

IPAM Workshop Explainable AI for the Sciences: Towards Novel Insights

# **Towards Higher-Order & Disentangled XAI**

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**Part 1: Motivations** 

P Keyl, M Bockmayr, D Heim, G Dernbach, G Montavon, KR Müller, F Klauschen Patient-level proteomic network prediction by explainable artificial intelligence NPJ Precis Oncol. 6(1):35, 2022

## **Example: Discovering Influencial Proteins**





#### Question:

Can we use ML/XAI to infer these networks (or aspects of them) directly from the data?

## Finding Influential Proteins with ML/XAI (Keyl et al. 2022)

### Step 1: From Data to ML

- Assemble a dataset.
- Build a ML model (neural network) that predicts proteins from other proteins with best possible accuracy.



## Finding Influential Proteins with ML/XAI (Keyl et al. 2022)

#### Step 2: From ML to XAI

Apply Explainable AI (here the LRP attribution technique) to identify to what extent proteins contribute to the expression of other proteins.



# Finding Influential Proteins with ML/XAI (Keyl et al. 2022)





- Generally consistent with existing knowledge, e.g. highlights mTOR pathway; correlates with entries in the reactome knowledgebase (https://reactome.org/).
- Provides cancer-specific (or even instance-specific) view of protein influences.



# Beyond 'Classical' Explainable AI



- Current explainable AI already provides single-instance nonlinear explanation capabilities that exceed by far classical statistical measures such as correlation.
- There is a potential demand for even more detailed explanations (e.g. *joint* features contributions, or latent concepts underlying features contributions).

### Part 2: Towards Higher-Order Explainable AI

T Schnake, O Eberle, J Lederer, S Nakajima, K T. Schütt, KR Müller, G Montavon Higher-Order Explanations of Graph Neural Networks via Relevant Walks IEEE TPAMI 44(11):7581-7596, 2022



#### **Observation:**

▶ Input of a GNN is not at layer one, but occurs (multiplicatively) at each layer.

## Limits of 'Classical' Attributions



Choice between first-order and higherorder is determined by the *model* rather than by the user.

GNN prediction (simplified):

$$\begin{split} h_j &= \rho \big( \sum_i \mathbf{1}_i \Lambda_{ij} w_j \big) & (\text{layer 1}) \\ h_k &= \rho \big( \sum_j h_j \Lambda_{jk} w_k \big) & (\text{layer 2}) \\ y &= \sum_k h_k & (\text{layer 3}) \end{split}$$



**Our approach:** computing  $R_{ijk}$  iteratively:

$$\begin{split} R_{jk} &= \mathcal{E}(y, \Lambda_{jk}) & (\text{step 1}) \\ R_{ijk} &= \mathcal{E}(R_{jk}, \Lambda_{ij}) & (\text{step 2}) \end{split}$$

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**Property:** For  $\rho$  linear, the iterative attribution produces the same result as identifying the summands in the expanded form:

$$y = \sum_{ijk} \underbrace{\Lambda_{ij} \Lambda_{jk} 1_i w_j w_k}_{R_{ijk}}$$



Model	Aggregate	Combine	GNN-LRP Rule
GCN [36]	$m{Z}_t = m{\Lambda} m{H}_{t-1}$	$m{H}_t =  ho(m{Z}_tm{W}_t)$	$R^{a}_{JKL\dots} = \sum_{b} \frac{\lambda_{JK} h^{a}_{J} w^{\dagger}_{ab}}{\sum_{J,a} \lambda_{JK} h^{a}_{J} w^{\dagger}_{ab}} R^{b}_{KL\dots} $ (11)
GIN [44]	$Z_t = \Lambda H_{t-1}$	$\boldsymbol{H}_t = (\mathrm{MLP}^{(t)}(\boldsymbol{Z}_{t,K}))_K$	$R^{a}_{JKL\dots} = \sum_{b} \frac{\lambda_{JK} h^{a}_{J}}{\sum_{J} \lambda_{JK} h^{a}_{J}} \text{LRP}(R^{b}_{KL\dots}, z^{a}_{K})  (12)$
Spectral [43], [45] (case $\lambda \ge 0$ )	$oldsymbol{Z}_{s,t} = oldsymbol{\Lambda}_s H_{t-1}$	$H_t =  ho(\sum_s {oldsymbol{Z}}_{s,t} {oldsymbol{W}}_{s,t})$	$R^{a}_{JKL} = \sum_{b} \frac{\sum_{s} \lambda^{s}_{JK} h^{a}_{J} w^{s\dagger}_{ab}}{\sum_{J,a} \sum_{s} \lambda^{s}_{JK} h^{a}_{J} w^{s\dagger}_{ab}} R^{b}_{KL} $ (13)

#### GNN-LRP at work:



#### Note:

- ▶ In vanilla form, GNN-LRP requires an LRP pass for each walk in the graph ( $\rightarrow$  expensive).
- Coarse-graining of the input graph can reduce computations.

## Evaluating Higher-Order Explanations (Schnake et al. 2022)

#### **Observation:**

XAI evaluation techniques such as 'Pixel-Flipping' require as input a sequence of features (e.g. nodes) from most to least relevant. However, Higher-Order XAI attributes to joint features.



#### Idea:

▶ From the given explanation, generalize relevance scores to subset of features S:

$$R_{\mathcal{S}} = \sum_{i \in \mathcal{S}} R_i \quad \text{(first-order XAI)} \qquad \qquad R_{\mathcal{S}} = \sum_{(ijk) \subseteq \mathcal{S}} R_{ijk} \quad \text{(higher-order XAI)}$$

Ask the explanation to produce an optimal sequence of nodes:

$$\mathcal{Q} = \operatorname*{argmax}_{\mathcal{S}_1 \subset \dots \subset \mathcal{S}_d} \left\{ \sum_{i=1}^d R_{\mathcal{S}_i} \right\}$$

▶ Finding Q is intractable ⇒ approximate it with greedy feature selection or randomization.

## Evaluating Higher-Order Explanations (Schnake et al. 2022)



#### **Results:**

- ► GNN-LRP achieves better performance than first-order explanations (LRP and GNNExpl).
- GNN-LRP is more robust than its simpler gradient-based counter part GNN-GI.

## Use Case: XAI for Quantum Chemistry

Decomposing molecular properties (predicted via a GNN) in terms of atom interactions of different order.



#### **Challenges:**

- $\blacktriangleright$  Larger explanations  $\rightarrow$  more difficult to comprehend for a human.
- General comment about XAI: Need to make a distinction between the strategy employed by the model to predict (dataset-specific) and the underlying physics (general).

### **Part 3: Towards Disentangled Explanations**

P Chormai, J Herrmann, KR Müller, G Montavon Disentangled Explanations of Neural Network Predictions by Finding Relevant Subspaces arXiv:2212.14855, 2022

## Limits of 'Classical' Explanations



### **Observation:**

 Several concepts (ball, player, etc.) are entangled in the same explanation.

## Limits of 'Classical' Explanations



#### **Observation:**

Several concepts (ball, player, etc.) are entangled in the same explanation.

#### Question:

Can we disentangle explanations into multiple distinct concepts so that they become more actionable?



### Disentangled Explanations (Chormai et al. 2022)



Forward pass:

 $egin{array}{ccc} m{x}\mapsto (h_k)_k & & ({
m input to subspaces}) \ (h_k)_k\mapsto y & & ({
m subspaces to output}) \end{array}$ 

Standard explanation

$$R_i = \mathcal{E}(y, x_i)$$

Disentangled explanation (ours):

$$\begin{split} R_k &= \mathcal{E}(y,h_k) & \text{(step 1)} \\ R_{ik} &= \mathcal{E}(R_k,x_i) & \text{(step 2)} \end{split}$$

## Extracting Relevant Subspaces (Chormai et al. 2022)

#### Notation:

a	Vector of activations
R	Vector of activation relevances
c	Vector such that $oldsymbol{R} = oldsymbol{a} \odot oldsymbol{c}$
$(U_k)_k$	Matrices that project activations to orthogonal subspaces.)



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### Key findings:

1. For a variety of methods (e.g. integrated gradients, LRP), the relevance score for subspace k can be expressed as:

$$R_k = (U_k^\top \boldsymbol{a})^\top (U_k^\top \boldsymbol{c})$$

2. We can find subspaces that *directly* maximize some statistic of  $R_k$ .







If setting  $c \leftarrow a$ , PRCA reduces to (uncentered) PCA.



$$\underset{U}{\text{maximize}}: \quad \overbrace{\mathsf{Tr}(U^{\top} \underbrace{\mathbb{E}[\boldsymbol{a}\boldsymbol{c}^{\top}]}_{\boldsymbol{\Sigma}_{\boldsymbol{a}\boldsymbol{c}}} U)}^{R}$$

If setting  $c \leftarrow a$ , PRCA reduces to (uncentered) PCA.

Disentangled Relevant Subspace Analysis (DRSA)

$$\underset{(U_k)_k}{\text{maximize}: } \mathbb{M}_k^{0.5} \mathbb{M}_n^2 \big\{ \big( \overbrace{(U_k^\top \boldsymbol{a}_n)^\top (U_k^\top \boldsymbol{c}_n)}^{R_{kn}} \big)^+ \big\}$$

If setting  $c \leftarrow a$  DRSA reduces to 'DSA'.





# PRCA/DRSA in Practice (Chormai et al. 2022)



### PRCA vs. Baselines (Chormai et al. 2022)



PRCA extracts much more strongly contributing subspaces than baseline methods.

	VGG16-TV	VGG16-ND	NFNet-F0	VGG16-TV	VGG16-ND
	+ LRP	+ LRP	+ LRP	+ Shapley	+ Shapley
Total ( $\sum_i R_{i,I}$ )	11.47	10.35	6.57	17.23	16.59
Random Subspace	0.02	0.00	0.01	0.02	0.02
Most Relevant FM [22]*	0.97	0.87	0.30	1.12	0.91
PCA	1.81	2.69	-2.22	21.81	18.98
PRCA (Ours)	<b>13.63</b>	<b>13.72</b>	<b>11.29</b>	<b>44.76</b>	<b>42.12</b>
Error bars (max)	$\pm 0.66$	$\pm 0.61$	$\pm 0.58$	± 1.78	$\pm$ 1.40

### DRSA vs. Baselines (Chormai et al. 2022)

Question: are the components of the explanation spatially disentangled?



#### Separability score:



## Use Case: Detecting and Removing Clever Hanses

#### **Current approaches:**

 Artifact models built from preliminarily detected Clever Hans instances.

### Our approach:

- Observe that Clever Hans strategies readily occur in distincts components of DRSA.
- 2. Identify these

components, and remove their contribution from the overall prediction.



Carton Training Samples with Their Standard and DRSA Subspace Heatmaps

## Use Case: Exploring Visual Relations between Classes



- Certain visual concepts are shared between classes (e.g. dotted pattern of 'admiral' and 'monarch' butterflies).
- ▶ This can be analyzed dataset-wide in a scatter plot (left).

## Summary

- Explanations should not only be faithful/understandable; they should also be informative & actionable by the user.
- This can be achieved by:
  - Ensuring the explanation reflects the use of **higher-order** feature interactions by the model (e.g. GNN-LRP).
  - Resolving the latent concepts attached to each feature contribution in order to produce a **disentangled** explanation (e.g. using PRCA / DRSA).
- Both approaches (higher-order & disentangled XAI) are not mutually exclusive. They could be combined in future work.

### References to presented works

#### XAI for Analyzing Protein Interactions

P Keyl, M Bockmayr, D Heim, G Dernbach, G Montavon, KR Müller, F Klauschen Patient-level proteomic network prediction by explainable artificial intelligence NPJ Precis Oncol. 6(1):35, 2022

#### Higher-Order XAI

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#### **Disentangled XAI**

P Chormai, J Herrmann, KR Müller, G Montavon Disentangled Explanations of Neural Network Predictions by Finding Relevant Subspaces arXiv:2212.14855, 2022

## Check our review paper on XAI

W Samek, G Montavon, S Lapuschkin, C Anders, KR Müller Explaining Deep Neural Networks and Beyond: A Review of Methods and Applications Proceedings of the IEEE, 109(3):247-278, 2021



### Visit our website



- Code/demos for our XAI methods
- ► Full list of papers