# Investigating the Intrinsic Geometry of the CelebA Dataset Using Diffusion Geometry Techniques

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September 2025

Undergraduate Summer Research Project

Durham University

Funded By The London Mathematical Society

## Abstract

In this project...

## Acknowledgements

I would like to thank Prof. Jeffrey Giansiracusa for his continued guidance and support during this project.

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## 1 Introduction

## 1.1 Motivation

The CelebFaces Attributes Dataset, commonly known as CelebA, is a facial attributes dataset comprising 202,599 images of celebrities [1]. CelebA is a popular benchmark dataset in machine learning, used for training and testing models for tasks such as facial recognition.

Julian Chang Jun Yeng discovered that hyperbolic variational autoencoders outperform Euclidean variational autoencoders when reconstructing the CelebA images [2]. This suggests that the CelebA dataset may have intrinsic hyperbolic geometry.

To further investigate the intrinsic geometry of the CelebA dataset, we employ novel tools developed by Dr Iolo Jones [3]. These tools utilise diffusion geometry techniques to estimate the dimension and curvature of data embedded in higher-dimensional manifolds.

## 1.2 Aims of the Project

In this summer research project, we first aim to understand relevent areas of differential geometry. We introduce surfaces in Euclidean space and study the second fundamental form and Gaussian curvature.

We will then study the theory of Diffusion Maps. We will build a Python algorithm that utilises diffusion maps to reduce the dimension of the CelebA images to three, in order to produce a visual representation of the entire dataset.

Finally, we implement the Python tools developed by Dr Iolo Jones in his paper [3]. We aim to investigate the intrinsic dimension and curvature of the CelebA dataset.

## 2 Mathematical Background

In this section, we present the neccessary background required to understand Gaussian and scalar curvature. We also explain the theory of diffusion maps and why they are useful for dimensionality reduction.

## 2.1 Surfaces in Euclidean Space and Gaussian Curvature

The information presented in this chapter is mainly adapted from Curves and Surfaces, by Montiel and Ros [4].

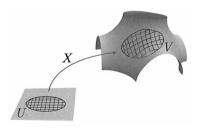
#### 2.1.1 Defining Surfaces and Curves

Informally, a set  $S \in \mathbb{R}^3$  is a surface if, at every point in S, S locally looks like a slightly deformed patch of  $\mathbb{R}^2$ . We rigourously define a surface below, and provide a visual representation in Figure 2.1.

**Definition 2.1.** Consider a non-empty set  $S \subset \mathbb{R}^3$ . We say S is a **surface** if for all  $p \in S$ , there exists an open neighbourhood  $V \subset S$  of p, an open set  $U \in \mathbb{R}^2$ , and a map  $X : U \to \mathbb{R}^3$ , where X has the following properties:

- i)  $X: U \to V$  is a homeomorphism.
- ii)  $(dX)_q: \mathbb{R}^2 \to \mathbb{R}^3$  is injective for all q in U.

We call each such  $X: U \to S$  a parametrisation of S.



**Figure 2.1:** Diagram of a parametrisation of some surface in  $\mathbb{R}^3$ .

**Remark 2.2.** Property (i) means X is continuous, bijective, and has a continuous inverse.

**Remark 2.3.** In property (i),  $(dX)_q$  is called a differential, and, in this case, can be defined as follows. If we write

$$X(u,v) = (X_1(u,v), X_2(u,v), X_3(u,v))$$

then

$$(dX)_q(a,b) = \begin{bmatrix} \frac{\partial X_1}{\partial u}(q) & \frac{\partial X_1}{\partial v}(q) \\ \frac{\partial X_2}{\partial u}(q) & \frac{\partial X_2}{\partial v}(q) \\ \frac{\partial X_3}{\partial u}(q) & \frac{\partial X_3}{\partial v}(q) \end{bmatrix} \begin{bmatrix} a \\ b \end{bmatrix}.$$

One can think of  $(dX)_q$  here as the best linear approximation of the function X at q. We will revisit differentials in Section 2.1.4.

Note 2.4. Surfaces can be embedded in higher dimensions than three, and we can also define hypersurfaces with dimensions greater than two. However, here we focus on 2-dimensional surfaces in  $\mathbb{R}^3$  to aid visualisation.

**Definition 2.5.** We define a **curve in**  $\mathbb{R}^3$  as a  $C^{\infty}$  map  $\alpha: I \to \mathbb{R}^3$  for some open  $I \subset \mathbb{R}$ . If  $\alpha(I) \subset S$  for some surface S, we say  $\alpha$  is a **curve on** S.

#### 2.1.2 Differentiable Functions on Surfaces

Consider a function defined on a surface. We must define what it means for such a function to be differentiable. To do this, we use parametrisations and inclusion maps to reduce the concept to our familiar notion of differentiability.

**Definition 2.6.** Given a surface  $S \subset \mathbb{R}^3$ , the **inclusion map**  $i: S \to \mathbb{R}^3$  is the identity map, regarded as having a different domain and codomain. That is,  $\forall p \in S$ , i(p) = p but we say  $i(p) \in \mathbb{R}^3$ .

**Definition 2.7.** Let S be a surface.

- (A) A function  $f: S \to \mathbb{R}^m$  is differentiable if, for every parametrisation  $X: U \to S$ , the composition  $f \circ X: U \to \mathbb{R}^m$  is differentiable (in the usual Euclidean sense).
- **(B)** Let  $O \subset \mathbb{R}^n$  be open. A map  $f: O \to S$  is **differentiable** if the composition  $i \circ f: O \to \mathbb{R}^3$  is differentiable, where  $i: S \to \mathbb{R}^3$  is the inclusion map.
- (C) If  $S_1$  is another surface and  $i_1: S_1 \to \mathbb{R}^3$  is its inclusion map, a map  $f: S \to S_1$  is **differentiable** if  $i_1 \circ f: S \to \mathbb{R}^3$  is differentiable in the sense of (A).

#### 2.1.3 Tangent Planes

Another important notion in differential geometry is that of the tangent plane to a surface at a given point.

**Definition 2.8.** Let S be a surface and  $p \in S$ . A vector  $v \in \mathbb{R}^3$  is **tangent to** S at p if there exists a curve on S

$$\alpha: (-\epsilon, \epsilon) \to S$$

for some  $\epsilon > 0$ , such that  $\alpha(0) = p$  and  $\alpha'(0) = v$ . The set of all vectors tangent to S at p is called the **tangent plane of** S at p and is denoted  $T_pS$ .

#### 2.1.4 The Differential of a Differentiable Map

The notion of the differential of a map between Euclidean spaces, aforementioned in Remark 2.3, can be extended to surfaces.

**Definition 2.9.** Let  $S_1$  and  $S_2$  be surfaces and  $f: S_1 \to S_2$  be a differentiable function. For a point  $p \in S_1$  the differential of f at p is a map

$$(df)_p: T_pS_1 \to \mathbb{R}^3$$

defined as follows: Given  $v \in T_pS_1$ , choose a curve  $\alpha: I \to S_1$  such that  $\alpha(0) = p$  and  $\alpha'(0) = v$ . Then

$$(df)_p(v) = \frac{d}{dt}\Big|_{t=0} (f \circ \alpha)(t) = (f \circ \alpha)'(0).$$

It can be shown that  $(df)_p(v)$  does not depend on the chosen curve,  $\alpha$ , hence is well defined. One can also prove that it is a linear map.

**Remark 2.10.** Since  $f: S_1 \to S_2$  and  $\alpha: I \to S_1$  are both differentiable functions, one can prove that  $f \circ \alpha: I \to S_2$  is a differentiable function. Hence, it is a curve on  $S_2$  with  $(f \circ \alpha)(0) = f(p)$  and  $(f \circ \alpha)'(0) = (df)_p(v)$ . This means  $(df)_p(v) \in T_{f(p)}S_2$ . So we can think of the differential of f at p as being a map

$$(df)_p: T_pS_1 \to T_{f(p)}S_2.$$

#### 2.1.5 The Gauss Map

**Definition 2.11.** Let  $S \subset \mathbb{R}^3$  be a surface. A **unit normal field** on S is a differentiable map  $N: S \to \mathbb{R}^3$  such that for all  $p \in S$  we have |N(p)| = 1 and for all  $v \in T_pS$  we have  $N(p) \perp v$ .

**Definition 2.12.** If one can define a single unit normal field on the entire surface, we call this unit normal field a **Gauss map**. In this case we call the surface **orientable**.

**Remark 2.13.** Let S be an orientable surface and  $N: S \to \mathbb{R}^3$  be a Gauss map. Since |N(p)| = 1 for all  $p \in S$  we have  $N(S) \subset \mathbb{S}^2$ , where  $\mathbb{S}$  is the unit sphere centered at the origin. Hence, we can consider the Gauss map as  $N: S \to \mathbb{S}^2$ .

## 2.1.6 Gaussian Curvature

the Gaussian curvature is well defined.

One can show that  $\mathbb{S}^2$  is a surface. Hence a Gauss map is a differentiable function between two surfaces. As a result, we can calculate its differential at a point.

Let  $S \in \mathbb{R}^3$  be an orientable surface,  $N: S \to \mathbb{S}^2$  be a Gauss map, and  $p \in S$ . The differential can be written  $(dN)_p: T_pS \to T_{N(p)}\mathbb{S}^2$ . However, note that  $T_{N(p)}\mathbb{S}^2$  is the set of all vectors orthogonal to N(p). Hence, it can be proven, and intuitively seen, that  $T_{N(p)}\mathbb{S}^2 = T_pS$ . This means  $(dN)_p$  is an endomorphism of  $T_pS$ .

**Definition 2.14.** Let  $S \in \mathbb{R}^3$  be an orientable surface with Gauss map  $N: S \to \mathbb{S}^2$ . Let  $p \in S$  and  $(dN)_p: T_pS \to T_pS$  be the differential of N at p. Since  $T_pS$  is 2-dimensional, given a basis of  $T_pS$  one can represent the endomorphism  $(dN)_p$  by a  $2 \times 2$  matrix. We define the **Gaussian curvature at** p to be the determinant of this matrix, and denote it by K(p). One can show the determinant does not depend on the chosen basis of  $T_pS$ , so

Note 2.15. To intuitively see why the Gaussian curvature relates to the shape of the surface, note that the Gauss map is a normal field on S. Hence, the differential of the Gauss map describes the variation of a normal field on the surface. It is not hard to see why this would give us information about the shape of the surface. The Gaussian curvature is just an associated invariant of this differential.

The sign of the Gaussian curvature at a point reveals important information about the geomtry of the surface at that point. Positive Gaussian curvature corresponds to an elliptic point, for example a bowl. Negative Gaussian curvature corresponds to a hyperbolic point, for example a saddle point. Zero Gaussian curvature corresponds to a point that is flat in at least one direction.

**Note 2.16.** In Section 3, we reference scalar curvature rather than Gaussian curvature. However, for 2-dimensional surfaces, the scalar curvature is just double the Gaussian curvature, so they have the same sign.

#### 2.1.7 The Second Fundamental Form

**Definition 2.17.** The second fundamental form of S at p is the bilinear form  $\sigma_p:T_p\times T_p\to\mathbb{R}$ , where

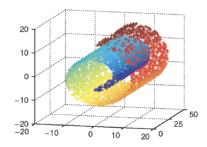
$$\sigma_p(v, w) = -\langle (dN)_p(v), w \rangle.$$

## 2.2 Diffusion Maps

## 2.2.1 Construction of a Random Walk on the Data

Suppose we have m data points, each of which is represented by a vector in  $\mathbb{R}^n$ . We will denote this dataset by X. We consider each data point as a node on a graph, and we aim to construct a random walk on this graph [5].

We define diffusion maps in general, however, it might be intuitively helpful to consider the following example given in Figure 2.2.



**Figure 2.2:** 5,000 points sampled from a Swiss roll in  $\mathbb{R}^3$  [6]

We first define a weight function  $k: X \times X \to \mathbb{R}$  by

$$k(x_i, x_j) = e^{\frac{-|x_i - x_j|}{\epsilon}},$$

where  $\epsilon > 0$  is some constant we discuss in Section 2.2.4.

This weight function allocates a larger weight between data points that are closer together in the ambient space  $\mathbb{R}^n$ .

Next, we define the degree function

$$d(x_i) = \sum_{x \in X} k(x_i, x).$$

We use this to normalise the weight function and create transition probabilities.

We define the matrix P with entries

$$P_{ij} = \frac{k(x_i, x_j)}{d(x_i)}.$$

P is a stochastic matrix as its rows sum to one. Hence, we can define a Markov process on our graph by setting the transition probability of going from  $x_i$  to  $x_i$  in one discrete time step to be  $P_{ij}$ .

**Note 2.18.** Since this Markov chain is clearly irreducible and aperiodic, it has a unique stationary distribution  $\pi \in [0, 1]^m$ . [7]

#### 2.2.2 Diffusion Distance and Diffusion Maps

**Definition 2.19.** The diffusion distance between  $x_i, x_i \in X$  is

$$D_t(x_i, x_j)^2 = \sum_{y \in X} \frac{(P_{x_i, y}^t - P_{x_j, y}^t)^2}{\pi(y)}$$
 (1)

where  $\pi$  is the unique stationary distribution of the Markov chain.

Note 2.20. Intuitively, the diffusion distance is a measure of how 'connected' two points are in the intrinsic geometry of the dataset. In some sense it is a measure of the probability of diffusing from one point to another along this Markov chain we have created. Points may be close together in the Euclidean sense but have a diffusion distance that is large. For example, the blue and red sections in Figure 2.2.

**Definition 2.21.** We define the diffusion map  $\Psi_t: X \to \mathbb{R}^n$  as

$$\Psi_t(x) = \begin{bmatrix} \lambda_1^t \psi_1(x) \\ \vdots \\ \lambda_n^t \psi_n(x) \end{bmatrix}$$

where  $\{\psi_l\}$  are the left eigenvectors of P corresponding to eigenvalues  $\{\lambda_l\}$ .

**Theorem 2.22.** Given  $x_i, x_i \in X$  we have

$$D_t(x_i, x_j)^2 = |\Psi_t(x_i) - \Psi_t(x_j)|^2$$
(2)

That is, the diffusion distance is equal to the Euclidean distance between the diffusion maps. The proof revolves around the eigenvector expansion of P [8].

**Remark 2.23.** This Theorem implies two points whose diffusion maps are close in the Euclidean sense have a strong 'connectivity'.

#### 2.2.3 Dimension Reduction

Consider the coordinates of the diffusion map corresponding to the smaller eigenvalues. When caluclating the Euclidean distance between diffusion maps, i.e. the right hand side of equation (2), these coordinates have less influence due to their smaller magnitude. This allows us to prove an approximation of Equation (2).

**Definition 2.24.** Firstly, we reorder the eigenvalue and eigenvector pairs so that  $\lambda_1 \leq \lambda_2 \leq ... \leq \lambda_n$ . We define then define the **truncated diffusion map** of order  $\mathbf{k}$ ,  $\Psi_t^{(k)}: X \to \mathbb{R}^k$ , as

$$\Psi_t^{(k)}(x) = \begin{bmatrix} \lambda_1^t \psi_1(x) \\ \vdots \\ \lambda_k^t \psi_k(x) \end{bmatrix}.$$

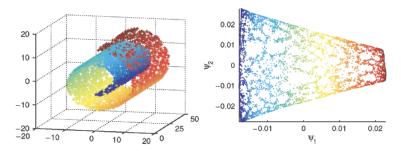
Equation (2) now becomes

$$D_t(x_i, x_j)^2 \approx |\Psi_t^{(k)}(x_i) - \Psi_t^{(k)}(x_j)|^2.$$

This is formally proven in [8].

By creating these truncated diffusion maps we can represent the data points of X, originally embedded in  $\mathbb{R}^n$ , with points now embedded in  $\mathbb{R}^k$  where k < n. The distance between the resulting points in  $\mathbb{R}^k$  is a measure of how 'connected' the points of the original dataset were.

We now revisit the exmple given in Figure 2.2.



**Figure 2.3:** 5,000 points sampled from a Swiss roll in  $\mathbb{R}^3$  (L). The truncated diffusion maps of order 2 (R). [6]

The data originally lived in  $\mathbb{R}^3$ , but when we plot the truncated diffusion maps of order 2 we obtain the plot given on the right of Figure 2.3. The diffusion maps have essentially uncovered the coordinates of the swiss roll and 'unravelled' it.

## 2.2.4 Revisiting the Parameter $\epsilon$ and Discussing t

When defining the random walk on our data in Section 2.2.1, we defined the weight function  $k: X \times X \to \mathbb{R}$  by

$$k(x_i, x_j) = e^{\frac{-|x_i - x_j|}{\epsilon}}.$$

The parameter  $\epsilon$  must be chosen when creating this kernel, and the value one picks is important.

A larger  $\epsilon$  means larger transition probabilities will be allocated to points father away from each other in the ambient Euclidean space. Conversely, a smaller  $\epsilon$  means points farther away will be allocated smaller transition probabilities.

The idea is that for  $\epsilon$  too large, the graph will be too 'connected' so the diffusion process will provide little information about the intrinsic geomtry of the data. Conversely, for epsilon too small, very little diffusion takes place, which again provides little information about the geometry of the surface.

There is a technique for estimating a good  $\epsilon$  to use, which we explore in Section 3.1.

#### DO YOU REALLY WANT TO DISCUSS T?

When defining the diffusion distance and diffusion maps, we have also seen the parameter t arise. The value of t we use when creating the diffusion maps decides how many steps we wish to take along our Markov chain. Adjusting this can change whether our diffusion maps pick up large scale or small scale features of the data.

An anology would be if we had data samples taken from two Swiss rolls living apart from each other in the ambient space  $\mathbb{R}^3$ . If we chose to pick a large t, we would not neccessarily uncover the small scale features of each swiss roll, but rather we would see two distinct clusters of points belonging to each Swiss roll.

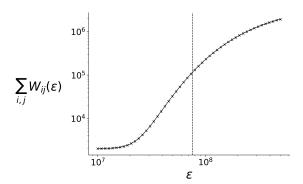
## 3 New Results

## 3.1 Estimating a Sensible $\epsilon$ for the CelebA Dataset

As previously mentioned in Section 2.2.4 there is a method for estimating an appropriate  $\epsilon$  to use when forming our diffusion maps. [9]

For a given  $\epsilon$  we calculate the weight matrix with entries  $W_{ij} = e^{\frac{-|x_i - x_j|^2}{\epsilon}}$ . We then sum over every entry of this matrix to obtain the sum of all the weights between our points,  $\sum_{i,j} W_{ij}$ .

The magnitude of this sum relates to how 'connected' the weighted graph we form is, and hence how much diffusion takes place in our final Markov chain. We plot the value of this sum against various  $\epsilon$  values, shown in Figure 3.1.



**Figure 3.1:** Logarithmic scale plot of  $\sum_{i,j} W_{ij}$  against  $\epsilon$  for 2,000 images of the CelebA dataset.

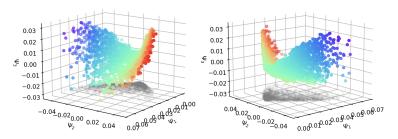
Coifman suggests that to ensure a meaningful amount of diffusion takes place, one should take an  $\epsilon$  at the center of the linear section of this logarithmic scale plot.

In the case of the CelebA dataset, this is about 76,000,000, so this is the value of  $\epsilon$  we shall use henceforth.

## 3.2 Reducing the Dimension of CelebA Images Using Diffusion Maps

By following the steps in Section 2.2 we can construct a visual 3-dimensional representation of the CelebA dataset by creating truncated diffusion maps of order 3.

The Python code used to do this is available at /GITHUB LINK



**Figure 3.2:** 2,000 images from the CelebA dataset reduced to 3-dimensions using diffusion maps.

In Figure 3.2 we obtain a curved 2-dimensional surface embedded in 3 dimensions. The fact that we obtain an obvious pattern rather than a random assortment of points suggests the diffusion maps have indeed uncovered some of the intrinsic geometry of the dataset.

## 3.3 Estimating the Dimension of the CelebA Dataset

Using Diffusion Geomtry techniques one can estimate the intrinsic dimension of a dataset. For example, the intrinsic dimension of the Swiss roll in Figure 2.2 is 2.

The techniques we will use in this section were developed by Dr Iolo Jones [3], and the code is available at GITHUB LINK.

By first reducing the dimensions to 5 and calculating dimension estimates we get a dimension estimate at each point. Plotting the reduced dimension graph we can plot a coloured graph inticating the dimension at each point, as shown in Fig...

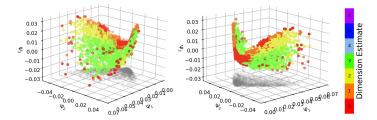


Figure 3.3: Cap

Figure 3.3 suggests that when we reduce the dimension of the ambient space to 5, our data primarily lives on a 3 dimensional space.

This could suggest that our diffusion maps are picking up three significant parameters from the images. More work is required to discover what these parameters may be, perhaps it is glasses, hats etc.

We also see clusters of different colours arise. This could perhaps mean that certain dimensions are not relevent in certain groupings. An analogy would be that if one parameter we are picking up is the length of someones hair, this parameter would be irrelevent in the images of people with bald hair. Again, more research is required to discover what the actual reason for these clusters is.

## 3.4 Estimating the Curvature of the CelebA Dataset

In Section 3.3 we discovered that when we reduce the dimension of the ambient space to five, our dataset lives in a 3-dimensional subset of this. We use Iolo's tools to compute the scalar curvature of the dataset in 5 dimensions, assuming it lives on a 3-dimensional subset.

Note 3.1. We cannot estimate the curvature of the dataset in the original high dimensional space because using diffusion geometry for geometric inference is only robust when the intrinsic dimension of the data is low [3].

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**Note 3.2.** The scalar curvature estimates shown in Figure 3.4 are calculated in five dimensions, despite the figure being represented in three dimensions.

Figure 3.4 suggests the CelebA dataset has negative curvature at almost all of

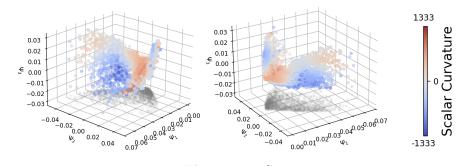


Figure 3.4: Cap

its points. This agrees with the hypothesis that the CelebA dataset has intrinsic hyperbolic geometry.

However, it is important to note that by reducing the dataset to five dimensions we are losing a large amount of information, as it is likely that the instrinsic dimension of CelebA is much larger than five.

Hence, whilst these results are a promising addition to the evidence that CelebA has hyperbolic geometry, they do not confirm this.

## 4 Conclusions

We have employed diffusion maps to reduce the dimension of the CelebA dataset and form a 3-dimensional plot. We can infer from our plot that diffusion maps successfully uncover some of the intrinsic geomtry of the CelebA dataset.

Using recently developed diffusion geometry tools we have attempted to estimate the dimension...

Using novel diffusion geometry tools we have found evidence supporting the original hypothesis of the paper, that the CelebA dataset has intrinsic hyperbolic geometry.

If more concrete evidence is found to further support this hypothesis, this ramifications in the field of

What could a future student do? We don't know what the 3 parameters we are uncovering are. We could try to work this out. Is it glasses is it hats?

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