

FELIX STÜRMER  
SKETCH-BASED IMAGE RETRIEVAL USING  
CURVELETS



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An Evaluation of Curvlet-Based Cross-Domain Descriptors for Sketch-Based Image  
Retrieval

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## ABSTRACT

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Short summary of the contents...



## ACKNOWLEDGMENTS

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## LISTINGS

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## ACRONYMS

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## INTRODUCTION

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### 1.1 MOTIVATION

Paragraph about increase in visual data, mobile cameras, medicine, etc...

At the core of the research into content-based image retrieval (CBIR) lies the need to be able to access the growing repositories of visual data in a convenient and efficient manner. In this context "convenient" describes the ability for the user to express the query without a complex reformulation of the intent to make it accessible to the query processor. At the same time the computational efficiency becomes more important as the amount of data to search grows. This issue becomes even more critical as the use of mobile, power-limited devices increases across many areas of application, such as autonomous vehicles or handheld augmented reality devices.

Research into text-based information retrieval has brought into existence many statistical methods to query a potentially large body of text using text as the query input. This preserves the close mapping of the intent of the user to the expression of the query and thereby makes the process accessible to users without knowledge about the internal workings of the retrieval system. Providing the means to access a large amount of visual data using a system with similar properties has turned out not to be an easy problem to solve. Using text-based querying for that purpose depends on the ability to reliably label visual data, which would require solving the general object recognition problem first [44]. To avoid that obstacle and to free the retrieval system from the requirement of translating between textual and visual information, many methods to search an image database using visual similarity have been developed.

While the goals of those systems are very similar, they differ considerably in many aspects of the processing pipeline. The query input ranges from example images over drawings to predicate describing color and shape distribution. Similarly, the structure and content of the databases and the means by which the systems query and rank the results vary significantly. This thesis focuses on evaluating a system that uses hand-drawn sketches as inputs to query databases of either full-color images or contour images. The fast discrete curvelet transform [9] is used to analyse image segments.

## 1.2 OUTLINE

[Chapter 2](#) presents the structure of the problem and prior solutions. The following [Chapter 3](#) proposes several variations of a particular solution using the Fast Discrete Curvelet Transform [\[9\]](#). The experimental setup and its results are documented in [Chapter 4](#) and discussed in [Chapter 5](#). In [Chapter 6](#) several possible conclusions are drawn and pointers towards future research are given.

## BACKGROUND & RELATED WORK

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### 2.1 GENERAL CHALLENGES OF COMPUTER VISION

#### 2.1.1 *The Semantic Gap*

One of the core insights of computer vision in general and content based image retrieval in specific probably is that human perception is inseparably linked to interpretation by the brain. As a human individual there is no way to directly access visual information without them having been filtered and weighted by one's personal experiences and cultural context. Therefore, when people talk about visual similarity of images, it usually includes a large degree of semantic similarity unconsciously added to the perception. The difference between that mode of perception and the current algorithmic ways to analyse visual data has been eloquently coined *the semantic gap* by [44]:

The semantic gap is the lack of coincidence between the information that one can extract from the visual data and the interpretation that the same data have for a user in a given situation.

Having had that realisation can guide the decision of a researcher or designer of such systems.

#### 2.1.2 *The Sensory Gap*

In addition to the semantic ambiguity described above, another major obstacle of computer vision impacts a CBIR system: *the sensory gap*. This term has also been coined by [44], where it's defined as follows:

The sensory gap is the gap between the object in the world and the information in a (computational) description derived from a recording of that scene.

That terse definition includes a multitude of conditions, that can affect an image, which a CBIR system operates on:

**ILLUMINATION** The brightness or direction of the illumination can hide or accent edges and texture properties in the scene. Similarly, the color of the illumination influences the recorded color information in the image.

**RESOLUTION** The imaging resolution sets a lower limit on the size of features that can be correctly recognised by any algorithm. As in all signal processing applications, aliasing of high frequency components of the image can introduce further ambiguities. [41]

**OCCCLUSION** Depending on the viewpoint of the recording and the composition of the scene, distinguishing parts of depicted objects may be occluded by other objects or objects may be only partially inside the recorded image.

**PERSPECTIVE** An object's proportions can be distorted by the imaging perspective.

An ideal CBIR system would use feature extraction and comparison methods that can account and correct for such conditions.

## 2.2 ANATOMY OF A CBIR SYSTEM

The inner workings of most CBIR systems can best be examined by looking at the processing pipeline each query has to go through. The coarse sequence of computational steps is almost the same in all such systems (Figure 1):

1. Acquire the image.
2. Extract the signature using a feature extraction algorithm.
3. Compare the signature to a database containing the signatures of the images to search within.
4. Rank the images by similarity using the comparison results.

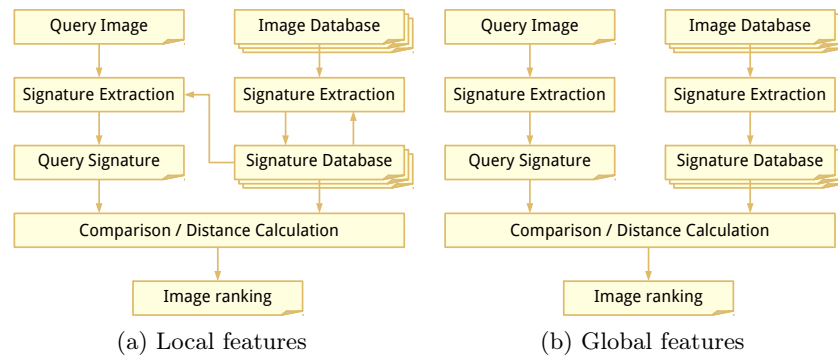


Figure 1: The processing pipeline for CBIR using both local and global features is very similar. The main difference is in the signature extraction step, in which local features are selected, weighted and/or compressed depending on the results of the signature extraction of the other images in the database.



### 2.2.1 Image Acquisition

The format in which the images are available to the system determines the maximum amount of information available to subsequent analysis steps.

A significant part of the preprocessing usually done after acquisition depends on the broadness of the image domain. The concept of the domain encompasses and describes the variability of many possible image parameters like illumination or composition and is therefore closely related to the sensory gap described above. The narrower the image domain is, the more assumptions the system can make about image from that domain. By their very nature, the domain of sketch based image retrieval systems is usually very broad. It contains the sketches create by the user to query the database as well as the images in the database itself, which can be of a completely different nature, e.g. photos or paintings.

Another factor usually is the accepted input format of the feature extraction algorithm. Many algorithms like SIFT [24] or SURF [5] are defined for single-channel data, but some have been specifically developed to operate on multi-channel images, like cSIFT [1] and Yang and Xiao's YIQ-based descriptor [51].

### 2.2.2 Signature Extraction

The signature of an image is its representation in the following comparison step. Therefore it should describe the image using its most discriminatory features compared to all other images in the database. Due to the effects of the *sensory gap* discussed above, there is, at the moment, no definitive way to determine the discriminatory power of features in general, even though knowledge about the image domain can guide the descisions. The signature composition depends on both, the kind of features extracted from the image and the way these features are encoded.

Over the last two decades, a wide variety of feature descriptors have been published, which mostly focus on specific types of features. Some techniques use color histograms [48], while others include spatial relations between colors in a region [45] [13] [23]. Many descriptors attempt to capture texture characteristics in an affine-invariant way [40]. Manjunath et al. [28] use Gabor wavelets for image retrieval, which Rubner and Thomasi [38] combine with the earth mover's distance to bypass segmentation. Another class of descriptors focuses on representing shapes using detection of edges and salient points. Lowe [24] developed the now widely adopted SIFT descriptor, that employs clustering of salient points. More recently, the SURF descriptor [5] uses Haar wavelets to deliver comparable performance. Several publications combine feature types to arrive at a more comprehensive descriptor. Oliva and Torralba [33] capture various scene properties like "roughness" and "openness".



Figure 2: Signature extraction in CBIR systems

#### 2.2.2.1 Global Features

Aside from the nature of the features captured by a descriptor, there are also differences in the geometrical scope the features are derived from. Global feature descriptors attempt to capture the structure of the whole scene or describe the distribution of properties across the image like the binary Haar color descriptor published by Utenpattanant et al. [48]. Some global algorithms subdivide the image into regular segments and derive the localized distribution of features for each division. Lazebnik et al. [21] [22] improves upon this concept by creating feature pyramids using iterative subdivision for multiscale analysis. While the computational complexity of those global approaches is usually quite low, they are especially susceptible to problems like partial occlusion or reflections within the scene. The spatial envelope descriptor [33] combines a global spectrogram with locally derived spectral information to produce an overall image descriptor.

#### 2.2.2.2 Local Features

In contrast to the global approach, many CBIR systems employ "bag-of-features" descriptors, that represent the image as an unsorted collection of local features extracted from small patches of the image. The unsorted nature of the feature collection leads to loss of large-scale geometric structures that can be counteracted by a suitable choice of the patch sizes. When the local feature descriptors are invariant to rotation, scale or similar deformations, the sensitivity to viewpoint variations or occlusions decreases. The prominent SIFT descriptor [24] achieves this by selecting the feature locations such that they can be normalized with respect to scale, orientation and limited 3D projections. The SURF descriptor by Bay et al [5] gives similar results, but has reduced computational requirements. The HOG descriptor [12] calculates histograms

of gradient directions on regular grid cells to describe the local angular distribution of edges.

### 2.2.2.3 Dimensionality Reduction

The signatures produced by local descriptors are often large sets of vector, that are themselves of considerable size. For example, the SIFT descriptor describes each image using about 1000 local feature vectors of 160 values each. Such large numbers of vectors are expensive to store and compare, so one of several data reduction methods is commonly used.

**PRINCIPAL COMPONENT ANALYSIS** The Principle Component Analysis (PCA) is a transformation, that computes the orthogonal basis best suited to describe the variance of the data. An  $n$ -dimensional data set is linearly mapped to a coordinate system, in which the direction of the first axis  $\mathbf{a}_1$  is the direction with the largest variance in the data. The following axes'  $\mathbf{a}_i$ ,  $i \in 2, \dots, n$  directions correspond to the orthogonal directions with the next-largest variances in descending order. By choosing the  $p$  largest component vectors and performing an inverse transformation of the PCA-transformed data, a projection of the original data in  $p$  dimensions can be obtained. Due to the choice of the vectors for the inverse transformation, the projection discards only the parts of each observation that vary the least between all observations.

PCA has been applied to the face recognition problem using intensity images (eigenfaces) [47], wavelets (waveletfaces) [15] and more recently curvelets (curveletfaces) [27]. Ke et al. [20] used it to improve the robustness of the SIFT [24] descriptor. To overcome the limited between-class discrimination of PCA, it has been combined with Linear Discriminant Analysis (LDA), yielding even better results [26].

**VISUAL WORDS AND CLUSTERING** Instead of reducing the size of the individual feature vectors, the bag-of-features approach collects the feature vectors into a single signature vector to represent the image. This is done by determining a codebook of representative feature vectors, the "visual words" [43], and assigning each local feature vector to the most similar visual word. A histogram of the distribution of visual words can then be calculated as a signature for each image.

To create the codebook, the large number of feature vectors extracted from local patches of each image in the database are grouped into clusters of similar vectors. The optimal number of clusters is usually determined experimentally and varies with other processing parameters such as the sampling strategy [32] [52].

The most common clustering method used in numerous publications [54] [43] [11] [16] [49] is k-means clustering. This algorithm uses the euclidean distance as a metric to assign each observation  $\mathbf{x}_p$  to the

nearest cluster  $S_i$  with mean  $m_i$ ,  $i \in 1, \dots, k$ . The goal is to minimize the variance within each cluster:

$$\sum_{i=1}^k \sum_{x_p \in S_i} \|x_p - m_i\|^2$$

Lloyd's algorithm is the usual way to calculate a k-means partition. It requires a set of  $k$  initial cluster centers, that are often randomly chosen from the dataset or randomly generated. The cluster centers  $m_i$  are then iteratively adjusted until no reassignment takes place in two consecutive iterations  $t$  and  $t + 1$ . In each iteration, each observation  $x_i$  is assigned to exactly one cluster  $S_i$  using

$$S_{i,t} = \{x_p : \|x_p - m_{i,t}\| \leq \|x_p - m_{j,t}\| \quad \forall j \in 1, \dots, k\}.$$

The centers are then recalculated as

$$m_{i,t+1} = |S_{i,t}|^{-1} \sum_{x_p \in S_{i,t}} x_p.$$

The greedy nature of the algorithm and the random initialization mean that it is merely a heuristic and can converge on a local minimum, that is not a global minimum. Other clustering methods are hierarchical agglomerative and divisive algorithms as used by Nister and Stewenius [31] and Philbin et al. [36], that recursively divide or merge partitions to optimize the vocabulary.

Once the clustering algorithm has converged, the cluster centers  $m_i$  will be used as the visual words of the codebook. To create the signature vector of an image, its feature vectors  $x_p$  will be quantized by grouping them into sets  $S_i$ ,  $i \in 1, \dots, k$  such that

$$S_i = \{x_p : \|x_p - m_i\| \leq \|x_p - m_j\| \quad \forall j \in 1, \dots, k\}.$$

The final image signature  $\tilde{I}$  then is the vector of the cardinality of these sets:

$$\tilde{I} = (|S_1|, |S_2|, \dots, |S_{k-1}|, |S_k|)$$

### 2.2.3 Comparison and Ranking

#### 2.2.3.1 Weighting

In the vocabulary of visual words created via clustering, a set of representatives for a group of features has been defined. How representative the presence of visual word in the signature of a specific image has not yet been taken into account though. To remedy this, weighting schemes are often used that modify the signatures to enhance distinguishing characteristics and make the ranking more accurate.

**LINEAR SUPPORT VECTOR MACHINES** The concept of support vector machines (SVMs) is usually employed when a feature needs to be classified into one or more classes. While such classification problem occur frequently in computer vision in the fields of object detection [37] [11], human detection [12] or scene classification [52], the class concept is too limited for general image retrieval. To achieve the classification, a SVM constructs a hyperplane, that optimally separates two sets of points  $\mathbf{x}_i \in \mathbb{R}^n$  with labels  $y_i \in \{-1, 1\}$  in a training dataset  $\{(\mathbf{x}_i, y_i)\}$ . The hyperplane can be defined using its normal vector  $\mathbf{w}$  and offset  $\mathbf{b}$  as

$$\mathbf{w} \cdot \mathbf{x} - \mathbf{b} = 0.$$

To turn the problem into an optimization problem, the plane is split up into two parallel hyperplanes described by

$$\mathbf{w} \cdot \mathbf{x} - \mathbf{b} = 1 \text{ and } \mathbf{w} \cdot \mathbf{x} - \mathbf{b} = -1.$$

The region between these two planes is characterized by their distance  $\frac{2}{\|\mathbf{w}\|}$ , which needs to be minimized while satisfying

$$\begin{aligned} \mathbf{w} \cdot \mathbf{x}_i - \mathbf{b} &\geq 1 & \text{if } y_i = 1 & \quad \text{or} \\ \mathbf{w} \cdot \mathbf{x}_i - \mathbf{b} &\leq -1 & \text{if } y_i = -1 \end{aligned}$$

for all  $i$ . It is possible to transform this into an equivalent quadratic optimization problem solvable by standard quadratic programming algorithms.

Even though image retrieval as described in this thesis is not a simple classification problem, linear SVMs can still be of use. Guyon et al. [19] showed, that the components of  $\mathbf{w}$  can be used as weights describing the discriminative power of feature vectors. This was applied to image retrieval by Shrivastava et al. [42]. There the authors trained a SVM for each query image  $I_q$  with the query image's signature constituting one class and the database images' signatures  $\mathbf{x}_i$  making up the other class. The pairwise similarity could then easily be obtained from the learned weights  $\mathbf{w}_q$  as

$$S(I_q, I_i) = \mathbf{w}_q^T \mathbf{x}_i.$$

That way, common features in the query signature are effectively downvoted, while unique features are assigned larger weights.

**TF-IDF** Along with the visual word analogy from the field of text retrieval, the statistical method of TF-IDF weighting [4] has been applied to CBIR [43]. It is a technique to weight features (terms) in a document in relation to their occurrence in the document and the whole database. The term frequency  $\text{tf}_{i,j}$  is the number of normalized occurrences of term  $t_i$  in document  $d_j$ . The normalization can be performed in different ways. The most common ones are dividing by the total number of words

$n_j$  in the document or the maximum term count  $\max\{tc_{i,j} : \forall t_i \in d_j\}$  of any term in the document  $j$ :

$$tf_{i,j} = \frac{tc_{i,j}}{n_j} \text{ or } tf_{i,j} = \frac{tc_{i,j}}{\max\{tc_{i,j} : \forall t_i \in d_j\}}$$

The occurrence of a term in the database  $D$  is measured as the logarithm of the quotient of the overall number of documents  $|D|$  and the number of documents  $m_i$  containing the term  $t_i$ :

$$idf_i = \log \frac{|D|}{m_i}$$

Therefore, the total weight  $w_{i,j}$  of a term is given as

$$w_{i,j} = tf_{i,j} \cdot idf_i = \frac{tc_{i,j}}{n_j} \cdot \log \frac{|D|}{m_i}$$

assuming the document length is used for normalization.

### 2.2.3.2 Distance Metrics

To retrieve the similar images from the database, a ranking of database image in respect to their similarity to the query image must be obtained. Since each image is represented by a signature vector, this means the pairwise distance between the query image's signature and each database image's signature can be used to sort the list of query results. This section will describe several distance metrics and similarity measure candidates, which can be used to that end.

**EUCLIDEAN DISTANCE** The simplest and probably most widely used distance metric is the euclidean distance. Its two-dimensional variant is derived from the formula of Pythagoras. Generalized to  $n$ -dimensional points  $p = (p_1, p_2, \dots, p_n)$  and  $q = (q_1, q_2, \dots, q_n)$ , it can be written as

$$d_{EUC}(p, q) = \sqrt{\sum_{i=1}^n (q_i - p_i)^2}.$$

**MAHALANOBIS DISTANCE** If the distance metric is used for clustering or classification of datasets with non-spherical within-class distributions, the euclidean distance will naturally not perform well. A common alternative is the Mahalanobis distance, that incorporates the correlation of the dataset into the result. The distance of a point  $p = (p_1, p_2, \dots, p_n)$  to a cluster with mean  $m = (m_1, m_2, \dots, m_n)$  is

$$d_{MAHA}(p, m) = \sqrt{(p - m)^T S^{-1} (p - m)},$$

where  $S$  is the cluster's covariance matrix. In practice, it has been used in [30] and [43] to compare feature vectors of the SIFT [24] descriptor.

**COSINE DISTANCE** A metric sometimes used in information retrieval applications is the cosine distance or cosine similarity. The similarity is defined as the cosine of the angle between two vectors  $\mathbf{p} = (p_1, p_2, \dots, p_n)$  and  $\mathbf{q} = (q_1, q_2, \dots, q_n)$  when interpreted geometrically:

$$\cos(\theta_{\mathbf{p}, \mathbf{q}}) = \frac{\mathbf{p} \cdot \mathbf{q}}{\|\mathbf{p}\| \|\mathbf{q}\|}$$

This means that the metric effectively normalizes the vectors in respect to their euclidean length. That effect is often used in text retrieval to achieve invariance regarding the document size.

**EARTH MOVER'S DISTANCE** The Earth Mover's Distance (EMD) is an application of the discrete transportation problem, which was first introduced into the field of computer vision by Peleg et al. [35]. The use of the algorithm as a signature comparison metric in image retrieval was published by Rubner et al. [39]. At the abstract level, the distance is calculated as the amount of work necessary to transform one signature into the other. A main advantage of the EMD over most other distance measures is that it accounts for inter-bin distances (ground distances) in binned distributions. Another useful property is the ability to compare distributions of different sizes, e.g. for partial matching. For signatures  $P = \{(p_1, w_{p_1}), \dots, (p_n, w_{p_n})\}$  and  $Q = \{(q_1, w_{q_1}), \dots, (q_m, w_{q_m})\}$  of bin centers  $p_i$  and  $q_i$  with bin sizes  $w_{p_i}$  and  $w_{q_i}$  the pairwise ground distances can be represented in a  $n \times m$  matrix  $D$ . These ground distances  $d_{i,j}$  between two bins is then interpreted as the costs of moving a unit of goods from one bin to the other. The optimal solution minimizes the overall costs by finding flow values  $f_{i,j}$  between each pair  $p_i$  and  $q_i$ , that satisfy the constraints

$$\begin{aligned} f_{i,j} &\geq 0 \\ \sum_{j=1}^m f_{i,j} &\leq w_{p_i} \\ \sum_{i=1}^n f_{i,j} &\leq w_{q_j} \\ \sum_{i=1}^n \sum_{j=1}^m f_{i,j} &= \min \left( \sum_{i=1}^n w_{p_i}, \sum_{j=1}^m w_{q_j} \right) \end{aligned}$$

for all  $1 \leq i \leq n$  and  $1 \leq j \leq m$ . The distance between signatures  $P$  and  $Q$  can then be calculated as

$$d_{\text{EMD}}(P, Q) = \frac{\sum_{i=1}^n \sum_{j=1}^m d_{i,j} f_{i,j}}{\sum_{i=1}^n \sum_{j=1}^m f_{i,j}}.$$

Rubner et al. [39] propose using a simplex-based algorithm to solve the transportation problem, that achieves good performance by exploiting the specific problem structure. Zhang et al. [53] use Rubner's implementation to train a support vector machine for image classification with Gaussian kernels scaled by the EMD.

**HISTOGRAM INTERSECTION** A very inexpensive way to compare two distributions has been shown by Swain and Ballard [46], namely the histogram intersection technique. While they used it to compare color histograms, it should work equally well for binned distributions generated using vector quantization. The normalized similarity measure between two histograms  $P = (p_1, \dots, p_n)$  and  $Q = (q_1, \dots, q_n)$  is

$$s_{HI}(P, Q) = \frac{\sum_{i=1}^n \min(p_i, q_i)}{\sum_{i=1}^n q_i}$$

A variation of the above definition treats partial matches as perfect matches by adjusting the denominator:

$$s_{HI}(P, Q) = \frac{\sum_{i=1}^n \min(p_i, q_i)}{\min\left(\sum_{i=1}^n p_i, \sum_{i=1}^n q_i\right)}$$

## 2.3 IMAGE TRANSFORMATIONS FOR FEATURE EXTRACTION

### 2.3.1 The Gabor Filter

In content based image retrieval, the inability of the Fourier transform to localize the frequency components in time disqualifies it as a suitable analysis method. Instead, wavelets are often used to obtain a time-frequency representation of the signal. A particularly successful application to CBIR was the use of gabor wavelets as proposed by Manjunath and Ma [28]. The mother Gabor wavelet  $g(x, y)$  with the sinus frequency  $W$  and gaussian scaling parameters  $\sigma_x, \sigma_y$  is given as

$$g(x, y) = \left(\frac{1}{2\pi\sigma_x\sigma_y}\right) \exp\left(-\frac{1}{2}\left(\frac{x^2}{\sigma_x^2} + \frac{y^2}{\sigma_y^2}\right) + i \cdot 2\pi Wx\right)$$

and thus its Fourier transform as

$$\hat{g}(u, v) = \exp\left(-\frac{1}{2}\left(\frac{(u-W)^2}{\sigma_u^2} + \frac{v^2}{\sigma_v^2}\right)\right), \sigma_u = \frac{1}{2}\pi\sigma_x \text{ and } \sigma_v = \frac{1}{2}\pi\sigma_y$$

The individual wavelets  $g_{m,n}(x, y)$  are generated from  $g(x, y)$  by scaling and rotation by  $\theta = \frac{n\pi}{K}$  for all  $0 < n < K$  orientations:

$$g_{m,n}(x, y) = a^{-m} g(a^{-m}(x \cos \theta + y \sin \theta), a^{-m}(-x \sin \theta + y \cos \theta)).$$



To avoid redundancy due to overlap, the scaling parameters  $\sigma_u$  and  $\sigma_v$  are chosen in a way that the frequency spectra can be tiled as in figure 3. A normalization to zero mean can be added to remove the influence of the input's intensity value scale.

The individual coefficients can be obtained by convolving each filter of the filter bank with the signal.

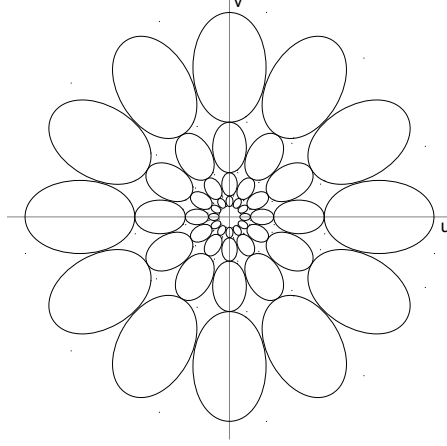


Figure 3: Tiling of Gabor wavelets. Note that due to the elliptical shape of the wavelets, there are some parts of the spectrum left uncovered.

### 2.3.2 The Continuous Curvelet Transform

The formulation of the continuous curvelet transform (CCT) by Candes and Donoho in [7] was based on Candes' previous definition and expansion of the ridgelet transform [6]. In that publication they looked at the state of research into efficient representations of edge discontinuities. They based their research on two realisations:

1. A nonadaptive approach of signal approximation can compete with many of the adaptive schemes prevalent in previous research. At the same time the non-adaptivity comes with a greatly reduced computational overhead and reduced requirements for a priori knowledge. Obtaining that knowledge in the presence of blurred or noisy data can sometimes be unfeasible.
2. Wavelet transforms can represent point singularities in a signal of up to two dimensions in a near-ideal manner, but fail to perform equally well on edges: Given a two-dimensional object in signal  $f$ , that is smooth except for discontinuities along a curve, a wavelet approximation  $\tilde{f}_m^W$  from the  $m$  largest coefficients exhibits an error of

$$\|f - \tilde{f}_m^W\|^2 \propto m^{-1}, \text{ for } m \rightarrow \infty$$

since up to  $O(2^j)$  localized wavelets are needed to represent the signal along the edge. That falls short of what an approximation  $\tilde{f}_m^T$  using a series of  $m$  adapted triangles could achieve:

$$\|f - \tilde{f}_m^T\|^2 \propto m^{-2}, \text{ for } m \rightarrow \infty$$

They showed that a similarly precise approximation can be achieved by combining Candes' ridgelet analysis [6] with smart windowing functions and bandpass filters. The steps of the transformation were as follows:

1. Decomposition of the signal into subbands of scale-dependent size
2. Partitioning of each subband into squares
3. Normalisation of each square to unit scale
4. Analysis of each square in an orthonormal ridgelet system

The result was a formulation of a decomposition that matched the parabolic scaling law  $\text{width} \propto \text{length}^2$  often observed in curves.

The above formulation became known as the curvelet gg transform when Candes and Donoho revised it soon after in [8]. The new version is not dependent on ridgelets and aims to remove some shortcomings of the curvelet gg transform, namely a simpler mathematical analysis, fewer parameters and improved efficiency regarding digital implementations, which will be described later.

The curvelet transform in  $\mathbb{R}^2$  works by localising the curvelet waveforms in the time domain. The "mother" curvelet waveform  $\varphi_j(x)$  is defined using two frequency domain windows  $W(r)$ , the "radial window" (Figure 4a), and  $V(t)$ , the "angular window" (Figure 4b). A combined frequency window  $U_j$  (Figure 4c) can then be defined as

$$U_j(r, \theta) = 2^{-\frac{3j}{4}} W(2^{-j}r) V\left(\frac{2^{\lfloor \frac{j}{2} \rfloor} \theta}{2\pi}\right).$$

The waveform  $\varphi_j$  can then be expressed as being the inverse Fourier transform of  $\hat{\varphi}_j = U_j$  and all curvelets of a scale  $2^{-j}$  can be derived by

ROTATING  $\varphi_j$  by a sequence of equispaced rotation angles  $\theta_l = 2\pi \cdot 2^{-\lfloor \frac{j}{2} \rfloor} \cdot l$  with  $l = 0, 1, \dots$  such that  $0 < \theta_l < 2\pi$  and

TRANSLATING  $\varphi_j$  by a sequence of offsets  $k = (k_1, k_2) \in \mathbb{Z}^2$ :

$$\varphi_{j,k,l}(x) = \varphi_j(R_{\theta_l}(x - x_k^{(j,l)})), \quad (1)$$

where  $x = (x_1, x_2)$ ,  $R_\theta$  the rotation matrix for angle  $\theta$  and  $x_k^{(j,l)} = R_{\theta_l}^{-1}(k_1 \cdot 2^{-j}, k_2 \cdot 2^{-\frac{j}{2}})$ .

Each curvelet coefficient  $c(j, l, k)$  can then be calculated as the inner product of  $f \in L^2(\mathbb{R}^2)$  and curvelet  $\varphi_{j,l,k}$ :

$$c(j, l, k) := \langle f, \varphi_{j,l,k} \rangle = \int_{\mathbb{R}^2} f(x) \overline{\varphi_{j,l,k}(x)} dx \quad (2)$$

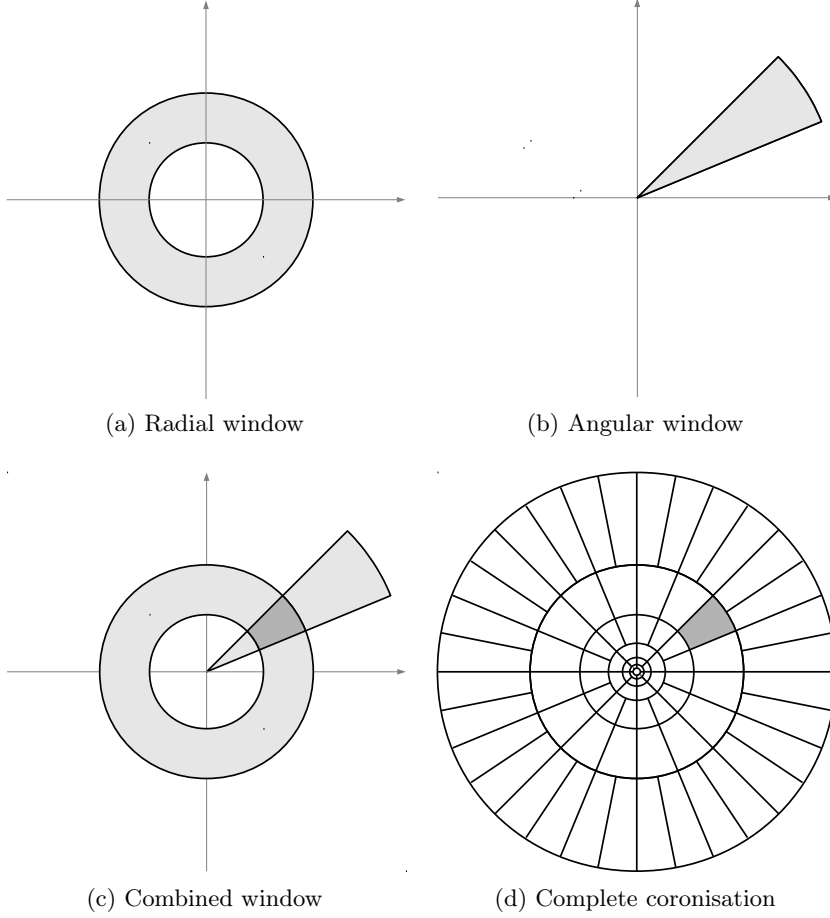


Figure 4: The window  $W(2^{-j}r)$  at scale  $2^j$  (a) is combined with the window  $V(t)$  (b) to form a support wedge for the curvelet (c). The wedge roughly obeys a  $\text{width} \propto \text{length}^2$  relation. (d) shows the wedge within a schema of the complete tiling in frequency domain.

As visible in figure 4d curvelets also have non-directional components at the coarsest scale, similar to those found in the wavelet transform. Those curvelets will be defined using a special low-pass filter window  $W_0$ , which is characterized as being the remainder of the tiling not covered by the previously described radial windows:

$$|W_0(r)|^2 + \sum_{j \geq 0} |W(2^{-j}r)|^2 = 1$$

Using the window defining the coarse scale curvelet  $\varphi_{j_0,k}$  via its Fourier transform is straightforward:

$$\varphi_{j_0,k}(x) = \varphi_{j_0}(x - 2^{-j_0}k), \quad \hat{\varphi}_{j_0}(\omega) = 2^{-j_0}W_0(2^{-j_0}|\omega|), \quad (3)$$

where  $k = (k_1, k_2) \in \mathbb{Z}^2$ .

Note that, in contrast to the Gabor wavelets (Figure 3), there is no gap in between the curvelets, so no information is lost.

### 2.3.3 The Fast Discrete Curvelet Transform

Based on the above definition of the continuous curvelet transform, a team around the authors of the original curvelet publication presented two digital, discrete implementations of the transform: the fast discrete curvelet transform (FDCT) [9]. The implementations have been described in 2D and 3D, but since this paper deals exclusively with 2D images, the explanation below will also be restricted to two dimensions.

The digital versions of the transforms operate on arrays  $f[t_1, t_2]$  with  $0 \leq t_1, t_2 < n$  to produce coefficients  $c^D(j, l, k)$  in a way consistent with the continuous version (Equation 2):

$$c^D(j, l, k) := \sum_{0 \leq t_1, t_2 < n} f[t_1, t_2] \overline{\varphi_{j,l,k}^D[t_1, t_2]}. \quad (4)$$

Since the windows used in the continuous form are based on rotations and dyadic coronae, they are not well suited for use with cartesian arrays. The discrete formulation substitutes them with appropriate concepts. Instead of concentric annuli, the window function  $W_j^D$  generates concentric squares "rings" using the square windows  $\Phi_j(\omega_1, \omega_2) = \phi(2^{-j}\omega_1)\phi(2^{-j}\omega_2)$ , with  $\phi$  being a low-pass 1D window:

$$W_j^D(\omega) = \sqrt{\Phi_{j+1}^2(\omega) - \Phi_j^2(\omega)}, \quad j \geq 0.$$

The rotation matrix  $R_\theta$  is replaced by the shear matrix  $S_\theta$  to create the combined window function

$$U_{j,l}^D := W_j^D(\omega) V_j(S_{\theta_l} \omega).$$

The sequence  $\theta_l$  is defined as a sequence of equispaced slopes  $\tan(\theta_l) := l \cdot 2^{-\lfloor \frac{j}{2} \rfloor}$  with  $l = -2^{\lfloor \frac{j}{2} \rfloor}, \dots, 2^{\lfloor \frac{j}{2} \rfloor} - 1$ . The construction of the corner windows are Figure 5 shows a tiling of all  $U_{j,l}^D$ .

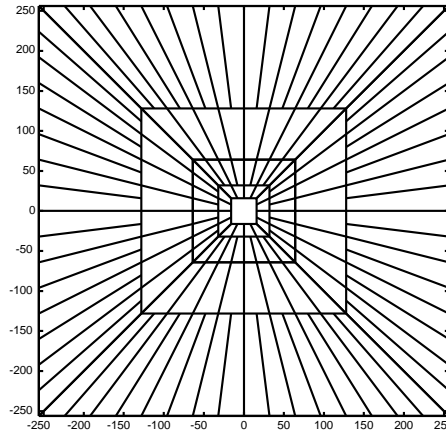


Figure 5: Discrete frequency tiling using concentric squares (from Candes et al. [9])

### 2.3.3.1 FDCT using unequispaced FFTs

The first implementation of the discrete curvelet transform transfers the input array  $f[t_1, t_2]$ ,  $0 \leq t_1, t_2 < n$  into the Fourier domain to obtain  $\hat{f}[n_1, n_2]$ :

$$\hat{f}[n_1, n_2] = \sum_{t_1, t_2=0}^{n-1} f[t_1, t_2] e^{-\frac{i2\pi(n_1 t_1 + n_2 t_2)}{n}}, \quad -\frac{n}{2} \leq n_1, n_2 < \frac{n}{2}$$

The obtained Fourier samples need to be interpolated for each pair of scale  $j$  and angle  $l$  to match the grid of the sheared support window  $U_j^D[n_1, n_2]$ . The authors achieve this by resampling  $\hat{f}$  on the grid implied by the sheared window for each angle via a series of 1D fast Fourier transforms. These transforms represent a polynomial interpolation of each "column" of the parallelogram  $P_j$  containing the sheared window (Figure 6a), that can be computed with a  $O(n^2 \log n)$  complexity in a sufficiently exact approximation.

This yields an object  $\hat{f}[n_1, n_2 - n_1 \tan \theta_l]$  for  $(n_1, n_2) \in P_j$ , that can be multiplied with the window  $U_j^D$  described above in order to create a localized "wedge" with the orientation  $\theta_l$ :

$$f_{j,l}^D[n_1, n_2] = \hat{f}[n_1, n_2 - n_1 \tan \theta_l] U_j^D[n_1, n_2]$$

The discrete curvelet coefficients  $c^D(j, l, k)$  can then be calculated by applying the inverse 2d Fourier transform:

$$c^D(j, k, l) = \sum_{n_1, n_2 \in P_j} \hat{f}[n_1, n_2 - n_1 \tan \theta_l] U_j^D[n_1, n_2] e^{i2\pi(\frac{k_1 n_1}{L_{1,j}} + \frac{k_2 n_2}{L_{2,j}})},$$

in which  $L_{1,j}$  and  $L_{2,j}$  are the length and width of the rectangle supporting  $U_j^D$ .

### 2.3.3.2 FDCT using wrapping

As before, FDCT using wrapping first calculates  $\hat{f}[n_1, n_2]$  with  $-\frac{n}{2} \leq n_1, n_2 < \frac{n}{2}$  as the Fourier transform of the input  $f$ . The sample is then localized by multiplying it with a window  $U_{j,l}^D$  for each angle  $j$  and scale  $l$ :

$$d_{j,l}[n_1, n_2] = U_{j,l}^D \hat{f}[n_1, n_2]$$

To avoid the computationally costly interpolation step required in the USFFT approach, this method keeps the rectangular grid of the input signal. Because an axis-aligned bounding box of the window  $U_j^D$  in Fourier domain cannot maintain the  $\text{width} \propto \text{length}^2$  proportions of the window, applying an inverse Fourier transform on such a bounding box in general would lead to significant oversampling of the coefficients and thereby increase the memory requirements for fine scale curvelets beyond that of the the USFFT approach. In order to circumvent that,

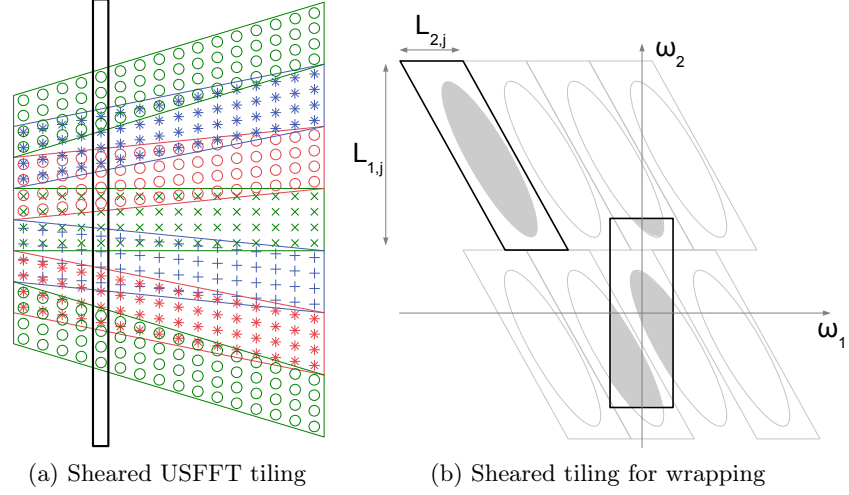


Figure 6: (a) illustrates the respective grid for each parallelogram containing the sheared support windows of the "east quadrant". The box highlights one of the columns, that represent one of the 1D polynomial interpolation problems solved for resampling. In (b) the parallelogram  $P_{j,l}$  is shown on top of the tilted tiling of a curvelet in frequency domain. Due to the periodicity the rectangle in the center contains the same curvelet, but has a much smaller axis aligned bounding box for the FFT to operate on. Both images have been adapted from Candes et al. [9].

the authors utilize the periodic nature of the Fourier transform and propose generating a periodically wrapped version of the fourier samples. For  $P_{j,l}$  as the bounding parallelogram of  $U_{j,l}^D$ ,  $L_{1,j}$  and  $L_{2,j}$  are the period lengths by which to translate  $P_{j,l}$  in the horizontal and vertical direction to produce a suitable tiling for each orientation  $\theta_l$  (Figure 6). Thus, the wrapped, localized data are

$$f_{j,l}^D[n_1, n_2] = Wd_{j,l}[n_1, n_2] = \sum_{m_1 \in \mathbb{Z}} \sum_{m_2 \in \mathbb{Z}} d_{j,l}[n_1 + m_1 L_{1,j}, n_2 + m_2 L_{2,j}]$$

with  $0 \leq n_1 < L_{1,j}$  and  $0 \leq n_2 < L_{2,j}$ , which gives a rectangle of size  $L_{1,j}$  times  $L_{2,j}$ .

Again, the discrete curvelet coefficients  $c^D(j, l, k)$  can then be collected using the inverse 2d Fourier transform:

$$c^D(j, k, l) = \sum_{n_1, n_2 \in P_j} f_{j,l}^D[n_1, n_2] e^{i2\pi(\frac{k_1 n_1}{L_{1,j}} + \frac{k_2 n_2}{L_{2,j}})}.$$

#### 2.3.4 *gPb Contour Detection*

In [25], Maire et al. describe an improvement of the contour detector published in [29] that includes global information in addition to local cues. The orientation-specific local parameters  $G_i$  extracted from a circular neighborhood around the location  $(x, y)$  are brightness, color and

texture gradients on three scales. They are summarized as a coefficient  $\mathbf{mPb}(x, y, \theta)$  using a weighted sum with weights  $\alpha_i$ :

$$\mathbf{mPb}(x, y, \theta) = \sum_{i=1}^9 \alpha_i G_i(x, y, \theta)$$

The global component  $\mathbf{sPb}(x, y, \theta)$  is the result of applying a filter-bank of directional gaussian derivatives to a set of  $k$  generalized eigenvectors  $v_j$ ,  $j \in 1, \dots, k$ . The linear system these eigenvectors are obtained from has an affinity matrix derived from the intervening contour cue [17]. The linear combination of the individual directional derivatives then represents the large-scale contours in the image:

$$\mathbf{sPb}(x, y, \theta) = \sum_{j=1}^k \frac{1}{\sqrt{\lambda_j}} \mathbf{sPb}_{v_j}(x, y, \theta)$$

A further linear combination of the local component  $\mathbf{mPb}$  and the global component  $\mathbf{sPb}$  with learned weights  $\alpha_i$  and  $\gamma$  provides a detailed map of contours in the image while limiting the amount of clutter compared to a purely local contour detector:

$$\mathbf{gPb}(x, y, \theta) = \sum_{i=1}^9 \alpha_i G_i(x, y, \theta) + \gamma \cdot \mathbf{sPb}(x, y, \theta)$$

From these directional contour maps, Arbeláez et al. derived a hierarchical contour detector [3], that conditionally joins adjacent regions to obtain closed-contour maps of high quality.





## PROPOSED SOLUTION

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The image processing pipeline described in this thesis aims to be suitable for content based image retrieval using hand-drawn sketches for querying. The main interest was to evaluate how well the fast discrete curvelet transform (FDCT) [9] is able to represent the lines in hand-drawn sketches as well as salient edges in photos or paintings. To explore the effects of preprocessing and signature extraction, several variations of the pipeline have been implemented. The used preprocessing steps include applying the sobel operator, extracting a canny edge map or determining segment borders using the gPb algorithm published in [2]. Signatures are constructed using both, global curvelet features, and a bag-of-features approach similar to what was described in [43] and [14].

The following sections will describe the variations of the processing stages *image acquisition*, *signature extraction*, and *ranking*. To reference the individual variations unambiguously, labels like LUMA will be introduced for each component.

### 3.1 IMAGE ACQUISITION

Since one premise of the system is that hand-drawn sketches are compared with a large body of images from various sources, a division into two input domains seems obvious. The first domain, the domain of query sketches, is quite narrow, because we can characterize its members as binary images with large, smooth areas separated by discontinuities along curves. The database images, that make up the second domain, are not subject to such assumptions. They may be color photographs (Figure 7a), paintings, computer renderings or black-and-white sketches.

**LUMA** Because the fast discrete curvelet transform used in every variant of the signature extraction step takes a single 2D matrix as input, images with more than one color channel need to be reduced to one channel. The RGB values from the benchmark dataset have therefore been converted to greyscale images using the definition of luma according to ITU standards [34] (Figure 7b). Each pixel with red, green and blue values (R, G, B) is mapped to a luminance value Y using

$$Y = \frac{299}{1000}R + \frac{587}{1000}G + \frac{114}{1000}B.$$

**SOBEL** In order to make comparing the query sketch to the database images more effective, an edge extraction algorithm can be applied to

each database image. The Sobel operator calculates horizontal and vertical gradients by convolving the image with the  $3 \times 3$  kernels

$$K_x = \begin{bmatrix} -1 & 0 & 1 \\ -2 & 0 & 2 \\ -1 & 0 & 1 \end{bmatrix} \text{ and } K_y = \begin{bmatrix} -1 & -2 & -1 \\ 0 & 0 & 0 \\ 1 & 2 & 1 \end{bmatrix}$$

to obtain the directional gradients  $G_x$  and  $G_y$ . The overall response is the gradient magnitude  $G$  at each pixel location (Figure 7c):

$$G = \sqrt{G_x^2 + G_y^2}$$

**CANNY** A slightly more complex way to extract edges is the Canny edge detector [10]. Initially, the image is smoothed via a convolution with a small gaussian kernel to reduce the susceptibility to noise, even though this increases the localization error of the edge detection. On the smoothed image the gradient magnitude is calculated using the Sobel operator described above. The angle of the gradient can be calculated from the directional gradients  $G_x$  and  $G_y$  using

$$\Theta = \arctan \frac{G_x}{G_y}$$

and quantized into bins for  $0^\circ$ ,  $45^\circ$ ,  $90^\circ$  and  $135^\circ$ . Thin edges can be obtained from the gradient magnitudes by performing non-maximum suppression along the direction perpendicular to the gradient direction, e.g. a pixel is marked as being on a  $90^\circ$  edge if its magnitude is larger than the magnitudes north and south of its location. To avoid lines being broken up by noisy fluctuations, the edges are traced along their direction and gaps are filled in if the signal within the gap is above a certain threshold. The result is a binary edge map of the whole image (Figure 7d).

**SEGMENT** What a human sketches as a line in an image is often a boundary between two regions with different color or texture characteristics. Therefore the output from image segmentation algorithms can also indicate to location of edges. The hierarchical segmentation algorithm gPb-owt-ucm published by Arbeláez et al. [3] [2] was chosen because it represents the most recent advances in contour detection and it incorporates both local and global image information. On the gPb contour detector described in 2.3.4 they apply the Oriented Watershed Transform, that merges adjacent regions of an over-segmented image. The criterion for merging is the strength of the boundary shared by the two regions.

### 3.2 SIGNATURE EXTRACTION

Each method of signature extraction described in this section has the fast discrete curvelet transform at its heart. The curvelet transform has

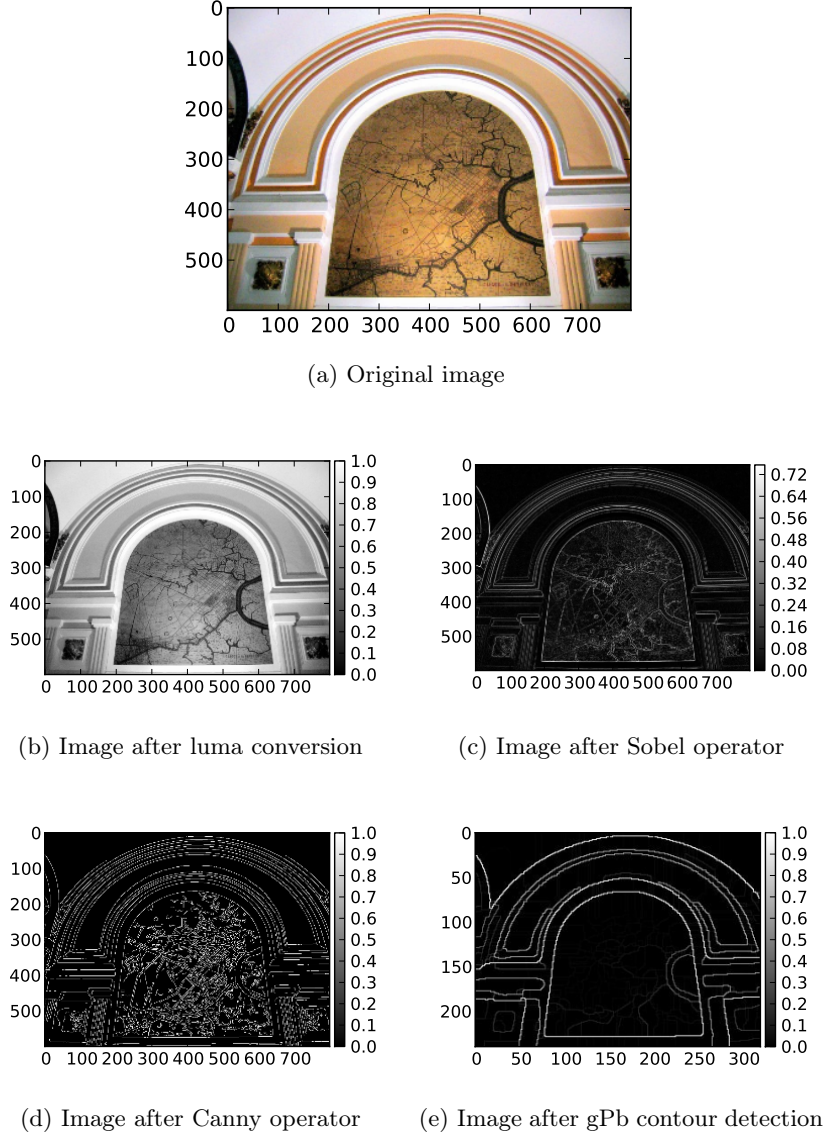


Figure 7: Image acquisition variants

two main parameters, that influence the result: The number of angles  $N_\theta$  used at the coarsest scale and the number of scales  $N_j$ , which corresponds to the number of concentric squares shown in 6. Experiments conducted to determine the optimal values of these parameters have shown that using more scales than 4 does not provide benefits that would justify the increased amount of processing time. Similar findings were reported by Mandal et al. [26] and Guha and Wu [18]. Furthermore, since the coarsest scale is non-directional, as explained in section 2.3.2, it is ignored in further computations. Finally, since the curvelet coefficients can have positive or negative sign, only the absolute value of the coefficient is used in the calculations of the mean values.

The response image generated by the FDCT for each pair of scale and angle is too large to be considered for the signature directly. Therefore

the response image  $C_{s,\theta}$  for scale  $s$  and angle  $\theta$  is subdivided into  $n^2$  equally sized grid cells  $G_{s,\theta,x,y}$  with  $x, y \in 1, 2, \dots, n$ :

$$C_{s,\theta} = \begin{bmatrix} G_{s,\theta,1,1} & G_{s,\theta,1,2} & \cdots & G_{s,\theta,1,n} \\ G_{s,\theta,2,1} & G_{s,\theta,2,2} & \cdots & G_{s,\theta,2,n} \\ \vdots & \vdots & \ddots & \vdots \\ G_{s,\theta,n,1} & G_{s,\theta,n,2} & \cdots & G_{s,\theta,n,n} \end{bmatrix}$$

For each of these grid cells, the mean  $\bar{C}_{s,\theta}$  is calculated:

$$\begin{aligned} \bar{C}_{s,\theta} &= \begin{bmatrix} \text{mean}(G_{s,\theta,1,1}) & \text{mean}(G_{s,\theta,1,2}) & \cdots & \text{mean}(G_{s,\theta,1,n}) \\ \text{mean}(G_{s,\theta,2,1}) & \text{mean}(G_{s,\theta,2,2}) & \cdots & \text{mean}(G_{s,\theta,2,n}) \\ \vdots & \vdots & \ddots & \vdots \\ \text{mean}(G_{s,\theta,n,1}) & \text{mean}(G_{s,\theta,n,2}) & \cdots & \text{mean}(G_{s,\theta,n,n}) \end{bmatrix} \\ &= \begin{bmatrix} \bar{c}_{s,\theta,1,1} & \bar{c}_{s,\theta,1,2} & \cdots & \bar{c}_{s,\theta,1,n} \\ \bar{c}_{s,\theta,2,1} & \bar{c}_{s,\theta,2,2} & \cdots & \bar{c}_{s,\theta,2,n} \\ \vdots & \vdots & \ddots & \vdots \\ \bar{c}_{s,\theta,n,1} & \bar{c}_{s,\theta,n,2} & \cdots & \bar{c}_{s,\theta,n,n} \end{bmatrix} \end{aligned}$$

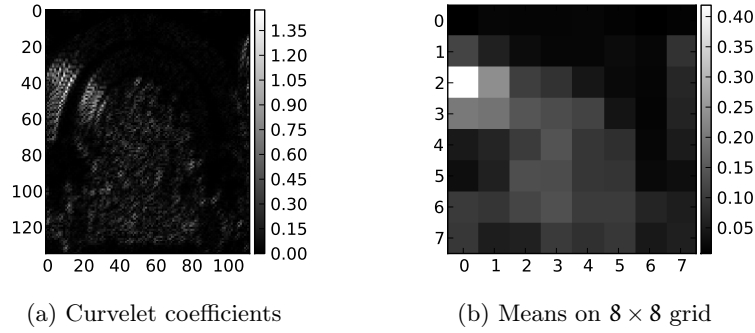


Figure 8: The curvelet coefficients for a specific scale and angle can be seen in (a). In (b) the response image has been subdivided into  $8 \times 8$  cells and the mean value of each cell are shown.

### 3.2.1 Global Features

**MEAN** The global approach to signature extraction simply takes the family of matrices  $\bar{C}_{s,\theta}$  and concatenates them as the image signature.

### 3.2.2 Local Features

The local feature extraction methods used here follow the bag-of-features approach, that aims to represent an image using a set of local feature descriptions or "visual words" similar to what was described by Sivic et al. [43].

### 3.2.2.1 Sampling

Both of the feature vector extraction methods described above use dense, overlapping sampling of a grid of mean values. By using  $m \times m$  windows, the small-scale spatial relationship between features can be captured. The overlap helps to avoid misinterpretation of features on grid boundaries that would occur with dense, non-overlapping sampling. Each window encodes the geometric relationships between cells within a neighborhood. Evaluations by Nowak et al. [32] have shown that random sampling, which dense sampling is a special case of, outperforms keypoint-based sampling for large enough numbers of samples. Dense sampling on grids has previously been successfully used by Lazebnik in [21] and [22]. The R-HOG descriptor [12] also uses a dense grid for sampling with overlapping windows to improve matching performance.

**PMEAN** Continuing from the set of matrices  $\bar{C}_{s,\theta}$ , this algorithm densely samples each matrix by sliding a window of size  $m \times m$ ,  $m < n$  across it (Figure 9). This results in  $n - m + 1$  parts  $\bar{W}_{s,\theta,u,v}$  with  $u, v \in 1, \dots, n - m + 1$ :

$$\bar{W}_{s,\theta,u,v} = \begin{bmatrix} \bar{c}_{s,\theta,u,v} & \bar{c}_{s,\theta,u,v+1} & \cdots & \bar{c}_{s,\theta,u,v+m} \\ \bar{c}_{s,\theta,u+1,v} & \bar{c}_{s,\theta,u+1,v+1} & \cdots & \bar{c}_{s,\theta,u+1,v+m} \\ \vdots & \vdots & \ddots & \vdots \\ \bar{c}_{s,\theta,u+m,v} & \bar{c}_{s,\theta,u+m,v+1} & \cdots & \bar{c}_{s,\theta,u+m,v+m} \end{bmatrix}$$

For each pair  $(u, v)$  these matrices are concatenated in a consistent way and stored as the feature vectors of the image. That way, the algorithm derives  $u \cdot v$  vectors of length  $N_s \cdot N_{\theta_s} \cdot m^2$  from each image, where  $N_{\theta_s}$  is the number of angles at the scale  $s$ .

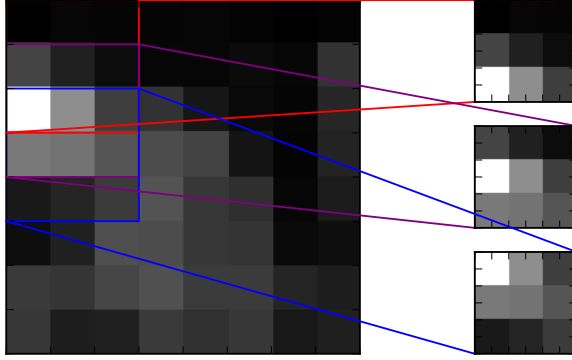


Figure 9: The  $8 \times 8$  mean coefficient grid  $\bar{C}_{s,\theta}$  is densely sampled using windows of  $3 \times 3$  grid cells to produce 36 patches per scale and angle.

**PMEAN2** In most parts, this variant is identical with the previously described PMEAN algorithm with the exception of the final signature feature vectors. Instead of concatenating all vectors  $\bar{W}_{s,\theta,u,v}$ , each scale

is handled separately. Therefore, each combination of  $(\mathbf{u}, \mathbf{v}, \mathbf{s})$  produces a vector of length  $N_{\theta_s} \cdot m^2$ .

### 3.2.2.2 *Clustering and Vector Quantization*

**CLUSTER** The set of feature vectors extracted from the images are numerous and large. In order to create meaningful image signatures from these vectors, the whole set is condensed into a dictionary using k-means clustering. As already discussed in section 2.2.2.3, the goal thereof is to derive a dictionary of predefined size that contains the visual words corresponding to the most discriminating features of the images in the image database. A universally optimal size for the code-book does not seem to exist, as Nowak et al. [32] observed an increase in accuracy up to 1000 words, but overfitting for some sampling algorithms beyond that. At the same time Yang et al. [52] report optimal sizes of 20000 to 80000 depending on the image database. The findings of Eitz et al. [14] agree with Nowak et al. in that a size of 1000 visual words was optimal for sketches, which is why that size was chosen for the following evaluation.

**VQ** The dictionary generated in the clustering step is then used to quantize the feature vectors extracted by the previous PMEAN and PMEAN2 procedures and record their number in signature vector as explained in section 2.2.2.3. To improve the extend in which this vector describes the image it was generated from, its components are then weighted. The TF-IDF statistic detailed in 2.2.3.1 is used to calculate a factor for each value that diminishes common, non-distinguishing visual words and boosts the importance of rare or unique words.

## 3.3 RANKING

As the last step of the pipeline, ranking operates on the signatures produced by prior extraction steps. It outputs a sequence of database images, sorted by the distance to the query image in ascending order. The distance can be determined using various metrics and similarity measures, which have been detailed in 2.2.3.2. This section lists the metrics used in the experiments and labels them for later reference.

**L<sub>2</sub>** The simplest and most widely used distance metric calculates the euclidean distance between the query image's signature and each database image's signature.

**HI** When comparing histograms of features, the histogram intersection measure  $s_{HI}$  has been shown to be superior to the euclidean distance in most of the cases [50]. It has the added benefit of allowing for partial matches in a signature. As can be seen from the definition

in 2.2.3.2, the result lies within  $[0, 1]$  with 1 being a perfect match. Therefore,  $1 - s_{HI}$  is used as a distance value to sort the result list.

**HIB** The binary variant of the histogram intersection measure converts the bin counts into a boolean representation, where a 1 indicates the presence of a codeword irrespective of the actual count, and a 0 denotes a codeword's absence. The resulting sequence of zeros and ones is then treated like in the HI measure.

**EMD** The Earth Mover's Distance is solved using a simplex algorithm variant, which has an exponential worst case complexity. That makes it computationally more expensive than the linear complexity measures described previously.





## EXPERIMENTAL RESULTS

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This chapter will present the benchmarking methods as well as the specific processing pipelines constructed from the steps explained in chapter 3. A detailed description of the experimental results for each pipeline variation will follow. Many pipelines were tested with varying parameters, some of which are common to all experiments and some of which are specific to the implementation.

### 4.1 BENCHMARKING METHOD

A usual way to evaluate the performance of retrieval systems is to calculate the ratio of true positive and false positive matches and visualize it in a receiver operating characteristic (ROC) curve. While that approach is well suited for benchmarking binary decision algorithms, it is not appropriate for retrieval problems, that don't feature a well-defined "correct" solution. An alternative approach is looking the recall and precision characteristics defined as

$$\text{recall} = \frac{\text{number of correct positive results}}{\text{total number of positives}}$$

$$1 - \text{precision} = \frac{\text{number of false positive results}}{\text{total number of results}}$$

Even though this metric works better for algorithms that return a set of results, it is still based on the notion of a "positive match", which requires an a-priori classification of the benchmark data. For systems that assign a degree of similarity to each item in the result set, this would require applying a threshold for classification. In addition to the distorting influence of an unsuitable thresholding choice, this method completely disregards the match quality within the positive result set, which can be important to the user of such a retrieval system.

Since sketch-based image retrieval systems are most likely to be used in interactive search applications of some form, it is desirable to assess the performance in relation to the results a human would achieve. Therefore the benchmark used to evaluate the retrieval pipelines corresponds to the method described by Eitz et al. [14], in which the authors create a benchmark dataset and perform a user study with 28 participants to define "ground truth" rankings. The dataset is divided into 31 groups of one sketch and 40 images each. Participants ranked the 40 images within each group by assigning scores indicating the similarity to the corresponding sketch in a controlled study environment. Each sketch/image pair's final ground truth ranking is calculated as the mean of the scores assigned by all participants.

To compare a ground truth ranking  $\mathbf{x} = (x_1, x_2, \dots, x_n)$  to a ranking  $\mathbf{y} = (y_1, y_2, \dots, y_n)$  produced by a retrieval system, the Kendall rank correlation coefficient  $\tau_B$  is used. It measures the similarity of the orderings by grouping all pairs  $\mathbf{p}_{i,j} = \{(x_i, y_i), (x_j, y_j)\}$ ,  $i, j \in 1, \dots, n$  into 5 sets:

$\mathbf{p}_{i,j} \in C$	if $x_i < x_j$ and $y_i < y_j$
$\mathbf{p}_{i,j} \in D$	if $x_i < x_j$ and $y_i > y_j$
$\mathbf{p}_{i,j} \in T_x$	if $x_i = x_j$ and $y_i \neq y_j$
$\mathbf{p}_{i,j} \in T_y$	if $x_i \neq x_j$ and $y_i = y_j$
$\mathbf{p}_{i,j} \in T_{xy}$	if $x_i = x_j$ and $y_i = y_j$

From that, the correlation value  $\tau_B$  in the interval  $[-1, 1]$  can be calculated as

$$\tau_B = \frac{|C| - |D|}{\sqrt{(|C| + |D| + |T_x|)(|C| + |D| + |T_y|)}}.$$

The higher  $\tau_B$  is, the more pairs in  $\mathbf{x}$  and  $\mathbf{y}$  have a similar ordering. Since the values are only compared within each ranking, the result is independent of each rankings' scaling, making it ideal for comparison of different distance metrics.

#### 4.2 CROSS-DOMAIN RESULTS

Some of the variations below include a Canny edge detector as a pre-processing step. The parameter  $\sigma$  determines the size of the Gaussian kernel used by the Canny algorithm. A value of  $\sigma = 1.5$  was determined to be appropriate for the image dataset used in the benchmark. Value larger than  $\sigma = 2$  tend not to detect any edges, while smaller values produced more "false" edges resulting from noise in the images.

The SEGMENT preprocessing step could unfortunately not use the same  $1024 \times 768$  pixel images as inputs as the other preprocessors, because the implementation provided by Arbelaez et al. [2] could not be run on any machine available due to extreme memory requirements. Therefore, the images in the database are rescaled to  $320 \times 240$  pixels for the evaluation of this algorithm. This might lead to loss of small features and puts limits on some parameters like the grid size  $G$ .

All pipeline variants utilize the FDCT to extract curve information. The number of scales and the number of angles at the coarsest scale will be called  $N_s$  and  $N_\theta$  respectively. Based on experimentation and other publications using the curvelet transform [26] [18], 4 scales and 12 angles are used in almost all cases, since larger values did not consistently lead to better results. The size of the codebook is set at 1000 visual words, as values beyond that did not show improvements in evaluations performed by Nowak et al. [32]. Eitz et al. [14] reported similar optimal values of 500 to 1000 visual words and attributed the low number to the sparsity of edge-like features in sketches.

Throughout the correlation coefficient graphs in the following sections, the red line denotes the best result obtained by Eitz et al. in [14] using the SHOG descriptor.

The first two sections present the results grouped by preprocessing steps and sampling method. Afterwards, the influence of parameter variations on selected pipelines are examined.

#### 4.2.1 Global Features

The processing steps applied to the database images and the query images in the pipelines based on global features are almost identical, with the exception of the CANNY, SOBEL and SEGMENT steps for the database images. The curvelet responses are averaged on a grid and the features are ranked using the  $L_2$  and the COS distance measures.

**LUMA+MEAN** The most straightforward combination of processing steps consists of a LUMA input image, on which the means of  $G \times G$  grid cells is computed for each scale and angle (Figure 10). Varying  $G$  determines the feature size that can be encoded best. Setting  $G = 12$  seems to yield the best correlation, although the advantage over  $G = 8$  and  $G = 16$  is below 0.01. The distance measure COS outperforms the  $L_2$  measure (Table 1). A possible explanation would be that the COS measure normalizes the magnitude of the feature vector as discussed in 2.2.3.2.

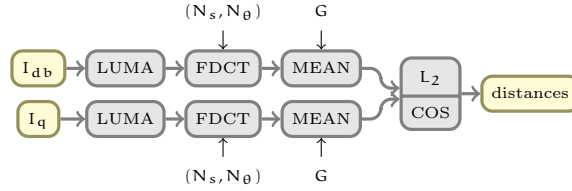


Figure 10: Global LUMA+MEAN Pipelines

$N_s$	$N_\theta$	$G$	Metric	
4	12	4	COS	0.177
4	12	8	$L_2$	0.14
4	12	8	COS	0.183
4	12	12	$L_2$	0.14
4	12	12	COS	0.191
4	12	16	COS	0.186

Table 1: Global LUMA+MEAN Results

**CANNY+MEAN** In addition to reading the images like in the previous LUMA+MEAN configuration, this pipeline applies a CANNY pro-

cessing step to the database images in an attempt to bring the query and database image domains closer together (Figure 11). Again, the COS distance measure produces the best rankings (Table 2). Surprisingly, the Canny edge detector does not lead to increased performance in comparison to plain the LUMA preprocessing step.

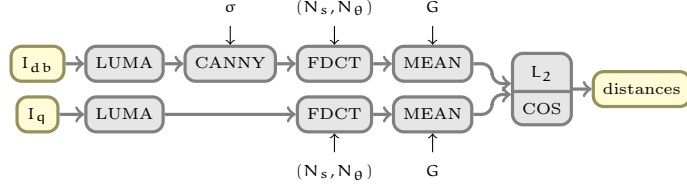


Figure 11: Global CANNY+MEAN Pipelines

$N_s$	$N_\theta$	G	$\sigma$	Metric	
4	12	4	1.5	COS	0.179
4	12	8	1.5	$L_2$	0.095
4	12	8	1.5	COS	0.188
4	12	12	1.5	$L_2$	0.096
4	12	12	1.5	COS	0.184
4	12	16	1.5	COS	0.182

Table 2: Global CANNY+MEAN Results

**SOBEL+MEAN** The SOBEL step used in this variant also attempts to bring the database images into the sketch domain (Figure 12). The results are slightly better than with the CANNY preprocessor (Table 3).

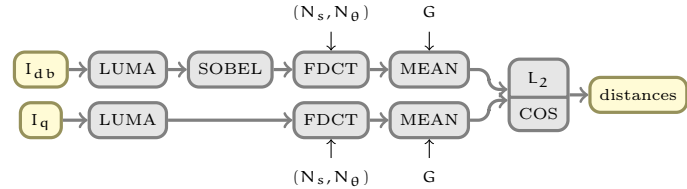


Figure 12: Global SOBEL+MEAN Pipelines

$N_s$	$N_\theta$	G	Metric	
4	12	8	$L_2$	0.15
4	12	8	COS	0.19
4	12	12	$L_2$	0.152
4	12	12	COS	0.2

Table 3: Global SOBEL+MEAN Results

**SEGMENT+MEAN** With the gPb contour detector in the SEGMENT step to find edges in the database images (Figure 13), the  $L_2$  distance metric produces results comparable to the COS metric in the CANNY+MEAN variant (Table 4). Unlike in the other cases, the COS distance measure performs worse than the  $L_2$  metric.

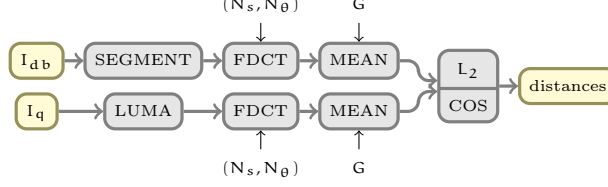


Figure 13: Global SEGMENT+MEAN Pipelines

$N_s$	$N_\theta$	G	Metric	
4	12	4	$L_2$	0.188
4	12	4	COS	0.168
4	12	8	$L_2$	0.171
4	12	8	COS	0.169

Table 4: Global SEGMENT+MEAN Results

#### 4.2.2 Local Features

The local feature extraction pipelines utilize the same preprocessing steps as the ones based on global features. The curvelet transform remains unchanged as well. The the curvelet coefficients are sampled using the PMEAN and PMEAN2 strategy to produce features for local patches. From these feature vectors, a codebook of size 1000 is generated and the feature vectors from both the database images and the query images are quantized and weighted using the TF-ITF method. The metrics used for ranking are the generic cosine and  $L_2$  distance measures as well as histogram intersection and the earth mover's distance.

**LUMA+PMEAN(2)** Analogous to the global LUMA+MEAN variant, this pipeline configuration directly uses the LUMA image representations as input for the curvelet transform (Figure 14). For both, the PMEAN (Table 5a) and PMEAN2 (Table 5b) sampling method, histogram intersection and the cosine measures are far superior to the  $L_2$  metric and even the EMD metric, although the results cannot compete with the global LUMA+MEAN version.

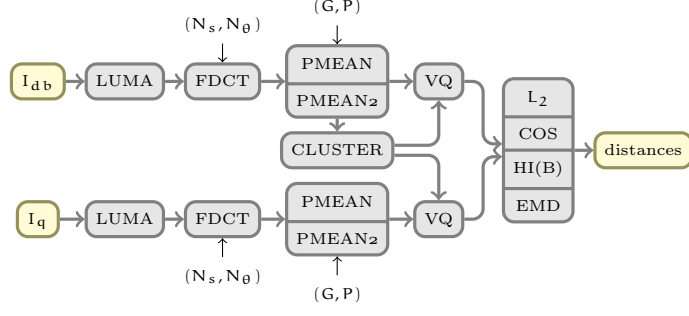
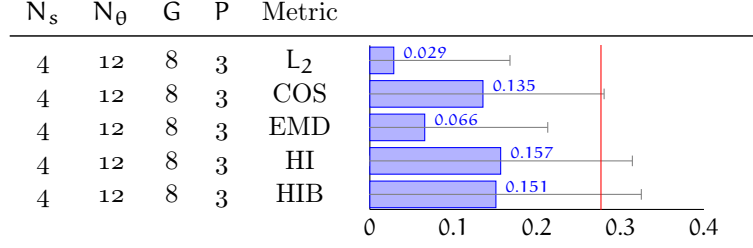


Figure 14: Local LUMA+PMEAN(2) Pipelines



(a) Local LUMA+PMEAN Results

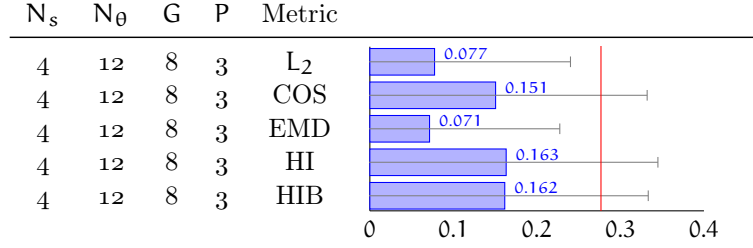
(b) Local LUMA+PMEAN<sub>2</sub> Results

Table 5: Local LUMA+PMEAN(2) Results

**CANNY+PMEAN(2)** Using the Canny edge detector and PMEAN sampling (Figure 15), the rank correlation coefficient for COS and HI exceed all previous results (Table 6a) with the highest value achieved is 0.22. Treating the coefficients on different scales separately using the PMEAN<sub>2</sub> sampling method is inferior to PMEAN in this case.

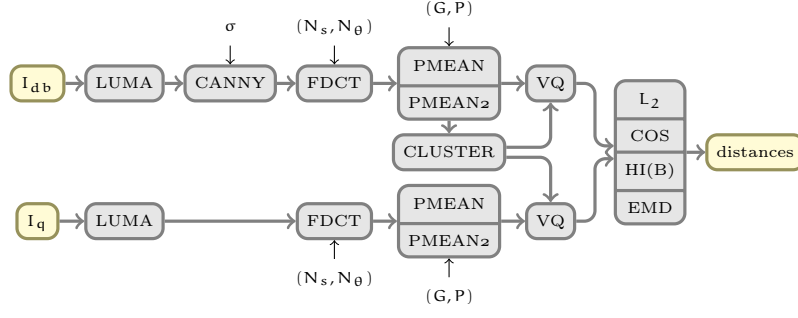
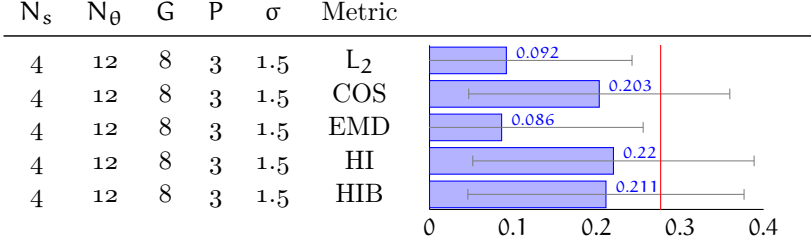


Figure 15: Local CANNY+PMEAN Pipelines



(a) Local CANNY+PMEAN Results

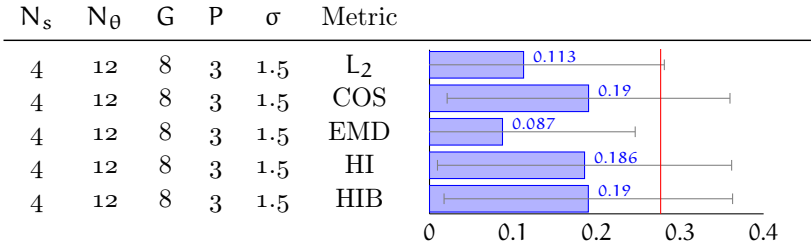
(b) Local CANNY+PMEAN<sub>2</sub> Results

Table 6: Local CANNY+PMEAN(2) Results

SOBEL+PMEAN(2) Preprocessing the database images with the Sobel operator results in slightly lower correlation coefficients. The PMEAN<sub>2</sub> sampling algorithm yields better results for the  $L_2$  and EMD metrics, but worse for COS, HI and HIB.

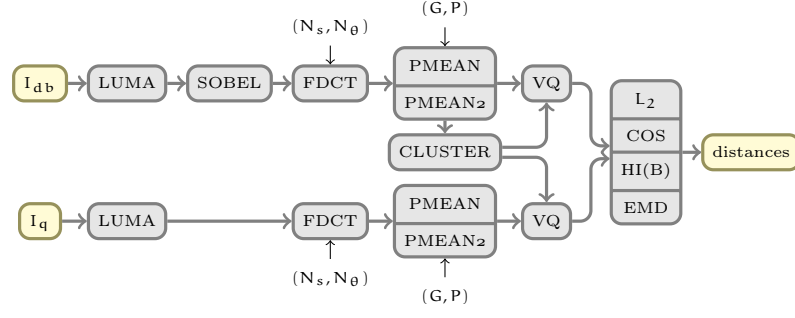
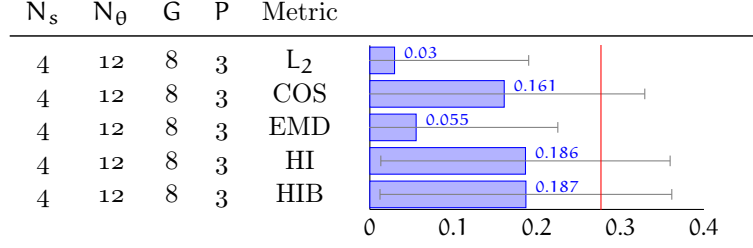


Figure 16: Local SOBEL+PMEAN Pipelines



(a) Local SOBEL+PMEAN Results

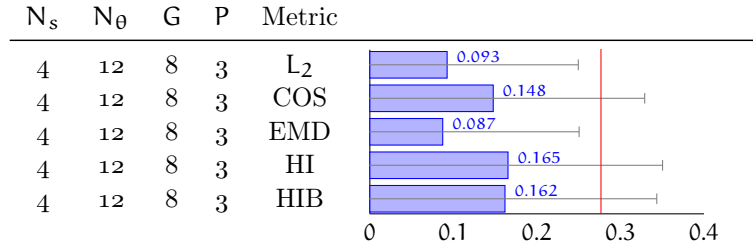
(b) Local SOBEL+PMEAN<sub>2</sub> Results

Table 7: Local SOBEL+PMEAN(2) Results

SEGMENT+PMEAN(2) The SEGMENT step leads to similar, although slightly lower, results than the Sobel operator. Neither PMEAN nor PMEAN<sub>2</sub> have a consistent advantage across the different distance measures.



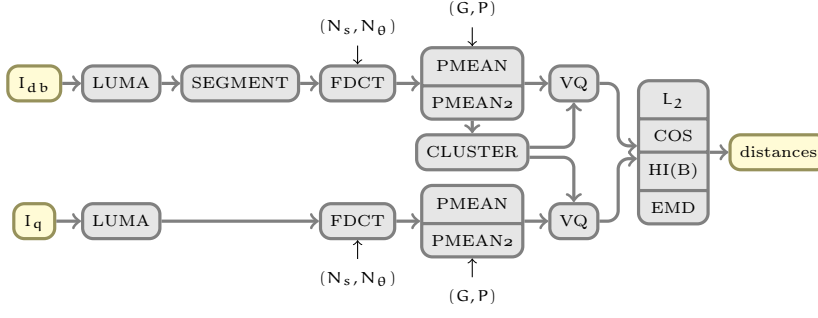
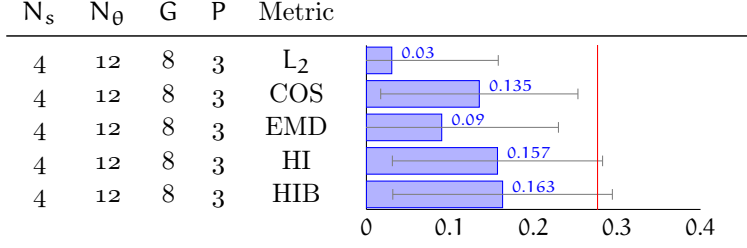


Figure 17: Local SEGMENT+PMEAN Pipelines



(a) Local SEGMENT+PMEAN Results

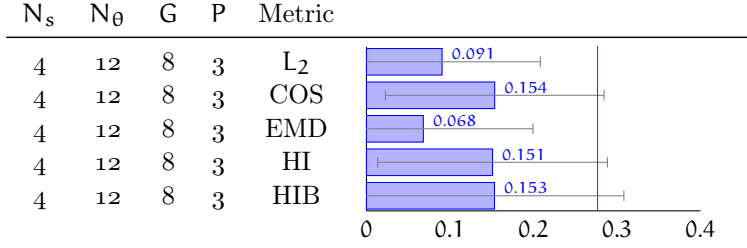
(b) Local SEGMENT+PMEAN<sub>2</sub> Results

Table 8: Local SEGMENT+PMEAN(2) Results

#### 4.2.3 Parameter Variations

In an attempt to improve the results, the best performing configurations presented in the previous sections will be re-used with varying parameter values (Table 9). The following sections compare the results of changing the parameters  $N_\theta$ ,  $P$ ,  $G$  and  $\sigma$ .

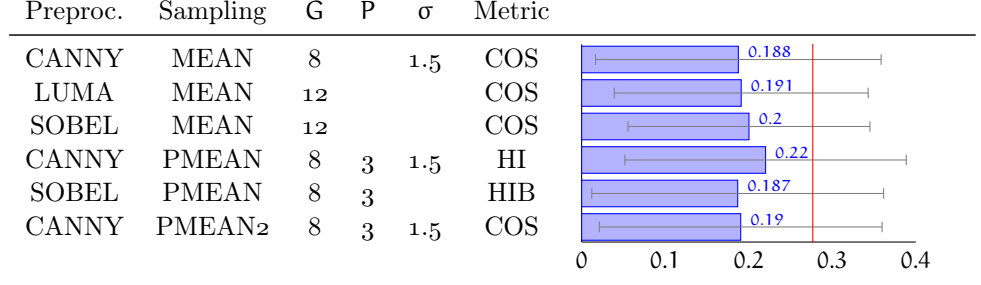


Table 9: Best Performing Configurations with default assumptions  $N_s = 4$  and  $N_\theta = 12$ .

#### 4.2.3.1 Curvelet Angles

The parameter  $N_\theta$  controls the number of angles, the curvelet coronization is divided into (see Section 2.3.3). Therefore, it determines how finely the angles of the lines are resolved and how sensitive to angular differences the descriptor is.

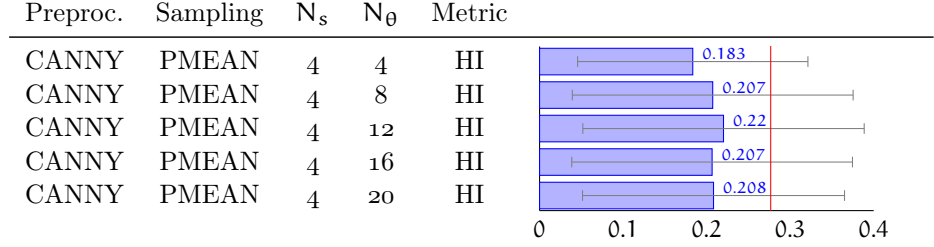


Table 10: Influence of  $N_\theta$  on the results of CANNY+PMEAN for  $G = 8$ ,  $P = 3$  and  $\sigma = 1.5$ .

#### 4.2.3.2 Grid and Patch Sizes

All of the MEAN, PMEAN and PMEAN<sub>2</sub> sampling methods use a regular grid to divide the curvelet coefficients into cells, in which the mean of the coefficients is calculated. Using a small number  $G$  of subdivisions means that smaller features might vanish within a large grid cell, unable to influence the mean value. A finer subdivision allows for smaller features to be represented at the risk of cutting apart larger features that lie on the grid lines. In addition to  $G$ , the local sampling methods PMEAN and PMEAN<sub>2</sub> are influenced by the number of grid cells that make up a patch. As explained in section 3.2.2, a patch captures the geometric relationships within a  $P \times P$  neighborhood of cells. It thus defines an upper limit on the size of a feature that can be represented atomically.

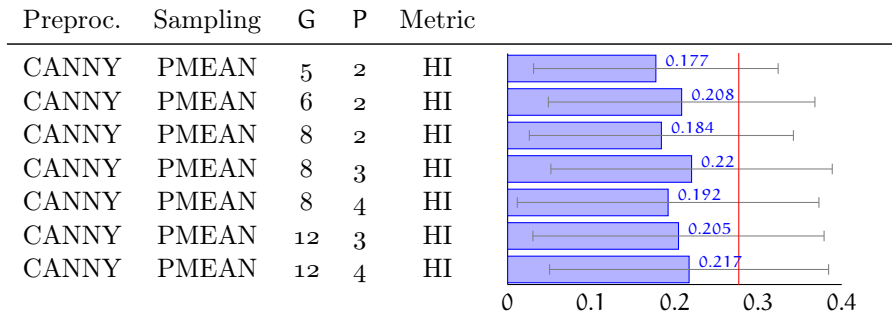


Table 11: Influence of grid parameters  $P$  and  $G$  on the results for  $N_s = 4$ ,  $N_\theta = 12$  and  $\sigma = 1.5$ .

#### 4.2.3.3 Canny Sigma

### 4.3 INTRA-DOMAIN RESULTS

TBD



## DISCUSSION

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Discussion of results goes here. . .



## CONCLUSION

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Conclusion goes here. . .





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*Berlin, January 2012*

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Felix Stürmer