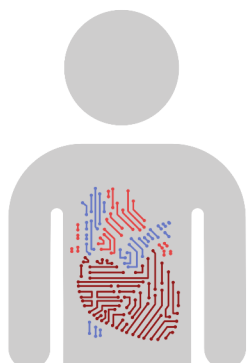


Integrative Scoring Approach

Network level interpretation



de novo
missense
variant

GENE LEVEL

Loss of Function tolerance scoring
Heart enriched expression
Mutation frequencies
Role in heart development

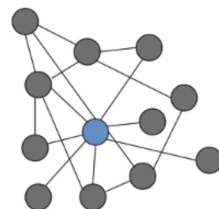
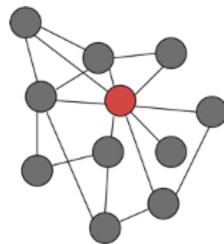
INDIVIDUAL LEVEL

Variant without other causal
mutations

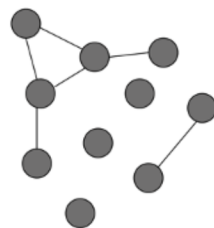
VARIANT LEVEL

Predicted pathogenicity
Variant location (domains)

Normal



Abnormal



Disrupted