FSHD Therapeutic Strategies: What Will It Take to Get to Clinic?

Charis L. Himeda and Peter L. Jones *

The Department of Pharmacology, University of Nevada, Reno School of Medicine, Reno, NV 89557, USA; chimeda@med.unr.edu
* Correspondence: peterjones@med.unr.edu

Abstract: Facioscapulohumeral muscular dystrophy (FSHD) is arguably one of the most challenging genetic diseases to understand and treat. The disease is caused by epigenetic dysregulation of a macrosatellite repeat, either by contraction of the repeat or by mutations in silencing proteins. Both cases lead to chromatin relaxation and, in the context of a permissive allele, pathogenic misexpression of DUX4 in skeletal muscle. The complex nature of the locus and the fact that FSHD is a toxic, gain-of-function disease present unique challenges for the design of therapeutic strategies. There are three major DUX4-targeting avenues of therapy for FSHD: small molecules, oligonucleotide therapeutics, and CRISPR-based approaches. Here, we evaluate the preclinical progress of each avenue, and discuss efforts being made to overcome major hurdles to translation.

Keywords: facioscapulohumeral muscular dystrophy; FSHD; DUX4; skeletal muscle; muscular dystrophy; gene therapy; AAV; CRISPR; antisense; therapeutics

1. Introduction

Facioscapulohumeral muscular dystrophy (FSHD) is one of the most prevalent myopathies that afflict males and females of all ages [1–4]. With the onset of clinical weakness typically appearing in the second or third decade of life and ~20% of patients ultimately using a wheelchair, the personal, social, and economic costs of this disease are enormous [5,6]. Although progress is being made, there are no cures or ameliorative treatments for FSHD, so an effective therapy is critically needed.

FSHD is caused by genetic mutations leading to epigenetic dysregulation of the D4Z4 macrosatellite repeat array at chromosome 4q35. FSHD1 is caused by contraction of the repeat, while FSHD2 is caused by mutations in silencing proteins (reviewed in [7]). Both cases lead to chromatin relaxation and, in the context of a permissive allele, misexpression of the DUX4 gene in skeletal muscle. DUX4 encodes a pioneer transcription factor that activates a program of gene expression during early human development [8–10], after which its expression is silenced in most somatic cells. When misexpressed in FSHD skeletal muscle, the DUX4 transcriptional program ultimately leads to myofiber degeneration with variable histopathologic features. Thus, in sharp contrast to most other genetic myopathies, which are caused by mutations in genes encoding muscle structural, regulatory, or signaling components, leading to a loss of function that needs to be fixed or replaced, FSHD is a toxic gain-of-function disease [11–15]. This necessitates the design of very different therapeutic strategies and, while avoiding some translational difficulties, introduces others.

While the most direct approach to therapy is eliminating DUX4 expression (Figure 1), both the gene and the D4Z4 repeat that encodes it present unique therapeutic challenges. Although highly similar D4Z4 arrays are found at multiple loci in the genome [16], DUX4 is only stably expressed from the distal-most repeat unit on a disease-permissive allele [17–20] due to the presence of a polyadenylation signal (PAS) in an exon distal to the array (Figure 2) [20,21]. Additionally, while all FSHD myocytes are epigenetically poised to express DUX4, the transcript and protein are only expressed rarely, in stochastic
bursts [21–24], consistent with the sporadic muscle involvement seen in FSHD patients. Moreover, while other mammals contain functional orthologs, the D4Z4 array and intact DUX4 gene are not well conserved outside of old-world primates and the Afrotherian lineage, and no natural animal models exist [25–28].

![Diagram of DUX4 regulation and downstream targets](image-url)

**Figure 1.** Incomplete silencing of the FSHD locus leads to misexpression of the pathogenic DUX4-fl transcript in skeletal muscles. DUX4-FL subsequently activates a program of embryonic gene expression, which triggers many aberrant downstream events. While some pathways downstream of DUX4 are amenable to targeting (orange), it is not clear which individual pathway, if any, is causal for pathology. Thus, serious efforts to develop specific FSHD therapeutics (green) have focused on restoring silencing to the locus or preventing expression of DUX4-fl mRNA or protein. Purple indicates DUX4-targeting strategies that are inherently promiscuous; asterisk indicates small molecules targeting DUX4-epigenetic regulation or transcription. Blocking DUX4-FL activity (blue) is a theoretically viable avenue, although still in the very early stages of development.

The creation of FSHD-specific therapeutics has been understandably hampered by these challenges, and in the interim, easier avenues have been explored, including compensatory treatments to improve muscle strength or health [29–36] (ClinicalTrials.gov Identifiers: NCT02927080, NCT02603562), immunomodulation (ClinicalTrials.gov Identifier: NCT02579239), and most recently, the repurposed FDA-approved drug losmapimod [37] (ClinicalTrials.gov Identifiers: NCT04003974, NCT04264442). Thus far, these treatments have either failed or yielded underwhelming results in clinical trials. This is not unprecedented; many treatments across numerous indications have been quite successful in preclinical studies only to fail during translation. Events in the field of myotonic dystrophy have underscored the importance of not abandoning alternative avenues of therapy even in the wake of a very promising treatment [38]. Fortunately, DUX4 misexpression offers numerous candidates for therapy, and the field has benefited from rapid advances in both targeting modalities and modes of delivery. Thus, there are now many FSHD-specific therapeutics in the developmental pipeline (Figures 1 and 2).
The bulk of work exploring FSHD therapeutic avenues has been recently and extensively reviewed elsewhere [39–41]. In this review, we evaluate the major avenues of therapy for FSHD in terms of their translational progress. We compare the advantages and disadvantages of each approach and take a hard look at the hurdles that must be overcome for each to reach the clinic.

2. Small-Molecule Drugs

These therapeutics are chemical compounds with a molecular weight in the range of 0.1–1 kDa that can target extracellular components or easily enter cells to affect intracellular molecules.

2.1. Progress in the Field

The discovery that FSHD pathogenesis is caused by misexpression of \textit{DUX4} has spurred the identification of numerous potential therapeutic targets, both upstream and downstream of the pathogenic gene. However, FSHD creates complications not present in traditional drug screening. To begin with, \textit{DUX4} is only expressed in a few primary human tissues, and differentiated FSHD skeletal myocytes—the therapeutic target in vivo—are typically used for identifying repressors of \textit{DUX4} expression. In addition to the relative difficulties of growing and differentiating myoblasts, FSHD cultures typically exhibit extremely low and highly variable levels of \textit{DUX4} mRNA and protein. Thus, one must look for a decrease in levels of a transcription factor that is already expressed at very low levels, which is not an ideal situation for high-throughput drug screening. To address these issues, many academic and corporate researchers identify lead compounds through indirect expression screens using \textit{DUX4} reporter constructs as opposed to directly measuring \textit{DUX4} expression levels.
transcript levels. This screening method is limited by the content of the chemical libraries screened, dosing, and modes of action. Highlighting these issues, despite the clear overlap in libraries, two published screens with similar approaches identified different molecules, targets, and pathways for DUX4 inhibition, even to the exclusion of other targets [42,43], which is cause for concern. Regardless, screens for compounds that repress DUX4 expression yielded bromodomain and extra-terminal (BET) inhibitors, agonists of the β2 adrenergic receptor [42,43], phosphodiesterase (PDE) inhibitors [43], p38 inhibitors [44,45], and Wnt agonists [46].

In contrast to chemical library screening, a candidate gene knockdown approach identified key regulators of DUX4 expression (ASH1L, BRD2, KDM4C, and SMARCA5) that would be prime therapeutic targets for small-molecule inhibition [47]. Other studies identified the NuRD and CAF-1 complexes [48], PARP1 [49], and SMCHD1 [50] as endogenous D4Z4 repressors. Boosting the expression or activity of such factors, potentially through small molecules, represents an alternative avenue for therapy. In a different approach, a genome-wide CRISPR-Cas9 screen identified DUX4 pathways that could potentially be targeted for therapy [51]. While these and other targets downstream of DUX4 activity have been identified [51–54], it is not clear which, if any, of these downstream pathways are causal for pathology, or if successful targeting of any would provide therapeutic benefit. Additionally, many downstream factors are part of ubiquitous and robust cellular pathways (e.g., p300) [54]; thus, drugs targeting them are likely to produce a host of unwanted effects.

To date, only one putative small-molecule inhibitor of DUX4 expression has reached clinical trials in the US and Europe. The p38 inhibitor losmapimod emerged as a repressor of DUX4 expression in two independent screens [44,45]. Subsequently, Fulcrum Therapeutics licensed losmapimod, originally developed for the treatment of cardiovascular disease, from GlaxoSmithKline. Following early trials ([37]; ClinicalTrials.gov Identifier: NCT04003974), losmapimod is currently scheduled to enter a Phase III clinical trial for FSHD in Q2 of 2022 (ClinicalTrials.gov Identifier: NCT04264442), but the results thus far are not encouraging. Data reported in June 2021 from the Phase Ib trial (ClinicalTrials.gov Identifier: NCT04003974) showed only nominal benefit at 48 weeks, and the primary end-point (reduction of DUX4 target gene expression) was not met. This indicates that any short-term functional improvements were DUX4-independent and, in the absence of DUX4 repression to halt new muscle pathology, casts doubt on the likelihood of long-term benefit.

2.2. Advantages

While other forms of therapy are gaining ground, small molecules still hold significant advantages and represent the gold standard for treatment of most disorders [55]. Compared to biologics, small molecules are easier and cheaper to synthesize and generally more accessible to patients. They are generally taken orally, readily taken up by cells, stable, and not immunogenic. Additionally, the modality is well-established and inherently safer than any other.

2.3. Disadvantages

Side-effects. Small molecules can cause both off-target and on-target toxicity, with the latter being a particular concern for targets that are widely expressed and play key cellular roles. Losmapimod serves to illustrate this point, since it targets p38α, a key regulator of muscle biology [56–58]. Chronic therapeutic targeting of p38 for FSHD is quite different from protocols used for other indications; it requires hitting a narrow dosing window to repress DUX4 expression without compromising important pathways such as muscle regeneration. Ultimately, the potential long-term negative effects of this dosing regimen are unknown. However, the D4Z4 repeat arrays are somewhat uniquely regulated in the human genome, and may offer more specific targets for therapy. Certain D4Z4/DUX4 epigenetic regulators are not global regulators, which reduces the likelihood of harmful on-target effects. Additionally, even small changes in the levels of these factors can lead to striking decreases in DUX4 expression [47]. This suggests that even incomplete target
inhibition (i.e., very low doses) may be therapeutic, which should reduce the likelihood of both on- and off-target toxicity.

Expensive over the long term. A traditional drug treatment for FSHD will require chronic, lifelong administration.

2.4. Path to Clinic

Fulcrum Therapeutics, Myocea, Altay Therapeutics, and Facio Therapies, along with a number of academic labs, have reported active programs for small-molecule discovery and development in FSHD. Drugs for some candidate therapeutic targets (e.g., BET inhibitors, losmapimod) are already FDA-approved, while other targets are still in the initial phases of drug development. The repurposing of FDA-approved drugs as potential therapeutics for FSHD is attractive from a business standpoint. However, while these have been shown to be safe for indications such as cancer, they are not necessarily the safest or most efficacious treatments for a muscle disease. Drugs targeting ubiquitous cellular effectors are likely to have serious side effects, particularly during the long-term dosing required for FSHD. As mentioned above, even selective p38 inhibition via losmapimod may have harmful long-term effects in skeletal muscle, since p38 and its targets regulate many stages of myogenesis [56–58]. In addition, while targeting kinase pathways is a common clinical path, they are notoriously compensatory, which casts doubt on the likelihood of long-term inhibition [59].

The alternative route is to screen and optimize compounds targeting more specific therapeutic targets for which no drugs exist. While this approach is more costly and time-consuming, it enables the design of more specific drugs to better targets. The limited efficacy of losmapimod in the Fulcrum trial highlights the disadvantage of taking the short road to the clinic. In some cases, the long road may be the fastest route to a safe and effective therapy.

The following two sections deal with biologic therapeutics: compounds of biological origin, typically larger than small molecules, including peptides, proteins, nucleic acid-based molecules (oligonucleotides and gene therapies), and antibodies.

3. Oligonucleotide Therapeutics

3.1. Progress in the Field

FSHD is a dominant gain-of-function disease ideally suited for antisense or RNAi-based approaches. Over the last decade, various antisense approaches (Figure 2) have been tested against the DUX4 gene and its pathogenic transcript, with considerable success in both in vitro and in vivo proof-of-principle studies.

The earliest antisense studies in the field focused on altering DUX4 pre-mRNA processing. In healthy cells, DUX4 can produce a short mRNA isoform that is translated into a nontoxic protein. In FSHD myocytes, there is a shift in mRNA splicing to generate the full-length, pathogenic DUX4 isoform (DUX4-fl). Initial studies geared toward therapy showed that siRNAs and antisense oligonucleotides targeting the DUX4 PAS and mRNA splice sites reduced levels of DUX4-fl mRNA and protein in FSHD myotubes [60,61]. When targeted to the DUX4 promoter, siRNAs induced a DICER/AGO-dependent epigenetic transcriptional silencing [62]. Phosphorodiamidate morpholino oligomers (PMOs) targeting the DUX4 PAS reduced levels of DUX4 and its targets in cultured FSHD myocytes [63,64] and following intramuscular delivery to a xenograft model [64]. LNA gapmers (against exon 3) and 2′-O-methoxyethyl (2′-MOE) gapmers (against the coding region) reduced DUX4 levels in immortalized FSHD myocytes and following intramuscular delivery to the FLexDUX4 FSHD mouse model [65,66]. Importantly, in a recent study, PAS-targeting PMOs delivered systemically reduced DUX4 and its targets, ameliorated pathology, and improved muscle function in the FLexDUX4 FSHD mouse model [67,68].

DUX4-targeting miRNAs were shown to reduce DUX4 mRNA and protein in a DUX4-overexpression mouse model [69,70], and U7 antisense DUX4 snRNAs reduced levels of DUX4 and its targets in cultured FSHD myocytes [71].
A different oligonucleotide approach utilizes DNA binding site decoys (DNA aptamers) to bind a transcription factor and block its ability to regulate target genes. Such decoys can bind recombinant DUX4-FL with high affinity [72], and others were demonstrated to block binding of DUX4 to its target genes following electroporation of DUX4 into wild-type mice [73].

3.2. Advantages
- ability to directly repress expression of DUX4, a difficult-to-drug transcription factor
- relatively straightforward to design, produce, and screen
- sequence-based targeting allows for precision/personalized medicine
- treatment can be stopped if adverse events occur

3.3. Disadvantages
- Off-target effects/cytotoxicity. All oligonucleotide drugs have the potential for sequence- and chemistry-dependent cytotoxicity and immunogenicity. Sequences that function efficiently in cell culture may have inherently bad toxicology profiles in non-human primates (NHPs) and humans. Oligonucleotide therapeutics that use RNAi are notoriously cytotoxic. miRNAs are inherently less specific than ASOs because they bind via incomplete base-pairing, increasing the risk of off-target effects. siRNAs are double-stranded, which increases the risk of an innate immune response.

- Dysregulation of endogenous RNA-processing pathways. All oligonucleotide drugs utilize endogenous cellular enzymes to mediate their effects. siRNAs, shRNAs, and miRNAs depend on RNAi, thus competing with normal substrates of the RISC complex and sometimes oversaturating the capabilities of this pathway [74]. Additionally, some ASOs that depend on RNase H for target degradation can induce “ASO tolerance” via increased expression of the target pre-mRNA [75].

- Inefficient uptake into muscle cells. In general, oligonucleotide drugs suffer from poor delivery to tissues other than liver.

- Requirement for chronic, lifelong administration. This requirement will likely exacerbate any toxicity caused by the treatment.

- Approved oligonucleotide drugs will likely be extremely expensive over a lifetime. To date, there is no publicly available data on durability and dosing for these compounds in FSHD.

3.4. Path to Clinic

Oligonucleotide drugs for multiple diverse disorders, including Duchenne Muscular Dystrophy (DMD), have recently been approved by the FDA [76], demonstrating that this technology has reached clinical maturity for some indications. For FSHD, it is encouraging that a number of biotech companies (Avidity Biosciences, Dyne Therapeutics, Arrowhead Pharmaceuticals, miRecule, and Armatusbio) are actively developing DUX4-targeting oligonucleotide therapeutics for FSHD.

Delivery challenges must be addressed. Fortunately, there is a major push to develop technologies improving the drug-like properties of nucleic acid therapeutics for multiple indications. These include chemical modifications, backbone modifications, conjugation to a carrier that facilitates uptake by the target tissue(s), and nanocarriers [76]. For some indications, nucleic acid oligomers (e.g., mRNA vaccines) are now in use, thanks in large part to efficient delivery by lipid nanoparticles (LNPs). Although targeting of LNPs to body-wide skeletal muscles poses different challenges and has yet to be optimized, cell-penetrating peptide conjugate oligonucleotides may improve delivery to muscle. Avidity Biosciences and Dyne Therapeutics, two companies with active FSHD programs, have reported the development of antibody conjugates that improve muscle uptake. Oligonucleotide drugs can also be delivered by gene therapy vectors (see below).

Specificity must be rigorously assessed for every candidate oligonucleotide drug at efficacious doses. Thus far, only two studies have reported RNA-seq data following treatment [64,66]. (Lim et al. focused only on the correction of FSHD signature gene expression, whereas
Chen et al. reported no significant off-target effects of their PMO treatment on the muscle transcriptome. Additionally, it is important to assess off-target effects in muscle cells that are not expressing DUX4, since these will be the overwhelming majority in a patient subjected to treatment, and the specificity of an oligonucleotide drug may decrease in the absence of a perfect target.

Potential immunogenicity/toxicity of any carriers/conjugates must be assessed at efficacious doses.

Long-term efficacy must be assessed. Thus far, the longest time point assessed in any FSHD study is following 4 weeks of treatment in a mouse [68]. The critical question is: can an oligonucleotide drug be administered systemically and chronically at a dose that provides functional benefit without toxic effects? Eteplirsen and golodirsen, FDA-approved exon-skipping drugs for DMD, have shown little clinical benefit, and it can be argued that the situation in FSHD is more challenging to treat. Treatment of DMD requires production of functional dystrophin at levels far below those found in healthy individuals and in only a fraction of myofibers, whereas treatment of FSHD may well require much higher doses to achieve DUX4 repression, which will likely be required in a majority of myofibers, since every muscle nucleus is poised to express DUX4 at any given time.

4. Gene Modification

4.1. Progress in the Field

Over the past few years, several groups have attempted CRISPR modulation of the FSHD locus, either by Cas9-mediated gene editing (CRISPRe) or dCas9-mediated transcriptional inhibition (CRISPRi) (Figures 2 and 3). Recently, the Dumonceaux lab demonstrated that editing the DUX4 exon 3 PAS (by Cas9 or TALENs) is highly inefficient [77]. Even in successfully edited clones, pathogenic DUX4 levels are reduced, but not silenced, likely due to compensation by a cryptic PAS; the authors concluded that this is not a viable therapeutic strategy for FSHD [77]. By contrast, another group attempting a similar approach reported that CRISPRe of the exon 3 PAS reduced D4Z4 transcripts and expression of DUX4 target genes in immortalized FSHD1 myoblasts [78]. RNA-programmable base editing targeting the DUX4 PAS (18% efficiency) reduced levels of DUX4 and its target genes in both FSHD1 and FSHD2 myoblasts, and a cursory assessment of specificity was encouraging [79]. Finally, CRISPRe correction of an SMCHD1 intronic variant restored SMCHD1 expression to healthy levels and reduced DUX4 expression [80].

Our group has taken a CRISPRi approach to mitigate the concerns related to genome editing, and because CRISPRi is ideally suited to correcting a disease caused by the pathogenic misexpression of a powerful transcription factor [81–83] (Figures 2 and 3). Targeting the enzymatically inactive dSaCas9 fused to one of several epigenetic repressor domains to the DUX4 promoter/exon 1 increases chromatin repression at the disease locus, specifically repressing DUX4 and its target genes in FSHD myocytes and in a mouse model of the disease [83]. We have recently re-engineered this platform to a single-vector system in which all therapeutic components can be packaged into recombinant AAV (rAAV) vectors for gene therapy.

4.2. Advantages

- Both CRISPRe and CRISPRi have the potential to achieve long-term or permanent correction of all forms of FSHD with a single treatment.
- In principle, CRISPRi is ideally suited to a dominant, gain-of-function disease such as FSHD. Since it does not involve cutting the genome, CRISPRi may also be safer than CRISPRe.
- In the long run, CRISPR therapies (and other gene therapy approaches) may be the most economical means of treatment, since they are delivered in a single dose that is meant to function over a lifetime.
Figure 3. CRISPR gene modification approaches to FSHD. (A) The Cas9 nuclease cuts DNA at sites targeted by single-guide RNAs (sgRNAs). DNA repair by nonhomologous end joining (NHEJ) mediates the disruption of genomic sequences by insertions/deletions (indels), whereas homology-directed repair (HDR) mediates precise editing in the presence of a donor template. CRISPR editing of the D4Z4 repeat (*) necessitates cutting the genome in hundreds of unintended places, as these repeats exist in many copies on the noncontracted 4q chromosome and both 10q chromosomes, in addition to polymorphic D4Z4s at other loci. Thus, groups attempting CRISPR editing have focused on the polyadenylation signal (PAS) in DUX4 exon 3, which provides a safer target. (B) RNA-programmable base editing utilizes a wild-type tRNA adenosine deaminase (TadA) and an engineered version of TadA (TadA*) fused as a dimer to a nicking version of Cas9 (nCas9). This enables adenine-to-guanine substitution without introducing double-stranded DNA breaks. (C-D) As an alternative strategy, enzymatically dead Cas9 (dCas9) is fused to the KRAB domain for transcription inhibition (panel C) or to chromatin-modifying proteins, which can act in a broader fashion across the locus (panel D). Fusing minimized versions of these regulators to dCas9 from Staphylococcus aureus allows packaging into rAAV vectors to silence pathogenic DUX4 expression in FSHD muscle. Additional abbreviation: PAM, protospacer-adjacent motif.

4.3. Disadvantages

Immunogenicity related to CRISPR components [84]. In the one in vivo clinical trial of CRISPR gene editing to date, it is encouraging that no significant safety concerns were reported in the six patients at 28 days post-treatment [85]. However, several studies have demonstrated that Cas9 evokes a host cellular and humoral immune response, and a large proportion of the population has pre-existing immunity to Cas9 [84]. Thus, developing strategies to avoid a potential Cas9 immune response is critical. These include: (1) removing immunogenic Cas9 epitopes (immunodominant epitopes have been removed from SpCas9 while preserving its cutting efficacy) [86], (2) interfering with processing and presentation of Cas9 antigens, (3) using orthologs from nonpathogenic bacteria, (4) inducing immune tolerance, and (5) transient immunosuppression [84]. miRNA-based detargeting has also emerged as a way to specifically repress expression of immunogenic transgenes in antigen-presenting cells by incorporating binding sites for macrophage-specific miRNAs into rAAV vectors [87,88].
Off-target effects. These are a bigger concern for CRISPRe than for CRISPRi approaches, since off-target cutting is likely to be more detrimental than off-target repression.

Genomic instability. This is only a concern for CRISPRe approaches, which involve genomic cutting.

4.4. Path to Clinic

At least one company (EpiSwitch Rx) has an active CRISPR program for FSHD.

- For CRISPRe, rigorous tests of specificity are required. Rapid advances in specific, programmable base-editing (e.g., prime-editing) [89] should greatly improve the safety of CRISPRe for FSHD. However, current base editors are much too large to be accommodated in a single-vector system for rAAV-mediated delivery [89]. These need to be better characterized and minimized to be therapeutically relevant.

- Stable repression mediated by CRISPRi remains to be demonstrated and is absolutely necessary to achieve, since current rAAV vectors for gene therapy can only be administered once. In brief, either the CRISPRi treatment must elicit a stable epigenetic change or the rAAV episome must persist indefinitely in treated fibers (discussed in more detail below).

- Both types of CRISPR approaches must be tested more rigorously in appropriate in vivo models, including a large animal model (see below). For CRISPRi, a human xenograft model [90,91] is ideal for assessing the persistence of epigenetic changes at the disease locus, as xenografted mice contain a full D4Z4 array from an FSHD patient.

5. Needs for Therapeutics Delivered by Gene Therapy (CRISPR Strategies, RNAi)

5.1. Delivery to Body-Wide Skeletal Muscles

Currently, rAAV vectors are the only practical way to deliver Cas9/dCas9 components to skeletal muscles throughout the body. Several AAV serotypes with high muscle tropism have been widely used in preclinical studies and in clinical trials for DMD and X-linked myotubular myopathy (XLM-TM) (clinicaltrials.gov identifiers: NCT03362502, NCT03368742, NCT03375164, and NCT03769116). Despite the utility of these vectors, the need remains to increase the efficiency of delivery, lower the high cost of therapy, and reduce the liver and immune toxicity associated with high viral doses. For these reasons, rAAV-mediated gene therapies should use therapeutic cassettes contained within single vectors [92].

Recently, two groups have reported engineered capsid variants (“AAVMYO” and “MyoAAV”) with greatly enhanced tropism for striated muscles combined with detargeting of the liver [93,94]. These capsids demonstrate greater efficiency and selectivity of muscle targeting than those currently being used in clinical trials and in comparable preclinical studies. The MyoAAV class was shown to be highly muscle-tropic in both mice and NHPs, with therapeutic efficacy demonstrated following systemic administration of a low dose in multiple disease models [94]. This ability to lower the effective vector dose by even several-fold represents a major safety and efficacy milestone for gene therapies of muscle diseases.

5.2. Overcoming rAAV-Related Limitations

Gene therapies using rAAV vectors have been used safely and successfully in many clinical trials, and three (Glybera, Luxturna, and Zolgensma) have been approved. Nonetheless, use of these delivery vectors has several limitations.

Neutralizing antibodies to the capsid prevent re-administration. Although efforts are being made to reduce the immunogenicity of rAAV vectors [95,96], gene delivery by rAAV is currently a single-dose injection; thus, stability of the treatment is key. For FSHD, this means that either the rAAV episome must persist indefinitely in infected fibers, or that the treatment itself must elicit a stable change (e.g., epigenetic repression at the disease locus). Whether the transduced vector will eventually be diluted past the point of therapeutic efficacy is unknown, but rAAV episomes have shown remarkable persistence in skeletal muscle, a largely quiescent tissue, assembling into active chromatin and mediating
transgene expression for several years in the muscles of NHPs [97]. In clinical trials for DMD, micro-dystrophin delivered by rAAVrh74 has persisted for over 3 years [98], and for spinal muscular atrophy, survival motor neuron delivered by AAV9 has proven efficacious for over 5 years [99].

Pre-existing immunity to AAV renders a certain percentage of the population untreatable [89]. Since neutralizing antibodies can cross-react with multiple AAV serotypes, switching serotypes is not a viable strategy to avoid neutralization of rAAV vectors [100]. More realistic strategies to circumvent this include: (1) transient or long-term immunosuppression [101], (2) plasmapheresis to deplete AAV-specific neutralizing antibodies [102], (3) treatment with IgG-cleaving endopeptidases to decrease anti-AAV antibody titers [103], and (4) chemical modification [104], capsid engineering [105], or use of empty capsid decoys [106] to protect rAAV vectors from neutralization. The capsid can also trigger clearance of infected cells by cytotoxic T lymphocytes [107], an event that can be mitigated by Treg cells [108]; thus, autologous Treg cells may prove to be a useful adjuvant to in vivo gene therapies [109]. For most of these strategies, effectiveness in clinical settings remains to be tested. Additionally, benefits may not be straightforward, as with the use of empty capsids [110].

High doses cause liver toxicity. Recent clinical trials for DMD and XLMTM (clinicaltrials.gov identifiers: NCT03368742 and NCT03199469) were put on hold by the FDA due to severe adverse events associated with liver toxicity [111]. Although the hold on the DMD trial has since been lifted, this highlights the urgent need to detarget the liver and lower efficacious doses, which is currently being met [93,94].

Potential genotoxicity caused by genomic integration. Predominantly episomal, rAAV vectors have been found to integrate in the host genome at very low frequencies. It is not known whether long-term persistence of rAAV vectors in largely quiescent tissues such as skeletal muscle might eventually result in genotoxicity due to vector genome integration; however, the evidence to this point has led to a general consensus within the gene therapy field that such events are extremely unlikely [112].

High cost of production. Approved gene therapies using rAAV vectors are astronomically expensive. Priced at USD 1.2 million, Glybera has already been withdrawn from the market after treatment of just a single patient. At USD 2.1 million, Zolgensma is now the most expensive drug in the world. While the economics of an effective single-dose gene therapy vs. a chronic, lifelong treatment are beyond the scope of this review, high manufacturing costs can be lowered by lowering the effective dose. This can be accomplished by: (1) reducing packaging of partial/truncated genomes and empty capsids that co-purify with the therapeutic vector [110,113–115], (2) reducing immunogenicity [95,96], and (3) detargeting liver while increasing delivery to skeletal muscle [93,94].

6. Needs for All DUX4-Targeted Therapeutics

6.1. A Large Animal Model of FSHD for Preclinical Testing of Therapeutics

It has long been established that mouse models do not accurately reflect important features of human disorders. In addition, the delivery, dosing, specificity, efficacy, and durability of treatments are highly variable between a mouse (~25–35 g) and an adult human (~45–135 kg). As a result of these differences, many treatments have been quite successful in mouse models only to fail during clinical trials. Testing candidate therapeutics in a large animal model with closer genetic, anatomical, physiological, and metabolic similarities to humans should go a long way toward solving this problem.

However, rising ethical concerns have cast disfavor on the use of NHPs and dogs in biomedical research, and when used for testing therapeutics, these studies are often underpowered due to their high costs. Minipigs have come to fill this gap and are becoming established as the large animal model of choice for translational research [116–118]. They share many similarities to humans in terms of size, physiology, anatomy, metabolic profile, and lifespan, while offering lower operating costs and the ability to perform properly powered experiments to determine both efficacy and toxicology [119–121]. To fill this
need, the first large animal model of FSHD, a Göttingen minipig based on conditional expression of DUX4-fl in skeletal muscles [122], was generated and is currently being validated with respect to FSHD gene expression signature and inducible myopathy. Most FSHD therapeutics under development require only a large animal model with consistent, inducible expression of DUX4-fl and its target genes in order to test target engagement, molecular efficacy, dosing, and delivery. Thus, while a phenotypic large animal model would be ideal, any model that produces mosaic DUX4 expression in skeletal muscle cells will still be extremely useful for the therapies described above.

6.2. DUX4-Responsive Circulating Biomarkers of Disease Progression

One of the greatest unmet needs in the FSHD field is a robust and reliable circulating biomarker for pathogenic gene expression or disease progression that can be used to readily monitor the efficacy of a treatment. Compared with muscle biomarkers, serum biomarkers are minimally invasive, inexpensive, and ideal for monitoring drug responsiveness over time. For drugs targeting DUX4 expression, a DUX4-responsive circulating biomarker is needed, since there is no consensus on which genes downstream of DUX4 reflect pathological changes in FSHD muscle.

Several studies have identified proteins, mRNAs, and miRNAs that are differentially expressed in the serum of FSHD patients vs. healthy controls [123–127]. Among these, miR-206 was shown to be DUX4-responsive, and thus, a strong candidate for a biomarker to assess DUX4-targeting treatments [127]. Levels of IL-6, a cytokine and myokine that is elevated in several neuromuscular diseases, were found to correlate reliably with clinical severity, making it a good potential biomarker for tracking disease progression [128]. All candidate biomarkers to date require independent validation in large patient cohorts, as well as longitudinal evaluation.

The current state of clinical outcome measures, including imaging biomarkers, has been reviewed in detail [129]. These authors call for the use of circulating biomarkers of active muscle disease in combination with whole-body MRI and comprehensive quantitative assessments of muscle function [129], all of which are undergoing development and optimization. In addition, standardization of patient procedures is absolutely critical when interpreting outcomes. Standard operating procedures regarding fasting, exercise, ambient temperature, etc. must be established and rigorously adhered to.

The development of biomarkers and clinical outcome measures is particularly challenging for FSHD, considering the heterogeneity of muscle involvement and the highly variable and unpredictable course of disease progression. The usefulness of any outcome measures depends on the establishment of proper enrollment criteria, which will vary depending on the treatment, and proper patient stratification. The latter cannot depend exclusively on repeat size, since this does not correlate with clinical severity [130–133]. Clinical severity, type of FSHD, presence of modifiers, degree of D4Z4 hypomethylation, and location of muscle involvement must all be considered during stratification.

7. DUX4-Independent Approaches

Incapacitating DUX4 and halting pathology are the primary goals of most therapeutic programs for FSHD. However, certain DUX4-independent interventions may prove useful as part of a combination therapy or adjuvant treatment.

Treatment with oral antioxidants yielded minimal benefit in a recent FSHD trial [35], but outcomes might be improved with mitochondria-targeted antioxidants [134] or by targeting multiple common pathways of mitochondrial dysfunction [135]. In this regard, aerobic exercise improved mitochondrial function and physical performance in DM1 patients [136], and might be helpful to FSHD patients, as well. Creatine supplementation has also improved muscle strength across multiple muscular dystrophies [137]. Regarding immune suppression, anti-IL-6 receptor monoclonal antibodies tocilizumab and sarilumab, approved for other indications [138], might be tested in FSHD, although on-target effects are likely to be an issue. While the immunomodulator ATYR1940 was unsuccessful in a
clinical trial for FSHD, it is feasible that an anti-inflammatory treatment in combination with a DUX4-targeting therapy might be effective at halting disease progression and restoring a healthy muscle environment.

Despite failure in a number of clinical trials, the book is not entirely closed on myostatin inhibition as a means of increasing muscle strength under the right conditions. The clear discrepancy between clinical trial results and effectiveness in mouse models highlights: (1) the difference between mouse and human, and the need for large animal models of disease, and (2) the need for properly stratified and controlled clinical trials. Often, a single trial does not provide a straightforward answer, as myostatin inhibition failed for different reasons in different trials [139]. Importantly, circulating baseline levels of myostatin are higher in FSHD than in DMD, and the ACE-083 trial demonstrated significant muscle hypertrophy (without an increase in strength) in treated FSHD patients [36]. Thus, it is conceivable that in combination with a DUX4-targeting therapy, myostatin inhibition might increase muscle function as well as size in FSHD.

The need for well-planned clinical trials becomes more pressing as a community grows increasingly desperate for a treatment, and is willing to grasp at untested straws. Two of these—stem cell therapy and exosome treatments—may be useful for providing local immune suppression, but their safety, efficacy, and durability are still untested for FSHD, even in preclinical models.

The potential benefits of stem cell implantation (generally used to repair or replace muscle in an otherwise healthy environment) are by no means straightforward in a dystrophic environment such as FSHD muscle. As the myostatin trials demonstrate, effects in healthy muscle are not necessarily predictive of effects in dystrophic muscle. Nonetheless, the efficacy of stem cell therapy has been well documented for other indications, and is worth testing in carefully planned, well-controlled, and properly powered clinical trials for FSHD. By contrast, the use of exosomes to carry cytokines or growth factors is promising, but these vesicles are just beginning to be characterized. Their use in a clinical setting is untested and unapproved, yet unregulated clinics have sprouted up in many countries, including the US, and are flourishing under the notion that the potential benefits are worth the risks. This is a serious concern because the risks are not trivial. Beyond the risks associated with the unregulated use of biologics from questionable sources, treatments that foster immune tolerance may only provide short-term benefit, while enabling the growth of early-stage tumors. Additionally, any treatment that stimulates muscle regeneration (as stem cell therapy is supposed to do) may deplete muscle satellite cells, thus hastening dystrophic progression.

None of these treatments will correct the underlying cause of FSHD, but one or more may ultimately prove useful, likely in combination with a DUX4-targeting therapy. When considering a general ameliorative treatment for FSHD, one must ask: (1) what are the likely benefits and risks in a dystrophic environment? (2) what are the probable effects in cells that are either expressing or poised to express DUX4? and (3) what are the best models for testing? While animal models are not always predictive of the response in humans, they have been indispensable for revealing the effects of perturbing conserved pathways and mechanisms.

8. Conclusions and Perspective

Despite the complexities of the disease and the locus, the FSHD field has made considerable progress since the pathogenic role of DUX4 was first identified. The fact that multiple modalities targeting DUX4 expression and activity are moving through the therapeutic pipeline is promising for the prospect of an effective treatment in the near future. With technology advancing rapidly on multiple fronts, it is hard to predict which avenue will be first to reach the clinic. Regardless, the first drug to market may not ultimately be the best; thus, it is important to continue developing other viable candidates while the long-term efficacy of any approved treatment is still in question.
Many tools have been developed and tested in proof-of-principle studies that do not always make the difficult leap to clinical relevance. Indeed, preclinical data must be especially robust and reliable for a rare disease where the number of participants is limited and clinical trial failure is highly damaging to the quest for any effective therapy [140]. Currently, there is a pressing need for studies that address the important translational questions and obstacles associated with each of the therapeutic avenues discussed here. Antisense approaches must demonstrate efficient uptake into muscle cells and both efficacy and lack of toxicity following long-term treatment. Similarly, CRISPR approaches must demonstrate long-term efficacy and safety, while mitigating Cas9 and AAV immunogenicity. One question that concerns all therapeutic approaches is: what level of DUX4 inhibition is required for functional benefit? While DUX4 is only expressed in rare myonuclei at any given time, every myonucleus in an FSHD patient is epigenetically poised to express DUX4. Thus, both antisense and CRISPR approaches will likely need to target a majority of myonuclei in a majority of muscle fibers throughout the body. Fortunately, all available data indicate that any reduction in DUX4 levels is likely to be therapeutic. In addition to the fact that asymptomatic individuals still express detectable levels of DUX4 [22,141], small increases have catastrophic effects in FSHD mouse models [122,142]. Thus, there is good reason to believe that even small decreases in DUX4 expression will be beneficial to patients.

FSHD is highly pleiotropic and the best treatment may prove to be a combinatorial or personalized approach that depends on disease type and severity. This may consist of a DUX4-targeted therapy in combination with a non-targeted approach, such as immune modulation or stem cell therapy. It is important to keep in mind that from a patient standpoint, simply halting disease progression does not constitute a real cure. To correct the primary defect in FSHD and also replace the loss of diseased muscle with healthy, functional muscle remains the ultimate goal. With continual progress in the many disciplines involved, such a prospect is no longer unfeasible to envision.

Author Contributions: Conceptualization and writing of the manuscript, C.L.H. and P.L.J. All authors have read and agreed to the published version of the manuscript.

Funding: This research received no external funding.

Acknowledgments: We thank Chip and Summer Wilson, the William R. Lewis family, the Chris Carrino Foundation for FSHD, the FSHD Canada Foundation, the FSHD Global Research Foundation, and Friends of FSH Research for supporting our FSHD research program. Funding provided by the Mick Hitchcock, Endowed Chair in Medical Biochemistry at UNR and the National Institute of Arthritis and Musculoskeletal and Skin Diseases, National Institutes of Health (1R01AR087984).

Conflicts of Interest: C.L.H. and P.L.J. are co-founders of EpiSwitch Rx, Inc., a company focused on bringing therapeutics for FSHD to the clinic. They are inventors on two U.S. patent applications pertaining to the use of CRISPR inhibition for FSHD and one pertaining to small molecule development for FSHD. P.L.J. is an inventor on a patent application identifying therapeutic targets for FSHD and on a patent for epigenetic diagnosis of FSHD.

References
1. Padberg, G.W. Facioscapulohumeral Disease. Ph.D. Thesis, Leiden University, Leiden, The Netherlands, 1982.
2. Deenen, J.C.; Arnts, H.; van der Maarel, S.M.; Padberg, G.W.; Verschuuren, J.J.; Bakker, E.; Weinreich, S.S.; Verbeek, A.L.; van Engelen, B.G. Population-based incidence and prevalence of facioscapulohumeral dystrophy. Neurology 2014, 83, 1056–1059. [CrossRef]
3. Wang, L.H.; Tawil, R. Facioscapulohumeral Dystrophy. Curr. Neurol. Neurosci. Rep. 2016, 16, 66. [CrossRef]
4. Orphanet. Prevalence and Incidence of Rare Diseases: Bibliographic Data. Orphanet Report Series: Rare Diseases Collection. 2021. Available online: http://www.orpha.net/orphacom/cahiers/docs/GB/Prevalence_of_rare_diseases_by_alphabetical_list.pdf (accessed on 23 May 2022).
5. Tawil, R.; Van Der Maarel, S.M. Facioscapulohumeral muscular dystrophy. Muscle Nerve 2006, 34, 1–15. [CrossRef]
6. Blokhuis, A.M.; Deenen, J.C.W.; Voermans, N.C.; van Engelen, B.G.M.; Kievit, W.; Groothuis, J.T. The socioeconomic burden of facioscapulohumeral muscular dystrophy. J. Neurol. 2021, 268, 4778–4788. [CrossRef]
7. Himeda, C.L.; Jones, P.L. The Genetics and Epigenetics of Facioscapulohumeral Muscular Dystrophy. *Annu. Rev. Genom. Hum. Genet.* 2019, 20, 265–291. [CrossRef]

8. De Iaco, A.; Planet, E.; Coluccio, A.; Verp, S.; Duc, J.; Trono, D. DUX-family transcription factors regulate zygotic genome activation in placental mammals. *Nat. Genet.* 2017, 49, 941–945. [CrossRef]

9. Hendrickson, P.G.; Dorais, J.A.; Grow, E.J.; Whiddon, J.L.; Lim, J.W.; Wike, C.L.; Weaver, B.D.; Pfliueger, C.; Emery, B.R.; Wilcox, A.L.; et al. Conserved roles of mouse DUX and human DUX4 in activating cleavage-stage genes and MERVL/HERVL retrotransposons. *Nat. Genet.* 2017, 49, 925–934. [CrossRef]

10. Whiddon, J.L.; Langford, A.T.; Wong, C.J.; Zhong, J.W.; Taptscott, S.J. Conservation and innovation in the DUX4-family gene network. *Nat. Genet.* 2017, 49, 935–940. [CrossRef]

11. Mbakam, C.H.; Lamothe, G.; Tremblay, J.P. Therapeutic Strategies for Dystrophin Replacement in Duchenne Muscular Dystrophy. *Front. Med.* 2022, 9, 859930. [CrossRef]

12. Pozsgai, E.; Griffin, D.; Potter, R.; Sahenk, Z.; Lehman, K.; Rodino-Klapac, L.R.; Mendell, J.R. Unmet needs and evolving treatment for limb girdle muscular dystrophies. *Neurolegener. Dis. Manag.* 2021, 11, 411–429. [CrossRef]

13. Soltanzadeh, P. Myotonic Dystrophies: A Genetic Overview. *Genes* 2022, 13, 367. [CrossRef]

14. Butterfield, R.J. Congenital Muscular Dystrophy and Congenital Myopathy. *Muscle Nerve* 2020, 61, 436–448. [CrossRef]

15. Heller, S.A.; Shih, R.; Kalra, R.; Kang, P.B. Emery-Dreifuss muscular dystrophy. *Nat. Rev. Neurol.* 2020, 16, 283–296. [CrossRef] [PubMed]

16. Potter, R.; Sahenk, Z.; Lehman, K.; Rodino-Klapac, L.R.; Mendell, J.R. Endogenous DUX4 expression in FSHD myotubes is sufficient to cause cell death and disrupts RNA splicing and cell migration pathways. *Hum. Mol. Genet.* 2015, 24, 5901–5914. [CrossRef] [PubMed]

17. Clapp, J.; Mitchell, L.M.; Bolland, D.J.; Fantes, J.; Corcoran, A.E.; Scouting, P.J.; Armour, J.A.; Hewitt, J.E. Evolutionary conservation of a coding function for D4Z4, the tandem DNA repeat mutated in facioscapulohumeral muscular dystrophy. *Am. J. Hum. Genet.* 2007, 81, 264–279. [CrossRef] [PubMed]

18. Leidenroth, A.; Clapp, J.; Mitchell, L.M.; Coneyworth, D.; Dearden, F.L.; Iannuzzi, L.; Hewitt, J.E. Evolution of DUX4 gene macrosatellites in placental mammals. *Chromosoma* 2012, 121, 489–497. [CrossRef] [PubMed]

19. Leidenroth, A.; Hewitt, J.E. A family history of DUX4: Phylogenetic analysis of DUXA, DUXB, and DUXC reveals the ancestral DUX4 gene. *BMC Evol. Biol.* 2010, 10, 364. [CrossRef] [PubMed]

20. Lek, A.; Rahimov, F.; Jones, P.L.; Kunkel, L.M. Emerging preclinical animal models for FSHD. *Trends Mol. Med.* 2015, 21, 295–306. [CrossRef] [PubMed]

21. Walter, M.C.; Lochmuller, H.; Reilich, P.; Klopstock, T.; Huber, R.; Hartard, M.; Hennig, M.; Pongratz, D.; Muller-Felber, W. Creatine monohydrate in muscular dystrophies: A double-blind, placebo-controlled clinical study. *Neurology* 2000, 54, 1848–1850. [CrossRef]

22. Kissel, J.T.; McDermott, M.P.; Mendell, J.R.; King, W.M.; Pandya, S.; Griggs, R.C.; Tawil, R.; Group, F.-D. Randomized, double-blind, placebo-controlled trial of albuterol in facioscapulohumeral dystrophy. *Neurology* 2001, 57, 1434–1440. [CrossRef]

23. van der Kooi, E.L.; de Greef, J.C.; Wohlgemuth, M.; Frants, R.R.; van Asseldonk, R.J.; Blom, H.J.; van Engelen, B.G.; van der Maarel, S.M.; Padberg, G.W. No effect of folic acid and methionine supplementation on D4Z4 methylation in patients with facioscapulohumeral muscular dystrophy. *Neuromuscul. Disord.* 2006, 16, 766–769. [CrossRef]
32. van der Kooi, E.L.; Kalkman, J.S.; Lindeman, J.C.; van Engelen, B.G.; Bleijenberg, G.; Padberg, G.W. Effects of training and albuterol on pain and fatigue in facioscapulohumeral muscular dystrophy. J. Neuro. 2007, 254, 931–940. [CrossRef]
33. Wagner, K.R.; Fleckenstein, J.L.; Amato, A.A.; Barohn, R.J.; Bushby, K.; Escolar, D.M.; Flanigan, K.M.; Pestronk, A.; Tawil, R.; Wolfe, G.I.; et al. A phase I/II trial of MYO-029 in adult subjects with muscular dystrophy. Ann. Neurol. 2008, 63, 561–571. [CrossRef] [PubMed]
34. Glasser, C.E.; Gartner, M.R.; Wilson, D.; Miller, B.; Sherman, M.L.; Attie, K.M. Locally acting ACE-083 increases muscle volume in healthy volunteers. Muscle Nerve 2018, 57, 921–926. [CrossRef] [PubMed]
35. Passerier, E.; Hayot, M.; Jaussent, A.; Carnac, G.; Gouzi, F.; Pillard, F.; Picot, M.C.; Bocker, K.; Hugon, G.; Pincemail, J.; et al. Effects of vitamin C, vitamin E, zinc gluconate, and selenomethionine supplementation on muscle function and oxidative stress biomarkers in patients with facioscapulohumeral dystrophy: A double-blind randomized controlled clinical trial. Free Radic. Biol. Med. 2015, 81, 158–169. [CrossRef] [PubMed]
36. Statland, J.M.; Campbell, B.; Desai, U.; Karam, D.; Diaz-Manara, J.; Guptill, J.T.; Korngut, L.; Genge, A.; Tawil, R.N.; Elman, L.; et al. Randomized Phase 2 Study of ACE-083, a Muscle-Promoting Agent, in Facioscapulohumeral Muscular Dystrophy. Muscle Nerve, 2022, in press. [CrossRef] [PubMed]
37. Mellion, M.L.; Ronco, L.; Berends, C.L.; Pagan, L.; Brooks, S.; van Esdonk, M.J.; van Brummelen, E.M.J.; Odueyungbo, A.; Thompson, L.A.; Hage, M.; et al. Phase 1 clinical trial of losmapimod in facioscapulohumeral dystrophy: Safety, tolerability, pharmacokinetics, and target engagement. Br. J. Clin. Pharmacol. 2021, 87, 4658–4669. [CrossRef]
38. Pascal-Gilbert, M.; Lopez-Castel, A.; Arteo, R. Myotonic dystrophy type 1 drug development: A pipeline toward the market. Drug Discov. Today. 2021, 26, 1765–1772. [CrossRef]
39. Le Gall, L.; Sidauskait, E.; Mariot, V.; Dumonceaux, J. Therapeutic Strategies Targeting DUX4 in FSHD. J. Clin. Med. 2020, 9, 2886. [CrossRef]
40. Wang, L.H.; Tawil, R. Current Therapeutic Approaches in FSHD. J. Neuromuscul. Dis. 2021, 8, 441–451. [CrossRef]
41. Cohen, J.; DeSimone, A.; Lek, M.; Lek, A. Therapeutic Approaches in Facioscapulohumeral Muscular Dystrophy. Trends Mol. Med. 2021, 27, 123–137. [CrossRef]
42. Campbell, A.E.; Oliva, J.; Yates, M.P.; Zhong, J.W.; Shadle, S.C.; Snider, L.; Singh, N.; Tai, S.; Hiramuki, Y.; Tawil, R.; et al. BET bromodomain inhibitors and agonists of the beta-2 adrenergic receptor identified in screens for compounds that inhibit DUX4 expression in FSHD muscle cells. Skelet. Muscle 2017, 7, 16. [CrossRef]
43. Cruz, J.M., Jr.; Hupper, N.; Wilson, L.S.; Concannon, J.B.; Wang, Y.; Oberhauser, B.; Patora-Komisarska, K.; Zhang, Y.; Glass, D.J.; Trendelenburg, A.U.; et al. Protein kinase A activation inhibits DUX4 gene expression in myotubes from patients with facioscapulohumeral muscular dystrophy. J. Biol. Chem. 2018, 293, 11837–11849. [CrossRef] [PubMed]
44. Rojas, L.A.; Valentine, E.; Accorsi, A.; Maglio, J.; Shen, N.; Robertson, A.; Kazmirski, S.; Rahl, P.; Tawil, R.; Cadavid, D.; et al. P38 alpha Regulates Expression of DUX4 in a Model of Facioscapulohumeral Muscular Dystrophy. J. Pharmacol. Exp. Ther. 2020, 374, 489–498. [CrossRef] [PubMed]
45. Oliva, J.; Galasinski, S.; Richey, A.; Campbell, A.E.; Meyers, M.J.; Modi, N.; Zhong, J.W.; Tawil, R.; Tapscott, S.J.; Sverdrup, F.M. Clinically Advanced p38 Inhibitors Suppress DUX4 Expression in Cellular and Animal Models of Facioscapulohumeral Muscular Dystrophy. J. Pharmacol. Exp. Ther. 2019, 370, 219–230. [CrossRef] [PubMed]
46. Block, G.J.; Narayan, D.; Amell, A.M.; Petek, L.M.; Davidson, K.C.; Bird, T.D.; Tawil, R.; Moon, R.T.; Miller, D.G. Wnt/beta-catenin signaling suppresses DUX4 expression and prevents apoptosis of FSHD muscle cells. Hum. Mol. Genet. 2013, 22, 4661–4672. [CrossRef]
47. Himeda, C.L.; Jones, T.I.; Virbasius, C.M.; Zhu, L.J.; Green, M.R.; Jones, P.L. Identification of Epigenetic Regulators of DUX4-fl for Targeted Therapy of Facioscapulohumeral Muscular Dystrophy. Mol. Ther. 2018, 26, 1797–1807. [CrossRef]
48. Campbell, A.E.; Shadle, S.C.; Jagannathan, S.; Lim, J.W.; Resnick, R.; Tawil, R.; van der Maarel, S.; Tapscott, S.J.; Sverdrup, F.M. Protein kinase A inhibition induces DUX4 expression in myoblasts of FSHD patients. J. Cell. Physiol. 2016, 231, 4658–4669. [CrossRef]
49. Sharma, V.; Pandey, S.N.; Khawaja, H.; Brown, K.J.; Hathout, Y.; Chen, Y.W. PARP1 Differentially Interacts with Promoter region of DUX4 Gene in FSHD Myoblasts. J. Genet. Syndr. Gene Ther. 2016, 7, 303. [CrossRef]
50. Bagar, J.; Thijsen, P.E.; Shadle, S.; Strasheijm, K.R.; van der Vliet, P.J.; Krom, Y.D.; van den Boogaard, M.L.; de Jong, A.; Lemmers, R.J.L.F.; Tawil, R.; et al. Increased DUX4 expression during muscle differentiation correlates with decreased SMCHD1 protein levels at D4Z4. Epigenetics 2015, 10, 1133–1142. [CrossRef]
51. Lek, A.; Zhang, Y.; Woodman, K.G.; Huang, S.; DeSimone, A.M.; Cohen, J.; Ho, V.; Conner, J.; Mead, L.; Kodani, A.; et al. Applying genome-wide CRISPR-Cas9 screens for therapeutic discovery in facioscapulohumeral muscular dystrophy. Sci. Transl. Med. 2020, 12, eay0271. [CrossRef]
52. Choi, S.H.; Bosnakovski, D.; Strasser, J.M.; Toso, E.A.; Walters, M.A.; Kyba, M. Transcriptional Inhibitors Identified in a 160,000-Compound Small-Molecule DUX4 Viability Screen. J. Biomol. Screen. 2016, 21, 680–688. [CrossRef]
53. Bosnakovski, D.; Choi, S.H.; Strasser, J.M.; Toso, E.A.; Walters, M.A.; Kyba, M. High-throughput screening identifies inhibitors of DUX4-induced myoblast toxicity. Skelet. Muscle 2014, 4, 4. [CrossRef] [PubMed]
54. Bosnakovski, D.; da Silva, M.T.; Sunny, S.T.; Ener, E.T.; Toso, E.A.; Yuan, C.; Cui, Z.; Walters, M.A.; Jadhav, A.; Kyba, M. A novel P300 inhibitor reverses DUX4-mediated global histone H3 hyperacetylation, target gene expression, and cell death. Sci. Adv. 2019, 5, eaaw7781. [CrossRef] [PubMed]
55. Makurvet, F.D. Biologics vs. small molecules: Drug costs and patient access. Med. Drug Discov. 2021, 9, 100075. [CrossRef]
56. Keren, A.; Tamir, Y.; Bengal, E. The p38 MAPK signaling pathway: A major regulator of skeletal muscle development. *Mol. Cell. Endocrinol.* 2006, 252, 224–230. [CrossRef]

57. Wissing, E.R.; Boyer, G.K.; Kwong, J.Q.; Sargent, M.A.; Karch, J.; McNally, E.M.; Otsu, K.; Molkentin, J.D. P38alpha MAPK underlies muscular dystrophy and myofiber death through a Bax-dependent mechanism. *Hum. Mol. Genet.* 2014, 23, 5452–5463. [CrossRef]

58. Segales, J.; Perdigueró, E.; Munoz-Canoves, P. Regulation of Muscle Stem Cell Functions: A Focus on the p38 MAPK Signaling Pathway. *Front. Cell Dev. Biol.* 2016, 4, 91. [CrossRef]

59. Rask-Andersen, M.; Zhang, J.; Fabbro, D.; Schioth, H.B. Advances in kinase targeting: Current clinical use and clinical trials. *Trends Pharmacol. Sci.* 2014, 35, 604–620. [CrossRef]

60. Vanderplanck, C.; Ansseau, E.; Charron, S.; Stricоварт, N.; Tassin, A.; Laurud-Chenivesse, D.; Wilton, S.D.; Coppee, F.; Belayew, A. The FSHD Atrophic Myotube Phenotype Is Caused by DUX4 Expression. *PLoS ONE* 2011, 6, e26820. [CrossRef]

61. Ansseau, E.; Vanderplanck, C.; Wauters, A.; Harper, S.Q.; Coppee, F.; Belayew, A. Antisense Oligonucleotides Used to Target the DUX4 mRNA as Therapeutic Approaches in Facioscapulohumeral Muscular Dystrophy (FSHD). *Genes* 2017, 8, 93. [CrossRef]

62. Lim, J.W.; Snider, L.; Yao, Z.; Tawil, R.; Van Der Maarel, S.M.; Rigo, F.; Bennett, C.F.; Filipova, G.N.; Tapscott, S.J. DICER/AGO-dependent epigenetic silencing of D4Z4 repeats enhanced by exogenous siRNA suggests mechanisms and therapies for FSHD. *Hum. Mol. Genet.* 2015, 24, 4817–4828. [CrossRef]

63. Marsollier, A.C.; Ciszewski, L.; Mariot, V.; Popplewell, L.; Voit, T.; Dickson, G.; Dumonceaux, J. Antisense targeting of 3′ end elements involved in DUX4 mRNA processing is an efficient therapeutic strategy for facioscapulohumeral dystrophy: A new gene-silencing approach. *Hum. Mol. Genet.* 2016, 25, 1468–1478. [CrossRef] [PubMed]

64. Chen, J.C.; King, O.D.; Zhang, Y.; Clayton, N.P.; Spencer, C.; Wentworth, B.M.; Emerson, C.P., Jr.; Wagner, K.R. Morpholino-mediated Knockdown of DUX4 Toward Facioscapulohumeral Muscular Dystrophy Therapies. *Mol. Ther.* 2016, 24, 1405–1411. [CrossRef] [PubMed]

65. Lim, K.R.Q.; Maruyama, R.; Echigoya, Y.; Nguyen, Q.; Zhang, A.; Khawaja, H.; Sen Chandra, S.; Jones, T.; Jones, P.; Chen, Y.W.; et al. Inhibition of DUX4 expression with antisense LNA gapmers as a therapy for facioscapulohumeral muscular dystrophy. *Proc. Natl. Acad. Sci. USA* 2020, 117, 16509–16515. [CrossRef]

66. Lim, K.R.Q.; Bittel, A.; Maruyama, R.; Echigoya, Y.; Nguyen, Q.; Huang, Y.; Dzierlega, K.; Zhang, A.; Chen, Y.W.; Yokota, T. DUX4 Transcript Knockdown with Antisense 2′-O-Methoxylthymidine Gapmers for the Treatment of Facioscapulohumeral Muscular Dystrophy. *Mol. Ther.* 2021, 29, 848–856. [CrossRef] [PubMed]

67. Lu-Nguyen, N.; Malerba, A.; Pineda, M.A.; Dickson, G.; Popplewell, L.J. Improving molecular and histopathology in diaphragm muscle of the double transgenic ACTA1-MCM/FLEExDUX4 mouse model of FSHD with systemic antisense therapy. *Hum. Gene Ther.* 2022; ahead of print. [CrossRef]

68. Lu-Nguyen, N.; Malerba, A.; Herath, S.; Dickson, G.; Popplewell, L. Systemic antisense therapeutics inhibiting DUX4 expression ameliorates FSHD-like pathology in an FSHD mouse model. *Hum. Mol. Genet.* 2021, 30, 1398–1412. [CrossRef]

69. Wallace, L.M.; Liu, J.; Domire, J.S.; Garwick-Coppens, S.; Guckes, S.M.; Mendell, J.R.; Flanigan, K.M.; Harper, S.Q. RNA interference inhibits DUX4-induced muscle toxicity in vivo: Implications for a targeted FSHD therapy. *Mol. Ther.* 2012, 20, 1417–1423. [CrossRef] [PubMed]

70. Wallace, L.M.; Saad, N.Y.; Pyne, N.K.; Fowler, A.M.; Eidahl, J.O.; Domire, J.S.; Griffin, D.A.; Herman, A.C.; Sahenk, Z.; Rodinoklucap, L.R.; et al. Pre-clinical Safety and Off-Target Studies to Support Translation of AAV-Mediated RNAi Therapy for FSHD. *Mol. Ther. Methods Clin. Dev.* 2018, 8, 120–131. [CrossRef] [PubMed]

71. Rashnoneyad, A.; Amini-Chermahini, G.; Taylor, N.K.; Wein, N.; Harper, S.Q. Designed U7 snRNAs inhibit DUX4 expression and improve FSHD-associated outcomes in DUX4 overexpressing cells and FSHD patient myotubes. *Mol. Ther. Nucleic Acids* 2021, 23, 476–486. [CrossRef] [PubMed]

72. Klingler, C.; Ashley, J.; Shi, K.; Stiefven, A.; Kyba, M.; Sinnenreich, M.; Aihar, H.; Kinter, J. DNA aptamers against the DUX4 protein reveal novel therapeutic implications for FSHD. *FASEB J.* 2020, 34, 4573–4590. [CrossRef]

73. Mariot, V.; Joubert, R.; Marsollier, A.C.; Houdré, C.; Voit, T.; Dumonceaux, J. A Deoxyribonucleic Acid Decoy Trapping DUX4 for the Treatment of Facioscapulohumeral Muscular Dystrophy. *Mol. Ther. Nucleic Acids* 2020, 22, 1191–1199. [CrossRef] [PubMed]

74. Grimm, D.; Streetz, K.L.; Jopling, C.L.; Storm, T.A.; Pandey, K.; Davis, C.R.; Marion, P.; Salazar, F.; Kay, M.A. Fatality in mice due to oversaturation of cellular microRNA/short hairpin RNA pathways. *Nature* 2006, 441, 537–541. [CrossRef] [PubMed]

75. Liang, X.H.; Nichols, J.G.; De Hoyos, C.L.; Crooke, S.T. Some ASOs that bind in the coding region of mRNAs and induce RNAse H1 cleavage can cause increases in the pre-mRNAs that may blunt total activity. *Nucleic Acids Res.* 2020, 48, 9840–9858. [CrossRef] [PubMed]

76. Roberts, T.C.; Langer, R.; Wood, M.J.A. Advances in oligonucleotide drug delivery. *Nat. Rev. Drug Discov.* 2020, 19, 673–694. [CrossRef]

77. Joubert, R.; Mariot, V.; Charpentier, M.; Concordet, J.P.; Dumonceaux, J. Gene Editing Targeting the DUX4 Polyadenylation Signal: A Therapy for FSHD? *J. Pers. Med.* 2020, 11, 7. [CrossRef]

78. Das, S.; Chadwick, B.P. CRISPR mediated targeting of DUX4 distal regulatory element represses DUX4 target genes dysregulated in Facioscapulohumeral muscular dystrophy. *Sci. Rep.* 2021, 11, 12598. [CrossRef]
83. Himeda, C.L.; Jones, T.I.; Jones, P.L. Targeted epigenetic repression by CRISPR/dSaCas9 suppresses pathogenic DUX4-fl expression in FSHD. Mol. Ther. Nucleic Acids 2021, 20, 298–311. [CrossRef] [PubMed]

84. Mehta, A.; Merkel, O.M. Immunogenicity of Cas9 Protein. J. Pharm. Sci. 2020, 109, 62–67. [CrossRef] [PubMed]

85. Gillimore, J.D.; Gane, E.; Taubel, J.; Kao, J.; Fontana, M.; Maitland, M.L.; Seitzer, J.; O’Connell, D.; Walsh, K.R.; Wood, K.; et al. CRISPR-Cas9 In Vivo Gene Editing for Translhyretin Amyloidosis. N. Engl. J. Med. 2021, 385, 493–502. [CrossRef] [PubMed]

86. Fardost, S.R.; Ewaisha, R.; Moghadam, F.; Krishna, S.; Park, J.G.; Ebrahimkhani, M.R.; Kiani, S.; Anderson, K.S. Multifunctional CRISPR-Cas9 with engineered immunosilenced human T cell epitopes. Nat. Commun. 2019, 10, 1842. [CrossRef]

87. Brown, B.D.; Venneri, M.A.; Zingale, A.; Sergi, L.; Naldini, L. Endogenous microRNA regulation suppresses transgene expression in hematopoietic lineages and enables gene stable transfer. Nat. Med. 2006, 12, 585–591. [CrossRef]

88. Muhuri, M.; Zhan, W.; Maeda, Y.; Li, J.; Lotun, A.; Chen, J.; Sylvia, K.; Dasgupta, I.; Arjomandnejad, M.; Nixon, T.; et al. Novel Combinatorial MicroRNA-Binding Sites in AAV Vectors Synergistically Diminish Antigen Presentation and Transgene Immunity for Efficient and Stable Transduction. Front. Immunol. 2021, 12, 674242. [CrossRef]

89. Kantor, A.; McClements, M.E.; MacLaren, R.E. CRISPR-Cas9 DNA Base-Editing and Prime-Editing. Int. J. Mol. Sci. 2020, 21, 6240. [CrossRef]

90. Sakellariou, P.; O’Neill, A.; Mueller, A.L.; Ladler, G.; Wright, W.E.; Roche, J.S.; Bloch, R.J. Neuromuscular electrical stimulation promotes development in mice of mature human muscle from immortalized human myoblasts. Skelet. Muscle 2016, 6, 14. [CrossRef]

91. Mueller, A.L.; O’Neill, A.; Jones, T.I.; Llach, A.; Rojas, L.A.; Sakellariou, P.; Ladler, G.; Wright, W.E.; Eyerman, D.; Jones, P.L.; et al. Muscle xenografts reproduce key molecular features of facioscapulohumeral muscular dystrophy. Exp. Neurol. 2019, 320, 113011. [CrossRef]

92. Hareendran, S.; Balakrishnan, B.; Sen, D.; Kumar, S.; Srivastava, A.; Jayandharan, G.R. Adeno-associated virus (AAV) vectors in gene therapy: Immune challenges and strategies to circumvent them. Rev. Med. Virol. 2013, 23, 399–413. [CrossRef]

93. Weinmann, J.; Weis, S.; Sippel, J.; Tulalamba, W.; Remes, A.; Chenuaud, P.; Schmidt, M.; von Kalle, C.; Rolling, F.; et al. Intronic SMCHD1 variants in FSHD: Testing the potential for CRISPR-Cas9 genome editing. J. Med. Genet. 2019, 56, 828–837. [CrossRef]

94. Tabebordbar, M.; Lagerborg, K.A.; et al. Directed evolution of a family of AAV capsid variants enabling potent muscle-directed gene delivery across species. Cell 2021, 184, 4919–4938.e22. [CrossRef]

95. Faust, S.M.; Bell, P.; Cutler, B.J.; Ashley, S.N.; Zhu, Y.; Rabinowitz, J.E.; Wilson, J.M. CpG-depleted adeno-associated virus vectors evade immune detection. J. Clin. Investig. 2013, 123, 2994–3001. [CrossRef]

96. Chan, Y.K.; Wang, S.K.; Chu, C.J.; Copland, D.A.; Letizia, A.J.; Verdera, H.C.; Chiang, J.J.; Sethi, M.; Wang, M.K.; Neidermyer, W.J.; Jr.; et al. Engineering adeno-associated viral vectors to evade innate immune and inflammatory responses. Sci. Transl. Med. 2021, 13, eabd3438. [CrossRef] [PubMed]

97. Penaud-Budloo, M.; Le Guiner, C.; Nowrouzi, A.; Toromanoff, A.; Chere, Y.; Chenuaud, P.; Schmidt, M.; von Kalle, C.; Rollings, R.; Mouliyer, P.; et al. Adeno-associated virus vector genomes persist as episomal chromatin in primate muscle. J. Virol. 2008, 82, 7875–7885. [CrossRef]

98. Mendell, J.R.; Sahenk, Z.; Lehman, K.; Nease, C.; Lowes, L.P.; Miller, N.F.; Iammarino, M.A.; Alfano, L.N.; Nicholl, A.; Al-Zaidy, S.; et al. Assessment of Systemic Delivery of rAAVrh74.MHCK7.micro-dystrophin in Children With Duchenne Muscular Dystrophy: A Nonrandomized Controlled Trial. JAMA Neurol. 2020, 77, 1122–1131. [CrossRef]

99. Al-Zaidy, S.A.; Kolb, S.J.; Lowes, L.; Alfano, L.N.; Shell, R.; Church, K.R.; Nagendran, S.; Sproule, D.M.; Feltner, D.E.; Wells, C.; et al. AVXS-101 (Onasemnogene Abeparvovec) for SMA1: Comparative Study with a Prospective Natural History Cohort. J. Neuromuscul. Dis. 2019, 6, 307–317. [CrossRef]

100. Bottin, S.; Monteilhet, V.; Veron, P.; Leborgne, C.; Benveniste, O.; Montus, M.F.; Masurier, C. Prevalence of serum IgG and neutralizing factors against adeno-associated virus (AAV) types 1, 2, 5, 6, 8, and 9 in the healthy population: Implications for gene therapy using AAV vectors. Hum. Gene Ther. 2010, 21, 704–712. [CrossRef]

101. Chu, W.S.; Ng, J. Immunomodulation in Administration of rAAV: Preclinical and Clinical Adjuvant Pharmacotherapies. Front. Immunol. 2021, 12, 658038. [CrossRef]
102. Bertin, B.; Veron, P.; Leborgne, C.; Deschamps, J.Y.; Moullec, S.; Fromes, Y.; Collaud, F.; Boutin, S.; Latournerie, V.; van Wittenberghe, L.; et al. Capsid-specific removal of circulating antibodies to adeno-associated virus vectors. Sci. Rep. 2020, 10, 864. [CrossRef]

103. Leborgne, C.; Barbon, E.; Alexander, J.M.; Hanby, H.; Delignat, S.; Cohen, D.M.; Collaud, F.; Murelaetharan, S.; Lupo, D.; Silverberg, J.; et al. IgG-cleaving endopeptidase enables in vivo gene therapy in the presence of anti-AAV neutralizing antibodies. Nat. Med. 2020, 26, 1096–1101. [CrossRef] [PubMed]

104. Lee, G.K.; Maheshri, N.; Kaspar, B.; Schaffer, D.V. PEG conjugation moderately protects adeno-associated viral vectors against antibody neutralization. Biotechnol. Bioeng. 2005, 92, 24–34. [CrossRef] [PubMed]

105. Huttner, N.A.; Giora, A.; Perabo, L.; Edbauer, D.; Kleinschmidt, J.A.; Buning, H.; Hallek, M. Genetic modifications of the adeno-associated virus type 2 capsid reduce the affinity and the neutralizing effects of human serum antibodies. Gene Ther. 2003, 10, 2139–2147. [CrossRef] [PubMed]

106. Mingozzi, F.; High, K.A. Immune responses to AAV vectors: Overcoming barriers to successful gene therapy. Blood 2013, 122, 23–36. [CrossRef]

107. Mingozzi, F.; Maus, M.V.; Hui, D.J.; Sabatino, D.E.; Murphy, S.L.; Rasko, J.E.; Ragni, M.V.; Manno, C.S.; Sommer, J.; Jiang, H.; et al. CD8(+) T-cell responses to adeno-associated virus capsid in humans. Nat. Med. 2007, 13, 419–422. [CrossRef]

108. Mueller, C.; Chulay, J.D.; Trapnell, B.C.; Humphries, M.; Carey, B.; Sandhaus, R.A.; McElvaney, N.G.; Messina, L.; Tang, Q.; Rouhani, F.N.; et al. Human Treg responses allow sustained recombinant adeno-associated virus-mediated transgene expression. J. Clin. Invest. 2013, 123, 5310–5318. [CrossRef]

109. Biswas, M.; Kumar, S.R.P.; Terhorst, C.; Herzog, R.W. Gene Therapy With Regulatory T Cells: A Beneficial Alliance. Front. Immunol. 2018, 9, 554. [CrossRef]

110. Wright, J.F. AAV empty capsids: For better or for worse? Mol. Ther. 2014, 22, 1–2. [CrossRef]

111. Morales, L.; Gambhir, Y.; Bennett, J.; Stedman, H.H. Broader Implications of Progressive Liver Dysfunction and Lethal Sepsis in Two Boys following Systemic High-Dose AAV. Mol. Ther. 2020, 28, 1753–1755. [CrossRef]

112. Wang, D.; Tai, P.W.L.; Gao, G. Adeno-associated virus vector as a platform for gene therapy delivery. Nat. Rev. Drug Discov. 2019, 18, 358–378. [CrossRef]

113. Xie, J.; Mao, Q.; Tai, P.W.L.; He, R.; Ai, J.; Su, Q.; Zhu, Y.; Ma, H.; Li, J.; Gong, S.; et al. Short DNA Hairpins Compromise Recombinant Adeno-Associated Virus Genome Homogeneity. Mol. Ther. 2017, 25, 1363–1374. [CrossRef]

114. Tran, N.T.; Heiner, C.; Weber, K.; Weiand, M.; Kaspar, B.; Schaffer, D.V. PEG conjugation weakly protects adeno-associated viral vectors from antibody neutralization. J. Virol. 2021, 95, e0077321. [CrossRef]

115. Echigoya, Y.; Trieu, N.; Duddy, W.; Moulton, H.M.; Yin, H.; Partridge, T.A.; Hoffman, E.P.; Kornegay, J.N.; Rohret, F.A.; Rogers, C.S.; et al. A Dystrophin Exon-52 Deleted Miniature Pig Model of Duchenne Muscular Dystrophy and Evaluation of Exon Skipping. Int. J. Mol. Sci. 2021, 22, 13065. [CrossRef]

116. Valles, A.; Evers, M.M.; Stan, A.; Sogorb-Gonzalez, M.; Brouwers, C.; Vendrell-Tornero, C.; Acrar-Broekmans, S.; Paerels, L.; Klima, J.; Bohuslavova, B.; et al. Widespread and sustained target engagement in Huntington’s disease minipigs upon intrastriatal microRNA-based gene therapy. Sci. Transl. Med. 2021, 13, eabb8920. [CrossRef] [PubMed]

117. Schachtschneider, K.M.; Schook, L.B.; Meudt, J.J.; Shanmuganayagam, D.; Zoller, J.A.; Haghani, A.; Li, C.Z.; Zhang, J.; Yang, A.; Raj, K.; et al. Epigenetic clock and DNA methylation analysis of porcine models of aging and obesity. Geronescience 2021, 43, 2467–2483. [CrossRef]

118. Swindle, M.M. The development of swine models in drug discovery and development. Future Med. Chem. 2012, 4, 1771–1772. [CrossRef] [PubMed]

119. Swindle, M.M.; Makin, A.; Herron, A.J.; Clubb, F.J.; Frazier, K.S. Swine as models in biomedical research and toxicology testing. Vet. Pathol. 2012, 49, 344–356. [CrossRef]

120. Gutierrez, K.; Dicks, N.; Glanzner, W.G.; Agellon, L.B.; Bordignon, V. Efficacy of the porcine species in biomedical research. Front. Genet. 2015, 6, 293. [CrossRef]

121. Jones, T.I.; Chew, G.L.; Barraza-Flores, P.; Schreier, S.; Ramirez, M.; Wuebbles, R.D.; Burkin, D.J.; Bradley, R.K.; Jones, P.L. Transgenic mice expressing tunable levels of DUX4 develop characteristic facioscapulohumeral muscular dystrophy-like pathophysiology ranging in severity. Skelet. Muscle 2020, 10, 8. [CrossRef]

122. Statland, J.; Donlin-Smith, C.M.; Tapscott, S.J.; van der Maarel, S.; Tawil, R. Multiplex Screen of Serum Biomarkers for FSHD Fetal Human Muscle Biopsies. PLoS ONE 2015, 10, e0116853. [CrossRef] [PubMed]

123. Portilho, D.M.; Alves, M.R.; Kratassiouk, G.; Roche, S.; Magdinier, F.; de Santana, E.C.; Polesskaya, A.; Harel-Bellan, A.; Moully, V.; Savino, W.; et al. miRNA Expression in Control and FSHD Fetal Human Muscle Biopsies. PLoS ONE 2014, 9, e0121397. [CrossRef] [PubMed]
126. Heier, C.R.; Zhang, A.; Nguyen, N.Y.; Tully, C.B.; Panigrahi, A.; Gordish-Dressman, H.; Pandey, S.N.; Guglieri, M.; Ryan, M.M.; Clemens, P.R.; et al. Multi-Omics Identifies Circulating miRNA and Protein Biomarkers for Facioscapulohumeral Dystrophy. *J. Pers. Med.* 2020, 10, 236. [CrossRef]

127. Nunes, A.M.; Ramirez, M.; Jones, T.I.; Jones, P.L. Identification of candidate miRNA biomarkers for facioscapulohumeral muscular dystrophy using DUX4-based mouse models. *Dis. Models Mech.* 2021, 14, dmm049016. [CrossRef]

128. Gros, M.; Nunes, A.M.; Daoudlarian, D.; Pini, J.; Martinuzzi, E.; Barbosa, S.; Ramirez, M.; Puma, A.; Villa, L.; Cavalli, M.; et al. Identification of Serum Interleukin 6 Levels as a Disease Severity Biomarker in Facioscapulohumeral Muscular Dystrophy. *J. Neuromuscul. Dis.* 2022, 9, 83–93. [CrossRef]

129. Ghasemi, M.; Emerson, C.P., Jr.; Hayward, L.J. Outcome Measures in Facioscapulohumeral Muscular Dystrophy Clinical Trials. *Cells* 2022, 11, 687. [CrossRef]

130. Butz, M.; Koch, M.C.; Muller-Felber, W.; Lemmers, R.J.; van der Maarel, S.M.; Schreiber, H. Facioscapulohumeral muscular dystrophy. Phenotype-genotype correlation in patients with borderline D4Z4 repeat numbers. *J. Neurol.* 2003, 250, 932–937. [CrossRef]

131. Salort-Campana, E.; Nguyen, K.; Bernard, R.; Jouve, E.; Sole, G.; Nadaj-Pakleza, A.; Niederhauser, J.; Charles, E.; Ollagnon, E.; Bouhour, F.; et al. Low penetrance in facioscapulohumeral muscular dystrophy type 1 with large pathological D4Z4 alleles: A cross-sectional multicenter study. *Orphanet J. Rare Dis.* 2015, 10, 2. [CrossRef]

132. He, J.J.; Lin, X.D.; Lin, F.; Xu, G.R.; Xu, L.Q.; Hu, W.; Wang, D.N.; Lin, H.X.; Lin, M.T.; Wang, N.; et al. Clinical and genetic features of patients with facial-sparing facioscapulohumeral muscular dystrophy. *Eur. J. Neurol.* 2018, 25, 356–364. [CrossRef]

133. Rodriguez, M.C.; MacDonald, J.R.; Mahoney, D.J.; Parise, G.; Beal, M.F.; Tarnopolsky, M.A. Beneficial effects of creatine, CoQ10, and lipoic acid in mitochondrial disorders. *Muscle Nerve* 2007, 35, 235–242. [CrossRef] [PubMed]

134. Willmann, R.; Lee, J.; Turner, C.; Nagaraju, K.; Aartsma-Rus, A.; Wells, D.J.; Wagner, K.R.; Csimma, C.; Straub, V.; Grounds, M.D.; et al. Improving translatability of preclinical studies for neuromuscular disorders: Lessons from the TREAT-NMD Advisory Committee for Therapeutics (TACT). *Dis. Models Mech.* 2020, 13, dmm042903. [CrossRef] [PubMed]

135. Jones, T.I.; King, O.D.; Himeda, C.L.; Homma, S.; Chen, J.C.; Beermann, M.L.; Yan, C.; Emerson, C.P., Jr.; Miller, J.B.; Wagner, K.R.; et al. Individual epigenetic status of the pathogenic D4Z4 macrosatellite correlates with disease in facioscapulohumeral muscular dystrophy. *Clin. Epigenet.* 2015, 7, 37. [CrossRef] [PubMed]