ARRIVE The ARRIVE Guidelines Checklist

Item	ltem No	RECOMMENDATION	Reported on Page Number/Line Number	Reported on Section/Paragraph
Title	1	Provide as accurate and concise a description of the content of the article as possible.	Page1/Line3-5	Title
Abstract	2	Provide an accurate summary of the background, research objectives, including details of the species or strain of animal used, key methods, principal findings and conclusions of the study.	Page2/Line35-58	Abstract
INTRODUCTIO	N			
Background	3	a. Include sufficient scientific background (including relevant references to previous work) to understand the motivation and context for the study, and explain the experimental approach and rationale.b. Explain how and why the animal species and model being used can address the scientific objectives and, where appropriate, the study's relevance to human biology.	Page2-3/Line67-103	Introduction/Paragraph1-2
Objectives	4	Clearly describe the primary and any secondary objectives of the study, or specific hypotheses being tested.	Page2-3/Line100-103	Introduction/Paragraph2
METHODS				
Ethical statement	5	Indicate the nature of the ethical review permissions, relevant licences (e.g. Animal [Scientific Procedures] Act 1986), and national or institutional guidelines for the care and use of animals, that cover the research.	Page11/Line759-762	Ethical Statement/Paragraph25
Study design	6	 For each experiment, give brief details of the study design including: a. The number of experimental and control groups. b. Any steps taken to minimise the effects of subjective bias when allocating animals to treatment (e.g. randomisation procedure) and when assessing results (e.g. if done, describe who was blinded and when). c. The experimental unit (e.g. a single animal, group or cage of animals). A time-line diagram or flow chart can be useful to illustrate how complex study designs were carried out. 	Page3-4/Line110-294	Methods/Paragraph3-5
Experimental procedures	7	 For each experiment and each experimental group, including controls, provide precise details of all procedures carried out. For example: a. How (e.g. drug formulation and dose, site and route of administration, anaesthesia and analgesia used [including monitoring], surgical procedure, method of euthanasia). Provide details of any specialist equipment used, including supplier(s). b. When (e.g. time of day). c. Where (e.g. home cage, laboratory, water maze). d. Why (e.g. rationale for choice of specific anaesthetic, route of administration, drug dose used). 	Page3-4/Line110-294	Methods/Paragraph3-5

8	 a. Provide details of the animals used, including species, strain, sex, developmental stage (e.g. mean or median age plus age range) and weight (e.g. mean or median weight plus weight range). b. Provide further relevant information such as the source of animals, international strain nomenclature, genetic modification status (e.g. knock-out or transgenic), genotype, health/immune status, drug or test naïve, previous procedures, etc. 	Page3-4/Line110-289	Methods/Paragraph3-4
9	 Provide details of: a. Housing (type of facility e.g. specific pathogen free [SPF]; type of cage or housing; bedding material; number of cage companions; tank shape and material etc. for fish). b. Husbandry conditions (e.g. breeding programme, light/dark cycle, temperature, quality of water etc for fish, type of food, access to food and water, environmental enrichment). c. Welfare-related assessments and interventions that were carried out prior to, during, or after the experiment. 	Page3-4/Line110-281	Methods/Paragraph3
10	a. Specify the total number of animals used in each experiment, and the number of animals in each experimental group.b. Explain how the number of animals was arrived at. Provide details of any sample size calculation used.c. Indicate the number of independent replications of each experiment, if relevant.	Page3-4/Line110-281	Methods/Paragraph3
11	 a. Give full details of how animals were allocated to experimental groups, including randomisation or matching if done. b. Describe the order in which the animals in the different experimental groups were treated and assessed. 	Page3-4/Line110-281	Methods/Paragraph3
12	Clearly define the primary and secondary experimental outcomes assessed (e.g. cell death, molecular markers, behavioural changes).	N/A	N/A
13	a. Provide details of the statistical methods used for each analysis.b. Specify the unit of analysis for each dataset (e.g. single animal, group of animals, single neuron).c. Describe any methods used to assess whether the data met the assumptions of the statistical approach.	Page6/Line344-350	Methods/Paragraph 10
14	For each experimental group, report relevant characteristics and health status of animals (e.g. weight, microbiological status, and drug or test naïve) prior to treatment or testing. (This information can often be tabulated).	N/A	N/A
15	 a. Report the number of animals in each group included in each analysis. Report absolute numbers (e.g. 10/20, not 50%²). b. If any animals or data were not included in the analysis, explain why. 	N/A	N/A
16	Report the results for each analysis carried out, with a measure of precision (e.g. standard error or confidence interval).	Page6-8/Line353-446	Result/Paragraph 11-15
17	a. Give details of all important adverse events in each experimental group.b. Describe any modifications to the experimental protocols made to reduce adverse events.	N/A	N/A
	9 10 11 12 13 14 15 16	plus age range) and weight (e.g. mean or median weight plus weight range). b. Provide further relevant information such as the source of animals, international strain nomenclature, genetic modification status (e.g. knock-out or transgenic), genotype, health/immune status, drug or test naïve, previous procedures, etc. 9 Provide details of: a. Housing (type of facility e.g. specific pathogen free [SPF]; type of cage or housing; bedding material; number of cage companions; tank shape and material etc. for fish). b. Husbandry conditions (e.g. breeding programme, light/dark cycle, temperature, quality of water etc for fish, type of food, access to food and wate, environmental enrichment). c. Welfare-related assessments and interventions that were carried out prior to, during, or after the experimental group. 10 a. Specify the total number of animals used in each experiment, and the number of animals in each experimental group. 11 a. Give full details of how animals were allocated to experimental groups, including randomisation or matching if done. 12 Clearly define the primary and secondary experimental outcomes assessed (e.g. cell death, molecular markers, behavioural changes). 13 a. Provide details of the statistical methods used for each analysis. 14 For each experimental group, report relevant characteristics and health status of animals (e.g. weight, microbiological status, and drug or test naive) prior to treatment or testing. (This information can often be tabulated). 15 a. Report the number of animals in each group included in each analysis. Report absolute numbers (e.g. 10/20, not	plus age range) and weight (e.g. mean or median weight plus weight range). b. Provide further relevant information such as the source of animals, international strain nomenclature, genetic modification status (e.g., knock-out or transgenic), genotype, health/immune status, drug or test naive, previous procedures, etc. Page3-4/Line110-281 9 Provide details of: a. Housing (type of facility e.g. specific pathogen free [SPF]; type of cage or housing; bedding material; number of cage companions; tank stape and material etc. for fish). b. Husbandry conditions (e.g., breeding programme, light/dark cycle, temperature, quality of water etc for fish, type of food, access to food and water, environmental anrichment). c. Welfare-related assessments and interventions that were carried out prior to, during, or after the experimental group. Page3-4/Line110-281 10 a. Specify the total number of animals used in each experiment, and the number of animals in each experimental could on used. Page3-4/Line110-281 11 a. Give full details of how animals were allocated to experiment, if relevant. Page3-4/Line110-281 12 Clearly define the primary and secondary experimental groups, including randomisation or matching if done. N/A 13 a. Brovide datails of the statistical methods used for each analysis. Specify the unit of analysis for each dataset (e.g. single animal, group of animals, e.g. weight, microbiological status, and drug or test naive) prior to treatment or testing. (This information can often be tabulated). N/A 13 a. Provide details of the statist

DISCUSSION				
Interpretation/ scientific implications	18	 a. Interpret the results, taking into account the study objectives and hypotheses, current theory and other relevant studies in the literature. b. Comment on the study limitations including any potential sources of bias, any limitations of the animal model, and the imprecision associated with the results². c. Describe any implications of your experimental methods or findings for the replacement, refinement or reduction (the 3Rs) of the use of animals in research. 	N/A	N/A
Generalisability/ translation	19	Comment on whether, and how, the findings of this study are likely to translate to other species or systems, including any relevance to human biology.	Page8-10/Line449-687	Discussion/Paragraph16-2 1
Funding	20	List all funding sources (including grant number) and the role of the funder(s) in the study.	Page11/Line753-754	Funding/Paragraph23

From:

Animal Research: Reporting In Vivo Experiments

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References:

1. Kilkenny C, Browne WJ, Cuthill IC, Emerson M, Altman DG (2010) Improving Bioscience Research Reporting: The ARRIVE Guidelines for Reporting Animal Research. PLoS Biol 8(6):

e1000412. doi:10.1371/journal.pbio.1000412

2. Schulz KF, Altman DG, Moher D, the CONSORT Group (2010) CONSORT 2010 Statement: updated guidelines for reporting parallel group randomised trials. BMJ 340:c332.

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*As the checklist was provided upon initial submission, the page number/line number reported may be changed due to copyediting and may not be referable in the published version. In this case, the section/paragraph may be used as an alternative reference.