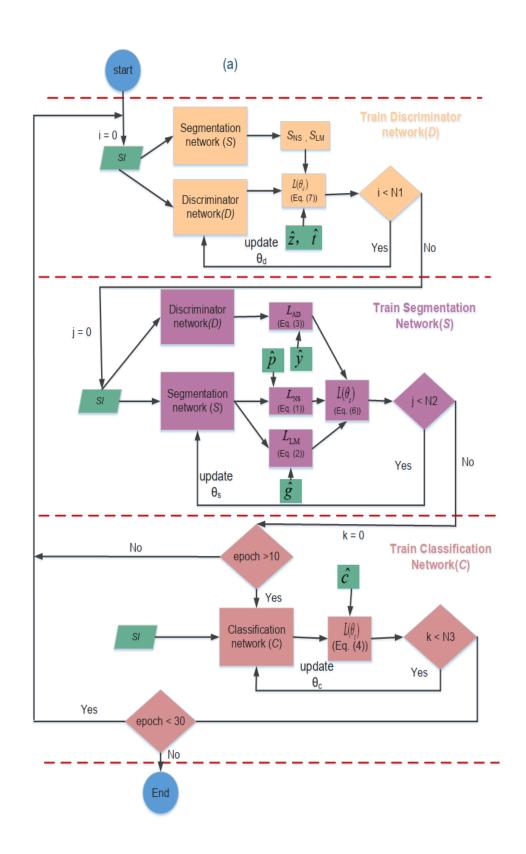
## Supplementary material



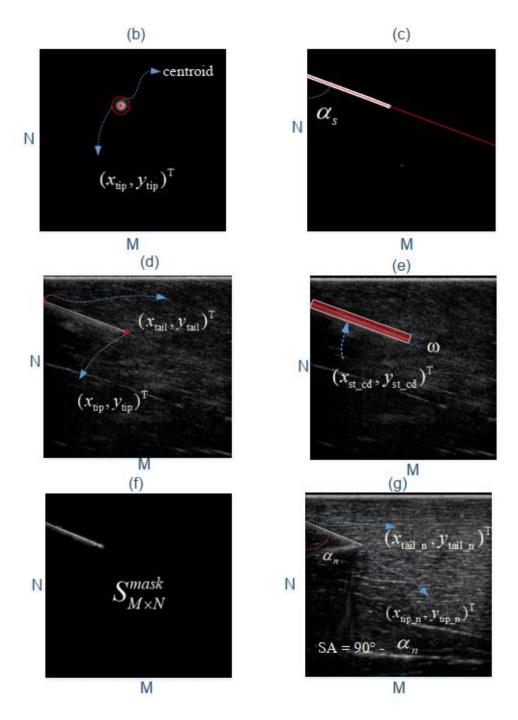
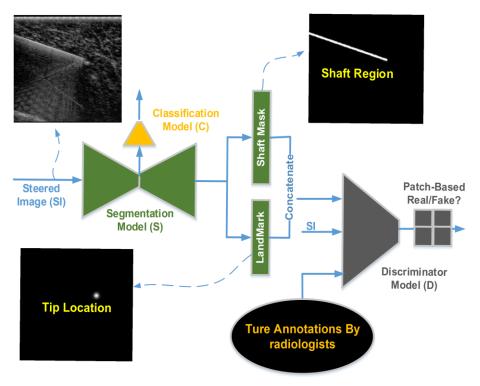


Fig. A1 (a) Training process of the MTL framework. N1, N2, N3 denote the number of iterations of discriminator network, segmentation network and classification network in each epoch respectively. (b) The image of needle tip landmark ( $S_{\rm LM}$ ). (c) The image of needle shaft segmentation ( $S_{\rm NS}$ ). (d) and (e) The steered image (SI). (f) The needle mask ( $S_{M\times N}^{mask}$ ) of normal-angled image, obtained by Eq. (10). (g) The final fusion image ( $S_{M\times N}^{Fusion}$ ). M is the width of the images. N is the height of images.



**Fig. A2.** Architecture of proposed MTL network. The MTL framework consists of a segmentation network (S), a discriminator network (D) and a classification network (C). Discriminator network (D) only be used in the training phase.

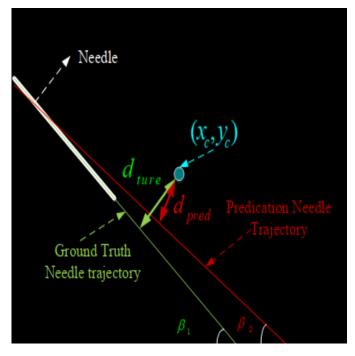
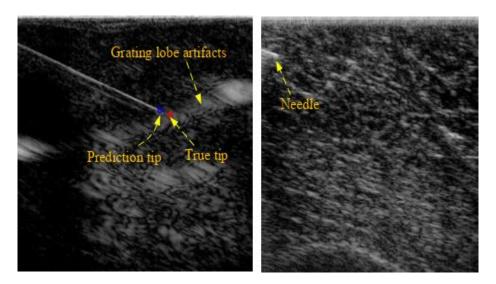
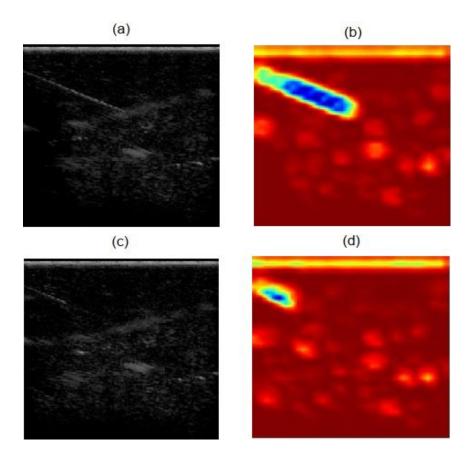


Fig. A3 Illustration of the targeting error (TE) and shaft localization error (SLE).



**Fig. A4** Illustration of failure cases. The needle tip is contaminated by grating lobe artifacts. The blue circle is the prediction tip location and the red circle is the true tip location.



**Fig. A5** Illustration of the feature map difference between short and long length samples for needle to insert. (a) The long length needle sample (b) The feature maps of (a), generating from the visualization of the last convolution layer of the classification network (C). (c) The short length needle sample. (d) The feature maps of (c), generating from the visualization of the last convolution layer of the classification network (C).

**Table A1.** Hyperparameters for training our MTL network

| Hyperparameters | Segmentation | Classification | Discriminator |  |
|-----------------|--------------|----------------|---------------|--|
|                 | network(S)   | network(C)     | network(D)    |  |
| Learning rate   | 2e-4         | 5e-4           | 1e-4          |  |
| Optimizer       | Adam         | Adam           | Adam          |  |
| Epoch           | 30           | 20             | 30            |  |
| Batch size      | 8            | 16             | 8             |  |
| $\mu_1$         | 0.5          |                |               |  |
| $\mu_2$         | 0.5          |                |               |  |
| $\mu_3$         | 0.2          |                |               |  |

**Table A2.** Comparing localization accuracy and overall processing time from the proposed NLEM method versus STL\_NLEM, WAT\_NLEM and NA\_NLEM methods.

| Method   | 1EDLS(%)            | ED(mm)             | SLE(°)             | TE (mm)            | OPT(s)                 |
|----------|---------------------|--------------------|--------------------|--------------------|------------------------|
| NLEM     | $95.63 \pm 1.00$    | $0.29 \pm 0.02$    | 0.27±0.02          | 0.46±0.01          | 0.0149±0.0001          |
| STL_NLEM | 92.64±2.63<br>(**)  | 0.31±0.22<br>(**)  | 0.33±0.02<br>(**)  | 0.48±0.01<br>(**)  | 0.0250±0.0003<br>(***) |
| WAT_NLEM | 86.44±7.36<br>(***) | 0.38±0.09<br>(***) | 0.35±0.05<br>(***) | 0.51±0.04<br>(***) | 0.0149±0.0001          |
| NA_NLEM  | 41.47±1.51<br>(***) | 0.67±0.03<br>(***) | 0.68±0.09<br>(***) | 0.78±0.15<br>(***) | 0.0149±0.0001          |

Note: The one tailed paired t-test is used to compare related methods with NLEM. \*\*And \* \*\* in the table represent p < 0.05 and p < 0.00001, respectively. A value of p < 0.05 is considered statistically significant.

**Table A3.** Comparing classification accuracy from the proposed NLEM method versus STL\_NLEM method.

| Method  | Accuracy (%)     | Precision (%) | Sensitivity (%)  | Specificity (%)  |
|---------|------------------|---------------|------------------|------------------|
| NLEM    | 97.37 ±0.71      | 97.70 ±1.01   | 99.71 ±0.31      | 96.82 ±1.25      |
| STL_NLM | 94.63 ±3.11 (**) | 98.49 ±2.39   | 93.60 ±4.38 (**) | 97.36 ±3.57 (**) |

Note: The one tailed paired t-test is used to compare STL\_NLEM method with NLEM. \*\* in the table represent p < 0.05. A value of p < 0.05 is considered statistically significant.