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題目

Analysis of Sports Movements Based on
Time-Series Pose Data
(時系列姿勢データに基づくスポーツ動作の解析)

主専攻 ソフトウェアサイエンス主専攻

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Abstract

A long standing goal in the field of classification is to develop effective techniques which are applicable to real life problems. With recent advancements in computer vision, significant strides towards this goal has been made due to the increased ability to capture richer and more fine-grained human motion data. For example, it is possible to capture three dimensional motion at a frame rate of 120 frames per second with little error using motion capture technology, as we demonstrate in domain of injury prevention.

In parallel, exponential growth in computer processing power has led to an incredible improvement in machine learning methods. For example there have been multiple breakthroughs in object detection, voice recognition and language modeling etc. Using pose estimators based on deep learning, it is now possible to extract three dimension motion data from video with the need of motion capture. Thus allowing a lightweight and easy-to-use application of motion analysis in environments where the setup cost of motion capture makes the use of it unrealistic.

Despite this progress, there are relatively few studies which effectively couple the advancements of machine learning technologies to human motion analysis. Therefore in this thesis, two studies that demonstrate the use of motion analysis and machine learning is introduced.

In the first study, we search for factors that differentiate the motion between athletes who are at risk of injury using motion capture and ground reaction force data of a commonly used sensorimotor performance indicator known as the single-leg drop jump.

In the second study, we propose an analysis framework for extracting important information from video and numerical data in fencing matches to assist experts in their analysis work.

We show that our approaches can provide useful information to assist those working in the domain of human motion analysis.

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Chapter 1

Analysis of Single-Legged Jumping Motion Based on 3D Time-Series Pose Data

There are a wide range of benefits can be received by progressing research in human motion analysis. One example we find could significantly benefit is the prevention of injury in sport. For professional athletes, sustaining an injury directly leads to an economic loss. For elite young athletes an injury has the potential to squander hopes and dreams of playing as a professional. For societies with healthcare, an injury for an can represent a significant economic cost to society, with studies showing the mean medical cost of a high school varsity athlete was \$709 per injury, \$2223 per injury in human capital costs, and \$10432 per injury in comprehensive costs [1].

Although there are many studies on the topic of injury prevention, these usually are from either a rigorously medical perspective [2] or a analysis made with a focus on machine learning with empirical (big) data [3]. This is a sizeable gap in which we try to bridge.

In this study, we aim to find factors that differentiate the motion between athletes who are at risk of injury. In order to do so, we develop models and techniques that allow us to analyse granular motion data of a well studied sensori-motor control task, the single-leg drop jump (SDJ).

We demonstrate that the methods introduced can be used to to detect anomalous motions. Moreover, we show that by interpreting the decision mechanism that led to the results can provide valuable practical information.

In we make the following contributions,

1. A Method to prevent injuries with computer vision and machine learning Explanation.
2. Novel research that intersects injury prevention, motion capture and machine learning.

1.1 Sensorimotor Performance Indictors

"Sensorimotor" indicates the involvement of both sensory and motor functions. The ability to execute sensorimotor tasks well is crucial for performance in athletic activity. Moreover, there is an abundance of research suggesting the connection between sensorimotor performance and injury. Therefore, great effort is directed to understanding these skills. In this section we will briefly review the literature intersecting sensorimotor performance and injury. In particular we will also introduce a method that involves an action known as the single leg drop jump (SLDJ) which is frequently used to measure sensorimotor performance.

SLDJ is a unilateral horizontal drop jump, in other words the subject hops off an elevated platform of usually 30cms. It has shown to be more reliable, in means of reproducibility, than other sensorimotor performance tasks and is known to show similarities to the bilateral drop jump which is used in many areas including athlete assessment, performance monitoring, talent identification and rehabilitation [4].

1.1.1 Ground Reaction Forces

There are largely two methods of measuring a SLDJ. The first is by measuring the ground reaction forces (GRFs) using a special measurement device known as a force plate. Several indicators such as Time to stabilization (TTS) [5], dynamic postural stability index (DPSI) [6] center of pressure (COP) [5] are then calculated and are assessed. It is important to note that there are multiple variations to calculate the above methods due to the possibility of different sample rate, filter settings or trial length. Such variances can cause a difference on outcome values and may lead to contradictory results [7].

A number of studies show that TTS, DPSI, COP can be used to differentiate participants between chronic ankle instability (CAI), functional ankle instability (FAI), anterior cruciate ligament etc [8]. Yet, to date, the interrelations among TTS, DPSI, and other indicators are largely unknown [6].

1.1.2 Pose Characteristics

The second method of measurement of SDJ is the measurement of pose characteristics using either manual notation or automated motion capture. Captured data is then analysed manually. It has been shown that women have a greater valgus knee angle at time of initial contact than men performing a SLDJ [9]. However, due to the high-dimensional and multivariate nature of this data, there is only a handful of research focusing on the measurement of pose characteristics.

1.2 Subspace Based Classification

Subspace analysis is a term used to describe a general framework used in computer vision, that is used for the comparison and classification of subspaces. In this section, we introduce typical approaches in subspace analysis, in particular the classification of subspaces, and describe how it can be extended to analyse shapes such as human poses.

1.2.1 Shape Subspace

1.2.2 Subspace Method

The Subspace method assumes an input vector \mathbf{x} and k -class subspaces. Each class subspace approximates a data distribution for a single class. This approximation is obtained by applying Principle Component Analysis (PCA) to each class. The similarity S of the input vector \mathbf{x} to the i^{th} class subspace \mathcal{P} is defined based on either:

- The length of the projection of \mathbf{x} to \mathcal{P} [10].
- The minimum angle between \mathbf{x} and \mathcal{P} [11].

The length of an input vector \mathbf{x} is often normalized to 1. In this case these two criteria are identical. Since they are the same, from here on we think of the angle-based similarity S defined by the following equation:

$$S = \cos^2 \theta = \sum_{i=1}^k \frac{(\mathbf{x}^\top \cdot \Phi_i)^2}{\|\mathbf{x}\|^2}$$

Φ_i is the i^{th} orthogonal normal basis vector of the class subspace \mathcal{P} , which are obtained from applying the PCA to a set of patterns of the class. In more rigorous terms, these orthonormal basis vectors can be obtained as the eigenvectors of the correlation matrix

$$\sum_{i=1}^l \mathbf{x}^{(i)} \mathbf{x}^{(i)\top} \text{ where } \mathbf{x}^{(i)} \in X$$

of the class (X is the training dataset).

Learning Phase

1. Generate k class subspaces from each class by using PCA.

Recognition Phase

1. Calculate S between \mathbf{x} and each subspace \mathcal{P} , \mathcal{Q} etc.
2. Classify the \mathbf{x} into the class of the subspace where S was calculated to be the highest.

1.2.3 Mutual Subspace Method

The Mutual Subspace Method (MSM) is an extension of the Subspace Method (SM), where instead of having an input vector x , we use an input subspace \mathcal{P} . MSM is commonly used for image set classification [12].

The Subspace method assumes an input subspace and k class subspaces. Let us define the input subspace to be a d_p -dimensional subspace \mathcal{P} and the class subspaces to be d_q -dimensional subspaces $\{\mathcal{P}, \mathcal{Q}, \mathcal{R}, \dots\}$.

The similarity S between, for example, \mathcal{P} and \mathcal{Q} was originally defined as the minimum canonical angle θ_1 . Canonical angles [13] are uniquely defined as:

$$\cos^2 \theta_i = \max_{u_i \perp u_j (=1, \dots, i-1) v_i \perp v_j (=1, \dots, i-1)} \frac{|(u_i, v_i)|^2}{\|u_i\|^2 \|v_i\|^2}$$

Where $u_i \in \mathcal{P}$, $v_i \in \mathcal{Q}$, $\|u_i\| \neq 0$, $\|v_i\| \neq 0$.

We can also include the remaining canonical angles when calculating the similarity.

$$\tilde{S} = \frac{1}{t} \sum_{i=1}^t \cos^2 \theta_i$$

This value \tilde{S} reflects the structural similarity between two subspaces. It is also defined on the t smallest canonical angles. For practical applications, the canonical angles between subspaces \mathcal{P}_1 and \mathcal{Q} are obtained by calculating the singular values $\{\lambda_j\}_{j=1}^m$ of the correlation matrix between their basis matrices $P, Q \in R^{d \times m}$, i.e. solving the SVD of $V \Lambda V^\top = P^\top Q$. This corresponds to finding the rotation of each basis that is closest to the opposing subspace. From solving this problem, the canonical angles can be obtained by $\theta_j = \cos^{-1}(\lambda_j)$, where $j = 1, \dots, m$.

Learning Phase

1. Generate k class subspaces from each class by using PCA.

Recognition Phase

1. Calculate \tilde{S} between the input subspace and each dictionary subspace.
2. Classify the input subspace into the class where the dictionary subspace was calculated to be the highest.

1.2.4 Grassmann Discriminant Analysis

1.2.5 Linear Discriminant Analysis

1.2.6 Grassmann Manifold

Now, we introduce here the concept of Grassmann manifold and how it is useful to correspond subspaces to vectors. The essence is to make a one-to-one mapping such that the similarities between subspaces is preserved, simplifying the representation and allowing general methods to be applicable.

Grassmann manifold $\mathcal{G}(m, d)$ is defined as the set of m -dimensional linear subspaces of R^d . It is an $m(d - m)$ -dimensional compact Riemannian manifold and can be derived as a quotient space of orthogonal groups $\mathcal{G}(m, d) = \mathcal{O}(d)/\mathcal{O}(m) \times \mathcal{O}(d - m)$, where $\mathcal{O}(m)$ is the group of $m \times m$ orthonormal matrices.

A Grassmann manifold can be embedded in a reproducing kernel Hilbert space by the use of a Grassmann kernel. In this case, the most popular kernel is the projection kernel k_p , which can be defined as $k_p(\mathcal{Y}_1, \mathcal{Y}_2) = \sum_{j=1}^m \cos^2 \theta_j$, which is homologous to the subspace similarity. We can measure the distance between two points on a Grassmann manifold by using this projection kernel [?], and a subspace \mathcal{Y} can be represented by a vector with regards to a reference subspace dictionary $\{\mathcal{Y}_q\}_{q=1}^N$ as $y = k_p(\mathcal{Y}, \mathcal{Y}_q) = [k_p(\mathcal{Y}, \mathcal{Y}_1), k_p(\mathcal{Y}, \mathcal{Y}_2), \dots, k_p(\mathcal{Y}, \mathcal{Y}_N)] \in R^N$.

We introduce a discriminatory mechanism to separate the classes of signals, which is the Grassmann discriminant analysis (GDA). Basically, GDA is conducted as kernel LDA with the Grassmann kernels. We first outline the algorithm of linear discriminant analysis (LDA) [?]. Let x_1, \dots, x_N be the data vectors and y_1, \dots, y_N ($y_i \in 1, \dots, C$) be the class labels. Each class c has N_c number of samples. Let $\mu_c = \frac{1}{N_c} \sum_{i|y_i=c} x_i$ be the mean of class c , and $\mu = \frac{1}{N} \sum_i x_i$ be the overall mean. LDA searches for the discriminant direction w which maximizes the Rayleigh quotient $Ra(w) = w^T S_b w / w^T S_w w$ where S_b and S_w are the between-class and within-class covariance matrices respectively:

$$S_b = \frac{1}{N} \sum_{c=1}^C N_c (\mu_c - \mu)(\mu_c - \mu)^T, \\ S_w = \frac{1}{N} \sum_{c=1}^C \sum_{i|y_i=c} (x_i - \mu_c)(x_i - \mu_c)^T.$$

The optimal w is obtained from the largest eigenvector of $S_w^{-1} S_b$. Since $S_w^{-1} S_b$ has rank $C - 1$, there are $C - 1$ optima $W = [w_1, \dots, w_{C-1}]$. By projecting data onto the space spanned by W , we achieve dimensionality reduction and feature extraction of data onto the most discriminant subspace.

Kernel LDA [?, ?, ?] can be formulated by using the kernel trick as follows. Let $\Gamma : R^d \rightarrow \mathcal{F}$ be a non-linear map from the input space R^d to a feature space \mathcal{F} , and $\Gamma = [\gamma_1, \dots, \gamma_N]$ be the feature matrix of the mapped training points γ_i . Assuming w is a linear combination of those feature vectors, $w = \Gamma \alpha$, we can use the kernel trick and rewrite the Rayleigh quotient in terms of α as:

$$\begin{aligned} Ra(\alpha) &= \frac{\alpha^T \Gamma^T S_b \Gamma \alpha}{\alpha^T \Gamma^T S_w \Gamma \alpha} = \\ &= \frac{\alpha^T K (V - e_N e_N^T / N) K \alpha}{\alpha^T (K (I_N - V) K + \sigma^2 I_N) \alpha} = \\ &= \frac{\alpha^T \Sigma_b \alpha}{\alpha^T (\Sigma_w + \sigma^2 I_N) \alpha}, \end{aligned} \tag{1.1}$$

where K is the kernel matrix, e_N is a vector of ones that has length N , V is a block-diagonal matrix whose c -th block is the matrix $e_{N_c}e_{N_c}^\top/N_c$, and $\Sigma_b = K(V - e_N e_N^\top/N)K$. For example, the kernel matrix, K , is calculated as the similarity matrix between subspaces Y_q and Y_w . The term $\sigma^2 I_N$ is used for regularizing the covariance matrix $\Sigma_w = K(I_N - V)K$. It is composed of the covariance shrinkage factor $\sigma^2 > 0$, and the identity matrix I_N of size N . The set of optimal vectors α are computed from the eigenvectors of $(\Sigma_w + \sigma^2 I_N)^{-1} \Sigma_b$.

We apply the GDA algorithm to the reference subspaces Y_i^c to generate reference vectors y_i^c . When given an unknown bioacoustic signal $x_{in}(t)$, we compute its SSA subspace \mathcal{Y}_{in} and map it onto the manifold to generate a vector y_{in} ; then we predict its corresponding bioacoustic class (e.g. species) based on the nearest reference vector (1-NN).

1.3 Proposed Method

1.4 Experiments

In this chapter we conduct two experiments to demonstrate that our proposed method outperforms other traditional methods that have been introduced in previous chapters. The implementation is available at the website [xxxx](#).

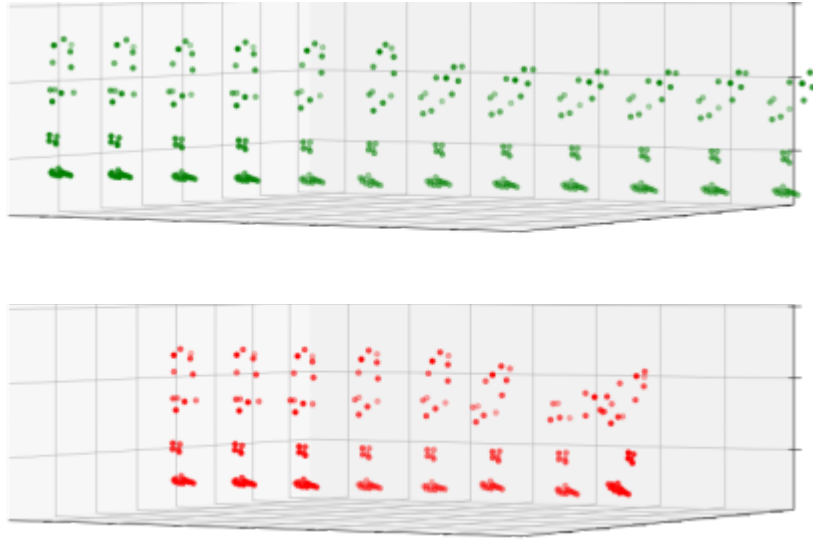


Figure 1.1: Two jump sequences sampled at 1fps. The successful jump is colored in green and the unsuccessful jump is colored in red.

1.5 Dataset

We collected data of 524 instances of a single-leg drop jump landing from 144 college football (soccer) players from the University of Tsukuba.

The dataset consists of the following attributes:

- 3 dimensional coordinates of 29 body markers capture by multiple infrared cameras.
- An integer (1 9) indicating the classification of ankle instability, labelled by an expert (label description: 1=Healthy, 2=Structural instability, 3=Subjective instability, 4=Sprained more than 3 times, 5=2 and 3, 6=2 and 4, 7=3 and 4, 8=2 and 3 and 4, 9=healthy but with a history of one or two sprains).
- Binary labels (0 or 1) indicating whether the single-leg drop jump was successful (0) or not (1). After consultation with an expert, we defined a successful jump as a jump where the subject was able to keep their balance on a single leg for more than 5 seconds after landing.
- Force plate data.
- Weight (kg) for each of the individual 144 athletes.

1.6 Evaluation Metrics

To evaluate the performance of different classification methods, we use a 5-fold cross-validation strategy to compute the classification accuracy, F1 score and the area under the receiver operating characteristic curve (AUC).

Although accuracy is a sufficient metric for performance in datasets with symmetric classes distribution and class value, it is an inadequate metric for datasets with class imbalance and datasets where values of false positive and false negatives are differ.

Therefore, we included the two additional metrics explained above. These metrics are frequently used in medical cases such as cancer detection from images [14][15].

1.6.1 Accuracy

Accuracy is computed as the fraction of correct predictions.

$$\text{accuracy}(y, \hat{y}) = \frac{1}{n_{\text{samples}}} \sum_{i=0}^{n_{\text{samples}}-1} 1(\hat{y}_i = y_i)$$

1.6.2 F1 Score

The F1 score can be interpreted as an average of the classifier's ability to:

- Not label as negative samples as a positive (i.e. Precision).
- Find all positive samples (i.e. Recall).

$$F_1 = 2 \times \frac{\text{precision} \times \text{recall}}{\text{precision} + \text{recall}}$$

1.6.3 Area Under the Curve

The receiver operating characteristic (ROC) plots the true positive rate (TPR) to the false positive rate (FPR) of a binary classifier with varying discrimination thresholds.

The area under the ROC curve (AUC) summarizes the information of the ROC in one number, between 0 and 1. AUC = 0.5 indicates an uninformative classifier (a classifier with uniform prediction for any sample) or random classifier if the classes are symmetric. Therefore no realistic classifiers should have an AUC < 0.5. AUC = 1 indicates a perfect classifier.

The algorithm to compute AUC with an in-depth explanation can be found in [16].

1.7 Experiment 1.

In this experiment we evaluate x classifiers accuracy on multiple binary classification problems regarding our single-leg drop jump dataset.

1.7.1 Experiment setup

We experiment with the following setup.

- (A) Pose data up until landing.
- (B) All pose data.
- (C) Pose data + Force Plate data.

1.7.2 Results

1.7.3

1.8 Discussion

Chapter 2

Development of an Analysis Framework for Fencing Based on 2D/3D Time-Series Pose Data

In many competitive sports, the analysis of video footage and numerical data can help gain an advantage in athletic performance. However, analysis requires not only specialized knowledge but also a great deal of time and effort. Therefore, it is very important to streamline and automate the process.

In recent years, advances in computer vision and machine learning have reduced the burden of these tasks in various fields. In this study, using such technology, we propose an analysis framework for extracting important information from video and numerical data to assist experts in their analysis work.

Specifically, we will

1. Create a panoramic image of the entire court from fencing footage captured by a Handycam
2. Extract skeletal information of each athlete and their position on the court
3. Examine methods for analyzing and visualizing matches and athletes using the skeletal and positional information.

2.1 Analytical Frameworks in Fencing

There are many studies that focus on applying computer vision to fencing. For example, Takahashi[17] et al. proposed a robust detection and tracking method to visualize the trajectory of a fencing sword for live TV programs. Since live TV programs require not only accuracy but also real-time computation, they successfully track fast-moving fencing swords by using supervised machine learning to detect the swords and using a particle filter to predict their positions in the next frame.

Athow[18] et al. proposed a computer vision-based scoring system to assist referees in foil events in fencing. They used color blob detection to track fencers who wore color patches on their hands.

In addition, there have been cases of computer vision and motion analysis: Makawaski et al. [19] created a new local trace image to represent fencing motions, and showed that the dynamics of the motion is useful for analyzing similar motion patterns. They showed that motion dynamics can be useful for analyzing similar motion patterns. In the same study, he also created a dataset for analyzing footwork, contributing to the development of the field.

However, there is still no research that proposes a comprehensive framework for extracting posture and positional information from video in fencing and performing tactical analysis to gain an advantage in competition.

In this study, we propose an analysis framework that uses computer vision and machine learning to extract posture and position information to assist experts in their analysis tasks.

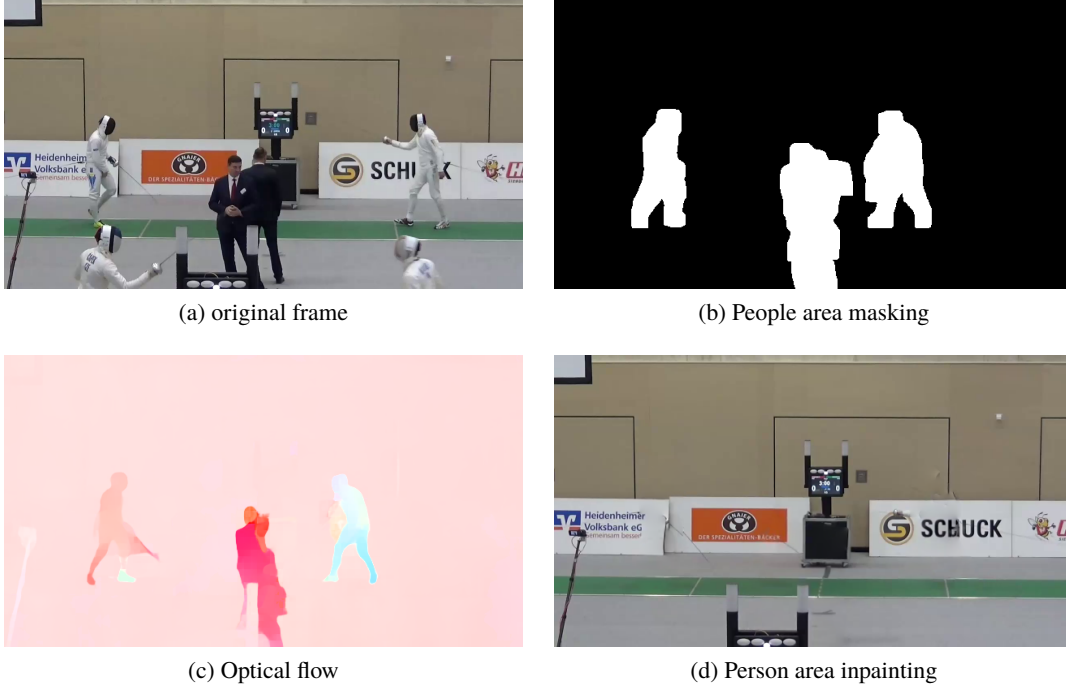


Figure 2.1: process until the person area is inpainted

2.2 Proposed Method

The framework proposed in this study can be divided into three main stages. In the first stage, a panoramic image is generated from a video camera image; in the second stage, the player's posture coordinates and absolute coordinates on the court are estimated from the panoramic image. In the second stage, we estimate the posture coordinates and the absolute coordinates of the player on the court from the panoramic images. In the last stage, we analyze and visualize the obtained skeletal and positional information of the player by dimensional compression and clustering. The details of each element are described below.

2.2.1 Generation of panoramic images

Since it is strategically useful to know where the players are on the court, it is useful to generate panoramic images showing the entire court. In addition, the position of the players on the pitch can be estimated from the panoramic image by using person detection, so that the positional information of the players can be obtained not only visually but also numerically.

2.2.2 extraction of person region

Usually, panoramic background images are generated using images of only stationary objects, so to use conventional methods, moving objects, i.e., humans, must be eliminated in each frame, and the

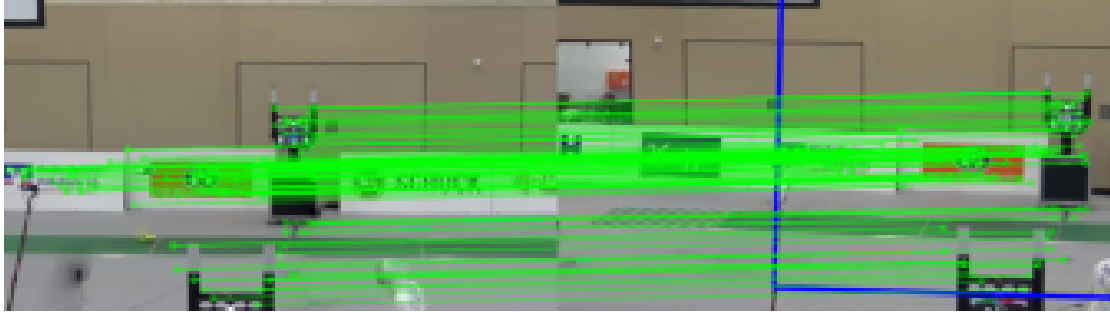


Figure 2.2: SIFT feature matching with RANSAC method

eliminated areas must be complemented in some way. In this study, this is done in two steps. For the extraction of human regions, we use segmentation with HR-NET[20], and label each pixel of each frame as human region ($=1$) or not human region ($=0$). As a post-processing step, we perform smoothing operations in the vertical, horizontal, and temporal directions to mask the human region more reliably in each frame. process until the person area is inpainted(b) shows the masking of the person area after post-processing.

2.3 Experiments

2.3.1 Dataset

公益社団法人日本フェンシング協会から提供された 100 本のフェンシング・エペ種目の試合映像からデータセットを作成した．試合映像は、ハンディカム 1 台で撮影したもので、1 秒あたり 60 フレームで約 15 分間ほどである．

2.4 Discussion

2.5 Results and Discussion

2.5.1 Generation of panoramic images

We were able to generate panoramic images and create video mosaics more quickly than conventional methods by devising our own method for obtaining posture and position information. For analysis using posture and position information, we will conduct more detailed analysis using the position and skeletal data of the players obtained by the above method.

2.5.2 clustering, visualization

As an example application of the skeletal information obtained by the proposed framework, clustering was performed. The results of clustering up to the second principal component of the skeletal information with a mixed Gaussian model are shown in Figure 2.3. The legend indicates each cluster. Some of the representative posture frames for each cluster are shown in Figure 2.4. A characteristic posture was observed in each cluster. The most common posture observed in cluster c2 was the most neutral posture, which can be shifted to any posture. Cluster c4, on the other hand, had a lower posture than the posture frames of the other clusters, with the epee sword sticking out in front of the body. The posture of cluster c4 was lower than that of the other clusters, and it was considered to be an attacking posture, based on the characteristics of fencing. Thus, clustering using the proposed framework may make it easier to classify the plays. In addition, the posture frames of clusters c8 and c13 are not directly related to the play. Clustering can also remove these data, which would otherwise have to be done manually.

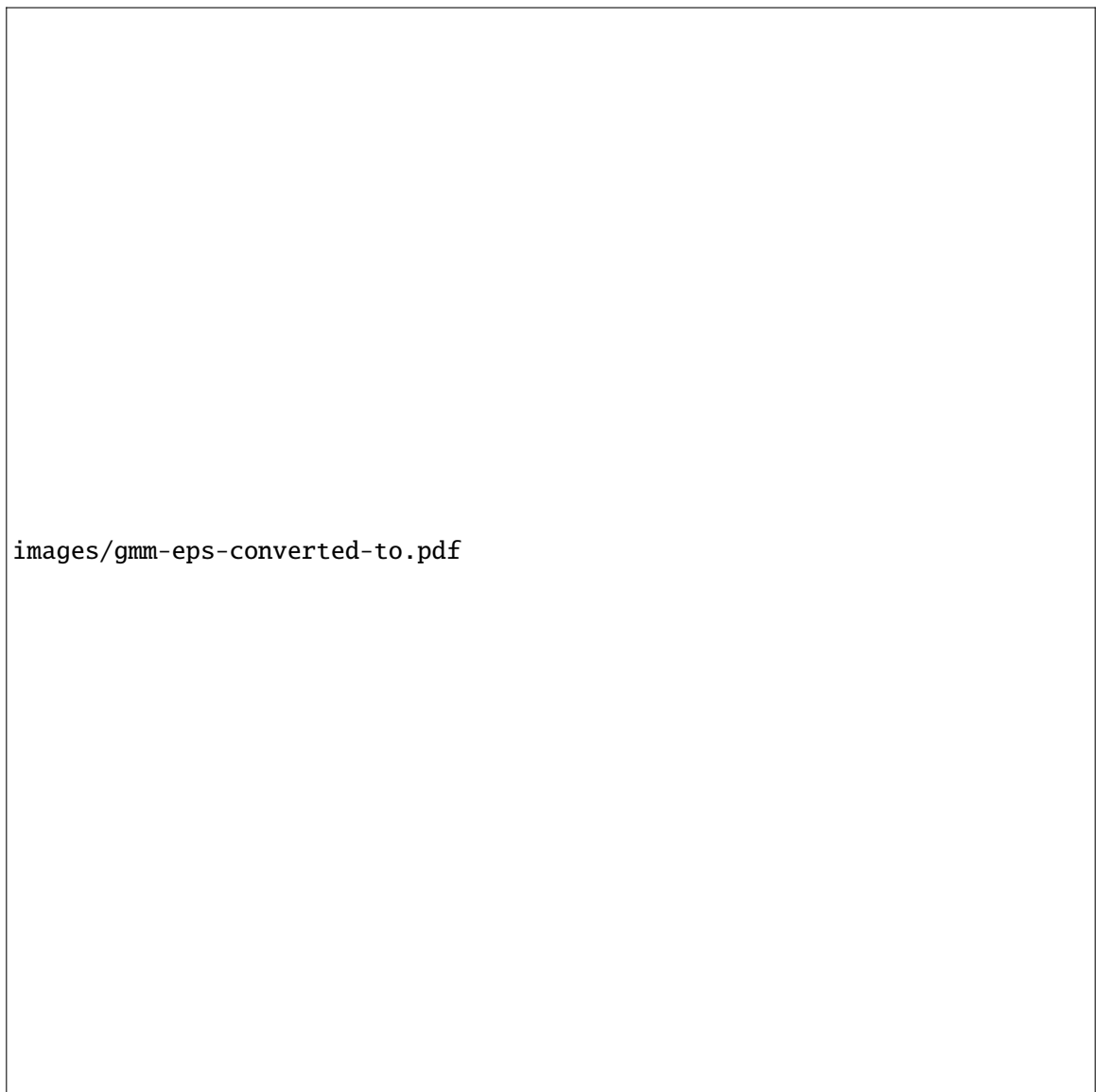


Figure 2.3: Clustering results. Up to the second principal component is illustrated.

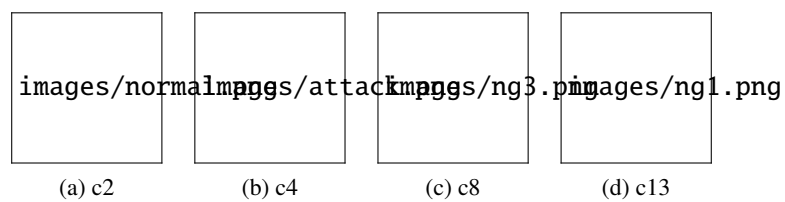


Figure 2.4: An example of a cluster posture frame. The label is the same as in Figure 2.3

2.6 conclusion

In this study, we proposed an analytical framework for extracting video and numeric values to assist experts in their analysis tasks using computer vision and machine learning in the epee fencing event. As a future prospect, we will conduct a more rigorous evaluation of the proposed method and try to extend the framework to other events.

Chapter 3

Conclusion

Appendices

Appendix A

Details regarding data

A.1 Details on the given data

Index	Name
0	Cervical vertebra 7
1	Calcaneal tuberosity
2	Fibula Head
3	Greater trochanter
4	Suprasternal notch
5	Anterior superior iliac spine (L)
6	lateral epicondyle
7	medial malleolus
8	Posterior superior iliac spine (L)
9	medial condyle
10	medial epicondyle
11	lateral malleolus
12	Base of first metatarsal bone
13	Base of second metatarsal bone
14	Base of fifth metatarsal bone
15	Head of first metatarsal bone
16	Head of second metatarsal bone
17	Head of fifth metatarsal bone
18	Fibular trochlea of calcaneus
19	First distal phalanges
20	Anterior superior iliac spine (R)
21	Posterior superior iliac spine (L)
22	Acromion (L)
23	Acromion (R)
24	Sustentaculum tali
25	Thoracic vertebrae 8
26	Tibial tuberosity
27	Scaphoid bone
28	Xiphoid process

A.2 Finding the time of landing

$$t = \sqrt{\frac{2h}{g}} = \sqrt{\frac{2 * 0.4441}{9.8}} \approx 0.39(s)$$

Therefore we find the frame where the average foot position on the y-axis is lowest, with the condition that the frame must be in within 39 frames after the highest point.

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