

Computational Study

1 Introduction

I am interested in the question of whether a melanopsin based signal might be useful to estimate the illuminant(s) in a scene.

Following initial psychophysical experiments, where results did not indicate a strong or simple relationship, I chose to take a step back and examine the problem that the HVS is faced with in a natural environment, which colour constancy solves. Through this route I hoped to answer the questions:

1. Would it be sensible for the HVS to use a melanopic signal to help solve this problem?
2. If so, in what way would it be used?

In this document I shall give an overview of the 4 main scripts I have written, going through the logic and results of each. All scripts are available at <https://github.com/da5nsy/Melanopsin-Computational>

2 melcomp_1

In melcomp_1 I considered three questions:

1. Considering only daylight spectra (excluding reflective surfaces for now), can a melanopic signal predict the chromaticity of daylight more precisely than signals provided by other retinal cell populations?
2. Now considering also object reflectances, does a melanopic signal provide a means of calculating a sign and weight of shift required to counteract the chromatic shift induced upon objects by a change in daylight conditions?
3. Are there objects, or luminance levels, or daylight chromaticities for which the melanopic signal is particularly effective or ineffective at performing the above task

The final question was asked with the hope that such limitations might provide nice tests to perform psychophysically.

Foundational data consisting of the Granada daylight dataset, the Stockman-Sharpe 10 degree cone fundamentals, the Lucas et al. melanopsin fundamental, and the a subset of the Vrhel et al. reflectances (consisting of the ‘natural’ reflectances) were used.

Tristimulus values (L,M,S) and analogous melanopic values (Mel) were calculated for each illuminant within the Granada dataset. The real-world correlate of this computation would be to observe a spectralon tile under each daylight condition.

MacLeod-Boynton chromaticity co-ordinates were calculated, and three-dimensional plots were made to display l_{MB} against s_{MB} against [L,M,S,Mel] in turn. Each plot was very visually similar to the Mel plot shown in 1. As can be seen, although there exists some relationship between the chromaticity values and the 'Mel' value it does not appear as though one could predict the other. All that can be said from this relationship is that if 'Mel' is above a certain threshold, it is likely to be in a group with a low s_{MB} value. The real-world correlate of this is to say - *if the day is bright enough, I can be fairly confident that the chromaticity of daylight will be relatively warm in colour*. This is as expected, since the brightest daylight conditions are likely to be those with unobstructed direct sunlight, which is warmer in colour than illumination provided by the blue sky. This relationship seems relatively weak however, and could not be used to accurately predict the chromaticity of an illuminant. The answer to the first question set out for the script then, is that a melanopic signal cannot predict chromaticity (though neither can any other signal).

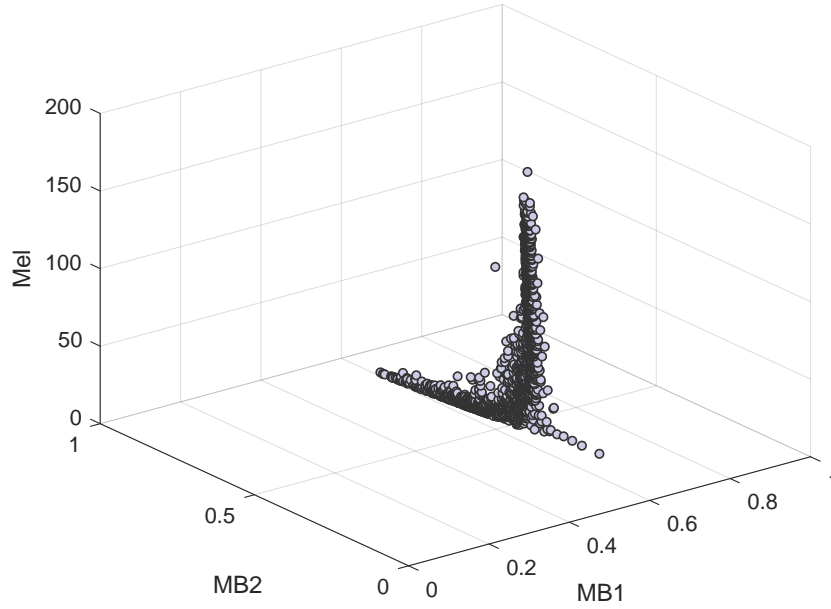


Figure 1: l_{MB} (MB1 above) against s_{MB} (MB2 above) against Mel

It should be noted that there was very high correlation between each signal. This can be considered as corresponding to the high importance of the first principal component of the spectral measurements in this dataset. Put another way, a sensor with

almost any spectral sensitivity could be used to estimate the magnitude of the daylight. I shall return to the discussion of principal components when discussing mel-comp_3.

corr (LMSM')

1.0000	1.0000	0.9993	0.9998
1.0000	1.0000	0.9995	0.9999
0.9993	0.9995	1.0000	0.9999
0.9998	0.9999	0.9999	1.0000

Following this, similar plots were made which plotted l_{MB} against s_{MB} as before, but now plotted the various combinations of [L,M,S,Mel] created by considering one signal over another (e.g. L/M). Following the terminology of [BC14] I shall refer to these signals as secondary signals (where the signals used previously will be referred to as primary signals). The twelve plots created all showed clear and simple relationships between chromaticity and these new derived signals.

For many of these signals however, such relationships are to be expected. For example, a relationship between l_{MB} and L/M should be expected, since l_{MB} is defined as the sum of weighted components of L and M. To confirm this, a null condition was performed, where [L,M,S,Mel] was replaced by randomly generated values, and the secondary signals were generated as before. Relationships between secondary signals derived from L, M or S signals still showed correlations with chromaticity (albeit now points fell on a plane, instead of a line), whereas signals with a Mel component showed only minimal coherence, forming a rough cloud in three-dimensional space.

This example shows that a melanopsin-based signal exhibits correlation with a chromatic signal *not* due to the underlying mathematics of how chromaticity is calculated, and that any relationship must follow from a genuine regularity in the data.

Following this, reflectances were introduced, so as to consider a slightly more realistic situation. Now the simulation considered the colour signals produced by each of the recorded reflectances, under each of the recorded illuminants. MacLeod-Boynton chromaticities, and an analogous melanopic signal, as defined by equation 1, were computed. Chromaticities are plotted in figure 2.

$$mel_{MB} = Mel / (L + M) \quad (1)$$

Now able to consider the second question, I considered whether it was possible to perform a conversion of these computed chromaticities into a hypothetical illuminant-independent space, using mel_{MB} as a corrective signal. It was found through an optimization procedure that through addition and subtraction of weighted mel_{MB} values, using the scaling factors shown in equation 2 such a conversion could approximately be performed. The results of applying equation 2 can be seen in 3.

$$l_{MB}^* = l_{MB} + 0.23mel_{MB} \quad (2a)$$

$$s_{MB}^* = s_{MB} - 1.70mel_{MB} \quad (2b)$$

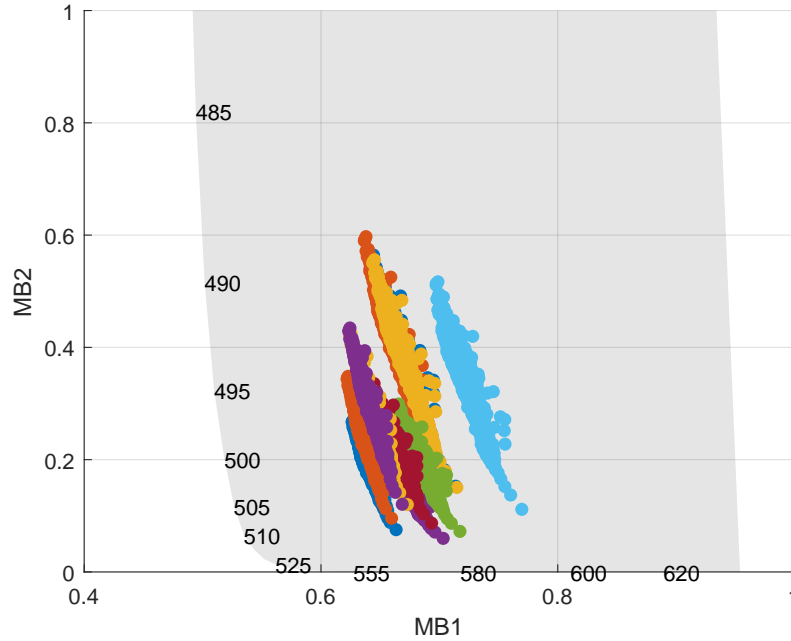


Figure 2: l_{MB} (MB1 above) against s_{MB} (MB2 above). MacLeod-Boynton chromaticities for 12 reflectances under 2600 daylight illuminants.

It can be seen that the points are no longer smeared by the impact of illumination, and instead roughly cluster together by object.

Following this, an additional optimization procedure was performed, to consider whether shifting the sensitivity of the melanopsin function along the wavelength spectrum affected the performance of such a signal in correcting these values. The optimization sought to minimise the overall spread of chromaticities. This had the unexpected and unfortunate effect of simply crushing all the chromaticities down onto a single point, without preserving any inter-object chromatic differences, by choosing spectral sensitivities which best overlapped with the cone fundamentals, thus essentially normalising each signal by a near-copy of itself, and returning near-unity for every situation. This is obviously not ideal.

References

- [BC14] Pablo A. Barrionuevo and Dingcai Cao. Contributions of rhodopsin, cone opsins, and melanopsin to postreceptoral pathways inferred from natural image statistics. *Journal of the Optical Society of America A*, 31(4):A131, April 2014. 00007.

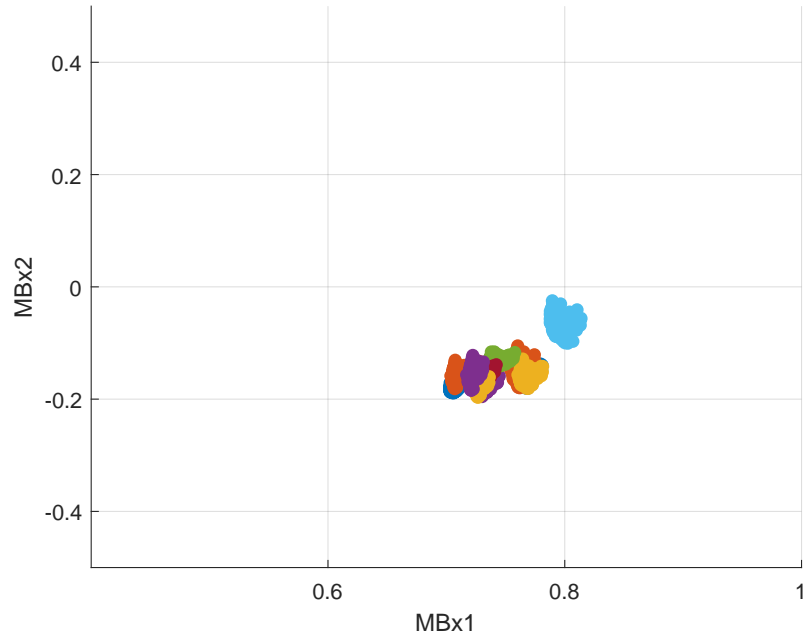


Figure 3: l_{MB}^* (MBx1 above) against s_{MB}^* (MBx2 above). MacLeod-Boynton chromaticities for 12 reflectances under 2600 daylight illuminants, corrected by the corresponding mel_{MB} value.