933622

5 5 6.03 5.920903 2.25 1.741651 0.14006697

ADULT 1 D - MAIN



Confesses Engagestial Model (Decreasing)

4 15 5 3.36 3.366941 1.00 1.021351 -0.01519591

BMD Computation

BMD = 1.287: BMDL = 0.6495

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.1094

se: 0.1072

var=se^2: 0.01149

Per cent. of background at unit dose: 90

Per cent. of background at the highest dose: 19

ED50 (95% CI): 6.336 (0.9286 , 43.24)

ln(Potency) -2.213

se[log(Potency)]: 0.9798
se[log(Potency)]^2: 0.96

DDVP:1-D:BRAIN:M:MAIN Tue Apr 25 16:46:28 2006 MRID: ADULT Guideline: 81-1

Continuous Exponential Model (Decreasing) Formula: $chei = B + (A-B) * exp(-(m*dose)^g)$

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik74.27587 78.25880 -33.13793

Coefficients:

Value Std.Error

A 7.50616158 0.77906555

B 1.53331933 0.55183572

m 0.09231263 0.05727478

Correlation:

Α \mathbf{B} A 1.0000000 0.5155696 0.6408690

В 0.5155696 1.0000000 0.8990115

m 0.6408690 0.8990115 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 6.02999496 7.50616158 9.343700 в 0.71757370 1.53331933 3.276414

m 0.02493219 0.09231263 0.341792

Residual standard error:

lower est. upper 1.628137 2.169729 3.252736

Degrees of freedom: 20 total; 17 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

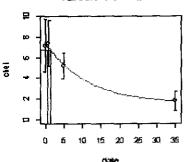
Pearson Chi-Square Statistic: 0.3867 with 1 degrees of freedom. P = 0.534

dose n chei Expected sd Exp.SD X2 Resid.

1 0 5 7.14 7.506162 2.05 2.1838492 -0.374916996

1 5 7.42 6.979477 1.81 2.0377088 0.483405815

5 5 5.22 5.297988 1.01 1.5652808 -0.111408487



ADULT 1 D - MAIN

Continuous Europerist Model (Decreasing)

4 35 5 1 77 1.769368 0.74 0.5147315 0.002747467

BMD Computation

BMD = 1.455: BMDL = 0.7623

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.09231

se: 0.05727

var=se^2: 0.00328

Per cent. of background at unit dose: 91
Per cent. of background at the highest dose: 4

ED50 (95% CI): 7.509 (2.226 , 25.33)

ln(Potency) -2.383

se[log(Potency)]: 0.6204
se[log(Potency)]^2: 0.385

DDVP:1-D:RBC:F:MAIN Tue Apr 25 16:46:26 2006

MRID: ADULT Guideline: 81-1

Continuous Exponential Model (Decreasing) Formula: $chei = B + (A-B) * exp(-(m*dose)^g)$

Variance Function: power

Highest 1 doses dropped from data set.

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

BIC logLik AIC 201.74977 203.87392 -97.87489

Coefficients:

Value Std.Error A 2.571638e+03 71.521128484 m 6.799292e-02 0.009431473

Correlation:

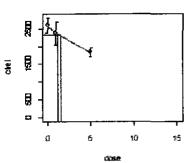
Α A 1.0000000 0.6800864 m 0.6800864 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 2.421676e+03 2.571638e+03 2.730886e+03 m 5.038701e-02 6.799292e-02 9.175059e-02

Residual standard error: lower est. upper 148.7009 205.1177 330.4533



ADULT 1 D - MAIN

Continuous Exponential Model (Decreasing)

Degrees of freedom: 15 total; 13 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.2042 with 1 degrees of freedom. P = 0.651

dose n chei Expected sd Exp.SD X2 Resid.

- 1 0 5 2597 2571.638 179 203.3146 0.27893149
- 1 5 2373 2402.597 264 189.8396 -0.34861343
- 5 5 1835 1830.480 101 144.2978 0.07003547

BMD Computation

BMD = 1.55: BMDL = 1.262

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.06799 se: 0.009431

var=se^2: 8.895e-05

Per cent. of background at unit dose: 93

Per cent. of background at the highest dose: 71

ED50 (95% CI): 10.19 (7.768 , 13.38)

ln(Potency) -2.688

se[log(Potency)]: 0.1387
se[log(Potency)]^2: 0.01924

DDVP:1-D:RBC:M:MAIN
Tue Apr 25 16:46:32 2006

MRID: ADULT Guideline: 81-1

Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

Highest 1 doses dropped from data set.

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 207.4049 209.5290 -100.7024

Coefficients:

Value Std.Error A 2.790461e+03 85.30415667 m 6.358638e-02 0.01043262

Correlation:

A 1.0000000 0.6773461 m 0.6773461 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 2.612126e+03 2.790461e+03 2.980971e+03 m 4.460953e-02 6.358638e-02 9.063597e-02

Residual standard error: lower est. upper 171.6033 236.7092 381.3486

Degrees of freedom: 15 total; 13 residual

Goodness of Fit

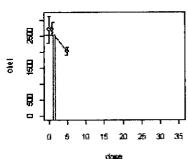
The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 1.633 with 1 degrees of freedom. P = 0.201

dose n chei Expected sd Exp.SD X2 Resid.
1 0 5 2705 2790.461 343 242.2597 -0.7888111
2 1 5 2719 2618.549 170 227.7068 0.9864189

3 5 5 2015 2030.486 94 177.7275 -0.1948314

ADULT 1 D - MAIN



Constructa Exponensial Model (Decreasing)

BMD Computation

BMD = 1.657: BMDL = 1.305

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)* of background activity

Potency: 0.06359

se: 0.01043

var=se^2: 0.0001088

Per cent. of background at unit dose: 94

Per cent. of background at the highest dose: 73

ED50 (95% CI): 10.9 (7.903 , 15.04)

ln(Potency) -2.755

se[log(Potency)]: 0.1641
se[log(Potency)]^2: 0.02692

DDVP: 1-PND: BRAIN: F: MAIN Tue Apr 25 16:46:45 2006 MRID: PND8 Guideline: 81-1

Continuous Exponential Model (Decreasing) Formula: $chei = B + (A-B)*exp(-(m*dose)^g)$

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 4.4493049 7.4365018 0.7753475

Coefficients:

Value Std.Error

A 3.11393414 0.096228698

m 0.05220586 0.003914497

Correlation:

Α

A 1.0000000 0.6615286

m 0.6615286 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper

A 2.91818820 3.11393414 3.32281030

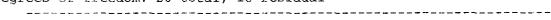
m 0.04459687 0.05220586 0.06111309

Residual standard error:

lower est. upper

0.2410419 0.3190020 0.4717480

Degrees of freedom: 20 total; 18 residual



Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.1003 with 2 degrees of freedom. P = 0.951

dose n chei Expected sd Exp.SD X2 Resid.

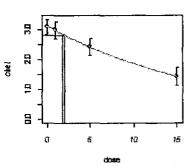
1 0 5 3.08 3.113934 0.21 0.3220624 -0.2356035

1 5 2.98 2.955539 0.23 0.3058005 0.1788627

5 5 2.41 2.398534 0.23 0.2485596 0.1031516

15 5 1.42 1.423045 0.24 0.1480511 -0.0459906





Continues Exponential Model (Decreasing)

BMD Computation

BMD = 2.018: BMDL = 1.797

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.05221 se: 0.003914

var=se^2: 1.532e-05

Per cent. of background at unit dose: 95

Per cent. of background at the highest dose: 46

ED50 (95% CI): 13.28 (11.46 , 15.38)

ln(Potency) -2.953

se[log(Potency)]: 0.07498
se[log(Potency)]^2: 0.005622

DDVP:1-PND:BRAIN:M:MAIN
Tue Apr 25 16:46:52 2006
MRID: PND8 Guideline: 81-1

Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B) *exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik -11.513330 -8.526133 8.756665

Coefficients:

Value Std.Error A 3.22799583 0.070577502 m 0.07048969 0.002815596

Correlation:

A m A 1.0000000 0.6557125 m 0.6557125 1.0000000

Approximate 95% confidence intervals

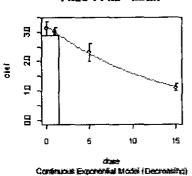
Coefficients:

lower est. upper A 3.08307201 3.22799583 3.37973197 m 0.06481574 0.07048969 0.07666033

Residual standard error:

lower est. upper 0.1743612 0.2307548 0.3412458

PND6 1 PND - MAIN



Degrees of freedom: 20 total; 18 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 1.551 with 2 degrees of freedom. P ≈ 0.46

 dose n
 chei
 Expected
 sd
 Exp.SD
 X2 Resid.

 1
 0
 5
 3.14
 3.227996
 0.20
 0.23552098
 -0.8354443

 2
 1
 5
 3.04
 3.008290
 0.10
 0.21999032
 0.3223142

 3
 5
 5
 2.33
 2.269167
 0.24
 0.16745554
 0.8123092

 4
 15
 5
 1.11
 1.121331
 0.09
 0.08465241
 -0.2992973

BMD Computation

BMD = 1.495: BMDL = 1.403

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.07049 se: 0.002816

var=se^2: 7.928e-06

Per cent. of background at unit dose: 93

Per cent. of background at the highest dose: 35

ED50 (95% CI): 9.833 (9.093 , 10.63)

ln(Potency) -2.652

se[log(Potency)]: 0.03994
se[log(Potency)]^2: 0.001595

DDVP:1-PND:RBC:F:MAIN
Tue Apr 25 16:46:50 2006
MRID: PND8 Guideline: 81-1

Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B) *exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 311.3597 315.3426 -151.6799

Coefficients:

Value Std.Error
A 3565.7686985 253.06695411
B 1304.7155991 653.72006213
m 0.1170090 0.08519761

Correlation:

A B m A 1.0000000 0.4552855 0.5739303 B 0.4552855 1.0000000 0.9594333 m 0.5739303 0.9594333 1.0000000

Approximate 95% confidence intervals

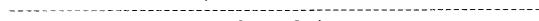
Coefficients:

lower est. upper A 3069.8953381 3565.7686985 4141.7393788 B 453.3342457 1304.7155991 3755.0280190 m 0.0251796 0.1170090 0.5437383

Residual standard error:

lower est. upper 553.7944 738.0112 1106.3850

Degrees of freedom: 20 total; 17 residual



Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

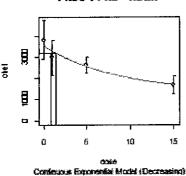
Pearson Chi-Square Statistic: 2.127 with 1 degrees of freedom. P = 0.145

dose n chei Expected sd Exp.SD X2 Resid. 1 0 5 3830 3565.769 733 696.0392 0.84885899

2 1 5 2991 3316.097 662 639.2002 -1.13726262 3 5 5 2636 2564.305 315 480.8800 0.33337969

D--- 00 -/

PND8 1 PND - MAIN



4 15 5 1689 1695.615 349 331.6280 -0.04460369

BMD Computation

BMD = 1.467: BMDL = 0.8317

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.117 se: 0.0852

var=se^2: 0.007259

Per cent. of background at unit dose: 89

Per cent. of background at the highest dose: 17

ED50 (95% CI): 5.924 (1.422 , 24.68)

ln(Potency) -2.146

se[log(Potency)]: 0.7281
se[log(Potency)]^2: 0.5302

DDVP:1-PND:RBC:M:MAIN
Tue Apr 25 16:46:56 2006
MRID: PND8 Guideline: 81-1

Continuous Exponential Model (Decreasing) Formula: $chei = B + (A-B) * exp(-(m*dose)^g)$

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 301.8481 304.8353 -147.9240

Coefficients:

Value Std.Error A 3.677296e+03 1.737759e+02 m 6.483032e-02 6.020933e-03

Correlation:

A 1.0000000 0.6594638 m 0.6594638 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 3.329745e+03 3.677296e+03 4.061124e+03 m 5.333839e-02 6.483032e-02 7.879824e-02

Residual standard error: lower est. upper 433.6720 573.9343 848.7482

Degrees of freedom: 20 total; 18 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.09684 with 2 degrees of freedom. P = 0.953

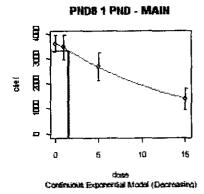
 dose
 n
 chei
 Expected
 sd
 Exp.SD
 X2 Resid.

 1
 0
 5
 3620
 3677.296
 276
 580.9977
 -0.22051497

 2
 1
 5
 3473
 3446.460
 411
 545.1029
 0.10887159

 3
 5
 5
 2693
 2659.202
 442
 422.3719
 0.17892749

 4
 15
 5
 1384
 1390.583
 359
 223.2216
 -0.06594177



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BMD Computation

BMD = 1.625: BMDL = 1.41

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.06483 se: 0.006021

var=se^2: 3.625e-05

Per cent. of background at unit dose: 94

Per cent. of background at the highest dose: 38

ED50 (95% CI): 10.69 (8.912 , 12.83)

ln(Potency) -2.736

se[log(Potency)]: 0.09287
se[log(Potency)]^2: 0.008625

Appendix 2: Benchmark dose runs for the repeated dosing (MRID no. 46433201, 46153304)

DDVP:7-D:BRAIN:F:WHOLE

Mon Apr 24 21:43:34 2006

MRID: MDAdconc Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 25.51746 29.50039 -8.75873

Coefficients:

Value Std.Error

A 5.4680592 0.21623032

B 1.1331342 0.27076016

m 0.1529701 0.03053395

Correlation:

A B r

A 1.0000000 0.1901865 0.3377826

B 0.1901865 1.0000000 0.9369459

m 0.3377826 0.9369459 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper

A 5.0303656 5.4680592 5.9438367

B 0.6844438 1.1331342 1.8759659

m 0.1003945 0.1529701 0.2330789

Residual standard error:

lower est. upper

0.5119114 0.6821960 1.0227101

Degrees of freedom: 20 total; 17 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.03915 with 1 degrees of freedom. P = 0.843

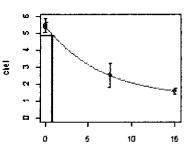
dose n chei Expected sd Exp.SD X2 Resid.

1 0.0 5 5.51 5.468059 0.34 0.6770768 0.138510867

2 0.1 5 5.36 5.402252 0.25 0.6689085 -0.141243990

3 7.5 5 2.51 2.509476 0.56 0.3104510 0.003776490

MDAdconc 7 D - WHOLE



Continuous Exponential Model (Decreasing)

4 15.0 5 1.57 1.570124 0.15 0.1945313 -0.001419862

BMD Computation

BMD = 0.8814: BMDL = 0.713

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.153 se: 0.03053

var=se^2: 0.0009323

Per cent. of background at unit dose: 86

Per cent. of background at the highest dose: 10

ED50 (95% CI): 4.531 (3.064 , 6.701)

ln(Potency) -1.878

se[log(Potency)]: 0.1996
se[log(Potency)]^2: 0.03984

DDVP:7~D:BRAIN:M:WHOLE Mon Apr 24 21:43:40 2006

MRID: MDAdconc Guideline: NONGUIDELINE Continuous Exponential Model (Decreasing) Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 38.65735 42.64027 -15.32867

Coefficients:

Value Std.Error

A 5.7226764 0.31678214

B 1.2589525 0.25158074

m 0.1917306 0.04343782

Correlation:

A B r

A 1.0000000 0.1403111 0.3067966

B 0.1403111 1.0000000 0.8887349

m 0.3067966 0.8887349 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 5.0918770 5.7226764 6.4316216

B 0.8258587 1.2589525 1.9191677

m 0.1188781 0.1917306 0.3092295

Residual standard error:

lower est. upper 0.7591362 1.0116588 1.5166222

Decrease of Empadam, 20 tetal 17 maridus

Degrees of freedom: 20 total; 17 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.1429 with 1 degrees of freedom. P = 0.705

dose n chei Expected sd Exp.SD X2 Resid.

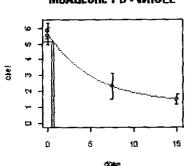
1 0.0 5 5.84 5.722676 0.40 0.9917110 0.264536182

2 0.1 5 5.52 5.637908 0.42 0.9768860 -0.269889484

3 7.5 5 2.32 2.318672 0.68 0.4003437 0.007418733

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MDAdconc 7 D - WHOLE



Communic Economial Model (Decreasing)

4 15.0 5 1.51 1.510537 0.26 0.2624115 -0.004578106

BMD Computation

BMD = 0.7156: BMDL = 0.551

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.1917

se: 0.04344

var=se^2: 0.001887

Per cent. of background at unit dose: 83

Per cent. of background at the highest dose: 5.6

ED50 (95% CI): 3.615 (2.319 , 5.636)

ln(Potency) -1.652

se[log(Potency)]: 0.2266
se[log(Potency)]^2: 0.05133

DDVP:7-D:BRAIN:F:WHOLE Mon Apr 24 21:43:45 2006

MRID: MDAdhist Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 64.04178 70.79730 -28.02089

Coefficients:

Value Std.Error

A 5.3368308 0.11905590

В 1.0997356 0.28281397

m 0.1465709 0.02915622

Correlation:

A 1.0000000 0.1150917 0.2036726 B 0.1150917 1.0000000 0.9481542 m 0.2036726 0.9481542 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 5.10097136 5.3368308 5.5835960 B 0.65311837 1.0997356 1.8517598 m 0.09794975 0.1465709 0.2193271

Residual standard error:

lower est. upper 0.5322331 0.6528354 0.8446040

Degrees of freedom: 40 total; 37 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.104 with 1 degrees of freedom. P = 0.747

 dose
 n
 chei
 Expected
 sd
 Exp.SD
 X2 Resid.

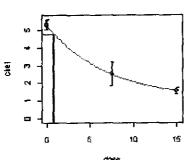
 1
 0.0
 25
 5.32
 5.336831
 0.60
 0.6499687
 -0.129474129

 2
 0.1
 5
 5.36
 5.275180
 0.25
 0.6424933
 0.295197982

 3
 7.5
 5
 2.51
 2.511155
 0.56
 0.3062988
 -0.008433535

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MDAdhist 7 D - WHOLE



Continuous Exponential Model (Decreasing)

4 15.0 5 1.57 1.569894 0.15 0.1909515 0.001242926

BMD Computation

BMD = 0.9185: BMDL = 0.7518

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.1466

se: 0.02916

var≈se^2: 0.0008501

Per cent. of background at unit dose: 86

Per cent. of background at the highest dose: 11

ED50 (95% CI): 4.729 (3.202 , 6.984)

ln(Potency) -1.92

se[log(Potency)]: 0.1989
se[log(Potency)]^2: 0.03957

DDVP:7~D:BRAIN:M:WHOLE Mon Apr 24 21:43:52 2006

MRID: MDAdhist Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 78.62698 85.38250 -35.31349

Coefficients:

Value Std.Error

A 5.4406635 0.14572755

в 1.2233684 0.23205212

m 0.1793503 0.03602974

Correlation:

A 1.00000000 0.08852252 0.1851917 B 0.08852252 1.00000000 0.9107214 m 0.18519170 0.91072141 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 5.1532609 5.4406635 5.7440950 B 0.8329963 1.2233684 1.7966830 m 0.1193784 0.1793503 0.2694501

Residual standard error:

lower est. upper 0.6580918 0.8072133 1.0443300

Degrees of freedom: 40 total; 37 residual

begrees of freedom; 40 cocar; 57 residuar

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.2307 with 1 degrees of freedom. P = 0.631

 dose
 n
 chei
 Expected
 sd
 Exp.SD
 X2 Resid.

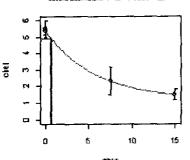
 1
 0.0
 25
 5.41
 5.440664
 0.61
 0.7954922
 -0.192733152

 2
 0.1
 5
 5.52
 5.365700
 0.42
 0.7846436
 0.439720961

 3
 7.5
 5
 2.32
 2.322001
 0.68
 0.3407920
 -0.013132451

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MDAdhist 7 D - WHOLE



Continuous Exponential Model (Decreasing)

4 15.0 5 1.51 1.509570 0.26 0.2200316 0.004374445

~_____

BMD Computation

BMD = 0.7701: BMDL = 0.6168

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.1794

se: 0.03603

var=se^2: 0.001298

Per cent. of background at unit dose: 84

Per cent. of background at the highest dose: 6.8

ED50 (95% CI): 3.865 (2.607 , 5.73)

ln(Potency) -1.718

se[log(Potency)]: 0.2009
se[log(Potency)]^2: 0.04036

DDVP:7-D:RBC:F:WHOLE Mon Apr 24 21:43:38 2006

MRID: MDAdconc Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 252.1003 256.0833 -122.0502

Coefficients:

Value Std.Error A 2727.000013 79.6172246 B 1258.499990 25.9812588 m 2.397083 0.8174793

Correlation:

A B m A 1.000000e+00 1.765012e-06 0.66321810 B 1.765012e-06 1.000000e+00 0.05862781 m 6.632181e-01 5.862781e-02 1.00000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 2564.091283 2727.000013 2900.259098 B 1204.860966 1258.499990 1314.526963 m 1.167355 2.397083 4.922246

Residual standard error: lower est. upper 133.5911 178.0295 266.8919

Degrees of freedom: 20 total; 17 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.1959 with 1 degrees of freedom. P = 0.658

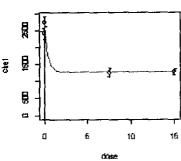
 dose n
 chei
 Expected
 sd
 Exp.SD
 X2 Resid.

 1
 0.0
 5
 2727
 2727.0
 150
 178.02953
 -1.663420e-07

 2
 0.1
 5
 2414
 2414.0
 158
 157.59563
 2.423883e-07

 3
 7.5
 5
 1247
 1258.5
 109
 82.15994
 -3.129848e-01

MDAdconc 7 D - WHOLE



Continuous Exponential Model (Decreasive)

4 15.0 5 1270 1258.5 70 82.15994 3.129847e-01

BMD Computation

BMD = 0.0857: BMDL = 0.05385

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 2.397

se: 0.8175

var=se^2: 0.6683

Per cent. of background at unit dose: 9.1

Per cent. of background at the highest dose: 2.4e-14

ED50 (95% CI): 0.2892 (0.1482 , 0.5642)

ln(Potency) 0.8743

se[log(Potency)]: 0.341
se[log(Potency)]^2: 0.1163

DDVP:7-D:RBC:M:WHOLE Mon Apr 24 21:43:43 2006

MRID: MDAdconc Guideline: NONGUIDELINE Continuous Exponential Model (Decreasing) Formula: chei = B + $(A-B)*exp(-(m*dose)^g)$

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 290.6427 294.6256 -141.3214

Coefficients:

Value Std.Error A 2681.7192647 145.3811253 В 1135.7992785 123.8320471 0.2702996 0.1094602

Correlation:

A В A 1.0000000 0.0618944 0.1785788 B 0.0618944 1.0000000 0.8143054 m 0.1785788 0.8143054 1.0000000

Approximate 95% confidence intervals

Coefficients:

est. lower A 2391.8831196 2681.7192647 3006.6762692 B 902.4076927 1135.7992785 1429.5534174 0.1150242 0.2702996 0.6351871

Residual standard error: lower est. upper

368.1789 490.6517 735.5575

Degrees of freedom: 20 total; 17 residual

Goodness of Fit

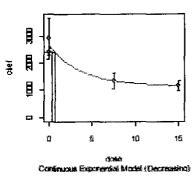
The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 2.552 with 1 degrees of freedom. P = 0.110

dose n chei Expected sd Exp.SD X2 Resid. 1 0.0 5 2908 2681.719 610 453.8838 1.11477671 2 0.1 5 2413 2640.493 111 445.6730 -1.14139604

3 7.5 5 1346 1339.393 236 220.5037 0.06699746

MDAdconc 7 D - WHOLE



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4 15.0 5 1159 1162.612 155 200.3797 -0.04030813

BMD Computation

BMD = 0.7048: BMDL = 0.4481

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.2703

se: 0.1095

var=se^2: 0.01198

Per cent. of background at unit dose: 76

Per cent. of background at the highest dose: 1.7

ED50 (95% CI): 2.564 (1.16 , 5.671)

ln(Potency) -1.308
se[log(Potency)]: 0.405
se[log(Potency)]^2: 0.164

DDVP:7-D:RBC:F:WHOLE Mon Apr 24 21:43:50 2006

MRID: MDAdhist Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

Highest 1 doses dropped from data set.

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 432.6351 436.8387 -213.3175

Coefficients:

Value Std.Error A 2.453625e+03 68.975570904 m 9.025628e-02 0.009178156

Correlation:

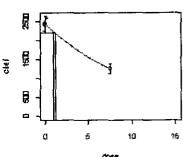
A m A 1.0000000 0.4137541 m 0.4137541 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 2.316326e+03 2.453625e+03 2599.0618980 m 7.328462e-02 9.025628e-02 0.1111583

Residual standard error: lower est. upper 273.4370 344.5616 466.0030 MDAdhist 7 D - WHÖLE



Confinues Exponential Model (Decressing)

Degrees of freedom: 30 total; 28 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.01654 with 1 degrees of freedom. P = 0.898

 dose
 n
 chei
 Expected
 sd
 Exp.SD
 X2 Resid.

 1
 0.0
 20
 2458
 2453.625
 399
 343.9574
 0.056889167

 2
 0.1
 5
 2414
 2431.579
 158
 340.8656
 -0.115315893

 3
 7.5
 5
 1247
 1246.880
 109
 174.7398
 0.001538003

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BMD Computation

BMD = 1.167: BMDL = 1

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.09026

se: 0.009178

var=se^2: 8.424e-05

Per cent. of background at unit dose: 91

Per cent. of background at the highest dose: 51

ED50 (95% CI): 7.68 (6.292 , 9.374)

ln(Potency) -2.405

se[log(Potency)]: 0.1017
se[log(Potency)]^2: 0.01034

DDVP:7-D:RBC:M:WHOLE Mon Apr 24 21:43:56 2006

MRID: MDAdhist Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B) *exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 506.1301 512.3515 -249.0650

Coefficients:

Value Std.Error
A 2610.5056190 77.5940879
B 1131.7044406 110.4118748
m 0.2605430 0.0942001

Correlation:

A B m A 1.00000000 0.04048633 0.1151000 B 0.04048633 1.00000000 0.8225566 m 0.11509996 0.82255665 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 2457.1412313 2610.5056190 2773.4423646 B 927.7403439 1131.7044406 1380.5101278 m 0.1247496 0.2605430 0.5441513

Residual standard error: lower est. upper 314.8342 391.4927 517.8247

Degrees of freedom: 35 total; 32 residual

Goodness of Fit

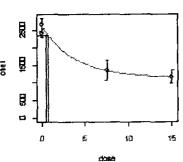
The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 1.095 with 1 degrees of freedom. P = 0.295

dose n chei Expected sd Exp.SD X2 Resid.
1 0.0 20 2650 2610.506 411 385.8967 0.45769820
2 0.1 5 2413 2572.474 111 379.8508 -0.93877615
3 7.5 5 1346 1341.244 236 195.8061 0.05430863

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MDAdhist 7 D - WHOLE



Continuous Esponential Model (Decreasing)

4 15.0 5 1159 1161.395 155 173.2784 -0.03091094

BMD Computation

BMD = 0.7455: BMDL = 0.4982

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.2605

se: 0.0942

var=se^2: 0.008874

Per cent. of background at unit dose: 77

Per cent. of background at the highest dose: 2

ED50 (95% CI): 2.66 (1.31 , 5.404)

ln(Potency) -1.345

se[log(Potency)]: 0.3616
se[log(Potency)]^2: 0.1307

DDVP:7-D:BRAIN:F:WHOLE Mon Apr 24 21:43:59 2006

MRID: MDPupconc Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

Highest 1 doses dropped from data set.

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 39.29316 41.41732 -16.64658

Coefficients:

Value Std.Error A 5.8381055 0.33369886 m 0.1111981 0.01312090

Correlation:

A 1.0000000 0.5883474 m 0.5883474 1.0000000

Approximate 95% confidence intervals

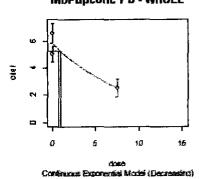
Coefficients:

lower est. upper A 5.15992657 5.8381055 6.6054187 m 0.08617663 0.1111981 0.1434846

Residual standard error:

lower est. upper 0.8565106 1.1814689 1.9033966

MDPupcone 7 D - WHOLE



Degrees of freedom: 15 total; 13 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 5.215 with 1 degrees of freedom. P = 0.0224

 dose n
 chei
 Expected
 sd
 Exp.SD
 X2 Resid.

 1
 0.0
 5
 6.59
 5.838105
 0.59
 1.0482540
 1.60389304

 2
 0.1
 5
 5.02
 5.773546
 0.46
 1.0365378
 -1.62558583

 3
 7.5
 5
 2.54
 2.535575
 0.54
 0.4511931
 0.02192763

BMD Computation

BMD = 0.9475: BMDL ≈ 0.7935

Potency Measures

A unit dose (1 mg/kg) would result in $100*\exp(-Potency)$ % of background activity

Potency: 0.1112

se: 0.01312

var=se^2: 0.0001722

Per cent. of background at unit dose: 89

Per cent. of background at the highest dose: 43

ED50 (95% CI): 6.233 (4.946 , 7.855)

ln(Potency) -2.196
se[log(Potency)]: 0.118
se[log(Potency)]^2: 0.01392

DDVP:7-D:BRAIN:M:WHOLE Mon Apr 24 21:44:06 2006

MRID: MDPupconc Guideline: NONGUIDELINE Continuous Exponential Model (Decreasing) Formula: chei = $B + (A-B)*exp(-(m*dose)^g)$

Variance Function: power

Highest 1 doses dropped from data set.

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 38.62180 40.74595 -16.31090

Coefficients:

Value Std.Error A 6.2176817 0.33406064 m 0.1201434 0.01231365

Correlation:

Α A 1.000000 0.589248 m 0.589248 1.000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 5.53629700 6.2176817 6.9829285 m 0.09628064 0.1201434 0.1499205

Residual standard error:

lower est. upper 0.8659536 1.1944947 1.9243816

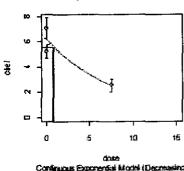
Degrees of freedom: 15 total; 13 residual

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Goodness of Fit

Pearson Chi-Square Statistic: 7.004 with 1 degrees of freedom. P = 0.00813

dose n chei Expected sd Exp.SD X2 Resid. 1 0.0 5 7.09 6.217682 0.64 1.0494073 1.85872816 2 0.1 5 5.27 6.143427 0.44 1.0367170 -1.88387271 3 7.5 5 2.53 2.525203 0.41 0.4213597 0.02545729 MDPupcone 7 D - WHOLE



Continuous Exponential Model (Decreasing)

BMD Computation

BMD = 0.877: BMDL = 0.7504

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.1201 se: 0.01231

var=se^2: 0.0001516

Per cent. of background at unit dose: 89

Per cent. of background at the highest dose: 41

ED50 (95% CI): 5.769 (4.719 , 7.053)

ln(Potency) -2.119

se[log(Potency)]: 0.1025
se[log(Potency)]^2: 0.0105

DDVP:7-D:BRAIN:F:WHOLE Mon Apr 24 21:44:12 2006

MRID: MDPuphist Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

MDPuphist 7 D - WHOLE

Continuous Engagemental Model (Decreasing)

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AIC BIC logLik 91.73904 98.07312 -41.86952

Coefficients:

Value Std.Error A 5.4535860 0.20843990 B 1.3874190 0.37759636

m 0.1686908 0.05235096

Correlation:

A B m A 1.0000000 0.0969725 0.1952096 B 0.0969725 1.0000000 0.9213612 m 0.1952096 0.9213612 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 5.04558081 5.4535860 5.8945841 B 0.79750815 1.3874190 2.4136825 m 0.08971903 0.1686908 0.3171746

Residual standard error:

lower est. upper 0.867120 1.075062 1.415080

Degrees of freedom: 36 total; 33 residual

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Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.7522 with 1 degrees of freedom. P = 0.386

 dose
 n chei
 Expected
 sd
 Exp.SD
 X2 Resid.

 1
 0.0
 21
 5.54
 5.453586
 1.18
 1.0587084
 0.374039299

 2
 0.1
 5
 5.02
 5.385569
 0.46
 1.0451764
 -0.782104088

 3
 7.5
 5
 2.54
 2.534843
 0.54
 0.4881430
 0.023622823

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4 15.0 5 1.71 1.711208 0.22 0.3342521 -0.008084159

BMD Computation

BMD = 0.8537: BMDL = 0.6228

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.1687 se: 0.05235

var=se^2: 0.002741

Per cent. of background at unit dose: 84
Per cent. of background at the highest dose: 8

ED50 (95% CI): 4.109 (2.237 , 7.549)

ln(Potency) -1.78

se[log(Potency)]: 0.3103
se[log(Potency)]^2: 0.09631

DDVP:7-D:BRAIN:M:WHOLE Mon Apr 24 21:44:18 2006

MRID: MDPuphist Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 90.63050 97.28474 -41.31525

Coefficients:

Value Std.Error

A 5.6883165 0.17732604

В 1.1782553 0.33427136

m 0.1610776 0.03905932

Correlation:

A B m A 1.0000000 0.1065217 0.2028358 B 0.1065217 1.0000000 0.9324765

m 0.2028358 0.9324765 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 5.33981535 5.6883165 6.059562 B 0.66276419 1.1782553 2.094690 m 0.09850411 0.1610776 0.263400

Residual standard error:

lower est. upper 0.7834716 0.9634302 1.2514578

Degrees of freedom: 39 total; 36 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.8159 with 1 degrees of freedom. P = 0.366

 dose
 n
 chei
 Expected
 sd
 Exp.SD
 X2 Resid.

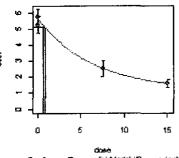
 1
 0.0
 24
 5.76
 5.688316
 1.05
 0.9517421
 0.36898250

 2
 0.1
 5
 5.27
 5.616251
 0.44
 0.9394402
 -0.82415231

 3
 7.5
 5
 2.53
 2.525726
 0.41
 0.4194838
 0.02278489

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MDPuphist 7 D - WHOLE



Confinence Exponential Model (Decreasing)

4 15.0 5 1.58 1.580839 0.21 0.2660983 -0.00704853

BMD Computation

BMD = 0.837: BMDL = 0.6514

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.1611 se: 0.03906

var=se^2: 0.001526

Per cent. of background at unit dose: 85

Per cent. of background at the highest dose: 8.9

ED50 (95% CI): 4.303 (2.675 , 6.921)

ln(Potency) -1.826

se[log(Potency)]: 0.2425
se[log(Potency)]^2: 0.0588

DDVP:7-D:RBC:F:WHOLE Mon Apr 24 21:44:03 2006

MRID: MDPupconc Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B) *exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 282.2902 286.2732 -137.1451

Coefficients:

Value Std.Error
A 3367.8252655 128.82707749
B 1182.3605015 84.42210176
m 0.2942281 0.06693016

Correlation:

A B m A 1.00000000 0.05726548 0.1789545 B 0.05726548 1.0000000 0.7844004 m 0.17895453 0.78440045 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 3106.7025997 3367.8252655 3650.8956538 B 1017.0123351 1182.3605015 1374.5913469 m 0.1820754 0.2942281 0.4754635

Residual standard error:

lower est. upper 305.9931 407.7801 611.3211

Degrees of freedom: 20 total; 17 residual

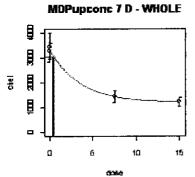
Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.1602 with 1 degrees of freedom. P = 0.689

dose n chei Expected sd Exp.SD X2 Resid.
1 0.0 5 3418 3367.825 470 401.9922 0.27909526
2 0.1 5 3254 3304.460 257 394.2670 -0.28617896
3 7.5 5 1424 1422.897 201 169.0092 0.01459009

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Continuous Exponental Model (Decreasing)

4 15.0 5 1208 1208.834 145 144.7334 -0.01289216

BMD Computation

BMD = 0.5688: BMDL = 0.427

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)* of background activity

Potency: 0.2942 se: 0.06693

var=se^2: 0.00448

Per cent. of background at unit dose: 75

Per cent. of background at the highest dose: 1.2

ED50 (95% CI): 2.356 (1.508 , 3.679)

ln(Potency) -1.223

se[log(Potency)]: 0.2275
se[log(Potency)]^2: 0.05175

DDVP:7-D:RBC:M:WHOLE Mon Apr 24 21:44:10 2006

MRID: MDPupconc Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B) *exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 284.3103 288.2933 -138.1552

Coefficients:

Value Std.Error A 3405.8244474 134.5531824 B 1265.5873297 82.6795659 m 0.3447276 0.0992573

Correlation:

A B m A 1.00000000 0.05821642 0.1595949 B 0.05821642 1.0000000 0.7517474 m 0.15959491 0.75174742 1.0000000

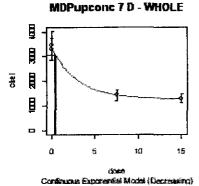
Approximate 95% confidence intervals

Coefficients:

lower est. upper A 3133.4511572 3405.8244474 3701.8736162 B 1102.6365176 1265.5873297 1452.6194839 m 0.1877810 0.3447276 0.6328497

Residual standard error: lower est. upper 317.7069 423.3904 634.7231

Degrees of freedom: 20 total; 17 residual



Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.6739 with 1 degrees of freedom. P = 0.412

dose n chei Expected sd Exp.SD X2 Resid.
1 0.0 5 3299 3405.824 353 418.5584 -0.57068917
2 0.1 5 3441 3333.302 484 410.1457 0.58715849

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3 7.5 5 1423 1426.879 186 177.3018 -0.04892138 4 15.0 5 1280 1277.743 146 155.8875 0.03238133

BMD Computation

BMD = 0.5028: BMDL = 0.3509

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.3447 se: 0.09926

var=se^2: 0.009852

var=se 2: 0.009652

Per cent. of background at unit dose: 71

Per cent. of background at the highest dose: 0.57

ED50 (95% CI): 2.011 (1.144 , 3.535)

ln(Potency) -1.065

se[log(Potency)]: 0.2879
se[log(Potency)]^2: 0.0829

DDVP:7-D:RBC:F:WHOLE Mon Apr 24 21:44:16 2006

MRID: MDPuphist Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

AIC BIC logLik 471.1821 477.1681 -231.5910

Coefficients:

Value Std.Error
A 3212.0118304 78.13348301
B 1175.9601007 84.40054910
m 0.2793221 0.06360987

Correlation:

A B m A 1.00000000 0.03961674 0.1212268 B 0.03961674 1.0000000 0.7972136 m 0.12122680 0.79721362 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 3056.3407841 3212.0118304 3375.6117944 B 1015.6286087 1175.9601007 1361.6022102 m 0.1754370 0.2793221 0.4447228

Residual standard error: lower est. upper 301.6668 377.5026 504.5976

Degrees of freedom: 33 total; 30 residual

Goodness of Fit

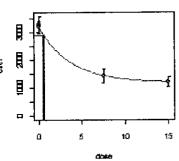
The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 0.4548 with 1 degrees of freedom. P = 0.5

dose n chei Expected sd Exp.SD X2 Resid.
1 0.0 18 3185 3212.012 377 372.5099 -0.30764682
2 0.1 5 3254 3155.927 257 366.1959 0.59885197
3 7.5 5 1424 1426.559 201 166.5420 -0.03435202

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MDPuphist ? D - WHOLE



Continues Exponential Model (Decreasing)

4 15.0 5 1208 1206.804 145 139.2642 0.01920492

BMD Computation

BMD = 0.6147: BMDL = 0.4647

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.2793

se: 0.06361

var=se^2: 0.004046

Per cent. of background at unit dose: 76

Per cent. of background at the highest dose: 1.5

ED50 (95% CI): 2.482 (1.588 , 3.878)

ln(Potency) -1.275

se[log(Potency)]: 0.2277
se[log(Potency)]^2: 0.05186

DDVP:7-D:RBC:M:WHOLE Mon Apr 24 21:44:21 2006

MRID: MDFuphist Guideline: NONGUIDELINE
Continuous Exponential Model (Decreasing)
Formula: chei = B + (A-B)*exp(-(m*dose)^g)

Variance Function: power

The BMD corresponds to a dose that results in a 10% reduction in the response relative to the control

Summary of Model Fitting Results

MDPuphist 7 D - WHOLE

Continuous Exponential Model (Decression)

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AIC BIC logLik 508.6161 514.8375 -250.3081

Coefficients:

Value Std.Error A 3229.7438903 83.54133448 B 1259.4597968 88.49275967 m 0.3240709 0.09828153

Correlation:

A B m A 1.00000000 0.03616319 0.1024154 B 0.03616319 1.0000000 0.7630263 m 0.10241540 0.76302628 1.0000000

Approximate 95% confidence intervals

Coefficients:

lower est. upper A 3063.9809484 3229.7438903 3404.4746923 B 1091.5109432 1259.4597968 1453.2506428 m 0.1747259 0.3240709 0.6010671

Residual standard error: lower est. upper

355.8975 442.5545 585.3637

Degrees of freedom: 35 total; 32 residual

Goodness of Fit

The chi-squared goodness-of-fit values should be taken as general indications of fit only. P-values are likely to be inaccurate to some degree

Pearson Chi-Square Statistic: 2.812 with 1 degrees of freedom. P = 0.0936

dose n chei Expected sd Exp.SD X2 Resid. 1 0.0 20 3162 3229.744 385 415.0112 -0.7300042

2 0.1 5 3441 3166.916 484 407.7138 1.5031865

3 7.5 5 1423 1432.825 186 187.6611 -0.1170705

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4 15.0 5 1280 1274.714 146 161.4604 0.0732030

BMD Computation

BMD = 0.5525: BMDL = 0.3825

Potency Measures

A unit dose (1 mg/kg) would result in 100*exp(-Potency)% of background activity

Potency: 0.3241 se: 0.09828

var=se^2: 0.009659

Per cent. of background at unit dose: 72

Per cent. of background at the highest dose: 0.77

ED50 (95% CI): 2.139 (1.18 , 3.876)

ln(Potency) -1.127

se[log(Potency)]: 0.3033
se[log(Potency)]^2: 0.09197



R128680

Chemical: Dichlorvos

PC Code: 084001

HED File Code: 13000 Tox Reviews

Memo Date: 6/9/2006 File ID: DPD328793 Accession #: 412-06-0192

HED Records Reference Center 7/12/2006