Contents lists available at ScienceDirect

Psychiatry Research

journal homepage: www.elsevier.com/locate/psychres

The potential role of oxytocin and perinatal factors in the pathogenesis of autism spectrum disorders – review of the literature



Psychiatry Research

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ARTICLE INFO

Keywords: Autism spectrum disorder, behavior Gene Cesarean section, oxytocin, oxytocin-receptor polymorphism

ABSTRACT

Autism Spectrum Disorders (ASD) are characterized by: social and communication impairments, and by restricted repetitive behaviors. The aim of the present paper is to review abnormalities of oxytocin (OXT) and related congenital malformations in ASD. A literature search was conducted in the PubMed database up to 2016 for articles related to the pathomechanism of ASD, abnormalities of OXT and the OXT polymorphism in ASD. The pathomechanism of ASD has yet to be. The development of ASD is suggested to be related to abnormalities of the oxytocin-arginin-vasopressin system. Previous results suggest that OXT and arginine vasopressin (AVP) may play a role in the etiopathogenesis of ASD.

1. Introduction

Autism spectrum disorder was previously perceived as a rare disorder, but increasing recognition, understanding, and awareness is changing this perception, and it is now considered to be a relatively common condition. Prevalence rate estimates vary between 1% and 2% in school age children (Ben-Ari, 2015).

ASD has a great impact on the quality of life in the affected children and their family. ASD seems to be a multifactorial disease with the interaction of genetic and environmental factors. Evidence supporting the hypothesis that perinatal outcome can play a role in the etiology of is ASD increasing. The present paper focuses on recent publications which have demonstrated the role of oxytocin, and prenatal and perinatal factors associated with an increased risk of ASD (Fakhoury, 2015; Muotri et al., 2016). In this paper, several methodological advantages have helped to support and extend this hypothesis.

2. Methods

2.1. Search strategy

Published data were used in our paper; therefore, no ethical approval was required.

The authors searched for the terms 'ASD' OR 'perinatal outcome' OR 'oxytocin' in the following databases: Medline, Pubmed, Scopus, Index Medicus, and Science Direct.

The following outcome measures were analyzed (1) oxytocin and oxytocin receptor defect in the development of ASD; (2) the relative risk of perinatal factors and congenital defects.

- Qualitative data: after screening the title and abstract, all texts not presenting data on oxytocin-receptor related to ASD were discarded.
- Quantitative data: only papers presenting original work with analysis on the relationship between oxytocin; cesarean section, perinatal outcome, and development of ASD were included. Reference lists from these papers were read to search for other possible relevant papers. Abstracts of scientific meetings, review papers, anecdotal case reports, duplicate papers, and editorials were excluded.

3. Results

3.1. The role of oxytocin

Social and communication difficulties are the most characteristic disturbances in ASD, and they are two of the core symptom domains. Repetitive behavior, and restricted activities and interests constitute the third domain of symptoms, as listed in the DSM-IV (American Psychiatric Association, 2000).

Oxytocin (OXT) is a nine-amino-acid peptide that is synthesized in the paraventricular and supraoptic nucleus of the hypothalamus and released into the bloodstream by axon terminals in the posterior pituitary (Lee et al., 2009). The best known effects in women of

http://dx.doi.org/10.1016/j.psychres.2016.12.007

Received 9 February 2016; Received in revised form 25 October 2016; Accepted 5 December 2016 Available online 06 December 2016 0165-1781/ © 2016 Elsevier Ireland Ltd. All rights reserved.

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reproductive age are in stimulating uterine contractions during labor and in regulating lactation. Furthermore, it is important to emphasize that OXT and the structurally similar peptide AVP are directly released within the brain, where they play a key role in regulating social affiliative behaviors, including sexual behavior, and social recognition (Lee et al., 2009; Harony, 2010; Caldwell, 2008). Besides these functions, the genes coding for the production of OXT and AVP are located on the same chromosome, 20p13, while the genes for oxytocin receptor (OXTR) are located on chromosome 3. OXT plays a role in increasing trust, empathy, eye contact, face memory, and social interaction (Donaldson and Young, 2008). AVP, which is well known for its antidiuretic effect, influences social behavior as well (Harony and Wagner, 2010). Repetitive behaviors are possibly influenced by OXT and AVP since they act as neuromodulators, targeting the hippocampus, amygdala, striatum, hypothalamus, and nucleus accumbens (Insel, 2010; Israel, 2008; Green, 2010). Lastly, it is through their neuromodulatory role that OXT and AVP are implicated in other behaviors such as stress-response, learning, and memory (Donaldson ZR, 2008). Overall, they play a critical role in regulating social behavior from the early phases of development, and thus are likely to be implicated in ASD unfolding. For more than a decade, several authors and teams have assessed the possibility that OXT may be influenced in the social symptoms of ASD (Insel et al., 2010; Green et al., 2010; Israel et al., 2008). Abnormalities in OXT/AVP in ASD might be due to different mechanisms, such as a reduction in hormonal levels as a result of diminished synthesis, an alteration in hormonal release, or insufficient amounts of these hormones as neuromodulators in the brain possibly because of receptor abnormalities (Israel et al., 2008; Gregory et al., 2009). Furthermore, it has been suggested that an alteration in hormonal release might be secondary to an accelerated synaptic pruning, and/or alteration of the glia (Wang and Hatton, 2009). Several neuronal network, neurotransmission and hormone systems are probably involved in ASD. Currently, neural systems of the amygdala, the mesocorticolimbic dopamine system and the OXT system are among the most important ones (Neuhaus et al., 2010). The relevance of OXT as implicated in the facilitation of social information processing is possibly obtained through auditory and visual pathways. Finally, OXT is involved in regulating stress and fear reactivity, possibly promoting social interaction in individuals with ASD that often experience anxiety associated with nuclear disturbances in social interaction (Bartz et al., 2008).

3.2. Genetic studies

Genetic studies also suggested a role of OXT in ASD. The region of chromosome 3, in which OXTR is located is now considered a susceptibility locus, as it has repeatedly been found to be implicated in ASD (Liu et al., 2016; Jansen et al., 2006; Ebstein et al., 2009; Andari et al., 2010; Green et al., 2001; Guastella et al., 2010; Wang and Hatton, 2009; Sala et al., 2011; Schmidt et al., 2012; Guastella et al., 2008; Kurth et al., 2011). Wu et al. have reported an association between two single nucleotide polymorphisms (SNP) in the OXTR related to the occurrence of ASD (Wu et al., 2005). Jacob et al. have also found an association between OXTR and autism; specifically, a significant association was detected at rs2254298, but in contrast with Wu et al., not at rs53576 (Jacob, 2007).

Similarly, Lerer et al. have shown that SNPs and haplotypes in the OXTR gene confer risk for ASD, and in addition, they have observed an association with IQ and adaptive functions measured with total Vineland Adaptive Behaviors Scales (communication, daily living skills, and socialization subdomains) (Lerer et al., 2008). Recently, a study with a large cohort of 2333 individuals with ASD has demonstrated an association between three markers implicated in ASD susceptibility, rs2268493, rs1042778, and rs7632287, further strengthening the hypothesis that functional polymorphisms of OXTR may contribute to ASD risk (Campbell et al., 2011). Similar conclusions have been

drawn in another study by Wertmer et al. observing a significant genetic variation in the OXTR gene in patients with ASD including atypical and high-functioning individuals (Wermter et al., 2010). These findings point to an epigenetic dysregulation of OXTR. How this process might unfold is currently unknown, but it is important to note that if further evidence is discovered, it might explain the failure to identify sequence alterations in many candidate genes of ASD, e.g., an epigenetic alteration would be implicated in ASD phenotype expression (Gurrieri et al., 2009).

3.3. Perinatal factors

ASD is influenced by a variety of genetic, environmental, and possibly immunological factors that act during critical periods to alter key developmental processes. Some pharmaceutical drugs, toxicants, and metabolic and nutritional factors have been identified in epidemiological studies to increase the risk for autism, especially during the prenatal period. Immunological risk factors including maternal infection during pregnancy, autoantibodies to fetal brain proteins, and familial autoimmune diseases have consistently been observed in multiple studies, as well as immune abnormalities in individuals with ASD.

Etiological hypotheses suggest a strong genetic component and a possible interaction between the epigenetic factors in early fetal development (Kolevzon et al., 2007).

The suggested environmental factors include nutritional and immune function related risk factors such as lack of vitamin D and folic acid supplementation, and metabolic syndrome (Gialloreti et al., 2014). It can be concluded that the rate of cesarean sections was also significantly higher in the case group than in the control group (Kolevzon et al., 2007; Gialloreti et al., 2014; Curran et al., 2015). Shieve et al. have concluded that three perinatal risk factors (preterm birth, small for gestational age, and cesarean section) notably contribute to ASD risk (Shieve et al., 2014). Nilsen et al. (2013) analyzed clinical data on a large cohort of pregnant women in 2014 concerning the perinatal outcomes and ASD. The results suggest that cesarean section, prenatal folic acid use, maternal smoking in the prenatal period, low birth weight and preterm birth have increased the risk of developing ASD.

3.4. Congenital anomalies

Earlier population-based register studies on the association between ASD and syndromes are scanty, as the syndromes are rare (Lee et al., 2015).

Kielinen et al. have reported a population based study of 187 ASD children with one Klinefelter syndrome (47, XXY), (0.5%) one 47/XYY (0.5%, one Down syndrome (0.5%), four Fragile X syndromes (2.1%) two fetal alcoholic syndrome (1.1%), and one Sotos syndrome (0.5%, one tuberous sclerosis (0.5%) (Kielinen et al., 2004).

4. Discussion

Considering the complex characteristics of ASD, which are a combination of biologic heterogeneity and phenotypic heterogeneity, it is difficult to choose and develop an effective treatment. Since oxytocin is responsible for not only maternal behavior but also core symptom domains of ASD, such as social interaction and repetitive behaviors, association with the pathogenesis of ASD is highly possible. There are various ongoing animal, human, and clinical studies assessing the potential disruption of oxytocin function or secretion leading to the etiology of ASD. SNP of the OXTR is the first step in the recognition of new treatment molecules. Through clinical trials applying oxytocin in patients with ASD, future clinical applications are promising. Therefore, further research should be carried out to develop treatment options for children with ASD (Ben-Ari, 2015; Curran et al.,

2015; Egawa et al., 2013).

Funding statement

N.a.

Conflict of interest

The authors have no conflict of interest.

Acknowledgement

No.

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