Prenatal and Perinatal Risk Factors for Autism at National Children's Hospital

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ABSTRACT

Background: Autism, or autism spectrum disorder, refers to a broad conditions characterized by challenges with social skills, repetitive behaviors, speech and nonverbal communication.

Objectives: To determine the demographic profile of patients diagnosed with ASD, determine the significant prenatal and perinatal risk factors associated with ASD.

Results: A total of 116 subjects were included in the study with 58 cases and 58 controls. They belong to the age ranging from 4 to 16 years old. Every case had a confirmed diagnosis of autism at NCH. There was a significant association noted between neonatal jaundice, nulliparity (OR=2.38; 95% CI, 0.85-6.8) and family history of autism (OR=5.30; 95% CI, 1.29-25.1) with ASD. Exposure to x-ray, medical problems, medicine intake and maternal complications during pregnancy were not significantly associated with ASD with OR 0.74; 95% CI, (0.12-4.15), OR 1.00; 95% CI (0.38-2.61), OR1.49; 95% CI, (0.63-3.53), and OR 1.27; 95% CI, (0.28-6.05), respectively.

Conclusion: The current study indicates that the only significant predictor of ASD is a family history of autism. However, neonatal jaundice, maternal age of >40 years old, smoking during pregnancy and nulliparity showed a trend towards being risk factors for ASD. None of the other prenatal and perinatal characteristics significantly predicts ASD.

Keywords Autism, Perinatal, Prenatal.

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INTRODUCTION

Autism is the prototypical form of a spectrum of related, complex neurodevelopmental disorders referred to as the autistic spectrum disorders¹. There is strong evidence from neuropathological studies that ASD has its origins in abnormal brain development early in prenatal life². Atypical neurodevelopment continues postnatal, with a unique pattern of acceleration in brain growth as measured by head circumference³, which correlates with enlarged grey matter volumes observed in MRI studies by two to three years of age⁴. Advances in neonatal intensive care have dramatically increased survival in preterm infants, most strikingly among the sickest and most preterm^{5,6}. Unfortunately, this decrease in mortality has not been matched by a comparable decrease in longterm neurodevelopmental morbidity⁷. Although autism is typically not diagnosed until late in the preschool years, there are marked neurodevelopmental abnormalities that are present at birth and continue to evolve from the earliest months of life. We now know that professionals can diagnose children with autism when they are as young as two years of age⁸. Screening and the role of the pediatrician have become even more critical as we have recognized the stability of early diagnosis over time and the importance of early intervention.

The literature is not always consistent in regards to which

with autism. Thus, this study is proposed to determine the prenatal and perinatal risk factors for autism at National Children Hospital, which may be helpful in identifying high-risk groups for ASD and will have significant role in early diagnosis and intervention.

METHODS

This was a case control study. The study was conducted at National Children's Hospital, a tertiary Government Hospital under the Department of Health. Cases were patients diagnosed to have Autism Spectrum Disorder at National Children Hospital. Controls were taken from patients with similar age and sex who visited the general OPD clinic. A standard pretested and validated questionnaire with presence or absence of important prenatal and perinatal risk factors was used. The questionnaire was prepared by conducting a focus group discussion with ten mothers of children diagnosed with ASD. This was further pretested and validated randomly with forty other mothers who visited our neurodevelopment and general OPD Clinic after signing of informed consent. The number of samples to be collected was computed using a 95% level of confidence and 80% power of the study. A sample size of at least 114 was reached to detect a 13% difference in the occurrence of prematurity among cases and controls. The questionnaire was distributed among at least 57 parents of patients with ASD and 57 parents of patients visiting in general OPD. Consent was provided for all parents who allowed their children to be part of the research. All information obtained from the participants in the study would be kept confidential. Data were encoded and tallied in SPPS version 17 for windows. For nominal data, frequency and percentage were generated. For numerical data, mean +/-SD was computed. Data was analyzed using chi-square, Fisher Exact test, and logistic regression analysis.

RESULTS

A total of 116 subjects were included in the study, with equal numbers of cases and controls. Table 1 shows the association of the demographic characteristics with autism spectrum disorder (ASD). The results showed that there was a significant association noted between the number of pregnancies and family history of autism with ASD as proven by the p values <0.05 and 0.007 respectively. For the number of pregnancies, nulliparity showed a higher

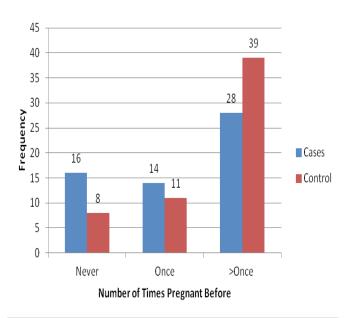
risk for ASD and a higher risk was noted among those with a family history of autism (OR=5. 30; 95% CI, 1.29-25.1). On the other hand, there were no significant associations noted between age and family history of psychiatric illness with ASD as proven by all p value>0.05.

Table 1: Association of the different demographic characteristics with autism spectrum disorder (ASD)

	Cases (n=58)	Controls (n=58)	OR (95% CI)	*p-value	
Number of times pregnant before					
Never Once >Once	16 (27.6%) 14 (24.1%) 28 (48.3%)	8 (13.8%) 11 (19.0%) 39 (67.2%)	2.38 (0.85-6.80) 0.64 (0.17-2.36) 0.36 (0.12-1.05)	0.04 (S) 0.44 (NS) 0.06 (NS)	
Age					
<40 y/o >40 y/o	54 (93.1%) 4 (6.9%)	57 (98.3%) 1 (1.7%)	0.24 (0.01-2.40) 4.22 (0.42- 102.4)	0.36 (NS) 0.36 (NS)	
Family history of autism					
Yes No	13 (22.4%) 45 (77.6%)	3 (5.2%) 55 (94.8%)	5.30 (1.29-25.1) 0.19 (0.04-0.78)	0.007 (S) 0.007(S)	
Family history of psych illness					
Yes No	6 (10.3%) 52 (89.7%)	5 (8.6%) 53 (91.4%)	1.22 (0.31-4.99) 0.82 (0.20-3.32)	1.00 (NS) 1.00 (NS)	

Chi-square test otherwise, Fisher Exact test *p-values >0.05 - Not significant; p-values ≤0.05 - Significant

Fig 2: Distribution of subjects with and without and according to the number of times pregnant before



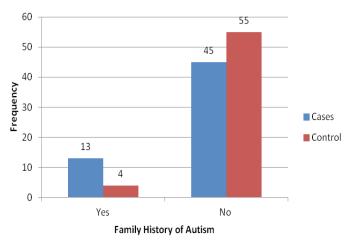


Fig 3: Distribution of subjects with and without ASD according family history of autism

Table 2, shows the association of the prenatal characteristics with ASD. Exposure to x-ray, medical problems, medicine intake, history of smoking and maternal complications during pregnancy were not significantly associated with ASD (p>0.05).

Table 2: Association of the prenatal characteristics withautism spectrum disorder (ASD)

	Cas (n=		Controls (n=58)	OR (95% CI)	*p-value	
Smoking exposure during pregnancy						
Yes No	17 (29 41 (70	'	21 (36.2%) 37 (63.8%)	0.73(0.31 – 1.71)	0.42 (NS)	
Smoking during pregnancy						
Yes No	5 (8.) 53 (9 [.]	'	1 (1.7%) 57 (98.3%)	5.38(0.58 –125.7)	0.20 (NS)	
Exposed to X-ray during pregnancy						
Yes No	3 (5. 55 (94	'	4 (6.9%) 54 (93.1%)	0.74(0.12 – 4.15)	1.00 (NS)	
Medical problems during pregnancy						
Yes No	· ·		13 (22.4%) 45 (77.6%)	1.00 (0.38 - 2.61)	1.00 (NS)	
Medication intake during pregnancy						
Yes No	· · ·	6.2%) 8.8%)	16 (27.6%) 42 (72.4%)	1.49(0.63 – 3.53)	0.32 (NS)	
Maternal complications during pregnancy						
Severe hyper-	F (0,0%)	4 (0.00())	4.07 (0)		4.00 (NIC)	
emesis	5 (8.6%)	4 (6.9%)	`	28 – 6.05)	1.00 (NS)	
Edema	11 (19.0%)	9 (15.5%)	`	44 – 3.72)	0.62 (NS)	
PROM	5 (8.6%)	11 (19.0%)	`	11 – 1.38)	0.10 (NS)	
Others	0	1 (1.7%)	0 (0 -	- 17.54)	1.00 (NS)	

Chi-square test otherwise, Fisher Exact test *P-values >0.05 - Not significant; p-values ≤ 0.05 - Significant

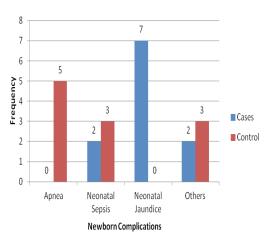
Table 3 shows the association of the perinatal characteristics with autism spectrum disorder (ASD). The results showed that there was a significant association noted between neonatal jaundice (p=0.01) with ASD. On the other hand, there were no significant associations noted between the other perinatal characteristics with ASD as proven by all p values >0.05.

Table 3: Association of the perinatal characteristics withautism spectrum disorder (ASD)

	Cases (n=58)	Controls (n=58)	OR (95% CI)	*p-value	
Gestational age					
37 – 42 weeks <37 weeks >42 weeks	42 (72.4%) 11 (19.0%) 5 (8.6%)	49 (84.5%) 8 (13.8%) 1 (1.7%)	0.48 (0.17 – 1.31) 1.60 (0.53 – 4.88) 5.83 (0.62-137.3)	0.11 (NS) 0.35 (NS) 0.10 (NS)	
Birth weight					
≥2.5 kg <2.5 kg <1.5 kg	50 (86.2%) 6 (10.3%) 2 (3.4%)	53 (91.4%) 5 (8.6%) 0	0.59 (0.15 – 2.17) 1.27 (0.32 – 5.20) 	0.38 (NS) 0.70 (NS) 0.24 (NS)	
Manner of delivery					
CS NSD	20 (34.5%) 38 (65.5%)	16 (27.6%) 42 (72.4%)	1.38 (0.58 – 3.30)	0.42 (NS)	
Exclusive br	reastfeeding	for 6 month	s		
Yes No	30 (51.7%) 28 (48.3%)	29 (50.0%) 29 (50.0%)	1.07 (0.48 – 2.37)	0.85 (NS)	
Newborn complications					
Apnea	0	5 (8.6%)	0 (0 – 1.12)	0.05 (NS)	
Neonatal Sepsis	2 (3.4%)	3 (5.2%)	0.65 (0.07 – 5.08)	1.00 (NS)	
Neonatal Jaundice	7 (12.1%)	0		0.01 (S)	
Others	2 (3.4%)	3 (5.2%)	0.65 (0.07 - 5.08)	1.00 (NS)	
Chi-square test otherwise. Fisher Exact test *p-values>0.05					

Chi-square test otherwise, Fisher Exact test *p-values>0.05 - Not significant; p-values ≤ 0.05 - Significant

Fig 4: Distribution of subjects with and without ASD according to the neonatal complications



In the univariate analysis, three variables were significantly associated with ASD, namely neonatal jaundice, nulliparity and family history of autism. However, in the multivariate analysis using logistic regression, only family history of autism was the significant predictor of ASD (p=0.02). The risk of subjects with a family history of autism for ASD was almost 5x higher than those without a family history of autism (OR=4. 72; 95% CI, 1.20- 18.54, p=0.02). In the univariate analysis, three variables were significantly associated with ASD, namely neonatal jaundice, nulliparity and family history of autism.

However, in the multivariate analysis using logistic regression, only family history of autism was the significant predictor of ASD (p=0.02). The risk of subjects with a family history of autism for ASD was almost 5x higher than those without a family history of autism (OR=4. 72; 95% CI, 1.20- 18.54, p=0.02).

Table 4: Predictors of ASD

Variable	OR	95% CI	P value
Smoking during pregnancy	7.28	0.81 - 65.90	0.08 (NS)
Premature rupture of membrane	0.39	0.10 - 1.48	0.16 (NS)
Abnormal AOG of baby	1.91	0.66 - 5.53	0.24 (NS)
Apnea	0	0-2.8E+19	0.76 (NS)
Neonatal jaundice	2542	0-4.9E+21	0.72 (NS)
Number of times pregnant before (Never)	2.04	0.74 – 5.65	0.17 (NS)
Family history of autism	4.72	1.20 - 18.54	0.02 (S)

Logistic Regression Analysis

p-values >0.05 - Not significant;

p-values ≤0.05 -Significant

DISCUSSION

A case-control study, frequency-matched on gender and birth year, to investigate prenatal and perinatal risk factors for autism in our hospital was conducted. The current study determines the association between the different demographic characteristics, prenatal and perinatal factors of autism. Most of the perinatal and prenatal factors examined in multiple studies have shown inconsistent results and the preponderance of findings overall have not been statistically significant. In previous studies, the factors with the strongest evidence for an association with autism risk included abnormal fetal presentation, umbilical-cord complications, fetal distress, birth injury or trauma, multiple birth, maternal haemorrhage, summer birth, low birth weight, small for gestational age, congenital malformation, low 5-minute Apgar score, feeding difficulties, meconium aspiration, neonatal anemia, ABO or Rh incompatibility, and hyperbilirubinemia. However, not all of these factors were examined in this study. In a previous study, the strongest prenatal factors included advanced maternal and paternal age at birth, maternal gestational bleeding, gestational diabetes, being first born versus third born or later, maternal prenatal medication use, and maternal birth abroad In the present study, aside from prenatal and perinatal characteristics, the demographic characteristics were also investigated. The results showed that in the univariate analysis, family history of autism was significantly associated with ASD. Nulliparity showed trend towards being a risk factor. Several previous studies have found that ASD individuals tend to be first or fourth born more commonly than controls¹⁷. Rather than having a role in the cause of autism, this phenomenon is widely believed to be a result of alterations in the reproductive behavior of parents in response to the birth of a handicapped child, also known as the "reproductive stoppage rules"¹⁸.

Among the perinatal characteristics, neonatal jaundice (p=0.01) was significantly associated with ASD. In previous studies, history of jaundice in neonates was associated with increased risk of disorders of psychological development for children born at term. The excess risk of developing a disorder in the spectrum of psychological development disorders among those who had neonatal jaundice was between 56% (HR: 1.56 [95% confidence interval [CI]: 1.05–2.30]) and 88% (HR: 1.88 [95% CI: 1.17–3.02]). The excess risk of infantile autism was 67% (HR: 1.67 [95% CI: 1.03–2.71])¹⁹.

Multivariate analysis using logistic regression showed only family history of autism as the significant predictor of ASD. However, the tendency of smoking during pregnancy and never been pregnant before as potential risk factors for ASD was also noted. This study suggests that only family history of autism is the strongest predictor of ASD. The risk of subjects with family history of autism for ASD was almost five times higher than those without family history of autism (OR=4.72; 95% CI, 1.20-18.54, p=0.02).

The finding in this study supports ASD to be a possible genetic problem. Previously, Genetic studies in the field of autistic disorder have mainly focused on molecular genetic studies, assessment of chromosomal abnormalities, twin studies and family studies. In families having an autistic child the recurrence rate has been reported as 3-8%²⁰.

The studies on twins and adopted children are important in identifying the actual importance of genetic factors. Concordance among twins enables to measure heritability, and thus to assess what percentage of the phenotype is affected by genetic factors. Monozygotic (identical) twins share 100% of the genetic material, whereas dizygotic (fraternal) twins share 50% of the genetic material. Monozygotic twins higher rate of concordance compared to dizygotic twins may be used for calculation of heritability. Twin studies generally showed a higher concordance rate for monozygotic twins compared to dizygotic twins. The concordance rate of monozygotic twins is at least 60% when diagnostic criteria for autism (DSM-IV) are used, whereas the number is as high as 71% for autism spectrum and 92% for a broader spectrum of verbal/social interaction disorders²¹. On the other hand, the concordance rate of dizygotic twins has been reported as 1-30%. Twin studies demonstrated an average autism inheritance of 90%. On the basis of these studies autism is considered to be among the most inherited psychiatric diseases.

The correlated occurrence of many of the complications limits the ability to determine which factors, if any, are independently associated with autism. For example, Cesarean deliveries are more common in pregnancies with abnormal fetal presentation, fetal distress, and multiple birth^{22,23}. Congenital malformations, low birth weight, abnormal presentation, and low Apgar score also are interrelated²⁴. In most studies, multivariate analyses were not used to simultaneously control for all obstetrical factors examined, and a different set of factors was examined in each study. It is possible that increasing rates of some obstetrical factors, such as Cesarean delivery, low birth weight, multiple birth, and neonatal resuscitation, may be contributing factors to the rising prevalence of autism²⁵. The obstetrical complications that have emerged as significant risk factors for autism in a meta-analysis study suggest a possible role of fetal and neonatal hypoxia. In particular, growth retardation, fetal distress, umbilical-cord wrapping around the neck, low Apgar score, respiratory distress, resuscitation, meconium aspiration, and Cesarean delivery are all potential risk factors that also may be associated with an increased risk of hypoxia²⁶. Although some brain abnormalities observed in individuals with autism may reflect a potential role of oxygen deprivation during development, this possibility requires additional examination. Hypoxia also has been shown to increase dopaminergic activity, and there is evidence for dopamine overactivation in autism²⁷.

However in this study related obstetrical problems like exposure to x-ray, medical problems, medicine intake and maternal complications during pregnancy were not significantly associated with ASD with OR 0.74; 95% CI, (0.12-4.15), OR 1.00; 95% CI (0.38-2.61), OR 1.49; 95% CI, (0.63-3.53), and OR 1.27; 95% CI, (0.28-6.05) respectively. Our study indicates that among the prenatal and perinatal risk factors only family history of autism is a significant predictor of ASD. None of the other prenatal and perinatal characteristics significantly predicts ASD.

CONCLUSION

The current study indicates that the only significant predictor of ASD is a family history of autism. However, neonatal jaundice, maternal age of >40 years old, smoking during pregnancy and nulliparity showed a trend towards being risk factors for ASD. None of the other prenatal and perinatal characteristics significantly predicts ASD.

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