Making a case for the role of DNA repair in autism Aditi Nadkarni, Ph.D

Department of Biology, New York University, 100 Washington Square East, 1009 Silver Center, New York, NY 10003-6688, USA Email: an60@nvu.edu

Abstract

Autism spectrum disorder (ASD) is a neurodevelopmental condition characterized by behavioral and social impairments and genetic predisposition. The genetic heterogeneity of autism and the complexity of the disease have made it difficult to delineate one mechanism or diagnostic biomarker that influences ASD risk. Mounting evidence points towards exposure to DNA damaging environmental pollutants as a major ASD risk factor. Studies have also identified genomic instability, defective DNA repair and greater oxidative DNA damage using ASD animal models and patient samples. This review highlights the salient findings that connect defective DNA repair with ASD risk and makes a case for the need of functional DNA repair assays characterizing the role of DNA repair mechanisms in autism etiology.

Keywords: autism, ASD, DNA damage, DNA repair, environmental pollutants, polycyclic aromatic hydrocarbons

Introduction

Autism spectrum disorder (ASD) is characterized complex neurodevelopmental by and behavioral dysfunction along with social and communication impairment [1]. While ASD is diagnosed by clinical, psychological and behavioral observation, there are no currently available biomarkers or genetic tests used for detection or confirmation of diagnosis [2]. Although a genetic basis of the disease has been widely accepted, the complexity of the disease poses a challenge in delineating the genetic components that contribute towards autism initiation and severity [3].

Several studies using linkage analysis, bioinformatics and high throughput genomic strategies have attempted to identify the neurobiological basis of this disease [4-6]. Although a majority of these early studies focused on genetic determinants associated with central nervous system defects, current research is veering focus towards factors such as exposure to environmental toxins and systemic genetic defects that may influence response to these toxins but may not

necessarily be directly involved in CNS development [7-12]. This increased association between autism and exposure to environmental pollutants bears striking resemblance to the influence of exposure to carcinogens on elevated cancer risk [13-16]. Moreover, a lot of these environmental pollutants such as cigarette smoke, exhaust fumes and oxidative stress elicit their cellular effects through DNA damage, which is also a mechanism of carcinogenesis [12, 15-19]. In the present article, I highlight studies that provide evidence for the association between DNA repair and ASD risk. This review aims to summarize the potential correlation between genome protective DNA repair mechanisms with the neurological and genetic observations made in ASD patients and to discuss further strategies in delineating this association for diagnostic or therapeutic purposes.

DNA repair in the brain

The structural effects of DNA damage are highly diverse and a single repair process cannot cope with the multiple types of DNA lesions [20, 21]. Hence, specific recognition and repair mechanisms operate based on the type of DNA damage. For instance, bulky adducts formed by a common class of environmental pollutants called polycyclic hydrocarbons is repaired by the nucleotide excision repair pathway (NER). Homologous recombination (HR) and nonhomologous end-joining (NHEJ) are the two main pathways involved in repairing doublestranded DNA damage and act differentially depending on cell cycle phase [22-26]. The MMR pathway plays an essential role in the correction of base-base mismatches that result from the misincorporation of nucleotides and BER targets small chemical alterations of the DNA bases, which generate miscoding lesions possibly resulting from cell metabolic events or single-strand breaks [21]. From the context of DNA repair mechanisms involved in neural development, the early cortical progenitors are hypersensitive to replication-associated DNA damage making mechanisms such as HR and MMR critical in repair during this stage [27, 28]. In the mature brain, on the other hand, cells are more susceptible to oxidative damage and may hence employ the BER or NER pathways [29-31]. The neurodevelopmental changes in autism correspond are thought to to early embryogenesis prenatal environmental or influences [32].

DNA repair defects, environmental pollutants and ASD risk

ASDs and intellectual disability often overlap: it is estimated that 70% of individuals with autism also have an ID. Both ASDs and ID are defined behaviorally and their onset is associated with neurological development. Additionally, both ASD and ID are associated with de novo copy number variants (CNV) a form of genetic instability that was recently associated with error-prone DNA repair mechanisms. In 2009, Dorus et al concluded that autosomal genes involved in DNA repair (FANCA, FANCC, NBS1, demonstrated and XRCC8), significantly elevated frequencies of positive Darwinian selection compared to other categories in ID [33]. Most recently, Maria Karayiorgou's group

at Columbia University used a family-based cluster detection approach designed to localize significant rare disease-risk variants clusters within the chromosome 15q13.3. Unlike other previously used methods which identify variants only within a specific gene, Karayiorgou's group used a scan statistic approach which identifies association and clustering of variants in a small window of a larger genetic region by analyzing the findings from two independent whole exome sequencing data sets. Using this rigorous methodology, they implicated a region mapping to a 20kb window of the Fanconi-associated nuclease 1 gene (FAN1) in autism susceptibility. FAN1 functions within the Fanconi Anemia pathway that is involved in DNA repair of interstrand crosslinks and replicative DNA damage. This latest finding is in line with a previous study by Peprah et al. which reported significant down regulation of DNA а damage/repair pathway transcripts in the expression profiles of autistic males due to a Fragile X mutation. Since Fragile X syndrome is currently the leading genetic cause of autism, this study was a significant indication of the relevance of DNA repair pathways in the genetics of autism. In 2011, Kinney et al discussed the puzzling association between high autism rates and hypomelanotic skin disorders suggesting that UV-B protection and vitamin D deficiency may play a large role in influencing autism risk [34]. While this review briefly mentioned the role of vitamin D in DNA repair, much remains to be understood about the precise pathway by which vitamin D in regulates DNA repair or nucleotide excision repair in particular since it is the primary pathway involved in repairing of UV-B induced DNA damage [35, 36]. This is especially important in the context of autism because of accumulating evidence that supports the association between vitamin D deficiency and ASD risk as well as at least one report of an autistic patient with an XPC splice site mutation [37, 38]. XPC is

required for genomic nucleotide excision repair and a long-term deficiency in XPC reportedly is also associated with suppressed double stranded break repair in combination with other genetic defects. A hallmark study published in Nature reported that that inhibition of topoisomerases adversely affects the expression of long genes, including numerous long genes associated with synaptic function and ASD [39]. These findings make it imperative to evaluate two alternative hypotheses 1) environmental toxins result in mutations within DNA repair genes such as topoisomerases causing widespread genome instability that influences specific genetic hotspots or 2) inherent repair deficiencies in autistic individuals makes them more susceptible to the DNA damage inflicted by environmental pollutants resulting in the downstream genetic mutations.

An analysis of 100 million US medical records recently revealed that autism and intellectual disability (ID) rates are correlated with an increased incidence of genital malformations in newborn males which is also an indicator of possible congenital exposure to harmful environmental factors [40]. Although DNA defects vary for ASD cases, Courchesne et al, using brain tissue samples from ASD samples discovered a shared set of overlapping gene pathways that included DNA damage detection and repair as one of the major mechanisms influencing dysregulation of early brain development [41]. This finding is further supported by Jeggo et al. where the authors theorize that high levels of oxidative DNA damage during neuronal development would be consistent with roles for HR, ATR-signaling and NHEJ proteins in protecting proliferating cells from damage [42]. In an excellent review published in Nature Neuroscience, Dr. Peter McKinnon makes the case for the role of various DNA repair pathways in maintaining neural homeostasis especially during early embryonic development when neurogenesis is driven by proliferation and cells are most susceptible to replicative or environmental DNA damage [27].

Perspective

Despite mounting evidence for the role of aberrant DNA repair mechanisms in autism, the general consensus within the field remains that autism is a highly complex disorder that may involve several gene loci, low penetrance alleles and gene-environment interactions that define its etiology thus complicating diagnosis. While the focus of current studies remains on the means to identify single genetic biomarkers or cytogenetic characteristics for diagnosis, there are no reports about the impact of DNA repair defects and exposure to environmental toxins using high-throughput functional assays on DNA repair capacity during neuronal development associated with ASD risk. Several genes are involved in the maintenance of genome integrity in response to environmental pollutants, and there is great variation amongst individuals with respect to common polymorphisms that impact on the activity of these enzymes. Hence there is a need for molecular diagnostics that can a) identify the aberrant DNA repair mechanism and b) identify the stage of neurodevelopment where the repair deficiency is most critical in influencing ASD risk. These functional repair assays can examine pre/peri or gestational exposure to DNA damaging environmental pollutants in place of biomarkers of these toxins which have thus far resulted in inconsistent findings [43]. While Fenech et al demonstrated that autistic individuals are not abnormally susceptible to oxidative stress compared to their non-autistic siblings, the authors did acknowledge that there was a need for more DNA repair markers and assays to test the differential effects of DNA damage in vivo on autistic neural cells [44].

It is also tempting to speculate if reduced maternal DNA repair capacities combined with environmental factors can increase autism risk. Studies using brain tissue or neurons obtained from patients can be subjected to highthroughput DNA repair assays that are currently being used in the cancer genetics field to assess repair defects that contribute towards cancer risk or therapeutic outcome [45-51]. Similarly, the impact of vitamin D or nutritional element on repair efficiency or protein kinetics is worth examining in cellular models of autism. The advent of somatic cell reprogramming now makes it possible to study ASD at a cellular level using pluripotent patient-derived cells [52-54]. This model offers an excellent tool to study the effects of various forms of DNA damage and the DNA repair capacity of cells derived from ASD versus normal cohorts. If a repair deficiency is identified, this cell model can also enable screening of phytochemicals or genes that can correct this defect.

Summary

In conclusion, the pattern of DNA repair pathways and defects emerging repeatedly in assessing autism susceptibility is highly suggestive of a role for DNA repair mechanisms in autism risk. The biological connection between DNA repair and autism deserves further analysis considering the now wellestablished link between environmental DNA damage, down-regulated DNA repair pathways and ASD risk in available literature and recent Since dysfunctional reports. cellular mechanisms of genomic stability during early embryogenesis and neural development can result in a plethora of genetic mutations and brain abnormalities, it is important to evaluate whether the link between environmental pollution and ASD is driven by DNA damage and exacerbated by reduced DNA repair capacity.

Acknowledgement

The author acknowledges Moinuddin Chowdhury for his critical reading of the review.

References

- Compart, P.J., *The Pathophysiology of Autism.* Glob Adv Health Med, 2013.
 2(6): p. 32-37.
- 2. Voineagu, I. and H.J. Yoo, *Current* progress and challenges in the search

for autism biomarkers. Dis Markers, 2013. **35**(1): p. 55-65.

- Veenstra-Vanderweele, J., S.L. Christian, and E.H. Cook, Jr., Autism as a paradigmatic complex genetic disorder. Annu Rev Genomics Hum Genet, 2004.
 5: p. 379-405.
- Yonan, A.L., et al., Bioinformatic analysis of autism positional candidate genes using biological databases and computational gene network prediction. Genes Brain Behav, 2003. 2(5): p. 303-20.
- Risch, N., et al., A genomic screen of autism: evidence for a multilocus etiology. Am J Hum Genet, 1999. 65(2): p. 493-507.
- Poot, M., Towards identification of individual etiologies by resolving genomic and biological conundrums in patients with autism spectrum disorders. Mol Syndromol, 2013. 4(5): p. 213-26.
- Reiss, A.L., C. Feinstein, and K.N. Rosenbaum, Autism and genetic disorders. Schizophr Bull, 1986. 12(4): p. 724-38.
- Chien, Y.L., et al., Association study of the CNS patterning genes and autism in Han Chinese in Taiwan. Prog Neuropsychopharmacol Biol Psychiatry, 2011. 35(6): p. 1512-7.
- 9. Gray, S.J., *Gene therapy and neurodevelopmental disorders.* Neuropharmacology, 2013. **68**: p. 136-42.
- McGinnis, W.R., Oxidative stress in autism. Altern Ther Health Med, 2004.
 10(6): p. 22-36; quiz 37, 92.
- 11. Melnyk, S., et al., *Metabolic imbalance* associated with methylation dysregulation and oxidative damage in children with autism. J Autism Dev Disord, 2012. **42**(3): p. 367-77.
- 12. Kinney, D.K., et al., Environmental risk factors for autism: do they help cause de novo genetic mutations that contribute to the disorder? Med

Hypotheses, 2010. **74**(1): p. 102-6.13. Crespi, B., *Autism and cancer risk*. Autism Res, 2011. **4**(4): p. 302-10.

- Windham, G.C., et al., Autism spectrum disorders in relation to distribution of hazardous air pollutants in the san francisco bay area. Environ Health Perspect, 2006. 114(9): p. 1438-44.
- 15. Kim, S.M., et al., *Exposure to environmental toxins in mothers of children with autism spectrum disorder.* Psychiatry Investig, 2010. **7**(2): p. 122-7.
- 16. Sheng, L., et al., *Prenatal polycyclic aromatic hydrocarbon exposure leads to behavioral deficits and downregulation of receptor tyrosine kinase, MET.* Toxicol Sci, 2010. **118**(2): p. 625-34.
- 17. Kalkbrenner, A.E., et al., *Perinatal exposure to hazardous air pollutants and autism spectrum disorders at age 8.* Epidemiology, 2010. **21**(5): p. 631-41.
- Rose, S., et al., Evidence of oxidative damage and inflammation associated with low glutathione redox status in the autism brain. Transl Psychiatry, 2012. 2: p. e134.
- Lu, J.M., et al., Chemical and molecular mechanisms of antioxidants: experimental approaches and model systems. J Cell Mol Med, 2010. 14(4): p. 840-60.
- Iyama, T. and D.M. Wilson, 3rd, DNA repair mechanisms in dividing and non-dividing cells. DNA Repair (Amst), 2013.
 12(8): p. 620-36.
- 21. Slupphaug, G., B. Kavli, and H.E. Krokan, *The interacting pathways for prevention and repair of oxidative DNA damage.* Mutat Res, 2003. **531**(1-2): p. 231-51.
- Pardo, B., B. Gomez-Gonzalez, and A. Aguilera, DNA repair in mammalian cells: DNA double-strand break repair: how to fix a broken relationship. Cell Mol Life Sci, 2009. 66(6): p. 1039-56.
- 23. Puchta, H., *The repair of double-strand breaks in plants: mechanisms and consequences for genome evolution.* J Exp Bot, 2005. **56**(409): p. 1-14.

- 24. Thacker, J. and M.Z. Zdzienicka, *The XRCC genes: expanding roles in DNA double-strand break repair.* DNA Repair (Amst), 2004. **3**(8-9): p. 1081-90.
- Pfeiffer, P., et al., Pathways of DNA double-strand break repair and their impact on the prevention and formation of chromosomal aberrations. Cytogenet Genome Res, 2004. 104(1-4): p. 7-13.
- Pastwa, E. and J. Blasiak, Nonhomologous DNA end joining. Acta Biochim Pol, 2003. 50(4): p. 891-908.
- McKinnon, P.J., Maintaining genome stability in the nervous system. Nat Neurosci, 2013. 16(11): p. 1523-9.
- Barzilai, A. and P.J. McKinnon, Genome maintenance in the nervous system; insight into the role of the DNA damage response in brain development and disease. DNA Repair (Amst), 2013.
 12(8): p. 541-2.
- Englander, E.W., DNA damage response in peripheral nervous system: coping with cancer therapy-induced DNA lesions. DNA Repair (Amst), 2013. 12(8): p. 685-90.
- Jaarsma, D., et al., Age-related neuronal degeneration: complementary roles of nucleotide excision repair and transcription-coupled repair in preventing neuropathology. PLoS Genet, 2011. 7(12): p. e1002405.
- Sykora, P., D.M. Wilson, 3rd, and V.A. Bohr, Base excision repair in the mammalian brain: implication for age related neurodegeneration. Mech Ageing Dev, 2013. 134(10): p. 440-8.
- Ploeger, A., et al., The association between autism and errors in early embryogenesis: what is the causal mechanism? Biol Psychiatry, 2010.
 67(7): p. 602-7.
- Crespi, B.J., The origins and evolution of genetic disease risk in modern humans. Ann N Y Acad Sci, 2010. 1206: p. 80-109.
- 34. Khan, S.G., et al., Xeroderma pigmentosum group C splice mutation

associated with autism and hypoglycinemia. J Invest Dermatol, 1998. **111**(5): p. 791-6.

- Nair-Shalliker, V., B.K. Armstrong, and M. Fenech, *Does vitamin D protect against DNA damage?* Mutat Res, 2012.
 733(1-2): p. 50-7.
- Kallay, E., et al., Characterization of a vitamin D receptor knockout mouse as a model of colorectal hyperproliferation and DNA damage. Carcinogenesis, 2001. 22(9): p. 1429-35.
- Cannell, J.J. and W.B. Grant, What is the role of vitamin D in autism? Dermatoendocrinol, 2013. 5(1): p. 199-204.
- 38. Cannell, J.J., *Autism and vitamin D.* Med Hypotheses, 2008. **70**(4): p. 750-9.
- King, I.F., et al., Topoisomerases facilitate transcription of long genes linked to autism. Nature, 2013.
 501(7465): p. 58-62.
- 40. Rzhetsky, A., et al., Environmental and state-level regulatory factors affect the incidence of autism and intellectual disability. PLoS Comput Biol, 2014.
 10(3): p. e1003518.
- 41. Courchesne, E., et al., *Mapping early brain development in autism*. Neuron, 2007. **56**(2): p. 399-413.
- 42. O'Driscoll, M. and P.A. Jeggo, *The role of the DNA damage response pathways in brain development and microcephaly: insight from human disorders.* DNA Repair (Amst), 2008. **7**(7): p. 1039-50.
- 43. Rossignol, D.A., S.J. Genuis, and R.E. Frye, *Environmental toxicants and autism spectrum disorders: a systematic review.* Transl Psychiatry, 2014. **4**: p. e360.
- 44. Main, P.A., et al., *Necrosis is increased in lymphoblastoid cell lines from children with autism compared with their non-autistic siblings under conditions of oxidative and nitrosative stress.* Mutagenesis, 2013. **28**(4): p. 475-84.

- Au, W.W., A.K. Giri, and M. Ruchirawat, Challenge assay: A functional biomarker for exposure-induced DNA repair deficiency and for risk of cancer. Int J Hyg Environ Health, 2010. 213(1): p. 32-9.
- 46. Forestier, A., et al., Functional DNA repair signature of cancer cell lines exposed to a set of cytotoxic anticancer drugs using a multiplexed enzymatic repair assay on biochip. PLoS One, 2012. **7**(12): p. e51754.
- 47. Chua, M.L., et al., DNA double-strand break repair and induction of apoptosis in ex vivo irradiated blood lymphocytes in relation to late normal tissue reactions following breast radiotherapy. Radiat Environ Biophys, 2014.
- Guidugli, L., et al., Functional assays for analysis of variants of uncertain significance in BRCA2. Hum Mutat, 2014. 35(2): p. 151-64.
- Birkelbach, M., et al., Detection of impaired homologous recombination repair in NSCLC cells and tissues. J Thorac Oncol, 2013. 8(3): p. 279-86.
- 50. Liu, R., et al., *Melatonin enhances DNA* repair capacity possibly by affecting genes involved in DNA damage responsive pathways. BMC Cell Biol, 2013. **14**: p. 1.
- 51. Wang, Y., et al., Modulation of DNA repair capacity by ataxia telangiectasia mutated gene polymorphisms among polycyclic aromatic hydrocarbonsexposed workers. Toxicol Sci, 2011. **124**(1): p. 99-108.
- 52. Aigner, S., et al., Human pluripotent stem cell models of autism spectrum disorder: emerging frontiers, opportunities, and challenges towards neuronal networks in a dish. Psychopharmacology (Berl), 2013.
- Yuan, S.H. and M. Shaner, Bioengineered stem cells in neural development and neurodegeneration research. Ageing Res Rev, 2013. 12(3): p. 739-48.

54. Chailangkarn, T., A. Acab, and A.R. Muotri, *Modeling neurodevelopmental disorders using human neurons.* Curr Opin Neurobiol, 2012. **22**(5): p. 785-90.