Abstract

Despite the benefits introduced by robotic systems in abdominal Minimally Invasive Surgery (MIS), major complications can still affect the outcome of the procedure, such as intra-operative bleeding. One of the causes is attributed to accidental damages to arteries or veins by the surgical tools, and some of the possible risk factors are related to the lack of subsurface visibility. Assistive tools guiding the surgical gestures to prevent these kind of injuries would represent a relevant step towards safer clinical procedures. However, it is still challenging to develop computer vision systems able to fulfill the main requirements: (i) long term robustness, (ii) adaptation to environment/object variation and (iii) real time processing.

The purpose of this paper is to develop computer vision algorithms to robustly track soft tissue areas (Safety Area, SA), defined intra-operatively by the surgeon based on the real-time endoscopic images, or registered from a pre-operative surgical plan. We propose a framework to combine an optical flow algorithm with a tracking-by-detection approach in order to be robust against failures caused by: (i) partial occlusion, (ii) total occlusion, and (iii) SA out of the field of view. A Bayesian inference-based approach is used to detect the failure of the tracker, based on online context information. A Model Update Strategy (MUpS) is also proposed to improve the SA re-detection after failures, taking into account the changes of appearance of the SA model due to contact with instruments or image noise. The performance of the algorithm was assessed on two datasets, representing ex-vivo organs and in-vivo surgical scenarios. Results show that the proposed framework, enhanced with MUpS, is capable of maintain high tracking performance for extended periods of time ($\approx 5\text{ min}$ - containing the aforementioned events) with high precision (0.85) and recall (0.6) values, and with a recovery time after a failure between 1 and 8 frames in the worst case.

Keywords: long-term tissue tracking, tracking failure detection, model update strategy, robotic minimally invasive surgery.

1. Introduction

The introduction of Robotics in Minimally Invasive Surgery (RMIS) allows overcoming many of the obstacles introduced by traditional laparoscopic techniques, by improving the surgeon dexterity and the ergonomics during the surgical procedure, and restoring the surgeon hand-eye coordination (Bravo et al., 2016; Forgione, 2009; Lanfranco et al., 2004). Despite these benefits, the outcome of the surgical procedure can still be compromised by adverse events occurring during the surgery. In robotic abdominal surgery, for example, one of the major complications is intra-operative bleeding due to injuries to vessels (Trinh et al., 2012; Kaouk et al., 2012; Sotelo et al., 2014). Main arteries or veins close to the surgical site can be accidentally damaged during the execution of a surgical procedure, being a major risk factor associated to the surgeon’s skill or robotic system reliability (Lorenzo et al., 2011). Vessel damage may also activate a chain of secondary effects, such as the switch to open-surgery approach, a longer anaesthesia time and post-operative bleeding, thus negatively affecting the surgical performance and leading, in the worst case scenario, to patient death (Opitz et al., 2005).

Computer-assisted technologies coupled with robotic surgical systems can enhance the surgeon capabilities and the control of the surgical tools by providing guidance to the surgical gestures. Specifically, these technologies could be used in abdominal robotic surgery to prevent vessel injury, by intra-operatively identifying and tracking a Region of Interest (ROI) bounding these delicate structures, which would work as active constraints to automatically prevent the robotic arms from touching this area. Intra-operative identification of structures of interest has been
Figure 1: An overview of the proposed framework for long term tracking of safety area identified on endoscopic images
can be applied to any other applications aiming at tracking structures of interest in the surgical field of view. The framework combines the advantages of an optical flow algorithm (Sec. 2.1) with a tracking-by-detection approach (Sec. 2.2), which exploits a novel Model Update Strategy (MUPS) for improving the identification of the Safety Area (SA). Since the optical flow methods are prone to failure, a bayesian network is built to detect possible failures, considering online context information (Sec. 2.3). An extensive quantitative analysis on ex-vivo and in-vivo video sequences is presented to demonstrate long-term achievement (Sec. 3). The results and discussion of this analysis are presented in Sec. 4 and conclusion in Sec. 5.

2. Methods

The workflow of the proposed framework for long-term soft tissue tracking on endoscopic images is shown in Fig. 1. We assumed that the Safety Area Definition, i.e. the identification of the structure to be preserved from injury during surgery on the endoscopic image, is done manually or registering a pre-operative model intra-operatively (Puerto-Souza et al., 2014). The basic steps for the Tracking of the SA along the video sequence consist in (i) detecting salient features inside the SA (ii) finding corresponding features in the successive frames, and exploiting the matched features to (iii) find the perspective transformation between them \((M)\) and used it to (iv) update the new position of the SA. Due to the presence of image noise, errors in the perspective transformation computation, or total occlusion of the SA, a tracking failure can occur. Failure Detection scheme is thus proposed, together with a Tracking Re-initialization strategy to re-detect the SA in the image when visible. The keyframe represent the reference frame, that is re-initialized every time a Tracking Re-initialization is performed. Fig. 2 shows more in detail the workflow of the proposed method, described in the following sections.

2.1. Tracking

The tracking of the SA is performed using a feature-based approach. In the first frame, a set of features \(f_{GFTT}\) are detected inside the SA contour \(SA_i\), using GFTT detector [Shi and Tomasi, 1994]. Kanade-Lucas-Tomasi Tracker (KLT) is then used for feature tracking since, as stated by [Tomasi and Kanade, 1991], it is fast and reliable in case of (i) small movements, (ii) constant brightness and (iii) constant flow in the local neighbourhood. The feature tracking is computed estimating a frame-by-frame feature translation. Since this approximation can lead to errors in tracking due to (i) image noise, (ii) intensity changes caused by illumination or camera exposure changes, (iii) artefacts of the image sensor and (iv) specular reflections, the following strategies were implemented to remove outliers:

1. In order to check the matching correctness, an affine consistency check is also performed between the features belonging to the keyframe and the features in frame \(i\), as stated by [Shi and Tomasi, 1994]; The estimation of the affine motion between local window around the feature is considered as a measure of dissimilarity to reject wrong matches.

2. Endoscopic images are usually affected by specular reflections due to the tissue characteristics and the proximity of the light source to the tissue. The specular reflections appear as bright regions in the images and are identified applying a thresholding operation on S and V channels and dilatation operations [Lehmann and Palm, 2001]. The features located close to specular highlights are discarded.

If \(f_{GFTT}\) is the set of features describing the \(SA_i\) in the keyframe and \(f_{GFTT}\) is the corresponding set of tracked features in the frame \(i\), the tracking of the SA is performed as follows:

\[
SA_i = M \cdot SA_k
\]
puted between $f_{GFTT_0}$ and $f_{GFTT_1}$. $M$ is computed and applied with respect to the keyframe and not with respect to the previous frame $i-1$ in order to avoid drifting and accumulating errors during tracking, as it is shown in Fig. 2. The perspective transform $M$ was computed using the RANSAC strategy [Fischler and Bolles 1981], which is robust in populations with an high number of outliers. Using these strategies in long video sequences, the number of matched features decreases in time, compromising the reliability of $M$ and thus, the tracking. In the proposed workflow, the re-detection of the features is performed each time the features number decreases below the 70% with respect to the features detected in the frame keyframe. The frame in which a re-detection is computed is considered as the new keyframe, i.e., the successive M transformations will be computed with respect to the set of features detected in this frame.

2.1.1. Foreground-Background Segmentation

A global Bayesian probabilistic model based on color histogram is implemented in order to constantly discriminate features describing the SA from the ones describing background or any other object occluding it, inspired by the work of Duffner and Garcia [2013] and Du et al. [2016]. A Probability Segmentation Map (PSM) is computed, representing for each pixel of the SA the probability of belonging to the background $p(c = 0)$ or to the foreground $p(c = 1)$. This map is used to keep only the features laying in a pixel with a foreground probability $p(c_{i} | y_{1:i}) > \tau_{\text{foreground}}$.

The initialization of the probabilistic model is done by computing the HSV histogram of the rectangular area fitting the SA. Assuming that the area of interest is completely visible in the SA, the foreground histogram is initialized considering the area of the image inside the convex hull of the features detected, as proposed by Du et al. [2016], while the background is initialized using the pixel values outside the convex hull, as shown in Fig. 3(a).

In the successive frames, in order to deal with appear–disappear changes of the tissue, the probabilistic model is updated as described by Eq. 2. The update is based on the segmentation of the previous frame $i-1$ and on transition probabilities for foreground and background $p(c_{i} | c_{i-1})$, empirically chosen.

$$p(c_{i} | y_{1:i}) = \frac{p(y_{i} | c_{i} = 1)}{Z}$$

$$= \sum_{c_{i-1}} p(c_{i} = 1 | c_{i-1}) p(c_{i-1} | y_{1:i-1})$$

(2)

where $c_{i}$ is the class of the pixel at frame $i$ (where $c \in \{0, 1\}$), $y_{1:i}$ is the pixel color from frame 1 to $i$, and $Z$ is a normalization constant to keep the probabilities sum to 1.

For not compromising the update of the model in case of partial occlusion of the SA, a clustering of the features inside the SA is computed using the \texttt{kmeans} OpenCV function, and only the pixels belonging to the convex hulls of the clustered features ($n_{\text{cluster}}$) are considered during the update of the foreground histogram (see Fig. 3(c)). An example of how the features are discarded depending on the computed PSM is illustrated in Fig. 3.

2.2. Failure Detection

A Bayesian network is used to estimate the joint failure probability of the tracker, defined as $P(F|A,B,C,D)$, and caused by a combination of the multi clues A, B, C and D.

- $A$ is the number of tracked features ($n_{\text{feat}}$) inside the SA, necessary for the computation of $M$. If $n_{\text{feat}}$ is less than 4, $M$ cannot be computed;
- $B$ is the percentage of features lost in frame $i$ with respect to the number of features in the keyframe ($p_{\text{lost}}$). A high percentage of lost features could indicate the presence of a partial occlusion or sudden changes in the scene;
- $C$ is the validity of the perspective transform $M$ computed between $f_{GFTT_0}$ and $f_{GFTT_1}$ ($v_{M}$), considered valid if: (i) the $z$ coordinate of the transformed points is positive, and (ii) the ninth element of the homography transformation is non-zero, which means a non-valid perspective matrix;
- $D$ is the standard deviation of the optical flow distribution ($std_{OF}$) in terms of image velocity directions. A wide distribution indicates errors in the matching stage due to a sudden change of the scene.
The conditional probability table was defined by assigning to each of the clues a probability distribution, as shown in Fig. 4. For the clues A, B and D, that are continuous variables, a Gaussian distribution was chosen, and for the clue C a probability cross table was used, since $\nu_M$ can assume only two values (0 or 1). If $P(F) > p_h$, the framework switch to the Tracking Re-initialization (Sec. 2.3).

In order to train the network and choose the best parameter set (made of the sigma values for each distribution and the probability values for C), a Monte Carlo sampling method was used. The parameters were sampled within pre-defined ranges, selected from experimental observations, and the cost function, used to determine the best parameter set, was defined as the weighted accuracy of the failure classification, as described by the following Equation:

$$\text{accuracy} = \frac{k_1TP + k_2TN}{k_1TP + FP + k_2TN + FN}$$

(3)

where TP, TN, FP and FN are respectively the number of true positive, true negative, false positive and false negative classification, and $k_1 = 0.8$, $k_2 = 0.2$ are the weights assigned to favour an high rate of TP.

The network was iteratively run, varying the randomly sampled parameters, on a subset of video sequences (training set), where the evidence values ($n_{feat}$, $p_{lost}$, $\nu_M$, std of $f$) were used as input, together with the ground truth information (manually defined when the failure of the tracking really occurred). The parameter set giving an accuracy higher than 0.9 was chosen after 1000 iterations.

2.3. Tracking Re-initialization

If a failure during the tracking is detected, a tracking-by-detection approach based on the generalized Hough transform [Ballard, 1981] is used to find the SA model in the current frame $i$, inspired by [Seib et al., 2012]. The re-detection of the SA is performed in three phases, as described in the following subsections.

2.3.1. Model Initialization

In the first keyframe, in which the SA is defined, a model of the SA is stored. SURF features ($\mathbf{f}_{SURF}$) and descriptors [Bay et al., 2006] are computed inside the SA, since its scale and rotation invariant characteristics are necessary to match features between non-consecutive frames, as in the case of Tracking Re-initialization. KLT initialized with GFTT would not be useful in this case, since it searches feature matches locally, without taking into account possible large displacements of the SA and being invariant to rotation and scale. The model is characterized by the feature position ($x$, $y$), scale ($\sigma$) and orientation ($\theta$), and the centroid ($c_0$) of the area. These feature descriptors, considered with respect to the centroid, uniquely characterise the SA, enabling the SA recognition at any frame.

2.3.2. Model Update

The model defined in the first frame is not always enough to re-detect the SA in long video sequences, since changes in the tissue appearance may occur. For this reason, we used multiple models chosen following a novel Model Update Strategy (MUP8). These models, stored in a buffer, should be different enough from the first model to represent small variations. However, in order to avoid the collection of erroneous models, a similarity with the first model should be ensured. As a measure of similarity we choose the Bhattacharyya distance ($BD$) between the color histogram of the model in the keyframe and in the current frame $i$, inspired by [Giannarou et al., 2013]. It is defined as:

$$BD(H_{first}, H_{curr}) = \sqrt{1 - \rho(H_{first}, H_{curr})}$$

(4)

where $H$ is the normalized histogram density defined as $H = \{h_{bin}\}_{bin=1..m}$, with $\sum_{bin=1}^{m} h_{bin} = 1$.

$\rho$ is the Bhattacharyya coefficient computed from the following Eq.:

$$\rho(H_{first}, H_{curr}) = \sum_{bin=1}^{m} \sqrt{h_{bin}^{first} h_{bin}^{curr}}$$

(5)

In order to make the similarity measure robust against illumination variations, we opted for using the combination of H and S channel from the HSV image instead of the RGB channels used by [Giannarou et al., 2013]. Thus, the strategy to update the model, also described in Alg. 1, can be explained in two steps:

- The model stored in the first frame is always kept fixed in order to always have a valid reference. The new models ($n_{model}$) are collected only if the $BD$ is inside the range $\delta_{PA} = [\delta_{min}, \delta_{max}]$, where $\delta_{min}$, and $\delta_{max}$ were chosen as the 10th and 90th percentile of the $BD$ distribution extracted from a training dataset;
Algorithm 1 Model Update Strategy

1: procedure UPDATE_WEIGHT(model, \(\beta_1, \beta_2\))
2: \(\text{model.weight} = \beta_1 \cdot \text{model.BD} + \beta_2 \cdot \frac{n\text{Times}_{\text{used}}}{n\text{Times}_{\text{notUsed}}}\)
3: end procedure

4: procedure UPDATE_MODEL(modelBuffer, modelNew, BDmax, BDmin, \(\beta_1, \beta_2\))
5: if modelNew.BD > BDmin and modelNew.BD < BDmax then
6: if size(modelBuffer) < \(n_{\text{models}}\) then
7: add modelNew to modelBuffer
8: UPDATE_WEIGHT(modelNew, \(\beta_1, \beta_2\))
9: else
10: modelWeakest = model with lowest weight in modelBuffer
11: if modelWeakest.nTimes_{\text{notUsed}} > \(m\) then
12: replace modelWeakest with modelNew in modelBuffer
13: UPDATE_WEIGHT(modelNew, \(\beta_1, \beta_2\))
14: end if
15: end if
16: end if
17: return modelBuffer
18: end procedure

- A weight is assigned to each model belonging to the buffer, as:

\[
w_j = \beta_1 \cdot BD_j + \beta_2 \cdot \frac{n\text{Times}_{\text{used}}}{n\text{Times}_{\text{notUsed}}}\]

where \(n\text{Times}_{\text{used}}\) is the number of times the model was previously used for the SA recognition, the number of times the model was not used is \(n\text{Times}_{\text{notUsed}}\), and \(j\) is the index of the model in the buffer. \(\beta_1\) and \(\beta_2\) are the weights assigned to the two parameters determining the goodness of the model \(w_j\).

- The model \(j\) with the minimum weight \(w_j\) will be replaced by a new model \(i\) only if:

\[
n\text{Times}_{\text{notUsed}} > m\]

where \(m\) was empirically chosen.

2.3.3. Model Recognition

SURF features are detected on the entire frame \(i\) and then are matched with the features belonging to the set of \(n_{\text{models}}\) using a nearest-neighbor matching. As a first outlier rejection stage, the wrong matches are rejected if the ratio between the closest and the second-closest descriptor distances is lower than a threshold \(\tau\) (Lowe 2004). The possible SA poses, represented by the position \((x, y)\), scale \((\sigma)\) and orientation \((\theta)\), are clustered in a multidimensional Hough-space accumulator, as shown in Fig. 5.

The coarse grid represents translation in \(x\) and \(y\) direction, while in each of these cells, the bins along \(x\) axis represent \(\sigma\) and the ones along the \(y\) axis represent \(\theta\). Each feature match independently votes for a possible SA position, orientation and scale, increasing the corresponding bin in the accumulator. The new centroid position \(c_i\) is estimated as:

\[
c_i = (c_{x_i}, c_{y_i}) = v + p_i\]

where \(p_i\) is the feature position in frame \(i\) and \(v\):

\[
v = \begin{pmatrix} \cos(\alpha) & -\sin(\alpha) \\ \sin(\alpha) & \cos(\alpha) \end{pmatrix} \left( c_0 - p_0 \frac{\sigma_s}{\sigma_0} \right)\]

\(v\) is the translation vector from the centroid of the model \(c_0\) to the position of a feature \(p_0\) in the model, normalized with the scale ratio of the feature in frame \(i\) \((\sigma_i)\) and of the feature of the model \((\sigma_0)\), and rotated depending on \(\alpha = |\theta_0 - \theta_i|\), i.e. the rotation angle between respectively the feature rotation of the model and of the frame \(i\).

The maximum in the Hough-space returns the set of features \(f_{\text{SURF}_j}\) of the model that best match the SA, a shown in Fig. 5.

![Figure 5: On the left, the process for the recognition of the SA in the new frame. The features detected in the current frame are matched with the ones belonging to the models (colored and black lines indicate respectively right and wrong matches). On the right, the Hough accumulator is shown: the two axes indicate the feature position, and, inside each cell, the horizontal and vertical translation encode the scale and rotation, respectively. Each feature match votes for a possible SA position, increasing the Hough space accumulator, represented on the right. Right matches increment the same Hough accumulator cells, leading to a maximum (white squares), while the wrong matches votes are scattered (gray squares).](image)

Every time a SA is recognised, the Tracking algorithm is re-initialized with the same workflow described in Sec. 2.1, establishing a new keyframe. In this phase, the probability segmentation map has a fundamental role, since the SA can still be partially occluded. Keeping only the features belonging to the object prevents from failure. If the SA is not recognised, the algorithm waits until it is visible again.

3. Experimental Evaluation

The evaluation is focused on demonstrating the robustness of the algorithm against: (i) partial occlusion, (ii) total occlusion and (iii) SA out of field of view, which are the main events, often happening during surgeries, that can affect the reliability of a tracker. In order to assess the performance of the algorithm against these events, we used ex-vivo and in-vivo datasets.

The ex-vivo dataset is made of endoscopic images of ex-vivo organs (goat kidney, pig liver). It was developed simulating surgical scenarios in a controlled way, recreating typical events happening during surgery. The videos were
recording using a da Vinci® stereo camera and the robotic system (Intuitive Surgical Inc., CA) at the Surgical Robot Vision group (University College London, London, UK). All the videos were recorded at 25fps with an image resolution of 720 × 576.

The in-vivo dataset consists of videos of real surgical operations performed at Ospedali Maggiore Ca’ Granda (Milan, Italy). The videos were captured with a monocular STORZ endoscope. All the data were anonymized.

Details of each video sequence in terms of duration, number of frames, and a brief description are presented in Fig. 6.

For each sequence, we created a Ground Truth (GT), in the form of a 2D polygon around the area of interest, with a frame of step size of 10. This was performed manually by an operator with the supervision of an expert surgeon.

For a more accurate evaluation, the same frames were also labeled with one of the following attributes: (i) Safe Area Visible (SAV), (ii) partial occlusion (PO), (iii) total occlusion (TO), (iv) out of field of view (OFV). These datasets are available online for the benefit of the community [1]. Tab. 1 shows the percentage of frames with SAV, PO, TO, and OFV for each video of the two datasets.

<table>
<thead>
<tr>
<th>in-vivo dataset</th>
<th>ex-vivo dataset</th>
</tr>
</thead>
<tbody>
<tr>
<td>EV1</td>
<td>EV2</td>
</tr>
<tr>
<td>SAV</td>
<td>67.71%</td>
</tr>
<tr>
<td>PO</td>
<td>7.43%</td>
</tr>
<tr>
<td>TO</td>
<td>7.86%</td>
</tr>
<tr>
<td>OFV</td>
<td>13.29%</td>
</tr>
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<td></td>
<td></td>
</tr>
</tbody>
</table>

Table 1: Percentage of frames with Safety Area Visible (SAV), Partial Occlusion (PO), Total Occlusion (TO), Out of Field of View (OFV) for each video of the two datasets.

The performance was assessed using precision and recall curves, and the F-measure [Wu et al., 2013]. For each video, the precision value $\alpha$ was computed as:

$$\alpha = \frac{TP}{TP + FP}$$

where $TP$ is the number of true positives of the SA tracked and $FP$ is the number of false positives of the SA tracked. The recall value $\beta$ is defined as:

$$\beta = \frac{TP}{TP + FN}$$

where $FN$ is the number of false negatives of the SA tracked. The F-measure $\gamma$ is the harmonic mean of precision and recall:

$$\gamma = 2 \cdot \frac{\alpha \cdot \beta}{\alpha + \beta}$$

The metrics used to define the true positives of the SA is the overlap ratio, measured in pixels, and defined as:

$$\varphi = \frac{|T \cap G|}{|T \cup G|}$$

where $T$ is the set of SA tracking results, and $G$ is the set of GT.

<table>
<thead>
<tr>
<th>Parameters</th>
<th>Definition</th>
<th>Values</th>
</tr>
</thead>
<tbody>
<tr>
<td>$t_{\text{segmentation}}$</td>
<td>threshold over which the features are considered belonging to the SA</td>
<td>0.7</td>
</tr>
<tr>
<td>$n_{\text{cluster}}$</td>
<td>number of feature clusters</td>
<td>8</td>
</tr>
<tr>
<td>$\tau_{\text{foreground}}$</td>
<td>foreground threshold</td>
<td>0.7</td>
</tr>
<tr>
<td>$\text{max} , \text{feat}$</td>
<td>maximum feature number for distribution A</td>
<td>$\frac{n_{\text{feat}A}}{n_{\text{feat}A}} \cdot 2$</td>
</tr>
<tr>
<td>$\sigma_{\text{feat}}(F = 0)$</td>
<td>non-failure probability</td>
<td>$\frac{\text{max} , \text{feat}}{3}$</td>
</tr>
<tr>
<td>$\sigma_{\text{feat}}(F = 1)$</td>
<td>probability distribution of A</td>
<td>6</td>
</tr>
<tr>
<td>$\sigma_{\text{fail}} F = 0$</td>
<td>non-failure probability of the failure event</td>
<td>19</td>
</tr>
<tr>
<td>$\sigma_{\text{fail}} F = 1$</td>
<td>distribution of B</td>
<td>53</td>
</tr>
<tr>
<td>$p_{\text{fail}} C = 0$</td>
<td>non-failure probability for C</td>
<td>0.1</td>
</tr>
<tr>
<td>$p_{\text{fail}} C = 1$</td>
<td>failure probability for C</td>
<td>0.47</td>
</tr>
<tr>
<td>$\sigma_{\text{std} A}$</td>
<td>sigma value of the failure event</td>
<td>43</td>
</tr>
<tr>
<td>$\sigma_{\text{std} D}$</td>
<td>probability distribution of D</td>
<td>165</td>
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<tr>
<td>$\rho_{\text{th}}$</td>
<td>failure threshold</td>
<td>0.4</td>
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<tr>
<td>$n_{\text{models}}$</td>
<td>number of models used by the MUpS</td>
<td>10</td>
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<tr>
<td>$m$</td>
<td>minimum number of times a model can be used before being replaced</td>
<td>5</td>
</tr>
<tr>
<td>$\delta_{\text{min}}$</td>
<td>10th percentile of BD distribution</td>
<td>0.1</td>
</tr>
<tr>
<td>$\delta_{\text{max}}$</td>
<td>90th percentile of BD distribution</td>
<td>0.5</td>
</tr>
<tr>
<td>$\beta_1$</td>
<td>model weight parameter</td>
<td>0.5</td>
</tr>
<tr>
<td>$\beta_2$</td>
<td>model weight parameter</td>
<td>0.5</td>
</tr>
</tbody>
</table>

Table 2: Summary of the algorithm parameters used or the evaluation of the framework.

The precision and recall curves were computed varying the overlap ratio threshold used to identify the TP values. The F-measure was computed considering $\varphi > (0.2, 0.5, 0.8)$.

In case of partial occlusion, the SA also included the occluding object. So, to take into consideration only the area
of interest, the probability segmentation map was used to discard the pixels belonging to the background (and thus the occluding object) and the overlap ratio was computed considering the foreground area.

The evaluation was performed with and without the MUps, in order to assess its contribution. Precision and recall curves where computed for both cases. In order to verify the behaviour of the framework against PO, these curves were also computed considering only the frames labelled with PO.

The recovery time after the failure was computed as the mean number of frames between the lost of the SA tracking and the correct re-detection ($\delta > 0.5$) for each video sequence, with and without MUps.

The code was implement in C++, using the OpenCV library for the management of the images and KLT library for KLT algorithm implementation, since the OpenCV version of the KLT tracker does not include the affine consistency check. The code released by Seib et al. (2012) was used for the tracking by detection approach. The program was running on a system with GNU/Linux operating system, and a CPU Intel Core i5-3230M with four cores.

The parameters used for this evaluation are summarized in Tab. 2.

Table 3: F-measure values (without/with MUps) for three different overall threshold (low = 0.2, medium = 0.5, high = 0.8) and Recovery Time [# frames] (without/with MUps)

<table>
<thead>
<tr>
<th>$\gamma_{\text{low}}$</th>
<th>EV1</th>
<th>EV2</th>
<th>EV3</th>
<th>IV1</th>
</tr>
</thead>
<tbody>
<tr>
<td>0.93/0.95</td>
<td></td>
<td></td>
<td></td>
<td>0.34/0.60</td>
</tr>
<tr>
<td>$\gamma_{\text{medium}}$</td>
<td>0.93/0.95</td>
<td>0.44/0.96</td>
<td>0.90/0.93</td>
<td>0.34/0.60</td>
</tr>
<tr>
<td>$\gamma_{\text{high}}$</td>
<td>0.80/0.71</td>
<td>0.30/0.45</td>
<td>0.20/0.38</td>
<td>0.22/0.32</td>
</tr>
<tr>
<td>$t_{\text{time}}$</td>
<td>0.84/0.50</td>
<td>37.50/0.88</td>
<td>7.00/2.00</td>
<td>16.00/8.04</td>
</tr>
</tbody>
</table>

4. Results and Discussion

In Fig. 6 example images from each video sequences of the in-vivo and ex-vivo dataset are shown. In the first column, the SA, as defined in the first frame, is shown. The second and third column show, respectively, an example of partial and total occlusion. There is a high percentage of frames with partial or total occlusion, where the target is not visible, as shown in Tab. 1 allowing the assessment of the algorithm under different conditions.

Fig. 7 shows an example of the trend of the variables representing the clues (A, B, C, D) used by the Bayesian network to estimate the joint failure probability P. The last row of the figure represents when the Tracking Re-initialization is active. From these data, we can observe different events (highlighted in Fig. 7 with numbered orange boxes) that trigger the tracking re-initialization:

1. The percentage of lost features drastically increases since the area is moving out of the camera field of view;
2. The number of features inside the SA decreases drastically due to an instrument occlusion. This event combined with an increase of the optical flow standard distribution (caused by wrong matches) leads to a failure of the KLT tracker;
3. The standard distribution of the optical flow increases due to a sudden movement of the camera; KLT tracker alone would fail since it is not robust to sudden movement of the scene;
4. The homography is invalid due to wrong feature matches caused by partial instrument occlusion. In this case, KLT would continue to work, however tracking wrong features (i.e. not belonging to the area of interest) and compromising the SA tracking.

The Bayesian network was adjusted to be very sensible to possible failures, because a false negative is not critical for our application. All the events aforementioned are examples of cases in which KLT fails, demonstrating the need of a failure detection and SA re-detection strategies. Moreover, if a failure is not detected at the first frame, the tracking shows a degradation and the Bayesian network will estimate the failure most probably after a few frames, re-initializing correctly the SA.

The precision and recall curves are shown in Fig. 8. These curves demonstrate the performance of the tracking and the effect of the MUps. Considering all frames, the precision and recall values are strong, as it is confirmed by the F-measure (Tab. 3). In case of partial occlusion, even if the values are lower, the algorithm still performs well. Regarding the effect of MUps, in most of the videos this strategy improves both the precision and the recall considering all frames. The same behaviour can be observed considering only the partially occluded frames.

Fig. 7 shows an example of the trend of the variables representing the clues (A, B, C, D) used by the Bayesian network to estimate the joint failure probability P. The

5. Conclusion

In this paper, we proposed a framework for Long-term Safety Area Tracking (LT-SAT), which aims to be used to
The video shows an exposed goat kidney and the SA is defined on the main vessel entering in the kidney.

The video shows a similar surgical field of view as in EV1 with a different kidney.

The video shows a pig liver and the SA is defined on a vessel.

This sequence was extracted from a video of a pancreatectomy procedure.

The overall results show that the framework fulfills the main requirements stated in Sec. 1, such as (i) the long-term robustness under complicated conditions (partial or total occlusion), thanks to the combination and improvement of state-of-the-art tracking strategies with a Bayesian-based failure detection scheme; and (ii) the adaptation to environment/object changes, thanks to the
novel strategy of updating the models used for the SA re-initialization. The hybrid combination of KLT tracker, based on GFTT, with tracking-by-detection approach, based on SURF features, aims at exploiting the strengths of the two approaches. The KLT tracker is robust, accurate and computationally cheap in case of small movements, while tracking-by-detection approach would be more computationally expensive and less accurate than KLT if used to track the SA frame-by-frame. Its strength, however, is the invariance to rotation and scale changes that allows to recover the SA, even if it disappear for many frames and reappears in a totally different pose, strengthened by a generalized Hough Transform approach to discard outliers.

The analysis of the ex-vivo and in-vivo videos show that the framework is capable of maintaining good tracking performance for extended periods of time (≈5 min), covering in some cases the entire video sequences in which the vessel had to be tracked. The high precision values confirm that the performance of the framework is within the specifications required by the surgeons.

In terms of precision, the performance of the proposed framework is comparable with state-of-the-art algorithms [Ye et al., 2016]. However, the real improvement given by the proposed framework with respect to state-of-the-art algorithms consists in the robustness, represented by recall values. Particularly, in contrast with the literature, we tested the algorithm on long video sequences (between 5000 and 7000 frame), simulating and considering many of the events happening in surgery and that can affect the performance of the tracking. This extensive evaluation on long video sequences allows to state that the algorithm is able to work properly and robustly in a real surgical scenario.

The weaknesses are (i) the difficulties in recovering the SA after a large deformation, due to the fact that the Model Re-initialization take into account only small deformation, and (ii) the computational time.

Future work will aim at addressing the issue of robust tracking under big deformations, exploiting deformation modelling techniques. Also since the current implementation of the algorithm is not able to run in real time, next steps will include software architecture improvements and code optimization, which should reduce considerably the computational time. At this point, authors aim at integrating this framework with a dense 3D reconstruction algorithm, already developed by [Penza et al., 2016], in order to obtain a 3D area tracking, and integrate the overall system in a robotic platform.

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