## **TEXT SEARCHABLE DOCUMENT - 2010**

# Data Evaluation Report on the Acute Toxicity of Dimethyl Disulfide to Terrestrial Vascular Plants: Seedling Emergence

PMRA Submiss	ion Number N/A		EPA MRID Number 47381001
Data Requireme	ent:	PMRA Data Code: EPA DP Barcode:	D351572
		OECD Data Point: EPA Guideline:	OPPTS 850.4100 and 850.4225 (123-1a)
Test material: Common name:	Dimethyl disulfi	de	<b>Purity:</b> 99.6%
	IUPAC: Not rep CAS name: Not CAS No.: Not re Synonyms: DMI	reported ported	
			Moncie V Wright
Primary Review Staff Scientist, C	ver: Moncie Cambridge Envir	Wright onmental, Inc.	Date: 05/16/08
Secondary Revi Senior Scientist	ewer: Teri S. , Cambridge Env	-	Signature: 6 S Mym  Date: 07/22/08  Date: (03/30/20/0
Primary Review {EPA/OECD/PM	ver: Valerie Wo MRA}	odard Woodard	Date: {03/30/20/0
Secondary Revi {EPA/OECD/PM	ewer(s): { MRA}		Date: {
Reference/Subn	nission No.: {	}	
Company Code Active Code Use Site Catego EPA PC Code	{}		
Date Evaluation	Completed: {dd	-mm-yyyy}	

<u>CITATION</u>: Porch, J.R., and H.O. Krueger. 2008. Dimethyl Disulfide: A Toxicity Test to Determine the Effects of the Test Substance on Seedling Emergence of Ten Species of Plants. Unpublished study performed by Wildlife International, Ltd., Easton, Maryland. Laboratory Project No.: 524-102. Study submitted by Arkema, Inc., Philadelphia, PA. Study completed March 17, 2008.

<u>DISCLAIMER</u>: This document provides guidance for EPA and PMRA reviewers on how to complete a data evaluation record after reviewing a scientific study concerning the acute toxicity of a pesticide to terrestrial vascular plants. It is not intended to prescribe conditions to any external party for conducting this study nor to establish absolute criteria regarding the assessment of whether the study is scientifically sound and whether the study satisfies any applicable data requirements. Reviewers are expected to review and to determine for each study, on a case-by-case basis, whether it is scientifically sound and provides sufficient information to satisfy applicable data requirements. Studies that fail to meet any of the conditions may be accepted, if appropriate; similarly, studies that meet all of the conditions may be rejected, if appropriate. In sum, the reviewer is to take into account the totality of



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factors related to the test methodology and results in determining the acceptability of the study.

#### **EXECUTIVE SUMMARY:**

The effect of Dimethyl disulfide applied as a gas on the seedling emergence of monocot (onion, *Allium cepa*; oat, *Avena sativa*; ryegrass, *Lolium perenne*; and corn, *Zea mays*) and dicot (sugarbeet, *Beta vulgaris*; cabbage, *Brassica oleracea*; cucumber, *Cucumis sativa*; soybean, *Glycine max*; lettuce, *Lactuca sativa*; and tomato, *Lycopersicon esculentum*) crops was studied at varying nominal application rates. All species were treated with nominal application rates of 0 (negative control), 37.5, 75, 150, 300, and 600 lbs a.i./A. The application rates were not analytically verified. Because DMDS is applied as a gas, the application rate also needs to be reported as a volume. Herbivory of seeds and plants and irrigation problems confounded the results for some species, particularly cucumber. Results for cucumber were not reliable and, so, are not considered in the reviewer's verification of the results.

The growth medium used in the seedling emergence test was natural soil, classified as a sandy loam, with an approximate organic matter content of 1.3%. On Day 21 the surviving plants per pot were recorded and cut at soil level for measuring the plant height and dry weight.

Dry weight and plant height was significantly affected in onion, ryegrass, sugarbeet, soybean, lettuce, and tomato. The percent inhibition in seedling emergence in the treated species as compared to the negative control ranged from - 12 to 23%.

Statistics are reported based on lb/A. The most sensitive monocot, based on dry weight, was ryegrass with NOAEC and EC<sub>25</sub> values of 75 and 290 lbs ai/A, respectively. The EC<sub>25</sub> for ryegrass should be interpreted with caution because it was not associated with a reliable 95% confidence interval. The most sensitive dicot, based on dry weight, was sugarbeet, with NOAEC and EC<sub>25</sub> values of 75 and 91 lbs ai/A, respectively.

The maximum phytotoxicity rating across all replicates and all crops on Day 21 was 100%, with means ranging from 0 to 55%, indicating that there dose-response related effects with respect to this endpoint.

Maximum Labeled Rate: 600 lbs ai/A

#### **Results Synopsis**

#### Monocot

 $EC_{05}/IC_{05}$ : <37.5 lbs ai/A 95% C.I.: Not reliable  $EC_{25}/IC_{25}$ : 290 lbs ai/A 95% C.I.: Not reliable  $EC_{50}/IC_{50}$ : >600 lbs ai/A 95% C.I.: N/A

NOAEC: 75 lbs ai/A

Slope: 0.734 Std err: 0.572

Most sensitive monocot: Ryegrass Most sensitive parameter: Dry weight

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**Dicot** 

EC<sub>05</sub>/IC<sub>05</sub>: 27 lbs ai/A 95% C.I.: 4.0-180 lbs ai/A EC<sub>25</sub>/IC<sub>25</sub>: 91 lbs ai/A 95% C.I.: 29-290 lbs ai/A EC<sub>50</sub>/IC<sub>50</sub>: 210 lbs ai/A 95% C.I.: 110-440 lbs ai/A

NOAEC: 75 lbs ai/A

Slope: 1.82 Std err: 0.573

Most sensitive dicot: Sugarbeet Most sensitive parameter: Dry Weight

This toxicity study is classified as supplemental and does not satisfy the guideline requirement for a Tier 2 terrestrial plant seedling emergence toxicity study.

Table 1. Summary of most sensitive parameters by species (lbs ai/A).

Species	Endpoint	NOAEC	EC <sub>05</sub>	EC <sub>25</sub>	EC <sub>50</sub>
Onion	Dry weight	150	<37.5	330	>600
Oat	Dry weight	600	<37.5	>600	>600
Ryegrass	Dry weight	75	<37.5	290	>600
Corn	Survival	600	<37.5	>600	>600
Sugarbeet	Dry weight	75	27	91	210
Cabbage	None	600	>600	>600	>600
Cucumber	ND	ND	ND	ND	ND
Soybean	Survival	300	310	530	>600
Lettuce	Dry weight	150	32	160	500
Tomato	Dry weight	300	180	400	>600

ND- Not Determined

### I. MATERIALS AND METHODS

#### **GUIDELINE FOLLOWED:**

This study was conducted following guidelines outlined in US EPA Series 850 – Ecological Effects Test Guidelines OPPTS Number 850.4225. The following deviations from OPPTS 850.4100 and 850.4225 were noted:

- 1. The historical % germination and storage of the seeds were not reported.
- 2. The moisture (% at 1/3 atm) of the soil was not reported.
- 3. All species were tested under similar environmental conditions instead of separating the warm-loving species from the cold-loving species.
- 4. The active ingredient was used to test the toxicity, rather than the recommended typical end-use product.
- 5. The effects of dimethyl disulfide on cucumber endpoints are unclear because of substantial herbivore damage to these seeds and plants, resulting in fewer than two replicates in at least one treatment level. The study author presents results for cucumber endpoints using both damaged and undamaged replicates and determines that toxicity estimates calculated excluding damaged replicates show this species to be one of the more sensitive dicots. The reviewer could not confidently verify these results because of the confounding factors.

These deviations do impact the acceptability of the study.

#### **COMPLIANCE:**

Signed and dated No Data Confidentiality, GLP and Quality Assurance statements were provided. This study was conducted in compliance with the requirements of 40 CFR Part 160, OECD Principles, and Japan MAFF

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with the exception of periodic analyses of well water and soil for contaminants.

A. MATERIALS:

1. Test Material

Dimethyl disulfide

**Description:** 

Liquid

Lot No./Batch No.:

05.03.06 (Batch Number)

**Purity:** 

99.6%

Stability of compound

under test conditions:

Samples were not analyzed for dimethyl disulfide due to its administration

as neat material.

(OECD recommends chemical stability in water and light)

Storage conditions of

test chemicals:

Stored at ambient room temperature.

Table 2. Physical/chemical properties of Dimethyl disulfide.

Parameter	Values	Comments
Water solubility at 20EC	Not Reported	
Vapor pressure	Not Reported	
UV absorption	Not Reported	
рКа	Not Reported	
Kow	Not Reported	

#### 2. Test organism:

Monocotyledonous species: Corn (Zea mays; Family Poaceae, Mandan Bride), Onion (Allium cepa, Family Liliaceae, WI 301), Ryegrass (Lolium perenne, Family Poaceae, Manhattan 4) and Oat (Avena sativa, Family Poaceae, Armor); EPA recommends four monocots in two families, including corn.

Dicotyledonous species: Cabbage (Brassica oleracea, Family Brassicaceae, Late Flat Dutch), Lettuce (Lactuca sativa, Family Asteraceae, Summertime), Soybean (Glycine max, Family Fabaceae, Williams 82), Sugarbeet (Beta vulgaris, Family Chenopodiaceae, none given), Tomato (Lycopersicon esculentum, Family Solanaceae, Rutgers) and Cucumber (Cucumis sativa, Family Cucurbitaceae, Straight Eight); EPA recommends six dicots in four families, including soybean and a root crop.

OECD recommends a minimum of three species selected for testing, at least one from each of the following categories: Category 1: ryegrass, rice, oat, wheat, and sorghum; Category 2: mustard, rape, radish, turnip, and Chinese cabbage; Category 3: vetch, mung bean, red clover, fenugreek, lettuce, and cress.

Seed source: Onion obtained from Wannamaker Seeds, St. Matthews, SC; oat obtained from Wilken Seed

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Grains, Inc., Pontiac, IL; ryegrass, cabbage, cucumber, and tomato obtained from The Meyer Seed Co., Baltimore, MD; corn obtained from Johnny's Selected Seeds, Winslow, ME; sugarbeet obtained from BetaSeed Inc., Shakopee, MN; soybean obtained from Missouri Foundation Seeds, Columbia, MO; and lettuce obtained from Territorial Seed Co., Cottage Grove, OR.

**Prior seed treatment/sterilization:** Seeds were not treated with fungicides, insecticides, or repellents prior to test initiation.

Historical % germination of seed: Not Reported

Seed storage, if any: Not Reported

#### **B. STUDY DESIGN:**

#### 1. Experimental Conditions

a. Limit test: N/A; this was a Tier II test.

b. Range-finding study: A range-finding study was not conducted.

c. Definitive Study

Table 3: Experimental Parameters - Seedling Emergence.

Parameters	Seedling Emergence				
	Details	Remarks			
		Criteria			
Duration of the test	21 Days				
		Recommended test duration is 14-21 days.			
	N. A.	OECD recommends that the test be terminated no sooner than 14 days after 50 percent of the control seedlings have emerged			
Number of seeds/plants/species/	All species were comprised of 10 seeds per replicate. Each replicate				
seeds/plants/species/ replicate	had a single pot, with 10 seeds per pot.	Ten seeds per replicate should be used.			
		OECD recommends a minimum of five seeds planted in each replicate within 24 hours of incorporation of the test substance. All seeds of each species for each test should be of the same size class. The seed should not be imbibed.			
Number of replicates Control:	4				

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Parameters	See	dling Emergence			
	Details	Remarks			
		Criteria			
Formulation control: Treated:	N/A 4	Four replicates per dose should be used.  OECD recommends a minimum of four replicates per treatment			
Test concentrations (lb ai/A or g ai/ha) Nominal: Measured:	0 (negative control), 37.5, 75.0, 150, 300, and 600 lbs a.i./A  Measured application rates could not be determined because analytical verification was not performed.	Five test concentrations should be used with a dose range of 2X or 3X progression  OECD recommends three concentrations, preferably with application rates equivalent to 0.0 (control), 1.0, 10.0 and 100 mg substance per kg of oven-dried soil.			
Method and interval of analytical verification	Not performed due to use of neat test substance.				
LOQ: LOD:	N/A N/A				
Adjuvant (type, percentage, if used)	N/A				
Test container (pot)		All pots contained drainage holes along the bottom.			
Size/Volume Material: (glass/polystyrene)	All species: 16 cm (diameter) x 12 cm (tall) Plastic	Non-porous containers should be used.  OECD recommends that non-porous plastic or glazed pot be used.			
Growth facility	Greenhouse; after initial 48-hour exposure period, treatment groups were taken out of the greenhouse, removed from treatment bags, and placed in subirrigation trays on greenhouse benches.				
Method/depth of seeding	Ryegrass, onion, tomato, cabbage, sugarbeet and lettuce planted at <i>ca</i> . 6 mm depth; corn, cucumber, soybean, and oat were planted at 20 mm.				

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<b>Parameters</b>	Seedling Emergence				
	Details	Remarks			
		Criteria			
Test material application Application time including the plant growth stage Number of application Application interval Method of application	Test material was applied on Day 0 to injection containers.  Each species was treated once N/A; single application  Test material was injected into dosing containers that were placed in bags along with the replicate pots containing the planted seeds. The bags were sealed for 48 hours so that the test substance could volatilize.				
Details of soil used Geographic location  Depth of soil collection Soil texture % sand % silt % clay pH: % organic carbon CEC Moisture at 1/3 atm (%)	Representative of a sandy loam soil Not Reported  75% 11% 14% Not reported. 1.3% organic matter Not reported. Not Reported	Soil mixes containing sandy loam, loam, or clay loam soil with no greater than 2% organic matter are preferable. Glass beads, rock wool, and 100% acid washed sand are not preferred.  OECD prefers the soil to be sieved (0.5 cm) to remove coarse fragments. Carbon content should not exceed 1.5% (3% organic matter). Fine particles (under 20um) makeup should be between 10 and 20%. The recommended pH is between 5.0 and 7.5.			
Details of nutrient medium, if used	N/A; a nutrient medium was not used				
Watering regime and schedules Water source/type:	Well water from the greenhouse facility. Not reported.	Following application, pots were top watered to initiate a continuous water column. Once this column was created, capillary action was used to water the soil.			
Volume applied: Interval of application: Method of application:	Intermittent; every 2 to 6 days. Sub-irrigation trays filled to a predetermined depth.	EPA prefers that bottom watering be utilized for seedling emergence studies so that the chemical is not leached out of the soil during the test.			
Any pest control	None				

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Parameters	See	Seedling Emergence				
	Details	Remarks				
		Criteria				
method/fertilization, if used						
Test conditions Temperature: Photoperiod: Light intensity and quality: Relative humidity:	8.49-29.64°C 16L:8D 1.3-16.9 moles photosynthetically active radiation. Natural light supplemented by artificial light. 13.83-74.70%	EPA prefers that the cold vs warm loving plants be tested in two separate groups to optimize plant growth.  OECD prefers that the temperature, humidity and light conditions be suitable for maintaining normal growth of each species for the test period.				
Reference chemical (if used) Name: Concentrations:	N/A N/A	A reference chemical was not used.				
Other parameters, if any	None					

#### 2. Observations:

Table 4: Observation Parameters - Seedling Emergence.

Parameters		Seedling Emergence
	Details	Remarks
Parameters measured (e.g., number of germinated seeds, emerged seedlings, plant height, dry weight or other endpoints)	-Emergence -Survival -Dry weight -Plant height	
Measurement technique for each parameter	Emergence and survival were assessed visually. Dry weight was determined by clipping seedlings at soil level, drying the shoots of all living seedlings in an oven, and then weighing. The total dry weights of the shoots within each replicate were measured and then divided by the	

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	number weighed. Height was measured to the nearest whole centimeter from the soil surface to the tip of the tallest leaf or apical meristem.	
Observation intervals	Emergence was assessed 7, 14 and 21 days after application. Height, survival, plant height, and dry weight were determined at test termination.	
Other observations, if any	None	
Were raw data included?	Replicate data were provided	
Phytotoxicity rating system, if used	0%- No injury/effect; 10-30%- slight plant effect (effects barely noticeable to not obviously detrimental; 40-60%-moderate effect involving the whole plant (recovery possible to doubtful); 70-90%- severe effect with recovery not possible (loss of some leaves to only a few surviving leaves); 100%- complete plant effect (death of plant)	

#### **II. RESULTS and DISCUSSION:**

#### A. INHIBITORY EFFECTS:

### 1. Seedling Emergence:

Negative control emergence ranged from 60% to 100% for all species tested and treatment group emergence ranged from 58% to 100%.

Percent survival in the negative control ranged from 97% to 100% while treatment group survival ranged from 80% to 100%.

The study author's results indicated that the level of inhibition in plant height across all plant treatments was low in half the treatments, with the exclusion of sugarbeet, cucumber, soybean, lettuce, and tomato. Sugarbeet exhibited the largest inhibition of plant height of up to 53%, and there was a clear dose-response relationship.

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The level of inhibition in dry weight across all treatments was moderate to high (up to 73%) with the exception of cabbage, corn, and oat suggesting that there were adverse effects on dry weight.

There was severe damage in the cucumber treatments due to mice which resulted in elimination of numbers of replicates across every concentration. The study authors reported that based on dry weight, the most sensitive dicot was cucumber, with reported NOAEC and EC25 values of 37.5 and 91.2 lbs a.i./A, respectively. Based on dry weight, the most sensitive monocot was ryegrass, with reported NOAEC and EC25 values of 75.0 and 286 lbs a.i./A.

The study authors used a standard phytotoxicity rating system. There were limited, scattered cases of necrosis, leaf curl, unshed seed coat, wilting and chlorosis across all species and concentrations.

#### **B. REPORTED STATISTICS:**

The study author analyzed phytotoxicity, survival, emergence, plant height and dry weight based on lb/A. Mean emergence, survival, dry weight, and height of the control and treatment groups were compared using a Dunnett's t-test, using the DUNNETT option of the GLM procedure of SAS. Data were evaluated using Dunnett's test to determine the NOEAC and LOAEC. The determination of effect concentrations and their confidence limits was performed using the non-linear regression analysis of Bruce and Versteeg when reductions in test endpoints among one or more treatment groups were 25% or more relative to control means. Analyses were conducted using the NLIN procedure of SAS version 8. Nominal application rates (lbs ai/A) were used for all analyses.

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Table 5: Reported effect of Dimethyl disulfide on Seedling Emergence

Species	Results summary for height (lbs ai/A)									
	cm*	NOAEC	EC <sub>05</sub>	95%CI	EC <sub>25</sub>	95%CI	EC <sub>50</sub>	95%CI	Slope	Std. Err.
Onion	7.6-10.0	150	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Oat	37.3-39.7	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Ryegrass	11.5-13.5	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Corn	42.7-47.3	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Sugarbeet	5.8-12.6	75	N.R.	N.R.	157	N.R.	448	N.R.	N.R.	N.R.
Cabbage	9.7-11.4	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Cucumber <sup>1</sup>	7.9-10.6	150	N.R.	N.R.	471	N.R.	>600	N/A	N.R.	N.R.
Soybean	12.6-18.5	300	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Lettuce	6.1-8.7	75	N.R.	N.R.	432	N.R.	>600	N/A	N.R.	N.R.
Tomato	6.2-8.1	150	N.R.	N.R.	473	N.R.	>600	N/A	N.R.	N.R.

<sup>\*</sup> range provided represents the range of the treatment means

N/A- Not Applicable

N.R.- Not Reported

<sup>1</sup> Only undisturbed replicates were evaluated.

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Table 5a: Reported effect of Dimethyl disulfide on Seedling Emergence

	Results summar	y for dry weigh	t (lbs ai/A)							
	g*	NOAEC	EC <sub>05</sub>	95%CI	EC <sub>25</sub>	95%CI	EC <sub>50</sub>	95%CI	Slope	Std. Err.
Onion	0.00554-0.00887	150	N.R.	N/A	334	N.R.	>600	N/A	N.R.	N.R.
Oat	0.129-0.159	600	N.R.	N/A	>600	N/A	>600	N/A	N.R.	N.R.
Ryegrass	8.98-14.83	75	N.R.	N/A	286	N.R.	>600	N/A	N.R.	N.R.
Corn	0.291-0.330	600	N.R.	N/A	>600	N/A	>600	N/A	N.R.	N.R.
Sugarbeet	0.0207-0.0910	75	N.R.	N/A	91.8	N.R.	215	N.R.	N.R.	N.R.
Cabbage	87.5-112.7	600	N.R.	N/A	>600	N/A	>600	N/A	N.R.	N.R.
Cucumber <sup>1</sup>	0.144-0.234	37.5	N.R.	N/A	91.2	N.R.	>600	N/A	N.R.	N.R.
Soybean	0.216-0.292	600	N.R.	N/A	>600	N/A	>600	N/A	N.R.	N.R.
Lettuce	20.4-44.4	75	N.R.	N/A	161	N.R.	498	N.R.	N.R.	N.R.
Tomato	26.4-52.4	150	N.R.	N/A	401	N.R.	>600	N/A	N.R.	N.R.

<sup>\*</sup> range provided represents the range of the treatment means

N/A- Not Applicable

N.R.- Not Reported

<sup>1</sup> Only undisturbed replicates were evaluated.

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Table 5b: Reported effect of Dimethyl disulfide on Seedling Emergence

Species	Results summ	ary for survival (	lbs ai/A)							
	%*	NOAEC	EC <sub>05</sub>	95%CI	EC <sub>25</sub>	95%CI	EC <sub>50</sub>	95%CI	Slope	Std. Err.
Onion	91-100	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Oat	98-100	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Ryegrass	100	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Corn	91-100	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Sugarbeet	88-100	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Cabbage	97-100	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Cucumber	N/A	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Soybean	80-98	300	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Lettuce	97-100	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.
Tomato	97-100	600	N.R.	N.R.	>600	N/A	>600	N/A	N.R.	N.R.

<sup>\*</sup> range provided represents the range of the treatment means

N/A- Not Applicable

N.R.- Not Reported

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Day 7 Er	mergence						_			
Control	Onion	Oat	Ryegrass	Corn	Sugarbeet	Cabbage	Cucumber	Soybean	Lettuce	Tomato
80 (10-100)	44 (25-70)	96 (90-100)	71 (60-88)	83 (68-95)	47 (30-70)	72 (55-87)	49 (0-100)	67 (40-85)	78 (63-95)	21 (0-38)

\* provide the mean and range

Day 14 E	mergence	e								
Control	Onion	Oat	Ryegrass	Corn	Sugarbeet	Cabbage	Cucumber	Soybean	Lettuce	Tomato
89 (60-100)	75 (68-80)	96 (90-100)	93 (88-98)	88 (78-95)	65 (58-73)	78 (73-87)	93 (75-100)	77 (40-93)	88 (78-98)	83 (73-90)

Day 21 F	Emergence	e								•
Control	Onion	Oat	Ryegrass	Corn	Sugarbeet	Cabbage	Cucumber	Soybean	Lettuce	Tomato
89 (60-100)	77 (70-83)	97 (90-100)	94 (88-98)	88 (78-95)	67 (58-75)	78 (73-87)	93 (75-100)	85 (70-93)	88 (78-98)	85 (73-93)

Plant Inj	jury Index	4			_		_		_	
Control	Onion	Oat	Ryegrass	Corn	Sugarbeet	Cabbage	Cucumber	Soybean	Lettuce	Tomato
0-100	0-100	0-100	0-20	0-100	0-100	0-100	0-100	0-100	0-100	0-100

0%- No injury/effect; 10-30%- slight plant effect (effects barely noticeable to not obviously detrimental; 40-60%- moderate effect involving the whole plant (recovery possible to doubtful); 70-90%- severe effect with recovery not possible (loss of some leaves to only a few surviving leaves); 100%- complete plant effect (death of plant)

### C. VERIFICATION OF STATISTICAL RESULTS BY THE REVIEWER:

Statistical Method(s): Any species exhibiting an inhibition of ≥5% in dry weight, plant height, or % survival relative to the negative control was analyzed; toxicity values for all other species and endpoints were determined visually. All analyses were conducted using the negative control only. The reviewer then tested each appropriate data set for normality using the Chi-square and Shapiro-Wilks tests and for homogeneity of variance using the Hartley and Bartlett's tests. All data sets with the exclusion of corn survival met these assumptions of ANOVA; therefore, the NOAEC values were determined using the parametric Dunnett's and Williams' tests via Toxstat statistical software. In the case of corn survival, the Steel's and Krusal-Wallis tests were used to determine the NOAEC. The reviewer then attempted to determine the ECx values, confidence intervals and probit slopes using the probit analysis via Nuthatch statistical software. All analyses were conducted using the nominal application rates in lbs a.i./A.

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Table 6: Reported effect of Dimethyl disulfide on Seedling Emergence

Species	Results summar	y for dry weigl	nt (lbs ai/A)							
	g*	NOAEC	EC <sub>05</sub>	95%CI	EC <sub>25</sub>	95%CI	EC <sub>50</sub>	95%CI	Slope	Std. Err.
Onion	0.00554-0.00887	150	<37.5	NR	330	59-1900	>600	N/A	0.719	0.434
Oat	0.129-0.159	600	<37.5	NR	>600	N/A	>600	N/A	0.315	0.427
Ryegrass	8.98-14.83	75	<37.5	NR	290	NR	>600	N/A	0.734	0.572
Corn	0.291-0.330	600	240	NR	>600	N/A	>600	N/A	0.785	1.02
Sugarbeet	0.0207-0.0910	75	27	4.0-180	91	29-290	210	110-440	1.82	0.573
Cabbage	87.5-112.7	600	>600	N/A	>600	N/A	>600	N/A	N/A	N/A
Cucumber <sup>1</sup>	0.144-0.234	ND	ND	N/A	ND	N/A	ND	N/A	N/A	N/A
Soybean	0.216-0.292	300	130	NR	>600	N/A	>600	N/A	1.28	0.937
Lettuce	20.4-44.4	150	32	2.8-360	160	49-540	500	260-990	1.38	0.546
Tomato	26.4-52.4	300	180	42-790	400	220-720	>600	N/A	2.83	1.70

<sup>\*</sup> range provided represents the range of the treatment means

NR-Not reliable; range fell outside the tested concentrations.

N/A- Not Applicable

N.D.- Not Determined

In the oat treatments, Replicate D from the Negative Control was dropped due to a problem with the irrigation tube, and Replicate B from the 37.5 lb/A treatment was dropped due to damage during handling

<sup>&</sup>lt;sup>1</sup>Cucumber could not be analyzed due to lack of replicate data resulting from widespread mice damage.

PMRA Submission Number N/A

EPA MRID Number 47381001

Table 6a: Reported effect of Dimethyl disulfide on Seedling Emergence

Species	Results summa	ary for height (lbs	s ai/A)							
	cm*	NOAEC	EC <sub>05</sub>	95%CI	EC <sub>25</sub>	95%CI	EC50	95%CI	Slope	Std. Err.
Onion	7.6-10.0	150	<37.5	NR	>600	N/A	>600	N/A	0.387	0.471
Oat <sup>1</sup>	37.3-39.7	600	320	NR	>600	N/A	>600	N/A	0.513	0.529
Ryegrass	11.5-13.5	600	>600	N/A	>600	N/A	>600	N/A	N/A	N/A
Corn	42.7-47.3	600	310	72-1300	>600	N/A	>600	N/A	1.09	0.890
Sugarbeet	5.8-12.6	75	35	9.6-130	160	82-300	450	310-640	1.48	0.322
Cabbage <sup>2</sup>	9.7-11.4	600	N.D.	N/A	N.D.	N/A	N.D.	N/A	N/A	N/A
Cucumber <sup>3</sup>	7.9-10.6	N.D.	N.D.	N.D.	N.D.	N/A	N.D.	N/A	N/A	N/A
Soybean	12.6-18.5	300	240	110-550	610	470-780	>600	N/A	2.44	1.03
Lettuce	6.1-8.7	75	28	1.9-410	430	170-1100	>600	N/A	0.815	0.323
Tomato	6.2-8.1	150	<37.5	NR	480	130-1800	>600	N/A	0.705	0.349

<sup>\*</sup> range provided represents the range of the treatment means

NR-Not reliable; range fell outside the tested concentrations.

N/A- Not Applicable

N.D.- Not Determined

<sup>1</sup> In the oat treatments, Replicate D from the Negative Control was dropped due to a problem with the irrigation tube.

<sup>2</sup> Cabbage didn't fit probit analysis; NOAEC and EC25 values had to be visually determined.

<sup>3</sup> Cucumber could not be analyzed due to lack of replicate data resulting from widespread mice damage.

PMRA Submission Number N/A

### EPA MRID Number 47381001

Table 6b: Reported effect of Dimethyl disulfide on Seedling Emergence

Species	Results summ	ary for survival (	lbs ai/A)							
	%*	NOAEC	EC <sub>05</sub>	95%CI	EC <sub>25</sub>	95%CI	EC <sub>50</sub>	95%CI	Slope	Std. Err.
Onion	91-100	600	51	. NR	>600	N/A	>600	N/A	0.501	0.712
Oat	98-100	600	>600	N/A	>600	N/A	>600	N/A	N/A	N/A
Ryegrass	100	600	>600	N/A	>600	N/A	>600	N/A	N/A	N/A
Corn	91-100	600	<37.5	NR	>600	N/A	>600	N/A	0.571	0.394
Sugarbeet	88-100	75	130	NR	580	180-1900	>600	N/A	1.51	1.73
Cabbage	97-100	600	>600	N/A	>600	N/A	>600	N/A	N/A	N/A
Cucumber <sup>1</sup>	95-100	-600	>600	N/A	>600	N/A	>600	N/A	N/A	N/A
Soybean	80-98	300	310	110-870	530	390-700	760	500-1100	4.30	3.18
Lettuce	97-100	600	>600	N/A	>600	N/A	>600	N/A	N/A	N/A
Tomato	97-100	600	>600	N/A	>600	N/A	>600	N/A	N/A	N/A

<sup>\*</sup> range provided represents the range of the treatment means

NR-Not reliable; range fell outside the tested concentrations.

N/A- Not Applicable

N.D.- Not Determined

<sup>1</sup> Cucumber replicates at each treatment level were damaged by mice. Study authors provided data for all replicates, and performed statistics on all replicates and on just undamaged replicates. The reviewer was unable to perform statistics, so NOAEC and EC25 values were visually determined.

### PMRA Submission Number N/A

EPA MRID Number 47381001

Day 7 E	mergence	_	_							
Control	Onion	Oat	Ryegrass	Com	Sugarbeet	Cabbage	Cucumber	Soybean	Lettuce	Tomato
80 (10-100)	44 (25-70)	96 (90-100)	71 (60-88)	83 (68-95)	47 (30-70)	72 (55-87)	49 (0-100)	67 (40-85)	78 (63-95)	21 (0-38)

Day 14 E	mergence	e								
Control	Onion	Oat	Ryegrass	Corn	Sugarbeet	Cabbage	Cucumber	Soybean	Lettuce	Tomato
89 (60-100)	75 (68-80)	96 .(90-100)	93 (88-98)	88 (78-95)	65 (58-73)	78 (73-87)	93 (75-100)	77 (40-93)	88 (78-98)	83 (73 <b>-</b> 90)

Day 21 E	Emergence	e								
Control	Onion	Oat	Ryegrass	Corn	Sugarbeet	Cabbage	Cucumber	Soybean	Lettuce	Tomato
89 (60-100)	77 (70-83)	97 (90-100)	94 (88-98)	88 (78-95)	67 (58-75)	78 (73-87)	93 (75-100)	85 (70-93)	88 (78-98)	85 (73-93)

Plant Inj	jury Inde	<u> </u>								
Control	Onion	Oat	Ryegrass	Corn	Sugarbeet	Cabbage	Cucumber	Soybean	Lettuce	Tomato
0-100	0-100	0-100	0-20	0-100	0-100	0-100	0-100	0-100	0-100	0-100

0%- No injury/effect; 10-30%- slight plant effect (effects barely noticeable to not obviously detrimental; 40-60%- moderate effect involving the whole plant (recovery possible to doubtful); 70-90%- severe effect with recovery not possible (loss of some leaves to only a few surviving leaves); 100%- complete plant effect (death of plant)

#### Monocot

 $EC_{05}/IC_{05}$ : <37.5 lbs ai/A 95% C.I.: Not reliable  $EC_{25}/IC_{25}$ : 290 lbs ai/A 95% C.I.: Not reliable  $EC_{50}/IC_{50}$ : >600 lbs ai/A 95% C.I.: N/A

NOAEC: 75 lbs ai/A

Slope: 0.734 Std err: 0.572

Most sensitive monocot: Ryegrass Most sensitive parameter: Dry weight

#### Dicot

EC<sub>05</sub>/IC<sub>05</sub>: 27 lbs ai/A 95% C.I.: 4.0-180 lbs ai/A EC<sub>25</sub>/IC<sub>25</sub>: 91 lbs ai/A 95% C.I.: 29-290 lbs ai/A EC<sub>50</sub>/IC<sub>50</sub>: 210 lbs ai/A 95% C.I.: 110-440 lbs ai/A

NOAEC: 75 lbs ai/A

Slope: 1.82 Std err: 0.573

Most sensitive dicot: Sugarbeet Most sensitive parameter: Dry Weight

#### D. STUDY DEFICIENCIES:

#### PMRA Submission Number N/A

EPA MRID Number 47381001

Herbivory (presumably by mice), resulting in seed and plant damage interfered with the ability to determine effects of treatment on cucumber, which may have been one of the most sensitive dicot species. While the study authors analyzed results for data including and excluding damaged plants, the reviewer did not believe that the results obtained from these analyses were reliable and suggests retesting this species, taking care to prevent the introduction of potentially confounding factors.

#### E. REVIEWER-S COMMENTS:

The results are based on the nominal test concentrations, because the application rates were not analytically verified.

The study authors did not report the confidence intervals or slopes for the results of statistical analyses. Therefore, the reviewer's results are reported in the Executive Summary and Conclusions sections of this DER using nominal test concentrations.

In the case of cabbage height, inhibitions were >5%, but the reviewer was unable to determine the  $EC_{05}$  value due to the lack of three or more distinct isotone means. Therefore, the reviewer reported "N.D." in the respective tables. In situations where inhibitions of  $\geq$ 5% were determined, but the NOAEC value was the highest nominal application rate, the reviewer indicated that the given endpoint exhibited no sensitivity.

Cucumber replicates suffered severe damage due to mice. The study authors reported data and statistics for all replicates, and for just the undamaged replicates. The reviewer used data for the undamaged replicates only, but was unable to generate statistics due to the lack of replicates. According to the statistics generated by the study authors, cucumber was the most sensitive dicot, but this could not be verified by the reviewer. Further, it is possible that the damage from the mice affected all of the replicates, not just the ones that appeared to be damaged.

The physiochemical properties of the test material were not reported.

In the oat treatments, Replicate D from the Negative Control was dropped due to a problem with the irrigation tube, and Replicate B from the 37.5 lb/A treatment was dropped due to damage during handling procedure.

The NOAEC for ryegrass dry weight and sugarbeet survival were determined visually by determination of the highest application rate with a % inhibition of <10% relative to the negative control.

#### F. CONCLUSIONS:

The study is calssified as supplemental. The most sensitive monocot, based on dry weight, was ryegrass with NOAEC and EC<sub>25</sub> values of 75 and 290 lbs ai/A, respectively. The most sensitive dicot, based on dry weight, was sugarbeet, with NOAEC and EC<sub>25</sub> values of 75 and 91lbs ai/A, respectively.

Most sensitive monocot and EC<sub>25</sub>: Ryegrass (Dry weight), 290 lbs ai/A Most sensitive dicot and EC<sub>25</sub>: Sugarbeet (Dry weight), 91 lbs ai/A

#### III. REFERENCES:

U.S. Environmental Protection Agency. 1996. Series 850- Ecological Effects Test Guidelines (draft), OPPTS Number 850.4225: Terrestrial Plant Toxicity, Tier II (Seedling Emergence).

Frans, Robert E. and Ronald E. Talbert. 1977. Design of Field Experiments and the Measurement and Analysis of Plant Responses. Pages 15-23 in B. Truelove, ed. Research Methods in Weed Science. Southern Weed Science

### PMRA Submission Number N/A

EPA MRID Number 47381001

Society, Auburn University, Alabama.

SAS Institute, Inc. 1999. SAS Proprietary Software Version 8, Cary, NC, SAS Institute, Inc.

Bruce, Robert D. and Donald J. Versteeg. 1992. A Statistical Procedure for Modeling Continuous Toxicity Data. *Environmental Toxicology and Chemistry*. 11: 1485-1494.

#### PMRA Submission Number N/A

EPA MRID Number 47381001

#### APPENDIX I. OUTPUT OF REVIEWER'S STATISTICAL VERIFICATION:

Dimethyl disulfide & Cabbage Height (cm) lbs ai/A File: 1001cah Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

INTERVAL <-1.5 -1.5 to <-0.5 -0.5 to 0.5 >0.5 to 1.5 >1.5

EXPECTED 1.541 5.566 8.786 5.566 1.541 OBSERVED 0

Calculated Chi-Square goodness of fit test statistic = 5.9331 Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Cabbage Height (cm) lbs ai/A Transform: NO TRANSFORMATION

Shapiro Wilks test for normality

D = 37.982

W = 0.973

Critical W (P = 0.05) (n = 23) = 0.914Critical W (P = 0.01) (n = 23) = 0.881

Data PASS normality test at P=0.01 level. Continue analysis.

Dimethyl disulfide & Cabbage Height (cm) lbs ai/A File: 1001cah Transform: NO TRANSFORMATION

Hartley test for homogeneity of variance

Calculated H statistic (max Var/min Var) = 10.71 Closest, conservative, Table H statistic = 184.0 (alpha = 0.01)

Used for Table H ==> R (# groups) = 6, df (# reps-1) = 3 Actual values ==> R (# groups) = 6, df (# avg reps-1) = 2.83 (average df used)

\_\_\_\_\_\_

Data PASS homogeneity test. Continue analysis.

NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used).

#### PMRA Submission Number N/A

EPA MRID Number 47381001

Dimethyl disulfide & Cabbage Height (cm) lbs ai/A File: 1001cah Transform: NO TRANSFORMATION

Bartletts test for homogeneity of variance

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Calculated B statistic = 5.43

Table Chi-square value = 15.09 (alpha = 0.01)

Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 2.83 Used for Chi-square table value ==> df (#groups-1) = 5

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Cabbage Height (cm) lbs ai/A File: 1001cah Transform: NO TRANSFORMATION

#### ANOVA TABLE

SOURCE	DF	SS	MS	F
Between	5	9.115	1.823	0.816
Within (Error)	17	37.982	2.234	
Total	22	47.097		

\_\_\_\_\_\_

Critical F value = 2.81 (0.05, 5, 17)

Since F < Critical F FAIL TO REJECT Ho: All groups equal

Dimethyl disulfide & Cabbage Height (cm) lbs ai/A File: 1001cah Transform: NO TRANSFORMATION

E	BONFERRONI T-TEST -	TABLE 1 OF 2	Ho: Contro	l <treatm< th=""><th>ent</th></treatm<>	ent
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg control	10.600	10.600		
2	37.5	10.433	10.433	0.146	
3	75	11.425	11.425	-0.781	
4	150	10.875	10.875	-0.260	
5	300	9.700	9.700	0.852	
6	600	9.700	9.700	0.852	

Bonferroni T table value = 2.57 (1 Tailed Value, P=0.05, df=17,5)

Dimethyl disulfide & Cabbage Height (cm) lbs ai/A File: 1001cah Transform: NO TRANSFORMATION

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#### PMRA Submission Number N/A

### EPA MRID Number 47381001

	BONFERRONI T-TEST -	TABLE	2 OF 2	Ho:Contr	ol <treatment< th=""></treatment<>
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg control	4			
2	37.5	3	2.930	27.6	0.167
3	75	4	2.713	25.6	-0.825
4	150	4	2.713	25.6	-0.275
5	300	4	2.713	25.6	0.900
6	600	4	2.713	25.6	0.900

Dimethyl disulfide & Cabbage Height (cm) lbs ai/A File: 1001cah Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg control	4	10.600	10.600	10.860
2	37.5	3	10.433	10.433	10.860
3	75	4	11.425	11.425	10.860
4	150	4	10.875	10.875	10.860
5	300	4	9.700	9.700	9.700
6	600	4	9.700	9.700	9.700

Dimethyl disulfide & Cabbage Height (cm) lbs ai/A File: 1001cah Transform: NO TRANSFORMATION

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg control	10.860	0.228		1.74	k= 1, v=17
75 150 300	10.860 10.860 9.700	0.246 0.246 0.852		1.82 1.85 1.87	k= 2, v=17 k= 3, v=17 k= 4, v=17
600	9.700	0.852		1.87	k = 5, v = 17

\_\_\_\_\_

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

1001CAH : Dimethyl disulfide & Cabbage Height (cm) lbs ai/A

Williams Test

[One-Sided Test for Decrease, alpha = 0.050000]

Dose	Isotone Means	T-bar P-	value	Significance
	means			
0	10.9			
37.5	10.9 -	0.2277	N.S.	

PMRA Submiss	sion Number	r_N/A			EPA_MRII	Number 47381001
75 150 300 600	10.9 10.9 9.7 9.7	-0.246 0.8515	N.S. N.S. N.S.			
"*"=Signifi	cant; "N.S	S."=Not Sign	ificant	•		
!!!Failure #	3: Data r	not suitable	for pr	obit model fit	t.	
Criterion is	3 or more	e distinct i	sotone	means.		
Dimethyl dis		Corn Height ansform: NO				
Chi-square to				d expected fre	equencies	
INTERVAL	<-1.5	-1.5 to <	0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
EXPECTED OBSERVED	1.608	5.808 11		9.168 5	5.808 8	1.608 0
	hi-Square uare value	goodness of e (alpha = 0	fit te.	st statistic = 13.277	≃ 10.5795	
Dimethyl dist File: 1001ch						
Shapiro Wilks	s test for	normality				
D = 143.660						
W = 0.962						
Critical W ()	P = 0.01)	(n = 24) = 0	0.884			
				. Continue ana		
Dimethyl dist File: 1001ch		Corn Height cansform: NO				
Hartley test	for homog	geneity of va	ariance			
Calculated H				= 17.87 = 184.0 (al	pha = 0.01)	
Used for Tabl Actual values	le H ==> S ==>	R (# groi R (# groi	ups) =	6, df (# 6, df (#	reps-1) = avg reps-1) =	3 3.00

#### PMRA Submission Number N/A

EPA MRID Number 47381001

Data PASS homogeneity test. Continue analysis.

NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used).

Dimethyl disulfide & Corn Height (cm) lbs ai/A File: 1001ch Transform: NO TRANSFORMATION

Bartletts test for homogeneity of variance

Calculated B statistic = 9.29

Table Chi-square value = 15.09 (alpha = 0.01) Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5

The state of the s

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Corn Height (cm) lbs ai/A File: 1001ch Transform: NO TRANSFORMATION

#### ANOVA TABLE

 SOURCE
 DF
 SS
 MS
 F

 Between
 5
 61.453
 12.291
 1.540

 Within (Error)
 18
 143.660
 7.981

 Total
 23
 205.113

Critical F value = 2.77 (0.05,5,18) Since F < Critical F FAIL TO REJECT Ho:All groups equal

Dimethyl disulfide & Corn Height (cm) lbs ai/A File: 1001ch Transform: NO TRANSFORMATION

	DUNNETTS TEST - T	ABLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	sig
1	Neg Control	45.975	45.975		
2	37.5	47.275	47.275	-0.651	
3	75	46.950	46.950	-0,488	
4	150	45.375	45.375	0.300	
5	300	44.000	44.000	0.989	
6	600	42.725	42.725	1.627	

#### PMRA Submission Number N/A

EPA MRID Number 47381001

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Corn Height (cm) lbs ai/A File: 1001ch Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)		DIFFERENCE FROM CONTROL
1	Neg Control	4			
2	37.5	4	4.814	10.5	-1.300
3	75	4	4.814	10.5	-0.975
4	150	4	4.814	10.5	0.600
5	300	4	4.814	10.5	1.975
6	600	4	4.814	10.5	3.250

Dimethyl disulfide & Corn Height (cm) lbs ai/A File: 1001ch Transform: NO TRANSFORMATION

	WILLIAMS TEST (Isotor	nic	regression model	) TABLE 1 O	F 2
GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	45.975	45.975	46.733
2	37.5	4	47.275	47.275	46.733
3	75	4	46.950	46.950	46.733
4	150	4	45.375	45.375	45.375
5	300	4	44.000	44.000	44.000
6	600	4	42.725	42.725	42.725

Dimethyl disulfide & Corn Height (cm) lbs ai/A File: 1001ch Transform: NO TRANSFORMATION

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg Control 37.5 75 150 300 600	46.733 46.733 46.733 45.375 44.000 42.725	0.380 0.380 0.300 0.989 1.627		1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 2.825

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

Estimates of EC%

PMRA Submission Number N/A

EPA MRID Number 47381001

Parameter	Estimate	95% Bot	ınds	Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EC5	3.1E+02	72.	1.3E+03	0.31	0.23
EC10	6.7E+02	2.8E+02	1.6E+03	0.18	0.42
EC25	2.4E+03	2.1E+02	2.7E+04	0.51	0.089
EC50	9.9E+03	86.	1.1E+06	0.99	0.0087

Slope = 1.09 Std.Err. = 0.890

Goodness of fit: p = 0.82 based on DF= 3.0 18.

1001CH : Dimethyl disulfide & Corn Height (cm) lbs ai/A

Observed vs. Predicted Treatment Group Means

 Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change
0.00	4.00	46.0	46.8	-0.792	100.	0.00
37.5	4.00	47.3	46.6	0.697	99.6	0.403
75.0	4.00	46.9	46.3	0.659	99.0	1.02
150.	4.00	45.4	45.7	-0.303	97.7	2.33
300.	4.00	44.0	44.5	-0.504	95.2	4.84
600.	4.00	42.7	42.5	0.241	90.8	9.16

!!!Warning: EC10 not bracketed by doses evaluated.

!!!Warning: EC25 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Lettuce Height (cm) lbs ai/A File: 1001lh Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

INTERVAL <-1.5 -1.5 to <-0.5 -0.5 to 0.5 >0.5 to 1.5 >1.5

EXPECTED	1.608	5.808	9.168	5.808	1.608
OBSERVED	0	8	9	7	0

Calculated Chi-Square goodness of fit test statistic = 4.2910

Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Lettuce Height (cm) lbs ai/A File: 1001lh Transform: NO TRANSFORMATION

Shapiro Wilks test for normality

D = 7.955

W = 0.976

Critical W (P = 0.05) (n = 24) = 0.916

PMRA Submission N			EP	A MRID Number
Critical W (P =				
Data PASS normal	ity test at P=0	.01 level. Conti	nue analysis.	
Dimethyl disulfic	de & Lettuce He Transform:	ight (cm) lbs ai NO TRANSFORMATIO	/A N	
Hartley test for	homogeneity of	variance		
Calculated H star Closest, conserva				01)
Used for Table H Actual values	==> R (# g ==> R (# g	roups) = 6, roups) = 6,	df (# reps-1) = df (# avg reps-	= 3 -1) = 3.00
Data PASS homoge		tinue analysis		
NOTE: This test : but do not	requires equal differ greatly	_	st may still be	
Dimethyl disulfic File: 1001lh Bartletts test fo	Transform: N or homogeneity	O TRANSFORMATION of variance		
Calculated B star Table Chi-square Table Chi-square	value = 15.0	9 (alpha = 0.01)	) )	
Average df used : Used for Chi-squa				
Data PASS homoger	neity test at 0	.01 level. Conti	nue analysis.	
		plicate sizes th tatistic (see ab		cate size is
Dimethyl disulfic	de & Lettuce He Transform:	ight (cm) lbs ai NO TRANSFORMATIC	/A N	
		ANOVA TABLE		
	DF	SS	MS	F
Between	5		4.805	10.871
Within (Error)	18	7.955	0.442	
Potal	23	31.978		

31.978 

#### PMRA Submission Number N/A

EPA MRID Number 47381001

Critical F value = 2.77 (0.05,5,18)
Since F > Critical F REJECT Ho:All groups equal

Dimethyl disulfide & Lettuce Height (cm) lbs ai/A File: 1001lh Transform: NO TRANSFORMATION

	DUNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	sig
1	Neg control	8.300	8.300		
2	37.5	8.675	8.675	-0.798	
3	75	8.275	8.275	0.053	
4	150	6.650	6.650	3.510	*
5	300	6.100	6.100	4.680	*
6	600	6.650	6.650	3.510	*

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Lettuce Height (cm) lbs ai/A File: 1001lh Transform: NO TRANSFORMATION

	DUNNETTS TEST - '	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg control	4			
2	37.5	4	1.133	13.7	-0.375
3	75	4	1.133	13.7	0.025
4	150	4	1.133	13.7	1.650
5	300	4	1.133	13.7	2.200
6	600	4	1.133	13.7	1.650

Dimethyl disulfide & Lettuce Height (cm) lbs ai/A File: 1001lh Transform: NO TRANSFORMATION

WILLIAMS	TEST	(Isotonic	regression	model)	TABLE	1 (	OF 2	
GROUP			ORIGINAI	J	TRANSFORM	ED	ISOTONI	ZED

GROUP	TDDDWTTTTALTTAL		ORIGINAL	TRANSFORMED	ISOTONIZED
	IDENTIFICATION	N 	MEAN	MEAN	MEAN
1	Neg control	4	8.300	8.300	8.488
2	37.5	4	8,675	8.675	8.488
3	75	4	8,275	8,275	8.275
4	150	4	6.650	6.650	6.650
5	300	4	6.100	6.100	6.375
6	600	4	6.650	6.650	6.375

Dimethyl disulfide & Lettuce Height (cm) lbs ai/A File: 1001lh Transform: NO TRANSFORMATION

#### PMRA Submission Number N/A

#### EPA MRID Number 47381001

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg control 37.5 75 150 300 600	8.488 8.488 8.275 6.650 6.375 6.375	0.399 0.053 3.510 4.095 4.095	* * *	1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 0.665

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

Estimates of EC%

					- <b></b>
Estimate	95% Bot	unds	Std.Err.	Lower Bound	
	Lower	Upper		/Estimate	
28.	1.9	4.1E+02	0.56	0.068	
78.	12.	5.2E+02	0.40	0.15	
4.3E+02	1.7E+02	1.1E+03	0.20	0.39	
2.9E+03	5.9E+02	1.4E+04	0.33	0.20	
	28. 78. 4.3E+02	Lower 28. 1.9 78. 12. 4.3E+02 1.7E+02	Lower Upper 28. 1.9 4.1E+02 78. 12. 5.2E+02 4.3E+02 1.7E+02 1.1E+03	Lower Upper 28. 1.9 4.1E+02 0.56 78. 12. 5.2E+02 0.40 4.3E+02 1.7E+02 1.1E+03 0.20	Lower Upper /Estimate 28. 1.9 4.1E+02 0.56 0.068 78. 12. 5.2E+02 0.40 0.15 4.3E+02 1.7E+02 1.1E+03 0.20 0.39

Slope = 0.815 Std.Err. = 0.323

!!!Poor fit: p = 0.0046 based on DF= 3.0 18.

1001LH : Dimethyl disulfide & Lettuce Height (cm) lbs ai/A

\_\_\_\_\_\_

Observed vs. Predicted Treatment Group Means

Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change
0.00 37.5 75.0 150. 300.	4.00 4.00 4.00 4.00 4.00 4.00	8.30 8.68 8.27 6.65 6.10	8.60 8.07 7.76 7.33 6.79 6.12	-0.298 0.609 0.518 -0.683 -0.686 0.530	100. 93.8 90.2 85.3 78.9 71.2	0.00 6.18 9.78 14.7 21.1 28.8

!!!Warning: EC5 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Oat Height (cm) lbs ai/A File: 1001oh Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
EXPECTED OBSERVED	1.541 0	5.566 9	8.786 7	5.566 7	1.541 0

Calculated Chi-Square goodness of fit test statistic = 5.9331

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PMRA Submission Number N/A
                                                     EPA MRID Number 47381001
Table Chi-Square value (alpha = 0.01) = 13.277
Data PASS normality test. Continue analysis.
Dimethyl disulfide & Oat Height (cm) lbs ai/A
File: 1001oh Transform: NO TRANSFORMATION
Shapiro Wilks test for normality
D = 85.555
W = 0.947
Critical W (P = 0.05) (n = 23) = 0.914
Critical W (P = 0.01) (n = 23) = 0.881
Data PASS normality test at P=0.01 level. Continue analysis.
Dimethyl disulfide & Oat Height (cm) lbs ai/A
File: 1001oh Transform: NO TRANSFORMATION
Hartley test for homogeneity of variance
______
Calculated H statistic (max Var/min Var) = 14.13
Closest, conservative, Table H statistic = 184.0 (alpha = 0.01)
Used for Table H ==> R (\# groups) = 6, df (\# reps-1) = Actual values ==> R (\# groups) = 6, df (\# avg reps-1) =
                                                              2.83
                                          (average df used)
______
Data PASS homogeneity test. Continue analysis.
NOTE: This test requires equal replicate sizes. If they are unequal
     but do not differ greatly, the Hartley test may still be used
     as an approximate test (average df are used).
Dimethyl disulfide & Oat Height (cm) lbs ai/A
File: 1001oh Transform: NO TRANSFORMATION
Bartletts test for homogeneity of variance
                          Calculated B statistic =
                        7.04
Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05)
Average df used in calculation ==> df (avg n - 1) = 2.83
Used for Chi-square table value ==> df (#groups-1) = 5
```

Data PASS homogeneity test at 0.01 level. Continue analysis.

#### PMRA Submission Number N/A

EPA MRID Number 47381001

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Oat Height (cm) lbs ai/A File: 1001oh Transform: NO TRANSFORMATION

#### ANOVA TABLE

\_\_\_\_\_

SOURCE	DF	SS	MS	F
Between	5	22.411	4.482	0.891
Within (Error)	17	85.555	5.033	
Total	22	107.966		

Critical F value = 2.81 (0.05,5,17) Since F < Critical F FAIL TO REJECT Ho:All groups equal

Dimethyl disulfide & Oat Height (cm) lbs ai/A File: 1001oh Transform: NO TRANSFORMATION

	BONFERRONI T-TEST -	TABLE 1 OF 2	Ho:Contro	l <treatm< th=""><th>ent</th></treatm<>	ent
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg control	39.800	39.800		
2 3	37.5 75	39.700 39.650	39.700 39.650	0.058 0.088	
4 5	150 300	38.150 37.325	38.150 37.325	0.963 1.444	
6	600	37.925	37.925	1.094	

Bonferroni T table value = 2.57 (1 Tailed Value, P=0.05, df=17,5)

Dimethyl disulfide & Oat Height (cm) lbs ai/A File: 1001oh Transform: NO TRANSFORMATION

E	CONFERRONI T-TEST -	TABLE	2 OF 2	Ho:Contr	ol <treatment< th=""></treatment<>
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg control	3			
2	37.5	4	4.398	11.1	0.100
3	75	4	4.398	11.1	0.150
4	150	4	4.398	11.1	1.650
5	300	4	4.398	11.1	2.475
6	600	4	4.398	11.1	1.875

Dimethyl disulfide & Oat Height (cm) lbs ai/A File: 1001oh Transform: NO TRANSFORMATION

#### PMRA Submission Number N/A

EPA	MRID	Number	47381001
E I A	MINI	rumber	<b>4/301001</b>

	WILLIAMS TEST (Isoto	nic	regression model	) TABLE 1 OF	7 2
GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg control	3	39.800	39.800	39.800
2	37.5	4	39.700	39.700	39.700
3	75	4	39.650	39.650	39.650
4	150	4	38.150	38.150	38.150
5	300	4	37.325	37.325	37.625
6	600	4	37.925	37.925	37.625

Dimethyl disulfide & Oat Height (cm) lbs ai/A File: 1001oh Transform: NO TRANSFORMATION

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg control 37.5 75 150 300 600	39.800 39.700 39.650 38.150 37.625 37.625	0.058 0.088 0.963 1.269 1.269		1.74 1.82 1.85 1.87 1.87	k= 1, v=17 k= 2, v=17 k= 3, v=17 k= 4, v=17 k= 5, v=17

s = 2.243

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

#### Estimates of EC%

Parameter	Estimate	95% Bot	ınds	Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EC5	3.2E+02	14.	7.4E+03	0.65	0.044
EC10	1.6E+03	76.	3.6E+04	0.64	0.046
EC25	2.5E+04	8.6	7.4E+07	1.7	0.00034
EC50	5.2E+05	0.32	8.5E+11	3.0	6.1E-07

Slope = 0.513 Std.Err. = 0.529

Goodness of fit: p = 0.73 based on DF= 3.0 17.

1001OH : Dimethyl disulfide & Oat Height (cm) lbs ai/A

Observed vs. Predicted Treatment Group Means

Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	
0.00 37.5 75.0 150. 300.	3.00 4.00 4.00 4.00 4.00	39.8 39.7 39.7 38.2 37.3 37.9	40.0 39.4 39.1 38.6 38.1	-0.240 0.335 0.590 -0.497 -0.776 0.529	100. 98.3 97.6 96.5 95.2 93.4	0.00 1.69 2.45 3.48 4.84 6.60	

<sup>!!!</sup>Warning: EC10 not bracketed by doses evaluated.

#### PMRA Submission Number N/A

EPA MRID Number 47381001

!!!Warning: EC25 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Onion Height (cm) lbs ai/A File: 1001onh Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

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\_\_\_\_\_\_

Calculated Chi-Square goodness of fit test statistic = 12.1805 Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Onion Height (cm) lbs ai/A File: 1001onh Transform: NO TRANSFORMATION

Shapiro Wilks test for normality

brapito with cost for notification

D = 12.365

W = 0.974

Critical W (P = 0.05) (n = 24) = 0.916 Critical W (P = 0.01) (n = 24) = 0.884

Data PASS normality test at P=0.01 level. Continue analysis.

Dimethyl disulfide & Onion Height (cm) lbs ai/A File: 1001onh Transform: NO TRANSFORMATION

Hartley test for homogeneity of variance

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Calculated H statistic (max Var/min Var) = 40.06 Closest, conservative, Table H statistic = 184.0 (alpha = 0.01)

Used for Table H ==> R (# groups) = 6, df (# reps-1) = 3 Actual values ==> R (# groups) = 6, df (# avg reps-1) = 3.00

Data PASS homogeneity test. Continue analysis.

NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used).

#### PMRA Submission Number N/A

EPA MRID Number 47381001

Dimethyl disulfide & Onion Height (cm) lbs ai/A File: 1001onh Transform: NO TRANSFORMATION

Bartletts test for homogeneity of variance

Calculated B statistic = 8.11

Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Onion Height (cm) lbs ai/A File: 1001onh Transform: NO TRANSFORMATION

#### ANOVA TABLE

SOURCE	DF	SS	MS	F
Between	5	15.953	3.191	4.645
Within (Error)	18	12.365	0.687	
Total	23	28.318		

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Critical F value = 2.77 (0.05,5,18)
Since F > Critical F REJECT Ho:All groups equal

Dimethyl disulfide & Onion Height (cm) lbs ai/A File: 1001onh Transform: NO TRANSFORMATION

	DUNNETTS TEST - TABLE 1 OF 2		Ho:Control <treatment< th=""></treatment<>		
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg control	9.275	9.275		
2	37.5	8.025	8.025	2.133	
3	75	10.025	10.025	-1.280	
4	150	8.275	8.275	1.706	
5	300	7.575	7.575	2.901	*
6	600	8.475	8.475	1.365	

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Onion Height (cm) lbs ai/A

#### PMRA Submission Number N/A

EPA MRID Number 47381001

File: 1001onh Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)		DIFFERENCE FROM CONTROL
1	Neg control	4			
2	37.5	4	1.412	15.2	1.250
3	75	4	1.412	15.2	-0.750
4	150	4	1.412	15.2	1.000
5	300	4	1.412	15.2	1.700
6	600	4	1.412	15.2	0.800

Dimethyl disulfide & Onion Height (cm) lbs ai/A File: 1001onh Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg control	4	9.275	9.275	9.275
2 3	37.5 75	4 4	8.025 10.025	8.025 10.025	9.025 9.025
4	150	4	8.275	8.275	8.275
5 6	300 600	4 4	7.575 8.475	7.575 8.475	8.025 8.025

Dimethyl disulfide & Onion Height (cm) lbs ai/A File: 1001onh Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

IDENTIFICATION	ISOTONIZED	CALC.	SIG	TABLE	DEGREES OF
	MEAN	WILLIAMS	P=.05	WILLIAMS	FREEDOM
Neg control 37.5 75 150 300 600	9.275 9.025 9.025 8.275 8.025 8.025	0.427 0.427 1.706 2.133 2.133	*	1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 0.829

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

Estimates of EC%

Parameter	Estimate	95% Bounds		Std.Err. Lower Bound	
		Lower	Upper		/Estimate
EC5	33.	0.0097	1.1E+05	1.7	0.00030
EC10	2.8E+02	3.2	2.5E+04	0.94	0.011
EC25	1.1E+04	1.6	7.0E+07	1.8	0.00015
EC50	5.8E+05	0.0062	5.5E+13	3.8	1.1E-08

10010NH : Dim	#Reps.  4.00 4.00 4.00 4.00 4.00	ulfide & Onio Treatment Gr Obs. P Mean M 9.28 8.03 10.0 8.28	n Heig  oup Me  red. ean 9.27 8.78	ObsPred. 0.00844 -0.756	Pred. %Control	%Change	
Dose  0.00 37.5 75.0 150. 300. 600.	ethyl dis  Predicted 	Ulfide & Onio Treatment Gr Obs. P Mean M 9.28 8.03 10.0 8.28	n Heig  oup Me  red. ean 9.27 8.78	ObsPred. 0.00844 -0.756	Pred. %Control	%Change	
Dose  0.00 37.5 75.0 150. 300. 600.	#Reps.  4.00 4.00 4.00 4.00 4.00 4.00	Treatment Gr Obs. P Mean M 9.28 8.03 10.0 8.28	oup Me  red. ean 9.27 8.78	Obs. -Pred. 0.00844 -0.756	Pred. %Control	*Change	
0.00 37.5 75.0 150. 300. 600.	#Reps.  4.00 4.00 4.00 4.00 4.00 4.00	Obs. P Mean M 9.28 8.03 10.0 8.28	red. ean 9.27 8.78	Obs. -Pred. 0.00844 -0.756	Pred. %Control	%Change	
37.5 75.0 150. 300. 600.	4.00 4.00 4.00 4.00	8.03 10.0 8.28	8.78	-0.756	100.	0.00	
75.0 150. 300. 600.	4.00 4.00 4.00	10.0 8.28	8.65		94.8	5.24	
	4.00 4.00		8.50	1.37 -0.228	93.4 91.8	6.61 8.24	
!!!Warning: E		7.58 8.47	8.33 8.12	~0.750 0.356	89.8 87.6	$10.2 \\ 12.4$	
	C5 not br	acketed by do	ses ev	aluated.		•	
IIIWarning. E	C25 not h	racketed by d	nses e	valuated			
Chi-square te	st for no	nsform: NO TR rmality: actu	al and	expected			
	1 600	F 000	_	0.160			1 600
EXPECTED OBSERVED	0.608	5.808 9		9.168	5	.808	0
BSERVED	0	5.808 9 goodness of f		9	6 		0

D = 54.142

W = 0.961

Critical W (P = 0.05) (n = 24) = 0.916 Critical W (P = 0.01) (n = 24) = 0.884

Data PASS normality test at P=0.01 level. Continue analysis.

### PMRA Submission Number N/A

EPA MRID Number 47381001

Dimethyl disulfide & Ryegrass Height (cm) lbs ai/A File: 1001rh Transform: NO TRANSFORMATION Hartley test for homogeneity of variance Calculated H statistic (max Var/min Var) = 3.75 Closest, conservative, Table H statistic = 184.0 (alpha = 0.01) Used for Table H ==> R (# groups) = 6, df (# reps-1) = 3 Actual values ==> R (# groups) = 6, df (# avg reps-1) = 3.00 Data PASS homogeneity test. Continue analysis. NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used). Dimethyl disulfide & Ryegrass Height (cm) lbs ai/A File: 1001rh Transform: NO TRANSFORMATION Bartletts test for homogeneity of variance Calculated B statistic = 1.49
Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05) Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5 Data PASS homogeneity test at 0.01 level. Continue analysis. NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above). Dimethyl disulfide & Ryegrass Height (cm) lbs ai/A File: 1001rh Transform: NO TRANSFORMATION ANOVA TABLE DF SS MS \_\_\_\_\_\_ 5 17.777 Between 3.555 Within (Error) 18 54.142 3.008 

Critical F value = 2.77 (0.05,5,18) Since F < Critical F FAIL TO REJECT Ho:All groups equal

71.920

23

Total

### PMRA Submission Number N/A

### EPA MRID Number 47381001

Dimethyl disulfide & Ryegrass Height (cm) lbs ai/A File: 1001rh Transform: NO TRANSFORMATION

	DUNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg Control	13.600	13.600		
2	37.5	13.450	13.450	0.122	
3	75	13.425	13.425	0.143	
4	150	11.750	11.750	1.509	
5	300	11.525	11.525	1.692	
6	600	12.175	12.175	1.162	

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Ryegrass Height (cm) lbs ai/A File: 1001rh Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)		DIFFERENCE FROM CONTROL
1	Neg Control	4			
2	37.5	4	2.956	21.7	0.150
3	75	4	2.956	21.7	0.175
4	150	4	2.956	21.7	1.850
5	300	4	2.956	21.7	2.075
6	600	4	2.956	21.7	1.425

Dimethyl disulfide & Ryegrass Height (cm) lbs ai/A File: 1001rh Transform: NO TRANSFORMATION

	WILLIAMS TEST (Isotor	nic	regression model	) TABLE 1 O	F 2
GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	13.600	13.600	13.600
2	37.5	4	13.450	13.450	13.450
3	75	4	13.425	13.425	13.425
4	150	4	11.750	11.750	11.817
5	300	4	11.525	11.525	11.817
6	600	4	12.175	12.175	11.817

Dimethyl disulfide & Ryegrass Height (cm) lbs ai/A File: 1001rh Transform: NO TRANSFORMATION

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 OF	F 2
IDENTIFICATION	ISOTONIZED MEAN		SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM

PMRA Submis	sion Numbe	r <u>N/A</u>			EPA MR	ID Number 47381001
Neg	Control	13.600				
	37.5	13.450	0.122		1.73 k=	= 1, v=18
	75	13.425	0.143		1.82 k=	= 2, V=18
	150	11.817	1.454			= 3, v=18
	300					= 4, v=18
	600	11.817	1.454		1.87 k=	= 5, v=18
					1.07 K-	
s = 1.734 Note: df use !!!Failure#	d for tab	le values ar		e when v :	> 20.	
Dimethyl dis File: 1001sb			ight (cm) lb TRANSFORMAT			
Chi-square t		•	_		quencies	
INTERVAL	<-1.5	-1.5 to <-	0.5 -0.5	to 0.5	>0.5 to 1.5	>1.5
EXPECTED	1 600	5.808		9.168	5.808	1.608
OBSERVED	0			6	8	0
OBBERVED	U	10		O	O	O
Data PASS no Dimethyl dis File: 1001sb	ulfide & S	Sugarbeet He ransform: NO				
D = 35.242						
W = 0.952						
Critical W (						
Data PASS no	rmality te	est at P=0.0	1 level. Con	tinue anal	lysis.	
Dimethyl dist			ight (cm) lb O TRANSFORMA			
Hartley test						
Calculated H Closest, con					pha = 0.01)	
Used for Tab Actual value	le H ==> s ==>	R (# gro R (# gro	ups) = 6, ups) = 6,	df (# r df (# a	reps-1) = avg reps-1) =	3 3.00

### PMRA Submission Number N/A

**EPA MRID Number 47381001** 

Data PASS homogeneity test. Continue analysis.

NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used).

Dimethyl disulfide & Sugarbeet Height (cm) lbs ai/A File: 1001sbh Transform: NO TRANSFORMATION

Bartletts test for homogeneity of variance

Calculated B statistic = 5.36

Table Chi-square value = 15.09 (alpha = 0.01)

Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5

\_\_\_\_\_\_

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Sugarbeet Height (cm) lbs ai/A File: 1001sbh Transform: NO TRANSFORMATION

### ANOVA TABLE

SOURCE	DF	SS	MS	F	
Between	5	164.687	32.937	16.822	-
Within (Error)	18	35.242	1.958		
Total	23	199.930			-

Critical F value = 2.77 (0.05, 5, 18)Since F > Critical F REJECT Ho: All groups equal

Dimethyl disulfide & Sugarbeet Height (cm) lbs ai/A File: 1001sbh Transform: NO TRANSFORMATION

	DUNNETTS TEST - TA	ABLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg Control	12.300	12.300		
2	37.5	12.600	12.600	-0.303	
3	75	11.775	11.775	0.531	
4	150	9.325	9.325	3.007	*
5	300	7.200	7.200	5.154	*

### PMRA Submission Number N/A

EPA MRID Number 47381001

6	600	5.775	5.775	6.595 *
Dunnett table value =	2.41	(1 Tailed Value,	P=0.05, df=18	,5)

Dimethyl disulfide & Sugarbeet Height (cm) lbs ai/A File: 1001sbh Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg Control	4			
2	37.5	4	2.385	19.4	-0.300
3	75	4	2.385	19.4	0.525
4	150	4	2.385	19.4	2.975
5	300	4	2.385	19.4	5.100
6	600	4	2.385	19.4	6.525

Dimethyl disulfide & Sugarbeet Height (cm) lbs ai/A File: 1001sbh Transform: NO TRANSFORMATION

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

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	12.300	12.300	12.450
2	37.5	4	12.600	12.600	12.450
3	75	4	11.775	11.775	11.775
4	150	4	9.325	9.325	9.325
5	300	4	7.200	7.200	7.200
6	600	4	5.775	5.775	5.775

Dimethyl disulfide & Sugarbeet Height (cm) lbs ai/A File: 1001sbh Transform: NO TRANSFORMATION

WILLIAMS	TEST	(Isotonic	regression	model)	TABLE	2	OF	2
***************************************		(				_		

IDENTIFICAT	ION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg C	ontrol 37.5 75 150 300 600	12.450 12.450 11.775 9.325 7.200 5.775	0.152 0.531 3.007 5.155 6.595	* *	1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 1.399

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

Estimates of EC%

<b>PMRA</b>	Subr	nission	Number	N/A

EPA MRID Number 47381001

Parameter	Estimate	95% Bo	unds	Std.Err.	Lower Bound	
		Lower	Upper		/Estimate	
EC5	35.	9.6	1.3E+02	0.27	0.28	
EC10	61.	22.	1.7E+02	0.22	0.35	
EC25	1.6E+02	82.	3.0E+02	0.14	0.52	
EC50	4.5E+02	3.1E+02	6.4E+02	0.075	0.70	

Slope = 1.48 Std.Err. = 0.322

Goodness of fit: p = 0.50 based on DF= 3.0 18.

1001SBH : Dimethyl disulfide & Sugarbeet Height (cm) lbs ai/A

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Observed vs. Predicted Treatment Group Means

 Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	
0.00	4.00	12.3	12.8	-0.499	100.	0.00	
37.5	4.00	12.6	12.1	0.510	94.5	5.54	
75.0	4.00	11.8	11.2	0.577	87.5	12.5	
150.	4.00	9.32	9.72	-0.396	75.9	24.1	
300.	4.00	7.20	7.71	-0.510	60.2	39.8	
600.	4.00	5.77	5.46	0.318	42.6	57.4	

!!!Warning: EC5 not bracketed by doses evaluated.

Dimethyl disulfide & Soybean Height (cm) lbs ai/A File: 1001sh Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
		·····			
EXPECTED OBSERVED	1.608 0	5.808 10	9.168 7	5.808 7	1.608 0

Calculated Chi Square goodness of fit test statistic - 6 9999

Calculated Chi-Square goodness of fit test statistic = 6.9989
Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Soybean Height (cm) lbs ai/A File: 1001sh Transform: NO TRANSFORMATION

Shapiro Wilks test for normality

D = 16.775

W = 0.976

Critical W (P = 0.05) (n = 24) = 0.916Critical W (P = 0.01) (n = 24) = 0.884

Data PASS normality test at P=0.01 level. Continue analysis.

### PMRA Submission Number N/A

EPA MRID Number 47381001

Dimethyl disulfide & Soybean Height (cm) lbs ai/A File: 1001sh Transform: NO TRANSFORMATION Hartley test for homogeneity of variance Calculated H statistic (max Var/min Var) = 6.46 Closest, conservative, Table H statistic = 184.0 (alpha = 0.01) Used for Table H ==> R (# groups) = 6,df (# reps-1) = R (# groups) = 6,df (# avg reps-1) = 3.00Actual values ==> Data PASS homogeneity test. Continue analysis. NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used). Dimethyl disulfide & Soybean Height (cm) lbs ai/A File: 1001sh Transform: NO TRANSFORMATION Bartletts test for homogeneity of variance Calculated B statistic = 3.63

Table Chi-square value = 15.09 (alpha = 0.01)

Table Chi-square value = 11.07 (alpha = 0.05) Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5 Data PASS homogeneity test at 0.01 level. Continue analysis. NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above). Dimethyl disulfide & Soybean Height (cm) lbs ai/A File: 1001sh Transform: NO TRANSFORMATION ANOVA TABLE \_\_\_\_\_\_ SS DF Between 5 75.703 15.141 Within (Error) 18 16.775 0.932 23 Total 92.478 

Critical F value = 2.77 (0.05,5,18)
Since F > Critical F REJECT Ho:All groups equal

### PMRA Submission Number N/A

EPA MRID Number 47381001

Dimethyl disulfide & Soybean Height (cm) lbs ai/A File: 1001sh Transform: NO TRANSFORMATION

D	UNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1 2 3 4 5	Neg control 37.5 75 150 300 600	16.850 15.500 18.525 15.550 15.725 12.600	16.850 15.500 18.525 15.550 15.725 12.600	1.978 -2.454 1.904 1.648 6.226	*

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Soybean Height (cm) lbs ai/A File: 1001sh Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg control	4			
2	37.5	4	1.645	9.8	1.350
3	75	4	1.645	9.8	-1.675
4	150	4	1.645	9.8	1.300
5	300	4	1.645	9.8	1.125
6	600	4	1.645	9.8	4.250

Dimethyl disulfide & Soybean Height (cm) lbs ai/A File: 1001sh Transform: NO TRANSFORMATION

	WILLIAMS TEST (Isoton	nic	regression model	l) TABLE 1 O	F 2
GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1 2 3 4 5	Neg control 37.5 75 150 300 600	4 4 4 4 4 4	16.850 15.500 18.525 15.550 15.725 12.600	16.850 15.500 18.525 15.550 15.725 12.600	16.958 16.958 16.958 15.638 15.638 12.600

Dimethyl disulfide & Soybean Height (cm) lbs ai/A File: 1001sh Transform: NO TRANSFORMATION

WILLIAMS		regression			_
	ISOTONIZED		SIG	TABLE	DEGREES OF

PMRA Submission Number N/A	j	EPA MRID Number 47381001

IDENTIFICATION	MEAN	WILLIAMS	P=.05	WILLIAMS	FREEDOM
Neg control	16.958				
37.5	16.958	0.159		1.73	k= 1, v=18
75	16.958	0.159		1.82	k = 2, v = 18
150	15.638	1.776		1.85	k = 3, v = 18
300	15.638	1.776		1.86	k = 4, $v = 18$
600	12.600	6.226	*	1.87	k = 5, v = 18

s = 0.965

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

### Estimates of EC%

						<b>-</b>
Parameter	Estimate	95% Bot	unds	Std.Err.	Lower Bound	
		Lower	Upper		/Estimate	
EC5	2.4E+02	1.1E+02	5.5E+02	0.17	0.45	
EC10	3.4E+02	2.0E+02	5.8E+02	0.11	0.59	
EC25	6.1E+02	4.7E+02	7.8E+02	0.053	0.78	
EC50	1.2E+03	6.0E+02	2.2E+03	0.13	0.52	

Slope = 2.44 Std.Err. = 1.03

!!!Poor fit: p = 0.0018 based on DF= 3.0 18.

1001SH : Dimethyl disulfide & Soybean Height (cm) lbs ai/A

Observed vs. Predicted Treatment Group Means

Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	
0.00 37.5 75.0 150.	4.00 4.00 4.00 4.00 4.00	16.9 15.5 18.5 15.5 15.7	16.7 16.7 16.7 16.5 15.5	0.113 -1.23 1.82 -0.931 0.274	100. 100. 99.8 98.5 92.3	0.00 0.0140 0.188 1.53 7.68	
600.	4.00	12.6	12.6	-0.0399	75.5	24.5	

!!!Warning: EC25 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Tomato Height (cm) lbs ai/A File: 1001th Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

eni square test for normarity, actuar and expected frequencies

INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
	<del></del>				
EXPECTED OBSERVED	1.608 0	5.808 7	9.168 9	5.808 8	1.608 0

Calculated Chi-Square goodness of fit test statistic = 4.2910 Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

### PMRA Submission Number N/A

EPA MRID Number 47381001

```
Dimethyl disulfide & Tomato Height (cm) lbs ai/A
File: 1001th Transform: NO TRANSFORMATION
Shapiro Wilks test for normality
______
D = 19.802
W = 0.972
Critical W (P = 0.05) (n = 24) = 0.916
Critical W (P = 0.01) (n = 24) = 0.884
Data PASS normality test at P=0.01 level. Continue analysis.
Dimethyl disulfide & Tomato Height (cm) lbs ai/A
                 Transform: NO TRANSFORMATION
File: 1001th
Hartley test for homogeneity of variance
Calculated H statistic (max Var/min Var) = 15.88
Closest, conservative, Table H statistic = 184.0 (alpha = 0.01)
Used for Table H ==> R (\# groups) = 6, df (\# reps-1) = 3
Actual values ==> R (\# groups) = 6, df (\# avg reps-1) = 3.00
Data PASS homogeneity test. Continue analysis.
NOTE: This test requires equal replicate sizes. If they are unequal
     but do not differ greatly, the Hartley test may still be used
     as an approximate test (average df are used).
Dimethyl disulfide & Tomato Height (cm) lbs ai/A
File: 1001th Transform: NO TRANSFORMATION
Bartletts test for homogeneity of variance
Calculated B statistic = 6.45
Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05)
Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (\#groups-1) = 5
Data PASS homogeneity test at 0.01 level. Continue analysis.
NOTE: If groups have unequal replicate sizes the average replicate size is
     used to calculate the B statistic (see above).
Dimethyl disulfide & Tomato Height (cm) lbs ai/A
```

### PMRA Submission Number N/A

EPA MRID Number 47381001

File: 1001th Transform: NO TRANSFORMATION

ANOVA TABLE

SOURCE	DF	SS	MS	F
Between	5	14.824	2.965	2.695
Within (Error)	18	19.803	1.100	
Total	23	34.626		

Critical F value = 2.77 (0.05, 5, 18)

Since F < Critical F FAIL TO REJECT Ho: All groups equal

Dimethyl disulfide & Tomato Height (cm) lbs ai/A File: 1001th Transform: NO TRANSFORMATION

D.	UNNETTS TEST - TAI	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1 2 3 4 5	Neg Control 37.5 75 150 300 600	8.375 8.150 7.125 7.350 6.500 6.225	8.375 8.150 7.125 7.350 6.500 6.225	0.303 1.685 1.382 2.528 2.899	*

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Tomato Height (cm) lbs ai/A File: 1001th Transform: NO TRANSFORMATION

	DUNNETTS TEST -	FABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	, IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg Control	4			
2	37.5	4	1.787	21.3	0.225
3	75	4	1.787	21.3	1.250
4	150	4	1.787	21.3	1.025
5	300	4	1.787	21.3	1.875
6	600	4	1.787	21.3	2.150

Dimethyl disulfide & Tomato Height (cm) lbs ai/A File: 1001th Transform: NO TRANSFORMATION

	WILLIAMS	TEST	(Isotonic	regression	model)	TABLE	1	OF 2		
					- <b></b>					
GROUP				ORIGINAI		TRANSFORM	ΞD	I	SOTONIZED	
	TDENTTE	FTCATT	DN N	MEAN		MEAN			MEAN	

PMRA S	ubmission Number N/A			EPA	A MRID Number 47381001  8.375	
	Neq Control	4	8.375	8.375	0 275	
2	37.5	4	8.150	8.150	8.150	
3	75	4	7.125	7.125	7.238	
4	150	4	7.350	7.350	7.238	
5	300	4	6.500	6.500	6.500	
6	600	4	6.225	6.225	6.225	

Dimethyl disulfide & Tomato Height (cm) lbs ai/A File: 1001th Transform: NO TRANSFORMATION

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg Control 37.5 75 150 300 600	8.375 8.150 7.238 7.238 6.500 6.225	0.303 1.534 1.534 2.528 2.899	* *	1.73 1.82 1.85 1.86 1.87	k= 1, V=18 k= 2, V=18 k= 3, V=18 k= 4, V=18 k= 5, V=18

s = 1.049

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

### Estimates of EC%

Parameter	Estimate	95% Bot	unds	Std.Err.	Lower Bound	
		Lower	Upper		/Estimate	
EC5	20.	0.51	7.9E+02	0.77	0.025	
EC10	66.	5.1	8.6E+02	0.54	0.077	
EC25	4.8E+02	1.3E+02	1.8E+03	0.27	0.27	
EC50	4.3E+03	3.7E+02	5.0E+04	0.51	0.086	

Slope = 0.705 Std.Err. = 0.349

Goodness of fit: p = 0.73 based on DF= 3.0 18.

1001TH : Dimethyl disulfide & Tomato Height (cm) lbs ai/A

Observed vs. Predicted Treatment Group Means

Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change
0.00 37.5 75.0 150. 300.	4.00 4.00 4.00 4.00 4.00 4.00	8.38 8.15 7.13 7.35 6.50 6.22	8.42 7.81 7.52 7.15 6.68 6.13	-0.0495 0.339 -0.399 0.200 -0.184 0.0935	100. 92.7 89.3 84.9 79.3 72.8	0.00 7.28 10.7 15.1 20.7 27.2

<sup>!!!</sup>Warning: EC5 not bracketed by doses evaluated.

<sup>!!!</sup>Warning: EC50 not bracketed by doses evaluated.

PMRA Subm		er N/A Corn Dry Weight (g	r) lbs ai/A	EPA MRII	O Number 4'
File: 1001		ransform: NO TRANSF			
-		normality: actual a	-	<b>-</b>	
INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
EXPECTED OBSERVED			9.168	5.808	1.608
		e goodness of fit t ue (alpha = 0.01) =		6.3102	
Data PASS r	normality (	test. Continue anal	ysis.		
		Corn Dry Weight (g			
		or normality			
					· <b></b> -
D = 0.01					
Critical W	(P = 0.05) (P = 0.01)	(n = 24) = 0.916 (n = 24) = 0.884			
					· <b>-</b>
Jata PASS I	iormaticy (	test at P=0.01 leve	I. Continue ana	Tysis.	
Dimethyl di File: 1001c	isulfide &	Corn Dry Weight (g Transform: NO TRANS	) lbs ai/A FORMATION		
Hartley tes	st for homo	ogeneity of varianc	e		
		ic (max Var/min Var e, Table H statisti		pha = 0.01)	
Used for Ta Actual valu	able H ==> ies ==>	R (# groups) = R (# groups) =	6, df (# 6, df (#	reps-1) = avg reps-1) =	3 3.00
Data PASS h	nomogeneity	/ test. Continue an	alysis.		
but d	do not diff	ires equal replicat Fer greatly, the Ha ate test (average d	rtley test may		
Dimethyl di File: 1001c		Corn Dry Weight (g			

### PMRA Submission Number N/A

### EPA MRID Number 47381001

Bartletts test for homogeneity of variance

Calculated B statistic = 8.55

Table Chi-square value = 15.09 (alpha = 0.01)

Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5

\_\_\_\_\_

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Corn Dry Weight (g) lbs ai/A File: 1001cw Transform: NO TRANSFORMATION

#### ANOVA TABLE

SOURCE	DF	SS	MS	F
Between	5	0.0038	0.0008	0.800
Within (Error)	18	0.0184	0.0010	
Total	23	0.0222		

Critical F value = 2.77 (0.05, 5, 18)Since F < Critical F FAIL TO REJECT Ho:All groups equal

Dimethyl disulfide & Corn Dry Weight (g) lbs ai/A File: 1001cw Transform: NO TRANSFORMATION

	DUNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg Control	0.312	0.312		~ ~ ~
2	37.5	0.330	0.330	-0.816	
3	75	0.308	0.308	0.179	
4	150	0.299	0.299	0.570	
5	300	0.298	0.298	0.626	
6	600	0.291	0.291	0.917	

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Corn Dry Weight (g) lbs ai/A File: 1001cw Transform: NO TRANSFORMATION

	DUNNETTS TE	ST -	TABLE 2	OF 2		Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
an arm			NUM OF			ig Diff		DIFFERENCE
GROUP	IDENTIFI	CATION	REPS	(I	N ORIG.	UNITS)	CONTROL	FROM CONTROL

PMRA Sul	bmission Number N/A			EPA :	MRID Number 47381001
1	Neg Control	4			
2	37.5	4	0.054	17.3	-0.018
3	75	4	0.054	17.3	0.004
4	150	4	0.054	17.3	0.013
5	300	4	0.054	17.3	0.014
6	600	4	0.054	17.3	0.021

Dimethyl disulfide & Corn Dry Weight (g) lbs ai/A File: 1001cw Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	0.312	0.312	0.321
2	37.5	4	0.330	0.330	0.321
3	75	4	0.308	0.308	0.308
4	150	4	0.299	0.299	0.299
5	300	4	0.298	0.298	0.298
6	600	4	0.291	0.291	0.291

Dimethyl disulfide & Corn Dry Weight (g) lbs ai/A File: 1001cw Transform: NO TRANSFORMATION

IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS		TABLE WILLIAMS	DEGREES OF FREEDOM
WILLIAMS TES	r (Isotonic	regression	model)	TABLE 2 O	F 2

IDENTIFICATION	MEAN	WILLIAMS	P=.05	WILLIAMS	FREEDOM
Neg Control	0.321				
37.5	0.321	0.404		1.73	k = 1, v = 18
75	0.308	0.177		1.82	k= 2, v≈18
150	0.299	0.565		1.85	k = 3, v = 18
300	0.298	0.620		1.86	k = 4, v = 18
600	0.291	0.908		1.87	k = 5, v = 18
				~	

s = 0.032

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

Estimates of EC%

						- <b></b> -
Parameter	Estimate	95% Bo	unds	Std.Err.	Lower Bound	
		Lower	Upper		/Estimate	
EC5	2.4E+02	7.5	7.5E+03	0.72	0.032	
EC10	6.9E+02	85.	5.6E+03	0.44	0.12	
EC25	4.1E+03	19.	8.6E+05	1.1	0.0047	
EC50	3.0E+04	0.84	1.0E+09	2.2	2.8E-05	

Slope = 0.785 Std.Err. = 1.02

Goodness of fit: p = 0.72 based on DF= 3.0 18.

1001CW : Dimethyl disulfide & Corn Dry Weight (g) lbs ai/A

### PMRA Submission Number N/A

### EPA MRID Number 47381001

Observed vs.	Predicted	Treatment	Group	Means			
Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	
0.00 37.5 75.0 150. 300. 600.	4.00 4.00 4.00 4.00 4.00	0.312 0.330 0.308 0.299 0.298	0.318 0.314 0.311 0.306 0.299	0.0158 -0.00349 -0.00747 -0.00142	100. 98.9 97.9 96.4 94.1	0.00 1.15 2.08 3.58 5.88 9.19	

!!!Warning: EC10 not bracketed by doses evaluated.

!!!Warning: EC25 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Lettuce Dry Weight (g) lbs ai/A File: 1001lw Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
EXPECTED OBSERVED	1.608 0	5.808 10	9.168 8	5.808 6	1.608 0

Calculated Chi-Square goodness of fit test statistic = 6.3968
Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Lettuce Dry Weight (g) lbs ai/A File: 1001lw Transform: NO TRANSFORMATION

Shapiro Wilks test for normality

\_\_\_\_\_\_

D = 1080.630

W = 0.933

Critical W (P = 0.05) (n = 24) = 0.916 Critical W (P = 0.01) (n = 24) = 0.884

Data PASS normality test at P=0.01 level. Continue analysis.

Dimethyl disulfide & Lettuce Dry Weight (g) lbs ai/A File: 1001lw Transform: NO TRANSFORMATION

Hartley test for homogeneity of variance

-----

Calculated H statistic (max Var/min Var) = 101.43

### PMRA Submission Number N/A

EPA MRID Number 47381001

Closest, conservative, Table H statistic = 184.0 (alpha = 0.01)

Used for Table H ==> R (# groups) = 6, df (# reps-1) = 3 Actual values ==> R (# groups) = 6, df (# avg reps-1) = 3.00

\_\_\_\_\_

Data PASS homogeneity test. Continue analysis.

NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used).

Dimethyl disulfide & Lettuce Dry Weight (g) lbs ai/A File: 1001lw Transform: NO TRANSFORMATION

Bartletts test for homogeneity of variance

\_\_\_\_\_\_

Calculated B statistic = 14.22

\_\_\_\_\_\_\_\_

Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Lettuce Dry Weight (g) lbs ai/A File: 1001lw Transform: NO TRANSFORMATION

### ANOVA TABLE

SOURCE	DF	SS	MS	F
Between	5	1769.070	353.814	5.893
Within (Error)	18	1080.630	60.035	
Total	23	2849.700		

Critical F value = 2.77 (0.05,5,18) Since F > Critical F REJECT Ho:All groups equal

Dimethyl disulfide & Lettuce Dry Weight (g) lbs ai/A File: 1001lw Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 1 OF 2 Ho:Control<Treatment

TRANSFORMED MEAN CALCULATED IN
GROUP IDENTIFICATION MEAN ORIGINAL UNITS T STAT SIG

PMRA Subi	mission Number N/A			EPA MRID Nu	mber 47381001
1	Neg Control	35.425	35.425		
2	37.5	44.375	44.375	-1.634	
3	75	36.175	36.175	-0.137	
4	150	26.750	26.750	1.583	
5	300	21.625	21.625	2.519	*
6	600	20.450	20.450	2.733	*
Dunnett ta	able value = 2.41	(1 Tailed V	alue, P=0.05, di	E=18,5)	

	DUNNETTS TEST - T	CABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg Control	4			
2	37.5	4	13.204	37.3	-8.950
3	75	4	13.204	37.3	-0.750
4	150	4	13.204	37.3	8.675
5	300	4	13.204	37.3	13.800
6	600	4	13.204	37.3	14.975

Dimethyl disulfide & Lettuce Dry Weight (g) lbs ai/A File: 1001lw Transform: NO TRANSFORMATION

	WILLIAMS TEST (Isotor	nic	regression model	) TABLE 1	OF 2
GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	35.425	35.425	39.900
2	37.5	4	44.375	44.375	39.900
3	75	4	36.175	36.175	36.175
4	150	4	26.750	26.750	26.750
5	300	4	21.625	21.625	21.625
6	600	4	20.450	20.450	20.450

Dimethyl disulfide & Lettuce Dry Weight (g) lbs ai/A File: 1001lw Transform: NO TRANSFORMATION

_	WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
_	IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
	Neg Control 37.5 75 150 300 600	39.900 39.900 36.175 26.750 21.625 20.450	0.817 0.137 1.583 2.519 2.733	*	1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18
-						

### PMRA Submission Number N/A

EPA MRID Number 47381001

s = 7.748

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

Estimates of EC%

 Parameter
 Estimate
 95% Bounds
 Std.Err.
 Lower Bound

 Lower
 Upper
 /Estimate

 EC5
 32.
 2.8
 3.6E+02
 0.51
 0.089

 EC10
 59.
 8.4
 4.1E+02
 0.41
 0.14

 EC25
 1.6E+02
 49.
 5.4E+02
 0.25
 0.30

 EC50
 5.0E+02
 2.6E+02
 9.9E+02
 0.14
 0.51

Slope = 1.38 Std.Err. = 0.546

Goodness of fit: p = 0.11 based on DF = 3.0 18.

1001LW : Dimethyl disulfide & Lettuce Dry Weight (g) lbs ai/A

Observed vs. Predicted Treatment Group Means

Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	
0.00	4.00	35.4	39.7	-4.30	100.	0.00	
37.5	4.00	44.4	37.3	7.06	93.9	6.06	
75.0	4.00	36.2	34.6	1.54	87.2	12.8	
150.	4.00	26.8	30.4	-3.63	76.5	23.5	
300.	4.00	21.6	24.7	-3.04	62.1	37.9	
600.	4.00	20.5	18.2	2.26	45.8	54.2	

!!!Warning: EC5 not bracketed by doses evaluated.

Dimethyl disulfide & Onion Dry Weight (g) lbs ai/A File: 1001onw Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

\_\_\_\_\_

Calculated Chi-Square goodness of fit test statistic = 5.3637 Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Onion Dry Weight (g) lbs ai/A File: 100lonw Transform: NO TRANSFORMATION

Shapiro Wilks test for normality

D = 33.592

PMRA Submission Nu	mber N/A		EPA MR	RID Number 47381001
W = 0.951				
Critical W (P = 0. Critical W (P = 0.				
Data PASS normalit	cy test at P=0.01	level. Continue	analysis.	
Dimethyl disulfide File: 1001onw		ght (g) lbs ai/A TRANSFORMATION		
Hartley test for h				
Calculated H stati Closest, conservat			(alpha = 0.01)	
Used for Table H = Actual values =	=> R (# grou => R (# grou	ups) = 6, df ups) = 6, df	(# reps-1) = (# avg reps-1) =	3 3.00
Data PASS homogene	eity test. Contin	nue analysis.		
	liffer greatly, t	plicate sizes. If the Hartley test mage df are used).	may still be used	
Dimethyl disulfide File: 1001onw	& Onion Dry Wei Transform: NO			
Bartletts test for				
Calculated B stati Table Chi-square v Table Chi-square v	istic = 6.60 value = 15.09	(alpha = 0.01)		
Average df used in Used for Chi-squar	re table value ==	<pre>=&gt; df (#groups-1</pre>	L) = 5	
Data PASS homogene	eity test at 0.01	level. Continue	analysis.	
NOTE: If groups ha		cate sizes the avistic (see above)		size is
Dimethyl disulfide File: 1001onw		ght (g) lbs ai/A TRANSFORMATION		
		IOVA TABLE		
SOURCE	DF	ss	MS	F
Between	 5	33.535	6.707	
Within (Error)		33.592	1.866	

### PMRA Submission Number N/A

EPA MRID Number 47381001

Total	23	67.126

Critical F value = 2.77 (0.05,5,18) Since F > Critical F REJECT Ho:All groups equal

Dimethyl disulfide & Onion Dry Weight (g) lbs ai/A File: 1001onw Transform: NO TRANSFORMATION

TRANSFORMED MEAN CALCULATED IN ORIGINAL UNITS T STAT  1 Neg Control 8.855 8.855 2 37.5 8.168 8.168 0.712 3 75 8.870 8.870 -0.016 4 150 7.150 7.150 7.150 1.765		DUNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
2     37.5     8.168     8.168     0.712       3     75     8.870     8.870     -0.016       4     150     7.150     7.150     1.765	GROUP	IDENTIFICATION			T STAT	SIG
6 600 6.938 6.938 1.985	3 4 5	37.5 75 150 300	8.168 8.870 7.150 5.543	8.168 8.870 7.150 5.543	-0.016 1.765 3.429	*

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Onion Dry Weight (g) lbs ai/A File: 1001onw Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <treat< th=""><th>ment</th></treat<>	ment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)		FFERENCE OM CONTROL
1	Neg Control	4		~===== ===	
2	37.5	4	2.328	26.3	0.687
3	75	4	2.328	26.3	-0.015
4	150	4	2.328	26.3	1.705
5	300	4	2.328	26.3	3.312
6	600	4	2.328	26.3	1.917

Dimethyl disulfide & Onion Dry Weight (g) lbs ai/A File: 1001onw Transform: NO TRANSFORMATION

GROUP IDEN	TIFICATION	N 	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1 2 3 4 5	Neg Control 37.5 75 150 300 600	4 4 4 4 4	8.855 8.168 8.870 7.150 5.543 6.938	8.855 8.168 8.870 7.150 5.543 6.938	8.855 8.519 8.519 7.150 6.240 6.240

### PMRA Submission Number N/A

EPA MRID Number 47381001

Dimethyl disulfide & Onion Dry Weight (g) lbs ai/A File: 1001onw Transform: NO TRANSFORMATION

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg Control	8.855				
37.5	8.519	0.348		1.73	k = 1, v = 18
75	8.519	0.348		1.82	k = 2, v = 18
150	7.150	1.765		1.85	k = 3, v = 18
300	6.240	2.707	*	1.86	k = 4, v = 18
600	6.240	2.707	*	1.87	k = 5, v = 18

s = 1.366

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

### Estimates of EC%

Parameter	Estimate	95% Bot	ınds	Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EC5	15.	0.12	1.8E+03	1.0	0.0081
EC10	48.	1.5	1.5E+03	0.72	0.031
EC25	3.3E+02	59.	1.9E+03	0.36	0.18
EC50	2.9E+03	2.1E+02	4.1E+04	0.55	0.071

Slope = 0.719 Std.Err. = 0.434

Goodness of fit: p = 0.077 based on DF= 3.0 18.

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1001ONW : Dimethyl disulfide & Onion Dry Weight (g) lbs ai/A

Observed vs. Predicted Treatment Group Means

 Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	
0.00	4.00	8.86	9.00	-0.146	100.	0.00	
37.5	4.00	8.17	8.21	-0.0474	91.3	8.73	
75.0	4.00	8.87	7.86	1.01	87.3	12.7	
150.	4.00	7.15	7.40	-0.253	82.2	17.8	
300.	4.00	5.54	6.85	-1.30	76.1	23.9	
600.	4.00	6.94	6.20	0.738	68.9	31.1	

!!!Warning: EC5 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Oat Dry Weight lbs ai/A File: 1001ow Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

INTERVAL <-1.5 -1.5 to <-0.5 -0.5 to 0.5 >0.5 to 1.5 >1.5

INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
EXPECTED	1.474	5.324	8.404	5.324	1.474
OBSERVED	0	7	8	7	0

### PMRA Submission Number N/A

EPA MRID Number 47381001

```
Calculated Chi-Square goodness of fit test statistic = 4.0226
Table Chi-Square value (alpha = 0.01) = 13.277
Data PASS normality test. Continue analysis.
Dimethyl disulfide & Oat Dry Weight lbs ai/A
File: 1001ow Transform: NO TRANSFORMATION
Shapiro Wilks test for normality
D = 0.006
W =
    0.961
Critical W (P = 0.05) (n = 22) = 0.911
Critical W (P = 0.01) (n = 22) = 0.878
Data PASS normality test at P=0.01 level. Continue analysis.
Dimethyl disulfide & Oat Dry Weight lbs ai/A
File: 1001ow Transform: NO TRANSFORMATION
Hartley test for homogeneity of variance
Calculated H statistic (max Var/min Var) = 6.15 Closest, conservative, Table H statistic = 184.0 (alpha = 0.01)
Used for Table H ==> R (\# groups) = 6, df (\# reps-1) = 3 Actual values ==> R (\# groups) = 6, df (\# avg reps-1) = 2.67
                                          (average df used)
_______
Data PASS homogeneity test. Continue analysis.
NOTE: This test requires equal replicate sizes. If they are unequal
     but do not differ greatly, the Hartley test may still be used
     as an approximate test (average df are used).
Dimethyl disulfide & Oat Dry Weight lbs ai/A
File: 1001ow Transform: NO TRANSFORMATION
Bartletts test for homogeneity of variance
Calculated B statistic = 2.31
Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05)
Average df used in calculation ==> df (avg n - 1) = 2.67
Used for Chi-square table value ==> df (\#groups-1) = 5
______
```

### PMRA Submission Number N/A

EPA MRID Number 47381001

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Oat Dry Weight lbs ai/A File: 1001ow Transform: NO TRANSFORMATION

### ANOVA TABLE

SOURCE	DF	SS	MS	F
Between	5	0.0025	0.0005	1.250
Within (Error)	16	0.0056	0.0004	
Total	21	0.0081		

Critical F value = 2.85 (0.05,5,16)
Since F < Critical F FAIL TO REJECT Ho:All groups equal

Dimethyl disulfide & Oat Dry Weight lbs ai/A File: 1001ow Transform: NO TRANSFORMATION

	BONFERRONI T-TEST -	TABLE 1 OF 2	Ho: Contro	l <treatm< th=""><th>ent</th></treatm<>	ent
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg control	0.154	0.154	<b>-</b>	
2	37.5	0.159	0.159	-0.265	
3	75	0.139	0.139	0.971	
4	150	0.131	0.131	1.528	
5	300	0.129	0.129	1.675	
6	600	0.143	0.143	0.775	

Bonferroni T table value = 2.58 (1 Tailed Value, P=0.05, df=16,5)

Direction of collection and the collection of th

Dimethyl disulfide & Oat Dry Weight lbs ai/A File: 1001ow Transform: NO TRANSFORMATION

I	BONFERRONI T-TEST -	TABLE	2 OF 2	Ho:Contr	ol <treatment< th=""></treatment<>
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg control	3			
2	37.5	3	0.042	27.3	-0.004
3	75	4	0.039	25.6	0.015
4	150	4	0.039	25.6	0.023
5	300	4	0.039	25.6	0.026
6	600	4	0.039	25.6	0.012

### PMRA Submission Number N/A

EPA MRID Number 47381001

Dimethyl disulfide & Oat Dry Weight lbs ai/A File: 1001ow Transform: NO TRANSFORMATION

WILLIAMS	TEST (	Tsotonic	regression	model'	TABLE	1	OF	2

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg control	3	0.154	0.154	0.157
2	37.5	3	0.159	0.159	0.157
3	75	4	0.139	0.139	0.139
4	150	4	0.131	0.131	0.134
5	300	4	0.129	0.129	0.134
6	600	4	0.143	0.143	0.134

Dimethyl disulfide & Oat Dry Weight lbs ai/A File: 1001ow Transform: NO TRANSFORMATION

WILLIAMS	TEST	(Isotonic	regression	model)	TABLE	2	OF	2
----------	------	-----------	------------	--------	-------	---	----	---

IDENTIFICATION	ISOTONIZED	CALC.	SIG	TABLE	DEGREES OF
	MEAN	WILLIAMS	P=.05	WILLIAMS	FREEDOM
Neg control 37.5 75 150 300 600	0.157 0.157 0.139 0.134 0.134	0.142 1.038 1.417 1.417		1.75 1.83 1.86 1.87	k= 1, v=16 k= 2, v=16 k= 3, v=16 k= 4, v=16 k= 5, v=16

s = 0.019

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

#### Estimates of EC%

Parameter	Estimate	95% Boi	 ınds	Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EC5	9.4	3.7E-05	2.4E+06	2.6	3.9E-06
EC10	1.3E+02	0.18	1.0E+05	1.4	0.0013
EC25	1.1E+04	0.30	4.2E+08	2.2	2.7E-05
EC50	1.6E+06	8.8E-05	2.8E+16	4.9	5.7E-11

Slope = 0.315 Std.Err. = 0.427

Goodness of fit: p =	0.26	based on DF=	3.0	16.	
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10010M . Dimothyl digulfide : Oat Dwy Weight lbg ci/h

10010W : Dimethyl disulfide & Oat Dry Weight lbs ai/A

Observed vs Predicted Treatment Group Means

Observed	v.	Fredreced	rreacment	Group	Means

Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	
0.00	3.00	0.154	0.155	-0.00115	100.	0.00	
37.5	3.00	0.159	0.144	0.0145	92.7	7.27	
75.0	4.00	0.140	0.142	-0.00250	91.3	8.67	

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PMRA Submiss	sion Numbe	r N/A			EPA_	MRID Number 473810	01
150.	4.00	0.131		-0.00852	89.7	10.3	
300.	4.00	0.129	0.137	-0.00797	87.9	12.1	
600.	4.00	0.143	0.134	0.00891	85.9	14.1	
!!!Warning:				valuated.			
!!!Warning:							
J.			2				
!!!Warning:	EC50 not	bracketed b	y doses	evaluated.			
Dimethyl dis File: 1001rw	ulfide & Tr	Ryegrass Dr ansform: NC	ry Weight TRANSFO	. (g) lbs ai/A RMATION			
				d expected fr			
INTERVAL	<-1.5	-1.5 to <	:-0.5	-0.5 to 0.5	>0.5 to	1.5 >1.5	
EXPECTED			1	9.168	5.80	8 1.608	
OBSERVED	0	7		11	6	0	
	hi-Square uare valu	goodness c e (alpha =	of fit te 0.01) =	est statistic 13.277			
Dimethyl dis File: 1001rw				(g) lbs ai/A	7		
Shapiro Wilk				MINITON			
SHapiro Wilk	s test IO		,				
D = 166.712							
W = 0.976							
Critical W (Critical W (	P = 0.01)	(n = 24) =	0.884				
Data PASS no	rmality t	est at P=0.	01 level	. Continue an	alysis.		
ماد الساطمات	7 5 2 3	D	70-3-1-1-4	· (\ 31   / 7			
File: 1001rw		ransform: N		: (g) lbs ai/A CORMATION	ı		
Hartley test	for homo	geneity of	variance				
Calculated H Closest, con				= 8.85 c = 184.0 (a	alpha = 0.01	)	
Used for Tab Actual value	le H ==>	R (# gr R (# gr	coups) =	6, df (# 6, df (#	reps-1) =	3 .) = 3.00	
		_	==		_ <del>_</del>		

### PMRA Submission Number N/A

EPA MRID Number 47381001

Data PASS homogeneity test. Continue analysis.

NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used).

Dimethyl disulfide & Ryegrass Dry Weight (g) lbs ai/A File: 1001rw Transform: NO TRANSFORMATION

Bartletts test for homogeneity of variance

Calculated B statistic  $\approx$  4.10 Table Chi-square value = 15.09 (alpha = 0.01) Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 3.00Used for Chi-square table value ==> df (#groups-1) = 5

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Ryegrass Dry Weight (g) lbs ai/A File: 1001rw Transform: NO TRANSFORMATION

### ANOVA TABLE

SOURCE	DF	SS	MS	F
Between	5	93.383	18.677	2.017
Within (Error)	18	166.712	9.262	
Total	23	260.095		

Critical F value = 2.77 (0.05, 5, 18)Since F < Critical F FAIL TO REJECT Ho: All groups equal

Dimethyl disulfide & Ryegrass Dry Weight (g) lbs ai/A File: 1001rw Transform: NO TRANSFORMATION

D,	UNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1 2 3 4 5	Neg Control 37.5 75 150 300 600	13.525 14.833 11.830 10.417 8.978 10.638	13.525 14.833 11.830 10.417 8.978 10.638	-0.608 0.788 1.444 2.113 1.342	

### PMRA Submission Number N/A

EPA MRID Number 47381001

Dunnett table value = 2.41

(1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Ryegrass Dry Weight (g) lbs ai/A File: 1001rw Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg Control	4			
2	37.5	4	5.186	38.3	-1.307
3	75	4	5.186	38.3	1.695
4	150	4	5.186	38.3	3.108
5	300	4	5.186	38.3	4.548
6	600	4	5.186	38.3	2.887

Dimethyl disulfide & Ryegrass Dry Weight (g) lbs ai/A File: 1001rw Transform: NO TRANSFORMATION

	WILLIAMS TEST (Isoton	nic	regression mode	1) TABLE 1 O	F 2
GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	13.525	13.525	14.179
2	37.5	4	14.833	14.833	14.179
3	75	4	11.830	11.830	11.830
4	150	4	10.417	10.417	10.417
5	300	4	8.978	8.978	9.808
6	600	4	10.638	10.638	9.808

Dimethyl disulfide & Ryegrass Dry Weight (g) lbs ai/A File: 1001rw Transform: NO TRANSFORMATION

WILLIAMS TES	T (Isotonic	regression	model)	TABLE 2 C	OF 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg Control 37.5 75 150 300 600	14.179 14.179 11.830 10.417 9.808 9.808	0.304 0.788 1.444 1.728		1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 3.043

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

Estimates	of EC%					
Parameter	Estimate	95% Bot	ınds	Std.Err.	Lower Bound	
		Lower	Upper		/Estimate	

<u>PMRA Submi</u>	ssion Numbe					<u>EPA MRID</u>	Number 4738
EC5	14.	0.024	7.7E+03	1.3	0.0018	}	
EC10	43.	0.42	4.4E+03	0.97	0.0098	}	
EC25 EC50	2.9E+02	28.	3.0E+03	0.49	0.097	,	
:C50	2.4E+03	99.	5.7E+04	0.66	0.042		
Sl	ope = 0	.734 Std.	Err. =	0.572			
Goodness of						18.	
LOO1RW : Di	methyl dis	ulfide & R	yegrass Dı	ry Weight	(g) lbs ai		· · ·
bserved vs	. Predicte	d Treatmen	t Group Me	eans			
Dose		Obs.					
Dobe	WKCPD.	Mean	Mean	-Pred.	%Control	·	
0.00	4.00	13.5	14.1	-0.532	100.	0.00	
37.5	4 00	14 8	12 7	2 08	90.7	9 31	
75.0	4 00	11 Ω	12.7	-0 333	26.7	13 6	
150.	4 00	13.5 14.8 11.8 10.4 8.98	11 /	-0.525	00.5	10 0	
	4.00	70.4	1 A E	- U • D / /	47 L	10.9	
300.	4.00	10.6	10.5	-T.50	74.5	33.1	
!Warning:	EC50 not	bracketed l	by doses e	evaluated.			
oimethyl di:	sulfide & ow T	Sugarbeet lansform: lormality:	Dry Weight	t (g) lbs a	frequenci	~	
imethyl di:	sulfide & ow T	Sugarbeet lansform: lormality:	Dry Weight	t (g) lbs a	frequenci	~	
Dimethyl di: File: 1001sl Chi-square	sulfide & Town Town Town Town Town Town Town Town	Sugarbeet I ransform: I ormality:	Dry Weight NO TRANSFO actual and	t (g) lbs a	frequenci .5 >0.5	~	>1.5
Calculated ( Fable Chi-So	sulfide & ow To test for no series for no se	Sugarbeet I ransform: I ormality: 6 -1.5 to 6 - 5.80 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	Dry Weight NO TRANSFO actual and <-0.5 8  of fit tes 0.01) = 1	c (g) lbs a DRMATION d expected -0.5 to 0 9.168 7 st statists	frequenci	to 1.5  5.808	>1.5 
Dimethyl dis File: 1001sl Chi-square d INTERVAL EXPECTED DBSERVED	sulfide & ow To test for no series for no se	Sugarbeet I ransform: I ormality: 6 -1.5 to 6 - 5.80 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9	Dry Weight NO TRANSFO actual and <-0.5 8  of fit tes 0.01) = 1	c (g) lbs a DRMATION d expected -0.5 to 0 9.168 7 st statists	frequenci	to 1.5  5.808	>1.5 
Dimethyl distile: 1001sl Chi-square to the control of the control	sulfide & ow The test for no section of the section	Sugarbeet I ransform: I ormality: 6	Dry Weight NO TRANSFO  actual and  <-0.5   8  of fit tes 0.01) = 1  nue analys	(g) lbs approximation of the control	frequenci .5 >0.5 — —	to 1.5  5.808	>1.5 
cimethyl disciple: 1001sl	sulfide & ow The test for no control of the square value or mality to sulfide & sow The sow The sulfide & sow The sow	Sugarbeet I ransform: I sugarbeet I sugarbeet I ransform: I r normality:	Dry Weight NO TRANSFO  actual and  <-0.5  8  of fit tes  0.01) = 1  nue analys  Dry Weight NO TRANSFO  y	(g) lbs approximation of the control	frequenci .5 >0.5 .ic = 6.3	to 1.5  5.808 8	>1.5 
Dimethyl disile: 1001sl Chi-square ( Chi-squ	sulfide & ow Titest for no control of the second of the se	Sugarbeet I ransform: I sugarbeet I sugarbeet I ransform: I r normality:	Dry Weight NO TRANSFO  actual and  <-0.5  8  of fit tes  0.01) = 1  nue analys  Dry Weight NO TRANSFO  y	g (g) lbs a DRMATION  d expected  -0.5 to 0  9.168 7  st statists 3.277  sis.  (g) lbs a DRMATION	frequenci .5 >0.5 .ic = 6.3	to 1.5  5.808 8	>1.5 
cimethyl disciple: 1001sl chi-square control chi-square control chi-square control chi-square control chi-square control calculated control calcul	sulfide & ow Titest for no control of the sulfide & sulfide & sow Tites test for no control of the sulfide & sow Tites	Sugarbeet I ransform: I sugarbeet I sugarbeet I ransform: I r normality:	Dry Weight NO TRANSFO  actual and  <-0.5  8  of fit tes  0.01) = 1  nue analys  Dry Weight NO TRANSFO  y	g (g) lbs a DRMATION  d expected  -0.5 to 0  9.168 7  st statists 3.277  sis.  (g) lbs a DRMATION	frequenci .5 >0.5 .ic = 6.3	to 1.5  5.808 8	>1.5 

Data PASS normality test at P=0.01 level. Continue analysis.

### PMRA Submission Number N/A

Critical F value =

EPA MRID Number 47381001

```
Dimethyl disulfide & Sugarbeet Dry Weight (g) lbs ai/A
File: 1001sbw Transform: NO TRANSFORMATION
Hartley test for homogeneity of variance
Calculated H statistic (max Var/min Var) = 10.16
Closest, conservative, Table H statistic = 184.0 (alpha = 0.01)
Used for Table H ==> R (\# groups) = 6, df (\# reps-1) = 3 Actual values ==> R (\# groups) = 6, df (\# avg reps-1) = 3.00
Data PASS homogeneity test. Continue analysis.
NOTE: This test requires equal replicate sizes. If they are unequal
    but do not differ greatly, the Hartley test may still be used
     as an approximate test (average df are used).
Dimethyl disulfide & Sugarbeet Dry Weight (g) lbs ai/A
File: 1001sbw Transform: NO TRANSFORMATION
Bartletts test for homogeneity of variance
Calculated B statistic = 3.95
Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05)
Average df used in calculation ==> df (avg n - 1) = 3.00
Used for Chi-square table value ==> df (#groups-1) = 5
Data PASS homogeneity test at 0.01 level. Continue analysis.
NOTE: If groups have unequal replicate sizes the average replicate size is
    used to calculate the B statistic (see above).
Dimethyl disulfide & Sugarbeet Dry Weight (g) lbs ai/A
File: 1001sbw Transform: NO TRANSFORMATION
                         ANOVA TABLE
______
              DF
                           SS
                                           MS
Between 5 15138.955 3027.791 8.704
Within (Error) 18
                         6261.590
                                   347.866
              23 21400.545
_______
```

2.77 (0.05,5,18)

Since F > Critical F REJECT Ho:All groups equal

### PMRA Submission Number N/A

EPA MRID Number 47381001

Dimethyl disulfide & Sugarbeet Dry Weight (g) lbs ai/A File: 1001sbw Transform: NO TRANSFORMATION

	DUNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg Control	76.675	76.675		
2	37.5	90.975	90.975	-1.084	
3	75	67.650	67.650	0.684	
4	150	47.975	47.975	2.176	
5	300	29.825	29.825	3.552	*
6	600	20.650	20.650	4.248	*

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Sugarbeet Dry Weight (g) lbs ai/A File: 1001sbw Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg Control	4			
2	37.5	4	31.784	41.5	-14.300
3	75	4	31.784	41.5	9.025
4	150	4	31.784	41.5	28.700
5	300	4	31.784	41.5	46.850
6	600	4	31.784	41.5	56.025

Dimethyl disulfide & Sugarbeet Dry Weight (g) lbs ai/A File: 1001sbw Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

GROUP ORIGINAL TRANSFORMED ISOTONIZE

IDENTIFICATION N MEAN MEAN MEAN MEAN MEAN

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	76.675	76.675	83.825
2	37.5	4	90.975	90.975	83.825
3	75	4	67.650	67.650	67.650
4	150	4	47.975	47.975	47.975
5	300	4	29.825	29.825	29.825
6	600	4	20.650	20.650	20.650

Dimethyl disulfide & Sugarbeet Dry Weight (g) lbs ai/A File: 1001sbw Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

ISOTONIZED CALC. SIG TABLE DEGREES OF

<b>PMRA</b>	Submission	Number N/A	A
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EPA MRID Number 47381001

IDENTIFICATION	MEAN	WILLIAMS	P=.05	WILLIAMS	FREEDOM
Neg Control	83.825				
37.5	83.825	0.542		1.73	k = 1, v = 18
75	67.650	0.684		1.82	k = 2, v = 18
150	47.975	2.176	*	1.85	k = 3, v = 18
300	29.825	3.552	*	1.86	k = 4, v = 18
600	20.650	4.248	*	1.87	k = 5, v = 18

s = 18.651

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

#### Estimates of EC%

Parameter	Estimate	95% Bounds		Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EC5	27.	4.0	1.8E+02	0.40	0.15
EC10	42.	8.4	2.1E+02	0.34	0.20
EC25	91.	29.	2.9E+02	0.24	0.32
EC50	2.1E+02	1.1E+02	4.4E+02	0.15	0.49

Slope = 1.82 Std.Err. = 0.573

Goodness of fit: p = 0.57 based on DF= 3.0 18.

1001SBW : Dimethyl disulfide & Sugarbeet Dry Weight (g) lbs ai/A

Observed vs. Predicted Treatment Group Means

Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change
0.00	4.00	76.7	84.9	-8.26	100.	0.00
37.5	4.00	91.0	77.8	13.2	91.6	8.40
75.0	4.00	67.7	67.7	-0.0472	79.7	20.3
150.	4.00	48.0	52.0	-3.98	61.2	38.8
300.	4.00	29.8	33,6	-3.82	39.6	60.4
600.	4.00	20.6	17.7	2.92	20,9	79.1

!!!Warning: EC5 not bracketed by doses evaluated.

Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A File: 1001sw Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
					<del></del>
EXPECTED OBSERVED	1.608 0	5.808 8	9.168 8	5.808 8	1.608 0

Calculated Chi-Square goodness of fit test statistic = 5.0194
Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

### PMRA Submission Number N/A

EPA MRID Number 47381001

```
Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A File: 1001sw Transform: NO TRANSFORMATION
Shapiro Wilks test for normality
0.024
W = 0.941
Critical W (P = 0.05) (n = 24) = 0.916
Critical W (P = 0.01) (n = 24) = 0.884
                                    ______
Data PASS normality test at P=0.01 level. Continue analysis.
Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A
File: 1001sw Transform: NO TRANSFORMATION
Hartley test for homogeneity of variance
Calculated H statistic (max Var/min Var) = 13.47
Closest, conservative, Table H statistic = 184.0 (alpha = 0.01)
Used for Table H ==> R (\# groups) = 6, df (\# reps-1) = Actual values ==> R (\# groups) = 6, df (\# avg reps-1) =
                                                                   3.00
______
Data PASS homogeneity test. Continue analysis.
NOTE: This test requires equal replicate sizes. If they are unequal
     but do not differ greatly, the Hartley test may still be used
     as an approximate test (average df are used).
Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A
File: 1001sw Transform: NO TRANSFORMATION
Bartletts test for homogeneity of variance
                                       Calculated B statistic = 7.91
Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05)
Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (\#groups-1) = 5
Data PASS homogeneity test at 0.01 level. Continue analysis.
NOTE: If groups have unequal replicate sizes the average replicate size is
     used to calculate the B statistic (see above).
Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A
File: 1001sw Transform: NO TRANSFORMATION
```

### PMRA Submission Number N/A

### EPA MRID Number 47381001

ANOVA TABLE								
SOURCE	DF	SS	MS	F				
Between	5	0.015	0.003	3.000				
Within (Error)	18	0.024	0.001					
Total	23	0.038						

Critical F value = 2.77 (0.05,5,18) Since F > Critical F REJECT Ho:All groups equal

Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A File: 1001sw Transform: NO TRANSFORMATION

]	DUNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg control	0.268	0.268		
2	37.5	0.269	0.269	-0.022	
3	75	0.292	0.292	-1.073	
4	150	0.239	0.239	1.286	
5	300	0.242	0.242	1.174	
6	600	0.216	0.216	2.337	
Dunnet	t table value = 2.41	(1 Tailed V	Value, P=0.05, df=18,	5)	

Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A File: 1001sw Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1	Neg control	4			
2	37.5		0.054	20.1	-0.000
3	75	4	0.054	20.1	-0.024
4	150	4	0.054	20.1	0.029
5	300	4	0.054	20.1	0.026
6	600	4	0.054	20.1	0.052

Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A File: 1001sw Transform: NO TRANSFORMATION

	IDENTIFICATIO	N N	MEAN	MEAN	MEAN
GROUP	,		ORIGINAI	TRANSFORMED	ISOTONIZED
	WILLIAMS TEST	(Isotonic	regression	model) TABLE 1	OF 2

PMRA Subm	nission Number N/A	EPA	MRID Number 47381001		
1	Neg control	4	0.268	0.268	0.276
2	37.5	4	0.269	0.269	0.276
3	75	4	0.292	0.292	0.276
4	150	4	0.239	0.239	0.241
5	300	4	0.242	0.242	0.241
6	600	4	0.216	0.216	0.216

Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A File: 1001sw Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

IDENTIFICATION	ISOTONIZED	CALC.	SIG	TABLE	DEGREES OF
	MEAN	WILLIAMS	P=.05	WILLIAMS	FREEDOM
Neg control 37.5 75 150 300 600	0.276 0.276 0.276 0.241 0.241 0.216	0.319 0.319 1.075 1.075 2.042	*	1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 0.036

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

### Estimates of EC%

Parameter	Estimate	95% Bounds		Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EC5	1.3E+02	9.8	1.6E+03	0.53	0.078
EC10	2.4E+02	47.	1.2E+03	0.34	0.19
EC25	7.2E+02	2.9E+02	1.8E+03	0.19	0.41
EC50	2.4E+03	2.5E+02	2.3E+04	0.47	0.10

Slope = 1.28 Std.Err. = 0.937

Goodness of fit: p = 0.41 based on DF= 3.0 18.

1001SW : Dimethyl disulfide & Soybean Dry Weight (g) lbs ai/A

\_\_\_\_\_\_

Observed vs. Predicted Treatment Group Means

 					<b></b>		
 Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	
0.00	4.00	0.268	0.274	-0.00636	100.	0.00	
37.5	4.00	0.268	0.272	-0.00304	99.0	1.03	
75.0	4.00	0.292	0.267	0.0250	97.3	2.68	
150.	4.00	0.239	0.258	-0.0183	93.9	6.12	
300.	4.00	0.242	0.241	0.00119	87.7	12.3	
600.	4.00	0.216	0.214	0.001.63	78.0	22.0	

!!!Warning: EC25 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Tomato Dry Weight (g) lbs ai/A

### PMRA Submission Number N/A

EPA MRID Number 47381001

Chi-square test for normality: actual and expected frequencies

Transform: NO TRANSFORMATION

INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
EXPECTED OBSERVED	1.608	5.808 7	9.168 9	5.808 8	1.608 0

Coloulated Chi Course goodness of fit test statistis - 4 2010

Calculated Chi-Square goodness of fit test statistic = 4.2910 Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Tomato Dry Weight (g) lbs ai/A File: 1001tw Transform: NO TRANSFORMATION

Shapiro Wilks test for normality

D = 1707.212

W = 0.959

Critical W (P = 0.05) (n = 24) = 0.916 Critical W (P = 0.01) (n = 24) = 0.884

-----

Data PASS normality test at P=0.01 level. Continue analysis.

Dimethyl disulfide & Tomato Dry Weight (g) lbs ai/A File: 1001tw Transform: NO TRANSFORMATION

Hartley test for homogeneity of variance

\_\_\_\_\_\_\_

Calculated H statistic (max Var/min Var) = 19.05Closest, conservative, Table H statistic = 184.0 (alpha = 0.01)

Used for Table H ==> R (# groups) = 6, df (# reps-1) = 3 Actual values ==> R (# groups) = 6, df (# avg reps-1) = 3.00

Data PASS homogeneity test. Continue analysis.

NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used).

Dimethyl disulfide & Tomato Dry Weight (g) lbs ai/A File: 1001tw Transform: NO TRANSFORMATION

Bartletts test for homogeneity of variance

#### PMRA Submission Number N/A

EPA MRID Number 47381001

Calculated B statistic = 9.31 Table Chi-square value = 15.09 (alpha = 0.01) Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Tomato Dry Weight (g) lbs ai/A File: 1001tw Transform: NO TRANSFORMATION

#### ANOVA TABLE \_\_\_\_\_\_

DF SS MS F \_\_\_\_\_ Between 5 1684.417 336.883 3.552 Within (Error) 18 1707.212 94.845 23 3391.630 

Critical F value  $\approx$  2.77 (0.05,5,18) Since F > Critical F REJECT Ho: All groups equal

Dimethyl disulfide & Tomato Dry Weight (g) lbs ai/A File: 1001tw Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 1 OF 2 Ho:Control<Treatment \_\_\_\_\_ TRANSFORMED MEAN CALCULATED IN GROUP IDENTIFICATION MEAN ORIGINAL UNITS ORIGINAL UNITS T STAT SIG Neg Control 44.850 37.5 44.800 75 38.875 150 52.375 300 34.100 600 26.375 44.850 2 44.800 0.007 38.875 0.868 3 52.375 34.100 1.561 26.375 2.683 \* \_\_\_\_\_\_

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Tomato Dry Weight (g) lbs ai/A File: 1001tw Transform: NO TRANSFORMATION

	DUNNETTS TEST	-			-	2 Ho:	Control <t< th=""><th></th></t<>	
						Minimum Sig Diff		
GROUP	IDENTIFICATION	1	REPS	3		(IN ORIG. UNITS)	CONTROL	FROM CONTROL

PMRA Sul	bmission Number N/A			EPA 1	MRID Number <u>47381001</u>
1	Neg Control	4			
2	37.5	4	16.596	37.0	0.050
3	75	4	16.596	37.0	5.975
4	150	4	16.596	37.0	-7.525
5	300	4	16.596	37.0	10.750
6	600	4	16.596	37.0	18.475

TITT T TANK	min Cm	/Tactonia		model	ייי דרד איייי ו	7	OB	~
MITTTAMS	1551	(TROCOUTE	regression	moder.	) TABLE	т.	O.F	4

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1 2 3 4 5	Neg Control 37.5 75 150 300 600	4 4 4 4 4	44.850 44.800 38.875 52.375 34.100 26.375	44.850 44.800 38.875 52.375 34.100 26.375	45.225 45.225 45.225 45.225 34.100 26.375

Dimethyl disulfide & Tomato Dry Weight (g) lbs ai/A File: 1001tw Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 2 OF 2

IDENTIFICATION	ISOTONIZED	CALC.	SIG	TABLE	DEGREES OF
	MEAN	WILLIAMS	P=.05	WILLIAMS	FREEDOM
Neg Control 37.5 75 150 300 600	45.225 45.225 45.225 45.225 34.100 26.375	0.054 0.054 0.054 1.561 2.683	*	1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 9.739

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

#### Estimates of EC%

Parameter	Estimate	95% Bounds		Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EC5	1.8E+02	42.	7.9E+02	0.31	0.23
EC10	2.4E+02	80.	7.5E+02	0.23	0.33
EC25	4.0E+02	2.2E+02	7.2E+02	0.12	0.56
EC50	6.9E+02	4.3E+02	1.1E+03	0.10	0.62

Slope = 2.83 Std.Err. = 1.70

Goodness of fit: p = 0.19 based on DF= 3.0 18.

1001TW : Dimethyl disulfide & Tomato Dry Weight (g) lbs ai/A  $\,$ 

#### PMRA Submission Number N/A

EPA MRID Number 47381001

							_
Observed vs.	Predicted	Treatment	Group	Means			
Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	•
0.00 37.5 75.0 150. 300.	4.00 4.00 4.00 4.00 4.00	44.8 44.8 38.9 52.4 34.1 26.4	44.8 44.8 44.7 43.5 38.0 25.6	0.00356 -5.79 8.91 -3.92	100. 100. 99.7 97.0 84.9 57.1	0.00 0.0168 0.313 2.99 15.1 42.9	

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Corn 21-day Survival lbs a.i./A File: 1001cs Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
		<del></del>			<del></del>
EXPECTED OBSERVED	1.608 0	5.808 5	9.168 16	5.808 3	1.608 0

\_\_\_\_\_

Calculated Chi-Square goodness of fit test statistic = 9.7772
Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Corn 21-day Survival lbs a.i./A File: 1001cs Transform: NO TRANSFORMATION

Shapiro Wilks test for normality

\_\_\_\_\_

D = 29.500

W = 0.912

Critical W (P = 0.05) (n = 24) = 0.916 Critical W (P = 0.01) (n = 24) = 0.884

Data PASS normality test at P=0.01 level. Continue analysis.

Dimethyl disulfide & Corn 21-day Survival lbs a.i./A File: 1001cs Transform: NO TRANSFORMATION

Hartley test for homogeneity of variance Bartletts test for homogeneity of variance

These two tests can not be performed because at least one group has zero variance

#### PMRA Submission Number N/A

EPA MRID Number 47381001

Data FAIL to meet homogeneity of variance assumption. Additional transformations are useless.

\_\_\_\_\_\_

Dimethyl disulfide & Corn 21-day Survival lbs a.i./A File: 1001cs Transform: NO TRANSFORMATION

	STEELS MANY-ONE RAI	NK TEST -	Но	:Control<	reatmen	t
GROUP	IDENTIFICATION	TRANSFORMED MEAN	RANK SUM	CRIT. VALUE	df	SIG
1	Neg Control	10.000				
2	37.5	8.500	10.00	10.00	4.00	*
3	75	9.500	16.00	10.00	4.00	
4	150	9.000	12.00	10.00	4.00	
5	300	7.750	12.00	10.00	4.00	
6	600	7.750	12.00	10.00	4.00	

Critical values use k = 5, are 1 tailed, and alpha = 0.05

Dimethyl disulfide & Corn 21-day Survival lbs a.i./A File: 1001cs Transform: NO TRANSFORMATION

KRUSKAL-WALLIS ANOVA BY RANKS - TABLE 1 OF 2

GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	RANK SUM
1	Neg Control	10.000	10.000	78.000
2	37.5	8.500	8.500	39.500
3	75	9.500	9.500	65,500
4	150	9.000	9.000	50.500
5	300	7.750	7.750	34.500
6	600	7.750	7.750	32.000

Calculated H Value = 9.331 Critical H Value Table = 11.070 Since Calc H < Crit H FAIL TO REJECT Ho: All groups are equal.

Dimethyl disulfide & Corn 21-day Survival lbs a.i./A File: 1001cs Transform: NO TRANSFORMATION

DUNNS MULTIPLE COMPARISON - KRUSKAL-WALLIS - TABLE 2 OF 2 ------

GROUP TRANSFORMED ORIGINAL 0 0 0 0 0 GROUP IDENTIFICATION MEAN MEAN 5 6 2 4 3 1 300 7.750 7.750 \
600 7.750 7.750 \
37.5 8.500 8.500 \
150 9.000 9.000 \
75 9.500 9.500 \
Neg Control 10.000 10.000 \\ 2 4

### PMRA Submission Number N/A

EPA MRID Number 47381001

\* = significant difference (p=0.05) . = no significant difference Table q value (0.05,6) = 2.936 SE = 4.771

Estimates of EC%

 Parameter
 Estimate
 95% Bounds Lower
 Std.Err. Lower Bound /Estimate

 EC5
 17.
 0.075
 3.7E+03
 1.1
 0.0045

 EC10
 72.
 2.1
 2.5E+03
 0.74
 0.029

 EC25
 8.4E+02
 97.
 7.2E+03
 0.45
 0.12

 EC50
 1.3E+04
 88.
 1.8E+06
 1.0
 0.0070

Slope = 0.571 Std.Err. = 0.394

Goodness of fit: p = 0.47 based on DF= 3.0 18.

1001CS : Dimethyl disulfide & Corn 21-day Survival lbs a.i./A

1001CS : Dimethyl disullide & Colh 21-day Sulvival ibs a.1./A

Observed vs. Predicted Treatment Group Means

			~			
Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change
0.00	4.00	10.0	9.93	0.0704	100.	0.00
37.5	4.00	8.50	9.19	-0.691	92.6	7.44
75.0	4.00	9.50	8.92	0.580	89.8	10.2
150.	4.00	9.00	8.58	0.417	86.4	13.6
300.	4.00	7.75	8.18	-0.426	82.3	17.7
600.	4.00	7.75	7.70	0.0508	77.5	22.5

!!!Warning: EC5 not bracketed by doses evaluated.

!!!Warning: EC25 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

Interpreted as Dose = 0

Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A File: 1001ls Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

-----

INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
EXPECTED OBSERVED	1.608	5.808 5	9.168 14	5.808 5	1.608 0

Calculated Chi-Square goodness of fit test statistic = 5.9875

Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A File: 1001ls Transform: NO TRANSFORMATION

PMRA Submission Number N/A	EPA MRID Number 47381001
Shapiro Wilks test for normality	
D = 15.750	
W = 0.927	
Critical W (P = 0.05) (n = 24) = 0.916 Critical W (P = 0.01) (n = 24) = 0.884	
Data PASS normality test at P=0.01 level. Continue analysis.	
Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A File: 1001ls Transform: NO TRANSFORMATION	
Hartley test for homogeneity of variance	
Calculated H statistic (max Var/min Var) = 9.00 Closest, conservative, Table H statistic = 184.0 (alpha =	0.01)
Used for Table H ==> R (# groups) = 6, df (# reps-1 Actual values ==> R (# groups) = 6, df (# avg re	1) = 3 eps-1) = 3.00
Data PASS homogeneity test. Continue analysis.	
	_
NOTE: This test requires equal replicate sizes. If they are but do not differ greatly, the Hartley test may still as an approximate test (average df are used).	
Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A File: 1001ls Transform: NO TRANSFORMATION	
Bartletts test for homogeneity of variance	
Calculated B statistic = 6.62  Table Chi-square value = 15.09 (alpha = 0.01)  Table Chi-square value = 11.07 (alpha = 0.05)	
Average df used in calculation $==>$ df (avg n - 1) = 3.0 Used for Chi-square table value $==>$ df (#groups-1) = 5	00
Data PASS homogeneity test at 0.01 level. Continue analysis	
NOTE: If groups have unequal replicate sizes the average repused to calculate the B statistic (see above).	olicate size is
Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A File: 1001ls Transform: NO TRANSFORMATION	
ANOVA TABLE	,

PMRA Submission Number N/A

EPA MRID Number 47381001

SOURCE	DF	SS	MS	F	
Between	5	12.875	2.575	2.943	
Within (Error)	18	15.750	0.875		
Total	23	28.625			

Critical F value = 2.77 (0.05, 5, 18)

Since F > Critical F REJECT Ho: All groups equal

Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A File: 1001ls Transform: NO TRANSFORMATION

DUNNETTS TEST - TABLE 1 OF 2 Ho:Control<Treatment

TRANSFORMED MEAN CALCULATED IN
ORIGINAL UNITS T STAT SIG

Neg Control 9.750 9.750

1 Neg Control 9.750 9.750
2 37.5 8.750 8.750 1.512
3 75 9.750 9.750 0.000
4 150 7.750 7.750 3.024 \*
5 300 8.250 8.250 2.268
6 600 9.000 9.000 1.134

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A File: 1001ls Transform: NO TRANSFORMATION

Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A File: 1001ls Transform: NO TRANSFORMATION

WILLIAMS TEST (Isotonic regression model) TABLE 1 OF 2

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	9.750	9.750	9.750
2	37.5	4	8.750	8.750	9.250
3	75	4	9.750	9.750	9.250
4	150	4	7.750	7.750	8.333

PMRA Submission	Number N/A			EPA	MRID Number 473810	)01
5	300	4	8.250	8.250	8.333	
6	600	4	9.000	9.000	8.333	

Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A File: 1001ls Transform: NO TRANSFORMATION

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg Control 37.5 75 150 300 600	9:750 9:250 9:250 8:333 8:333 8:333	0.756 0.756 2.142 2.142 2.142	* * *	1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 0.935

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

#### Estimates of EC%

Parameter	Estimate	95% Bounds		Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EC5	0.38	6.9E-14	2.1E+12	6.1	1.8E-13
EC10	79.	0.0020	3.2E+06	2.2	2.5E-05
EC25	5.8E+05	9.8E-10	3.4E+20	7.1	1.7E-15
EC50	1.2E+10	2.1E-23	6.5E+42	16.	1.8E-33

Slope = 0.157 Std.Err. = 0.321

!!!Poor fit: p = 0.040 based on DF= 3.0 18.

1001LS : Dimethyl disulfide & Lettuce 21-day Survival lbs ai/A

Observed vs. Predicted Treatment Group Means

			<u> </u>				
Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	_
0.00 37.5 75.0 150. 300.	4.00 4.00 4.00 4.00 4.00	9.75 8.75 9.75 7.75 8.25	9.76 8.86 8.79 8.70 8.62	-0.00665 -0.115 0.963 -0.954 -0.366	100. 90.9 90.1 89.2 88.3	0.00 9.14 9.94 10.8 11.7	
600.	4.00	9.00	8.52	0.477	87.4	12.6	

!!!Warning: EC5 not bracketed by doses evaluated.

!!!Warning: EC25 not bracketed by doses evaluated.

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Onion 21-day Survival lbs ai/A File: 1001os Transform: NO TRANSFORMATION

Chi-square	nission Numbe			EPA MRID	Number 4
		ormality: actual a			
INTERVAL	<-1.5	-1.5 to <-0.5	-0.5 to 0.5	>0.5 to 1.5	>1.5
a A DECarro	1 600	5.808	9.168	E 000	1 600
DBSERVED	1.608 0	6	13	5.808 5	0
		goodness of fit to e (alpha = 0.01) =	test statistic =		
ata PASS 1	normality t	est. Continue anal	lysis.		
Dimethul d	ieulfide (	Onion 21-day Surv	ival lhe ai/A		
		ansform: NO TRANSI			
Shapiro Wil	lks test fo	r normality			
D = 32.50	00				
W = 0.94	46				
		(m. 04)			
Critical W	(P = 0.01)	(n = 24) = 0.916 (n = 24) = 0.884			
Data PASS 1	normality t	est at P=0.01 leve	el. Continue anal	lysis.	
Jimathul J.	iaulfiao c	Onion 21-days Gurri	ival lbc of/A		
File: 1001	os T	Onion 21-day Survi ransform: NO TRANS	SFORMATION		
Hartlev te	st for homo	geneity of variand	ce		
	H statisti	c (max Var/min Var			
Calculated			c) = 4.38		
Calculated Closest, co	onservative able H ==>	c (max Var/min Var , Table H statisti R (# groups) =	f) = 4.38 Lc = 184.0 (alg	pha = 0.01) reps-1) =	3
Calculated Closest, co	onservative able H ==>	c (max Var/min Va , Table H statisti	f) = 4.38 Lc = 184.0 (alg	pha = 0.01) reps-1) =	3
Calculated Closest, co Used for Ta Actual valu	onservative able H ==> ues ==>	c (max Var/min Var , Table H statisti R (# groups) =	f) = 4.38 Lc = 184.0 (alg = 6, df (# a = 6, df (# a	pha = 0.01) reps-1) = avg reps-1) =	3 3.00
Calculated Closest, co Used for Ta Actual valu	onservative able H ==> ues ==>	c (max Var/min Var , Table H statisti R (# groups) = R (# groups) =	f) = 4.38 Lc = 184.0 (alg = 6, df (# a = 6, df (# a	pha = 0.01) reps-1) = avg reps-1) =	3 3.00
Calculated Closest, co Used for Ta Actual valu Control	onservative able H ==> ues ==> homogeneity	c (max Var/min Var, Table H statisti  R (# groups) =  R (# groups) =	f) = 4.38 lc = 184.0 (algorithm of the control of t	pha = 0.01) reps-1) = avg reps-1) =	3 3.00
Calculated Closest, co Used for Ta Actual valu  Data PASS h NOTE: This	onservative  able H ==> ues ==>  homogeneity  test requi do not diff	c (max Var/min Var, Table H statisti  R (# groups) =  R (# groups) =  test. Continue ar  res equal replicat er greatly, the Ha	f) = 4.38 ic = 184.0 (algorithm (algorithm) = 6, df (# algorithm) = 6, df (# algorithm) algorithm (algorithm) artley test may are also artley test may are arranged to the arranged test may are arranged	pha = 0.01) reps-1) = avg reps-1) =	3 3.00
Calculated Closest, co Used for Ta Actual valu  Data PASS h NOTE: This	onservative  able H ==> ues ==>  homogeneity  test requi do not diff	c (max Var/min Var, Table H statisti  R (# groups) =  R (# groups) =  test. Continue ar	f) = 4.38 ic = 184.0 (algorithm (algorithm) = 6, df (# algorithm) = 6, df (# algorithm) algorithm (algorithm) artley test may are also artley test may are arranged to the arranged test may are arranged	pha = 0.01) reps-1) = avg reps-1) =	3 3.00
Calculated Closest, co Used for Ta Actual valu  Data PASS h NOTE: This	onservative  able H ==> ues ==>  homogeneity  test requi do not diff	c (max Var/min Var, Table H statisti  R (# groups) =  R (# groups) =  test. Continue ar  res equal replicat er greatly, the Ha	f) = 4.38 ic = 184.0 (algorithm (algorithm) = 6, df (# algorithm) = 6, df (# algorithm) algorithm (algorithm) artley test may are also artley test may are arranged to the arranged test may are arranged	pha = 0.01) reps-1) = avg reps-1) =	3 3.00
Calculated Closest, co Used for Ta Actual valu  Data PASS h  NOTE: This but o as ar	onservative  able H ==> ues ==>  homogeneity  test requi do not diff n approxima  isulfide &	c (max Var/min Var, Table H statisti  R (# groups) =  R (# groups) =  test. Continue ar  res equal replicat er greatly, the Ha te test (average of	f) = 4.38 ic = 184.0 (algorithm) = 6, df (# algorithm) algorithm algorithm algorithm artley test may algorithm are used).	pha = 0.01) reps-1) = avg reps-1) =	3 3.00
Calculated Closest, co Used for Ta Actual valu  Data PASS h NOTE: This but o	onservative  able H ==> ues ==>  homogeneity  test requi do not diff n approxima  isulfide &	c (max Var/min Var, Table H statisti  R (# groups) =  R (# groups) =  test. Continue ar  res equal replicat er greatly, the Hat te test (average of	f) = 4.38 ic = 184.0 (algorithm) = 6, df (# algorithm) algorithm algorithm algorithm artley test may algorithm are used).	pha = 0.01) reps-1) = avg reps-1) =	3 3.00

#### PMRA Submission Number N/A

### EPA MRID Number 47381001

Calculated B statistic = 2.61
Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Onion 21-day Survival lbs ai/A File: 100los Transform: NO TRANSFORMATION

#### ANOVA TABLE

SOURCE	DF	SS	MS	F
Between	5	5.500	1.100	0.609
Within (Error)	18	32.500	1.806	
Total	23	38.000		

Critical F value = 2.77 (0.05,5,18) Since F < Critical F FAIL TO REJECT Ho:All groups equal

Dimethyl disulfide & Onion 21-day Survival lbs ai/A File: 1001os Transform: NO TRANSFORMATION

GROUP         IDENTIFICATION         TRANSFORMED MEAN         MEAN CALCULATED IN ORIGINAL UNITS         T STAT         SIG           1         Neg Control         8.000         8.000         -0.263         <		DUNNETTS TEST - TA	Ho:Control <treatment< th=""></treatment<>			
2     37.5     8.250     8.250     -0.263       3     75     7.250     7.250     0.789       4     150     7.000     7.000     1.052       5     300     7.500     7.500     0.526	GROUP	IDENTIFICATION			T STAT	SIG
	3 4 5	37.5 75 150 300	8.250 7.250 7.000 7.500	8.250 7.250 7.000 7.500	0.789 1.052 0.526	

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Onion 21-day Survival lbs ai/A File: 1001os Transform: NO TRANSFORMATION

ים	UNNETTS TEST	- TA	BLE 2	OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATIO		NUM O	F	Minimum Sig Diff (IN ORIG. UNITS)	% of CONTROL	DIFFERENCE FROM CONTROL
1 2	Neg Con	trol	4 4		2.290	28.6	-0.250

PMRA Submission	on Number N/A			EPA N	MRID Number 47381001
3	75	4	2.290	28.6	0.750
4	150	4	2.290	28.6	1.000
5	300	4	2.290	28.6	0.500
6	600	4	2.290	28.6	1.000

Dimethyl disulfide & Onion 21-day Survival lbs ai/A File: 1001os Transform: NO TRANSFORMATION

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

GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	8.000	8.000	8.125
2	37.5	4	8.250	8.250	8.125
3	75	4	7.250	7.250	7.250
4	150	4	7.000	7.000	7.250
5	300	4	7.500	7.500	7.250
6	600	4	7.000	7.000	7.000

Dimethyl disulfide & Onion 21-day Survival lbs ai/A File: 1001os Transform: NO TRANSFORMATION

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2	OF	2
---------------	-----------	------------	--------	---------	----	---

IDENTIFICATION	ISOTONIZED	CALC.	SIG	TABLE	DEGREES OF
	MEAN	WILLIAMS	P=.05	WILLIAMS	FREEDOM
Neg Control 37.5 75 150 300 600	8.125 8.125 7.250 7.250 7.250 7.000	0.132 0.789 0.789 0.789 1.052		1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 1.344

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

#### Estimates of EC%

	<b></b>				
Parameter	Estimate	95% Bot	ınds	Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EÇ5	51.	0.013	2.1E+05	1.7	0.00025
EC10	2.7E+02	2.4	3.1E+04	0.99	0.0087
EC25	4.4E+03	2.9	6.7E+06	1.5	0.00066
EC50	9.8E+04	0.014	6.7E+11	3.3	1.5E-07

Slope = 0.501 Std.Err. = 0.712

Goodness of fit: p = 0.69 based on DF= 3.0 18.

10010S : Dimethyl disulfide & Onion 21-day Survival lbs ai/A

Observed vs. Predicted Treatment Group Means

PMRA Submi			<u>_</u>			EPA MRID	Number
Dose	#Reps.	Obs. Mean	Pred. Mean		Pred. %Control	_	
0.00	4.00	8.00	8.06	-0.0630	100.	0.00	
37.5				0.537			
	4.00	7 25	7.71	0.337	04 1		
75.0	4.00	7.45	7.59	-0.336 -0.425	94.1	3.91	
150.	4.00	7.00	7.43	-0.425	92.1	7.91	
300.	4.00	7.50	7.23	0.274 0.0141	89.6	10.4	
600.	4.00	7.00	6.99	0.0141	86.6	13.4	
!Warning:	EC25 not 1	oracketed b	y doses	evaluated.			
!Warning:	EC50 not 1	oracketed b	y doses	evaluated.			
		Sugarbeet 2 ansform: NC		irvival lbs ORMATION	ai/A		
				nd expected			
TERVAL	<-1.5	-1.5 to <	-0.5	-0.5 to 0.	5 >0.5 — —	to 1.5	>1.5
	1.608			9.168 10		5.808 7	1.608
		Sugarbeet 2 ansform: NC		urvival lbs ORMATION	ai/A		
		r normality					
= 57.75	0						
= 0.95	1						
ritical W		(n = 24) = (n = 24) =					<b></b> -
ata PASS n	ormality to	est at P=0.	01 leve	l. Continue	analysis.		
imethyl di ile: 1001s		Sugarbeet 2 ransform: N		urvival lbs FORMATION	ai/A		
artley tes	t for homo	geneity of	variance				
				) = 10.37 c = 184.0	(alpha =	0.01)	

Used for Table H ==> R (# groups) = 6, df (# reps-1) = 3 Actual values ==> R (# groups) = 6, df (# avg reps-1) = 3.00

### PMRA Submission Number N/A

EPA MRID Number 47381001

Data PASS homogeneity test. Continue analysis.

NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used).

Dimethyl disulfide & Sugarbeet 21-day Survival lbs ai/A File: 1001sb Transform: NO TRANSFORMATION

Bartletts test for homogeneity of variance

-----

Calculated B statistic = 3.86
Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05)

Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5

Data PASS homogeneity test at 0.01 level. Continue analysis.

NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above).

Dimethyl disulfide & Sugarbeet 21-day Survival lbs ai/A

File: 1001sb Transform: NO TRANSFORMATION

#### ANOVA TABLE \_\_\_\_\_\_

SOURCE	DF	ss	MS	F
Between	5	11.875	2.375	0.740
Within (Error)	18	57.750	3.208 '	
Total	23	69.625		

Critical F value = 2.77 (0.05,5,18) Since F < Critical F FAIL TO REJECT Ho: All groups equal

Dimethyl disulfide & Sugarbeet 21-day Survival lbs ai/A File: 1001sb Transform: NO TRANSFORMATION

	DUNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg control	7.000	7.000		
2	37.5	7.000	7.000	0.000	
3	75	6.750	6.750	0.197	
4	150	6.000	6.000	0.790	

PMRA Submission N	lumber N/A		EI	PA MRID Number 4	7381001
5	300	6.500	6,500	0.395	
6	600	5.000	5.000	1.579	
Dunnett table val	lue = 2.41	(1 Tailed Va	lue, P=0.05, df=1	.8,5)	

Dimethyl disulfide & Sugarbeet 21-day Survival lbs ai/A File: 1001sb Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)		DIFFERENCE FROM CONTROL
1	Neg control	4			
2	37.5	4	3.052	43.6	0.000
3	75	4	3.052	43.6	0.250
4	150	4	3.052	43.6	1.000
5	300	4	3.052	43.6	0.500
6	600	4	3.052	43.6	2.000

Dimethyl disulfide & Sugarbeet 21-day Survival lbs ai/A File: 1001sb Transform: NO TRANSFORMATION

	WILLIAMS TEST (Isotor	nic	regression model	.) TABLE 1 OF	7 2
GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg control	4	7.000	7.000	7.000
2	37.5	4	7.000	7.000	7.000
3	75	4	6.750	6.750	6.750
4	150	4	6.000	6.000	6.250
5	300	4	6.500	6.500	6.250
6	600	4	5.000	5.000	5.000

Dimethyl disulfide & Sugarbeet 21-day Survival lbs ai/A File: 1001sb Transform: NO TRANSFORMATION

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 C	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg control 37.5 75 150 300 600	7.000 7.000 6.750 6.250 6.250 5.000	0.000 0.197 0.592 0.592 1.579		1.73 1.82 1.85 1.86 1.87	k= 1, v=18 k= 2, v=18 k= 3, v=18 k= 4, v=18 k= 5, v=18

s = 1.791

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

#### PMRA Submission Number N/A

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Estimates of EC%

Estimate	95% Bounds		Std.Err.	Lower Bound	
	Lower	Upper		/Estimate	
1.3E+02	3.0	5.7E+03	0.79	0.023	
2.3E+02	18.	2.9E+03	0.53	0.078	
5.8E+02	1.8E+02	1.9E+03	0.25	0.31	
1.6E+03	1.1E+02	2.4E+04	0.56	0.069	
	1.3E+02 2.3E+02 5.8E+02	Lower 1.3E+02 3.0 2.3E+02 18. 5.8E+02 1.8E+02	Lower Upper 1.3E+02 3.0 5.7E+03 2.3E+02 18. 2.9E+03 5.8E+02 1.8E+02 1.9E+03	Lower Upper 1.3E+02 3.0 5.7E+03 0.79 2.3E+02 18. 2.9E+03 0.53 5.8E+02 1.8E+02 1.9E+03 0.25	Lower Upper /Estimate 1.3E+02 3.0 5.7E+03 0.79 0.023 2.3E+02 18. 2.9E+03 0.53 0.078 5.8E+02 1.8E+02 1.9E+03 0.25 0.31

Slope = 1.51 Std.Err. = 1.73

Goodness of fit: p = 0.87 based on DF= 3.0 18.

1001SBS: Dimethyl disulfide & Sugarbeet 21-day Survival lbs ai/A

Observed vs. Predicted Treatment Group Means

Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change	
0.00	4.00	7.00	6.93	0.0681	100.	0.00	
37.5	4.00	7.00	6.88	0.116	99.3	0.686	
75.0	4.00	6.75	6.78	-0.0282	97.8	2.22	
150.	4.00	6.00	6.52	-0.518	94.0	5.98	
300.	4.00	6.50	6.00	0.504	86.5	13.5	
600.	4.00	5.00	5.14	-0.142	74.2	25.8	

!!!Warning: EC50 not bracketed by doses evaluated.

Dimethyl disulfide & Soybean 21-day Survival lbs ai/A File: 1001ss Transform: NO TRANSFORMATION

Chi-square test for normality: actual and expected frequencies

EXPECTED 1.608 5.808 9.168 5.808 1.608 OBSERVED 0 7 12 5 0

Calculated Chi-Square goodness of fit test statistic = 4.4479
Table Chi-Square value (alpha = 0.01) = 13.277

Data PASS normality test. Continue analysis.

Dimethyl disulfide & Soybean 21-day Survival lbs ai/A File: 1001ss Transform: NO TRANSFORMATION

FILE: 100185 ITAMSFORMATIO

Shapiro Wilks test for normality

D = 29.000

W = 0.950

Critical W (P = 0.05) (n = 24) = 0.916 Critical W (P = 0.01) (n = 24) = 0.884

### PMRA Submission Number N/A

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Total

EPA MRID Number 47381001

Data PASS normality test at P=0.01 level. Continue analysis. Dimethyl disulfide & Soybean 21-day Survival lbs ai/A Transform: NO TRANSFORMATION File: 1001ss Hartley test for homogeneity of variance Calculated H statistic (max Var/min Var) = 19.67 Closest, conservative, Table H statistic = 184.0 (alpha = 0.01) Used for Table H ==> R (# groups) = 6, df (# reps-1) = 3 Actual values ==> R (# groups) = 6, df (# avg reps-1) = 3.00 Data PASS homogeneity test. Continue analysis. NOTE: This test requires equal replicate sizes. If they are unequal but do not differ greatly, the Hartley test may still be used as an approximate test (average df are used). Dimethyl disulfide & Soybean 21-day Survival lbs ai/A File: 1001ss Transform: NO TRANSFORMATION Bartletts test for homogeneity of variance Calculated B statistic = 8.75
Table Chi-square value = 15.09 (alpha = 0.01)
Table Chi-square value = 11.07 (alpha = 0.05) Average df used in calculation ==> df (avg n - 1) = 3.00 Used for Chi-square table value ==> df (#groups-1) = 5 Data PASS homogeneity test at 0.01 level. Continue analysis. NOTE: If groups have unequal replicate sizes the average replicate size is used to calculate the B statistic (see above). Dimethyl disulfide & Soybean 21-day Survival 1bs ai/A File: 1001ss Transform: NO TRANSFORMATION ANOVA TABLE \_\_\_\_\_\_ DF SS MS \_\_\_\_\_\_\_ 5.767 Between 5 28.833 Within (Error) 18 29.000 1.611

57.833

#### PMRA Submission Number N/A

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Critical F value = 2.77 (0.05,5,18)
Since F > Critical F REJECT Ho:All groups equal

Dimethyl disulfide & Soybean 21-day Survival lbs ai/A File: 1001ss Transform: NO TRANSFORMATION

	DUNNETTS TEST - TA	BLE 1 OF 2	Ho:Control <tr< th=""><th>eatment</th><th></th></tr<>	eatment	
GROUP	IDENTIFICATION	TRANSFORMED MEAN	MEAN CALCULATED IN ORIGINAL UNITS	T STAT	SIG
1	Neg Control	8.750	8.750		
2	37.5	9.000	9.000	-0.279	
3	75	8.000	8.000	0.836	
4	150	8.750	8.750	0.000	
5	300	8.250	8.250	0.557	
6	600	5.750	5.750	3.343	*

Dunnett table value = 2.41 (1 Tailed Value, P=0.05, df=18,5)

Dimethyl disulfide & Soybean 21-day Survival lbs ai/A File: 1001ss Transform: NO TRANSFORMATION

	DUNNETTS TEST -	TABLE 2 OF	2 Ho:	Control <t< th=""><th>reatment</th></t<>	reatment
GROUP	IDENTIFICATION	NUM OF REPS	Minimum Sig Diff (IN ORIG. UNITS)		DIFFERENCE FROM CONTROL
1	Neg Control	4			
2	37.5	4	2.163	24.7	-0.250
3	75	4	2.163	24.7	0.750
4	150	4	2.163	24.7	0.000
5	300	4	2.163	24.7	0.500
6	600	4	2.163	24.7	3.000

Dimethyl disulfide & Soybean 21-day Survival lbs ai/A File: 1001ss Transform: NO TRANSFORMATION

	WILLIAMS TEST (Isoto	nic	regression model	l) TABLE 1 O	F 2
GROUP	IDENTIFICATION	N	ORIGINAL MEAN	TRANSFORMED MEAN	ISOTONIZED MEAN
1	Neg Control	4	8.750	8.750	8.875
2	37.5	4	9.000	9.000	8.875
3	75	4	8.000	8.000	8.375
4	150	4	8.750	8.750	8.375
5	300	4	8.250	8.250	8.250
6	600	4	5.750	5.750	5.750

Dimethyl disulfide & Soybean 21-day Survival lbs ai/A File: 1001ss Transform: NO TRANSFORMATION

### PMRA Submission Number N/A

### EPA MRID Number 47381001

WILLIAMS TEST	(Isotonic	regression	model)	TABLE 2 O	F 2
IDENTIFICATION	ISOTONIZED MEAN	CALC. WILLIAMS	SIG P=.05	TABLE WILLIAMS	DEGREES OF FREEDOM
Neg Control 37.5 75 150 300 600	8.875 8.875 8.375 8.375 8.250 5.750	0.139 0.418 0.418 0.557 3.343	*	1.73 1.82 1.85 1.86 1.87	k= 1, V=18 k= 2, V=18 k= 3, V=18 k= 4, V=18 k= 5, V=18

s = 1.269

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

#### Estimates of EC%

Parameter	Estimate	95% Bot	ınds	Std.Err.	Lower Bound
		Lower	Upper		/Estimate
EC5	3.1E+02	1.1E+02	8.7E+02	0.21	0.36
EC10	3.8E+02	1.8E+02	7.9E+02	0.15	0.48
EC25	5.3E+02	3.9E+02	7.0E+02	0.061	0.75
EC50	7.6E+02	5.0E+02	1.1E+03	0.087	0.66

Slope = 4.30 Std.Err. = 3.18

Goodness of fit: p = 0.77 based on DF= 3.0 18.

1001SS : Dimethyl disulfide & Soybean 21-day Survival lbs ai/A

Observed vs. Predicted Treatment Group Means

Dose	#Reps.	Obs. Mean	Pred. Mean	Obs. -Pred.	Pred. %Control	%Change
0.00 37.5 75.0 150.	4.00 4.00 4.00 4.00	8.75 9.00 8.00 8.75	8.63 8.63 8.63 8.61	0.125 0.375 -0.625 0.136	100. 100. 100. 99.9	0.00 1.00e-06 0.000792 0.126
300. 600.	4.00 4.00 4.00	8.25 5.75	8.26 5.75	-0.0112 0.00114	95.8 66.7	4.22

<sup>!!!</sup>Warning: EC50 not bracketed by doses evaluated.

PMRA Submission Number N/A

EPA MRID Number 47381001