ADENINE — A Data Exploration plpeline

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Abstract

Abstract here.

Keywords: Exploratory data analysis, unsupervised learning, dimensionality reduction, clustering

1. Introduction

2. Implementation

From an algorithmic standpoint, adenine is built upon the concept of *pipeline*, that is a sequence of four fundamental steps: (i) missing values imputing, (ii) data preprocessing, (iii) dimensionality reduction and (iv) clustering. For each step, a fair number of off-the-shelf algorithms implementations are available (see Table 2). None of such steps are mandatory. The vast majority of the algorithm implementation is inherited from the machine learning Python library scikit-learn (Pedregosa et al., 2011; Buitinck et al., 2013). {except KNN imputing}.

3. Experiments and results

To assess the quality of the obtained results, we tested adenine on a set of synthetic and real dataset.

{parla qui dei test synth} {TGCA}

4. Conclusions

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Table 1: Pipelines building blocks and relative references (which are not reported when the definition is given in Section 2).

Step	Algorithms	Ref.
Imputing	Mean Median KNN	(Troyanskaya et al., 2001)
Preprocessing	Recentering Standardize Normalize MinMax	
Dimensionality reduction	Principal component Analysis (PCA) Incremental PCA Randomized PCA Kernel PCA Isomap Locally linear embedding Spectral embedding Multidimensional scaling t-Distributed Stochastic Neighbor Embedding (t-SNE)	(Jolliffe, 2002) (Ross et al., 2008) (Halko et al., 2011) (Schölkopf et al., 1997) (Tenenbaum et al., 2000) (Roweis and Saul, 2000) (Ng et al., 2002) (Borg and Groenen, 2005) (Van der Maaten and Hinton, 2008)
Clustering	K-means Affinity propagation Mean Shift Spectral Hierarchical	(Bishop, 2006) (Frey and Dueck, 2007) (Comaniciu and Meer, 2002) (Shi and Malik, 2000) (Friedman et al., 2001)

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