

RYR1-related myopathies, the most common non-dystrophic muscle disease subtype, are a heterogeneous group of muscle disorders united by the presence of disease-causing mutations in the skeletal muscle ryanodine receptor (*RYR1*). Affected individuals can exhibit severe disabilities that include the need for a wheelchair for ambulation, severe spine curvature requiring surgery, breathing difficulties necessitating the use of a ventilator, and in some cases, can result in premature death in childhood. Despite great advances in our understanding of *RYR1* function and the pathologic mechanisms associated with disease causing mutations in *RYR1*, there are currently no effective treatments for these devastating myopathies and there is an alarming paucity of candidate therapies in the development pipeline. We recently showed that homology-independent targeted integration (HITI) mediated CRISPR/Cas9 gene editing allows for targeted knock-in within the context of both dividing and non-dividing cells *in vitro* and *in vivo*. This approach employs a single double-strand break at a specific genomic locus induced by the CRISPR/Cas9 system, while the desired knock-in fragment flanked with inverted target sequence is cut from donor vector at the same time. Oriented integration at a specific genomic locus occurs based on non-homologous end joining (NHEJ) repair machinery. As homology-directed repair (HDR) only occurs in dividing cells, post-mitotic tissues like skeletal muscle are difficult to adapt to using HDR to repair a genomic mutation. Our central hypothesis is that HITI-mediated gene editing can be used to correct *RYR1* mutations, and thus, restore normal *RYR1* channel function and localization. Therefore, successful optimization of HITI-mediated CRISPR/Cas9 gene repair is an essential preclinical step needed in order to justify future studies designed to use this approach to treat genetically-inherited skeletal muscular diseases. **The overall goal of this proposal is to develop and validate the *in vitro* efficacy of homology-independent targeted integration (HITI) CRISPR/Cas9 gene editing to correct mutations that result in *RYR1* myopathy.**

IV. Specific Aims and Rationale

RYR1 Research Priority Area:

- Utilization of CRISPR/Cas9 as a Therapeutic Strategy for RYR-1 Myopathy and Related Diseases