

Comvi - Comparative Visualization of Molecular Surfaces using Similarity-based Clustering

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Abstract— The goal of this paper is to show the reader the abstract methods and concrete applications that were used to extract and compare features and rank the similarity of the molecular protein maps. Further we present a new method of how the won data can be visualized on high resolution and large displays with a The paper describes the process and the approaches that were taken to solve this task.

Index Terms—Clustering, Similarity, feature extraction, Visualization, high-resolution display, Powerwall, MegaMol, VISUS

1 INTRODUCTION

Helvetica

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Figure 1: lorem ipsum rasterized plus a longish caption spanning two lines

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2 RELATED WORK

TODO: cite kolezar for clustering

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2.1 Initial Challenges and encountered problems

It is clear that the task required from us that we learn how to compare the images, measure the distances between the images, and cluster these images. The given task required that

We could chose to find similarities and cluster the proteins in the .pdb format or given as bitmap image generated by **TODO: cite megamol protein image**

Table 1: lorem ipsum tabulated

dataset	full performance (fps)	half performance (ms)
balls	1,243	0.1
buckets	23	23
bolts	23,312,134.3	22.1

2.2 Approaches to the Clustering-Problem

Right of the start we had several ideas of how we could approach this problem. With the recent trend in machine learning we had a couple of ideas of how we could determine a similarity metric between two images or classify an image into a more usable vector of data.

We ended up using a higher dimensional feature vector ... to determine the similarity between two protein maps because we didnt manage to train a custom model in the given timeframe, due to

But our initial results with a pretrained Imagenet **TODO: cite imagenet TODO: cite darknet publication** model let us to believe that it should be definetly possible for this specified task to find a machine learning solution.

2.3 Approaches to the Visualization-Task

Our approaches to visualizing the given clusters were the following, the reader is reminded that we are not just visualizing the clusters on a "normal machine" but rather the POWERWALL, a projected display with effectively 6-24 times the resolution of a consumer grade display. Details on the POWERWALL can be found **TODO: cite powerwall publicatoin here if available**.

2.3.1 Approaches to the Interaction with the Powerwall
TODO: cite correct pub and use correct name

2.4 Finding a feature vector to cluster the images

2.5 Finding the best performing similarity measure

2.6 Finding the best performing clustering algorithm

2.7 Finding the

2.8 Discussion of results

2.8.1 Dolor

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