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Male Reproduction in Spinal Muscular Atrophy (SMA) and the Potential Impact of Oral Survival of Motor Neuron 2 (*SMN2*) Pre-mRNA Splicing Modifiers

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ABSTRACT

Spinal muscular atrophy (SMA) is a neuromuscular disease caused by deletions or mutations in the survival of motor neuron 1 (*SMN1*) gene resulting in reduced levels of SMN protein. SMN protein is produced by cells throughout the body, and evidence suggests that low SMN protein can have systemic implications, including in male reproductive organs. However, a paucity of research exists on this important topic. This article will discuss findings from non-clinical studies on the role of SMN in the male reproductive system; additionally, real-world observational reports of individuals with SMA will be examined. Furthermore, we will review the non-clinical reproductive

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Department of Neurology, The Ohio State University Wexner Medical Center, Columbus, OH, USA findings of risdiplam, a small-molecule *SMN2* splicing modifier approved for the treatment of SMA, which has widespread distribution in both the central nervous system and peripheral organs. Specifically, the available non-clinical evidence of the effect of risdiplam on male reproductive organs and spermatogenesis is examined. Lastly, the article will highlight available capabilities to assess male fertility as well as the advanced reproductive technologies utilized to treat male infertility. This article demonstrates the need for further research to better understand the impacts of SMA on male fertility and reproduction.

Keywords: Male fertility; Spinal muscular atrophy; Survival of motor neuron; Risdiplam

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Key Summary Points

Animal studies have shown that survival of motor neuron (SMN) protein plays a key role in spermatogenesis; however, clinical data on the reproductive system of men with spinal muscular atrophy (SMA) are lacking

Men with Type 1 and 2 SMA have a significant rate of unilateral and bilateral undescended testis

Negative reproductive effects of *SMN2* splicing modifiers on spermatogenesis were stage specific to the postpubertal event of meiosis 1 and were reversible in male animals

These effects are expected to be translatable to men with SMA treated with *SMN2*-splicing modifiers

Advancements in assisted reproductive technologies will aid men with SMA who also have male factor infertility in achieving parenthood

INTRODUCTION

Spinal muscular atrophy (SMA) is a progressive neuromuscular disease that affects individuals with a broad age range and spectrum of disease severity [1]. SMA is caused by reduced levels of survival of motor neuron (SMN) protein due to deletions and/or loss of function mutations in the SMN1 gene [1, 2]. Most humans carry a second gene, SMN2, which has a single-base substitution that can cause exclusion of exon 7 during splicing, leading to the production of SMN Δ 7 [3], a shortened version of SMN protein which is unstable and rapidly degrades [4]. Approved disease-modifying therapies (DMTs) for SMA aim to increase the level of SMN protein by either increasing the amount of functional SMN protein produced by SMN2 or delivering a copy of SMN1 [5-7].

SMN protein is ubiquitously produced by human cells, and reduced levels of SMN protein throughout the body are thought to play a vital role in the disease pathophysiology of SMA [8, 9]. While the effect of reduced SMN protein on motor neurons is well established, SMA is considered to be a systemic disease with widespread implications [10, 11]. Non-neuromuscular phenotypes have been observed in individuals with SMA, including those specific to the cardiovascular, gastrointestinal, metabolic, and reproductive systems [11, 12]. It is important to note that high levels of SMN protein are produced in the male reproductive system of mice [13–15] and humans [16, 17].

With more patients living longer because of the availability of DMTs for SMA, the number of individuals considering family-building options will likely increase [18]. How SMA may affect fertility, particularly in male individuals, is not well understood, nor has it been thoroughly investigated. A variety of genetic and acquired variables, as well as lifestyle and environmental risk factors, may impact male fertility [19]. A recent meta-analysis reported the worldwide prevalence of infertility in individuals of reproductive potential across the general population as ranging from 12.6 to 17.5% [20]. The limited data on the fertility status of men with SMA make it difficult to understand the impact that DMTs may have on the male reproductive system.

The authors assess the available published literature on the potential effects of SMA on male fertility, the non-clinical effects of selected oral *SMN2* splicing modifiers on male fertility, and the tools and technologies available for fertility assessment and treatment.

Compliance with Ethics Guidelines

This article is based on previously conducted studies and does not contain any new studies with human participants or animals performed by any of the authors.

SMA AND MALE FERTILITY

Experimental Studies

Although the relationship between SMA and male fertility in humans is not well established,

animal models suggest that SMA does negatively impact male reproductive competency. In experimental mouse models that exhibit a deficiency in SMN protein, a broad range of negative reproductive consequences have been observed [13, 14, 21]. In the most severe animal models, low SMN expression was linked to impairments in male reproductive organ development (reduced testis size), defective sperm maturation, degenerated seminiferous tubules, and a reduction in sperm count [15]. Reduced male fecundity was observed, with only two out of eight male SMA mice able to successfully sire a litter compared with all wild-type male mice mated [15]. It has been suggested that high expression of SMN protein in the testis is due to its critical role in the development and maintenance of male germ cells, in particular for spermatogonia and the initial stages of sperm development [14, 15].

Experiments conducted in a mouse model of SMA in which mice have a knockout of the smn gene but express a human SMN2 transgene reported higher expression of full-length SMN2 mRNA in testis compared with other tissues. indicating there may be a specific mechanism facilitating the inclusion of exon 7 in SMN2 mRNA in the testis [15, 22]; however, this has not been confirmed in men with SMA. The presence of a specific mechanism for SMN2 splicing in testes might indicate that the SMN2 gene and the SMN protein produced from SMN2 may have a unique and pivotal role in human spermatogenesis. This specific SMN gene is evolutionarily isolated to humans and shows high expression in the testes [16, 17].

In humans, an in vitro analysis of pluripotent stems cells derived from men with the most severe form of infertility, azoospermia (i.e., a sperm count of zero), found low expression of *SMN1* compared with stem cells from healthy controls [21], identifying a link between *SMN1* expression and azoospermia. Expression of SMN protein in pluripotent stem cells derived from patients with azoospermia resulted in upregulation of germ-cell markers and induced differentiation into primordial germ cell-like cells [21]. This in vitro human study provides additional evidence that SMN protein plays an important role in human spermatogenesis.

Observational Studies

An analysis of insurance claims in the US healthcare system has identified a higher prevalence of testicular hypofunction and male infertility in individuals with adult-onset SMA compared with matched controls [23]. Lipnick et al. reported that males living with SMA who were diagnosed between the ages of 21 and 65 (N=196) (suggestive of milder disease) were more likely to have had a diagnosis of testicular hypofunction (odds ratio=2.4, p=0.02) or male infertility (odds ratio=5.1, p=0.01) [23]. This analysis was based upon insurance claims covering a period (January 2008 to October 2015) primarily before the approval of DMTs.

In a recent retrospective study by Ribault et al. from the Lyon Neuromuscular Disease Center, patients with SMA were followed across a period covering the availability of DMTs (2012–2022) [24]. Among seven adult males who attempted to undergo sperm cryopreservation prior to initiating risdiplam therapy, three (43%) had a complete absence of sperm in the ejaculate, a condition termed azoospermia. This extraordinarily high prevalence rate of azoospermia (43%) is far beyond the incidence of 1% for the male general population or the rate of 10% observed in men with infertility [25].

Two real-world observational reports of individuals with SMA have reported high rates of cryptorchidism (undescended testes) in patients with Type 1 or 2 SMA [26, 27]. Brener et al. assessed 27 male patients with Types 1-3 SMA, reporting bilateral cryptorchidism in 60% of patients with Type 1 SMA (n=10) and 30% in patients with Type 2 SMA (n=10), with a mean age at diagnosis of cryptorchidism of 6.4 ± 4.4 (range 1.2–14.3) years [26]. All patients with Type 3 SMA (n=7) had descended testes. In a study of patients with Type 1 SMA, Bach et al. reported bilateral cryptorchidism in 52% (13 of 25) of male patients whose testes were examined, and unilateral cryptorchidism was observed in an additional two male patients [27]. The prevalence of cryptorchidism in males with SMA in observational studies is significantly higher than the incidence (2-4%) reported in the general population for full-term males [28].

Testicular descent during embryonic development requires intra-abdominal muscles to provide adequate pressure to facilitate the migration of the testes from the abdomen into the scrotum [28, 29]. It is suspected that weakness in the intra-abdominal muscles could inhibit or prevent the testes from properly descending, thus leading to cryptorchidism. As muscle weakness is more profound in severe forms of SMA, the risk of cryptorchidism is significantly higher in males with early-onset SMA.

Effects of Cryptorchidism on Fertility

Properly descended testes maintained in a cooler environment within the scrotum (~4 °C below body temperature) are necessary for optimal spermatogenesis [28, 29]. Elevated testes temperatures are associated with male infertility and impaired semen quality [28, 29]. In patients with cryptorchidism, the undescended testes remain near or at body temperature. If the testes do not descend by 6 months of age, then surgical intervention to bring the testis into the scrotum is ideally recommended within 12-18 months [30]. Without surgical correction, children with bilateral cryptorchidism will become permanently sterile. Therefore, in children with uncorrected bilateral cryptorchidism who have become sterile, the impact of any SMA DMT on male fertility may not be relevant. Cryptorchidism is also associated with a significantly increased risk (3.7-7.5 times higher) for testicular cancer [31], and surgical placement of the undescended testis in the scrotum is often performed to preserve testosterone production as well as providing access for the ongoing physical examinations required for long-term cancer screening. As men with SMA are expected to live longer with DMTs, it is imperative that urological surveillance and monitoring of undescended male gonads for testis cancer with advanced imaging be undertaken.

Summary of SMA and Male Fertility

These studies collectively demonstrate the critical nature of SMN protein in the male

reproductive system. In animal studies, low levels of SMN protein in the testis showed a detrimental effect on the development of male organs, a reduction in fertility, and diminution of the spermatogonia germ-cell population [13–15, 21].

In a study of healthcare usage, significantly higher rates of infertility and testicular hypofunction were noted in men with adult-onset SMA where symptoms are usually milder compared with infantile-onset SMA. Furthermore, in a small cohort of seven adult men with SMA who attempted sperm cryopreservation prior to initiating risdiplam therapy, 43% had no sperm in the ejaculate. In more severe forms of SMA, there is also a markedly increased prevalence of cryptorchidism. Collectively, these findings demonstrate the potential for a direct and negative impact of SMA on the male reproductive system, and men with SMA are at a higher risk for reduced fertility.

SMN2 SPLICING MODIFIERS AND MALE FERTILITY

Risdiplam and Male Fertility

Prior to the development of DMTs, treatment of SMA was focused primarily on the management of disease symptoms driven by the loss of motor neurons and improvements in standard of care [32]. Risdiplam is one of three DMTs available for SMA and is approved for the treatment of pediatric and adult patients with SMA [5, 33]. Risdiplam is an oral pre-mRNA splicing modifier that promotes the inclusion of exon 7 in SMN2 mRNA to produce stable SMN protein [34, 35]. As a small molecule, risdiplam was specifically designed to distribute evenly in the body, including the central nervous system. This leads to increased levels of functional SMN protein in tissues throughout the body [36], i.e., in the central nervous system and elsewhere, including the testes. Another small molecule with a similar mechanism of action, known as RG7800, was evaluated in patients with SMA when risdiplam was still in preclinical development [37]. In trials of healthy adults and patients with SMA, RG7800 increased SMN protein levels; however, studies in patients were put on hold because of safety findings in animal toxicology studies [38]. Due to improvements in drug metabolism (suitable half-life and wide tissue distribution), improved in vitro potency on *SMN2* splicing, and a favorable preclinical safety profile, risdiplam was selected for subsequent clinical development [35, 39]. Risdiplam shows high selectivity to two binding sites in exon 7 of *SMN2* pre-mRNA, namely the exonic splicing enhancer 2 and a 5' splice site [39, 40]. The combination of binding to exonic splicing enhancer 2 and the 5' splice site provides risdiplam high selectivity to *SMN2* pre-mRNA [40].

During the drug development process, nonclinical toxicologic findings were reported in male germ cells of animals exposed to risdiplam and RG7800. An analysis of risdiplam, RG7800, and related SMN2 gene splicing compounds identified splicing events in genes other than SMN2 [35] as animals do not have the SMN2 gene. In a few genes, similar exon inclusion events were seen in the mRNA transcripts as with SMN2 in human cells at drug concentrations relevant for the male germ-cell target organ and other tissues with toxicologically relevant findings in animal studies. Although it remains difficult to attribute such tissue-specific effects to any single off target, it was plausible to focus on Forkhead Box M1 (FOXM1) and MAP kinaseactivating death domain protein (MADD) among the affected genes, as the toxicologically relevant features were seen exclusively in proliferating and/or self-renewing tissues [35, 39]. Of the affected genes and their known exon inclusion variants, only FOXM1 and MADD are associated with regulation of the cell cycle and apoptosis [35, 39]. *FOXM1* can be considered of particular relevance to male fertility due to its expression pattern in male reproductive tissues and across a specific stage of spermatogenesis [41, 42].

Effects of *SMN2* Splicing Modifiers on Secondary Splice Targets

The FOXM1 gene produces several splice variants, which are present in animals and humans (see Table 1). These include a transcriptionally inactive FOXM1a variant with exon A2 and transcriptionally active variants FOXM1b and FOXM1c, which lack exon A2 [43-45]. FOXM1 is a transcription factor that regulates genes that control G1/S (interphase of the cell cycle). Thus, the FOXM1b/c isoforms promote the cell cycle, whereas the FOXM1a variant results in a stop of the cell cycle. Depending on the stage of the cell cycle, interference of *FOXM1* or these splice variants can impact mitosis and meiosis. Accordingly, changes in FOXM1 splice variants are expected to interfere with spermatogenesis. An additional splice target, MADD, is likely responsible for the induction of apoptosis observed as cytoplasmic vacuoles because of increased expression of a pro-apoptotic splice variant known as IG20 [35, 46].

Small-molecule *SMN2* pre-mRNA splicing modifiers can interact with the *FOXM1* mRNA transcript by upregulating the *FOXM1a* variant via exon inclusion with concomitant downregulation of the *FOXM1b/c* variants. In vitro experiments of *SMN2* splicing modifiers using mouse cell lines and human cells derived from patients with SMA showed evidence of an increased frequency of micronucleated cells and an increase

Gene	Splice variant	Activity	Potential impact on cells
FOXM1	FOXM1a	Transcriptionally inactive	Stoppage of cell cycle [45]
	FOXM1b	Transcriptional activator	Promote cell cycle activity [45]
	FOXM1c		
MADD	IG20	Pro-apoptotic	Induction of apoptosis [35, 46]

 Table 1 Impact of key secondary splice targets

FOXM1 Forkhead Box M1, MADD MAP kinase-activating death domain protein

in apoptosis (manifested as cells with large cytoplasmic vacuoles potentially due to incomplete apoptotic processes), which likely indicated changes in the expression of *FOXM1* and *MADD* splice variants [35]. Experiments using cells derived from patients with SMA demonstrated splicing impacts in the mRNA transcripts of secondary splice targets in human genes after treatment with *SMN2* splicing modifiers [35].

In non-clinical in vivo experiments of rats and mice with both RG7800 and risdiplam, similar concomitant up- and downregulation of different FOXM1 splice variants were observed which caused mitotic arrest and appeared as micronucleated cells [35]. In monkey studies with RG7800, FOXM1b/c was downregulated in a single monkey at the highest dose; however, no effects were observed at lower doses [47]. In another study, FOXM1b/c was downregulated in five of seven monkeys after treatment with RG7800, and FOXM1a was upregulated in the remaining two monkeys [47]. These changes in the expression of splice variants of this specific cell cycle gene likely resulted in damage to sperm-producing tissues of rats and monkeys in non-clinical experiments with RG7800 and risdiplam [47], with stage-specific and reversible effects as outlined below.

Effects of *SMN2* Splicing Modifiers on Spermatogenesis

SMN2 splicing modifiers can impact spermatogenesis, which is the process where new sperm are produced in the seminiferous tubules of the testes [48, 49]. Spermatogenesis can be divided into three stages: the mitotic stage, where spermatogonia (2n) undergo mitosis to maintain the stem cell pool and differentiate into primary spermatocytes; the meiotic stage (which begins at puberty), where primary spermatocytes (2n) divide into secondary spermatocytes (n); and spermiogenesis, where mature sperm are produced [48]. SMN1 is critical for the preservation of spermatogonia and paramount for prepubertal and postpubertal germ cell survival. In contrast, FOXM1 is highly expressed in spermatocytes during meiosis 1, a process which only occurs postpuberty, where primary spermatocytes differentiate into secondary spermatocytes (Fig. 1). Interactions of *SMN2* splicing modifiers with *FOXM1* during meiosis 1 can interrupt the development of mature sperm and thus negatively impact male fertility [50].

Evidence of Stage-specific Degeneration During Spermatogenesis

A staging study of monkeys exposed to RG7800 was conducted to determine whether treatment affected specific cells and/or specific stages of spermatogenesis [35]. Microscopic analysis of testicular tissue samples showed evidence of stage-specific degeneration in the seminiferous tubules and the presence of micronucleated cells and cytoplasmic vacuoles but mature sperm cells were still observed (Fig. 2) [47]. The presence of these micronucleated cells (and vacuoles) were similar to those described in in vitro experiments reported by Ratni et al. [35]. Testicular degeneration was observed to be specific to germ-cell maturation, and the arrest occurred during stages of spermatogenesis after meiosis 1 (postpuberty), with no impact on spermatogonia observed [35, 47]. More specifically, germ-cell degeneration was observed almost exclusively in spermatocytes during meiosis 1, a stage when FOXM1 is highly expressed, and in tissue where alternative splicing of *FOXM1* was observed [47].

Evidence of Reversibility of *SMN2* **Splicing Modifiers**

The impact of RG7800 and risdiplam on male reproductive tissues was further investigated in non-clinical studies in which animals were exposed to the drug and then assessed after a drug-free recovery period. In non-clinical studies of rats, after a 28- to 56-day recovery period, full reversibility of germ-cell degeneration was observed in half of the rats assessed following exposure to either risdiplam or RG7800 [47]. Consistent with the reversibility of germ-cell degeneration, male rats exposed to risdiplam for 13 weeks and allowed to recover for 8 weeks were able to breed successfully and did not show impaired fertility when paired with non-treated females [47]. Monkeys dosed with RG7800 and



Fig. 1 Proposed mechanism of off-target effects of SMN2 splicing modifiers on spermatogenesis. Spermatogenesis takes place over three stages: the mitotic stage, where spermatogonia (2n) undergo mitosis to maintain the stem cell pool and differentiate into primary spermatocytes; the meiotic stage (which beings during puberty), where primary spermatocytes (2n) divide into secondary spermatocytes

then allowed to recover for at least 55 days did not show any impaired testicular findings and did not demonstrate alternative splicing of *FOXM1* and MADD [47]. No testicular degeneration was observed in a longer-term study where monkeys were exposed with RG7800 for 39 weeks and allowed to recover for 22 weeks [47]. No testicular findings were reported in any studies involving immature prepubertal monkeys [47].

The period of recovery following drug withdrawal occurs over a short timescale relative to the length of the spermatogenic cycle (56 days in rats [51]; 42 days in monkeys [52]). This contrasts with the long-lasting or at times irreversible effects on fertility observed following exposure to cytotoxic agents used in chemotherapy or radiotherapy [53]. The short recovery period observed with *SMN2* splicing modifiers is consistent with the proposed stage-specific

(*n*); and spermiogenesis, where mature sperm are produced. SMN protein is highly expressed in spermatogonia and is essential for normal spermatogenesis. The impact of *SMN2* pre-mRNA splicing modifiers is proposed to impact spermatocytes during meiosis 1 as a result of splicing changes in *FOXM1*. *FOXM1* Forkhead Box M1, *SMN* survival of motor neuron

mechanism of action, which implies minimal or no damage to the primary germ-cell population of spermatogonia. Damage induced by cytotoxic agents, such as chemotherapy or radiotherapy, usually damages all stages of spermatogenesis including spermatogonia and thus results in long-term or permanent damage that can reduce sperm counts, often to azoospermic levels [54]. The time course of damage from cytotoxic agents depends on the cell types impacted, with the spermatogonia stem cells being most sensitive to damage and their loss resulting in the most severe and long-lasting damage [54]. It is important to note that damage to spermatogonia was not evident in any of the non-clinical studies of SMN2 splicing modifiers reported by Mueller et al., including studies in prepubertal monkeys [47].



Fig. 2 Histology of testicular tissue in a non-clinical monkey study. a A schematic of the process of spermatogenesis within the seminiferous tubules. *Image adapted from Allais-Bonnet A and Pailhoux E (2014) Role of the prion protein family in the gonads. Front. Cell Dev. Biol. 2:56.* b Testis sample from a control animal; note a layer of spermatocytes (arrows). c Testis tissue from a cynomolgus monkey dosed at 6 mg/kg/day of RG7800. The seminiferous tubules depict an insult to the specific spermatocyte layer;

vacuoles at a germinal epithelial location compatible with spermatocytes. Note that mature sperm can be observed in the center of the tubule in both images. *Histology images* reproduced from Reproductive Toxicology, Vol 118, Mueller L. et al. Reproductive findings in male animals exposed to selective survival of motor neuron 2 (SMN2) gene splicingmodifying agents. Copyright 2023, with permission from Elsevier. FOXM1 Forkhead Box M1

arrows denote germ cells with abnormal morphology and

Summary of Findings from Non-clinical Animal Studies

These findings provide evidence that the effects of oral *SMN2* pre-mRNA splicing modifiers were reversible in non-clinical animal experiments. As *SMN2* is absent in animals, the effects are attributed to off-target splicing events. The stage-specific nature of the damage to spermatocytes in meiosis 1, with no evidence of damage to spermatogonia, suggests the drug exposure did not impact the spermatogonia stem cell line. Following cessation of drug exposure, the impact on *FOXM1* splicing is removed, and spermatogonial division and sperm maturation appear to resume unimpaired.

ASSESSMENT AND PRESERVATION OF MALE FERTILITY

Treatment and Technological Advancements

With the availability of multiple therapies for SMA, many patients are now living longer and have or will consider family-building opportunities. As the effects of SMA disease progression or *SMN2* splicing modifiers are not fully elucidated, it has been recommended for male patients to consider options to screen and preserve fertility. It is also important to highlight that infertility in the general population has been estimated as high as 17.5% worldwide [20]. This section provides an overview of several advancements, recommendations, and options to assess and optimize fertility, particularly in male patients

who may be on treatment or have experienced infertility.

Infertility is defined by the failure to achieve pregnancy following a 12-month period of regular unprotected sexual intercourse [55], which can be caused by a variety of factors in either the male or female reproductive system. The American Urological Association/American Society for Reproductive Medicine recommends evaluating both partners concurrently per their guidelines for infertility [56]. For males, the American Urological Association/American Society for Reproductive Medicine evaluation incorporates both a male reproductive history and a semen analysis that measures key parameters including semen volume and sperm concentration, motility, and morphology. This initial assessment should guide the physician on determining baseline male fertility status, consider additional testing if needed, and provide the most appropriate and targeted therapy.

Men with male factor infertility now have many options for assessing and managing their fertility [56]. Historically, a standard semen analysis would require the patient to attend a fertility clinic and produce an ejaculate onsite. For many men this activity was psychologically challenging and created a barrier and delay for a proper fertility evaluation [57]. This coupled with financial constraints and lack of adequate access to fertility care can make navigating infertility challenging for couples. Male infertility has traditionally been overlooked and often stigmatized and may be a barrier to individuals accessing available resources [58, 59]. In addition, people with disabilities may experience greater difficulties in accessing reproductive services, and specifically for men with SMA upper limb weakness may limit their ability to produce an ejaculate.

Fortunately, recent advances in fertility telehealth, home male fertility testing, and sperm cryopreservation techniques have emerged, and their utilization and acceptance have dramatically accelerated. It should be noted that there are wide differences in the availability of and access to services both within and between countries. These advances have included convenient technologies to test semen parameters at home which can provide a preliminary general assessment of sperm concentration as well as newer technologies that provide a real-time video and determine motile sperm concentration [60]. Home-collected samples can also be maintained using optimized containers and buffered with media which then enables them to be shipped to a centralized andrology laboratory for a complete semen analysis and possible cryopreservation [61]. These diagnostic advances have been complemented by the broad acceptance and proven benefit of fertility telehealth services. These developments are uniquely applicable to the specific population of men with SMA where mobility and access to fertility care are relevant and emerging issues.

For male patients who find it difficult to produce a semen sample, particularly among men who are experiencing physical weakness, techniques such as vibratory stimulation can be utilized [62]. For patients who are unable to ejaculate or upon semen analysis have no viable or detectable sperm in their semen, numerous surgical sperm retrieval techniques can be used to extract mature sperm which can then be utilized in conjunction with in vitro fertilization (IVF) and intracytoplasmic sperm injection (ICSI) [63].

When appropriate, sperm cryopreservation is considered the gold standard for male fertility preservation [64], and options for semen samples to be collected at home and then transported to centralized facilities are available. Cryopreservation of multiple vials as well as a post-thaw test of the sample is recommended at the time of initial cryopreservation to establish confidence that the sperm will survive future cryo-thaw cycles. Testing for communicable diseases is performed at the time of cryopreservation to enable the sperm to be utilized in the future with assisted reproductive technologies (ARTs). A discussion or meeting between the patient and a fertility specialist is recommended to discuss fertility preservation options and implications as well as review the current quality and viability of the cryopreserved semen samples.

ARTs are the mainstay and have been successfully utilized to treat male factor infertility. These include intrauterine inseminations and IVF with ICSI, which can be used to markedly improve the chances of successful conception. ICSI has become the gold standard for treating male factor infertility during in vitro fertilization as it offers the profound benefit of requiring only a single viable sperm cell to be directly injected under a microscope and using advanced reproductive technologies into every retrieved oocyte [65].

Depending on the resources available to patients and healthcare professionals, in this population of men with SMA, a pre-pregnancy genetic carrier screening panel for both partners and genetic counseling has been recommended for couples considering pregnancy [66]. If both partners have mutations in SMN1 and IVF is utilized, the embryos created can subsequently undergo preimplantation genetic testing for monogenic/single gene defects (PGT-M). Embryos that are homozygous or compound heterozygous for SMN1 mutations and at risk of developing SMA can be screened out [67]. PGT-M for single gene disorders is limited by the incredibly small amount of DNA obtained in an embryo biopsy where the usual techniques for analysis of the SMA gene are not possible. As a result, PGT-M for SMA requires the creation of a custom genetic test using linkage analysis prior to beginning IVF treatment. To establish informative linkage markers, DNA from both the sperm and oocyte contributors and their biological parents (or offspring, if any) is typically required to facilitate test creation. It is now also more common for couples to consider pre-emptively undergoing IVF/ICSI, PGT testing, and embryocryopreservation to optimize current and future family-building opportunities [68, 69].

DISCUSSION

There are robust animal data highlighting the importance of SMN protein for normal spermatogenesis, male reproductive organ development, and male fertility [13–15, 21]. Evidence from animal models of SMA demonstrates abnormalities in male reproduction which affect both sperm development and fertility [15]. There is limited information on how SMA may impact the male reproductive system in humans. Higher rates of male infertility have been reported among healthcare claims from patients with SMA [23],

Effects of SMN2 splicing modifiers on male sperm cell production were identified in nonclinical toxicologic studies in animals [35, 47]. Impacts on spermatogenesis isolated to postpubertal sperm maturation differentiation are proposed to be the result of off-target effects from oral SMN2 splicing modifiers primarily via a cell cycle controlling gene, FOXM1, during spermatogenesis. This insult appears to be cell stage specific to the postpubertal maturation event of meiosis 1 and does not appear to impair the spermatogonia germ-cell line [47]. Consistent with the proposed mechanism of action, these effects were observed to be reversible following cessation of exposure to SMN2 splicing modifiers. Furthermore, in non-clinical experiments, exposure to SMN2 splicing modifiers did not result in a complete arrest of sperm production. Further research is needed to explore if there is a benefit from restoration of and/or increasing the levels of SMN protein in testicular tissues by SMN2 splicing modifiers that can reach the male testis.

Implications for Human Patients

In the absence of any clinical data from human patients, the effects of SMN2 splicing modifiers on fertility in male patients are not fully understood. Due to the conserved nature of these secondary splice targets between species, the testicular impacts and the reversible nature observed from the animal studies would be expected to be translatable to humans [47]. It is important to note these effects of risdiplam are relevant to post-pubescent males with SMA, as the observed impairments in spermatogenesis are specific to postpubertal meiosis 1. Although these data suggest normal fertility function might be restored within 4 months (encompassing the length of the human sperm cycle, drug transit time, and six half-lives of the drug), the US Prescribing Information and European Summary of Product Characteristics contain recommendations that male patients with SMA who desire to father a child may consider sperm preservation before commencing risdiplam treatment [5, 33]. Physicians should be aware of these recommendations and refer to the following section for a discussion of procedures that can assess or preserve sperm in patients with SMA. Although the full impact of *SMN2* splicing modifiers on the male reproductive system in humans is unknown, at the time of this writing the authors have been made aware of three reports of men with SMA who have conceived while treated with risdiplam.

CONCLUSIONS

Current understanding of how SMN protein impacts the male reproductive system in individuals with SMA is limited by the lack of research into this aspect of the disease; however, there are robust animal data demonstrating the importance of SMN protein in spermatogenesis and testicular function. Evidence on the impact of SMA on male fertility in humans primarily comes from observational reports that did not seek to identify the underlying mechanistic reasons. Yet, some additional credible mechanistic/ experimental evidence with human male germ cells in vitro is available to attribute a significant role of SMN protein in human spermatogenesis.

Male patients of reproductive age should be counseled about the potential effects of treatment and may consider sperm preservation prior to starting treatment or after a sufficient treatment-free period of 4 months [33]. With increased access to male fertility testing via telehealth and home-collection capabilities, there are more options for men with SMA to explore their fertility status. Significant advances in assisted reproductive technologies such as IVF, ICSI, and PGT-M will enable many men with SMA and male factor infertility to achieve their hope of parenthood.

Further research is needed to better understand the effects of SMA on the human male reproductive system and male fertility and the potential impact resulting from available DMTs.

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Declarations

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Ethical Approval. This article is based on previously conducted studies and does not

contain any new studies with human participants or animals performed by any of the authors.

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