



GRACE: Generating Concise and Informative Contrastive Sample to Explain Neural Network Model's Prediction

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- ① ✓ Introduction
- ② Motivation & Challenges
- ③ Contrastive Explanation
- ④ GRACE Algorithm
- ⑤ Experiments
- ⑥ Conclusion

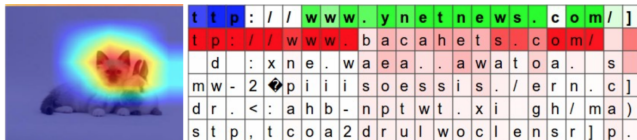
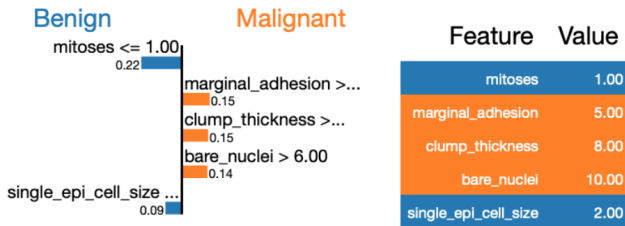


Figure: Example of Highlighting Spans of Words/Phrases. Source: Google Image



Text: "if bare_nuclei is less than or equal 6.0, on average, this prediction would be 0.14 less Malignant. etc.,"

Figure: Example of Lime [Ribeiro et al., 2016] Method on Tabular Data Instance



1. **Tabular data** is prominent in many important fields.
2. Explanation for tabular data: high-dimensional inter-correlated features
 - Which key features to select?
 - If top K, what if features are highly correlated? (E.g. "Frequency of !" and "Frequency of !!!")
3. How to present the explanation to the end-users?
 - Highlighting a batch of an image, a span of words in a sentence?
4. **End-users** might have different interests than researchers, developers
5. **GOAL: Develop an algorithm to explain neural network models' predictions on tabular datasets to end-users**



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Feature	freq_now	freq_credit	freq_!!!	freq_!	class
\mathbf{x}_1	0.1	0.0	0.0	0.0	Ham
$\tilde{\mathbf{x}}_1$	0.1	0.0	0.3	0.453	Spam

Feature	freq_you	freq_direct	avg_longest_capital	class
\mathbf{x}_2	0.68	0.34	158.0	Spam
$\tilde{\mathbf{x}}_2$	0.68	0.34	1.0	Ham

Table: Examples of original samples \mathbf{x}_i and contrastive samples $\tilde{\mathbf{x}}_i$ on *spam* dataset.)



1. End-users are interested in explanation: "Why X rather than Y?"

Feature	freq_you	freq_direct	avg_longest_capital	class
x	0.68	0.34	158.0	Spam
\tilde{x}	0.68	0.34	1.0	Ham

Examples

Explanation *"Had the message had no words written in all capital letters, it would have been classified as **ham** rather than spam."*



Problem Statement

Given \mathbf{x} and neural network model $f(\cdot)$, our goal is to generate new contrastive sample $\tilde{\mathbf{x}}$ to provide concise and informative explanation for the prediction $f(\mathbf{x})$.



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Table: Examples of original samples \mathbf{x}_i and contrastive samples $\tilde{\mathbf{x}}_i$

1. Constraint on the contrastive class:

$$\operatorname{argmax}(f(\tilde{\mathbf{x}})) \neq \operatorname{argmax}(f(\mathbf{x})) \quad (1)$$

2. Constraint on the # of key features:

$$|\mathcal{S}| \leq K \quad (2)$$

3. Constraint on the mutual information:

$$\operatorname{SU}(\mathcal{X}^i, \mathcal{X}^j) \leq \gamma \quad \forall i, j \in \mathcal{S} \quad (3)$$

4. Constraint on the domain:

$$\tilde{\mathbf{x}} \in \operatorname{dom}(\mathcal{X}) \quad (4)$$



Objective Function

Given \mathbf{x} , hyperparameter K, γ , our goal is to generate new contrastive sample $\tilde{\mathbf{x}}$ to explain the prediction $f(\mathbf{x})$ by solving the objective function:

$$\begin{aligned} \min_{\tilde{\mathbf{x}}} \quad & \text{dist}(\tilde{\mathbf{x}}, \mathbf{x}) \\ \text{s.t.} \quad & \text{argmax}(f(\mathbf{x})) \neq \text{argmax}(f(\tilde{\mathbf{x}})), \quad |\mathcal{S}| \leq K \\ & \text{SU}(\mathcal{X}^i, \mathcal{X}^j) \leq \gamma \quad \forall i, j \in \mathcal{S}, \quad \tilde{\mathbf{x}} \in \text{dom}(\mathcal{X}) \end{aligned} \quad (5)$$

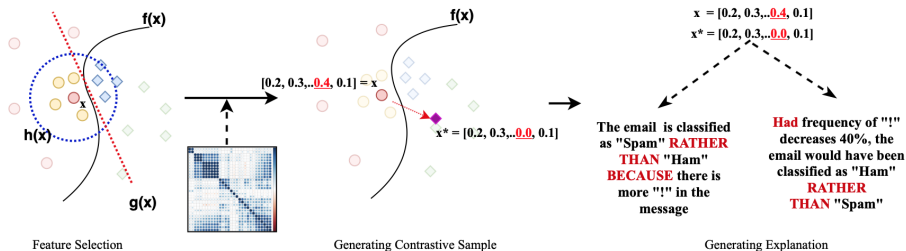


Figure: Grace Algorithm - Local-Based Feature Selection

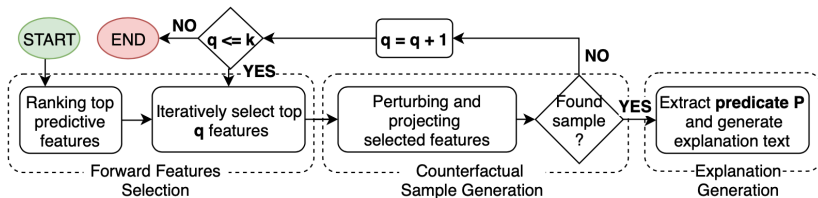


Figure: Grace Algorithm



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Table: Dataset statistics and prediction performance

Dataset	#Class	#Feat.	#Data	Acc.*	F1*
eegeye	2	14	14980	0.858	0.858
diabetes	2	8	768	0.779	0.777
cancer95	2	9	699	0.963	0.963
phoneme	2	5	5404	0.774	0.772
segment	7	19	2310	0.836	0.817
magic	2	10	19020	0.862	0.859
biodeg	2	41	1055	0.853	0.851
spam	2	57	4601	0.932	0.932
cancer92	2	30	569	0.958	0.958
mfeat	10	216	2000	0.943	0.936
musk	2	166	476	0.783	0.789

(*) Accuracy and F1 scores are averaged across 10 different runs.

1. **NearestCT**: Select nearest contrastive sample from the training set
2. **DeepFool** [Moosavi-Dezfooli et al., 2016]: Generate adversarial samples with $\min_{\tilde{x}} \|\tilde{x} - \mathbf{x}\|_2$
3. **Lime** [Ribeiro et al., 2016]: Instance-based explanation method



1. Conciseness:

$$\mathbf{R}_{\text{fidelity}} = \frac{1}{|\tilde{\mathcal{X}}|} \sum_{(\tilde{\mathbf{x}}, \tilde{\mathbf{y}}) \in \tilde{\mathcal{X}}} \mathbb{1}(\tilde{\mathbf{y}} == \operatorname{argmax}(f(\tilde{\mathbf{x}}))) \quad (6)$$

$$\mathbf{R}_{\text{avg\#Feats}} = \frac{1}{|\tilde{\mathcal{X}}|} \sum_{\tilde{\mathbf{x}} \in \tilde{\mathcal{X}}} |\mathcal{S}_{\tilde{\mathbf{x}}}| \quad (7)$$

2. Info-Gain:

$$\mathbf{R}_{\text{info-gain}} = 1 - \frac{1}{|\tilde{\mathcal{X}}|} \sum_{\tilde{\mathbf{x}} \in \tilde{\mathcal{X}}} \sum_{i \in \mathcal{S}_{\tilde{\mathbf{x}}}} \sum_{j \in \mathcal{S}_{\tilde{\mathbf{x}}}} \frac{\text{SU}(\mathcal{X}^i, \mathcal{X}^j)}{|\mathcal{S}_{\tilde{\mathbf{x}}}|^2} \quad (8)$$

3. Influence:

$$\mathbf{R}_{\text{domain}} = \frac{1}{|\tilde{\mathcal{X}}|} \sum_{\tilde{\mathbf{x}} \in \tilde{\mathcal{X}}} \mathbb{1}(\tilde{\mathbf{x}} \in \operatorname{dom}(\mathcal{X})) \quad (9)$$

$$\mathbf{R}_{\text{influence}} = \frac{\mathbf{R}_{\text{fidelity}} \times \mathbf{R}_{\text{info-gain}} \times \mathbf{R}_{\text{domain}}}{\mathbf{R}_{\text{avg\#Feats}}} \quad (10)$$



		biodeg	spam	cancer92	mfeat	musk
$R_{\text{avg}\#Feats}$	NearestCT	20.53	17.50	29.97	204.22	147.86
	DeepFool	41.00	57.00	30.00	216.00	166.00
	GRACE-Local	<u>3.07</u>	<u>2.95</u>	3.95	<u>3.28</u>	<u>3.74</u>
	GRACE-Gradient	1.93	1.09	<u>4.5</u>	2.76	2.85
$R_{\text{info-gain}}^*$	NearestCT	0.44	<u>0.62</u>	0.02	<u>0.58</u>	0.28
	DeepFool	<u>0.58</u>	0.53	0.01	0.59	0.29
	GRACE-Local	0.46	0.47	0.13	0.34	<u>0.3</u>
	GRACE-Gradient	0.76	0.95	<u>0.04</u>	0.50	0.4
$R_{\text{influence}}$	NearestCT	0.02	0.04	0.00	0.00	0.00
	DeepFool	0.01	0.01	0.00	0.00	0.00
	GRACE-Local	<u>0.15</u>	<u>0.16</u>	0.04	<u>0.1</u>	<u>0.08</u>
	GRACE-Gradient	0.4	0.88	<u>0.01</u>	0.18	0.14

Table: All results are averaged across 10 different runs. The best and second best results are highlighted in **bold** and underline.

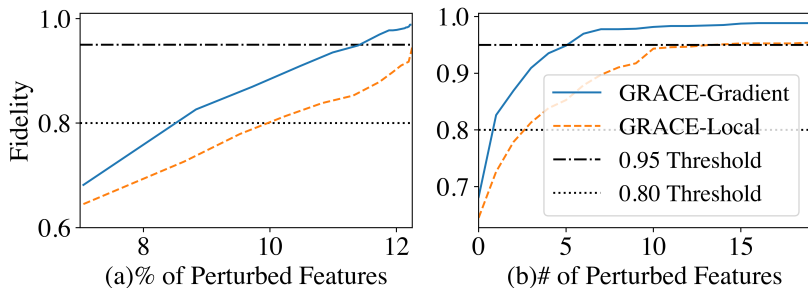


Figure: Percentage of Perturbed Features v.s. Fidelity



1. Comparison between Grace and Lime
 - Hypothesis \mathcal{H}_1 : **more intuitive and friendly**
 - Hypothesis \mathcal{H}_2 : **more comprehensible**
 - Hypothesis \mathcal{H}_3 : **leads to better post-explanation decisions**
2. Recruit Amazon MTurk workers as general end-users
3. No assumptions on workers' prior knowledge in machine learning



Feature	Bare_nuclei	MarAdh	CluThic	mitoses	CelSizUni	CelShaUni	NorNuc	SinEpCeSi	Model Prediction
Value	10.0	5.0	8.0	1.0	8.0	5.0	3.0	2.0	Malignant

Explanation

*"Had **bare_nuclei** been 3.0 point lower and **CluThic** been 7.0 point lower, the patient would have been diagnosed as Benign rather than Malignant"*

Q1: Given a scale from 1 to 10, "how intuitive and friendly is the explanation to you?" (1 is least preferable, 10 is most preferable)

0 0

Q2: Given a scale from 1 to 10, "how understandable is the explanation to you?" (1 is least preferable, 10 is most preferable)

0 0

Figure: Example of an User-study Task for \mathcal{H}_1 , \mathcal{H}_2



Explanation

*"If **Bare_nuclei** is 3.0 point lower and **CluThic** is 7.0 point lower, while keeping other features the same, the patient would be diagnosed as Benign rather than Malignant"*

Q2: Below is the current value for each features of **PATIENT 1**. Following the **explanation** displayed, please **ADJUST** (increase, decrease, do not change) these values such that the computer model will change the prediction for this patient to **BENIGN**

Bare_nuclei:



BlaChr:



Figure: Example of an User-study Task for \mathcal{H}_3

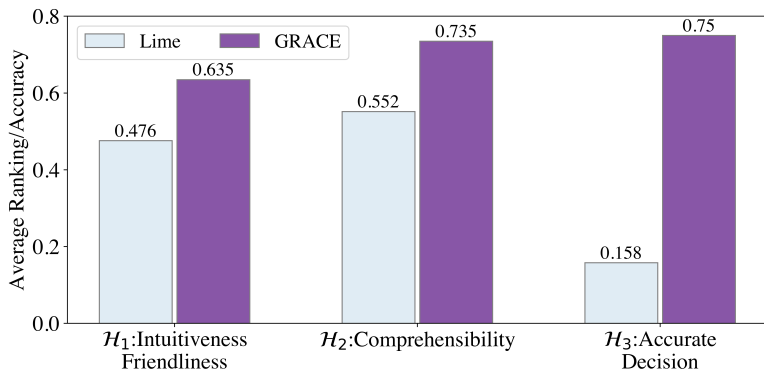


Figure: Comparison of generated explanation: GRACE v.s. Lime. Scores are normalized to $[0,1]$. All results are statistically significant ($\mathcal{H}_1 : p\text{-value} < 0.05$, $\mathcal{H}_2, \mathcal{H}_3 : p\text{-value} < 0.01$)

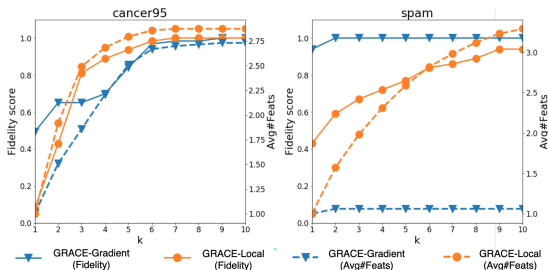


Figure: Sensitivity of K on Fidelity

Table: Effects of entropy threshold γ on Info-Gain

Dataset	Method	1.0	0.7	0.5	0.3
musk	GRACE-Gradient	0.51	0.51	0.58	0.58
	GRACE-Local	0.36	0.36	0.54	0.54
segment	GRACE-Gradient	0.57	0.57	0.59	0.59
	GRACE-Local	0.79	0.79	0.84	0.84



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1. **GRACE**: A novel instance-based algorithm that provides end-users with simple natural text explaining neural network models' predictions in a contrastive "*Why X rather than Y*" fashion.
2. **GRACE**: more intuitive, friendly, comprehensible and leads to more accurate decisions than Lime



1. Source Code and Slides:
https://github.com/lethaiq/GRACE_KDD20
2. Pike Group at Penn State:
<http://pike.psu.edu>



- [Moosavi-Dezfooli et al., 2016] Moosavi-Dezfooli, S.-M., Fawzi, A., and Frossard, P. (2016).
Deepfool: a simple and accurate method to fool deep neural networks.
In *Proceedings of the 2016 IEEE CVPR*, pages 2574–2582.
- [Ribeiro et al., 2016] Ribeiro, M. T., Singh, S., and Guestrin, C. (2016).
"why should I trust you?": Explaining the predictions of any classifier.
In *KDD*, pages 1135–1144.