

Automated neuron tracing using probability hypothesis density filtering

Miroslav Radojević and Erik Meijering

Supplementary Information

Algorithm 1 Neuron tracing

- 1: $k = 0$ ▷ Initialize
 - 2: $\{\omega_{0|0}^n, \mathbf{x}_{0|0}^n\}_{n=1}^{\rho N_0}$ ▷ Initial particle and observation set
 - 3: $\{\hat{\mathbf{x}}_{0,i}\}_{i=1}^{N_0}$ ▷ Initial estimate
 - 4: **repeat**
 - 5: $k = k + 1$
 - 6: $\mathbf{p}_i^n \sim h(\mathbf{p}|\hat{\mathbf{x}}_{k-1,i}) \quad n \in [1, \rho N_{k-1}]$ ▷ Draw observation particles
 - 7: $\mathbf{p}_{i,j}^n \in \mathcal{C}_j, \quad j \in [1, M_k], \quad n \in [1, |\mathcal{C}_j|]$ ▷ Cluster observation particles
 - 8: $\mathbf{z}_{k,j} = [\mathbf{p}_{i,j}^{\hat{n}}, \tau(\mathbf{p}_{i,j}^{\hat{n}})]$ ▷ Select representative sample
 - 9: $\mathbf{Z}_k = \{\mathbf{z}_{k,j}, \dots, \mathbf{z}_{k,M_k}\}$ ▷ Construct observations
 - 10: $\{\omega_{k|k}^n, \mathbf{x}_{k|k}^n\}_{n=1}^{\rho N_k}, \nu_k, \{\hat{\mathbf{x}}_{k,i}\}_{i=1}^{N_k} \leftarrow \text{SMC-PHD}(\{\omega_{k-1|k-1}^n, \mathbf{x}_{k-1|k-1}^n\}_{n=1}^{\rho N_{k-1}}, \mathbf{Z}_k)$ ▷ Algorithm 2
 - 11: **until** $[\nu_k] = 0$ ▷ $[\cdot] \equiv$ nearest integer
-

Algorithm 2 SMC-PHD filtering

- 1: **Input:** $\{(\omega_{k-1|k-1}^n, \mathbf{x}_{k-1|k-1}^n)\}_{n=1}^{\rho N_{k-1}}, \{Z_{k,j}\}_{j=1}^{M_k}$ $\triangleright D_{k-1}(\mathbf{x})$ approx. observation Z_k
 - 2: **for** $n = 1, \dots, \rho N_{k-1}$ **do**
 - 3: **for** $m = 1, \dots, \eta$ **do**
 - 4: $i = (n - 1)\eta + m$
 - 5: **Draw:** $\mathbf{x}_{k|k-1,p} \sim \pi_{k|k-1}(\mathbf{x}|\mathbf{x}_{k-1|k-1}^n) \rightarrow \mathbf{x}_{k|k-1,p}^i$ \triangleright Persistent object particles
 - 6: **Compute:** $\omega_{k|k-1,p}^i = p_S \frac{1}{\eta} \omega_{k-1|k-1}^n$
 - 7: **Draw:** $\mathbf{x}_{k|k-1,s} \sim \beta_{k|k-1}(\mathbf{x}|\mathbf{x}_{k-1|k-1}^n) \rightarrow \mathbf{x}_{k|k-1,s}^i$ \triangleright Spawning object particles
 - 8: **Compute:** $\omega_{k|k-1,s}^i = p_S \frac{1}{\eta} \omega_{k-1|k-1}^n$
 - 9: **end for**
 - 10: **end for**
 - 11: $\{(\omega_{k|k-1}^n, \mathbf{x}_{k|k-1}^n)\}_{n=1}^{S_k} = \{(\omega_{k|k-1,p}^n, \mathbf{x}_{k|k-1,p}^n)\}_{n=1}^{\rho\eta N_{k-1}} \cup \{(\omega_{k|k-1,s}^n, \mathbf{x}_{k|k-1,s}^n)\}_{n=1}^{\rho\eta N_{k-1}}$ \triangleright Union of particle sets
 - 12: **for** $n = 1, \dots, S_k$ **do**
 - 13: **Update:** $\omega_{k|k}^n = (1 - p_D)\omega_{k|k-1}^n + \sum_{z \in Z_k} \frac{p_{Dgk}(z|\mathbf{x}_{k|k-1}^n)\omega_{k|k-1}^n}{C_k(z) + \sum_{n=1}^{S_k} p_{Dgk}(z|\mathbf{x}_{k|k-1}^n)\omega_{k|k-1}^n}$
 - 14: **end for**
 - 15: $\nu_k = \sum_{n=1}^{S_k} \omega_{k|k}^n$ \triangleright Cardinality calculation
 - 16: **Estimate:** $\hat{\mathbf{x}}_{k,i} \leftarrow \{\omega_{k|k}^n, \mathbf{x}_{k|k-1}^n\}_{n=1}^{S_k}$ \triangleright Mean-shift clustering
 - 17: **Resample:** $N_k = \lceil \nu_k \rceil, \{\omega_{k|k}^n, \mathbf{x}_{k|k-1}^n\}_{n=1}^{S_k} \rightarrow \{\omega_{k|k}^n, \mathbf{x}_{k|k}^n\}_{n=1}^{\rho N_k}, \omega_{k|k}^n = \nu_k / (\rho N_k)$
 \triangleright Systematic resampling with ρ particles per object
-

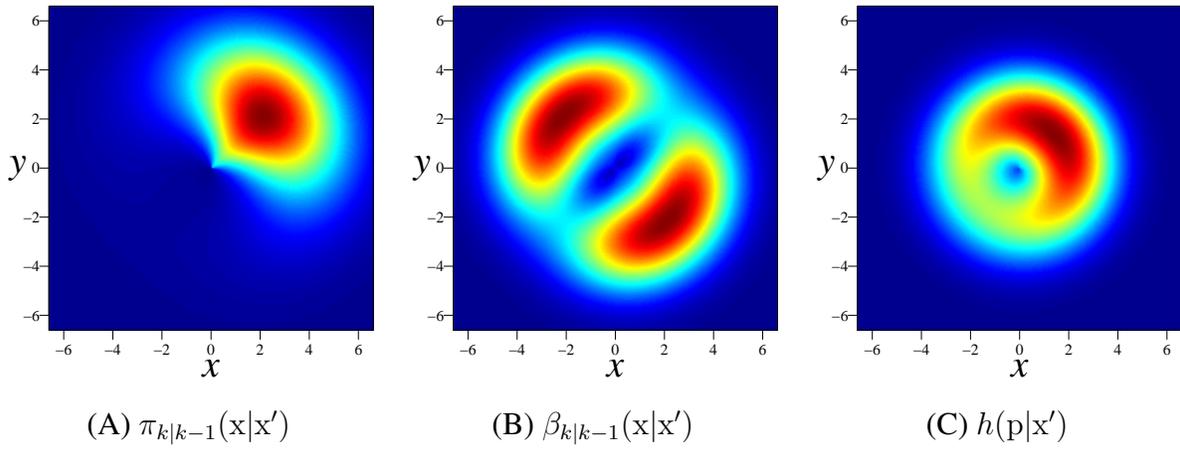


Figure S1: Transition densities (2D examples) for persistent (A) and spawned (B) objects with $z = 0$, $\mathbf{x}' = \left[0, 0, 0, \frac{1}{\sqrt{2}}, \frac{1}{\sqrt{2}}, 0\right]$, $\kappa = 2$, and $r_k = 3$. (C) Importance sampling used in the observation model without the tubularity component, $\tau(\mathbf{p}) = 1$, and $\kappa = 0.5$. Rainbow color coding is used running from blue (indicating low values) to red (indicating high values).

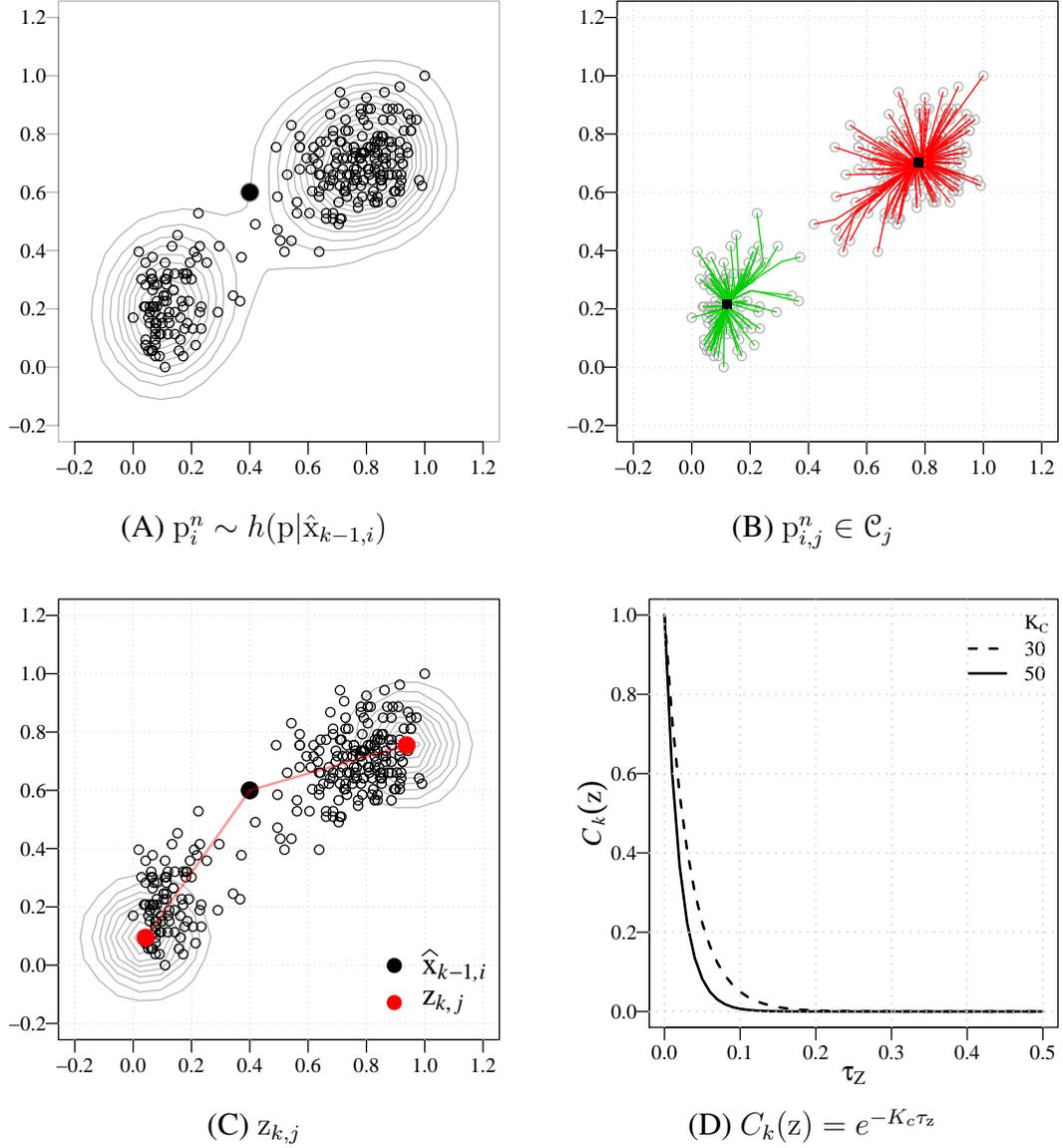


Figure S2: Formation of the observations (2D example). (A) For each object i from iteration $k - 1$, particles p_i^n are sampled from the importance sampling function h , using the state estimate $\hat{x}_{k-1,i}$. The solid dot indicates the location of $\hat{x}_{k-1,i}$ and the contours represent lines of equal particle weight. (B) The particles are processed by mean-shifting resulting in clusters \mathcal{C}_j whose labeled particles are denoted as $p_{i,j}^n$. (C) Each observation $z_{k,j}$ is obtained from the representative cluster particle $p_{i,j}^n$ as described in the main text. Contours represent lines of equal observation likelihood. (D) The clutter intensity function.

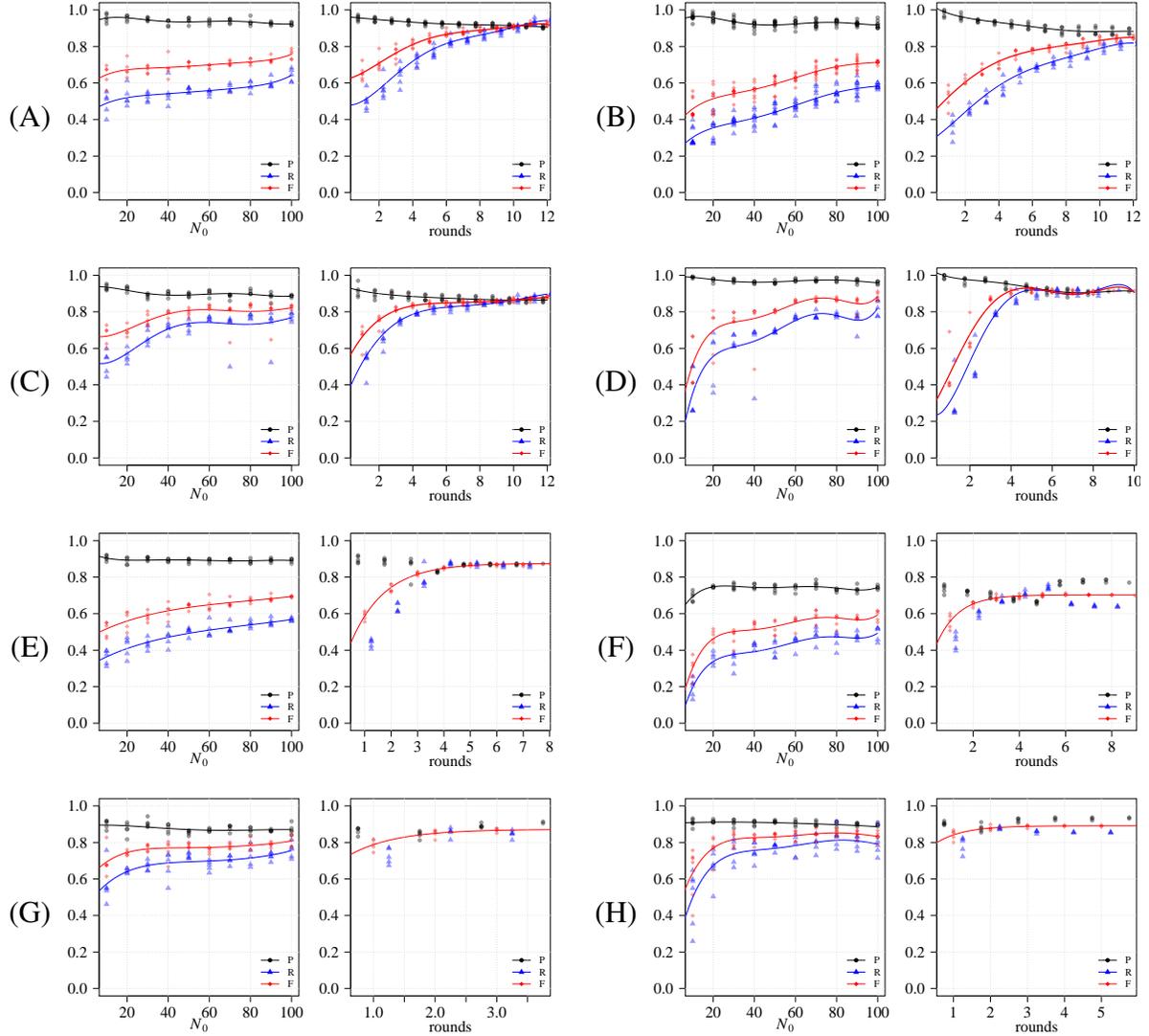


Figure S3: Performance as a function of numbers of seeds and rounds for four example cases from the OPF (A-D) and the HCN (E-H) data set. Similar trends were observed for all cases in the respective data sets. Left panel per case: Precision (P), recall (R), and F-score (F) after one round initialized with different numbers of seeds (N_0). Right panel per case: The scores after multiple rounds with a fixed number of seeds ($N_0 = 40$). Fifth-order polynomial curves were fit to the data to show approximate trends.

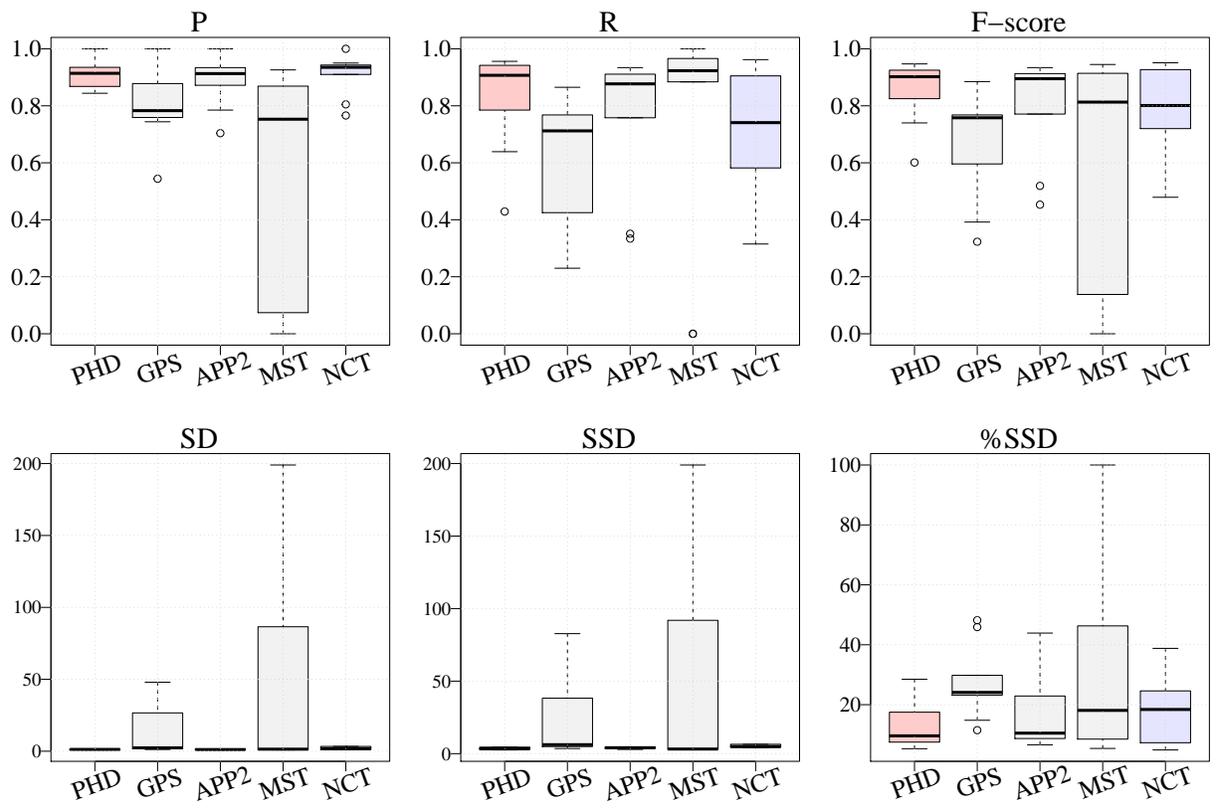


Figure S4: Performance comparison of our method with several other methods on the OPF data set. For each method and each measure, the plotted box indicates the 25-75 percentile, the horizontal bar indicates the median score, and the whiskers and outliers are drawn using the default settings of R.

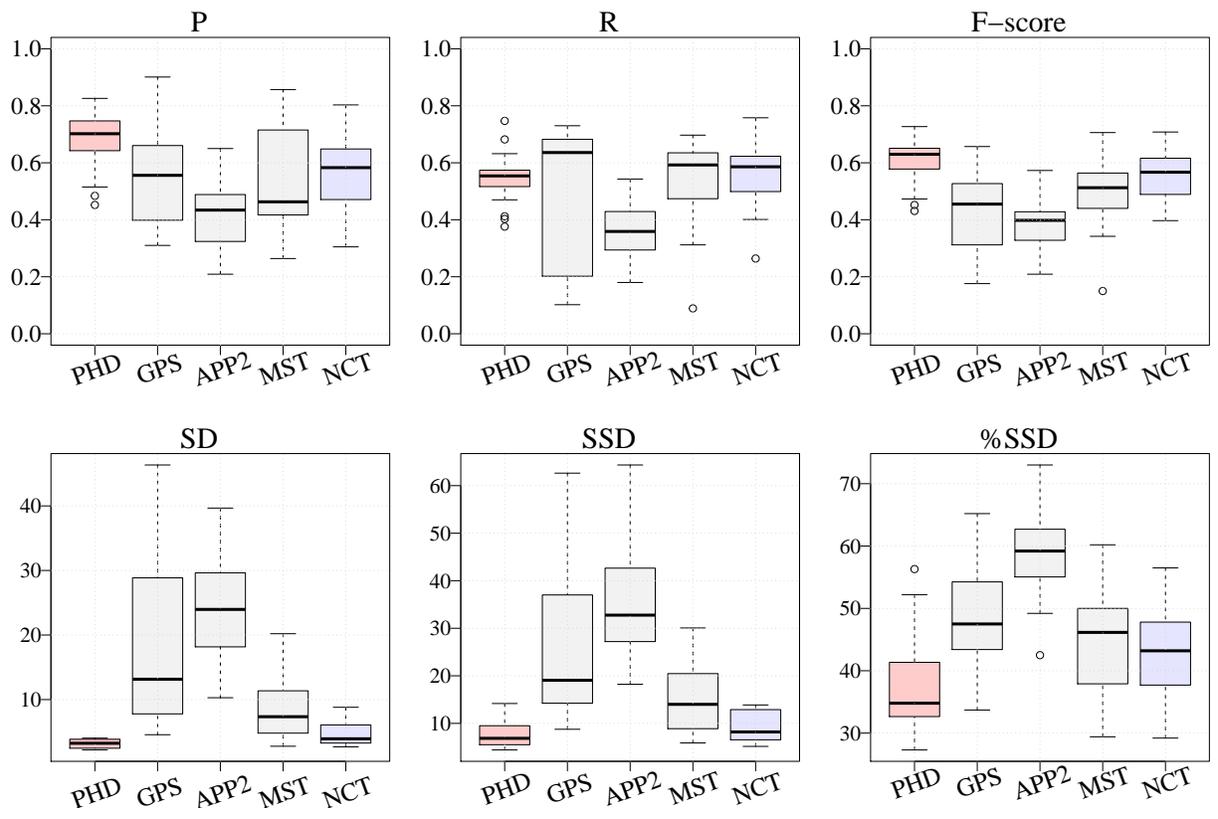


Figure S5: Performance comparison of our method with several other methods on the HCN data set. For each method and each measure, the plotted box indicates the 25-75 percentile, the horizontal bar indicates the median score, and the whiskers and outliers are drawn using the default settings of R.

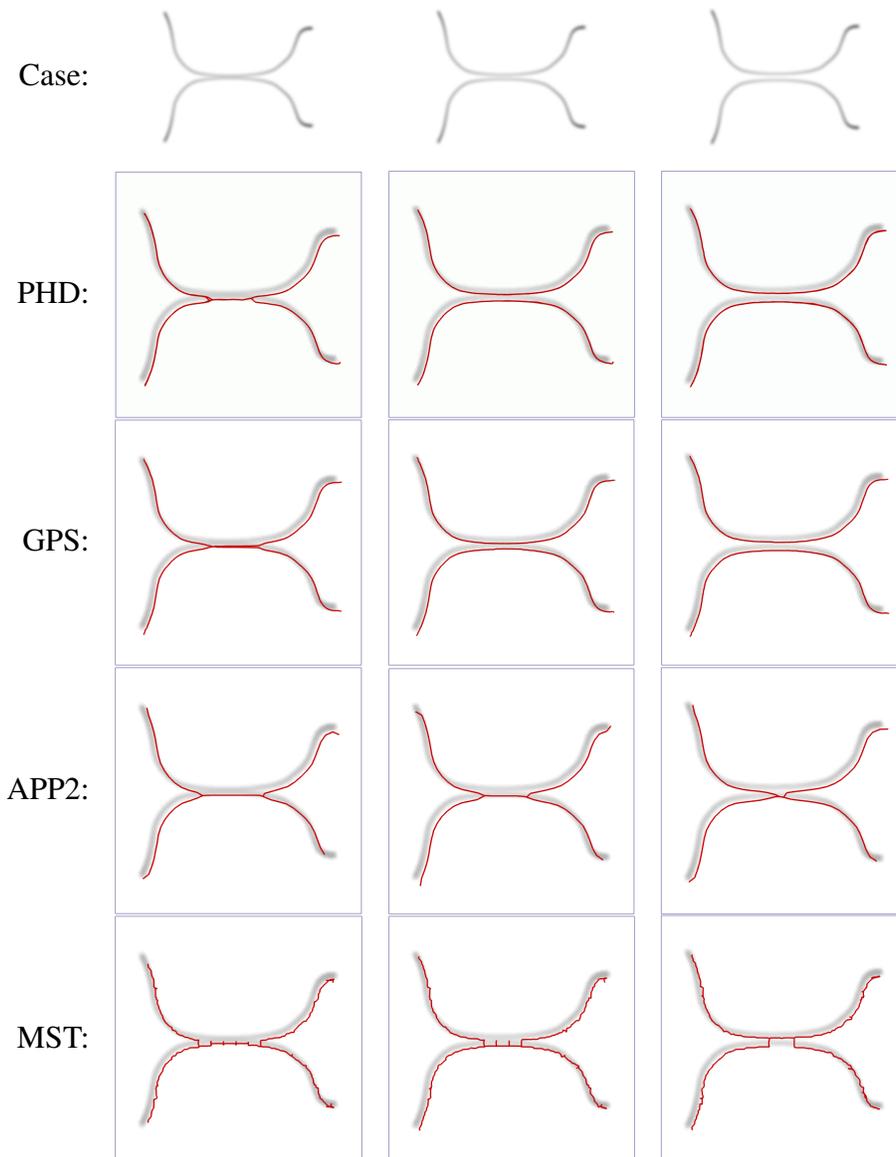


Figure S6: Ability of the tested methods to separate two fibers of similar intensity and scale running closely in parallel. The examples show cases with gradually increasing distance between the fibers: overlap (left column), just separated (middle column), and clearly separated (right column). The tracing results of PHD, GPS, APP2, MST are overlaid (with slight offset) in red color.

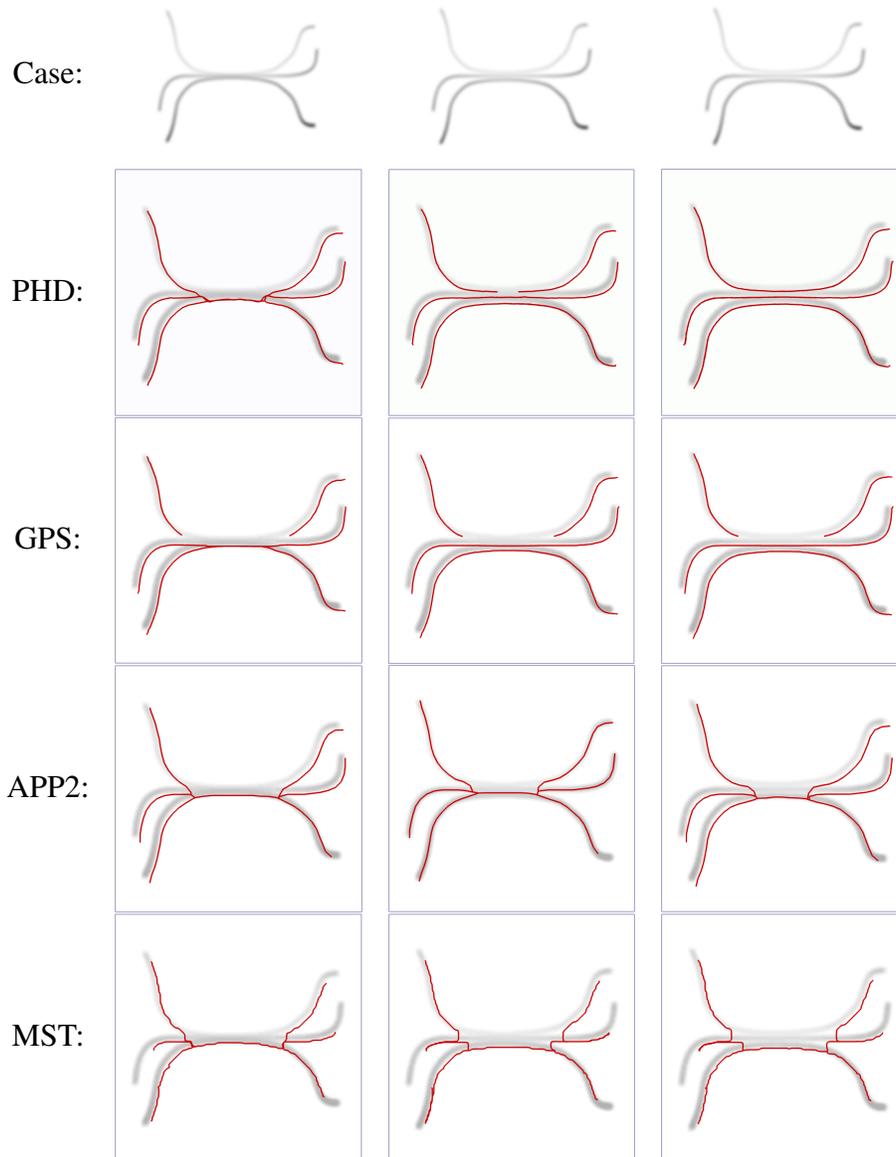


Figure S7: Ability of the tested methods to separate three fibers with different intensity and scale running closely in parallel. The examples show cases with gradually increasing distance between the fibers: overlap (left column), just separated (middle column), and clearly separated (right column). The tracing results of PHD, GPS, APP2, MST are overlaid (with slight offset) in red color.