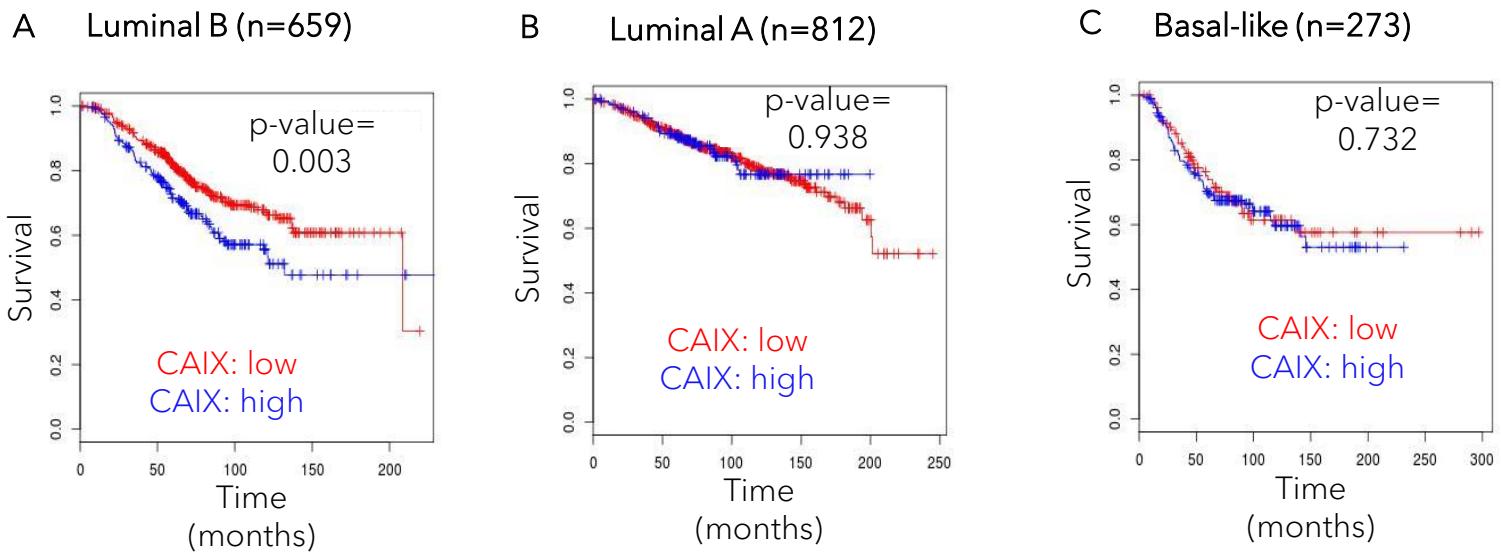
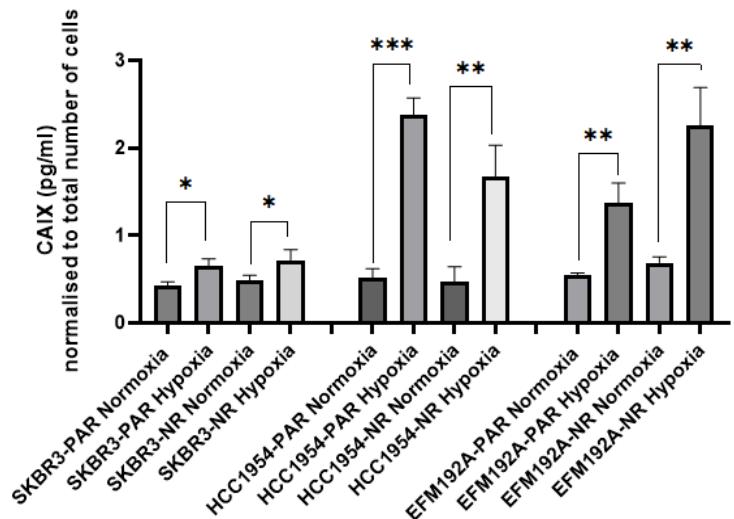


Supplemental Information

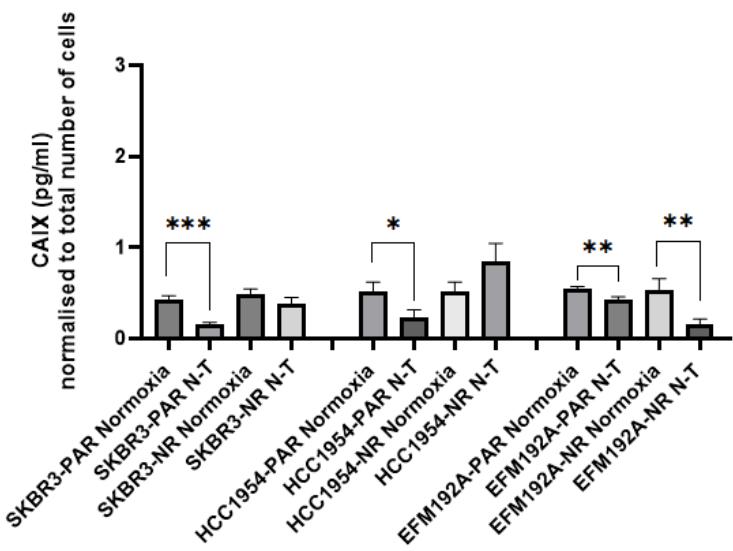


1 Supplementary Figure 1. CAIX is prognostic for poor outcome for patients with luminal B breast cancer, but is not significantly associated with outcome from either luminal A or basal-like tumours.

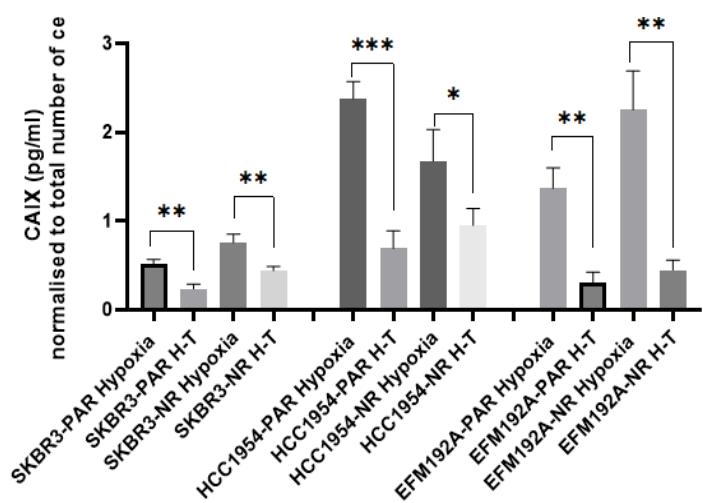
A Hypoxia-induced CAIX



B Cellular CAIX under normoxia ± S4



C Cellular CAIX under hypoxia ± S4



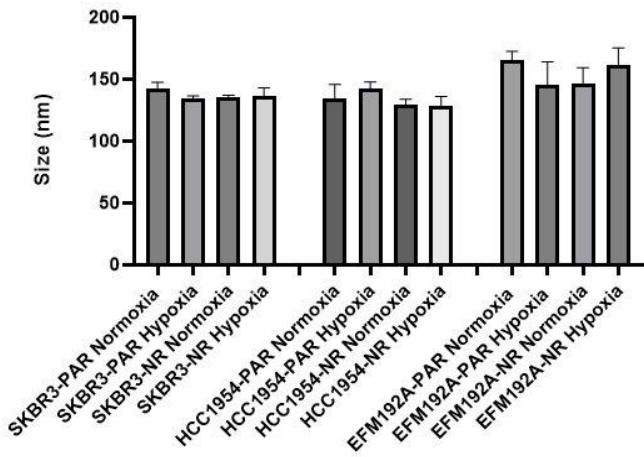
3 Supplemental Figure 2. CAIX protein expression, and the effect of S4 treatment on CAIX protein expression when cell culture 4 under normoxia or hypoxia conditions.

5 A CAIX protein expression was quantified by ELISA in whole cell lysates of SKBR3, HCC1954, EFM192A and their neratinib-6 resistant (NR) variants, following 48 hours of culture under normoxic or hypoxic conditions.

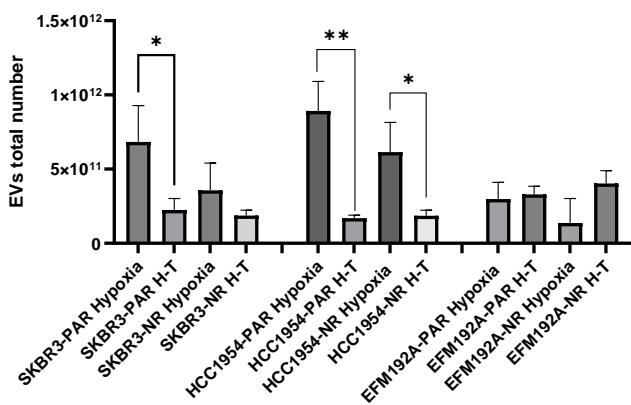
7 CAIX protein quantified with S4 (a CAIX inhibitor) of the same cell lines under **B** normoxic, or **C** hypoxic condition. Data are presented as mean \pm SEM of n=3 independent biological experiments. Statistical significance was assessed using T-tests.

9 *P < 0.05, **P < 0.01, ***P < 0.001, ****P < 0.0001. N-T = normoxia with S4 treatment; H-T = hypoxia with S4 treatment.

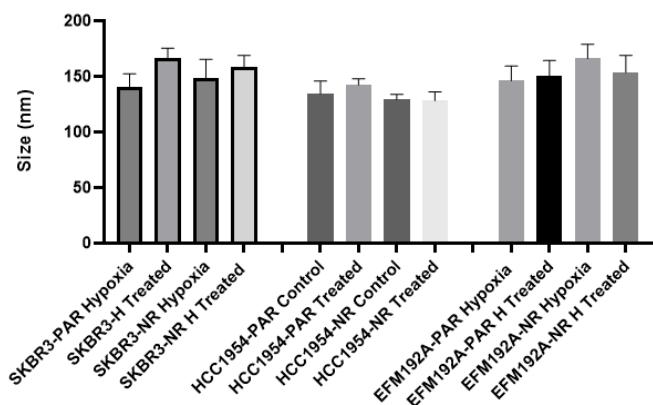
A



B



C



10 Supplemental Figure 3. Effect of S4 treatment on EVs release and size under hypoxia.

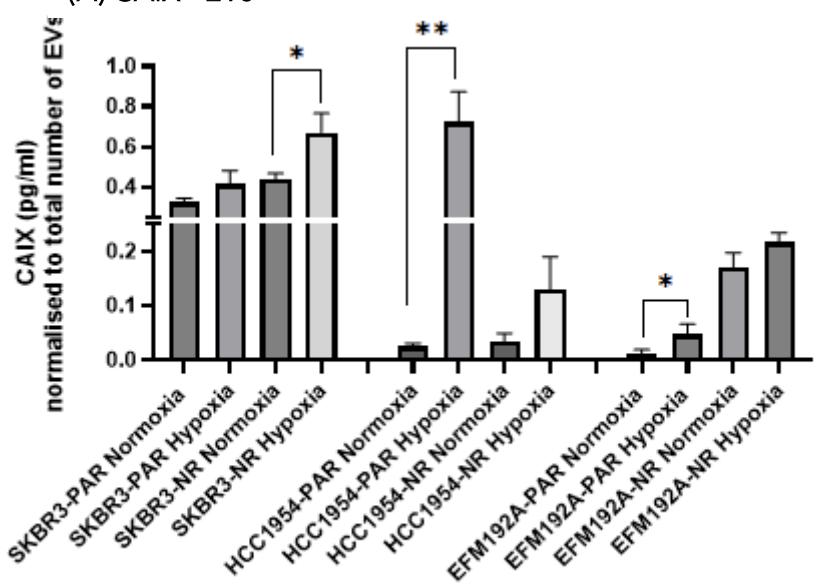
11 A. Particle size mode estimations of EVs released under both normoxic and hypoxic conditions.

12 B. The total numbers of EVs released and **C.** their sizes after treating their cells of origin with S4.

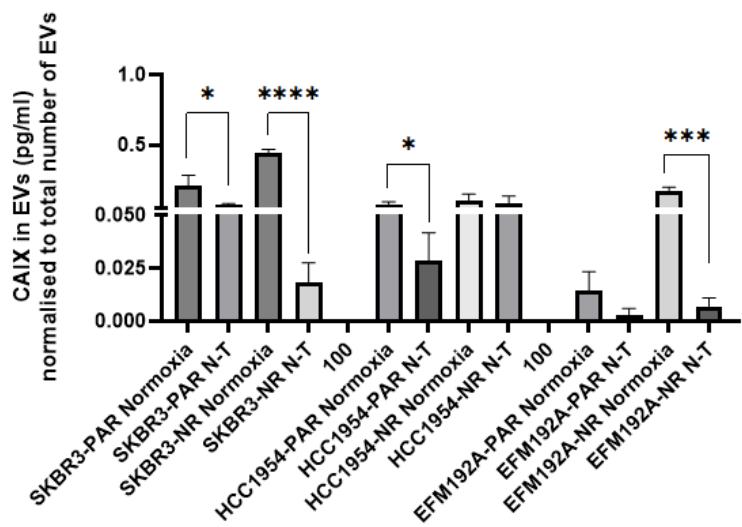
13 Data are presented mean \pm SEM of n=3 independent experiments. Statistical significance was assessed using T-tests.

14 *p<0.05, **p < 0.01. N-T = normoxia with S4 treatment; H-T = hypoxia with S4 treatment.

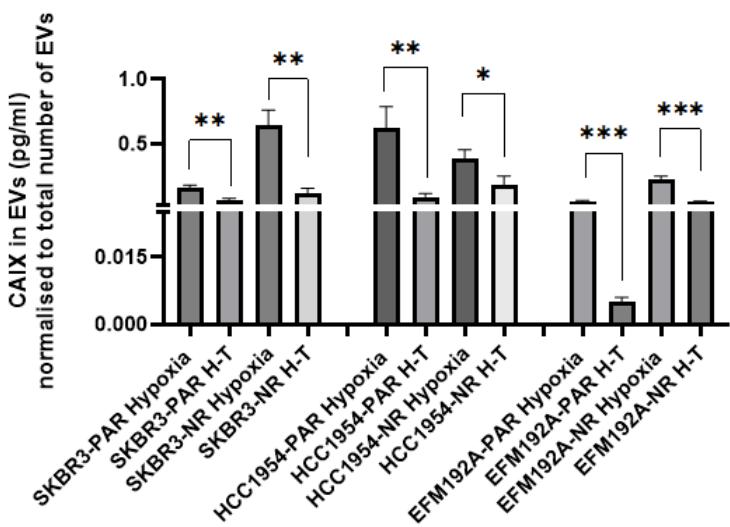
(A) CAIX - EVs



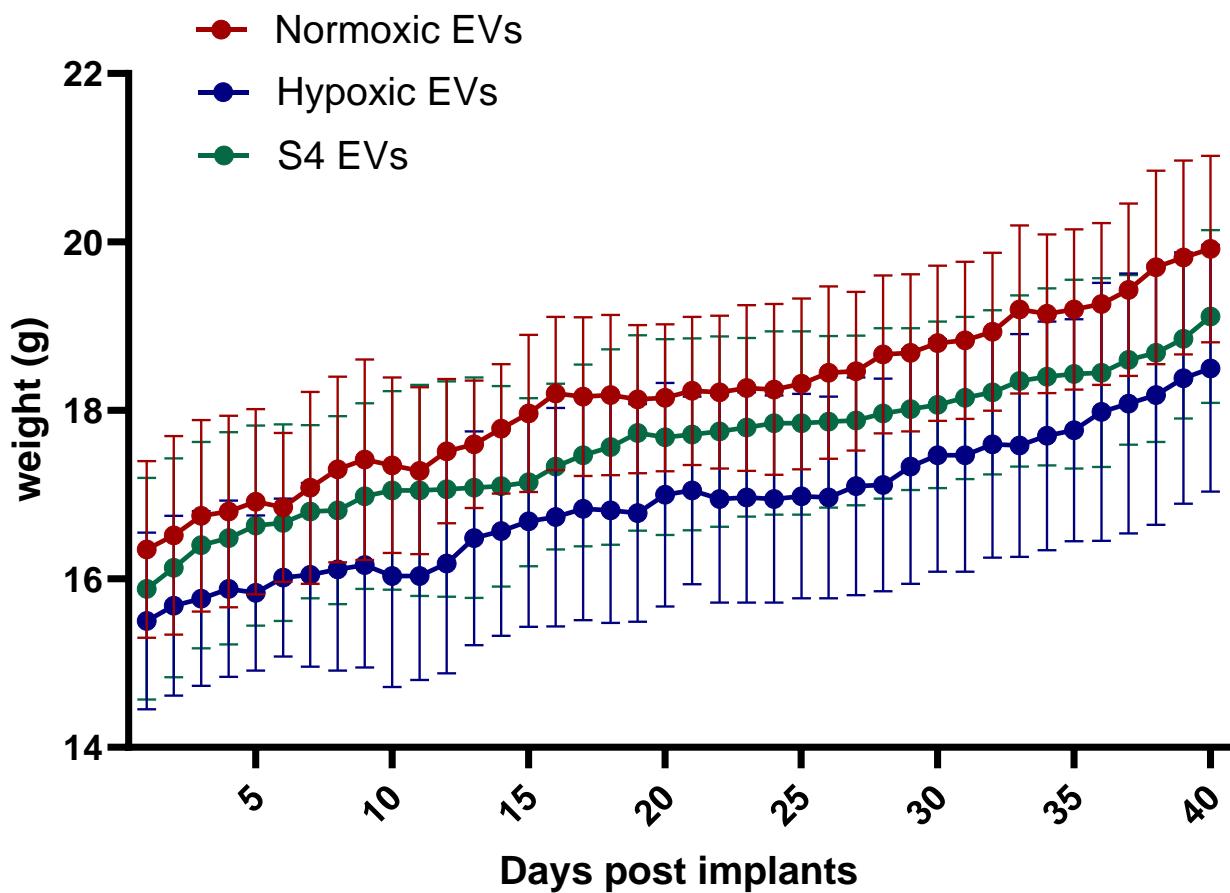
B CAIX-EVs under normoxia ± S4



C CAIX-EVs under hypoxia ± S4



15 Supplemental Figure 4. EVs CAIX cargo (CAIX-EVs). **A** CAIX protein levels was quantified by ELISA in EVs collected from 16 SKBR3, HCC1954, EFM192A and their NR variants following 48 hours of culture under normoxic or hypoxic conditions. CAIX 17 protein cargo of EVs collected from the same cell lines under **B** normoxic condition or **C** hypoxic condition after 48 hours of 18 S4 treatment were quantified by ELISA. Data are presented mean \pm SEM of n=3 independent experiments (n=3). Statistical 19 significance was assessed using T-tests. *P < 0.05, **P < 0.01, ***P < 0.001, ****P < 0.0001. N-T = normoxia with S4 20 treatment; H-T = hypoxia with S4 treatment.



21 Supplemental Figure 5. Body weight monitoring of mice following implantation of EVs pretreated HCC1954-LUC⁺ cells.

22 Mice were weighed daily for 40 days following orthotopic implantation of HCC1954-LUC⁺ cells pretreated with EVs from normoxic,

23 hypoxic, or hypoxic + S4-treated cultures. Data represent mean \pm SD for all animals (n = 6 mice per group). All groups showed progressive

24 weight gain, with no signs of systemic toxicity.

The ARRIVE Guidelines Checklist

Animal Research: Reporting In Vivo Experiments

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¹The National Centre for the Replacement, Refinement and Reduction of Animals in Research, London, UK, ²School of Veterinary Science, University of Bristol, Bristol, UK, ³School of Biological Sciences, University of Bristol, Bristol, UK, ⁴National Heart and Lung Institute, Imperial College London, UK, ⁵Centre for Statistics in Medicine, University of Oxford, Oxford, UK.

ITEM	RECOMMENDATION	Section/ Paragraph
Title	1 Provide as accurate and concise a description of the content of the article as possible.	
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.	
INTRODUCTION		
Background	3 <ol style="list-style-type: none"> 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. 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. 	
Objectives	4 Clearly describe the primary and any secondary objectives of the study, or specific hypotheses being tested.	
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.	
Study design	6 For each experiment, give brief details of the study design including: <ol style="list-style-type: none"> The number of experimental and control groups. 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). The experimental unit (e.g. a single animal, group or cage of animals). <p>A time-line diagram or flow chart can be useful to illustrate how complex study designs were carried out.</p>	
Experimental procedures	7 For each experiment and each experimental group, including controls, provide precise details of all procedures carried out. For example: <ol style="list-style-type: none"> 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). When (e.g. time of day). Where (e.g. home cage, laboratory, water maze). Why (e.g. rationale for choice of specific anaesthetic, route of administration, drug dose used). 	
Experimental animals	8 <ol style="list-style-type: none"> 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). 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. 	

Housing and husbandry	9	<p>Provide details of:</p> <ol style="list-style-type: none"> 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). 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). Welfare-related assessments and interventions that were carried out prior to, during, or after the experiment. 	
Sample size	10	<ol style="list-style-type: none"> Specify the total number of animals used in each experiment, and the number of animals in each experimental group. Explain how the number of animals was arrived at. Provide details of any sample size calculation used. Indicate the number of independent replications of each experiment, if relevant. 	
Allocating animals to experimental groups	11	<ol style="list-style-type: none"> Give full details of how animals were allocated to experimental groups, including randomisation or matching if done. Describe the order in which the animals in the different experimental groups were treated and assessed. 	
Experimental outcomes	12	Clearly define the primary and secondary experimental outcomes assessed (e.g. cell death, molecular markers, behavioural changes).	
Statistical methods	13	<ol style="list-style-type: none"> Provide details of the statistical methods used for each analysis. Specify the unit of analysis for each dataset (e.g. single animal, group of animals, single neuron). Describe any methods used to assess whether the data met the assumptions of the statistical approach. 	
RESULTS			
Baseline data	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).	
Numbers analysed	15	<ol style="list-style-type: none"> Report the number of animals in each group included in each analysis. Report absolute numbers (e.g. 10/20, not 50%). If any animals or data were not included in the analysis, explain why. 	
Outcomes and estimation	16	Report the results for each analysis carried out, with a measure of precision (e.g. standard error or confidence interval).	
Adverse events	17	<ol style="list-style-type: none"> Give details of all important adverse events in each experimental group. Describe any modifications to the experimental protocols made to reduce adverse events. 	
DISCUSSION			
Interpretation/scientific implications	18	<ol style="list-style-type: none"> Interpret the results, taking into account the study objectives and hypotheses, current theory and other relevant studies in the literature. Comment on the study limitations including any potential sources of bias, any limitations of the animal model, and the imprecision associated with the results². Describe any implications of your experimental methods or findings for the replacement, refinement or reduction (the 3Rs) of the use of animals in research. 	
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.	
Funding	20	List all funding sources (including grant number) and the role of the funder(s) in the study.	

References:

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