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UNITED STATES ENVIRONMENTAL PROTECTION AGENCY

WASHINGTON, D.C. 20460

MEMORANDUM

OFFICE OF PREVENTION, PESTICIDES AND TOXIC SUBSTANCES

SUBJECT: Review of Fish Life Cycle Study for Tebuconazole (MRID No. 430096-01)

FROM:

Anthony F. Maciorowski, Chief

Ecological Effects Branch

7507C

TO:

Denise Greenway Registration Division

7505C

The Ecological Effects Branch completed the review of a fish life cycle study. The study was submitted to support the registration of tebuconazole. The study is the following:

Wheat, J., 1993. HWG 1608 (tebuconazole): Life-Cycle Chronic Toxicity to the Sheepshead Minnow (*Cyprinodon variegatus*) Under Flow-Through Conditions. MRID No. 430096-01.

The study is scientifically sound however, it is classified as supplemental because (1) water quality was variable throughout the study and (2) solvent control, survival was very low and length was different from the control. A new study will not be required. The results of the fish early life stage for the rainbow trout (MRID No. 407009-14) and the fathead minnow (MRID No. 420382-02) and this fish life cycle study provide sufficient information to characterize the chronic effects of tebuconazole on fish. In addition a mesocosm is in the process of review. The mesocosm will provide additional information on the effects to fish. The rainbow trout fish early life stage study showed that this species is more sensitive than the sheepshead and the fathead minnow.

The attached Data Evaluation Record will provide the necessary information for the classification of the study. If you have any questions please contact Harry Craven (305-5320) or Conchi Rodríguez (308-2805).



DP Barcode PC Code No EEB Out

To:

Denise Greenway

Product Manager

Registration Division

From: Anthony F. Maciorowski, Chief

Ecological Effects Branch/EFED (H7507C)

Attached, please find the EEB review of...

Reg./File #	: Tebuçonazole
Type Product	: Fungicide
Product Name	: Folicur
Company Name	: Miles Inc.
Purpose	: Review of fish life cycle study
Action Code	: 116 Date Due : 3/24/94

Reviewer : Conchi Rodríguez Date In EEB: 11/30/93

gdln no	MRID NO	CAT GDLN NO	MRID NO	CAT	GDLN NO	MRID NO	CAT
71-1 (A)		72-2 (A)			72-7 (A)		
71-1(B)		72-2 (B)			72-7 (B)		
71-2 (A)		72-3 (A)			122-1 (A)		
71-2(B)		72-3 (B)			122-1 (B)	,	
71-3		72-3 (C)			122-2		
71-4(A)		72-3 (D)			123-1(A)		
71-4 (B)		72-3 (B)			123-1(B)		
71-5 (A)		72-3 (P)			123-2		
71-5(B)		72-4(A)			124-1		
72-1(A)		72-4(B)			124-2		
72-1 (B)		72-5	430096-01	s	141-1		
72-1(C)		72-6			141-2		
72-1(D)				١,	141-5		

Y=Acceptable (Study satisfied Guideline)/Concur
P=Partial (Study partially fulfilled Guideline but
additional information is needed
S=Supplemental (Study provided useful information but Guideline was
not satisfied)
M=Unacceptable (Study was rejected)/Nonconcur

DATA EVALUATION RECORD

- 1. CHEMICAL: Tebuconazole. Shaughnessey No. 128997.
- 2. TEST MATERIAL: BAY HWG-1608 (Folicur technical); α -[2-(4-Chlorophenyl)ethyl]- α -(1,1-dimethylethyl)-1H-1,2,-triazole-1-ethanol; Batch No. 0-79-01530; 96.4% purity; a white powder.
- 3. <u>STUDY TYPE</u>: 72-5. Fish Full Life-Cycle Toxicity Test. Species Tested: Sheepshead Minnow (*Cyprinodon variegatus*).
- 4: CITATION: Wheat, J. 1993. HWG 1608 (tebuconazole): Life-Cycle Chronic Toxicity to the Sheepshead Minnow (Cyprinodon variegatus) Under Flow-Through Conditions. Laboratory ID No. J9104001. Prepared by Toxikon Environmental Sciences, Jupiter, FL. Submitted by Miles Incorporated, Kansas City, MO. EPA MRID No. 430096-01.
- 5. REVIEWED BY:

Conchi Rodríguez

Biologist

Ecological Effects Branch

Date: 12/16/94

Environmental Fate and Effects Division

6. APPROVED BY:

Harry Craven
Supervisory Biologist
Ecological Effects Branch
Environmental Fate and Effects Division

Signature: 7/2/16/44

Date:

- 7. <u>CONCLUSIONS</u>: This study is scientifically sound. It is classified as supplemental. The information may be useful in risk assessment. A new study is not required. Water quality parameters (i.e., salinity and temperature) were highly variable throughout the test period. Solvent control survival was low (24%) in one of the solvent control replicates. The MATC of BAY HWG-1608 for *Cyprinodon variegatus* is >19.0 and <43.6 μ g ai/l based mean measured concentrations (geometric mean MATC = 28.8 μ g ai/l). The most sensitive endpoint was fish length.
- 8. RECOMMENDATIONS: N/A.

9. BACKGROUND:

10. DISCUSSION OF INDIVIDUAL TESTS: N/A.

11. MATERIALS AND METHODS:

- A. Test Animals: Sheepshead minnow (Cyprinodon variegatus) embryos were obtained by stripping eggs from 25 adults females and fertilizing the eggs with the sperm of 5 adult males. The adults were obtained from two commercial suppliers (Aquatic BioSystems, Fort Collins, CO and TRAC Labs, Gulf Breeze, FL) and were maintained at the testing facility for 43 days before hormone injection. During the first four weeks, the fish were treated with a dilute formalin solution and nitrofurazone. No treatment was administered during the 15 days before injection. The embryos were less than 24 hours old post-fertilization at test initiation.
- Test System: Two proportional vacuum-siphon diluter в. systems each with a 50% dilution factor were used to deliver the test solutions to the test vessels. A 2-1 diluter system with 24-1 glass tanks was used for the F_0 and F_1 phases of the study. The test vessels had a solution volume of 15 l and a solution depth of 13 cm. The diluter provided 6.5 volume additions per day to each test vessel during the F₀ phase and 6.3 volume additions per day during the F, phase of the study. 4-1 system with 53-1 glass tanks was used for the reproductive phase of the study. The test vessels had a solution volume of 40 1 and a solution depth of 23 The diluter provided 4.1 volume additions per day to each test vessel.

For the incubation period of the F_0 and F_1 phases of the study, incubation cups were used to contain the embryos. The cups were 60-mm diameter glass tubes with 355- μ m mesh screen and were stationary in the replicate tanks. A self-starting siphon provided a fluctuating level of test solution.

After hatching, all fish were transferred to screen retention chambers within each test vessel. The retention chambers were constructed of a petri dish bottom (150 mm x 10 mm) with a Nitex screen collar (20-cm high).

For the reproductive phase of the study, the spawning chambers were each constructed of two, 3-gallon high

density polyethylene, white pails with screen bottoms. The pail containing adult fish had a 7-mm plastic screen mesh bottom and was placed into the second pail which had $355-\mu m$ screen mesh bottom. The spawning chambers were suspended by the sides of the tanks above the bottom to allow passage of incoming test solutions.

The test chambers were randomly positioned in a water bath. The test system was maintained under a 16-hour light/8-hour dark photoperiod with 15-minute dawn/dusk simulation periods. Light intensity was 430-957 lux. From test day 43 to test termination, all test solutions were aerated.

The dilution water was natural saltwater from a shallow well. The saltwater was carbon treated and adjusted to a salinity of 20 parts per thousand (ppt) with carbon-treated, aerated laboratory freshwater. The dilution water was aerated and filtered (20 $\mu\text{m})$ prior to use. The water was also UV sterilized prior to use in the F_0 and F_1 phases of the study.

Stock solutions were prepared monthly by weighing 8.16 g of BAY HWG-1608 into 100 ml of triethylene glycol (TEG). An appropriate amount of stock solution was pumped to the chemical mixing chamber of each diluter at each cycle to provide 200 μg ai/l, the highest nominal concentration tested. This test solution was proportionally diluted to provide the remaining test concentrations.

- C. <u>Dosage</u>: Two-hundred-three day, flow-through, full life-cycle toxicity test. Based on the results of previous toxicity testing, five nominal concentrations (12.5, 25.0, 50.0, 100, and 200 µg ai/l) were selected for this study. A dilution water control and a solvent control were also included. The concentration of TEG in the solvent control and all exposure solutions was 20.55 µl/l.
- Design: Two replicate aquaria were provided per treatment. For the F₀ phase of the study, each aquarium contained two replicate incubation/retention chambers (i.e., 4 chambers/treatment). Embryos (<24 hours post-fertilization) were impartially placed, five at a time, into each incubation cup until each cup contained 50 embryos (i.e., 200 eggs/treatment). Survival of embryos was recorded daily until hatching was complete.</p>

After hatching, all fish were transferred into retention chambers within the same test chamber. Survival and sublethal effects were monitored daily. On day 33, all fish were photographed for length measurements. On day 58, all fish in the replicates of each treatment were pooled and 25 fish were indiscriminately selected, photographed for length measurements and released into their respective tank. The fish were maintained in this system until day 113 when the reproductive phase of the study began. The remaining fish were sacrificed for length and weight measurements and then shipped frozen to Miles Incorporated for residue analysis.

For the reproduction phase of the study, fish were transferred from the first diluter system into the corresponding replicate tank of the second diluter system. The first reproductive session was initiated with the selection of sexually mature adult fish (generally 2 males and 5 females) and placement of these fish into a spawning chamber. Four spawning sessions were conducted, each lasting 14 days. The first pail of the spawning chamber contained the adult fish with mesh screen which allowed the eggs to pass through to the second pail with mesh screen to retain the spawned eggs. Spawning chambers were removed daily and checked for newly spawned eggs. Any eggs present were removed and counted. After each spawning session, the fish were removed and returned to the first diluter system for holding. In an attempt to lower the variability in egg production, it was decided to run the fourth spawning session using fish from previous spawnings. At the end of the spawning sessions, the fish were sacrificed for length and weight measurements and were submitted for residue analysis.

The F_1 phase of the study was initiated on day 163 in the first diluter system. Where possible, 25 embryos, obtained during the period between spawning sessions 3 and 4, were distributed to each of two embryo incubation chambers per replicate tank (i.e., 50 embryos/replicate, 100 embryos/treatment). For three days following initiation, each embryo chamber was placed in a solution of 10% formalin in seawater for 10 minutes. It was decided that this treatment was necessary to control disease. Survival was recorded until hatching was complete. Approximately two weeks after hatching, all fish were transferred into screen retention chambers within the same test chamber. Survival and sublethal effects were monitored daily.

On day 28, all fish were sacrificed for length and weight measurements and then shipped frozen to Miles Incorporated for residue analysis.

During the F_0 and F_1 phases of the study, fish were fed live, fatty-acid supplemented brine shrimp nauplii, two to three times daily. During the reproduction phase of the study, the fish were fed frozen adult brine shrimp at least twice daily.

Temperature was measured hourly in one replicate of the dilution water control in the first diluter system. In addition, the diurnal temperature range of the water bath was measured. The diurnal temperature range of both replicates of the dilution water control was measured in the second diluter system. Salinity was measured daily in the dilution water control of both systems. Dissolved oxygen concentrations (DO) and pH were measured at the initiation of each phase and weekly thereafter in all test solutions.

One day prior to test initiation and weekly thereafter, water samples were collected from all replicates and analyzed using high pressure liquid chromatography to measure actual exposure concentrations.

E. Statistics: Several statistical analyses (two-tail Fisher's exact test, two-tail Chi-square test of independence, nested analysis of variance (ANOVA), or a Student's t-test) were conducted to compare dilution water control and solvent control responses, responses between chambers within each replicate tank, or responses between replicates within each treatment group. The decision to pool control data for a given variable was based on consistent pattern of biological and statistical findings.

Dichotomous data were analyzed using a one-tailed Fisher's exact test and a one-tailed Chi-Square test of independence. Where there was sufficient variability across chambers, dichotomous data were analyzed using a nested analysis of variance (ANOVA). Continuous data were analyzed using an ANOVA with Dunnett's comparison test. Homogeneity of variance across treatments was compared using Levene's test at p<0.01. Arcsine or log transformations were applied, as necessary, to the dichotomous or continuous data, respectively, to improve the homogeneity of variance. If the p value was still less than 0.01, a nonparametric Dunnett's

test was conducted, otherwise a parametric analysis was performed on the transformed data.

12. REPORTED RESULTS: Mean measured concentrations were 9.2, 19.0, 43.6, 96.7, and 196.9 μg ai/1 for the F₀ phase; 9.2, 21.9, 49.0, 98.6, and 196.0 μg ai/1 for the reproductive phase, and 9.6, 18.4, 46.9, 97.8, and 192 μg ai/1 for the F₁ phase. These values represent 73-99% of nominal concentrations (Tables 2, 16, and 24, attached). No undissolved test material was observed in the test chambers during the study.

Hatching success (%) of F0 embryos exposed to concentrations of BAY HWG-1608 was not significantly different from that of the pooled control (Table 3, attached). Hatching success of the control and the solvent control was 86.5% and 89.0%, respectively. F0 survival on day 58 (number surviving on day 58 divided by total hatched) was significantly reduced at 196.9 μ g ai/l when compared to the pooled control (Table 4, attached). Survival of the control and solvent control survival on day 58 was 93.6% and 89.3%, respectively. F0 survival on day 177 (number of fish alive on day 177 divided by the number of fish present after culling on day 58) was significantly reduced at 196.9 μ g ai/l when compared to the pooled control (Table 5, attached). Survival of the control and solvent control on day 177 was 76% and 56%, respectively.

Due to the significant survival effects, the 196.9 μg ai/l exposure group was not included in the growth analyses. Fo length on day 33 was significantly reduced at 96.7 μg ai/l when compared to the pooled control (Table 6, attached). Fo growth (i.e., length and weight) on days 58 and 177 was not significantly reduced at any exposure concentration when compared to the pooled control or to the solvent control (Tables 7, 8, 9, and 10, attached).

The residue in tissues of F₀ sheepshead minnow generally increased with concentrations of test material in water. The mean tissue residue ranged from 0.12 mg/kg for fish in the lowest test level to 12.87 mg/kg for fish in the highest test level.

The exposure solutions showed no significant effect on the reproductive parameters when compared to the control data (Tables 17 and 18, attached). For spawning sessions 1 and 2, reproductive data at 9.2 μg ai/l was substantially lower than the pooled control. However, this difference was not statistically significant and was not confirmed by the results at higher exposure concentrations.

Hatching success (%) of F_1 embryos exposed to 18.4 μg ai/l was significantly reduced when compared to that of the pooled control and the solvent control (Table 25, attached). However, this difference was regarded as spurious. There were no significant differences at any other exposure concentration. Hatching success of the control and the solvent control was 93.7% and 100%, respectively. F_1 survival at the exposure concentrations were higher than the solvent control at 28 days post-hatch (Table 26, attached). Survival of the control and solvent control on day 28 post-hatch was 95.5% and 86.0%, respectively. F_1 growth (i.e., length and weight) at 28 days post-hatch was significantly reduced at 97.8 μg ai/l when compared to the pooled control (Tables 27 and 28, attached).

The residue in F_1 sheepshead minnow tissues generally increased with concentrations of the test material in water. The mean tissue residue ranged from 0.31 mg/kg for fish in the lowest test level to 2.96 mg/kg for fish in the highest . test level.

During the F_0 phase of the study, the test solutions had a temperature range of 26.1-33.6°C, a pH range of 7.9-8.4, a salinity range of 16-22 ppt, and a DO range of 2.4-6.6 mg/l. During this phase, the DO was reduced in the solvent control and all BAY HWG-1608 solutions due to bacterial growth as a result of the solvent. Aeration was initiated on day 43 to maintain acceptable DO levels.

During the reproduction phase of the study, the test solutions had a temperature range of 22.0-29.9°C, a pH range of 8.0-8.3, a salinity range of 16-26 ppt, and a DO range of 4.8-6.9 mg/l.

During the F_1 phase of the study, the test solutions had a temperature range of 20.6-30.7°C, a pH range of 8.1-8.3, a salinity range of 15-20 ppt, and a DO range of 5.9-7.7 mg/l.

Based upon growth (length at day 33) of the F_0 and F_1 sheepshead minnows as the most sensitive biological endpoint, the maximum acceptable toxicant concentration (MATC) was >43.6 and <96.7 μg ai/l with a geometric mean of the MATC of 64.9 μg ai/l. The NOEC was 43.6 μg ai/l.

"Although there was some variability in results such as the percent survival for control groups at day 177 of the F_0 generation, reproduction and wet weight of the F_0 and F_1 generation, the study clearly shows that BAY HWG-1608 is

neither cumulatively nor reproductively toxic at concentrations tested in this study."

Quality Assurance and Good Laboratory Practice Compliance Statements were included in the report, indicating that the study was conducted in accordance with USEPA Good Laboratory Practice Standards (40 CFR Part 160). The dates and types of quality assurance audits performed were also included in the report.

14. REVIEWER'S DISCUSSION AND INTERPRETATION OF STUDY RESULTS:

A. <u>Test Procedure</u>: The test procedures were generally in accordance with the SEP except for the following:

The SEP states that 50 embryos should be distributed to each of four replicate chambers at test initiation. In this test, two egg cups were used per aquarium (two aquaria per concentration) and therefore constitutes a nested design rather than individual replicates.

Water quality parameters were highly variable during the study. Throughout the study, the test solutions had a salinity of 15-26%, a temperature of 20.6-33.6°C, and a DO range of 2.4-7.7 mg/l.

 F_{o} survival in the solvent control was 56% at the end of F_{o} phase. Only 6 out of 25 fish (24%) survived in replicate B of the solvent control. In addition, F_{o} growth was substantially reduced in replicate B, of the solvent control, at test termination.

The light intensity used during the test (430-957 lux) was greater than recommended by the SEP (10-100 lux).

- B. Statistical Analysis: Hatchability, survival, and reproduction data were not analyzed since significant reductions were visually evident. The reviewer used an ANOVA and Bonferroni's test to analyze juvenile and adult growth.
- C. <u>Discussion/Results</u>: Water quality parameters (i.e., salinity and temperature, oxygen) were highly variable throughout the test period. Mean temperature for the F₀ generation, ranged from 28.5 (day 103) to 32.44 °C (day 50) during the period of the study. Mean temperature, for the F₁ generation, ranged from 25.65 (day 35) to 28.88 °C (day 16). Salinity, in the F₀ generation, generally ranged from 17 to 22%. During the reproduction session there was a big jump in

salinity because of a malfunction in the saltwater system. Dissolved oxygen was low at day 43 and aeration was initiated. Since this is a long study, the deviations in the water quality parameters might not have an influence in the overall results of the study.

Survival in one replicate of the solvent control was very low at the end of the study (24%). In a period of 83 days the survival was reduced to 24 %. The mortality in the solvent control does not have a big influence in the results of the study since the solvent control mortality in the F1 generation was not different from the control.

During the reproduction session, female survival was low. This lead to variability in the number of eggs produced per female reproductive day.

Residues of tebuconazole were found in the fish in all the concentrations. The mean residues at the lowest concentration (9.6 $\mu g/l$) was 0.12 mg/kg. The mean residue at the highest concentration (196.9 $\mu g/l$) was 12.87 mg/kg.

For all analyzed parameters, the reviewer obtained results similar to those of the author (pages 1, 8, 15, 21, and 22 of printouts, attached), with the following exceptions: 1) F_1 weight at test termination was statistically reduced at 9.2 and 96.7 μ g ai/l when compared to the solvent control (page 29 of printouts, attached); and 2) F_1 length at test termination was statistically reduced at 43.6 and 96.7 μ g ai/l when compared to the solvent control (page 30 of printouts, attached). The reduction in F_1 weight at 9.2 μ g ai/l is probably not biologically significant. The results of the reviewer's verification of the author's conclusions gives the MATC as >19.0 and <43.6 μ g ai/l mean measured concentrations which is more conservative than the author's.

This study is scientifically sound. It is classified as supplemental. A new is not required. The results of the fish early life stage for the rainbow trout (MRID No. 407009-14) and the fathead minnow (MRID No. 420382-02) and this fish life cycle study provide sufficient information to characterize the effects of tebuconazole on fish. Based on the conditions of this study the MATC of BAY HWG-1608 for Cyprinodon variegatus was >19.0 and <43.6 μg ai/l mean measured

concentrations (geometric mean MATC = 28.8 μg ai/1). The most sensitive parameter was length.

D. Adequacy of the Study:

- (1) Classification: Supplemental
- (2) Rationale: 1) Water quality parameters were highly variable (temperature, salinity and DO).
 2) Survival of F₀ fish in replicate B of the solvent control was extremely low (24%) at terminal sacrifice. However, survival of the fish in the solvent control of the F1 generation was not low (86.0%).
 3) Solvent control growth of F₀ fish at test termination was highly variable.
- (3) Repairability: No.
- 15: COMPLETION OF ONE-LINER FOR STUDY: Yes; 22 March 1994.

Table 2. Mean Measured Concentrations of BAY HWG-1608 in Saltwater During the F0 Phase of a Full Life Cycle Toxicity Test with Sheepshead Minnow, Cyprinodon variegatus

Nominal Test Concentration	Rep	Test	Mean Measured Test Concentration (µq/L; ppb)				
(µg/L; ppb)	_		icate		reatment	Nominal	
Control	○ - A \)	<1.3					
	A) B	<1.3		<1.3			
Solvent Control	A B	<1.3					
	В	<1.3		<1.3			
12.5	A		(1.2)				
	В	9.0	(1.3)	9.2	(1.2)	74	
25.0	A	19.1	(2.1)			•	
	В	18.9	(2.4)	19.0	(2.2)	76	
50.0	A	42.8	(3.6)			200	
•	В		(4.1)	43.6	(3.9)	87	
100	A	96.0	(11.7)				
	В		(11.8)	4 96.7	(11.6)	97	
200	A	195.6	(15.7)				
•	В		(17.2)	196.9	(16.3)	99	

NOTE: Values in parentheses are the standard deviations of the means.

Table 3. Hatching Success (Recent Hatch) of F0 Sheepshead Minnows, <u>Cyprinodon variegatus</u>, Exposed to BAY HWG-1608 During a Full Life Cycle Toxicity Test

Treatment Group Mean Measured	-	Numbe:	r of Fished	l Hato	hed (1	Hatch) [NI	OF ¹]
Concentration		Tank_	Α		Tank	В		
$(\mu g/L; ppb)$	Cham			Cha	mber		Tre	eatment
Group		2	Mean	1	2	Mean		Mean
Control	84)	86	85.0 [1]	94	82	88.0	[0]	86.5
Sol. Control	96 🗇	82	89.0 [1]	86	92	89.0	[1]	89.0
Pooled Control	,							87.8
9.2	94	98∌	96.0 [2]	88	86	87.0	[0]	91.5
19.0	82	86	84.0 [3]	98	84	91.0	[0]	87.5
43.6	88	86	87.0.[1]	96	88	92.0	[1]	89 .5 °
96.7	92	88	90.0 ^G 0]	94	84	89.0	[1]	89.5
196.9	82	90	86.0 [3]	94	92	93.0	[1]	89.5

Note: Initial number = 50 for all chambers.

There were no significant differences from the pooled control using a one-tailed Chi-square, Fisher's exact test, or parametric Dunnett's means comparison procedure $(\underline{P}>.05)$.

¹NDF = Number of deformed fish in the replicate.

Table 4. Survival (Percent Survival) of F0 Sheepshead Minnows, <u>Cyprinodon variegatus</u>, Exposed to BAY HWG-1608 on Day 58 (Date Population Culled to 25 Fish per Replicate) During a Full Life Cycle Toxicity Test

Treatment Gro Mean Measure Concentration	ed .		of Fi	nt Surviva sh Alive/			
$(\mu g/L; ppb)$	Cham	ber		Chamb	er	Tr	eatment
	11	22	Mean	1	2	Mean	Mean
	<i>,</i>		•				
Control	95.2	. 97.7	96.5	95.7	85.4	90.9	93.6
	(40/42)	(42/43)		(45/47) ((35/41)		
Sol. Control	93.8	90.2	92.1	93.0	80.4	86.5	89.3
	(45/48)	(37/41)		(40/43)	(37/46)		
			,				•
Pooled Contro	ol		. · · -		* ***		91.4
	•						
	89.4		89.6	93.2	88.4	90.8	90.2
	(42/47)	(44/49)	\cup	(41/44)((38/43)		
			. 5	`			
19.0	95.1	90.7	92.8	85.7	95.2	90.1	91.4
	(39/41)	(39/43)		(42/49)((40/42)		
				2			
43.6	100.0	95.3	97.7	89.6	95.5 (42/44)	92.4	95.0
	(44/44)	(41/43)		(43/48) ((42/44)		
•				€ 6			
96.7	89.1	88.6	88.9	91.5	81.0	86.5	87.7
	(41/46)	(39/44)		(43/47)	(34/42)		
	•	•		• • •	· · ·		•
196.9	22.0	0.0	10.5	23.4	4.3	14.0	12.3*
	(9/41)	(0/45)		(11/47)	(2/46)		
		•			1		

Note: *Significantly different (P<.05) from the pooled control using a one-tailed Chi-square test, a Fisher's exact test, and a nonparametric Dunnett's means comparison procedure.

Table 6. Mean (Standard Deviation and n) Growth (Measured Photographically as Length in mm) of F0 Sheepshead Minnows, <u>Cyprinodon variegatus</u>, Exposed to BAY HWG-1608 on Day 33 During a Full Life Cycle Toxicity Test

Treatment Gro Mean Measure Concentration (µg/L: ppb)	ed on			Chamb		Treatment	
	<u> </u>	2	Mean	1	2	Mean	Mean
Control	11.8 (1.8) (n=40)	11.1 (1.5) (n=42)	11.4		11.2 (1.8) (n=36)	11.0	11.2
Sol. Control		11.8 (1.6)	11.4	(2.0)	11.8 (1.9) (n=38)	11.5	11.4
Pooled Contro	ol						11.3
9.2	10.9 (1.1) (n=42)	(1.1)		11.6 (1.2) (n=41)		11.9	11.4
19.0		11.0 (1.3) (n=43)	11.1	(1.5)	11.3 (1.5) (n=40)	11.1	11.1
43.6	10.4 (1.4) (n=44)	11.4 (1.0) (n=41)	10.9	(1.2) (n=43)	11.2 (1.2) (n=42)		
96.7		10.2 (1.8) (n=41)	10.5	(2.1)	10.7 (1.8) (n=38)		10.4*
196.9		(2.6)	9.6	11.3 (3.2) (n=11)		11.0	10.3#

Note:

*Significantly different (P<.05) from the pooled control using a one-tailed parametric Dunnett's means comparison procedure.

#Treatment level not included in statistical comparison to control because of survival effects.

Table 5. Survival (Percent Survival) of F0 Sheepshead Minnows, <u>Cyprinodon variegatus</u>, Exposed to BAY HWG-1608 on Day 177 (Date Population F0 Population Terminated and Based Upon Culled Population) During a Full Life Cycle Toxicity Test

Treatment Group - Mean Measured Concentration		Hatch)			
(μg/L; ppb)	Tank A		Tank B		Treatment Mean
Control	68.0 (17/25)	84.0	(21/25)	76.0
Sol. Control	88.0 (22/25)	24.0	(6/25)	56.0
Pooled Control	\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\\				66.0
9.2	56.0 (14/25)	52.0	(13/25)	54.0
19.0	96.0 (24/25)	72.0	(18/25)	84.0
43.6	92.0 (23/25)	64.0	(16/25)	78.0
96.7	36.0 (9/25)	92.0	(23/25)	64.0
196.9	11.1 (1/9)	1, 7.7	(1/13)	9.1*

Note: Initial number = 25 per chamber, except for 196.9 μ g/L, tank A, n=9 and tank B, n=13.

*Significantly different (\underline{P} <.05) from the pooled control using a one-tailed Chi-square test and a Fisher's exact test.

Table 7. Mean (Standard Deviation and n) Growth (Measured as Length in mm) of F0 Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 on Day 58 During a full Life Cycle Toxicity Test

Treatment Group	o -			Length			
Mean Measured		Tank A			Tank B		
Concentration (µg/L; ppb)	Char 1	mber 2	Mean	Chaml 1	per 2	Mean	Treatment <u>Mean</u>
Control	(2.4)	16.4 (2.8) (n=42)	16.8	(2.8)	16.5 (2.4) (n=35)	17.4	17.1
Sol. Control		17.0 (3.1) (n=37)	18.0	(2.8)	17.8 (2.6) (n=37)	18.7	18.3
Pooled Control							17.7
9.2		17.3 (2.3) (n=44)	0	(2.0)	17.4 (2.2) (n=38)	18.2	18.1
19.0	(1.7)	17.3 (2.3) (n=39)	18.3		16.9 (3.3) (n=40)	17.6	17.9
43.6	(2.5)	17.0 (2.4) (n=41)		(14)	17.0 (2.2) (n=42)	17.6	17.8
96.7	(2.5)	16.0 (2.7) (n=39)		(2.4)	16.1 (3.2) (n=34)	17.5	17.6
196.9	17.7 (5.3) (n=9)		17.7		19.5 (7.8) (n=2)	15.6	16.5#

Note: There were no significant differences from the pooled control or the solvent control using a one-tailed parametric Dunnett's means comparison procedure (P>.05).

#Treatment level not included in statistical comparison to control because of survival effects.

Table 8. Mean (Standard Deviation and n) Growth (Measured as Wet Weight in mg) of F0 Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 on Day 58 During a Full Life Cycle Toxicity Test

Treatment Green Mean Measure	-	Mean Length Will Tank B					
Concentration		mber		Chamber			Treatment
(µg/L; ppb)) 1	2	Mean	11	2	Mean	Mean
Control	(57.1)	142.1 (54.9) (n=17)		174.3 (92.7) (n=45)		172.4	158.9
Sol. Control	(68.1)	180.2 (96.2) (n=12)		216.6 (100.7) (n=40)	(73.8)		194.3
Pooled Contro	ol	, ***	X				176.3
9.2	(75.7)	177.3 (82.8) (n=19)		(71.0)	189.2 (85.2) (n=15)	201.5	190.1
19.0	(68.8)	173.8 (65.4) (n=14)		173.0 (73.1) (n=42)	(97.6)	173.7	184.6
43.6	(90.3)	173.1 (54.6) (n=16)	186.4			164.8	175.6
96.7	(88.4)	158.8 (91.1) (n=14)			190.3 (102.1) (n=9)	192.0	193.9

Note: There were no significant differences from the pooled control or the solvent control using a parametric one-tailed Dunnett's means comparison procedure (P>.05).

Treatment level 5 not included because no fish were available for wet weights.

Table 9. Mean (Standard Deviation and n) Growth (Measured as Length in cm) of F0 Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 on Day 177 During a Full Life Cycle Toxicity Test

Treatment Group	_		
Mean Measured		<u>Length</u>	•
Concentration (µq/L: ppb)	Tank A	Tank B	Treatment Mean
Control	3.41 (0.32) (n=17)	3.56 (0.52) (n=21)	3.49
Sol. Control	3.56 (0.44) (n=22)	2.60 (0.68) (n=6)	3.35
Pooled Control	3.0	•	3.43
9.2	2.91 (0.50) (n=14)	2.97 (0.43) (n=13)	2.94
19.0	3.33 (0.43) (n=24)	○3.12 (0.42) ○ (n=18)	3.24
43.6	3.50 (0.33) (n=23)	2.98 (0.44) (n=16)	3.28
96.7	2.97 (0.49) (n=9)	3.24 (0.44) (n=23)	3.17
196.9	3.10 () (n=1)	4.00 (√) (n=1) ∪	3.55#

Note:

There were no significant differences from the pooled control using a one-tailed parametric Dunnett's means comparison procedure (P>.05).

#Treatment level not included in statistical comparison to control because of survival effects.

Table 10. Mean (Standard Deviation and n) Growth (Measured as Wet Weight in grams) of F0 Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 on Day 177 During Full Life Cycle Toxicity Test

Treatment Group Mean Measured Concentration (µg/L; ppb)	- Tank A	Tank B	Treatment Mean
Control	1.65 (0.49) (n=17)	1.89 (0.83) (n=21)	1.78
Solv.Control	1.95 (0.73) (n=22)	0.74 (0.63) (n=6)	1.69
Pooled Control			1.74
9.6	1.08 (0.65) (n=14)	1.07 (0.65) (n=13)	1.08
19.0	1.64 (0.83) (n=24)	1.19 (0.53) (n=18)	1.45
43.6	1.53 (0.50) (n=23)	1.10 (0.57) (n=16)	1.35
96 . 7	0.90 (0.58) (n=9)	1.52 (0.59) (n=23)	1.34
196.9	0.70 () (n=1)	2.41 (1.56#

Note: There were no significant differences from the pooled control using a one-tailed parametric bunnett's means comparison procedure (P>.05).

#Treatment level not included in statistical comparison to control because of survival effects.

Table 16. Mean Measured Concentrations of BAY HWG-1608 in Saltwater During the Reproduction Phase of a Full Life Cycle Exposure of the Sheepshead Minnow, Cyprinodon variegatus, Under Flow-Through Conditions

Nominal Test Concentration (µg/L; ppb)	Rep				l ug/L; ppb) reatment	Percent of Nominal
Control	a)	<1.3				
	B	<1.3		<1.3		
Solvent Control		<1.3				
	В	<1.3		<1.3		
12.5	A		(1.1)			,
	В	8 • 8	(1.4)	9.2	(1.3)	74
25.0	A		(2.0)		** va ##*.	
	В	22.0	(2.1)	21.9	(2.0)	88
50.0	A	49.3	(3,1)			
	В	48.6	(4.1)	49.0	(3.6)	98
100	A	98.8	(7.2)			
	В		(5.9)	4 98.6	(6.4)	99
200 .	A	190.5	(16.5)	1		
	В		(35.9)	196.0	(27.7)	98

NOTE: Values in parentheses are the standard deviations of the means.

Table 17. Number of Eggs and Number of Female Reproductive Days of F0 Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 in a Full Life Cycle Toxicity Test

Treatment Group	-						
Mean Measured Concentration			c	ession	•		
(µq/L; ppq)	Tan	k.	1 1	2	3	4	Total
	1 (1)	<u> </u>	-				IOCAL
Control	A	EGGS	2329	1764	1011	171	5275
		REPR. DAYS	70	61	59	37	227
	$\frac{1}{2}$					•	
	В	EGGS	1306	1871	29	809	4015
	•	REPR.DAYS	70	70	4	42	186
Sol. Control	A	EGGS	2046	1657	148	3	3854
		REPR.DAYS	70	70	9	4	153
		40					
	В	EGGS	729	927	130	76	1862
		REPR. DAYS	45	52	11	. 5	113
Pooled Control		EGGS	6410	6210	1210	1050	15006
FOOTER COUCTOI		REPR.DAYS	255	6219 253	1318 83	1059 88	15006 679
		REFR. DAIS	O 255	253	83	88	6/9
9.2	A	EGGS	<^ 571	230	470	698	1969
		REPR.DAYS	49	38	17	42	146
			, et				
	В	EGGS	25	106	185	0	316
		REPR.DAYS	26	19	21	3	69
21.9	A	EGGS	1059	2417	1937	813	6226
		REPR. DAYS	70	70	70	30	240
					1		~.0
	В	EGGS	441	503 ⁾	944	684	2572
	•	REPR.DAYS	26	42	51	37	156
40.0		77.00.00					
49.0	A	EGGS	2322	2062	1284	590	6258
		REPR.DAYS	70	70	84	31	255
	В	EGGS	307	1668	184	513	2672
	_	REPR.DAYS	70	56	24	23	173
			, ,		~ -	2.0	173
98.6	A	EGGS	a	13		1190	1203
		REPR.DAYS	0	8	0	42	50
	В	EGGS	2662	1500	2602	1105	70.15
	D	REPR.DAYS	2662 70	1502 70	2603 70	1195	7962
·		ICHE ICO DATO	, 0	70	70	42	252

Note: Treatment level 5 not included because of survival effects in the F0 generation.

a Insufficient number of sexually mature fish with which to form a spawning group. Not included in analysis.

Table 18. Reproductive Growth (Measured as Eggs per Female Reproductive Day) of F0 Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 in a Full Life Cycle Toxicity Test

Treatment Gro	oup -		Eggs Per		Reproductive Session	Day Overall
Group	Tank	1	2	3	4	Mean
Control	A	33.3	28.9	17.	1 4.6	23.2
COLLCTOT	В	18.7	26.7	7.		21.6
,	MEAN)	26.0	27.7	16.		22.5
•	MEAN	20.0	27.7	11,	19 11.95	22.5 22.4
Sol. Control	A	29.2	23.7	16.	4 0.8	25.2
	В	160.2	17.8	11.	8 15.2	16.5
	MEAN	24.1	21.2	13.		21.5 - 20.
		33.4	20.75	14.	.l 8	·
Pooled Contro	1	25.1	24.6	15.	9 12.0	22.1
9.2	A	11.7	6.1	27.	6 16.6	13.5
J • 4	В	1.0	5.6	8.		4.6
	MEAN	7.9	5.9	17.		10.6
	*****	, , ,	J.,		2 23.3	10.0
21.9	A	15.1	34.5	27.	7 27.1	25.9
		17.0	12.0			16.5
	MEAN	15.6	26.1	23.		22.2
49.0	A	33.2	29.5	15.	3 19.0	24.5
43.0	В	4.4	29.8	7.		15.4
	MEAN	18.8	29.6	13.		20.9
•	LITTURAL	10.0	23.0		~ 20.7	20.3
98.6	A	3	1.6		-*<^ 28.3	24.1
	В	38.0	21.5	37.	2 ് എ 28.5	31.6
	MEAN	38.0	19.4	37.	\ 1	30.3

Note: There were no significant differences from the pooled control using a one-tailed Dunnett's means comparison procedure (P>.05).

Treatment level 5 not included because of survival effects in the FO generation.

Insufficient number of sexually mature fish with which to form a spawning group. Not included in analysis.

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Table 24. Mean Measured Concentrations of BAY HWG-1608 in Saltwater During the F1 Phase of a Full Life Cycle Toxicity Test with Sheepshead Minnow, Cyprinodon variegatus

Nominal Test Concentration	Rep	Test	Mean Measured Test Concentration (μg/L; ppb)						
(μg/L; ppb)		Repl:	icate	T	Nominal				
Control	A)	<1.3							
	В .	<1.3	,	<1.3					
Solvent Control	A	<1.3							
	В	<1.3		<1.3					
12.5	A	9).8	(1.6)						
	В		(1.4)	9.6	(1.5)	77			
25.0	A	18.5	(2.3)		• •• ••				
	В		(2.2)	18.4	(2.2)	73			
50.0	A	49.3	(17 🕞)						
	В	44.6	(3.7)	46.9	(12.8)	94			
100	A	99.3	(8.8)						
	В		(8.2)	97.8	(8.4)	98			
200	A	194.7	(25.6)	1					
•	В		(14.7)	192.0	(20.6)	96			

NOTE: Values in parentheses are the standard deviations of the means.

Table 25.

Hatching Success (Percent Hatch) of F1 Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 During a Full Life Cycle Toxicity Test

Treatment Gro		Tank		t Hatch	Tank 1	2	
Concentratio		mber		Cham	ber		Treatment
(µg/L; ppb)	1	2	Mean	1	2	Mean	Mean .
Control	7540	96.0	86.7	100.0	100.0	100.0	93.7
Sol. Control	100.0	100.0	100.0	100.0	100.0	100.0	100.0
Pooled Contro	1						96.9
9.6	100.0	100.0	100.0				100.0
18.4	76.0	72.0	74.0	100.0	84.0	92.0	83.0*
46.9	100.0	100.0	100.0	92.0	96.0	94.0	97.0
97.8	100.0	96.0	98.0	100.0	100.0	100.0	99.0

Note:

Initial number = 25 per chamber, except for control, tank A, chamber 1, n=20 and level 1, tank B, for which there were no data available.

*Significantly different (P<.05) from the pooled control using a one-tailed Chi-square and Fisher's exact test.

Treatment level 5 not included because of survival effects in the F0 generation.

Table 26.

Survival of F1 (Day 28 Post-Hatch) Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 at the Termination of a Full Life Cycle Toxicity Test

Treatment Gro	oup - (1		nt Surv		tal Hat	ch)	-
Mean Measu	red	Tank I	A		Tank I	3	
Concentrat:	ion Chai	mber		Cham	ber		Treatment
Group	1	22	Mean	11	2	Mean	Mean
•	36 B						
Control	100.0	91.7	94.9	100.0	92.0	96.0	95.5
	(15/15)	(22/24)		(25/25)(23/25)		
		•					
Sol. Control	72.0	92.0	82.0	84.0	96.0	90.0	86.0
	(18/25)	(23/25)		(21/25)	(24/25)		
		4					•
Pooled Contro	ol	(i)					90.5
9.6	100.0	100.0	100.0	ª			100.0
	(25/25)	(25/25)		(0/0)	(0/0)		•
18.4	94.7	100.0	97.3	92.0	90.5	91.3	94.0
	(18/19)	(18/18)	\bigcirc	(23/25)	(19/21)		
			, ·	<u> </u>			
46.9	96.0	96.0	96.0	100.0	100.0	100.0	97.9
	(24/25)	(24/25)		(23/23)	(24/24)		
				96.0 (24/25)			
97.8	100.0	83.3	91.8	96.0	100.0	98.0	94.9
•	(25/25)	(20/24)		(24/25)	(25/25)		
•	•				•		
•					A		

Note: There were no significant differences from the pooled control using a one-tailed test.

Treatment level 5 not included because of survival effects in the FO generation.

^{*} No eggs were produced for this replicate.

Table 27. Mean (Standard Deviation and n) of Growth (Measured as Length in mm) of F1 Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 at Termination of a Full Life Cycle Toxicity Test

			Mean 1	Length			
Treatment Gro	∍d	Tank A	<u> </u>	www.th.co.co.co.co.co.co.co.co.co.co.co.co.co.	Tank B		
Concentration	on Cha	mber		Char	nber		Treatment
(#a/L: ppb)	11_	2	Mean	1	2	Mean	Mean
Control	(1.9)	12.6 (1.6) (n=22)		(1.2)	14.2 (1.4) (n=23)	14.0	13.5
Sol. Control	(1.2)	13.6 (1.3) (n=23)		(0.9)		14.2	14.0
Pooled Contro	o 1					. .	13.7
9 . 6	13.5 (1.0) (n=25)	(1.0)	13.3	()	a () (n=0)	a	13.3
18.4		13.8 (1.2) (n=18)	13.9	(1.0)	14.7 (0.9) (n=19)	14.4	14.2
46.9	13.3 (1.3) (n=24)	(0.8)	13.2	(1.3)	13.3 (1.3) (n=24)	13.4	13.3
97.8	13.2 (1.3) (n=25)		13.3	(1.3)	12.5 (1.1) (n=25)	12.6	12.9*
						Lan.	

Note: *Significantly different (P<.05) from the pooled control using a one-tailed parametric Dunnett's means comparison procedure.

Treatment level 5 not included because of survival effects in the FO generation.

* Tank B - There were no data available; see Table 26.

Table 28. Mean (Standard Deviation and n) of Growth (Measured as Wet Weight in mg) of F1 Sheepshead Minnows, Cyprinodon variegatus, Exposed to BAY HWG-1608 at Termination of a Full Life Cycle Toxicity Test (28 Days Post-Hatch)

Treatment Group		mamir 3		Wet Weigh			
Mean Measured Concentration				Cham	<u>Tank B</u> ber		Treatment
Concentration	1	2	Mean	1	2	Mean	Mean
Control	(28.8)	53.8 (17.4) (n=22)		(22.6)	83.6 (23.5) (n=23)	78.4	70.1
Sol. Control		64.0 (18.0) (n=23)		(21.9)		80.6	75.2
Pooled Control		•			* ma		72.6
9.6	63.0 (17.6) (n=25)	55.9 (14.7) (n=25)		()	a () (n=0)	ā	59 .5
18.4	(36.5)	80.0 (19.7) (n=18)		87.5 (19.3) (n=23)	(23.9)	93.8	88.9
46.9		62.8 (14.7) (n=24)		(1848)	66.7 (22.2) (n=24)	69.0	67.7
97.8	(19.7)	71.1 (21.1) (n=20)		(15.1)	(12.9) (n=25)	42.2	54.7*

Note: *Significantly different (P<.05) from the pooled control using a one-tailed parametric Dunnett's means comparison procedure.

Treatment level 5 not included because of survival effects in the FO generation.

^{*} Tank B - There were no data available; see Table 26.

TITLE: Sheepshead Minnow Survival Day 33 FILE: a:\surv.d33

TRANSFORM: ARC SINE(SQUARE ROOT(Y)) NUMBER OF GROUPS: 6

GRP	IDENTIFICATION	REP	VALUE	TRANS VALUE
1	GRPS 1&2 POOLED	1	0.8800	1.2171
ī	GRPS 1&2 POOLED		0.7600	1.0588
1	GRPS 1&2 POOLED	3	0.8000	1.1071
1	GRPS 1&2 POOLED .	2 3 4 5	0.7600	1.0588
1	GRPS 1&2 POOLED	5	0.8000	1.1071
1	GRPS 1&2 POOLED	6	0.8400	1.1593
1	GRPS 1&2 POOLED	7	0.9000	1.2490
1	GRPS 1&2 POOLED	8	0.7200	1.0132
2	10.2 ppb	1	0.8400	1.1593
2	10.2 ppb	2	0.8800	1.2171
2 2 3 3 3	10.2 ppb	3	0.8200	1.1326
2	10.2 ppb	4	0.7600	1.0588
3	20.7 ppb	1	0.7600	1.0588
3	20.7 ppb	2	0.8000	1.1071
3	20.7 ppb	3	0.8400	1.1593
	20.7 ppb	4	0.8000	1.1071
4	37.2 ppb	1	0.8800	1.2171
4	37.2 ppb	2	0.8200	1.1326 1.1873
4	37.2 ppb	7 8 1 2 3 4 1 2 3 4 1 2 3 4 1 2 3 4 1 2 3 4 1 2 3	0.8600 0.8400	1.1593
4	37.2 ppb	1± 1	0.8200	1.1326
5 5 5	81.05 ppb 81.05 ppb	2	0.8200	1.1326
5	81.05 ppb 81.05 ppb	3	0.8600	1.1873
5	81.05 ppb 81.05 ppb	4	0.7600	1.0588
6	170.5 ppb	1	0.2000	0.4636
6	170.5 ppb	2	0.1700	0.4250
6	170.5 ppb	3	0.2200	0.4882
6	170.5 ppb	4	0.1800	0.4381
-	FF			

Sheepshead Minnow Survival Day 33

File: a:\surv.d33 Transform: ARC SINE(SQUARE ROOT(Y))

SUMMARY STATISTICS ON TRANSFORMED DATA TABLE 1 of 2

GRP	IDENTIFICATION	N	MIN	MAX	MEAN
1	GRPS 1&2 POOLED 10.2 ppb	8	1.013	1.249 1.217	1.121
3	20.7 ppb	4	1.059	1.159 1.217	1.108 1.174
4 5	37.2 ppb 81.05 ppb	$rac{4}{4}$	1.133 1.059	1.187	1.128
6	170.5 ppb	4	0.425	0.488	0.454

File: a:\surv.d33 Transform: ARC SINE(SQUARE ROOT(Y))

SUMMARY STATISTICS ON TRANSFORMED DATA TABLE 2 of 2

 GRP IDENTIFICATION
 VARIANCE
 SD
 SEM

 1 GRPS 1&2 POOLED
 0.007
 0.082
 0.029

 2 10.2 ppb
 0.004
 0.066
 0.033

 3 20.7 ppb
 0.002
 0.041
 0.021

 4 37.2 ppb
 0.001
 0.036
 0.018

 5 81.05 ppb
 0.003
 0.053
 0.026

 6 170.5 ppb
 0.001
 0.028
 0.014

Sheepshead Minnow Survival Day 33

File: a:\surv.d33 Transform: ARC SINE(SQUARE ROOT(Y))

WILLIAMS T	EST (Isotonic	regression	model)	TABLE	1	OF	2
------------	-------	----------	------------	--------	-------	---	----	---

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1 2 3	GRPS 1&2 POOLED 10.2 ppb 20.7 ppb 37.2 ppb	8 4 4	0.808 0.825 0.800 0.850	1.121 1.142 1.108 1.174	1.133 1.133 1.133 1.133
5	81.05 ppb 170.5 ppb	4	0.815 0.193	1.128	1.128

Sheepshead Minnow Survival Day 33

File: a:\surv.d33 Transform: ARC SINE(SQUARE ROOT(Y))

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
GRPS 1&2 POOLED 10.2 ppb 20.7 ppb 37.2 ppb 81.05 ppb 170.5 ppb	1.133 1.133 1.133 1.133 1.128 0.454	0.327 0.327 0.327 0.178 18.144	*	1.72 1.80 1.83 1.84 1.85	k= 1, v=22 k= 2, v=22 k= 3, v=22 k= 4, v=22 k= 5, v=22

s = 0.060

Note: df used for table values are approximate when v > 20.

TITLE: Sheepshead Minnow Hatching Success FILE: a:\hatch

NUMBER OF GROUPS: 6 TRANSFORM: SQUARE ROOT(Y)

GRP	IDENTIFICATION	REP	VALUE	TRANS VALUE
1	GRPS 1&2 POOLED 10.2 ppb 10.2 ppb 10.2 ppb 20.7 ppb 20.7 ppb 20.7 ppb 20.7 ppb 37.2 ppb 37.2 ppb 37.2 ppb 37.2 ppb 37.2 ppb 37.2 ppb 81.05 ppb 81.05 ppb 81.05 ppb 81.05 ppb 81.05 ppb 81.05 ppb 170.5 ppb 170.5 ppb	1	0.9600	0.9798
ī	GRPS 1&2 POOLED	2	0.8200	
1	GRPS 1&2 POOLED	3	0.8600	0.9274
ī	GRPS 1&2 POOLED	4	0.9200	0.9592
1	GRPS 1&2 POOLED	5	0.8400	
1	GRPS 1&2 POOLED	6		0.9274
1	GRPS 1&2 POOLED	7	0.9400	0.9695
1	GRPS 1&2 POOLED	8	0.8200	0.9055
2	10.2 ppb	1	0.9400	0.9695
2	10.2 ppb	2	0.9800	0.9899
2 2 3 3 3 4	10.2 ppb	3	0.8800 0.8600	0.9381
2	10.2 ppb	4	0.8600	0.9274
3	20.7 ppb	1	0.8200	0.9055
3 .	20.7 ppb	2		0.9274
3	20.7 ppb	3	0.9800	0.9899
3	20.7 ppb	4	0.8400 0.8800 0.8600	0.9165
4	37.2 ppb	1	0.8800	0.9381
4	37.2 ppb	2	0.8600	0.9274
4	37.2 ppb	3	0.9600	0.9/98
4	37.2 ppb	4	0.8800	0.9381
5	81.05 ppb	1	0.9200	0.9592
5 5	81.05 ppb	2	0.8800	0.9381
5	81.05 ppb	3	0.9400	0.9695
5	81.05 ppb	4		0.9165
6	170.5 ppb	1	0.8200	0.9055
6	170.5 ppb	2	0.9000	0.9487
6	170.5 ppb	3	0.9400 0.9200	0.9695
6	170.5 ppb	4	0.9200	0.9592

Sheepshead Minnow Hatching Success

File: a:\hatch Transform: SQUARE ROOT(Y)

SUMMARY STATISTICS ON TRANSFORMED DATA TABLE 1 of 2

GRP	IDENTIFICATION	N	MIN	MAX	MEAN	
1 2 3 4 5	GRPS 1&2 POOLED 10.2 ppb 20.7 ppb 37.2 ppb 81.05 ppb 170.5 ppb	8 4 4 4 4	0.906 0.927 0.906 0.927 0.917 0.906	0.980 0.990 0.990 0.980 0.970	0.936 0.956 0.935 0.946 0.946	

File: a:\hatch Transform: SQUARE ROOT(Y)

SUMMARY STATISTICS ON TRANSFORMED DATA TABLE 2 of 2

GRP	IDENTIFICATION	VARIANCE	SD	SEM
1	GRPS 1&2 POOLED	0.001	0.029	0.010
2	10.2 ppb	0.001	0.029	0.014
. 3	20.7 ppb	0.001	0.038	0.019
4	37.2 ppb	0.001	0.023	0.012
5	81.05 ppb	0.001	0.024	0.012
6	170.5 ppb	0.001	0.028	. 0.014

Sheepshead Minnow Hatching Success
File: a:\hatch Transform: SQUARE ROOT(Y)

	WILLIAMS TEST (Isotor	nic	regression model) TABLE 1 OF	7 2
. GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1 2 3 4 5	GRPS 1&2 POOLED 10.2 ppb 20.7 ppb 37.2 ppb 81.05 ppb 170.5 ppb	8 4 4 4 4	0.878 0.915 0.875 0.895 0.895 0.895	0.936 0.956 0.935 0.946 0.946	0.936 0.946 0.946 0.946 0.946

Sheepshead Minnow Hatching Success
File: a:\hatch Transform: SQUARE ROOT(Y)

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIÄMS	DEGREES OF FREEDOM
GRPS 1&2 POOLED 10.2 ppb 20.7 ppb 37.2 ppb 81.05 ppb 170.5 ppb	0.936 0.946 0.946 0.946 0.946 0.946	0.518 0.518 0.532 0.532 0.532		1.72 1.80 1.83 1.84 1.85	k= 1, v=22 k= 2, v=22 k= 3, v=22 k= 4, v=22 k= 5, v=22

s = 0.029

Note: df used for table values are approximate when v > 20.

```
TRT 1 = Dilution Water Control
 TRT 2 = Solvent Control
 TRT 3 = 9.21 \, \text{ug/l}
 TRT 4 = 19.0 \text{ ug/l}
 TRT 5 = 43.6 \text{ ug/l}
 TRT 6 = 97.0 \text{ ug/l}
 TRT 7 = 197 \text{ ug/l}
 FO LENGTH AT 33 DAYS
 ANOVA on Lengths
 LEVELS ENCOUNTERED DURING PROCESSING ARE:
 TRT
        1.0000
                                                  4.0000
                                                               5.0000
                      2.0000
                                    3.0000
                                                                             6.0000
        7.0000
 REP
        1,0000
                      2,0000
DEP VAR: LENGTH
                             1028 MULTIPLE R: 0.245 SQUARED MULTIPLE R: 0.060
                         ANALYSIS OF VARIANCE
              SUM-OF-SQUARES
 SOURCE
                                DF MEAN-SQUARE
                                                      F-RATIO
                    130.7688
                                                      7.9581
 TRT
                                 6
                                        21.7948
                                                                  0.0000
 REP
                     17.9259
                                 1
                                        17.9259
                                                      6.5455
                                                                  0.0107
 TRT*REP
                     41.2929
                                 6
                                         6.8822
                                                      2.5129
                                                                  0.0203
ERROR 2777.0240 1014
DURBIN-WATSON D STATISTIC 1.8
                                         2.7387
                               1.818
FIRST ORDER AUTOCORRELATION
                                .089
 Post-hoc pairwise comparison of length/Bonferroni.
USING LEAST SQUARES MEANS.
POST HOC TEST OF
                   LENGTH
MATRIX OF PAIRWISE MEAN DIFFERENCES:
                                                   3
                                                                4
                                                                             5
                         0.0000
              2
                         0.1978
                                      0.0000
                         0.1648
                                     -0.0330
                                                  0.0000
                        -0.1290
                                     -0.3268
                                                  -0.2938
                                                               0.0000
                        -0.2197
                                     -0.4175
                                                  -0.3845
                                                              -0.0907
                                                                            0.0000
                        -0.7904
                                     -0.9882
                                                  -0.9552
                                                              -0.6614
                                                                           -0.5707
                        -0.9197
                                     -1.1175
                                                  -1.0845
                                                              -0.7907
                                                                           -0.7000
                                       7
                          6
                         0.0000
              6
              7
                        -0.1293
                                      0.0000
BONFERRONI ADJUSTMENT.
MATRIX OF PAIRWISE COMPARISON PROBABILITIES:
                          1
                                       2
                                                   3
                                                                             5
                         1.0000
                         1.0000
                                      1.0000
              2
              3
                         1.0000
                                      1.0000
                                                  1.0000
                         1.0000
                                      1.0000
                                                               1.0000
                                                   1.0000
              5
                         1.0000
                                      0.4624
                                                  0.7100
                                                               1.0000
                                                                            1.0000
                         0.0004
                                      0.0000
                                                  0.0000
              6
                                                               0.0064
                                                                            0.0358
                         0.0353
                                      0.0029
                                                  0.0045
                                                               0.1425
                                                                            0.3415
                          6
                         1.0000
                         1.0000
                                      1.0000
```

1

TRT 1.0000

TOTAL OBSERVATIONS: 163

LENGTH

N OF CASES 163 MINIMUM 7.0000 16.0000 11.2209 MAXIMUM MEAN STANDARD DEV 1.6998

THE FOLLOWING RESULTS ARE FOR: .

2.0000 TRT

TOTAL OBSERVATIONS: 161

LENGTH

N OF CASES 161 MINIMUM 6.0000 16.0000 11.4161 1.7908 MAXIMUM MEAN STANDARD DEV

THE FOLLOWING RESULTS ARE FOR:
TOT = 3.0000

TOTAL OBSERVATIONS: 165

LENGTH

N OF CASES 165 8.0000 MINIMUM MAXIMUM 14.0000 11.3697 MEAN STANDARD DEV 1.2308

THE FOLLOWING RESULTS ARE FOR:

TOT = 4.0000

TOTAL OBSERVATIONS: 166

LENGTH

N OF CASES 166 6.0000 MINIMUM MAXIMUM 15.0000 MEAN 11.0904 STANDARD DEV 1.3699

TRT = 5.0000

TOTAL OBSERVATIONS: 170

LENGTH

N OF CASES 170 MINIMUM 6,0000 MAXIMUM 14.0000 11.0000 MEAN STANDARD DEV 1.2593

THE FOLLOWING RESULTS ARE FOR:

TRT = 6.0000

TOTAL OBSERVATIONS: 163

LENGTH

N OF CASES 163 5.0000 15.0000 MUNIMUM MUNIXAM MEAN 10.4294 STANDARD DEV 1.9115

THE FOLLOWING RESULTS ARE FOR:
TOT = 7.0000

TOTAL OBSERVATIONS: 40

LENGTH

N OF CASES 40 MINIMUM 5.0000 MAXIMUM 19.0000 MEAN 10.3000 STANDARD DEV 3.3066

SUMMARY STATISTICS FOR LENGTH

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE = 127.4388 DF= 6 PROBABILITY = 0.0000

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES DF MEAN SQUARE F PROBABILITY

BETWEEN GROUPS 129.1288 6 21.5215 7.7766 0.0000

WITHIN GROUPS 2825.59881021 2.7675

1.0000 TRT = 1.0000 REP

TOTAL OBSERVATIONS:

LENGTH

N OF CASES MINIMUM 7.0000 16.0000 MAXIMUM MEAN 11.4024 STANDARD DEV 1.6841

THE FOLLOWING RESULTS ARE FOR:

TRY = 1.0000

2.0000 REP

TOTAL OBSERVATIONS:

LENGTH

N OF CASES MINIMUM 7.0000 15.0000 MAXIMUM MEAN 11.0370 STANDARD DEV 1.7062

THE FOLLOWING RESULTS ARE FOR:

TPT = 2.0000

REP 1.0000

TOTAL OBSERVATIONS:

LENGTH

N OF CASES 6.0000 MINIMUM 16.0000 MAXIMUM MEAN 11.3735 STANDARD DEV 1.6507

THE FOLLOWING RESULTS ARE FOR:

TRT = 2.0000

2.0000

TOTAL OBSERVATIONS: 78

LENGTH

H OF CASES 78 8.0000 16.0000 MINIMUM MAXIMUM MEAN 11.4615 STANDARD DEV 1.9386

THE FOLLOWING RESULTS ARE FOR:
TRT = 3.0000

= 1.0000 REP

TOTAL OBSERVATIONS: 86

LENGTH

N OF CASES MINIMUM 9.0000 13.0000 MUMIXAM 11.0349 MEAN STANDARD DEV 1.1214

THE FOLLOWING RESULTS ARE FOR:

= 3.0000 TRT 2.0000 REP

TOTAL OBSERVATIONS:

LENGTH

N OF CASES MINIMUM 8.0000 MUNIXAM 14.0000 11.7342 1.2477 MEAN STANDARD DEV

THE FOLLOWING RESULTS ARE FOR:

TRT = 4.0000

REP = 1.0000

TOTAL OBSERVATIONS:

LENGTH

N OF CASES MINIMUM 6.0000 MUMIXAM 13.0000 11:0595 MEAN STANDARD DEV 1.2834

THE FOLLOWING RESULTS ARE FOR:

= 4.0000 TRT REP 2.0000

TOTAL OBSERVATIONS: 82

LENGTH

N OF CASES 82 MINIMUM 7.0000 15.0000 11.1220 MAXIMUM MEAN STANDARD DEV 1.4605

5.0000 TRT = 1.0000 = REP

TOTAL OBSERVATIONS: 85

LENGTH

N OF CASES 85 6.0000 MINIMUM MAXIMUM 10.8706 MEAN STANDARD DEV 1.3255

THE FOLLOWING RESULTS ARE FOR:
TRT = 5.0000

REP 2.0000

85 TOTAL OBSERVATIONS:

LENGTH

N OF CASES 85 8.0000 14.0000 MINIMUM MAXIMUM 11.1294 MEAN STANDARD DEV 1.1831

THE FOLLOWING RESULTS ARE FOR:

6.0000 TRT REP

1.0000

TOTAL OBSERVATIONS: 82

LENGTH

N OF CASES 82 6.0000 15.0000 MINIMUM MAXIMUM 10.4512 MEAN STANDARD DEV 1.8735

THE FOLLOWING RESULTS ARE FOR:

= 6.0000 TRT

REP 2.0000

TOTAL OBSERVATIONS: 81

LENGTH

N OF CASES 81 5.0000 MINIMUM MAXIMUM 10.4074 MEAN STANDARD DEV 1.9607

THE FOLLOWING RESULTS ARE FOR:
TOT = 7.0000

1.0000 REP

TOTAL OBSERVATIONS:

20

LENGTH

N OF CASES 20 MINIMUM 5.0000 MAXIMUM 19.0000 MEAN 9.6000 STANDARD DEV 3.5452

THE FOLLOWING RESULTS ARE FOR: 7.0000

2.0000 REP

TOTAL OBSERVATIONS: 20

LENGTH

N OF CASES 20 MINIMUM 7.0000 MAXIMUM 18.0000 MEAN 11.0000 STANDARD DEV 2.9736

SUMMARY STATISTICS FOR LENGTH

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

135.3557 DF= 13 PROBABILITY = CHI-SQUARE = 0.0000

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES OF MEAN SQUARE PROBABILITY

BETWEEN GROUPS 177.7036 13 13.6695 4.9913 0.0000

2777.02401014 WITHIN GROUPS 2.7387

KOLMOGOROV-SMIRNOV ONE SAMPLE TEST USING STANDARD NORMAL DISTRIBUTION

VARIABLE N-OF-CASES MAXDIF PROBABILITY (2-TAIL)

LENGTH 1028.0000 1.0000 0.0000

```
TRT 1 = Dilution Water Control
TRT
     2 = Solvent Control
TRT 3 = 9.21 ug/L
TRT 4 = 19.0 ug/l
TRT 5 = 43.6 ug/l
TRT 6 = 97.0 \, \text{ug/l}
TRT 7 = 197 \text{ ug/l}
FO LENGTH AT 58 DAYS
ANOVA on Lengths
LEVELS ENCOUNTERED DURING PROCESSING ARE:
TRT
       1.0000
                     2.0000
                                   3.0000
                                                 4.0000
                                                               5.0000
                                                                            6.0000
       7.0000
REP
       1.0000
                     2.0000
DEP VAR: LENGTH
                               997 MULTIPLE R: 0.191 SQUARED MULTIPLE R: 0.037
                        ANALYSIS OF VARIANCE
             SUM-OF-SQUARES DF MEAN-SQUARE
SOURCE
                                                     F-RATIO
                                                                    P
                                                     4.2354
                                                                  0.0003
                                       31.5496
                   189, 2975
TRT
                                6
REP
                     9.5438
                                1
                                        9.5438
                                                     1.2812
                                                                  0.2579
TRT*REP
                    78.9553
                                6
                                       13.1592
                                                     1.7666
                                                                  0.1028
                  7322.4497
                             983
                                        7.4491
DURBIN-WATSON D STATISTIC
                               1.665
FIRST ORDER AUTOCORRELATION
                                .162
Post-hoc pairwise comparison of length/Bonferroni.
USING LEAST SQUARES MEANS.
POST HOC TEST OF LENGTH
MATRIX OF PAIRWISE MEAN DIFFERENCES:
                                                   3
                                                                            5
                        0.0000
             2
                        1.2724
                                     0.0000
                                    -0.1972
                                                  0.0000
             3
                        1.0752
                        0.8864
                                    -0.3860
                                                 -0.1888
                                                              0.0000
             5
                        0.7173
                                    -0.5551
                                                 -0.3579
                                                              -0.1691
                                                                           0.0000
                                    -0.7096
                        0.5629
                                                 -0.5124
                                                              -0.3236
                                                                           -0.1545
             6
                                                              -1.3046
                                    -1.6905
                                                                          -1.1354
             7
                        -0.4181
                                                 -1.4933
                         6
                        0.0000
             6
7
                        -0.9810
                                     0.0000
BONFERRONI ADJUSTMENT.
MATRIX OF PAIRWISE COMPARISON PROBABILITIES:
                                      2
                                                   3
                                                                            5
                        1.0000
             2
                        0.0007
                                     1.0000
             3
                        0.0078
                                     1.0000
                                                  1.0000
                        0.0768
                                     1.0000
                                                  1.0000
                                                               1.0000
                                     1,0000
                                                  1,0000
                                                               1.0000
                                                                           1,0000
             5
                        0.3542
                                                               1.0000
                                                                           1.0000
                                     0.4431
                                                  1.0000
                        1,0000
             6
7
                                     0.1559
                                                  0.3709
                                                               0.8108
                                                                           1.0000
                        1,0000
                         6
             6
                        1.0000
                        1.0000
                                     1.0000
```

9

THE FOLLOWING RESULTS ARE FOR:

TRT = 1.0000

TOTAL OBSERVATIONS: 162

LENGTH

N OF CASES 162 10.0000 MUNIMUM MAXIMUM 27.0000 17.0556 MEAN 2.6616 STANDARD DEV

THE FOLLOWING RESULTS ARE FOR:
TRT = 2.0000

TOTAL OBSERVATIONS: 159

LENGTH

N OF CASES MINIMUM 8.0000 **MAXIMUM** 25.0000 18.3208 MEAN STANDARD DEV 2.9324

THE FOLLOWING RESULTS ARE FOR:
TRT = 3.0000

TOTAL OBSERVATIONS: 167

LENGTH

N OF CASES 167 MINIMUM 12.0000 23.0000 HUMIXAM MEAN 18.1317 STANDARD DEV 2.2643

THE FOLLOWING RESULTS ARE FOR:

= 4.0000 TRT

TOTAL OBSERVATIONS: 160

LENGTH

N OF CASES 160 10.0000 MINIMUM MAXIMUM 24.0000 MEAN-17.9375 2.6006 STANDARD DEV

TRT = 5.0000

TOTAL OBSERVATIONS: 170

LENGTH

 N OF CASES
 170

 MININUM
 12.0000

 MAXINUM
 25.0000

 MEAN
 17.7765

 STANDARD DEV
 2.2968

THE FOLLOWING RESULTS ARE FOR:

TRT = 6.0000

TOTAL OBSERVATIONS: 157

LENGTH

 N OF CASES
 157

 MINIMUM
 9.0000

 MAXIMUM
 24.0000

 MEAN
 17.6242

 STANDARD DEV
 3.0727

THE FOLLOWING RESULTS ARE FOR:

TRT = 7.0000

TOTAL OBSERVATIONS: 22

LENGTH

 N OF CASES
 22

 MINIMUM
 10.0000

 MAXIMUM
 27.0000

 MEAN
 16.4545

 STANDARD DEV
 5.3960

SUMMARY STATISTICS FOR LENGTH

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE = 58.8147 DF= 6 PROBABILITY = 0.0000

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES DF MEAN SQUARE F PROBABILITY

BETWEEN GROUPS 198.6053 6 33.1009 4.4275 0.0002 WITHIN GROUPS 7401.4068 990 7.4762

THE FOLLOWING RESULTS ARE FOR:

TRT = 1.0000

1.0000

82 TOTAL OBSERVATIONS:

LENGTH

N OF CASES 82 MINIMUM 10.0000 27,0000 MAXIMUM MEAN 16.7683 STANDARD DEV 2,5834

THE FOLLOWING RESULTS ARE FOR:

TPT = 1.0000

2.0000

TOTAL OBSERVATIONS: 80

LENGTH

N OF CASES 80 MINIMUM 11.0000 MAXIMUM 25.0000 MEAN 17.3500 STANDARD DEV 2.7240

THE FOLLOWING RESULTS ARE FOR:

TRT = 2.0000 REP 1.0000

TOTAL OBSERVATIONS: 82

LENGTH

N OF CASES 82 MINIMUM 8.0000 MAXIMUM 25.0000 MEAN 17.9878 STANDARD DEV 3.0449

THE FOLLOWING RESULTS ARE FOR:

= 2.0000 TRT REP 2.0000

TOTAL OBSERVATIONS: 77

LENGTH

N OF CASES 77 MINIMUM 13.0000 MAXIMUM 25.0000 MEAN. 18.6753 STANDARD DEV 2.7836

3.0000 TRT REP 1.0000

TOTAL OBSERVATIONS:

LENGTH

N OF CASES MINIMUM 12.0000 MAXIMUM 23,0000 MEAN 18.0465 2.3106 STANDARD DEV

REP 2.0000

TOTAL OBSERVATIONS: 81

LENGTH

N OF CASES 81 12.0000 23.0000 MINIMUM MAXIMUM MEÁN 18.2222 STANDARD DEV 2.2249

THE FOLLOWING RESULTS ARE FOR:

4.0000 TRT 1.0000

REP

TOTAL OBSERVATIONS: 78

LENGTH

N OF CASES 78 11.0000 24.0000 MINIMUM MAXIMUM MEAN 18.2692 STANDARD DEV 2.2371

THE FOLLOWING RESULTS ARE FOR:

4.0000 TRT

2.0000

TOTAL OBSERVATIONS: 82

LENGTH

N OF CASES 82 MINIMUM 10,0000 23.0000 17.6220 MAXIMUM MEAN STANDARD DEV 2.8831

THE FOLLOWING RESULTS ARE FOR:
TRT = 5.0000 REP 1.0000

TOTAL OBSERVATIONS: 85

LENGTH

N OF CASES MINIMUM 12.0000 MAXIMUM 25.0000 MEAN 17.9294 STANDARD DEV 2.5856

THE FOLLOWING RESULTS ARE FOR: TRT = 5.0000

2.0000

TOTAL OBSERVATIONS: 85

LENGTH

N OF CASES 85 MINIMUM 12.0000 MAXIMUM 22.0000 MEAN 17.6235 STANDARD DEV 1.9699

THE FOLLOWING RESULTS ARE FOR:

6.0000 TRT REP 1.0000

TOTAL OBSERVATIONS: 80

LENGTH

N OF CASES MUNIMUM 12.0000 MAXIMUM 24.0000 17.7375 MEAN STANDARD DEV 3.0883

THE FOLLOWING RESULTS ARE FOR:

TRT 6.0000 REP 2.0000

TOTAL OBSERVATIONS: 77

LENGTH

N OF CASES MINIMUM 9.0000 MAXIMUM 22.0000 MEAN 17.5065 STANDARD DEV 3.0720

THE FOLLOWING RESULTS ARE FOR:

TRT = 7.0000 REP = 1.0000

TOTAL OBSERVATIONS:

LENGTH

N OF CASES 9
MINIMUM 10.0000
MAXIMUM 25.0000
MEAN 17.6667
STANDARD DEV 5.2915

THE FOLLOWING RESULTS ARE FOR:

TRT = 7.0000

REP = 2.0000

TOTAL OBSERVATIONS: 13

LENGTH

N OF CASES 13
MINIMUM 10.0000
MAXIMUM 27.0000
MEAN 15.6154
STANDARD DEV 5.5157

SUMMARY STATISTICS FOR LENGTH

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE = 69.4818 DF= 13 PROBABILITY = 0.0000

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES DF MEAN SQUARE F PROBABILITY

BETWEEN GROUPS 277.5624 13 21.3510 2.8663 0.0004 WITHIN GROUPS 7322.4497 983 7.4491

KOLMOGOROV-SMIRNOV ONE SAMPLE TEST USING STANDARD NORMAL DISTRIBUTION

VARIABLE N-OF-CASES MAXDIF PROBABILITY (2-TAIL)

LENGTH 997.0000 1.0000 0.0000

```
TRT 1 = Dilution Water Control
TRT 2 = Solvent Control
TRT
     3 = 9.21 \, \text{ug/l}
     4 = 19.0 \text{ ug/l}
TRT
    5 = 43.6 \text{ ug/l}
TRT
TRT 6 = 97.0 \text{ ug/l}
TRT 7 = 197 \, \text{ug/l}
FO WEIGHT AT 58 DAYS
ANOVA on Weight
LEVELS ENCOUNTERED DURING PROCESSING ARE:
TRT
        1.0000
                     2.0000
                                   3.0000
                                                 4.0000
                                                              5.0000
                                                                            6.0000
REP
       1.0000
                     2.0000
                               675 MULTIPLE R: 0.203 SQUARED MULTIPLE R: 0.041
DEP VAR: WEIGHT
                       N:
                        ANALYSIS OF VARIANCE
SOURCE
              SUM-OF-SQUARES
                              DF
                                   MEAN-SQUARE
                                                     F-RATIO
                102865.8475
                                    20573.1695
                                                     3.3722
                                                                  0.0051
TRT
                                    1568.9061
13907.4474
                  1568.9061
REP
                                                     0.2572
                                                                  0.6122
TRT*REP
                 69537.2372
                                                                  0.0453
                                                     2.2796
ERROR
                .404479E+07
                              663
                                     6100.7379
                               1.986
DURBIN-WATSON D STATISTIC
FIRST ORDER AUTOCORRELATION
                                .005
Post-hoc pairwise comparison of length/Bonferroni.
USING LEAST SQUARES MEANS.
POST HOC TEST OF WEIGHT
MATRIX OF PAIRWISE MEAN DIFFERENCES:
                                                   3
                                                                            5
                                                                4
                        0.0000
             2
                       35.1969
                                     0.0000
             3
                       31.0524
                                    -4.1444
                                                  0.0000
             4 5
                                                              0.0000
                       25.4257
                                    -9.7712
                                                 -5.6268
                       16.1627
                                   -19.0341
                                                -14.8897
                                                              -9.2630
                                                                           0.0000
             6
                       34.3659
                                    -0.8310
                                                  3.3134
                                                              8.9402
                                                                          18.2031
                         6
             6
                        0.0000
BONFERRONI ADJUSTMENT.
MATRIX OF PAIRWISE COMPARISON PROBABILITIES:
                                                   3
                                                                4
                                                                            5
                         1
                                      2
                        1.0000
             2
                        0.0129
                                     1.0000
             3
                                     1.0000
                        0.0412
                                                  1.0000
                        0.2343
                                                  1.0000
                                     1.0000
                                                              1.0000
             5
                        1.0000
                                     0.9916
                                                  1.0000
                                                              1.0000
                                                                           1.0000
             6
                        0.0179
                                     1.0000
                                                  1.0000
                                                              1.0000
                                                                           1.0000
                         6
             6
                        1.0000
```

THE FOLLOWING RESULTS ARE FOR:

TOT = 1.0000

TOTAL OBSERVATIONS: 112

WEIGHT

N OF CASES 37.0000 MINIMUM MAXIMUM 459.0000 159.1696 MEAN 74.4602 STANDARD DEV

THE FOLLOWING RESULTS ARE FOR:

TOT = 2.0000

TOTAL OBSERVATIONS: 109

WEIGHT

N OF CASES 109 MINIMUM 51.0000 466.0000 MAXIMUM 194.2110 MEAN STANDARD DEV 86.1174

THE FOLLOWING RESULTS ARE FOR:

TDT = 3.0000

TOTAL OBSERVATIONS: 117

WEIGHT

117 N OF CASES MINIMUM 51.0000 431.0000 MAXIMUM 190.0000 MEAN STANDARD DEV 76.4768

THE FOLLOWING RESULTS ARE FOR:

TOT = 4.0000

110 TOTAL OBSERVATIONS:

WEIGHT

N OF CASES 110 38.0000 387.0000 MINIMUM MAXIMUM 184.4273 MEAN STANDARD DEV 74.8994

TRY = 5.0000

TOTAL OBSERVATIONS: 120

WEIGHT

N OF CASES 120 MINIMUM 49.0000 MAXIMUM 473.0000 MEAN 175.5750 STANDARD DEV 71.5292

THE FOLLOWING RESULTS ARE FOR:

TRT = 6.0000

TOTAL OBSERVATIONS: 107

WEIGHT

N OF CASES 107
MININUM 29.0000
MAXIMUM 410.0000
MEAN 193.8318
STANDARD DEV 86.9095

SUMMARY STATISTICS FOR WEIGHT

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE = 7.4541 DF= 5 PROBABILITY = 0.1890

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES DF MEAN SQUARE F PROBABILITY

BETMEEN GROUPS 102368.8109 5 20473.7622 3.3279 0.0056 WITHIN GROUPS .411580E+07 669 6152.1691

THE FOLLOWING RESULTS ARE FOR:

TRT = 1.0000

REP = 1.0000

TOTAL OBSERVATIONS: 57

WEIGHT

N OF CASES 57
MINIMUM 67.0000
MAXIMUM 314.0000
MEAN 145.8246
STANDARD DEV 56.0131

TRT = 1.0000 = 2.0000 REP

TOTAL OBSERVATIONS: 55

WEIGHT

55 37.0000 N OF CASES MINIMUM MUMIXAM 459.0000 173.0000 MEAN 88.0919 STANDARD DEV

THE FOLLOWING RESULTS ARE FOR: TRT = 2.0000 REP \cdot = 1.0000

TOTAL OBSERVATIONS:

WEIGHT

57 N OF CASES 51.0000 MINIMUM 376.0000 MUMIXAM 185.9298 MEAN 74.0265 STANDARD DEV

THE FOLLOWING RESULTS ARE FOR:
TRT = 2.0000

REP 2.0000

TOTAL OBSERVATIONS: 52

WEIGHT

N OF CASES 52 MINIHUM 55.0000 466.0000 MUMIXAM MEAN 203.2885 STANDARD DEV 97.6067

THE FOLLOWING RESULTS ARE FOR: TRT = 3.0000

REP 1.0000

TOTAL OBSERVATIONS: 61

WEIGHT

N OF CASES 61 51.0000 MINIMUM 406.0000 MUMIXAM 179.5902 MEAN STANDARD DEV 77.2969

3.0000 TRT = 2.0000 REP

TOTAL OBSERVATIONS:

WEIGHT

N OF CASES MINIMUM 65.0000 MAXIMUM 431.0000 201.3393 MEAN STANDARD DEV 74.6081

THE FOLLOWING RESULTS ARE FOR:

TRT = 4.0000 1.0000 REP

TOTAL OBSERVATIONS:

WEIGHT

N OF CASES MINIMUM 38.0000 387.0000 MAXIMUM 196.1321 MEAN STANDARD DEV 68.6129

THE FOLLOWING RESULTS ARE FOR:

TRT = 4.0000 REP 2.0000

TOTAL OBSERVATIONS: 57

WEIGHT

N OF CASES 39.0000 329.0000 MINIMUM MAXIMUM MEAN 173.5439 STANDARD DEV 79.3575

THE FOLLOWING RESULTS ARE FOR:

5.0000 TRT = REP 1.0000

TOTAL OBSERVATIONS:

WEIGHT

N OF CASES 60 MINIMUM 49.0000 MAXIMUM 473.0000 MEAN 186,3500 STANDARD DEV 82.2002

5.0000 TRT = = REP 2.0000

TOTAL OBSERVATIONS: 60

WEIGHT

N OF CASES MINIMUM 68.0000 MAXIMUM 298.0000 164.8000 MEAN STANDARD DEV 57.6764

THE FOLLOWING RESULTS ARE FOR:

6.0000 TRT REP 1.0000

TOTAL OBSERVATIONS: 55

WEIGHT

N OF CASES MINIMUM 29.0000 MAXIMUM 410.0000 195.6909 MEAN STANDARD DEV 90.8888

THE FOLLOWING RESULTS ARE FOR:

6.0000 TRT =

REP 2.0000

TOTAL OBSERVATIONS: 52

WEIGHT

N OF CASES MINIMUM 52.0000 MAXIMUM 346.0000 191.8654 MEAN STANDARD DEV 83.3340

SUMMARY STATISTICS FOR WEIGHT

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

32.0878 DF= 11 PROBABILITY = 0.0007 CHI-SQUARE =

ANALYSIS OF VARIANCE

SUM OF SQUARES OF MEAN SQUARE SOURCE

PROBABILITY

173380.7350 11 15761.8850 BETWEEN GROUPS .404479E+07 663 6100.7379 WITHIN GROUPS

2.5836 0.0032

KOLMOGOROV-SMIRNOV ONE SAMPLE TEST USING STANDARD NORMAL DISTRIBUTION

VARIABLE N-OF-CASES MAXDIF PROBABILITY (2-TAIL)

675.0000 1.0000 0.0000 WEIGHT

```
21
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```
TRT 1 = Dilution Water Control
TRT 2 = Solvent Control
TRT 3 = 9.21 \text{ ug/l}
TRT 4 = 19.0 \text{ ug/l}
TRT 5 = 43.6 \text{ ug/l}
TRT 6 = 97.0 ug/l
TRT 7 = 197 ug/l
FO LENGTH AT DAY 177
ANOVA on Lengths
LEVELS ENCOUNTERED DURING PROCESSING ARE:
TRT
       1.0000
                     2.0000
                                   3.0000
                                                 4.0000
                                                              5.0000
                                                                            6.0000
REP
       1.0000
                     2.0000
                               206 MULTIPLE R: 0.507 SQUARED MULTIPLE R: 0.257
DEP VAR: LENGTH
                        ANALYSIS OF VARIANCE
SOURCE
              SUM-OF-SQUARES
                               DF MEAN-SQUARE
                                                     F-RATIO
                                        1.0973
                                                     5,6393
                                                                 0.0001
TRT
                     5.4865
                                                                 0.0039
REP
                     1.6583
                                        1.6583
                                                     8.5226
TRT*REP
                     6.3963
                                5
                                        1.2793
                                                     6.5746
                                                                 0.0000
ERROR
                    37.7482
                                        0.1946
                               1.730
DURBIN-WATSON D STATISTIC
FIRST ORDER AUTOCORRELATION
                               .132
Post-hoc pairwise comparison of length/Bonferroni.
USING LEAST SQUARES MEANS.
POST HOC TEST OF
                   LENGTH
MATRIX OF PAIRWISE MEAN DIFFERENCES:
                                                   3
                                                               4
                                                                            5
                        0.0000
             2
                       -0.4049
                                     0.0000
                       -0.5427
                                    -0.1378
                                                  0.0000
             4
5
                       -0.2615
                                                  0.2812
                                     0.1434
                                                              0.0000
                                     0.1830
                                                              0.0396
                                                                           0.0000
                       -0.2220
                                                  0.3207
                       -0.3794
                                     0.0255
                                                                          -0.1574
             ٠6
                                                  0.1633
                                                              -0.1178
                         6
                        0.0000
             6
BONFERRONI ADJUSTMENT.
MATRIX OF PAIRWISE COMPARISON PROBABILITIES:
                                                   3
                                                                            5
                         1
                                      2
             1
                        1.0000
             2
3
                        0.0202
                                     1.0000
                        0.0000
                                     1.0000
                                                  1.0000
             4
                        0.1393
                                     1.0000
                                                  0.1627
                                                              1.0000
             5
                        0.4530
                                     1.0000
                                                  0.0656
                                                              1.0000
                                                                           1.0000
                        0.0138
             6
                                     1.0000
                                                  1.0000
                                                              1.0000
                                                                           1.0000
                        1.0000
             6
```

ANOVA on Weight LEVELS ENCOUNTERED DURING PROCESSING ARE: TRT					
1.0000 REP	2.0000	3.0000	4.0000	5.0000	6.0000
1.0000	2.0000				
DEP VAR: WEIGH		06 MULTIPLE R: S OF VARIANCE	0.474 SQU	ARED MULTIPLE	R: 0.224
SOURCE SUI	M-OF-SQUARES	DF MEAN-SQUARE	F-RATI	0 P	
TRT	9.1649	5 1.8330	4.2985	0.0010	
REP	1.8895	1 1.8895	4.4311		
TRT*REP	12.2392	5 2.4478	5.7404	0.0001	
ERROR	82.7255 19	4 0.4264	,		
DURBIN-WATSON D	STATISTIC 1	.824			
FIRST ORDER AUTO	OCORRELATION	.085			
Post-hoc pairwis USING LEAST SQUA POST HOC TEST OF	ARES MÉANS.	f Weight/ Bonfe	rroni.		
MATRIX OF PAIRW	ISE MEAN DIFFER	ENCES:			
	1	2	3	4	5
1	0.0000				
2	-0.4244	0.0000			
3 4	-0.6931	-0.2687	0.0000		•
4	-0.3539	0.0705	0.3391	0.0000	
5	-0.4528	-0.0284	0.2403	-0.0989	0.0000
6	-0.5602	-0.1358	0.1328	-0.2063	-0.1074
	6				
6	0.0000				
BONFERRONI ADJUS MATRIX OF PAIRW		PROBABILITIES:			
	1	2	3	4	5
4	1.0000				
1	0.3351	1 0000			
2		1.0000	1 0000		
3 4	0.0006	1.0000	1.0000 0.5608	1.0000	
4	0.2587	1.0000	1.0000	1.0000	1.0000
5	0.0445	1.0000			
6	0.0142	1.0000	1.0000	1.0000	1.0000
	6		•		
6	1.0000				

1.0000 TRT =

TOTAL OBSERVATIONS: 38

LENGTH WEIGHT N OF CASES 38 38 0.6300 3.5000 1.7811 MINIMUM 2.7000 MUMIXAM 4.5000 MEAN 3.4921 STANDARD DEV 0.4420 0.7015

THE FOLLOWING RESULTS ARE FOR:

2.0000 TRT . =

TOTAL OBSERVATIONS: 28

LENGTH WEIGHT N OF CASES 28 28 1.9000 0.2200 MINIMUM 4.2000 MAXIMUM 3.1500 MEAN 3.3536 1.6879 STANDARD DEV 0.6280 0.8627

THE FOLLOWING RESULTS ARE FOR:
TRT = 3.0000

TOTAL OBSERVATIONS: 27

LENGTH N OF CASES 27 27 2.2000 0.2800 MINIMUM MAXIMUM 3.9000 2.2900 2.9407 MEAN 1.0759 STANDARD DEV 0.4627 0.6374

THE FOLLOWING RESULTS ARE FOR:

4.0000 TRT

TOTAL OBSERVATIONS:

LENGTH WEIGHT N OF CASES MINIMUM 2.3000 0.4200 MUMIXAM 4.2000 3.8000 MEAN 3.2381 1.4474 STANDARD DEV 0.4333 0.7438

TRT = 5.0000

TOTAL OBSERVATIONS: 3

LENGTH WEIGHT N OF CASES 39 39 MINIMUM 2.3000 0.4400. 4.1000 2.7200 MUMIXAM 3.3051 1.3549 MEAN STANDARD DEV 0.4436 0.5625

THE FOLLOWING RESULTS ARE FOR:

TRT = 6.0000

TOTAL OBSERVATIONS: 32

N OF CASES

STANDARD DEV

MINIMUM MUMIXAM

MEAN

LENGTH WEIGHT

32 32
2.0000 0.2800
4.0000 2.7600
3.1656 1.3441
0.4674 0.6440

SUMMARY STATISTICS FOR LENGTH

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE = 6.4456 DF= 5 PROBABILITY = 0.2652

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES DF MEAN SQUARE F PROBABILITY

BETWEEN GROUPS 5.4269 5 1.0854 4.7822 0.0004 WITHIN GROUPS 45.3927 200 0.2270

SUMMARY STATISTICS FOR WEIGHT

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE = 6.9227 DF= 5 PROBABILITY = 0.2265

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES OF MEAN SQUARE F PROBABILITY

BETWEEN GROUPS 10.2212 5 2.0442 4.2397 0.0011 WITHIN GROUPS 96.4328 200 0.4822

TRT = 1.0000 REP = 1.0000

TOTAL OBSERVATIONS: 17

LENGTH WEIGHT N OF CASES 17 17 2.9000 0.9700 MINIMUM 4.0000 MAXIMUM 2.6500 MEAN 3.4118 1.6524 STANDARD DEV 0.3180 0.4939

THE FOLLOWING RESULTS ARE FOR:

TRT = 1.0000 REP = 2.0000

TOTAL OBSERVATIONS: 21

LENGTH WEIGHT N OF CASES 21 MINIMUM 2.7000 0.6300 MAXIMUM 4.5000 3.5000 MEAN 3.5571 1.8852 STANDARD DEV 0.5202 0.8306

THE FOLLOWING RESULTS ARE FOR:

TRT = 2.0000 REP = 1.0000

TOTAL OBSERVATIONS: 22

LENGTH WEIGHT N OF CASES 22 MINIMUM 2.7000 0.8600 4.2000 3.1500 MAXIMUM 3.5591 MEAN 1.9455 STANDARD DEV 0.4382 0.7329

THE FOLLOWING RESULTS ARE FOR:

TRT = 2.0000

REP = 2.0000

TOTAL OBSERVATIONS: 6

 LENGTH
 WEIGHT

 N OF CASES
 6
 6

 MINIMUM
 1.9000
 0.2200

 MAXIMUM
 3.7000
 1.8400

 MEAN
 2.6000
 0.7433

 STANDARD DEV
 0.6753
 0.6327

3.0000 TRT = 1.0000 REP

TOTAL OBSERVATIONS:

LENGTH WEIGHT N OF CASES MINIMUM 2.2000 0.2800 MAXIMUM 3.9000 2.2900 1.0807 MEAN 2.9143 STANDARD DEV 0.5021 0.6476

THE FOLLOWING RESULTS ARE FOR:

3.0000 TRT × REP 2.0000

TOTAL OBSERVATIONS: 13

LENGTH WEIGHT N OF CASES 13 2.4000 MINIMUM 0.2900 MAXIMUM 3.6000 2.0000 MEAN 2.9692 1.0708 STANDARD DEV 0.4347 0.6527

THE FOLLOWING RESULTS ARE FOR:

TRT 4.0000 1.0000 REP

TOTAL OBSERVATIONS: 24

LENGTH WEIGHT N OF CASES . 24 2.7000 MUNIMUM 0.4200 4.2000 3.8000 **MAXIMUM** 3.3292 1.6425 MEAN STANDARD DEV 0.8294 0.4278

THE FOLLOWING RESULTS ARE FOR:

TDY = 4.0000

2.0000 REP

TOTAL OBSERVATIONS:

LENGTH WEIGHT N OF CASES 18 18 2.3000 MINIMUM 0.4200 MAXIMUM 3.8000 2.4400 MEAN 3.1167 1.1872 STANDARD DEV 0.5274 0.4218

THE FOLLOWING RESULTS ARE FOR:

TRT = 5.0000 1.0000

TOTAL OBSERVATIONS: 23

,	LENGTH	WEIGHT
N OF CASES	23	23
MINIMUM	2.9000	0.8400
MAXIMUM	4.1000	2.7200
MEAN	3.5000	1.5326
STANDARD DEV	0.3303	0.4979

THE FOLLOWING RESULTS ARE FOR:
TRT = 5.0000
REP = 2.0000

TOTAL OBSERVATIONS: 16

	LENGTH	WEIGHT
N OF CASES	16	16
MINIMUM	2.3000	0.4400
MAXIMUM	3.8000	2.3400
MEAN	3.0250	1.0994
STANDARD DEV	0.4435	0.5656

THE FOLLOWING RESULTS ARE FOR:

TRT = 6.0000

REP = 1.0000

TOTAL OBSERVATIONS: 9

	LENGTH	WEIGHT
N OF CASES	9	9
MINIMUM	2.3000	0.2900
MAXIMUM	3.8000	2.0700
MEAN	2.9667	0.8989
STANDARD DEV	0.4924	0.5752

THE FOLLOWING RESULTS ARE FOR:

TRT = 6.0000REP 2.0000

TOTAL OBSERVATIONS: 23

	LENGTH	WEIGHT
N OF CASES	23	23
MINIMUM	2.0000`	0.2800
MAXIMUM	4.0000	2.7600
MEAN	3.2435	1.5183
STANDARD DEV	0.4440	0.5928

SUMMARY STATISTICS FOR LENGTH

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE =

10.0582 DF= 11 PROBABILITY =

0.5252

- ANALYSIS OF VARIANCE

1.1883

0.1946

SOURCE

SUM OF SQUARES DF MEAN SQUARE

PROBABILITY

BETWEEN GROUPS WITHIN GROUPS

13.0714 11 37.7482 194 6.1071

0.0000

SUMMARY STATISTICS FOR WEIGHT

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE =

13.4413 DF= 11 PROBABILITY =

0.2655

5.1014

ANALYSIS OF VARIANCE

SOURCE

SUM OF SQUARES OF MEAN SQUARE

PROBABILITY

BETWEEN GROUPS WITHIN GROUPS

23.9286 11 82.7255 194 2.1753 0.4264

0.0000

KOLMOGOROV-SMIRNOV ONE SAMPLE TEST USING STANDARD NORMAL DISTRIBUTION

VARIABLE

N-OF-CASES MAXDIF PROBABILITY (2-TAIL)

WEIGHT LENGTH 206.0000 206.0000 0.6434 0.9724

0.0000 0.0000

```
TRT 1 = Dilution Water Control
```

TRT 2 = Solvent Control

TRT $3 = 9.21 \, \text{ug/l}$

TRT 4 = 19.0 ug/l

TRT 5 = 43.6 ug/l

TRT 6 = 97.0 ug/l

TRT 7 = 197 ug/l

1.0000

F1 LENGTH AND WEIGHT AT TEST TERMINATION ANOVA on Weights of F1 fish at test termination

2.0000

6 1.0000

LEVELS ENCOUNTERED DURING PROCESSING ARE:

TRT

DEP VAR:	WEIGHT		38 MULTIPLE R: S OF VARIANCE	0.449 SQ	UARED MULTIP	LE R: 0.202
SOURCE	SUM-OF	-SQUARES [F MEAN-SQUARE	F-RAT	10 P	
TRT	578	396.3560	11579.2712	24.382	1 0.000	0
	2289 Atson D Sta Der Autocoa		2 474.9077 .799 .098			
	ST SQUARES		f weight/Bonfer	roni.		
MATRIX OF	PAIRWISE	MEAN DIFFERE	NCES:			
		1	2	3	4	5
	1	0.0000				
	2	5.0453	0.0000			
	2 3	-10.6502	-15.6955	0.0000		
	4	18.7738	13.7286	29.4241	0.0000	
	5	-2.42 3 0	-7.4682	8.2273	-21.1968	0.0000
	6	-15.3614	-20.4067	-4.7112	-34.1352	-12.9385
		6				
	6	0.0000				
BONFERRON	I ADJUSTME					
			ROBABILITIES:			
		1	2	3	4	5
	1	1.0000				
	2	1.0000	1.0000			
	3	0.0950	0.0009	1.0000		
	4	0.0000	0.0010	0.0000	1.0000	
	5	1.0000	0.3261	0.4680	0.0000	1.0000
	6	0.0000	0.D000	1.0000	0.0000	0.0008
		-	*			· · -

3.0000

4.0000

5.0000

6.0000

ANOVA on Lengths of F1 fish at test termination LEVELS ENCOUNTERED DURING PROCESSING ARE:

TRT 2.0000 3.0000 4.0000 5.0000 6.0000 1.0000 DEP VAR: LENGTH 488 MULTIPLE R: 0.330 SQUARED MULTIPLE R: 0.109 ANALYSIS OF VARIANCE SUM-OF-SQUARES OF MEAN-SQUARE Ρ SOURCE F-RATIO 93.9243 18.7849 0.0000 TRT 5 11.7577 ERROR 770.0737 482 1.5977 DURBIN-WATSON D STATISTIC FIRST ORDER AUTOCORRELATION 1.859 .066 Post-hoc pairwise comparison of length/Bonferroni. USING LEAST SQUARES MEANS. POST HOC TEST OF LENGTH MATRIX OF PAIRWISE MEAN DIFFERENCES: 3 4 5 1 0.0000 0.4476 0.0000 23456 -0.2459 -0.6935 0.0000 0.6608 0.2132 0.9067 0.0000 -0.2427 0.0032 -0.9035 0.0000 -0.6903 -0.3589 -0.6016 -1.0492 -0.3557 -1.2624 6 6 BONFERRONI ADJUSTMENT. MATRIX OF PAIRWISE COMPARISON PROBABILITIES: 5 3 2 1.0000 2 3 0.3152 1.0000 1.0000 0.0323 1.0000 0.0138 4 1.0000 0.0013 1.0000 0.0001 0.0041 1.0000 1.0000 1.0000 0.0000 0.0000 0.0235 0.7732 6 6 1.0000 6

TOTAL OBSERVATIONS: 85

	WEIGHT	LENGTH
N OF CASES	85	85
MINIMUM	13.4000	9.0000
MAXIMUM	136.8000	17.0000
MEAN	70.1082	13.5059
STANDARD DEV	25.0776	1,6008

THE FOLLOWING RESULTS ARE FOR: TRT = 2.0000

TOTAL OBSERVATIONS: 86

	WEIGHT	LENGTH
N OF CASES	86	86
MINIMUM	28.8000	11.0000
MAXIMUM	126.6000	16.0000
MEAN	75.1535	13.9535
STANDARD DEV	19.4249	1.1051

THE FOLLOWING RESULTS ARE FOR:
TRT = 3.0000

TOTAL OBSERVATIONS: 50

	WE I GHT	LENGTH
N OF CASES	50	50
MINIMUM	10.8000	10.0000
MAXIMUM	96.6000	15.0000
MEAN	59.4580	13.2600
STANDARD DEV	16.4066	1.0461

THE FOLLOWING RESULTS ARE FOR:
TRT = 4.0000

TOTAL OBSERVATIONS: 78

	WEIGHT	LENGTH
N OF CASES	78	78
MINIMUM	28.5000	11,0000
MAXIMUM	171.8000	16.0000
MEAN	88.8821	14.1667
STANDARD DEV	26.1082	1.2105

TRT = 5.0000

95

TOTAL OBSERVATIONS:

WEIGHT LENGTH

N OF CASES 95 95
MINIMUM 17.5000 10.0000
MAXIMUM 112.6000 16.0000
MEAN 67.6853 13.2632
STANDARD DEV 19.2520 1.2047

THE FOLLOWING RESULTS ARE FOR:

rrt = 6.0000

TOTAL OBSERVATIONS: 9

WEIGHT LENGTH

N OF CASES 94 94
MINIMUM 15.0000 10.0000
MAXIMUM 112.8000 15.0000
MEAN 54.7468 12.9043
STANDARD DEV 21.6249 1.2621

SUMMARY STATISTICS FOR WEIGHT

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE = 20.5860 DF= 5 PROBABILITY = 0.0010

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES OF MEAN SQUARE F PROBABILITY

BETWEEN GROUPS 57896.3560 5 11579.2712 24.3821 0.0000 WITHIN GROUPS 228905.5283 482 474.9077

SUMMARY STATISTICS FOR LENGTH

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE = 17.5673 DF= 5 PROBABILITY = 0.0035

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES OF MEAN SQUARE F PROBABILITY

BETWEEN GROUPS 93.9243 5 18.7849 11.7577 0.0000 WITHIN GROUPS 770.0737 482 1.5977

TRT 1,0000 = 1.0000 = REP

TOTAL OBSERVATIONS:

WEIGHT LENGTH N OF CASES MINIMUM 13.4000 9.0000 111,6000 MAXIMUM 16.0000 12.8919 MEAN 59.3378 STANDARD DEV 23.3474 1.7286

THE FOLLOWING RESULTS ARE FOR:
TET = 1.0000

2.0000 REP

TOTAL OBSERVATIONS:

WEIGHT LENGTH N OF CASES 48 48 MINIMUM 32.3000 10.0000 136.8000 17.0000 MAXIMUM MEAN 78.4104 13.9792 STANDARD DEV 23.3395 1.3287

THE FOLLOWING RESULTS ARE FOR:

TRT 2.0000 REP 1.0000

TOTAL OBSERVATIONS:

WEIGHT LENGTH N OF CASES 28.8000 MINIMUM 11.0000 MUMIXAM 102.6000 15.0000 MEAN 69.1878 13.7317 STANDARD DEV 18.0889 1.2252

THE FOLLOWING RESULTS ARE FOR:

TDY = 2.0000

2.0000 REP

TOTAL OBSERVATIONS: 45

WEIGHT LENGTH N OF CASES 45 45 36.8000 12.0000 16.0000 MINIMUM 126.6000 MAXIMUM MEAN 80.5889 14.1556 STANDARD DEV 19.1857 0.9524

TRT 3.0000 REP 1.0000

TOTAL OBSERVATIONS: 50

WEIGHT LENGTH N OF CASES 50 50 MINIMUM 10.8000 10.0000 MAXIMUM 96.6000 15.0000 MEAN 59.4580 13.2600 STANDARD DEV 16.4066 1.0461

THE FOLLOWING RESULTS ARE FOR:

= 4.0000 = 1_0000

REP

TOTAL OBSERVATIONS: 36

WEIGHT LENGTH N OF CASES 36 36 MINIMUM 28.5000 11.0000 16.0000 171.8000 MAXIMUM 83.0333 13.8611 MEAN 1.3764 STANDARD DEV 29.0958

THE FOLLOWING RESULTS ARE FOR:

TPT = 4.0000

REP 2.0000

TOTAL OBSERVATIONS: 42

WEIGHT LENGTH N OF CASES 42 MINIMUM 49.0000 12.0000 143.6000 16.0000 MUMIXAM 93.8952 MEAN 14.4286 STANDARD DEV 22.3984 0.9913

THE FOLLOWING RESULTS ARE FOR: TRT = 5.0000

REP 1.0000

TOTAL OBSERVATIONS: 48

WEIGHT LENGTH N OF CASES 48 48 MINIMUM 22.6000 10.0000 112.6000 16.0000 MAXIMUM 66.4438 13,1667 MEAN STANDARD DEV 18.0753 1.0980

= 5.0000 TRT REP 2.0000

TOTAL OBSERVATIONS:

47

	WEIGHT	LENGTH
N OF CASES	47	47
MINIMUM	17.5000	10.0000
MAXIMUM	103.8000	15.0000
MEAN	68.9532	13.3617
STANDARD DEV	20.5018	1.3093

THE FOLLOWING RESULTS ARE FOR:

TRT 6.0000 = =

1.0000 REP

TOTAL OBSERVATIONS:

	WEIGHT	LENGTH
N OF CASES	45	45
MINIMUM	26.1000	10.0000
MAXIMUM	112.8000	15.0000
MEAN	68.4333	13.2667
STANDARD DEV	20.2256	1,2321

THE FOLLOWING RESULTS ARE FOR:

= 6.0000 TRT

2.0000 REP

TOTAL OBSERVATIONS: 49

WEIGHT LENGTH N OF CASES 49 15.0000 MINIMUM 10.0000 MUNIXAM 75.5000 15.0000 12.5714 MEAN 42.1776 STANDARD DEV 13.9345 1.2076

SUMMARY STATISTICS FOR WEIGHT

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

32.8193 DF= 10 PROBABILITY = 0.0003

ANALYSIS OF VARIANCE

SOURCE SUM OF SQUARES DF MEAN SQUARE F PROBABILITY

BETWEEN GROUPS 86892.8228 10 8689.2823 20.7334 0.0000 WITHIN GROUPS 199909.0615 477 419.0966

SUMMARY STATISTICS FOR LENGTH

BARTLETT TEST FOR HOMOGENEITY OF GROUP VARIANCES

CHI-SQUARE =

24.0000 DF= 10 PROBABILITY = 0.0076

ANALYSIS OF VARIANCE

SOURCE

SUM OF SQUARES OF MEAN SQUARE

PROBABILITY

BETWEEN GROUPS

140.9623 10

14.0962 1.5158 0.0000

WITHIN GROUPS

723.0356 477

9.2995

KOLMOGOROV-SMIRNOV ONE SAMPLE TEST USING STANDARD NORMAL DISTRIBUTION

VARIABLE

N-OF-CASES MAXDIF PROBABILITY (2-TAIL)

LENGTH WEIGHT 488.0000 488.0000

1.0000

0.0000 0.0000

Ecological Effects Branch One-Liner Data Entry Form

Pesticide Use Shaughnessy No. 128997 Chemical Tebuconazole

AQUATIC VERTEBRATE TOX.	% AI	LC ₅₀ LC ₅₀	HRB/ TYPE	NOEC	STUDY/RE VIEW DATES	MRID/ CATEGORY	LAB	RC
1.								
2.								
3.								
CHRONIC TOX.	% AI	MATC	DAYS	AFFECTED PARA.	STUDY/ REVIEW DATES	MRID/ CATEGORY	LAB	RC RC
1. Cyprinodon variegatus	96.4	>18.4 <46.9 (geo mean MATC = 29.4)	203	F, length at test termination	1993/ 1994	430096- TES I	TES	RG M
2.			·			h 4 c	2	
3.								

COMMENTS: Results based on mean measured concentrations. TES=Toxikon Environmental Sciences,