Sex Differences in White Matter Microstructure in the Human Brain Predominantly Reflect Differences in Sex Hormone Exposure

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Abstract
Sex differences have been described regarding several aspects of human brain morphology; however, the exact biological mechanisms underlying these differences remain unclear in humans. Women with the complete androgen insensitivity syndrome (CAIS), who lack androgen action in the presence of a 46,XY karyotype, offer the unique opportunity to study isolated effects of sex hormones and sex chromosomes on human neural sexual differentiation. In the present study, we used diffusion tensor imaging to investigate white matter (WM) microstructure in 46,XY women with CAIS (n = 20), 46,XY comparison men (n = 30), and 46,XX comparison women (n = 30). Widespread sex differences in fractional anisotropy (FA), with higher FA in comparison men than in comparison women, were observed. Women with CAIS showed female-typical FA throughout extended WM regions, predominantly due to female-typical radial diffusivity. These findings indicate a predominant role of sex hormones in the sexual differentiation of WM microstructure, although sex chromosome genes and/or masculinizing androgen effects not mediated by the androgen receptor might also play a role.

Key words: androgens, CAIS, DTI, sexual differentiation, white matter

Introduction
Sex differences have been described regarding several aspects of human brain morphology, although the specific biological mechanisms driving neural sexual differentiation remain to be determined. Numerous neuroimaging studies have focused on macro- and mesoanatomical sex differences, such as in overall or regional gray and white matter (WM) volume derived from structural magnetic resonance imaging (MRI) scans (for meta-analysis see Ruigrok et al. 2014). More recently, diffusion tensor imaging (DTI), an MRI technique that measures characteristics of water molecule diffusion (Basser et al. 1994), has been widely used to study sex differences in WM microstructure. Quantitative measures that can be derived from the diffusion tensor include fractional anisotropy (FA), axial (AD), and radial diffusivity (RD) (Pierpaoli and Basser 1996). FA provides information about the degree of diffusion anisotropy. A low FA value reflects isotropic diffusion, that is, equal diffusion in all directions (e.g., in cerebrospinal fluid). A high degree of anisotropy is found in WM fiber bundles, in which water diffusion is restricted in the direction perpendicular to the axon, and relatively unrestricted along the axon. AD, which equals the largest eigenvalue of the diffusion tensor, and RD, the mean of the two smaller eigenvalues, are...
interpreted as the diffusivity parallel and perpendicular to the WM tract, respectively, and are typically assessed to further characterize FA.

There is evidence for sex differences in WM microstructure as assessed by diffusivity measures. The majority of whole-brain DTI studies have found higher FA in men in major WM regions (Chou et al. 2011; Inano et al. 2011; Menzler et al. 2011; Rametti et al. 2011; Den Braber et al. 2013; Kanaan et al. 2014; Schoonheim et al. 2014; Takao et al. 2014). While sex effects in other diffusivity measures are less frequently studied, and results are less consistent compared with FA, a study in a large sample of 547 men and 310 women, showed that a higher FA in men is generally associated with a higher AD and a lower RD than in women (Inano et al. 2011).

At present, the exact factors and mechanisms underlying the sexual differentiation of WM microstructure are unknown. The classical theory of the sexual differentiation of the brain emphasizes a pivotal role for steroid hormones (Phoenix et al. 1959). Exposure to high levels of androgens during the pre- and/or early postnatal period of neural development is thought to have a permanent, that is, organizing, effect on the brain. Human brain structure, specifically gray matter morphometry, has for instance been associated with fetal testosterone levels in boys at age 8–11 years in some, but not all, sexually differentiated brain regions (Lombardo et al. 2012), and with androgens (with unknown timing) in a study in boys with early androgen excess due to familial male precocious puberty (Mueller et al. 2011). In several species androgens masculinize and feminize the brain after being aromatized to estrogens (reviewed in Baum 1979), whereas studies in human primates and rare clinical conditions in humans suggest that androgens act directly, that is, by activation of the androgen receptor (AR), in the development of the human brain (Cohen-Bendahan et al. 2005; Wallen 2005; Baum 2006; Hamann et al. 2014; van Hemmen et al. 2016). The absence of androgens during early development would induce female–typical neural development, although in mice a role for estrogens in feminizing the brain has recently been suggested (Brock et al. 2011). High sex hormone levels during adolescence and adulthood are thought to have activational, that is, transient, effects on sexually differentiated neural characteristics, while more recently puberty has been proposed as a second sensitive period for organizational sex hormone effects on brain structure (Sisk and Zehr 2005; Schulz et al. 2009).

Even though sex hormones have received most attention from researchers, animal studies have shown that sex chromosome composition might also directly influence neural and behavioral sexual differentiation (e.g., McCarthy and Arnold 2011). It is, however, difficult to disentangle potential direct sex chromosomal effects from sex hormone actions, since sex chromosomes determine gonadal development and are therefore intrinsically related to gonadal hormone levels. While these factors can be manipulated and separately studied in animal models, we depend on naturally occurring disorders/differences of sex development (DSDs) when studying humans. Studies of human brain structure have been performed in men with Klinefelter syndrome (KS), which is characterized by a Y- and two or more X-chromosomes, and women with Turner syndrome (TS), who have one X-chromosome and lack (all or part of) the second sex chromosome. These sex chromosome aneuploidies usually result in a sex hormone deficiency; reduced androgen levels in KS and reduced estrogen levels in TS. Neuroanatomical differences between these groups and comparison men and women include increased or decreased regional volumes (for reviews, see Savic 2012; Hong and Reiss 2014) as well as aberrant WM microstructure, including lower FA throughout major WM tracts in TS (Holzapfel et al. 2006; Yamagata et al. 2012; Villalon-Reina et al. 2013, but see Molko et al. 2004), and a lower FA within the bilateral anterior cingulate and the left internal capsule and arcuate bundle in KS (DeLisi et al. 2005). The exact implications of these findings for the relative influence of sex hormones and sex chromosome genes on the sexual differentiation of human brain structure are, however, unclear, because it is difficult to establish whether these findings represent direct genetic factors related to sex chromosome gene dosage, sex hormone effects, or a combination of these factors.

The complete androgen insensitivity syndrome (CAIS) is a DSD that provides a unique opportunity to assess the relative role and importance of sex hormones and sex chromosomes in human neural sexual differentiation. This rare condition is characterized by a 46,XY karyotype and normally functioning abdominal testes in combination with a complete androgen resistance. The underlying cause is a mutation in the X-linked AR gene, resulting in a nonfunctional AR (Oakes et al. 2008; Hughes et al. 2012). Despite testosterone concentrations within or above the male range (Melo et al. 2003; Doehehnert et al. 2015), CAIS results in a female phenotype and nearly all women with CAIS are androphilic (sexually attracted to men), have a female gender identity, and show female–typical gender role behavior (Masica et al. 1971; Wiśniewski et al. 2000; Hines et al. 2003). Two functional MRI (fMRI) studies in women with CAIS revealed a female–typical activation pattern in response to sexual images (Hamann et al. 2014) and during mental rotation (van Hemmen et al. 2016). This indicates that sex differences in regional brain function related to these tasks are most likely not directly driven by genetic sex, but rather reflect differences in sex hormone exposure.

Measures of brain structure have not yet been studied in women with CAIS. Therefore, in the present study, we acquired DTI scans in 46,XY women with CAIS, as well as 46,XY comparison men and 46,XX comparison women, to determine the relative role of sex hormones versus sex chromosomes in the sexual differentiation of WM characteristics.

Materials and Methods

Participants

From a total of 85 participants, the DTI data from 5 participants had to be excluded from the present study because of anatomical abnormalities (3 comparison women), excessive head movement (1 comparison man) and technical failure (1 woman with CAIS). The final sample consisted of 20 women with CAIS (46,XY), 30 comparison men (46, XY), and 30 comparison women (46, XX). Groups were matched for age and level of education. Participants reported no history of a serious medical, neurological or psychiatric disease, or MRI contraindications. All participants had a right-hand preference for writing and accordingly, with 1 exception (score + 3, i.e., ambidexter), had scores indicating "extreme right-handedness" (+8 to +10) as determined by the Dutch Handedness inventory (van Strien 1992). All participants were heterosexual, that is, women with CAIS were androphilic, and had a sex-typical gender identity and gender role behavior (see van Hemmen et al. 2016 for further details regarding gender-related psychological functioning questionnaires). In this context, "sex-typical" refers to gender-of-rearing, which is male in comparison men and female in comparison women and women with CAIS.

The diagnosis CAIS was based on clinical characteristics in all women. In addition, mutation analyses of the AR gene were
performed using genomic DNA, showing a mutation of the AR gene confirmative of the clinical diagnosis in 13 women, an unclassified variant of the AR gene mutation in 6 women and an inconclusive result in 1 participant. All women with CAIS were gonadectomized and therefore, with one exception, used hormone replacement (estrogens n = 15, combined estrogens and progesterins n = 4) to compensate for the lack of gonadal sex hormone production. Comparison women were not using hormonal contraceptives. Venous blood samples were obtained in all participants to assess circulating levels of estradiol and total (TT) and free testosterone (FT) (see van Hemmen et al. 2016 for further details on hormone assessment).

Women with CAIS were recruited from the databases of the VU University Medical Center Amsterdam and the Erasmus University Medical Center—Sophia Children’s Hospital Rotterdam and from the support group DSDNederland. Comparison men and women were recruited using flyers and advertisements in a local newspaper. The study was approved by the Medical Ethics Committee of the VU University Medical Center Amsterdam (application number NL32740.029.10) and the Erasmus University Medical Center (application number MEC-2010-350). All participants gave their written informed consent according to the Declaration of Helsinki.

MRI data Acquisition

MRI data were acquired at 3T (Signa HDxt, General Electric, Milwaukee, WI, USA). To reduce head motion, foam padding was used in the 8-channel head coil. For DTI, single-shot echo-planar imaging (EPI) was used to acquire 5 nondiffusion weighted (b0) volumes and 30 volumes with noncollinear diffusion gradients (b = 1000 s/mm², repetition time [TR] 13 s, echo time [TE] 86 ms, field of view [FoV] 256 mm, matrix size 128 × 128, 45 axial slices, 2 × 2 × 2.4 mm³ voxels). T₁-weighted anatomical images were acquired using a 3D fast spoiled gradient echo sequence (TR 7.8 ms, TE 3.0 ms, inversion time [TI] 450 ms, 1 mm isotropic resolution).

Data Analysis

Sample Characteristics

Statistical analyses of sample characteristics were performed with IBM SPSS Statistics for Windows, Version 20.0 (IBM Corp., Armonk, NY, USA). For between-group analyses, one-way analyses of variance (ANOVA) with post hoc pairwise tests or, when the assumptions for the use of parametric tests were violated, Kruskal–Wallis and post hoc Mann–Whitney U-tests were used. For all analyses, a P < 0.05 was considered significant and all post hoc test statistics were Bonferroni-corrected for multiple comparisons.

<table>
<thead>
<tr>
<th>Sample Characteristics</th>
<th>Comparison men</th>
<th>Comparison women</th>
<th>CAIS</th>
<th>F/γ²</th>
<th>P-value</th>
</tr>
</thead>
<tbody>
<tr>
<td>n</td>
<td>30</td>
<td>30</td>
<td>20</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Age</td>
<td>31.7 (9.5)</td>
<td>32.0 (9.7)</td>
<td>31.0 (11.0)</td>
<td>0.26</td>
<td>0.878</td>
</tr>
<tr>
<td>Level of education (C)</td>
<td>5.5 (1.8)</td>
<td>5.8 (1.5)</td>
<td>5.8 (1.6)</td>
<td>0.52</td>
<td>0.770</td>
</tr>
<tr>
<td>Level of education (E)</td>
<td>6.3 (1.9)</td>
<td>6.3 (1.7)</td>
<td>6.0 (1.6)</td>
<td>0.69</td>
<td>0.708</td>
</tr>
<tr>
<td>TWMV (mL)</td>
<td>567.8 (70.0)</td>
<td>490.6 (55.3)</td>
<td>541.5 (47.3)</td>
<td>12.88*</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Estradiol (pmol/L)</td>
<td>82.3 (18.2)</td>
<td>391.8 (305.5)</td>
<td>243.3 (151.6)</td>
<td>37.74</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Total testosterone (nmol/L)</td>
<td>13.3 (6.0)</td>
<td>0.9 (3.0)</td>
<td>0.3 (0.1)</td>
<td>68.72</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Free testosterone (pmol/L)</td>
<td>297.6 (128.4)</td>
<td>14.7 (5.2)</td>
<td>3.7 (1.7)</td>
<td>69.08</td>
<td>&lt;0.001</td>
</tr>
</tbody>
</table>

Mean and (SD) per group. Scores for the level of education range from 1 (primary school) to 8 (university degree). Bold P-values represent a significant (P < 0.05) main effect of group.

MRI data

DTI data were preprocessed using FSL 5.04 (Functional Magnetic Resonance Imaging of the Brain Software Library, http://fsl.fmrib.ox.ac.uk/fsl/, last accessed 15 May, 2015, Smith et al. 2004). After motion and eddy current correction and reorientation of gradient vectors, the tensor model was fitted. Three volumes revealing motion-related artifacts were deleted from the raw data in one participant, after which the previous steps were repeated. Further processing and voxel-wise analysis of FA, AD, and RD data were performed with tract-based spatial statistics (TBSS; Smith et al. 2006). TBSS projects all subjects’ FA data onto a mean FA tract skeleton, thresholded at 0.2, before applying voxel-wise cross-subject statistics. The registration and projection parameters used for the FA data were also applied to the AD and RD images.

The skeletonized FA, AD, and RD images were used for whole-brain between-group analyses. First, data from comparison men and women were compared in order to verify sex differences. Subsequently, whole-brain comparisons between these groups and women with CAIS were conducted. A non-parametric permutation method was used (FSL’s randomize; Winkler et al. 2014) with 5000 permutations for each between-group comparison, using threshold-free cluster enhancement (TFCE; Smith and Nichols 2009) and a family-wise error (FWE) corrected P < 0.05 threshold. In addition, between-group analyses of mean FA, AD, and RD, extracted from the regions of the skeleton showing sex differences in FA, were performed to further characterize the overall differences in FA and the contribution of AD and RD to these effects. Although groups were successfully matched for age, the age range was large (18.1–52.7 years). Therefore, age was added as a covariate to all analyses to ensure that any observed differences between the groups were independent of age-related effects.

Total white matter volume (TWMV) was calculated from T1 images after brain extraction using BET (implemented in FSL) with image bias and residual neck voxel reduction (−8) and a 0.1 fractional intensity threshold, and segmentation using the voxel-based morphometry toolbox (VBM; http://dbm.neuro.uni-jena. de/vbm8/, last accessed 15 May, 2015) for statistical parametric mapping (SPM8; Wellcome Trust Center for Neuroimaging, Institute of Neurology at UCL, UK) with a medium bias regularization.

Results

Sample Characteristics

Sample characteristics are summarized in Table 1. The three groups did not differ in age and level of education. Serum
hormone levels obtained at the day of testing showed significant between-group differences. Estradiol levels in comparison women and women with CAIS were not significantly different, whereas both groups of women had higher estradiol levels than men (P < 0.001). TT and FT levels were highest in comparison men, followed by comparison women, and women with CAIS having the lowest TT and FT levels (all P < 0.001). TWMV showed a significant effect of group, with larger TWMV in men and women with CAIS than comparison women (P < 0.001 and 0.012, respectively).

**Diffusion Tensor Imaging (TBSS)**

**Fractional Anisotropy**

Whole-brain between-group TBSS analyses revealed widespread sex differences in FA (Fig. 1). Comparison men showed higher FA than comparison women in a single cluster covering a large part of the skeleton, including major WM tracts, subcortical regions such as the bilateral thalamus and basal ganglia, and the brainstem. Comparison women did not show any regions with higher FA than comparison men. Similar differences in FA were found between comparison men and women with CAIS (Fig. 1); comparison men had higher FA in a large part of the skeleton, whereas the reverse contrast did not show any significant differences. No differences in FA were observed between comparison women and women with CAIS.

Similar between-group effects, with large effect sizes for the differences between comparison men and both comparison women and women with CAIS, were observed for the mean FA values extracted from the FA sex difference cluster (Fig. 2 and Table 2).

**Axial, Radial Diffusivity**

Whole-brain TBSS analyses of AD and RD also revealed between-group differences (Fig. 1). Comparison men showed higher AD values than comparison women throughout extensive parts of the skeleton, without differences in the reverse contrast. Women with CAIS showed differences in AD relative to both comparison groups: lower AD than comparison men in a left hemisphere cluster, including (parts of) the internal and external capsule, sagittal stratum, superior longitudinal fasciculus, and posterior thalamic radiation, and higher AD than comparison women in a right hemisphere cluster, comprising (parts of) the internal and external capsule, corona radiata, superior longitudinal and fronto-occipital fasciculus, corpus callosum (splenium), cerebral peduncle, and posterior thalamic radiation.

A sex difference in RD, with higher RD values in comparison women than in men, was observed in a cluster covering (parts of) the left internal and external capsule, corona radiata, cerebral peduncle, uncinate and superior fronto-occipital fasciculus, pallidum, and amygdala. Women with CAIS showed higher RD values than comparison men in a small cluster of 25 voxels located within the right anterior corona radiata. Women with CAIS also showed higher RD than comparison women in a cluster covering part of the right superior longitudinal fasciculus.

Mean AD and RD (Fig. 2 and Table 2), extracted from the FA sex difference region, showed sex differences with large effect sizes. In comparison men, a higher AD and lower RD relative to comparison women was found. Compared with women with CAIS, a tendency towards higher AD and a statistically significantly lower RD, with a large effect size, was found in comparison men. Comparison women did not differ from women with CAIS.

**Discussion**

The present study investigated the relative role of sex hormones versus sex chromosome genes on the sexual differentiation of human WM microstructure. This question was addressed by comparing DTI data from a unique group, that is, 46,XY women with CAIS, to 46,XY comparison men and 46,XX comparison women. Anisotropy findings suggest a more important role for sex hormones, most likely masculinizing/defeminizing androgen and/or feminizing estrogen effects, than for genetic effects related to sex chromosome genes in the sexual differentiation of WM microstructure. Furthermore, our findings regarding the underlying DTI metrics showed that these sex hormone influences are most pronounced in RD, while AD, to some extent, might also reflect other factors directly related to sex chromosome composition and/or masculinizing androgen actions that are not mediated by the AR.

Widespread sex differences in FA were found, showing higher FA in comparison men than women, in line with the majority of findings in DTI studies using TBSS to investigate sex differences (e.g., Inano et al. 2011). Similar to comparison women, women with CAIS had lower FA than comparison men throughout extensive regions of the skeleton, showing great overlap with sex difference regions. Furthermore, their FA values did not differ from those of comparison women. These findings were also substantiated by the mean FA values in the region demonstrating a sex difference in FA.

In addition to FA, we analyzed the underlying DTI metrics AD and RD, to assess their contribution to the differences observed in FA. Based on the whole-brain voxel-wise analyses of AD and RD, higher FA in men appeared to be predominantly associated with higher AD. However, the mean AD and RD in areas with FA sex differences showed that the sex difference in FA was due to both higher AD and lower RD in men than in women, with equal large effect sizes. Although results from previous studies on sex effects in these DTI measures are less consistent than in FA, the current results are generally in accordance with a study in a large sample of 857 subjects (Inano et al. 2011).

In the whole-brain voxel-wise comparisons to women with CAIS, higher AD contributed significantly to higher FA in men. This contribution of AD was, however, observed in a considerable smaller area than in the comparison groups, only partially covering the areas where FA differences were found. In addition, a difference relative to comparison women was found, with higher AD in women with CAIS in part of the right hemisphere skeleton. In accordance with these voxel-wise results, the mean AD in the FA sex difference region showed a tendency, but did not reach statistical significance, toward a female-typical AD in women with CAIS. While a higher RD in women with CAIS relative to both comparison groups was apparent in relatively small regions within the skeleton, the mean RD in the part of the skeleton showing FA sex differences was female-typical in women with CAIS, that is, higher than in comparison men. Within the FA sex difference region, both the sex difference in mean RD, and the female-typical mean RD in women with CAIS, appears to result from a widespread, subtle effect, which is only detected in the mean RD calculated from all individual voxel values and, apart from some small clusters, is not detected in the whole-brain voxel-wise analysis.

Taken together, the female-typical FA in women with CAIS is in agreement with previous findings in neural activity patterns in response to visual sexual stimuli and mental rotation performance (Hamann et al. 2014; van Hemmen et al. 2016). The present findings thus argue against a dominant role for sex chromosome
Figure 1. Whole-brain between-group differences (P < 0.05 FWE-corrected) in FA, AD, and RD. Clusters of significant voxels are projected on the mean FA skeleton (green), using the “TBSS fill” procedure in FSL, and the MN152 T1-weighted standard brain (FSL). Slice labels refer to MNI z-coordinates. MvsW, comparison men versus comparison women; MvsC, comparison men versus women with CAIS; WvsC, comparison women versus women with CAIS. For comparison XvsY, X > Y effects are displayed in red-yellow, X < Y effects are displayed in blue-green. Left is left.
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genes and rather indicate that the sexual differentiation of human WM microstructure is primarily associated with differences in sex hormone exposure. Due to the complete androgen resistance in CAIS, it is likely that these sex hormone effects reflect feminizing actions through direct activation of the AR. An important role for andro-
genome acting on the AR would indeed be consistent with findings from animal studies, showing the presence of ARs in cortical axons, dendrites, and glial cells (Sarkey et al. 2008). Furthermore, an association between testosterone and WM development in human adolescents was modulated by functional AR gene var-

Table 2 Between-group statistics and effect sizes for mean FA, AD, and RD within FA sex difference regions

<table>
<thead>
<tr>
<th></th>
<th>F</th>
<th>P-Value</th>
<th>Effect size</th>
</tr>
</thead>
<tbody>
<tr>
<td>FA</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Main effect of group</td>
<td>25.435</td>
<td>&lt;0.001</td>
<td>0.401</td>
</tr>
<tr>
<td>PH: M versus W</td>
<td>&lt;0.001</td>
<td>2.01</td>
<td></td>
</tr>
<tr>
<td>PH: M versus C</td>
<td>&lt;0.001</td>
<td>1.28</td>
<td></td>
</tr>
<tr>
<td>PH: W versus C</td>
<td>0.126</td>
<td>0.64</td>
<td></td>
</tr>
<tr>
<td>AD (×10⁻³ mm²/s)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Main effect of group</td>
<td>8.247</td>
<td>0.001</td>
<td>0.178</td>
</tr>
<tr>
<td>PH: M versus W</td>
<td>&lt;0.001</td>
<td>1.04</td>
<td></td>
</tr>
<tr>
<td>PH: M versus C</td>
<td>0.073</td>
<td>0.66</td>
<td></td>
</tr>
<tr>
<td>PH: W versus C</td>
<td>0.592</td>
<td>0.38</td>
<td></td>
</tr>
<tr>
<td>RD (×10⁻³ mm²/s)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Main effect of group</td>
<td>10.861</td>
<td>&lt;0.001</td>
<td>0.222</td>
</tr>
<tr>
<td>PH: M versus W</td>
<td>&lt;0.001</td>
<td>0.19</td>
<td></td>
</tr>
<tr>
<td>PH: M versus C</td>
<td>0.028</td>
<td>0.77</td>
<td></td>
</tr>
<tr>
<td>PH: W versus C</td>
<td>0.451</td>
<td>0.42</td>
<td></td>
</tr>
</tbody>
</table>

Bold and italic P-values represent a significant (P < 0.05) effect and a trend (P < 0.1), respectively. For post hoc (PH) tests a Bonferroni-correction for multiple comparisons was applied. Effect sizes are expressed in partial η² for the main effect of group and Cohen’s d for pairwise comparisons.

M, comparison men; W, comparison women; CAIS, women with CAIS.

studies investigating an association between circulating estradiol levels and FA have not shown consistent results (Herting et al. 2011; Peper et al. 2013; Kranz et al. 2014).

The female-typical FA in women with CAIS was predominantly due to a female-typical RD, although a contribution of AD, at least in some of the WM tracts with a sex difference in FA, cannot be ruled out. Consequently, these results indicate that the proposed sex hormone effects are most pronounced in WM microstructural characteristics related to RD. The absence of a compelling female-typical AD in women with CAIS was unexpected based on previous neuroimaging studies in these women (Hamann et al. 2014; van Hemmen et al. 2016), although a recent behavioral study on spatial learning in women with CAIS also did not observe an overall female-typical pattern (Mueller et al. 2016); women with CAIS did not differ significantly from comparison men and women on any of the variables measuring spatial learning, and effect size analyses revealed a more female-like pattern in women with CAIS on some, but not all variables. Regarding the sexual differentiation of the microstructural WM characteristics associated with AD, the present findings are suggestive of a contribution of other factors than the previously proposed sex hormone effects. These factors may include direct effects of sex chromosome genes, or other feminizing/defeminizing androgen actions that are not mediated by the AR. It is important to note that AD in women with CAIS differed from comparison women in only a small part of the skeleton, while the effect size of the summarizing mean AD within the FA sex difference region was in the female direction. Combined with the predominant female-typical RD and FA, this indicates that the contribution of these other factors is less pronounced and less extensive than the hypothesized masculinizing/defeminizing androgen or feminizing estrogen effects.

An interesting finding in light of the female-typical FA in women with CAIS is that their TWMV is more similar to that in comparison men. A potential explanation is that overall WM volume is susceptible to direct sex chromosomal effects or androgen actions that are not mediated by the AR. Nonetheless, the present findings in WM microstructure are not likely to result from differences in TWMV for 2 reasons. First, susceptibility to partial volume effects (PVEs), known to affect the accuracy of DTI mea-

Figure 2. Mean corrected for age (±SEM) FA, AD, and RD within FA sex difference regions. M, comparison men; W, comparison women; CAIS, women with CAIS.
be ruled out for some aspects of WM microstructure. After aromatization to estrogens, acting on the estrogen receptor, differences in sex chromosome composition, or androgenic effects feminizing estrogen effects, or a combination of both. Furthermore, masculinizing and/or defeminizing androgen effects through the AR, supportive of an important role for sex hormones, most likely mas-

ual differentiation of human WM microstructure. Our unique insights into the biological mechanism underlying the sex-

sexual differentiation of human WM microstructure. In conclusion, this DTI study in women with CAIS presents remarkable and

ber tracts, that are wider than the voxel size, are unaffected by PVEs. Second, the overall female-typical findings in women with CAIS are in the opposite direction as would be expected when TWMV drives the between-group differences.

The tracts showing sex differences in WM microstructure connect extensive parts of the cerebral cortex and subcortical brain regions. The widespread differences in FA were found throughout major WM tracts, including projection fibers such as the internal capsule and the corona radiata, containing axons going to and from the cerebral cortex, association fibers such as the external capsule, connecting different cortical regions within a hemisphere, and parts of the corpus callosum, which is responsible for interhemispheric connections. These findings might reflect general differences in structural connectivity of the male and female brain, which does not show a clear association with regional gray matter sex differences, which have generally been found to be limited to certain brain regions (for meta-analysis see Ruigrok et al. 2014). Based on the present study it is, therefore, difficult to draw conclusions about the exact functional implications of these findings with regard to specific behavioral and cognitive sex differences.

It is also difficult to interpret the current results with respect to the underlying physiological characteristics. FA is a sensitive, but rather nonspecific measure, since differences in FA could reflect several physiological differences. Even though AD and RD provide more specific information about FA, the exact physiological correlates of these measures are still under investigation (Jones et al. 2013). The predominant contribution of RD to the female-typical FA in women with CAIS in sexually differentiated WM regions might, for instance, reflect sex hormone effects on axonal density, diameter or the degree of myelination.

A limitation of the present study is that no inferences can be made about the exact timing of the proposed sex hormone effects. Since women with CAIS are insensitive to androgens throughout life, the current findings might reflect organizational and/or activational sex hormone effects during the pre-/neonatal phase, adolescence and/or adulthood. Longitudinal studies in women with CAIS, starting before adolescence, could provide information regarding the timing of sex hormone effects on the sexual differentiation of human WM microstructure.

In conclusion, this DTI study in women with CAIS presents unique insights into the biological mechanism underlying the sexual differentiation of human WM microstructure. Our findings are supportive of an important role for sex hormones, most likely masculinizing and/or defeminizing androgen effects through the AR, feminizing estrogen effects, or a combination of both. Furthermore, current findings argue against substantial genetic effects related to differences in sex chromosome composition, or androgenic effects after aromatization to estrogens, acting on the estrogen receptor, although a more subtle contribution of these factors should not be ruled out for some aspects of WM microstructure.

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