

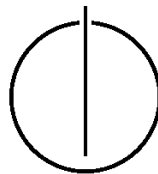
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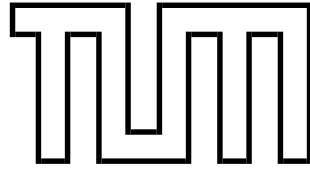
DER TECHNISCHEN UNIVERSITÄT MÜNCHEN

Master's Thesis in Robotics, Cognition, Intelligence

Online Activity Recognition from skeletal features through Kernel Methods

Evgeni Pavlidis





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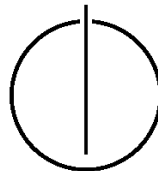
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Online Activity Recognition from skeletal features
through Kernel Methods

Echtzeit Aktivitätserkennung durch Skelettmerkmale
mittels Kernelmethoden

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Date: October 8, 2014



Abstract

English abstract

Zusammenfassung

German Abstract

Ich versichere, dass ich diese Masterarbeit selbständig verfasst und nur die angegebenen Quellen und Hilfsmittel verwendet habe.

München, den October 8, 2014

Evgeni Pavlidis

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1 Introduction

1.1 Motivation

Activity recognition is a big research field in machine learning and robotics. Being able to infer what human actors are doing helps in many practical robotic tasks. An example is doing short-time prediction for collision-avoidance. Moreover in social robotics it is crucial to know what humans are doing when reasoning about the current state of the robot's environment.

To do activity recognition the human pose has to be inferred for each frame. With the advent and further development of RGBD (color and depth) sensors it is now possible to perform skeletal tracking of persons. This allows us to decouple pose estimation and activity learning, which make the problem a bit easier.

For real robotic tasks it is very important that the activity recognition is online i.e. runs in real-time.

Advantages of skeletal features are that no person sensitive information has to be processed by the learning algorithm. In the context of the SPENCER project this is an important prerequisite. Another advantage is also that the skeletal features are very informative for activity recognition. Also beginning to concept an algorithm that builds on top of robust pose estimation reduces the complexity, as we can fully ignore the pose estimation problem.

1.1.1 The SPENCER project

With the modern technologies (ms Kinect SDK, primesense ...) it is possible to decouple skeleton tracking and learning

1.2 Problem statement

cite:software packages and tools used

cite:datasets (mocap, daily activities, ms activities)

Check bibliography style and data!!!

define simple variables mathematically eg. feature sequence etc.

make a distinction between action and activity

make a distinction between online recognition and online learning !!! maybe change online to real-time

The biggest challenge in activity recognition is classification of time series data. In contrast to the simpler sample model, here we have to classify sequences. Therefore appropriate models have to be implemented that take the dynamics into account.

On top of that the problem is ill-posed.

Also for online activity recognition we have to classify incomplete data consisting of subsequences. This makes the problem more challenging and reduces the pool of methods that can be used for the classification.

We identify three requirements for a practical activity recognition system:

1. Online recognition

The algorithm has to be able to do real-time classification as the sequence is progressing.

2. Classification of incomplete sequences

This follows directly from the first requirement. There should be no assumptions about the activities regarding completeness, length or periodicity.

3. Novelty detection

The algorithm should be able to recognize unobserved activities.

elaborate
on this

Optimally it should be possible to also recognize when an unknown activity started and finished. This way automatic segmentation will be possible and will considerably reduce the supervised learning time. In combination with active learning this will greatly reduce training time for practical applications.

Nice to have... online learning??

1.2.1 Online learning of human activities

Use Gaussian processes to learn new activities in real time

1.2.2 Evaluate Gaussian processes against different ml algorithms for activity recognition

Evaluate the performance of GPs in relation to the other solutions

1.2.3 Scope

1.3 Prerequisites and notation

We assume a basic understanding in *Linear Algebra* and *Probability theory*. Although a high-level overview on machine learning is given in chapter 2 deeper knowledge in this field will help understand the *Related Works* chapter better.

1.3.1 Mathematical notation

- Matrices uppercase
- Vectors lowercase bold
- Constants lowercase
- Parameters lowercase greek letters

1.4 Outline

Introduction This chapter introduced the topic of this work. The motivation and the scope is explained.

Background The second chapter summarizes some basic concepts and models that are prerequisites for our approach. It begins with an overview of machine learning and introduces the multivariate Gaussian distribution. Then an emphasis is led on Gaussian Process Regression and Gaussian Process - Latent Variable Models, which is an unsupervised learning method for dimensionality reduction. Last the Dynamic Time Warping algorithm, which is used for sequence alignment, is explained.

Related Work The third chapter gives an overview of methods used in similar approaches and then analyses strength and weaknesses of these methods in regards to online activity recognition.

Approach The fourth chapter presents two approaches to online activity recognition and their implementations. The first one is an implementation of "Discriminative Sequence Back-constrained {GP}-{LVM} for {MOCAP} based Action Recognition" [1]. The second one is a novel approach which learns a dense motion flow field in latent space through Gaussian Process Regression.

Evaluation In the fifth chapter the two approaches are being evaluated and discussed.

Results and Outlook The last chapter summarizes the results of the two approaches and gives a brief outlook of future improvements.

2 Background

This chapter introduces some basic concepts needed to understand the proposed approaches. First a high-level overview is given on machine learning and its terminology. Then the Kernel function is explained along with the *Support Vector Machine* - a kernelized learning method. Following is an explanation of *Gaussian Processes*, their different interpretations and properties. After that the *Gaussian Process - Latent Variable Model* is being introduced along with some extensions for learning a backward mapping and optimizing it for discrimination in the case of multiple classes. Last two *Sequence similarities measures* are presented which are used in our implementations.

2.1 Machine Learning

Machine Learning is a discipline where one makes inference on real world data.

2.1.1 Supervised learning

Supervised learning is the task of classification or regression when the data is labeled i.e. we have the ground truth of every sample. The algorithm then takes the labeled samples (and maybe some confidence values) and infers the model parameters (or hyperparameters) accordingly.

There are two distinct cases in supervised learning:

1. **Classification**

Classification is the task of learning which category a sample belongs to. A prominent example is Spam filtering. By taking a large number of emails which are labeled either as spam or as ham (regular email), the algorithm deduces a model which can classify unknown samples into these two categories.

2. **Regression**

Regression is a terminus in machine learning and means function approximation. Here the domain of the sample's label is continuous. An example would be ...

In most cases we search for a good model that explains the data we have. Parametric models, for example, try to learn the ... When searching for an appropriate model it is also important that we try to capture the underlying relationship without compromising the generalization property, which is the ability of the model to

correctly predict unseen samples. The case that an algorithm learns the relationship of the data that is used to train the model (training data) but poorly predicts new samples is called overfitting.

Very often the parameter search is done by maximizing the probability of the data given the model parameters.

$$\arg \max_{\theta} p(\mathbf{X}|\theta) = \arg \max_{\theta} \frac{p(\theta|\mathbf{X})p(\mathbf{X})}{p(\theta)}$$

where θ are the model parameters and \mathbf{X} is the data.

2.1.2 Unsupervised learning

In contrast to supervised learning in unsupervised learning we have no labeled data i.e. there is no supervisor giving each sample a category (classification) or a value (regression). In this case we can only derive properties of the generation process. Therefore we try to detect patterns in the unlabeled data. These pattern may be clusters of similarity or a lower dimensional generative manifold from which the samples are generated. The last one is called *Dimensionality Reduction* which will be also a subject in this work. [2]

K-means algorithm

An example of an unsupervised learning method for finding a given number of clusters k in given data is the *k-means* method. The idea is that we first determine the number of clusters and choose k points randomly in the space, which represent a guess of the cluster means (center of mass). After that we try to move these points, such that they align with the real data's k centers of mass. This is done by iterating between two steps:

1. Assign each point x to the closest centroid (cluster mean)
2. Find new centroids by computing the mean of all assigned points for each cluster k

Doing so it is guaranteed that the algorithm will converge, although it could be a local minimum.

[2]

2.1.3 Generative models

generative
model ex-
ample

Generative methods model the underlying process which generates the data. In Bayesian terms we model the likelihood and the. Thus more data is needed to find an appropriate model. On the other side the model is very flexible and many attributes have a natural interpretation. An example of this is

2.1.4 Discriminative models

A discriminative model is only concerned with modeling the actual posterior. This way fewer samples are needed to find an appropriate model. On the other hand by not taking the prior into account the model's ability to generalize unseen data is worse. For this reason discriminative methods are more susceptible to overfitting.

2.1.5 Online learning

Algorithms which can be gradually optimized towards a good solution using streaming batches of samples are considered to do online learning. In contrast to online learning online recognition means that the algorithm works in real-time and fast recognition is possible.

2.1.6 Active learning

Very often the bottleneck of powerful supervised learning techniques is that they rely on a large number of correctly labeled data. Since labeling has to be performed by a human it is very difficult and costly to label large amount of data. By identifying more important samples by their information ability of selecting a good model, it is possible to achieve good results with fewer samples. Letting the algorithm select such samples and query only their labels from a human, who is now actively participating in the learning loop, is called active learning.

Active learning is in practice a convenient way to acquire new informative samples without letting someone go over a huge amount of data to label.

2.2 Kernel methods

Many machine learning algorithms work not with the features directly but instead use only the dot product between features. The dot product between two vectors can be seen as a measure of similarity.

2.2.1 A space defined by sample similarity

Suppose we have n sample points x_i of dimensionality d : $x_i \in \mathcal{R}^d$. When extracting features we try to capture the most characteristic properties of the data for each sample. Let us say that we want to extract m features. Then we have a vector $z_i \in \mathcal{R}^m$ which represents each sample. This means that learning is done in a feature space of dimensionality m . Another space, where we can reason about the data is a similarity space. Suppose we have a function $k(x, y)$ which measures the similarity between point x and point y , then we can define a vector s of similarities for a new point

This similarity measure is also called a *kernel function*. We can also define some properties of the kernel function resulting of the informal introduction as a similarity measure.

2.2.2 The Kernel trick

A kernel defines a similarity measure between two points \mathbf{x} and \mathbf{y} . The kernel function can be defined as the dot product between two feature vectors.

$$k(\mathbf{x}, \mathbf{y}) = \phi(\mathbf{x})^T \phi(\mathbf{y})$$

where $\phi(\mathbf{x})$ is a mapping from the input space (raw data) to a feature space.

If a machine learning algorithm is formulated only in terms of the dot product of two feature vectors, it this term can be exchanged with a kernel. As the kernel defines the feature space, we can work in feature space which are high- and even infinite-dimensional. This is called the kernel trick.

2.2.3 The Radial Basis Function

2.2.4 Support Vector Machines

Suppose we have data which is linearly separable. If we have only two features we can draw all samples in a 2D plot. This is shown in Figure 2.1. In this case the *best* line that can separate both classes should be as far apart from all samples as possible. This line can be defined by the samples that are nearest to it. These samples are called support vectors as they are sufficient to span the boundary. For this reason *SVM* is also called a sparse method as one only needs the support vectors to define the classification boundary. For higher dimensional feature spaces the same idea holds, but instead of having a line we have a plane (a hyperplane) which dissects the space in two parts. As the *SVM* models the boundary between each class without considering any generative process it is a discriminative model.

The assumption that the data is linearly separable can be relaxed in two ways:

Instead of finding a boundary in the feature space we can use the kernel trick to project the data into a kernel space. This way the data may not be linearly separable in the feature space, but instead could be linearly separated in some kernel space. If we take the *Radial Basis Function* for example the kernel space has infinite dimensions and thus the data can be linearly separated.

We can also allow for a small subset of samples to cross the boundary without compromising its discriminative properties. This is called the *soft-margin SVM*.

The theory behind *SVM* and the fact that the support vectors can be found by optimizing a convex function make this method a very robust way to do classification. For this reason there are multiple implementations of *SVMs* which are very popular and are used very often in practical applications.

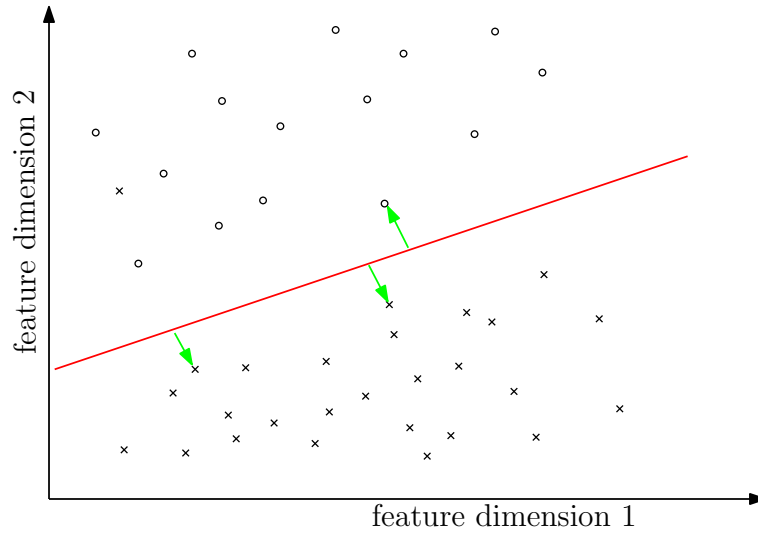


Figure 2.1: SVM decision boundary (red) between two classes (cross, circle). The support vectors are indicated in green.

2.3 Gaussian Processes

Consider the multivariate Gaussian distribution above. If we want to model the distribution of a discrete function defined over a finite interval, we can treat each element of the vector x as a point of the function. Thus we can view the multivariate Gaussian distribution as a probability density function over the function space. Letting the dimensionality d go to infinity (the distance between each point goes to zero) we can model continuous functions.

In this case the mean is a point in function space, thus a function $E[x] = f(x)$. And because of the fact that we now have infinite dimensions the covariance can be seen as an "infinite matrix/", thus a function of two elements: $Cov(x, y)$. This can be also seen as a kernel as discussed in Kernel Methods. Therefore it can be seen as a Gaussian distribution over function space.[14]

The marginalization property is what makes Gaussian Processes feasible as it lets us compute likelihoods with a finite part of the covariance function – which can be seen as a covariance matrix.

A Gaussian process can be also seen as the bayesean posterior consisting of the product of the (Gaussian) functional prior and the observed samples. Another view is a kernelized regression with infinite parameters. [14]

A Gaussian process is a non-parametric model and is governed by the hyperparameters of the used kernel. This also means that the model is less prone to overfitting which is an important property as it not needed to perform cross validation.

2.3.1 The Gaussian distribution

1. **Univariate Gaussian distribution** In the one dimensional case the Gaussian distribution is well known and understood. Moreover many processes in nature can be modeled with this distribution and for this reason it is also called the Normal distribution. The probability of an event is very high on a certain "point" (its mean value μ) and it drops quickly on each side with the standard deviation σ .

$$\mathcal{N}(\mu, \sigma^2) = \frac{1}{\sigma\sqrt{2\pi}} e^{-\frac{x-\mu}{2\sigma^2}}$$

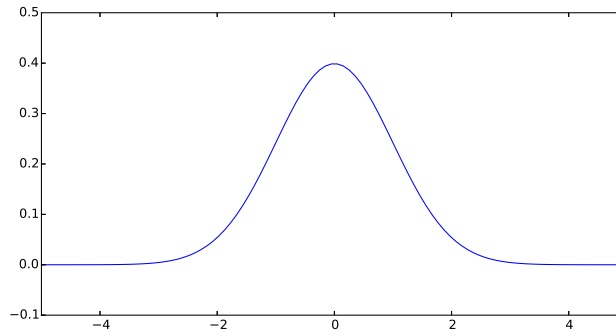


Figure 2.2: The univariate Gaussian distribution with mean $\mu = 0$ and variance $\sigma^2 = 1$

As you can see in Figure 2.2

One disadvantage of this distribution which we can see from the above formula is that it can model only one hypothesis. This is also the case for the Gaussian distributions of multiple (multivariate Gaussian distribution) and infinite (Gaussian process) dimensions.

2. **Multivariate Gaussian distribution** The multivariate Gaussian distribution is the generalization of the Gaussian distribution in higher dimensions.

$$\mathcal{N}(\boldsymbol{\mu}, \boldsymbol{\Sigma}) = \frac{1}{\sqrt{(2\pi)^d |\boldsymbol{\Sigma}|}} e^{-\frac{1}{2}(\mathbf{x}-\boldsymbol{\mu})^T \boldsymbol{\Sigma}^{-1}(\mathbf{x}-\boldsymbol{\mu})}$$

The two parameters of the distribution are:

mean $\boldsymbol{\mu} = E[\mathbf{x}]$ Representing the most probable vector

covariance $\boldsymbol{\Sigma}$ Representing the mutual variance for each pair of the elements of the random vector: $\boldsymbol{\Sigma}_{ij} = Cov[x_i, x_j]$

The exponent is mahalanobis distance, which measures the distance of a point to the ellipsoid defined by the covariance matrix.

3. Properties of Gaussian distributions Aside for being an appropriate model for many processes occurring in nature, Gaussian distributions are also very nice to work with. The marginal and conditional of two Gaussian distributions are also Gaussian.

One reason GPs are straightforward and work is the math behind them. It is just linear algebra operations.

Linear maps for Gaussian distributions:

Product of two multivariate Gaussian distributions:

$$\mathcal{N}(\mathbf{x}; \boldsymbol{\mu}_x, \boldsymbol{\Sigma}_x) \mathcal{N}(\mathbf{y}; \boldsymbol{\mu}_y, \boldsymbol{\Sigma}_y) = \frac{1}{\sqrt{(2\pi)^d |\boldsymbol{\Sigma}|}} e^{-\frac{1}{2}(\mathbf{x}-\boldsymbol{\mu}_x)^T \boldsymbol{\Sigma}^{-1}(\mathbf{x}-\boldsymbol{\mu}_x)} \frac{1}{\sqrt{(2\pi)^d |\boldsymbol{\Sigma}|}} e^{-\frac{1}{2}(\mathbf{x}-\boldsymbol{\mu}_y)^T \boldsymbol{\Sigma}^{-1}(\mathbf{x}-\boldsymbol{\mu}_y)}$$

Marginal of a multivariate Gaussian:

Conditional of a multivariate Gaussian:

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2.3.2 Kernels

The most used kernel when using Gaussian Process is the *Radial Basis Function (RBF)*.

1. Effects of the hyper-parameters

2.3.3 Regression

With *Gaussian Processes* we don't learn a model, but instead we have a probability over infinitely many models with the mean being the most probable one.

2.3.4 Learning


In the case of a GP the learning phase is different than in parametric models, where the model parameters are inferred from the data. Training in the case of GPs means finding good hyperparameter for the kernel, by reducing the log-likelihood by variational optimization (data fit term + cov. regularizer)

In contrast to parametric models Gaussian processes are less prone to overfitting because of the covariance regularizer term.

$$E(\theta) = \frac{1}{2} \log(K) - \frac{y^T K^{-1} y}{2}$$

We see that we have to invert the covariance matrix, which is of dimensions $n \times n$. Therefore this operation has a runtime complexity of $\mathcal{O}(n) = n^3$ which is also the bottleneck of the whole algorithm and a serious drawback of Gaussian Processes.

2.3.5 Classification

 Classifying with GPs is a little more involved, because of the discriminative function and the fact that the likelihood explain problems of GP classification right} is not a Gaussian. For this reason different models exist which try to approximate this likelihood.

2.3.6 Advantages

1. non parametric When using a parametric model one has to make sure that the chosen model is sufficiently complex to fit the data but at the same time is not too complex that it will overfit the training data. This is a very hard task and is in most cases done through cross-validation of the model with an independent validation set. As discussed above GPs are less prone to overfitting and therefore we do not need to reduce the training data to create a validation set.
2. probabilistic Being a model which has a Bayesian interpretation GP The hyper-parameters can be interpreted. The lengthscale controls how much neighboring points contribute to the covariance of the function.
3. generative
4. nice for Bayesian
5. linear algebra operations (marginals and conditionals)

2.3.7 Disadvantages

1. susceptible to outliers One big problem of the Gaussian distribution is that it has the assumption that the noise is Gaussian. When this assumption does not hold and we have several an outlier it either shift the mean un-proportionally to itself or raise the variance. Both cases are The student-t distribution, for exmaple, is robust against outliers but is much harder to deal with.
2. Unimodal Since the Gaussian distribution is concave it can model only one hypothesis. This a curse but also a blessing since the math behind it is simple and unambiguous.

3. high computational complexity $\mathcal{O}(n^3)$
4. non-convex optimization of the hyper-parameters

2.3.8 Sparse Methods

As the computation cost for inverting the covariance matrix is cubic, there are some methods which approximate the solution. One of these methods is the /Informative Vector Machine/[9] where a subset of samples is selected by maximum entropy. This way this active set can explain the data rest of the data, and using it will still result in a good model.

This reduce the complexity to $\mathcal{O}(d^2n)$ where d is the number of chosen samples. There is also an *IVM* method which works for multiple classes.[15]

2.4 Gaussian Process - Latent Variable Model

The GP-LVM is an unsupervised learning model to perform a non-linear dimensionality reduction from an observed space X to a latent space Y . It does this by maximizing the likelihood

$$p(Y|X) = p(Y|f)p(f|X)$$

using a Gaussian prior for the mapping f . Technically a GP-LVM is a product of Gaussian Processes which model a regression of the mapping from latent space to observed space. This means also that if we want to compute the latent position of a new observed sample we have to compute the Using a linear kernel the model generalizes to PCA. By using a non linear kernel a non-linear mapping is inferred making it a non-linear latent variable model.[8]

formulas
etc.
elaborate
GP-LVM
PCA

Analogy LVM j-ç marionettes

2.4.1 Dual Probabalistic PCA

Tipping and Bishop, Journal of the Royal Statistical Society (1999)

Assuming X has a Gaussian prior $P(X) = \mathcal{N}(X|0, I)$

$$P(Y|X) = \prod_{j=1}^p \mathcal{N}(\dagger_j | \iota, \mathcal{K})$$

Where $K = XX^T + \sigma^2 I$ is the covariance matrix. Lawrence noted that this can be interpreted as a product of gaussian processes where the covariance matrix represents

the linear kernel [8]. By exchanging the linear kernel with a non-linear one, we automatically have a technique for non-linear dimensionality reduction.

Using the trace properties

$$\text{tr}(a) = a, \text{ } a \text{ is a scalar}$$

$$\text{tr}(AB) = \text{tr}(BA)$$

cite some
source for
this

we can change the mahalanobis distance term $\mathbf{x}_i^T \mathbf{K}^{-1} \mathbf{x}_i = \text{tr}(\mathbf{x}_i^T \mathbf{K}^{-1} \mathbf{x}_i) = \text{tr}(\mathbf{K}^{-1} \mathbf{x}_i \mathbf{x}_i^T)$

The log likelihood is:

As there is no linear kernel any more this equation cannot be solved in closed form. Therefore we have to do gradient based optimization. But marginalizing over all latent space samples means that we have to include these in the optimization. This fact makes the optimization problem very hard as the dimensionality is high-dimensional – number of samples n + hyperparameters – and has many local minima.

As initially proposed the standard way of initializing the latent space is using *Principal Components Analysis*.

2.4.2 Back-constraints GP-LVM

One problem with this model is that it does not preserve local distances in the latent space. This is because it tries to explain the data by moving distant samples from the observed space also far apart in the latent space. This problem is addressed by Lawrence et al. in the back-constrained GP-LVM [10]. A mapping $g_i(y_i) = x_i$ is introduced which constrains the points in latent space to be more near if they are also near in observed space. Instead of optimizing directly on X the back-constrained GP-LVM optimizes the mapping $X = f(Y)$ instead. This back-constrained mapping

Having this back-constraints also gives us a mapping from observed space to latent space which can be used to project a new sample into the latent space without costly maximum likelihood estimates.

2.4.3 Discriminative GP-LVM

Another improvement in the context of classification in latent space is the Discriminative GP-LVM [21]. Using a *General Discriminant Analysis* criterion a prior is being enforced on the latent space which ensures that samples from one class are more clustered and different classes are more separated. This is done by maximizing the between-class separability and minimizing the within-class variability while optimizing the log likelihood of the GP-LVM.[21]

2.4.4 Other variants

1. Bayesian GP-LVM An interesting approach for computing the likelihood of the latent variable mapping was proposed in [19]. By using a variational method it becomes possible to marginalize over X . Doing so the mapping can be learned together with an ARD kernel. This way the dimensionality of the manifold can be learned from the data.

explain
ARD

2. Subspace GP-LVM

3. Manifold Relevance Determination Combining the Subspace GP-LVM with the variational approach and the ARD kernel it is possible to learn the manifold .[4]

explain
MRD

2.4.5 Advantages

1. interpolation Because of its probabilistic nature GP-LVM interpolation between two data sample is very natural. [13]
2. probabalistic
3. Generative: it can generalize beyond training data
4. non-linear mapping

2.4.6 Disadvantages

1. No mapping from observation space to latent space The idea of the GP-LVM is to learn a mapping from latent space to observation space by marginalization over the latent space. Resulting from this is that we do not have an inverse mapping into the latent space. This fact may be of no importance for character modeling and motion interpolation but in our case it is crucial. An inverse mapping can be computed by using the Back-constrained GP-LVM described above. However one should also keep in mind that using back-constraints inherently changes the latent space as employs an additional constraint on the mapping.
2. Very hard optimization problem Resulting from the disadvantages of Gaussian Process regarding the optimization of the hyper-parameters the GP-LVM is also very hard to optimize as its objective function is non-convex. But in the case of GP-LVM we have a much larger optimization space due to the fact the we do not optimize only the hyper-parameters, of the mapping Gaussian Process, but also the latent space itself which is of dimensionality n .

This in fact is the biggest problem as it limits its use on real world data, because for more complex manifold structures there will likely be many local minima. For

this reason it is crucial to choose a good initialization. Examples are PCA, Local Linear Embedding or ISOMAP.

2.5 Sequence similarity measures

2.5.1 Longest Common Subsequence

2.5.2 Dynamic Time Warping

The *Dynamic Time Warping* is an algorithm which tries to find a minimal path between two sequences where the path can be warped in the time dimension. The sequences can be of arbitrary length.

The recursive definition – excluding some corner cases – reveals the workings of this method.

$$\text{dtw}_{x,y}(i, j) = \text{dist}(x_i, y_j) + \min \begin{cases} \text{dtw}_{x,y}(i-1, j) \\ \text{dtw}_{x,y}(i, j-1) \\ \text{dtw}_{x,y}(i-1, j-1) \end{cases}$$

Where $\text{dist}(x, y)$ is a distance function which tells how close two points are, and i and j are the element indices for the first and second sequence. The DTW can be computed with dynamic programming and has a runtime complexity of $\mathcal{O}(nm)$ where n, m are the lengths of the two sequences.

It is closely relates to the *Longest Common Subsequence* where, but instead of maximizing a common subsequence, we minimizing the total warping cost between both sequences.

Since we are not interested in the path itself but in the cost of the minimal path we define the DTW as a mapping from two time series to an real value. We consider DTW to be a distance which is not entirely correct as the triangle inequality does not hold. Nevertheless it gives us a notion of how similar two time series are and since it is non-negative ($d(x, y) \geq 0$), symmetric ($d(x, y) = d(y, x)$) and respects the identity property ($d(x, x) = 0$) it can be used to define a meaningful, be it not formally correct, kernel. [16]

3 Related work

This chapter will introduce some models and their corresponding algorithms for activity recognition. An emphasis is led on methods which work with skeleton data. In the last part a short analysis is done on these methods and some observations are discussed.

3.1 Overview

Activity recognition is a difficult task as we have to make sure our algorithm will discriminate between different classes – activities – but also will leave room for inner class variations. These variations are the result of different persons performing activities differently. A simple example is walking, where different person has a different walking style – also called gait. Also different environments will result in actions to be performed slightly differently. [11]

There are many methods which learn from videos and try to explain. This approach is very flexible but also has several drawbacks. One of which is that it is very hard to achieve scale and view-invariance. Furthermore inferring the human pose is very difficult and ambiguous.

For these reasons we will consider only data with pose information in this thesis.

3.1.1 machine vision for human activities: a survey [20]

Generative models such as HMM Discriminative models such as CRF

Survey on Time-Series Data for classification

3.2 Histogram based approaches

3.2.1 Motion history image

3.2.2 Motion energy image

3.3 Dynamic time warping

3.4 A class of space-varying parametric motion fields for human activity recognition

3.5 Action Recognition Based on A Bag of 3D Points

action graph - nodes are shared poses

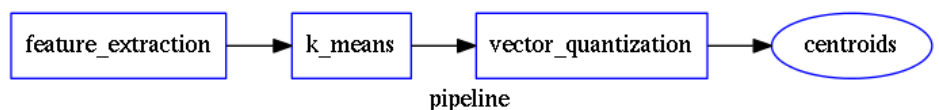
3.6 Methods using skeleton features

3.6.1 Gaussian Mixture Based HMM for Human DailyActivity Recognition Using 3D Skeleton Features

3.6.2 Sung et al. [17]

1. Features: Skeleton data + HOG features of RGBD image and depth image
2. Naive classification: SVM
3. Maximum entropy markov model Solved via max-flow/min-cut

3.6.3 RGB-D Camera-based Daily Living Activity Recognition [27]



1. Bag of Features
See ¹
2. Features: Structural and Spatial motion Feature capturing transition between two frames
3. Bag of Features approach (histogram of features)
4. Other: People identification (reidentification)

¹RGB-D Camera-based Daily Living Activity Recognition - Chenyang Zhang, Student Member, IEEE and Yingli Tian, Senior Member, IEEE

3.6.4 View Invariant Human Action Recognition Using Histograms of 3D Joints

3.6.5 Learning Human Activities and Object Affordances from RGB-D Videos

1. Learning both: activities and object detection/affordance
2. Using Markov Random Field and SVM for learning

3.6.6 Eigenjoints [26]

3.6.7 Gaussian Process - Latent Conditional Random Field (GP-L CFR)

[6] use GP-LVM to reduce dimensionality of human motion. (earlier approach was Gibbs sampling)

3.6.8 Modeling Human Locomotion with Topologically Constrained Latent Variable Models

3.6.9 GPDM

In [24] the dynamics of the latent space is being modeled from time series data. In [25] this model is being used to model human motion by applying a GP-LVM to the high-dimensional mocap data and simultaneously learning the dynamic transition in the latent space:

$$x_{t_{k+1}} = f(x_k)$$

$f(x)$ is being modeled by a Gaussian process.

This model was applied for activity recognition in [5] where the classification is done through an SVM in the hyperparameter space. (only 2? features)

3.6.10 Joint Gait Pose Manifold

The Joint Gait Pose Manifold models the activity and the gait in an common latent space. This way several samples from different persons are modeled with the addition of the gait and do not corrupt the class learning. Each activity is mapped to an toroidal structure where the length represents the activity dynamics and the width represents the gait variation.

3.6.11 Human Action Recognition Using a Temporal Hierarchy of Covariance Descriptors on 3D Joint Locations

3.7 Analysis

Skeleton features are sufficient but other features can be useful:

- hand
- head pose recognition
- situation awareness ...

3.7.1 Observations

cite action
graph

- One observation one can make is that activities are represented by the dynamics of the poses, and thus we try to capture this dynamic model. Several options exist. One way is to use popular graph based probability models, such as Hidden Markov Models, Conditional Random Fields or Action Graphs. Another option is to try to capture the dynamics by appropriate feature extraction.
- Difference between activity and action Activities are composed of actions
- Context information can tremendously help in classification of activities (e.g. object detection and human anticipation)
- Skeleton data is sufficient for classification ([?].????) and also robust to changes in appearance (most state-of-the-art methods work with visual features) and also unobtrusive and sensible data doesn't need to be stored (like face features etc.)
- hierarchical learning: Some methods learn the actions that a activity is composed of. This practice is also very common in HMM models as they model discrete states and their temporal dependencies
- DTW is a good measure but has several drawbacks, such as in cyclic activities where some motions can be repeated several times
- LLE is not generative therefore LL GP-LVM to preserve smooth map also in latent space

3.7.2 Approaches

1. Discriminative Sequence BCGPLVM Use this to find the activity
 - a) DTW between walking and walking backwards very big ...
 - b) not taking temporal dimension into account

2. GPDM

- a) approach to classify by hyperparameters not optimal

3. Classify by dynamics of the skeleton (this should bring good classification)

- a) GPDM can model the dynamics of the movement
- b) has good properties (Gaussian processes)
- c) has intrinsic dim reduction
- d) ?? shared GP-LVM to model different activities in the same latent manifold
??

3.7.3 Problems and solutions

1. limited sample data - probabilistic model + discriminative Probabilistic (and generative ??) models are more accurate using fewer samples, because they model the probability directly ...
2. high dimensional - dim reduction(gp-lvm)
3. classification - BC GP-LVM + discriminative
4. time series data - GPDM An can be modeled as a sequence of consecutive poses. Hence a dynamical model. By using a dynamical model classification becomes more discriminative.
5. confidence is important !!! Using a probabilistic model (especially Gaussian processes) we also get a confidence which in turn can be used for active learning
6. high dim. noise = GP-LVM is very robust because of the nature of optimization (distance is preserved instead of locality)

3.7.4 Assumptions

1. Skeleton tracking is correct and stable For the algorithm we assume that the skeleton extraction from RGBD data works as expected. This is far from the truth with current skeleton tracking algorithms but we also get confidences of the poses. This way we can prune a large number of incorrect poses and because we model the dynamics and do not compare poses this is not a big problem.
2. Smooth skeleton transition !!!
3. Correctly labeled samples (no outliers)

3.7.5 Ideas

1. Use hand and/or head features
 - a) Head direction is important
 - b) Hand structure is very important for most tasks
 - c) Object interrelation ???
 - d) Use HOG for hand features only
2. bag of features
 - no time dependency
 - no online capable because of k-means clustering

3.7.6 GP-LVM for human motion

As the space of human motion is high-dimensional (spatio-temporal) dimensionality reduction is crucial for a number of models dealing with human motion (e.g. [?]). The GP-LVM preserve the distances in the mapping and are therefore suitable to model human motion with high noise of the poses see Urtasun DGPLVM Newest addition is [6]

4 Approach: KMeans clustering approach

In consequence of the discussed problems above and time constraints we choose to implement another model to have a working system. For this we choose an existing algorithm base on *bag-of-features* approach published in 2012 [27].

5 Approach: Discriminative Sequence Back-Constrained GP-LVM

5.1 Feature extraction

Regardless of the chosen algorithm the features used for learning will have a big impact on the performance of the model. Therefore it is imperative to extract discriminative features from the skeleton data.

We get the joint positions and the angles between them in the camera frame defined by the used depth camera (e.g. Microsoft's Kinect). When extracting features we have to make sure that we have view invariant features of the skeleton. We want these data in the frame of the skeleton.

One way to achieve scale invariance is to normalize all link lengths in respect to the torso link. This corrects for variances of skeleton lengths in different persons. To make the pose view invariant we have to define a local skeleton frame which captures the skeleton's *orientation* in the world coordinate system.

Another way to achieve view invariance is to not consider the 3D points of the joints all together but instead to take only relative features. These can be, for example the angles or distances between two adjacent joints. An interesting approach is used in [18], which is to define a polar coordinate frame for each joint and use only two angles, which define the orientation of the joint in a polar coordinate frame, as features. This way we also reduce the observation space.

As discussed in *Related Work* many methods also make the extracted temporal features (e.g. Eigenjoints). However since we want to include the dynamics in our model we do not extract such features explicitly.

We selected a 3D point cloud of the joints in the skeleton's own coordinate frame as features. The reason for this is that we believe the 3D point cloud to be more linear than relative features, which in turn will help when optimizing the model. Figure 5.1 shows this approach. We chose the two vectors – torso to right hip and torso to left hip – to define our local coordinate system. By normalizing and computing the cross product we have also the third vector which points to the walking direction of the skeleton.

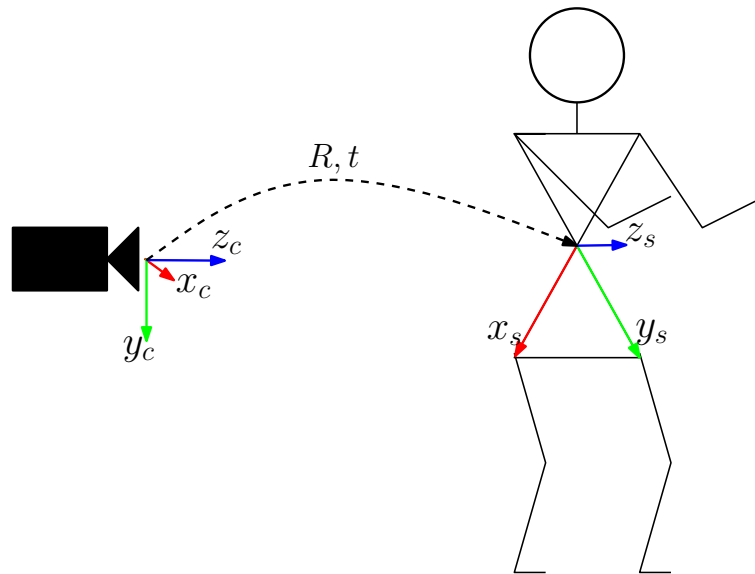


Figure 5.1: Sketch of the local skeleton frame inside the camera frame. The rotation matrix R and the translation vector t define the needed transformation to change from camera coordinates to the local skeleton coordinates

5.2 Dynamic time warping with mahalanobis distance

The Dynamic Time Warping algorithm is a prominent and very effective choice for computing similarity between two sequences. The problem with this approach, in the context of activity recognition, is how to define the distance metric between two poses.

really ???

Popular choices for the distance function is the euclidean distance, if 3D points are used as features, and the geodesic distance for angles. The problem with these two distances is that they are just the sum of the individual feature differences. As the dimensionality grows this metric becomes less informative.

In the case of human poses we have a certain notion of which poses are similar and which are far apart. Maybe this is due to the fact that we inherently know – or classify – to which activity the pose corresponds to and have therefore some notion of closeness with respect to an activity which cannot be approximated with the euclidean distance. Poses from different activities will most likely also seem to be more or less similar depending on how similar the actions are.

One idea to transfer this knowledge is by using the Mahalanobis distance instead of the euclidean distance when computing the similarity of two pose sequences. By computing the covariance for each activity we have some notion of the variance across all feature dimensions for a specific class. This way we can capture – to some extent – the variability for each class. Now we can compute a similarity measure with a new

sequence x_{new} for each class and each sample of this class. Thus we can define a notion of measure between a class and a new sample by:

$$s(j, x_{new}) = \frac{1}{|C_j|} \sum_{x \in C_j} \frac{\text{DTW}_{\text{mahalanobis}(\Sigma_j^{-1})}(x, x_{new})}{\min(|x_i|, |x_{new}|)}$$

where C_j is the set containing all class sequences and $|C_j|$ is the number of sequences in class j . The normalization factor $\min(|x_i|, |x_{new}|)$ makes sure that the minimum cost computed by the DTW is proportional to the smallest sequence.

This way the distance error is distributed by a way defined by the variance across each dimension.

A similar idea was also proposed in the context of handwritten signature verification in [12], which uses just one covariance matrix. The covariance matrix is determined such that, just like in the case of Discriminant GP-LVM, it maximizes the variability between classes and minimizes the difference for samples in the same class. In contrast to our approach the overall covariance matrix may define a more meaningful and discriminative measure but it is also more difficult to update when performing online learning and when learning a new class (novelty detection).

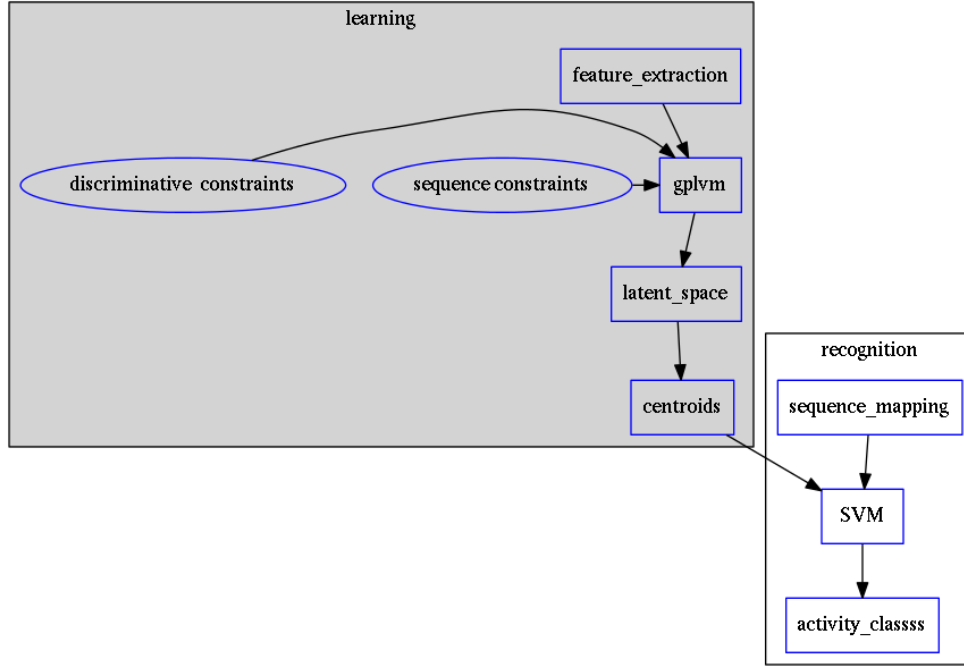
5.2.1 Implementation

We wrote a simple version of the Dynamic Time Warping in Python using dynamic programming and following the recursive definition in chapter 2.5.2. As the variance for some feature dimensions can be zero the constructed covariance matrix does not have full rank and thus cannot be inverted. We mitigate this problem with an approximation of the inverse by computing the pseudoinverse.

5.3 Discriminative Sequence Back-Constrained GP-LVM

At first we concentrated our efforts for learning with the MOCAP data. In theory the data collected data from the kinect should be equivalent. One difference is the high noise in the pose estimation, but due to the fact that the GP-LVM preserves distances rather than locality this problem is mitigated to a certain degree.

In the paper "Discriminative Sequence Back-Constrained GP-LVM for MOCAP Based Action Recognition"[1] the authors propose a method for classifying MOCAP actions.



Sequence back-constrained GP-LVM pipeline ... CITATION

By using a similarity feature for the sequences in the observed space and constraining the optimization to preserve this measure the local distances between the sequences are transferred into the latent space. This has two advantages. First of all the sequences have a meaningful clustering in the latent space. Second by also learning the back-constraint it is possible to calculate the centroid of a sequence in the latent space directly without maximizing a likelihood. This in turn is being used to do real-time classification for actions. The mapping is defined as a linear combination of the DTW distance between every other sequence. For every latent dimension q we have:

$$g_q(Y_s) = \sum_{m=1}^S a_{mq} k(Y_s, Y_m)$$

where the similarity measure is $k(Y_s, Y_m) = \gamma e^{DTW(Y_s, Y_m)}$. This measure is to be preserved in the latent spaces.

$$g_q(Y_s) = \mu_{sq} = \frac{1}{L_s} \sum_{n \in J_s} x_{nq}$$

This constraints are being enforced in the optimization by adding Lagrangians to the objective function.

Furthermore, by applying the Discriminative GP-LVM we ensure that poses of different activities are separated from each other and poses from similar activities are located

closer together. This ensures that the centroid of an activity is more informative and thus discriminative. The Discriminative GP-LVM works by also maximizing the between class variance and minimizing the in-class similarity [21] Also by applying the Discriminative GP-LVM the clustering of similar actions and the distances of different actions is enhanced which allows for a better classification. Recognition is being done by applying the mapping above to the new sequence and using a SVM in the latent space.

expain D
GP-LVM
properly

5.3.1 Advantages

Recognition can be done in real time by using the learned back constrained. The centroid in the latent space is being calculated for the whole sequence and classified by the SVM. Also incomplete trajectories can be classified.

5.3.2 Shortcomings

As the optimization for GP-LVM is determined by the above similarity measure and the discriminative criterion online optimization is very difficult. It is thus highly likely that performing a gradient online optimization will be stuck in an local minimum.

Also one problem with the real-time recognition is that determining when a activity has ended/begun is very difficult. Also as we do not know how long a sequence is we have to calculate the centroid for several time frames.

5.3.3 Implementation

As there was no publicly available source code we choose to re-implement this method. As it was planned to implement a *ROS (Robot Operating System)* module for online activity recognition we choose the Python platform which can be easily integrate with *ROS*. We used the *GP_y* library from the ... Sheffield University . We ported the Discriminative GP-LVM constraints code from Prof. Urtasun and integrated it with *GP_y*. To implement the sequence back-constraints we performed a constrained optimization using Lagrangians.

cite GP_y

5.3.4 Extensions:

1. Learn poselets (pose and velocities) to capture dynamics The GP-LVM learns a mapping for each pose but does not consider velocities and accelerations. If we take a pose along with its first and second moments (let us call them poselets) as the high-dimensional space we allow for the temporal displacements to be also modeled. The latent space represents the poselet and the DTW kernel in the constraint captures also the motion of the activity.

2. Use mahalanobis for the DTW

5.4 Evaluation

As the idea was to implement the algorithm in an language that can be easily integrated into the ROS infrastructure we implemented the model in Python. Unfortunately we were not able to perform an appropriate dimensionality reduction. We believe that the many constraints on the optimization and the highly different data is very hard to optimize. For this reason we choose to implement a new model basing on motion flow fields. [23] [3] [22]

The author in Exploring model selection techniques for nonlinear dimensionality reduction also suggest to use ISOMAP or LLE to initialize the GPLVM and argues that direct optimization of the GP-LVM is very difficult.

Probabilistic Feature Extraction from Multivariate Time Series using Spatio-Temporal Constraints

6 Approach: GP-Latent Motion Flow Field (based on the gp regreesion flow)

It can be argued that the mean poses computed in the *bag-of-features* method capture the most probable motion tendencies of an activity. The good performance of the algorithm can be attributed to this fact.

Many models which use GP-LVM to reduce the high dimensional space into fewer dimension. These approaches make the problem more feasible but the problem remains how to do classification for time-series data. Human motions are mostly characterized by the dynamics of the model (temporal dimension). So we have to compare trajectories in the latent space. One idea is to use GPRF as classification can be done using second order dynamics which should give better results. Going further the activity itself is characterized by the first and second moments of the trajectory function. By explicitly modeling the velocity of the trajectory we can take changes in the joint movement into account.

6.0.1 The Gaussian Process Regression Flow

[7] can be used to model the trajectories in the latent space.

explain
GPRF

6.0.2 GP-Latent Motion Flow

The GP-LMF method is inspired by this model. The difference being that in the case of activity recognition we do not know the starting position and also the trajectories can have significantly different lengths. For this reason it is very difficult to normalize with respect to the time dimension. Nevertheless, resulting from the properties of Gaussian Process regression, we have also a dense mean flow field and dense variances. This allows us perform efficient and robust online recognition in the latent space.

This model is attractive for two reasons. First real-time classification of incomplete trajectories is possible. Incomplete not only in the sense of the first part of an activity but any interval of an activity, which could be also somewhere in the middle of the sequence. Second it is possible to do online learning by simply adding the new class as a new flow field to the pool of GPs. It is very difficult to adjust the other models for online learning, because of the problem that we can get stuck in a local minimum when optimizing the parameters of the GP.

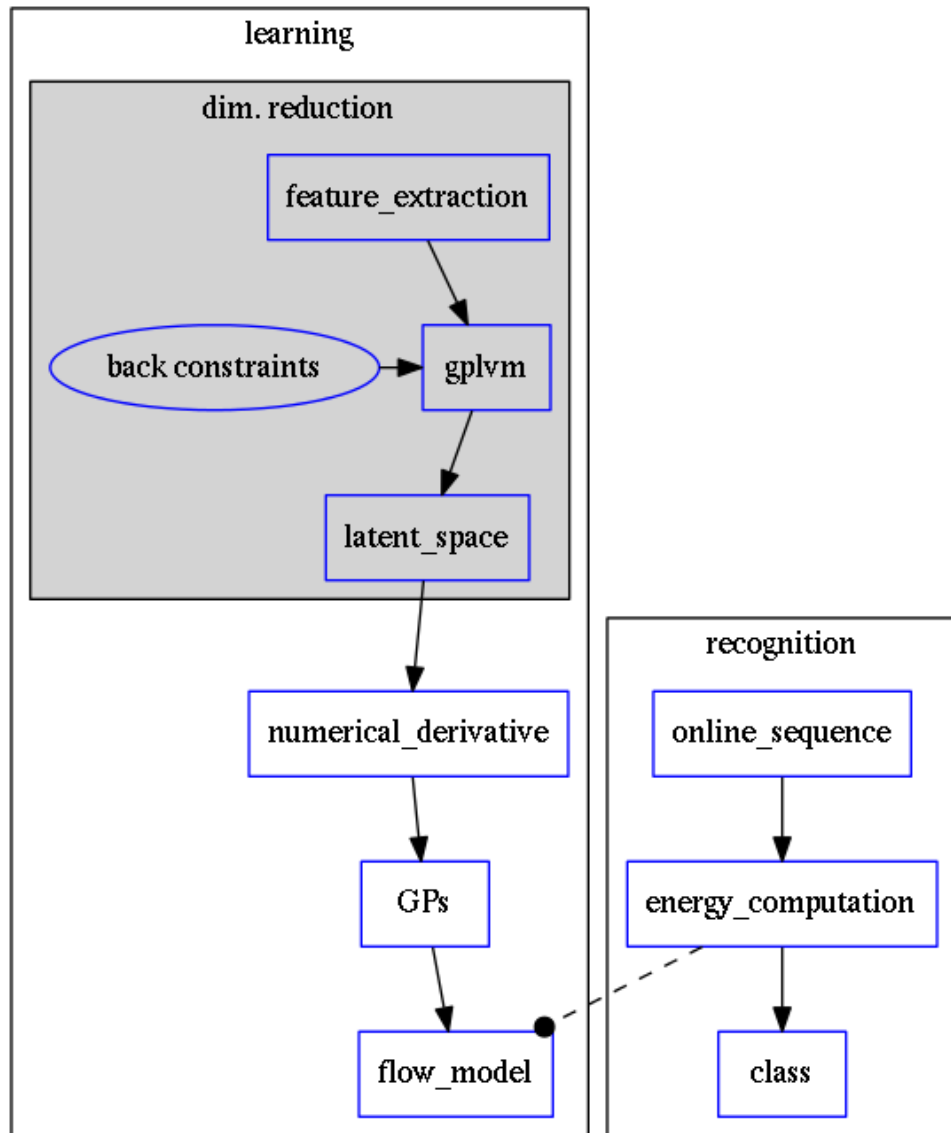
The idea is to learn a motion field in the latent space for each activity. This can be achieved by learning the velocity function of the latent point just like in the GPRF model presented above. With the difference that we do not use the spatio-temporal domain but spatial domain of the latent space. The reason being that we do not have starting and ending positions for each activity and also the lengths can be variable. On top of that we also want to recognize an activity which is being interrupted by another activity, so we can't fix the lengths of the trajectories.

Each activity has its own flow field. Recognition and prediction is done by calculating the energy of the currently moving point with each different field. The field with the minimum energy represents the most probable activity as the point follows more closely its "current" of motion.

Variances in the speed of performing an activity can be modeled by giving the point in the latent space a mass which can be adjusted in real time. When a point has greater mass then it needs more energy to be propagated through the flow field (the overall activity is slower) and vice versa.

This way we have two indicators for recognizing unobserved data. The first one is the variance of the back-constraint mapping. If it is high we know that current sample is far apart from the observed ones. The second is the variance of the *Gaussian Process Regression*. If this value is high we know that we didn't see any sample in the latent space with the current motion. Therefore, with this two indicators, we have a notion of how new a sample and its current motion are.

An advantage of this method is that activities with repetitive motions, such as walking or running, can be learned without using periodic kernels or without resorting to model them explicitly. Repetitive motions can be seen as just multiple samples of the same motion which define the flow field.



Gaussian Process - Latent Motion Flow

6.0.3 Learning the flow field

The initial idea was to learn a general dimensionality reduction for a high number of varying activities and work with only one latent space. The problem is that it is very difficult to learn a smooth mapping in the latent space. This is described more deeply in [23] where the authors try to incorporate the optimization criterion of Locally Linear Embedding together with the a back-constrained Gaussian Process Dynamical Model.

As this approach needs also prior knowledge and is very complex we decided to learn each activity separately. Future work should deal with the possibilities of learning a unified latent space as it will allow us to learn different flow fields in the same space and we will not have to perform a heuristic normalization.

We deploy GP for learning the flow field which gives us several advantages.

One problem we encounter by learning the motion flow field from several samples is complexity of the Gaussian Process. There are two solutions for this. The first one is to use a sparse GP model. The second one is to sample points from all samples and use only those that are most suitable for the regression. If we take IVM as the sparse GP model both approaches can be seen as equivalent as the IVM will automatically take the most informative samples.

6.0.4 Interpretation

The proposed model has a natural interpretation. A point represents a pose in latent space and an activity is a trajectory in time inside the same space. With the flow field we learn the motion tendencies for each pose. When performing recognition we let the current point traverse each separate flow and compute the needed energy. If we consider that the point has a mass we can model the speed at which activities are being done. This way we can recognize when a point leaves an activity, which represents a *motion current*, and passes over to some other activity.

The model captures the changes in velocity which is comparable to the motion history images...

6.0.5 Advantages

1. Recognition The current activity is being mapped into the latent space. Through the learned back-constrained. The recognition is being performed solely in the latent space. By propagating the current position by each flow field we can calculate the next possible pose. By comparing the similarity considering the variances we have a measure of how well the current activity resamples each flow field e.g. learned activity.
2. Prediction If we have detected the activity predicting is simply a matter of propagating the pose through the flow field by taking the mean of the GP.
3. Online learning
4. Natural interpretation
5. Novelty detection (anomaly detection) In [7] the authors present the ability of the GPRF model for anomaly detection. This approach is also suitable for finding new classes as the above energy value can be used to recognize novel activities. The reasoning is that if we cannot find an flow field with a small energy the activity has to be unobserved.

-
6. Active learning
 7. Multiple Hypothesis Prediction Since we have a GP representing our flow field we can predict future point positions with the mean value. Moreover also having informative variances we can sample several possible trajectories. This can be accomplished using an particle filter. Hence we can have multi-hypothesis predictions along with their probabilities.
 8. In comparison to the GPDM it can model cyclic activities

6.0.6 Problems

1. Dimensionality reduction Performing a non-linear dimensionality reduction is no easy task. Testing was done with only two dimensions as it easier to visualize the latent space and the resulting flow fields. A latent space with higher dimension will naturally make the reduction more robust and the field will have a more natural interpretation. . . .
2. Stable class mean flow field When learning a stable flow field from several samples the field can degenerate with the inclusion of strong variable paths. Therefore it is important to ensure that the algorithm learns stable paths. This can be achieved by sampling uniform random sampling from all samples of the same activity.

active
learning
- problem
??

6.0.7 Recognition

Energy minimization:

$$E_{t_1, t_n} = \sum_{i \in \mathcal{T}, j \in \mathcal{T}} E_{i,j} + \text{regularizer}$$

where the regularizer ensures that we do not change flow fields often. . .

7 Conclusions and Outlook

7.1 Summary

7.1.1 Dimensionality reduction for all activities is very difficult (also with extra constraints)

7.1.2 Dynamics is a good measure for classification of human activities

7.1.3 Contributions

1. Advantages and Disadvantages of dimensionality reduction with GP-LVM for human motion in the context of activity recognition
2. Implementation of the Discriminative GP-LVM with python We ported the matlab code provided by Prof. Urtasun into python and integrated it with the GPy library
3. Implementation of the Sequence Back-constraints We used Lagrangians to implement a constrained optimization of the likelihood function
4. Improvement of the DTW measure with the mahalanobis distance ???????
5. A novel approach for activity recognition (prediction??)
6. Introduction of an energy minimization approach for online recognition of complex activities

7.2 Outlook

7.2.1 Energy minimization evaluation

7.2.2 Semi-supervised activity learning by automatic segmentation of activities !!!

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