



Fragile X mental retardation protein (FMRP) and metabotropic glutamate receptor subtype 5 (mGlu5) control stress granule formation in astrocytes

B. Di Marco^{a,1}, P. Dell'Albani^a, S. D'Antoni^a, M. Spatuzza^{b,2}, C.M. Bonaccorso^b, S. A. Musumeci^b, F. Drago^c, B. Bardoni^d, M.V. Catania^{a,b,*}

^a Institute for Biomedical Research and Innovation – The National Research Council of Italy (IRIB-CNR), Catania, Italy

^b Oasi Research Institute – IRCCS, Troina, Italy

^c Department of Biomedical and Biotechnological Sciences, University of Catania, Italy

^d Université Côte d'Azur, Inserm, CNRS UMR7275, Institute of Molecular and Cellular Pharmacology, Valbonne 06560, France

ARTICLE INFO

Keywords:

Stress granules
mGlu5 receptor
FMRP
Fragile X syndrome
eIF2 α
Astrocytes

ABSTRACT

Fragile X syndrome (FXS) is a common form of intellectual disability and autism caused by the lack of Fragile X Mental Retardation Protein (FMRP), an RNA-binding protein involved in RNA transport and protein synthesis. Upon cellular stress, global protein synthesis is blocked and mRNAs are recruited into stress granules (SGs), together with RNA-binding proteins including FMRP. Activation of group-I metabotropic glutamate (mGlu) receptors stimulates FMRP-mediated mRNA transport and protein synthesis, but their role in SGs formation is unexplored. To this aim, we pre-treated wild type (WT) and *Fmr1* knockout (KO) cultured astrocytes with the group-I-mGlu receptor agonist (S)-3,5-Dihydroxyphenylglycine (DHPG) and exposed them to sodium arsenite (NaAsO₂), a widely used inducer of SGs formation. In WT cultures the activation of group-I mGlu receptors reduced SGs formation and recruitment of FMRP into SGs, and also attenuated phosphorylation of eIF2 α , a key event crucially involved in SGs formation and inhibition of protein synthesis. In contrast, *Fmr1* KO astrocytes, which exhibited a lower number of SGs than WT astrocytes, did not respond to agonist stimulation. Interestingly, the mGlu5 receptor negative allosteric modulator (NAM) 2-methyl-6-(phenylethynyl)pyridine (MPEP) antagonized DHPG-mediated SGs reduction in WT and reversed SGs formation in *Fmr1* KO cultures. Our findings reveal a novel function of mGlu5 receptor as modulator of SGs formation and open new perspectives for understanding cellular response to stress in FXS pathophysiology.

1. Introduction

Fragile X syndrome (FXS) is the most common form of inherited intellectual disability (ID) and a leading genetic cause of autism. FXS patients suffer from moderate to severe cognitive impairment, and can also exhibit autistic behaviour, increased susceptibility to seizures, hyperactivity, anxiety, and hypersensitivity to sensory stimulation (Hagerman and Hagerman, 2002; Hagerman et al., 2017). FXS is caused by the amplification of CGG trinucleotide repeat in the 5'UTR of the

Fragile X Mental Retardation gene 1 (*FMR1*). In patients this mutation is associated with the methylation of the *FMR1* gene resulting into the transcriptional silencing of this gene (Verkerk et al., 1991; Pieretti et al., 1991; Devys et al., 1993) and the lack of the Fragile X Mental Retardation Protein (FMRP), an RNA binding protein involved in the regulation of translation and transport of its target mRNAs (Maurin et al., 2014; Maurin et al., 2018). FMRP acts mainly as a negative regulator of translation, although recent evidence indicates that it can also function as enhancer of translation (Bechara et al., 2009; Darnell et al., 2011;

Abbreviations: DHPG, Dihydroxyphenylglycine; EDTA, ethylenediaminetetraacetic acid; FMR1, Fragile X Mental Retardation gene 1; FMRP, Fragile X Mental Retardation Protein; FXS, Fragile X syndrome; ID, intellectual disability; mGlu receptors, metabotropic glutamate receptors; MPEP, 2-methyl-6-(phenylethynyl)pyridine; mRNPs, messenger ribonucleoprotein particles; NaAsO₂, sodium arsenite; NAM, negative allosteric modulator; PMSF, phenylmethanesulfonyl fluoride; SGs, stress granules.

* Corresponding author at: Institute for Biomedical Research and Innovation, National Research Council (IRIB-CNR), Via Paolo Gaufami n° 18, 95126 Catania, Italy. E-mail address: mariavincenza.catania@cnr.it (M.V. Catania).

¹ Current address: Department of Clinical Neurobiology, German Cancer Research Center (DKFZ), Im Neuenheimer Feld 280, 69120 Heidelberg, Germany.

² Current address: Institute for Biomedical Research and Innovation – IRIB, the National Research Council of Italy, Catania, Italy.

<https://doi.org/10.1016/j.nbd.2021.105338>

Received 2 October 2020; Received in revised form 12 March 2021; Accepted 17 March 2021

Available online 26 March 2021

0969-9961/© 2021 The Author(s). Published by Elsevier Inc. This is an open access article under the CC BY license (<http://creativecommons.org/licenses/by/4.0/>).

Greenblatt and Spreadling, 2018; Maurin and Bardoni, 2018; Liu et al., 2018; Shah et al., 2020; Richter and Zhao, 2021). In neurons, FMRP is also implicated in mRNAs transport along dendrites and axons being a component of RNA granules, the messenger ribonucleoprotein particles (mRNPs) that escort mRNAs in repressed conditions from soma to synapses, where mRNAs can be locally translated upon specific signals (Ling et al., 2004; Antar et al., 2005; Dichtenberg et al., 2008; Christie et al., 2009; Khayachi et al., 2018; Maurin et al., 2018). FMRP interacts with a plethora of different proteins, expanding the range of cellular functions potentially deregulated in FXS (Bardoni et al., 2006; Pasciuto and Bagni, 2014; Ferron, 2016; Davis and Broadie, 2017).

A new aspect of FMRP function in the cytoplasm is related to its presence in peculiar structures called stress granules (SGs), cytoplasmic aggregates that are formed only under stress conditions, such as exposure to heat, oxidative agents, UV irradiation (Anderson and Kedersha, 2002). SGs are dynamic membrane-less structures composed of stalled preinitiation complexes, RNAs and proteins, including initiation factors and RNA-binding proteins that scaffold untranslated mRNAs and interact with each other (Anderson and Kedersha, 2002; Buchan and Parker, 2009; Protter and Parker, 2016). SGs are reversible aggregates where mRNAs are recruited and temporarily stored during stress, and are dispersed upon stress resolution (Anderson and Kedersha, 2002). They are thought to redirect protein translation during stress by limiting global protein synthesis while allowing the translation of stress-induced mRNAs. FMRP has been found to be associated with the pool of mRNAs that go into SGs upon cellular stress and can be involved in the inhibition of protein synthesis occurring during stress (Kim et al., 2006). Lack of FMRP in mouse fibroblasts has been reported to impair SGs formation (Didiot et al., 2009), although FMRP appears to be dispensable in *Drosophila* (Gareau et al., 2013).

Several FMRP-mediated functions, such as mRNPs transport and protein synthesis, are crucially regulated by activation of group-I metabotropic glutamate (mGlu) receptors (mGlu1 and mGlu5 receptor subtypes) (Nicoletti et al., 2011; D'Antoni et al., 2014). Activation of group-I mGlu receptors increases the rapid translation of pre-existing mRNAs, including the *Fmr1* mRNA (Weiler et al., 1997; Weiler et al., 2004). This mechanism underlies mGlu-mediated Long-Term Depression, a form of protein synthesis-dependent synaptic plasticity, which is abnormally exaggerated in the hippocampus and cerebellum of the *Fmr1* knock out (KO) brain (Huber et al., 2000; Huber et al., 2002; Koekkoek et al., 2005). Furthermore, the activation of mGlu5 receptors is necessary for FMRP-containing mRNPs trafficking from the cell body into dendrites (Antar et al., 2004; Dichtenberg et al., 2008). However, the involvement of mGlu5 receptors in SGs formation has never been investigated.

FMRP is highly expressed in neurons, but is also expressed in glial cells although at lower extent (Bonaccorso et al., 2015; Gholizadeh et al., 2015). Accordingly, a growing number of recent studies highlights the contribution of astrocytes to synaptic defects in FXS and subsequently to the pathophysiology of this disorder (Pacey and Doering, 2007; Cheng et al., 2012; Cheng et al., 2016; Higashimori et al., 2016; Wallingford et al., 2017; Hodges et al., 2017). Importantly, mGlu5 receptor-mediated signaling in astrocytes modulates specific functions involved in synaptic transmission and may also directly participate to pathological events in different neurological disorders, including neurodevelopmental disorders (D'Antoni et al., 2008; Petrelli and Bezzi, 2018).

Based on the premise that regulation of mRNA metabolism via mGlu5 receptors in astrocytes may give an insight into the mechanisms of contribution of this cell type to FXS pathophysiology, we report that upon stress primary cultured astrocytes from *Fmr1* KO mice exhibit less SGs than wild type (WT) astrocytes. More importantly, the activation of mGlu5 receptors reduces the formation of SGs in WT, but has no effect in *Fmr1* KO astrocytes, highlighting a link between mGlu5 receptor and translational regulation during stress in the presence and in the absence of FMRP.

2. Materials and methods

2.1. Astroglial cell cultures

Primary astroglial cultures were prepared from cortex of P0-P1 newborn pups of litters obtained from WT or *Fmr1* KO mice (Dutch-Belgian Fragile X Consortium, 1994). All experiments were performed without previous knowledge of the genotype of the culture being tested. The mice genotype was defined by PCR (Musumeci et al., 2007). Cortices were dissected at stereomicroscope (STEMI DV4 ZEISS) and tissues were enzymatically dissociated into nutrient medium. The basal nutrient medium consisted of Dulbecco modified Eagle's medium (DMEM, Sigma), containing 10% heat inactivated fetal bovine serum (FBS, Sigma), 1% penicillin and streptomycin. Cells were seeded into plastic flasks of 25 cm² at a plating density of 0.5×10^5 cells/cm² (one hemisphere/flask). Cultures were incubated at 37 °C in a humidified 5% CO₂ / 95% air atmosphere. After 10–12 days, cultures were treated with 5 mM leucine methyl ester (Sigma) and shaken (for 6–8 h, 180 rpm) to purify cell cultures from oligodendrocytes and microglia. Subsequently, 35 mm Ø dishes were prepared with 2000 cells/dishes to perform immunocytochemistry. To evaluate specific proteins by Western Blot analysis, cultures were seeded onto 100 mm Ø dish at density of 8×10^5 cells/dish.

2.2. Treatments

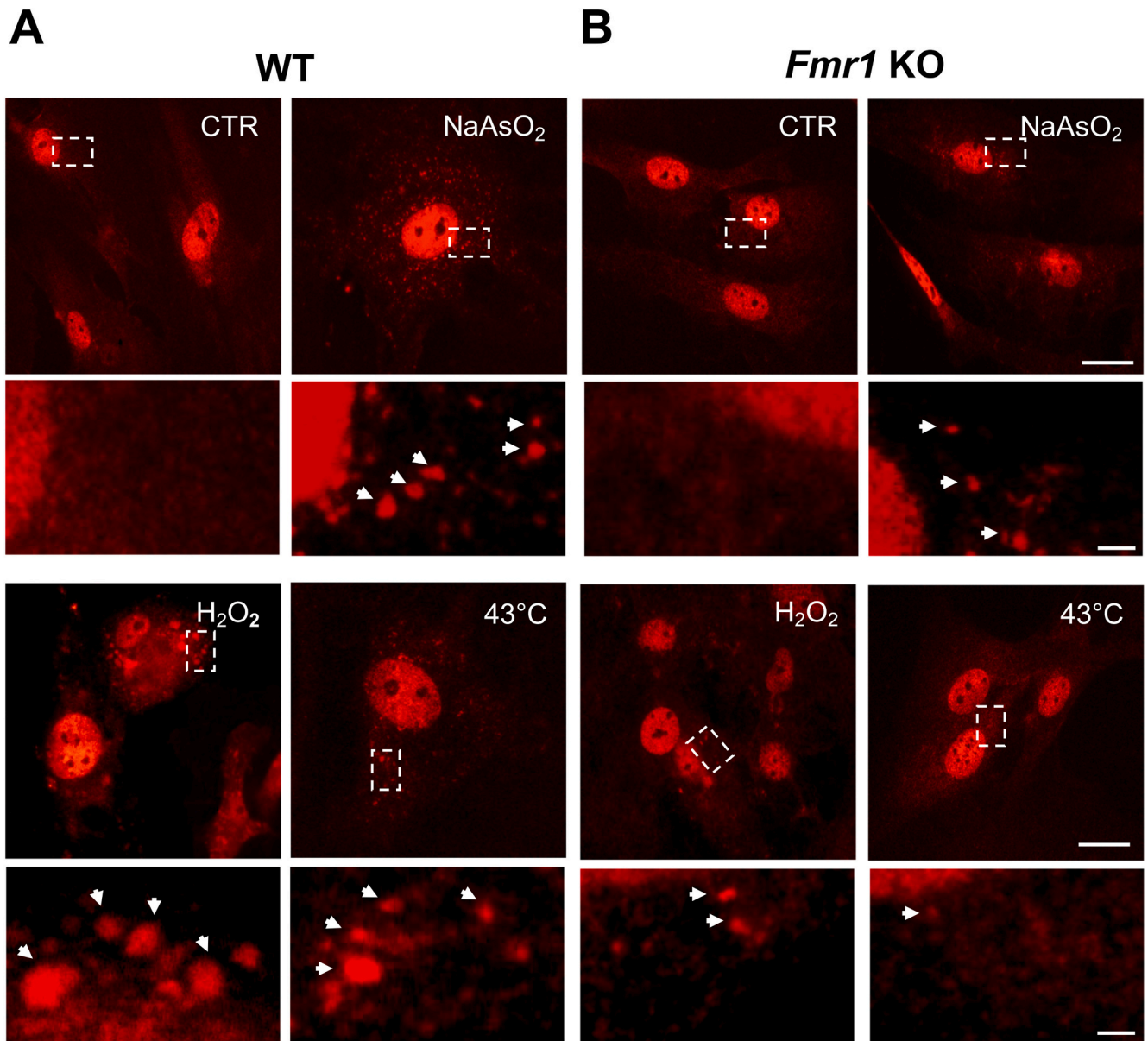
To induce oxidative stress, astrocytes were first shifted in serum-free media for 16 h, then treated with sodium arsenite (500 µM NaAsO₂, Carlo Erba) or with hydrogen peroxide (500 µM H₂O₂, Fluka) for 60 or 90 min, and kept at 37 °C in a humidified 5% CO₂ / 95% air atmosphere. To induce heat shock, cells were maintained at 43 °C for 1 h in a humidified mix 5% CO₂ / 95% air atmosphere. A 5 min pre-treatment with the orthosteric group-I mGlu receptor agonist (S)-3,5-Dihydroxyphenylglycine (100 µM DHPG, Tocris) was carried out before the exposure to stress-inducing agents. The negative allosteric modulator (NAM) of mGlu5 receptor 2-methyl-6-(phenylethynyl)pyridine (3 µM MPEP, Tocris) was applied 10 min before DHPG and during DHPG treatment. In the absence of DHPG, MPEP treatment was carried out 10 min before and during NaAsO₂ exposure. Cycloheximide (30 µg/ml, Tocris) or puromycin (20 µg/ml, Tocris) were added 30 min after NaAsO₂ and maintained for the whole duration of stress exposure.

2.3. Immunocytochemistry and Image analysis

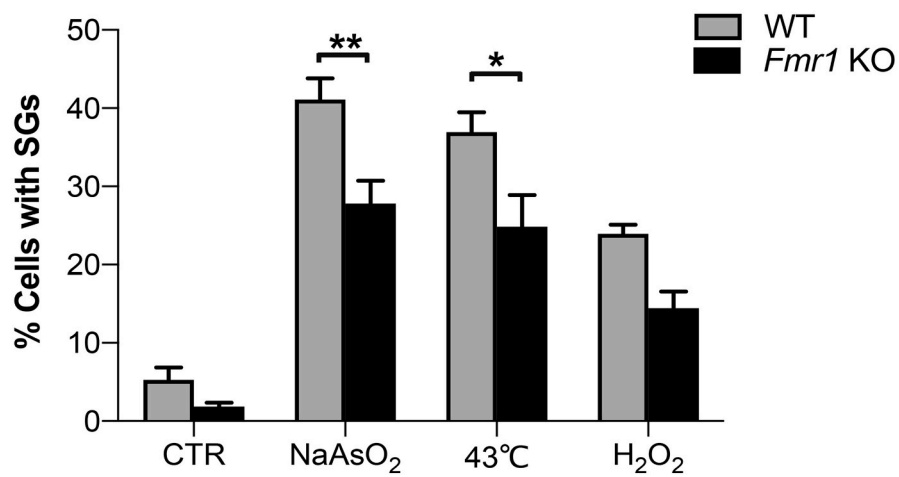
After treatments, cultures were fixed with 4% paraformaldehyde for 15 min at R.T. followed by additional 10 min with cold methanol. Then, cultures were permeabilized in PBS containing 0.2% Triton for 10 min, incubated for 20 min at R.T. with blocking solution containing 4% donkey or goat serum and subsequently with anti-TIA-1 antibody (goat, 1:250, Santa Cruz Biotechnology Cat#sc-1751) for 2 h at R.T. The expression of FMRP in astrocytes was studied by double-labelling immunocytochemistry using the anti-FMRP antibody (rabbit 1:50, Cell Signaling Cat#4317). After washing, cultures were incubated with donkey anti-goat Cy3-conjugated and goat anti-rabbit DyLight488 fluorescent secondary antibodies (Jackson ImmunoResearch). Images were obtained with the Axio Imager. D2 (Zeiss) or LSM-510 Meta Confocal microscopes (Zeiss), and analyzed using the AxioVision Imaging System and the ImageJ softwares. A negative control (no primary antibodies) was used to properly set acquisition parameters.

We identified cells displaying SGs by observing cytoplasmic TIA-1 positive granular spots, which were clearly detectable above a diffuse background (See Fig. 1). We considered SG positive a cell displaying at least 2 TIA-1 positive cytoplasmic spots.

We used the ImageJ software for quantitative analysis of images. To quantify SG numbers and size, images were converted in binary format and processed by the Analyze Particles function of Image J. In order to



C



(caption on next page)

Fig. 1. Stressed *Fmr1* KO astrocytes exhibit a reduced number of cells showing TIA-1⁺ SGs compared to stressed WT astrocytes. (A, B) Cultured astrocytes from WT (A) and *Fmr1* KO (B) were stained with the anti-TIA-1 primary antibody to detect SG formation. The selected cells are representative of the majority of cells expressing high (WT) or low levels of SGs (*Fmr1* KO) after treatments. Scale bar 10 μ m. Small panels show magnifications of the dashed-line boxed areas; arrows indicate TIA-1 positive SGs, scale bar 1 μ m. (C) Graph showing the percentage of cells bearing SGs after exposure to different stress-inducing agents namely 500 μ M NaAsO₂, 43 °C or 500 μ M H₂O₂ for 1 h. Values represent mean \pm S.E.M; n = number of dishes from 1 to 4 cultures, 10 (CTR WT), 12 (NaAsO₂, WT), 13 (43 °C WT), 4 (H₂O₂ WT), 12 (CTR *Fmr1* KO), 13 (NaAsO₂, *Fmr1* KO), 11 (43 °C WT *Fmr1* KO), 4 (H₂O₂ *Fmr1* KO). 115–1235 and 133–1707 cells per dish were analyzed in WT and *Fmr1* KO, respectively. **p* < 0,05; ***p* < 0,01 by Multifactorial Two-Way ANOVA followed by Holm-Sidak method.

exclude nuclei, the size parameters of Analyze Particles were set on 0–1000 pixels. Then, we obtained the masks showing SGs that were used for SG quantification and figure preparation. For TIA-1 and FMRP SG colocalization, images were splitted in order to obtain an image per antibody. Subsequently, images were processed with JACoP colocalization tool using Objects Based Methods set on Centre of Mass with a range of 0–1000 pixel. SGs numbers resulting from the TIA-1/FMRP colocalization mask were used for quantification.

2.4. Western Blotting

Astroglial cell cultures (80% confluence) were harvested by scraping them on ice. Cells were homogenated in lysis buffer [Tris-HCl 40 mM pH 6,8, 1 \times Protease Inhibitor Cocktail-Roche, 1 \times Phosphatase Inhibitor Cocktail-Roche, 100 μ M phenylmethanesulfonyl fluoride (PMSF), 1 mM ethylenediaminetetraacetic acid (EDTA), 5 mM EGTA and 2% SDS], centrifuged for 10 min at 1000 g at 4 °C. Cortices obtained from WT, *Fmr1* KO and mGlu5 KO mice were allowed to thaw on ice, weighed, and homogenized in 10% (w/v) Lysis buffer containing EDTA (1 mM, Sigma), Tris-HCl (10 mM, pH 7.4, Sigma), PMSF (0.5 mM, Sigma), sodium chloride (150 mM, Sigma) and protease inhibitor cocktail (Roche). Supernatant were collected and protein concentration was determined by using the bicinchoninic acid method (BCA kit; Pierce Rockford, IL). Then, proteins were denaturated and loaded onto 8% SDS-polyacrylamide gels. Gels were electroblotted to nitrocellulose membranes (Amersham Biosciences). Filters obtained were processed as indicated by the manufacturer of WesternBreeze Chemiluminescent Immunodetection System kit (INVITROGEN). In brief, filters were blocked for 30 min with blocking solution and incubated O/N with the following primary antibodies: anti-mGlu5 receptor (rabbit 1:6000, Upstate N. 06–451), anti-eIF2 α and anti-phospho-eIF2 α (rabbit 1:1000, Cell Signaling Cat# 9722 and 9721, respectively), anti-GAPDH (rabbit 1:1000, Cell Signaling Cat#2118) and anti- β -Tubulin (rabbit 1:1000, Cell Signaling Cat#2146). Alkaline phosphatase-conjugated secondary anti-rabbit antibodies from INVITROGEN kit were used. Chemiluminescence was detected and quantified by VersaDoc™ 4000 Imaging System (BIORAD).

2.5. Statistical analysis

Statistical analysis was carried out by GraphPad Prism version 8.1.1 (GraphPad Software, USA). All datasets were tested for normal distribution between groups. We applied either parametric or non-parametric tests according to the results of Shapiro-Wilk and Kolmogorov-Smirnov tests.

Most data were analyzed using multifactorial ANOVA (One-Way or Two-Way analysis of variance) followed by appropriate post-hoc comparisons (Tukey's, Holm-Sidak or Dunn's methods). Unpaired *t*-test was used to quantify the differences shown in the graphs of Figs. 4E and 5C.

3. Results

3.1. *Fmr1* KO astrocytes show less SGs than WT astrocytes

To induce formation of SGs, WT and *Fmr1* KO cultured astrocytes were exposed to different stressors such as sodium arsenite (NaAsO₂, 500 μ M for 60 min), heat (43 °C for 60 min), and hydrogen peroxide (H₂O₂, 500 μ M for 60 min). SGs were studied by means of

immunocytochemistry aimed at revealing the TIA-1 protein, a known marker of SGs, which has a nuclear localization under control condition, whereas it accumulates in the cytoplasm and takes part in SGs formation upon stress induction (Kedersha et al., 1999). As expected, exposure to both oxidative stress and heat significantly increased the formation of SGs in both WT and *Fmr1* KO cultured astrocytes, as revealed by the increased number of cells bearing TIA-1 positive SGs (Fig. 1). However, in *Fmr1* KO astrocytes we detected a significantly lower number of cells with SGs compared to WT astrocytes after exposure to stress (Fig. 1). Among different stressors we decided to expose astrocytes to NaAsO₂ only, a widely used and well characterized inducer of SGs formation.

3.2. Activation of mGlu5 receptors before exposure to stress reduces SGs in WT, but has no effect in *Fmr1* KO astrocytes

We examined the expression of mGlu5 receptors in primary cultures of WT and *Fmr1* KO cortical mouse astrocytes (Fig. S1) and found that their expression levels were comparable in cultures from WT and *Fmr1* KO mice. As expected from previous published results (Janssens and Lesage, 2001; Aronica et al., 2003; D'Antoni et al., 2008), we found that the expression levels of mGlu5 receptors in cultured astrocytes were low, as compared with those found in the cortex of mice at post-natal day 7, when the expression of mGlu5 receptors in the brain is maximal (Catania et al., 1994; Catania et al., 2007).

Activation of mGlu5 receptors with the group-I mGlu receptor agonist (S)-3,5-Dihydroxyphenylglycine (DHPG, 100 μ M for 5 min) before exposure of astrocytes to NaAsO₂ induced a significant reduction in the number of SGs per cell in WT cultures (Fig. 2A, B), but had no effect in *Fmr1* KO astrocytes (Fig. 2C, D). Quantification of the size of SGs also revealed that SGs were smaller in *Fmr1* KO than in WT cultures and that DHPG treatment before exposure to NaAsO₂ reduced SGs size in WT astrocytes, at similar levels as SGs size observed in *Fmr1* KO astrocytes; however, DHPG treatment before exposure to NaAsO₂ did not modify the size of SGs in *Fmr1* KO astrocytes (Fig. S2).

We also quantified the percentage of cells bearing SGs in both WT and *Fmr1* KO stressed cultures and found that the pre-treatment with DHPG induced a significant reduction of SGs positive cells in stressed WT cultures only, with no effect in *Fmr1* KO cultures (Fig. 3A, B). The DHPG-induced effect in WT cultures was antagonized by the highly selective mGlu5 receptor NAM 2-methyl-6-(phenylethynyl)pyridine (MPEP, 3 μ M), clearly indicating an mGlu5 receptor – mediated effect (Fig. 3A). Interestingly, the stress-induced increase in the percentage of SG positive cells was not affected by application of MPEP alone before and during stress in WT astrocytic cultures, whereas it was further increased in *Fmr1* KO cultures (Fig. 3B).

To get an insight into the mechanisms underlying the effects of mGlu5 receptor activation on the modulation of SGs formation, we exposed both WT and *Fmr1* KO cell cultures to NaAsO₂ and then treated them with the protein synthesis inhibitors puromycin and cycloheximide, which have different mechanisms of action. Puromycin destabilizes polysomes and facilitates SGs formation by making mRNAs available, while cycloheximide freezes ribosomes on translating mRNAs and therefore inhibits SGs formation (Kedersha et al., 2000). Indeed, puromycin induced a significant increase of cells bearing SGs, whereas cycloheximide induced a drastic reduction of SGs formation in both WT and *Fmr1* KO astrocytes (Fig. 3). This result suggests that the basic mechanisms underlying SGs formation are not disrupted in the absence of FMRP. Interestingly, the exposure to DHPG for 5 min before stress

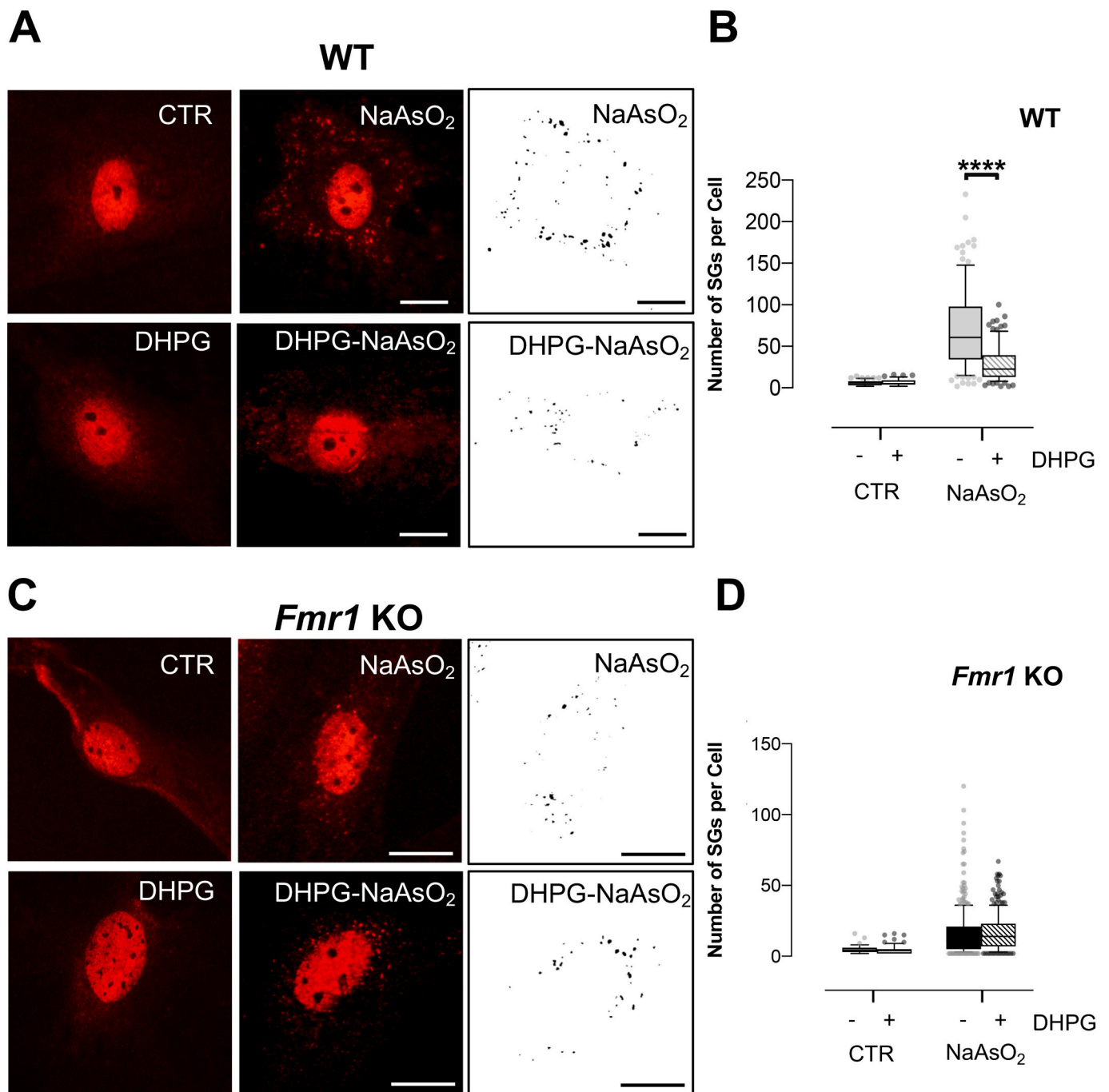


Fig. 2. Activation of mGlu5 receptor reduces the number of SGs in WT but does not affect SGs formation in *Fmr1* KO astrocytes.

(A, C) Left and middle panels show WT (A) and *Fmr1* KO (C) astrocytes immunostained with anti-TIA-1 antibody. Astrocytes were untreated (CTR), exposed to 100 μ M DHPG for 5 min (DHPG), treated with 500 μ M NaAsO₂ for 1 h (NaAsO₂), or exposed to DHPG for 5 min and then to NaAsO₂ for 1 h. Selected cells are representative of the majority of cells that do not show SGs (CTR, DHPG) and that do express SGs after treatments (NaAsO₂, DHPG-NaAsO₂). Right panels show SG masks as revealed by the Analyze Particles module of Image J. Scale bar 10 μ m. (B, D) Box and whisker plots show quantitative analysis of the number of SGs per SG-positive cell in WT (B) and *Fmr1* KO (D) cultures after treatments. (B) n = cells from 3 to 4 different cultures, 89 (CTR), 99 (DHPG), 115 (NaAsO₂), 106 (DHPG NaAsO₂), ****p < 0,0001 by One-Way Anova with Dunn's multiple comparisons test. (D) n = cells from 3 to 4 different cultures, 200 (CTR), 172 (DHPG), 391 (NaAsO₂), 314 cells (DHPG NaAsO₂).

induction reduced the number of cells with SGs also in puromycin-treated WT cells (Fig. 3A), but had no effect in *Fmr1* KO astrocytes (Fig. 3B). This would indicate that activation of mGlu5 receptors before stress induction counteracts the formation of SGs despite the availability of mRNAs in WT cultures, whereas does not trigger a similar mechanism in *Fmr1* KO astrocytes (Fig. 3B).

3.3. Activation of mGlu5 receptors before stress induction reduces phosphorylation of translation initiation factor eIF2 α in WT but not in *Fmr1* KO astrocytes

Since the stress-induced phosphorylation of eIF2 α factor is a major trigger of SGs formation, (Kedersha et al., 1999, see discussion), we tested if mGlu5 receptor activation affects NaAsO₂-induced eIF2 α

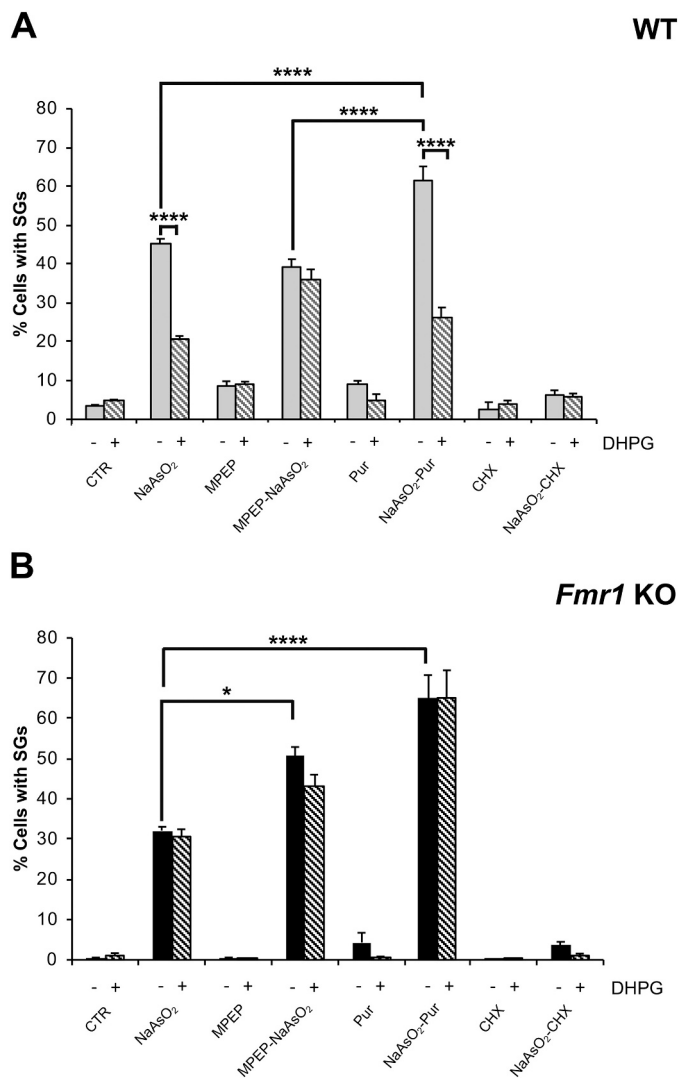


Fig. 3. Pharmacological blockade of mGlu5 receptors and treatment with puromycin rescue SGs formation in *Fmr1* KO astrocytes.

Graphs show the percentage of cells bearing TIA-1-positive SGs in WT (A) and *Fmr1* KO (B) cultures. Astrocytes were exposed to 5 min pre-treatment with/without DHPG followed by NaAsO₂ in the presence of mGlu5 NAM MPEP or protein synthesis inhibitors such as puromycin (Pur) and cycloheximide (CHX). To antagonize the effect of DHPG, MPEP was added 10 min before and during DHPG exposure. Astrocyte samples are shown as following: untreated (CTR); treated with 500 μM NaAsO₂ for 90 min (NaAsO₂); treated with 3 μM MPEP for 10 min (MPEP); treated with 3 μM MPEP for 10 min followed by 500 μM NaAsO₂ for 90 min (MPEP-NaAsO₂); treated with 20 μg/ml Puromycin (Pur) or 30 μg/ml Cycloheximide (CHX) for 60 min; treated with NaAsO₂ for 90 min during which Pur (NaAsO₂-Pur) or CHX (NaAsO₂-CHX) were added 60 min before the end of NaAsO₂ treatment. (A): Values represent mean ± S.E.M. n = number of dishes from 1 to 2 different WT cultures, 5 (CTR), 5 (DHPG), 9 (NaAsO₂), 6 (DHPG NaAsO₂), 4 (CHX), 3 (DHPG CHX), 5 (NaAsO₂ CHX), 4 (DHPG NaAsO₂ CHX), 6 (Pur), 3 (DHPG Pur), 5 (Pur NaAsO₂), 5 (NaAsO₂ DHPG Pur), 3 (MPEP), 3 (MPEP NaAsO₂), 4 (MPEP DHPG), 4 (MPEP DHPG NaAsO₂). 54–1109 cells per group were analyzed ****p < 0,0001 by One-Way ANOVA with Tukey's multiple comparisons test. (B) Values represent mean ± S.E.M. n = dishes from 1 to 2 different *Fmr1* KO cultures 6 (CTR), 6 (DHPG), 6 (NaAsO₂), 6 (DHPG NaAsO₂), 4 (CHX), 3 (DHPG CHX), 4 (NaAsO₂ CHX), 4 (DHPG NaAsO₂ CHX), 4 (Pur), 3 (DHPG Pur), 4 (Pur NaAsO₂), 4 (NaAsO₂ DHPG Pur), 4 (MPEP), 4 (MPEP NaAsO₂), 4 (MPEP DHPG), 4 (MPEP DHPG NaAsO₂). 86–1986 cells per group were analyzed. ****p < 0,0001, *p < 0,05 by One-Way ANOVA with Tukey's multiple comparisons test.

phosphorylation. Western Blot analyses showed that eIF2α was highly phosphorylated under stress condition, as expected, in both WT and *Fmr1* KO astrocytes (Fig. 4A, B). Interestingly, while exposure to NaAsO₂ induced a robust phosphorylation of eIF2α in both WT and *Fmr1* KO (Fig. 4A, D), a 5 min pre-treatment with DHPG before stress induction differently affected eIF2α phosphorylation in the two genotypes. Semi-quantitative analysis of phosphorylated-eIF2α revealed lower levels of eIF2α phosphorylation upon stress in WT than in *Fmr1* KO cultures after activation of mGlu5 receptors (Fig. 4C, D, E).

3.4. Activation of mGlu5 receptors reduces recruitment of FMRP in SGs

Double-labelling immunocytochemistry and confocal microscopy revealed a remarkable co-localization of FMRP and TIA-1 protein in WT astrocytes exposed to NaAsO₂ (Fig. 5A) indicating that FMRP is recruited in SGs as observed in other cell types. Astrocytes exposed to DHPG before NaAsO₂ showed a lower amount of FMRP co-localization in TIA positive SGs than cells exposed to NaAsO₂ only (Fig. 5A, C).

4. Discussion

Within the Central Nervous System, the function of FMRP has been principally investigated in neurons, whereas the biological significance of FMRP in other cell types has received scant attention until recently. Emerging evidence suggests that lack of FMRP in astrocytes contributes to FXS phenotype, i.e. abnormal dendritic spine morphology/dynamics and synapse development, through mechanisms that involve neuron-glia interaction (Cheng et al., 2016; Higashimori et al., 2016; Hodges et al., 2017; Wallingford et al., 2017). This can occur because the FMRP-regulated synthesis of both resident and secretory astrocytic proteins is disrupted in FXS. Therefore, SGs formation and its modulation in astrocytes is an important yet unexplored aspect of mRNA metabolism in FXS. Here we report that the activity of mGlu5 receptors, which regulate FMRP-dependent mRNA transport and translation in neurons, can also modulate SGs formation in astrocytes.

The activation of mGlu5 receptors reduced SGs formation in WT to a similar extent as in *Fmr1* KO astrocytes. However, in the absence of FMRP, mGlu5 receptor activation did not further reduce SGs formation. In contrast, the NAM MPEP, which is known to inhibit the constitutive activity of mGlu5 receptors (Pagano et al., 2000), did not have any effect in WT cultures, but reversed the reduced SGs formation in *Fmr1* KO astrocytes. These results resemble several observations reporting that activation of mGlu5 receptors mediates effects in WT, i.e. mRNA transport and translation, whereas it has no effect in *Fmr1* KO cells (reviewed in Bassell and Warren, 2008). Indeed, activation of mGlu5 receptors triggers protein translation in hippocampal slices of WT mice, but does not further increase the constitutively elevated protein synthesis in *Fmr1* KO mice, which, in contrast, is strikingly reversed by the pharmacological blockade of mGlu5 receptors or its genetic down-regulation (Dolen et al., 2007; Michalon et al., 2012).

To deepen the relationship between activation of mGlu5 receptors, SGs formation and mRNA translation, we carried out stress inducing experiments with/without DHPG in the presence of cycloheximide or puromycin. Using these drugs, it was established that SGs-associated mRNAs are in a dynamic equilibrium with polyribosomes (Kedersha et al., 2000). In line to what previously reported in other cell types, we observed that in both WT and *Fmr1* KO astrocytes puromycin increased SGs formation upon stress, while cycloheximide completely reversed SGs formation. Interestingly, we observed that in the presence of puromycin SGs formation occurred in *Fmr1* KO astrocytes to a similar extent as in WT, indicating that destabilization of polyribosomes makes available the initiation complex and mRNAs for SGs formation both in the presence and in the absence of FMRP. The reduced SGs formation in *Fmr1* KO astrocytes is also restored by MPEP suggesting that this molecular phenotype could be due to an increased rate of mRNA recruitment in polyribosomes in the absence of FMRP. This is in agreement

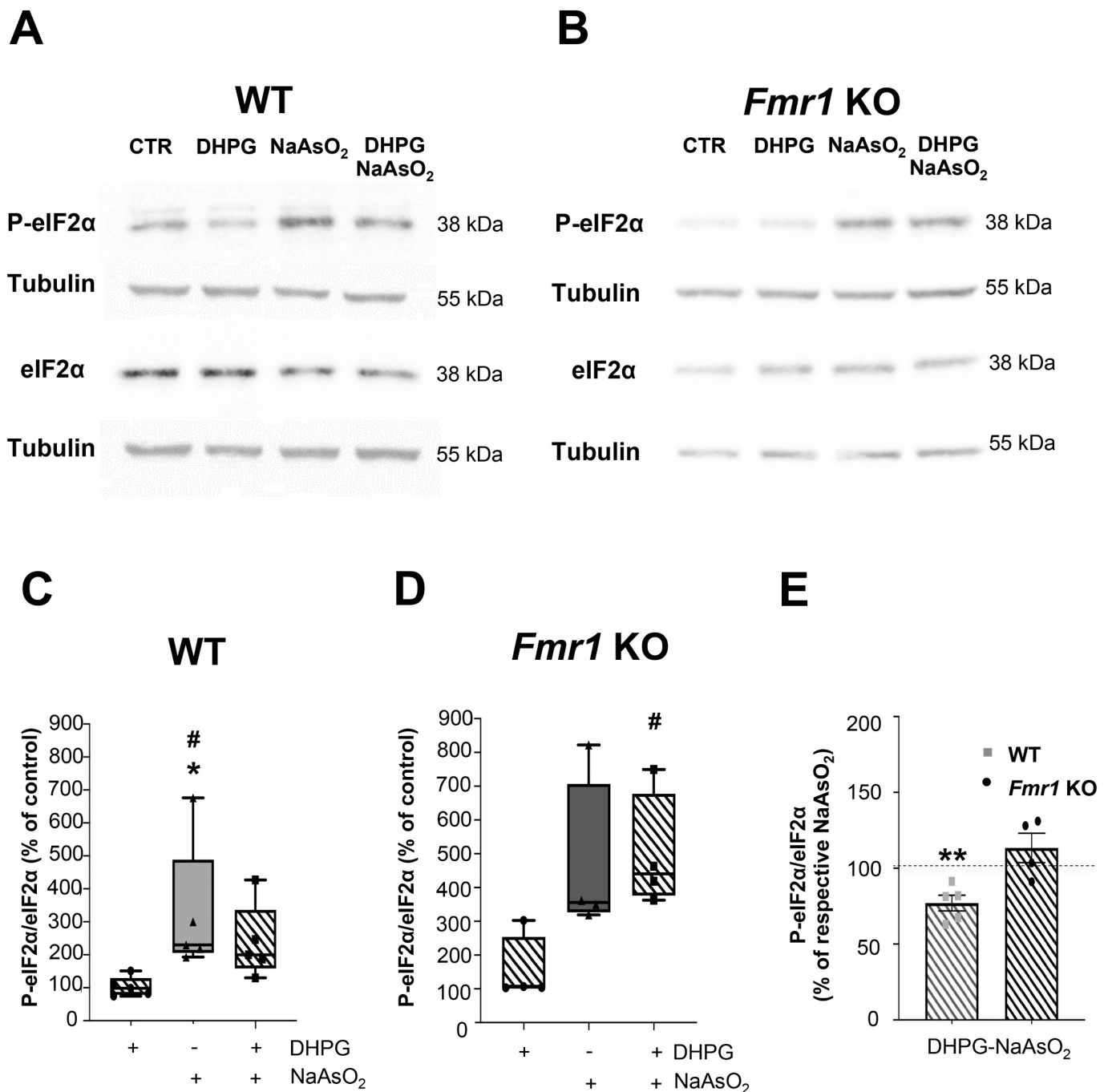


Fig. 4. Activation of mGlu5 receptors before stress reduces phosphorylation of eIF2α in WT astrocytes, but has no effect in *Fmr1* KO astrocytes.

(A,B) Representative immunoblots showing the levels of total and phosphorylated-eIF2α protein in WT (A) and *Fmr1* KO astrocytes (B). Cultures were untreated (CTR), treated with 100 μM DHPG for 5 min (DHPG), treated with 500 μM NaAsO₂ for 90 min (NaAsO₂) and treated with 100 μM DHPG for 5 min followed by 500 μM NaAsO₂ for 90 min (DHPG- NaAsO₂). 50 μg of proteins were loaded. Tubulin was used as loading control. (C-E) Semiquantitative analysis of phosphorylated eIF2α vs. total eIF2α. The expression levels of phospho-eIF2α and eIF2α were quantified by densitometry and normalized first against the respective tubulin and then calculated as ratio of total eIF2α signal. Relative optical density is presented as percentage of control. Data are presented as box and whisker plots (C, D) and mean ± S.E.M. (E) of five (C, E) and four (D, E) separate experiments (C) *p < 0,05 vs. ctr; # p < 0,05 vs. DHPG by One Way ANOVA with Tukey's multiple comparisons test. (D) # p < 0,05 vs. DHPG by One Way ANOVA with Dunn's multiple comparisons test. (E) **p = 0,0022 vs. respective NaAsO₂ by unpaired t-test.

with the notion that the absence of FMRP leads to a constitutive mGlu5-dependent increased rate of protein synthesis (Dolen et al., 2007; Michalon et al., 2012). In other words, an altered balance between polyribosomes and SGs is possibly responsible for the reduced SGs formation in *Fmr1* KO cells rather than the absence of the shuttling action of FMRP between the two ribonucleoproteic structures.

We also found that mGlu5 receptor activation differently affected eIF2α phosphorylation in stressed WT astrocytes and *Fmr1* KO cultures,

with lower levels in WT astrocytes. In stressed cells, activation of one or more eIF2α kinases (e.g. PKR, PERK/PEK, GCN2, HRI) results in the phosphorylation of eIF2α, an essential subunit of the eIF2-GTP-tRNA^{Met} ternary complex required to initiate protein synthesis. Once phosphorylated eIF2α is no longer available to the canonical assembly of the translation initiation complex, and favours the formation of an abnormal 48S complex carrying mRNAs that were destined for translation and that take part in SGs (Anderson and Kedersha, 2002; Kedersha and Anderson,

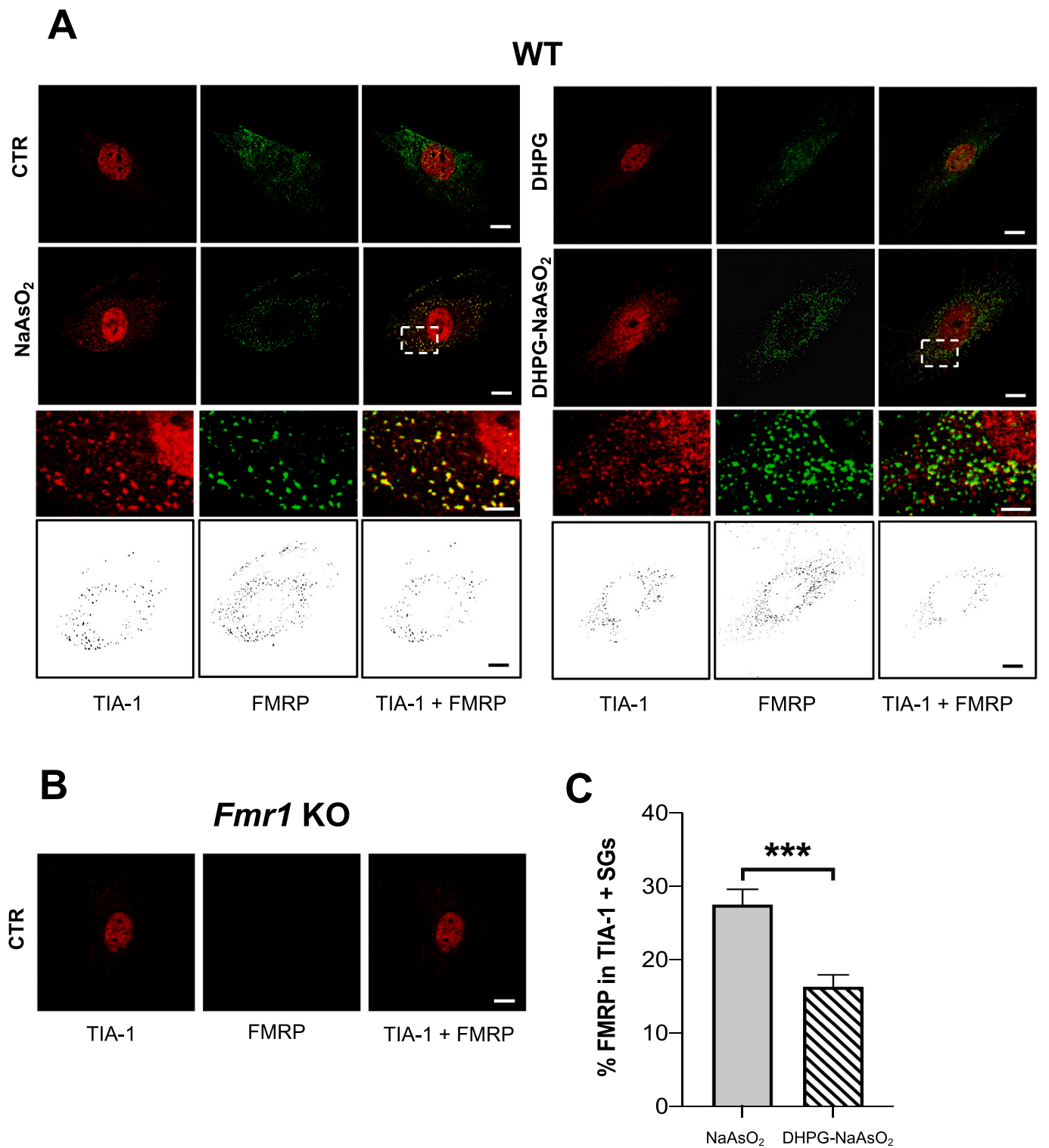


Fig. 5. Activation of mGlu5 receptors reduces FMRP recruitment into SGs.

(A) Images show WT astrocytes stained with anti-TIA-1 and anti-FMRP primary antibodies. Drawings show TIA-1 positive SGs, FMRP positive SGs and double TIA/FMRP SGs as revealed by masks generated by the Analyze Particles module of Image J. Astrocytes were untreated, exposed to DHPG (100 μ M for 5 min), treated with NaAsO₂ (500 μ M for 30 min) or exposed to DHPG for 5 min and then to NaAsO₂ for thirty minutes. TIA-1 staining is shown in red and FMRP in green. Scale bar = 20 μ m. Small panels show magnifications of the dashed-line boxed areas, scale bar 20 μ m. (B) *Fmr1* KO astrocytes stained with anti-TIA-1 and anti-FMRP primary antibodies as a negative control. (C) The graph represents the percentage of FMRP co-localization in TIA positive SGs calculated by JACoP colocalization plugin of Image J software *** $p < 0,0003$ unpaired t-test. $n = 27$ cells (NaAsO₂) and 16 cells (DHPG NaAsO₂) from 1 to 2 cultures.

2009). After this crucial initial event, TIA-1 and then other RNA binding proteins including FMRP are recruited to SGs. Thus, mGlu5 receptor activation in WT may impair SG formation by reducing the number of abnormal pre-initiation complexes which represent the core of SGs essential for the subsequent recruitment of TIA-1 and FMRP. Accordingly, in WT astrocytes the activation of mGlu5 receptors before stress counteracts SGs formation even in the presence of puromycin, whereas this did not occur in *Fmr1* KO cultures. On the other hand, it is also possible that mGlu5 activation may favour the rapid synthesis or post-translational modification of other proteins interfering with the subsequent aggregation of interacting RNA-binding proteins in SGs.

Intriguingly, eIF2 α phosphorylation was also increased upon stress in *Fmr1* KO astrocytes, although it was not reduced by activation of mGlu5 receptors. This is not in contrast with our observation that *Fmr1* KO astrocytes exhibit an impaired SGs formation. In fact, SGs formation is abolished even in the presence of continued phosphorylation of eIF2 α when the availability of free mRNAs is reduced by drugs such as cycloheximide or emetine (Kedersha et al., 2000, Fig. 4). The lack of a DHPG-induced effects on eIF2 α dephosphorylation and SGs formation might rather indicate that in the absence of FMRP mGlu5 receptors are insensitive to the orthosteric agonists and/or uncoupled from downstream signaling, as shown for DHPG stimulated mRNA translation of FMRP targets (Dolen et al., 2007; Bassell and Warren, 2008; Michalon et al., 2012).

Despite recent advancements in elucidating the SGs composition and mechanisms underlying their formation, the biological significance of SGs remains undefined. By providing a sink for pro-apoptotic signaling molecules SGs may play a role in promoting cell survival upon stress (Arimoto et al., 2008; Eisinger-Mathason et al., 2008). Therefore, the reduced SGs formation in the absence of FMRP argues for a further vulnerability of FXS phenotype in coping with different stressors, including oxidative stress. Several pieces of evidence indicate that oxidative stress is indeed increased in the *Fmr1* KO mouse model and may play a role in FXS pathophysiology (El Bekay et al., 2007; Bechara et al., 2009; Davidovic et al., 2011; D'Antoni et al., 2020). The restored formation of SGs by MPEP suggests that antagonism of mGlu5 receptors could be a protective therapeutic strategy against the deleterious consequences of stress in FXS. Besides the pathophysiological relevance of our data, we believe that, highlighting the role of FMRP in SG formation and its modulation by mGlu5 receptors, our study contributes to a further understanding of the function of FMRP in the control of RNA metabolism.

To our knowledge, this is the first report that the activation of a neurotransmitter receptor has an impact on SGs formation, revealing a novel function of mGlu5 receptors in astrocytes. Our study adds relevant information to a complex biological problem involved in the mechanisms of cellular response to stress and may have critical implication for FXS pathophysiology. Furthermore, considering a possible link between SGs formation and cell survival (Arimoto et al., 2008; Eisinger-Mathason et al., 2008), our study may open new perspectives for pharmacological modulation of SGs in neurological disorders in which oxidative stress and endoplasmic reticulum stress contribute to cell death.

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.nbd.2021.105338>.

Author contribution

Barbara Di Marco: Conceptualization, Validation, Formal Analysis, Investigation, Writing – Original Draft, Writing – Review & Editing, Visualization. **Paola Dell'Albani:** Formal Analysis, Investigation, Writing – Review & Editing. **Simona D'Antoni:** Formal Analysis, Investigation, Writing – Review & Editing, Visualization. **Michela Spatuzza:** Formal Analysis, Investigation. **Carmela M. Bonaccorso:** Investigation. **Salvatore A. Musumeci:** Funding Acquisition. **Filippo Drago:** Funding Acquisition. **Barbara Bardoni:** Conceptualization, Resources, Writing – Review & Editing. **Maria Vincenza Catania:**

Conceptualization, Formal Analysis, Resources, Supervision, Writing – Original Draft, Writing – Review & Editing, Funding Acquisition.

Ethical approval

Animal care and handling were carried out in compliance with the European Council Directive (86/609/EEC) and the Italian Animal Welfare Act for the use and care of laboratory animals. All efforts were made to minimize animal suffering and to reduce the number of animals used.

Funding

This work was supported by the International Ph.D. Program in Neuropharmacology, Medical School, University of Catania, Catania, Italy; IRCCS Oasi Research Institute, Troina, Italy; Programma Operativo Obiettivo Convergenza 2007–2013, FSE, Regione Siciliana, avviso n. 1 2012. B.D.M. was supported by PON01_00110 and Programma Operativo Obiettivo Convergenza 2007–2013, FSE, Regione Siciliana, avviso n. 1 2012. B.B. is supported by the "Agence Nationale de la Recherche" ANR-20-CE16-0016-01.

Declaration of Competing Interest

The authors declare that they have no conflict of interest.

Acknowledgements

The authors are grateful to Giuseppina Barrancotto (IRCCS Oasi Research Institute, Troina) and Frédéric Brau (Institute of Molecular and Cellular Pharmacology, Valbonne) for technical support. We thank Prof. Umberto Spampinato for helpful discussion and insightful comments.

References

- Anderson, P., Kedersha, N., 2002. Stressful initiations. *J. Cell Sci.* 115, 3227–3234.
- Antar, L.N., Afroz, R., Dichtenberg, J.B., Carroll, R.C., Bassell, G.J., 2004. Metabotropic glutamate receptor activation regulates fragile X mental retardation protein and FMR1 mRNA localization differentially in dendrites and at synapses. *J. Neurosci.* 11, 2648–2655. <https://doi.org/10.1523/JNEUROSCI.0099-04.2004>.
- Antar, L.N., Dichtenberg, J.B., Plociniak, M., Afroz, R., Bassell, G.J., 2005. Localization of FMRP-associated mRNA granules and requirement of microtubules for activity-dependent trafficking in hippocampal neurons. *Genes Brain Behav.* 6, 350–359. <https://doi.org/10.1111/j.1601-183X.2005.00128.x>.
- Arimoto, K., Fukuda, H., Imajoh-Ohmi, S., Saito, H., Takekawa, M., 2008. Formation of stress granules inhibits apoptosis by suppressing stress-responsive MAPK pathways. *Nat. Cell Biol.* 11, 1324–1332. <https://doi.org/10.1038/ncb1791>.
- Aronica, E., Gorter, J.A., Ijlst-Keizers, H., Rozemuller, A.J., Yankaya, B., Leenstra, S., Troost, D., 2003. Expression and functional role of mGluR3 and mGluR5 in human astrocytes and glioma cells: opposite regulation of glutamate transporter proteins. *Eur. J. Neurosci.* 17 (10), 2106–2118. <https://doi.org/10.1046/j.1460-9568.2003.02657.x>.
- Bardoni, B., Davidovic, L., Bensaid, M., Khandjian, E.W., 2006. The fragile X syndrome: exploring its molecular basis and seeking a treatment. *Expert Rev. Mol. Med.* 8 (8), 1–16. <https://doi.org/10.1017/S1462399406010751>.
- Bassell, G.J., Warren, S.T., 2008. Fragile X syndrome: loss of local mRNA regulation alters synaptic development and function. *Neuron* 2, 201–214. <https://doi.org/10.1016/j.neuron.2008.10.004>.
- Bechara, E.G., Didiot, M.C., Melko, M., Davidovic, L., Bensaid, M., Martin, P., Castets, M., Pogoniec, P., Khandjian, E.W., Moine, H., Bardoni, B., 2009. A novel function for fragile X mental retardation protein in translational activation. *PLoS Biol.* 7, 16. <https://doi.org/10.1371/journal.pbio.1000016>.
- Bonaccorso, C.M., Spatuzza, M., Di Marco, B., Gloria, A., Barrancotto, G., Cupode, A., Musumeci, S.A., D'Antoni, S., Bardoni, B., Catania, M.V., 2015. Fragile X mental retardation protein (FMRP) interacting proteins exhibit different expression patterns during development. *Int. J. Dev. Neurosci.* 42, 15–23. <https://doi.org/10.1016/j.ijdevneu.2015.02.004>.
- Buchan, J.R., Parker, R., 2009. Eukaryotic stress granules: the ins and out of translation. *Mol. Cell* 6, 932–941. <https://doi.org/10.1016/j.molcel.2009.11.020>.
- Catania, M.V., Landwehrmeyer, G.B., Testa, C.M., Standaert, D.G., Penney Jr., J.B., Young, A.B., 1994. Metabotropic glutamate receptors are differentially regulated during development. *Neuroscience* 61 (3), 481–495. [https://doi.org/10.1016/0306-4522\(94\)90428-6](https://doi.org/10.1016/0306-4522(94)90428-6).
- Catania, M.V., D'Antoni, S., Bonaccorso, C.M., Aronica, E., Bear, M.F., Nicoletti, F., 2007. Group I metabotropic glutamate receptors: a role in neurodevelopmental disorders? *Mol. Neurobiol.* 35 (3), 298–307. <https://doi.org/10.1007/s12035-007-0022-1>.

- Protter, D.S.W., Parker, R., 2016. Principles and properties of stress granules. *Trends Cell Biol.* 26 (9), 668–679. <https://doi.org/10.1016/j.tcb.2016.05.004>.
- Richter, J.D., Zhao, X., 2021 Feb 19. The molecular biology of FMRP: new insights into fragile X syndrome. *Nat. Rev. Neurosci.* <https://doi.org/10.1038/s41583-021-00432-0>. Epub ahead of print. 33608673.
- Shah, S., Molinaro, G., Liu, B., Wang, R., Huber, K.M., Richter, J.D., 2020. FMRP control of ribosome translocation promotes chromatin modifications and alternative splicing of neuronal genes linked to autism. *Cell Rep.* 30 (13), 4459–4472 e6. <https://doi.org/10.1016/j.celrep.2020.02.076>.
- Verkerk, A.J., Pieretti, M., Sutcliffe, J.S., Fu, Y.H., Kuhl, D.P., Pizzuti, A., Reiner, O., Richards, S., Victoria, M.F., Zhang, F.P., 1991. Identification of a gene (FMR-1) containing a CGG repeat coincident with a breakpoint cluster region exhibiting length variation in fragile X syndrome. *Cell* 65 (5), 905–914. [https://doi.org/10.1016/0092-8674\(91\)90397-h](https://doi.org/10.1016/0092-8674(91)90397-h).
- Wallingford, J., Scott, A.L., Rodrigues, K., Doering, L.C., 2017. Altered developmental expression of the astrocyte-secreted factors Hevin and SPARC in the fragile X mouse model. *Front. Mol. Neurosci.* 10, 268. <https://doi.org/10.3389/fnmol.2017.00268>.
- Weiler, I.J., Irwin, S.A., Klintsova, A.Y., Spencer, C.M., Brazelton, A.D., Miyashiro, K., Comery, T.A., Patel, B., Eberwine, J., Greenough, W.T., 1997. Fragile X mental retardation protein is translated near synapses in response to neurotransmitter activation. *Proc. Natl. Acad. Sci. U. S. A.* 105395–105400. <https://doi.org/10.1073/pnas.94.10.5395>.
- Weiler, I.J., Spangler, C.C., Klintsova, A.Y., Grossman, A.W., Kim, S.H., Bertina-Anglade, V., Khaliq, H., de Vries, F.E., Lambers, F.A., Hatia, F., et al., 2004. Fragile X mental retardation protein is necessary for neurotransmitter-activated protein translation at synapses. *Proc. Natl. Acad. Sci. U. S. A.* 5017504–5017509. <https://doi.org/10.1073/pnas.0407533101>.