

# ToxGenie Two-Group Comparison Report

Report Date: 2026-06-29 13:39:23

<b>Study Title:</b> In Vitro Mammalian Chromosomal Aberration Test	
<b>Analysis Method:</b> Student's t-test for % Tail DNA	
<b>Study No.:</b> Test-135	<b>Test Material:</b> Test Chemicals
<b>Exposure Time:</b> 24-hours	<b>Test Medium:</b> Not Applicable

**Table 1. Raw Data Summary for % Tail DNA**

(Unit: mg/kg)

Dose	1	2	3	4	5
Negative Control	4.1000	6.2000	5.5000	3.8000	4.9000
Positive Control	54.1000	58.2000	55.9000	52.8000	56.5000

## Step 1. Outlier Detection (Data QA/QC) Analysis

No statistical outliers were detected in the raw data based on Tukey's fences.

*Tukey's fences: Values outside  $Q1 - 1.5 \cdot IQR$  or  $Q3 + 1.5 \cdot IQR$  (IQR method).*

## Step 2. Descriptive Statistics

**Table 2. Descriptive Statistics for % Tail DNA**

(Unit: mg/kg)

Dose	N	Mean	SD	SE	CV (%)	Median
Negative Control	5	4.90000	0.98742	0.44159	20.15145	4.90000
Positive Control	5	55.50000	2.10357	0.94074	3.79021	55.90000

*Abbreviations & Explanations:*

- N: Number of replicates.
- Mean: Average response value.
- SD: Standard Deviation of the mean.
- SE: Standard Error of the mean.
- CV (%): Coefficient of Variation.
- Median: 50th percentile of observed values.

**Table 3. Shapiro-Wilk's test for normality on the % Tail DNA**

Dose	Statistic	P-value	Result
Negative Control	0.9581	0.7948	Normal Distribution
Positive Control	0.9816	0.9432	Normal Distribution

*$P > 0.05$  indicates Normal Distribution.*

*Final decision: The % Tail DNA data follows a normal distribution.*

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**Table 4. Levene's test for homogeneity on the % Tail DNA**

Statistic	P-value	Result
1.74292	0.22329	Equal Variances

*P*>0.05 indicates Equal Variances.

Final decision: The % Tail DNA data shows equal variances.

## Statistical Analysis Procedure & Decision Tree

The statistical analysis for '% Tail DNA' was conducted based on the following decision steps:

### 1. Assumption Verification (Normality & Homogeneity):

- Since the raw data satisfied both the Shapiro-Wilk test (Normality) and Levene's test (Homogeneity of Variance), no data transformation was required.
- Consequently, the analysis proceeded directly to Parametric Tests.

### 2. Data Transformation Strategy:

- Not applicable (The raw data already satisfied statistical assumptions).

### 3. Final Selected Test Method:

- Because the data satisfied the required statistical assumptions (normality and homogeneity), the Student's t-test (Parametric) was selected as the final analytical method.
- This method was chosen for its high statistical power to detect differences between two groups, ensuring a reliable comparison between the control and treatment groups.

**Table 5. Student's t-test on the % Tail DNA data**

Control vs Dose	P-value	Result
Negative Control vs Positive Control	0.0000	Difference from Control

Note: *p*<0.05 indicates a statistically significant difference from the control group.

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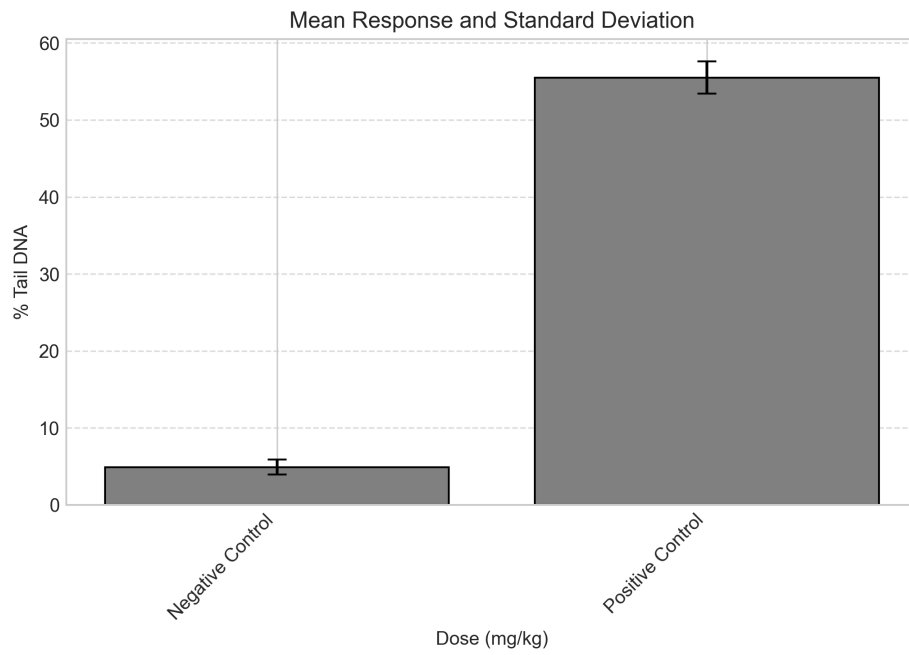


Figure 1. Dose-response trend for % Tail DNA. Bars represent the mean and Standard Deviation (SD).

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