RECONNECT (ZYN2-CL-033): Design of a Phase 3 Trial of ZYN002 Cannabidiol Transdermal Gel in Children and Adolescents with Fragile X Syndrome Based Upon Learnings from CONNECT-FX (ZYN2-CL-016) Completed During SARS-COV-2 Pandemic

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BACKGROUND

- Fragile X syndrome (FXS) is a rare genetic condition involving a range of developmental, neuropsychiatric, and behavioral symptoms that are insufficiently managed by the current standard of care (including behavioral and educational interventions, dietary modifications, and off-label prescription therapies)¹⁻⁵
- Patients with complete/near complete methylation of their FMR1 gene are believed to be most likely to have silencing of the FMR1 gene^{1,6}
- Disruption in the endocannabinoid system is one of the proposed mechanisms for the loss of synaptic plasticity and the deficits in emotional responsivity observed in FXS^{7,8}
- Cannabidiol is a negative allosteric modulator at presynaptic CB₁ receptors, a 5HT_{1A} agonist, and a D₂ partial agonist⁹⁻¹¹
- ZYN002 is a pharmaceutically produced cannabidiol transdermal gel in clinical development for the treatment of behavioral symptoms associated with FXS⁵
- RECONNECT (ZYN2-CL-033) is a pivotal, randomized, doubleblind, Phase 3 trial to assess the efficacy and safety of ZYN002 in children and adolescents with FXS aged 3 through 17 years
- RECONNECT is designed based on learnings from CONNECT-FX (ZYN2-CL-016), a randomized, double-blind Phase 3 trial, which was completed during the SARS-CoV-2 (COVID-19) pandemic, in 212 patients with FXS¹³

OBJECTIVE

 To describe the learnings from CONNECT-FX, the first randomized, double-blind, placebo-controlled trial of ZYN002 in the treatment of FXS, and how those learnings shaped the design of the follow-on trial, RECONNECT

METHODS

- The primary endpoint in CONNECT-FX was change in severity of Social Avoidance (SA) as measured by the SA subscale of the Aberrant Behavior Checklist–Community FXS (ABC-C_{FXS} SA)
- Secondary endpoints included Caregiver Clinical Global Impression of Change (CaGI-C) for Irritability/Disruptive behaviors and Social Interactions
- A pre-planned ad hoc analysis of patients having ≥90% methylation of the FMR1 gene^a was conducted
- A post hoc analysis of patients with complete methylation (100%) of the FMR1 gene^a was conducted
- The results and conduct of CONNECT-FX were reviewed to identify key learnings to design RECONNECT

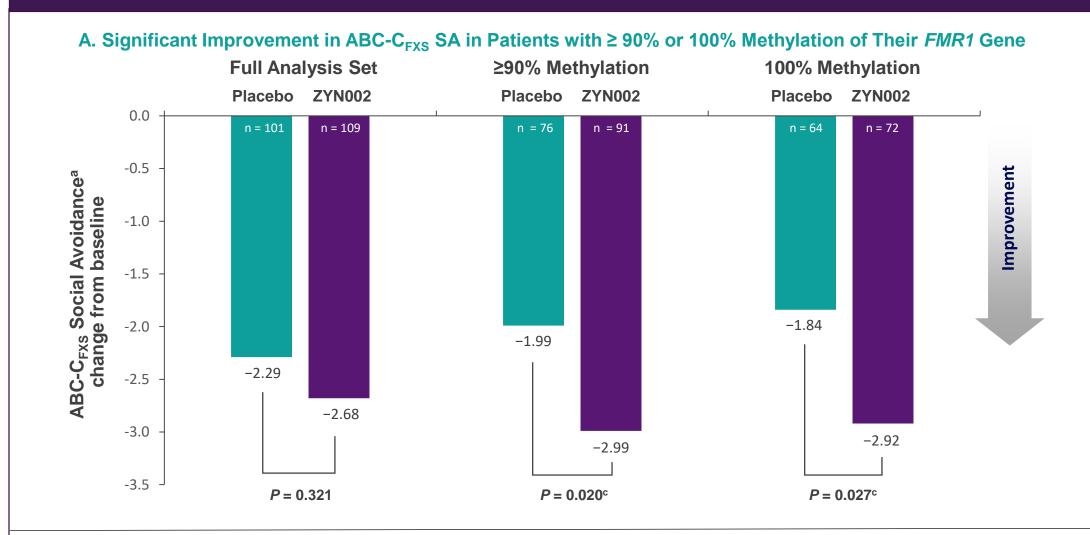
RESULTS

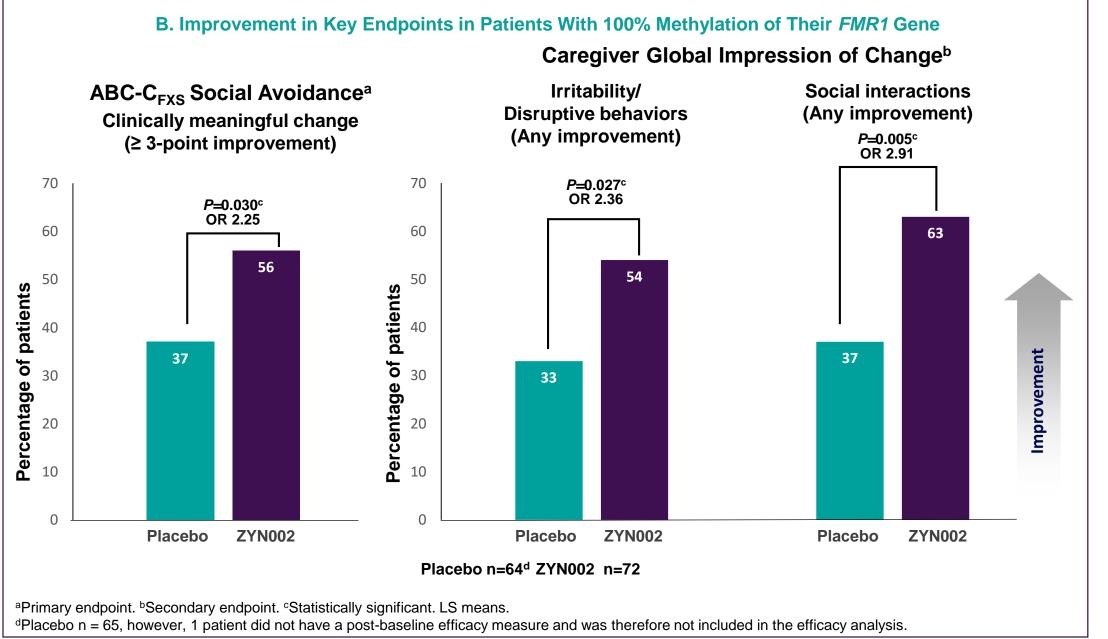
- In CONNECT-FX, improvements were seen in SA behaviors as measured by ABC-C_{FXS} SA in the full population but did not reach statistical significance.
- Significant improvements were seen in patients with ≥90% or 100% methylation of their *FMR1* gene (**Figure 1A**)
- ZYN002 was superior to placebo in multiple analyses in the groups of patients with either ≥90% methylation or complete methylation (100%, **Figure 1B**) of their *FMR1* gene, which represented 80% and 65% of patients, respectively

Lessons Learned From CONNECT-FX

- Treatment response to ZYN002 was greatest in patients with ≥90% methylation of the FMR1 gene
- Qualitative studies¹⁴ support that the ABC-C_{FXS} is fit for purpose in FXS. Meaningful within-patient change thresholds (MCT) were estimated in CONNECT-FX for the SA, Irritability and Socially Unresponsive/Lethargic subscales¹⁵
- Caregiver Global Impression of Severity/Change (CaGI-S/C) was responsive to change to treatment with ZYN002, with greatest change on the Social Interactions subscale
- Virtual visits were successfully incorporated as a result of COVID-19 restrictions

Figure 1. CONNECT-FX Efficacy Results That Informed the Design of RECONNECT





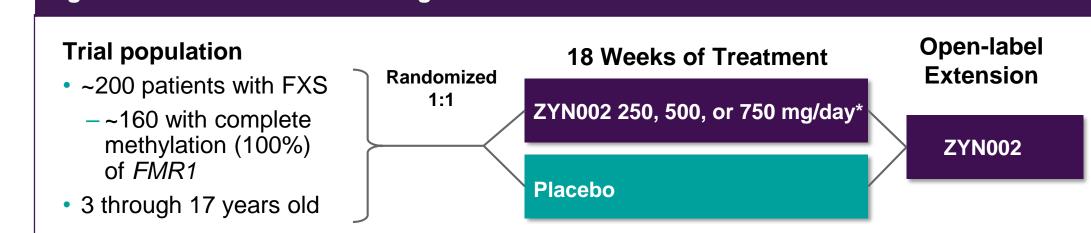
RECONNECT Trial Design Considerations and Trial Design

- Based on the learnings from the CONNECT-FX trial, several optimizations were made in the design of the RECONNECT trial (Table 1)
- These optimizations included a strategic enrichment of the trial population by enrolling a primary cohort of patients with complete methylation (100%) of their *FMR1* gene and a secondary cohort with partial methylation (<100%) to allow confirmation of the impact of *FMR1* methylation on treatment response based upon the data from CONNECT-FX and input from the FDA
 - Approximately 160 patients with complete methylation (100%) of FMR1 will be enrolled
 - Approximately 40 patients with partial methylation (<100%) of FMR1 will be enrolled
- The duration of the trial was lengthened from 14 weeks in CONNECT-FX to 18 weeks in RECONNECT to provide more time for elucidation of treatment effect
- Other key elements of CONNECT-FX were maintained, including the primary and several key secondary endpoints
- The trial design for RECONNECT is shown in Figure 2

Table 1. RECONNECT Trial Design Optimizations

- Population enrichment: Patients with complete methylation (100%) of their FMR1 gene as the primary analysis population; patients with partial methylation (<100%) as a secondary cohort
- **Dosing:** 3 vs 2 weight-based dose strata (250, 500 and 750 mg/day vs 250 and 500 mg/day) incorporated, with the highest dose of 750 mg/day for patients > 50 kg to optimize dosing across weight strata
- FXS-anchored Clinical Global Impression of Severity/Change (CGI-S/I): Anchored on 3
 core behavioral symptoms of FXS: Social Avoidance/Isolation, Social Interactions, and
 Irritability
- Longer trial duration: 18 weeks of treatment vs 14 weeks in CONNECT-FX
- Virtual visits: Virtual visits incorporated (4 of 8) to reduce burden for families and provide flexibility in the event of ongoing challenges with COVID-19

Figure 2. RECONNECT Trial Design



*Weight-based dosing: ≤ 30 kg = 250 mg/day; > 30 kg = 500 mg/day; > 50 kg = 750 mg/day.

Endpoints

Primary endpoint

 Change from baseline in the ABC–C_{FXS} SA subscale at week 18 in patients with complete methylation (100%) of the *FMR1* gene

Key secondary endpoints

Complete methylation (100%) population

- Change from baseline in ABC-C_{FXS} Irritability subscale
- Improvement in CaGI-C Social Interactions
- Improvement in FXS-specific CGI-I

Total population

Change from baseline in ABC- C_{FXS} SA in the full population (complete and partial methylation)

CONCLUSIONS

- The design of RECONNECT was optimized from learnings from CONNECT-FX and will provide key data to determine the effectiveness of ZYN002 in FXS, while reducing burden of participation in a clinical trial for families with children and adolescents with FXS
- Results from RECONNECT, if successful, could serve as the basis for ZYN002 approval for use in patients with FXS
- RECONNECT will be conducted in 29 centers in the US, Australia, the UK, and Ireland
- US RECONNECT study locations include:

California Minnesota
Florida Mississippi
Georgia New Jersey
Illinois New York
Maryland North Carolina
Massachusetts Ohio

Oklahoma
Pennsylvania
South Carolina
Texas
Utah
Washington
Washington DC

 For more information on RECONNECT, visit www.fragilexhelp.com

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ACKNOWLEDGEMENTS / DISCLOSURES

Acknowledgements

Editorial/medical writing support under the guidance of the authors was provided by *p*-value communications, and was funded by Zynerba Pharmaceuticals, Devon, PA, USA.

Disclosures

NT, TS, and SO are employees of Zynerba Pharmaceuticals. TD is a contractor for Zynerba Pharmaceuticals. EM is an employee of Labcorp Drug Development, which has received research funding from Zynerba. CB is an employee of VeraSci, which has received research funding from Zynerba.

The trial was funded by Zynerba Pharmaceuticals.

Presented at the American Society of Clinical Psychopharmacology 2022 Annual Meeting; May 31-June 3, 2022; Scottsdale, Arizona.