# An Online Robot Collision Detection and Identification Scheme by Supervised Learning and Bayesian Decision Theory

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Abstract-This article is dedicated to developing an online collision detection and identification (CDI) scheme for human-collaborative robots. The scheme is composed of a signal classifier and an online diagnosor, which monitors the sensory signals of the robot system, detects the occurrence of a physical human-robot interaction, and identifies its type within a short period. In the beginning, we conduct an experiment to construct a data set that contains the segmented physical interaction signals with ground truth. Then, we develop the signal classifier on the data set with the paradigm of supervised learning. To adapt the classifier to the online application with requirements on response time, an auxiliary online diagnosor is designed using the Bayesian decision theory. The diagnosor provides not only a collision identification result but also a confidence index which represents the reliability of the result. Compared to the previous works, the proposed scheme ensures rapid and accurate CDI even in the early stage of a physical interaction. As a result, safety mechanisms can be triggered before further injuries are caused, which is quite valuable and important toward a safe human-robot collaboration. In the end, the proposed scheme is validated on a robot manipulator and applied to a demonstration task with collision reaction strategies. The experimental results reveal that the collisions are detected and classified within 20 ms with an overall accuracy of 99.6%, which confirms the applicability of the scheme to collaborative robots in practice.

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Note to Practitioners-This article is intended to provide a novel online collision event handling scheme for robots in industrial environments. This scheme is designed to quickly and accurately detect an accidental collision and distinguish it from the intentional human-robot interaction. The method takes the raw signals from external torque sensors and provides a collision diagnosis result with a reliability index. The simple structure makes it easy to be implemented as a regular fault monitoring routine for collaborative robots. Different from the conventional methods, the proposed collision identification scheme in this article especially focuses on overcoming the following two challenges in practice: first, to timely and accurately report a collision within its early stage, and second, to ensure a high identification accuracy in a complicated environment, where ubiquitous disturbance and noise are unneglectable. The experimental validation at the end of this article confirms its promising application value in human-robot collaboration.

*Index Terms*—Anomaly monitoring, collision detection and identification (CDI), collision event pipeline, fault detection and isolation, human–robot interaction, robot safety, supervised learning.

#### I. INTRODUCTION

S AFETY is always a critical issue for human–robot collab-orative tasks, especially when physical human–robot interaction (pHRI) is demanded [1], [2]. Different types of pHRI exert various impacts on humans. The intentional contacts desired by the cooperative tasks, such as the human teaching processes, are usually quite safe. On the contrary, the accidental collisions, which may lead to unexpected injuries or damages, are often dangerous to humans. As pointed out by many surveys on the safety of human-robot collaboration [1], [3], the accidental collisions are inevitable in pHRI and should be carefully handled. Especially, the reaction strategy for the accidental collisions should be distinguished from the intentional contacts since they tend to cause opposite consequences, which motivates the research on collision event handling. In the literature, a typical collision event handling pipeline usually contains two procedures, namely, the precollision and the postcollision ones [4], [5]. The development of such a pipeline involves various topics, including collision avoidance [6], [7], collision force estimation [8], [9], collision detection and identification (CDI) [10], [11], and collision reaction strategy design [12], [13]. These topics correspond to different components in the pipeline.

This article is specifically concerned with the CDI problem of robot systems. As part of the postcollision procedure, it is intended to monitor the sensory signals, detect the occurrence of pHRI, and identify whether it is an accidental collision or an intentional contact. The results of CDI will be given to the collision reaction procedures, where the corresponding safe strategy is enabled to avoid further harms [5]. As claimed in [5], accidental collisions and intentional contacts possess distinguishable properties, which are reflected in the features of the sensory signals. By considering this, one can formulate robot CDI as a multiclass classification problem, with collision, contact, and free, respectively, denoting the accidental collisions, intentional contacts, and nominal cases where no pHRI occurs. Then, the main target of CDI becomes to develop a pHRI signal classifier using a data set that contains the pHRI waveform. In general, this proposes a classification problem for time series.

For such a problem, expertise can be found in related work, which will be discussed in detail in Section II. In general, an important technique in the classification problem for time series is the segmentation of raw signals since classifiers cannot process the series with infinite length. In a conventional paradigm of classifier development, the raw signals are segmented before predictions are conducted, even for the online applications. In the context of CDI, the conventional solutions typically design some detection logics, such as a signal threshold, to approximately mark the starting instant of a pHRI and segment the online signals after observing the entire pHRI waveform, before producing a prediction result using a trained classier. As a result, an accurate prediction result can only be produced when the pHRI almost vanishes. By almost, we mean that the pHRI waveform is contained in the signal segments as much as a correct identification result can be produced. Such a method, however, usually leads to a useless result in the context of safe human-robot collaboration since a collision or a contact has already occurred and negative consequences may have already been caused. Intuitively speaking, a prediction result of the classifier does not really predict the occurrence of a collision. The reason for this issue is that the signal values in the future are not available for the segment at the current time, due to the causality of online time series.

Nevertheless, this is a significant issue for safe human-robot collaboration since a CDI scheme that can be practically applied is expected to accurately report the pHRI in its early stage, definitely before it vanishes, such that potential injuries can be prevented in advance. Unfortunately, such a valuable topic has attracted very little attention, probably due to the following two challenges. First, accurate prediction results should be produced even when the segmented pHRI waveform is incomplete. From a general perspective, this introduces inconsistency between the distribution of the training data and that of the predicted samples, if the classifier is developed on the signal segments with complete pHRI waveform. This indicates that the identification result in the early stage of a pHRI is very likely to be incorrect. Second, an accidental collision is usually referred to as an *instantaneous anomaly* [14], which occurs unexpectedly and only lasts for a short period

of time. Thus, it is more difficult to be identified than the other anomalies that are featured with low bandwidth and steady changes [15], [16]. The main target of this article is to fill these gaps by proposing a novel online CDI scheme that ensures both high accuracy and fast response.

The remaining content of this article is organized as follows. Section II summarizes the related work and clarifies the main contributions of this article. Section III introduces the overall structure of the scheme and the collection process of the pHRI data set. The development of two important components of the scheme, the signal classifier and the online diagnosor, is, respectively, presented in Sections IV and V. In Section VI, the proposed CDI scheme is validated on a seven-degree-offreedom (DoF) KUKA robot. The decent performance of the CDI scheme revealed by the experimental validation confirms its applicability in practice. Finally, Section VII concludes this article.

#### II. RELATED WORK

Conventionally, CDI for robot systems has been recognized as an engineering-oriented problem and is mainly solved by simple threshold logic or hypothesis testing-based methods [11], [17], where the filter-based techniques are widely applied [18]. More recent work tends to formulate CDI as a classification problem for segmented time series. Similar work can be found in relevant fields, such as speech recognition [19], snoring identification [20], bird sound recognition [21], and fault identification of mechatronic systems [22]–[24]. The common ground of these applications is that the segmented time series, either acoustic signals or sensory streams, is expected to be classified as a certain type. In [25], the classification of time series is generally investigated by applying various data sets. In [26], a deep neural network is constructed to detect the system faulty status. In [27], various classifiers are developed and tested using the recorded signal samples. Similar work also includes [28] and [29], where neural network is constructed to monitor the grasping slippages and colliding torques, and [30] and [31], where SVM classifiers are developed to detect external collisions. Based on the abovementioned work, a mature development framework for time series has been well-formed. However, such a framework is confronted with challenges when applied to CDI for robot systems with pHRI tasks since the prediction results are only available after the collision vanishes or the entire pHRI waveform is segmented, which has been explained in Section I. Therefore, these methods are not suitable for human-robot collaboration with high requirements for human safety.

As alternatives, probabilistic series models, such as the hidden Markov model (HMM) [32] and the Gaussian mixture model (GMM) [33], are also applied to CDI by exploiting the dependence properties of time series. In [34], an HMM is developed to detect exceptional events in an object-alignment robot task, where the measured torques and their derivatives are used. The work in [35] also develops an HMM model to realize a fault detection scheme for a feeding-assistant robot using multisensor signals. In [36], probabilistic support vector machine is used to detect online anomalies. Nevertheless, these

series models require artificially assigned prior knowledge on the distributional dependence of the raw signals. In most practical scenarios, such knowledge is challenging to acquire, especially in a complicated environment where the distribution of the disturbance and noise is not treated as Gaussian.

The main contribution of this article is to develop an online CDI scheme for safe human-robot collaboration. The major advantage of this novel scheme over the conventional methods is to produce fast and accurate CDI in the early stage of pHRI. Apart from the identification result, it also provides a confidence index to indicate how much the result can be trusted, which benefits the design of a flexible collision handling pipeline in future work. Moreover, the scheme does not require extra detection logic and any prior knowledge on the dependence of pHRI signals. Therefore, no heuristic thresholds or distribution assumptions are needed. Another contribution of this article is to provide a pHRI data set with manual labels specifically for the CDI usage, which is still lacking in the literature. Different from most of the previous work where the torque sensors are installed on the end-effectors, the signal data we used in this article are from the torque sensors on the robot joints, which can even detect pHRIs on robot links. On the other hand, the data set contains more interfering signals due to coupled mechanical vibrations and strong current noise on the sensors, which leads to a larger challenge to the development of the CDI scheme. From another perspective, however, this article reflects the complicated conditions in practice, which provides an outcome truly with application values.

## III. SCHEME OVERVIEW AND DATA COLLECTION

In this section, we introduce the overall structure of the proposed CDI scheme. To develop this scheme, we conducted a data collection experiment with manual collisions and contacts to construct a pHRI data set. The data come from the segmentation of the external torques that are measured by shaft-mounted torque sensors installed on the robot joints.

#### A. Overview of the CDI Scheme

The general structure of the proposed online CDI scheme and the development flowchart of each component are illustrated in Fig. 1. During an online execution, the CDI routine takes the torque signal  $\tau$  and segment it recurrently. Then, the features extracted from the signal segment  $\mathcal{T}$  are given to the classifier, which produces a prediction result *r* among *collision, contact,* and *free.* Based on a series of prediction results, the online diagnosor offers a CDI diagnosis result  $C^*$ and a confidence index  $\mathcal{I}$ . In this framework, no heuristic threshold values are needed to mark the occurrence time of the pHRI.

## B. Data Collection

To construct the pHRI data set for the development of signal classifier, we conduct a collection experiment using a seven-DoF KUKA LWR4+ robot manipulator [37], which is firmly mounted onto a fixed platform. During the experiment, different types of physical interaction in various strengths



Fig. 1. Structural overview of the proposed online CDI scheme and the corresponding development flowcharts, where  $\tau$  is the measured torque signal,  $\mathcal{T}$  is the signal segment, r is the prediction result of the signal classifier, and  $C^*$  and  $\mathcal{I}$  are, respectively, the ultimate diagnosis result and its confidence index.

and directions are manually exerted on the robot end-effector by seven subjects hired for this experiment, thus aiming to include individual uncertainties in the collected data, to ensure the generalization ability of the data set. In the experiment, accidental *collisions* are created using a soft hammer with fast hand speed and tough strength, while intentional *contacts* are made by gloved hands with compliant forces. A spherical plastic end-effector is specially designed to bear the contact forces from arbitrary directions.

Note that the pHRI is exerted during the movement of the robot, such that the measured signals contain drifts and noise caused by the joint motions. It is worth mentioning that the drifts and noise are also affected by the direction of the joint motion. By considering this, we apply several different robot trajectories that are assigned with the same starting position but different ending positions, such that both positive and negative motion directions are covered for each joint. The trajectories are designed in the joint space and interpolated by sinusoidal functions to ensure smooth motion. All these experimental configurations are intended to ensure a representative data distribution, which is important to ensure a high generalization ability to the classifier.

The motion of the robot is controlled by a trajectory tracking program that implements a continuous reciprocating motion of the manipulator between the starting position and each ending position. The program also reads the measured torque signals and records them as 7-D time series with a sampling period of 1 ms. During the motion of the robot, one experimenter exerts an accidental *collision* or an intentional *contact* on the end-effector. At the same time, the other experimenter notes down the occurrence instant and the type of pHRI. To obtain a balanced data set between *collisions* and *contacts*, we try to produce an equal amount of samples for these two classes.

#### C. Signal Segmentation and Sample Labeling

After collecting the raw pHRI signals, we conduct segmentation to split the signals into segments with a fixed length. Let us denote the width of the segments as l and the bias as b, with a unit of ms. By bias, we mean the time period between the occurrence instant of a pHRI and the ending instant of the segment, where we have  $0 \le b \le l$ . In general, l determines how much signal information is included in the segments,



Fig. 2. Waveform of signals (external toques on seven joints), respectively, with classes of (a) *collision*, (b) *contact*, and (c) *free*. The segmented partitions are marked with colored areas, where b = l = 300 (ms).

while *b* adjusts the proportion of the pHRI waveform in the segment. If the ending instant of the segment represents the current time, *b* denotes the period after the occurrence of the corresponding pHRI. Therefore, the segmentation scheme is well-determined by the two parameters *l* and *b*. For the data set, the segment bias is typically set as b = l since the torque values before pHRIs are irrelevant to the classification. The determination of segment width *l* is mainly based on the engineering experience. Usually, it should be large enough to contain sufficient information of the pHRI waveform. On the other hand, an overlarge width may involve irrelevant signals and lead to poor generalization ability.

To determine the value of l, we inspect the waveform of the pHRI signals presented in Fig. 2. We notice that the average width of a *contact* is approximately 400 ms while that of a *collision* is less than 200 ms. This also applies to most of the recorded pHRI signals. Therefore, we set the segment width as l = 300 (ms), as a balanced result. Then, we label the two types of segment samples, respectively, as *collision* [Fig. 2(a)] and *contact* [Fig. 2(b)]. The segmented parts of the raw signals are marked by colored areas in Fig. 2. During the segmentation, invalid data due to storage damages or unrecognizable pHRI waveforms are eliminated. As a result, we obtain 6718 collisions and 7346 contacts, which are approximate to a balanced radio 1:1. We also create signal segments without any pHRI waveform and label them as *free*. However, the number of *free* samples should not be equal to the other two classes since the frequency of contact-less cases is usually higher than pHRI. In practice, both collisions and contacts are positive instances which will call the corresponding safety mechanism, while *frees* are negative instances which do not trigger the safety mechanism. Therefore, a reasonable idea would be to keep a balance between the *free* samples and the summary of all pHRI samples, i.e., a scale ratio 1:1:2 between collision, contact, and free. Therefore, we take the free segments, respectively, in the front, rear, and middle of the pHRI segments in the raw signals, and finally obtain 13633 of them.

Until now, a data set containing the pHRI signal segments is constructed. We randomly shuffle all the samples and split them into a training set and a test set with the partition ratio 3:1. The training set will be applied to feature engineering and model selection, while the test set will be used to evaluate the trained model.

## IV. DEVELOPMENT OF THE SIGNAL CLASSIFIER

In this section, we develop the signal classifier component in the CDI scheme. As explained in Section II, we do not consider the probabilistic series models to avoid prior knowledge on the distributional dependence of the raw signals. In addition, although the end-to-end learning mechanisms, such as the convolutional neural network (CNN) and recurrent neural network (RNN), have drawn much attention and made great achievements in image recognition and model-free planning, they still suffer from the lack of explainability and high dependence on complex manually designed structures. Therefore, in this article, we develop the signal classifier based on the paradigm of supervised learning, by which we assume that the samples in the data set are independent of each other.

### A. Feature Extraction

To form a feature set that benefits the signal classification, we consider both the properties of pHRI signals and the successful experience in the previous work [27], [38]. From Fig. 2, we can conclude that the waveform of *collisions* [Fig. 2(a)] has sharp shapes and fast amplitude changes, while the waveform of contacts [Fig. 2(b)] changes more gently and lasts for a longer time. We also investigate the spectrum of the signal segments in Fig. 2 and shown it in Fig. 3. Compared to the free sample, the *collision* and *contact* possess more components in all frequency ranges. Especially, the *collision* shows a large amplitude in the frequency interval 10-20 Hz, while the *contact* shows that in the frequency interval 0-10 Hz. Therefore, distinguishable properties between the two classes are found in both time and frequency domains. Based on this, we initially extract 18 features in both domains. Most of these features also have achieved success in the previous work [38].

1) Features in the Time Domain: The time-domain features are frequently used in the fields like signal processing and pattern recognition [39], [40]. Since each sample is naturally a segmented time series  $\mathcal{T} = \{\tau_1, \tau_2, \ldots, \tau_l\}$ , the timedomain features can be represented as the functions of  $\mathcal{T}$ . Here, we mainly select features that are concerned with the amplitude changes of the signals. First of all, the first- to fourth-order statistical moments, namely the mean value  $M_{\tau}$ , the variance  $V_{\tau}$ , the skewness  $S_{\tau}$ , and the kurtosis  $K_{\tau}$  of samples  $\mathcal{T}$ , are applied to depict the stochastic properties



Fig. 3. Spectrum of signal segments (external toques on seven joints), respectively, with classes of (a) collision, (b) contact, and (c) free using FFT.

of the signal segment, where  $M_{\tau} = (1/l) \sum_{i=1}^{l} \tau_i$ ,  $K_{\tau} = (1/l) V_{\tau}^{-2} \sum_{i=1}^{l} (\tau_i - M_{\tau})^4 - 3$ ,  $V_{\tau} = (1/l) \sum_{i=1}^{l} (\tau_i - M_{\tau})^2$ , and  $S_{\tau} = (1/l) V_{\tau}^{-(3/2)} \sum_{i=1}^{l} (\tau_i - M_{\tau})^3$ . The median value  $\tilde{m}_{\tau}$ , the extreme range  $R_{\tau}$ , and the extreme deviation  $D_{\tau}$  of  $\mathcal{T}$  are also used as supplementary, where  $R_{\tau} = \max_i |\tau_i| - \min_i |\tau_i|$  and  $D_{\tau} = \max_i |\tau_i| - M_{\tau}$ , for  $1 \leq i \leq l$ . In addition, we propose the energy increasing rate  $E_I = (1/2) \lg(\sum_{i=[n/2]}^{n} \tau_i^2 / \sum_{i=1}^{[n/2]} \tau_i^2)$  to represent the temporal change of signal energy within a segment, where the ratio is made between the squared signal sums within the two halves of the segment. Here,  $[\cdot]$  rounds a real number to integer.

2) Features in the Frequency Domain: The frequencydomain features are commonly applied to depict the spectral profiles of signals [11], [41], which are mostly calculated by fast Fourier transformation (FFT). The FFT of a signal sequence  $\mathcal{T}$  includes a sampled frequency sequence  $\mathcal{F} = \{f_1, f_2, \dots, f_m\}$  with the frequency sampling interval  $f_s/m$  and the corresponding complex spectrum sequence  $S = \{S(f_1), S(f_2), \dots, S(f_m)\}$ . The length of the spectrum sequence m is usually equal to l. The frequency-domain features used in this article are mainly concerned with the amplitudes and phase angles of the signal spectrum in different frequency intervals, such as the mean frequency f, the fundamental frequency  $f^*$ , and their corresponding spectral amplitudes  $|S(\bar{f})|$ ,  $|S(f^*)|$  and phase angles  $\phi(\bar{f}), \phi(f^*)$ , where  $|\cdot|: \mathbb{C} \to \mathbb{R}$  and  $\phi(\cdot): \mathbb{C} \to \mathbb{R}$ , respectively, denote the amplitudes and phase angles. The mean frequency  $\bar{f}$ is defined as  $\bar{f} = \sum_{i=1}^{m} f_i |S(f_i)| / \sum_{i=1}^{m} |S(f_i)|$ , and the fundamental frequency  $f^*$  is the frequency point such that  $S(f^*)$  is the spectrum summit. The crest factor  $F_{\text{crest}}$ , the average signal energy  $E_{\rm rms}$ , and the subband energy ratio  $E_r^{f_c}$  are, respectively, defined as  $F_{\text{crest}} = |S(f^*)|/((1/m)\sum_{i=1}^m |S(f_i)|^2)^{1/2}$ ,  $E_{\text{rms}} = ((1/m)\sum_{i=1}^m |S(f_i)|^2)^{1/2}$ ,  $E_r^{f_c} = (1/2)\lg(\sum_{f_i=f_c}^{f_s} |S(f_i)|^2/\sum_{f_i=0}^{f_c} |S(f_i)|^2$  for  $f_c = 10$ , 20 Hz. These features are also widely used in [38], [39], and [42] to depict the energy properties of signals. The crest factor  $F_{\text{crest}}$ , also known as peak-to-average ratio, represents the significance of the signal peak. The average energy  $E_{\rm rms}$ denotes the average signal power level, and the subband energy ratio  $E_r^{f_c}$  reflects the relative energy proportion on the two sides of the subband frequency  $f_c$ . Here, we define two subband energy indexes. The corresponding subband frequencies 10 and 20 Hz are determined according to the distinguishable spectral features of *collisions* and *contacts*.

In summary, we extract the following features  $\mathcal{X}^+ = \{M_{\tau}, V_{\tau}, S_{\tau}, K_{\tau}, \bar{m}_{\tau}, R_{\tau}, D_{\tau}, E_I, f^*, |S(f^*)|, \phi(f^*), \bar{f}, |S(\bar{f})|, \phi(\bar{f}), F_{\text{crest}}, E_{\text{rms}}, \text{ and } E_r^{10}, E_r^{20}\}$ , where  $\mathcal{X}^+$  denotes a *full* feature set. For brevity, we order these features and number them from #1 to #18. Note that the signals have seven dimensions corresponding to the seven robot joints. Therefore, in total, 126 features are generated.

### **B.** Feature Selection

To select the beneficial features for the classification, we use the Spearman correlation analysis and the Relieff algorithm to evaluate the 126 features in  $\mathcal{X}^+$ . The former inspects the dependence between any two features, while the latter calculates weights for the features that are recognized as their *importance* to the classification. Features with high dependence on the others or low *importance* are eliminated after the analysis. Note that we only use the data in the training set during the entire feature selection procedure.

1) Dependence Analysis: High dependence among features brings redundancy to the model training, which potentially causes a poor generalization ability to the classifier. Therefore, we analyze the Spearman correlation between the features to investigate their dependence. The Spearman rank correlation analysis is an efficient tool to describe the general monotonic relationship between two stochastic variables, which applies to not only the linear dependence but also the nonlinear cases. Given the values of two features  $X = \{X_1, X_2, \dots, X_m\}$  and  $Y = \{Y_1, Y_2, \dots, Y_m\}$ , where *m* is the size of the data set, the Spearman correlation coefficient  $\rho_{X,Y}$  is calculated by  $\rho_{X,Y} = \sum_{i=1}^{m}$  $(x_i - \bar{x})(y_i - \bar{y})/(\sum_{i=1}^m (x_i - \bar{x})^2 \sum_{i=1}^m (y_i - \bar{y})^2)^{1/2}$ , where  $x = \{x_1, x_2, \dots, x_m\}$  and  $y = \{y_1, y_2, \dots, y_m\}$  are the ranks of features X and Y, and  $\overline{\cdot}$  denotes the mean value [43]. Ranging from -1 to 1, the coefficient  $\rho_{X,Y}$  depicts the dependence between X and Y, of which the sign denotes whether the relationship of the two variables is proportional (when positive) or inversely proportional (when negative), and the absolute value indicates the extent of dependence. We apply the Spearman correlation analysis to all 126 features in  $\mathcal{X}^+$ . The results are illustrated in Fig. 4 by colored matrices.

Fig. 4(a) displays the dependence between all 126 features that are grouped by the joint numbers. Each square block (surrounded by black dashed lines) on the diagonal represents the self-feature dependence of each joint, i.e., the dependence



Fig. 4. Spearman correlation analysis of the features. Each microsquare block represents the correlation coefficients between the corresponding two features, of which the values are marked by colors. A darker block indicates higher dependence between the corresponding features. (a) Spearman correlation between the 126 features. (b) Spearman correlation between the 18 features on joint #1.

between the features of the same joint. Meanwhile, the blocks off the diagonal indicate the dependence among the features of different joints. It is noticed that the blocks on the diagonal are highly similar to each other, which indicates that the features on all robot joints possess similar self-dependence properties. Therefore, it is reasonable to only investigate the analysis result of a single joint and selected or eliminated the features in batches in the unit of joint.

The correlation analysis of all 18 features on joint #1, which corresponds the block in the down-left corner in Fig. 4(a), is illustrated in Fig. 4(b). It can be seen that #1( $M_\tau$ ) is highly dependent on #5( $\bar{m}_\tau$ ). Meanwhile, #2( $V_\tau$ ), #6( $R_\tau$ ), #7( $D_\tau$ ), and #16( $E_{\rm rms}$ ) also indicate high dependence on each other. Therefore, these features are preferably considered to be eliminated from  $\mathcal{X}^+$  yet their *importance* to the classification still needs to be considered.

2) Importance Analysis: After analyzing the dependence of the features, we apply the Relieff algorithm to the 126 features in  $\mathcal{X}^+$  to evaluate their *importance*. As a popular feature engineering technique used in the previous work [38], it provides a weight for each feature to depict its *importance* to the

classification, which is independent of the classifier models. The predecessor of Relieff, known as Relief, is originally proposed in [44] for binary classification problems. Its main idea is to recursively and randomly select a sample x from the data set and find out its closest samples with the same and different labels (respectively, referred to as the near-hit  $z^+$  and the near-miss  $z^-$ ). For the *i*th feature of sample  $x, x_i$ , its weight  $w_i$  is iteratively updated by the following law:

$$w_i = w_i - (x_i - z_i^+)^2 + (x_i - z_i^+)^2$$

where  $z_i^+$  and  $z_i^-$  are the *i*th features of  $z^+$  and  $z^-$ . The weight  $w_i$  depicts the reward (if  $w_i$  is positive) or punishment (if  $w_i$  is negative) of the *i*th feature to the classification, and a larger  $w_i$  indicates higher *importance*. Relieff is adapted from Relief as the extension to multiclass problems [45].

On the training set, we calculate the weights or importance of all 126 features using Relieff and illustrate the results in Fig. 5, grouping them, respectively, by the feature number [see Fig. 5(a)] and joint number [see Fig. 5(b)]. The results indicate that the feature importance varies among both features and joints. In Fig. 5(a), the feature #2 ( $D_\tau$ ) shows the lowest *importance* (0.49), followed by #4( $K_\tau$ ), #5( $\bar{m}_\tau$ ), #9( $f^*$ ), and #13( $S|(\bar{f})|$ ). In Fig. 5(b), the feature weights on joints #1, #2, and #4 are higher than the other joints.

Since the classifier is supposed to function in an online scheme, we are quite concerned with the computational load of the features. Therefore, we tend to form a small feature set while maintaining a decent classification accuracy. Considering both high dependence and low *importance*, we eliminate features #2  $(D_{\tau})$ , #4 $(K_{\tau})$ , and #5 $(\bar{m}_{\tau})$ . Features #6 $(R_{\tau})$  and  $#16(E_{\rm rms})$  are also eliminated due to their strong relationship with  $\#7(D_{\tau})$ , as well as  $\#18(E_r^{20})$  due to its redundancy to  $\#17(E_r^{10})$ . In addition, we remove the features of joints #3, #5, #6, and #7 due to their low importance. Therefore, we determine the feature set for the development of the signal classifier as  $\mathcal{X}^* = \{M_\tau, S_\tau, D_\tau, E_I, f^*, |S(f^*)|, \phi(f^*), \bar{f}, \}$  $|S(\bar{f})|, \phi(\bar{f}), F_{\text{crest}}, E_r^{10}\}_{1,2,4}$  that contains 36 features, where the subscript denotes the joints. To evaluate  $\mathcal{X}^*$ , we also define a minimum set  $\mathcal{X}^- = \{M_\tau, S_\tau, D_\tau, E_I, f^*, |S(f^*)|, \phi(f^*), \}$  $\bar{f}$ ,  $|S(\bar{f})|$ ,  $\phi(\bar{f})$ ,  $F_{\text{crest}}$ ,  $E_r^{10}$ }, which only includes 12 features on joint #2. We will evaluate the three feature sets  $\mathcal{X}^+$ ,  $\mathcal{X}^*$ , and  $\mathcal{X}^{-}$  and select the best when determining the classifier models. Note that such a development procedure is justified since the entire feature selection process is independent of the classification models.

## C. Model Validation

After determining the feature sets  $\mathcal{X}^+$ ,  $\mathcal{X}^*$ , and  $\mathcal{X}^-$ , we are ready to select the model for the signal classifier. Since a large number of models are used in the literature for the classification of time series, enumerating all of them for the best one is a tedious and unnecessary process. Instead, we assign several popular models with representative properties as candidates and validate these candidate models using the training set, before selecting the best one. In this article, we determine the candidate models based on the three factors, namely, the pHRI signal properties, engineering experience in the previous work,



Fig. 5. Weights of the features obtained by Relieff algorithm. (a) Feature weights grouped by feature numbers. The average weight over all seven joints is marked for each feature. (b) Feature weights grouped by joint numbers. The average weight over all features is marked for each joint.

and the values of models in practice. Thus, four classifier models, namely, a linear discriminant analysis (LDA) model, a k-nearest neighbors (kNN) model, an SVM model, and a feedforward neural network (FNN) model, are chosen as candidate models, due to their simple structures and successful applications in the previous work [38]. These four models are featured with various properties and essentially cover most representative classifier structures. Specifically, the LDA model is well-known for its high efficiency and stable performance and is able to achieve high prediction accuracy for the linearly classifiable data sets. The kNN model is supported by well-developed theoretical foundations and is easy to be extended to a larger scale database in future work. The SVM model shows the excellent ability of generalization. The NN model, as a powerful and comprehensive approximator, has strong adaptability and flexibility for various data sets. In this article, we only consider the NN model with one hidden layer. Note that simple structures are always preferred in practice since the reliability of the system tends to decrease when the system becomes more complicated. To evaluate the four classifier models on the training set with the feature sets  $\mathcal{X}^+$ ,  $\mathcal{X}^*$ , and  $\mathcal{X}^-$ , we use a tenfold cross-validation method and

#### TABLE I

HYPERPARAMETERS FOR GRID-SEARCHING VALIDATION

Models	Hyper-params	Grid values			
LDA	solver	'svd', 'lsqr', 'eigen'			
kNN	k value	$2, 3, \cdots, 80$			
	distance motion	'canberra', 'chebyshev', 'euclidean',			
	distance metrics	'manhattan', 'minkowski'			
SVM	C value	$10^{-3}, 10^{-2.8}, \cdots, 10^{3}$			
	kernel function	'linear', 'rbf', 'sigmoid'			
FNN	hidden neurons	$2, 4, \cdots, 100$			
	activation function	'identity', 'logistic', 'tanh', 'relu'			

#### TABLE II

MODEL CONFIGURATIONS WITH BEST VALIDATION SCORES

Models	Hyper-parameters	$score(\mathcal{X}^+)$	$score(\mathcal{X}^*)$	$score(\mathcal{X}^{-})$
LDA	'svd'	94.8%	92.4%	82.2%
kNN	5, 'canberra'	97.5%	96.3%	88.0%
SVM	63.1, 'linear'	97.3%	96.0%	86.4%
FNN	48, 'logistic'	96.8%	96.6%	89.7%

calculate the mean values of the ten validation scores which are briefly referred to as *score*. We also apply a grid search method to seek for the best hyperparameters of the models. The models and the corresponding grid values of their hyperparameters are shown in Table I.

The four models and the corresponding hyperparameter values achieving the best scores in the validation are listed in Table II, where the hyperparameter values are in the same order as the hyperparameter values in Table I. The validation scores of the models trained with various feature sets are presented in the *score* columns,  $\mathcal{X}^+$ ,  $\mathcal{X}^*$ , and  $\mathcal{X}^-$ , respectively, which show that all the four models achieve satisfactory scores, with the FNN model showing the highest (96.6%). Thus, we select FNN with 48 hidden-layer neurons, with "logistic" activation function as the model for the signal classifier of the CDI scheme. In the meantime, the results also reveal that the scores using  $\mathcal{X}^*$  are close to that using the full feature set  $\mathcal{X}^+$  for all models, despite slight decreases. On the contrary, the scores of the minimum feature set  $\mathcal{X}^-$  drastically drop compared to  $\mathcal{X}^+$   $\mathcal{X}^*$ . Therefore, we determine  $\mathcal{X}^*$  as the feature set to train the FNN model since it achieves similar scores to the full feature set  $\mathcal{X}^+$  with a greatly reduced number of features.

## D. Model Test

After determining the model for the signal classifier (FNN, 48 hidden-layer neurons, "logistic" activation function), we train the model using all samples in the training set with the feature set  $\mathcal{X}^*$  and evaluate its prediction accuracy using the test set. The tested accuracy score, 96.5%, is generally comparable with the score 96.6% in the model



Fig. 6. Confusion matrix in the test of the well-trained FNN classifier.

validation in Section IV-C, which indicates that the signal classifier possesses a satisfactory ability of generalization. The confusion matrix of the test is presented in Fig. 6, where the prediction accuracy for *collisions*, *contacts*, and *free*, respectively, achieves as high as 95.6%, 93.7%, and 98.4%, thus showing a decent performance of the classifier. Especially, the higher accuracy rate of *collisions* than *contacts* reveals that the classifier is more sensitive to accidental *collisions*, which is reasonable since *collisions* are more dangerous to humans than *contacts*. The high-accuracy score for *free* means that the classifier hardly produces false-positive results.

## V. DESIGN OF THE ONLINE DIAGNOSOR

Although the signal classifier shows high accuracy for samples with complete pHRI waveform, the same performance is not likely to be ensured in the early stage of a pHRI in online applications since the segmented waveform is incomplete. As discussed in Section I, the main reason is the distribution inconsistency between the training set and the segmented online signals, such that a single prediction result of the classifier does not guarantee a reliable CDI before the pHRI vanishes. To resolve this issue, we design an additional online diagnosor for the CDI scheme to produce online decisions based on a series of prediction results, which is proven to be more reliable than using a single result.

## A. Prediction Analysis of Incomplete pHRI Waveform

As the first step in designing the online diagnosor, we investigate the prediction accuracy of the signal classifier for signal samples with incomplete pHRI waveform. We create several test sets from the raw signals with various b values, which represent the segmented online signals with a different proportion of pHRI waveform, and use them to evaluate the prediction accuracy of the signal classifier. Note that all the test sets are of the same size as the one in Section IV-D. The accuracy scores with different b values are illustrated in Fig. 7.

As a general tendency, the prediction accuracy increases when the segment bias b becomes larger. An intuitive explanation is that more useful information on the pHRI waveform



Fig. 7. Accuracy scores of the signal classifier when predicting samples with various b values.

naturally leads to higher accuracy scores. Within the interval  $b \leq 50$ , the prediction accuracy is lower than 77.3%, which indicates that a prediction result within 50 ms after the pHRI is very likely to be incorrect. In contrast, when b > 200, the accuracy is higher than 93.0%, which reveals that an accurate result is only possible after 200 ms. Apparently, a single prediction result of the signal classifier does not offer a reliable CDI decision in the early stage of pHRI.

### B. Diagnosis Using Confidence Indexes

Our solution to this problem is to extend the decision horizon, such that the CDI result  $C^*$  is produced based on a series of successive prediction results. Moreover, a quantified metric is need to depict the reliability of the CDI result. Let us assume that the pHRI occurs at time  $t_0$ . At a certain time instant  $t_1 = t_0 + b$ , a segmented torque signal  $\mathcal{T}_1$  is given to the signal classifier which then reports a prediction  $r_1$ . The value of the prediction result  $r_1$  can be -1, 0, or 1, which corresponds to collision, contact, and free. Here,  $r_1$  is not directly used to make the decision  $C^*$ . Instead, we postpone s sampling intervals until we obtain s signal segments  $T_1, T_2, \ldots, T_s$  and a series of prediction results  $\mathcal{R} =$  $\{r_1, r_2, \ldots, r_s\}$ . Here, we refer to  $\mathcal{R}$  as an *observation* series with the window size s. According to the Bayesian decision theory, the posterior probability of the class  $C = \{-1, 0, 1\}$ based on the observation series  $\mathcal{R}$  with parameter b reads

$$p(C|\mathcal{R}, b) = \frac{p(C|b)p(\mathcal{R}|C, b)}{p(\mathcal{R}|b)}$$
(1)

where p(C|b) is the prior knowledge of proportions of the data set,  $p(\mathcal{R}|C, b)$  is the likelihood of class *C*, and  $p(\mathcal{R}|b)$  is the evidence that adjusts the weight of the posterior probabilities, where  $p(\mathcal{R}|b) = \sum_{C} p(C|b)p(\mathcal{R}|C, b)$ . As mentioned in the beginning of Section IV, the signal segments  $\mathcal{T}_1, \mathcal{T}_2, \ldots, \mathcal{T}_s$  are independent of each other, which leads to a series of independent prediction results  $r_1, r_2, \ldots, r_s$ . Therefore, the likelihood  $p(\mathcal{R}|C, b)$  can be calculated by the following product:

$$p(\mathcal{R}|C,b) = \prod_{j=1}^{s} p(r_j|C,b_j)$$
(2)

where for each  $r_j$ , j = 1, 2, ..., s,  $p(r_j|C, b_j)$  represents the likelihood of a single prediction result  $r_j$  with given pHRI class *C*. The parameter  $b_j$  is the bias of segment  $\mathcal{T}_j$ , reflecting a hidden condition that  $r_j$  is the *j*th element of  $\mathcal{R}$ . The likelihood  $p(r_j|C, b_j)$  depicts the probability that the classifier predicts a sample labeled *C* as  $r_j$ , which depends on the capability of the classifier.

Here, the posterior probability  $p(C|\mathcal{R}, b)$  serves as a confidence index for decision *C* given observation series  $\mathcal{R}$ . It indicates the belief on the classifier to produce an accurate diagnosis result within *b* ms after the occurrence of the pHRI. A larger value of  $p(C|\mathcal{R}, b)$  corresponds to a more trustful decision *C*. Therefore, the decision law for the online diagnosor can be designed as

$$C^* = i$$
, if  $p(C^{[i]}|\mathcal{R}, b) = \max_{C} p(C|\mathcal{R}, b)$ ,  $i = -1, 0, 1$  (3)

where  $C^*$  is the diagnosis result ( $C^* = -1$  for *contacts*,  $C^* = 1$  for *collisions* and  $C^* = 0$  for *free*) and  $C^{[i]}$  represents the event C = i for brevity. The confidence index for decision  $C^*$  is  $\mathcal{I}_{\mathcal{R}}^{C^*} = p(C^*|\mathcal{R}, b)$ .

To calculate the posterior probabilities, the likelihoods  $p(r_i|C, b_i)$  for all  $r_i = \{-1, 0, 1\}$  and  $C = \{-1, 0, 1\}$  can be approximated by experimental methods similar to that in Section V-A. The difference is that the calculation of the likelihoods requires the values in the confusion matrices, rather than the overall accuracy scores. Such an approximation is justified by the large number principle. The observation window size s can be determined accordingly. A wider window leads to a higher resolution of the confidence index and provides more flexibility to the collision handling pipeline, which, on the other hand, involves more computational load. Note that the value b, as a parameter to be assigned, represents a conservative estimation, rather than the true diagnosis delay. A reliable CDI diagnosis is guaranteed in the early stage of a pHRI, as long as a high confidence index is obtained for even small values of b. Therefore, in the practical design of the CDI scheme, b can be heuristically determined as a reference reaction time. By designing the decision law (3) with the confidence index  $\mathcal{I}$ , the collision handling pipeline can be implemented in a more flexible manner.

## C. Fast Online Diagnosis Algorithm

The decision law in (3) requires the calculation of the posterior probabilities for every observation series  $\mathcal{R}$ , which is not suitable for online applications due to the large computational load. Here, we propose a fast online diagnosis algorithm (as Algorithm 1) that only involves simple comparison logic but ensures high reliability. In the algorithm, the observation window size is set as s = 5. The main routine of the algorithm is to compare the observation  $\mathcal{R}$  with two reference events  $\mathcal{R}_l^4$  and  $\mathcal{R}_t^2$ , where  $\mathcal{R}_l^4$  means that at least 4 *collisions* are observed in  $\mathcal{R}$ , and  $\mathcal{R}_t^2$  represents at least two *contacts* are witnessed. To evaluate the reliability of Algorithm 1, we calculate its posterior probabilities,  $p(C^{[1]}|\mathcal{R}_l^4, b)$ ,  $p(C^{[1]}|\mathcal{R}_t^2, b)$  and  $p(C^{[1]}|\mathcal{R}_f, b)$  with b = 50, where  $\mathcal{R}_f$  denotes  $\overline{\mathcal{R}}_l^4 \vee \mathcal{R}_t^2$ , i.e., neither a *collision* nor a *contact* occurs. Here,  $\vee$  denotes the union of two events and  $\overline{(\cdot)}$  represents the compliment

Algorithm 1 Fast Online Diagnosor

**Input:**  $\mathcal{R} = \{r_1, r_2, r_3, r_4, r_5\}$  **Output:**  $C^*$ 1: if  $\mathcal{R} \in \mathcal{R}^4_{cls}$  then 2:  $C^* = 1$ ; 3: else if  $\mathcal{R} \in \mathcal{R}^2_{ctc}$  then 4:  $C^* = -1$ ; 5: else 6:  $C^* = 0$ ; 7: end if

of an event. Therefore, the posterior  $p(C^{[1]}|\mathcal{R}_l^4, b)$  represents the confidence index of Algorithm 1 when making a decision C = 1 given  $\mathcal{R} \in \mathcal{R}_l^2$ ,  $p(C^{[1]}|\mathcal{R}_l^4, b)$  is for decision C = -1with  $\mathcal{R}_t^2$ , and the else is depicted by  $p(C^{[0]}|\mathcal{R}^f, b)$ , which describes the reliability of the algorithm in *free* cases.

To calculate the posteriors, we first use (2) to calculate the following likelihoods:

$$p(\mathcal{R}_{l}^{4}|C^{i}, b_{j}^{[50]}) = \sum_{\mathcal{R}\in\mathcal{R}_{l}^{4}} \prod_{j=1}^{s} p(r_{j}|C^{i}, b_{j}^{[50+j]}), \quad i = [1], \overline{[1]}$$

$$p(\mathcal{R}_{l}^{2}|C^{i}, b_{j}^{[50]}) = \sum_{\mathcal{R}\in\mathcal{R}_{l}^{2}} \prod_{j=1}^{s} p(r_{j}|C^{i}, b_{j}^{[50+j]}), \quad i = [-1], \overline{[-1]}$$

$$p(\mathcal{R}_{f}|C^{i}, b_{j}^{[50]}) = \sum_{\mathcal{R}\in\mathcal{R}_{f}} \prod_{j=1}^{s} p(r_{j}|C^{i}, b_{j}^{[50+j]}), \quad i = [0], \overline{[0]}$$

where  $b^{[50]}$  means b = 50 and  $C^{\overline{[1]}}$  denotes the event that  $C \neq 1$ , i.e.,  $C^{\overline{[1]}} = C^{[-1]} \vee C^{[0]}$ , while similar is  $C^{\overline{[-1]}}$ . Such a compact form is for the purpose of brevity. We assume that  $p(r_j|C^{[1]}, b_j^{[50+j]}) \approx p(r_j|C^{[1]}, b_j^{[50]})$ , for all j = 1, 2, ..., 5, considering a high sampling rate of the system. Therefore, the likelihoods are calculated as follows:

$$p(\mathcal{R}_{l}^{4}|C^{i}) = p(r^{[1]}|C^{i})^{5} + 5 \ p(r^{[1]}|C^{i})^{4} \ p(r^{\overline{[1]}}|C^{i}), \qquad (4a)$$

$$p(\mathcal{R}_{l}^{2}|C^{i})$$

$$= 1 - p(r^{\overline{[-1]}}|C^{i})^{5} - 5 \ p(r^{\overline{[-1]}}|C^{i})^{4} \ p(r^{[-1]}|C^{i})$$
(4b)

$$p(\mathcal{R}_{f}|C^{i}) = p(r^{[0]}|C^{i})^{5} + 5p(r^{[0]}|C^{i})^{4}p(r^{[1]}|C^{i}) + 10p(r^{[0]}|C^{i})^{3}p(r^{[1]}|C^{i})^{2} + 10p(r^{[0]}|C^{i})^{2}p(r^{[1]}|C^{i})^{3} + 5p(r^{[-1]}|C^{i}) \times (p(r^{[0]}|C^{i})^{4} + 4p(r^{[0]}|C^{i})^{3}p(r^{[1]}|C^{i}) + 6p(r^{[0]}|C^{i})^{2}p(r^{[1]}|C^{i})^{2} + 4p(r^{[0]}|C^{i})p(r^{[1]}|C^{i})^{3})$$
(4c)

respectively, for  $i = [1], \overline{[1]}, i = [-1], \overline{[-1]}$ , and  $i = [0], \overline{[0]}$ , where  $b^{[50]}$  and the subscript j for  $r_j$  are omitted for brevity. Similar to above, we use  $r^{[1]}$  and  $r^{\overline{[1]}}$  to represent r = 1and  $r \neq 1$ . To obtain the values of the single-prediction likelihoods on the right sides of (4), we evaluate the signal classifier using the test set with b = 50 (ms) and calculate the confusion matrix which is decomposed to three matrices in Table III, respectively, for the calculation of (4a)–(4c). The rows of Table III  $C^{[i]}$ , i = -1, 0, 1 represent the ground truth,

#### TABLE III

Confusion Matrices of Signal Classifier With Test Set b = 50 ms

p(r C)	$r^{[1]}$	$r^{\overline{[1]}}$	p(r C)	$r^{[-1]}$	$r^{[-1]}$	p(r C)	$r^{[0]}$	$r^{[-1]}$	$r^{[1]}$
$C^{[1]}$	1207 (72.5%)	458 (27.5%)	$C^{[-1]}$	593 (31.9%)	1268 (68.1%)	$C^{[0]}$	3376 (99.4%)	20 (5.89%)	3 (0.09%)
$C^{\overline{[1]}}$	13 (0.25%)	5247 (99.7%)	$C^{\overline{[-1]}}$	419 (8.27%)	4645 (91.8%)	$C^{\overline{[0]}}$	1317 (37.4%)	992 (28.2%)	1217 (34.6%)

while the columns  $r^{[i]}$  denote the predicted classes. In each block, both the number of samples and the accurate scores are presented, which explicitly correspond to the values of the likelihoods p(r|C). For example, the value of  $p(r^{[1]}|C^{[1]})$  is approximated as 0.73 corresponding to the value in  $C^{[1]}$ ,  $r^{[1]}$ .

Checking the values in Table III and using (4), we calculate the values of the likelihoods as

$$p(\mathcal{R}_{l}^{4}|C^{[1]}) = 0.87, \quad p(\mathcal{R}_{l}^{4}|C^{\overline{[1]}}) = 9.80 \times 10^{-5}$$
$$p(\mathcal{R}_{l}^{2}|C^{[-1]}) = 0.51, \quad p(\mathcal{R}_{l}^{2}|C^{\overline{[-1]}}) = 6.74 \times 10^{-2}$$
$$p(\mathcal{R}_{f}|C^{[0]}) = 0.99, \quad p(\mathcal{R}_{f}|C^{\overline{[0]}}) = 0.52.$$

Here, we set the prior probabilities as  $p(C^{[0]}) = 0.5$ ,  $p(C^{[1]}) = p(C^{[-1]}) = 0.25$ , according to the proportion of samples in the data set. Therefore, according to the Bayes law in (2), we calculate the confidence indexes for Algorithm 1 as follows:

$$p(C^{[1]}|\mathcal{R}_{l}^{4}) = \frac{p(C^{[1]})p(\mathcal{R}_{l}^{4}|C^{[1]})}{\sum_{i=[1]}^{\overline{[1]}} p(C^{i})p(\mathcal{R}_{l}^{4}|C^{i})}$$

$$p(C^{[-1]}|\mathcal{R}_{t}^{2}) = \frac{p(C^{[-1]})p(\mathcal{R}_{t}^{2}|C^{[-1]})}{\sum_{i=[-1]}^{\overline{[-1]}} p(C^{i})p(\mathcal{R}_{t}^{2}|C^{i})}$$

$$p(C^{[0]}|\mathcal{R}_{f}) = \frac{p(C^{[0]})p(\mathcal{R}_{f}|C^{[0]})}{\sum_{i=[0]}^{\overline{[0]}} p(C^{i})p(\mathcal{R}_{f}|C^{i})}$$

which leads to  $p(C^{[1]}|\mathcal{R}_l^4) = 0.99$ ,  $p(C^{[-1]}|\mathcal{R}_t^2) = 0.72$ , and  $p(C^{[0]}|\mathcal{R}_f) = 0.66$ .

The results of the posterior probabilities indicate that Algorithm 1 produces high confidence on its diagnosis for *collisions* ( $\mathcal{I}_{\mathcal{R}}^1 = 0.99$ ) and *contacts* ( $\mathcal{I}_{\mathcal{R}}^{-1} = 0.72$ ) with given observation series  $\mathcal{R}$ . A larger value of  $p(C^{[1]}|\mathcal{R}_l^4)$  than  $p(C^{[-1]}|\mathcal{R}_l^2)$  means that the diagnosor is more sensitive to *collisions* than *contacts*. The confidence index  $\mathcal{I}_{\mathcal{R}}^0 = 0.66$  for *free* reveals that the diagnosor is also trustful for avoiding false detection. Due to the fact that  $p(C^{[1]}|\mathcal{R}_l^4) = \max_C p(C|\mathcal{R}_l^4)$ ,  $p(C^{[-1]}|\mathcal{R}_l^2) = \max_C p(C|\mathcal{R}_l^2)$ , and  $p(C^{[0]}|\mathcal{R}_f) =$  $\max_C p(C|\mathcal{R}_f)$ , Algorithm 1 is consistent with the decision law in (3). Therefore, the confidence index  $\mathcal{I}$  can be explicitly obtained without calculation, which leads to a faster and easier implementation for online applications.

## D. Comparison With the Single Prediction

To justify the advantage of the online diagnosor using observation series instead of a single prediction result, we also calculate the confidence indexes for the latter for comparison. The confidence index for class  $C^i$ , with a single observation  $r^i$ , is calculated by

$$p(C^{i}|r^{i}) = \frac{p(C^{i})p(r^{i}|C^{i})}{\sum_{k=i}^{\overline{i}} p(C^{k})p(r^{k}|C^{k})}, \quad i = [-1], [0], [1].$$

Checking the likelihoods in Table III, we obtain  $p(C^{[-1]}|r^{[-1]}) = 0.57$ ,  $p(C^{[0]}|r^{[0]}) = 0.75$ ,  $p(C^{[1]}|r^{[1]}) = 0.99$ . The results show that such a scheme ensures a low reliability on identifying *contacts* (0.57), i.e., the diagnosor is sensitive to *collisions* but quite dull to *contacts*, which shows an inferior performance than Algorithm 1.

#### VI. EXPERIMENTAL VALIDATION

In this section, we evaluate the performance of the proposed CDI scheme in terms of both the identification accuracy and the responsiveness by applying it to an online validation on the recorded raw signals. These signals contain 242 collisions and 225 contacts and have never been used during the development of the signal classifier. The torque signal is sequentially sampled at the rate of 1 kHz and recursively segmented as new samples for the signal classifier. The online diagnosor is equipped with the fast online diagnosor in Algorithm 1 to provide online diagnosis. Note that the evaluation method and metrics in this experiment are different from the model test in Section IV-D. For the classifier development, the accuracy scores are calculated based on the number of signal segments containing the pHRI waveform, while in a human-robot collaboration, a CDI scheme focuses more on the number of correctly identified collisions or contacts. The main difference is that the pHRI, in an online application, usually produces several segments with various b values, corresponding to the signals segmented in different sampling instants. Therefore, for the experimental validation, we must clarify that we evaluate the performance of the CDI scheme with respect to the entire pHRI, rather than its segments. We recognize that the pHRI is accurately identified, if the scheme proposes a correct diagnose before it vanishes.

During the experiment, 240 out of 242 collisions and all 225 interactions are correctly identified, with two collisions misclassified as one contact and one free, achieving an overall accuracy 99.6%. Some identification instances are shown in Fig. 8, which illustrates the accuracy and responsiveness of the scheme. It is noticed that most of the diagnosis results are correctly produced in the early stages of the pHRI, which confirms the high accuracy of the online CDI scheme. The fast response of the scheme is also clearly reflected in Fig. 9 with both instances of a *collision* and a *contact*. The two pHRIs





Fig. 8. Experimental results of the online CDI scheme, respectively, for collisions and contacts. (a) CDI diagnosis of collisions. Red solid lines: raw signals. Black dotted lines: occurrence instants of the collisions. Gray regions: time instants when collisions are detected and identified. (b) CDI diagnosis of contacts. Red solid lines: raw signals. Black dotted lines: occurrence instants of the contacts. Blue regions: time instants when contacts are detected and identified.



Fig. 9. Diagnosis results of a collision and a contact. (a) CDI diagnosis of a collision. Red solid lines: raw signals. Black dotted lines: occurrence instants of the collisions. Gray regions: time instants when collisions are detected and identified. (b) CDI diagnosis of a contact. Red solid lines: raw signals. Black dotted lines: occurrence instants of the contacts. Blue regions: time instants when contacts are detected and identified.

are all correctly identified within 20 ms after their occurrence, which confirms the responsiveness of the scheme.

To demonstrate the applicability of the proposed online CDI scheme in a practical task, we implement it on the

Fig. 10. Application demonstration of the online CDI diagnosis to the KUKA manipulator platform. The CDI scheme identifies collisions and interactions, such that different reaction routines are activated. (a) Collision is exerted. (b) Emergency stop. (c) Interaction is exerted. (d) Guidance mode.

Kuka LWR4+ robot platform (as shown in Fig. 10) together with a collision reaction routine. The task is intended to survey the external torques on the robot joints, detect any pHRI, and activate the corresponding reaction strategies. During a robot task routine, if a *collision* is detected [as Fig. 10(a)], an emergency stop is triggered and the robot holds on the current position until the safety alarm is deactivated [as Fig. 10(b)]. If a *contact* is detected [as Fig. 10(c)], the robot enables its gravity compensation mode such that the human can conduct active guidance on the robot [as Fig. 10(d)].

During the operation of the robot, the CDI scheme correctly detects and identifies most of the pHRI and enables the corresponding reaction procedure although some week contacts and collisions fail to be detected. When the robot is smoothly moving, the false diagnosis of collisions and contacts is hardly witnessed. Nevertheless, a false diagnosis occurs when the robot motion shows large vibrations due to fierce motion or bad controller design. Therefore, a smooth functioning condition is still necessary. As for the responsiveness, the diagnosis delay is basically not obviously perceivable experimenters, which is acceptable in a typical human–robot collaborative task. Therefore, the demonstration confirms the performance of the proposed CDI and indicates its applicability to human–robot collaboration tasks in practice.

### VII. CONCLUSION

In this article, we develop a novel online CDI scheme for robot manipulators using supervised learning and the Bayesian decision theory. During the data collection process, we try to cover various execution conditions of the robot to ensure a representative data set. After carefully investigating the properties of the pHRI signals, we extract and select the features by analyzing their dependence and importance. As a result, the signal classifier presents excellent predictability and generalization ability. Based on this, we propose a fast and simply implementable online diagnosor with a quantified confidence index to depict the reliability of the diagnosis result based on the Bayesian decision theory. Examined by both theoretical analysis and experimental validation, the proposed CDI scheme shows a promising value to improve the safety of human–robot collaboration systems.

In general, our work in this article has achieved the target raised in the beginning, i.e., to rapidly and correctly detect and identify collisions in their early stages. As a result, a collision is *predicted* before it vanishes, such that further damages or injuries can be avoided. The common ground of our work and the probabilistic-series-model-based methods is to make a decision using the observation series. Nevertheless, the proposed CDI scheme in this article is more suitable to be widely applied in practice since it does not require prior knowledge of the signal dependence and assumptions on the data distribution.

It should be noted that the development of the CDI scheme, including the procedures of the data collection, feature engineering, and the model training, is conducted on a specific robot platform. Thus, the applicability of the developed CDI scheme to different platforms needs further investigations, which motivates our future work on the adaptation of the CDI scheme to a wider range of platforms. Another interesting topic is to compensate for the effects of the varying robot loads, which can be solved by dynamic-model-based disturbance estimation methods [9]. To investigate the influence of the individual uncertainties on the classification accuracy, we will also collect collision data with more experimental subjects in future work.

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