Myotonic

ASSESSMENT OF MYOTONIC DYSTROPHY TYPE 2 RESEARCH AND DRUG DEVELOPMENT WITH RECOMMENDATIONS FOR INVESTMENT

PROJECT BACKGROUND AND GOALS

Myotonic commissioned the development of a landscape assessment of the status of research and drug development in myotonic dystrophy type 2 (DM2) in Spring 2019. The goal of this project was to understand current opportunities and constraints in the research and drug development pipelines in order to implement projects designed to increase academic and industry interest in, and incentivize therapy development for, DM2. The below results of this assessment have been reviewed by the Myotonic Scientific Advisory Committee for veracity and scope.

Next steps will include selecting key projects to drive research and drug development for DM2, identifying and confirming funding partners, and subsequently releasing Requests for Application to engage leading investigators to carry out high-impact, time-sensitive projects.

The application of tools to study and treat DM2 has lagged significantly behind the development and application of tools for DM1, although many of the DM1 tools, and drug development efforts are likely to be applicable to DM2 as well. The DM2 challenges besides adequate funding, include the technical difficulties involved in working with the large DM2 repeat expansions, difficulty pinpointing distinct phenotype characteristics for which to monitor therapy-driven changes, the lack of biomarkers to serve as surrogate endpoints and identify early target engagement, and an understanding of the disease itself, particularly as it relates to CNS involvement. As one researcher noted, "the headwinds against DM2 have been strong."

However, research has clearly identified the predominant disease process: a gain of function in the *CNBP* gene leading to the build-up of toxic RNA which causes the sequestration of MBNL proteins. This sequestration prevents MBNL from binding with its normal RNA targets and leads to downstream consequences of alternative splicing and the diffuse, highly variable symptomatology. The below provides a review and assessment of the status of key basic and translational research efforts and opportunities to accelerate therapy development for DM2.

CURRENT RESEARCH

Current DM2 research includes both bench and lab research and clinical studies involving affected patients.

• Bench and lab research studies:

Lab-based studies of disease/protein/genetic modifiers and molecular interactions of DM2 are currently being conducted by several international investigators:

- Repeat Associated Non-ATG (RAN) translation, RNA gain of function and protein gain of function in repeat expansion disorders including amyotrophic lateral sclerosis (ALS), spinocerebellar ataxia (SCA) types 5 and 8, myotonic dystrophy (DM) types 1 and 2 and Huntington's disease (HD). Laura Ranum, PhD, at the University of Florida is building from her work in ALS to demonstrate the role of Repeat Associated Non-ATG (RAN) translation, RNA gain of function, and protein gain of function in repeat expansion disorders including (DM) types 1 and 2.
- How do simple repeat expansions in non-coding regions result in disease? Maurice Swanson, PhD. Maury Swanson has generated a number of knock-out, knock-in and transgenic mouse models to investigate the roles of specific RNA-binding proteins in disease pathogenesis. His lab is studying an RNA-mediated disease model in which mutant DM1 and DM2 mRNAs are trapped in the nucleus and sequester (C)CUG repeat expansion binding proteins that are essential for normal tissue development and maintenance. His lab identified the sequestered factors as the muscleblind-like (MBNL) proteins. Ongoing efforts are focused on elucidating the normal functions of these proteins as well as investigating whether RNA toxicity has a pathogenic role in other hereditary disorders.
- Intron retention induced by microsatellite expansions as a disease biomarker: Maurice Swanson, PhD, along with Myotonic post-doctoral fellow Dr. Łukasz Sznajder, PhD, have developed a diagnostic assay using a person's peripheral blood to assess intron retention events as disease-specific diagnostic biomarkers for microsatellite expansions such as those found in DM1 and DM2 patients. Using DM2 as a model, they showed that CNBP i1 retention occurs even with expansion sizes in the low pathogenic range, which suggests that this assay may be informative for pre-symptomatic patients.
- Myotonic Dystrophy Population-based Prevalence Study. Myotonic-funded study led by Dr. Nicholas Johnson, MD, Virginia Commonwealth University, reporting new data regarding the population-based (as opposed to diagnosed) prevalence of both DM1 and DM2. The study results, which are expected to publish in Summer/Fall 2019, will support Myotonic efforts to make the case to industry for DM2 as a desirable drug development.

Clinical studies:

There are three clinical research studies currently underway to identify biomarkers and phenotypic changes in DM2 patients. They are longitudinal and prospective without the ability to yield outcomes and information for \sim 3 years.

 STOPP DM2. A natural history study with a target enrollment of 50 DM2 patients is being conducted at the University of Rochester. Dr. Charles Thornton, MD, Dr. Joanna Hamel, MD, Dr. Chad Heatwole MD, and others are looking at muscle biopsies, patient-reported outcomes of disease burden, and mechanistic changes over three years of observation. Currently 12 DM2 patients have completed the 3 years of observations. There are 28 additional DM2 patients enrolled past baseline; participant selection is designed to include DM2 patients with broad symptom burden, age, etc. to reduce bias in the study results. If the additional 10 patients can be enrolled within a year, cross-sectional data regarding baseline readings can initially be made available. No other outcome or publication is planned until all enrolled patients complete the 3-year observational period and the data is analyzed against the full cohort of participants.

The single-site STOPP DM2 study will then move to a multi-site, multi-year study conducted via the Myotonic Dystrophy Clinical Research Network (DMCRN), likely in the next 3 years.

- Clinical Outcome Measures in Myotonic Dystrophy Type 2 (COMEDY-2). Dr. Benedikt Schoser, MD, at Ludwig-Maximilian Universitat Munchen will be recruiting at least 60 DM2 patients to be evaluated through a battery of patientreported outcomes (PROs) and clinical outcome measures in order to define suitable measurement strategies for DM2, and to propose a disease-specific severity scale. Patients will be re-evaluated after 6 months. An age and gendermatched control cohort will be assessed.
- Identifying practical and reliable endpoints to quantitate lean tissue mass in Myotonic Dystrophy Type 1 and 2. Dr. Araya Puwanant, M.D. at Wake Forest, is using Dual Energy X-ray Absorptiometry (DEXA) scans as a simple imaging technique, to test whether DEXA measurements in affected muscles may provide a sensitive endpoint that can be bridged to meaningful outcome measures that are sensitive and appropriate to monitor disease progression and therapeutic response in the clinical trial setting of DM2. This work was started at the University of Pittsburgh and will continue at Wake Forest.
- Assessing Clinical Endpoints and Biomarkers in Myotonic Dystrophy Type-1 and Type-2 (ASCEND-DM). Dr. Ami K Mankodi, M.D., at the National Institute of Neurological Disorders and Stroke (NINDS), is using results of the Myotonic Dystrophy Clinical Research Network (DMCRN) 100-patient longitudinal study data to continue work on biomarker and endpoint development in both DM1 and DM2. Her current project will recruit 50 DM2 patients at NINDS as a single site study. It will include efforts to validate custom devices she has developed for both muscle endpoint assessment and non-muscle endpoints and biomarkers, including respiratory and central nervous system (CNS) function. She will also assess selected tests, imaging, and patient-reported outcomes for their ability to quantify disease burden, detect disease progression and predict changes. Dr. Mankodi will study DM2 disease progression over 2 years. She will also examine muscle and cerebrospinal fluid (CSF) for RNA alternative splicing events that may be used as potential biomarkers of DM2 severity, and genetic modifiers of DM2 severity by genome-wide association (GWA).

RESEARCH TOOLS

- Cell lines:
 - DM2 cell lines commissioned by Myotonic are in progress and should be ready in Fall 2019 via a partnership with Rutgers University and NINDS. The 4 DM2 patient-derived cell lines will create induced pluripotent stem cells (iPSCs) derived from erythroblasts extracted from DM2 blood samples. The goal with the new cell lines offered by Myotonic will be to offer readily available, IP-free, at-cost

tissue-derived models to industry and academia in an effort to further encourage research and development for DM2 treatments.

 DM2 cell lines published in 2019 by Dr. Denis Furling, PhD, Institut de Myologie, and Dr. Elizabeth McNally MD, PhD, Northwestern University, were produced from muscle biopsies (myoblasts) and urine-derived cells reprogrammed into myotubes, respectively. These cell lines do not show missplicing events expected in DM2 patient-derived cells. As opposed to DM1, the DM2 field remains divided on the degree to which RNA splicing changes are related to pathogenesis.

The need for human iPSC for DM2 cannot be overstated. In the absence of human-derived DM2 cell lines, researchers have been working with and transfecting the mouse myoblast C2C12 cell line. This cell line is a subclone of myoblasts that were originally obtained by Yaffe and Saxel at the Weizmann Institute of Science in Israel in 1977. The cell line is widely available through commercial vendors such as ATCC.

- Mouse models:
 - Dr. Charles Thornton, MD, University of Rochester, has been working for several years to create a DM2 mouse based on the technology he used to develop the HSA^{Ir} DM1 mouse. Significant technical challenges arise in making a DM2 transgenic mouse because of the expansion size and amount of toxic RNA involved. A successful DM2 mouse is likely in the next few years.
 - Dr. Maurice Swanson, PhD, University of Florida, is working on a DM2 mouse model using CRISPR-Cas9 technology to impact the germline through expression of the DM2 repeats in the mutant gene in endogenous DM2 intron 1 of CNBP.
 - Dr. Kiruphagaran Thangaraju, PhD, University of Florida, a MYOTONIC-funded Fellow in the lab of Dr. Laura Ranum, generated a patient-derived BAC library and screened ~92,000 clones. Dr. Thangaraju has identified several clones with the CNBP gene. One clone in particular contains a substantial flanking sequence and a large (~700 CCTGs) repeat expansion. Pronuclear injections were performed but genotyping did not identify any transgenic founders. This work-inprogress continues as Dr. Thangaraju continues to purify additional BAC DNA with the expansion for another round of pronuclear injections. The goal is to identify transgenic founders and build colonies
 - Dr. Ralf Krahe, PhD, University of Texas MD Anderson Cancer Center, developed an HSA transgenic DM2 mouse model. The model includes a (CCTG)121 expansion in intron 1 of the human skeletal actin gene (DM2-HSATG) with a small number of repeats. The model does not demonstrate the mis-splicing events many believe to be associated with DM2. According to Dr. Krahe, the mice do display phenotypic changes associated with DM2.
 - Dr. Lubov Timchenko PhD, Cincinnati Children's Hospital, recently published a cnbp-KO mouse created at CCH. Her focus is on downstream genetic events resulting from the role of CNBP, such as the reduction in ZNF9. Dr. Timchenko proposes that heterozygous cnbp-KO mice could be a model for DM2 because patients with DM2 have only partial reduction of ZNF9, similar to that seen in the CCH heterozygous mice.

More mouse, and possibly other animal, models are needed to demonstrate a comprehensive picture of DM2 with multiple clinical-like phenotypic changes, as well as the molecular and mis-splicing events associated with the biology and pathogenesis of DM2.

• Other models for DM2 study:

- Dr. Andrew Berglund, PhD, The RNA Institute, collaborating with Dr. Karen Guillemin's lab at the University of Oregon, are developing and characterizing zebrafish models of myotonic dystrophy. The main focus of the zebrafish project is to the study the mechanisms underlying DM-related changes in gut motility and the microbiome due to GI issues suffered by DM patients.
- Teams including Dr. Ralf Krahe, PhD, at The University of Texas and Dr. Giovanni Meola, MD, at Università degli Studi di Milano are working with Drosophila models of DM2.
- Drug development efforts:
 - Dr. Andrew Berglund, PhD, RNA Institute, is evaluating the feasibility of conducting a clinical trial using a repurposed antimalarial for DM2 patients. He is also developing novel small molecules around unique chemistry scaffolds to target toxic RNA in DM2.
 - Dr. Matthew Disney, PhD, Scripps Research Institute, is working on early-stage small molecule compounds with the potential to treat patients with DM2, although his lead indication at this time is for DM1. His recently-formed biotech, Expansion Therapeutics, has formally expressed interest in developing DM2 therapies.

Potential areas for study and funding in DM2 (as of 8/2019):

There are no active NIH grants working exclusively on DM2, although 2 out of the 3 aims of the NIH 5 year, \$5M Wellstone grant to the University of Rochester are DM2-focused. Further, there are currently no DOD Peer-reviewed Medical Research Program grants funding DM2, although that funding stream has only been available for one year. Myotonic intends to launch a multi-year initiative to attract researchers and industry to DM2 discovery and drug development, assuming funds can be raised. Areas of potential solicitation across the spectrum of DM2 are included below.

The below were suggested by interviewees who provided content for this DM2 landscape review.

- 1. Longitudinal DM2 single and multi-site studies to drive disease understanding, and clinical trial outcome measures and design
- 2. Publication and availability of animal models and cell lines
- 3. Research into modifier genes and their potential role in DM2 disease progression
- 4. **Exploration of DM2 CNS symptoms** and mechanisms behind "brain fog" reported by many DM2 patients.

5. **Research into DM2 repeat size and incomplete penetrance,** collecting data from more patients with "small" DM2 expansions and known repeat tract structure to better estimate penetrance and provide for better genetic counseling for DM2 patients

6. Research examining age and sex differences in DM2

- 7. Studies of cellular nucleic acid-binding protein (CNBP) as a transcription regulators: Such studies may determine whether CNBP acts as a transcription regulator required for activating the innate immune response. Such studies may lead to the identification of specific binding motifs present in the promoter region of inflammatory cytokines that induce activity of the CNBP gene.
- 8. **Cancer and DM2** to explore earlier study results suggesting that cancer should be considered in the overall pathogenesis of disease manifestation in DM2 patient
- 9. **Overexpression of rbFOX1**: rbFOX1 has been identified as a possible molecular mechanism in DM2. More studies looking at its overexpression may lead to a greater understanding of the role it plays in releasing MBNL1 from sequestration within CCUG RNA foci and muscle atrophy.

RECENT PUBLICATIONS - 2015-2019

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