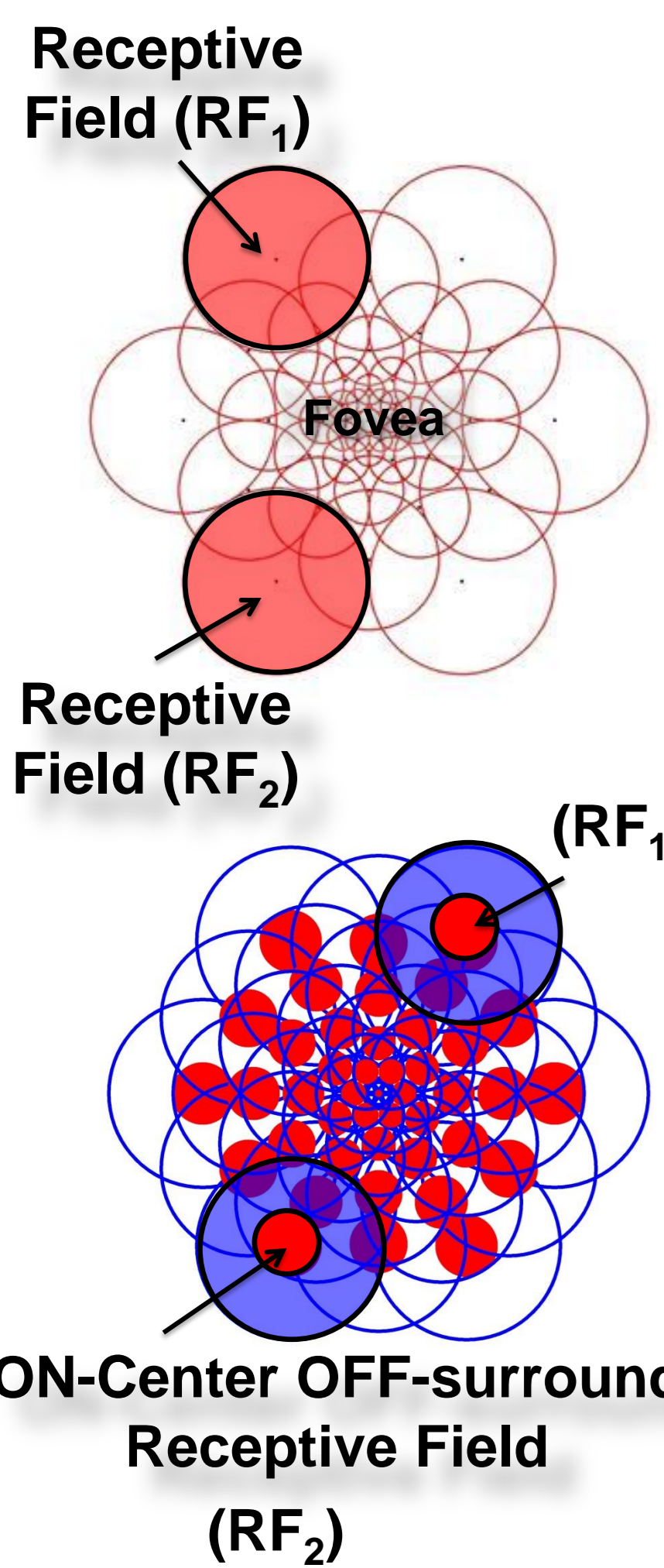


Abstract

- Retina extracts many features from visual stimuli with a high efficiency. Our goal is to build a **bio-inspired descriptor for scene categorization based on low-level processing** performed by the retina.
- We propose a **new image descriptor based on the center-surround organization of the ganglion receptive fields**. Each receptive field is described with a **linear-nonlinear model (LN)** taking into account the inhibitory surrounds.
- FREAK¹ descriptor** is used as baseline, which is enriched sticking to biological data and models of retina. We define a **set of FREAK-variant descriptors** and test them on the problem of scene classification.

Proposed Method

1. Retinal Sampling Pattern



FREAK:

- Circular grid** of concentric distribution of overlapping receptive fields (RFs)
- Higher density** of RFs near the **fovea**
- Considers the difference in intensity between **pairs of receptive fields**:

$$\text{binary descriptor} = \begin{cases} 1 & \text{if } (\text{intensity}(\text{RF}_1) - \text{intensity}(\text{RF}_2)) > 0 \\ 0 & \text{otherwise} \end{cases}$$

Our model:

- Circular center-surround** organization of RFs; the radius of the **surround** is calculated using the retinal eccentricity² ρ :

$$r_S(\rho) = 0.203\rho^{0.472}$$

- ON-center OFF-surround** processing is mimicked using a **difference of Gaussians (DoG)**:

$$K(x, y) = w_C G_{\sigma_C}(x, y) - w_S G_{\sigma_S}(x, y)$$

$$\sigma_S = 3\sigma_C$$

2. Approach

- Retinal activity** is defined by a **linear-nonlinear (LN) model**, where the **activity A** of a cell is defined by:

$$A = f(\text{RF}) \quad \text{where } \text{RF}: \text{Receptive field of the cell}$$

$$f: \text{Nonlinear function}$$

- Each **binary descriptor** is defined as the **difference in activity** between **pairs of RFs**:

$$\text{binary descriptor} = \begin{cases} 1 & \text{if } f(\text{RF}_1) - f(\text{RF}_2) > 0 \\ 0 & \text{otherwise} \end{cases}$$

Descriptor 1: Consider the activity in the center of the RF

$$A = f(I * K_C(x, y)) \quad \text{where } K_C(x, y) = w_C G_{\sigma_C}(x, y)$$

I: Static Image

$$\text{binary descriptor} = \boxed{0101100100\dots 1}$$

Center RF

Descriptor 2: Add the surround response (sign of the DoG)

$$A = \text{sign}(I * K(x, y)) \quad \text{where } K(x, y) = w_C G_{\sigma_C}(x, y) - w_S G_{\sigma_S}(x, y)$$

$$\text{binary descriptor} = \boxed{0101100100\dots 1} \quad \boxed{0101\dots 1}$$

Center RF Sign DoG

Descriptor 3: Enrich the descriptor adding ON and OFF cell responses

$$A_{+1} = f(+ (I * K(x, y))) \quad A_{-1} = f(- (I * K(x, y)))$$

$$\text{binary descriptor} = \boxed{0101100100\dots 1} \quad \boxed{0101\dots 1} \quad \boxed{0101100100\dots 1} \quad \boxed{0101100100\dots 1}$$

Center RF Sign DoG ON cells OFF cells

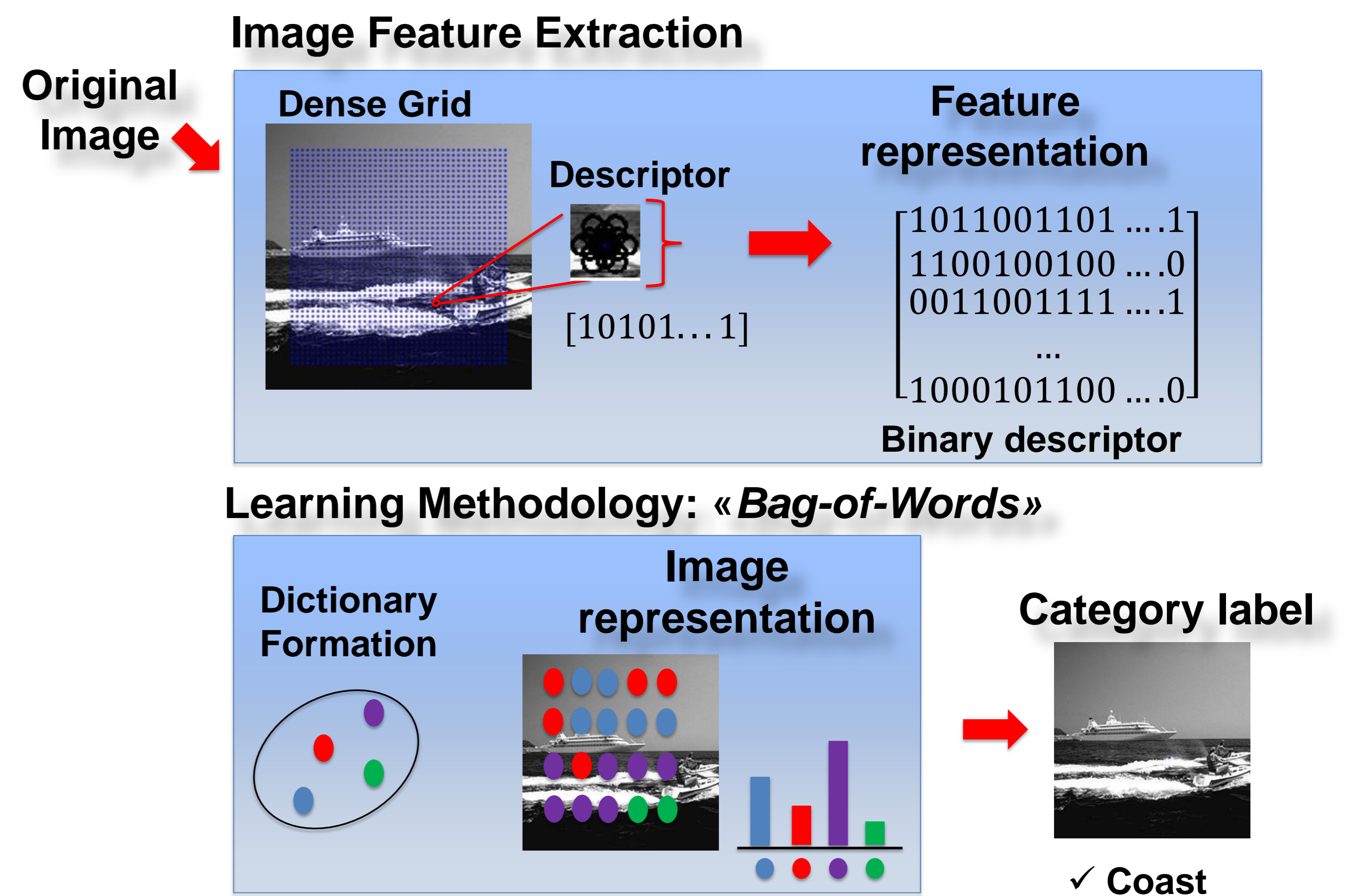
References

- Alahi A., Ortiz R. and Vandergheynst P. **FREAK: Fast retina keypoint**. CVPR 2012, pages 510-517.
- Croner L.J. and Kaplan E. **Receptive fields of P and M ganglion cells across the primate retina**. Vision Research, 35(1):7-24, 1995

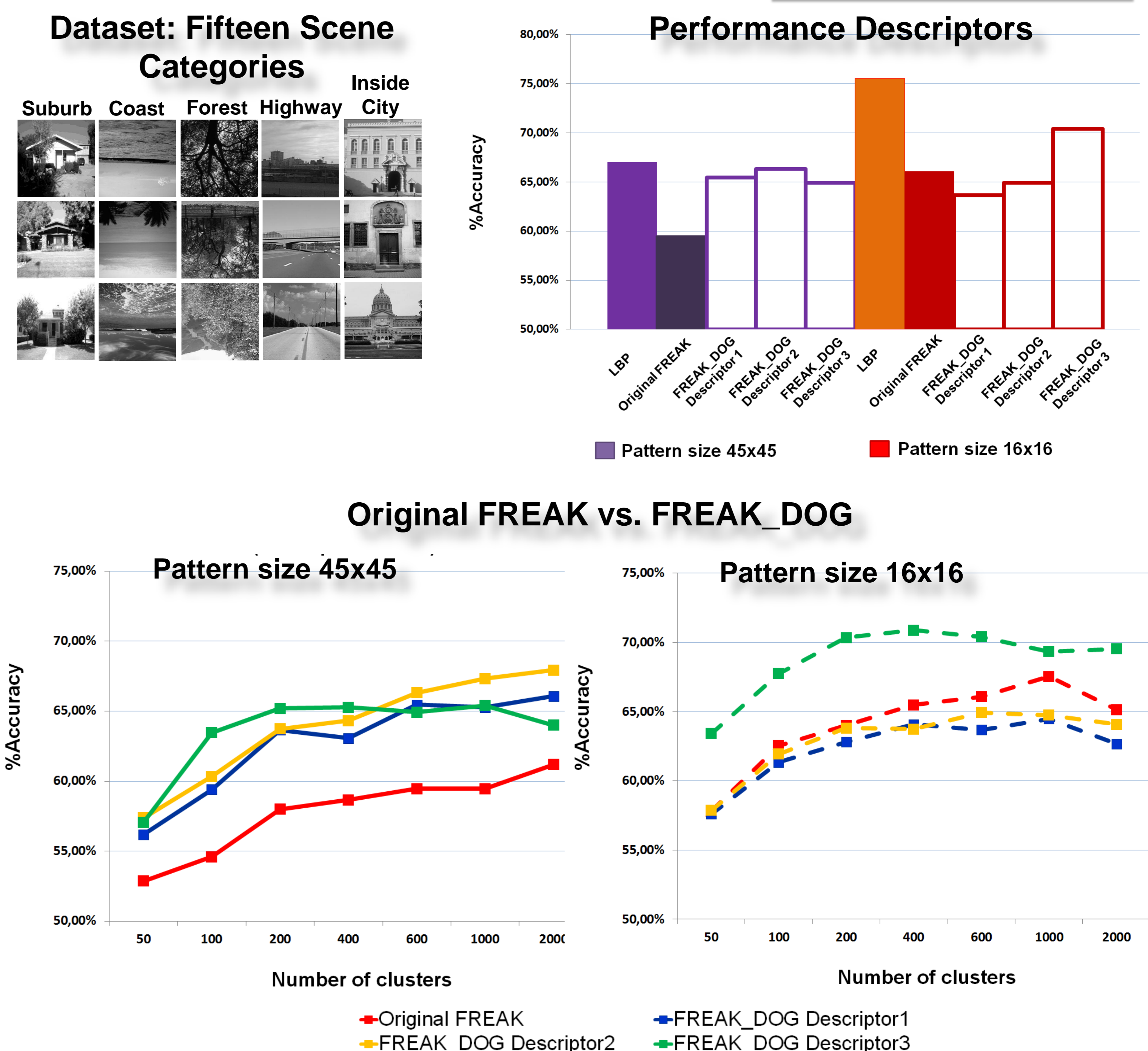
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3. Pipeline



4. Results



- Our preliminary results shows that **enriching FREAK with ON and OFF RFs** improves the performance of the original FREAK.
- In order to evaluate the impact of retinal geometry, our model had been compared to **LBP**. From our data, it seems that the **size of the pattern** makes a difference in the results, more than the organization of the RFs.
- In the future, the model will be enriched considering **other processing performed by the retina**.