

Genetic mechanism of rare neurological disorders



최 무 림

서울대학교 의과학과

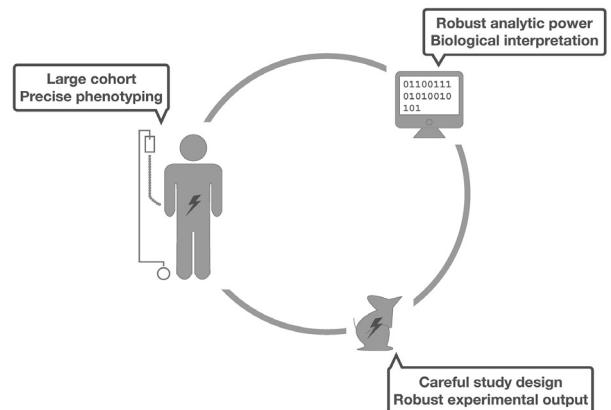
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Overview

- Neurodevelopment disease cohort ($n = 553$)
- Neuromuscular disease cohort ($n = 107$)

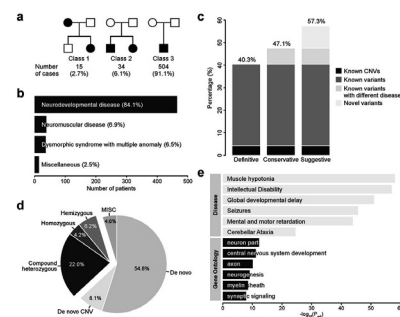
Overall approach



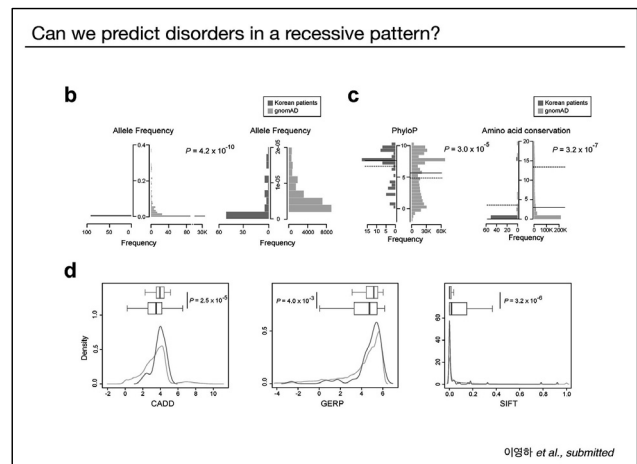
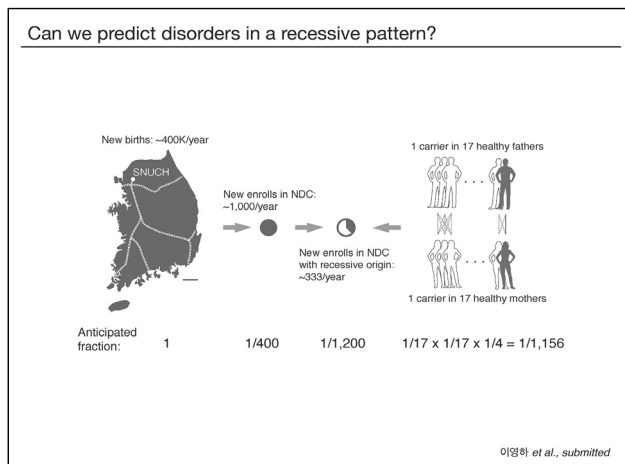
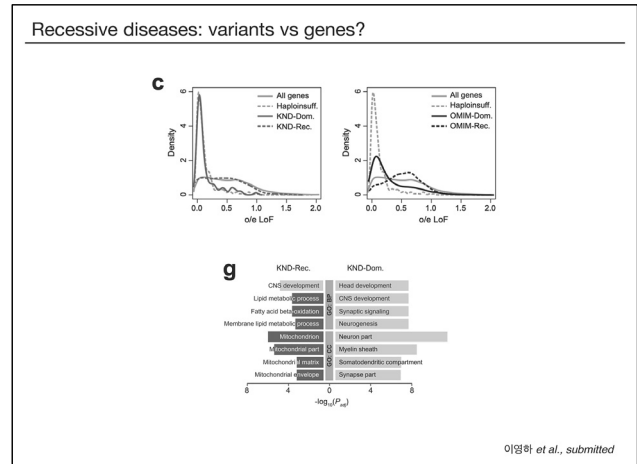
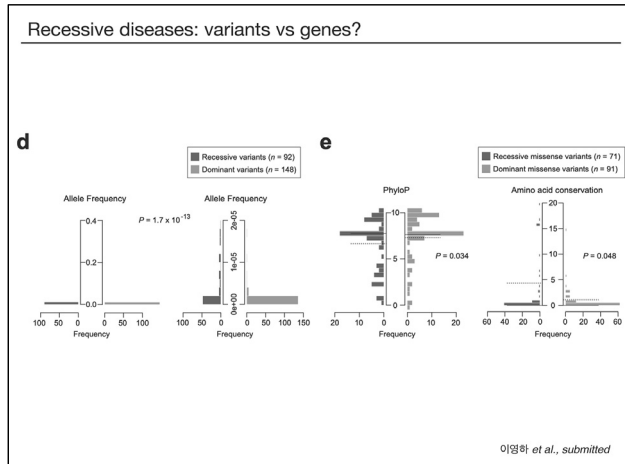
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What we've learned so far - 553 patients with ND problems



이영하 et al., submitted



Neuromuscular diseases

Jana Kneissl
(석사과정)

채종희
김수연

조안나

Study Hypothesis

- Can we increase diagnostic rate and disease mechanism by combining genome and transcriptome data?

Pros and Cons

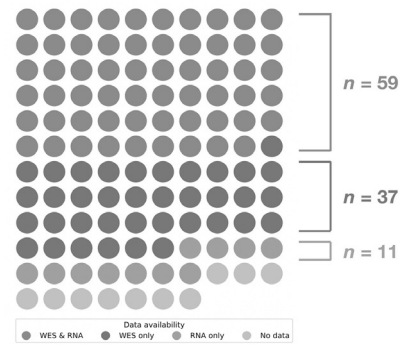


1. Tissue accessible
2. Support from histology
3. Recessive variants
4. Understudied

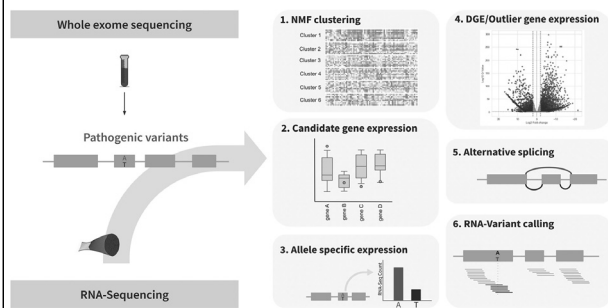


1. Rare
2. Heterogeneous
3. Structural genes
4. Understudied

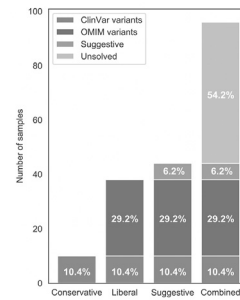
Study Overview



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Genome analysis



- 54% of the samples still do not have readily usable variants after WES

Overall summary

1. WES based rare disease analysis shed new insights into disease mechanism and variant profile
2. RNA-seq and WGS augment WES-based analysis
3. Novel genes provide new revenue for additional benefit to the researchers and patients