Fetal testosterone and autistic traits

Citation for published version:
https://doi.org/10.1348/000712608X311731

Digital Object Identifier (DOI):
10.1348/000712608X311731

Link:
Link to publication record in Edinburgh Research Explorer

Document Version:
Peer reviewed version

Published In:
British Journal of Psychology

Publisher Rights Statement:

General rights
Copyright for the publications made accessible via the Edinburgh Research Explorer is retained by the author(s) and / or other copyright owners and it is a condition of accessing these publications that users recognise and abide by the legal requirements associated with these rights.

Take down policy
The University of Edinburgh has made every reasonable effort to ensure that Edinburgh Research Explorer content complies with UK legislation. If you believe that the public display of this file breaches copyright please contact openaccess@ed.ac.uk providing details, and we will remove access to the work immediately and investigate your claim.
Foetal testosterone and autistic traits

Bonnie Auyeung¹, Simon Baron-Cohen¹, Emma Ashwin¹, Rebecca Knickmeyer¹,², Kevin Taylor³ and Gerald Hackett⁴

¹Autism Research Centre, Department of Psychiatry, University of Cambridge, Douglas House, 18B Trumpington Rd, Cambridge, CB2 8AH, UK
²Department of Psychiatry, University of North Carolina at Chapel Hill, CB # 7160, Chapel Hill, NC 27599-7160, USA.
³ Department of Clinical Biochemistry, Addenbrooke’s Hospital, Cambridge, CB2 2QQ, UK
⁴ Department of Foetal Medicine, Rosie Maternity Hospital, Robinson Way, Cambridge, CB2 2SW, UK

Acknowledgments. This work was supported by the Nancy Lurie-Marks Family Foundation and the Medical Research Council (MRC). Bonnie Auyeung was supported by a scholarship from Trinity College, Cambridge. We are grateful to the families who have taken part in this longitudinal study over many years and to Melissa Hines, Rosa Hoekstra, Bhismadev Chakrabarti, Malcolm Bang, Svetlana Lutchmaya, Greg Davis, and Ieuan Hughes for valuable discussions.
Abstract

Studies of amniotic testosterone in humans suggest that foetal testosterone (fT) is related to specific (but not all) sexually dimorphic aspects of cognition and behaviour. It has also been suggested that autism may be an extreme manifestation of some male-typical traits, both in terms of cognition and neuroanatomy. In this paper, we examine the possibility of a link between autistic traits and fT levels measured in amniotic fluid during routine amniocentesis. Two instruments measuring number of autistic traits (the Childhood Autism Spectrum Test (CAST) and the Child Autism Spectrum Quotient (AQ-Child)) were completed by these women about their children (n=235), ages 6-10 yrs. IQ was measured in a subset of these children (n=74). fT levels were positively associated with higher scores on the CAST and AQ-Child. This relationship was seen within sex as well as when the sexes were combined, suggesting this is an effect of fT rather than of sex per se. No relationships were found between overall IQ and the predictor variables, or between IQ and CAST or AQ-Child. These findings are consistent with the hypothesis that prenatal androgen exposure is related to children exhibiting more autistic traits. These results need to be followed up in a much larger sample to test if clinical cases of ASC have elevated fT.
Introduction

Many clinical conditions occur in males more often than females, including autism, dyslexia, specific language impairment, attention-deficit hyperactivity disorder (ADHD), and early-onset persistent antisocial behaviour (Rutter, Caspi & Moffitt, 2003). Depression, anorexia, and the anxiety disorders do not show this male bias in sex ratio, raising the question of whether there are sex-linked or sex-limiting factors involved in the aetiology of those conditions that do exhibit this male bias. Autism in particular has been described as an extreme manifestation of certain sexually dimorphic traits or as a consequence of an “extreme male brain” (EMB) (Baron-Cohen, 2002). Autism, high-functioning autism, Asperger syndrome, and Pervasive Developmental Disorder (not otherwise specified; PDD/NOS) will be referred to as Autism Spectrum Conditions (ASC), and are considered to lie on the same continuum. Individuals with an ASC diagnosis are impaired in reciprocal social interaction and communication, and show strongly repetitive behaviours and unusually narrow interests (APA, 1994). A recent epidemiological study in the United Kingdom reports that 1% of people have a diagnosis of ASC (Baird et al., 2006).

The strong bias of ASC towards males has been well established (Bryson & Smith, 1998; Fombonne, 2005; Tidmarsh & Volkmar, 2003). Children with ASC have a sex ratio of 4:1 (male: female) across the full IQ range (Chakrabarti & Fombonne, 2005) and the ratio is as high as 9:1 for Asperger Syndrome (Scott et al., 2002a; Wing, 1981), the subgroup in which
individuals have intact IQ and language development. The strikingly higher incidence of ASC in males might provide important clues to the aetiology of the condition (Baron-Cohen, 2002; Baron-Cohen, Knickmeyer & Belmonte, 2005). ASC have a strong neurobiological and genetic component (Stodgell, Ingram & Hyman, 2001), however the specific factors (hormonal, genetic, or environmental) that are responsible for the higher male incidence in ASC are unclear. Recent evidence is consistent with the idea that male sex hormones, and in particular, prenatal exposure to testosterone may be related to the development of autistic traits (Baron-Cohen, Lutchmaya & Knickmeyer, 2004).

It has been proposed that some observable sex differences in human behaviour and cognition may be accounted for by sex differences in cerebral lateralisation (Hines & Shipley, 1984). Geschwind and Galaburda (1985) hypothesised that foetal testosterone (fT) exposure facilitates the growth of certain areas in the right hemisphere of the brain while simultaneously inhibiting the growth of the same areas in the left hemisphere of the brain (Geschwind & Galaburda, 1985). The Geschwind and Galaburda hypothesis was questioned because of a lack of supportive evidence (Bryden, McManus & Bulman-Fleming, 1994), though it should be noted that the majority of studies which aimed to test the hypothesis did not directly test whether fT affects sexually dimorphic neurobiology. Instead studies used indirect tests, such as examining associations between left-handedness and strongly sex dependent conditions like autism.
In support of the Geschwind and Galaburda hypothesis, studies investigating body asymmetry found that left-handedness and asymmetrical lateralisation were associated with both being male, and with autism (Fein et al., 1985; McManus et al., 1992; Satz et al., 1985; Soper et al., 1986). Analogies can also be drawn between sex differences in brain development and neuroanatomical characteristics found in autism. The typical male brain is heavier than the female brain and individuals with autism have heavier brains than typical males (Harden et al., 2001). The amygdala is also disproportionately large in boys compared to girls (Giedd et al., 1996) and children with autism have enlarged amygdalae (Hazlett et al., 2005). These findings are consistent with the EMB theory.

The ratio between the length of the 2nd and 4th digit (2D:4D) is sexually dimorphic, being lower in males than in females, and may be a useful proxy measure for fT production in humans during the first trimester of gestation (Manning et al., 1998). Studies researching foetal hand development have observed the sex difference in 2D:4D ratio in foetuses between 9-40 weeks of gestation (Malas et al., 2006). 2D:4D ratio has been found to be negatively associated with the ratio of fT to foetal oestrogen (Lutchmaya et al., 2004). Lower (i.e., hyper-masculinised) digit ratios have been found in children with autism compared to typically developing children. This pattern was also found in the siblings and parents of children with autism, suggesting genetically-based elevated fT levels in autism (Manning et al., 2001; Milne et al., 2006). If 2D:4D ratio does reflect prenatal exposure to testosterone, this evidence
suggests children with ASC may have been exposed to higher than average levels of fT.

The direct manipulation of fT levels is not possible in humans for ethical reasons. The investigation of prenatal hormone exposure and its relation to development in humans has therefore been investigated in naturally occurring abnormal environments such as in individuals with Congenital Adrenal Hyperplasia (CAH) which is a genetic disorder that causes excess adrenal androgen production beginning prenatally in both males and females (New, 1998). Studies of individuals with CAH have generally found that girls with CAH show masculinisation of performance in activities typically dominated by males such as spatial orientation, visualisation, targeting, personality, cognitive abilities and sexuality (Hampson, Rovet & Altmann, 1998; Hines et al., 2003; Resnick et al., 1986). Results from one study of girls with CAH suggest that they exhibit more autistic traits, measured using the Adult version of the Autism Spectrum Quotient (AQ), compared to their unaffected sisters (Knickmeyer et al., 2006b). Whilst CAH provides an interesting opportunity to investigate the effects of additional androgen exposure, the relatively rare occurrence of CAH in conjunction with ASC makes it difficult to obtain large enough sample sizes for generalisation of research findings to the wider population. Researchers have also suggested that CAH-related disease characteristics, rather than prenatal androgen exposure, could be responsible for the atypical cognitive profiles observed in this population (Fausto-Sterling, 1992; Quadagno, Briscoe & Quadagno, 1977).
The relationship between fT exposure and post-natal development have also been examined using measures of fT levels in amniotic fluid, obtained during amniocentesis performed for other clinical reasons. In animal models, the critical period for sexual differentiation of the brain occurs when differences in serum testosterone are highest between sexes (Smith & Hines, 2000). The human fT surge is thought to occur between weeks 8 to 24 of gestation (Baron-Cohen et al., 2004; Collaer & Hines, 1995; Hines, 2004). A major advantage of amniocentesis for examining hormone-behaviour relations is that it is typically performed during the second trimester of pregnancy, during a relatively narrow window of time (usually 14-20 weeks of gestation) that coincides with the serum testosterone peak period in male foetuses.

Although the origins of hormones (and in particular androgens) found in amniotic fluid are not fully understood, the main source seems to be the foetus itself (Cohen-Bendahan, van de Beek & Berenbaum, 2005). Hormones can enter the amniotic fluid in two ways: via diffusion through the foetal skin in early pregnancy, and via foetal urine in later pregnancy (Judd et al., 1976; Schindler, 1982). Thus, testosterone levels measured in amniotic fluid might be expected to be a good reflection of the levels in the foetus, providing an alternative to direct assay of foetal serum (Finegan, Bartleman & Wong, 1991). A limitation is that amniocentesis is an invasive procedure conducted only when foetal anomalies are suspected (such as chromosomal trisomy 21 (Down Syndrome)). This invasive nature and the small risk of miscarriage that amniocentesis entails, means that samples cannot be specifically collected for research purposes. As a result, women who undergo amniocentesis (about
6% of pregnancies) are not a random selection of all pregnant women. They tend to be above average in age (because maternal age is a risk factor for having a child with Down Syndrome) and above average in education, which may limit generalisability. Nevertheless, amniocentesis itself does not appear to have any negative effects on offspring behaviour (Finegan et al., 1990). Maternal age can be studied as a variable in its own right (since women may have an amniocentesis at any age), as can maternal educational level (since this is measurable and can be entered as a covariate in any later analysis). Furthermore, studies can opt to include only those pregnancies that were normal and where the baby was born healthy and with no cytogenetic abnormalities. As such, the measurement of testosterone levels in amniotic fluid presents a unique opportunity for exploration of the foetal hormonal environment, and arguably the only ethical option for conducting such research in humans.

The EMB theory is an extension of the Empathising-Systemising (E-S) theory of typical psychological sex differences (Baron-Cohen, 2002, 2003) which proposes that females on average have a stronger drive to empathise (to identify another person’s emotions and thoughts, and to respond to these with an appropriate emotion), while males have a stronger drive to systemise (to analyse or construct rule-based systems, whether mechanical, abstract, natural, etc.) (Baron-Cohen, 2003; Baron-Cohen et al., 2005). Evidence has been reported showing a female advantage in empathy and a male advantage in systemising (Baron-Cohen et al., 1999; Baron-Cohen, Wheelwright & Hill, 2001a; Davis, 1994; Lawrence et al., 2004). Consistent with the EMB theory,
individuals with ASC have been shown to have a stronger drive to systemise (Baron-Cohen et al., 2003), but show impairment on tests of empathising (Baron-Cohen et al., 1999; Baron-Cohen et al., 2001a). It has been found that in typically developing children whose mothers had undergone amniocentesis, fT levels show a positive association with systemising (Auyeung et al., 2006), and a negative association with empathising (Chapman et al., 2006; Knickmeyer et al., 2006a).

Empathy and systemising are difficult to measure in infancy. Developmental precursors of empathy include social interest, indexed by behaviours such as eye contact and attention to faces (Baron-Cohen, 1995; Johnson, Cheek & Smither, 1983) and language development (Baron-Cohen, Baldwin & Crowson, 1997). Developmental precursors to systemising include attention to detail and narrow interests (Baron-Cohen, 2002). Reduced eye contact, delayed language development, social difficulties and narrow interests are also characteristic of children with ASC (APA, 1994; Lutchmaya, Baron-Cohen & Raggatt, 2002a; Rutter, 1978; Swettenham et al., 1998). Results from the Cambridge fT Project showed that in typically developing children whose mothers had undergone amniocentesis, fT was inversely associated with frequency of eye contact in males at 12 months old (Lutchmaya et al., 2002a), and inversely predicted vocabulary development in children between ages 18 to 24 months (Lutchmaya, Baron-Cohen & Raggatt, 2002b). At four years of age, high levels of fT were associated with poorer quality of social relationships and more narrow interests (Knickmeyer et al., 2005). These studies provide evidence that prenatal fT levels are associated with individual
differences in sexually dimorphic behavioural traits relevant to the features of ASC.

In this paper we report the most recent study from the Cambridge Foetal Testosterone (fT) Project, an ongoing longitudinal study investigating the relationship between fT levels and the development of behaviours relating to ASC (Baron-Cohen et al., 2004; Knickmeyer & Baron-Cohen, 2006). This is a unique longitudinal study, having measured fT levels in amniotic fluid for a sample of children whose behaviour has been examined post-natally at 1, 2, 4, and 8 years of age, using a variety of age appropriate tests. This project is ongoing and it is intended to follow these children through puberty and into adulthood. The key question behind this project is whether there is any relationship between individual variation in fT levels and later phenotypic traits.

The aim of the present study is to examine the relationship between autistic traits and fT exposure in this cohort of typically developing children who are between six and ten years old. Two measures of autistic traits were employed: the Childhood Autism Spectrum Test (CAST) (Scott et al., 2002b; Williams et al., 2005), (formerly known as the Childhood Asperger Syndrome Test, renamed because it can be used for all subgroups on the autistic spectrum (Baron-Cohen et al., submitted)) and the Child Autism Spectrum Quotient (AQ-Child) (Auyeung et al., 2008). The CAST was administered because it is an established measure of autistic traits that has been validated on a large population in the United Kingdom (Williams et al., 2005) and has been shown
to be heritable (Ronald et al., 2006). The AQ-Child was administered because research suggests that the AQ is a good measure of autistic traits across the lifespan. The AQ also predicts clinical diagnosis in adults (Woodbury-Smith et al., 2005) and shows strong cross-cultural consistency (Hoekstra et al., 2008; Wakabayashi et al., 2006) and substantial heritability in the general population (Hoekstra et al., 2007). We also investigate the relationship between intelligence (IQ) and fT level in a subset of this typically developing sample, in order to explore if fT is related to general cognitive ability.

Methods

Participants

Participants were recruited from a longitudinal study on the effects of fT (Baron-Cohen et al., 2004). Medical records of 950 patients who had undergone amniocentesis in the Cambridge (UK) region between 1996 and 2001 were examined. Women were excluded if: (a) the amniocentesis revealed a chromosomal abnormality; (b) there was a twin pregnancy; (c) the pregnancy ended in miscarriage, termination or significant medical problems after birth; (d) relevant information was absent from medical records; or (e) medical practitioners indicated that contacting the family would be inappropriate.

The AQ-Child and CAST were sent to all mothers meeting inclusion criteria, resulting in 452 mothers contacted; 261 mothers completed the CAST and 248 mothers completed the AQ-Child, resulting in a total of 235 (118 boys, 117 girls) children with complete data for both questionnaires. Intelligence
Quotient (IQ) (measured using the WASI (Wechsler, 1999)) was completed for a subgroup of these children (n=74, 43 boys, 31 girls), whose mothers consented to bring them in for cognitive assessment.

**Outcome variables**

*The Child Autism Spectrum Quotient (AQ-Child).* This is a 50-item parent-report questionnaire developed to detect autistic traits in children 4-11 years of age (Auyeung et al., 2008). Higher scores indicate a greater number of autistic traits. AQ-Child items are answered in a Likert format (definitely agree, slightly agree, slightly disagree and definitely disagree). The AQ-Child has shown good test-retest reliability (r=0.85, p<0.001), high sensitivity (95%) and high specificity (95%) (Auyeung et al., 2008). The AQ-Child was originally designed with five subscales to assess various domains of functioning: social skills, attention to detail, attention switching, communication and imagination. Each domain is assessed by ten questions, giving a maximum attainable score of 150. Principal components analysis suggests the AQ-Child has four empirically derived subscales: mind-reading (16 items), attention to detail (9 items), social skills (15 items) and imagination (7 items), resulting in a new maximum score of 141. A score of 76 or above indicates a risk for ASC. These four subscales were found to be respectively highly correlated with the original AQ subscales of communication (r=0.97, p<0.001), attention to detail (r=0.95, p<0.001) social skills (r=0.97, p<0.001) and imagination (r=0.97, p<0.001) (Auyeung et al., 2008). Cronbach’s α was calculated for total score, and results for this sample demonstrated good internal consistency (α=0.91).
The internal consistency of the AQ-Child subscales were also satisfactory (Mindreading=0.80; Attention to Detail=0.80; Social Skills=0.87; and Imagination=0.75).

The Childhood Autism Spectrum Test (CAST). This 37-item parent-report questionnaire was developed to detect ASC in 4-11 year old children (Scott et al., 2002b). CAST items require a binary response (‘yes/no’) to 37 questions, 31 of which are scored (maximum score of 31). A validation study suggested that a score of 15 or above should be used to indicate risk for ASC (Scott et al., 2002b; Williams et al., 2005). The CAST has good test-retest reliability (Spearman’s rho=0.83, p<0.001) (Williams et al., 2006), good positive predictive value (50%) and high specificity (97%) and sensitivity (100%) for ASC (Williams et al., 2005). For the current sample, Cronbach’s α showed acceptable internal consistency (α=0.85).

The Wechsler Abbreviated Scale of Intelligence (WASI). This scale was used to measure IQ (Wechsler, 1999). The WASI provides scores for Verbal IQ, Performance IQ and Full Scale IQ. We also examined the relationship between fT and the Block Design component of the Wechsler Abbreviated Scale of Intelligence (WASI). Block Design performance shows sexual dimorphism in adulthood (male advantage) (Lynn, 1998; Lynn et al., 2005; Rönnlund & Nilsson, 2006) and individuals with ASC show superior performance on this subtest (Happe, 1994; Shah & Frith, 1993).
**Predictor variables**

*fT levels*. The major predictor in this study is fT level in amniotic fluid, measured by radioimmunoassay. Amniotic fluid was extracted with diethyl ether. The ether was evaporated to dryness at room temperature and the extracted material re-dissolved in an assay buffer. Testosterone was assayed by the DPC ‘Count-a-Coat’ method (Diagnostic Products Corp, Los Angeles, CA 90045-5597), which uses an antibody to testosterone coated onto propylene tubes and a 125-I labelled testosterone analogue. The detection limit of the assay using the ether-extraction method is approximately 0.05 nmol/L. The coefficient of variation (CV) for between batch imprecision is 19% at a concentration of 0.8 nmol/L and 9.5% at a concentration of 7.3 nmol/L. The CV’s for within batch imprecision are 15% at a concentration of 0.3 nmol/L and 5.9% at a concentration of 2.5 nmol/L. This method measures total extractable testosterone.

The following control variables were also included in our study.

**Gestational age at amniocentesis.** The amniocentesis procedure generally occurs between weeks 14 and 22. Therefore it is important to determine whether fT is related to gestational age. No significant relationships were found between fT levels and gestational age when both sexes were combined (\(r=-0.03, p>0.05\)), as well as when examining boys (\(r=-0.10, p>0.05\)) and girls (\(r=0.07, p>0.05\)) separately.
Maternal age. Maternal age was included because women undergoing amniocentesis have a higher mean age than the general childbearing population.

Level of education obtained by the parents. Parental education level was measured according to a 5-point scale: 1=no formal qualifications, 2=typical 16-year-old qualification, 3=typical 18-year-old qualification, 4=university degree and 5=postgraduate qualification. The average of the level of education obtained by the mother and father was computed to obtain parental education level.

Presence of older siblings. Older siblings have been found in previous research to have an impact on the social environment and influence child development (Nystul, 1981). Since the outcome of interest was sexually dimorphic, this variable was further split into two variables: older brothers present in the home (or not) and older sisters present in the home (or not).

Child’s Age. The children included in the analyses were between six and ten years of age. Child age was therefore included as a control variable.

Results

Examination of the univariate distributions revealed that foetal testosterone level was positively skewed, and was the only predictor variable with a distribution that deviated significantly from the Gaussian distribution. Four female outliers in fT levels (individuals who scored three or more standard
deviations from the mean) were observed. These outlying values were replaced using a winsorizing procedure, where the extreme values are replaced by the highest observed level within three standard deviations from the mean (0.80 nmol/L). The winsorizing procedure was chosen because it is a compromise between the two goals of eliminating the strong influence of extreme values while at the same time utilizing all of the information. Winsorized fT levels showed no outliers and acceptable (skewness<1) skewness statistics for boys and girls separately and combined, and are used in subsequent analyses.

No significant sex differences were found for any of the predictor variables except fT level. As expected, results showed that boys ($M=0.84, SD=0.41$) had higher fT levels than girls ($M=0.32, SD=0.20$), $t(167.97)=12.92, p<0.001$, equal variances not assumed). Table 1 presents the means and standard deviations for predictor variables, AQ-Child and CAST scores.

**AQ-Child Scores**

AQ-Child Total. For AQ-Child total score, examination of the univariate distribution revealed that it was not skewed (skewness<1) for all cases together as well as in boys and girls separately. Figure 1 shows the raw distribution of total AQ-Child scores. AQ-Child scores are normally distributed, as shown in Figure 1.
Scores on the AQ-Child showed significant sex-differences, \( t(233)=6.64, p<0.001 \) (equal variances assumed), with boys \((M=48.75, SD=17.96)\) scoring higher than girls \((M=34.42, SD=15.01)\).

A hierarchical multiple regression analysis was conducted. In the first block, any predictor variable that showed a significant correlation with AQ-Child scores at the \( p<0.20 \) was entered into the analysis (Altman, 1991). In addition, the influence of suppressor variables (predictors that are highly correlated with other predictors in the model at \( p<0.01 \)) was investigated. The main effects of fT level and child sex were tested for inclusion in the second block using the stepwise method (entry criterion \( p<0.05 \), removal criterion \( p>0.10 \)). The interaction between child sex and fT level was tested for inclusion in the third block using the stepwise method. Table 2 shows the correlation coefficients for both the predictor and outcome variables for boys and girls together.

The predictor variables that correlated with total AQ-Child scores at \( p<0.20 \) were presence of older sisters \((r=-0.19, p<0.01)\) and presence of older brothers \((r=-0.14, p<0.05)\). The inclusion of fT level in the second block produced a significant F-change \((F=46.35, p<0.001, R^2=0.21)\). Inclusion of child sex in the final regression model also produced a significant F-change \((F=7.99, p<0.05)\). The interaction of sex and fT level was excluded as a predictor from the final regression model. See Figure 2 for a visual representation of the relationship between fT level and AQ-Child scores for males and females combined.
Within sex analyses were conducted to further investigate scoring patterns in boys and girls separately. For girls only, parent education level (r=-0.14, p<0.01), presence of older sisters (r=-0.17, p<0.01) and older brothers (r=-0.23, p<0.05) showed correlations at the p<0.20 level and were entered into the first block using the enter method. FT level (r=0.27, p<0.001) was included in the second block using the stepwise method. A significant F-change (F=4.12, p<0.01) was observed when FT was entered into the regression in the second block. The predictor variables that correlated with AQ-Child scores at the p<0.20 level for boys were presence of older sisters (r=-0.19, p<0.01) and brothers (r=-0.14, p<0.05). Presence of older sisters and brothers were included in the first block using the enter method. FT level (r=0.22, p<0.001) was entered in the second block using the stepwise method. The final model for boys included FT level, which showed a significant F-change (F=5.13, p<0.05). Residual analysis revealed acceptable plots and no outliers.

A score of 76 or higher was considered to indicate risk for ASC (Auyeung et al., 2008), and the above analyses were repeated excluding those who scored at or above this cut-off (n=10, 1 girl, 9 boys) to again ensure that the results were not affected by the high scorers on this measure. See Table 3 for AQ-Child regression results with and without the high scorers. The first block included presence of older sisters (r=-0.18, p<0.01) and brothers (r=-0.13, p=0.06). The second block tested for the inclusion of FT level (r=0.43, p<0.001) and child sex (r=0.37, p<0.001). Results show that, when excluding the high scorers, FT level (F=48.69, p<0.001) and sex (F=4.20, p<0.05) both
produce a significant F-change, and is consistent with results including the high scorers. Residual analysis revealed acceptable plots and no outliers.

Insert Table 3 here

**AQ-Child Subscales.** The AQ-Child subscales were next examined to test if the fT relationships were consistent across all four subscales. Due to the uneven number of items per subscale, a raw score was calculated for each of the four subscales and then divided by the number of items, allowing for comparison between subscales.

Sex differences were explored among the four empirical AQ-Child subscales (see Table 4). All four subscales showed significant sex differences (all \( p<0.001 \)) with boys scoring higher than girls.

Insert Table 4 here

Mindreading, Attention to Detail, Social Skills and Imagination were significantly associated with fT level and sex. These results are consistent with those observed in AQ-Child Total when all participants are examined together (see Table 5).

Insert Table 5 here

**CAST Scores**

Examination of univariate distributions indicated that the distribution for CAST scores was positively skewed. Figure 3 shows the raw distribution of CAST
scores. CAST scores were transformed by adding one and taking the square root of each score, resulting in a distribution that was not significantly skewed.

Transformed CAST scores showed significant sex-differences, $t(226.55)=2.12, p<0.05$, equal variances not assumed, with boys ($M=2.36, SD=0.82$) scoring higher than girls ($M=2.15, SD=0.69$).

For the hierarchical regression analysis the predictor variables that correlated with CAST scores at the $p<0.20$ level, were sex ($r=0.14, p<0.05$), fT ($r=0.25, p<.001$) and maternal age ($r=-0.13, p=0.06$). No suppressor variables were observed. Inclusion of fT level in the second block produced a significant F-change ($F=10.72, p<0.01, R^2=0.07$). The main effect of sex was excluded as a predictor. Residual analysis showed no outliers and acceptable plots. See Figure 4 for a visual representation of the relationship between fT level and CAST scores.

In addition, to further investigate whether the results might be due to a sex difference (not necessarily involving fT), we analysed the relationship between these scores and fT within each sex. For boys, maternal age ($r=-0.15, p=0.12$) and presence of older brothers ($r=0.14, p=0.15$) met criteria for entry into the analysis ($r=0.14, p<0.001$). Presence of older sisters was included as a suppressor variable due to its high correlation with the presence of older brothers ($r=0.32, p<0.001$). Inclusion of fT level in the second block produced
a significant F-change ($F=6.57$, $p<0.05$, $R^2=0.12$). For girls alone, no significant relationship was found between CAST scores and fT levels, therefore regression analyses were not conducted.

A raw score of 15 or higher was considered to indicate risk for ASC (Scott et al., 2002b). As for the AQ-Child, the above analyses were repeated excluding those who scored at or above this cut-off ($n=7$, 1 girl, 6 boys). Table 6 shows regression results for the CAST with and without high scorers. Maternal age ($r=-0.12$, $p<0.20$) and presence of older brothers ($r=-0.10$, $p<0.20$) were entered into the first block using the enter method. Presence of older sisters was also included in the first block due to its positive relationship with older brothers ($r=0.37$, $p<0.001$). The second block included fT level ($r=0.32$, $p<0.001$) and child sex ($r=0.32$, $p<0.001$). Inclusion of fT level in the second block produced a significant F-change ($F=4.29$, $p<0.05$, $R^2=0.05$), while child sex and the interaction of sex and fT level were excluded as predictors from the final regression model. These results suggest that children with higher fT levels score higher on the CAST.

Insert Table 6 here

**IQ Scores**

Table 7 shows the means, standard deviations and t statistics for all variables for this subset of children. No significant sex-differences were found between boys and girls for Full Scale IQ, Performance IQ, Verbal IQ or Block Design scores (using the Bonferroni correction for multiple comparisons). No
significant correlations were found between any of the predictor variables and Full Scale IQ, Performance IQ, Verbal IQ and Block Design scores. Regression analyses were not conducted for these variables (see Table 8).

Discussion

This study examines the relationship between foetal testosterone (fT) levels and the later development of autistic traits as measured by the Childhood Autism Spectrum Test (CAST) and the Child Autism Spectrum Quotient (AQ-Child). fT levels were found to be positively associated with number of autistic traits. The significant positive relationship between fT levels was observed across CAST total score, AQ-Child total score, as well as in the four subscales of the AQ-Child. Results remained consistent when excluding individuals who scored above the established cut-offs for the CAST and AQ-Child.

The finding that boys scored higher on both the AQ-Child and CAST coupled with the significant positive associations observed between fT levels and these measures provides support for the Extreme Male Brain (EMB) theory at both psychometric and biological levels. While this dual-level convergence of evidence has been reported previously using indirect biological measures such as the 2D:4D ratio (assumed to be a proxy for fT) (Manning et al., 2001), or brain activity using fMRI (Baron-Cohen et al., 2006b), this is first time that such a relationship has been reported using direct measures. We can assume
that the observed positive association between fT levels and autistic traits may reflect a direct causal effect of fT on neural development, but this remains speculative due to the correlational design of this study. It is for example possible that fT is serving as an index for an unknown third variable.

Scores from the AQ-Child showed a significant positive relationship with fT levels when the sexes were combined as well as when they were examined independently. The CAST, however, was found to have a significant positive relationship with fT levels when the sexes were combined and in boys only. The relationship was not found between CAST and fT levels when girls were examined alone. fT levels predicted about 7% of the variation in CAST scores, and 20% of the variation in AQ-Child scores. These results suggest that the AQ-Child is a better instrument for measuring the variance in autistic traits and its association with fT levels. This may be because the AQ-Child is a wider scale with a less skewed distribution, relative to the CAST.

Although the correlation between the CAST and AQ-Child ($r=0.25$, $p<0.001$) was statistically significant, the low coefficient does not suggest high convergent validity. This may be because of the differing response format of each questionnaire. The CAST requires a binary response, where parents indicate whether their child presents a particular behaviour. Naturally, one would expect to see many children who have low scores, because the majority of children are not at risk for ASC. In contrast, the AQ-Child is answered in Likert format, and was designed to measure the extent to which
the parents agree their child exhibits certain behaviours, resulting in a larger variation in scores.

It is of interest to note that most other research examining the role of testosterone in human psychosexual development has produced more supportive evidence in females than in males (Hines, 2004; Hines, 2006). In the current study, we found a significant relationship between fT levels and AQ-Child scores in boys and girls separately. For CAST scores, a significant correlation with fT levels in boys was observed, but no significant relationship was seen with fT levels in girls. However, compared to boys, girls showed less variability in CAST scores as well as fT levels possibly limiting the ability to detect associations.

No relationships were found between full scale IQ and the predictor variables (including fT) or the CAST or AQ-Child scores. Our prediction that fT level would be significantly positively related to Block Design scores was not supported ($r=0.16$, $p>0.05$). Our results suggest that fT level does not account for individual variation in Full Scale IQ, Verbal IQ, Performance IQ or Block Design scores. No sex differences were found between these variables. It is possible that testosterone only plays a role in a sub-set of behaviours where sex differences are found. However, an alternative explanation could be that the strength of the relationship between fT levels and outcome may vary for different behaviours. If the effect of normal variation in fT level on IQ is small, then only studies with large sample sizes will reveal it. It is also possible that fT levels contribute to IQ but does so at a different time period than that
examined in this study. However, the lack of significant sex differences in IQ scores in the current study suggest that even in a larger sample, no relationship between fT levels and IQ would have been observed. It would nevertheless be beneficial for future studies to examine the relationship between fT levels and IQ scores (in particular Block Design performance) in a larger sample of children. In addition, it would be interesting to investigate if these relationships remain consistent throughout adolescence and adulthood, since sex differences in Block Design performance have been seen in adults (Lynn, 1998; Lynn et al., 2005; Rönnlund & Nilsson, 2006).

Whilst we have obtained evidence that higher fT levels are associated with a child exhibiting more autistic traits, it remains unknown if postnatal testosterone (pT) level would also show such an association. Addressing this question would help to narrow down the window of time in which testosterone exerts its effects on neural development. It is possible that for testosterone to have such effects, these need to occur within a “critical period” (during the second trimester). In addition to the surge that occurs in the second trimester of pregnancy, a second peak in circulating testosterone occurs in human male infants a few days after birth. The levels of pT are strikingly high: they remain in a pubertal range for a few months, and then usually drop to barely detectable levels by 4-6 months of age (Smail et al., 1981). It has been proposed that pT levels in humans influence later spatial skills such as mental rotation ability (Hines et al., 2003), but there is a clear need for further research into the roles of fT and pT in relation to behavioural development before conclusions can be reached.
It is known that hormones fluctuate during the day and across days, even in foetuses (Seron-Ferre, Ducsay & Valenzuela, 1993; Walsh, Ducsay & Novy, 1984). It is ethically unacceptable to attempt to obtain repeated samples of fT during one day or across days, during gestation. Therefore the representativeness of a single sample of fT remains unclear. Given the reported time course of testosterone secretion (Smail et al., 1981), it is likely the most promising time to be measuring fT is prenatal weeks 8 to 24 (Baron-Cohen et al., 2004; Collaer & Hines, 1995; Hines, 2004), but this is still a relatively wide range. Since the decision to perform amniocentesis is understandably based on clinical-medical factors, rather than purely scientific ones, the inferences we can draw about the single measure of fT will necessarily be limited.

While the factors contributing to individual variation in fT levels are not fully understood, adolescent and adult levels of testosterone are heritable (Harris, Vernon & Boomsma, 1998; Hoekstra, Bartels & Boomsma, 2006). There is also significant heritability of autism in clinical samples (Rutter, 2000) and of autistic traits measured in the general population (Hoekstra et al., 2007; Ronald et al., 2006). It may therefore be likely that genetic effects underlie the relationship between fT levels and number of autistic traits. A recent study found both diagnosis of ASC, and number of autistic traits, are significantly associated with specific candidate genes that regulate testosterone production and synthesis (Chakrabarti et al., submitted). It is not yet clear though if the effects reported here are purely genetic, if high fT levels also serve as an endocrine environmental risk factor for autism, or if there are gene-
environment interaction effects, such that high fT only increases risk for ASC in genetically susceptible individuals.

Our studies suggest that variations in fT levels are related to autistic traits as well as other specific aspects of sexually dimorphic behaviour and cognition in typically developing children (Auyeung et al., 2006; Baron-Cohen et al., 2004; Chapman et al., 2006; Grimshaw, Sitarenios & Finegan, 1995; Knickmeyer et al., 2005; Knickmeyer et al., 2006a; Lutchmaya et al., 2002a; Lutchmaya et al., 2002b), but caution needs to be taken when extrapolating these results to individuals with a formal diagnosis of autism spectrum conditions (ASC). The current study is too small a sample to be able to test if fT levels are elevated in formally diagnosed cases of ASC, since these have a prevalence rate of about 1% (Baird et al., 2006). We are currently conducting a large-scale collaboration with the Danish Biobank so as to increase our sample size sufficiently to compare fT levels of cases vs. controls.

Conclusions

There is converging evidence from a range of methods showing the masculinising effects of fT on individual differences in the following sexually dimorphic traits: social skills, language development, empathy, systemising, and visual-analytic skills. If, according to the Extreme Male Brain theory, ASC is an extreme of male-typical behaviour, exposure to elevated levels of fT could be one important factor that is involved with the development of the condition. This study was conducted to examine the effects of fT levels on the development of autistic traits. fT levels were positively associated with higher
scores (indicating greater number of autistic traits) on the CAST and the AQ-Child. No relationships were found between IQ and the predictor variables. Further work is needed to test for any link between fT and later development of ASC, focusing on individuals with a clinical diagnosis.
References


Table 1. Means, standard deviations, and ranges for each sex separately as well combined

<table>
<thead>
<tr>
<th>Variable</th>
<th>Combined Group (n=235)</th>
<th>Girls (n=117)</th>
<th>Boys (n=118)</th>
<th>t</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>M</td>
<td>SD</td>
<td>Range</td>
<td>M</td>
</tr>
<tr>
<td>fT level (nmol/L)**</td>
<td>0.58</td>
<td>0.42</td>
<td>0.05-2.05</td>
<td>0.32</td>
</tr>
<tr>
<td>Gestational Age</td>
<td>16.49</td>
<td>1.44</td>
<td>13-22</td>
<td>16.57</td>
</tr>
<tr>
<td>Child Age</td>
<td>8.91</td>
<td>0.95</td>
<td>6.97-10.68</td>
<td>8.80</td>
</tr>
<tr>
<td>Maternal Age</td>
<td>35.77</td>
<td>4.40</td>
<td>23.68-45.90</td>
<td>35.88</td>
</tr>
<tr>
<td>Parental Education</td>
<td>3.24</td>
<td>1.01</td>
<td>1-5</td>
<td>3.18</td>
</tr>
<tr>
<td>CAST Total**</td>
<td>4.65</td>
<td>3.87</td>
<td>0-22</td>
<td>4.08</td>
</tr>
<tr>
<td>AQ-Child Total**</td>
<td>41.62</td>
<td>18.02</td>
<td>6-103</td>
<td>34.42</td>
</tr>
</tbody>
</table>

^Square-root transformation carried out prior to analyses

* p<0.05
** p<0.01
Table 2. Correlation matrix showing relationships between predictor variables and CAST and AQ-Child Total scores

<table>
<thead>
<tr>
<th></th>
<th>Sex Level</th>
<th>Sex Age</th>
<th>Gest. Age</th>
<th>Child Age</th>
<th>Maternal Age</th>
<th>Parent Education</th>
<th>Older Sister</th>
<th>Older Brother</th>
<th>CAST Total</th>
<th>AQ-Child Total</th>
</tr>
</thead>
<tbody>
<tr>
<td>Sex</td>
<td>.63**</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Gestational Age</td>
<td>-.04</td>
<td>-.06</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Child Age</td>
<td>.03</td>
<td>.12</td>
<td>-.05</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Maternal Age</td>
<td>-.02</td>
<td>-.03</td>
<td>-.28**</td>
<td>-.06</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Parent Education</td>
<td>.07</td>
<td>.06</td>
<td>-.09</td>
<td>-.05</td>
<td>.16*</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Older Sister</td>
<td>-.07</td>
<td>-.08</td>
<td>-.11</td>
<td>-.10</td>
<td>-.03</td>
<td>-.07</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Older Brother</td>
<td>-.04</td>
<td>-.12</td>
<td>-.10</td>
<td>.01</td>
<td>.09</td>
<td>-.15*</td>
<td>.36**</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>CAST Total</td>
<td>.25**</td>
<td>.14*</td>
<td>.03</td>
<td>.08</td>
<td>-.13</td>
<td>-.02</td>
<td>-.01</td>
<td>-.07</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>AQ-Child Total</td>
<td>.41**</td>
<td>.40**</td>
<td>.01</td>
<td>-.01</td>
<td>-.01</td>
<td>-.06</td>
<td>-.19**</td>
<td>-.14*</td>
<td>.25**</td>
<td></td>
</tr>
</tbody>
</table>

*p < 0.05

**p < 0.01
### Table 3. Final Regression Model for the AQ-Child

<table>
<thead>
<tr>
<th>Outcome</th>
<th>Predictor Variable</th>
<th>R²</th>
<th>Δ R²</th>
<th>B</th>
<th>SE</th>
<th>β</th>
</tr>
</thead>
<tbody>
<tr>
<td>AQ-Child Total*</td>
<td>Constant</td>
<td>0.04</td>
<td>0.04</td>
<td>36.02</td>
<td>2.19</td>
<td>0.13*</td>
</tr>
<tr>
<td></td>
<td>Older Sister</td>
<td></td>
<td></td>
<td>7.36</td>
<td>3.45</td>
<td>0.13*</td>
</tr>
<tr>
<td></td>
<td>Older Brother</td>
<td></td>
<td></td>
<td>3.53</td>
<td>4.03</td>
<td>0.06</td>
</tr>
<tr>
<td></td>
<td>fT level</td>
<td>0.20</td>
<td>0.16</td>
<td>11.61</td>
<td>3.23</td>
<td>0.27**</td>
</tr>
<tr>
<td></td>
<td>Child Sex</td>
<td>0.23</td>
<td>0.03</td>
<td>3.82</td>
<td>1.35</td>
<td>0.21**</td>
</tr>
<tr>
<td>AQ-Child Totalb</td>
<td>Constant</td>
<td>0.04</td>
<td>0.04</td>
<td>33.86</td>
<td>1.92</td>
<td>0.13*</td>
</tr>
<tr>
<td></td>
<td>Older Sister</td>
<td></td>
<td></td>
<td>5.91</td>
<td>2.99</td>
<td>0.13*</td>
</tr>
<tr>
<td></td>
<td>Older Brother</td>
<td></td>
<td></td>
<td>2.81</td>
<td>3.48</td>
<td>0.05</td>
</tr>
<tr>
<td></td>
<td>fT level</td>
<td>0.21</td>
<td>0.17</td>
<td>11.92</td>
<td>2.84</td>
<td>0.32**</td>
</tr>
<tr>
<td></td>
<td>Child Sex</td>
<td>0.22</td>
<td>0.01</td>
<td>2.44</td>
<td>1.19</td>
<td>0.16*</td>
</tr>
</tbody>
</table>

*aAQ-Child regression results including all participants.
bAQ-Child regression results excluding individuals scoring >76.

* p<0.05
** p<0.01
Table 4. Examination of weighted AQ-Child subscale scores by sex

<table>
<thead>
<tr>
<th>Variable</th>
<th>Girls (n=117)</th>
<th>Boys (n=118)</th>
<th>t</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>M</td>
<td>SD</td>
<td>M</td>
</tr>
<tr>
<td>Mindreading</td>
<td>0.83</td>
<td>0.46</td>
<td>1.09</td>
</tr>
<tr>
<td>Attention to Detail</td>
<td>1.05</td>
<td>0.56</td>
<td>1.37</td>
</tr>
<tr>
<td>Social Skills</td>
<td>0.58</td>
<td>0.39</td>
<td>0.89</td>
</tr>
<tr>
<td>Imagination</td>
<td>0.43</td>
<td>0.43</td>
<td>0.82</td>
</tr>
<tr>
<td>AQ-Child Total</td>
<td>34.42</td>
<td>15.01</td>
<td>48.75</td>
</tr>
</tbody>
</table>

* p<0.05
** p<0.01
Table 5. Correlations for fT level and AQ-Child Subscales

<table>
<thead>
<tr>
<th></th>
<th>fT Level</th>
<th>Mindreading</th>
<th>Attn. Detail</th>
<th>Social Skills</th>
<th>Imagination</th>
</tr>
</thead>
<tbody>
<tr>
<td>Mindreading</td>
<td>.30**</td>
<td>--</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Attention to Detail</td>
<td>.27**</td>
<td>.35**</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Social Skills</td>
<td>.33**</td>
<td>.62**</td>
<td>.31**</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Imagination</td>
<td>.38**</td>
<td>.38**</td>
<td>.17**</td>
<td>.38**</td>
<td>--</td>
</tr>
<tr>
<td>AQ-Child Total</td>
<td>.41**</td>
<td>.86**</td>
<td>.61*</td>
<td>.83**</td>
<td>.57**</td>
</tr>
</tbody>
</table>
Table 6. Final Hierarchical Regression Model for the CAST

<table>
<thead>
<tr>
<th>Outcome</th>
<th>Predictor Variable</th>
<th>$R^2$</th>
<th>$\Delta R^2$</th>
<th>B</th>
<th>SE</th>
<th>$\beta$</th>
</tr>
</thead>
<tbody>
<tr>
<td>CAST&lt;sup&gt;a&lt;/sup&gt;</td>
<td>Constant</td>
<td>0.02</td>
<td>0.02</td>
<td>2.76</td>
<td>0.42</td>
<td>2.76</td>
</tr>
<tr>
<td></td>
<td>Mother Age</td>
<td>0.02</td>
<td>0.01</td>
<td>11.61</td>
<td>3.23</td>
<td>0.13</td>
</tr>
<tr>
<td></td>
<td>fT level</td>
<td>0.07</td>
<td>0.05</td>
<td>2.76</td>
<td>0.42</td>
<td>0.22**</td>
</tr>
<tr>
<td>CAST&lt;sup&gt;b&lt;/sup&gt;</td>
<td>Constant</td>
<td>0.03</td>
<td>0.03</td>
<td>2.60</td>
<td>0.38</td>
<td>2.60</td>
</tr>
<tr>
<td></td>
<td>Mother Age</td>
<td>0.02</td>
<td>0.01</td>
<td>11.61</td>
<td>3.23</td>
<td>0.13</td>
</tr>
<tr>
<td></td>
<td>Older Sister</td>
<td>0.25</td>
<td>0.17</td>
<td>0.17</td>
<td>0.11</td>
<td>0.11</td>
</tr>
<tr>
<td></td>
<td>Older Brother</td>
<td>0.15</td>
<td>0.14</td>
<td>0.14</td>
<td>0.08</td>
<td>0.08</td>
</tr>
<tr>
<td></td>
<td>fT level</td>
<td>0.05</td>
<td>0.02</td>
<td>0.25</td>
<td>0.12</td>
<td>0.14*</td>
</tr>
</tbody>
</table>

Note: Square-root transformation was conducted before analysis

<sup>a</sup>CAST regression results including all participants.

<sup>b</sup>CAST regression results excluding individuals scoring >15.

* $p<0.05$

** $p<0.01$
Table 7. Means and standard deviations for subset of children by sex

<table>
<thead>
<tr>
<th>Variable</th>
<th>Girls (n=31)</th>
<th>Boys (n=43)</th>
<th>t</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>M</td>
<td>SD</td>
<td>M</td>
</tr>
<tr>
<td>ft level (nmol/L)**</td>
<td>0.35</td>
<td>0.23</td>
<td>0.79</td>
</tr>
<tr>
<td>Gestational Age</td>
<td>16.44</td>
<td>1.25</td>
<td>16.15</td>
</tr>
<tr>
<td>Child Age</td>
<td>9.23</td>
<td>0.80</td>
<td>9.16</td>
</tr>
<tr>
<td>Maternal Age</td>
<td>34.79</td>
<td>4.70</td>
<td>35.71</td>
</tr>
<tr>
<td>Parental Education</td>
<td>3.30</td>
<td>0.81</td>
<td>3.60</td>
</tr>
<tr>
<td>CAST Total^</td>
<td>3.84</td>
<td>3.31</td>
<td>4.70</td>
</tr>
<tr>
<td>AQ-Child Total**</td>
<td>32.35</td>
<td>13.03</td>
<td>46.67</td>
</tr>
<tr>
<td>Full Scale IQ</td>
<td>100.61</td>
<td>13.86</td>
<td>108.95</td>
</tr>
<tr>
<td>Verbal IQ</td>
<td>94.90</td>
<td>12.82</td>
<td>100.26</td>
</tr>
<tr>
<td>Performance IQ</td>
<td>108.13</td>
<td>17.39</td>
<td>116.26</td>
</tr>
<tr>
<td>Block Design</td>
<td>16.26</td>
<td>10.08</td>
<td>19.14</td>
</tr>
</tbody>
</table>

^Square-root transformation carried out prior to analyses
* p<0.05
** p<0.01
Table 8. Correlations for IQ scales and fT level

<table>
<thead>
<tr>
<th></th>
<th>fT Level</th>
<th>Full IQ</th>
<th>Verbal IQ</th>
<th>Performance IQ</th>
</tr>
</thead>
<tbody>
<tr>
<td>Full IQ</td>
<td>.13</td>
<td>--</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Verbal IQ</td>
<td>.02</td>
<td>.83**</td>
<td>--</td>
<td>--</td>
</tr>
<tr>
<td>Performance IQ</td>
<td>.19</td>
<td>.84***</td>
<td>.41**</td>
<td>--</td>
</tr>
<tr>
<td>Block Design</td>
<td>.16</td>
<td>.68**</td>
<td>.27*</td>
<td>.85**</td>
</tr>
</tbody>
</table>

* $p<0.05$
** $p<0.01$
Figure 1. Distribution of AQ-Child scores

Note: Maximum score=141, cut-off score=76
Figure 2. Relationship between fT level and AQ-Child scores
Figure 3. Distribution of CAST scores

Note: Maximum score=31, cut-off score=15
Figure 4. Relationship between fT level and CAST scores