

Fuzzy Set Theory in Computer Vision: **Example 7**

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Linear Convex Sum

In most works, we assume that a kernel \mathcal{K} is composed by a weighted combination of pre-computed kernel matrices, i.e.,

$$\mathcal{K} = \sum_{k=1}^m \sigma_k K_k,$$

where there are m kernels and σ_k is the weight applied to the k th kernel. The domain of σ is very important and many MKL implementations only work for a single domain. For example, $\Delta_2 = \{\sigma \in \mathcal{R}^m : \|\sigma\|_2 = 1, \sigma_k \geq 0\}$ is the ℓ_2 -norm MKL.

Geometric way to think about LCS

$$\mathcal{K}_{ij} = \langle \phi_\sigma(\mathbf{x}_i), \phi_\sigma(\mathbf{x}_j) \rangle = \sum_{k=1}^m \sigma_k (K_k)_{ij} =$$

$$\begin{pmatrix} \sqrt{\sigma_1} \phi_i^1 \\ \dots \\ \sqrt{\sigma_m} \phi_i^m \end{pmatrix}^t \begin{pmatrix} \sqrt{\sigma_1} \phi_j^1 \\ \dots \\ \sqrt{\sigma_m} \phi_j^m \end{pmatrix}$$

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3. Collect sorting indices π , such that
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4. Use a GA to learn the FM g , such that the classification accuracy of a learner (e.g., SVM) on \mathcal{K} is maximized

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- ▶ The genes of the GAMKL_p algorithm are the values of the m weights, i.e., $\{\sigma_1, \sigma_2, \dots, \sigma_m\}$
- ▶ To ensure the GAMKL_p genes lie in the ℓ_p -norm domain Δ_p , all candidate genes $\tilde{\sigma}$ are ℓ_p -norm normalized to form σ as

$$\sigma = \frac{\tilde{\sigma}}{\sqrt[p]{\sum_{i=1}^m |\tilde{\sigma}_i|^p}}$$

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- ▶ Of course, this are FI centric questions