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1 **Clarity or confusion? – problems in attributing large-scale ecological changes to**
2 **anthropogenic drivers**

3

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14 **Abstract**

15 Ways of reducing the drivers of global biodiversity loss and degradation of ecosystem

16 services are needed more than ever before. Policy options must be based on the best

17 evidence of the role of multiple driving forces. Increasingly, a significant part of the

18 evidence base comes from attributing signals of biological change detected in large-

19 scale analytical surveys to a range of possible causal factors. We highlight a number

20 of subtle difficulties that can beset the challenge of detecting such correlative

21 relationships. These are as follows: 1. The Modifiable Area Unit Problem. 2.

22 Incomplete explanatory variable data. 3. Lack of control over the replication and

23 crossing of driving variables. In most cases these problems can be avoided by more

24 careful specification of the scientific question and application of relatively new

25 analytical techniques. Ignoring them can lead to mis-specification of hypothesised

1 driver-state-impact relationships and flawed conclusions as to the most important
2 causes of change.

3

4 **1. Introduction**

5 On 29th October 2010, the parties to the UN Convention on Biological Diversity
6 agreed a new 10 year Strategic Plan. This followed failure to meet the Rio Earth
7 Summit goal “to achieve by 2010 a significant reduction of the current rate of
8 biodiversity loss at the global, regional and national level as a contribution to poverty
9 alleviation and to the benefit of all life on Earth” (Butchart *et al.* 2010). Action is
10 needed now more than ever to halt the direct drivers of biodiversity loss and
11 degradation of ecosystem services. This in turn means understanding how the relative
12 importance of different drivers varies across a heterogenous planet. Increasingly, this
13 understanding comes from analysis of the relationship between potential causal
14 factors, such as atmospheric pollutant deposition, climate change and land-use and
15 ecological state indicators (Sala *et al.* 2000; MA 2005). Such research takes time yet,
16 in a rapidly changing world, answers are needed quickly. Policy makers therefore find
17 themselves under pressure to infer possible causes from indicator change in the
18 absence of the evidence base that might emerge from time-consuming statistical and
19 model-based attribution analyses supported by experimentation to establish the
20 plausibility of the underlying mechanisms (Gadbury & Schreuder 2003). Sidestepping
21 the analytical attribution phase is attractive because changes in indicators can appear
22 to be superficially consistent with known changes in driving variables. This invites
23 common-sense interpretations; if on average, the ecological indicator has increased,
24 decreased or remained stable and, on average, the driver has intensified, reduced or
25 not changed, then the two are likely to be causally linked (Bellamy *et al.* 2005;

1 Butchart *et al.* 2010). While demonstrations of these simple and consistent
2 relationships are vital warning signs in the assessment of possible driver-state-impact
3 relationships, in themselves they offer incomplete evidence if the goal is to unravel
4 the relative contributions of multiple drivers across a domain of interest. Although the
5 apparent clarity and policy resonance of the message increases with the size of the
6 spatial units across which indicators are averaged, such high-level reporting can
7 obscure the existence and directions of ecosystem-specific trajectories of change
8 linked to different human-induced drivers (Jelinski & Wu 1996; Anderson *et al.*
9 2009). A step toward quantification of these relationships comes from analysis of
10 change in state indicators recorded from large-scale ecological surveillance programs.
11 Whilst such analyses maximise the realism and policy relevance of results, they must
12 also contend with a number of problems that serve to separate the detection and
13 attribution of signals in surveillance data from partitioning of the variation in a
14 designed experiment where the observer has control over the identity and arrangement
15 of treatments and covariates (Stow *et al.* 1998; Biggs *et al.* 2009; Wintle *et al.* 2010).
16
17 Here we illustrate some of the dangers of over-simplification in the presentation,
18 interpretation and analysis of large-scale ecological change and discuss a number of
19 related problems that are particularly relevant to the identification of cause-effect
20 hypotheses in ecological surveillance data. As a new round of global target setting,
21 indicator development and monitoring begins it is timely to highlight the influence of
22 these problems on the analysis of the impacts of anthropogenic stressors on
23 biodiversity and ecosystem services. By ecological surveillance data we mean large-
24 scale surveys repeated or not, where the primary motivation of the sampling design is
25 to quantify the range of ecological or biological variation in a region and the way it

1 may change over time (Gadbury & Schreuder 2003; Haughland *et al.* 2009). We
2 exclude long-term experiments where ecological change results from designed
3 arrangements of treatments and controls (Lindenmayer & Likens 2009).

4

5 **2. Three important issues**

6 *2.1 The Modifiable Area Unit Problem*

7 Across a heterogenous sampling domain, the relationships between variables can
8 change simply by altering the orientation and size of the areal units measured. In their
9 classic demonstration, Openshaw & Taylor (1977) found that correlations between
10 proportions of elderly voters and Republican voters across the state of Iowa ranged
11 from highly positive to highly negative simply by changing the number and
12 orientation of areal units used to divide up the state. Likewise, by changing the scale
13 over which indicators are averaged, the same phenomenon can readily influence the
14 apparent direction of ecological driver-state-impact relationships (Armhein 1995;
15 Stein *et al.* 2001). We illustrate the problem using recently published ecological
16 surveillance data for Great Britain.

17

18 In Britain, national changes in a range of biodiversity indicators have been reported
19 for the 29 year period from 1978 to 2007 based on large-scale ecological surveys
20 repeated at roughly decadal intervals (Carey *et al.* 2008). Each survey collects a large
21 number of biophysical measurements and these have been published as indicators of
22 change in extent and condition of major habitats, freshwaters, soils and plant species
23 (Carey *et al.* 2008 and available on-line at www.countrysidesurvey.org.uk). Following
24 the latest survey in 2007, the results were summarised by ecosystem type and at the
25 national-scale (Carey *et al.* 2008; CEH, 2008). Particular interest focussed on changes

1 in plant species composition that could be interpreted as shifts along the nutrient
2 availability gradient and therefore as a signal of eutrophication. Attribution analyses
3 of previous surveillance data highlighted the separate impacts of intensive agriculture,
4 nutrient surpluses and atmospheric nitrogen deposition on characteristic species
5 diversity between 1978 and 1998 (Firbank *et al.* 2008; Smart *et al.* 2006a; Smart *et al.*
6 2004). Hence, key questions were whether a signal of increasing eutrophication had
7 continued across British terrestrial ecosystems up to 2007, and if so, what were the
8 likely drivers? Expressing between-survey change in the nutrient status indicator as
9 the cross-ecosystem national average gave the simplest possible answer to the first
10 question: The relative contribution of the more nutrient-demanding plant species
11 increased and then decreased over the 29 year period (Fig. 1). At this scale, the
12 indicator has maximum impact with policy makers because it is communicated as one
13 simple, spatially unified message (CEH 2008). Within-ecosystem changes were also
14 published separately (Carey *et al.* 2008). These showed overall reductions in the
15 indicator in woodland and intensive farmland, but net increases in less fertile semi-
16 natural habitats such as acid, calcareous and neutral grasslands (Fig 2). Because the
17 time series is based on a representative, stratified-random sample of land cover, the
18 national trend was a weighted average of the directions and sizes of change observed
19 across the mosaic of habitats in Britain (Fig 1). The national indicator was correct. It
20 reflected the aggregate pattern of change along the inferred nutrient availability axis
21 but did not convey the net increase in the indicator in heath & bog and the semi-
22 natural grasslands (Fig 2). The national average trend also diverted attention from the
23 fact that different drivers were likely to have impacted different ecosystems and that
24 these different ecosystems vary in conservation value and response to perturbation.
25 For example, the extensification signal on arable land (Fig 2) appeared to reflect a

1 substantial transfer from cultivation to fallow in response to the domestic
2 implementation of Europe-wide setaside mechanisms designed to reduce crop
3 production (Boatman *et al.* 2009). In these highly responsive systems post-disturbance
4 colonisation is rapid (Critchley & Fowbert, 2000). However, the duration of any
5 positive ecosystem effects on soil protection, carbon sequestration and resource
6 availability for invertebrates and farmland birds are dependent on changes in crop
7 prices and can be abruptly reduced if fallow land is returned to intensive arable
8 cropping as happened in Britain in 2008 (DEFRA 2008; Boatman *et al.* 2009). In less
9 resilient semi-natural vegetation, eutrophication can be a much more persistent effect;
10 hard to reverse and cumulative in its impact (Strengbom *et al.* 2001; Dupouey *et al.*
11 2002). Only when the national trend was disaggregated was it possible to discriminate
12 between divergent ecosystem-specific trends, each associated with different suites of
13 known or possible drivers (Fig 2). Rather than concluding that the same pattern of
14 change occurred across the British landscape and that the most recent period saw
15 widespread recovery, a more detailed assessment revealed an ecosystem-specific
16 mixture of extensification, stability and eutrophication over the 29 year period (Fig 2).
17
18 Decisions about the definition and size of sampling units also impact greatly on the
19 detection and meaningfulness of patterns that purport to help understand ecosystem
20 service delivery and the spatial partitioning of biodiversity. Analyses that have sought
21 to identify trade-offs in ecosystem service provision have, for example, shown marked
22 scale-dependence in the direction of correlations (Anderson *et al.* 2009; Naidoo *et al.*
23 2008). In addition, the measurement of change in species diversity is highly scale-
24 dependent since increasing the size of the units censused confounds beta-diversity
25 with alpha-diversity (Buckland *et al.* 2005; Huston 1999). Partitioning diversity at

1 appropriate scales is especially important if the processes and drivers of change in
2 biodiversity are to be properly understood (Huston 1999; Green *et al.* 2005; Gabriel *et*
3 *al.* 2010). For example, regional to larger-scale reductions in rare species (gamma
4 diversity) have often occurred in parallel with local increases in small-scale alpha-
5 diversity (Sax & Gaines 2003). While both patterns can be reconciled with the effect
6 of human activities that increase the range of generalists and exotics but reduce the
7 range of scarce specialists, (Olden & Poff 2003; Smart *et al.* 2006b) analysis is
8 required at more than one scale to fully characterise these inter-related trends (Weber
9 *et al.* 2004; Stohlgren *et al.* 2002). Summing changes over fewer larger areas risks
10 averaging out opposing directions of ecological change whose individual trajectories
11 may be linked to local ecosystem-specific sets of drivers and starting conditions
12 (Wright & Jones 2004; Smart *et al.* 2006b).

13

14

15 *2.2 Incomplete or missing explanatory variables*

16 Averaging across areal units of different sizes is not the only way in which important
17 relationships with driving variables can be concealed. Previous attribution analyses of
18 British data have shown that large-scale ecosystem impacts can be correlated with
19 spatial gradients of multiple global change phenomena (e.g. Firbank *et al.* 2008;
20 Maskell *et al.* 2010). Yet, if a driver has increased in one place but decreased to an
21 equal extent in another then a cross-ecosystem average could convey no significant
22 mean change in the indicator rather than significant but location-dependent
23 correlation. The same could potentially happen where the impacts of two drivers are
24 negatively correlated in their operation within an ecosystem. This is the same as the
25 absence of significant main effects in an experiment because of the presence of a

1 negative interaction between levels of those main effects (Underwood 2005; Biggs *et*
2 *al.* 2009a). Such averaging problems will arise where explanatory variables have been
3 omitted from analyses. Therefore lack of an observed change in an indicator averaged
4 across a domain does not mean the absence of regional driver-state-impact dynamics.
5 The challenge is to incorporate appropriate data on covariates and driving variables
6 sufficient to isolate important interactions and main effects for example reflecting
7 geographical variation in starting conditions, land-use and socio-economic context
8 (Van Buskirk & Willi, 2004, 2005; Kleijn & Báldi, 2005) or across species' centres of
9 distribution versus range margins (Oliver *et al.* 2009; Warren *et al.* 2001). An ongoing
10 problem is that in regions where biodiversity is high yet ecosystem degradation
11 severe, adequate explanatory variable data may be sparse or non-existent (MA, 2005).

12
13 Analysis of recent large-scale changes in disease prevalence provide some of the best
14 examples of how spatial disaggregation of data can reveal smaller scale trends each
15 associated with different suites of social and ecological covariates. Increased risk of
16 tick-borne encephalitis (TBE) to humans has occurred across parts of Europe in the
17 last thirty years. Increased disease risk was thought to have resulted from improved
18 conditions for natural transmission cycles resulting in higher densities of infected
19 ticks or from changes in human behaviour resulting in greater exposure to ticks
20 (Randolph 2008a). Since both of these factors are climate dependent, TBE is
21 commonly listed amongst those vector-borne pathogens anticipated to become more
22 of a threat to humans as the climate warms (e.g. Lindgren, 1998). Lindgren &
23 Gustafson (2001) were the first to attribute the upsurge in TBE incidence over the past
24 two decades to warming temperatures – namely milder winters and early arrival of
25 spring (conditions favouring early development and extended autumn activity of

1 ticks). However, this study considered only climatic drivers and was restricted to
2 Stockholm county, Sweden, close to the northern range margin of this pathogen. In
3 fact, when a regional level investigation at a pan-european scale was conducted,
4 considerable variation was revealed, both within and between countries, in the timing
5 and extent of changes in TBE incidence. Though biologically-relevant climate
6 warming had indeed occurred since the early 1990s, and may have enhanced
7 transmission, the pattern of these changes was too uniform across the continent to
8 account for the extreme spatio-temporal heterogeneity in the upsurge of TBE (Sumilo
9 *et al.* 2006; 2007; Randolph 2008a; 2008b). Instead a network of interacting biotic
10 and socio-political drivers affecting both risk of infection and exposure of humans,
11 differing in force in space and time, was involved (Randolph 2008a).

12

13 A similarly comprehensive analysis invoking multiple and interacting drivers was also
14 necessary to understand the increasing incidence of American cutaneous leishmaniasis
15 (ACL), (a zoonotic vector-borne disease, caused by several species of *Leishmania* and
16 transmitted by sandflies) in the neotropics. In this example inclusion of another
17 covariate revealed an interaction with the primary driver that reversed the apparent
18 direction of the initial driver-state-impact relationship. The emergence of ACL has
19 been associated with changes in the interactions between people and forests. The
20 association between outbreaks and higher rates of infection found in populations
21 living close to forests and lower incidence in urban areas led to the proposal that
22 deforestation could reduce re-emergence of disease. These analyses ignored socio-
23 economic factors. Chaves *et al.* (2008) examined county level incidence rates as a
24 function of social and environmental variables. In common with other infectious
25 diseases, socially excluded populations were most affected by the disease. Once social

1 marginality was taken into account, living close to the forest could actually diminish
2 the risk of ACL infection. In addition, impacts of climate change on disease, due to El
3 Nino Southern Oscillation events (initially described in Chaves *et al.* 2006), interacted
4 with forest cover since resulting increases in incidence were exacerbated by higher
5 levels of deforestation. Including socio-economic covariates thus reversed the original
6 expectation. Poorer citizens were more likely to contract ACL whether in urban or
7 rural areas but were more likely to live closer to forest edges. While overall incidence
8 increased with climate effects, incidence declined with greater remaining forest cover.

9

10 Complex relationships between multiple driving variables and responses are likely to
11 be common. Unravelling them may face a simple problem of lack of data that track
12 drivers of interest at the study scale. This can be an insurmountable obstacle (e.g. MA,
13 2005) in the face of which, the identification of causal hypotheses based on those data
14 that are available, must consider the possibility that other drivers could be equally
15 important and that, if included, they may even alter the direction of the relationship
16 with the principal driver (see Supplementary Material).

17

18 *2.3 Arrangement of drivers is beyond the control of the observer*

19 A number of analytical and interpretative problems can also arise because of an
20 under-appreciation of the differences between analysis of correlative/causal
21 relationships in ecological surveillance data versus partitioning the variation in a
22 designed experiment (Stow *et al.* 1998). Unlike a designed experiment, the identity,
23 crossing, replication and interspersion of driving variables is, by definition, outside
24 the control of the observer (Stow *et al.* 1998; Gadbury & Schreuder 2003). This
25 makes attribution an analytical challenge since we must apply strong inference (*sensu*

1 Platt 1964) to weak data. That is we apply the steps involved in inductive inference
2 especially hypothesis construction and testing but to data that reflects accidental
3 arrangements of possible explanatory variables. This creates difficulties in testing
4 competing hypotheses and in gaining clear understanding from the analyses.
5
6 Analyses of large-scale survey and monitoring data face well known issues such as
7 regression to the mean (Palmer 1993; Smart & Scott 2004; Biggs *et al.* 2009), pseudo-
8 replication (Hurlbert 1984; Cottenie & De Meester 2003) and spatial autocorrelation
9 (Beale *et al* 2010; Hawkins *et al* 2007). Less widely discussed is the influence of
10 differences in the range of variation in driver intensity. This can profoundly affect the
11 conclusions drawn from analyses that seek to detect driver-state-impact relationships.
12 In a large-scale analytical survey potential drivers may show considerable spatial
13 variation in their intensity. Analyses may have high realism but the apparent
14 importance of one driver versus another in driving change, maybe as much to do with
15 the fact that the severity of a driver has been consistently high in most places, while
16 the severity of another driver varies being high in some places but being absent or at
17 low severity in others. If the range of variation in a driver is relatively small despite its
18 severity being high everywhere, then variance partitioning techniques have a lesser
19 chance of detecting a significant effect of the driver over and above residual variation
20 in the response. We carried out a simulation exercise to demonstrate this effect (Fig
21 3). The driving variable ranged between 1 and 40, approximating the current range of
22 total atmospheric nitrogen deposition ($\text{Kg N ha}^{-1} \text{ yr}^{-1}$) across Britain as a realistic
23 example of a national-scale driver of ecological change. A realistic relationship with a
24 response variable was specified and the power of detecting a significant relationship
25 tested across different ranges of variation in the driver (Fig 3). The conclusion from

1 such a test is not necessarily that the driver has not or could not induce an ecological
2 impact. If the regression slope coefficients of two driver-impact relationships were
3 equivalent, then their potential for forcing ecological change is estimated to be
4 equivalent if they were applied with equal crossing and replication. The problem can
5 be made clearer by acknowledging that two questions are involved; 1) what is the
6 inherent potency of the driver as a cause of ecological change relative to other
7 potential drivers (best answered via experiment and up-scaled via a model), and, 2)
8 which drivers happen to be most strongly correlated with observed ecological change
9 across the entirety of a region of interest? Statistical attribution relies on being able to
10 evaluate the size of a driver's impact by comparing ecological changes at high and
11 low levels of a driver; either along a gradient or by classifying the sampling domain
12 into control and impacted regions. Taking into account the relative range of variation
13 in possible drivers across the sampling domain is therefore of equal importance in
14 answering the second question as is quantifying variation in the ecological impact. If a
15 driver had operated with constant and high severity everywhere and an associated
16 magnitude of impact had occurred everywhere then the only deviations from the mean
17 of the response will be random about the mean overall level of the explanatory
18 variable. This rules out the possibility of a systematic association between the two and
19 hence a significant detected effect of the driver on the response (Fig 3). It also means
20 that a highly significant average change in a state indicator may defy statistical
21 attribution based on variance partitioning techniques even if the potential driver seems
22 very obvious. Analysis of the ecosystem impacts of reduced versus oxidised
23 atmospheric nitrogen deposition provides a topical example of the importance of
24 separating assessment of the inherent potency of two drivers from an assessment of
25 their relative importance as an additional consequence of their spatial variation in

1 severity. At the resolution of larger grid cells, reduced nitrogen deposits over a much
2 greater range of estimable values across Europe than oxidised. This means that
3 attribution of large-scale change in state indicators to reduced nitrogen is more likely
4 than to oxidised (e.g. Smart *et al.* 2004; McClean *et al.* 2011) even though the
5 biogeochemical mechanisms underlying their impacts, indicate that both can be
6 significant pollutants (Stevens *et al.* 2011). Reanalysis of datasets based on equalising
7 gradient lengths is possible but the constrained subsample of a large-scale survey
8 would no longer represent the realistic, un-designed variation in each nitrogen form as
9 a function of the human activities that control the deposition of each.

10

11 **3. Conclusions**

12 While the principal objective of many surveillance and monitoring programs is to
13 detect change, the question, why has change occurred?, inevitably follows. Yet, just
14 because schemes are designed to detect change across a region does not mean they are
15 optimised to attribute changes to potential drivers. The basic difference between
16 controlled experimentation and passive ecological surveillance contributes to all three
17 of the problems we have illustrated. We make the following recommendations:

18

19 Be wary of narratives that analyse large-scale change in indicators and then invoke
20 drivers operating across large areas but without quantitatively linking them: The
21 statistical power of such an analysis can be high leading to a powerfully simple
22 message but one that may conceal important local differences. High-level results
23 should therefore, be accompanied by an appropriate decomposition into disaggregated
24 trends and a health warning that the simple picture is simple at the expense of
25 potentially critical complexity. Techniques such as random forests (Bradter *et al.*

1 2011), locally-weighted regression and CART modelling can be used to detect and
2 characterise these variations in ecosystem response thus highlighting the locations of
3 different driver-state-impact dynamics for further hypothesis testing (Smart *et al.*
4 2003). Techniques such as hierarchical Bayesian modelling (Biggs *et al.* 2009) allow
5 joint analyses of these substrata enabling the detection of overall correlative/causal
6 relationships between drivers and state indicators but conditioned on environmental
7 heterogeneity. Such approaches help avoid excessive belief in a single simplistic
8 unified message about the causes of ecological change. Similar issues of scale-
9 dependence and aggregation or averaging effects on the clarity of higher level
10 messages have also been discussed regarding the calculation and dissemination of
11 environmental sustainability indicators (Morse & Fraser 2005; Parris & Kates 2003).

12

13 Analyses of driver-state-impact relationships from large-scale ecological surveillance
14 have high realism and policy relevance. However, the ‘accidental’ way in which
15 drivers are spatially organised means that detection of relationships can be as much an
16 indication of the lack of equal crossing and replication of possible causal factors as of
17 the inherent potency of a driver to cause change. For example to what extent does one
18 driver emerge as more important than another just because it had operated with
19 greater variation in severity across the study region?

20

21 With sufficiently large datasets it may be possible to analyse subsets of replicated data
22 where variation in a driver is maximised along a gradient or between impacted and
23 control groups. Other drivers could be equivalently crossed and replicated or held
24 constant along with other additional covariates, such as ecosystem type. However,
25 such designed sub-sampling changes the question. Results will no longer realistically

1 represent the accidental interplay of the range of drivers across the entire sampling
2 domain. Such studies provide the basis for quantifying the impact of a driver but over
3 a domain deliberately engineered to maximise the magnitude of the driver-state-
4 impact signal. These approaches are therefore most useful in deriving realistic
5 statistical models of driver-impact relationships (Maskell *et al.* 2010; Stevens *et al.*
6 2009). Predicted impacts can then be usefully compared with observations in other
7 parts of the sampling domain where other drivers may be operating (e.g. Diekman &
8 Falkengren-Grerup 2002; Stevens *et al.* 2004).

9

10 Finally, we should be aware that the emerging importance of a driver may simply
11 reflect data availability: If data is only available that tracks one driver but an
12 ecological impact could be more fully explained by the main or interacting effect of
13 other drivers then conclusions need to be accompanied by a qualification that analysis
14 is incomplete. Differences in data quality can also be influential. For example more
15 coarsely resolved explanatory variables cannot explain as much variation as those
16 more closely matched to the resolution of the ecological response data (Smart *et al.*
17 2006a). Again, this may mean that some drivers appear more important in explaining
18 change in ecological state indicators than others.

19

20 Whilst the problems we have described are simple to understand, each demands a
21 cautionary perspective which may be easily overlooked in the rush to interpret
22 indicator change in the light of apparently obvious changes in the most likely drivers.
23 Reducing the decline in global biodiversity and the degradation of ecosystem services
24 is both costly and vital. If cost-effective responses are to be formulated they must be
25 based on an accurate understanding of the role of multiple drivers and the way they

1 interact to cause change. Analyses of ecological surveillance data should support the
2 elucidation of causal-correlative links between drivers and state variables. To do this
3 we believe that scientists and policy consumers need to be more aware of the
4 implications of the simple yet subtle problems we have highlighted.

5

6

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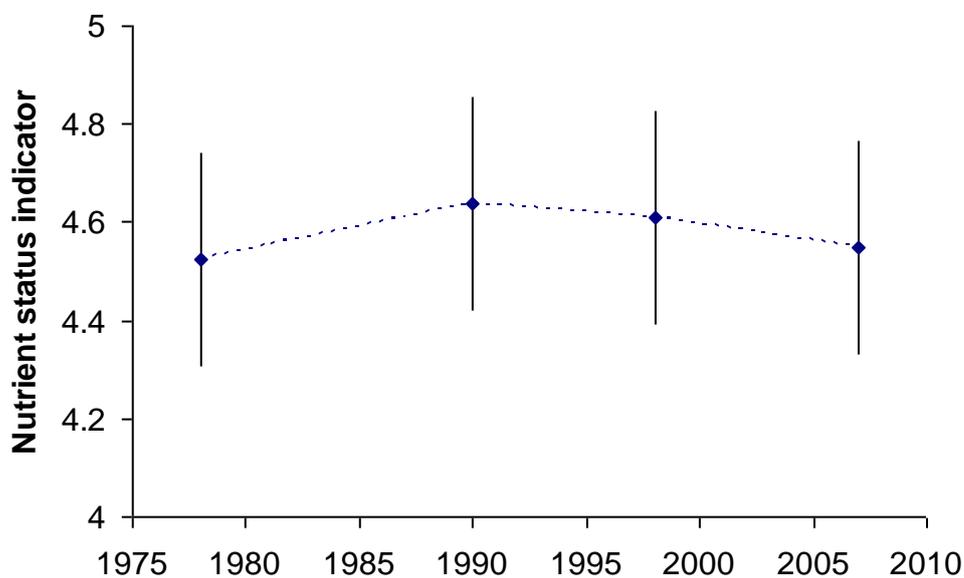
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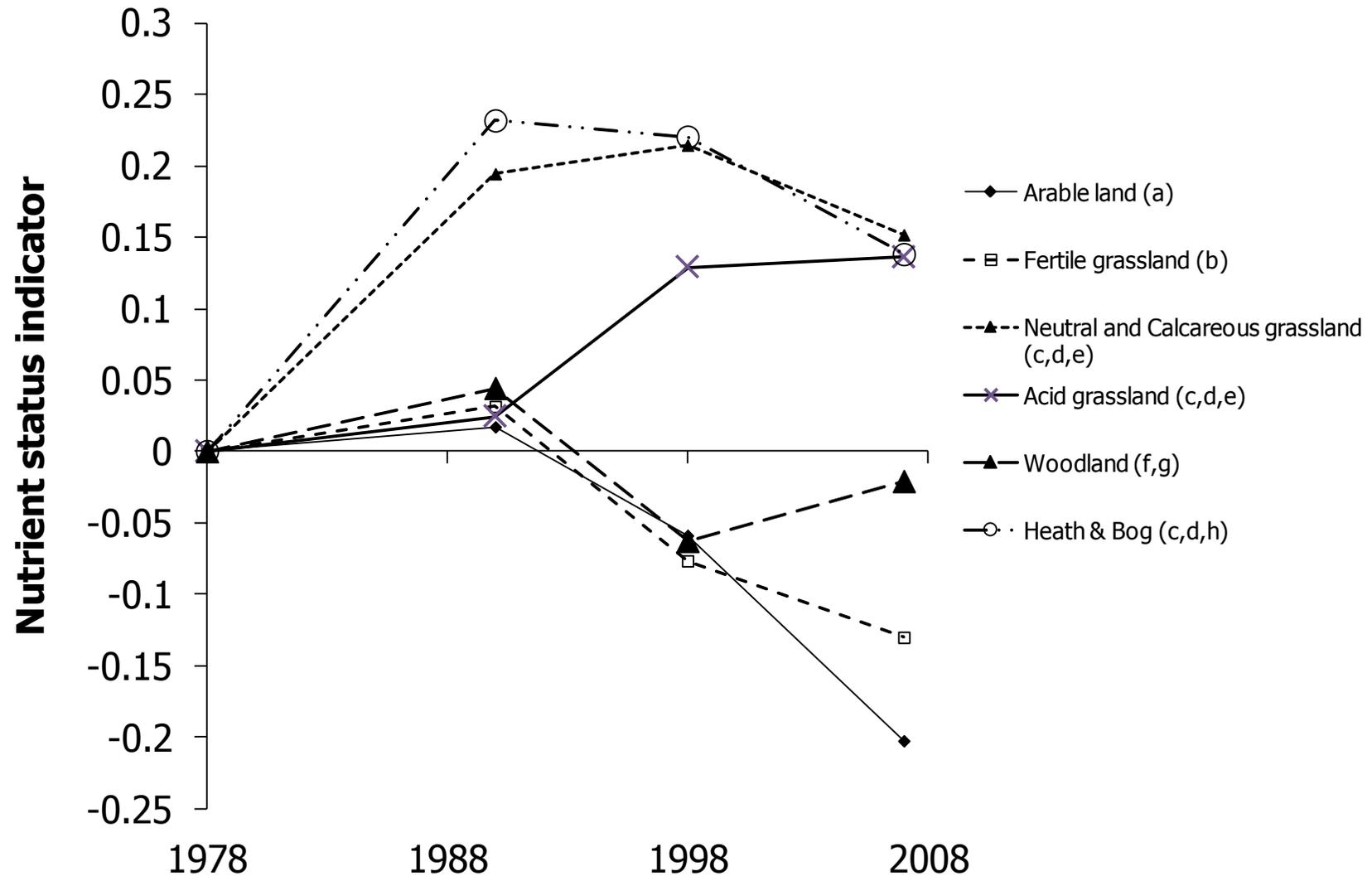
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1 Figure 1. Change in the vegetation nutrient status indicator across British ecosystems
2 between the Countryside Surveys of Great Britain in 1978, 1990, 1998 and 2007. The
3 score was derived from the mean of individual indicator (Ellenberg N) values
4 assigned to species in the British flora, for species that were found in fixed sampling
5 plots in each survey year. Error bars are the 95% confidence intervals on the within-
6 year means (n=729 repeat plots). Statistically significant changes in mean score
7 occurred between all pairs of years except 1978 to 2007 and 1990 to 1998, based on
8 an autoregressive, generalised linear mixed model. An increase in the index indicates
9 greater representation of more nutrient-demanding species. The scale is such that 1
10 indicates species associated with the least, and 9, with the most productive conditions.
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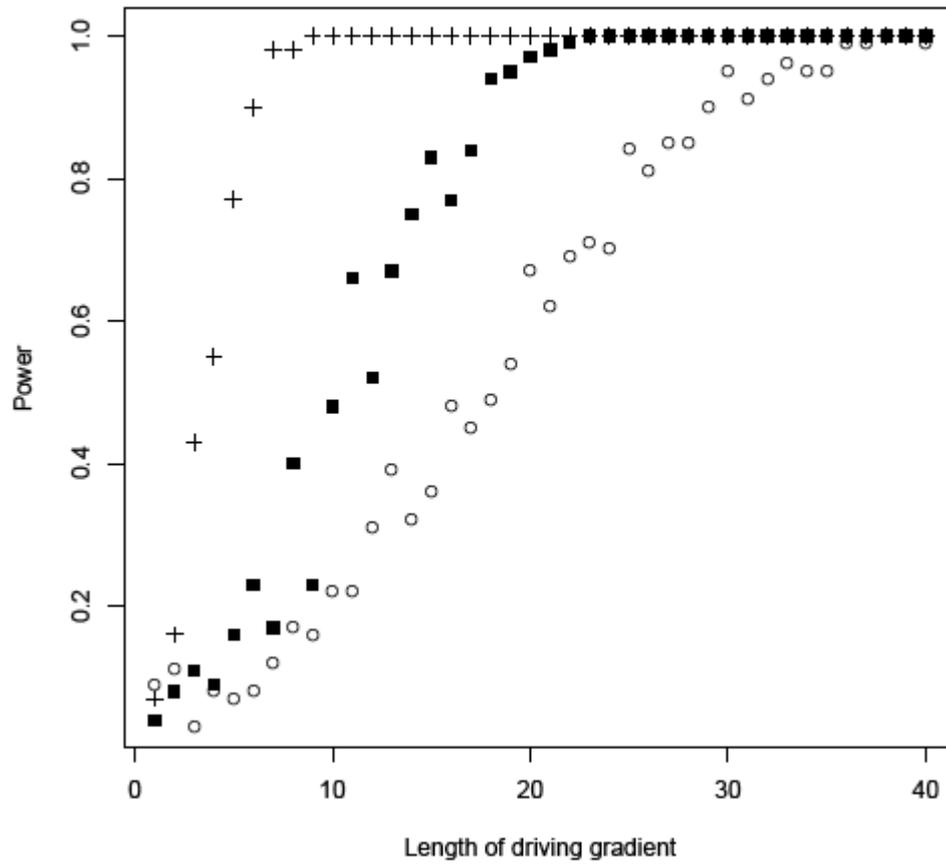


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1 Figure 2. Change in the vegetation nutrient status indicator (see Fig 1) decomposed by ecosystem type. Values are the difference in the indicator
2 from the starting value in the 1978 survey where all ecosystem types start at zero. Letters after each ecosystem type refer to known or
3 hypothesised large-scale drivers of change in the ecosystem type over the survey period as follows; a) removal of land from production post-
4 1988 (Carey *et al.* 2008; Critchley & Fowbert 2000); b) reduced management intensity post-1990 (Carey *et al.* 2008; Smart *et al.* 2005); c)
5 atmospheric nitrogen deposition (Smart *et al.* 2004; Maskell *et al.* 2010); d) abandonment (Sketch 1995); e) agricultural intensification (Smart *et*
6 *al.* 2006a; Chamberlain *et al.* 2000), f) woodland succession (Kirby *et al.* 2005), g) lack of traditional management (Kirby *et al.* 2005), h) over-
7 grazing (Fuller & Gough 1999).



1 Figure 3. Effect of the range of variation in a driver of ecological change on the
2 probability of detecting a significant correlation with an ecological indicator. Data
3 were simulated based on the linear regression model $y=0.7*x$, with random residual
4 error $e \sim N(0, \sigma^2)$ where σ^2 was set to each of three values; + = 5, ■ = 15, ○ = 25.
5 For each error variance, forty datasets were generated each of which consisted of 200
6 values but with x made to vary in range within each dataset between 40 and 1 but
7 always with a maximum value of 40. The slope parameter was therefore the same in
8 each dataset but the range of variation over which non-random deviations from the
9 mean response were quantified, varied from large to small. The power of detecting a
10 significant x parameter ($p<0.05$) was expressed as the number of significant linear
11 regression results out of 100 random draws from each error distribution for each
12 dataset. Power was then plotted against the range of the driving variable x . As the
13 range of x was reduced the chance of detecting a significant regression of y on x also
14 reduced. Power declined more quickly if the response variable had greater residual
15 error. Note that since the maximum value of x was always 40, low values on the X
16 axis represent situations where the driver operated with high severity everywhere.



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