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; Rozman et al., 2024; 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. Across participants, the phase of the fitted sinusoid aligned with stimulus hue such

that the maximum correction was applied to the stimuli closest to the L-M isolating axis, as

expected for a correction to adjust for individual variability in the luminance mechanism. The

maximum amplitude of this achromatic correction (L=M=S) varied across individuals (range 0.8-

4.9%, mean 2.9% cone contrast).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

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

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

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

193　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):

$$d' = norminv(HR) - norminv(FA)$$

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

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

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. Trials were sampled without replacement, to ensure that each pseudo-trial

was comprised of independent data. 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)

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: as Euclidian distance between stimuli 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 fit a series

of general linear models (GLMs). For each GLM, we used the 3 models as regressors, applying

spm_orth.m (from SPM12, *www.fil.ion.ucl.ac.uk/spm/doc*) to perform recursive Gram-Schmidt

orthogonalization (Golub & Loan, 1996) of these regressors so that the final regressor would only

capture variance that is predicted by the final model and orthogonal to the other two models. We

then converted each data matrix to ranked classification accuracy values and fit these ranks with a

GLM comprised of the orthogonalized regressors, using the MATLAB function *fitglm.m*. We

repeated this process three times, so that each model was the final regressor in one GLM,

providing the parameter estimates (beta weights) which indicated any unique variance captured

by this 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 5.

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*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 parameter estimates (beta weights) 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).

Results

We recorded MEG data while participants viewed a series of colored stimuli and performed either a categorization task or a discrimination task. During both tasks, reconstructed sources were focussed in visual cortex, and showed similar average event-related fields over time (**Figure 3**). To quantify the stimulus-related information present in the pattern of response across cortex, and to explore how neural representations of stimulus hue and luminance unfolded over time, we used a series of classification analyses.

*Figure 3: Estimated distributed source amplitudes (RMS values), averaged (n=8) for category (**A**) and discrimination (**B**) tasks, projected onto a partly inflated template cortical surface, viewed from above (left) and below (right). In **C** and **D**, the average response (event related fields) across 8 participants and all sensors is plotted, grouped by the 14 stimulus hues (**C**) or 3 luminance levels (**D**). Thin black lines in the upper part of each plot in **C** and **D**, along with the shaded grey background, indicate the temporal envelope of the stimulus, relative to trial onset*

Classification of stimulus hue and luminance

In **Figure 4** 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

407 slightly later than reported previously (e.g. Hermann et al., 2022). This later onset is likely because

408 our stimuli did not onset abruptly but ramped on gradually, as shown in Figure 4.

Figure 4: 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 5**, insets). Representational

Similarity Analyses of the luminance decoding data are shown in **Figure 5**.

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 5. The average

DSMs in Figure 5 are shown for four time windows to match those used for the hue analysis (see

next section). Across these time windows, the average DSMs are consistent with the contrast

model rather the intensity model across the first three time windows 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 5 with the average decoding of

luminance in Figure 4). 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 5: 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 time windows
(using the same time windows 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 4.

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 time windows (see **Figure 6**). To better visualise the stimulus representations suggested by

these DSMs, we also include MDS solutions based on each of these DSMs. When the DSMs are

reduced with MDS, the resultant 2D solutions do not capture all information that may be present

in the DSM. However, there is a reasonable *a priori* prediction that the neural representation of

hue will be close to 2D, and the 14 stimulus hues can be represented in a number of different 2D

spaces (such as the a*b* plane of L*a*b* space). In this way, the MDS solutions provide a useful

insight into how the neural representations are structured if we assume that they are 2D. To test

how well each MDS solution captured the corresponding DSM, in each case we rank-correlated

(Spearman's rho) the original DSM with a distance matrix defined by the best-fitting 2D MDS

solution (Mantel test). These values were similarly high across time windows (see Figure 6).

We selected the four time windows in Figure 6 (also used in Figure 5) as equal-duration periods

that together span the times with above-chance classification of stimulus hue. We chose to divide

this period into four time windows in order to visualise the dynamics indicated by the RSA results

(reported below), particularly the suggestion of a reorganisation at around 300-350ms after

stimulus onset. For an alternate visualization, with shorter time windows that span the entire

timecourse, we also include movies of the MDS solutions for each 50ms time window in the

Appendix.

In comparing the DSMs and MDS solutions in Figure 6 with the model predictions in Figure 2, none

of the time windows 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 time windows in Figure 6. The

most striking feature of the data in Figure 6 is that this evolution across time windows is

remarkably consistent for data from different tasks, particularly across the first 3 time windows.

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 time windows of Figure 6, 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 time window of Figure 6. This pattern is

replicated across both tasks. In the third time window, MDS solutions from both tasks include a

more substantial deviation from a circular arrangement, forming a 'figure of 8' with a crossing

point around blue/yellow. Although the exploratory nature of these analyses precludes strong

conclusions, they present interesting avenues for future work, as discussed in the general

discussion below.

Figure 6: *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 time windows 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 4, and grey shaded regions indicate the time windows corresponding to the four DSMs. The colormap used to plot each DSM is scaled by the maximum d' of each matrix, indicated above each DSM. In the upper right of each MDS plot, rho values indicate the DSM/MDS correlation (Mantel test, see text for details).

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 7**) for data from both the discrimination and category tasks. All three models performed well overall (Figure 7A), 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 7A) 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 6).

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 used a GLM analysis (detailed in the methods), with results shown in Figure 7**B**. The cone contrast model, which we expected to perform best for the earliest cortical responses, accounted for some unique variance in the first half of the response, but this was not clearly restricted to the response onset for either task. 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 6) that the early colour responses included clustering of some hues of the same category, most notably the greens. For data from the discrimination task, 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. This indication of a transition from the category to hue model better accounting for unique variance occurs at 300-350ms after stimulus onset, which is around the boundary between the second and third time windows in Figure 6. Together, these results 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.

Despite very similar overall patterns of effects, the GLM analyses (Figure 7B) also suggest some task dependent effects, with the hue model accounting for unique variance at ~400ms after stimulus onset 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 7: 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**: Parameter estimates from GLM, quantifying the unique variance captured*
by each model, beyond the variance shared by other models (see methods for details) . Colored
*regions of the Venn diagrams in the top right of **A** and **B** illustrate the variances captured by these*

Discussion

570 In this paper, we performed a series of classification analyses on the MEG recordings to explore

571 how the neural response to colour unfolds over time, using a combination of exploratory analyses

572 (MDS), and hypothesis-driven RSA. We chose stimuli with a relatively fine sampling of the hue

573 space (14 hue directions) with the aim of revealing distortions of the neural representation of

574 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 the GLM analyses to compare the models in terms of the unique variance they capture (Figure 7B) which revealed differences between the three models to be seen that were invisible previously (Figure 7A). 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.

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 5) 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. Although our source reconstruction indicated that sources were primarily located in visual cortex (Figure 3A-B), we found similar patterns of classifier 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 6 or movies in the Appendix), and in the GLM analyses (Figure 7B). The GLM analyses, 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 cone contrast model, potentially reflecting responses that modulate with cone

contrast, and also with unique variance captured by 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 for the discrimination task.

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. Instead, we found early contributions of the

category model, meaning it is very unlikely that these reflect a higher-level representation where

hues of the same semantic label are coded in a more similar way than hues of different names.

Instead, we interpret these findings as suggesting that during earliest cortical responses there are

hues which evoke very similar responses, which are then differentiated later in the response, but

that this early clustering is not related to the influence of language. The '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 evidence in favor of the hue model for

data from the discrimination task than for the category task at corresponding times (Figure 7B).

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 time windows in Figure 6), 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 time

window, 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

GLM analyses 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 (https://doi.org/10.17605/OSF.IO/KMFZ7). This online repository includes deidentified MEG data from the MEG experiments and details of the stimulus timing for each participant.

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1006