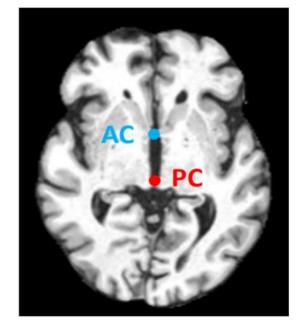
# Scaling deep RL on landmark localization

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### Background

- The field of radiology is moving towards a collaborative space between human experts and Al
- Automatic anatomical localization is an integral part of an AI radiology framework
- Anatomical localization has diverse application cross multiple task such as image segmentation, registration, and classification
- Deep reinforcement learning (RL) has emerged as one of the best techniques for landmark localization.



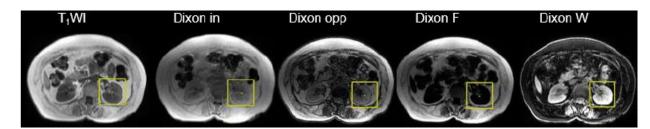
The anterior commissure (AC) and posterior commissure (PC) points in brain MRI.



the single 2D agent locating different landmarks in 2D slices

### Deep RL setup

Source: https://github.com/bocchs/MIDRL-2D

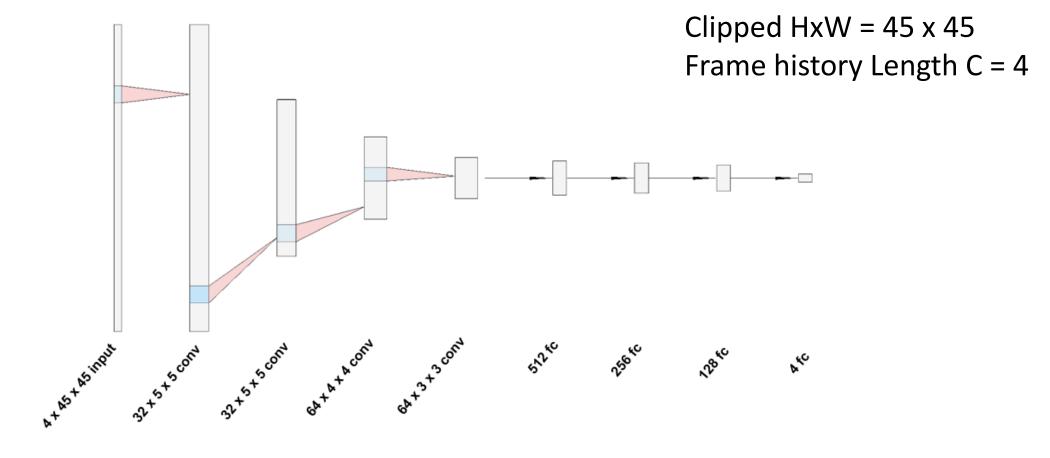


- Environment: Radiological image
- State: Sequence of areas within the image (bounding box)
- Actions: move bounding box in one direction (+/-)
- Reward: change in Euclidean distance to landmark
  - + if moved closer to landmark
  - - if moved away
  - Clipped between 1 and 1
- Method: Q learning with experience replay



the single 2D agent locating different landmarks in 2D slices

### 2D DQN single agent



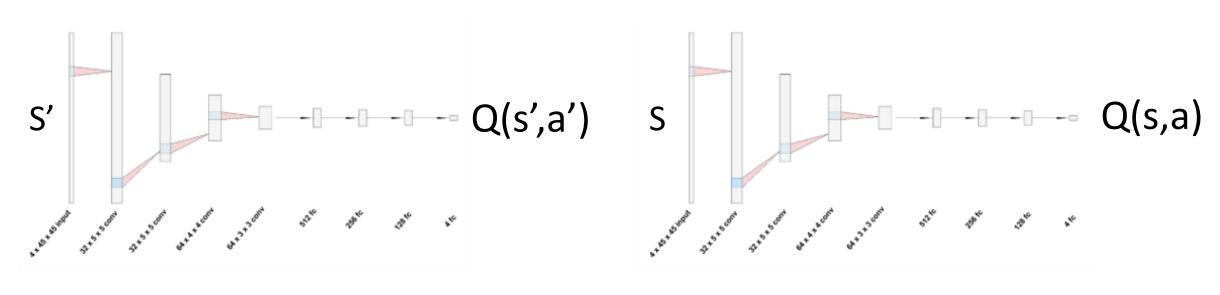
- Input: bounding box regions from last 4 time steps
- Output: Q-value for each action (x++, x--, y++, y--)

### Training

Target Network DQN\_T

Gamma = 0.9, batch =48, lr = 0.001, Optimizer: Adam Gradient Clipping, lr scheduler e-greedy policy with decreasing e

#### Policy Network DQN\_P



Target = r + gamma \* Q(s',a') Loss = Huber loss (Target, Q(S,a))

### Experience replay

- Experience: (S, A, R, S')
- TD or Q-learning problems:
  - Waste of Experience
  - Correlated Updates
- Replay Buffer

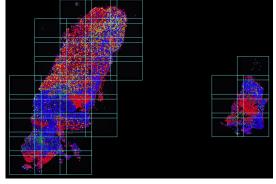


Examples of single 2D agent locating different landmarks in 2D slices. Red is the target bounding box, yellow is the agent bounding box.



## Challenges in Medical Decision Support System

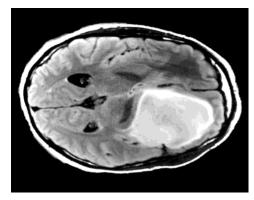
- 1. Data sets with very large cardinalities and high dimensionalities are emerging (100M+ to few B, 10~35 dim)
- 2. Memory problem with replay buffer
- 3. Existing dimensional reduction tools have suboptimal scaling
- 4. Data geo-distributed. direct aggregation impossible
- 5. Free movement of clinic data restricted



Billions of pixels from one set of tumor biopsies



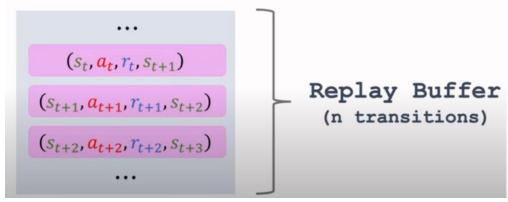
Geo-distributed



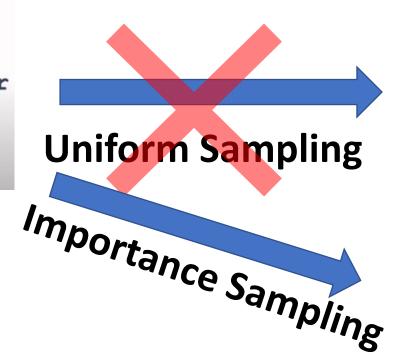
Private and sensitive healthcare data

### Experience replay

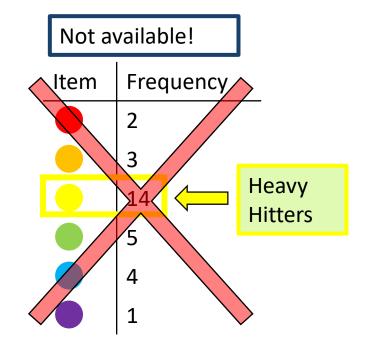
- Stream of Experience: (S0, A0, R0, S1), (S1,A1,R1,R2)....
- Replay Buffer



- 1. Giving high priority to high TD error experiences
  - (S,A,R,S,delta) tuple
- 2. Sketching the Experience



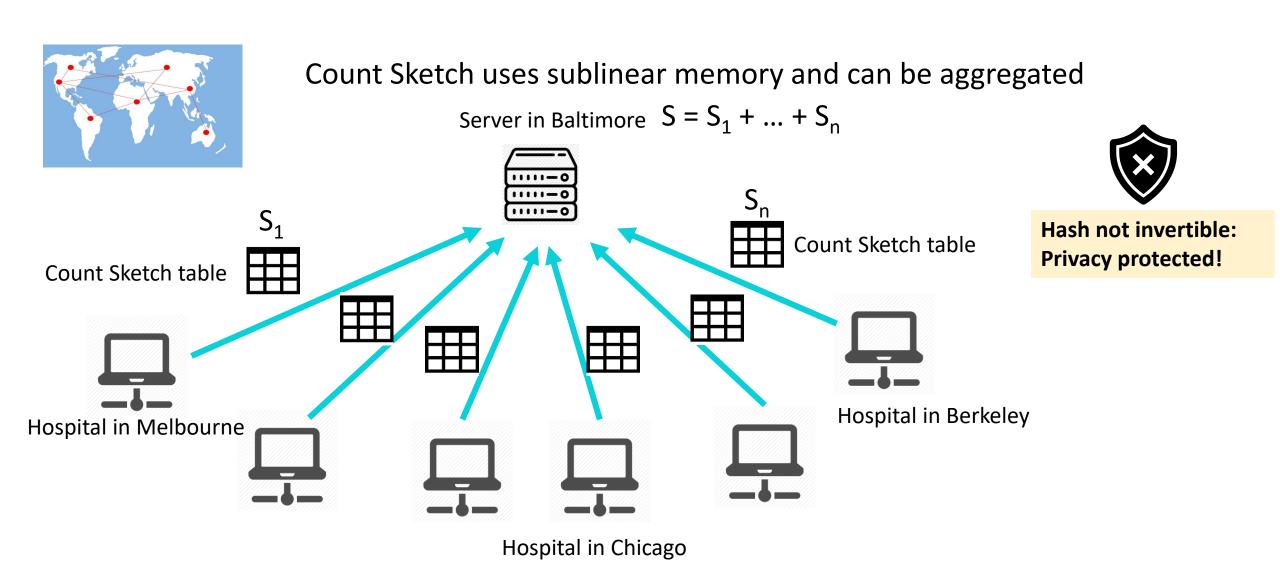
#### Count Sketch Algorithm Count Sketch Hash Table C1 C2 C3 C4 **C5** Stream: ID = 3 -7 R1+1 +6 Freq = 12-2 0 ID = 16 -9 -2 +1+1 Freq = 0+3



R1	sign	bucket
	+1	3
	-1	2
	-1	4
	-1	5
	+1	4
	-1	1

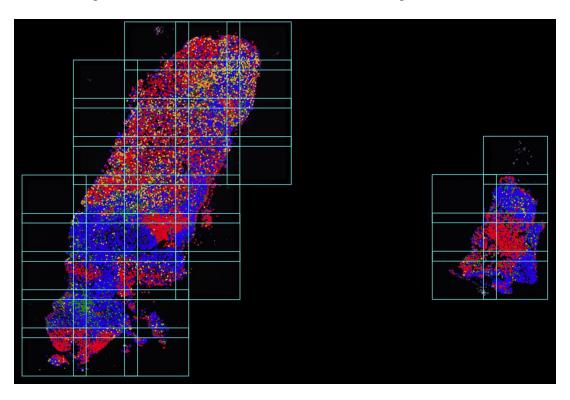
R2	sign	Bucket
	-1	2
	+1	5
	-1	1
	+1	3
	+1	2
	-1	1

#### Count Sketch Algorithm



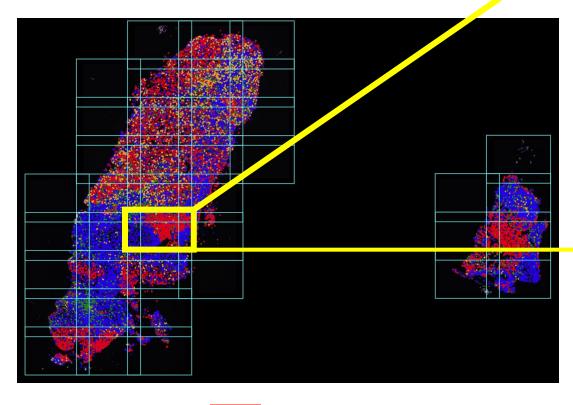
### Quick check on Count Sketch performance

Cancer Immunotherapy
54M pixels in 35 bands ~ 2B pix

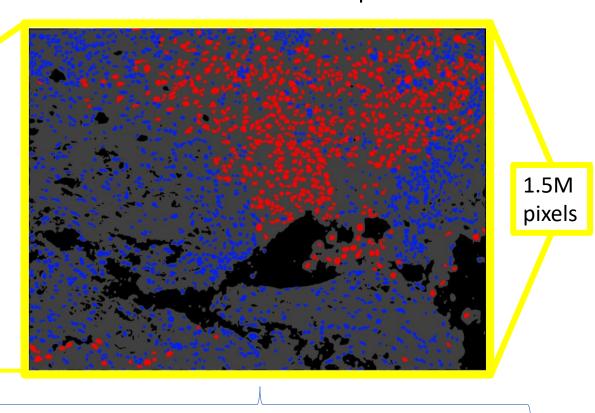


## Cancer Imaging Data

Data: Multiplex image mosaic of a melanoma biopsy with multiple fluorescent markers



40 Images, each is 35 Layers  $\mathbf{x}$  1.5M pix Total = 40 x 1.33M x 35  $\approx$  2B pix

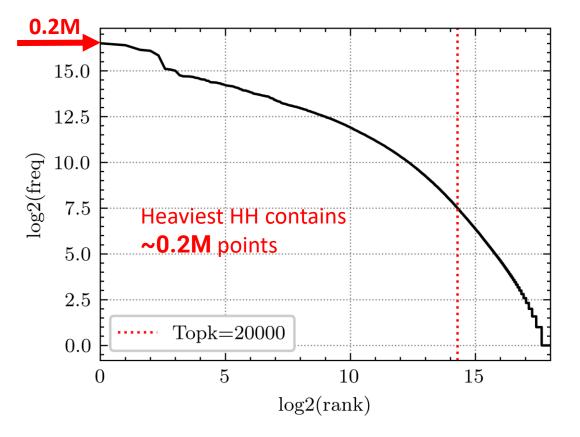




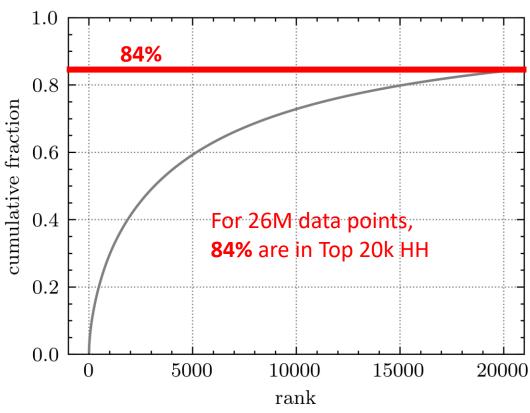
Tumor

35 Layers

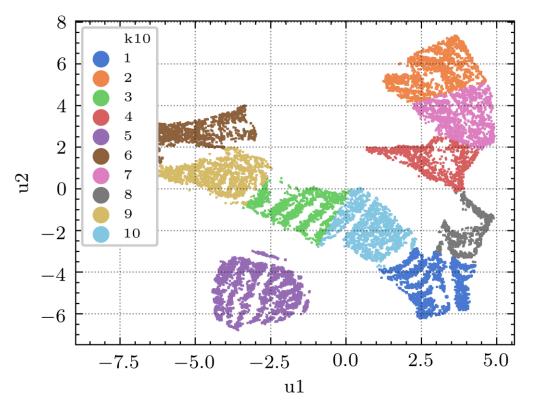
#### **HH cardinality vs Rank (first is top)**



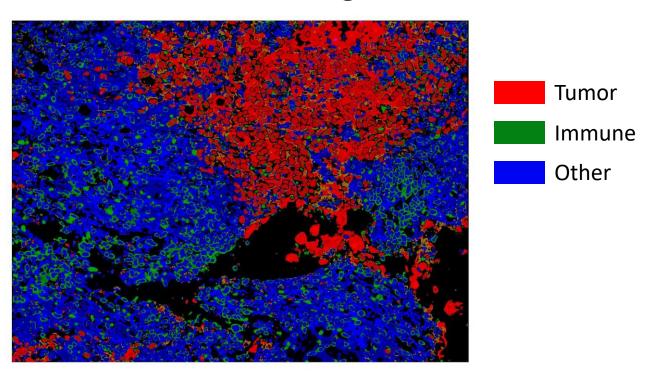
#### **HH Cumulative Fraction vs Rank**



#### **UMAP + KMEANS (20K HH)**



#### **Clusters shown on one image**



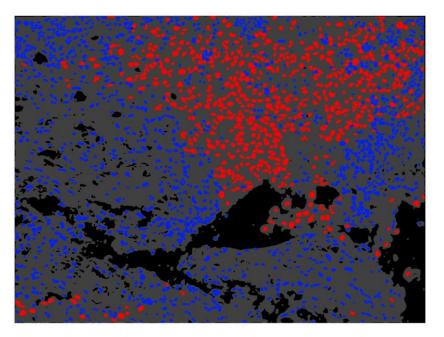
#### Results

Our pixel-based clustering is in excellent agreement with results derived from state-of-theart cell segmentation using also shape information

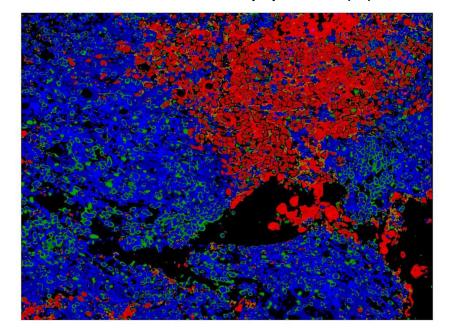
	Tumor (A)	Other (A)
Tumor (B)	95792	3675
Other (B)	6795	108630

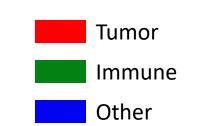
Tumor Misclassification Rate: 3.5%

Label image from cell segmentation software (A)

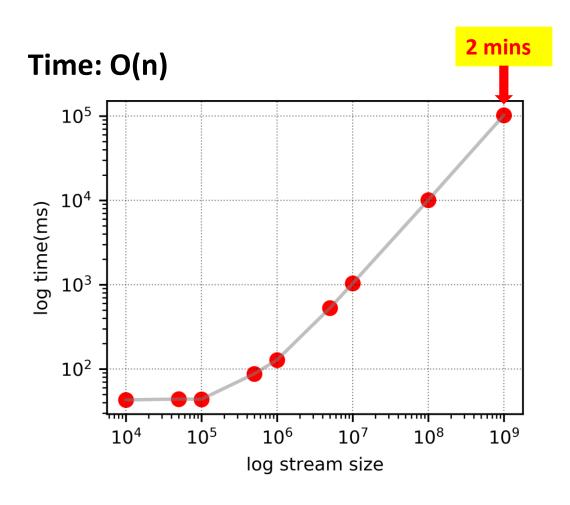


Pixel classifications from our Sketch & Scale pipeline (B)





## Scaling Performance



#### Memory: O(log(n))

- Sketch Table Parameters:
  - R = 16
  - C = 100000
  - Top k = 20000

Count Sketch Preprocessor can find low-dimensional embedding of state-action space in a time and memory efficient way.

#### Conclusion: Count Sketch Preprocessor can scale up deep Q learning capacity

Quantization

(S,A,R,S')-tuple

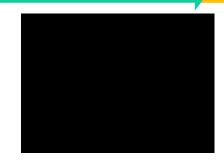
Encoded Stream

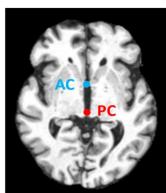
**Count Sketch** 

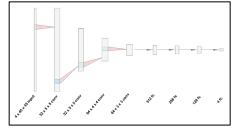
Coreset w/
Frequency List

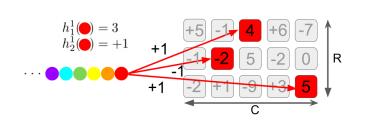
DQN

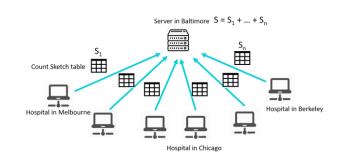
- 1. Deep RL for medical decision support system
- 2. Adding Count Sketch preprocessor can find real clustering in large data sets, using streaming
- The sketches stored in logarithmic space can be easily moved, shared and aggregated.
- 4. Privacy preserved as re-indentification impossible

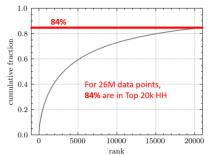


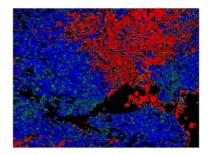












#### Reference

- Amir Alansary, ...Daniel Rueckert. Evaluating Reinforcement Learning Agents for Anatomical Landmark Detection. Medical Image Analysis, 2019.
- https://github.com/bocchs/MIDRL-2D
- Wei, V., Ivkin, N., Braverman, V., and Szalay, A., "Sketch and Scale: Geo-distributed tSNE and UMAP", IEEE BigData 2020.
- https://github.com/ViskaWei/sketch-scale