

# Random Forests on Hierarchical Multi-Scale Supervoxels for Liver Tumor Segmentation in Dynamic Contrast-Enhanced CT Scans

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## Context

We tackle **multi-label tissue classification** covered through supervised ensemble learning towards accurate **tumor segmentation** :

- high diversity in shape, location and size
- wide appearance heterogeneity
- severe class overlap in feature space
- ambiguous boundaries

Recent approaches capture long-range spatial context but are limited in their ability to deal with **spatial adaptivity & appearance heterogeneity**

**Our motivation.** Enable random forest (RF) [1] to find itself the best data sampling by:

- combining RF and **hierarchical multi-scale** tree resulting from recursive supervoxel decomposition
- describing each leaf supervoxel as a sequence of supervoxels belonging to its ascendant hierarchy

## Application

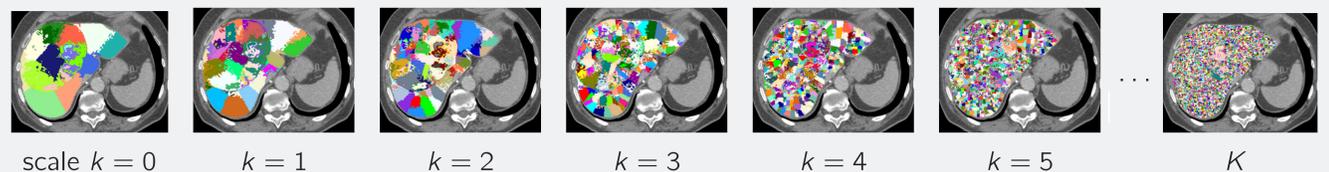
Clinical management of *hepatocellular carcinoma* (HCC) [2], most common type of liver cancer :

- requires the segmentation of **healthy liver**, **tumoral active** and **necrotic** tissues
- from dynamic contrast-enhanced (DCE) CT scans: HCC characterized by arterial enhancement followed by venous washout in response to contrast agent injection



## Contributions

**Hierarchical multi-scale supervoxel representation** | following [3]

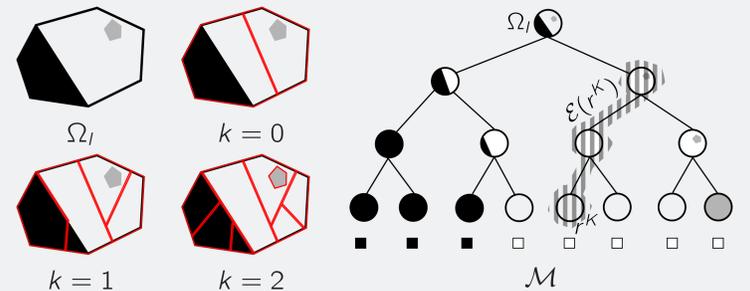


- liver area  $\Omega_l$  decomposed into a set of  $K + 1$  partitions  $\mathcal{P}^k$

-  $\mathcal{P}^k$  is a collection of SLIC [4] compact 3D supervoxels  $\{\mathbf{r}_i^k\}$  such that  $\mathbf{r}_i^k \cap \mathbf{r}_{j \neq i}^k = \emptyset$  and  $\bigcup_i \mathbf{r}_i^k = \Omega_l$

-  $\{\mathcal{P}^k\}$  encoded in the layers of a multi-resolution tree  $\mathcal{M} = \{\mathcal{M}^k\}$

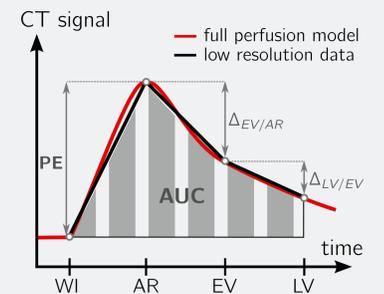
-  $\mathcal{M}^k$  maps each supervoxel  $\mathbf{r}_i^k \in \mathcal{P}^k$  to a set of child supervoxels  $\{\mathbf{r}_j^{k+1}\} \subset \mathcal{P}^{k+1}$  s.t.  $\mathbf{r}_i^k = \bigcup_j \mathbf{r}_j^{k+1}$



**Hierarchical multi-scale supervoxel-based RF** | extends [5] from single to multi-scale since intrinsic tissue properties may emerge at different scales for different tissues

- ① Build hierarchical multi-scale tree  $\mathcal{M}$
- ② Assign visual features  $\theta(\mathbf{r}^k)$  to all supervoxels  $\mathbf{r}^k$  in each partition  $\mathcal{P}^k$  with  $k \in \{0, \dots, K\}$

Related to	Features	Nb
Intensity	mean intensity + std dev.	4 + 4
Gradient	mean gradient magnitude + std dev.	4 + 4
Multi-phase	peak enhancement (PE)	1
	inter-phase diff. $\Delta_{EV/AR}$ , $\Delta_{LV/EV}$	2
	area under enhancement curve (AUC)	1



③ Associate to each supervoxel  $\mathbf{r}^k$  at finest scale  $K$  all the supervoxels of decreasing scale belonging to its ascendant hierarchy including itself:  $\mathcal{E}(\mathbf{r}^k) = \{\mathbf{r}^k\}_{k \in \{0, \dots, K\}}$

④ Define a new feature vector  $\gamma(\mathbf{r}^k)$  associated to each  $\mathbf{r}^k \in \mathcal{P}^k$  as the concatenation of all visual features assigned to supervoxels of  $\mathcal{E}(\mathbf{r}^k) \Rightarrow$  **powerful multi-scale description of finest scale supervoxels**

⑤ Tissue classification based on supervoxels of scale  $K$  carried out via standard RF [1]

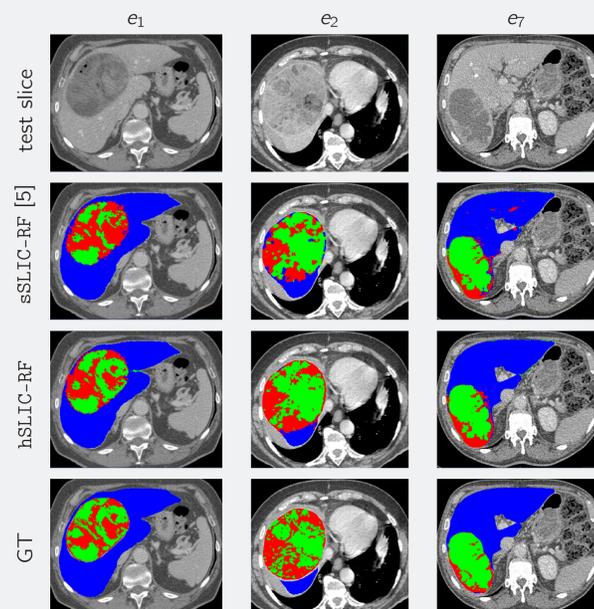
## Results

**Dataset.** 8 examinations  $\{e_1, \dots, e_8\}$  stemming from patients with HCC with 6 equally reparted 2D axial slices labeled by 4 experts in hepato-digestive surgery.

**Experiments.** Comparison between hierarchical multi-scale supervoxel-based RF (hSLIC-RF) and single-scale supervoxel-based RF [5] (sSLIC-RF) with optimal shared and  $e_j$ -dependent supervoxel resolutions.

methods	optimized sSLIC-RF [5]		hSLIC-RF
	shared	$e_j$ -dependent	multi-scale
DICE <sub>activ</sub>	76.5 ± 10.1	78.7 ± 9.18	<b>80.4 ± 8.81</b>
DICE <sub>necro</sub>	85.3 ± 12.5	<b>86.9 ± 9.51</b>	<b>86.9 ± 10.5</b>
DICE <sub>prcm</sub>	94.3 ± 4.12	94.9 ± 3.85	<b>95.5 ± 3.56</b>
DICE <sub>tumor</sub>	88.9 ± 8.51	89.4 ± 6.12	<b>91.0 ± 6.99</b>

- significant impact of scale selection in single-scale context
- hSLIC-RF outperforms the upper bound reachable by sSLIC-RF  $\Rightarrow$  confirms the benefits of our **adaptive data sampling scheme**
- stronger spatial regularization inherited from the capacity of multi-scale SLIC supervoxels to adhere to image boundaries



## Further work

- multi-examination training to make our strategy becoming fully automatic
- HCC management  $\Rightarrow$  correlation between tumor necrosis rate and survival rates
- longitudinal liver tumor study
- extension to other tumor types, organs and modalities

## References

- [1] Breiman L. Random Forests. *Machine learning*, 45(1):5–32, 2001.
- [2] Forner, A., Llovet, J. M and Bruix, J. Hepatocellular carcinoma. *The Lancet*, 2012
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- [4] Radhakrishna A. et al., SLIC superpixels compared to state-of-the-art superpixel methods. *IEEE Transactions on Pattern Analysis and Machine Intelligence*, 34(11):2274–2282, 2012.
- [5] Conze, P.-H., Rousseau, F., Noblet, V., Heitz, F., Memeo, R. and Pessaux, P. Semi-automatic Liver Tumor Segmentation in DCE-CT Scans Using Random Forests and Supervoxels. *Machine Learning in Medical Imaging*, 2015, 212–219, 9352