Understanding Neural Networks for Image Segmentation

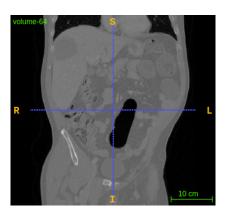
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Target application: medical image segmentation



Abdominal CT scan



Liver and tumor segmentation

Why automate?

- · needed for treatment plans
- · costly (time+effort) to perform by hand
- less interobserver variability → better accuracy

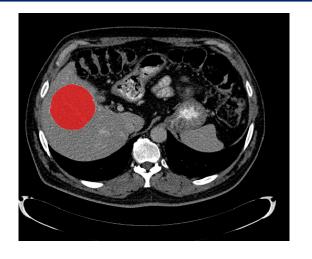
errors in segmentation = errors in radiation treatment

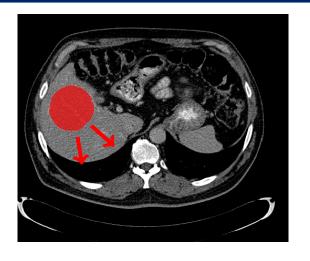
Goal: understand why CNNs work so well

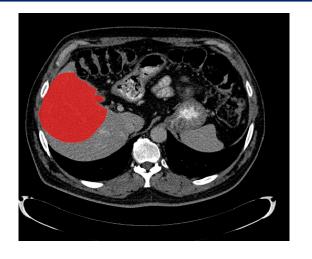
- Compare PDEs to CNNs
 - Aside: Upwind schemes
- Build CNNs like PDEs
- 3 Analyze kernels and explain performance
 - Entrywise comparison
 - SVD comparison

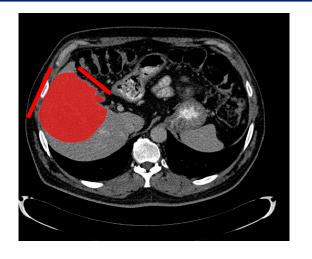
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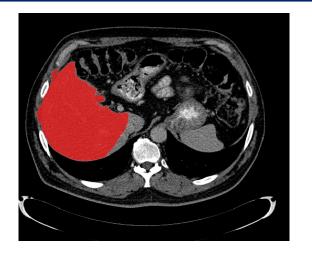
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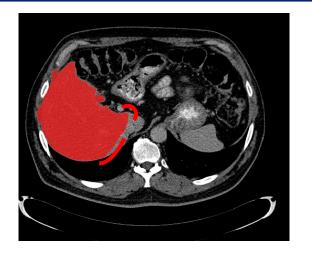


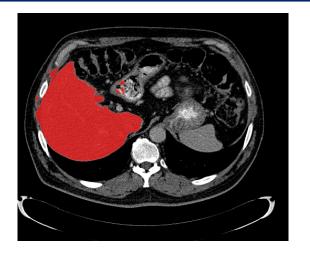












PDE of choice: Level Set Equation

$$\partial_t u - \alpha \underbrace{g_l}_{\text{balloon force}} - \beta \underbrace{g_l \kappa(u)}_{\text{mean curvature}} - \gamma \underbrace{\nabla g_l \cdot \nabla u}_{\text{convection}} = 0$$

- Well-established theory to analyze approximation, stability
- Upwind finite differences + fast marching method
- · Semiautomated: requires initialization by user
- Works for simple problems only: relies on edge information

Why upwind schemes?

- convergence like finite differences
- improved stability

M-Matrices

Definition

A matrix $A \in \mathbb{R}^{n \times n}$ is a *M-Matrix* if:

- $a_{ij} \leq 0$ for all $i \neq j$
- $\mathbf{Q} A^{-1}$ exists
- 3 $A^{-1} \geq 0$

Checking for M-matrices

Theorem

The matrix $A \in \mathbb{R}^{n \times n}$ is a M-matrix if and only if:

- $\mathbf{0}$ $a_{ij} \leq \mathbf{0}$ for $i \neq j$
- **2** $\exists \mathbf{x} \in \mathbb{R}^n$ such that $A\mathbf{x} > 0$.

The vector **x** is called the majorizing element of A.

M-matrices provide stable numerical methods

An example: convection

Consider on
$$\Omega = [0, 1]$$
 the convection equation

$$\partial_t u = a(x)\partial_x u$$

+ Dirichlet BC's

convection a both positive and negative

Discretization

Discretize in space with a finite difference stencil:

$$\partial_x u_i \approx \frac{1}{h} \sum_{j=-1}^1 w_{i,j} u_{i+j}$$

Examples:

- $w_{i,-1} = -1$, $w_{i,0} = 1$, $w_{i,1} = 0$: forward FD
- $w_{i,-1} = \frac{-1}{2}$, $w_{i,0} = 0$, $w_{i,1} = \frac{1}{2}$: centered FD

Discretization

$$\partial_t u = a(x)\partial_t u$$

$$\partial_t u_i \approx a_i \sum_{j=-1}^1 \frac{1}{h} w_{i,j} u_{i+j}$$

$$\partial_t \mathbf{u} = L \mathbf{u}$$

Upwind scheme

Choose our upwind scheme:

$$w_{i,-1} = -\max\{0, sgn(a_i)\}$$

 $w_{i,0} = 1$
 $w_{i,1} = \min\{0, sgn(a_i)\}$

Upwind scheme

$$L_{i,i-1} = -\frac{1}{h} \max\{0, a_i\}$$
 $L_{i,i} = -\frac{1}{h} |a_i|$
 $L_{i,i+1} = -\frac{1}{h} \min\{0, a_i\}$

Stability of our discretization

Lemma

L is an M-matrix.

Proof.

- $L_{i,j} \leq 0$ for $i \neq j$
- The vector **x** with $x_i = ih$ is a majorizing element.

Comparison: PDE vs CNN

	Can analyze?	Accurate?
PDE	yes	sometimes
CNN	no	yes

Goal: accurate method we can analyze

Similarities between LSE and CNN

	LSE	CNN		
Convolution	finite difference kernel	learned kernel		
	$\frac{1}{h^2} \begin{bmatrix} 0 & 1 & 0 \\ 1 & -4 & 1 \\ 0 & 1 & 0 \end{bmatrix}$	$\begin{bmatrix} k_{11} & k_{12} & k_{13} \\ k_{21} & k_{22} & k_{23} \\ k_{31} & k_{32} & k_{33} \end{bmatrix}$		
ReLU	upwind scheme	activation function		
	$ \max(0, D^+ * u) + \min(0, D^- * u) $	$\max(0,K*x+b)$		
		4.0		

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Build a CNN like a LSE solver

Level Set Equation → Level Set Network

- ① Discretize Level Set Equation
 - Explicit forward Euler in time
 - Upwind finite differences in space
- ② Forward Euler → residual skip connections
- ③ Upwind finite differences → convolutions and ReLU

Forward Euler time discretization

$$\frac{u^{(t+1)} - u^{(t)}}{\delta_t} - \alpha g_I - \beta g_I \kappa(u^{(t)}) - \gamma \nabla g_I \cdot \nabla u^{(t)} = 0$$

Edge indicator g_l

$$g_I(x) = \frac{1}{1 + \|\nabla I(x)\|^2}$$

becomes

$$g_I(x) = \frac{1}{1 + \sum_{j=1}^{N_{conv}} (\sigma_j * I(x))^2}$$

Convection term: Upwind Finite Differences

$$\begin{split} \nabla g_I \cdot \nabla u^{(k)} &= \max\{0, \partial_X g_{\mathcal{I}}\} D_X^+ * u^{(k)} - \max\{0, -\partial_X g_{\mathcal{I}}\} D_X^- * u^{(k)} \\ &+ \max\{0, \partial_y g_{\mathcal{I}}\} D_y^+ * u^{(k)} - \max\{0, -\partial_y g_{\mathcal{I}}\} D_y^- * u^{(k)} \end{split}$$

$$&= \text{ReLU}(\partial_X g_{\mathcal{I}}) D_X^+ * u^{(k)} - \text{ReLU}(-\partial_X g_{\mathcal{I}}) D_X^- * u^{(k)} \\ &+ \text{ReLU}(\partial_y g_{\mathcal{I}}) D_y^+ * u^{(k)} - \text{ReLU}(-\partial_y g_{\mathcal{I}}) D_y^- * u^{(k)} \end{split}$$

Curvature term $\kappa(u)$

$$\kappa(u^{(k)}) = \nabla \cdot \left(\frac{\nabla u^{(k)}}{\|\nabla u^{(k)}\|} \right)$$

becomes

$$\nabla \hat{u}_{i}^{(k)} = \rho_{i} * u^{(k)}$$

$$\kappa(u^{(k)}) = \sum_{j=1}^{N_{conv}^{2}} \sigma_{j} * \left(\sum_{i=1}^{N_{conv}^{1}} \frac{\nabla \hat{u}_{i}^{(k)}}{\left\| \nabla \hat{u}_{i}^{(k)} \right\| + \varepsilon} \right)$$

Implementation

- If convolution kernels are finite difference kernels, we recover LSE
- If convolution kernels are learned, we construct LSN
- Python + Tensorflow / Keras implementation
- Our LSE agrees with ITK-SNAP, a common LSE-segmentation program

LSN: Results

K-Fold	LSE	LSN Test	LSN Validation	UNet
0	0.736	0.837	0.619	0.912
1	0.600	0.847	0.729	0.919
2	0.483	0.116	0.005	0.874
3	0.730	0.827	0.606	0.895
4	0.643	0.831	0.596	0.915
Avg	0.604	0.692	0.511	0.903
Avg $-\{2\}$	0.640	0.837	0.638	0.911

Table: DSC scores for each fold, from training the level set network.

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Identification of Kernels

- Are CNN convolution kernels finite difference stencils?
- Are they close to finite difference stencils?
- What about other standard image processing kernels?

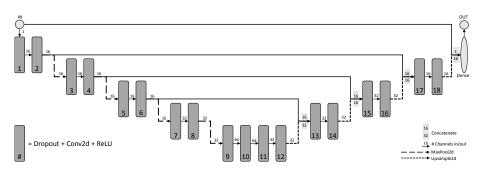
Numerical analysis kernels

- Laplacian
- Edge detection
- Identity

Image processing kernels

- Gaussian blur
- Local mean
- Sharpen

Setup of CNN



Trained on MICCAI LiTS 2017 dataset for liver segmentation

Kernel Analysis: Comparing entries

- For each layer, separate each channel's 3×3 convolution kernel
- Flatten each 3 \times 3 kernel into a vector $\in \mathbb{R}^9$
- Cluster with k-means
- Project down using PCA
- Project known numerical analysis and image processing kernels

Kernel Clustering Results

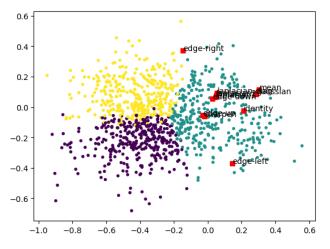


Figure: Convolution Layer 11 (encoder)

Kernel Clustering Results

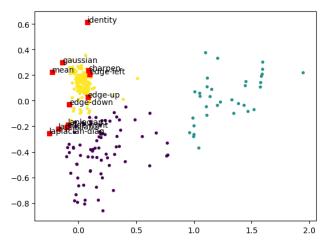


Figure: Convolution Layer 14 (decoder)

- **1** Construct matrix $A_{[K]} \in \mathbb{R}^{n_x n_y \times n_x n_y}$ describing convolution with K
- 2 Compute singular values of linear operator
- 3 Compute singular values of clinical image processing kernels
- Assign closest clinical feature F that has smallest spectral distance to K

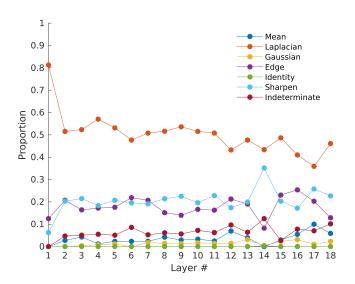
Kernel
$$K = \begin{bmatrix} k_{-1,-1} & k_{-1,0} & k_{-1,1} \\ k_{0,-1} & k_{0,0} & k_{0,1} \\ k_{1,-1} & k_{1,0} & k_{1,1} \end{bmatrix}$$

For
$$i \in \{-1, 0, 1\}$$
,

$$\begin{split} A_{[K]} &= U_K \Sigma_K V_K^T & \forall K \in \{ \text{layers} \} \\ A_{[F]} &= U_F \Sigma_F V_F^T & \forall F \in \{ \text{features} \} \end{split}$$
 find $\underset{F \in \{ \text{features} \}}{\text{arg min}} \| \Sigma_K - \Sigma_F \|_1$

Label as 'indeterminate' if not within 10% of largest singular value

Kernel Analysis Results



Conclusions

- Level Set Equation ≠ Level Set Network ≠ UNet
- Framework for using same operations (convolutions + ReLU) for both NNs and PDEs
- Framework for GPU-supported finite differences in Tensorflow
- Examine how learned CNN kernels change across different layers

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