# Hierarchical clustering of the pre-computed signals database to solve the parametric inverse light-scattering problem

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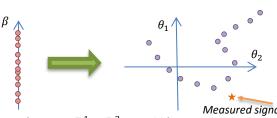
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## Parametric inverse problem

- Non-invasive characterization of single particles through the solution of the inverse light-scattering problem(LSP)
- Particle models with several free parameters [1]
- Preliminary computed database of simulated signals [2]

# Process experimental signal: Traverse the tree starting from the root (top) node. For each node (cluster) k do the following (D – current upper bound for $\min d_i$ ): Compute $d_k' \stackrel{\text{def}}{=} ||x - c_k||$ , $d_{\min} \stackrel{\text{def}}{=} d_k' - r_k$ and test $d_{\min} > D$ No Yes Discard the cluster and all its descendants

### **Problem statement**



- Illustration is for a map  $\mathbb{R}^1 \to \mathbb{R}^2$ . In real-life examples the direct map is, e.g.,  $g \colon \mathbb{R}^4(\beta) \to \mathbb{R}^{61}(y)$  if the signal is an angle-resolved light-scattering pattern.
- Inverse problem is reduced to (slow) global optimization [1]:  $\beta_*(x) \stackrel{\text{\tiny def}}{=} \operatorname{argmin} \|x g(\beta)\| \, .$ 
  - Further approximated by nearest-neighbor interpolation using precomputed database  $Y=\{\beta_i,y_i\stackrel{\mathrm{def}}{=}g(\beta_i)\}$  [2]:

$$\beta_*(x) pprox \beta_{i_*}$$
,  $i_* \stackrel{\text{def}}{=} \operatorname{argmin} d_i$ ,  $d_i \stackrel{\text{def}}{=} \|x - y_i\|$ .

 Statistical quantities (math. expectations, SDs, marginal probabilities, etc.) are obtained in the Bayesian framework

$$< f(\beta) > \stackrel{\text{def}}{=} \int_{\beta} \mathrm{d}\beta f(\beta) P(\|x - g(\beta)\|) \approx \sum_{i} f(\beta_{i}) P(d_{i})$$

where f – any function, P – conditional probability.

 Evaluating all distances (each ~ 100 FLOPs) is still slow for real-time applications (100 signals/second). Need to be accelerated!

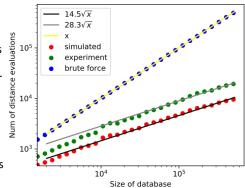
# **Calculation of statistical quantities**

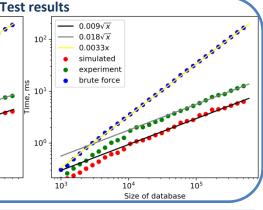
- Each discarded cluster decreases the number of distance evaluations ⇒ decreases the computational time
- Statistical quantities can still be estimated. For each discarded cluster  $A_k$  with  $N_k$  last-level descendants (elements of the original database)

$$\sum_{i \in A_k} f(\beta_i) P(d_i) \approx M_k f(\beta_k') P(d_k'), \qquad \beta_k' \stackrel{\text{def}}{=} \frac{1}{M_k} \sum_{i \in A_k} \beta_i$$

• Error should not be large, since larger  $M_k$  are for larger  $d_k'$ , hence smaller  $P(d_k')$ .

- Database of  $5 \times 10^5$  LSP of blood platelets, characterized by 4 parameters (randomly distributed). LSP has 61 values in the range [10,70], corresponding to the measurements using the scanning flow cytometer.
- Test data are 1000 of either experimental or separately simulated LSPs. Results are averaged.
- Smaller databases are obtained as a subset of the original one.
- Both distance evaluations and whole time were measured in comparison with previous brute-force search.





# **Conclusion and plans**

Presented algorithm significantly accelerates finding the nearest neighbor:

Additionally update  $D \rightarrow d'_k + r_k$  if beneficial

- Number of distance evaluations scales with square root of database size
- Up to 49/25 times decrease in this number for simulated/experimental data
- Total wall times decrease up to 25/13 times (cache issues?)

### Further research:

- Optimal clustering of points (metric k-center problem)
- Other tree structures (e.g., cover tree)
- Open-source library, including computation of statistical quantities

### References

- 1. Strokotov DI et al. J. Biomed. Opt. 2009;14:064036
- 2. Moskalensky AE et al. J. Biomed. Opt. 2013;18:017001

