# #2 Explainable AI for DMD Drug-Repurposing

Obtaining drug candidates that can treat symptoms observed in Duchenne muscular dystrophy disease

\* Indicates required question

This questionnaire is part of a Leiden University master's thesis project performed by Rosa Zwart in collaboration with the LUMC BioSemantics group. This project aims to investigate whether the explainability of explainable AI algorithms can be improved by using Foundational Ontologies.

#### Introduction

This questionnaire is meant for obtaining feedback from users with knowledge and experience in drug repurposing and/or the rare disease DMD.

A knowledge graph is created containing phenotypic and genetic information directly and indirectly associated with Duchenne muscular dystrophy (DMD). Subsequently, data have been incorporated into the graph that relate to drugs and their associated protein targets as well as information regarding the symptoms and diseases for which a drug is used during treatment. Using this knowledge graph, a deep learning model is trained which results in predictions of new links between drugs and phenotypes directly associated with DMD. Due to the black-box nature of deep learning algorithms, it is difficult for human decision-makers to assess the reliability of these predictions. To enable decision-makers to subject the predictions obtained by the model to their own reasoning, interpretable explanations are added to the link predictions of the trained graph neural network (GNN) model. By adding an explanation-generating method to the deep learning algorithm, we arrive at the field of explainable Al that already shows promising developments into the direction of applying trustworthy deep learning techniques to important medical decisions.

During this research project, we want to investigate whether the explanations generated for the predictions by the trained model can be improved. To attempt this, we tested two different approaches, and we are now running a blind assessment of the results. In this questionnaire, you got an explanation generated by one of the two approaches.

The explanations are subgraphs that explain the predicted existence of a certain link between a drug and DMD-related symptom node by showing the most impactful links that lead to this prediction.

## **Background Information**

In order to acquire an idea about the background of all participants, some questions will be asked regarding your research experience.

1.	How many years of experience do you have in the field of drug repurposing and/or rare diseases research?	<b>*</b>
2.	Do you conduct research on the rare disease Duchenne muscular dystrophy? *  Mark only one oval.	
	Yes No	

# **Explanation**

To assess the explainability of the given explanation, Likert scaled scores need to be assigned for each explainability aspect.

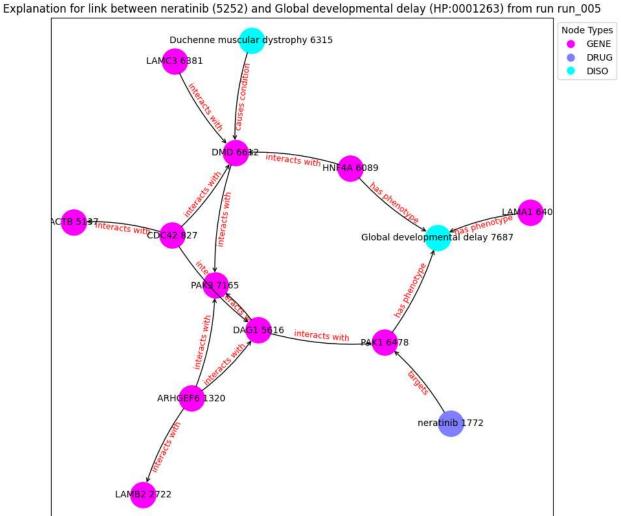
A caption is added to the subgraph to show the meaning of the most relevant paths for explaining the prediction and to provide additional information about the instances found in the graphs.

### **Prediction**

It has been predicted that neratinib treats global developmental delay.

## **Explanation (Check the graph before reading the text below)**

Neratinib targets the gene PAK1. PAK1 interacts with DAG1. DAG1 interacts with PAK3 and CDC42. Both CDC42 and PAK3 interact with DMD. DMD interact with HNF4A. Mutations in HNF4A and PAK1 can cause the phenotype global developmental delay.



#### Clarity \* 3.

I believe that the explanation is unambiguous regarding the use of concepts and relations.

Mark only one oval.

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4. Parsimony \*

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