Effect of a context shift on the inverse base-rate effect

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Abstract

The Inverse Base Rate Effect (IBRE; Medin and Edelson (1988)) is a non-rational behavioural phenomenon in predictive learning. In the IBRE, participants learn that a stimulus compound AB leads to one outcome and that another compound AC leads to a different outcome. Importantly, AB and its outcome are presented three times as often as AC (and its outcome). On test, when asked which outcome to expect on presentation of the novel compound BC, participants preferentially select the rarer outcome, previously associated with AC. This is irrational because, objectively, the common outcome is more likely. Usually, the IBRE is attributed to greater attention paid to cue C than to cue B, and so is an excellent test for attentional learning models. The current experiment tested a simple model of attentional learning proposed by Le Pelley, Mitchell, Beesley, George, and Wills (2016) where attention paid to a stimulus is determined by its associative strength. This model struggles to capture the IBRE, but a potential solution suggested by the authors appeals to the role of experimental context. In the present paper, we derive three predictions from their account concerning the effect of changing to a novel experimental context at test, and examine these predictions empirically. Only one of the predictions was supported, concerning the effect of a context shift on responding to a novel cue, was supported. In contrast, Kruschke (2001b)’s EXIT model, in which attention and associative strength can vary independently, captured the data with a high degree of quantitative accuracy.

Keywords: IBRE, EXIT, predictive learning, categorisation

1 Introduction

The Inverse Base Rate Effect (IBRE; Medin & Edelson [1988]) is a non-rational learning phenomenon that has generated considerable debate within the literature (Bohil, Markman, & Maddox, 2005; Juslin, Wennerholm, & Winman, 2001; Kruschke, 1996, 2001b, 2003; Winman, Wennerholm, & Juslin, 2003). In its canonical form, participants are asked to diagnose fictitious patients under a simulated medical diagnosis procedure. They are initially presented with a patient showing one of two different symptom pairs, which can be considered abstractly as AB and AC. They are then asked to decide which of two fictitious diseases that patient has. For example, a participant might be presented with a patient suffering from a rash and nausea (AB), where the correct diagnosis is Jominy Fever. Then they might see a patient suffering from a rash and back pain (AC), where the correct diagnosis is Phipps Syndrome. In this example nausea (B) is perfectly predictive of Jominy Fever, while back pain (C) is perfectly predictive of Phipps Syndrome. The rash (A) is uninformative. Participants see patients for whom the correct diagnosis is Jominy Fever three times as often as those for whom the correct diagnosis is Phipps Syndrome. In other words, Jominy Fever is a common disease, while Phipps Syndrome is a rare disease. Participants are then presented with both perfectly predictive symptoms together, nausea (B) and back pain (C). If participants correctly make use of the base rate of the two diseases, they should make the rational diagnosis of the more common disease (Jominy Fever in our example). However, the majority of participants preferentially diagnose the patient with the rarer disease. This pattern of responding is called the IBRE.

The IBRE sometimes co-occurs with another response pattern. Specifically, common-disease responding to B is sometimes observed to be greater than rare-disease responding to C, when these stimuli are presented individually at test. This is surprising; it suggests that B is more strongly associated with the common outcome than C is with the rare outcome, while C dominates responding when they are presented in conjunction. This response pattern has been reported in a number of studies (e.g. Bohil et al., 2005; Winman, Wennerholm, Juslin, & Shanks, 2005), with Wills, Lavric, Hemmings, and Surrey (2014) first confirming that the difference was statistically significant; a finding that has recently been replicated (Inkster, 2019, Exp.3). We refer to
the co-occurrence of the IBRE and this response pattern as the Compound versus Features Dissociation (COFED). We describe the effect as a dissociation because the response to the compound BC is opposite to what one would expect from the summation of the responses to the individual cues.

Although several potential explanations of the IBRE and the COFED exist, the EXIT model (Kruschke, 2001a) is particularly strong, having previously accounted for both the IBRE and the COFED in other experiments (Kruschke, 2001a, 2003; Kruschke, Kappenman, & Hetrick, 2005). EXIT assumes that the IBRE is driven by an error-driven learning attentional effect, where the participant learns to direct their attention away from cues that lead to prediction errors. Specifically, early in training, participants make many errors on rare AC trials due to their similarity to common AB trials. EXIT assumes that people learn to avoid those errors by directing their attention on AC trials away from A and towards C. Crucially, this attentional weighting of C is assumed to be persistent. Hence, when presented with BC, C can dominate responding, even in cases where B is more strongly associated to the common outcome than C is to the rare outcome (a COFED).

Recently, Le Pelley et al. (2016) argued that much of the data relating to human attentional learning could be accounted for by a simpler model that EXIT and its relatives (e.g., Mackintosh, 1975). In this model, the attention that a stimulus demands is a simple function of the associative strength of that stimulus. Although consistent with much of the attentional learning literature, this simple model appears to be undermined by the IBRE; here, the cue that appears to possess the greatest associative strength (B) does not attract more attention than the weaker cue (C). Le Pelley et al. (2016) offered a potential cue to this, appealing to the role of experimental context. Experimental context refers to the procedural context of the experiment. In a medical diagnosis task, examples of experimental context include the patients being diagnosed or where patients are diagnosed.

In Le Pelley et al. (2016)'s context explanation the experimental context is represented as a cue that is present on every trial; cue X. In the IBRE procedure, X becomes more strongly associated with the common outcome (also associated with B) than the rare outcome (also associated with C), due to the greater frequency of the common outcome. Figure 1 illustrates how the COFED can be explained from this assumption, plus the assumption that the associative strength from C to the rare outcome is greater than the associative strength from B to the common outcome (we leave aside the issue of why C might have greater associative strength than B at this point). In Le Pelley et al.’s account, responding to cue compounds at test is then predicted by summing the associative strengths of the cues they contain. As can be observed, XBC results in rare responding (an IBRE), while at the same time there is more common responding to XB than there is rare responding to XC (a COFED).

Le Pelley et al.’s account of the COFED leads to three predictions, which are illustrated in Figure 2. These predictions involve switching to a novel context (Y) at test. Y has no associative strength to either the common or the rare outcome. The first prediction of the Le Pelley et al. account is that YBC produces a larger proportion of rare responding than XBC—in other words, a context shift should enhance the size of the IBRE due to the novel context having no association to either outcome. The second prediction is that rare responding to the YC will be greater than common responding to YB—a context shift will reverse the COFED. The third prediction is that while a novel cue presented in the same context as training (XN) should produce common responding, a novel cue presented in a novel context (YN) should not produce preferential responding to either the common or the rare outcome. In other words, a shift to context Y will bias responding away from the common outcome (with which context X is associated) and towards the rare outcome. In the experiment reported below, we test these three predictions.

Figure 1: Informal example of the context explanation of the COFED. Orange bars represent possible learned associative strengths of individual cues at the end of training. Blue bars represent predicted associative strengths for test stimuli.
2 Method

2.1 Participants

Participants were all undergraduate students from the University of Plymouth, completing the experiment for partial course credit. They were a range of ages, genders, and ethnicities. Ninety four participants were tested. Assuming participant exclusion rates similar to Inkster (2019), this sample size provides adequate power to detect a small-to-medium-sized effect of context shift \( d = .31 \) at 80% power. It further provides over 99% power to detect the IBRE and the COFED, at the effect sizes observed in Inkster (2019), \( d = .46 \) and .56, respectively.

2.2 Stimuli and apparatus

The stimuli (see Figure 3) were abstract shapes, red and yellow in colour and \( 30 \times 30 \) pixels in size; previously used in Wills et al. (2014). They were displayed on 22-inch flat-screen monitors using PsychoPy (Peirce, 2007). Participants sat approximately 50 cm from the screen, giving each cell a visual angle of approximately 2 degrees. Responses were collected using a standard PC keyboard.

Table 1 shows the combinations of abstract cues and diseases presented in the training and test phases of the experiment. The stimuli were assigned at random to one of 7 abstract cues (A-G) for each participant for the training phase. As in Wills et al. (2014), each abstract cue had 3 stimuli assigned to it. A subset of possible cell combinations was used for the compound cue trials, for example on AB trials the cells presented were: \( A_1B_1, A_2B_2 \) or \( A_3B_3 \). The cue compounds FD and GE represent disjoint cue trials, where the proportion of outcomes is the same as AB and AC but there is no shared cue. During the test phase one more abstract cue was presented, the novel cue, N. N had six stimuli assigned to it, rather than the three assigned to the other abstract cues. This is due to the novel nature of the cue and the fact that there are two test phases; resulting in three of the cues assigned to N being used in the first test phase and the other three in the second. In total, 27 different cells were used in this experiment. Two diseases were used as outcomes: “Jominy Fever” and “Phipps Syndrome”. Diseases were mapped to the abstract disease types in both possible ways, across participants, as were the response keys mapped to those diseases.

![An example compound stimulus.](image)

**Figure 3** An example compound stimulus.

2.3 Procedure

The procedure was closely based on Wills et al. (2014), with the addition of a context manipulation. Participants were tasked with diagnosing patients with one of two diseases, Jominy Fever or Phipps Syndrome, on the basis of the “cells” they were presented with. Two contexts were used, one where participants diagnosed human patients and one where they diagnosed orcs. Participants always diagnosed humans during training, but completed two test phases, one with orcs and one with humans. The order of the test phases was counterbalanced between participants. In each of the three phases (one training, two test) the trial order was randomised between participants.

The training phase comprised 20 blocks of 18 trials, 360 trials in total. Each trial began with a 1000 ms presentation of a grey viewbox, which indicated where the cells would appear. The cells then appeared, centralised horizontally, but towards the top and bottom of the viewbox vertically. The cell presented to the top and to the bottom of the viewbox in the training phase and participants continue to receive feedback at test for these trials to maintain learning.
and what the correct diagnosis was. Feedback was presented on screen for 1500 ms before a new trial began. If a response was not made during the 2000 ms the cells were on screen, participants instead received a time-out message, displayed for the same duration as the feedback.

In the “orcs” test phase, participants were told they would be completing a medical placement in a different dimension and would now be diagnosing orcs. This was further emphasized on every trial by surrounding the viewbox with a large green outline of an orc’s face. In the “humans” test phase, participants were told they would continue to diagnose humans. Each test phase consisted of 216 trials, with the same trial structure as in the training phase, and single cells being presented in the middle of the viewbox. Feedback continued to be presented for stimuli that were presented during the training phase, in order to maintain learning. For the novel cues and compounds, no feedback was presented; participants instead received the message “data missing”.

3 Results

Raw data, analysis and modelling scripts are available at https://osf.io/8p42b/ Analysis was conducted using R (R Core Team, 2018), with packages, ez (Lawrence, 2016), tidyr (Wickham & Henry, 2019), dplyr (Wickham, François, Henry, & Müller, 2019) and pwr (Champely, 2018). Null-hypothesis significance tests were conducted at an alpha level of .05.

Bayesian tests were also conducted, with Bayes Factors less than one third interpreted as substantial evidence for the null, and Bayes Factors greater than 3 were interpreted as substantial evidence for the alternative. Prior effect sizes were drawn from previous related experiments reported by Inkster (2019). Following Dienes (2011), these effect sizes were used to construct a half-normal prior with a mean of 0.26 and standard deviation of 0.05.

In the test phase, all participants received both a same-context test phase (humans), and a different-context test phase (orcs). In the following analyses, context (same vs. different) is treated as a within-subjects factor. Table 2 shows the response proportions for each stimulus under both same-context and different-context conditions. Cues that are abstractly identical have been combined in this Table. For example, “A” represents responses to $A_1$, $A_2$, and $A_3$. In the following analyses, the data were first analysed within the same and different conditions, and then between the two conditions.

**Same-context test.** The proportion of rare-outcome responding to BC was greater than .5, $BF_{10} = 21$, indicating the presence of an IBRE. Common-outcome responding to B was greater than rare-outcome responding to C, $BF_{10} = 33644$; together with the IBRE, this demonstrates the presence of the COFED in this condition. The proportion of common-outcome responses to the novel stimulus, N, was greater than .5, $BF_{10} = 3.66$.

**Different-context test.** The IBRE was again observed, $BF_{10} = 11$, as was greater common-outcome responding to B than rare-outcome responding to C, $BF_{10} = 31269$, again demonstrating the COFED. Common-outcome responding to N did not differ from .5, $BF_{10} = .09$.

**Effect of changing context.** The proportion of rare-outcome responding to BC, and hence the size of the IBRE, was unaffected by the change in context, with substantial evidence for the null, $BF_{10} = .03$. Similarly, neither common-outcome responding to B, nor rare-outcome responding to C, was affected, again with substantial evidence for the null; $BF_{10} = .02$ and $BF_{10} = .02$, respectively. Taken with the evidence suggesting that context shift has no effect on the IBRE, this suggests that context shift has no effect on the

![Figure 4: Participants’ accuracy on the abstract training trial types at different levels of training. The error bars represent within-subject Cousineau-Morey 95% confidence intervals](image-url)
Table 2: Proportion of common and rare responses to each of the stimulus types presented under different-context and same-context conditions. Bold type highlights the results of primary theoretical interest. Values within brackets represent response proportions from the simulation of this experiment using the EXIT model with optimised parameters.

<table>
<thead>
<tr>
<th>Stimulus type</th>
<th>Common Same</th>
<th>Diff</th>
<th>Rare Same</th>
<th>Diff</th>
</tr>
</thead>
<tbody>
<tr>
<td>A</td>
<td>.72(.72)</td>
<td>.69(.66)</td>
<td>.28(.28)</td>
<td>.31(.34)</td>
</tr>
<tr>
<td>AB</td>
<td>.89(.86)</td>
<td>.87(.83)</td>
<td>.11(.14)</td>
<td>.13(.17)</td>
</tr>
<tr>
<td>AC</td>
<td>.30(.19)</td>
<td>.30(.20)</td>
<td>.70(.81)</td>
<td>.70(.80)</td>
</tr>
<tr>
<td>B</td>
<td>.87(.90)</td>
<td>.88(.88)</td>
<td>.13(.10)</td>
<td>.12(.12)</td>
</tr>
<tr>
<td>BC</td>
<td>.42(.41)</td>
<td>.42(.45)</td>
<td>.58(.59)</td>
<td>.58(.55)</td>
</tr>
<tr>
<td>C</td>
<td>.26(.20)</td>
<td>.26(.23)</td>
<td>.74(.80)</td>
<td>.74(.77)</td>
</tr>
<tr>
<td>D</td>
<td>.90(.86)</td>
<td>.88(.82)</td>
<td>.10(.14)</td>
<td>.12(.18)</td>
</tr>
<tr>
<td>DE</td>
<td>.47(.50)</td>
<td>.47(.52)</td>
<td>.53(.50)</td>
<td>.53(.48)</td>
</tr>
<tr>
<td>E</td>
<td>.25(.34)</td>
<td>.26(.36)</td>
<td>.75(.66)</td>
<td>.74(.64)</td>
</tr>
<tr>
<td>FD</td>
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<td>.92(.88)</td>
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<td>.08(.12)</td>
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<td>.14(.14)</td>
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</tr>
<tr>
<td>N</td>
<td>.57(.50)</td>
<td>.50(.50)</td>
<td>.43(.50)</td>
<td>.50(.50)</td>
</tr>
</tbody>
</table>

4 Discussion

4.1 Summary and interpretation of findings

We observed both the inverse base-rate effect (IBRE) and the compound versus features dissociation (COFED) in this experiment, replicating the results of Wills et al. (2014) and Inkster (2019) in this regard. However, the context-based explanation of these effects, recently offered by Le Pelley et al. (2016), was largely not supported by the current experiment. Their explanation predicted an increase in the size of the IBRE with a change to a novel context at test, while in our experiment this had no effect, with Bayesian evidence for the null. Similarly, the context explanation predicted a reversal of the COFED with a change of context, while we observed no effect, again with Bayesian evidence for the null. It would be difficult to argue that the context manipulation went unnoticed by the participants, because it affected the proportion of common-outcome responding to a novel cue, in the manner predicted by the context explanation.

4.2 The EXIT model

The EXIT model was able to capture all the main results of our experiment, as shown in Table 2, and it does so with an excellent level of quantitative fit, \( R^{2} = .97 \). For technical details of our simulation methodology, see the Appendix.

Interestingly, the EXIT model can capture both responding to a novel cue and the effect of context shift on responding to a novel cue, both effects that EXIT has not been tasked with capturing previously. It’s ability to do so is because, like Le Pelley et al.’s explanation, it assumes a context cue that is present on every trial. However, unlike the Le Pelley et al. explanation, it is not the presence of this context cue that drives the IBRE in EXIT. Rather, the context is seen as a relatively low salience stimulus in the EXIT account (the COFED is due to differences in two factors — strength and attention/salience). This then allows EXIT to capture both the IBRE and the COFED, as detailed in the introduction. It further allows EXIT to capture the response pattern to the novel cue. On a novel cue trial, the context is the only cue that has any salience, and so it is the only basis on which a prediction can be made.

4.3 Future research

The COFED, although robust within the procedures of the current experiment, appears to be somewhat procedure dependent. Specifically, although we have now found a significant COFED on three separate occasions with the current procedure, changes in procedure can result in either an IBRE without a COFED, or a COFED without an IBRE (Inkster, 2019). One possible explanation is that, with insufficient training, an IBRE is not observed but, with too much training, performance on both B and C are at ceiling, precluding the ability to observe a COFED. This possibility merits further investigation.

A second possible topic for future research concerns the responding to a novel cue at test. In the current experiment, we reported common-outcome responding when the context was the same as in training. Interestingly, this result is in contrast to Juslin et al. (2001) who reported that, under some conditions, a novel cue receives preferentially rare-outcome responding. Two other studies of the IBRE have also presented a novel cue at test. Johansen, Fouquet, and Shanks (2007) reported preferentially-rare responding, but did not statistically analyze those data, and used a procedure quite unlike those of other IBRE experiments. Don and Livesey (2017) reported preferentially-common responding, but they presented the novel cue in compound with familiar cue A, which makes their result hard to interpret. Again, further research is merited.
4.4 Conclusion

The current study raised some new questions for future research, but it also provided a clear answer to the question we set out to investigate. Specifically, the results of the current experiment largely do not support Le Pelley et al.’s explanation of the IBRE and COFED. As such, the results cast doubt on the generality of the rather simple model of the relationship between attention and learning outlined by Le Pelley et al.. In contrast, the slightly more complex theory of the relationship between learning and attention provided by the EXIT model (Kruschke 2001b) provides an excellent account of the current results.

References


**Appendix: Modelling**

These simulations were conducted using `slpEXIT`, part of the *catlearn* R package (Wills et al., 2019). This implementation of EXIT is based on the model as described in Kruschke (2001b), with the inclusion of a bias cue that was later implemented in Kruschke (2003). The bias cue was assumed to be analogous to the experimental context in Le Pelley et al.’s explanation. As such, two bias cues were implemented, one for each context, with the salience of these cues represented by the $\sigma$ parameter.

The EXIT model was applied to simulated training and test trials that replicated the details of experimental procedure, generating response patterns for each simulated trial. The values of the free parameters given to the model were varied using the `optim` function in R (R Core Team, 2018). The goal of this variation was to optimise the free parameters given to the model: in order to find the parameter set that when given to the model gave the closest approximation to the behavioural data. This was accomplished by calculating the sum of squared errors (SSE) between the response patterns generated by the model under a specific parameter set and the behavioural response patterns; `optim` was used to find the parameter set that minimised the SSE.

The method used for optimisation within `optim` was the limited memory Broyden-Fletcher-Goldfarb-Shanno (BFGS) algorithm (Byrd, Lu, Nocedal, & Zhu, 1995). As `optim` requires an initial set of parameters to vary, each free parameter within the EXIT model was initially set to one of two values. As there are 7 free parameters, this resulted in a total of $2^7$ or 128 sets of parameter values. Each of these starting parameter sets were supplied to `optim` individually. `optim` then used the BFGS algorithm to perform a hill-climbing optimisation and arrive at an optimised parameter set for each individual starting set. This produced 128 sets of optimised parameter values. These sets of optimised parameters values were compared in terms of the SSE generated when they were given to EXIT, in order to identify the set that produced the lowest SSE. The parameter values within this final optimised set for the experiment we report were: $c = .004$, $P = 4.490$, $\phi = 2.731$, $\lambda_g = 1.577$, $\lambda_w = .456$, $\lambda_x = .059$, $\sigma = .818$. 

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