1   Temporal evolution of color representations measured with MEG reveals a 'coarse to fine'

2   dynamic

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Abstract

Color perception is based on the differential spectral responses of the L, M and S-cones, subsequent subcortical and cortical computations, and may include the influence of higher order factors such as language. Although the early subcortical stages of color vision are well characterised, the organization of cortical representations of color remain elusive, despite numerous models based on discrimination thresholds, appearance and categorization. An underexplored aspect of cortical color representations is their dynamic evolution. Here we compare the evolution of three different color representations over time using magnetoencephalography (MEG). We measured neural responses to 14 hues at each of 3 achromatic offsets (increment, isoluminant and decrement) while participants attended either to the exact color of the stimulus or its color category. We used a series of classification analyses, combined with multidimensional scaling (MDS) and Representational Similarity Analysis, to ask how cortical representations of color unfold over time from stimulus onset. We compared the performance of 'higher order' models based on hue and color category with a model based simply on stimulus cone contrast and found that all models had significant correlations with the data. However, the unique variance accounted for by each model revealed a dynamic change in hue responses over time, that was consistent with a 'coarse to fine' transition from a broad clustering into categorical groups to a finer within category representation. Notably, these dynamics were replicated across datasets from both tasks, suggesting they reflect a robust reorganization of cortical hue responses over time.

Introduction

The first stage of human color vision is based on the combination of the differential responses of the L, M and S-cones to the visual stimulus. In this first postreceptoral stage, comparisons of the cone responses are initially made within two distinct cone opponent streams, an L/M cone opponent response (loosely called 'red-green') and one based on the opponent combination of S-cones with the other two (loosely called 'blue-yellow'). Based on physiological and behavioural data, these two pathways are thought to remain segregated within the subcortical processing stages, using different anatomical cell types and pathways in the retina, LGN, and with distinct V1 cortical inputs (Derrington et al., 1984; Lennie & Movshon, 2005). There is also evidence for a behavioural separation at detection threshold (Eskew, 2009; Mullen & Sankeralli, 1999). These two low-level pathways, however, do not provide adequate accounts of color appearance or the perceptual organization of the different colors. For example, colors that are equally spaced in a representation based on perceptual differences/similarity (e.g. using a CIE L*a*b* space), vary in a non-uniform way in a space based on the relative cone responses (e.g. a cone contrast space). Furthermore, it has been widely observed that stimulating the cardinal directions of a cone contrast space, which uniquely activate one type of cone opponent process at a time, do not generate the color appearance of a unique hue. Hence, the unique hues are not aligned with the cone opponent axes and further computations are required for cone opponency to generate the full range of color appearances (Conway et al., 2023; de Valois & de Valois, 1993; Li et al., 2022; Stockman & Brainard, 2010). Embedded within this perceptual color space is a categorical organization, in which colors can be placed into groups according to their color appearance, such as the 11 basic color names of Berlin and Kay (1969). Whether there are any categorical effects in color perception, including potential interaction with language, remains controversial, but evidence is clearer for 'categorical facilitation' effects, likely mediated by a shift of attention to the linguistic distinction between color categories (for a review, see Witzel, 2019). So far, the computational nature of the processes underlying categorical color grouping and subsequent color naming also remains unknown (Siuda-Krzywicka et al., 2019).

Imaging approaches provide an important route into the understanding of how colors are represented in the cortex. fMRI can be used to decode stimulus color using a multivariate approach to analyse spatially distributed patterns of voxel activity in cortex (Brouwer & Heeger, 2009, 2013; Goddard et al., 2010; Goddard & Mullen, 2020; Parkes et al., 2009). Although colors

can be decoded in early visual cortical areas (V1, V2, V3, hV4 & VO), the ventral areas hV4 and VO showed a stronger perceptual relationship than earlier areas such as V1 and V2, with perceptually similar colors evoking the most similar responses (Brouwer & Heeger, 2009). Brouwer & Heeger (2013) also found that areas hV4 and VO1 showed evidence of categorical clustering, especially when participants were engaged in a color categorization task, compared to a diverted attention condition. There is also evidence of further areas of high color responsiveness more anterior in the ventral visual cortex (Komatsu et al., 1992; Stoughton & Conway, 2008; Zeki & Bartels, 1999), including adjacent to areas associated with higher-order responses such as to objects, faces and places (Lafer-Sousa et al., 2016; Lafer-Sousa & Conway, 2013).

MEG signals can be used to decode stimulus color with the advantage of allowing the precise timing of the stimulus response to be determined (Hermann et al., 2022; Rosenthal et al., 2021). This previous work, using 4 stimulus colors and two luminance levels, has shown that MEG signals can be used to decode stimulus color for over 300ms after stimulus onset (Rosenthal et al., 2021), and provides a proof of concept that similarity relationships can be used to test the geometry of stimulus color representations. Other MEG studies have also demonstrated color decoding from MEG signals (Goddard, Carlson, et al., 2022; Goddard, Shooner, et al., 2022; Sandhaeger et al., 2019; Teichmann et al., 2019, 2020), and task-based changes in color coding consistent with sharpening of color tuning with attention (Bartsch et al., 2017; Goddard, Carlson, et al., 2022). In addition, there are many examples of decoding stimulus color from EEG signals (Bae & Chen, 2024; Chauhan et al., 2023; Grootswagers et al., 2024; Hajonides et al., 2021; Retter et al., 2023; Sutterer et al., 2021; Wu et al., 2022).

Here we recorded MEG signals while participants viewed a set of 42 stimuli that varied in hue as well as achromatic offset and compared the timing of the emergence of neural representations of chromatic and achromatic information. We included two task conditions, both involving attention to the stimulus color but only one requiring color naming (categorization), to test whether the task of color naming influences the way color is represented in the cortex.

To test how neural color representations evolve over time, we first used an exploratory analysis of the MEG data (multi-dimensional scaling, MDS), and second, a model-driven Representational Similarity Analysis (RSA) (Kriegeskorte, Mur, & Bandettini, 2008; Kriegeskorte, Mur, Ruff, et al., 2008; Popal et al., 2019). In the RSA, we compared three different theoretical (model) response

spaces, reflecting different factors that could be influential in the neural representation of color: the cone contrast differences between the stimuli; the perceptual differences between stimuli based on color discriminability; and color category, in which within-category pairs of stimuli are predicted to produce more similar responses than cross-category ones. Our results show an orderly representation of hue across the stimulus-induced response that is broadly consistent with each of the three *a priori* models above. Interestingly, we find a non-linear warping of the representational space that is dynamic, changing over time from the stimulus onset. These nonlinearities suggest a 'coarse to fine' dynamic, in which the cortical hue representation includes an initial clustering of similar hues before a later separation of these clusters to represent the finer differences between stimuli. We replicated the dynamics of these hue representations across the data from the two different tasks, and finding only a subtle indication of possible task-mediated differences in the later part of the response.

Materials and methods

Participants

We collected MEG data on 8 participants (5 female, 3 male, aged 21-34 years, mean=26.4 years). Each participant had no history of neurological and/or psychiatric disorders, normal or corrected-to-normal visual acuity, and normal color vision as assessed with Ishihara plates (Ishihara, 1990) and the Farnsworth-Munsell 100-hue test (Farnsworth, 1957). All participants provided informed consent prior to taking part, and all experimental procedures were approved by the Ethics Review Board of the McGill University Health Centre and were conducted in accordance with the Declaration of Helsinki. We collected MEG data for each participant during a single session, of approximately 2 hours. Each participant was familiarised with all tasks prior to the day of the MEG data collection, but all behavioural data reported here are from the MEG session.

Our sample size (n=8) is smaller than previous work testing for color representations with MEG (n=18, Hermann et al., 2022; Rosenthal et al., 2021), but larger than earlier work using fMRI (n=5, Brouwer & Heeger, 2009, 2013), and larger than our recent MEG work which included similar analysis approaches (n=6, Goddard, Shooner, et al., 2022). Teichmann et al. (2022) used real and simulated data to test the effect of participant and trial numbers on statistical power for classification analyses of MEG responses to visual stimuli. They report that sufficient data at a

subject level (i.e. enough trials) is likely more important for statistical power than sample size,

giving the empirical example of a data set with 9 participants completing 1600 trials having

equivalent power to data from 18 participants who completed 400 trials. In this study, we

prioritised many trials (>3,800 trials per participant) to increase this subject level power.

Visual stimuli

All visual stimuli were large circular patches of uniform color, serially presented. Each circular

stimulus was 40 degrees in diameter, presented in a raised-cosine envelope, and outside the

stimulus area the screen was set to a grey of mean luminance, as illustrated in **Figure 1.** We used

14 hues, chosen to be equally spaced around a circle of diameter 40 in the isoluminant plane of

CIE Lab space. The saturation of the stimuli was chosen to be close to the maximum achievable

within the gamut of the display.

Figure 1: *Spatial and temporal stimulus properties. **A**: Stimuli were 14 hues, equally spaced around*
a circle in the isoluminant plane of Lab space, presented at 3 luminance levels (not shown here).
*The background was held at mean grey (a=b=0). **B**: Each stimulus was a color presented in a*
*circular raised-cosine spatial envelope. **C**: Example timecourse of 2 trials: each stimulus was*
presented in a temporal raised-cosine envelope.

We determined the isoluminant plane of CIE Lab space for each subject individually using

heterochromatic flicker photometry (He et al., 2020; Smith & Pokorny, 1975) . At the start of the

experimental session, participants viewed a sinusoidally flickering stimulus, alternating between a

frame of a colored stimulus (as in **Figure 1B**), and an achromatic stimulus of the same spatial

properties. Participants used a mouse to adjust the luminance of the achromatic stimulus until

their impression of flicker was minimised, indicating their selection with a mouse click. After

performing at 3 settings for each of the 14 hues, we averaged the selected luminances for each

hue. We fit a sinusoid to these average offsets across hue direction, with amplitude and phase as

fitted parameters, and used the fitted values to adjust the experimental stimuli for perceptual

isoluminance.

In addition to the 14 hues that were isoluminant with the background, we used another 28 hues

that were of the same hue direction but with a 12.5% luminance increment or decrement, giving a

total of 42 unique stimuli.

After the stimuli were calibrated for their individual perceptual isoluminance, participants viewed

all stimuli sequentially, and verbally reported the color category of each stimulus, which were

recorded by the experimenter. Participants were restricted to using the labels 'red/pink', 'orange',

'yellow', 'green', 'blue' and 'purple'. Each stimulus was presented 3 times, totalling 126 trials. The

modal color labels for each unique stimulus were used to define each participant's color

categories, used in the category task (described under 'participant task' below) and the category

model (used in the Representational Similarity Analysis, described below).

No MEG data were collected during these preliminary tasks.

MEG methods: acquisition protocols

MEG data were collected with a whole-head MEG system (CTF OMEGA System) consisting of 275

axial gradiometers. For each MEG session we first collected 5 minutes of empty room recordings

which we used to estimate noise covariance of the sensors (see below). Prior to the participant

entering the magnetically shielded room, three marker coils were placed on the participant's

head. Marker positions, nasion, left and right pre-auricular points, and the participant's head

shape were recorded with a pen digitizer (Polhemus Isotrak, Colchester, VT), using a minimum of

500 points. Two EOG electrodes were places above and below the left eye, to record eye blinks

and eye movements during the MEG session. Two electrodes were placed across the plane of the

chest to collect electrocardiographic (ECG) signals, and a reference electrode was placed below

the participant's collarbone. Each participant's MEG data and simultaneous EOG and ECG signals

were collected at a sampling frequency of 2400Hz. In conjunction with these data, we collected

participants button responses using a VPixx ResponsePixx button box system (VPixx Technologies,

Canada).

MEG methods: display apparatus and calibrations

We displayed stimuli using a PROPixx DLP LED projector (VPixx Technologies, Canada, resolution

1920x1080), located outside the magnetically shielded room, to back-project images onto a

custom screen via two mirrors. Participants, lying supine in the MEG system, viewed the custom

screen, located above them, from a distance of 45cm. We used a Windows PC (Windows 7)

running MATLAB (R2017a) in conjunction with routines from Psychtoolbox 3.0 (Brainard, 1997;

Kleiner et al., 2007; Pelli, 1997) to generate and project the stimuli (refresh rate 60Hz, mean

luminance 106 cd/m2). The PROPixx DLP LED projector has a linear gamma, and was color

calibrated as described previously (Michna et al., 2007; Mullen et al., 2007, 2008). We precisely

aligned stimulus presentation times with the recorded MEG data using the VPixx 'Pixel Mode' to

record the output of a single pixel along with the MEG data.

MEG methods: experimental design and participant's task

Each participant completed the MEG data collection in a single session, which was divided into ten

blocks of 5-7 minutes each, with breaks between blocks. On each block of trials, participants saw

the same stimulus set (summarised in Figure 1) and performed a 1-back task requiring color

categorization or discrimination. In both tasks, the spatial and temporal properties of the stimuli

were identical, with each trial comprising a single color presented in a temporally raised-cosine

envelope (300ms in total, with 100ms onset, 100ms sustained presentation, and 100ms offset).

Across both tasks, the trials were presented in a continuous stream, and participants responded

with a keypress whenever 2 consecutive stimuli were an exact match (discrimination task), or a

color category match (category task, where same category was defined as stimuli that were given

the same label in the color naming task).

For both tasks, we used at least 42 trials of each of the 42 unique stimuli (1764 trials). These were

presented in a counterbalanced order where each stimulus was preceded by each of 42 stimuli at

least once. Since category matches were more common than exact stimulus matches, we added

extra exact match trials to the discrimination task blocks so that the average rate of match trials

was the same across the two tasks. The total number of trials in the discrimination task ranged

from 2066 to 2130 across participants, since the match rate in the category task varied with their

individual color labels.

In pilot testing, participants were faster and more accurate for the discrimination task than for the

category task: to approximately equate the tasks for difficulty in the experiment we used a longer

inter-stimulus interval (ISI) in the category task (700ms) than in the discrimination task (450ms),

giving stimulus onsets that were separated by 1000ms and 750ms respectively.

For both the category and the discrimination tasks we evaluated participants' performance using

their hit rate (HR: proportion of match trials with a button press) and false alarm rate (FA:

proportion of non-match trials with a button press) to calculate sensitivity (d') using the MATLAB

function *norminv* (inverse of the cumulative normal distribution):

228   d' = *norminv*(HR) – *norminv*(FA)

229   All participants performed these tasks well above the chance rate of d'=0. Across participants, the

230   average d' = 2.7 (range d'=1.8-4.1) on the category task, and d' = 3.3 (range d'=2.4-4.2) on the

231   discrimination task.

MRI methods: retinotopic and functional localisers

Each participant completed an MRI session in which we acquired high-resolution anatomical

images of their brains and functional data used to define regions of interest. All magnetic

resonance imaging took place at the McConnell Brain Imaging Centre, McGill University, Montreal,

Canada. For each participant we acquired two high-resolution three-dimensional whole head T1

images using an MP-RAGE sequence (TI=900ms, TR = 2300ms, TE = 3.41ms, 1.0mm^3 resolution),

and averaged these two images to generate the participant's anatomical template. Functional T2*

MR images were acquired on a 3T Siemens MAGNETOM Prisma system with 32-channel head coil.

Gradient-echo pulse sequences were used to measure blood oxygenation level-dependent (BOLD)

signal as a function of time. We identified the visual cortical regions V1, V2, V3, V3A/B, LO1/LO2

and hV4 for each participant using rotating wedge stimuli and expanding and contracting

concentric rings (Engel et al., 1994; Sereno et al., 1995), standard definitions of these areas

(Brewer et al., 2005; Goddard et al., 2011; Larsson & Heeger, 2006), and the foveal confluence

(Schira et al., 2009). To localize areas VO1, VO2 and hMT+ we used data from the retinotopic

mapping scans in conjunction with functional localisers for VO (Mullen et al., 2007) and hMT+ (Huk

et al., 2002). Full details of our retinotopic mapping procedures, including scanning protocols, data

preprocessing and area definition have been described previously (Goddard et al., 2019).

MEG data analysis: preprocessing and source reconstruction

Preprocessing, forward modeling and source reconstruction of MEG data were performed using

Brainstorm (Tadel et al., 2011, http://neuroimage.usc.edu/brainstorm). For each participant's

template anatomical (a high-resolution MRI image), we used the automatic segmentation

processes from Freesurfer 6.0 (Dale et al., 1999; Fischl et al., 1999) to define the grey/white

matter and pial/grey matter boundaries. Using Brainstorm, we imported the output of Freesurfer

and created a 15,000-vertex model of each participant's cortical surface. For each block of trials in

the MEG data, we aligned the participant's cortical surface model to the median measured marker

coil locations for that block by aligning the head shape data from the MRI with the head shape

relative to the marker coils, as recorded with the pen digitiser. For each functional run, we

generated a forward model for each model by applying a multiple spheres model (Huang et al.,

1999) to the participant's cortical surface model at this measured head location.

Functional data were preprocessed in Brainstorm with notch filtering (60, 120 and 180Hz),

followed by bandpass filtering (0.2-200Hz, using the Brainstorm default of an even-order linear

phase FIR filter). We preprocessed data from the empty room recording using identical protocols,

then used the output to estimate the noise covariance for the session. Cardiac and eye blink

artifacts were removed from functional data using signal space projection (SSP): cardiac and eye

blinks events were identified using default filters in Brainstorm, manually verified, then used to

estimate a small number of basis functions corresponding to these noise components, which were

removed from the recordings (Uusitalo & Ilmoniemi, 1997). From these functional data we

extracted an epoch of data for each trial: from -100 to 1000ms relative to stimulus onset, and

downsampled to 200Hz. Using the noise covariance estimate, regularized using the median

eigenvalue, we applied a minimum norm source reconstruction to this trial data.

For classification analyses we generated three datasets using each participant's functionally

defined cortical areas, as in previous work (Goddard, Shooner, et al., 2022). In the 'early visual

cortex' (EVC) region of interest, we included data from all vertices located within areas V1, V2 and

V3, while the 'ventral visual cortex' (VVC) region of interest included areas hV4, VO1 and VO2, and

'dorsolateral visual cortex' (DVC) included areas V3A/B, LO1, LO2 and hMT+. We found very similar

effects across these ROIs, which we believe most likely reflects the failure of the source

reconstruction to completely isolate responses from these adjoining cortical regions. Since there is

little difference across ROIs, below we show data from EVC only, but equivalent figures for VVC

and DVC are included in the Appendix.

MEG data analysis: classification-based analyses

We used a series of classification analyses to measure the amount of information about stimulus

color and luminance that was available in the neural responses, as measured with MEG. We

repeated these analyses for each 5ms bin to capture how this information changed over time. Out

of the total 15,000 sources in each participant's head model, the EVC ROI included an average of

731 sources (*SD* = 165), the VVC ROI included an average of 396 sources (*SD* = 67), and the DVC

ROI included an average of 434 sources (*SD* = 106). We reduced these with PCA and retained data

from the first *n* components, which accounted for 99.9% of the variance, for the classification

analysis (mean = 285.9, *SD* = 13.7 for EVC; mean = 283.6, *SD* = 17.4 for VVC; mean = 283.6, *SD* =

20.5 for DVC).

We conducted a series of pairwise classifications based on either stimulus hue or stimulus

luminance, with separate analyses for data from the category and discrimination tasks, and for

each ROI. In each case, we trained classifiers to discriminate between two categories of trial (i.e.

two hues or two luminance levels) and tested on held-out data. We report results obtained with a

linear support vector machine (SVM) classifier, using the MATLAB function *fitcsvm* with

'*KernelFunction*' set to *'linear'*. For all analyses we expressed average classifier accuracy in *d'* (a

unit-free measure of sensitivity). Chance classification performance yields *d'* = 0.

For classification analyses, we generated 'pseudo-trials' to increase signal-to-noise ratio along the

dimension of interest, and to reduce data variability due to stimulus dimensions orthogonal to the

dimension of interest (e.g. Goddard & Mullen, 2021). When training classifiers to discriminate

stimulus hue, we used pseudo-trials that were each the average of 12 trials of the same hue: 4 of

each luminance level. For data from a single task, this yielded 11 pseudo-trials for each of the 14

hues. For each pair of hues, we used these pseudo-trials in an 11-fold cross-validation, in each

case training on data from 20 pseudo-trials (10 from each hue) then testing the classification rule

on its classification on the held-out pair of pseudo-trials. We repeated this entire process 10 times,

with different partitions of the data into pseudo-trials. We performed an equivalent procedure

when classifying stimulus luminance: generating pseudo-trials that were each the average of 14

trials of the same luminance, 1 of each stimulus hue. This yielded 42 pseudo-trials for each of the 3

luminance levels, and for each pair of luminance levels we used a 42-fold cross-validation.

For both hue and luminance, we performed these classification analyses on data from within each

5ms time bin, to see how information about these stimulus features in the neural responses

unfolded over time. For decoding of stimulus hue in the EVC ROI, we additionally tested for cross-

temporal generalization: training classifiers on data from one time bin and then testing this

classification rule on data from each later time bin in held-out pseudo-trials.

MEG data analysis: Representational Similarity Analysis (RSA) and Multi-dimensional Scaling (MDS)

MEG data analysis: Representational Similarity Analysis (RSA) and Multi-

dimensional Scaling (MDS)

When decoding stimulus hue, pairwise comparisons of the 14 unique hue directions resulted in a

14x14 dissimilarity matrix (DSM) for each time bin where each cell of the matrix was defined by

classifier accuracy. To visualise how the neural representations of color were unfolding over time,

we used multidimensional scaling (MDS) to reduce these matrices to the best-fitting two-

dimensional representation of the 14 hues. Each MDS was completed using the MATLAB function

mdscale with '*criterion*' set to '*metricstress*' (minimising the stress normalised with the sum of

squares of the dissimilarities).

We also used Representational Similarity Analysis (Kriegeskorte, Mur, Ruff, et al., 2008; Nili et al.,

2014) to compare these matrices with three *a priori* models, based on different representations of

stimulus hue (**Figure 2**). Each of these models predicted that the dissimilarity of the neural

responses would be proportional to the stimulus difference, but with stimulus difference

quantified in three possible ways: in cone contrast (Figure 2A), CIE L*a*b* space (Figure 2B), or in

color category (Figure 2C). For each time bin, we rank-correlated each participants' observed DSM

with each of these three models. We compared model performance against the benchmark of the

'noise ceiling', which estimates the expected correlation of the observed data with a hypothetical

(unknown) 'true' model, given the noise in the data (Nili et al., 2014). The upper-bound of the

noise ceiling is calculated by correlating the group average DSM at each time bin with each

individual's DSM, which, due to overfitting, will overestimate the true model's correlation. The

lower-bound of the noise ceiling is calculated by correlating each individual's data with a group

average that excludes their own data, avoiding overfitting, and providing a lower bound on the

expected correlation with the true model.

Since our *a priori* models are positively correlated with each other (see legend of Figure 2 for

correlation values), they will tend to co-vary in their correlation with the data. To estimate the

extent to which these models predicted unique variance in classifier performance, we also

conducted a series of partial correlations, in each case testing the rank partial correlation between

the observed data and one model, controlling for correlation with a second model.

When decoding stimulus luminance, the pairwise comparisons resulted in a 3x3 DSM. We

conducted an RSA on these data, testing the data on two models, based either on luminance

intensity or luminance contrast. Both models are shown in insets in Figure 4.

Figure 2: Model dissimilarity matrices (DSMs) based on hue for Representational Similarity

Analysis (RSA). *We tested classifier data against 3 models of dissimilarity for the 14 stimulus hues,*

*based on stimulus cone-contrast (**A**), hue direction in CIE L*a*b* space (**B**) or color category (**C**). In*

each case, the upper plot indicates the stimulus representation according to the relevant model,

and the low plot shows the corresponding model DSM. The hues were selected to be equally

*spaced in CIE L*a*b* space (**B**); when replotted in a cone contrast space (**A**), the hues are not*

*equally spaced, but elongated along the S-cone contrast axis, while the category model (**C**) treats*

within-category hues as identical, grouped into 6 clusters equally spaced around a circle. The

*category model shown above (**C**) is an example model for one participant; in the analysis, each*
participant's color naming data were used to generate an individual category model. These three
models were positively correlated with one another, Spearman's rho r = 0.72 (cone contrast vs
hue), r = 0.25-0.44 (cone-contrast vs category), r = 0.68-0.73 (hue vs category), with variation in
correlation with the category model due to inter-participant variation in color naming.

Statistical analyses

To test whether classifier accuracy was above chance, and to test whether model correlations and
partial correlations were above zero, we used Bayes Factor analyses, an alternative to the
traditional frequentist approach (Kass & Raftery, 1995; Morey & Wagenmakers, 2014). A Bayes
Factor compares evidence for competing hypotheses; here we report where there is moderate
(BF>3) or strong (BF>10) evidence in favour of the alternate hypothesis, or at least moderate
(BF<1/3) evidence in favour of the null hypothesis. We implemented all Bayes Factor analyses
using a MATLAB package (Krekelberg, 2021).

To test times where cross-temporal decoding was significantly different from chance, we used an
alternative approach, adopting the strategy of Hermann et al (2022), based on a 2D cluster
threshold (Maris & Oostenveld, 2007). We used permutation testing to obtain *p*-values for each
train/test time, permuting the sign of the classifier accuracy data on a participant basis 1000
times, and taking the p-value as twice the smaller of the proportion of average values above or
below the observed average (2-sided test). We then used a cluster-threshold to find clusters of
times that were significantly above or below 0 (*p*<0.05) with a cluster defining threshold of *p*<0.01.

Results

We recorded MEG data while participants viewed a series of colored stimuli and performed either
a categorization task or a discrimination task. We used a series of classification analyses to
measure how neural representations of stimulus hue and luminance unfolded over time.

Classification of stimulus hue and luminance

In **Figure 3** we show the average classifier accuracy, collapsed across all pairwise comparisons,
when decoding hue and luminance. Across the two tasks, we found qualitatively similar
timecourses of average classifier decoding for both stimulus features, with a trend toward higher
classifier accuracy in the category task. This tendency for higher classifier accuracy in the category
task may reflect the longer inter-stimulus interval for the category task (700ms) compared with
the discrimination task (450ms), which we included to better match the tasks for difficulty (see

Methods for details). The onset of above-chance decoding was approximately 100-120ms after stimulus onset in each case, which is slightly later than reported previously (e.g. Hermann et al., 2022). This later onset is likely because our stimuli did not onset abruptly but ramped on gradually, as shown in Figure 3.

Figure 3: Decoding of stimulus hue (upper) and luminance (lower)*. Average classifier accuracy (n=8) across the two tasks is shown for classification of hue and luminance. Error bars are standard errors of the between-subject means. Thin black lines in the upper part of each plot, along with the shaded grey background, indicate the temporal envelope of the stimulus, relative to trial onset. Dots below the x-axis indicate the results of the Bayes Factor analyses. Colored dots show times when the classifier performance was above chance (small dots indicate moderate evidence BF>3, large dots indicate strong evidence, BF>10). Small grey dots indicate times where there was at least moderate evidence in favor of chance performance (BF<0.3).*

Trend towards achromatic contrast, rather than intensity, in luminance decoding

Our stimuli were presented at three luminance levels: luminance decrements, stimuli isoluminant with the background, and luminance increments. If the neural responses were primarily driven by

the luminance intensity differences between these, then discriminating increments vs decrements

should lead to the highest classifier performance. However, if the neural responses to these

stimuli were driven by luminance contrast more than intensity, then classifier performance should

be highest when responses to the isoluminant stimuli are compared with either increments or

decrements. These predictions are shown in the model DSMs (**Figure 4**, insets). Representational

Similarity Analyses of the luminance decoding data are shown in **Figure 4**.

For both the category and the discrimination task, the data tended to be more consistent with the

model based on achromatic contrast, rather than intensity. However, these effects were not

particularly robust, as seen in the results of the Bayes Factor analysis in Figure 4. The average

DSMs in Figure 4 are shown for four epochs to match those used for the hue analysis (see next

section). Across these epochs, the average DSMs are consistent with the contrast model rather the

intensity model across the first three epochs shown (110-455ms), with the lowest classifier

performance for increments vs decrements in each case, but there was moderate variability across

participants, which is seen in the standard error of the model correlations. The inter-participant

variability is also seen in the noise ceiling, where the lower bound is not consistently above zero at

time bins where there was above-chance classifier performance (as seen by comparing the dotted

lines in Figure 4 with the average decoding of luminance in Figure 3). In these ways, these data are

consistent with neural responses being driven by achromatic contrast, rather than intensity, but

they provide only modest support for an effect in this direction.

Figure 4: Representational Similarity Analysis of stimulus luminance, for the category (upper) and discrimination (lower) tasks. *Average correlations (n=8) with the achromatic contrast and intensity models, along with the average dissimilarity matrices (DSMs) for four epochs (using the same epochs as in the reporting of hue RSA, below). Note that the achromatic contrast versus intensity models (shown in insets at the left) are perfectly negatively correlated, so the model correlations are mirror symmetrical about the x-axis (time axis) and positive correlation with one model will drive negative correlation with the other. The lower bound of the noise ceiling (Nili et al., 2014; see methods) is indicated by the dashed black line. Error bars are standard errors of the between-subject means, and dots below the x-axis indicate the results of the Bayes factor analysis, with plotting conventions as in Figure 3.*

Dynamics of hue representations are consistent across task

Since each participant viewed stimuli of 14 different hues, the pairwise classifications of stimulus hue included 84 different comparisons. We include the average DSMs based on hue decoding for four epochs (see **Figure 5**). To better visualise the stimulus representations suggested by these DSMs, we also include MDS solutions based on each of these DSMs. We selected the four epochs in Figure 5 (also used in Figure 4) as equal-duration periods that together span the times with above-chance classification of stimulus hue. For an alternate visualization, with shorter epochs that span the entire timecourse, we also include movies of the MDS solutions for each 50ms epoch in the Appendix.

In comparing the DSMs and MDS solutions in Figure 5 with the model predictions in Figure 2, none of the epochs in either task provide a clear match to any of the 3 *a priori* models. Nonetheless, there are several features of the data that suggest that they reflect meaningful structure and can provide insight into how the neural representation of hue unfolds over time, including features that concur with one or more of the *a priori* models. For instance, all three models predict that classifier performance will tend to increase as hue difference increases, and this is consistent with all observed DSMs, which have relatively low classifier performance (close to grey) around their negative diagonals. In the MDS solutions, this is seen in tendency for hues to be arranged in order according to physical similarity (albeit with some clustering and/or folding).

For both the category task and discrimination task data, there is an evolution of the hue representation over time, as seen in the differences across the 4 epochs in Figure 5. The most striking feature of the data in Figure 5 is that this evolution across epochs is remarkably consistent for data from different tasks, particularly across the first 3 epochs. This is notable given that data for the two tasks were collected across different MEG recordings. This replication indicates a reliable but dynamic structure in the neural representations of hue, reflecting a reorganization of hue responses over time.

In the first two epochs of Figure 5, there is relatively low classifier performance amongst green hues: this is seen in the squares of grey in the DSMs, and the clustering of the greens in the MDS solutions. Interestingly, the greens do not remain clustered across the entire stimulus-induced response but show greater spacing in the third epoch of Figure 5. This pattern is replicated across

470   both tasks. In the third epoch, MDS solutions from both tasks include a more substantial deviation

471   from a circular arrangement, forming a 'figure of 8' with a crossing point around blue/yellow.

472   Although the exploratory nature of these analyses precludes strong conclusions, they present

473   interesting avenues for future work, as discussed in the general discussion below.

Figure 5: *Average dissimilarity matrices (DSMs) and corresponding MDS solutions for data from the category task (upper) and discrimination task (lower). In each case, we considered data from four epochs of 115ms each, spanning the times that included above-chance classification of hue for both tasks. In both cases, the average decoding of hue (d') is replotted from Figure 3, and grey shaded regions indicate the epochs corresponding to the four DSMs.*

RSA suggests a 'coarse to fine' transition in color representations

To better characterise how these hue representations change over the course of the stimulus-induced response, we performed a Representational Similarity Analysis, comparing the data with models based on cone contrast differences, hue difference, and color category (**Figure 6**) for data from both the discrimination and category tasks. All three models performed well overall (Figure 6A), with similar timing, which likely reflects the substantial overlap in their predictions based on their inter-model correlations (see Figure 2). Bayes Factors revealed moderate or strong effects across most times when classification of hue was above chance, and average model correlations that approached the lower-bound of the noise ceiling in each case. The strong performance of all three models we considered (Figure 6A) suggest a possible ceiling effect, where the shared predictions of these correlated models may account for performance approaching the lower-bound of the noise ceiling across the stimulus-induced response. These correlations offer limited additional insight into the dynamics of color evident in the dissimilarity matrices and MDS solutions (Figure 5).

To better isolate the predictions of each model from the others and test whether the unique variance captured by each model varies over time, we performed partial correlations across each pair of models. The most informative of these comparisons were partial correlations based on the hue and category models, as shown in Figure 6**B**. Interestingly, although we initially included the 'category' model to potentially capture later, higher-order color representations associated with color naming/categorization, we instead found that the unique variance captured by the category was most evident in the earlier part of the stimulus-induced response. This complements observations from the exploratory analysis (Figure 5) that the early colour responses included clustering of some hues of the same category, most notably the greens. We also found that the hue model captured variance beyond that of the category model later in the response. This suggests that later in the response, representations of similar hues in the same colour category are more distinct, reflecting their different appearance more than their category. Together, these partial correlations suggest that the dynamic reorganization of colour representations could include a transition from a coarse, early response, with greater clustering of hues of the same category, to a representation that accentuates the finer appearance differences between within-category hues.

Partial correlations between the cone-contrast model and these models are shown in the

Appendix (Figure A9) and in each case reflect a similar trend to that seen in Figure 6B, where the

category model tends to capture more unique variance in the earlier part of the stimulus-induced

response, whereas the hue model tends to emerge later. Together, the partial correlation data

indicate that there is a transition at 300-350ms after stimulus onset from the category to hue

model better accounting for unique variance. This pattern was replicated across data sets from

both tasks.

Despite very similar overall patterns of effects, the partial correlations also reveal a trend towards

task dependence, with the hue model tending to account for more unique variance in when

participants were performing the discrimination task, rather than the category task. The direction

of this effect is consistent with the task requirements, since the discrimination task, but not the

category task, required participants to pay attention to finer differences in hue, including within a

colour category.

Figure 6: Representational similarity analysis. **A**: *Average correlations between individual participants' dissimilarity matrices for hue, and the cone contrast, hue and category models (see Figure 2 for models). **B**: Partial correlations between the hue and category models, showing the unique variance captured by these two models. In **A**, the lower bound of the noise ceiling (Nili et al., 2014; see methods) is indicated by the dashed black line. Error bars are standard errors of the between-subject means, and dots below the x-axis indicate the results of the Bayes factor analysis, with plotting conventions as in Figure 3.*

As a final analysis of the dynamics of hue representations, we performed a cross-classification of stimulus hue over time (**Figure 7**). We trained classifiers to discriminate each pair of hues using data from one time bin and tested the predictions of this classifier for data from all later time bins. When there are sustained representations of hue, we expect that classifier performance would show above chance performance for temporally offset test data. Conversely, if there were a dynamic reorganization of hue representations, then classifiers trained on data from an earlier time might show chance or even significantly below chance performance when trained on data from after such a reorganization.

Consistent with previous reports (e.g. Hermann et al., 2022), we found that decoding of stimulus hue was highest for adjacent time bins (close to the negative diagonal of the heatmaps in Figure 7), but also with some significant clusters of off-diagonal cross-decoding, particularly from approximately 150-350ms, across data from both tasks. Since the RSA results (partial correlations in Figure 6B) suggest that there may be a transition in the neural representation of hue at around 300-350ms post-stimulus onset, we were interested in whether the cross-decoding of hue would also suggest a reorganization at this time. If there were positive classifier performance within the heatmap regions highlighted in orange in the insets of Figure 7, and low or negative classifier performance within the regions highlighted in teal in the same insets, this would provide additional support for a reorganization in hue representations at around 340ms. While we found some evidence of cross-temporal generalization before this time, we found weaker evidence of cross-temporal decoding after this time, and no clusters of classifier performance that was significantly below-chance. Overall, while the data in Figure 7 show some tendency towards the predicted pattern, these data do not provide as clear support for a reorganization in hue representation as the RSA results.

Figure 7: Cross-temporal decoding of hue over time for category task (left) and discrimination

task (right). Average accuracy across all pairwise classifications of stimulus hue, when trained on

data from one time bin and tested on data from each later time bin. To the top and right of each

heatmap, the average classifier accuracy within each time bin is replotted from Figure 3 for

comparison with cross-decoding. Within the heatmap, clusters of significant cross-temporal

decoding (meeting a permutation test threshold of p<0.05, and cluster threshold p<0.01) are

outlined with a thin black line. For comparison with the epochs in Figure 5, dashed lines on the

heatmaps indicate 110ms, 340ms and 570ms post stimulus onset. The insets in the lower-left of

each heatmap highlight regions where we would expect to see good generalization (red or yellow)

and poor or below-chance generalization (grey or teal) if there were a reorganization of the neural

representation of hue at around 340ms.

Discussion

In this paper, we performed a series of classification analyses on the MEG recordings to explore how the neural response to colour unfolds over time, using a combination of exploratory analyses (MDS), and hypothesis-driven RSA. We chose stimuli with a relatively fine sampling of the hue space (14 hue directions) with the aim of revealing distortions of the neural representation of colour that could distinguish between different models of colour representation.

The importance of comparing multiple models of colour representation

In the RSA, we tested three different representations (models) of stimulus colour, against the human cortical MEG data, with the aim of asking which model, if any, best captures the MEG-generated neural code. All models are based on human behavioural data. One representation of our chromatic stimuli is in a CIE L*a*b* space, where the fourteen suprathreshold colours are equally spaced around a colour circle in terms of their perceived colour differences ('hue model'). In a second representation, these same stimuli are placed in a space based on the relative responses of the three different cone types to the stimuli ('cone contrast model'). In this space, stimuli vary in a non-uniform way although retaining the overall colour stimulus ordering. The distortions in the cone contrast space reflect, at least in part, the differential contrast sensitivities of the two underlying cone opponent mechanisms, with the S-cone opponent direction elongated due to its poor contrast sensitivity compared to the highly sensitive L/M opponent mechanism (Cole et al., 1993; Sankeralli & Mullen, 1996). We selected a cone contrast model as it represents lower-level cortical responses in the visual system and provides a useful benchmark against which to compare other models. The simplest account of cortical colour representations would be that they reflect their subcortical inputs, so a robust test of higher-level models is whether they account for colour representations better than this lower-level one. The third colour representation we tested is based on the categorical groupings of our 14 colours, measured individually for each participant's naming data, which produces a re-arrangement of the 'hue model' stimuli into clusters, predicting high within-category similarity as well as an orderly progression of categories based on their hue similarity ('category model').

The challenge of overlapping models

These three models are conceptually distinct, and were selected as biologically plausible color spaces that may best capture the neural code. However, it is important to note that they are also

correlated and so make substantially overlapping predictions in accounting for the observed data (e.g. the same ordering of hues). We overcome this issue by using partial correlations to make comparisons between the models. For example, partial correlations in the hue and category models in Figure 6B reveal the variance uniquely captured by these two models, allowing differences to be seen that were invisible previously (Figure 6A). As mentioned above, comparing higher-level models with one based on a lower level of processing provides a valuable benchmark when considering what constitutes evidence in favor of a higher-level model, and for both hue and category models we find evidence that they significantly outperform the cone contrast model at different times (see Appendix).

Robust decoding of luminance and hue

Across the stimulus-induced response, we found robust decoding of luminance and hue for both tasks, consistent with previous evidence that stimulus color can be decoded based on a topographic cortical representation of color in the MEG data (e.g. Goddard, Carlson, et al., 2022; Goddard, Shooner, et al., 2022; Hermann et al., 2022; Rosenthal et al., 2021; Teichmann et al., 2019, 2020). Decoding of both luminance and hue reached significance from approximately 100ms after initial stimulus onset and is optimal between 200 to 400ms. These onsets are slightly later than previously reported, and, unlike previous studies, we did not find evidence for decoding of luminance preceding that of hue (Hermann et al., 2022; Rosenthal et al., 2021). However, we note that our gradual onset stimuli may account for the later onset of the responses and are not optimised for comparing the onset of hue and luminance responses.

Patterns of luminance decoding (Figure 4) tended to indicate a neural representation based on luminance contrast, rather than luminance intensity. That is, accuracy was generally higher when the classifier was trained to discriminate between the isoluminant stimuli and those with luminance contrast, rather than between luminance increments and decrements, even though the latter have a greater difference in intensity. However, while decoding of luminance was highly robust, the pattern of decoding across luminance levels was variable across participants, and hence yielding only modest support for a neural representation of based on luminance contrast rather than intensity.

Where in cortex are these hue and luminance representations located?

The spatial resolution of MEG source reconstruction does not allow us to make strong inferences

about which visual cortical areas are contributing to the color representations. We found similar

patterns of results across early, ventral, and dorsal visual cortex (see Appendix). This could reflect

dynamics that are propagating across these interconnected areas and/or that signals from these

nearby regions were not well separated in our analyses. For this reason, the timing of the

observed effects provides the clearest constraints on where these neural representations are

located, which we consider below.

Orderly hue representations, with minimal differences across task

RSA revealed hue representations that were less variable across participants than the luminance

ones, possibly because our experimental design included larger variation in stimulus hue than

luminance. From the onset of above-chance decoding, our data were consistent with all three of

the models tested. This suggests that, from the earliest cortical responses to color, there is an

orderly representation of hue. If the earliest stimulus-induced responses found here reflect the

feedforward information in lower visual areas, then our data differ from fMRI studies showing that

the responses of areas such as V1 and V2 lack an orderly representation of hue, with orderly

responses emerging only in higher areas such as hV4 and VO (Brouwer & Heeger, 2009). This

discrepancy could be related to the fact that, in the fMRI study, participants' attention was

diverted from the colored stimuli. Alternatively, it could be that any orderly representation of hue

in V1 was relatively transient in the fMRI study, and beyond the temporal resolution of fMRI. The

data presented here are consistent with more recent evidence from optical imaging suggesting a

'pinwheel' spatial organization of hue responses in V1 (Li et al., 2022).

The hue representations found here showed interesting temporal dynamics, considered below,

but only modest variation with task. Using fMRI, Brouwer and Heeger (2013) demonstrated that

areas hV4 and VO1 showed evidence of categorical clustering in hue representations, but this was

present only when participants were engaged in a color naming task, and not when their attention

was diverted. We were interested in whether the categorical clustering reported by Brouwer and

Heeger (2013) was specific to a task requiring categorization, or a more general effect of attending

to stimulus color. For this reason, we included a task requiring attention to color category (color-

name 1-back task) as well as a task that required attention to color but without requiring

categorization (exact-match 1-back task). The MDS and RSA analyses suggested that hue

representations were largely consistent across the two tasks, but there was one noticeable trend

in the RSA results in the direction of a task-specific effect, which we consider in the following

section.

Dynamics of hue representations, and a 'coarse to fine' transition

Our results show some interesting dynamics in the cortical response to color that are robust and

replicated across data from both tasks. These dynamics are evident both in the MDS solutions (see

Figure 5 or movies in the Appendix), and in the RSA partial correlations (Figure 6B). The partial

correlations in the RSA, key to comparing the different models, indicate that there is a

reorganization of the representation of color at around 300-350ms after stimulus-onset. Prior to

this time, the data tend to be more consistent with the category model, supporting a

representation based on a clustering of within-category hues. In the group data, this is seen in the

MDS solutions primarily as a clustering of the neural responses to the green hues, although the

dominance of the greens here may be driven by the fact that in our stimulus set, green was the

largest category. After this reorganization, there is a more even separation of hues within the

same category, as seen in the unique variance captured by the hue model in the latter part of the

response. Partial correlations between the cone contrast model and the other models (see

Appendix) are also consistent with these trends: the category model tends to capture unique

variance before 300-350ms after stimulus onset, and the hue model is more likely to capture

unique variance after this time of reorganization.

Inspection of the MDS solutions suggests that the neural data has systematic deviations from each

of the three models we considered. Of course, it is possible that there are other color

representations that would more closely match the observed data than the models we

considered. Although we used the CIE L*a*b* space as an approximate model of color appearance

in our 'hue' model, there are numerous models of color appearance based on apparent color

difference or discriminability (Fairchild, 2005), or on natural image statistics (Smet et al., 2016)

which would make similar predictions. However, given the substantial reorganization of these

representations over time it is unlikely that there is any single (static) model that could capture

the neural representational space across time. For instance, our 'hue' model may not be an

accurate model of perceptual difference, since although CIE L*a*b* space is designed to be

approximately perceptually uniform, it contains known deviations from a truly Euclidian space

(Ennis & Zaidi, 2019). However, we find more pronounced deviations from the model based on CIE

L*a*b* space than predicted by these known nonlinearities, and more importantly, the MDS

results suggest that the neural data deviate from the CIE L*a*b* space in different ways over time: while greens are initially more clustered together than predicted by the hue model, at later times they are relatively more separated than other hues. Overall, the most pronounced feature of the hue representations in our data is their dynamic reorganization over time.

It is interesting that the category model performs best in the earlier part of the neural response. Since higher-level areas like hV4 and VO are associated with higher-level responses to color, including categorical representations (Brouwer & Heeger, 2013), we might expect that any categorical effects would emerge later in the stimulus-induced response, possibly incorporating feedback effects related to task and/or language. This 'earlier' timeframe of the categorial model includes responses from response onset, likely driven by areas such as V1 and V2, up to 300ms after stimulus onset, by which time the data would also include responses from higher-order areas such as hV4 and VO, and more anterior color centres (Lafer-Sousa et al., 2016; Zeki & Bartels, 1999). For instance, subdural electrode recordings in human ventral visual cortex, anterior to hV4, showed neuronal responses to color from ~100ms after stimulus onset, peaking at ~150ms after stimulus onset (Murphey et al., 2008).

After ~300ms post stimulus-onset, there is no evidence of increasing categorical clustering. Hence, our data argue against the idea that any feedback or recurrent processing within cortex is driving hue representations to become more categorical over time. Instead, we find that hue representations of finer, within-category differences become stronger over time. This transition may echo the 'coarse to fine' strategy in different aspects of spatial vision, with the coarse color structure processed before the fine within-color detail. While coarse-to-fine processing typically refers to neural responses shifting from lower to higher spatial scale information over time (e.g. Goffaux et al., 2011; Watt, 1987), equivalent coding principles have been implicated in orientation coding (Ringach et al., 1997, 2003; Shapley et al., 2003), and face processing (Dobs et al., 2019), where initial responses are more coarsely tuned, preceding representations of finer differences (i.e. wider then narrower tuning for orientation responses, face age and gender preceding representations that differentiate between face identities).

The late time at which we observe this transition to an enhanced representation of finer hue differences (~300ms) suggests that these effects could be at least partly driven by top-down effects of attention, which have previously been reported to produced effects consistent with

sharpening of population tuning for color (Bartsch et al., 2017; Goddard, Carlson, et al., 2022). In

both conditions, participants were attending to color, and across both tasks we saw similar

dynamics, consistent with a transition from coarser to finer hue representations. However, at later

stages of the stimulus induced response there was stronger partial correlation with the hue model

for data from the discrimination task than for the category task at corresponding times (Figure

6B). This task-specific effect is much subtler than the fMRI differences when comparing a

categorical task with diverted attention (Brouwer & Heeger, 2013), suggesting that the effect of

attending to a colored stimulus, rather than specifically attending to its category, is of greater

relevance.

A final notable feature of the hue representations in our data is in the latter part of the neural

response (e.g. the 340-455ms and 455-570ms epochs in Figure 5), the hue closest to yellow is also

close to one or more blue hues. This is clearest in the MDS results in the 340-455ms epoch, where

the hue circle is represented more as a 'figure of 8', with crossing at yellow/blue, but within

greenish and reddish hues there is greater separation. This is not accounted for by the S-cone

inputs being slower to reach cortex than red-green signals carried by the parvocellular pathway

(e.g. Cottaris & De Valois, 1998), since the effect we observe is well after the initial cortical

response, and because the effect is in the opposite direction, with a relatively late encoding of

red-green differences. Although these observations are still speculative and based on an

exploratory analysis, we note that in both tasks, the yellow/blue crossing occurred not for the

extremes along the S-cone isolating axis, but closer to unique yellow/blue, and the daylight axis

(Delahunt & Brainard, 2004). We did not design this work to test for such asymmetries, but these

observations could motivate future work to characterise these.

Conclusion

In summary, in this MEG study we reveal a dynamic reorganization of the cortical response to

color over time, which is remarkably consistent across our two tasks. We compared three

conceptually distinct, biologically plausible models that may best capture the neural code, using

partial correlations to isolate the unique data variance accounted for by each model. The resulting

model dynamics reflects a 'coarse to fine' transition over time from an early categorial grouping of

colors in the cortical representation evolving into to a finer, more even separation of the hues.

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Data Availability

Data from MEG experiments are freely available online from the Open Science Framework (doi: *Will be provided upon acceptance for publication*). This online repository includes deidentified raw data from the MEG experiments and details of the stimulus timing for each participant.

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