

Title: A hands-on tutorial on a modelling framework for projections of climate change impacts on health.

Authors: Ana M. Vicedo-Cabrera^a, Francesco Sera^a, Antonio Gasparri^{a,b}.

- a. Department of Public Health, Environments and Society, London School of Hygiene and Tropical Medicine, London, United Kingdom.
- b. Centre for Statistical Methodology, London School of Hygiene and Tropical Medicine, London, United Kingdom

Corresponding author:

Ana M. Vicedo-Cabrera, PhD

Department of Public Health, Environments and Society, London School of Hygiene and Tropical Medicine.

15-17 Tavistock Place, London WC1H 9SH. United Kingdom

ana.vicedo-cabrera@lshtm.ac.uk

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1 **Abstract**

2 Reliable estimates of future health impacts due to climate change are needed to inform
3 and contribute to the design of efficient adaptation and mitigation strategies. However,
4 projecting health burdens associated to specific environmental stressors is a challenging
5 task, due to the complex risk patterns and inherent uncertainty of future climate
6 scenarios. These assessments involve multi-disciplinary knowledge, requiring expertise
7 in epidemiology, statistics, and climate science, among other subjects. Here, we present
8 a methodological framework to estimate future health impacts under climate change
9 scenarios based on a defined set of assumptions and advanced statistical techniques
10 developed in time-series analysis in environmental epidemiology. The proposed
11 methodology is illustrated through a step-by-step hands-on tutorial structured in well-
12 defined sections that cover the main methodological steps and essential elements. Each
13 section provides a thorough description of each step, along with a discussion on
14 available analytical options and the rationale on the choices made in the proposed
15 framework. The illustration is complemented with a practical example of study using real-
16 world data and a series of R scripts included as Supplementary Digital Content, which
17 facilitates its replication and extension on other environmental stressors, outcomes,
18 study settings, and projection scenarios. Users should critically assess the potential
19 modelling alternatives and modify the framework and R code to adapt them to their
20 research on health impact projections.

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33 **Background**

34 Climate change is one of the most important environmental challenges that humanity will
35 face in the coming decades. Quantifying future health burdens associated to global
36 warming is therefore a major priority for the scientific community, as attested by the
37 increasing number of publications on health impact projections. Several studies have
38 focused on direct impacts of environmental stressors, such as non-optimal temperature
39 and air pollution.¹⁻⁵ Generally, these projection studies follow a common methodological
40 scheme. The basic idea consists in applying risk functions on simulated future exposure
41 distributions generated by climate change models under specific emissions scenarios.
42 However, this scheme entails important methodological challenges due, for instance, to
43 the complex patterns of health risks associated with environmental stressors, the
44 inherent uncertainty of potential future climate change processes, and the set of (rarely
45 stated) assumptions.⁶ A wide variety of data sources, statistical approaches and
46 assumptions have been applied so far, as summarized and discussed in previous
47 reviews.⁶⁻⁸ However, a structured illustration that covers the important steps and discuss
48 the most recent statistical developments is still lacking.

49 Here, we illustrate a methodological framework to estimate health impact projections
50 under climate change scenarios, built on clearly defined assumptions and state-of-the-
51 art statistical methodologies developed in time-series analysis in environmental
52 epidemiology. This contribution extends a methodology previously presented to project
53 temperature-related excess mortality in climate change scenarios.^{5,9} The proposed
54 framework is illustrated through a hands-on tutorial, structured in well-differentiated steps
55 that cover each of the methodological issues and the essential elements. Each section
56 provides a detailed description of the methodology and a discussion on the potential
57 assumptions and limitations, compared to other available choices. The text is
58 complemented with a practical illustration of a projection study using real-world data, and
59 a series of R scripts included as Supplementary Digital Content, with updated versions
60 available in the personal website and GitHub repository of the last author. The
61 methodological framework and R code can be modified and adapted to a broad range of
62 health impact projection studies, optionally assessing different environmental stressors
63 and health outcomes, and with different study settings.

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65 **Illustrative example**

66 The practical example consists of a projection study on temperature-related mortality
67 impacts in the city of London, United Kingdom. The dataset includes observed daily

68 mean temperature and total number of deaths in London between 1990 and 2012. This
69 is part of the large database collected within the Multi-City Multi-Country (MCC) network
70 (<http://mccstudy.lshhtm.ac.uk/>), and has been previously used as example in other
71 manuscripts.¹⁰ We complement these observed data with daily-modelled temperature
72 series for historical (1950-2005) and future (2006-2100) periods, projected under
73 scenarios defined within the Coupled Model Intercomparison Project Phase 5 of
74 Intergovernmental Panel on Climate Change (IPCC).¹¹ Climate data was obtained,
75 processed and made available by the Inter-Sectoral Impact Model Intercomparison
76 Project (ISI-MIP, <https://www.isimip.org/>).¹² Further details on the modelled data is
77 provided in the Section 2 of the tutorial.

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79 **Tutorial on the modelling framework**

80 1. Estimation of exposure-response associations

81 One critical step in health impact projection studies is to appropriately define the
82 relationship between the exposure to the environmental stressor of interest and the
83 health outcome. While this information can be based on association estimates reported
84 in the literature,^{13,14} this often requires strong assumptions due to extrapolation across
85 geographical areas, and simplification of usually complex relationships.

86 A more appropriate approach is to directly estimate the relationship using actual
87 epidemiological data, for which several statistical methods are available.^{15,16} Among
88 these, time series analysis using aggregated data has been shown to be ideal to assess
89 short-term associations in environmental epidemiology,¹⁷ and often applied in climate
90 change projection studies.^{1,18,19}

91 A representation of the standard time series regression model is provided by the
92 following equation:

$$93 \quad \log[E(Y_t)] = \alpha + f(x_t; \boldsymbol{\theta}) + s(t; \boldsymbol{\beta}) + \sum_{p=1}^P h_p(z_{pt}; \boldsymbol{\gamma}_p) \quad (1)$$

94 where typically the outcome Y_t corresponds to daily counts assumed to follow a Poisson
95 distribution with overdispersion, the function $f(x_t; \boldsymbol{\theta})$ specifies the association with the
96 environmental exposure of interest x at time t , $s(t; \boldsymbol{\beta})$ represents the baseline trend
97 which captures the effect of confounders changing slowly over time (i.e., seasonal and
98 long-term trends), and $h_p(z_{pt}; \boldsymbol{\gamma}_p)$ models the contribution of other confounders varying
99 on a daily basis.

100 The exposure-response association can be modelled using different types of function f ,
101 ranging from simple indicators for extreme exposure events, to linear or linear-threshold

102 shapes, to distributed lag non-linear models (DLNMs) representing complex exposure-
103 lag-response surfaces.²⁰ The selection of the function depends on the environmental
104 stressor, for instance measured as a continuous exposure (e.g., temperature, rain fall)
105 or defined extreme event (e.g., heat wave, floods), and the assumed dependency with
106 the health outcome. As shown below, wrong assumptions on the shape of the
107 dependency can introduce important biases in estimates and projections.

108 In our example, the environmental stressor and the outcome corresponds to historical
109 series of daily mean temperature and death counts (T_{obs} and D_{obs}). Our main choice for
110 the exposure-response function $f(x_t)$ is represented by a DLNM through a bi-
111 dimensional cross-basis term, using flexible natural cubic spline functions to model both
112 exposure-response and lagged-response dimensions, accounting for 21 days of lag,
113 following previous work.¹⁰ As further described in Section 4 of this tutorial, the choice of
114 natural splines allows the log-linear extrapolation of the function beyond the boundaries
115 of the observed series, a step needed to project the risk using the modelled temperature.
116 Figure 1A shows the resulting 3-D plot of the estimated exposure-lag-response
117 association, and Figure 1B represents the overall cumulative exposure-response
118 association across up to 21 days of lag. As expected, we observe a non-linear
119 temperature-mortality relationship, with increases in relative risk (RR) above and below
120 the minimum mortality temperature (T_{mm}) that correspond to heat and cold associations,
121 respectively. At the same time, risks are distributed differently across time, with
122 immediate heat-mortality and more delayed cold-mortality associations (Figure 1A).

123 Alternative models with different specifications of the exposure-response association,
124 such as linear or double-threshold parameterizations, are shown in Figure 1C. While
125 simpler, however these choices seem less ideal for modelling the mortality risk of non-
126 optimal temperature, highlighting the importance of the selection of suitable functions to
127 represent the association of interest, and the potential bias of inappropriate
128 simplifications.

129

130 2. Projected temperature and mortality series

131 Two additional essential elements needed in health impact projection studies are the
132 information on future climatic and population scenarios.

133 Data on future distribution of the environmental stressor (e.g., temperature, precipitation,
134 air pollution levels) are commonly based on specific scenarios that account for changes
135 in multiple and often inter-related factors. For instance, socioeconomic and technological
136 changes, population growth and land use changes can affect pathways of greenhouse

137 gases emissions or atmospheric concentrations of other pollutants, which in turn will
138 determine trends in global warming and potential levels of specific environmental
139 exposures.²¹ Under each scenario, these trends can be generated from general
140 circulation models (GCMs), which offer projections of future conditions based on specific
141 and simplified assumptions.²¹ To have a better representation of future trends, the usual
142 approach is to combine impact estimates obtained either using more than one model per
143 scenario or using ensemble members output from multiple runs of the same climate
144 model, but with different initial conditions.^{6,7}

145 In our worked example, we applied the first approach by including modelled temperature
146 data from 5 different GCMs for two climate change scenarios, defined as representative
147 concentration pathways 4.5 and 8.5 (RCP4.5 and RCP8.5).^{22,23} Figure 2 shows the
148 temporal trends in temperature for the historical (1971-2005) and future (2006-2100)
149 periods projected in London under the two scenarios, depicted as GCM-ensemble
150 averages (solid lines) and associated variability (shaded areas). As discussed later in
151 Section 6, the availability of exposure trends from multiple models can be used to
152 determine the related uncertainty of the projected health impacts.

153 Projection exercises also depend on representations of future mortality trends,
154 determined by the demographic structure and outcome baseline rates. Data on these
155 population scenarios can be built following different approaches based on the adopted
156 assumptions. The simplest procedure consists in assuming that populations and
157 outcome rates will remain constant in the future, thus isolating the climate effect from
158 other important trends.²⁴⁻²⁶ However, other studies relied on population projections
159 derived from predictive models under varying levels of future fertility, mortality and
160 migration,²⁷⁻²⁹ a procedure that requires additional assumptions.

161 In our example, we illustrate an application of the former method. First, we compute an
162 annual series of total mortality counts as the average for each day of the year from
163 observed daily deaths, thus keeping into account the seasonal structure of the observed
164 mortality series (Figure 3). The annual series is then replicated along the whole
165 projection period. The extension to more complex scenarios requires the derivation of
166 age-specific mortality series, obtained using projection methods that model changes in
167 the demographic structure and baseline rates, as further explained in Section 7 of this
168 tutorial.

169

170 3. Downscaling and calibration

171 Climate simulations of historical periods usually show systematic deviations from the
172 real-world observations. This can be explained by real differences due to the different
173 geographical resolution of the data (gridded versus point-source), or to biases due to
174 poor performance of climate models, occurring in areas with sparse information from
175 meteorological stations. These deviations should be carefully considered in climate
176 change projection studies, as the predicted impacts will depend on the alignment of
177 observed and modelled series.^{30,31} Corrections of biases related to these two aspects
178 have been defined separately as downscaling and calibration, although in most cases
179 they rely on similar analytic procedures. Downscaling refers to the process of obtaining
180 location-specific climate information from global or regional models that provide data at
181 a larger geographical resolution, and is based on either dynamical or statistical methods.⁷
182 Conversely, calibration is a more general concept of re-aligning two series of data, in this
183 case observed and modelled series.

184 Bias correction methods have been proposed for both statistical downscaling and
185 calibration, and encompass various different techniques with varying degree of
186 complexity, ranging from basic statistical approaches (i.e., use of additive or
187 multiplicative corrections, shifted distribution), to more complex statistical procedures.³¹
188 However, limited evidence exists about the potential impact of the choice of method on
189 the estimated projections.

190 In the present tutorial, the model outputs from the GCMs are firstly downscaled through
191 bi-linear interpolation at a $0.5^\circ \times 0.5^\circ$ spatial resolution and linear interpolated by day of
192 the year. The resulting series are then calibrated with the observed data using the bias-
193 correction method developed within ISI-MIP.³² This ensures that the trend and variability
194 of the original data are preserved by adjusting the cumulative distribution of the simulated
195 data to the observed one. In detail, the monthly variability and mean are corrected only
196 using a constant offset or multiplicative correction factor that corrects for long-term
197 differences between the simulated and observed monthly mean data in the historical
198 period.³² Figure 4 shows a comparison between the modelled series from a specific GCM
199 (T_{mod} , green area and line), and the observed series (T_{obs} , black area and line), in terms
200 of their overall and cumulative distribution (left and right panels, respectively). It can be
201 noted that the modelled series is shifted towards colder ranges, likely for the reasons
202 mentioned above. As discussed, this would create a bias in the future projections. The
203 bias-correction procedure described above calibrates the modelled series (T_{mod}^* , green
204 dashed line), re-aligning it to the observed one (Figure 4, right panel).

205

206 4. Extrapolation of exposure-response curves

207 Risk estimates obtained over historical periods do not automatically apply to future
208 scenarios, due to several reasons. For instance, it is possible that the estimated
209 exposure-response association will be different in the future, due to for example
210 adaptation or changes in vulnerability of the population. However, even when assuming
211 no changes in risk, the future distribution of a specific environmental stressor is likely to
212 be different to that observed in the present days, and can extend further than the region
213 of the estimated exposure-response curve. Thus, we need to perform an additional step
214 consisting in the extrapolation of the exposure-response beyond the observed
215 boundaries. This, however, implies the adoption of additional assumptions on the
216 hypothetical shape of the association over the unobserved range.

217 As shown in Figure 5 (top panel), a viable method is based on a log-linear extrapolation
218 of the curve beyond the observed boundaries. The use of a natural cubic spline function
219 to model the exposure-response dimension ensures this non-linear extrapolation,
220 although this step can be more problematic when applying different functions.
221 Nonetheless, this entails a series of strong assumptions on the future risk associated to
222 environmental factors. The first assumption, mentioned above, is that the exposure-
223 response association estimated on the currently observed range will not change in the
224 future, for instance as a result of changes in susceptibility of the population, as discussed
225 in Section 7. The second assumption is that the extrapolation represents appropriately
226 the risk over the unobserved range. In addition, due to the nature of the epidemiological
227 approaches, the extrapolation of the curve over un-observed ranges constitutes an
228 important source of uncertainty to our projection estimates. This last issue will be further
229 described in Section 6.

230

231 5. Projection and quantification of the impact

232 The next step of the proposed analytical framework consists in estimating the projected
233 health impacts estimates by applying the exposure-response association estimates over
234 the modelled series of the specific environmental stressor and outcome. Previous studies
235 reported measures of impact using various measures, for instance in terms of percent
236 changes in the rate of the outcome, excess mortality or morbidity, or attributable
237 fractions.^{5,18,33} Our framework incorporates the procedure previously developed to
238 estimate the impacts in terms of attributable fractions within in time series analysis,
239 applicable either with the DLNM framework or with simpler exposure-response
240 dependencies.³⁴

241 In brief, the method consists in computing for each day of the series the number of cases
 242 attributed to a specific environmental stressor based on the estimated risk and the level
 243 of exposure in that specific day. Then daily attributable numbers are aggregated by
 244 defined intervals of time in the future period. It can be also expressed in terms of
 245 attributable fraction computed as the ratio with the corresponding total number of cases.
 246 Finally, projection studies are mostly interested in obtaining comparative measures of
 247 impact between climate change scenarios or timeframes, which can be easily computed
 248 as differences in attributable numbers or fractions.

249 In the specific setting of the example of study, we estimate the attributable number of
 250 deaths D_{attr} due to non-optimal temperatures using the calibrated temperature series
 251 T_{mod}^* following:

$$252 \quad D_{attr} = D \cdot \left(1 - e^{-\left(f^*(T_{mod}^*; \theta_b^*) - s^*(T_{mm}; \theta_b^*) \right)} \right) \quad (2)$$

253

254 where f^* and θ^* represents the uni-dimensional overall cumulative exposure-response
 255 curves with reduced lag dimension, derived from the bi-dimensional term estimated in
 256 Section 1 of the tutorial. In Eq.2, we can also separate components due to heat and cold
 257 by summing the subsets corresponding to days with temperatures higher or lower than
 258 T_{mm} .¹⁰ The same computation can be used with simpler exposure-response functions,
 259 and the equation simplifies to the usual (RR-1)/RR in the case of linear or binary
 260 unlagged relationships.

261 The selection of the T_{mm} is a critical step in the quantification of the attributable mortality.
 262 While this step has been shown to have little impact in well-powered multi-location
 263 studies relying on best linear unbiased predictions, this choice can be problematic in
 264 single-location analyses that can be affected by highly imprecise exposure-response
 265 curves.^{10,35}

266 Figure 5 (mid and bottom panels) shows the distributions of temperatures and estimated
 267 attributable mortality, respectively, for the historic and future period in London under the
 268 assumption of stable populations and no changes in vulnerability. We can observe that
 269 the mortality burden due to cold temperatures is currently much larger than for heat,
 270 especially across the moderate cold temperatures. However, if we compare the
 271 estimates between each of the two periods, we can see that heat-attributable mortality
 272 will substantially increase in the future by 4.0% (95% empirical confidence interval (eCI):
 273 0.7-6.8), while mortality due to cold will be reduced by 3.3% (95% eCI: 4.3-1.9). A
 274 description on the computation of the eCI is provided in the following section. The same

275 methodological procedure can be applied to derive attributable mortality for more
276 complex scenarios, as illustrated in Section 7.

277

278 6. Ensemble estimates and quantification of uncertainty

279 A key methodological issue in projection studies is to properly identify and deal with the
280 different sources of uncertainty involved in the projection of impacts in future scenarios.
281 These include those related to purely statistical aspects, such as the imprecision of the
282 estimated exposure-response function, and the inherent uncertainty of the exposure
283 simulations obtained from the climate and circulation models.⁶

284 Based on the proposed framework, uncertainty arises mainly from two main sources: the
285 estimation of the exposure-response function, especially regarding the range over which
286 we extrapolated the curve, and climate projections. These are represented by the
287 covariance matrix $V(\theta_b)$ of the model coefficients estimated in Equation 1 defining the
288 exposure-response function, and the variability of the modelled series generated in each
289 GCM (Figure 2), respectively. In the tutorial, we quantify this uncertainty by generating
290 1000 samples of the coefficients through Monte Carlo simulations, assuming a
291 multivariate normal distribution for the estimated spline model coefficients, and then
292 generating results for each of the five GCMs.³⁴ We report the results as point estimates,
293 using the average across climate models (GCM-ensemble) obtained by the estimated
294 coefficients, and as eCI, defined as the 2.5th and 97.5th percentiles of the empirical
295 distribution of the attributable mortality across coefficients samples and GCMs. These
296 eCIs account for both sources of uncertainty.

297 As briefly mentioned before, we did not account for additional uncertainty derived from
298 the estimation of T_{mm} . If desired, it is possible to quantify it using probabilistic methods
299 showed in recent publications.^{35,36} Likewise, other sources of uncertainty can arise in
300 more complex projection scenarios, such as those assuming changes in vulnerability
301 (adaptation) and population structure. However, these can be more difficult to integrate
302 quantitatively in the overall estimate of uncertainty.

303

304 7. Accounting for complex scenarios: demographic changes and adaptation

305 The example illustrated so far is built under the assumptions of no-adaptation and stable
306 populations. Findings from this exercise can answer the question: “What will the
307 temperature-related impact be in the future if the current population would be exposed
308 to warmer temperatures projected in the future?”. However, there is a growing interest in

309 assessing environmental impacts under more complex scenarios that account for
310 changes in both future risks and baseline population, which could *a priori* approximate
311 more realistically future health impacts. This additional section aims at describing these
312 potential extensions.

313 As mentioned before in the Section 2 of the tutorial, changes in size and population
314 structure may have a strong influence on future health impacts, both by increasing the
315 population at risk and by shifting it toward more susceptible groups with higher
316 associated risks. Some studies have accounted for this using age-specific risks and
317 outcome rates derived from socio-economic trajectories,^{18,19,27,37} defined for example in
318 the so-called Shared Socio-economic Pathways (SSP).³⁸ This can be incorporated in this
319 framework by replicating the proposed procedure by each age category. This step
320 requires the estimation of age-specific exposure-response associations, as shown in
321 Figure 6A, and their application over the corresponding future age-specific outcome
322 series built under a specific SSP. These modelled outcome series can be derived by re-
323 scaling the observed seasonal counts in the current period using age-specific baseline
324 populations and rates projected in the future under a specific SSP. However, it should
325 be noted that, while the “stable populations” approach is built on simplistic assumptions
326 and cannot provide a realistic representation of future excess burdens, it offers a more
327 straightforward interpretation as it separates the impact of global warming from other
328 changes, such as those related to demographic variations, that would occur anyway
329 even in a stable climate.

330 Another important issue to be considered in health projection studies is the potential
331 changes in susceptibility to specific environmental stressors. For example, evidence
332 obtained so far indicates that populations have partly adapted to heat stress in the last
333 decades, with related risks showing an attenuation along this period.³⁹ Under these
334 assumptions, exposure-response associations obtained on historical data would not be
335 representative of future risks, and several methods have been proposed to address this
336 issue. These include the analogue city approach,^{14,40} which makes use of exposure-
337 response estimates from a location with a climate similar to that projected in the future,
338 or methods that allows direct changes in the estimated exposure-response function^{41–44}
339 Both approaches can be incorporated into the proposed framework by replacing or
340 modifying the estimated exposure-response function. As an illustrative example, Figure
341 6B shows the modified temperature-mortality curve for London, assuming a decrease in
342 30% in the mortality log-RR associated with heat only, obtained by applying a scaling
343 factor to the related part of the curve. However, one should take into account that this
344 approach, while potentially more realistic, often implies simplistic assumptions on the

345 form of the future exposure-response shape and its changes due to adaptation (e.g.,
346 linear-threshold shapes, or shifts). In addition, while few studies have used empirical
347 evidence from historical data,⁴³ most of them have defined an arbitrary set of parameters
348 to model the extent and timing of adaptation mechanisms.⁴² A recent publication has
349 discussed problems and limitations of existing methods for modelling adaptation, also
350 showing how the choice greatly influences the estimated health impacts, and discussing
351 the difficulties in defining and quantifying valid adaptation mechanisms.⁴⁵ Thus, further
352 implications on the potential limitations of the applied method should be considered and
353 clearly discussed when assuming hypothetical changes in vulnerability.

354

355 **Overview and final remarks**

356 In this contribution, we have presented a well-structured and flexible methodological
357 framework, based on cutting-edge statistical techniques and clearly defined
358 assumptions, to obtain health impact projections under climate change scenarios of
359 variable complexity. Shaped as a hands-on tutorial, this article describes the key
360 methodological steps through a practical example of an applied analysis, complemented
361 with real data and R code. While the analytical approaches described in the example are
362 tailored to the specific study settings, and should not be uncritically applied in a 'cut-and-
363 paste' approach, this tutorial offers the reader the opportunity to advance through general
364 methodological steps, following how different statistical choices and assumptions have
365 been translated in the analysis and code. At the same time, it enables the reader to
366 replicate, adapt and potentially extend the proposed modelling framework by applying
367 alternative modelling choices using other environmental stressors, outcomes, study
368 settings, and more complex climate change scenarios. In a more general context, this
369 tutorial highlights the need of multi-disciplinary knowledge and skills for projecting health
370 impacts under climate change scenarios, involving experts working in different research
371 areas, such as epidemiology, statistics, and climate science, among other subjects. This
372 contribution clearly advocates for collaborative research and emphasizes the benefits of
373 reproducibility and transparency in science.

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543 **Figure legends**

544 **Figure 1. Temperature-related mortality in London (1990-2012).** Left panel: three-
545 dimensional plot showing the estimated exposure-lag-response association between
546 temperature and mortality. Mid panel: overall cumulative mortality risk (and 95%
547 confidence interval). Right panel: comparison between the exposure-response shapes
548 estimated using three modelling approaches.

549 **Figure 2. Temporal trends in projected temperature in London (1971 - 2099).** Solid
550 lines correspond to the mean annual temperature estimated across the 5 GCMs-specific
551 modelled series. The shaded area shows its variability, corresponding to the range for
552 each year. The two horizontal bars in the right correspond to the average annual
553 maximum and minimum for each modelled temperature series.

554 **Figure 3. Seasonal mortality trends in London.** Grey dots correspond to the observed
555 daily mortality counts registered in each day of the year between 1990 and 2012. The
556 blue line depicts the mean number of deaths per day of the year.

557 **Figure 4. Bias-correction of the modelled temperature series.** Comparison between
558 the distribution (left panel) and cumulative distribution (right panel) of the raw and bias-
559 corrected modelled temperature (T_{mod} , T_{mod}^*), and the observed temperature series
560 (T_{obs}).

561 **Figure 5. Temperature and excess mortality in London for present and future**
562 **periods.** Top panel: exposure-response curve represented as mortality relative risk (RR)
563 across the temperature (°C) range, with 95% empirical confidence intervals (grey area).
564 The dotted vertical line corresponds to the minimum mortality temperature (T_{mm}) used
565 as reference, which defines the two portions of the curve related to cold and heat (blue
566 and red, respectively). The dashed part of the curve represents the extrapolation beyond
567 the maximum temperature observed in 2010-19 (dashed vertical line). Mid panel:
568 distribution of T_{mod}^* for the current (2010-19, grey area) and at the end of the century
569 (2090-99, green area), projected using a specific climate model (NorESM1-M) and
570 scenario (RCP8.5). Bottom panel: the related distribution of excess mortality, expressed
571 as the fraction of additional deaths (%) attributed to non-optimal temperature compared
572 with T_{mm} .

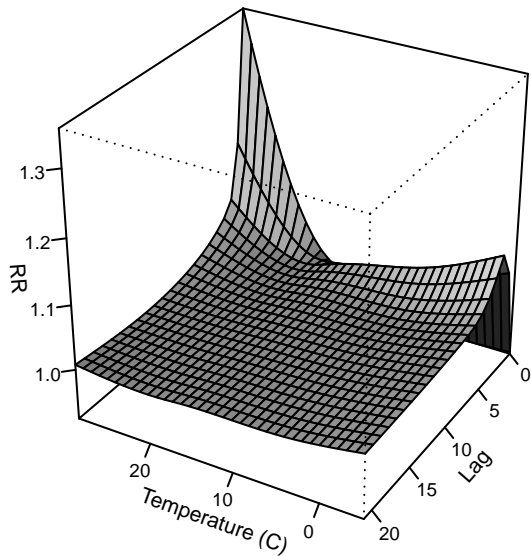
573 **Figure 6. Accounting for complex scenarios accounting for socio-demographic**
574 **changes and adaptation.** Right panel: age-specific exposure-response curves,
575 applicable to project health impact separately for each age category, thus potentially
576 accounting for demographic changes by using differential baseline mortality trends. Left
577 panel: comparison between the exposure-response curves under scenarios of no

578 adaptation (continuous line) and adaptation (dashed line), the latter under the (simplistic)
579 assumption of an hypothetical attenuation of 30% in risk associated to heat.

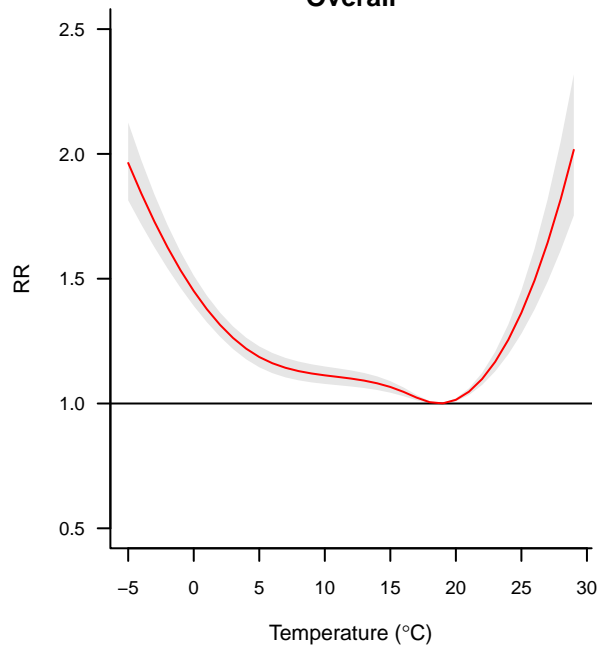
London

1990–2012

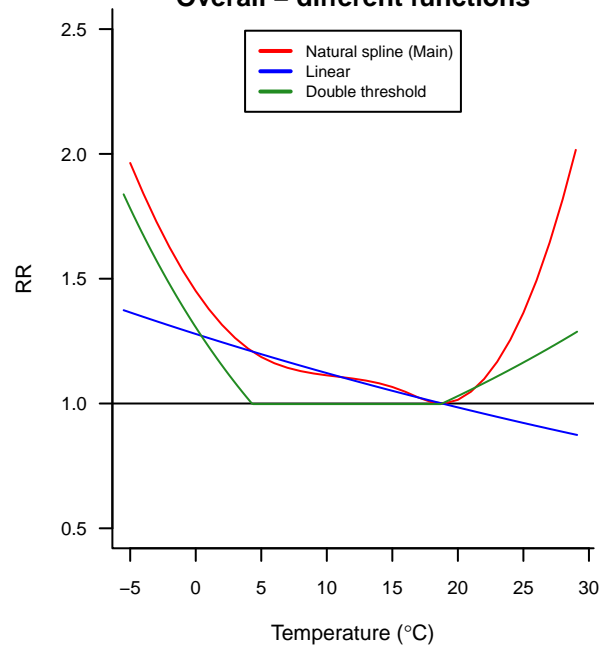
Exposure-lag-response



Overall

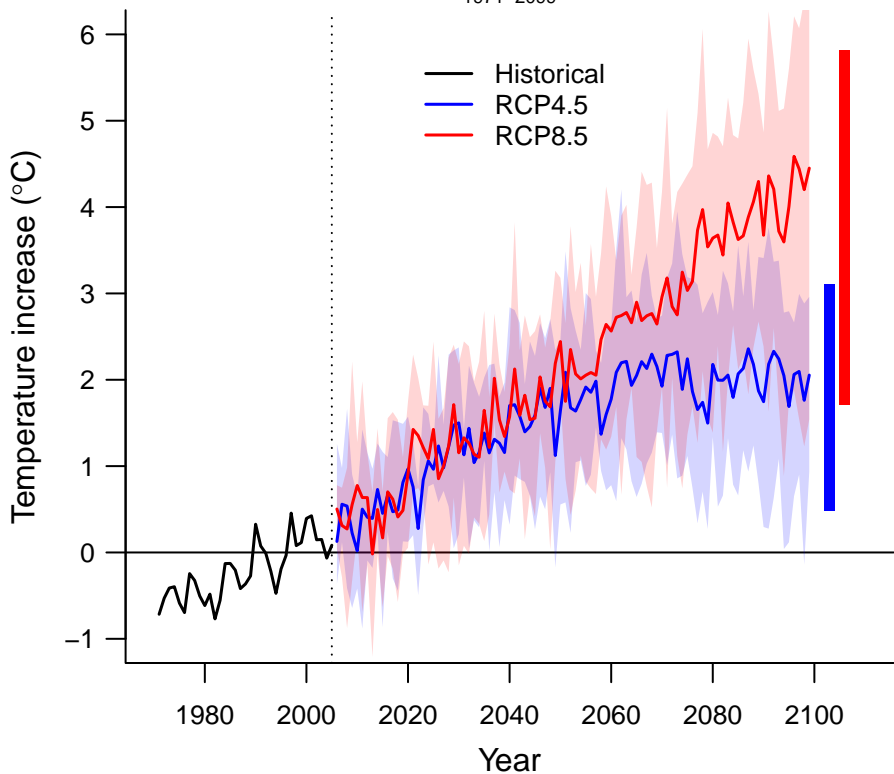


Overall – different functions



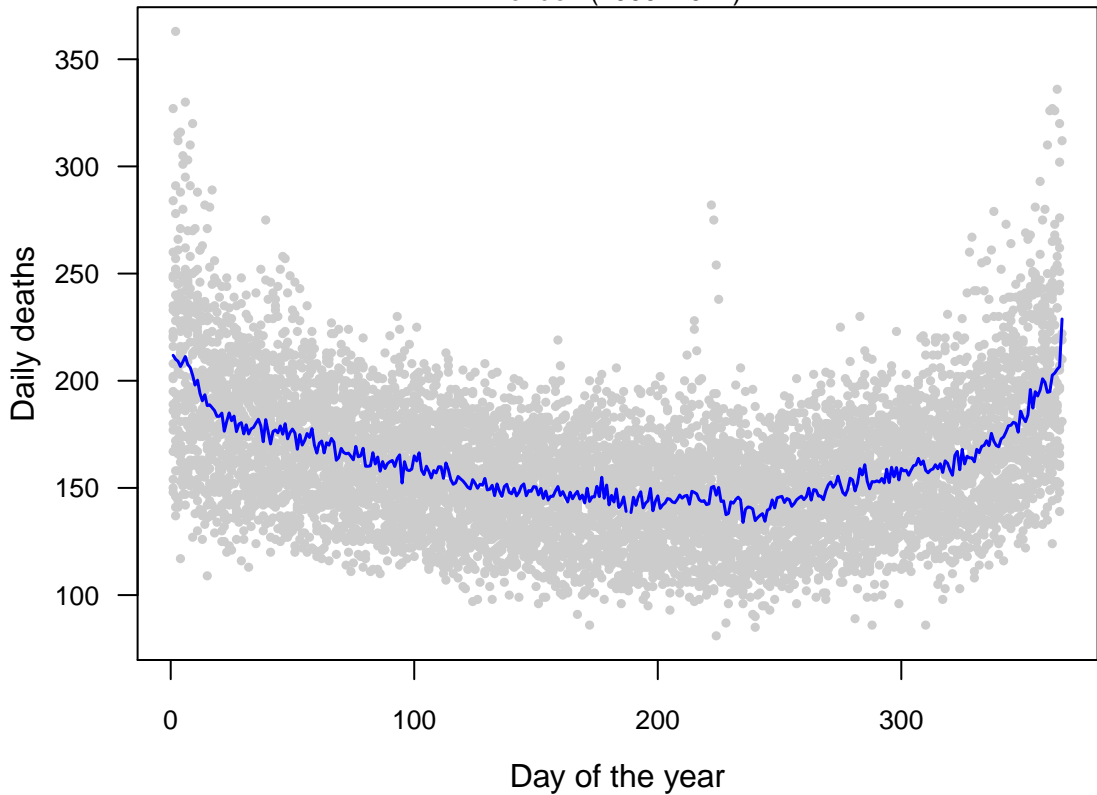
London

1971–2099



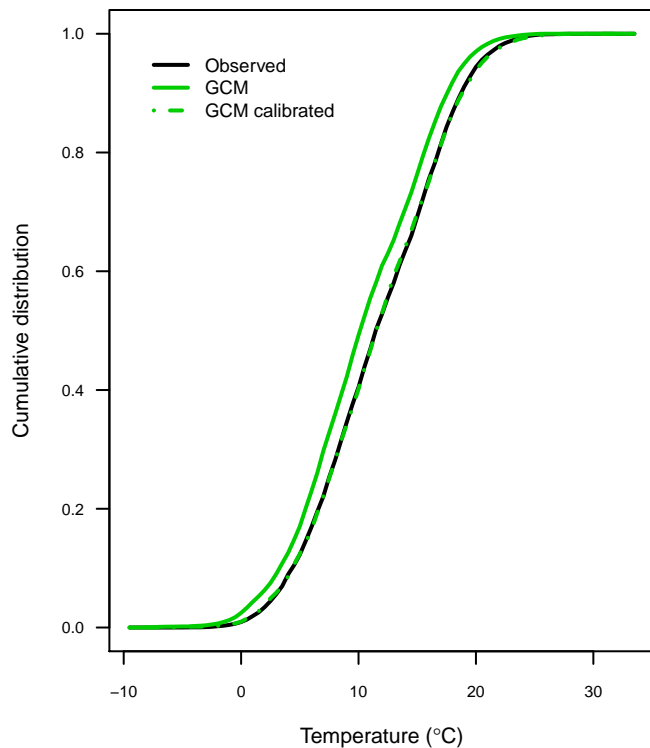
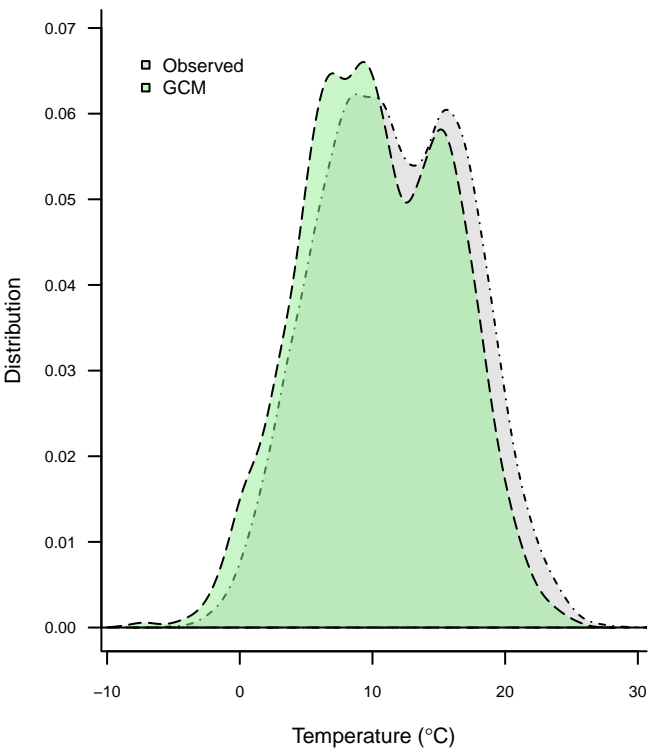
Observed Mortality

London (1990–2012)



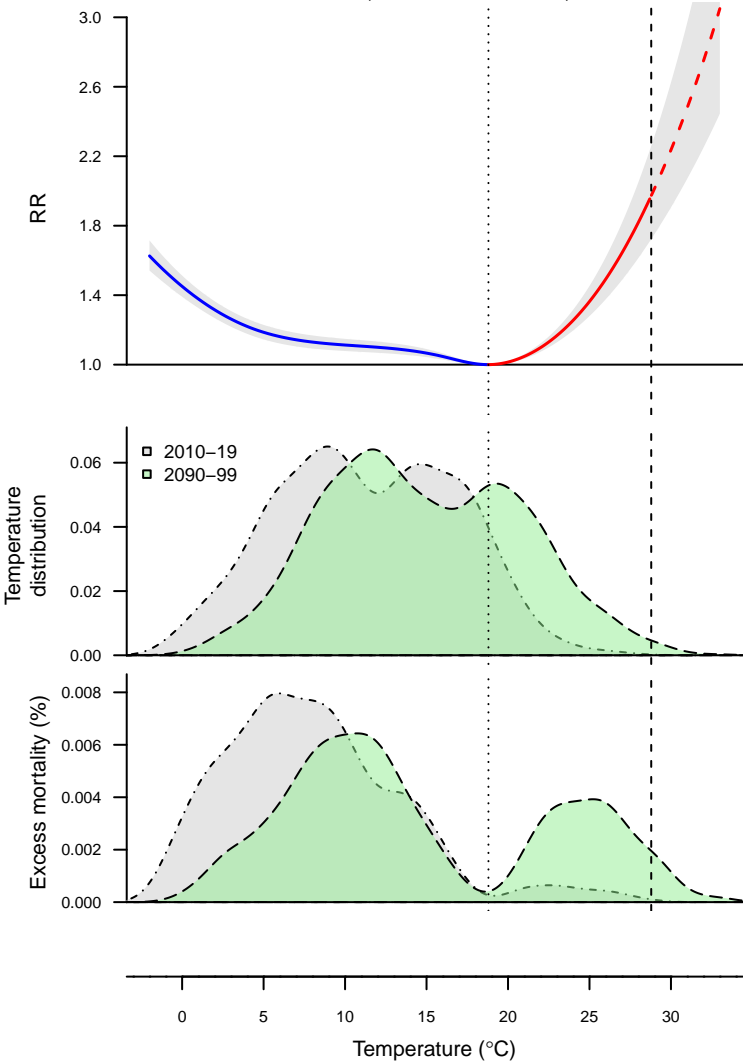
Calibration

London (1990–2012)

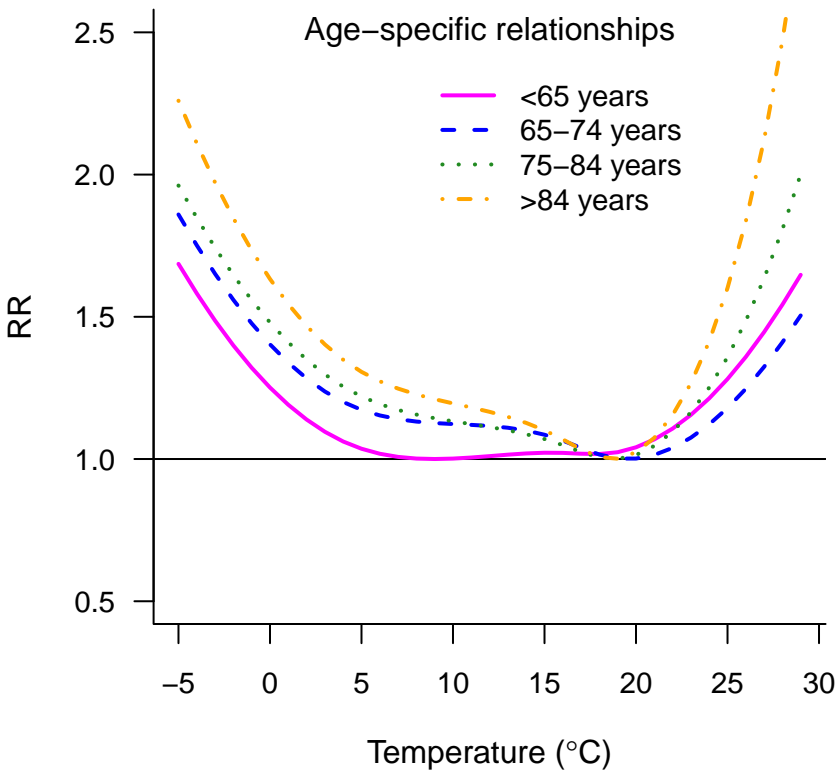


Temperature-related mortality impacts

London (NorESM1-M – RCP8.5)



Demographic changes



Adaptation

