

P.09

Comprehensive Database for Ryanodine Receptor Type-1 Related Disorders: Concept and Progress Update

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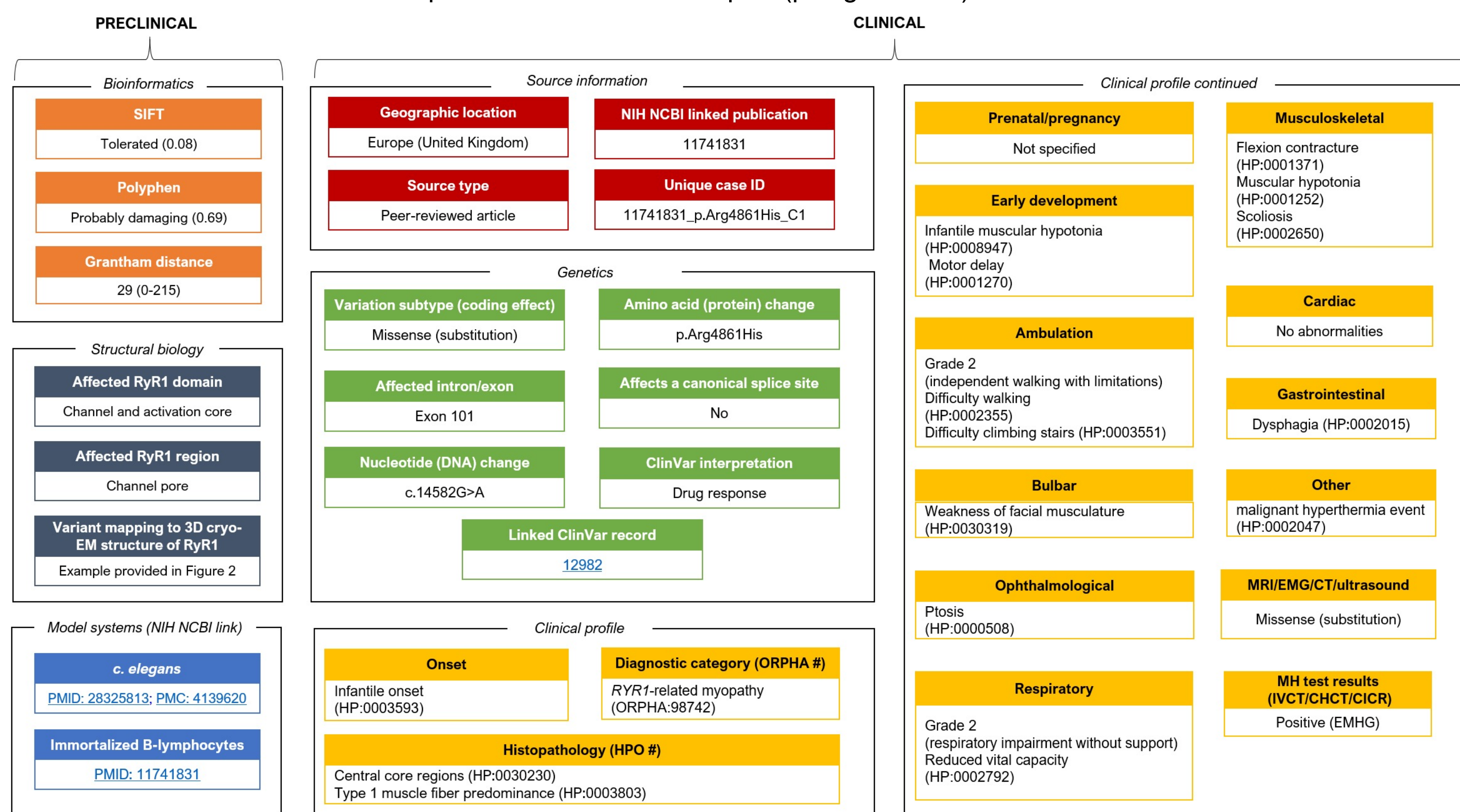
Background:

RYR1 gene encodes the largest ion channel in humans, the skeletal muscle ryanodine receptor (RyR1). RyR1 gates calcium in the sarcoplasmic reticulum and is critical to excitation-contraction coupling. Pathogenic *RYR1* variations cause a diverse spectrum of rare neuromuscular disorders including congenital and late-onset myopathies, rhabdomyolysis-myalgia syndrome, and are linked to malignant hyperthermia (MH) susceptibility. There is no approved treatment for *RYR1*-related disorders. >1000 *RYR1* variations have been reported. A majority are classified as variants of uncertain significance (VUS) for which interpretation remains challenging for researchers, clinicians, and affected individuals. We therefore developed two datasets as the foundation for a dedicated database for *RYR1*-related disorders:

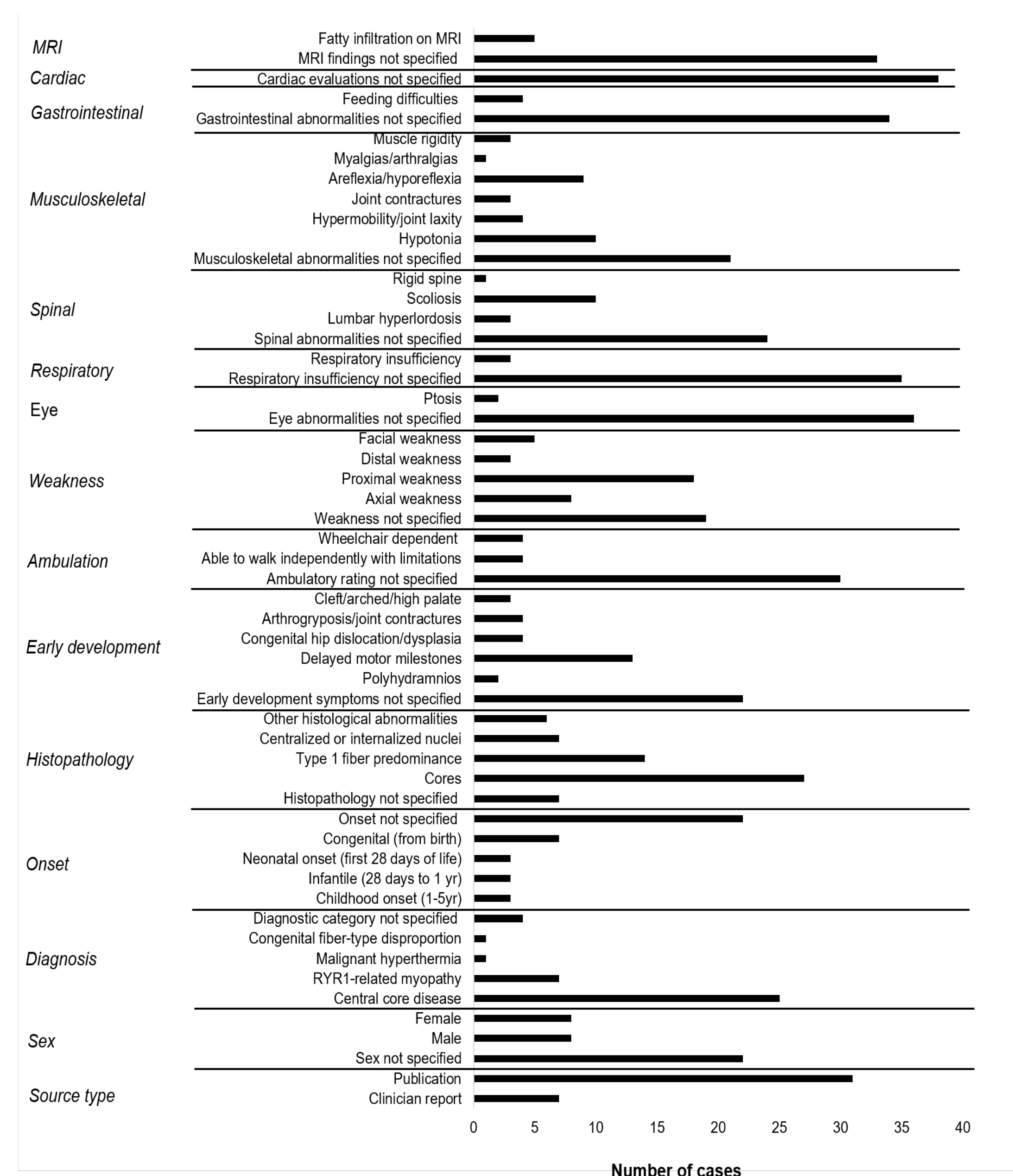
- (1) Clinical dataset: genotype-phenotype data on >2500 patients
- (2) Preclinical dataset: analyses on >200 published *RYR1* variations

Aim 1: Database Taxonomy

Example Individual Variant Report (p.Arg4861His)



Example Aggregate Clinical Report (p.Arg4861His)

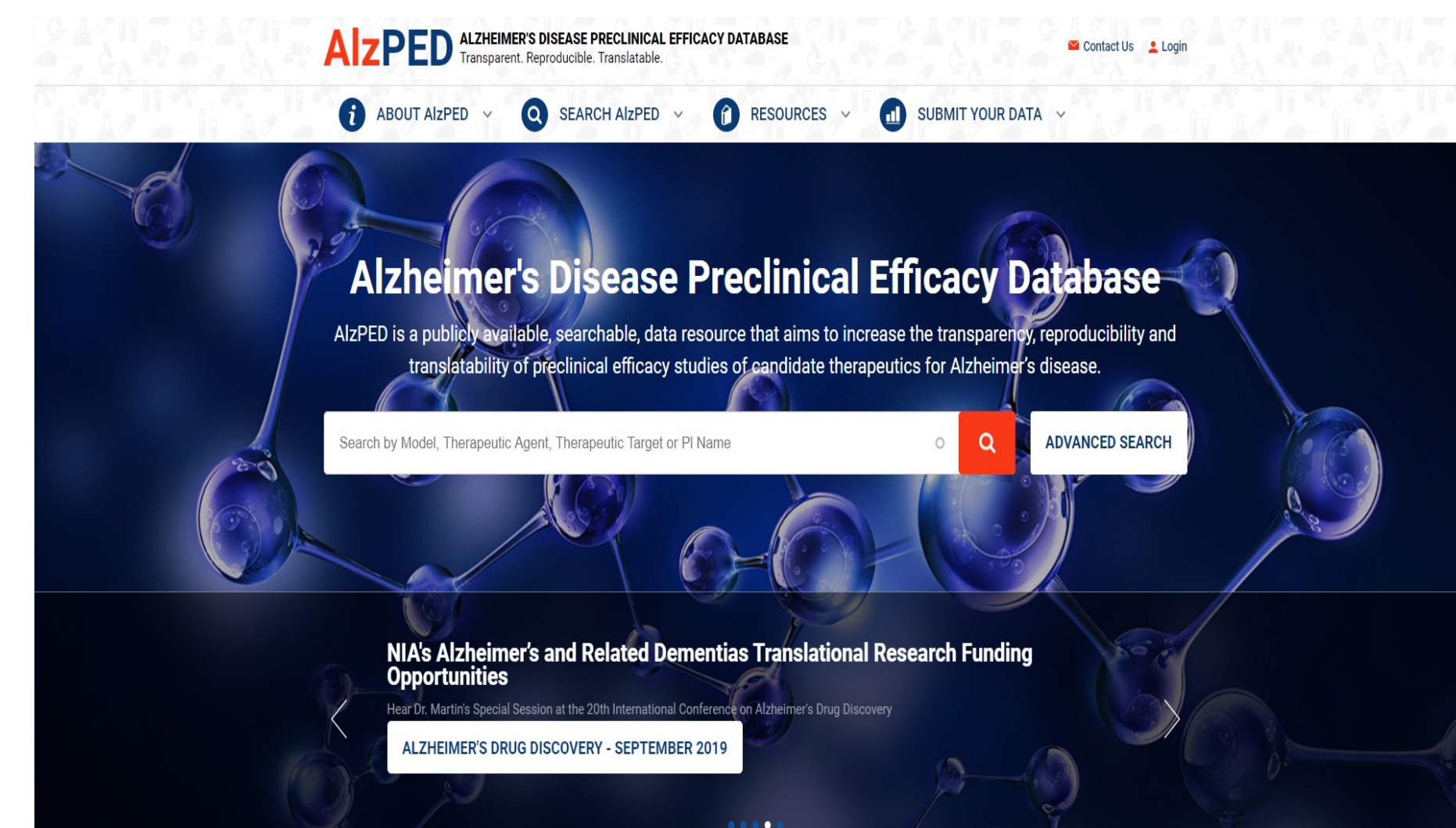


Aim 2: Engage the patient community, research scientists, and medical professionals to refine and optimize database interface

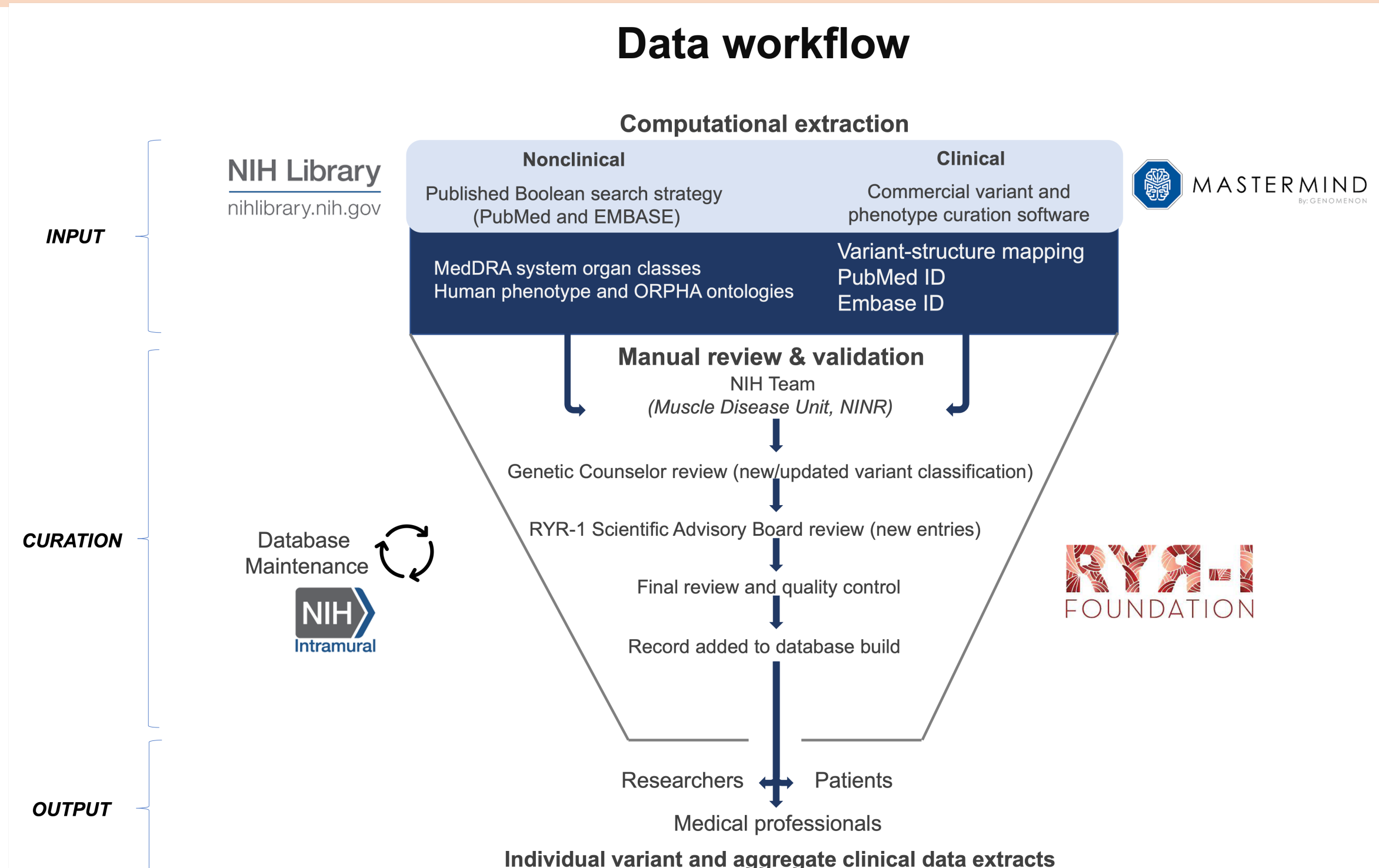
- Semi-structured survey of stakeholders (minimal risk IRB approved protocol)
- Recruitment strategies: existing patient registries (RYR-1 Foundation, Congenital Muscle Disease International Registry), patient advocacy groups and research consortia (Muscular Dystrophy Association, Treat-NMD, European Neuromuscular Centre, EMHG), and medical/research institutions

Aim 3: Launch database for *RYR1*-related disorders

- The database will be built by contracted information architects in Drupal
- Hosted on secure NIH Intramural Program cloud servers
- Interface will be based on existing NIH-led Alzheimer's disease database



Aim 4: Establish a mechanism for long-term curation and funding



Progress Update

- On-going manual review & validation by NIH team
- Structural biology team identified
- Submission of minimal risk protocol for IRB approval in 2023
- Submission of Infrastructure Grant application to the Muscular Dystrophy Association (MDA) in 2023



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