Post-mortem Iris Recognition with Deep-Learning-based Image Segmentation

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Abstract

This paper proposes the first known to us iris recognition methodology designed specifically for post-mortem samples. We propose to use deep learning-based iris segmentation models to extract highly irregular iris texture areas in post-mortem iris images. We show how to use segmentation masks predicted by neural networks in conventional, Gabor-based iris recognition method, which employs circular approximations of the pupillary and limbic iris boundaries. As a whole, this method allows for a significant improvement in post-mortem iris recognition accuracy over the methods designed only for ante-mortem irises, including the academic OSIRIS and commercial IriCore implementations. The proposed method reaches the EER less than 1% for samples collected up to 10 hours after death, when compared to 16.89% and 5.37% of EER observed for OSIRIS and IriCore, respectively. For samples collected up to 369 hours post-mortem, the proposed method achieves the EER 21.45%, while 33.59% and 25.38% are observed for OSIRIS and IriCore, respectively. Additionally, the method is tested on a database of iris images collected from ophthalmology clinic patients, for which it also offers an advantage over the two remaining methods. This work is the first step towards post-mortem-specific iris recognition, which increases the chances of identification of deceased subjects in forensic investigations. The new database of post-mortem iris

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images acquired from 42 subjects, as well as the deep learning-based segmentation models are made available along with the paper, to ensure all the results presented in this manuscript are reproducible.

Keywords: biometrics, iris recognition, post-mortem, image segmentation

1. Introduction

Post-mortem iris recognition has recently gained attention from the biometrics community, starting with a Master's thesis from Boston University [1], and studies on *ex vivo* pig eyes in 2015 [2]. Since then, some research groups have been hard at work evaluating how this biometrics works with irises of decased subjects [3, 4, 5, 6, 7], and proposing PAD (Presentation Attack Detection) techniques aimed at detecting cadaver iris presentations to the sensor [8, 9]. Post-mortem iris recognition is also discussed as a potentially useful method for forensic proceedings in addition to typical forensic biometric characteristics such as fingerprints, if the method could be applied reliably [7].

Despite this attention, concluded with many claims that post-mortem iris biometrics, however challenging, may still be possible in favorable conditions, no specific, cadaver-aware methods for improving the recognition reliability have been proposed, apart from the work of Trokielewicz *et al.* [10], who introduced a data-driven semantic segmentation model for iris localization within images collected from cadaver eyes, yet without evaluating how the newly created segmentation algorithm improves the recognition accuracy.

In this work, we build upon the findings presented in [10], by training several additional models with larger amounts of post-mortem data and different ground truth mask creation rules, together with constructing an iris image normalization method based on the circular Hough transform (CHT). This end-to-end segmentation algorithm is then coupled with the well known open source iris recognition method OSIRIS [11], and evaluated against the baseline OSIRIS performance and against the commercial Iri-Core method, which presented the best performance of post-mortem iris matching in past studies [7]. **The contributions that this study makes towards the state-of-the-**

art in post-mortem iris recognition are thus the following:

- introduction of two new, deep convolutional neural network-based (DCNN-based)
 post-mortem iris image segmentation methods, trained with different ground
 truths, namely coarse iris approximation, and fine-grained, post-mortem specific
 masking of the regions affected by tissue decay,
- an end-to-end iris localization and image segmentation model that can be used as
 a drop-in replacement for any iris recognition methodology that uses normalized
 images with iris boundary approximated by circles,
- a new dataset of cadaver iris images collected from 42 subjects over a time period of up to 369 hours post-mortem,
- experiments showing a considerable improvement in the matching performance of a method employing the proposed segmentation and Gabor wavelet-based encoding,
- source codes and neural network models' weights, which allow for full reproducibility of the results, as well as incentive for further research.

This article is laid out as follows. After the introductory part, Sec. 2 discusses the historical assumptions and the current course of post-mortem iris recognition research. Biometric databases used for training and estimation, as well as evaluation of the proposed approach are introduced in Sec. 3, whereas in Sec. 4 we describe the methodology of our solution and ways for its evaluation, which is performed and reported in Sec. 5. Finally, Sec. 6 contains relevant conclusions.

2. Related work

Even before any meaningful experimental studies regarding post-mortem iris recognition have come to fruition, both the research community and the industry were sceptical about the feasibility of post-mortem iris recognition [12, 13, 14].

The first systematic study in this field was Sansola's MSc thesis at Boston University [1], in which the IriShield M2120U iris recognition camera and IriCore matching

software were employed to photograph irises of 43 deceased subjects at different postmortem time intervals. The method yielded 19-30% of false non-matches and no false matches, depending on the time since death involved. Later on, Saripalle *et al.* [2] studied ex-vivo eyes of domestic pigs and their biometric capability during the degradation of tissues after they are taken out of the cadaver. They found that the irises lose their biometric capabilities 6 to 8 hours after death, proving the fast rate of ex-vivo eye degradation.

In our previous work we refuted most of the popular claims, showing that the iris can still successfully serve as a biometric identifier for 27 hours after death [3], with the pupil typically remaining mid-sized without any excessive dilation or constriction, with well visible iris structure. More than 90% of the irises were correctly recognized when photographed a few hours after death, including 100% accuracy obtained for the IriCore method (which comes from the same vendor as the camera used in this study). As time after death increases, the equal error rate increases to 13.3% when images captured approximately 27 hours after death are compared against those obtained 5h after demise. Later, we showed that correct matches can still be expected even after 17 days [4] and offered the first known to us database of 1330 NIR and VIS post-mortem iris images acquired from 17 cadavers [15].

A study tracking biometric capabilities of both the typically used forensic modalities, namely fingerprints and faces, as well as iris is presented by Bolme *et al.* [5]. The authors placed twelve deceased subjects in outdoor conditions to assess how the environment and time affect the biometric performance. Irises were found to be degrading very fast and become unusable as shortly as few days after the body placement. When the bodies were kept outside for 14 days, the correct verification rate reported by the authors decreased to 0.6%, but the real-life chance of recognizing an iris is estimated to be less than 0.1%. Fingerprints and faces, on the other hand, remained readable for much longer periods of time.

Sauerwein *et al.* [6] showed that irises stay readable for up to 34 days after death, when cadavers were kept in outdoor conditions during winter. This readability parameter, however, was assessed by human experts acquiring the samples and no automatic iris recognition algorithms were used.

The most recent paper in this study by Trokielewicz *et al.* [7] included the most comprehensive experiments in this field to date, and introduces the new dataset of cadaver iris images [16], which in combination with the first part of the Warsaw corpus aggregates 1,200 near-infrared and 1,787 visible light samples collected from 37 deceased subjects kept in mortuary conditions. In the foretold paper, we employed four iris recognition methods to analyze the genuine and impostor comparison scores and evaluate the dynamics of iris decay over a period of up to 814 hours. The findings show that iris recognition may be close-to-perfect 5 to 7 hours after death and occasionally is still viable even 21 days after death. We also demonstrated that false-match probability is significantly higher when live iris images are compared with post-mortem samples than when only live samples are used in comparisons.

3. Databases of iris images

For the purpose of this work, we use four different databases of iris images, including three subject-disjoint sets of post-mortem iris images, namely the:

- Warsaw-BioBase-Postmortem-Iris-v1.1, comprising 574 near-infrared (NIR) and 1023 visible light (VIS) images collected from 17 cadavers over a period of up to 34 days [15, 4],
- Warsaw-BioBase-Postmortem-Iris-v2, a subject-disjoint extension of the v1.1 of the database, adding 626 NIR and 764 VIS data from 20 more cadavers [16, 7],
- Warsaw-BioBase-Postmortem-Iris-v3, a new set of images collected for the purpose of this study, adding data from 40 more subjects with a total of 1094 NIR images and 785 VIS images, collected over a time horizon of up to 369 hours since demise.

The first two datasets are aggregated together and used as training data, whereas the third set is employed for evaluation purposes. This way, the training and testing data are subject-disjoint, *i.e.*, data from subjects 'participating' in the training of the segmentation models and estimation of the normalization algorithm's parameters is not repeated in the evaluation experiments.

In addition to the post-mortem data, we use another database of challenging iris images for supplemental testing of the proposed approach:

• Warsaw-BioBase-Disease-Iris-v1, comprising iris images collected mostly from elderly patients of an ophthalmology clinic, including subjects with various ophthalmic conditions; we use a subset of 551 NIR images from 76 randomly chosen eyes [17, 18].

Following the findings reported in [7] that strongly favor the NIR images for postmortem iris recognition purposes, we only employ this type of images in all experiments conducted in this work. Example images from each of the evaluation databases are shown in Fig. 1.



Figure 1: Example images from the test databases used in this work. Warsaw-BioBase-Postmortem-Iris-v3 (**left**) and Warsaw-BioBase-Disease-Iris-v1 (**right**.)

4. Proposed methodology

4.1. Baseline iris recognition method

OSIRIS, or *Open Source for IRIS*, is an academic solution developed as a part of the BioSecure EU project [11] and implements the original iris recognition concept of John Daugman, including iris image segmentation and normalization to dimensionless polar coordinate system, followed by the calculation of a binary iris code using phase quantization of the Gabor wavelet filtering outcomes. Fractional Hamming distance calculated using the XOR operation between the two codes is used as a comparison

score, with values close to zero expected for two same-eye images, whereas scores oscillating around 0.5 should be produced when two different-eye images are compared. Due to the additional compensation of the eyeball rotation, different-eye score distributions are typically skewed toward 0.4 - 0.45 range.

The original score calculation method can be complemented by score normalization penalizing comparisons with small number of compared bits, as compared to the *typical* (here: within a given dataset) number of compared bits, as proposed by Daugman [19]:

$$HD_{norm} = 0.5 - (0.5 - HD_{raw})\sqrt{\frac{n}{N}}$$

This transforms the samples of scores obtained when comparing different eyes into samples drawn from the same binomial distribution, as opposed to drawing sample scores from different binomial distributions with σ dependent on the number of bits n that were available for comparison (commonly unmasked bits). N is the 'typical number of bits compared (unmasked) between two different irises,' which Daugman states to be 911 or 960, depending on the data. The N parameter is estimated for a particular database of iris images. Since this work uses different datasets of iris images and different subsets of these datasets (*i.e.*, post-mortem samples with varying time horizon), the N is calculated separately for each experiment (cf. description in Sec. 5). The total number of bits in the OSIRIS code is 1536.

In the following Sections we will evaluate the proposed approach and discuss whether applying this normalization procedure is beneficial for the overall system's performance.

4.2. Data-driven image segmentation models

The first stage of the proposed scheme consists of employing a data-driven segmentation model based on a deep convolutional neural network (DCNN) for the purpose of localizing the iris within an image. Our models are based on the solution described in one of our previous papers [10], which employs a re-trained off-the-shelf SegNet model [20], and offers a large improvement over the OSIRIS segmentation method measured with the IoU (Intersection over Union) metric. We fine-tuned it with a database of cadaver iris images and their corresponding ground truth binary masks. The masks used

for training were annotated in a fine-grained manner, which has a purpose of teaching the network to only localize iris regions that are not affected by post-mortem changes, and represent the well-visible iris texture.

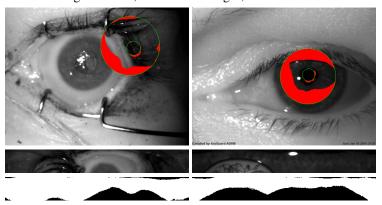
In addition to the model described above, in this work we have implemented and trained two additional models, ending up with three models in total, which are detailed below and their example predictions given in Fig. 2

- fine segmentation model: the one published in [10], trained with data from the Warsaw-Biobase-Postmortem-Iris-v1 database with fine-grained ground truth masks and yielding predictions of 120 × 160 pixels,
- 2. fine v2highres segmentation model: similar to the fine model, but trained with data from Warsaw-Biobase-Postmortem-Iris-v1 database together with NIR samples from the Warsaw-Biobase-Postmortem-Iris-v2 database for twice as many epochs (120 vs 60), also with fine-grained ground truth masks, but yielding higher-resolution predictions, namely 240 × 320 pixels,
- 3. *coarse* segmentation model: trained with both NIR and VIS data from the v1 and v2 versions of the Warsaw-Biobase-Postmortem-Iris database for 120 epochs, but this time with **coarse ground truth masks**, denoting only the inner and the outer iris boundary and eyelids, also producing masks in 240 × 320 size.

4.3. Iris and mask normalization

For the proposed method to serve as a drop-in replacement for the Daugman method recognition pipeline, the prediction obtained from the DCNN-based segmentation has to be correctly normalized onto a dimensionless polar-coordinate rectangle. For this stage, a method for localizing pupillary and limbic iris boundaries has been devised, which employs a Hough transform that is applied to the prediction obtained from the *coarse* segmentation model, as this model yields the smoothest prediction. These boundary parameters are then used in all subsequent experiments, including those involving the *fine* and *fine v2highres* models. The segmentation methodology employing the *coarse* segmentation model is visualized in Fig. 2. Similar results are then presented in Fig. 3 for the two remaining fine-grained models. Notably, the original *fine*

OSIRIS segmentation, normalized images, and normalized masks



coarse CNN binary predictions with fitted Hough circles, segmented images, normalized images, and normalized masks

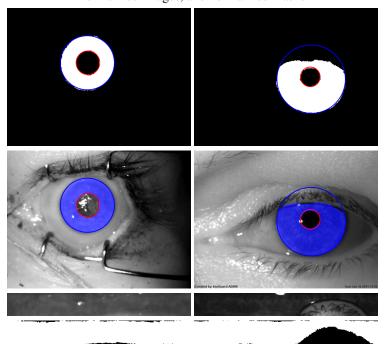
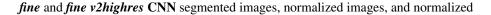


Figure 2: Segmentation results for samples from Fig. 1, obtained from the unmodified OSIRIS method and the *coarse* segmentation model. Warsaw-BioBase-Postmortem-Iris-v3 (**left**), Warsaw-BioBase-Disease-Iris-v1 (**right**).



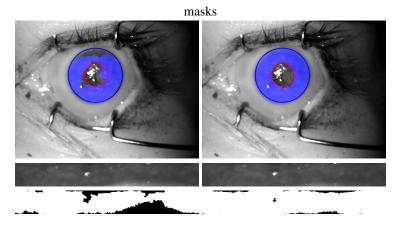


Figure 3: Segmentation results for post-mortem sample from Fig. 2, obtained from the *fine* and *fine v2highres* segmentation models.

model from [10] seems to over-aggressively mask out some portions of the iris, while at the same time mistakenly denoting some portions of the pupil as iris. This is fixed in the *fine v2highres* model, which was trained with twice the iterations count, cf. right side of Fig. 3.

4.4. IriCore: a commercial benchmark method

In addition to evaluating our approach against the unmodified OSIRIS algorithm, we also choose to compare it with one of the leading iris matchers from commercial vendors, namely the IriCore [21], which offered the best performance out of four matchers in our previous evaluations of post-mortem decay impact on various iris recognition methodologies [7]. IriCore also comes from the same manufacturer as the IriShield camera used to collect post-mortem samples in our studies – and therefore it may have an additional advantage in this particular task. Since IriCore follows an undisclosed iris recognition algorithm, and can be only used as a black box taking two images and producing a dissimilarity score between them. The distributions of these dissimilarity scores, which are expected to be ranging from 0 to 1.1 for genuine comparisons and from 1.1 to 2.0 for impostor ones, are then used to construct ROC curves which can be compared against those obtained from original and modified OSIRIS. No

additional score normalization is introduced for IriCore.

5. Iris recognition accuracy evaluation

5.1. Comparison score generation

In this Section we evaluate the new segmentation approach introduced in Sec. 4 by injecting the segmentation results obtained from the three DCNN-based models into the OSIRIS recognition pipeline, and also comparing their performance with unmodified OSIRIS algorithm, as well as with the accuracy offered by the IriCore. The evaluation is performed on two test databases described in Sec. 3. Generation of genuine and impostor comparison score distributions was done by performing comparisons between all possible iris image pairs. One comparison per a given image pair was performed, *i.e.*, if image A was matched against image B, then image B is not matched against image A.

For the post-mortem data, eight separate score distributions are generated for both the genuine and impostor comparisons, taking into account samples collected during different, progressing observation horizons in respect to time that has elapsed since a subject's death. These include scores obtained from samples collected fewer than 10 hours after death, fewer than 24 hours, 48 hours, 60 hours, 110 hours, 160 hours, 210 hours, and finally 370 hours (all available samples). This is to evaluate the recognition accuracy in respect to the increasing post-mortem interval. All possible comparison pairs were evaluated here as well.

5.2. Recognition accuracy: post-mortem dataset

The results are presented in the form of receiver operating characteristic (ROC) curves, which plot the correct match rates against false match rates. In addition, equal error rates (EER) are calculated and denoted in the plot legend. Fig. 6 shows ROC curves denoting the performance of the baseline methods (OSIRIS and IriCore) performance compared against performance obtained by employing image segmentation results from the three DCNN-based models for the post-mortem data in four shortest post-mortem time horizons, comprising samples collected earlier than 60 hours after

the demise of each subject. Raw Hamming distance scores are plotted in solid lines, whereas the scores after score normalization (cf. Sec. 4.1) are plotted with dotted lines. Fig. 7 contains the analogous graphs, but for the remaining four longer time horizons, namely those with samples collected up to 369 hours post-mortem.

Notably, very good performance can be achieved using each one of the proposed models for samples collected up to 10 hours after death, with EER as low as 0.96%, which only slightly increases to 1.36% when samples collected up to 24 hours postmortem are added to the database. For samples collected up to 48 hours (two days) after death, the lowest EER offered by the fine v2highres model is 3.38%, which means that the method can still be usable. As more difficult samples are being added to the database, the errors increase, reaching 21.95% for the best performing fine v2highres model. Although the fine model given better EER, it also presents a risk of higher false match rates than the coarse model (see the region of the ROC curve near the Y axis). This can be alleviated by introducing score normalization (cf. Sec. 4.1), but this in turn increases the EER by almost 3%. The fine v2highres model, on the other hand, seems to offer similar risk of false matches as the fine model with score normalization, but at the same time preserving most of the overall performance with EER only slightly higher than this of the *fine* model. Original OSIRIS, on the other hand, is incomparable to the proposed approach from the very beginning, yielding EER of almost 17% for the 'easiest' samples, and the EERs from this method are increasing steadily up to 33.59% for the entire database (samples collected up to 370 hours post-mortem). The IriCore's performance starts with EER=5.37% for the easiest samples, which is still more than 4 percentage points behind our proposed solution, and decreases steadily with the increasing post-mortem time horizon, reaching 25.38% for the most challenging set of samples (all images). The only advantage the IrisCore may have over our approach is better performance in selected time post-mortem horizons (samples collected up to 60, 110 and 160 hours) in very low FMR registers.

5.3. Recognition accuracy: elderly/ophthalmic conditions dataset

Fig. 4 presents ROC curves obtained when matching samples from the Warsaw-BioBase-Disease-Iris-v1 databases using a baseline OSIRIS approach and the proposed

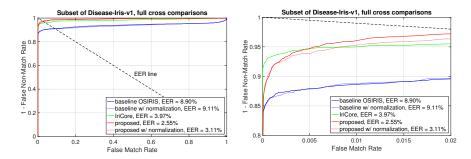


Figure 4: ROC curves obtained for the Warsaw-BioBase-Disease-Iris-v1 dataset. Original (solid lines) and normalized (dotted lines) scores are shown. The plot on the right is a close-up of the one on the left.

segmentation approach based on the *coarse* model. Because this dataset does not contain any post-mortem samples, there is no justification to apply post-mortem-aware fine-grained segmentation model here.

Because this test database contains images collected from elderly people, often with various ophthalmic conditions – also challenging ones, the advantage of the proposed approach is evident, as it offers EER=2.55% compared to 8.9% yielded by OSIRIS and 3.97% obtained from IriCore.

5.4. Discussion of score normalization

As argued in Sec. 4.1, score normalization can help rectify significant portions of false matches that were caused by comparing small numbers of commonly unmasked bits in two iris codes. As seen in Fig. 6, employing this normalization for the very selective segmentation model, namely the old *fine* model (red plots) can vastly reduce the false matches at a little EER-related cost. With less selective segmentation, however, the normalization cannot offer any advantage over lack of such, *e.g.*, in the best performing *fine v2 highres model* (black plots). With increasing time horizons, as in Fig. 7, even the less selective models start to mask out iris portions more aggressively, hence the score normalization can improve the performance in low FMR registers of most of the models.

5.5. False Non-Match Rate dynamics

In addition to the ROCs, we have also calculated the False Non-Match Rate (FNMR) values at acceptance thresholds which allow the False Match Rate (FMR) values to stay below 1%. This is done for the post-mortem data to reveal the dynamics of the FNMR as a function of post-mortem sample capture horizon, and therefore to know the chances for a false non-match as time since death progresses. We plot this dynamics for the two baseline methods: OSIRIS and IriCore, as well as the best performing fine v2highres DCNN-based modification, Fig. 5. Notably, for each moment during the increasing post-mortem sample capture time horizon, our proposed approach consistently offers an advantage over the other two algorithms, allowing to reach nearly perfect recognition accuracy for samples collected up to a day after a subject's death.

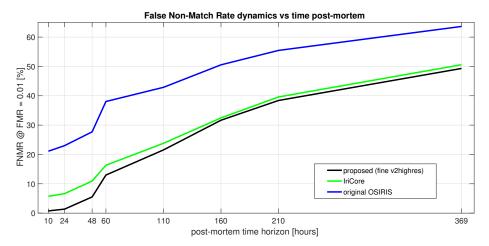


Figure 5: Dynamics of False Non-Match Rates in the function of post-mortem sample capture horizon, plotted for two baseline iris recognition methods, and the approach proposed in this paper.

6. Conclusions

This paper introduces the first known to us attempt at improving iris recognition reliability for post-mortem samples, by proposing a robust image segmentation method that can be used as a drop-in replacement for the OSIRIS image segmentation. The proposed solution, employing DCNN-based semantic segmentation, allows to achieve close-to-perfect recognition accuracy on the subject-disjoint evaluation database of post-mortem samples, as well as on another challenging database of iris images, collected from elderly individuals with various ophthalmic conditions. This proves the flexibility of the proposed approach in solving problems associated with atypical iris images, which the models introduced in this work can reliably and consistently localize and segment in most of the cases. Despite the expected drop in recognition accuracy with the increasing time horizon of post-mortem sample collection, the proposed approach consistently outperforms both the academic and the commercial, state-of-the-art iris recognition methods.

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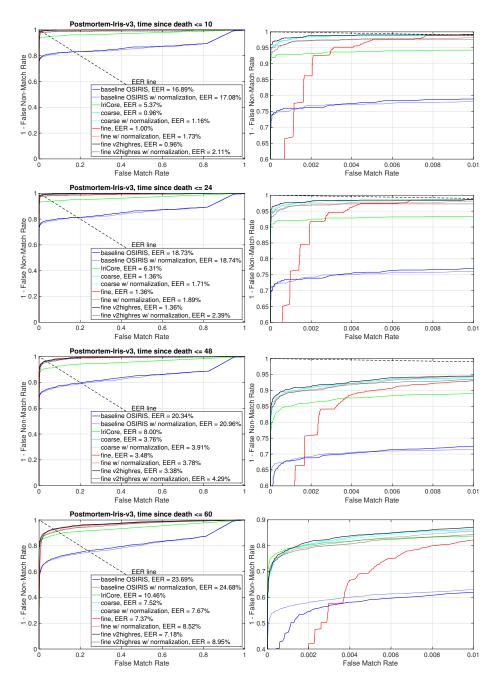


Figure 6: Receiver Operating Characteristic curves obtained when comparing post-mortem samples with different observation time horizons. Results for the baseline OSIRIS performance, IriCore benchmark method, and three modifications introduced by the Authors are presented. Original (solid lines) and normalized (dotted lines) scores are shown. Plots on the right are close-ups of the ones on the left in each pair.

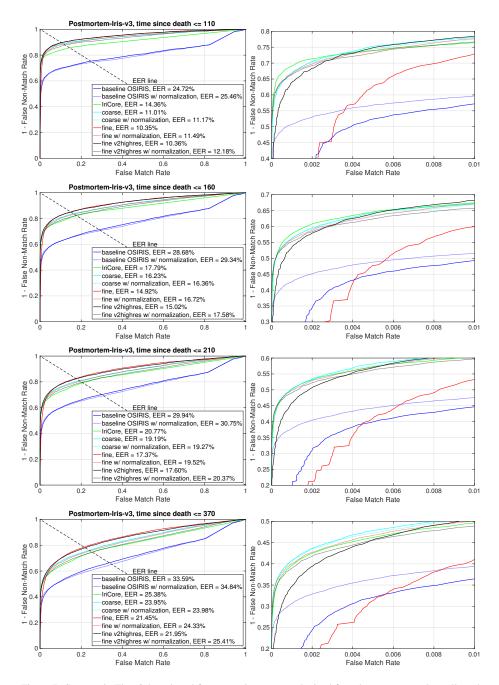


Figure 7: Same as in Fig. 6, but plotted for comparison scores obtained from long-term samples collected from 210 up to 369 hours post-mortem.