

Interpretability

April 18, 2023



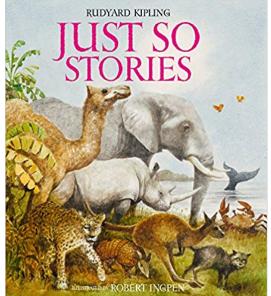


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Interpretability Issues

- People understand simple models
 - George Miller, 7±2: "There seems to be some limitation built into us either by learning or by the design of our nervous systems, a limit that keeps our channel capacities in this general range."
 - "... the number of chunks of information is constant for immediate memory. The span of immediate memory seems to be almost independent of the number of bits per chunk ..."
 - Not surprising that one cannot "keep in mind" complex models
- What leads to complex models? And what to do about it?
 - Overfitting
 - Restrict model complexity; e.g., regularization
 - True complexity
 - Make up "just-so" stories that give a simplified explanation of how the complex model applies to specific cases
 - Trade off lower performance for simplicity of model



Trust

- Critical for adoption of ML models
 - Case-specific prediction Local Interpretability
 - Clinical decision support
 - Confidence in model Global Interpretability
 - Population health
- Recall my critique of randomized controlled trials
 - Simplest cases (no comorbidities), smallest sample needed for significance test, shortest follow-up time
 - · Results applied to very different populations
- Same concerns for ML models
 - Train and test samples often drawn from same population
 - Are results applicable elsewhere?

Explanation — Not a New Idea! Mycin, 1975

- Mycin (1974) used <u>backward-chaining</u> rules to determine whether a patient had a bacterial infection that needed to be treated, and how best to treat
- Collection of several hundred rules, each of which encoded a relatively independent fact
- <u>Certainty factors</u> encoded a theory of uncertain reasoning (tantamount to very strong independence assumptions, leading to problems)
- <u>Context</u> mechanism to fill in implicit clauses in rules;
 patient→site→infection→culture→organism→ drug

RULE092

IF we have identified organisms for which treatment is indicated THEN select a treatment that covers those organisms

RULE037

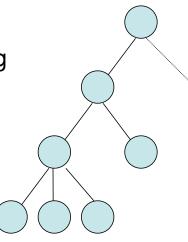
- IF the organism
 - I) stains gram positive
 - 2) has a coccus shape
 - 3) grows in chains

THEN

There is suggestive evidence (.7) that the identify of the organism is streptococcus

How Mycin Works

- · Dynamically generates an and/or tree via backward chaining
- To find out a fact
 - If there are rules that can conclude it, run them
 - · Otherwise, ask the user
- To run a rule
 - Find out if the facts in the premises are true (enough)
 - If they all are, then assert the conclusion (with suitable certainty factor)
- This traces out the equivalent of a flowchart, but by generating it on the fly from underlying rules
 - Knowledge is always applied when relevant
 - Can answer questions about how/why it is working



Explanations from a Backward-Chaining Rule Interpreter

- In light of the site from which the culture was obtained, and the method of collection, do you feel that a significant number of ORGANISM-1 were obtained?
 **WHY
- [1.0] It is important to find out whether there is therapeutically significant disease associated with this occurrence of ORGANISM-1 It has already been established that:
 - [1.1] the site of the culture is not one of those which are normally sterile, and
 - [1.2] the method of collection is sterile
- Therefore, if:
 - [1.3] the organism has been observed in significant numbers
- Then: there is strongly suggestive evidence (.9) that there is therapeutically significant disease associated with this occurrence of the organism
- [Also : there is strongly suggestive evidence (.8) that the organism is not a contaminant]

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• **WHY

- [2.0] . . . in order to find out a locus of infection in PATIENT-1 It has already been established that:
 - [2.1] the site of the culture is known, and
 - [2.2] the number of days since the specimen for the culture was obtained is less than 7
- Therefore, if:
 - [2.3] there is therapeutically significant disease associated with this occurrence of the organism
- Then: it is definite (1.0) that the site of the culture is a locus of infection in the patient

- ****WHY** [i.e., WHY is it important to find out a locus of infection in PATIENT-1?]
- [3.0] . . . in order to find out about an organism or class of organisms for which therapy should cover

It has already been established that:

- [3.1] the site of the culture is blood, and
- [3.2] the stain of the organism is grampos, and
- [3.3] the portal of entry of the organism is GI
- Therefore, if:
 - [3.4] [A- abdomen is a locus of infection in the patient, or B- pelvis is a locus of infection in the patient]
- Then : there is strongly suggestive evidence (.9) that enterobacteriaceae is a class of organisms for which therapy should cover

- ** **HOW 3.0** [i.e., HOW were you finding out about an organism or class of organisms for which therapy should cover?]
- [4.0] At that point RULE095 was being used. In addition :
 - [4.1] RULE162 was used to conclude that bacteroides is an organism for which therapy should cover, and
 - [4.2] RULE115 will be tried later, as well.
- ** HOW 4.1 [i.e., HOW was RULE162 used?]
- [5.0] We have already established that:
 - [5.1] the site of the culture is blood, and
 - [5.2] the portal of entry of the organism is GI, and
 - [5.3] the patient is a compromised host
- Therefore : it is definite (1.0) that bacteroides is an organism for which therapy should cover (RULE162)

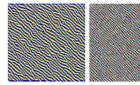
- ** DO YOU EVER PRESCRIBE CARBENICILLIN FOR PSEUDOMONAS INFECTIONS?
- RELEVANT RULES: RULE064 RULE119 RULE120 WHICH ONES DO YOU WISH TO SEE?
 ** RULE064 IF: 1) IT IS KNOWN WITH CERTAINTY THAT THE IDENTITY OF THE ORGANISM IS PSEUDOMONAS, AND
 2) THE DRUG UNDER CONSIDERATION IS GENTAMICIN
- THEN: RECORD THE FOLLOWING AS A MORE APPROPRIATE THERAPY: GENTAMICIN-AND-CARBENICILLIN

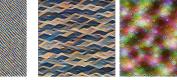
Local vs. Global interpretability

- Global interpretability understand model as a whole
 - Will it work prospectively as intended?
 (dataset shift, label misspecification, label leakage)
 - What data was most useful? (find more signal of a similar type, form causal hypotheses, figure out how to simplify for deployment purposes)

Local vs. Global interpretability

- Global interpretability understand model as a \bullet whole
 - For **any model**: do feature ablation. How does performance on held-out data change?
 - Ex. **linear models**: look at largest positive and negative weight features
 - Ex: decision trees: look at the top few splits
 - Ex. **deep models**: visualize specific filters













Edges (layer conv2d0)

Textures (layer mixed3a)

Patterns (layer mixed4a)

Parts (layers mixed4b & mixed4c) Objects (layers mixed4d & mixed4e)

Olah et al., Feature Visualization: How neural networks build up their understanding of images, Distill 2017 https://distill.pub/2017/feature-visualization/

Increasingly more difficult as models become more complex...

Predicting Psychiatric Readmission with Bag-of-Words NLP model

- 470 of patient cohort (4687) were readmitted within 30 days for psychiatric diagnoses
 - 2977 readmitted with a nonpsychiatric diagnosis
 - 1240 not readmitted (only 26%)
- We built a model using demographics, comorbidity + 75 topics from LDA on notes
 - Top 1000 TF/IDF words from each patient's notes
 - Considerable overlap, but total vocabulary ~66K words
 - SVM models to predict readmission
 - Baseline features
 - Baseline + top-1000 bag-of-words (really 66K)
 - Baseline + 75 topics

Table 3. Comparison of models with topics	and witho	out inclusio	n of LDA
Configuration	AUC	Sensitivity	Specificity
Baseline = age/gender/insurance/ Charlson	0.618	0.979	0.104
Baseline+top-1 words	0.654	—	_
Baseline+top-10 words	0.676	—	
Baseline+top-100 words	0.682	—	_
Baseline+top-1000 words	0.682	0.213	0.945
Baseline+75 topics (no words)	0.784	0.752	0.634
Abbreviations: AUC, area under the curve; LDA, Latent Dirichlet Allocation.			

Example topics for MDD patients readmitted with a psychiatric diagnosis within 30 days

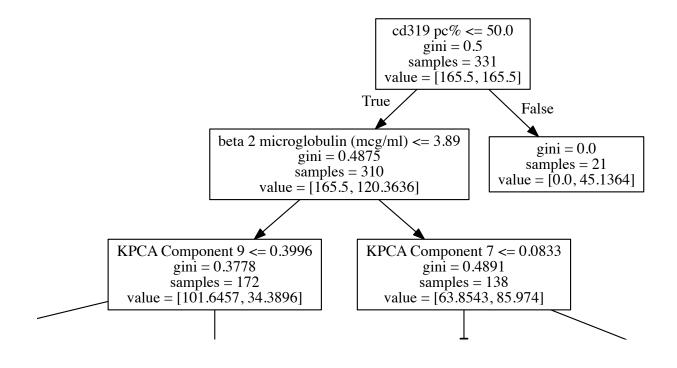
Terms	Topic annotation
*patient <i>alcohol withdrawal</i> depression <i>drinking</i> end ativan <i>etoh drinks</i> medications clinic inpatient diagnosis days hospital <i><substance name="" program="" treatment="" use=""> use abuse</substance></i> problem number	<mark>Alcohol</mark>
*mg daily discharge <i>anxiety klonopin seroquel clonazepam</i> admission wellbutrin given md lexapro date b signed night low admitted sustained hospitalization	<mark>Anxiety</mark>
* <i>ideation suicidal mood decreased</i> hallucinations history <i>depressed depression thought</i> psychiatric energy denied sleep auditory appetite homicidal symptoms increased speech <i>thoughts</i>	Suicidality
* <i>ect depression</i> treatment treatments dr mg course <i><ect name="" physician=""></ect></i> symptoms received medications prior improved decreased medication md trials tsh continued qhs	<mark>ECT</mark>
* <i>weight eating</i> admission discharge hospital intake <i>loss</i> date hospitalization day dr week physical months <i>prozac food</i> increased md did <i>anorexia</i>	Anorexia
* <i>seizure seizures</i> intact <i>eeg neurology</i> normal <i>temporal dilantin</i> head <i>bilaterally events activity</i> weakness sensation disorder tongue <i>neurologist brain</i> loss <i>tegretol</i>	Seizure
* <i>therapist mother</i> program <i>father</i> disorder age school parents brother <i>abuse</i> treatment <i>relationship</i> outpatient college behavior partial plan currently <i>group personality</i>	Psychotherapy
*psychiatry <i>suicide overdose attempt transferred</i> depression <i>transfer</i> level <i>tylenol</i> hospital service <i>unit</i> normal floor screen <i>tox</i> room admission medical general	Overdose
*baby delivery bleeding vaginal breast feeding cesarean weight ibuprofen maternal newborn available p fever pregnancy sex estimated danger gp	Postpartum
* <i>psychotic thought</i> features <i>paranoid psychosis paranoia</i> symptoms psychiatric dose continued treatment mental cognitive memory <i>risperidone</i> people th somewhat interview affect	Psychosis

 In 2018, Sontag's group submitted a paper using the Multiple Myeloma Research Foundation's IA9 data release. *Great results*

Table 3: Predicting Mortality

Method	1 Yr Full	1 Yr ISS-FISH	2 Yr Full	2 Yr ISS-FISH
LR	0.66 ± 0.1	0.62 ± 0.14	0.8 ± 0.08	0.69 ± 0.1
LR-B-PCA	0.66 ± 0.1	0.61 ± 0.13	0.79 ± 0.08	0.65 ± 0.11
LR-T-PCA	0.68 ± 0.1	0.61 ± 0.14	0.8 ± 0.08	0.65 ± 0.11
\mathbf{RF}	0.65 ± 0.09	0.63 ± 0.12	0.82 ± 0.08	0.73 ± 0.09
RF-B-PCA	0.69 ± 0.11	0.63 ± 0.12	0.83 ± 0.08	0.73 ± 0.09
RF-T-PCA	0.72 ± 0.1	0.64 ± 0.12	0.85 ± 0.08	0.72 ± 0.09

 Curious to see why "full" feature set with random forests so much better, so looked at one decision tree:



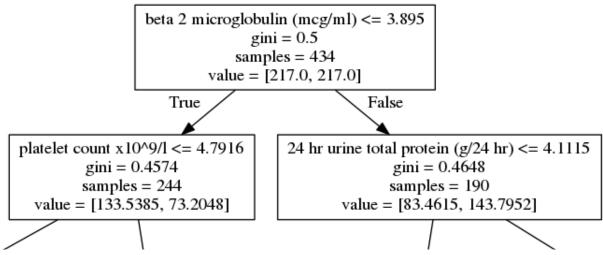
 Surprised to see cd319% at the top, but after discussing with clinical collaborator, concluded it is reasonable

 3 months later, new release of data (IA11) is available and I ask students to reproduce results Big differences!

	Method	1 Yr Full	1 Yr ISS-FISH	2 Yr Full	2 Yr ISS-FISH
	LR	0.66 ± 0.1	0.62 ± 0.14	0.8 ± 0.08	0.69 ± 0.1
Old results	LR-B-PCA	0.66 ± 0.1	0.61 ± 0.13	0.79 ± 0.08	0.65 ± 0.11
(IA9):	LR-T-PCA	0.68 ± 0.1	0.61 ± 0.14	0.8 ± 0.08	0.65 ± 0.11
(173).	\mathbf{RF}	0.65 ± 0.09	0.63 ± 0.12	0.82 ± 0.08	0.73 ± 0.09
	RF-B-PCA	0.69 ± 0.11	0.63 ± 0.12	0.83 ± 0.08	0.73 ± 0.09
	RF-T-PCA	0.72 ± 0.1	0.64 ± 0.12	0.85 ± 0.08	0.72 ± 0.09
	Models	1 Yr Full	1 Yr ISS-FISH	2 Yr Full	2 Yr ISS-FISH
	LR	0.68 ± 0.09	0.65 ± 0.14	0.76 ± 0.08	0.7 ± 0.09
New results	LR-B-PCA	0.68 ± 0.1	0.65 ± 0.13	0.75 ± 0.08	0.67 ± 0.09
	LR-T-PCA	0.69 ± 0.09	0.64 ± 0.13	0.77 ± 0.07	0.66 ± 0.09
IA11):	\mathbf{RF}	0.63 ± 0.1	0.63 ± 0.11	0.75 ± 0.08	0.73 ± 0.08
	RF-B-PCA	0.66 ± 0.1	0.64 ± 0.11	0.76 ± 0.08	0.72 ± 0.08
	RF-T-PCA	0.78 ± 0.08	0.64 ± 0.11	0.77 ± 0.08	0.72 ± 0.08

Ν

 3 months later, new release of data (IA11) is available and I ask students to reproduce results



- Cd319% no longer shows up as a top predictor!
- What happened!?
- After digging deeper, they realized that what was predictive originally was the feature Cd319% being missing, and moreover that this was correlated with the outcome (i.e. label leakage!)

What are other ways to learn models that have "good" global interpretability?

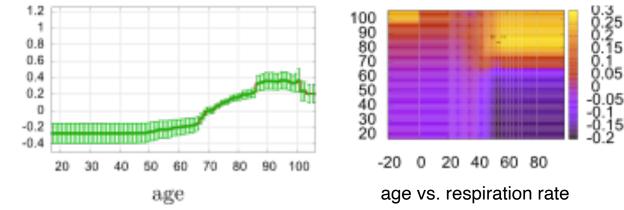
Generalized additive models (GAMs)

• GAMs with pairwise interactions have the form:

$$g(E[y]) = \beta_0 + \sum_j f_j(x_j) + \sum_{i \neq j} f_{ij}(x_i, x_j)$$

g is the link function (e.g. logistic, for binary data), and
 E[f] = 0.

Model	Pneumonia	Readmission
Logistic Regression	0.8432	0.7523
GAM	0.8542	0.7795
$GA^{2}M$	0.8576	0.7833
Random Forests	0.8460	0.7671
LogitBoost	0.8493	0.7835



[Caruana et al., KDD '15]

Falling rule lists

- Ordered list of if-then rules where:
 - 1. It is a decision list, i.e. order matters
 - 2. Probability of outcome decreases monotonically

	Conditions		Probability	Support
IF	IrregularShape AND Age ≥ 60	THEN malignancy risk is	85.22%	230
ELSE IF	Spiculated Margin AND Age ≥ 45	THEN malignancy risk is	78.13%	64
ELSE IF	IllDefinedMargin AND Age ≥ 60	THEN malignancy risk is	69.23%	39
ELSE IF	IrregularShape	THEN malignancy risk is	63.40%	153
ELSE IF	LobularShape AND Density ≥ 2	THEN malignancy risk is	39.68%	63
ELSE IF	RoundShape AND Age ≥ 60	THEN malignancy risk is	26.09%	46
ELSE		THEN malignancy risk is	10.38%	366

Table 1: Falling rule list for mammographic mass dataset.

Empirical Test: 30-Day Hospital Readmission

- 8,000 patients
- Features: "impaired mental status," "difficult behavior," "chronic pain," "feels unsafe" and over 30 other features
- Mined rules with support $\geq 5\%$, no more than two conditions
- Expected length of decision list = 8
- Compared to SVM, Random Forest, Logistic Regression, CART, Inductive Logic Programming

Method	Mean AUROC (STD)
FRL	.80 (.02)
NF_FRL	.75(.02)
NF_GRD	.75 (.02)
RF	.79 $(.03)$
SVM	.62 (.06)
Logreg	.82 (.02)
Cart	.52 (.01)

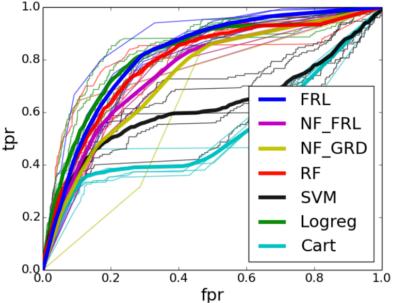


Figure 2: ROC curves for readmissions prediction. 22

Readmission Rule List

	Conditions		Probability	Support
IF	BedSores AND Noshow	THEN readmissions risk is:	33.25%	770
ELSE IF	PoorPrognosis AND MaxCare	THEN readmissions risk is:	28.42%	278
ELSE IF	PoorCondition AND Noshow	THEN readmissions risk is:	24.63%	337
ELSE IF	BedSores	THEN readmissions risk is:	19.81%	308
ELSE IF	NegativeIdeation AND Noshow	THEN readmissions risk is:	18.21%	291
ELSE IF	MaxCare	THEN readmissions risk is:	13.84%	477
ELSE IF	Noshow	THEN readmissions risk is:	6.00%	1127
ELSE IF	MoodProblems	THEN readmissions risk is:	4.45%	1325
ELSE		Readmissions risk is:	0.88%	3031

Table 2: Falling rule list for patients with no multiple readmissions history.

Supersparse linear integer models

- Learn linear model where:
 - 1. Coefficients are all integer
 - 2. As sparse as possible (training objective):

$$\min_{\lambda} \frac{1}{N} \sum_{i=1}^{N} \mathbf{1}[y_i \lambda^T x_i \le 0] + C_0 \|\lambda\|_0 + \epsilon \|\lambda\|_1$$

PREDICT PATIENT HAS OBSTRUCTIVE SLEEP APNEA IF SCORE > 1

1.	$age \ge 60$	4 points		
2.	hypertension	4 points	+	••••
3.	body mass index ≥ 30	$2 {\rm points}$	+	••••
4.	body mass index ≥ 40	$2 {\rm points}$	+	••••
5.	female	-6 points	+	••••
	ADD POINTS FROM ROWS 1 – 5	SCORE	=	••••

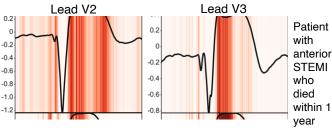
[Ustun & Rudin, ML '16]

Local vs. Global interpretability

- Local interpretability understand predictions for individual data points (i.e., patients)
 - Build trust in predictions; recognize errors due to model being poor, data point being an outlier, or engineering problems
 - Provide guidance to decision makers who may have additional information
 - *Explanations* that we described earlier, for Mycin, are an example of this

Local vs. Global interpretability

- Local interpretability understand predictions for individual data points (i.e., patients)
 - Ex: linear (bag of words) models: look at highest weighted non-zero feature
 - Ex: decision trees: look at path to prediction for this patient
 - Ex: deep models: saliency maps and GradCAM

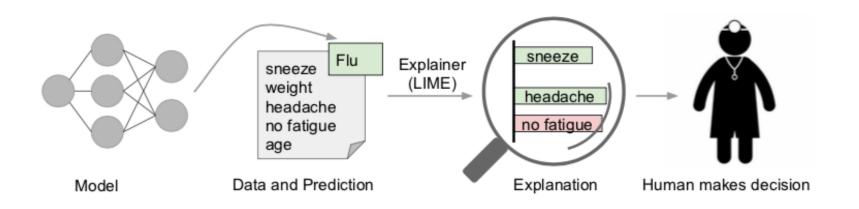


Gradient-CAM (Selvaraju et al., IJCV '19)

[Raghunath et al., Prediction of mortality from 12-lead electrocardiogram voltage data using a deep neural network, Nature Medicine 2020]

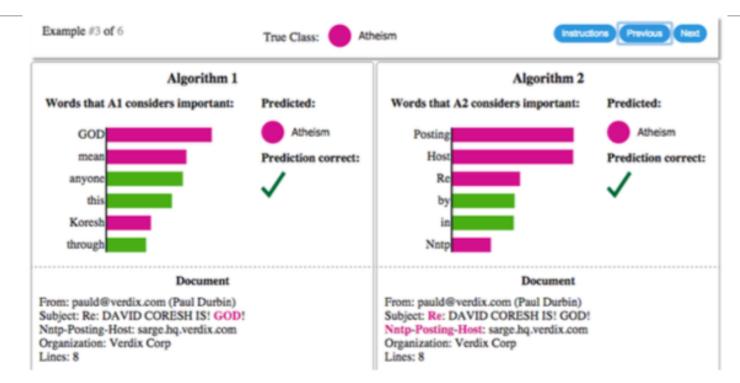
- How can we do this more generally?

Model-agnostic Explanations



- A model predicts that a patient has the flu, and LIME highlights:
 - Sneeze and headache are portrayed as contributing to the "flu" prediction
 - "no fatigue" is evidence against it.
- With these, a doctor can make an informed decision about whether to trust the model's prediction.
- Approach helps detect data leakage, data set shift, using human expertise

Explanation of Cases May be Useful to Compare Models



- · Predict whether a post is about "Christianity" or "Atheism"
- Algorithm 2 may be overall more accurate, but Algorithm 1 makes more sense, at least on this example.
- Again, relies on human expertise, which is much broader than any of our models

Desiderata for Explanations

- Interpretable "provide qualitative understanding between the input variables and the response"
 - depends on audience
 - requires sparsity
 - features must make sense
 - e.g., eigenvectors in principal component analysis are not explainable features
- Local fidelity "it must correspond to how the model behaves in the vicinity of the instance being predicted"
- Model-agnostic "treat the original model as a black box"
 - Is this really a good idea for all models?

Sparse Linear Explanation

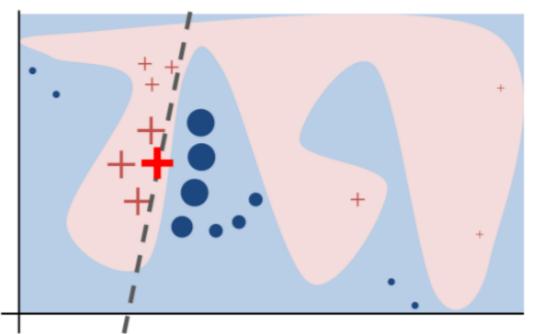
- Choose G to be the class of linear models such that $g(z') = w_g \cdot z'$

Algorithm 1 Sparse Linear Explanations using LIME

Require: Classifier f, Number of samples N **Require:** Instance x, and its interpretable version x' **Require:** Similarity kernel π_x , Length of explanation K $\mathcal{Z} \leftarrow \{\}$ for $i \in \{1, 2, 3, ..., N\}$ do $z'_i \leftarrow sample_around(x')$ $\mathcal{Z} \leftarrow \mathcal{Z} \cup \langle z'_i, f(z_i), \pi_x(z_i) \rangle$ end for $w \leftarrow \text{K-Lasso}(\mathcal{Z}, K) \triangleright$ with z'_i as features, f(z) as target return w

- Let $\pi_x(z) = \exp(-D(x,z)^2/\sigma^2)$ be an exponential kernel on some distance function *D* with width σ
 - E.g., cosine distance for bag-of-words, L2 distance or DICE for images

 $\mathcal{L}(f, g, \pi_x) = \sum_{z, z' \in \mathcal{Z}} \pi_x(z) (f(z) - g(z'))^2$



Toy example to present intuition for LIME. The black-box model's complex decision function f (unknown to LIME) is represented by the blue/pink background, which cannot be approximated well by a linear model. The bold red cross is the instance being explained. LIME samples instances, gets predictions using f, and weighs them by the proximity to the instance being explained (represented here by size). The dashed line is the learned explanation that is locally (but not globally) faithful.

How to Make Interpretable Models

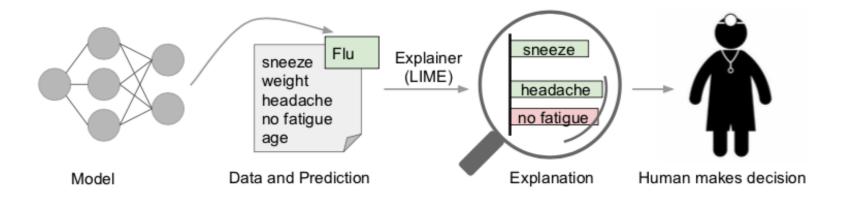
- If the original data are $x \in \mathbb{R}^d$, define a new set of variables, $x' \in \{0,1\}^{d'}$ that can serve as the interpretable representation of the data
- An *explanation* is a model $g \in G$ where G is the class of interpretable models
 - E.g., linear models, additive scores, decision trees, falling rule lists, ...
 - The domain of g is $\{0,1\}^{d'}$, i.e., the interpretable representation of the data
- The complexity of a model is $\Omega(g)$
 - E.g., depth of a decision tree, number of non-zero weights in a linear model
- The full model is $f:\mathbb{R}^d\to\mathbb{R}$
 - E.g., for classification, *f* is probability that *x* belongs to a certain class
- $\pi_x(z)$ is a proximity measure of how close z is to x, thus defining a locality around x
- Let $\mathcal{L}(f, g, \pi_x)$ be a measure of how *un*faithful g is to f in the locality defined by π_x
- Then

$$\xi(x) = \operatorname{arg\,min}_{g \in G} \mathcal{L}(f, g, \pi_x) + \Omega(g)$$

is the best explanatory model for *x* given our choices for $\{\mathcal{L}, \pi_x, \Omega\}$

Apply to Text Classification

- Bag of words, cosine distance for π_x
- · Choose K as a limit on the number of words in an explanation



Apply to Image Interpretation

- Superpixel is a group of connected pixels with similar colors or gray levels
 - Image is segmented into super pixels
 - K is chosen as the number of superpixels to represent
- K-LASSO predicts label from superpixels, to select which K of them to use for explanation
- with N=5000, scikit-learn random forests with 1000 trees \Rightarrow 3 sec
- explaining Inception network results \Rightarrow ~10 min



(a) Original Image (b) Explaining *Electric guitar* (c) Explaining *Acoustic guitar* (d) Explaining *Labrador*

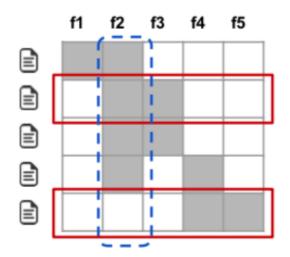
Figure 4: Explaining an image classification prediction made by Google's Inception neural network. The top 3 classes predicted are "Electric Guitar" (p = 0.32), "Acoustic guitar" (p = 0.24) and "Labrador" (p = 0.21)

Choosing a Suite of Examples to Explain

- Choose a diverse, comprehensive set of B examples to explain
 - WHY?
- Given explanations for a set of instances X(|X| = n), consider the $n \times d'$ explanation matrix \mathcal{W} whose rows are examples and columns are features
 - Each entry gives the local importance of that feature \overline{for}^{u} that example
 - For linear models, for instance $x_i, g_i = \xi(x_i)$, set $\mathcal{W}_{ij} = |w_{g_{ij}}|$
 - recall that $g(z') = w_g \cdot z'$
 - I_i is a measure of *global* importance of that feature

•
$$I_j = \sqrt{\sum_{i=1}^n \mathcal{W}_{ij}}$$
 for text

more difficult for superpixels because they don't recur over different instances



LIME Experiments

- Two sentiment analysis datasets (2000 instances, each; used 1600/400 test/train)
- Bag-of-words as features
- Models:
 - Decision Trees
 - Logistic Regression with L2 regularization
 - Nearest Neighbors
 - Support Vector Machines with RBF kernels
 - Random Forest (1000 trees) with word2vec embeddings
- *K* = 10

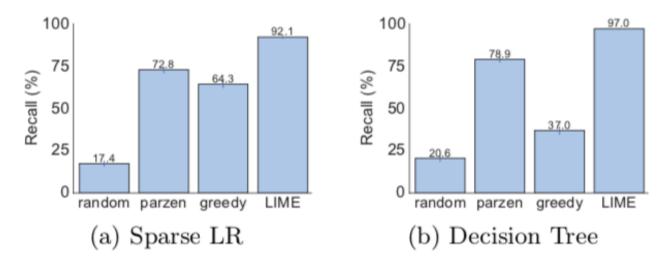


Figure 6: Recall on truly important features for two interpretable classifiers on the books dataset.

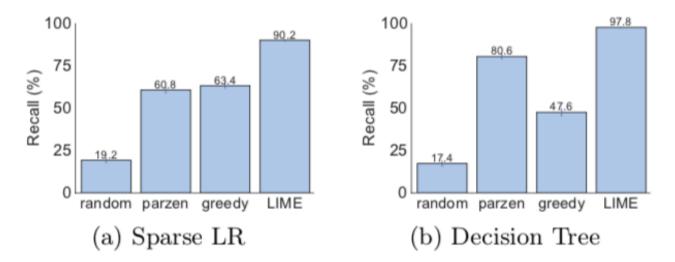


Figure 7: Recall on truly important features for two interpretable classifiers on the DVDs dataset.

Human Experiments

- Questions:
 - · Can users choose which of two classifiers generalizes better
 - Based on the explanations, can users perform feature engineering to improve the model
 - Are users able to identify and describe classifier irregularities by looking at explanations
- "Christianity" vs. "Atheism" from 20-newsgroups dataset
 - known problems of data leakage from headers, ...
 - trained original and "cleaned" classifiers for comparison
 - test set accuracy favors the "wrong" classifier!!!
- Separate test set of 819 web pages about these topics from http://dmoz-odp.org
- SVM with RBF kernels, trained on the 20-newsgroup data
- Mechanical Turk, 100 users, *K*=6 words, *B*=6 documents/Turk
 - in 2nd experiment, they are asked to remove word features they believe inappropriate

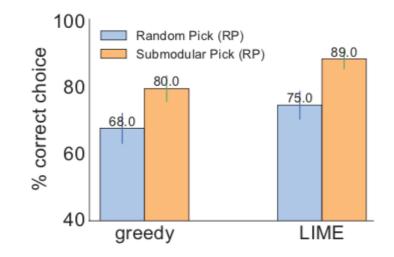


Figure 9: Average accuracy of human subject (with standard errors) in choosing between two classifiers.

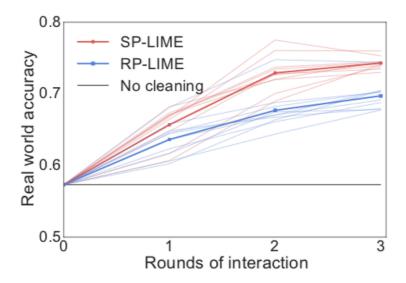
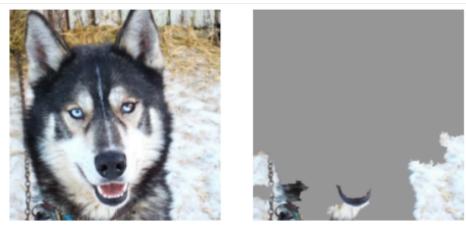


Figure 10: Feature engineering experiment. Each shaded line represents the average accuracy of subjects in a path starting from one of the initial 10 subjects. Each solid line represents the average across all paths per round of interaction.

Can People Gain Insight from these Explanations?

- Trained a deliberately bad classifier between Wolf and Husky
 - All wolves in training set had snow in the picture, no huskies did
- Presented cases to graduate students with ML background
 - 10 balanced test predictions, with one husky in snow, one wolf not in snow
- Comparison between pre- and postexperiment trust and understanding



(a) Husky classified as wolf

(b) Explanation

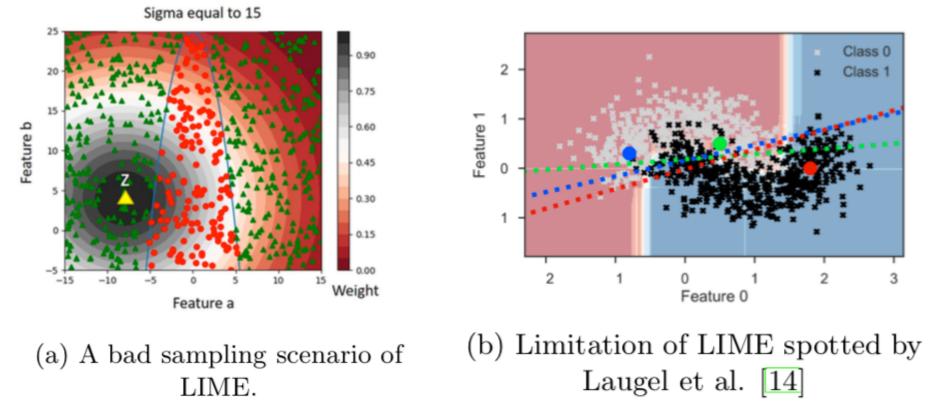
Figure 11: Raw data and explanation of a bad model's prediction in the "Husky vs Wolf" task.

fore After	
	t of 27 $3 \text{ out of } 2$ t of 27 $25 \text{ out of } 2$

Table 2: "Husky vs Wolf" experiment results.

Critique of LIME

- Choice of σ is arbitrary and can lead to bad sampling
 - in implementation, often set to $0.75\sqrt{d}$
- it is important to tune the size of the neighbourhood according to how far z is to the closest decision boundary



Counterfactual explanations

- Why did the treatment not work on the patient?
- Why was my loan rejected?
- Simplest approach:
 - Find the *smallest* change to the features that would change the prediction from rejected to approved
 - Note: (a) there may be many, (b) should be *realistic*

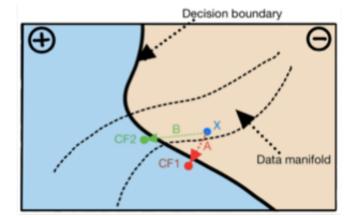


Figure 1: Two possible paths for a datapoint (shown in blue), originally classified in the negative class, to cross the decision boundary. The end points of both the paths (shown in red and green) are valid counterfactuals for the original point. Note that the red path is the shortest, whereas the green path adheres closely to the manifold of the training data, but is longer.

[Figure from: Verma et al., Counterfactual Explanations for Machine Learning: A Review, arXiv:2010.10596, 2020]

See also:

Karimi, Scholkopf, Valera. Algorithmic Recourse: from Counterfactual Explanations to Interventions. FAccT '21

Can Attention Models in Deep Learning Serve as Explanations?

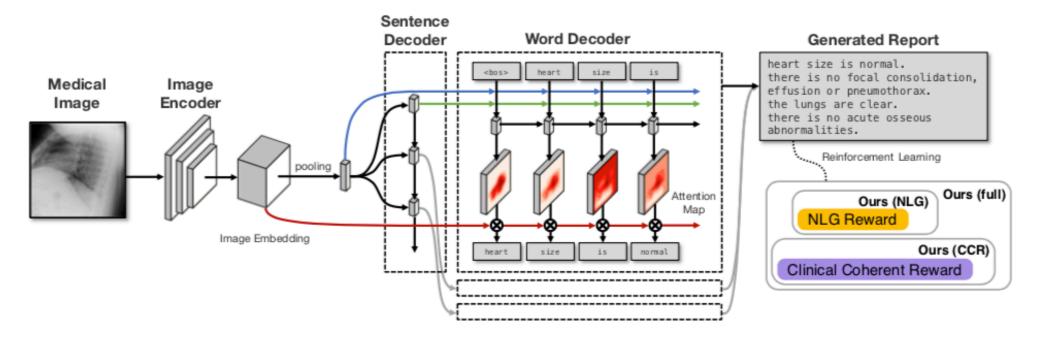
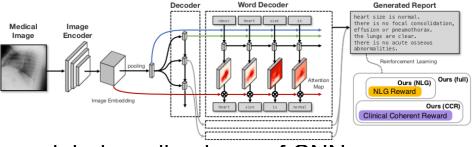


Figure 2: The model for our proposed *Clinically Coherent Reward*. Images are first encoded into image embedding maps, and a sentence decoder takes the pooled embedding to recurrently generate topics for sentences. The word decoder then generates the sequence from the topic with attention on the original images. NLG reward, clinically coherent reward, or combined, can then be applied as the reward for reinforcement policy learning.

- Image encoder (CNN)
 - Spacial image features $V = \{v\}_{k=1}^{K}$
 - computed by fully connected layer on pre-global-pooling layer of CNN
- Sentence decoder (RNN/LSTM) uses image features
 - $h_i, m_i = \text{LSTM}(\bar{v}; h_{i-1}, m_{i-1})$
 - topic vector and stop signal $\boldsymbol{\tau}_i = \operatorname{ReLU}(\boldsymbol{W}_{\tau}^T \boldsymbol{h}_i + \boldsymbol{b}_{\tau}), \ u_i = \sigma(\boldsymbol{w}_u^T \boldsymbol{h}_i + b_u)$
- Word decoder (RNN/LSTM)
 - Uses \bar{v} , $\tau\!\!$, and embedding of previous word generated
 - Word is sampled from either conditional probability or overall corpus probability
- Reinforcement learning to favor most readable and clinically correct output
 - Use CheXpert annotations for 12 diagnoses: pos, neg, uncertain, absent
- Hack: remove duplicate generated sentences



Sentence

Ground Truth



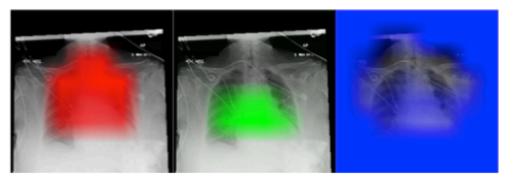
cardiomegaly is moderate. bibasilar atelectasis is mild. there is no pneumothorax. a lower cervical spinal fusion is partially visualized. healed right rib fractures are incidentally noted.

TieNet

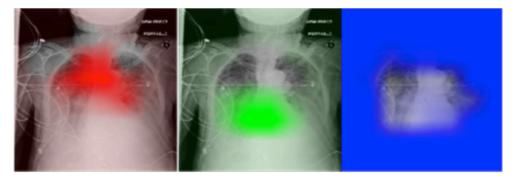
ap portable upright view of the chest. there is no focal consolidation, effusion, or pneumothorax. the cardiomediastinal silhouette is normal. imaged osseous structures are intact. Ours (full)

pa and lateral views of the chest. there is mild enlargement of the cardiac silhouette. there is no pleural effusion or pneumothorax. there is no acute osseous abnormalities.

Attention Map Identified Relevant Parts of the Image



ap upright and lateral views of the <u>chest</u>. there is moderate <u>cardiomegaly</u>. there is no pleural <u>effusion</u> or pneumothorax. there is no acute osseous abnormalities.



as compared to the previous radiograph, there is no relevant change. <u>tracheostomy</u> tube is in place. there is a layering pleural effusions. NAME bilateral pleural effusion and compressive <u>atelectasis</u> at the right base. there is no <u>pneumothorax</u>.

(a)

(b)

Figure 3: Visualization of the generated report and image attention maps. Different words are underlined with its corresponding attention map shown in the same color.

Attention is not Explanation

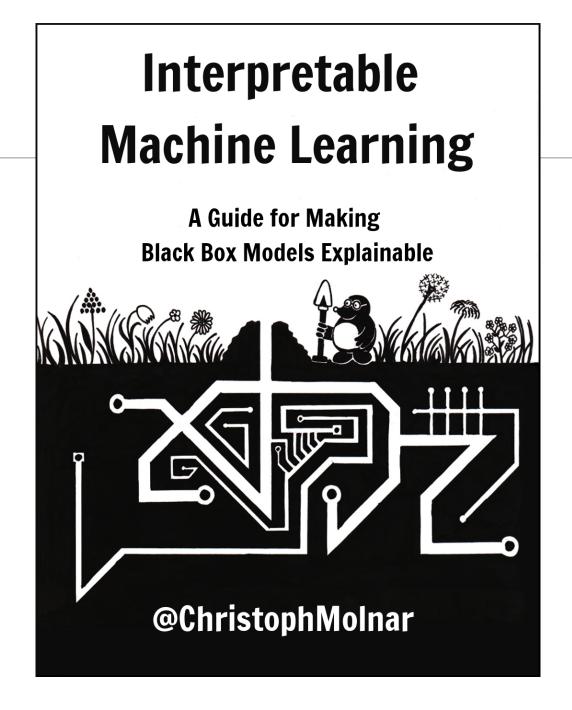
But

Sarthak Jain Northeastern University jain.sar@husky.neu.edu Byron C. Wallace Northeastern University b.wallace@northeastern.edu

- "assumption that the input units (e.g., words) accorded high attention weights are responsible for model outputs"
- Desiderata if attention actually is to give insight into how a DNN operates
 - Attention weights should correlate with feature importance measures (e.g., gradient-based measures)
 - Alternative (or counterfactual) attention weight configurations ought to yield corresponding changes in prediction
- Mixed results, though the study has been criticized for methodology
 - "evidence that correlation between intuitive feature importance measures (including gradient and feature erasure approaches) and learned attention weights is weak"
 - counterfactual attention distributions which would tell a different story about why a model made the prediction that it did — often have no effect on model output

Achieving Interpretability for Humans

- Why: Incompleteness in problem formalization
 - Scientific understanding
 - Safety
 - Ethics
 - Indirect objectives
 - Competing objectives
- How: Methods
 - Application-grounded; in the context of its end-task
 - Compare to value of human-generated explanation to help other people
 - Human-grounded; simplified tasks
 - Choose better explanation; predict model outcome based on inputs and explanation; counterfactual (what input must change to change output)
 - Functionally-grounded; formal definition of interpretability
 - Posit certain classes of models to be interpretable; e.g., decision lists



Also, see work by faculty here in Boston....

Hima Lakkaraju (Harvard)

Finale Doshi (Harvard)

Manish Raghavan (MIT)

Byron Wallace (Northeastern)

https://christophm.github.io/interpretable-ml-book/