

1 **Hierarchical Bayesian Augmented Hebbian Reweighting Model of Perceptual Learning**

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## Abstract

The Augmented Hebbian Reweighting Model (AHRM) has been effectively utilized to model the collective performance of observers in various perceptual learning studies. In this work, we have introduced a novel hierarchical Bayesian Augmented Hebbian Reweighting Model (HB-AHRM) to simultaneously model the learning curves of individual participants and the entire population within a single framework. We have compared its performance to that of a Bayesian Inference Procedure (BIP), which independently estimates the posterior distributions of model parameters for each individual subject without employing a hierarchical structure. To cope with the substantial computational demands, we developed an approach to approximate the likelihood function in the AHRM with feature engineering and linear regression, increasing the speed of the estimation procedure by 20,000 times. The HB-AHRM has enabled us to compute the joint posterior distribution of hyperparameters and parameters at the population, observer, and test levels, facilitating statistical inferences across these levels. While we have developed this methodology within the context of a single experiment, the HB-AHRM and the associated modeling techniques can be readily applied to analyze data from various perceptual learning experiments and provide predictions of human performance at both the population and individual levels. The likelihood approximation concept introduced in this study may have broader utility in fitting other stochastic models lacking analytic forms.

**Keywords:** Perceptual Learning, Augmented Hebbian Reweighting Model, Hierarchical Bayesian Model, Pytensor, Likelihood Approximation

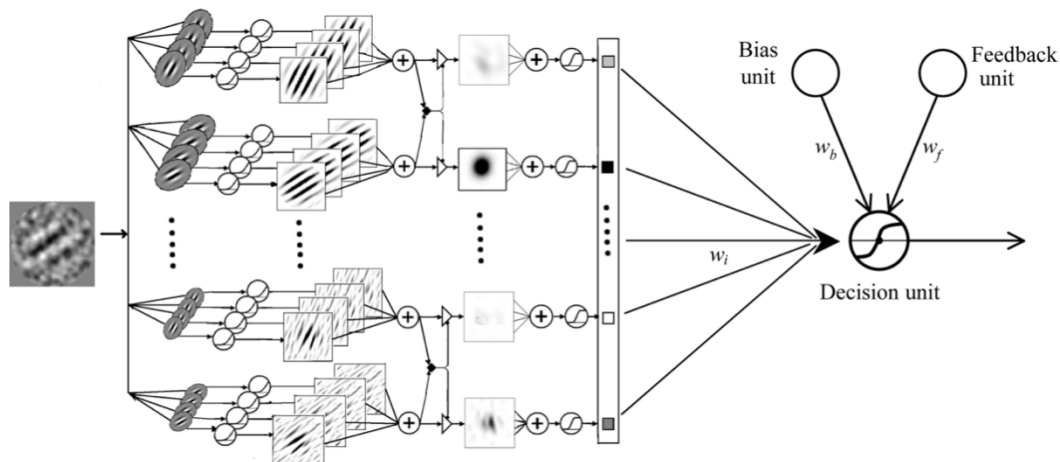
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## INTRODUCTION

41 Perceptual learning is a powerful process that can significantly enhance human performance  
42 in various perceptual tasks (Doshier & Lu, 2020; Fahle & Poggio, 2002; Green, Banai, Lu, &  
43 Bavelier, 2018; Lu & Doshier, 2022; Lu, Hua, Huang, Zhou, & Doshier, 2011; Sagi, 2011; Seitz,  
44 2017; T. Watanabe & Sasaki, 2015). It can lead to improvements in tasks such as orientation,  
45 spatial frequency, and motion direction judgements, taking performance from near chance to high  
46 proficiency (Ball & Sekuler, 1982; Fiorentini & Berardi, 1980; Poggio, Fahle, & Edelman, 1992).  
47 Contrast sensitivity can increase by over 100% (Doshier & Lu, 1998; Huang, Zhou, & Lu, 2008),  
48 and response times can decrease by approximately 40% (Petrov, Van Horn, & Ratcliff, 2011).  
49 Perceptual learning is increasingly being applied in rehabilitation and the development of  
50 perceptual expertise (Cavanaugh, 2015; L. Gu et al., 2020; Hess & Thompson, 2015; Huang et al.,  
51 2008; Huxlin et al., 2009; Levi, 2020; Lu, Lin, & Doshier, 2016; Maniglia, Visscher, & Seitz, 2021;  
52 Roberts & Carrasco, 2022; F.-F. Yan et al., 2015).

53 Two main theories, representation enhancement and selective reweighting, have been proposed  
54 to explain performance improvements in visual perceptual learning (Ahissar & Hochstein, 2004;  
55 Doshier & Lu, 1998, 2009b; Fahle, 1994; Karni & Sagi, 1991; Mollon & Danilova, 1996;  
56 Sotiropoulos, Seitz, & Seriès, 2011; Talluri, Hung, Seitz, & Seriès, 2015; T. Watanabe et al., 2002;  
57 T. Watanabe & Sasaki, 2015; Zhang et al., 2010). Representation enhancement suggests that  
58 perceptual learning improves performance by altering the responses or tuning characteristics of  
59 neurons in early visual cortical areas. On the other hand, selective reweighting involves the up-  
60 weighting of relevant and down-weighting of irrelevant representations from early visual cortical  
61 areas during perceptual decision without changing the representations themselves. While both  
62 processes can contribute to perceptual learning (Kourtzi, Betts, Sarkheil, & Welchman, 2005;

63 Roelfsema, van Ooyen, & Watanabe, 2010; Seitz & Watanabe, 2005; T. Watanabe & Sasaki,  
64 2015), selective reweighting appears to be the dominant component (Doshier & Lu, 2020). This  
65 conclusion is also supported by physiological and brain imaging studies, which indicate that early  
66 sensory representation enhancement accounts for only a small fraction of behavioral performance  
67 improvements (Ghose, Yang, & Maunsell, 2002; Schoups, Vogels, Qian, & Orban, 2001), while  
68 evidence of neural plasticity is strongest in higher visual areas (Adab & Vogels, 2011; Law &  
69 Gold, 2008; Y. Yan et al., 2014). Notably, representation enhancement remains primarily a verbal  
70 theory, and most existing computational models of visual perceptual learning are based on  
71 selective reweighting. These models aim to enhance the readout of sensory information during  
72 perceptual decision making (Doshier, Jeter, Liu, & Lu, 2013; Doshier & Lu, 1998; Jacobs, 2009;  
73 Law & Gold, 2009; Petrov, Doshier, & Lu, 2005; Poggio et al., 1992; Sotiropoulos et al., 2011;  
74 Vaina, Sundaeswaran, & Harris, 1995; Weiss, Edelman, & Fahle, 1993; Zhaoping, Herzog, &  
75 Dayan, 2003).



76  
77 Figure 1. The Augmented Hebbian Reweighting Model (AHRM) with representation, bias,  
78 feedback, and decision modules.  
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80 The Augmented Hebbian Reweighting Model (AHRM; Figure 1) is the first full computational  
81 process model of perceptual learning (Petrov et al., 2005). It comprises representation, bias,

82 feedback, and decision modules. The representation module computes activations in multiple  
83 orientation- and frequency-selective channels from stimulus images. The decision module weights  
84 and sums activations along with a bias module and yields a response on each trial. The learning  
85 module updates the weights to the decision module on every trial using augmented Hebbian  
86 learning, which moves the “late” post-synaptic activation in the decision module towards the  
87 correct response when feedback is available, and operates on the early decision activation when  
88 there is no feedback (Doshier & Lu, 2009a; Petrov et al., 2005; Petrov, Doshier, & Lu, 2006).

89 The AHRM has successfully modeled various phenomena in perceptual learning, including  
90 perceptual learning in nonstationary environments with and without feedback (Petrov et al., 2005,  
91 2006), basic mechanisms of perceptual learning, asymmetric transfer of learning in high and low  
92 external noise, and effects of pretraining mechanisms (Lu, Liu, & Doshier, 2010), co-learning of  
93 multiple tasks (Huang, Lu, & Doshier, 2012), interaction between training accuracy and feedback  
94 (Liu, Lu, & Doshier, 2010; Liu, Lu, & Doshier, 2012), and trial-by-trial and block feedback (Liu,  
95 Doshier, & Lu, 2014). It has also led to the development of several related models (Doshier et al.,  
96 2013; Jacobs, 2009; Law & Gold, 2009; Sotiropoulos et al., 2011; Talluri et al., 2015).

97 Despite its success, fitting the AHRM to data presents a significant challenge. The AHRM is  
98 a sequential stochastic learning model, that, with a given set of parameters, must be simulated to  
99 generate performance predictions with sequential trial-by-trial updates of the decision weights.  
100 Simulations typically involve running the model hundreds to thousands of times to generate  
101 average predictions and confidence bands for a given set of parameter values. For a fixed set of  
102 parameter values, each run of the model leads to a different sequence of responses and somewhat  
103 different weight changes due to stochastic trial-by-trial variations resulting from internal and  
104 external noises and different random trial sequences. Fitting the AHRM with typical curve fitting

105 procedures (e.g., maximum likelihood, least squares, Bayesian) is not feasible because the fitting  
106 process requires simulations of many potential parameter sets (tens to hundreds of thousands).  
107 Instead, estimation of the AHRM parameters is generally done using hierarchical grid-search  
108 methods. These methods evaluate a matrix of spaced parameter values and then narrow down  
109 regions of the parameter space that are more promising, making it difficult to obtain the optimal  
110 solutions.

111 In this study, we introduce three modeling technologies to facilitate AHRM fitting:

- 112 1) *A Hierarchical Bayesian AHRM (HB-AHRM)*: This approach incorporates  
113 population, subject, and test levels to estimate the joint posterior hyperparameter and  
114 parameter distribution across all subjects while considering covariance within and  
115 between subjects.
- 116 2) *Vectorization with PyTensor*: Leveraging PyTensor library and GPU acceleration,  
117 these techniques drastically speed up simulations by optimizing the computation of  
118 mathematical expressions involving multi-dimensional arrays.
- 119 3) *Likelihood function approximation*: We developed an approach to approximate the  
120 likelihood function in the AHRM with feature engineering and linear regression.  
121 Based on simulated predictions of the AHRM over a large parameter grid, we  
122 encoded the functional relationship between the likelihood and parameters, greatly  
123 accelerating model computations.

124 Hierarchical models enable effective combination of information across subjects and groups  
125 while preserving heterogeneity (Kruschke, 2014; Rouder & Lu, 2005). These models typically  
126 consist of sub-models and probability distributions at multiple levels of the hierarchy and can  
127 compute the joint posterior distributions of the parameters and hyperparameters using Bayes'

128 theorem based on all available data (Kruschke, 2014; Kruschke & Liddell, 2018). Hierarchical  
129 models are valuable for reducing the variance of estimated posterior distributions by decomposing  
130 variabilities from different sources using parameters and hyperparameters (Song et al., 2020) and  
131 shrinking estimated parameters at lower levels towards the modes of higher levels when there is  
132 insufficient data at the lower level (Kruschke, 2014; Rouder et al., 2003; Rouder & Lu, 2005).

133 The HB-AHRM consists of three levels: population, subject, and test. In this framework, all  
134 subjects belong to a population and may, in principle, run the same experiment (called “test”)  
135 multiple times. The distributions of AHRM parameters at the test level are conditioned on the  
136 hyperparameter distributions at the subject level, which, in turn, are conditioned on the  
137 hyperparameter distribution at the population level. The HB-AHRM also includes covariance  
138 hyperparameters at the population and subject levels to capture the relationship between and within  
139 subjects.

140 PyTensor is a Python library used to define, optimize, rewrite, and evaluate mathematical  
141 expressions, particularly those involving multi-dimensional arrays. It combines elements of a  
142 computer algebra system and an optimizing compiler. PyTensor is particularly useful for tasks  
143 where complex mathematical expressions need repeated evaluation, and speed is critical. The  
144 library provides a loop mechanism called *scan*, which can process inputs efficiently. We used  
145 PyTensor to represent all variables in the HB-AHRM and applied the *scan* function, significantly  
146 speeding up simulations from 22.2 to 1.6 seconds for 300 repeated runs of the experiment in Petrov  
147 et al. (2005) based on one set of AHRM parameters.

148 Although PyTensor improved simulation speed, computing the HB-AHRM still involves  
149 evaluating of hundreds of thousands of parameter sets. Because of the tremendous computational  
150 load, we developed a method to approximate the likelihood function in the AHRM with feature

151 engineering and linear regression. It involves simulating AHRM predictions in a large parameter  
152 grid using parallel computing with GPU processors, taking <24 hours for a 64,000 mesh grid. We  
153 then employed feature engineering and linear regression to learn the relationship between the  
154 likelihood function and AHRM parameters, which took about 30 minutes. The differentiable  
155 functional relationship enabled efficient exploration of a large parameter space in fitting the  
156 models (<1 ms per sample).

157 In this paper, we provide an overview of the AHRM as a generative model of trial-by-trial  
158 human performance in perceptual learning. We also introduce a Bayesian inference procedure  
159 (BIP) used to independently estimate the posterior distribution of AHRM parameters for each  
160 subject. Subsequently, we present the HB-AHRM, designed to collectively estimate the joint  
161 posterior distribution of hyperparameters and parameters at multiple levels of the hierarchy. We  
162 discuss the simulation technologies, including PyTensor, and the method for likelihood function  
163 approximation. These procedures are applied to data from Petrov et al. (2005). Our analysis  
164 involves comparing the goodness of fit the BIP and HB-AHRM, and evaluating the variability of  
165 estimated AHRM parameters, learning curves and weight structures. In addition, we conducted a  
166 simulation study to evaluate parameter recovery and HB-AHRM's ability in predicting the  
167 performance of a new simulated observer with no or limited training data.

## 168 THEORETICAL DEVELOPMENT

### 169 *The Augmented Hebbian Reweighting Model (AHRM)*

170 In this section, we briefly describe the augmented Hebbian reweighting model (AHRM). More  
171 details of the model can be found in the original paper (Petrov et al., 2005).

172 Representation module. For subject  $i$  in test  $j$ , the stimulus image consists of a signal image  $S_{ijt}$   
173 and an external noise image  $N_{ijt}$  in each trial  $t$ . The representation module encodes the stimulus



174 image into expected activations over a bank of orientation and spatial-frequency channels tuned  
 175 to different orientations  $\varphi$  and spatial frequencies  $f$ ,  $E(\varphi, f | S_{ijt}, N_{ijt})$ , through convolution,  
 176 halfwave rectification, contrast normalization and pooling over phase and space (Petrov et al.,  
 177 2005). We consider 35 *channels* centered at 7 orientations ( $\varphi \in [0^\circ, \pm 15^\circ, \pm 30^\circ, \pm 45^\circ]$ ) and 5  
 178 spatial frequencies ( $f \in [1, 1.4, 2, 2.8, 4]$  cycles per degree). The expected activation is then  
 179 combined with internal noise  $\varepsilon_{\varphi, f}$  (with mean 0 and standard deviation  $\sigma_r$ ) and passed to a  
 180 saturating non-linearity to compute the activation in each channel:

$$181 \quad A'(\varphi, f | S_{ijt}, N_{ijt}) = E(\varphi, f | S_{ijt}, N_{ijt}) + \varepsilon_{\varphi, f}, \quad (1)$$

$$182 \quad A(\varphi, f | S_{ijt}, N_{ijt}) = A_{max} \begin{cases} \frac{1 - e^{-\gamma A'(\varphi, f | S_{ijt}, N_{ijt})}}{1 + e^{-\gamma A'(\varphi, f | S_{ijt}, N_{ijt})}}, & \text{if } A'(\varphi, f | S_{ijt}, N_{ijt}) \geq 0 \\ 0, & \text{otherwise} \end{cases} \quad (2)$$

183 *Task-specific decision module.* The decision module weighs the evidence in the noisy activations  
 184 from the representation module to generate a response in each trial. Specifically, it first  
 185 aggregates the activation pattern  $A(\varphi, f | S_{ijt}, N_{ijt})$  over the orientation and spatial-frequency  
 186 channels using current weights  $w_{\varphi, f}(t)$ , a top-down bias  $b(t)$ , and a Gaussian decision noise  $\varepsilon$   
 187 (mean 0 and standard deviation  $\sigma_d$ ):

$$188 \quad u(t) = \sum_{\varphi, f} w_{\varphi, f}(t) A(\varphi, f | S_{ijt}, N_{ijt}) - b(t) + \varepsilon. \quad (3)$$

189 and then computes its output  $o(t)$  as a sigmoidal function of  $u(t)$ :

$$190 \quad o(t) = G(u(t)), \quad (4)$$

$$191 \quad G(u(t)) = \frac{1 - e^{-\gamma u(t)}}{1 + e^{-\gamma u(t)}} A_{max}. \quad (5)$$

192 The decision variable  $o(t)$  saturates at  $\pm A_{max} = \pm 0.5$ ; the model responds left or  
 193 counterclockwise if  $o(t) < 0$ , and right or clockwise otherwise.

194 The weights are initiated to be proportional to the preferred orientation of the representation  
195 module relative to the vertical:  $w_{\varphi,f}(0) = (\varphi/30^\circ)w_{init}$ .

196 Learning module. The weights between the representation and decision modules are updated on  
197 each trial using an augmented Hebbian reweighting rule, in which feedback,  $F(t) = \pm 0.5$ , is  
198 used as the correct output of the decision module. The change of weight  $w_{\varphi,f}(t)$  in each channel  
199 depends on the activation of the representation  $A(\varphi, f|S_{ijt}, N_{ijt})$ , the correct output of the  
200 decision module and the internal learning rate  $\eta$ :

$$201 \quad \delta_{\varphi,f} = \alpha A(\varphi, f|S_{ijt}, N_{ijt})F(t), \quad (6a)$$

$$202 \quad \Delta w_{\varphi,f}(t) = (w_{\varphi,f}(t) - w_{min})[\delta_{\varphi,f}]_- + (w_{max} - w_{\varphi,f}(t))[\delta_{\varphi,f}]_+, \quad (6b)$$

203 where  $w_{min} = -1$  and  $w_{max} = 1.0$ .

204 Adaptive criterion control. The adaptive criterion control module shifts the decision variable on  
205 each trial to compensate for biases in the immediate history of responses by adding a bias  
206 correcting term to the activation at the decision module. It assumes that observers monitor their  
207 own responses and seek to equalize response frequencies—trying to match stimulus probabilities  
208 that are often balanced in experiments. A weighted running average of recent responses  
209 exponentially discounts the distant past response history:

$$210 \quad r(t+1) = \rho R_{ijt} + (1 - \rho)r(t), \quad (7a)$$

$$211 \quad b(t+1) = \beta r(t), \quad (7b)$$

212 where  $R(t)$  is the response in the current trial, and  $r(t)$  is the weighted running average responses,  
213 and  $b(t)$  is the bias. Following Petrov et al. (2005), we set  $\rho=0.02$ .

214 In summary, the AHRM has six free parameters (Table 1), internal learning rate  $\alpha$ , bias  
215 strength  $\beta$ , activation function gain  $\gamma$ , standard deviation of decision noise  $\sigma_d$ , standard deviation

216 of representation noise  $\sigma_r$ , and initial weight scaling factor  $w_{init}$ . Additional parameters, including  
 217 maximum activation level, weight bounds, orientation spacing, and spatial frequency spacing, are  
 218 fixed.

219 **Table 1:** AHRM parameters and their corresponding symbols in the BIP and HB-AHRM

Parameters	AHRM	BIP/HB-AHRM	Values
Learning rate	$\alpha$	$\theta_{ij1}$	
Bias strength	$\beta$	$\theta_{ij2}$	
Activation function gain	$\gamma$	$\theta_{ij3}$	
Decision noise	$\sigma_d$	$\theta_{ij4}$	
Representation noise	$\sigma_r$	$\theta_{ij5}$	
Initial weight scaling factor	$w_{init}$	$\theta_{ij6}$	
Maximum activation level	$A_{max}$		0.5
Weight bounds	$w_{min/max}$		$\pm 1$
Orientation spacing	$\Delta\varphi$		$15^\circ$
Spatial frequency spacing	$\Delta f$		0.5 oct

220  
 221 To simplify notations, we use  $\theta_{ij}$  to denote the AHRM parameters for subject  $i$  in test  $j$  (see  
 222 Table 1 for the correspondence with the original AHRM parameters). For a given subject  $i$  in test  
 223  $j$  with AHRM parameters  $\theta_{ij}$ , we can compute the probability of obtaining a correct response in  
 224 trial  $t$ ,  $p(R_{ijt} = 1 | \theta_{ij}, S_{ijt}, N_{ijt})$ , and the probability of obtaining an incorrect response in trial  $t$ ,  
 225  $p(R_{ijt} = 0 | \theta_{ij}, S_{ijt}, N_{ijt})$ , from repeated simulations of the AHRM. The two probabilities define  
 226 the likelihood function for each of  $T$  trials. The likelihood of obtaining all the observed responses  
 227 of subject  $i$  in test  $j$  is the product of all the trial-by-trial likelihoods:

$$228 \quad p(R_{ij1:T} | \theta_{ij}) = \prod_{t=1}^T p(R_{ijt} | \theta_{ij}, S_{ijt}, N_{ijt}), \quad (8)$$

### 229 **Bayesian Inference Procedure**

230 The Bayesian Inference Procedure (BIP) is used to estimate the posterior distribution of  $\theta_{ij}$   
 231 from the trial-by-trial data  $Y_{ij} = \{(R_{ij1:T}, S_{ij1:T}, N_{ij1:T})\}$  of subject  $i$  in test  $j$  via Bayes' rule  
 232 (Figure 2a):

$$p(\theta_{ij}|Y_{ij}) = \frac{\prod_{t=1}^T p(R_{ijt}|\theta_{ij}, S_{ijt}, N_{ijt}) p_0(\theta_{ij})}{\int \prod_{t=1}^T p(R_{ijt}|\theta_{ij}, S_{ijt}, N_{ijt}) p_0(\theta_{ij}) d\theta_{ij}}. \quad (9)$$

Here,  $p(\theta_{ij}|Y_{ij})$  is the posterior distribution of AHRM parameters,  $\theta_{ij}$ , given the trial-by-trial data  $Y_{ij}$ ,  $p(R_{ijt}|\theta_{ij}, S_{ijt}, N_{ijt})$  is the likelihood term, which quantifies the probability of observing responses  $R_{ijt}$  given  $\theta_{ij}$ ,  $S_{ijt}$ , and  $N_{ijt}$ ,  $p_0(\theta_{ij})$  is the prior probability distribution of  $\theta_{ij}$ .

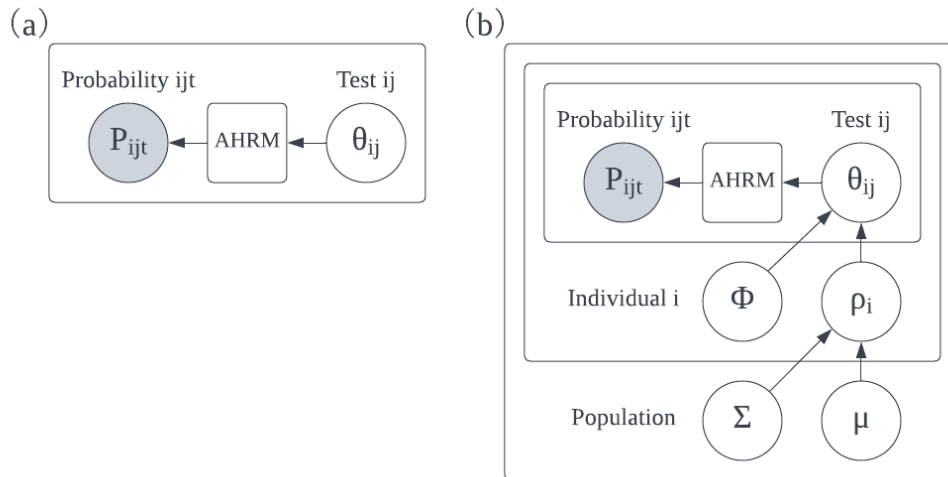


Figure 2. (a) The Bayesian inference procedure (BIP). For a given subject  $i$  in test  $j$  with parameters  $\theta_{ij}$ , the likelihood of obtaining response  $p(R_{ijt})$  in trial  $t$  is computed from the AHRM. (b) The HB-AHRM is a three-level hierarchical Bayesian model in which the population level hyperparameter  $\eta$  is modeled as a mixture of Gaussian distributions with mean  $\mu$  and covariance  $\Sigma$ , hyperparameter  $\tau_i$  at the subject level is modeled as a mixture of Gaussian distributions with mean  $\rho_i$  and covariance  $\phi$ , and the probability distribution of parameters  $\theta_{ij}$  is conditioned on  $\tau_i$ .

The prior of  $\theta_{ij}$  is set as a uniform distribution in all its dimension:

$$p_0(\theta_{ijk}) = \mathcal{U}(\theta_{0k,min}, \theta_{0k,max}), \quad (10)$$

where  $\theta_{0k,min}$  and  $\theta_{0k,max}$  are the lower and upper bounds of the uniform distribution for dimension  $k$  (Table 2), which are set based on observed values in prior applications of the model.

The denominator of equation (9) is an integral across all possible values of  $\theta_{ij}$ .

In the BIP, the AHRM parameters are estimated independently for each subject.

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**Table 2.** Lower and upper bounds of the priors.

$k$	1	2	3	4	5	6
$\theta_{0k,min}$	0	0	0	0	0	0
$\theta_{0k,max}$	0.003	3	3	0.3	0.1	0.3

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256 ***Hierarchical Bayesian Augmented Hebbian Reweighting Model (HB-AHRM)***

257 The HB-AHRM is a three-level hierarchical Bayesian model used to estimate the joint  
 258 posterior hyperparameter and parameter distribution in all levels, considering covariance within  
 259 and between subjects (Figure 2b). The HB-AHRM includes probability distributions at the  
 260 population, subject, and test levels.

261 Population level. The probability distribution of the six-dimensional hyperparameter  $\eta$  of the  
 262 AHRM parameters (Table 1) at the population level is modeled as a mixture of six-dimensional  
 263 Gaussian distributions with mean  $\mu$  and covariance  $\Sigma$ , which have distributions  $p(\mu)$  and  $p(\Sigma)$ :

$$264 \quad p(\eta) = \mathcal{N}(\eta, \mu, \Sigma)p(\mu)p(\Sigma). \quad (11)$$

265 Subject level. The probability distribution of hyperparameter  $\tau_i$  for subject  $i$  at the subject level is  
 266 modeled as a mixture of 6-dimensional Gaussian distributions with mean  $\rho_i$  and covariance  $\phi$ ,  
 267 with distributions  $p(\rho_i|\eta)$  and  $p(\phi)$ :

$$268 \quad p(\tau_i|\eta) = \mathcal{N}(\tau_i, \rho_i, \phi) p(\rho_i|\eta)p(\phi), \quad (12)$$

269 in which  $\rho_i$  is conditioned on  $\eta$ .

270 Test level.  $p(\theta_{ij}|\tau_i)$ , the probability distribution of parameters  $\theta_{ij}$  is conditioned on  $\tau_i$ . The  
 271 probability of obtaining the entire dataset is computed using probability multiplication, which  
 272 involves all levels of the model and the likelihood function based on the trial-by-trial data:

$$273 \quad p(Y_{1:l,1:j}|X) = \prod_{i=1}^l \prod_{j=1}^J \prod_{t=1}^T p(R_{ijt}|\theta_{ij}, S_{ijt}, N_{ijt})p(\theta_{ij}|\tau_i) p(\tau_i|\eta)p(\eta)$$

$$274 \quad = \prod_{i=1}^l \prod_{j=1}^J \prod_{t=1}^T p(R_{ijt}|\theta_{ij}, S_{ijt}, N_{ijt})p(\theta_{ij}|\tau_i) \mathcal{N}(\tau_i, \rho_i, \phi) p(\rho_i|\eta)p(\phi)\mathcal{N}(\eta, \mu, \Sigma)p(\mu)p(\Sigma), \quad (13)$$

275 where  $X = (\theta_{1:l,1:j}, \rho_{1:l}, \mu, \phi, \Sigma)$  are all the parameters and hyperparameters in the HB-AHRM.

276 Bayes' rule is used to compute the joint posterior distribution of  $X$ , which includes all HB-  
277 AHRM parameters and hyperparameters:

$$278 \quad p(X|Y_{1:l,1:j}) = \frac{\prod_{i=1}^l \prod_{j=1}^J \prod_{t=1}^T p(R_{ijt}|\theta_{ij}, S_{ijt}, N_{ijt}) p(\theta_{ij}|\tau_i) \mathcal{N}(\tau_i, \rho_i, \phi) p(\rho_i|\eta) p_0(\phi) \mathcal{N}(\eta, \mu, \Sigma) p_0(\mu) p_0(\Sigma)}{\int \prod_{i=1}^l \prod_{j=1}^J \prod_{t=1}^T p(R_{ijt}|\theta_{ij}, S_{ijt}, N_{ijt}) p(\theta_{ij}|\tau_i) \mathcal{N}(\tau_i, \rho_i, \phi) p(\rho_i|\eta) p_0(\phi) \mathcal{N}(\eta, \mu, \Sigma) p_0(\mu) p_0(\Sigma) dX}, \quad (14)$$

279 where the denominator is an integral across all possible values of  $X$  and is a constant for a given  
280 dataset and HB-AHRM;  $p_0(\mu)$ ,  $p_0(\Sigma)$ , and  $p_0(\phi)$  are the prior distributions of  $\mu$ ,  $\Sigma$ , and  $\phi$ , with

$$281 \quad p_0(\mu) = \mathcal{U}_k(\theta_{0k,min}, \theta_{0k,max}), \quad (15)$$

282 where  $\mathcal{U}_k(\theta_{0k,min}, \theta_{0k,max})$  denotes a uniform distribution between  $\theta_{0k,min}$  and  $\theta_{0k,max}$  in each  
283 of the six dimensions, with  $\theta_{0k,min}$  and  $\theta_{0k,max}$  defined in Table 2. Both  $p_0(\Sigma)$  and  $p_0(\phi)$  are  
284 set with the LKJ distribution with a shape parameter of 2.0 (Lewandowski, Kurowicka, & Joe,  
285 2009).

286 Equation 14 allows us to estimate the joint posterior distribution of HB-AHRM parameters and  
287 hyperparameters across all tests and subjects. Unlike the BIP, the HB-AHRM hyperparameters  
288 and parameter estimates mutually constrain each other across tests and subjects via the joint  
289 distribution. This allows for more robust and interconnected estimates of HB-AHRM parameters  
290 and hyperparameters.

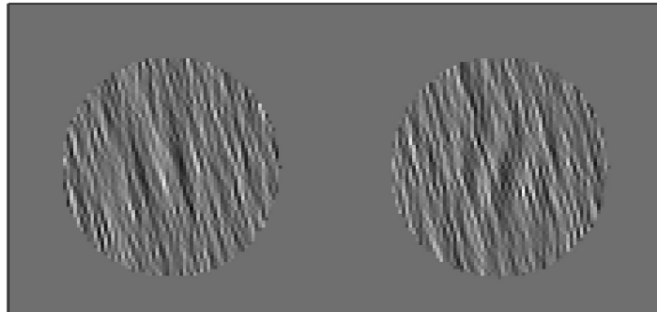
## 291 **STUDY 1. RE-ANALYSIS OF Petrov et al. (2005)**

### 292 ***Methods***

293 Data. Petrov et al. (2005) investigated perceptual learning in an orientation identification task  
294 involving 13 adult subjects with normal or corrected-to-normal vision. Subjects judged the  
295 orientation ( $\pm 10^\circ$  from vertical) of Gabor patches (windowed sinusoidal gratings, peak spatial  
296 frequency=2 c/d) in each trial. The experiment included a nonstationary context where external

297 noise images, predominantly oriented left in context L and right in context R, were superimposed  
298 on the target stimuli (Figure 3).

299 The study consisted of eight daily sessions, each with four blocks of 300 trials, totaling 9600  
300 trials. Subjects were trained in block sequences of either L-8R-8L-8R-6L-R (7 subjects) or R-8L-  
301 8R-8L-6R-L (6 subjects) contexts. Context congruency was randomly selected, with the target  
302 Gabor and context either congruent or incongruent in orientation. Gabor contrast was randomly  
303 selected from three fixed levels (0.106, 0.160, and 0.245). The resulting behavioral data shows a  
304 complex pattern related to congruency and contrast.



305  
306 Figure 3. An illustration of left and right titled Gabors in context L.  
307

308 The study adhered to ethical standards, with written consent obtained from all subjects prior to  
309 the experiment. The research protocol received approval from the institutional review board for  
310 human subject research at the University of California, Irvine, and complied with the principles of  
311 the Declaration of Helsinki.

312 Likelihood function approximation. Fitting the BIP and HB-AHRM to the data involves using  
313 simulations to evaluate the likelihood of a vast set of model parameters. Due to the significant  
314 computation time, previous studies relied on grid search methods for AHRM evaluation (Petrov,  
315 et al., 2005; 2006). To reduce the computational cost, we approximated the likelihood function  
316 by learning its functional relationship with AHRM parameters based on simulated predictions  
317 over a large parameter grid. This facilitated fitting the BIP and HB-AHRM models.

318 We constructed a 6-dimensional mesh grid  $\Theta$  (Table 3) to train the functional relationship  
 319 between AHRM parameters and the likelihood function. The mesh grid contained  $8 \times 8 \times 8 \times 5 \times$   
 320  $5 \times 5 = 64,000$  sets of model parameters. The ranges and values of the AHRM parameters were  
 321 chosen based on an exploration of model predictions with various parameters.

322  
 323

**Table 3.** Mesh grid used in the simulation.

Parameter		Values							
$\alpha$	$\theta_{ij1}$	0.0001	0.0004	0.0008	0.0012	0.0016	0.0020	0.0024	0.0028
$\beta$	$\theta_{ij2}$	0.1	0.4	0.8	1.2	1.6	2.0	2.4	2.8
$\gamma$	$\theta_{ij3}$	0.1	0.3	0.6	0.9	1.2	1.5	1.8	2.1
$\sigma_d$	$\theta_{ij4}$	0.05	0.10	0.15	0.20	0.25			
$\sigma_r$	$\theta_{ij5}$	0.025	0.050	0.075	0.100	0.125			
$w_{init}$	$\theta_{ij6}$	0.05	0.10	0.15	0.20	0.25			

324

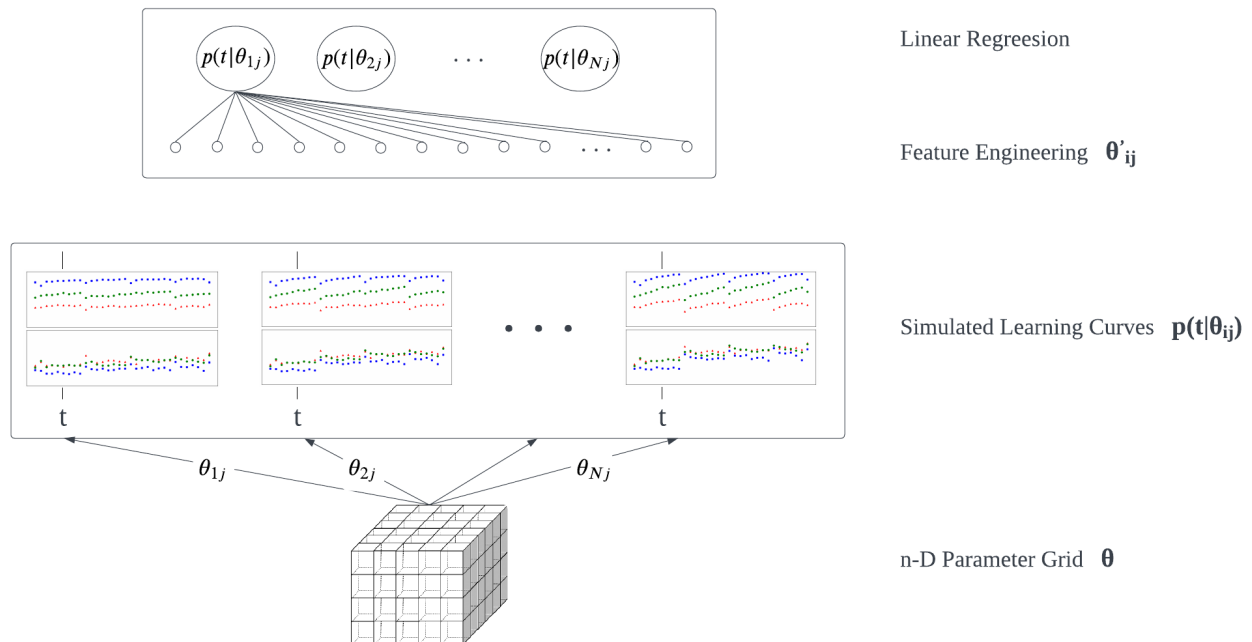
325 We calculated the likelihood, representing the trial-by-trial probability of a correct response,  
 326 for each set of AHRM parameters  $\theta_{ij}$  across six stimulus conditions over 9600 trials. This  
 327 computation was based on the average of five simulations, each comprising 300 repeated runs with  
 328 the same AHRM parameters and a different trial sequence.

329 The AHRM was used to generate trial-by-trial response based on the set of parameters and the  
 330 stimulus sequence, using Pytensor library's *scan* function. Because the exact external noise image  
 331 on each trial was not available, we obtained a cache of 1200 expected 35-dimensional activations,  
 332  $E(\varphi, f | S_{ijt}, N_{ijt})$ , consisting of 100 random samples of the 12 combinations of 2 (context)  $\times$  3  
 333 (Gabor contrast)  $\times$  2 (Gabor orientation). In each of the 300 runs, the AHRM starts with the same  
 334 initial weights and no decision bias, generating orientation judgements in eight sessions with four  
 335 blocks of 300 trials each. The contexts of the blocks were arranged in terms of L-8R-8L-8R-6L-R  
 336 or R-8L-8R-8L-6R-L, totaling 9600 trials. Each block of 300 trials consisted of 50 trials in each



337 of the 2 (congruency)  $\times$  3 (Gabor contrast) conditions, with a random permutation of the trial  
 338 types.

339 Each simulation took 1.6 seconds, in contrast to 22.2 seconds when using a for loop.  
 340 Additionally, we averaged the likelihoods within each block of 300 trials, resulting in one  
 341 likelihood for each of the six conditions per block for each set of parameters, and 64,000  
 342 likelihoods for the 64,000 sets of AHRM parameters in each of the six conditions per block.



343 Figure 4. Approximating the likelihood function with feature engineering and linear regression.  
 344 The simulated learning curves show data for incongruent (top) and congruent (bottom) trials at  
 345 three contrast levels (colors) over training blocks for each parameter combination.  
 346  
 347

348 In each block of the six experimental conditions, we computed the functional relationship  
 349 between the likelihood and AHRM parameters using feature engineering and linear regression in  
 350 the Scikit-learn library. Across blocks and experimental conditions, we established a total of 192  
 351 functional relationships.

352 For each relationship, the 64 predictors  $\theta'_{ij}$  included an intercept, the six AHRM parameters  
 353  $\theta_{ij}$ , and 57 additional features obtained through comprehensive feature engineering by exploring

354 all 21 quadratic and 36 cubic terms created from the six AHRM parameters. For each block of  
355 trials  $t^1$ , the linear regression is expressed as:

356 
$$p(t|\theta_{ij})=a(t) + \sum_{l=1}^{64} b_k(t)\theta'_{ijk}. \quad (16)$$

357 The feature engineering and linear regression step took about half an hour.

358 The functional relationship for the six conditions in 32 blocks is encoded in a 192 x 64 matrix,  
359 where each row represents the coefficients in one experimental condition in one block. This  
360 coefficient matrix allows us to calculate the likelihood function  $p(R_{ijt} = 1|\theta_{ij}, S_{ijt}, N_{ijt})$  for any  
361 AHRM parameters within the mesh grid range, not just the parameter sets in the mesh grid.  
362 Moreover, the likelihood functions are differentiable, facilitating various inference functions in  
363 *PYMC* (Abril-Pla et al., 2023).

364 *Estimating the Posterior Distributions.* We utilized the Automatic Differentiation Variational  
365 Inference (ADVI) method in the *PYMC* library to generate representative samples of the  
366 posterior distributions in the BIP and HB-ARHM. In this method, the variational posterior  
367 distribution is assumed to be spherical Gaussian without correlation between parameters and fit  
368 to the true posterior distribution. The means and standard deviations of the variational posterior  
369 are referred to as variational parameters.

370 We ran ADVI optimization for 300,000 iterations in the BIP and HB-AHRM to ensure good  
371 approximations of the posterior distributions. To generate representative samples of the posterior  
372 distribution in the BIP, we used the ADVI method to generate 100,000 samples for each subject  $i$ .  
373 Similarly, we computed 100,000 representative samples of the joint posterior distribution of  $\mu$  (6

---

<sup>1</sup> Although the BIP and HBM are formulated to model trial-by-trial data, we use  $t$  to denote each block of 300 trials in the rest of the paper.

374 parameters),  $\Sigma$  (21 parameters),  $\rho_{ik}$  ( $6 \times 13 = 78$  parameters),  $\phi$  (21 parameters), and  $\theta_{ijk}$  ( $6 \times$   
375  $13 = 78$  parameters). A model is considered “converged” when the Evidence Lower Bound  
376 (ELBO) stabilizes during iterations, indicating that the variational posterior has adequately  
377 approximated the true posterior distribution.

378 Goodness of Fit. We used the Watanabe–Akaike information criterion (WAIC) to compare the  
379 BIP and HB-AHRM fits. WAIC quantifies the likelihood of the observed data based on the joint  
380 posterior distribution of model parameters while penalizing for model complexity (S. Watanabe  
381 & Opper, 2010). Additionally, we assessed the accuracy of model predictions with *RMSE*, the  
382 proportion of variance in the observed data explained by the model ( $R^2$ ), and the uncertainty of  
383 the parameter estimates and model predictions with estimated credible intervals.

384 The *RMSE* between the predicted and observed quantities is defined as:

$$385 \quad \text{RMSE} = \sqrt{\sum (y_m - \hat{y}_m)^2 / M}, \quad (17)$$

386 where  $y_m$  is the observed value,  $\hat{y}_m$  is the predicted value,  $\bar{y}$  is the mean of all the observations,  
387 and  $M$  is the total number of observations.

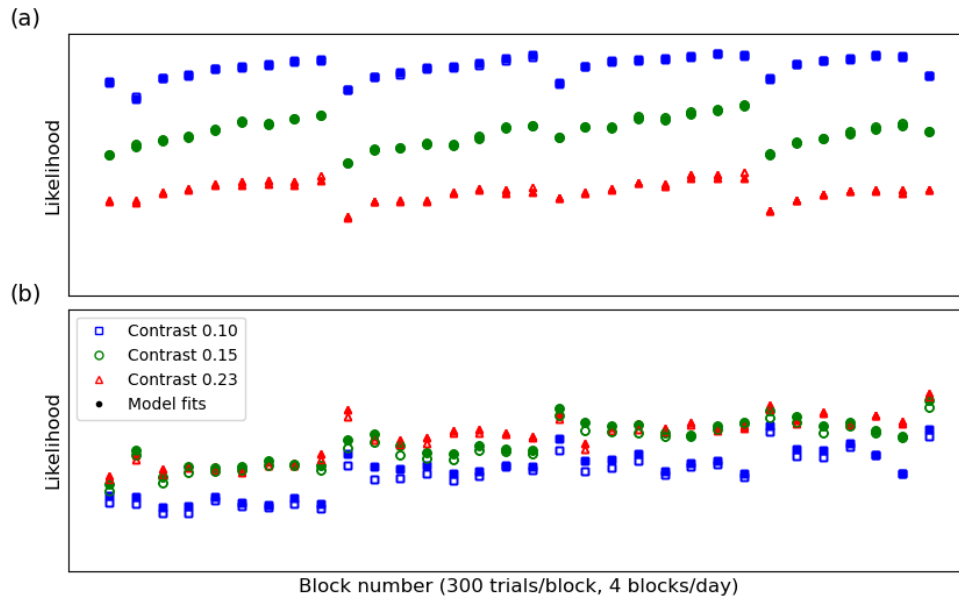
388 The proportion of variance accounted for,  $R^2$ , is defined as:

$$389 \quad R^2 = 1 - \frac{\sum (y_m - \hat{y}_m)^2 / M}{\sum (y_m - \bar{y})^2 / M}, \quad (18)$$

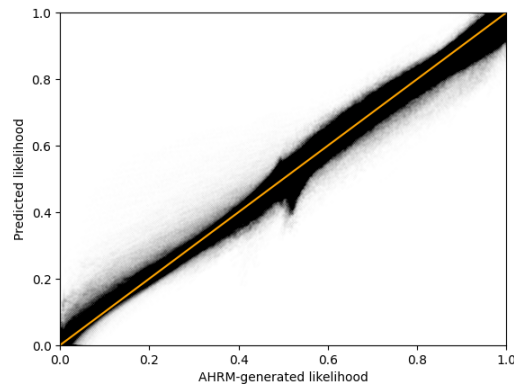
## 390 **Results**

391 Likelihood function approximation. Figure 5 shows the predicted likelihood function of the  
392 AHRM for one set of parameters:  $\alpha = 0.0008$ ,  $\beta = 1.8$ ,  $\gamma = 1.2$ ,  $\sigma_d = 0.2$ ,  $\sigma_r = 0.1$ , and  
393  $w_{init} = 0.17$ . The model predictions exhibit characteristic patterns observed in Petrov et al.  
394 (2005): general learning, persistent switch costs, and within context rapid relearning.

395 Figure 6 shows a scatter plot of the approximate likelihoods from feature engineering and linear  
396 regression against likelihoods generated from the AHRM across all 64,000 sets of AHRM  
397 parameters in the mesh grid, with an  $R^2$  of 0.991 and an RMSE of 0.016, indicating excellent  
398 approximation of the likelihoods by the linear model (Eq. 16).



399 Figure 5. Predicted likelihood function of the AHRM for one set of AHRM parameters ( $\alpha =$   
400  $0.0008, \beta = 1.8, \gamma = 1.2, \sigma_d = 0.2, \sigma_r = 0.1, w_{init} = 0.17$ ) from the simulations.  
401  
402



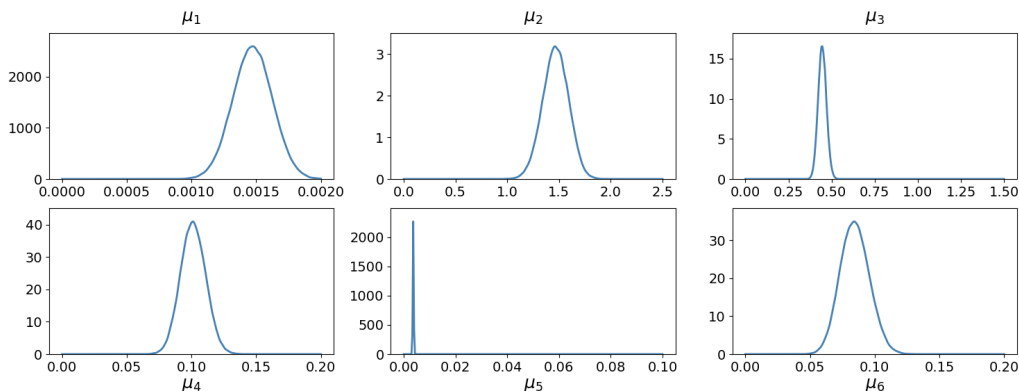
403  
404 Figure 6. A scatter plot of the approximate likelihoods from feature engineering and linear  
405 regression against likelihoods generated from the AHRM.

406 BIP and HB-AHRM Comparison. Both the BIP and HB-AHRM converged, indicated by the  
407 stabilization of ELBO.  
408

409 The WAIC values of the BIP and HB-AHRM were  $-7908.7 \pm 92.5$  and  $-8754.1 \pm 153.3$ ,  
 410 respectively, with a difference of  $-845.4 \pm 179.0$ . The HB-AHRM provided a significantly better  
 411 fit to the data. We will focus on the results from the HB-AHRM in the main body of the paper.  
 412 Detailed results from the BIP can be found in Supplementary Materials A.

413 Posterior Distributions (HB-AHRM). The marginal posterior distributions of the population-level  
 414 hyperparameter  $\eta$  are depicted in Figure 7. The mean and 94% half width credible interval  
 415 (HWCI) of these distributions are listed in Table 4. For most  $\eta$  components, except  $\eta_5$   
 416 (representation noise), the HWCI was quite small relative to their respective mean.

417 The expected correlation matrix derived from the expected covariance hyperparameter  $\Sigma$  is  
 418 shown in Table 5. The expected between-subject correlations among  $\eta$  components were quite  
 419 small, and none of them was significantly different from zero, due to the relatively small range of  
 420 performance variations across the 13 subjects.



421  
 422 Figure 7: Marginal posterior distributions of the population-level hyperparameter  $\eta$ .  
 423

424 **Table 4.** Mean and 94% HWCI of the marginal distributions of  $\eta$ .

	$\eta_1$	$\eta_2$	$\eta_3$	$\eta_4$	$\eta_5$	$\eta_6$
Mean	0.0014	1.71	0.46	0.10	0.004	0.08
HWCI	0.0001	0.13	0.02	0.01	0.002	0.01

425

426

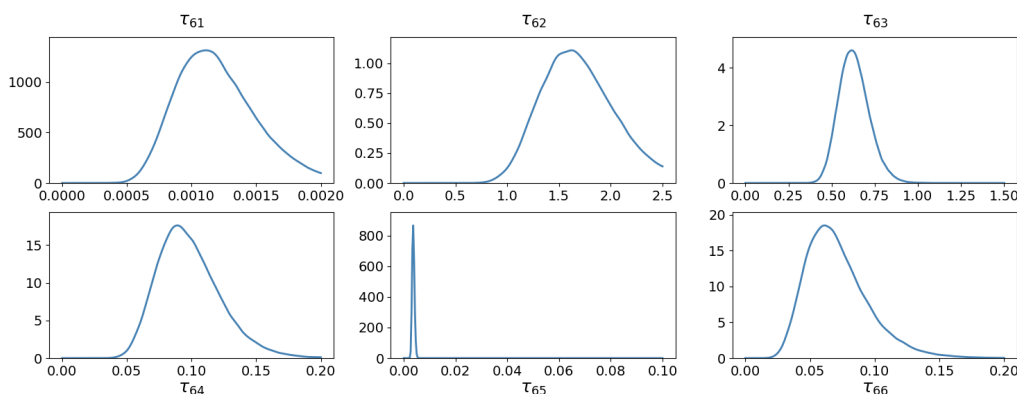
**Table 5.** Expected correlation matrix at the population level.

1	0.037	-0.006	-0.01	0.036	-0.05
0.037	1	-0.031	0.006	0.034	-0.024
-0.006	-0.031	1	0.023	-0.044	-0.006
-0.01	0.006	0.023	1	0.024	0.02
0.036	0.034	-0.044	0.024	1	0.029
-0.05	-0.024	-0.006	0.02	0.029	1

427

428 The marginal posterior distributions of the subject-level hyperparameter  $\tau_i$  for a typical subject  
 429 ( $i = 6$ ) are depicted in Figure 8. The mean and 94% HWCI of these distributions are listed in Table  
 430 6. Compared to  $\eta$ ,  $\tau_6$  components exhibited higher uncertainties, a pattern held across all subjects.  
 431 The full table with the mean and HWCI of the marginal distributions for all 13 subjects is available  
 432 in Supplementary Materials B.

433 The expected correlation matrix derived from the expected covariance hyperparameter  $\Phi$  is  
 434 shown in Table 7. The expected correlations between  $\tau_{i2}$  (bias strength) and  $\tau_{i4}$  (decision noise)  
 435 was -0.201 and between components  $\tau_{i4}$  (decision noise) and  $\tau_{i6}$  (initial weight scaling factor) was  
 436 0.215, both were however not significantly different from zero. The lack of significant within-  
 437 subject correlations among  $\tau_i$  components suggests that they are essentially independent.



438

439 Figure 8: Marginal posterior distributions of the subject-level hyperparameter  $\tau_i$  for subject  $i =$   
 440 6.

441  
442

**Table 6.** Mean and 94% HWCI of the marginal posterior distributions of  $\tau_6$ .

	$\tau_{61}$	$\tau_{62}$	$\tau_{63}$	$\tau_{64}$	$\tau_{65}$	$\tau_{66}$
Mean	0.0012	1.71	0.63	0.10	0.004	0.07
HWCI	0.0003	0.38	0.05	0.02	0.002	0.03

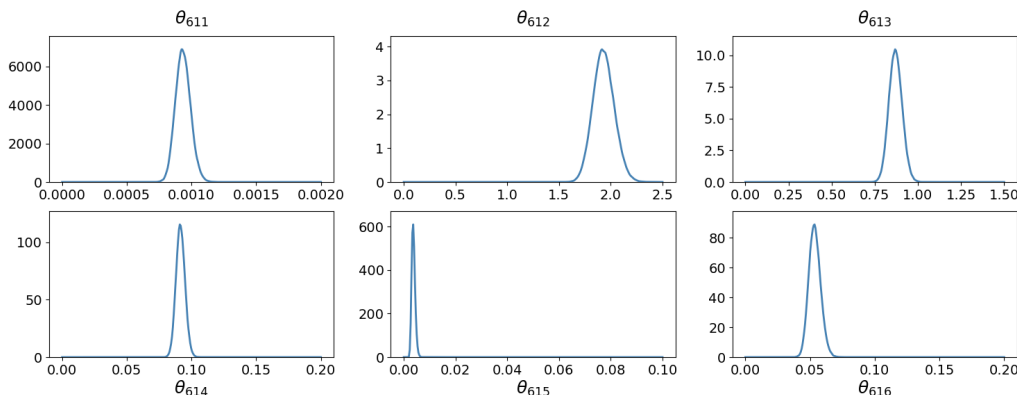
443

**Table 7.** Expected correlation matrix at the subject level.

1	-0.085	0.075	0.093	-0.008	0.052
-0.085	1	0.074	-0.201	-0.029	0.023
0.075	0.074	1	-0.014	0.006	-0.095
0.093	-0.201	-0.014	1	0.004	0.215
-0.008	-0.029	0.006	0.004	1	-0.004
0.052	0.023	-0.095	0.215	-0.004	1

445

446 The marginal posterior distributions of the test-level parameter  $\theta_{i1}$  for subject 6 are depicted in  
 447 Figure 9. The mean and 94% half width credible interval (HWCI) of these distributions are listed  
 448 in Table 8. Compared to  $\eta$  and  $\tau_6$ ,  $\theta_{61}$  components exhibited much lower uncertainties because  
 449 these test-level parameters are constrained by the experimental data directly. The pattern held  
 450 across all subjects. The full table with the mean and HWCI of the test-level marginal distributions  
 451 for all 13 subjects is available in Supplementary Materials B.



452  
453 **Figure 9.** Marginal posterior distributions of the test-level parameter  $\theta_{i1}$  for subject  $i = 6$ .  
454

455 **Table 8.** Mean and 94% HWCI of the marginal posterior distributions of  $\theta_{61}$ .

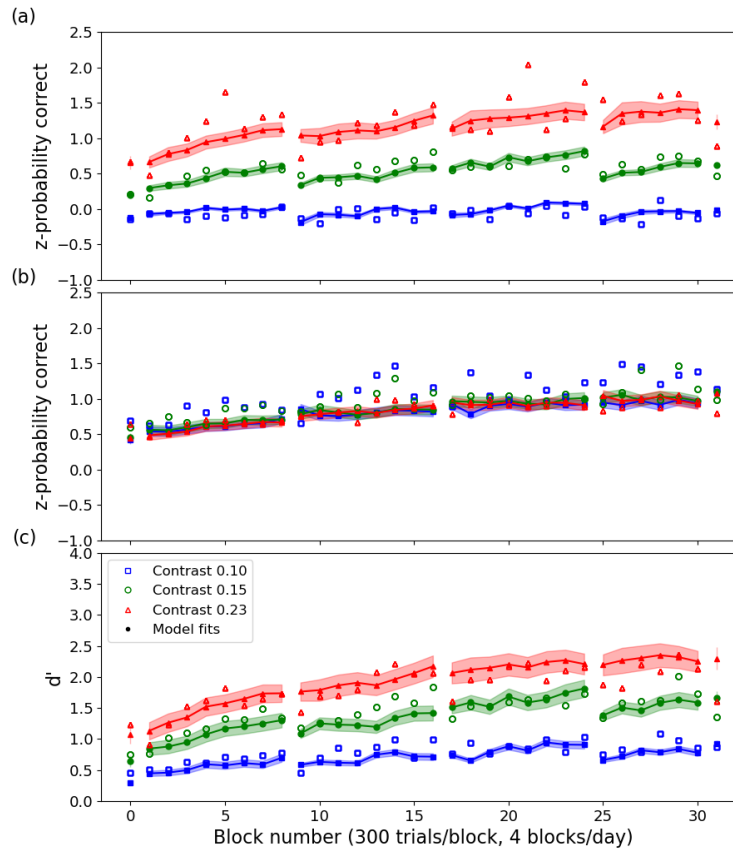
	$\theta_{611}$	$\theta_{612}$	$\theta_{613}$	$\theta_{614}$	$\theta_{615}$	$\theta_{616}$
Mean	0.0009	1.93	0.87	0.09	0.004	0.05
HWCI	0.0001	0.10	0.04	0.00	0.001	0.00

456

457 Predicted learning curves and weight structures (HB-AHRM). Figure 10 depicts the observed  
458 and predicted population-level z-score learning curves in the incongruent and congruent  
459 conditions, as well as the derived  $d'$  learning curves. For the z-score learning curves, the average  
460  $RMSE$  was  $0.173 \pm 0.071$  and  $0.175 \pm 0.090$  and the average 94% HWCI was  $0.064 \pm 0.037$  and  
461  $0.071 \pm 0.001$  in the congruent and incongruent conditions, respectively, with a  $R^2$  of 0.835.

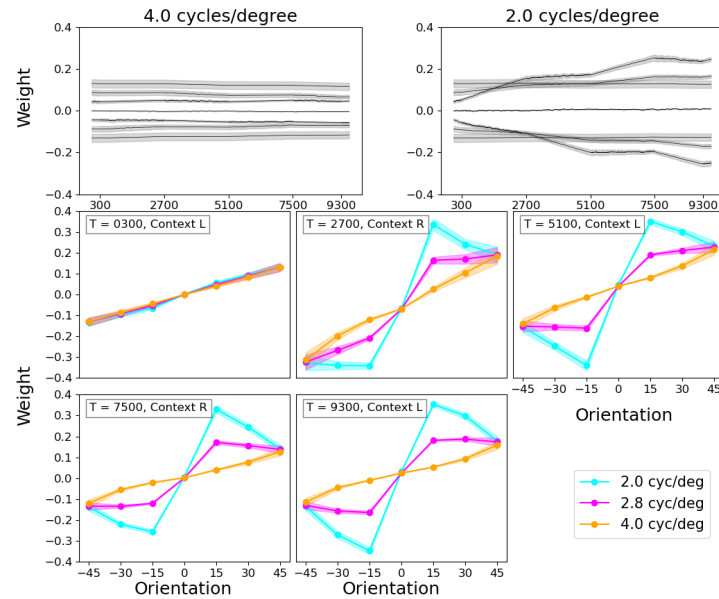
462 For the  $d'$  learning curves, the average  $RMSE$  was  $0.199 \pm 0.074$  and average 94% HWCI was  
463  $0.105 \pm 0.051$ , with a  $R^2$  of 0.887. In comparison, the AHRM with parameters from grid search in  
464 Petrov et al. (2005) accounted for 88.2% of the variance of the average  $d'$  learning curve across  
465 all the subject, with a  $RMSE$  of 0.209. The results suggest that the HB-AHRM provided a largely  
466 comparable solution as the original AHRM at the population level. A table that details  $RMSE$  and  
467 94% HWCI in each of the six experimental conditions is available in Supplementary Materials B.





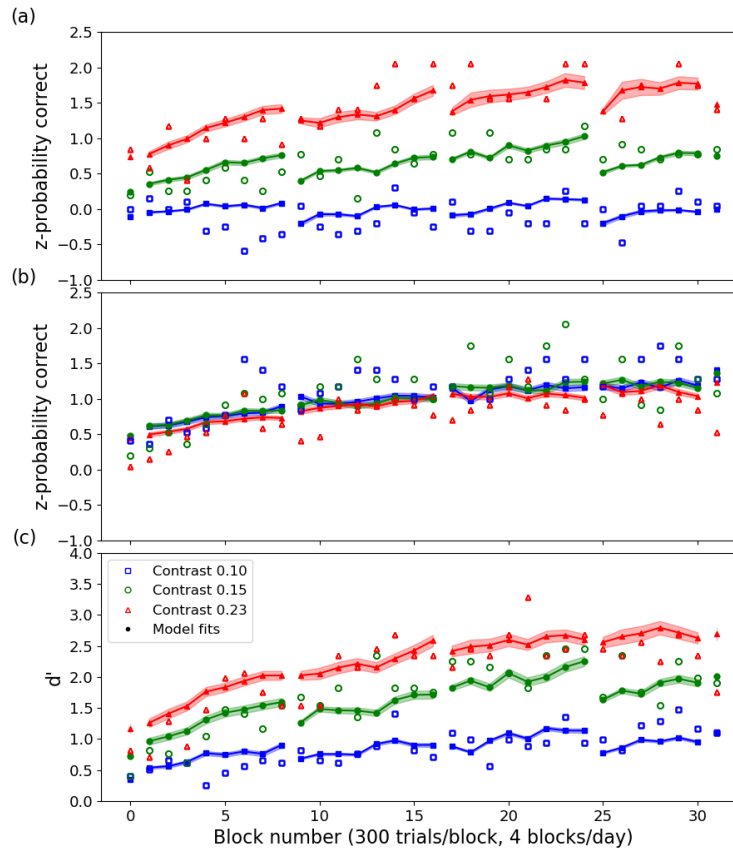
468  
469 Figure 10: Observed and predicted population-level z-score learning curves in the incongruent  
470 (a) and congruent (b) conditions, and the derived  $d'$  learning curves (c). Data points represent the  
471 average learning curves across all 13 subjects.  
472

473 Figure 11 depicts population-level weight structure evolution over the course of training.  
474 Similar to Petrov et al. (2005), the weights for task-relevant channels (e.g., 2 c/d) increased over  
475 the course of training, while the weights for task-irrelevant channels (e.g., 4 c/d) stayed more or  
476 less the same. The HB-AHRM also allowed us to estimate the uncertainties of the weights. For the  
477 channels at 2 c/d, the average HWCI of the weights was 0.008 after 300 trials of training, 0.011  
478 after 2700 trials of training, and 0.011 after 9300 trials of training. For the channels at 4 c/d, the  
479 average HWCI of the weights was 0.008 after 300 trials of training, 0.007 after 2700 trials of  
480 training, and 0.006 after 9300 trials of training.



481  
482 Figure 11: Population-level weight structure evolution over the course of training. Top row:  
483 Longitudinal weight traces for units tuned to the target frequency (2.0 c/d) and an irrelevant  
484 frequency (4.0 c/d). Each trace corresponds to a particular orientation. Middle and bottom rows:  
485 Cross-sections of the weights at 2.0 c/d, 2.8 c/d and 4.0 c/d at the end of each epoch.  
486  
487

488 Figure 12 depicts the observed and predicted test-level z-score learning curves in the  
489 incongruent and congruent conditions, as well as the derived  $d'$  learning curves for subject 6. For  
490 the z-score learning curves, the average  $RMSE$  was  $0.423 \pm 0.204$  and  $0.456 \pm 0.227$  and the average  
491 94% HWCI of  $0.035 \pm 0.001$  and  $0.032 \pm 0.015$  in the congruent and incongruent conditions,  
492 respectively, with a  $R^2$  of 0.675. For the  $d'$  learning curves, the average  $RMSE$  was  $0.276 \pm 0.071$   
493 and the average 94% HWCI was  $0.031 \pm 0.014$ , with a  $R^2$  of 0.744. A table that details  $RMSE$  and  
494 94% HWCI in each of the six experimental conditions for all the subjects is available in  
495 Supplementary Materials B.

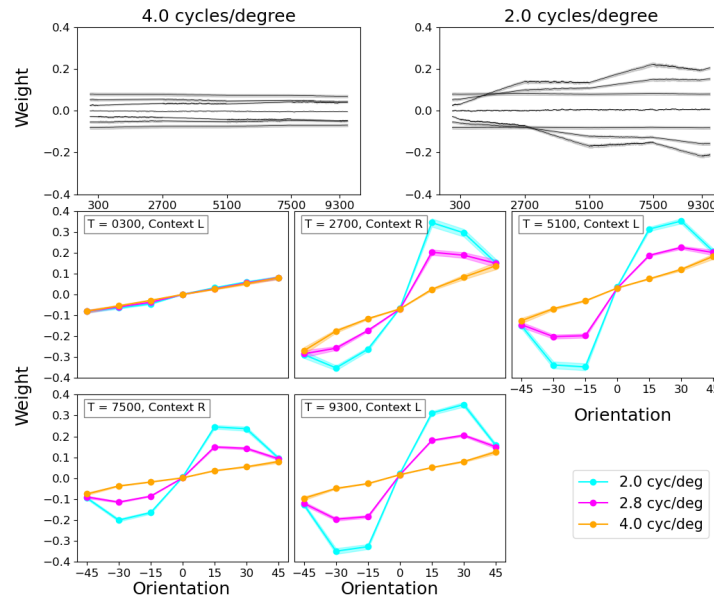


496  
497 Figure 12: Observed and predicted test-level z-score learning curves in the incongruent (a) and  
498 congruent (b) conditions, and the derived  $d'$  learning curves (c) for subject 6.  
499

500 Figure 13 depicts test-level weight structure evolution over the course of training for subject  
501 6. The pattern of results is very similar to what happened at the population level, although the  
502 quantitative weights were different. For the channels at 2 c/d, the average HWCI of the weights  
503 was 0.003 after 300 trials of training, 0.005 after 2700 trials of training, and 0.006 after 9300 trials  
504 of training. For the channels at 4 c/d, the average HWCI of the weights was 0.003 after 300 trials  
505 of training, 0.002 after 2700 trials of training, and 0.001 after 9300 trials of training.

506

507



508  
 509 Figure 13: Test-level weight structure evolution over the course of training for subject 6. Top  
 510 row: Longitudinal weight traces for units tuned to the target frequency (2.0 c/d) and an irrelevant  
 511 frequency (4.0 c/d). Each trace corresponds to a particular orientation. Middle and bottom rows:  
 512 Cross-sections of the weights at 2.0 c/d, 2.8 c/d and 4.0 c/d at the end of each epoch.  
 513

514  
 515

## STUDY 2: SIMULATIONS

516 In study 2, we conducted a simulation study to evaluate parameter recovery and HB-AHRM's  
 517 ability in predicting the performance of a new simulated observer with no or limited training data.

### 518 *Methods*

519 Simulated Datasets. We created simulated dataset1 with 13 simulated observers to evaluated  
 520 parameter recovery. For each simulated observer  $i$ , we set their AHRM parameters by drawing a  
 521 random sample from the six-dimensional test-level HB-AHRM posterior distribution  $\theta_{i1}$  (see  
 522 Supplementary Materials C for a table of all the parameters). We then created a single trial  
 523 sequence with 9600 trials based on Petrov et al. (2005) and simulated these observers'  
 524 performance using the AHRM with the same trial sequence 300 times. Simulated dataset1  
 525 therefore consisted of performance in six experimental conditions, averaged every 300 trials,

526 from each of the 13 simulated observers from the 300 repeated simulations. The structure of the  
527 data was identical to that in Petrov et al. (2005).

528 To evaluate HB-AHRM's ability in predicting the performance of a new simulated observer  
529 with no or limited training data, we created three additional simulated datasets by deleting some  
530 training data for a randomly selected subject 13 in simulated dataset1, while keeping all the data  
531 from the other 12 simulated observers. Specifically, simulated dataset2, simulated dataset3, and  
532 simulated dataset4 comprised 9600 trials of subjects 1-12 and 0 trials, the first 300 trials, and the  
533 first 2700 trials of training data for subject 13, respectively.

534 *HB-AHRM fitting; statistical evaluation.* We fit the HB-AHRM to each of the four simulated  
535 datasets and computed the predicted learning curves from the joint posterior distributions of the  
536 HB-AHRM hyperparameters and parameters. The procedure was identical to that of STUDY 1.

537 For simulated dataset1, we evaluated parameter recovery by comparing the mean of the  
538 posterior distributions of the AHRM parameters with those of the simulated observers ("the truth")  
539 and computed the 94% HWCI for each of the parameters. We also evaluated the goodness of fit  
540 using *RMSE* and  $R^2$ .

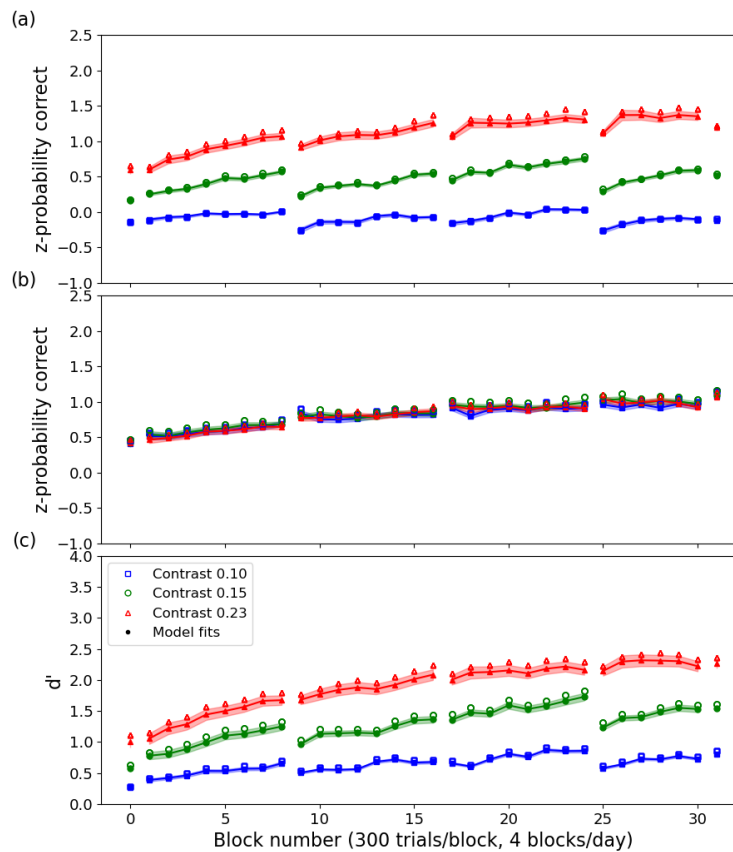
541 For simulated dataset2, dataset3, and dataset4, we focused on the predicted learning curves of  
542 simulated observer 13 and compared them with the simulated learning curves of the same observer  
543 in simulated dataset1 ("the truth").

## 544 ***Results***

545 *Model recovery.* As shown in Figures 14 and 15, the HB-ARHM provided excellent fits to  
546 simulated dataset1 at both the population and test levels.

547 At the population level, for the z-score learning curves, the average RMSE was  $0.014 \pm 0.001$   
548 and  $0.005 \pm 0.002$  and the average 94% HWCI was  $0.037 \pm 0.001$  and  $0.031 \pm 0.015$ , in the congruent

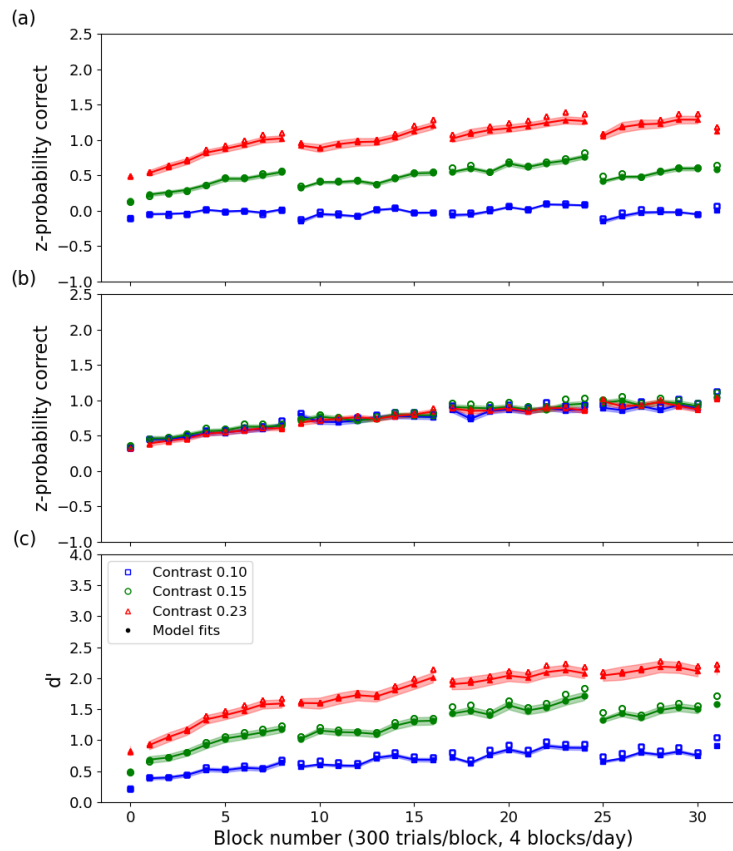
549 and incongruent conditions, respectively, with a  $R^2$  of 0.994. For the  $d'$  learning curves, the  
550 average RMSE was  $0.078 \pm 0.038$  and the average 94% HWCI was  $0.052 \pm 0.031$ , with a  $R^2$  of  
551 0.981. A table that details  $RMSE$  and 94% HWCI in each of the six experimental conditions is  
552 available in Supplementary Materials C.



553  
554 Figure 14: Simulated and predicted population-level z-score learning curves in the incongruent  
555 (a) and congruent (b) conditions, and the derived  $d'$  learning curves (c). Data points represent the  
556 average simulated learning curves across all 13 simulated observers.  
557

558 At the test level, for the z-score learning curves, the average RMSE was  $0.010 \pm 0.000$  and  
559  $0.010 \pm 0.000$  and the average 94% HWCI was  $0.034 \pm 0.001$  and  $0.030 \pm 0.018$  for simulated  
560 observer 13 in the congruent and incongruent conditions, respectively, with a  $R^2$  of 0.994. For the  
561  $d'$  learning curves, the average RMSE was  $0.068 \pm 0.023$  and the average 94% HWCI was

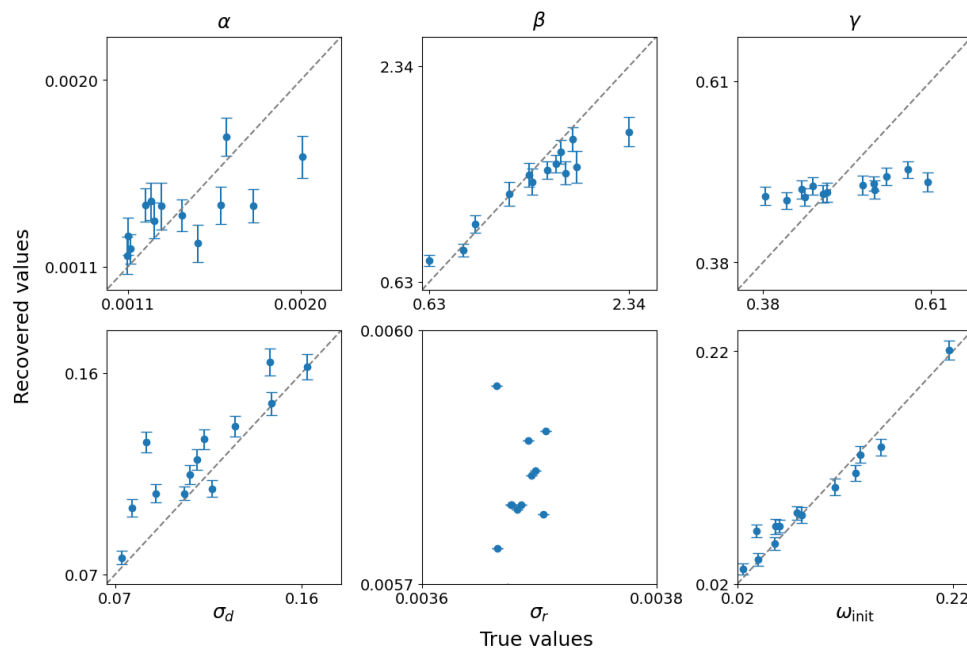
562  $0.048 \pm 0.028$ , with a  $R^2$  of 0.987. A table that details  $RMSE$  and 94% HWCI in each of the six  
563 experimental conditions for all 13 observers is available in Supplementary Materials C.  
564



565  
566 Figure 15. Simulated and predicted test-level z-score learning curves in the incongruent (a) and  
567 congruent (b) conditions, and the derived  $d'$  learning curves (c) for simulated observer 13. Data  
568 points represent the simulated learning curves.  
569

570 Figure 16 shows scatter plots of the recovered AHRM parameters against their true values in  
571 simulated dataset1. The  $RMSE$  between the recovered and true AHRM parameters were 0.00019,  
572 0.308, 0.061, 0.017, 0.002, and 0.011 for the learning rate ( $\alpha$ ), bias strength  
573 ( $\beta$ ), activation function gain ( $\gamma$ ), decision noise ( $\sigma_d$ ), representation noise ( $\sigma_r$ ), and initial weight  
574 scaling factor ( $w_{init}$ ), respectively. The 94% HWCI for the recovered parameters were  $9e-5 \pm 2e-$   
575  $5$ ,  $0.091 \pm 0.040$ ,  $0.012 \pm 0.002$ ,  $0.005 \pm 0.001$ ,  $7e-17 \pm 2e-17$ , and  $0.007 \pm 0.001$ , respectively. For the  
576 learning rate, bias strength, decision noise, and initial weight scaling factor, the recovered

577 parameters exhibited excellent correlations with their true values, with Pearson's correlation  
578 coefficients of 0.690, 0.956, 0.877 and 0.980, respectively. For activation function gain, the true  
579 values ranged from 0.384 to 0.606, but the recovered values all fell within a narrow range between  
580 0.459 and 0.497, suggesting that the model was not very sensitive to activation function gain. For  
581 representation noise, the true values were in a very narrow range (0.00366 to 0.00371), and the  
582 recovered values were also in a very narrow range (0.00567 to 0.00593), although with a slightly  
583 higher mean. This is because representation noise was very small relative to the external noise in  
584 this experiment; it didn't have much impact on model performance. Overall, these results indicate  
585 that the HB-ARHM exhibited very good model recovery.



586 Figure 16. Scatter plots of the recovered AHRM parameters against their true values in simulated  
587 dataset1. Each panel shows one AHRM parameter and each point represent one simulated  
588 observer. Error bars represent 94% HWCI.

590 Model predictions. Figure 17 shows the predicted learning curves of subject 13 with no data, 30  
591 trials of data, 2700 trials of data, and 9600 trials of data. We compared the predictions with the  
592 simulated performance of the subject in simulated dataset1.



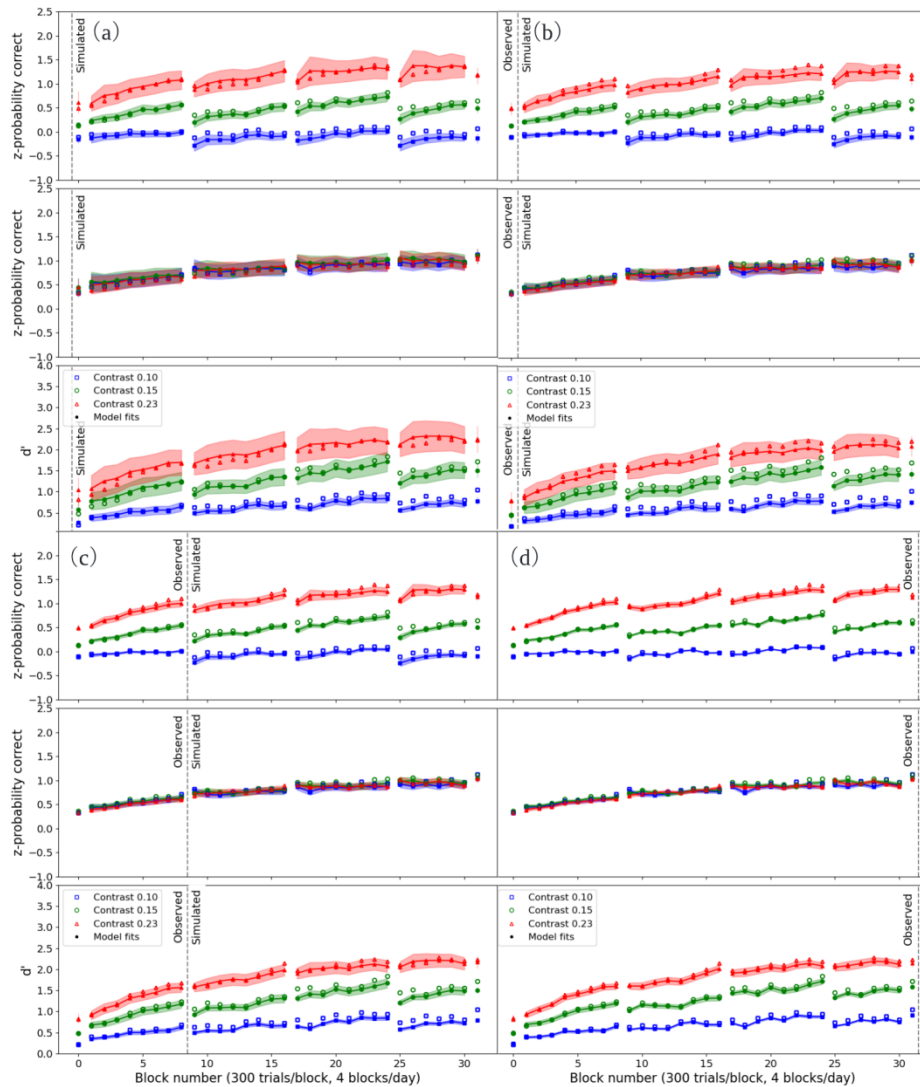
594 With no data, for the z-score learning curves, the average RMSE was  $0.011 \pm 0.002$  and  
595  $0.027 \pm 0.009$  and the average 94% HWCI was  $0.146 \pm 0.003$  and  $0.128 \pm 0.068$  in the congruent and  
596 incongruent conditions, respectively, with a  $R^2$  of 0.972. For the  $d'$  learning curves, the average  
597 RMSE was  $0.104 \pm 0.007$  and the average 94% HWCI was  $0.193 \pm 0.115$ , with a  $R^2$  of 0.960.

598 With 300 trials of data, for the z-score learning curves, the average RMSE was  $0.018 \pm 0.006$   
599 and  $0.025 \pm 0.004$  and the average 94% HWCI was  $0.098 \pm 0.002$  and  $0.088 \pm 0.041$  in the congruent  
600 and incongruent conditions, respectively, with a  $R^2$  of 0.968. For the  $d'$  learning curves, the  
601 average RMSE was  $0.132 \pm 0.016$  and the average 94% HWCI was  $0.146 \pm 0.078$ , with a  $R^2$  of 0.942.

602 With 2700 trials of data, for the z-score learning curves, the average RMSE was  $0.012 \pm 0.004$   
603 and  $0.022 \pm 0.006$  and the average 94% HWCI was  $0.062 \pm 0.002$  and  $0.055 \pm 0.022$  in the congruent  
604 and incongruent conditions, respectively, with a  $R^2$  of 0.980. For the  $d'$  learning curves, the  
605 average RMSE was  $0.101 \pm 0.017$  and the average 94% HWCI was  $0.084 \pm 0.044$ , with a  $R^2$  of 0.966.

606 Overall, the HB-AHRM made excellent predictions with no data, 300 trials of data, and 2700  
607 trials of data for simulated observer 13. A table that details RMSE and 94% HWCI in each of the  
608 six experimental conditions is available in Supplementary Materials C.

609



610  
 611 Figure 17. Simulated and predicted z-score and  $d'$  learning curves for subject 13 with no data (a),  
 612 300 trials of data (b), 2700 trials of data (c), and 9600 trials of data (d). Data points represent the  
 613 simulated learning curves for the subject in simulated dataset1.  
 614  
 615

## DISCUSSION

616 In this study, we developed the HB-AHRM and new modeling technologies to address the  
 617 challenge in fitting the AHRM, a very successful model in visual perceptual learning. A  
 618 combination of feature engineering and linear regression provided a high-quality approximation  
 619 of the likelihood function. This approach allowed us to drastically reduce the computation time for  
 620 fitting the HB-AHRM, estimated to be over 125 days for the dataset in Petrov et al. (2005), which  
 621 is practically infeasible. The HB-AHRM produced significantly better fits than the BIP, enabling

622 fitting at both the group level with comparable goodness of fit as Petrov et al. (2005), and at the  
623 individual level. In stimulation studies, the HB-AHRM along with the new modeling technologies  
624 demonstrated robust model recovery properties, accurately predicting simulated observer  
625 outcomes with minimal or no initial data.

626 The AHRM generates trial-by-trial responses based on its parameters and the stimulus  
627 sequence. As the exact external noise image for each trial in Petrov et al. (2005) was unavailable,  
628 we adopted a procedure similar to the original study, sampling from a cache of 1200 expected  
629 activations. The impact of the mismatch with the exact stimulus sequence used in the experiment  
630 appeared to be minimal at the group level due to cross-subject averaging, but it may have  
631 influenced the fits at the individual subject level because each subject in the original study used a  
632 unique random trial sequence. Therefore, it is crucial to accurately record the exact stimulus  
633 sequences in future studies.

634 Many traditional statistical methods assume homogeneity or complete independence across  
635 subjects and tests. In contrast, hierarchical modeling (HB) integrates heterogeneous information  
636 across multiple levels, using Bayes' theorem to combine sub-models and probability distributions  
637 from all observed data in a study (Kruschke, 2014; Kruschke & Liddell, 2018; Rouder & Lu,  
638 2005). This yields updated joint posterior distributions of hyperparameters and parameters,  
639 enhancing accuracy compared to methods that treat each individual independently (H. Gu et al.,  
640 2016; Kruschke, 2014). Previous studies, including this one, have shown that HBM provides more  
641 precise estimates of parameters of interest compared to traditional methods like the BIP, in  
642 estimating contrast sensitivity functions (Zhao, Lesmes, Hou, & Lu, 2021), visual acuity  
643 behavioral functions (Zhao, Lesmes, Dorr, & Lu, 2021), and learning curves in perceptual learning  
644 (Zhao, Liu, Doshier, & Lu, 2024a, 2024b).

645        Moreover, HBM offers a robust framework for predictions, treating to-be-predicted  
646 performance as missing data to compute their posterior distributions based on available  
647 information. Leveraging conditional dependencies across the hierarchy and between- and within-  
648 subject covariances, HBM facilitates constructing digital twins and making highly accurate and  
649 reasonably precise predictions (Zhao, Lesmes, Dorr, & Lu, 2024).

650        This study introduces the concept of likelihood function approximation and demonstrates its  
651 application in fitting the HB-AHRM. This approach may have broader utility in fitting other  
652 stochastic models lacking analytic forms, such as the perceptual template model (Lu & Doshier,  
653 2008), integrated reweighting theory (Doshier et al., 2013), and various response time models  
654 (Ratcliff, Smith, Brown, & McKoon, 2016), as well as complex models that requires extensive  
655 computations to generate predictions, such as the population receptive field model in retinotopic  
656 map studies (Dumoulin & Wandell, 2008). In this particular application, feature engineering and  
657 linear regression were employed to generate the approximate likelihood function. Alternatively,  
658 other methods, including non-linear regression and machine learning, could be utilized to derive  
659 the approximate likelihood function.

660        In conclusion, we successfully developed and implemented the HB-AHRM using newly  
661 developed modeling technologies. These advancements hold promise for widespread applications  
662 in fitting stochastic models.

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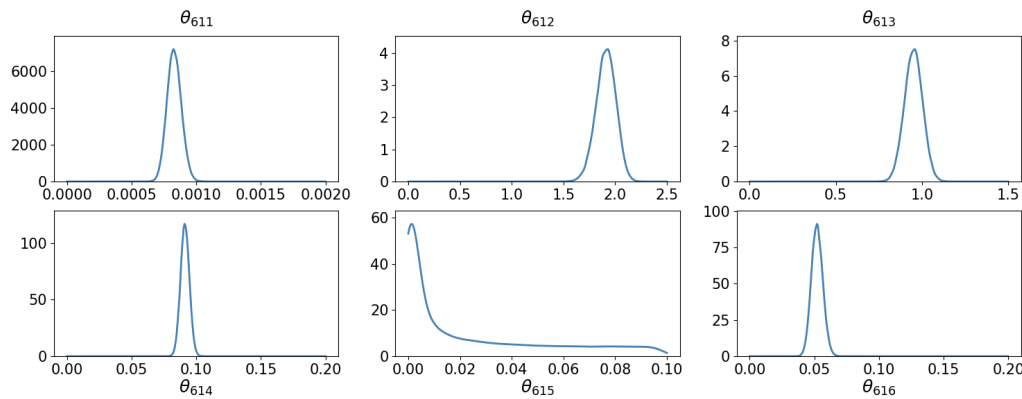
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### Supplementary Materials A. BIP Results.

The marginal posterior distributions of parameter  $\theta_{i1}$  for subject 6 from the BIP are depicted in Figure S1. The mean and 94% half width credible interval (HWCI) of these distributions are listed in Table S1.

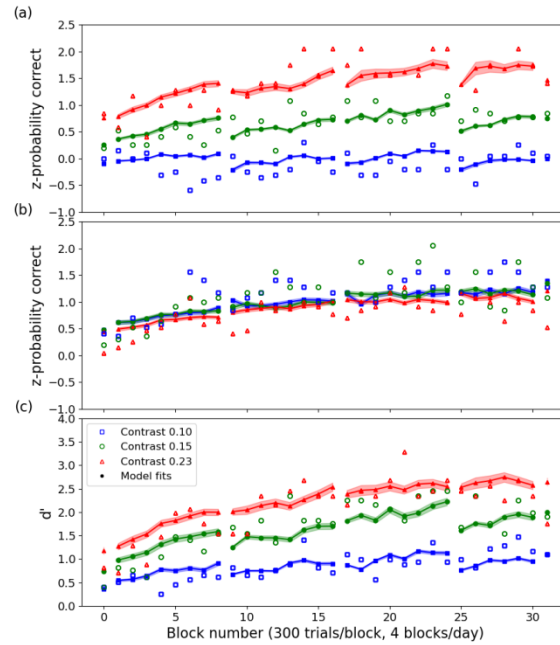


**Figure S1.** Marginal posterior distributions of parameter  $\theta_{i1}$  for subject  $i = 6$  from the BIP.

**Table S1.** Mean and 94% HWCI of the marginal posterior distributions of  $\theta_{61}$ .

	$\theta_{611}$	$\theta_{612}$	$\theta_{613}$	$\theta_{614}$	$\theta_{615}$	$\theta_{616}$
<i>Mean</i>	<i>0.0008</i>	<i>1.91</i>	<i>0.95</i>	<i>0.09</i>	<i>0.054</i>	<i>0.05</i>
<i>HWCI</i>	<i>0.0001</i>	<i>0.10</i>	<i>0.05</i>	<i>0.00</i>	<i>0.496</i>	<i>0.00</i>

Figure S2 depicts the observed and predicted test-level z-score learning curves in the incongruent and congruent conditions, as well as the derived  $d'$  learning curves from the BIP for subject 6. For the z-score learning curves, the average  $RMSE$  was  $0.403 \pm 0.092$  and  $0.389 \pm 0.158$  and the average 94% HWCI was  $0.039 \pm 0.001$  and  $0.037 \pm 0.016$  in the congruent and incongruent conditions, respectively, with a  $R^2$  of  $0.535 \pm 0.090$  across all 13 subjects. For the  $d'$  learning curves, the average  $RMSE$  was  $0.366 \pm 0.057$  and the average 94% HWCI was  $0.052 \pm 0.021$ , with a  $R^2$  of  $0.604 \pm 0.135$  across all 13 subjects. Tables S2 and S3 detail  $RMSE$  and 94% HWCI in each of the six experimental conditions.



**Figure S2:** Observed and predicted test-level z-score learning curves in the incongruent (a) and congruent (b) conditions, and the derived  $d'$  learning curves (c) for subject 6.

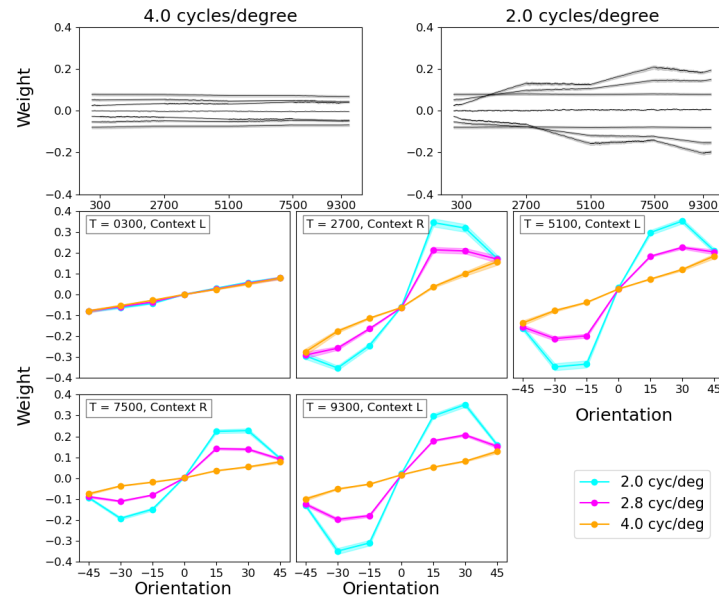
**Table S2.** RMSE, 94% HWCI and  $R^2$  for the z-score learning curves across 13 subjects (BIP).

	RMSE	HWCI	$R^2$
Congruent, low contrast	0.526 (0.267)	0.039 (0.010)	0.535 (0.090)
Incongruent, low contrast	0.261 (0.040)	0.021 (0.007)	
Congruent, medium contrast	0.379 (0.153)	0.040 (0.009)	
Incongruent, medium contrast	0.296 (0.064)	0.030 (0.005)	
Congruent, high contrast	0.304 (0.039)	0.038 (0.007)	
Incongruent, high contrast	0.613 (0.386)	0.059 (0.014)	

**Table S3.** RMSE, 94% HWCI and  $R^2$  for the  $d'$  learning curves across 13 subjects (BIP).

	RMSE	CI of $d'$	$R^2$
Low contrast	0.289	0.025	0.604
Medium contrast	0.383	0.054	(0.135)
High contrast	0.425	0.077	

Figure S3 depicts weight structure evolution over the course of training from the BIP for subject 6. For the channels at 2 c/d, the average HWCI of the weights was 0.003 after 300 trials of training, 0.004 after 2700 trials of training, and 0.006 after 9300 trials of training. For the channels at 4 c/d, the average HWCI of the weights was 0.003 after 300 trials of training, 0.002 after 2700 trials of training, and 0.002 after 9300 trials of training.



**Figure S3.** Weight change over time for subject 6 (BIP). Top row: Longitudinal weight traces for units tuned to the target frequency (2.0 c/d) and an irrelevant frequency (4.0 c/d). Each trace corresponds to a particular orientation. Middle and bottom rows: Cross-sections of the weights at 2.0 c/d, 2.8 c/d and 4.0 c/d at the end of each epoch.

**Supplementary Materials B. Additional HB-AHRM Results.**

**Table S4.** Mean and 94% HWCI of the marginal  $\tau_{ik}$  distributions (HB-AHRM).

$i$	$\tau_{i1}$	$\tau_{i2}$	$\tau_{i3}$	$\tau_{i4}$	$\tau_{i5}$	$\tau_{i6}$
<b>1</b>	0.0011 (0.0003)	1.63 (0.36)	0.45 (0.06)	0.03 (0.01)	0.004 (0.000)	0.02 (0.01)
<b>2</b>	0.0012 (0.0003)	1.88 (0.42)	0.47 (0.07)	0.10 (0.03)	0.004 (0.000)	0.10 (0.03)
<b>3</b>	0.0017 (0.0015)	1.87 (0.42)	0.53 (0.08)	0.08 (0.02)	0.004 (0.000)	0.09 (0.03)
<b>4</b>	0.0022 (0.0006)	1.07 (0.24)	0.40 (0.06)	0.11 (0.03)	0.004 (0.000)	0.13 (0.05)
<b>5</b>	0.0010 (0.0003)	1.98 (0.45)	0.35 (0.05)	0.10 (0.03)	0.004 (0.000)	0.15 (0.05)
<b>6</b>	0.0012 (0.0003)	1.71 (0.38)	0.63 (0.05)	0.10 (0.02)	0.004 (0.000)	0.07 (0.03)
<b>7</b>	0.0019 (0.0005)	0.94 (0.31)	0.48 (0.09)	0.17 (0.04)	0.004 (0.000)	0.16 (0.06)
<b>8</b>	0.0022 (0.0006)	1.92 (0.43)	0.42 (0.07)	0.09 (0.02)	0.004 (0.000)	0.11 (0.04)
<b>9</b>	0.0018 (0.0005)	0.79 (0.17)	0.40 (0.06)	0.16 (0.04)	0.004 (0.000)	0.06 (0.02)
<b>10</b>	0.0007 (0.0002)	1.54 (0.34)	0.38 (0.05)	0.11 (0.03)	0.004 (0.000)	0.10 (0.03)
<b>11</b>	0.0022 (0.0016)	1.88 (0.32)	0.47 (0.06)	0.09 (0.02)	0.004 (0.000)	0.14 (0.05)
<b>12</b>	0.0014 (0.0004)	1.24 (0.28)	0.44 (0.06)	0.12 (0.03)	0.004 (0.000)	0.09 (0.03)
<b>13</b>	0.0023 (0.0006)	1.67 (0.37)	0.52 (0.07)	0.14 (0.04)	0.004 (0.006)	0.05 (0.02)

**Table S5.** Mean and 94% HWCI of the marginal  $\theta_{ij}$  distributions (HB-AHRM).

$i$	$\theta_{ij}(1)$	$\theta_{ij}(2)$	$\theta_{ij}(3)$	$\theta_{ij}(4)$	$\theta_{ij}(5)$	$\theta_{ij}(6)$
<b>1</b>	0.0008 (0.0001)	1.74 (0.05)	0.46 (0.01)	0.01 (0.00)	0.004 (0.001)	0.01 (0.00)
<b>2</b>	0.0010 (0.0001)	2.30 (0.13)	0.49 (0.01)	0.11 (0.00)	0.004 (0.001)	0.11 (0.01)
<b>3</b>	0.0018 (0.0001)	2.25 (0.07)	0.62 (0.01)	0.07 (0.00)	0.004 (0.001)	0.10 (0.01)
<b>4</b>	0.0031 (0.0003)	0.77 (0.03)	0.36 (0.01)	0.11 (0.00)	0.004 (0.001)	0.18 (0.02)
<b>5</b>	0.0006 (0.0001)	2.61 (1.14)	0.27 (0.01)	0.09 (0.00)	0.004 (0.091)	0.24 (0.01)
<b>6</b>	0.0009 (0.0001)	1.93 (0.10)	0.87 (0.04)	0.09 (0.00)	0.004 (0.001)	0.05 (0.00)
<b>7</b>	0.0023 (0.0002)	0.59 (0.03)	0.51 (0.01)	0.28 (0.01)	0.004 (0.001)	0.29 (0.02)
<b>8</b>	0.0030 (0.0003)	2.45 (0.11)	0.37 (0.01)	0.09 (0.00)	0.004 (0.001)	0.13 (0.02)
<b>9</b>	0.0021 (0.0002)	0.41 (0.04)	0.35 (0.01)	0.23 (0.01)	0.004 (0.001)	0.04 (0.01)
<b>10</b>	0.0003 (0.0001)	1.61 (1.10)	0.32 (0.01)	0.11 (0.00)	0.004 (0.001)	0.10 (0.01)
<b>11</b>	0.0031 (0.0002)	2.36 (0.06)	0.48 (0.01)	0.07 (0.00)	0.004 (0.001)	0.20 (0.02)
<b>12</b>	0.0012 (0.0001)	1.02 (0.06)	0.43 (0.01)	0.14 (0.01)	0.004 (0.001)	0.10 (0.01)
<b>13</b>	0.0035 (0.0002)	1.84 (0.13)	0.59 (0.02)	0.19 (0.01)	0.004 (0.001)	0.04 (0.01)

**Table S6.** RMSE, HWCI and  $R^2$  for the z-score learning curves at the population and test levels.

		RMSE	HWCI	$R^2$
Population	Congruent, low contrast	0.303	0.069	0.835
	Incongruent, low contrast	0.088	0.029	
	Congruent, medium contrast	0.165	0.072	
	Incongruent, medium contrast	0.119	0.047	
	Congruent, high contrast	0.103	0.071	
	Incongruent, high contrast	0.262	0.115	
Test	Congruent, low contrast	0.532	0.034	0.533 (0.093)
	Incongruent, low contrast	0.268	0.018	
	Congruent, medium contrast	0.377	0.036	
	Incongruent, medium contrast	0.307	0.026	
	Congruent, high contrast	0.300	0.035	

**Table S7.** RMSE, HWCI and  $R^2$  for the  $d'$  learning curves at the population and test level.

		RMSE	HWCI	$R^2$
Population	Low contrast	0.135	0.043	0.887
	Medium contrast	0.157	0.103	
	High contrast	0.305	0.168	
Test	Low contrast	0.287	0.023	0.615 (0.115)
	Medium contrast	0.303	0.050	
	High contrast	0.418	0.075	

**Supplementary Materials C. Additional Results in the Simulation Study.**

**Table S8.** Parameters of the Simulated Observers

$i$	$\theta_{i1}(1)$	$\theta_{i1}(2)$	$\theta_{i1}(3)$	$\theta_{i1}(4)$	$\theta_{i1}(5)$	$\theta_{i1}(6)$
1	0.0011	1.48	0.41	0.11	0.004	0.15
2	0.0014	1.31	0.47	0.11	0.004	0.02
3	0.0016	1.89	0.52	0.15	0.004	0.08
4	0.0015	1.02	0.38	0.14	0.004	0.13
5	0.0012	0.92	0.53	0.16	0.004	0.08
6	0.0011	1.86	0.46	0.09	0.004	0.13
7	0.0020	1.79	0.55	0.12	0.004	0.11
8	0.0012	0.63	0.45	0.09	0.004	0.04
9	0.0016	1.71	0.58	0.07	0.004	0.06
10	0.0011	1.75	0.44	0.08	0.004	0.05
11	0.0013	1.64	0.53	0.10	0.004	0.22
12	0.0017	1.51	0.43	0.13	0.004	0.04
13	0.0013	2.34	0.61	0.11	0.004	0.06

**Table S9.** Mean and 94% HWCI of the  $\eta$  distributions.

	$\eta(1)$	$\eta(2)$	$\eta(3)$	$\eta(4)$	$\eta(5)$	$\eta(6)$
Mean	0.0014	1.37	0.48	0.12	0.006	0.08
HWCI	0.0003	0.08	0.01	0.01	0.000	0.01

**Table S10.** Mean and 94% HWCI of the  $\tau_{ik}$  distributions (Simulation study).

$i$	$\tau_{i1}$	$\tau_{i2}$	$\tau_{i3}$	$\tau_{i4}$	$\tau_{i5}$	$\tau_{i6}$
<b>1</b>	0.0012 (0.0001)	1.44 (0.21)	0.47 (0.01)	0.13 (0.02)	0.006 (0.000)	0.11 (0.03)
<b>2</b>	0.0013 (0.0001)	1.37 (0.21)	0.47 (0.01)	0.12 (0.01)	0.006 (0.000)	0.05 (0.02)
<b>3</b>	0.0014 (0.0001)	1.46 (0.21)	0.48 (0.01)	0.13 (0.02)	0.006 (0.000)	0.08 (0.03)
<b>4</b>	0.0014 (0.0001)	1.24 (0.19)	0.47 (0.01)	0.14 (0.02)	0.006 (0.000)	0.10 (0.03)
<b>5</b>	0.0013 (0.0001)	1.13 (0.16)	0.47 (0.01)	0.14 (0.02)	0.006 (0.000)	0.09 (0.02)
<b>6</b>	0.0013 (0.0001)	1.55 (0.23)	0.48 (0.01)	0.11 (0.01)	0.006 (0.000)	0.11 (0.03)
<b>7</b>	0.0015 (0.0001)	1.44 (1.22)	0.49 (0.01)	0.12 (0.01)	0.006 (0.000)	0.10 (0.03)
<b>8</b>	0.0014 (0.0001)	1.06 (0.15)	0.47 (0.01)	0.13 (0.02)	0.006 (0.000)	0.08 (0.02)
<b>9</b>	0.0015 (0.0001)	1.48 (0.22)	0.48 (0.01)	0.10 (0.01)	0.006 (0.000)	0.08 (0.02)
<b>10</b>	0.0013 (0.0001)	1.51 (0.22)	0.47 (0.01)	0.11 (0.01)	0.006 (0.000)	0.07 (0.02)
<b>11</b>	0.0014 (0.0001)	1.46 (0.22)	0.47 (0.01)	0.11 (0.01)	0.006 (0.000)	0.14 (0.04)
<b>12</b>	0.0014 (0.0001)	1.40 (0.22)	0.48 (0.01)	0.13 (0.02)	0.006 (0.000)	0.06 (0.02)
<b>13</b>	0.0014 (0.0001)	1.61 (0.25)	0.47 (0.01)	0.12 (0.01)	0.006 (0.000)	0.08 (0.02)



**Table S11.** Mean and 94% HWCI of the  $\theta_{ij}$  distributions (Simulation study).

<i>i</i>	$\theta_{ij}(1)$	$\theta_{ij}(2)$	$\theta_{ij}(3)$	$\theta_{ij}(4)$	$\theta_{ij}(5)$	$\theta_{ij}(6)$
<b>1</b>	0.0012 (0.0001)	1.48 (0.09)	0.46 (0.01)	0.13 (0.00)	0.006 (0.000)	0.14 (0.03)
<b>2</b>	0.0013 (0.0001)	1.33 (0.09)	0.47 (0.01)	0.12 (0.00)	0.006 (0.000)	0.03 (0.03)
<b>3</b>	0.0014 (0.0001)	1.53 (0.12)	0.48 (0.01)	0.15 (0.00)	0.006 (0.000)	0.08 (0.01)
<b>4</b>	0.0014 (0.0001)	1.09 (0.07)	0.46 (0.01)	0.17 (0.00)	0.006 (0.000)	0.11 (0.01)
<b>5</b>	0.0013 (0.0001)	0.88 (0.05)	0.47 (0.01)	0.16 (0.01)	0.006 (0.000)	0.08 (0.01)
<b>6</b>	0.0013 (0.0001)	1.77 (0.10)	0.47 (0.01)	0.11 (0.01)	0.006 (0.000)	0.13 (0.01)
<b>7</b>	0.0016 (0.0001)	1.49 (0.09)	0.49 (0.01)	0.11 (0.01)	0.006 (0.000)	0.10 (0.01)
<b>8</b>	0.0014 (0.0001)	0.80 (0.04)	1.48 (0.01)	0.14 (0.00)	0.006 (0.000)	0.07 (0.01)
<b>9</b>	0.0017 (0.0001)	1.57 (0.07)	1.50 (0.01)	0.08 (0.00)	0.006 (0.000)	0.07 (0.01)
<b>10</b>	0.0012 (0.0001)	1/66 (0.09)	0.46 (0.01)	0.10 (0.00)	0.006 (0.000)	0.05 (0.01)
<b>11</b>	0.0014 (0.0001)	1.51 (0.07)	0.48 (0.01)	0.11 (0.00)	0.006 (0.000)	0.22 (0.01)
<b>12</b>	0.0014 (0.0001)	1/43 (0.11)	0.47 (0.01)	0.14 (0.00)	0.006 (0.000)	0.04 (0.01)
<b>13</b>	0.0014 (0.0001)	1.82 (0.12)	0.48 (0.01)	0.11 (0.00)	0.006 (0.000)	0.07 (0.01)

**Table S12.** RMSE, HWCI and  $R^2$  for the z-score learning curves at the population and test levels.

		RMSE	HWCI	$R^2$
Population	Congruent, low contrast	0.014	0.036	0.994
	Incongruent, low contrast	0.003	0.019	
	Congruent, medium contrast	0.015	0.038	
	Incongruent, medium contrast	0.006	0.022	
	Congruent, high contrast	0.014	0.036	
	Incongruent, high contrast	0.007	0.051	
Test	Congruent, low contrast	0.010	0.033	0.994 (0.003)
	Incongruent, low contrast	0.007	0.017	
	Congruent, medium contrast	0.011	0.035	
	Incongruent, medium contrast	0.012	0.022	
	Congruent, high contrast	0.010	0.034	
	Incongruent, high contrast	0.012	0.052	

**Table S13.** RMSE, HWCI and  $R^2$  for the  $d'$  learning curves at the population and test levels.

		RMSE	HWCI	$R^2$
Population	Low contrast	0.037	0.021	0.981
	Normal contrast	0.077	0.050	
	High contrast	0.120	0.084	
Test	Low contrast	0.041	0.020	0.986 (0.003)
	Normal contrast	0.073	0.046	
	High contrast	0.090	0.077	

**Table S14.** RMSE, HWCI and  $R^2$  for the z-score learning curves in the prediction tasks.

		RMSE	HWCI	$R^2$
No data	Congruent, low contrast	0.010	0.142	0.972
	Incongruent, low contrast	0.037	0.076	
	Congruent, medium contrast	0.010	0.150	
	Incongruent, medium contrast	0.026	0.089	
	Congruent, high contrast	0.014	0.146	
	Incongruent, high contrast	0.017	0.218	
300 trials	Congruent, low contrast	0.022	0.095	0.968
	Incongruent, low contrast	0.028	0.051	
	Congruent, medium contrast	0.020	0.100	
	Incongruent, medium contrast	0.028	0.072	
	Congruent, high contrast	0.012	0.100	
	Incongruent, high contrast	0.019	0.142	
2700 trials	Congruent, low contrast	0.016	0.060	0.980
	Incongruent, low contrast	0.028	0.042	
	Congruent, medium contrast	0.013	0.063	
	Incongruent, medium contrast	0.023	0.042	
	Congruent, high contrast	0.007	0.064	
	Incongruent, high contrast	0.014	0.083	
9600 trials	Congruent, low contrast	0.014	0.033	0.994
	Incongruent, low contrast	0.009	0.012	
	Congruent, medium contrast	0.012	0.034	
	Incongruent, medium contrast	0.010	0.025	
	Congruent, high contrast	0.006	0.034	
	Incongruent, high contrast	0.012	0.051	

**Table S15.** RMSE, 94% HWCI and  $R^2$  for the  $d'$  learning curves in the prediction tasks.

		RMSE	HWCI	$R^2$
No data	Low contrast	0.112	0.077	0.960
	Medium contrast	0.096	0.184	
	High contrast	0.105	0.317	
300 trials	Low contrast	0.132	0.065	0.942
	Medium contrast	0.150	0.146	
	High contrast	0.115	0.226	
2700 trials	Low contrast	0.111	0.038	0.966
	Medium contrast	0.114	0.083	
	High contrast	0.078	0.133	
9600 trials	Low contrast	0.059	0.025	0.987
	Medium contrast	0.065	0.055	
	High contrast	0.067	0.081	