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Beta-2-Microglobulin in Autism Spectrum Disorders

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Abstract: Autism spectrum disorders (ASD) are heterogeneous neurodevelopmental diseases of unknown etiology. There are no biological markers for ASD and current diagnosis is based on behavioral criteria. Recent data has shown that MHC I, a compound involved in adaptive immune function, is also involved in neurodevelopment, synaptic plasticity and behavior. It has been suggested that altered MHC I expression could play a part in neurodevelopmental diseases like ASD. To address this possibility, we measured plasma levels of beta-2-microglobulin (β 2m), a molecule that associates with MHC I and is indicative of MHC I expression, in 36 children with autism, 28 typically developing controls and subjects with developmental disabilities (n=16) but not autism. The age range of our study population was 17-120 months. We found no statistically significant differences in plasma β 2m levels between groups. Therefore, plasma levels of β 2m measured in early childhood in autism may not reflect changes in MHC class I in autism.

Key words: MHC class I, autism, immune system

INTRODUCTION

Autism Spectrum Disorders (ASD) are a group of neurodevelopmental diseases characterized by a range of social difficulties, deficits in communication and restricted stereotypic behaviors and interests^[1, 2]. The etiology of ASD is unknown and may involve genetic, environmental and immune factors. There are no known definitive biological markers for ASD or its phenotypic variations and current diagnosis is based on behavioral criteria. In the United States, the incidence of ASD has risen to an estimated 1:150 children^[3]; a finding which has hastened the search for relevant biological markers associated with these disorders.

Several studies have suggested a potential link between ASD and immune system dysfunction (reviewed in^[4]. Issues such as increased inflammation in the brain^[5,6], autoantibodies to brain tissues (reviewed in^[14, 7-11]) and maternal/perinatal infections (^[12,13], reviewed in^[14]) have been associated with the development of ASD. Immune system related genes have also been linked to ASD, including the complement component C4B gene and the extended HLA haplotype B44-S30-DR4 (^[15-16]). Due to the extensive crosstalk between the immune system and central nervous system (CNS) during development and beyond^[17-19], it is conceivable that aberrant immune activity could be detrimental to CNS function.

The major histocompatibility complex (MHC) encodes several genes that play a central role in adaptive immune responses. In addition to its immunological functions, recent evidence suggests that MHC I expressed on neuronal cells may also be important for brain development and function (reviewed in^{[18, 20])}. It has been shown in several studies that MHC I is crucial for synaptic plasticity in the adult mammalian brain^[21-23]. developing and Furthermore, it has been suggested that deficiencies in MHC I expression may play a role in neurological diseases including schizophrenia (reviewed in^[24] and autism^[18]. B-2- microglobulin (B2m) is an essential component of MHC I and can be detected in the serum in lieu of directly measuring cellular expression of MHC I^[25]. In the current study, we measured levels of β 2m in the sera of subjects with ASD compared to children with developmental delays but not autism (DD) and typically developing controls from the general population (TD). We further divided the ASD group based on phenotypic differences in the onset of the disease (early onset-EO and regressive)^[26], since differing forms of ASD may have distinct biomarkers. We hypothesized that children with autism would have

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lower plasma levels of $\beta 2m$; a finding which may be responsible for aspects of the disorder.

MATERIALS AND METHODS

Subjects and samples: This study examined 79 children enrolled through the M.I.N.D. (Medical Investigations of Neurodevelopmental Disorders) Institute Clinic as part of the ongoing CHARGE Autism Risk from Genetics (Childhood and Environment) study at UC Davis^[27]. The sample population consisted of children with early onset autism (AU-EO; n=19, 2 females, 17 males, ages 17-63 months), regressive autism (AU-reg; n=17, 3 females, 14 males, ages 28-60 months), a control group of agematched typically developing children (TD; n=28, 10 females, 18 males, ages 17-120 months) and children with developmental disabilities (DD) without autism (n=16; 5F, 11M, ages 12-54 months).

All children were assessed at the UC Davis M.I.N.D. Institute. Autism was confirmed using the Autism Diagnostic Interview-Revised (ADI-R)^[28, 29] and the Autism Diagnostic Observation Schedule, modules 1, 2 and 3 (ADOS) (^[30-32]). The ADI-R is a standardized semi-structured interview that provides a diagnostic algorithm for the DSM-IV^[33] and the ICD-10 definitions of $autism^{[34, 35]}$. The ADOS is a standardized, semi-structured assessment in which the researcher observes the social behavior, communication and imaginative use of materials for children suspected of having ASD. A diagnosis of autism was defined as meeting criteria on the communication, social and repetitive behaviors domains of the ADI-R and scoring at or above the cut off for autistic disorder on the ADOS module 1 or 2.

The Social Communication Questionnaire was used to screen for behavioral and developmental characteristics of ASD among subjects with developmental disabilities and typically developing controls. Children who scored above the screening cutoff were fully assessed using the ADI-R and ADOS.

The children with autism were further subdivided into children who initially developed normally, reaching typical developmental milestones before regressing and losing language and social skills and those who had early impairments in the development of language and social skills. A classification of regression was based on clinical characteristics using both parental reporting and answers to questions regarding language loss (Q11) and social skills (Q25) of the ADI-R. Our autism study population could be classified into 19 subjects with early onset autism (classical) and 17 subjects with delayed-onset autism (regression). The study protocol followed the ethical guidelines of the most recent Declaration of Helsinki and was approved by the Institutional Review Boards of the UC Davis School of Medicine and the State of California and all subjects enrolled in the study had written informed consent provided by their parents and assented to participate if developmentally able.

Blood samples were collected into yellow top citrate-containing tubes and plasma was isolated by centrifugation at 2300 rpm for 10 minutes. Plasma aliquots were stored at -80° C until use.

Levels of β-2-microglobulin were measured using ELISA kits (Bioquant) according to the manufacturer's protocols. Briefly, samples were diluted 1:100 in the provided diluent. 20 µL of sample or standard and 200 uL of diluent were pipetted in duplicate into the appropriate pre-coated wells and incubated at 37°C for 30 minutes. Plates were washed 5 times before addition of 200 µL of enzyme conjugate reagent. Following a 30-minute incubation, the wash step was repeated and 100 µL of TMB was added. After 20 minutes, a stop solution was added to the wells and absorbance was read at 450nm. Concentrations were calculated using a standard curve plotting absorbance versus standard concentration. Sample concentrations were calculated with the equation derived from the curve. Statistical differences between groups were determined using Student's two-tailed t-test for samples with unequal variances. Values were considered significant if p was less than 0.05.

RESULTS AND DISCUSSION

We found no statistically significant differences in β -2-microglobulin levels between AU (median value: 1.071 µg/ml, interquartile range: 0.830-1.311), age matched DD (1.312, 0.793-1.943) and age matched TD controls (0.952, 0.826-1.387) (Fig. 1a). Based on autism onset patterns, we further divided the AU group into early onset (EO) (0.908, 0.722-1.509) and regressive (0.841, 0.718-1.322) phenotypes^[26] and similarly found no difference between groups (Fig. 1b).

The current study explores a possible connection between ASD and MHC I expression as represented by serum levels of β -2-microglobulin (β 2m). We chose to investigate this potential link in response to research suggesting that neuronal MHC I expression is required for proper neurodevelopment and synaptic plasticity (reviewed in ^[18, 19]).

For years it was thought that neuronal cells did not express MHC I under normal conditions. This was one



Fig. 1a: Beta-2-microglobulin levels in AU, TD and DD subjects (A) Scatter plot representing levels of β-2-microglobulin in subjects with AU compared to TD and DD, bars represent median values

of many factors believed to contribute to the "immuneprivileged" status of the brain. However, several studies have now shown that MHC I is in fact expressed by healthy neurons^[21,22,36]</sup>. Furthermore, MHC I has</sup>several non-immunological roles and is involved in several aspects of neuronal development and function. First, MHC I gene expression levels have been linked to activity-dependent plasticity in the visual system^[21]. This finding was corroborated in a study with MHC I deficient mice that demonstrated the necessity of MHC I for the development of visual projections and longterm potentiation in the adult hippocampus^[22]. Additionally, a 2004 study showed that MHC I was vital for the selective maintenance of neuronal synapses after induction of axonal lesions^[23]. Together, these studies suggest that low levels of neuronal MHC I expression may contribute to a loss of synaptic plasticity and axon regeneration.

In addition to its involvement in synaptic maintenance in the brain, MHC I has also been linked to behavioral traits^[37]. MHC I molecules have been shown to associate with certain pheromone receptors and aid their expression on the surface of neurons in the vomeronasal organ in mice^[38]. ß2m deficient mice showed decreased expression of certain pheromone receptors and exhibit decreased male-male aggression^[38].

Subjects with ASD have abnormal brain circuitry, which may include high local synaptic connectivity and low long-range synaptic connectivity^[39]. Abnormalities in synaptic and columnar structure, as well as increased



Fig. 1b: Comparison of ß-2-microglobulin levels in children with early onset (EO or classic) autism, regressive (reg) autism and TD controls

cell density in the cerebral cortex have also been demonstrated in individuals with ASD^[40-44]. In addition, children with ASD have been shown to have larger brain volumes than typically developing children^[45]. MHC I deficient mice exhibit many similar structural features in several brain regions^[22]. Thus, in addition to genetic linkages, the involvement of MHC I in brain formation and behavior make it an ideal candidate to study in relation to ASD.

In our cross-sectional analysis, we found there to be no significant difference in plasma ß2m between subjects with ASD and TD controls. However, the negative results reported herein do not exclude the possibility that MHC I may play a role in ASD development. First, due to the clinical nature of our study, it was not possible to directly measure the expression of MHC I by neuronal cells. It is possible that serum ß2m levels are not fully representative of MHC expression in the brain. Perhaps a better source would be cerebral spinal fluid (CSF). Second, MHC I expression on neurons is developmentally regulated and is highest during perinatal periods^[21, 22]. A longitudinal study encompassing several stages of development may render more accurate results and future research should explore this possibility.

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