# **Clinical Decision Making Using Deep Learning**

Bradley J Erickson, MD PhD Mayo Clinic

#### Disclosures

Grant Funding: DK-90728
 Commercial: FlowSIGMA, VoiceIT, OneMedNet

There are known knowns; there are things we know that we know.

There are known unknowns; that is to say, there are things that we now know we don't know.

But there are also unknown unknowns - there are things we do not know we don't know.



There are known knowns; there are things we know that we know.

There are known unknowns; that is to say, there are things that we now know we don't know.

But there are also unknown unknowns - there are things we do not know we don't know.

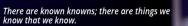


There are known knowns; there are things we know that we know.

There are known unknowns; that is to say, there are things that we now know we don't know.

But there are also unknown unknowns - there are things we do not know we don't know.





There are known unknowns; that is to say, there are things that we now know we don't know.

But there are also unknown unknowns - there are things we do not know we don't know.



### **Clinical Decision Making**

- General Decision Making Requires
- Data
- Knowledge: meaning of data
- Judgement: meaning of data where knowledge is not decisively clear
- Trust: experience that judgements are defendable

#### **Clinical Decision Making**

- Decision Making Requires
- Data
- Knowledge: meaning of data
- Judgement: meaning of data where knowledge is not decisively clear
  Trust: experience that judgements are defendable
- Clinical Decision Making also Requires
- Data Interpolation: This patient versus trial group
- Relevant research is often gray (no trial has 100% response rate)
- Relationship between research outcomes and causality can be difficult to understand

Anything you can do, AI can do better



## **Clinical Decision Making and DL**

DL requires much data

### **Clinical Decision Making and DL**

DL requires much dataDL requires much *annotated* data

## **Clinical Decision Making and DL**

- DL requires much data
- DL requires much *annotated* data
- DL algorithms find associations, not proof

#### **Clinical Decision Making and DL**

- DL requires much data
- DL requires much *annotated* data
- DL algorithms find associations, not proof
- DL algorithms have poor explain-ability (so far)

#### **Clinical Decision Making and DL**

- DL requires much data
- DL requires much *annotated* data
- DL algorithms find associations, not proof
- DL algorithms have poor explain-ability (so far)
- Proposed solutions must have clinical value

#### DL and Clinical Diagnosis

DL tools that increase 'attentiveness'
 Traditional role for screening CAD
 Usually most helpful to trainees or generalists



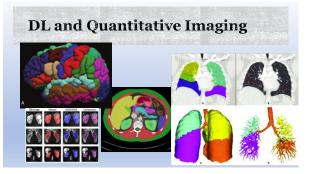




#### **DL and Image Reconstruction**

Several papers have shown DL can reconstruct high quality ('normal dose') images with significantly reduced dose
 PET tracer dose

- CT dose
   MR acquisition time
- This is likely to be readily accepted in medical imaging



#### **Current Results: Brain** - Fully automatic segmentation of Brain 127 parts of the brain semi-automatically labeled with human expert verification • 2000 train, 350 test • 30 seconds to segment! MRI Truth Our Segmentation Dice scores: Mean: 0.954 Range: 0.852 - 0.989

#### **Current Results: CT Abdomen**

- 100 CT abdomens for range of diseases

- 80 traced by 1 human, used for training • 20 were used as test/truth
- Hand labeled by 5
- STAPLE used to create 'truth'
- Dice scores range from 0.98 for liver to 0.7 for adrenals and renal veins
  Now using machine labeled as starting point for human correction

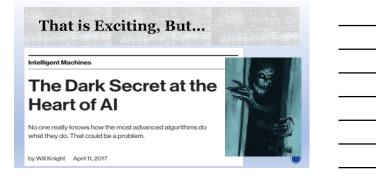


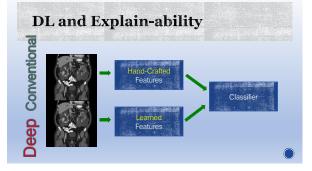
#### **DL Enables Radiogenomics**

N=498 subjects suing T2-weighted images from Mayo, UCSF, TCIA 398 for training, 100 for testing 50 layer ResNet, VGGNet, Inception

IDH1         0.95         0.95         0.95           1p19q Co-Del         0.91         0.85         0.87           ATRX         0.93         0.89         0.91           MGMT Methylation         0.95         0.95         0.95	Marker	Sens	Spec	Accuracy
ATRX 0.93 0.89 0.91	IDH1	0.95	0.95	0.95
	1p19q Co-Del	0.91	0.85	0.87
MGMT Methylation 0.95 0.95 0.95	ATRX	0.93	0.89	0.91
	MGMT Methylation	0.95	0.95	0.95

\*Korfiatis, submitted

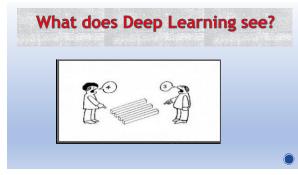


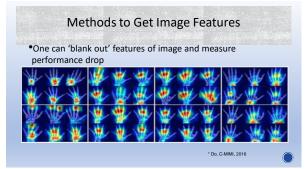


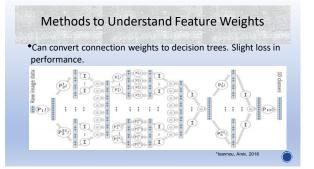
#### How are Features Determined?

Many traditional radiology features start with little physical basis

- Calcification
- Breast cancers have micro-calcifications (most calcification is not cancer)
- $\ensuremath{\cdot}$  Gliomas usually have no calcifications, but if they do, they are coarse
- Lymphomas don't calcify until they are treated
- The pathological correlate can be seen, but not always fully explained.
- Expected imaging methods based on pathology sometimes don't turn out due to complexity
  of the living human environment or because we don't fully understand the biology (which can
  lead to new insights)







#### **Implementation Issues**

What is the impact of acquisition technique?
 Some diseases will be impacted more

Device makers are best positioned to address this

#### **Implementation Issues**

Who is responsible for results / reliability
 Device manufacturers
 Medical Doctor

#### **Implementation Issues**

#### Informatics Challenges

- Must assure all and only the correct image(s) are sent to analytic • May need combinations of images (e.g. prior exam)
- How to capture, represent, transmit, store the output in an informative and computable fashion
- Clinical workflow: much like 3D lab renderings, these results must be produced reliably on a recognizable set of images, and results must either be completed before scanning is done, or process for notifying radiologist that results are ready must be put in place

#### Conclusions

- DL has tremendous potential to increase value of medical imaging Quantitative results
- New diagnostic capabilities
- This will demand
- Careful assessment of imaging technique impact on DL
   Increased use of DICOM SR/AIM to convey results efficiently
   New workflow methods to implement these tools in efficient way