

Title	Adaptive responses of animals to climate change are most likely insufficient
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Give P values as exact values whenever suitable.
- ☒ ☐ For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
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Software and code

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Data collection

WebPlotDigitiser, metagear package in R software

Data analysis

R 3.5.0

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- Accession codes, unique identifiers, or web links for publicly available datasets
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The PRC dataset, containing metadata on each study as well as coefficients estimated for each step of the framework (Fig. 1) per each study is available as Supplementary Data 4. We also make available the raw data for a subset of studies for which we received the consent of data owners (4819 out of 4835 studies). These data are available as part of the R package 'adRes' (www.github.com/radchukv/adRes), which implements the complete workflow of this study and provides functions to be used to conduct similar analyses on new data in the future. The source data underlying Figs 1b-c, 1e-f, 2, 3, 4, and 5a-c are provided as a Source Data file.

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Ecological, evolutionary & environmental sciences study design

All studies must disclose on these points even when the disclosure is negative.

Study description	This study is a meta-analysis based on a systematic literature review. We addressed a question of to what extent the trait changes in animals in response to climate change are adaptive.
Research sample	We focused on adaptive responses of animals, and therefore our search targeted a wide range of animal taxa, from insects to mammals. As a result of our literature search we assembled a dataset consisting of 4835 studies (representing 1413 non-aquatic species in 23 countries). This 'Phenotypic responses to Climate' (PRC) dataset contained information on phenotypic responses to climate change. Out of these 4835 studies a subset of 71 studies (representing 17 species in 13 countries) contained all the information required to assess whether responses were adaptive and was used for the analyses for which selection differentials were necessary.
Sampling strategy	Our sample size was a result of what was available on this topic in the published literature.
Data collection	The data were collected by conducting a systematic literature review. We identified the potentially suitable papers by key words search on Web of Knowledge. In the next step, the 10090 identified abstracts were skimmed and the papers that seemed to satisfy all the necessary requirements were retained. Finally, after the full read of the subselected 180 papers, 58 of them were retained to assemble the dataset.
Timing and spatial scale	The key word search was conducted on May 23rd 2016 in Berlin and the abstract skimming took place in the following months.
Data exclusions	No data were excluded from the analysis
Reproducibility	Not applicable because this is not an experimental study
Randomization	Not applicable. We did not experiment with animals. Instead, our dataset was based on the studies that were published in the literature
Blinding	Blinding is not relevant to our study because we do not experiment with animals but collected the data from studies conducted previously and published in literature.
Did the study involve field work?	<input type="checkbox"/> Yes <input checked="" type="checkbox"/> No

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Materials & experimental systems

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<input checked="" type="checkbox"/>	<input type="checkbox"/> Palaeontology
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<input checked="" type="checkbox"/>	<input type="checkbox"/> Human research participants
<input checked="" type="checkbox"/>	<input type="checkbox"/> Clinical data

Methods

n/a	Involved in the study
<input checked="" type="checkbox"/>	<input type="checkbox"/> ChIP-seq
<input checked="" type="checkbox"/>	<input type="checkbox"/> Flow cytometry
<input checked="" type="checkbox"/>	<input type="checkbox"/> MRI-based neuroimaging