CGA15324/Lambda-cyhalothrin

CGA15324/Lambda-cyhalothrin EC (A13735F) - *Salmonella Typhimurium* and *Escherichia Coli* Reverse Mutation Assay

Final Report

TEST GUIDELINE(S): OECD 471 (2020)

AUTHOR(S): Dr. Steffi Chang

COMPLETION DATE: 07 July 2021

PERFORMING LABORATORY: ICCR-Roßdorf GmbH
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SPONSOR(S): Syngenta Ltd.
Jealott's Hill International Research Centre
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This study performed in the test facility of ICCR-Roßdorf GmbH, In den Leppsteinswiesen 19, 64380 Rossdorf, Germany was conducted in compliance with Good Laboratory Practice Regulations:

Chemikaliengesetz (Chemicals Act) of the Federal Republic of Germany, "Anhang 1" (Annex 1), in its currently valid version

OECD Principles of Good Laboratory Practice, (as revised in 1997), ENV/MC/CHEM(98)17

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These procedures are compatible with Good Laboratory Practice regulations specified by regulatory authorities throughout the European Community, the United States (EPA and FDA), and Japan (MHW, MAFF, and METI), and other countries that are signatories to the OECD Mutual Acceptance of Data Agreement.

There were no circumstances that may have affected the quality or integrity of the study.

Dr. Steffi Chang
Study Director Bacterial Systems

.....
Date: 07 July 2021

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QUALITY ASSURANCE STATEMENT

ICCR Study Number: 2165900
Test substance: CGA15324/lambda-cyhalothrin EC (A13735F)
Study director: Dr. Steffi Chang
Study Title: CGA15324/Lambda-cyhalothrin EC (A13735F) - *Salmonella Typhimurium* and *Escherichia Coli* Reverse Mutation Assay

Study based activities at the Test Facility ICCR-Roßdorf GmbH were audited and inspected. The details of these audits and inspections are given below.

Type of Inspection	Date(s) of Inspection	Date Reporting to Study Director, Test Facility Management
Study Plan Verification	30 April 2021	30 April 2021
Study – based Test Item preparation	05 May 2021	05 May 2021
Report Audit	24 June 2021	25 June 2021

General facilities and activities where this study was conducted were inspected on an annual basis and results are reported to the relevant responsible person and Management.

The statement is to confirm, that this report reflects the raw data.

S. Ebert

Sabine Ebert

Quality Assurance Auditor
ICCR-Roßdorf GmbH

07 July 2021

Date

PROJECT STAFF SIGNATURE

Study Director Dr. Steffi Chang

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Date: 07 July 2021

GENERAL INFORMATION

Contributors
The following contributed to this report in the capacities indicated:

Name	Title
Dr. Steffi Chang	Study Director
Dr. Markus Schulz	Test Facility Management
Frauke Hermann	Head of Quality Assurance Unit
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Study Dates
Study initiation date: 03 May 2021
Experimental start date: 05 May 2021
Experimental completion date: 25 May 2021

Deviations from the Guidelines
None

Retention of Samples
None

Performing Laboratory Test Substance Reference Number
S 2156011

Other
ICCR-Roßdorf GmbH will archive:
Records and documentation relating to this study will be maintained in the archives of ICCR-Roßdorf GmbH for a period of 4 years from the date on which the Study Director signs the final report. This will include electronic and paper raw data, and report that support the reconstruction of the study.

At termination of the aforementioned period, the records and documentation will be transferred to the GLP compliant Archive of Rhenus Archiv Services GmbH, Frankfurt am Main for further archiving up to a total archiving period of 15 years.

A sample of the test substance will not be archived.

ICCR Roßdorf GmbH will retain in its archive a copy of the study plan and final report, and any amendments indefinitely.

Deviations from the study plan
There were no deviations (unplanned changes) from the study plan.

Distribution of the report

Sponsor	2 × electronic copy (1 × pdf-file, 1 × Word-file)
Study Director	1 × (original)

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1.0 EXECUTIVE SUMMARY

1.1 Study Design

This study was performed to investigate the potential of CGA15324/lambda-cyhalothrin EC (A13735F) to induce gene mutations in the plate incorporation test (Experiment I) and the pre-incubation test (Experiment II) using the *Salmonella typhimurium (S. typhimurium)* strains TA1535, TA1537, TA98, and TA100, and the *Escherichia coli (E. coli)* strains WP2 *uvrA* (pKM101) and WP2 (pKM101).

1.2 Results

The plates incubated with the test substance showed normal background growth up to the maximal concentration of 5000 µg/plate with and without S9 mix in all strains used.

Cytotoxic effects, evident as a reduction in the number of revertants (below the indication factor of 0.5), occurred in Experiment I in strain WP2 (pKM101) in the absence of S9 mix and in strains TA1535 and TA100 in the presence of S9 mix, and in Experiment II in strain TA1537 in the absence of S9 mix and in strains TA98, TA100 and WP2 (pKM101) in the presence and absence of S9 mix.

No relevant increase in revertant colony numbers of any of the six tester strains was observed following treatment with CGA15324/lambda-cyhalothrin EC (A13735F) at any concentration, neither in the presence nor absence of metabolic activation (S9 mix). There was also no observed tendency of higher mutation rates with increasing concentrations in the range below the generally acknowledged border of biological relevance.

Appropriate reference mutagens were used as positive controls, which showed a distinct increase of induced revertant colonies consistent with the laboratory's historical control data demonstrated the sensitivity of the test system and the efficacy of the S9 mix. Each batch of S9 was also tested with 2 pro-mutagens, benzo(a)pyrene and 2-aminoanthracene.

1.3 Conclusion

In conclusion, it can be stated that during the described mutagenicity tests and under the experimental conditions reported, CGA15324/lambda-cyhalothrin EC (A13735F) did not induce gene mutations by base pair changes or frameshifts in the genome of the strains used.

Therefore, CGA15324/lambda-cyhalothrin EC (A13735F) is considered to be non-mutagenic in the *Salmonella typhimurium* and *Escherichia coli* reverse mutation assay.

2.0 INTRODUCTION

2.1 Purpose

These experiments were performed to assess the potential of the test substance to induce gene mutations by means of the *S. typhimurium* and *E. coli* reverse mutation assay. Experiment I was performed as a plate incorporation assay. Since a negative result was obtained in this experiment, Experiment II was performed as a pre-incubation assay.

The most widely used assays for detecting gene mutations are those using bacteria (1). They are relatively simple and rapid to perform, and give reliable data on the ability of an agent to interact with DNA and produce mutations.

Reverse mutation assays determine the frequency with which an agent reverses or suppresses the effect of the forward mutation. The genetic target presented to an agent is therefore small, specific and selective. Several bacterial strains, or a single strain with multiple markers are necessary to assure reliable detection of mutagens that may be specific to one tester strain or locus. The reversion of bacteria from growth-dependence on a particular amino acid to growth in the absence of that amino acid (reversion from auxotrophy to prototrophy) is the most widely used marker.

The S. *typhimurium* histidine (his) and the *E. coli* tryptophan (trp) reversion system measures $his^- \rightarrow his^+$ and $trp^- \rightarrow trp^+$ reversions, respectively. The *S. typhimurium* and *E. coli* strains are constructed to differentiate between base pair (TA1535, TA100, WP2 *uvrA* (pKM101), and WP2 (pKM101)) and frameshift (TA1537, TA98) mutations.

According to the direct plate incorporation and pre-incubation method the bacteria are exposed to the test substance with and without metabolic activation and plated on selective medium. After a suitable period of incubation, revertant colonies are counted.

To establish a concentration response effect at least six concentrations with adequately spaced intervals were tested. The maximum concentration was 5000 µg/plate.

To validate the test, reference mutagens were tested in parallel to the test substance.

2.2 Test Guideline(s)

This study followed the procedures indicated by the following internationally accepted guideline and recommendations:

"Ninth Addendum to OECD Guidelines for Testing of Chemicals", Section 4, No. 471: "Bacterial Reverse Mutation Test", corrected June 26, 2020

3.0 MATERIALS AND METHODS

3.1 Test Substance

Information as provided by the Sponsor.

Identification:	CGA15324/lambda-cyhalothrin EC (A13735F)
Batch:	SMU1BL001
Content of Profenofos:	27.3% w/w corresponding to 300 g/L
Content of Lambda-Cyhalothrin:	1.37% w/w corresponding to 15.1 g/L
Appearance:	brown liquid
Recertification Date:	28 February 2023
Storage Conditions:	At room temperature
Stability in Solvent:	Not indicated by the Sponsor

The test substance concentrations were not adjusted for the content of CGA15324 or lambda-cyhalothrin.

On the day of the experiment (immediately before use), the test substance was dissolved in dimethylsulfoxide (DMSO, purity > 99%). The solvent was chosen as the most suitable solvent compared to water and ethanol, according to its solubilisation properties and its relative non-toxicity to the bacteria (2).

All formulations were prepared freshly before treatment and used within two hours of preparation. The formulation was assumed to be stable for this period unless specified otherwise by the Sponsor.

3.2 Controls

3.2.1 Negative controls

Concurrent untreated and solvent controls were performed.

3.2.2 Positive control substances

Without metabolic activation

Strains:	TA1535, TA100
Name:	Sodium azide, (NaN_3)
Supplier:	SERVA, 69042 Heidelberg, Germany
Batch No.:	STBJ7813
Purity:	≥ 99%
Dissolved in:	Deionised water
Concentration:	10 µg/plate

Strains:	TA1537, TA98
Name:	4-nitro-o-phenylene-diamine, (4-NOPD)
Supplier:	Sigma-Aldrich, 82024 Taufkirchen, Germany
Batch No.:	MKBM 5257V
Purity:	≥ 98%
Dissolved in:	DMSO (purity >99 %, Fisher Leics LE11 5RG, United Kingdom)
Concentration:	10 µg/plate in strain TA 98, 50 µg/plate in strain TA 1537

Strains:	WP2 *uvrA* (pKM101), WP2 (pKM101)
Name:	Methyl methane sulfonate, (MMS)
Supplier:	Sigma-Aldrich, 82024 Taufkirchen, Germany
Batch No.:	MKCL 6261
Purity:	≥ 99%
Dissolved in:	Deionised water
Concentration:	2.0 µL/plate

With metabolic activation

Strains:	TA1535, TA1537, TA98, TA100, WP2 *uvrA* (pKM101), WP2 (pKM 101)
Name:	2-aminoanthracene, (2-AA)
Supplier:	Sigma-Aldrich, 82024 Taufkirchen, Germany
Batch No.:	STBG 0630V
Purity:	≥ 96%
Dissolved in:	DMSO (purity > 99 %, Fisher Leics LE11 5RG, United Kingdom)
Concentration:	2.5 µg/plate (TA1535, TA1537, TA98, TA100), 10 µg/plate (WP2 *uvrA* (pKM101), WP2 (pKM101))

The stability of the positive control substances in solution is unknown but a mutagenic response in the expected range is sufficient evidence of biological activity.

3.3 Experimental Design

3.3.1 Characterisation of the *Salmonella typhimurium* and *E. coli* strains

The histidine dependent strains are derived from *S. typhimurium* strain LT2 through mutations in the histidine locus. Additionally, due to the "deep rough" (*rfa*$^-$) mutation they possess a faulty lipopolysaccharide envelope which enables substances to penetrate the cell wall more easily. A further mutation causes a reduction in the activity of an excision repair system. The last alteration includes mutational processes in the nitrate reductase and biotin genes produced in a UV-sensitive area of the gene named *uvrB*$^-$. In the strains TA98 and TA100 the R-factor plasmid pKM101 carries the ampicillin resistance marker (3).

Strain WP2 (4) and its derivatives all carry the same defect in one of the genes for tryptophan biosynthesis. Tryptophan-independent (*Trp*$^+$) mutants (revertants) can arise either by a base change at the site of the original alteration or by a base change elsewhere in the chromosome so that the original defect is suppressed. This second possibility can occur in several different ways so that the system seems capable of detecting all types of mutagen which substitute one base for another. Additionally, the *uvrA* derivative is deficient in the DNA repair process (excisable repair damage). Such a repair-deficient strain may be more readily mutated by agents. The *E. coli* strains WP2 *uvrA* (pKM101) and WP2 (pKM101) are constructed by introduction of the R-factor plasmid pKM101.

When summarized, the mutations of the *S. typhimurium* and *E. coli* strains used in this study can be described as follows:

Strains	**Genotype**	**Type of mutations indicated**
Salmonella typhimurium		
TA1537	*his* C 3076; *rfa*$^-$; *uvrB*$^-$	frame shift mutations
TA98	*his* D 3052; *rfa*$^-$; *uvrB*$^-$; R-factor	" "
TA1535	*his* G 46; *rfa*$^-$; *uvrB*$^-$	base-pair substitutions
TA100	*his* G 46; *rfa*$^-$; *uvrB*$^-$; R-factor	" "
Escherichia coli		
WP2 *uvrA* (pKM101)	*trp* E 56 *uvrA*$^-$; R-factor	base-pair substitutions and others
WP2 (pKM101)	*trp* E 56; R-factor	" "

Regular checking of the properties of the *S.typhimurium* and *E. coli* strains regarding the membrane permeability and ampicillin resistance; UV sensitivity, and amino acid requirement as well as normal spontaneous mutation rates is performed by ICCR-Roßdorf GmbH according to Ames *et al.* (5), Maron and Ames (3), and Mortelmans and Riccio (7). In this way it is ensured that the experimental conditions set down by Ames are fulfilled.

The bacterial strains TA1535, TA1537, TA98, TA100, WP2 *uvrA* (pKM101), and WP2 (pKM101) were obtained from Trinova Biochem GmbH (35394 Gießen, Germany).

3.3.2 Storage

The strain cultures were stored as stock cultures in ampoules with nutrient broth + 5 % DMSO (Fisher Leics, LE11 5RG, United Kingdom) in liquid nitrogen.

3.3.3 Precultures

The thawed bacterial suspension was transferred into 250 mL Erlenmeyer flasks containing nutrient medium (50 mL). A solution of ampicillin (50 µL, 25 µg/mL) was added to the strains TA98, TA100, WP2 *uvrA* (pKM101), and WP2 (pKM101). This nutrient medium contains per liter:

8 g Nutrient Broth (MERCK, 64293 Darmstadt, Germany)
5 g NaCl (MERCK, 64293 Darmstadt, Germany)

The bacterial cultures were incubated in a shaking water bath for 4 hours at 37 °C. The optical density of the bacteria was determined by absorption measurement and the obtained values indicated that the bacteria were harvested at the late exponential or early stationary phase (10^8-10^9 cells/mL).

3.3.4 Selective agar

Plates with selective agar (without Histidine/Tryptophan) were used.

3.3.5 Overlay agar

The overlay agar contained per litre:

for *Salmonella* strains:	for *Escherichia coli* strains:
7.0 g Agar Agar*	7.0 g Agar Agar*
6.0 g NaCl*	6.0 g NaCl*
10.5 mg L-Histidine×HCl×H_2O*	10.2 mg Tryptophan*
12.2 mg Biotin*	

* (MERCK, 64293 Darmstadt, Germany)

Sterilisations were performed at 121 °C in an autoclave.

3.4 Mammalian Microsomal Fraction S9 Mix

The bacteria used in this assay do not possess the enzyme systems which, in mammals, are known to convert promutagens into active DNA damaging metabolites. In order to overcome this major drawback an exogenous metabolic system is added in the form of mammalian microsome enzyme activation mixture.

3.4.1 S9 (Preparation by ICCR-Roßdorf GmbH)

Phenobarbital/ß-naphthoflavone induced rat liver S9 was used as the metabolic activation system. The S9 was prepared from male Wistar rats (RjHan:WI; weight approx. 220 – 320 g,

Janvier Labs, 53941 Saint-Berthevin Cedex, France) induced by peroral administration of 80 mg/kg b.w. phenobarbital (Sigma-Aldrich Chemie GmbH, 82024 Taufkirchen, Germany) and by peroral administrations of ß-naphthoflavone (Acros Organics, 2440 Geel, Belgium) each, on three consecutive days. The livers were prepared 24 hours after the last treatment. The S9 fractions were produced by dilution of the liver homogenate with a KCl solution (1+3 parts) followed by centrifugation at 9000 g. Aliquots of the supernatant were frozen and stored in ampoules at –80 °C. Small numbers of the ampoules can be kept at –20 °C for up to one week. Each batch of S9 mix is routinely tested with 2-aminoanthracene as well as benzo[a]pyrene (Appendix 3).

The protein concentration in the S9 preparation was 31.2 mg/mL (lot no. 291020D, recertified 10 May 2021) in both experiments.

3.4.2 S9 mix

Before the experiment an appropriate quantity of S9 supernatant was thawed and mixed with S9 cofactor solution. The amount of S9 supernatant was 10% v/v in the S9 mix. Cofactors were added to the S9 mix to reach the following concentrations in the S9 mix:

8	mM	$MgCl_2$
33	mM	KCl
5	mM	Glucose-6-phosphate
4	mM	NADP

in 100 mM sodium-ortho-phosphate-buffer, pH 7.4.
During the experiment the S9 mix was stored in an ice bath. The S9 mix preparation was performed according to Ames *et al.*(5).

3.5 Pre-Experiment for Cytotoxicity

To evaluate the cytotoxicity of the test substance a pre-experiment was performed with all strains. Eight concentrations were tested for cytotoxicity and mutation induction each with three replicate plates. The experimental conditions in this pre-experiment are described in section 3.7 (plate incorporation test).

Cytotoxicity of the test substance results in a reduction in the number of spontaneous revertants (below a factor of 0.5) or a clearing of the bacterial background lawn.

The pre-experiment is reported as the Main Experiment I since the criteria mentioned in Section 3.8.2 Acceptability of the Assay were met.

3.6 Concentration Selection

In the pre-experiment the concentration range of the test substance was 3 - 5000 µg/plate. The pre-experiment is reported as Experiment I. Since minor cytotoxic effects were

observed in Experiment I, six concentrations were tested in all strains without S9 mix and seven concentrations in all strains with S9 mix. 5000 µg/plate was chosen as the maximal concentration in Experiment II.

The concentration range included two logarithmic decades. The following concentrations were tested in Experiment II:
All strains without S9 mix: 33; 100; 333; 1000; 2500; and 5000 µg/plate
All strains with S9 mix: 10; 33; 100; 333; 1000; 2500; and 5000 µg/plate

3.7 Experimental Performance

For each strain and concentration including the controls, three plates were used.

The following materials were mixed in a test tube and poured onto the selective agar plates:
100 µL Test solution at each concentration, solvent (negative control) or reference mutagen solution (positive control),
500 µL S9 mix (for test with metabolic activation) or S9 mix substitution buffer* (for test without metabolic activation),
100 µL Bacteria suspension (cf. test system, pre-culture of the strains; OD = 1.0 - 1.2; wavelength = 500 nm; approx. $8x10^8$ cells/mL),
2000 µL Overlay agar

For the pre-incubation method test solution (100 µL) (solvent or reference mutagen solution (positive control)), S9 mix / S9 mix substitution buffer* (500 µL) and bacteria suspension (100 µL) were mixed in a test tube and incubated at 37 C ± 1.5° C for 60 minutes. After pre-incubation overlay agar (2.0 mL, 45 °C) was added to each tube. The mixture was poured on selective agar plates.

After solidification the plates were incubated upside down for 72 hours at 37 C ± 1.5° C in the dark, plates were then stored at 4 °C until counted (6).

In parallel to each test a sterile control of the test substance was performed and documented in the raw data. Therefore, stock solution (100 µL) and S9 mix / S9 mix substitution buffer* (500 µL) were mixed with overlay agar (2.0 mL) and poured on minimal agar plates.

* Substitution buffer: 7 parts of the 100 mM sodium-ortho-phosphate-buffer pH 7.4 with 3 parts of KCl solution 0.15 M

3.8 Data Evaluation

3.8.1 Data recording

The colonies were counted using a Petri Viewer with the software program Ames Study Manager (see section 3.9, Major computerized systems). The evaluation unit was connected to a PC with printer to print out the individual values, the means from the plates for each

concentration together with standard deviations and enhancement factors as compared to the spontaneous reversion rates (see tables of results). The print outs are kept with the raw data. Due to precipitation of the test item some test groups were scored manually (as indicated on data tables).

3.8.2 Acceptability of the assay

The *Salmonella typhimurium* and *Escherichia coli* reverse mutation assay is considered acceptable if it meets the following criteria:

- regular background growth in the negative and solvent control
- the spontaneous reversion rates in the negative and solvent control are in the range of the historical data
- the positive control substances should produce an increase above the threshold of twofold (strains TA 98, TA 100, WP2 uvrA (pKM101, and WP2 (pKM101))) or threefold (strains TA 1535 and TA 1537) the revertant colony count of the corresponding solvent control;
- a minimum of five analysable concentrations should be present with at least four showing no signs of toxic effects, evident as a reduction in the number of revertants below the indication factor of 0.5.

3.8.3 Evaluation of results

A test substance is considered as a mutagen if a biologically relevant increase in the number of revertants of twofold or above (strains TA 98, TA 100, WP2 uvrA (pKM101), and WP2 (pKM101)) or of threefold or above (strains TA 1535 and TA 1537) the spontaneous mutation rate of the corresponding solvent control is observed.

A concentration dependent increase is considered biologically relevant if the threshold is reached or exceeded at more than one concentration (6).

An increase of revertant colonies equal or above the threshold at only one concentration is judged as biologically relevant if reproduced in an independent second experiment.

A concentration dependent increase in the number of revertant colonies below the threshold is regarded as an indication of a mutagenic potential if reproduced in an independent second experiment. However, whenever the colony counts remain within the historical range of negative and solvent controls, such an increase is not considered biologically relevant.

3.8.4 Biometry

According to the OECD guideline 471, a statistical analysis of the data is not mandatory.

3.9 Major Computerized System

Petri Viewer Sorcerer Colony Counter 3.0 (Instem, Suffolk IP33 3TA, UK) with the software program Ames Study Manager (v1.24) and Ames Archive Manager (v1.01).

4.0 RESULTS AND DISCUSSION

The test substance, CGA15324/lambda-cyhalothrin EC (A13735F), was assessed for its potential to induce gene mutations in the plate incorporation test (Experiment I) and the pre-incubation test (Experiment II) using *S. typhimurium* strains TA1535, TA1537, TA98, and TA100, and the *E. coli* strains WP2 (pKM101) and WP2 *uvrA* (pKM101).

In the pre-experiment the concentration range of the test substance was 3 - 5000 µg/plate. The pre-experiment is reported as Experiment I. Since minor cytotoxic effects were observed in Experiment I, six concentrations were tested in all strains without S9 mix and seven concentrations in all strains with S9 mix. 5000 µg/plate was chosen as the maximal concentration in Experiment II. This is the maximum concentration recommended in the OECD test guideline.

The assay was performed with and without liver microsomal activation. Each concentration, including the controls, was tested in triplicate. The concentration range included two logarithmic decades. The test substance was tested at the following concentrations:
Pre-Experiment/Experiment I: 3; 10; 33; 100; 333; 1000; 2500; and 5000 µg/plate
Experiment II:
All strains without S9 mix: 33; 100; 333; 1000; 2500; and 5000 µg/plate
All strains with S9 mix: 10; 33; 100; 333; 1000; 2500; and 5000 µg/plate

The test substance precipitated in the overlay agar in the test tubes from 1000 to 5000 µg/plate. Precipitation of the test item in the overlay agar on the incubated agar plates was observed at 5000 µg/plate in the absence of S9 mix and from 2500 to 5000 µg/plate in the presence of S9 mix. The undissolved particles had no influence on the data recording.

The plates incubated with the test substance showed normal background growth up to the maximal dose of 5000 µg/plate with and without S9 mix in all strains used.

Cytotoxic effects, evident as a reduction in the number of revertants (below the induction factor of 0.5), were observed at the following concentrations (µg/plate):

Strain	Experiment I		Experiment II	
	without S9 mix	with S9 mix	without S9 mix	with S9 mix
TA1535	/	5000	/	/
TA1537	/	/	5000	/
TA98	/	/	2500 – 5000	5000
TA100	/	5000	2500 – 5000	5000
WP2 (pKM101)	5000	/	2500 – 5000	5000
WP2 *uvrA* (pKM101)	/	/	/	/

/ = no cytotoxic effects, evident as a reduction in the number of revertants (below the induction factor of 0.5)

No substantial increase in revertant colony numbers in any of the six tester strains was observed following treatment with CGA15324/lambda-cyhalothrin EC (A13735F) at any concentration, neither in the presence nor absence of metabolic activation (S9 mix). There was also no tendency of higher mutation rates with increasing concentrations in the range below the generally acknowledged border of biological relevance.

Appropriate reference mutagens were used as positive controls. They showed a distinct increase in induced revertant colonies.

5.0 CONCLUSIONS

In conclusion, it can be stated that during the described mutagenicity tests and under the experimental conditions reported, CGA15324/lambda-cyhalothrin EC (A13735F) did not induce gene mutations by base pair changes or frameshifts in the genome of the strains used.

Therefore, CGA15324/lambda-cyhalothrin EC (A13735F) is considered to be non-mutagenic in the *Salmonella typhimurium* and *Escherichia coli* reverse mutation assay.

6.0 REFERENCES

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TABLES SECTION

TABLE 1 Summary of Results Pre-Experiment/Experiment I

Study Name: 2165900
Experiment: 2165900 VV Plate
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 05.05.2021
Date Counted: 10.05.2021

Metabolic Activation	Test Group	Concentration (per plate)	Revertant Colony Counts (Mean ±SD)					
			TA 1535	TA 1537	TA 98	TA 100	WP2 pKM101	WP2 uvrA pKM101
Without Activation	DMSO		12 ± 4	12 ± 4	29 ± 4	97 ± 9	253 ± 12	328 ± 30
	Untreated		11 ± 4	12 ± 2	27 ± 2	118 ± 17	291 ± 17	381 ± 37
	CGA15324/lambda-cyhalothrin EC (A13735F)	3 µg	12 ± 3	10 ± 2	34 ± 8	105 ± 5	282 ± 4	365 ± 54
		10 µg	13 ± 2	10 ± 3	34 ± 2	110 ± 13	254 ± 14	330 ± 23
		33 µg	10 ± 2	10 ± 4	31 ± 8	104 ± 8	254 ± 17	333 ± 4
		100 µg	13 ± 2	10 ± 4	25 ± 4	88 ± 18	233 ± 36	292 ± 15
		333 µg	11 ± 3	12 ± 2	20 ± 6	80 ± 21	204 ± 17	331 ± 54
		1000 µg	12 ± 4	11 ± 3	24 ± 5	85 ± 13	193 ± 11	283 ± 28
		2500 µg	10 ± 3	13 ± 3	17 ± 4	70 ± 10	142 ± 12	244 ± 16
		5000 µg	14 ± 4 P	9 ± 3 P	16 ± 5 P	54 ± 13 P M	105 ± 6 P	205 ± 13 P
	NaN3	10 µg	1110 ± 37			1793 ± 55		
	4-NOPD	10 µg			436 ± 44			
	4-NOPD	50 µg		62 ± 3				
	MMS	2.0 µL					2333 ± 159	2097 ± 232
With Activation	DMSO		13 ± 4	11 ± 3	43 ± 2	111 ± 26	274 ± 31	374 ± 20
	Untreated		13 ± 3	13 ± 3	47 ± 12	113 ± 13	299 ± 14	422 ± 21
	CGA15324/lambda-cyhalothrin EC (A13735F)	3 µg	11 ± 2	14 ± 3	34 ± 7	100 ± 2	279 ± 16	416 ± 5
		10 µg	12 ± 3	12 ± 4	40 ± 3	107 ± 9	243 ± 17	391 ± 8
		33 µg	11 ± 1	12 ± 3	40 ± 2	112 ± 12	216 ± 28	364 ± 4
		100 µg	12 ± 4	9 ± 3	33 ± 4	104 ± 6	231 ± 20	352 ± 17
		333 µg	10 ± 1	14 ± 3	33 ± 8	113 ± 8	249 ± 13	422 ± 24
		1000 µg	10 ± 0	13 ± 3	30 ± 5	79 ± 12	220 ± 17	356 ± 6
		2500 µg	9 ± 1 P	16 ± 4 P	21 ± 7 P	58 ± 8 P	185 ± 5 P	304 ± 24 P
		5000 µg	5 ± 1 P	14 ± 2 P	24 ± 4 P	41 ± 4 P M	127 ± 3 P	281 ± 31 P
	2-AA	2.5 µg	244 ± 4	464 ± 7	2703 ± 182	3676 ± 710		
	2-AA	10.0 µg					953 ± 33	1334 ± 15

Key to Positive Controls

NaN3	sodium azide
2-AA	2-aminoanthracene
4-NOPD	4-nitro-o-phenylene-diamine
MMS	methyl methane sulfonate

Key to Plate Postfix Codes

P	Precipitate
M	Manual count

TABLE 2 Summary of Results Experiment II

Study Name: 2165900
Experiment: 2165900 HV2 Pre
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 19.05.2021
Date Counted: 25.05.2021

Metabolic Activation	Test Group	Concentration (per plate)	Revertant Colony Counts (Mean ±SD)					
			TA 1535	TA 1537	TA 98	TA 100	WP2 pKM101	WP2 uvrA pKM101
Without Activation	DMSO		10 ± 4	10 ± 1	29 ± 5	95 ± 6	227 ± 21	357 ± 37
	Untreated		13 ± 4	10 ± 1	28 ± 9	107 ± 3	289 ± 20	370 ± 3
	CGA15324/lambda-cyhalothrin EC (A13735F)	10 µg						
		33 µg	10 ± 1	12 ± 4	25 ± 5	92 ± 3	194 ± 9	382 ± 66
		100 µg	6 ± 2	9 ± 0	20 ± 3	78 ± 7	160 ± 7	341 ± 24
		333 µg	9 ± 2	10 ± 1	16 ± 6	72 ± 12	109 ± 21	305 ± 8
		1000 µg	6 ± 1	8 ± 3	15 ± 4	79 ± 16	106 ± 21	293 ± 15
		2500 µg	10 ± 3	10 ± 3	12 ± 2	12 ± 1	87 ± 13	222 ± 23
		5000 µg	7 ± 1 P	3 ± 1 P M	4 ± 1 P	1 ± 1 P	45 ± 8 P	166 ± 7 P
	NaN3	10 µg	1174 ± 32			1684 ± 110		
	4-NOPD	10 µg			491 ± 4			
	4-NOPD	50 µg		83 ± 16				
	MMS	2.0 µL					3287 ± 411	2818 ± 126
With Activation	DMSO		11 ± 1	16 ± 1	34 ± 5	95 ± 8	248 ± 41	443 ± 38
	Untreated		15 ± 4	18 ± 2	40 ± 9	111 ± 6	267 ± 33	405 ± 29
	CGA15324/lambda-cyhalothrin EC (A13735F)	10 µg	13 ± 3	18 ± 2	34 ± 12	116 ± 14	246 ± 18	528 ± 38
		33 µg	13 ± 3	15 ± 5	32 ± 5	108 ± 13	237 ± 12	437 ± 21
		100 µg	12 ± 4	15 ± 0	33 ± 4	103 ± 10	232 ± 4	408 ± 39
		333 µg	13 ± 4	14 ± 4	33 ± 6	86 ± 15	198 ± 9	469 ± 21
		1000 µg	10 ± 1	17 ± 2	25 ± 3	65 ± 9	173 ± 31	368 ± 20
		2500 µg	10 ± 2 P	17 ± 6 P	22 ± 1 P	49 ± 9 P M	122 ± 3 P	299 ± 33 P
		5000 µg	8 ± 2 P	12 ± 4 P M	12 ± 3 P M	25 ± 4 P M	88 ± 5 P	233 ± 10 P
	2-AA	2.5 µg	231 ± 30	391 ± 13	2136 ± 186	3222 ± 150		
	2-AA	10.0 µg					954 ± 32	1462 ± 159

Key to Positive Controls

NaN3	sodium azide
2-AA	2-aminoanthracene
4-NOPD	4-nitro-o-phenylene-diamine
MMS	methyl methane sulfonate

Key to Plate Postfix Codes

P	Precipitate
M	Manual count

TABLE 3 Pre-Experiment and Experiment I: 2165900 VV Plate Incorporation Without Metabolic Activation

Study Name: 2165900
Experiment: 2165900 VV Plate
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 05.05.2021
Date Counted: 10.05.2021

Without metabolic activation

Strain	Compound	Concentration per plate	Mean revertants per plate	Standard Deviation	Ratio treated / solvent	Individual revertant colony counts
TA 1535	**CGA15324/ lambda-cyhalothrin EC (A13735F)**	3 µg	*11.7*	*2.9*	*1.0*	10, 15, 10
		10 µg	*12.7*	*2.1*	*1.1*	15, 12, 11
		33 µg	*10.3*	*1.5*	*0.9*	10, 12, 9
		100 µg	*13.0*	*1.7*	*1.1*	12, 12, 15
		333 µg	*11.0*	*2.6*	*0.9*	9, 10, 14
		1000 µg	*11.7*	*3.8*	*1.0*	9, 16, 10
		2500 µg	*9.7*	*2.5*	*0.8*	7, 12, 10
		5000 µg	*13.7*	*4.2*	*1.1*	17 P, 15 P, 9 P
	DMSO		*12.0*	*4.4*		14, 7, 15
	Untreated		*10.7*	*4.0*		7, 15, 10
TA 1537	**CGA15324/ lambda-cyhalothrin EC (A13735F)**	3 µg	*10.0*	*1.7*	*0.8*	12, 9, 9
		10 µg	*10.0*	*2.6*	*0.8*	12, 7, 11
		33 µg	*10.3*	*3.5*	*0.9*	14, 7, 10
		100 µg	*10.0*	*3.6*	*0.8*	7, 14, 9
		333 µg	*12.0*	*2.0*	*1.0*	14, 12, 10
		1000 µg	*10.7*	*2.9*	*0.9*	14, 9, 9
		2500 µg	*12.7*	*2.9*	*1.1*	11, 11, 16
		5000 µg	*9.3*	*3.1*	*0.8*	6 P, 12 P, 10 P
	DMSO		*12.0*	*4.4*		17, 9, 10
	Untreated		*11.7*	*2.1*		11, 14, 10
TA 98	**CGA15324/ lambda-cyhalothrin EC (A13735F)**	3 µg	*33.7*	*7.6*	*1.1*	27, 32, 42
		10 µg	*33.7*	*2.3*	*1.1*	35, 31, 35
		33 µg	*31.3*	*7.6*	*1.1*	38, 33, 23
		100 µg	*24.7*	*4.0*	*0.8*	27, 27, 20
		333 µg	*20.3*	*6.1*	*0.7*	19, 15, 27
		1000 µg	*24.3*	*4.7*	*0.8*	19, 26, 28
		2500 µg	*17.0*	*4.4*	*0.6*	19, 20, 12
		5000 µg	*16.3*	*5.1*	*0.6*	22 P, 15 P, 12 P
	DMSO		*29.3*	*4.0*		30, 33, 25
	Untreated		*27.0*	*1.7*		28, 28, 25

Key to Plate Postfix Codes

P Precipitate

Study Name: 2165900
Experiment: 2165900 VV Plate
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 05.05.2021
Date Counted: 10.05.2021

Without metabolic activation

Strain	Compound	Concen-tration per plate	Mean revertants per plate	Standard Deviation	Ratio treated / solvent	Individual revertant colony counts
TA 100	**CGA15324/**	3 µg	*105.3*	*4.5*	*1.1*	110, 105, 101
	lambda-	10 µg	*110.3*	*13.4*	*1.1*	120, 95, 116
	cyhalothrin	33 µg	*103.7*	*7.6*	*1.1*	109, 95, 107
	EC (A13735F)	100 µg	*87.7*	*18.2*	*0.9*	91, 68, 104
		333 µg	*79.7*	*20.5*	*0.8*	80, 100, 59
		1000 µg	*85.0*	*13.0*	*0.9*	85, 72, 98
		2500 µg	*70.0*	*9.6*	*0.7*	59, 77, 74
		5000 µg	*54.3*	*13.3*	*0.6*	63 P M, 39 P M, 61 P M
	DMSO		*96.7*	*9.3*		94, 89, 107
	Untreated		*118.3*	*17.0*		101, 119, 135
WP2	**CGA15324/**	3 µg	*282.0*	*3.6*	*1.1*	285, 278, 283
pKM101	**lambda-**	10 µg	*253.7*	*13.6*	*1.0*	269, 243, 249
	cyhalothrin	33 µg	*254.3*	*17.1*	*1.0*	274, 246, 243
	EC (A13735F)	100 µg	*233.0*	*36.4*	*0.9*	272, 227, 200
		333 µg	*203.7*	*17.2*	*0.8*	211, 184, 216
		1000 µg	*193.3*	*11.2*	*0.8*	189, 185, 206
		2500 µg	*142.0*	*11.5*	*0.6*	131, 141, 154
		5000 µg	*104.7*	*6.1*	*0.4*	98 P, 106 P, 110 P
	DMSO		*253.3*	*11.5*		265, 253, 242
	Untreated		*291.0*	*17.3*		295, 306, 272
WP2	**CGA15324/**	3 µg	*365.0*	*54.1*	*1.1*	427, 341, 327
uvrA	**lambda-**	10 µg	*330.0*	*22.6*	*1.0*	351, 333, 306
pKM101	**cyhalothrin**	33 µg	*333.3*	*4.2*	*1.0*	332, 338, 330
	EC (A13735F)	100 µg	*292.3*	*15.3*	*0.9*	310, 283, 284
		333 µg	*331.0*	*53.7*	*1.0*	382, 336, 275
		1000 µg	*282.7*	*27.7*	*0.9*	279, 257, 312
		2500 µg	*244.0*	*15.5*	*0.7*	259, 228, 245
		5000 µg	*204.7*	*13.1*	*0.6*	190 P, 215 P, 209 P
	DMSO		*328.0*	*29.6*		342, 348, 294
	Untreated		*380.7*	*37.1*		419, 378, 345
TA 1535	**NaN3**	10 µg	*1109.7*	*37.2*	*92.5*	1142, 1118, 1069
TA 1537	**4-NOPD**	50 µg	*62.3*	*2.9*	*5.2*	59, 64, 64
TA 98	**4-NOPD**	10 µg	*436.3*	*44.2*	*14.9*	483, 431, 395
TA 100	**NaN3**	10 µg	*1793.0*	*54.6*	*18.5*	1761, 1856, 1762
WP2 pKM101	**MMS**	2.0 µL	*2333.0*	*158.8*	*9.2*	2194, 2299, 2506
WP2 uvrA pKM101	**MMS**	2.0 µL	*2097.0*	*231.8*	*6.4*	1909, 2026, 2356

Key to Positive Controls

NaN3	sodium azide
4-NOPD	4-nitro-o-phenylene-diamine
MMS	methyl methane sulfonate

Key to Plate Postfix Codes

P	Precipitate
M	Manual count

TABLE 4 Pre-Experiment and Experiment I: 2165900 VV Plate Incorporation With Metabolic Activation

Study Name: 2165900
Experiment: 2165900 VV Plate
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 05.05.2021
Date Counted: 10.05.2021

With metabolic activation

Strain	Compound	Concen-tration per plate	Mean revertants per plate	Standard Deviation	Ratio treated / solvent	Individual revertant colony counts
TA 1535	**CGA15324/**	3 µg	*10.7*	*1.5*	*0.8*	9, 11, 12
	lambda-	10 µg	*11.7*	*2.5*	*0.9*	12, 9, 14
	cyhalothrin	33 µg	*11.3*	*0.6*	*0.9*	11, 11, 12
	EC (A13735F)	100 µg	*12.3*	*4.0*	*1.0*	10, 17, 10
		333 µg	*9.7*	*0.6*	*0.8*	10, 10, 9
		1000 µg	*10.0*	*0.0*	*0.8*	10, 10, 10
		2500 µg	*9.3*	*0.6*	*0.7*	9 P, 10 P, 9 P
		5000 µg	*5.0*	*1.0*	*0.4*	5 P, 6 P, 4 P
	DMSO		*12.7*	*3.8*		10, 17, 11
	Untreated		*13.3*	*3.1*		10, 16, 14
TA 1537	**CGA15324/**	3 µg	*13.7*	*2.5*	*1.2*	11, 16, 14
	lambda-	10 µg	*11.7*	*4.0*	*1.0*	14, 7, 14
	cyhalothrin	33 µg	*12.0*	*2.6*	*1.1*	15, 10, 11
	EC (A13735F)	100 µg	*9.3*	*2.5*	*0.8*	9, 7, 12
		333 µg	*14.3*	*3.1*	*1.3*	15, 17, 11
		1000 µg	*12.7*	*3.2*	*1.1*	9, 15, 14
		2500 µg	*15.7*	*3.5*	*1.4*	19 P, 16 P, 12 P
		5000 µg	*14.0*	*2.0*	*1.2*	14 P, 12 P, 16 P
	DMSO		*11.3*	*2.5*		11, 9, 14
	Untreated		*12.7*	*3.1*		10, 12, 16
TA 98	**CGA15324/**	3 µg	*33.7*	*7.4*	*0.8*	31, 28, 42
	lambda-	10 µg	*40.3*	*3.1*	*0.9*	41, 37, 43
	cyhalothrin	33 µg	*39.7*	*1.5*	*0.9*	40, 38, 41
	EC (A13735F)	100 µg	*32.7*	*3.8*	*0.8*	37, 30, 31
		333 µg	*32.7*	*8.0*	*0.8*	25, 41, 32
		1000 µg	*30.3*	*4.6*	*0.7*	25, 33, 33
		2500 µg	*21.0*	*7.0*	*0.5*	14 P, 21 P, 28 P
		5000 µg	*23.7*	*4.2*	*0.6*	27 P, 25 P, 19 P
	DMSO		*43.0*	*1.7*		41, 44, 44
	Untreated		*47.3*	*11.8*		40, 61, 41

Key to Plate Postfix Codes

P Precipitate

Study Name: 2165900
Experiment: 2165900 VV Plate
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 05.05.2021
Date Counted: 10.05.2021

With metabolic activation

Strain	Compound	Concen-tration per plate	Mean revertants per plate	Standard Deviation	Ratio treated / solvent	Individual revertant colony counts
TA 100	**CGA15324/**	3 µg	*99.7*	*2.1*	*0.9*	98, 99, 102
	lambda-	10 µg	*107.3*	*8.5*	*1.0*	107, 116, 99
	cyhalothrin	33 µg	*111.7*	*11.9*	*1.0*	117, 98, 120
	EC (A13735F)	100 µg	*104.3*	*5.7*	*0.9*	106, 109, 98
		333 µg	*113.3*	*8.0*	*1.0*	105, 121, 114
		1000 µg	*79.3*	*11.5*	*0.7*	68, 79, 91
		2500 µg	*58.3*	*8.1*	*0.5*	62 P, 64 P, 49 P
		5000 µg	*41.0*	*3.6*	*0.4*	44 P M, 37 P M, 42 P M
	DMSO		*111.3*	*25.9*		121, 131, 82
	Untreated		*113.3*	*12.7*		106, 106, 128
WP2	**CGA15324/**	3 µg	*279.0*	*15.5*	*1.0*	278, 264, 295
pKM101	**lambda-**	10 µg	*243.3*	*16.8*	*0.9*	247, 225, 258
	cyhalothrin	33 µg	*216.0*	*28.2*	*0.8*	237, 227, 184
	EC (A13735F)	100 µg	*231.3*	*19.5*	*0.8*	209, 240, 245
		333 µg	*249.3*	*13.2*	*0.9*	261, 252, 235
		1000 µg	*219.7*	*16.8*	*0.8*	238, 205, 216
		2500 µg	*185.0*	*5.2*	*0.7*	188 P, 188 P, 179 P
		5000 µg	*127.3*	*2.5*	*0.5*	130 P, 125 P, 127 P
	DMSO		*274.0*	*30.8*		282, 300, 240
	Untreated		*299.3*	*13.8*		294, 289, 315
WP2	**CGA15324/**	3 µg	*416.0*	*5.3*	*1.1*	422, 412, 414
uvrA	**lambda-**	10 µg	*390.7*	*8.1*	*1.0*	387, 385, 400
pKM101	**cyhalothrin**	33 µg	*364.0*	*4.4*	*1.0*	362, 369, 361
	EC (A13735F)	100 µg	*352.0*	*16.5*	*0.9*	369, 336, 351
		333 µg	*422.3*	*23.5*	*1.1*	424, 445, 398
		1000 µg	*356.3*	*6.4*	*1.0*	349, 359, 361
		2500 µg	*304.0*	*24.3*	*0.8*	291 P, 289 P, 332 P
		5000 µg	*281.3*	*31.2*	*0.8*	293 P, 246 P, 305 P
	DMSO		*374.3*	*20.0*		375, 354, 394
	Untreated		*421.7*	*21.1*		446, 410, 409
TA 1535	**2-AA**	2.5 µg	*244.3*	*4.2*	*19.3*	249, 241, 243
TA 1537	**2-AA**	2.5 µg	*464.3*	*6.7*	*41.0*	466, 457, 470
TA 98	**2-AA**	2.5 µg	*2702.7*	*182.2*	*62.9*	2496, 2772, 2840
TA 100	**2-AA**	2.5 µg	*3676.0*	*710.2*	*33.0*	4136, 2858, 4034
WP2 pKM101	**2-AA**	10.0 µg	*953.0*	*33.0*	*3.5*	987, 921, 951
WP2 uvrA pKM101	**2-AA**	10.0 µg	*1334.3*	*14.6*	*3.6*	1332, 1350, 1321

Key to Positive Controls

2-AA	2-aminoanthracene

Key to Plate Postfix Codes

P	Precipitate
M	Manual count

TABLE 5 Experiment II: 2165900 HV2 Pre Incubation Without Metabolic Activation

Study Name: 2165900
Experiment: 2165900 HV2 Pre
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 19.05.2021
Date Counted: 25.05.2021

Without metabolic activation

Strain	Compound	Concen-tration per plate	Mean revertants per plate	Standard Deviation	Ratio treated / solvent	Individual revertant colony counts
TA 1535	**CGA15324/**	10 µg				
	lambda-	33 µg	*10.0*	*1.0*	*1.0*	9, 11, 10
	cyhalothrin	100 µg	*6.3*	*2.3*	*0.6*	9, 5, 5
	EC (A13735F)	333 µg	*9.3*	*2.1*	*0.9*	7, 10, 11
		1000 µg	*6.3*	*0.6*	*0.6*	6, 7, 6
		2500 µg	*10.0*	*2.6*	*1.0*	11, 7, 12
		5000 µg	*6.7*	*0.6*	*0.7*	7 P, 7 P, 6 P
	DMSO		*10.0*	*3.6*		14, 7, 9
	Untreated		*12.7*	*3.8*		17, 11, 10
TA 1537	**CGA15324/**	10 µg				
	lambda-	33 µg	*12.3*	*3.5*	*1.2*	9, 16, 12
	cyhalothrin	100 µg	*9.0*	*0.0*	*0.9*	9, 9, 9
	EC (A13735F)	333 µg	*9.7*	*0.6*	*0.9*	9, 10, 10
		1000 µg	*8.3*	*3.2*	*0.8*	6, 12, 7
		2500 µg	*10.3*	*2.9*	*1.0*	12, 12, 7
		5000 µg	*3.0*	*1.0*	*0.3*	4 P M, 2 P M, 3 P M
	DMSO		*10.3*	*1.2*		11, 11, 9
	Untreated		*10.0*	*1.0*		11, 10, 9
TA 98	**CGA15324/**	10 µg				
	lambda-	33 µg	*25.3*	*4.9*	*0.9*	31, 23, 22
	cyhalothrin	100 µg	*19.7*	*3.2*	*0.7*	22, 21, 16
	EC (A13735F)	333 µg	*16.3*	*5.9*	*0.6*	12, 14, 23
		1000 µg	*15.3*	*3.5*	*0.5*	19, 15, 12
		2500 µg	*12.3*	*1.5*	*0.4*	11, 12, 14
		5000 µg	*4.3*	*0.6*	*0.2*	4 P, 4 P, 5 P
	DMSO		*28.7*	*5.1*		30, 23, 33
	Untreated		*27.7*	*8.6*		20, 26, 37
TA 100	**CGA15324/**	10 µg				
	lambda-	33 µg	*91.7*	*3.2*	*1.0*	88, 94, 93
	cyhalothrin	100 µg	*78.0*	*7.0*	*0.8*	75, 73, 86
	EC (A13735F)	333 µg	*72.3*	*11.9*	*0.8*	86, 64, 67
		1000 µg	*79.0*	*15.9*	*0.8*	61, 91, 85
		2500 µg	*11.7*	*0.6*	*0.1*	12, 11, 12
		5000 µg	*1.3*	*0.6*	*0.0*	2 P, 1 P, 1 P
	DMSO		*95.3*	*5.9*		93, 102, 91
	Untreated		*106.7*	*2.5*		107, 104, 109

Key to Plate Postfix Codes

P	Precipitate
M	Manual count

Study Name: 2165900
Experiment: 2165900 HV2 Pre
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 19.05.2021
Date Counted: 25.05.2021

Without metabolic activation

Strain	Compound	Concen-tration per plate	Mean revertants per plate	Standard Deviation	Ratio treated / solvent	Individual revertant colony counts
WP2	**CGA15324/**	10 µg				
pKM101	**lambda-**	33 µg	*194.0*	*8.7*	*0.9*	198, 184, 200
	cyhalothrin	100 µg	*159.7*	*7.4*	*0.7*	154, 157, 168
	EC (A13735F)	333 µg	*109.0*	*21.2*	*0.5*	117, 125, 85
		1000 µg	*106.0*	*21.2*	*0.5*	130, 98, 90
		2500 µg	*87.3*	*13.2*	*0.4*	90, 99, 73
		5000 µg	*44.7*	*7.6*	*0.2*	43 P, 38 P, 53 P
	DMSO		*227.0*	*21.4*		251, 220, 210
	Untreated		*289.3*	*20.3*		312, 283, 273
WP2	**CGA15324/**	10 µg				
uvrA	**lambda-**	33 µg	*381.7*	*66.3*	*1.1*	451, 375, 319
pKM101	**cyhalothrin**	100 µg	*341.3*	*24.4*	*1.0*	336, 320, 368
	EC (A13735F)	333 µg	*304.7*	*8.1*	*0.9*	296, 312, 306
		1000 µg	*293.0*	*14.7*	*0.8*	306, 296, 277
		2500 µg	*222.3*	*22.7*	*0.6*	248, 214, 205
		5000 µg	*166.0*	*6.6*	*0.5*	172 P, 167 P, 159 P
	DMSO		*356.7*	*37.1*		395, 354, 321
	Untreated		*370.0*	*2.6*		373, 369, 368
TA 1535	**NaN3**	10 µg	*1173.7*	*31.6*	*117.4*	1210, 1153, 1158
TA 1537	**4-NOPD**	50 µg	*83.0*	*15.6*	*8.0*	73, 101, 75
TA 98	**4-NOPD**	10 µg	*491.3*	*3.5*	*17.1*	488, 491, 495
TA 100	**NaN3**	10 µg	*1683.7*	*110.0*	*17.7*	1687, 1792, 1572
WP2 pKM101	**MMS**	2.0 µL	*3286.7*	*410.7*	*14.5*	3014, 3087, 3759
WP2 uvrA pKM101	**MMS**	2.0 µL	*2818.0*	*125.6*	*7.9*	2941, 2690, 2823

Key to Positive Controls		Key to Plate Postfix Codes	
NaN3	sodium azide	P	Precipitate
4-NOPD	4-nitro-o-phenylene-diamine		
MMS	methyl methane sulfonate		

TABLE 6 Experiment II: 2165900 HV2 Pre Incubation With Metabolic Activation

Study Name: 2165900
Experiment: 2165900 HV2 Pre
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 19.05.2021
Date Counted: 25.05.2021

With metabolic activation

Strain	Compound	Concen-tration per plate	Mean revertants per plate	Standard Deviation	Ratio treated / solvent	Individual revertant colony counts
TA 1535	**CGA15324/**	10 µg	*13.3*	*3.2*	*1.2*	11, 17, 12
	lambda-	33 µg	*13.0*	*2.6*	*1.2*	16, 12, 11
	cyhalothrin	100 µg	*11.7*	*3.8*	*1.1*	10, 16, 9
	EC (A13735F)	333 µg	*13.0*	*3.6*	*1.2*	17, 12, 10
		1000 µg	*9.7*	*1.2*	*0.9*	9, 9, 11
		2500 µg	*10.3*	*1.5*	*0.9*	12 P, 10 P, 9 P
		5000 µg	*8.0*	*1.7*	*0.7*	6 P, 9 P, 9 P
	DMSO		*11.0*	*1.0*		11, 10, 12
	Untreated		*14.7*	*4.0*		14, 11, 19
TA 1537	**CGA15324**	10 µg	*17.7*	*2.1*	*1.1*	20, 16, 17
	/lambda-	33 µg	*15.0*	*5.0*	*0.9*	20, 10, 15
	cyhalothrin	100 µg	*15.0*	*0.0*	*0.9*	15, 15, 15
	EC (A13735F)	333 µg	*14.0*	*3.6*	*0.9*	15, 10, 17
		1000 µg	*17.0*	*2.0*	*1.0*	19, 17, 15
		2500 µg	*17.3*	*5.5*	*1.1*	23 P, 17 P, 12 P
		5000 µg	*12.3*	*3.5*	*0.8*	16 P M, 9 P M, 12 P M
	DMSO		*16.3*	*1.2*		17, 17, 15
	Untreated		*17.7*	*2.3*		19, 15, 19
TA 98	**CGA15324/**	10 µg	*34.0*	*12.2*	*1.0*	26, 48, 28
	lambda-	33 µg	*31.7*	*4.5*	*0.9*	32, 27, 36
	cyhalothrin	100 µg	*33.0*	*3.6*	*1.0*	37, 32, 30
	EC (A13735F)	333 µg	*33.3*	*5.7*	*1.0*	27, 35, 38
		1000 µg	*25.3*	*2.5*	*0.7*	23, 25, 28
		2500 µg	*22.0*	*1.0*	*0.6*	23 P, 22 P, 21 P
		5000 µg	*12.0*	*3.0*	*0.4*	9 P M, 12 P M, 15 P M
	DMSO		*34.0*	*5.3*		30, 32, 40
	Untreated		*39.7*	*8.5*		43, 30, 46
TA 100	**CGA15324/**	10 µg	*116.3*	*14.2*	*1.2*	100, 126, 123
	lambda-	33 µg	*108.3*	*13.2*	*1.1*	94, 120, 111
	cyhalothrin	100 µg	*103.0*	*10.1*	*1.1*	114, 94, 101
	EC (A13735F)	333 µg	*86.3*	*15.0*	*0.9*	72, 85, 102
		1000 µg	*65.3*	*8.7*	*0.7*	75, 58, 63
		2500 µg	*49.0*	*8.7*	*0.5*	59 P M, 45 P M, 43 P M
		5000 µg	*25.0*	*4.4*	*0.3*	23 P M, 30 P M, 22 P M
	DMSO		*94.7*	*8.4*		85, 100, 99
	Untreated		*111.3*	*6.4*		104, 115, 115

Key to Plate Postfix Codes

P	Precipitate
M	Manual count

Study Name: 2165900
Experiment: 2165900 HV2 Pre
Assay Conditions:

Study Code: ICCR 2165900
Date Plated: 19.05.2021
Date Counted: 25.05.2021

With metabolic activation

Strain	Compound	Concen-tration per plate	Mean revertants per plate	Standard Deviation	Ratio treated / solvent	Individual revertant colony counts
WP2	**CGA15324/**	10 µg	*245.7*	*18.0*	*1.0*	263, 247, 227
pKM101	**lambda-**	33 µg	*236.7*	*11.6*	*1.0*	249, 235, 226
	cyhalothrin	100 µg	*232.3*	*4.0*	*0.9*	230, 237, 230
	EC (A13735F)	333 µg	*198.3*	*9.0*	*0.8*	199, 189, 207
		1000 µg	*172.7*	*31.2*	*0.7*	146, 207, 165
		2500 µg	*121.7*	*2.9*	*0.5*	120 P, 120 P, 125 P
		5000 µg	*88.0*	*5.2*	*0.4*	94 P, 85 P, 85 P
	DMSO		*248.0*	*41.0*		248, 207, 289
	Untreated		*267.3*	*33.3*		278, 230, 294
WP2	**CGA15324/**	10 µg	*527.7*	*38.1*	*1.2*	545, 554, 484
uvrA	**lambda-**	33 µg	*437.3*	*20.6*	*1.0*	453, 414, 445
pKM101	**cyhalothrin**	100 µg	*408.0*	*39.1*	*0.9*	382, 453, 389
	EC (A13735F)	333 µg	*469.0*	*20.8*	*1.1*	457, 493, 457
		1000 µg	*368.3*	*19.6*	*0.8*	387, 348, 370
		2500 µg	*299.3*	*32.7*	*0.7*	337 P, 282 P, 279 P
		5000 µg	*232.7*	*10.4*	*0.5*	221 P, 241 P, 236 P
	DMSO		*443.3*	*38.1*		454, 475, 401
	Untreated		*404.7*	*29.5*		394, 438, 382
TA 1535	**2-AA**	2.5 µg	*231.0*	*30.1*	*21.0*	238, 257, 198
TA 1537	**2-AA**	2.5 µg	*391.3*	*12.5*	*24.0*	391, 404, 379
TA 98	**2-AA**	2.5 µg	*2135.7*	*186.2*	*62.8*	1934, 2301, 2172
TA 100	**2-AA**	2.5 µg	*3222.3*	*149.7*	*34.0*	3376, 3214, 3077
WP2 pKM101	**2-AA**	10.0 µg	*954.3*	*32.1*	*3.8*	951, 924, 988
WP2 uvrA pKM101	**2-AA**	10.0 µg	*1462.3*	*158.6*	*3.3*	1307, 1456, 1624

Key to Positive Controls		Key to Plate Postfix Codes	
2-AA	2-aminoanthracene	P	Precipitate

APPENDICES SECTION

APPENDIX 1 Historical Control Data

These data represent the laboratory´s historical control data from July 2018 until July 2020 representing approx. 600 experiments (WP2 pKM101, WP2 uvrA pKM101 the historical data are based on approx. 80 experiments).

The positive controls that used to compile the historical positive control data correspond to the positive control substances described in Methods; section 3.2.2 (Positive control substances).

Strain		without S9 mix				with S9 mix			
		Mean	SD	Min	Max	Mean	SD	Min	Max
TA 1535	Solvent control	12	2.6	7	22	13	2.5	7	24
	Untreated control	12	2.9	6	26	13	2.8	7	23
	Positive control	1116	141.3	340	1612	346	72.1	170	736
TA1537	Solvent control	11	2.4	6	20	14	2.8	7	28
	Untreated control	11	2.8	5	22	14	3.2	7	30
	Positive control	83	22.1	48	400	286	98.7	82	630
TA 98	Solvent control	28	4.9	13	46	38	6.4	12	62
	Untreated control	29	5.0	14	48	41	6.8	14	64
	Positive control	421	91.2	216	1218	3275	774.9	322	5699
TA 100	Solvent control	127	30.7	63	214	131	30.0	72	214
	Untreated control	135	35.7	64	233	140	34.4	68	217
	Positive control	1759	273.4	511	2588	3566	837.6	553	5444
WP2 pKM 101	Solvent control	248	31.7	171	299	266	33.0	205	315
	Untreated control	269	26.6	212	346	299	28.2	233	345
	Positive control	3343	428.4	2332	4653	1092	257.8	933	2781
WP2uvrA pKM 101	Solvent control	322	31.6	248	388	375	38.5	287	466
	Untreated control	346	28.2	279	403	393	32.6	313	480
	Positive control	3176	468.5	2021	4717	1897	183.2	1270	2464

Mean = mean value of revertants/plate
SD = standard deviation
Min = minimal value
Max = maximal value

APPENDIX 2 Copy of GLP Certificate

HESSEN

Gute Laborpraxis/Good Laboratory Practice

GLP-Bescheinigung/Statement of GLP Compliance
(gemäß/according to § 19b Abs. 1 Chemikaliengesetz)

Eine GLP-Inspektion zur Überwachung der Einhaltung der GLP-Grundsätze gemäß Chemikaliengesetz bzw. Richtlinie 2004/9/EG wurde durchgeführt in

Assessment of conformity with GLP according to Chemikaliengesetz and Directive 2004/9/EEC at:

☒ Prüfeinrichtung/Test facility ☐ Prüfstandort/Test site

ICCR-Roßdorf GmbH
Institute for Competent Contract Research
In den Leppsteinswiesen 19
64380 Roßdorf

(Unverwechselbare Bezeichnung und Adresse/Unequivocal name and adress)

Prüfungen nach Kategorien/Areas of Expertise
(gemäß/according ChemVwV-GLP Nr. 5.3/OECD guidance)

2 Prüfungen zur Bestimmung der toxikologischen Eigenschaften	**2** Toxicity studies
3 Prüfungen zur Bestimmung der erbgutverändernden Eigenschaften (in vitro und in vivo)	**3** Mutagenicity studies
8 Analytische Prüfungen an biologischen Materialien	**8** Analytical and clinical chemistry testing

22.11.2018, 21.02.2019, 12. bis 14.03.2019
Datum der Inspektion/Date of Inspection
(Tag Monat Jahr/day month year)

Die genannte Prüfeinrichtung befindet sich im nationalen GLP-Überwachungsverfahren und wird regelmäßig auf Einhaltung der GLP-Grundsätze überwacht.

The above mentioned test facility is included in the national GLP Compliance Programme and is inspected on a regular basis.

Auf der Grundlage des Inspektionsberichtes wird hiermit bestätigt, dass in dieser Prüfeinrichtung die oben genannten Prüfungen unter Einhaltung der GLP- Grundsätze durchgeführt werden können.

Based on the inspection report it can be confirmed, that this test facility is able to conduct the aforementioned studies in compliance with the Principles of GLP.

Im Auftrag

Brandt

Dr. Astrid Brandt, Referentin, Wiesbaden, den **23. Oktober 2019**
(Name und Funktion der verantwortlichen Person/
Name and function of responsible person)

HESSISCHES MINISTERIUM FÜR UMWELT, KLIMASCHUTZ, LANDWIRTSCHAFT UND VERBRAUCHERSCHUTZ

Hessisches Ministerium für Umwelt, Klimaschutz, Landwirtschaft und Verbraucherschutz, Mainzer Straße 80, D 65189 Wiesbaden
(Name und Adresse der GLP-Überwachungsbehörde/Name and address of the GLP Monitoring Authority)

English name and address of the GLP Monitoring Authority: Hessian Ministry for Environment, Climate Protection, Agriculture and Consumer Protection; Department II 10; P.O. Box 31 09; 65189 Wiesbaden
Translation of seal inscription: Hessian Ministry for Environment, Climate Protection, Agriculture and Consumer Protection

APPENDIX 3 Certificate of S9

ICCR ROßDORF

CERTIFICATE

ICCR-Roßdorf S9 Preparation Lot No. 291020D

Date of preparation: October 29, 2020
Release date: November 11, 2020

Protein assay: 31.2 mg protein / ml S9

Sterility: 0 colonies / ml S9 on glucose-minimal-agar

Salmonella typhimurium assay (AMES-test)

Treatment	µl S9 / plate	number of revertants in TA 98
negative	0	29
control	100	34
10 µg/plate	0	84
2-Aminoanthracene	100	2898
10 µg/plate	0	30
Benzo(a)pyrene	100	118

The S9 was obtained from the livers of male Wistar rats which received triple treatments of 80 mg / kg body weight Phenobarbital and ß-Naphthoflavone orally on consecutive days. The livers were prepared 24 hours after the last treatment.

Quality Assurance Auditor
ICCR-Roßdorf GmbH — H. Pilawa — 17. NOV. 2020 — Date

Dr. Steffen Naumann
Study Director
ICCR-Roßdorf GmbH — 18. NOV. 2020 — Date

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Registergericht Darmstadt, HRB 6837, USt.-ID DE812333696
Geschäftsführer: Dr. Markus Schulz

SOP Origin TS-SOP S9_23

CERTIFICATE

ICCR-Roßdorf S9 Preparation Lot No. 291020D

Date of preparation: October 29, 2020
Recertification date: May 10, 2021

Protein assay: 31.2 mg protein / ml S9

Sterility: 0 colonies / ml S9 on glucose-minimal-agar

Salmonella typhimurium assay (AMES-test)

Treatment	µl S9 / plate	number of revertants in TA 98	number of revertants in TA 98 (Recertification)
negative	0	29	30
control	100	34	34
10 µg/plate 2-Aminoanthracene	0	84	86
	100	2898	1978
10 µg/plate Benzo(a)pyrene	0	30	25
	100	118	106

The S9 was obtained from the livers of male Wistar rats which received triple treatments of 80 mg / kg body weight Phenobarbital and ß-Naphthoflavone orally on consecutive days. The livers were prepared 24 hours after the last treatment.

S. Ebert Sabine Ebert — 20. MAI 2021

Quality Assurance Auditor
ICCR-Roßdorf GmbH — Date

20. MAI 2021

Dr. Steffen Naumann
Study Director
ICCR-Roßdorf GmbH — Date

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Geschäftsführer: Dr. Markus Schulz

SOP Origin TS-SOP S9_23

APPENDIX 4 Certificate of Analysis

Syngenta Crop Protection AG
GLP Testing Facility WMU
Analytical Development & Product Chemistry
Breitenloh 5
4333 Münchwilen, Switzerland

Certificate of Analysis

A13735F
CGA15324/lambda-cyhalothrin EC (300/015)
SMU1BL001

Batch Identification	**SMU1BL001**
Other Batch ID	1182111
Product Code	**A13735F**
Other Product Code(s)	CGA15324/lambda-cyhalothrin EC (300/015)
Chemical Analysis (Active Ingredient content)	
– **Identity of the Active Ingredient(s)***	**confirmed**
– **Content of profenofos***	**27.3 % w/w corresponding to 300 g/l**
– **Content of lambda-cyhalothrin***	**1.37 % w/w corresponding to 15.1 g/l**
	The Active Ingredient(s) content is within the FAO limits.
Methodology used for Characterization / Recertification	GC, oscillating density meter
Physical Analysis	
– **Appearance**	**brown liquid**
– **Density***	**1099 kg/m³**
Stability:	
– **Storage Temperature**	**< 30 °C**
– **Recertification Date**	**End of February 2023**

If stored under the conditions given above, this test substance can be considered stable until the recertification date is reached.
This Certificate of Analysis summarizes data which originates either from a single study or from several individual studies. Tests marked with an asterisk (*) have been conducted in compliance with GLP.
Raw data, documentation, study plans, any amendments to study plans and reports pertaining to this/these study/studies are stored under the study number(s) referenced below within the archives of the GLP Testing Facility WMU at Syngenta Crop Protection AG, Switzerland.

Study number of batch characterization:	CHMU210204
Study number(s) of batch recertification:	---

Authorization: 01 APR 2021

Daniel Jenniches
Analytical Development & Product Chemistry

Page 1 of 1