

# Rett Syndrome: Reaching for Clinical Trials

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**Abstract** Rett syndrome (RTT) is a syndromic autism spectrum disorder caused by loss-of-function mutations in *MECP2*. The methyl CpG binding protein 2 binds methylcytosine and 5-hydroxymethylcytosine at CpG sites in promoter regions of target genes, controlling their transcription by recruiting co-repressors and co-activators. Several pre-clinical studies in mouse models have identified rational molecular targets for drug therapies aimed at correcting the underlying neural dysfunction. These targeted therapies are increasingly translating into human clinical trials. In this review, we present an overview of RTT and describe the current state of preclinical studies in methyl CpG binding protein 2-based mouse models, as well as current clinical trials in individuals with RTT.

**Keywords** Rett syndrome · Rare disease · X-linked dominant · *MECP2* · Clinical trials · Drug targets

## Background

Rett syndrome (RTT) was first noted by Andreas Rett in Vienna, Austria, and virtually simultaneously by Bengt Hagberg in Göteborg, Sweden, about 50 years ago [1]. However, it did not become widely known until the report of Hagberg et al. in

1983 [2]. Thereafter, general awareness expanded rapidly through the 1980s, spurred by symposia in Vienna under the guidance of Andreas Rett, and meetings organized through the International Rett Syndrome Association in the USA. Diagnostic criteria were generated and continuously updated [3–7], clinical studies were organized, and a search for possible causation was initiated.

Neuropathologic examinations confirmed the clinical finding of reduced brain growth, and identified specific cellular abnormalities, including reduced neuronal size and simplified dendritic arborizations. These also failed to identify any evidence of progressive neural pathologies that would indicate a neurodegenerative condition [8–10]. Instead, RTT had all the features of a neurodevelopmental disorder. Owing to the overwhelming predominance of RTT among females, a genetic etiology was proposed, directed at the X chromosome [11]. Eventually, attention was focused on the Xq28 locus [12]. In 1999, Amir et al. [13] described mutations in *MECP2*, the gene encoding methyl CpG binding protein 2 (MeCP2). The MeCP2 protein binds methylcytosine and 5-hydroxymethylcytosine at CpG sites in promoter regions of target genes, controlling their transcription by recruiting co-repressors and co-activators. *Mecp2* knockout mouse models demonstrated that MeCP2 is critical for the maturation and maintenance of neurons and glial cells [14, 15]. Thereafter, *Mecp2* knockout and knockin mice carrying mutations observed in RTT individuals allowed fundamental research to advance our understanding of the molecular, cellular, and systems-level pathology [16], and to begin addressing possible therapeutic interventions. These strategies have proved disease modifying, and have even resulted in full reversal of the underlying abnormalities [17]. At the same time, human clinical studies expanded rapidly worldwide to obtain natural history information that is essential for the design of appropriate clinical trials [18–21] (see also <https://www.clinicaltrials.gov/ct2/show/NCT00299312>). This combination of fundamental and preclinical research in mouse models with

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human clinical research has fostered a vibrant level of translational and clinical trial planning and execution that is described in detail in the following sections.

## Clinical Features

RTT is a unique neurodevelopmental disorder with an incidence of about 1 : 10,000 female births [22], and multiple systemic issues appearing first in young females after a period of apparently normal postnatal development. Detailed evaluation of the first 6 months of life established that developmental skills do not follow a typical timeline when viewed critically [23–25]. These findings of delayed or arrested developmental are sufficiently subtle that specific delays are not often suspected until the second year of life when frank regression, accompanied in some by autistic-like social avoidance, is noted. Other clues may also be present, including abnormal deceleration of head circumference increase (a barometer of appropriate brain growth; [26–28]), reduction in muscle tone, and an overall appearance of being too quiet or “good”. This is punctuated in some individuals by periods of inappropriate and unexplained screaming spells. The most critical feature is the appearance of stereotypic movements during wakefulness, most prominent in the hands, but also seen in oro-motor regions and the lower extremities.

The initial period of regression is followed by a period of stabilization beginning by 30 months with improved eye contact and socialization that remains throughout life [29]. However, motor development may transition as muscle tone begins to increase, first with a period of apparently normal tone transitioning ultimately to an increase in tone or rigidity. This may be coupled with the appearance of dystonic positions, particularly at the ankles and wrists. Multiple neurologic or systemic issues often arise: 1) epilepsy beginning typically by the third year of life [30–32]; 2) periodic breathing consisting of breath-holding or hyperventilation or both [33–35]; 3) prominent gastrointestinal (GI) issues from top to bottom, including poor chewing and swallowing, gastroesophageal reflux, delayed stomach emptying, and constipation [36–39]; 4) decline in growth parameters in most, including height, weight, hands, and feet [28, 40]; 5) scoliosis in most with up to 13% requiring surgical correction of the deformity [41]; 6) intolerance of warm temperatures; and 7) cool hands and feet. Ambulation is achieved initially in as many as 80% but is typically broad-based and nonpurposeful. It is often accompanied by retropulsion (first steps are backwards), and may include prominent toe walking. Approximately one-third of this group will stop walking. Overall, about 50% maintain independent gait and another 20% ambulate with some level of assistance.

## Genetics

*MECP2* mutations were identified in 1999 as causative for RTT [13]. Presently, up to 96% of individuals in the US Natural History Study (NHS) have such mutations [42, 43]. Nevertheless, a *MECP2* mutation does not indicate the clinical diagnosis of RTT. Individuals with mutations may not have features of RTT syndrome. Conversely, up to 4% have RTT but do not have a *MECP2* mutation [43]. More than 200 pathologic mutations have been identified. However, 8 point mutations (4 missense and 4 nonsense) account for nearly 60% of all affected individuals. Deletions and insertions account for nearly 20%, such that a large number of the remaining mutations occur privately or in only a few individuals. Among the most common mutations, definite phenotype–genotype correlations have been established [42–44]. Based on the Clinical Severity Score employed in the NHS, individuals with R133C, R294X, R306C, and 3′-truncations are significantly less involved than those with the 5 other common point mutations and with deletions of whole exons. However, those with exactly the same mutation may have widely differing clinical abilities. This is thought to represent differences in X-chromosome inactivation (XCI) and overall genetic background. One factor that has not received major attention is the clonal distribution of normal and mutant X chromosomes in brain as differences in expression patterns may lead to markedly different clinical phenotypes. LaSalle et al. noted this in 2001 [45], but scarce mention has been made in recent years. Attempts at identifying variations in XCI in blood have indicated skewing in > 25% of RTT individuals, which is approximately double the occurrence in the general population. Highly skewed (>90%) XCI was noted in > 10%, more than twice that in the general population [unpublished data from the US NHS (NCT00299312)]; in this group, the skewing was roughly equal in favor of the mutant or normal chromosome. These data should be interpreted cautiously, as XCI in brain may differ from that in blood. In addition, environmental background may be critically important to the overall level of clinical involvement.

As an X-linked dominant disorder, RTT is thought to arise from spontaneous or *de novo* mutations in rapidly dividing germinal cells. As such, the majority of *de novo* mutations occur in the sperm [46, 47]. In some instances, the mutation is transmitted by a female carrying a specific mutation but, owing to favorable XCI, she is unaffected or has mild learning disability or cognitive impairment [48, 49]. In offspring resulting from affected women, a girl may have classic or typical RTT, but boys are likely to have much greater involvement resulting in infantile encephalopathy and a very much shortened life expectancy (~1 year) owing mainly to failure to maintain normal respiratory function [50]. In other instances, mutations in the 3′-truncation region in boys produce a progressive dystonia against a background of severe cognitive

delay [49]. However, in families that do not have an affected female, recognition of such features in a male may not raise suspicion of a *MECP2* mutation and hence go undetected.

Males with seemingly typical RTT have been reported in a small number under 2 scenarios: one being co-occurrence of Klinefelter syndrome (47 XXY) [51–54] and the other being somatic mosaicism [55]. In each instance, 2 populations of cells exist: 1 with a normal and 1 with a mutant X chromosome.

More recently, mutations in other genes have been associated with atypical RTT. These include children with mutations in *CDKL5*, *FOXG1*, and other associated genes [56–60].

## Therapeutic Targets for Clinical Trials

At present, clinical trials are limited to RTT individuals with *MECP2* mutations. In addition, careful consideration is given to establish inclusion and exclusion criteria in order to ensure similar groups in the design of placebo-controlled, double-blind studies. Suitable end points are also essential. RTT offers several highly relevant and important clinical end points for which suitable outcome measures exist [61]. These include communication, epilepsy, motor function, GI parameters, and periodic breathing, either breath holding or hyperventilation. Current studies using eye-tracking technology coupled with evoked response potentials are leading the way to assess communication. Video-electroencephalography recordings can provide both spectral analysis of the various waveforms and time-locked assessments of epileptiform features. Coupled with plethysmographic recordings, periodic breathing can also be captured to provide evidence of altered breathing patterns. For those individuals who are ambulatory or have improvement in walking capabilities, gait analysis paradigms provide objective evidence of motor performance. Improvements in muscle tone or dystonia can also be assessed objectively. Actigraphy provides an excellent measure of both purposeful and stereotypic movements. Evaluation of GI function is more varied. Potentially useful targets are objective improvements in chewing and swallowing and better GI motility in terms of improved gastric emptying and reduced constipation.

In terms of effective clinical outcome measures, efforts are ongoing to improve the acute responsiveness of clinical severity measures, communication assessment, and neurophysiologic parameters. What is lacking is an appropriate behavioral measure. This deficiency is actively being pursued. Quality of life (QoL) using standardized instruments (not specific to RTT) has been addressed in both those with RTT [62] and their caregivers (work in progress). QoL for those with RTT has a distinct pattern of involvement. Individuals with greater clinical severity demonstrate more severe motor impairment, and those with lower clinical severity have fewer motor

impairments. However, those with greater clinical severity have better behavioral function than those with lower clinical severity. This raises the possibility that modest improvement in motor function could have an adverse effect on behavior. QoL has also been assessed in the primary caregiver, generally the mother, and is currently undergoing analysis. In addition, a separate caregiver inventory is being collected and standardized specifically among families managing an individual with RTT (data from NCT00299312).

At present, no relevant laboratory biomarker is known. A metabolomics pilot study has started with the current NHS; however, the results are not anticipated before 2016.

## Repertoire of Clinical Trials

Therapeutic targets for RTT are either aimed at reversing the loss-of-function mutation in *MECP2* or modifying its downstream pathways, including neurotransmitter receptor systems, neurotrophins, and their intracellular signaling pathways (Tables 1 and 2) [63–65]. We follow the standard convention: “preclinical” is used for experiments done in experimental animal models, and “clinical” for tests performed in humans.

## *MECP2* Gene Therapy and “Read-through” Compounds

Preclinical studies have demonstrated that restoring *Mecp2* gene expression in mice improves RTT-like neurological symptoms [17, 66]. However, translating this gene therapy approach to clinical trials requires overcoming multiple challenges, including developing safe vectors that allow effective *in vivo* gene delivery and avoiding overexpression of exogenous *MECP2* in wild-type cells in the mosaic brain of individuals with RTT. Among different gene vectors, the serotype 9 of adeno-associated virus is the most promising owing to its ability to cross the blood–brain barrier (BBB) and to infect neurons efficiently, yielding long-term transgene expression in male *Mecp2* knockout mice and in female *Mecp2* heterozygous mice [67, 68]. It should be noted that no deleterious consequences of the overexpression of *Mecp2* in wild-type cells were noted in the mosaic female brain [69]. At present, there are no clinical trials of *MECP2* gene therapy in individuals with RTT.

Owing to a premature STOP codon, a nonsense mutation (e.g., R168X, R255X) observed in approximately one-third of RTT individuals causes the termination of translation and may destabilize mRNA molecules [70, 71]. These individuals may have a more severe clinical presentation than those with missense *MECP2* mutations that result in single amino acid substitutions, except, perhaps, for the R294X allele. Aminoglycoside antibiotics allow ribosomal read-through of premature

**Table 1** Current clinical trials

Agent	Results	Trial outcome	Sponsor	Type
<b>Growth factors</b>				
IGF-1 (Mecasermin)	Phase II	Safety confirmed Ongoing Scheduled through Fall 2015	Approved drug in children with growth failure	Double-blind, placebo-controlled, injection; patients aged 3–10 years
NNZ-2566	Phase II	Safety confirmed Short-term trial positive	Neuren Pharmaceuticals	Double-blind; placebo-controlled; oral; patients aged 16–45 years
<b>BDNF boosters</b>				
Copaxone (Glatiramer acetate)	Phase II	Improved ambulation	Approved drug	Open-label; injection
Fingolimod	Phase I	Ongoing; scheduled through August 2017	Novartis Pharmaceuticals; approved drug in adults	Open-label; oral
<b>NMDA antagonist</b>				
Dextromethorphan	Phase II	Ongoing; scheduled through June 2015	Approved drug in cough suppressant	Double-blind; placebo-controlled
<b>Norepinephrine reuptake inhibitor</b>				
Desipramine	Phase II	Ongoing; scheduled through December 2014	Approved drug in Europe	Double-blind; placebo-controlled
<b>Mitochondrial effector</b>				
EPI-743	Phase II	Safety confirmed Improved head circumference growth	Edison Pharmaceuticals	Double-blind; placebo-controlled; oral

IGF-1 = insulin-like growth factor-1; BDNF = brain-derived neurotrophic factor; NMDA = *N*-methyl-D-aspartate

STOP codons during translation, yielding a full-length functional protein. Such aminoglycosides and novel nonaminoglycoside “read-through” compounds have been tested for therapeutic efficacy in Duchenne muscular dystrophy and cystic fibrosis [72]. Gentamycin or geneticin allowed translation of full-length MeCP2 protein in a lymphocyte cell line derived from an individual with a R255X nonsense mutation [73], and in fibroblasts from knock-in mice expressing an R168X nonsense mutation [69]. More recently, embryonic fibroblasts from a mouse model with the R255X nonsense mutation were treated with gentamycin and were able to express full-length MeCP2 [74]. Gentamycin also increased dendritic spine density in neurons derived from induced pluripotent stem cells obtained from a RTT individual with a Q244X nonsense mutation [75]. However, the renal and auditory toxicity, as well as poor central nervous system penetration of aminoglycoside antibiotics, limits their applicability. New “read-through” compounds with better safety profiles are currently being developed.

### Neurotrophins and Growth Factors: Brain-derived Neurotrophic Factor and Insulin-like Growth Factor-1

One of the most prominent targets of MeCP2 transcriptional regulation is the gene encoding brain-derived neurotrophic factor (BDNF) [76, 77], a neurotrophin well known for its critical role in neuronal growth, synapse formation, and activity-dependent plasticity through activation of its selective tropomyosin receptor kinase B (TrkB) receptors [78]. MeCP2

controls BDNF expression through complex interactions [79, 80]. Knockout of *Bdnf* results in neurologic phenotypes reminiscent of the deficits observed in *Mecp2*-deficient mice, suggesting that impaired *Bdnf* transcription contributes to RTT pathophysiology. Furthermore, conditional overexpression of BDNF in excitatory forebrain neurons of *Mecp2*-deficient mice—achieved by crossing BDNF<sup>STOP</sup> mice with *Mecp2*;cre mice—led to the improvement of some of their RTT-like neurologic phenotypes [81]. In addition, *in vitro* expression of green fluorescent protein-tagged BDNF from a cytomegalovirus-driven plasmid rescued the dendritic atrophy caused by short hairpin RNA-mediated knockdown of *Mecp2* in cultured rat hippocampal neurons [82], and the dendritic atrophy and lower spine density in cultured hippocampal neurons from *Mecp2* knockout mice (Xu and LP-M, in preparation). Unfortunately, BDNF has very low BBB permeability, which limits the bioavailability of peripherally administered BDNF as a potential therapy. Hence, BDNF “boosters” or mimetics with sufficient bioavailability in brain are being developed for therapy.

Currently, there are 2 clinical trials in RTT individuals, testing compounds that boost BDNF levels: a Phase I open-label trial of fingolimod and a Phase II open-label trial of glatiramer acetate (Copaxone; Teva Pharmaceuticals, Petah Tikva, Israel), both Food and Drug Administration (FDA)-approved drugs for the treatment of multiple sclerosis. Fingolimod modulates the sphingosine-1 phosphate receptor and increases BDNF levels in neurons in an activity-dependent fashion through the mitogen-activated protein

**Table 2** Current translational trials

Agent or approach	Model	Sponsor	Action	Effect
BDNF mimetic or booster				
LME2A-4	Mouse	Sigma-Aldrich; RSRT	Partial agonist of TrkB receptors	Improves breathing
RP103 (Cysteamine)	Mouse	Raptor Pharmaceuticals	Increases BDNF expression and release	Improves breathing
5-HT1a receptor agonists				
Sarizotan	Mouse	Newron Pharmaceuticals	Agonist of 5-HT1 receptors	Improves breathing
NLX-101	Mouse	Neurolix	Agonist of 5-HT1 receptors	Improves breathing
NMDA receptor modulators				
GluN2A negative allosteric modulator	Mouse	Mnemosyne Pharmaceuticals	Negative allosteric modulator of GluN2A-containing NMDA receptors	Improved visual cortical function
Ketamine	Mouse	Approved drug	Noncompetitive antagonist of NMDA receptors	Improved breathing and locomotor function
Antidepressant				
REV-003 (Tianeptine)	Mouse	Revive Therapeutics	Atypical tricyclic antidepressant	Improves breathing
Gene therapy and "read-through" drugs				
Gene replacement	Mouse	Research Laboratories	Improvement of symptoms	Improvement of lifespan and behavioral symptoms
Read-through agents	<i>In vitro</i> (lymphocyte cell line from RTT; fibroblasts from knockin mice)	Research Laboratories	Skip premature STOP codon in nonsense mutations	<i>In vitro</i> expression of full length MeCP2 protein

BDNF = brain-derived neurotrophic factor; TrkB = tropomyosin receptor kinase B; NMDA = N-methyl-D-aspartate; RTT = Rett syndrome; MeCP2 = methyl CpG binding protein 2



kinase pathway [83]. Glatiramer acetate is an immunomodulatory agent based on the amino acid structure of myelin basic protein that is currently used for the treatment of relapsing–remitting multiple sclerosis. One of the proposed mechanisms of action of Copaxone (Teva Pharmaceuticals) is the increased expression and release of BDNF by autoreactive T cells [84].

Additional compounds have been described that increase BDNF levels and improve RTT-like phenotypes in *Mecp2*-deficient cells and mice. Ampakines are fast-acting molecules that prevent the desensitization of  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxalepropionic acid-type glutamate receptors, which results in higher activity-dependent expression of BDNF, both *in vivo* and *in vitro* [85]. Peripheral treatment with ampakines significantly improved respiratory function in male *Mecp2* knockout mice [86], which phenocopies the recurrent apneas suffered by individuals with RTT. BDNF loop domain mimetics are small molecules designed *in silico* to interact with the BDNF binding pocket in the TrkB receptor [87]. One of them, LM22A-4, restored respiratory regularity in female *Mecp2* heterozygous mice [88, 89], increased dendritic spine density in CA1 pyramidal neurons in organotypic slice cultures of male *Mecp2* knockout mice (Miller, Longo, and LP-M, in preparation), and restored long-term potentiation at CA3→CA1 excitatory synapses in hippocampal slices from symptomatic female *Mecp2* heterozygous mice (Li, Longo, and LP-M, in preparation). Finally, cysteamine (and its FDA-approved dimer cystamine) were shown to increase BDNF release by increasing heat shock DnaJ-containing protein 1b levels and inhibiting transglutaminase [90], which supports the Phase II/III clinical trial of delayed-release cysteamine (RP103) for Huntington disease. None of these compounds has reached human clinical trials for RTT.

Unlike BDNF, insulin-like growth factor-1 (IGF-1) and its active peptide (1–3)IGF-1 cross the BBB and activate intracellular signaling cascades similar to those triggered by BDNF activation of TrkB receptors, which includes phosphatidylinositol-3 kinase–Akt and mitogen-activated protein kinase [91]. Intraperitoneal injection of (1–3)IGF-1 (glypromate) in male *Mecp2* knockout mice improved survival and locomotor activity, as well as social and anxiety behaviors [92]. Recombinant full-length IGF-1 (mecasermin) is already approved by the FDA for the treatment of growth failure in children, and improves survival and RTT-like phenotypes in male *Mecp2* knockout mice and in female *Mecp2* heterozygous mice [93]. The fact that full-length IGF-1 worsened the metabolic syndrome of *Mecp2*-deficient mice should be considered [94]. The effects of full-length IGF-1 are owing to the direct activation of IGF-1 receptors and its downstream signaling cascades. However, the (1–3)IGF-1 tripeptide may increase expression of IGF-1 [95], although its molecular mechanism of action is currently unknown. Based on these promising leads, a Phase II double-blind, placebo-controlled clinical trial involving children aged 3–10 years is underway to

treat patients with RTT with full-length IGF-1 [96]. More recently, a Phase II double-blind, placebo-controlled clinical trial involving older teenagers and adults aged 16–45 years has been initiated to test the efficacy of NNZ-2566, a protease-resistant analogue of (1–3)IGF-1. The results of this short-term trial were sufficiently positive to promote consideration of further study of this agent.

### Neurotransmitter Systems: Monoamines, Glutamate and $\gamma$ -Aminobutyric Acid

Monoamine neurotransmitters like dopamine, serotonin, and noradrenaline have been found to be reduced in autopsy of RTT brains and in *Mecp2*-deficient mice [97–100]. As these monoamines are associated with the regulation of breathing patterns in the brainstem, augmenting their levels may be therapeutic for breathing dysfunction in RTT. Desipramine is an antidepressant that blocks the uptake of noradrenaline, and has been shown to reverse the breathing dysregulation in male *Mecp2* knockout mice [101, 102]. Desipramine is currently in a Phase II double-blind, placebo-controlled clinical trial for RTT. The atypical tricyclic antidepressant tianeptine (REV-003) also improved respiratory activity in *Mecp2*-deficient mice, although this effect may reflect modulation of monoamine levels, glutamate receptor function, or BDNF levels [103]. Sarizotan and NLX-101 are selective agonists of the 5-HT $1\alpha$  receptor that significantly improved breathing patterns and reduced the frequency of apneas in *Mecp2*-deficient mice [104].

$\gamma$ -Aminobutyric acid (GABA)-ergic and glutamatergic systems maintain the fine balance between excitation and inhibition during neuronal maturation. Dysfunctions in these neurotransmitter systems have been described in preclinical models of RTT [105–107]. Dextromethorphan is a *N*-methyl-D-aspartate (NMDA) receptor antagonist currently in a double-blind, placebo-controlled trial in individuals with RTT; to date, its efficacy has not been noted. The NMDA receptor antagonist ketamine improved some RTT-like phenotypes in *Mecp2* knockout mice [108]; based on these encouraging results, a clinical trial of low-dose ketamine is currently planned. A delay of the developmental switch in the expression of GluN2 subunits of the NMDA receptors in the visual cortex contributes to visual acuity deficits in *Mecp2*-deficient mice, which were improved by genetic deletion of the GluN2A subunit [109]; a negative allosteric modulator selective for GluN2A-containing NMDA receptors is currently in preclinical trials in *Mecp2*-deficient mice. Selective deletion of *Mecp2* in GABAergic neurons caused impaired GABAergic transmission, cortical hyperexcitability, and several neurologic features of RTT and autism spectrum disorders [110]. Vigabatrin is an antiepileptic drug that irreversibly inhibits GABA transaminase, inhibits GABA catabolism, and thereby increases GABA levels [111]. The drug is already

FDA-approved for use in epilepsy syndromes. Planning for a clinical trial in RTT is underway. However, retinal toxicity may limit the chronic use of this medication.

### Mitochondrial Function: EPI-743

RTT is associated with high levels of systemic oxidative stress and alteration in mitochondrial morphology, while plasma levels of oxidative stress biomarkers correlate with disease severity and progression [112]. Based on these observations, augmenting glutathione synthesis is thought to be potentially beneficial. The structure of the small molecule EPI-743 is based on vitamin E and its proposed mechanisms of action include augmenting glutathione synthesis and acting at the mitochondrial level to regulate electron transport. An exploratory Phase II placebo-controlled trial of EPI-743 in individuals with RTT revealed improvement in head growth but not in other core features of RTT.

### Conclusion

The discovery of loss-of-function *MECP2* mutations in individuals with RTT, and the generation of experimental cell culture and animal models has provided detailed information of the underlying pathophysiology at the molecular, cellular, and systems levels. In turn, this information prompted preclinical studies of rationally designed therapies aimed at identified molecular targets, some with known mechanism of action. Thus, exciting new opportunities are emerging for the discovery and development of new pharmacotherapies for RTT and other related autism spectrum disorders. However, the complexity of these disorders requires more extensive interdisciplinary collaborations following the highest standards for the validation of animal models, outcome measures, and study design to yield robust and reproducible preclinical information relevant to the human condition. Only then can the preclinical studies provide the strong foundation needed for the success of human clinical trials.

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