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Visual phenotype in Williams-Beuren syndrome challenges magnocellular theories explaining human neurodevelopmental visual cortical disorders

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Williams-Beuren syndrome (WBS), a neurodevelopmental genetic disorder whose manifestations include visuospatial impairment, provides a unique model to link genetically determined loss of neural cell populations at different levels of the nervous system with neural circuits and visual behavior. Given that several of the genes deleted in WBS are also involved in eye development and the differentiation of retinal layers, we examined the retinal phenotype in WBS patients and its functional relation to global motion perception. We discovered a low-level visual phenotype characterized by decreased retinal thickness, abnormal optic disk concavity, and impaired visual responses in WBS patients compared with age-matched controls by using electrophysiology, confocal and coherence in vivo imaging with cellular resolution, and psychophysics. These mechanisms of impairment are related to the magnocellular pathway, which is involved in the detection of temporal changes in the visual scene. Low-level magnocellular performance did not predict high-level deficits in the integration of motion and 3D information at higher levels, thereby demonstrating independent mechanisms of dysfunction in WBS that will require remediation strategies different from those used in other visuospatial disorders. These findings challenge neurodevelopmental theories that explain cortical deficits based on low-level magnocellular impairment, such as regarding dyslexia.

Introduction

Williams-Beuren syndrome (WBS) is a genetic neurodevelopmental disorder characterized by a well-defined hemideletion on chromosome 7q11.23 and predominant visuospatial impairment with relatively preserved auditory and verbal processing (1, 2).

We and others have previously found consistent impaired performance in high-level structure from motion (3D object perception from motion cues) perceptual tasks (3–5), which corroborates previously established anatomical and functional cortical–dorsal stream abnormalities in WBS (6). The cortical–dorsal stream pathway is involved in motion perception and action planning, as opposed to the ventral stream pathway, which is involved in pattern and object recognition and is believed to be less impaired in WBS. The cortical–dorsal stream pathway receives largely predominant input from the retinal magnocellular pathway (see ref. 3 and references therein). This pathway responds with high fidelity to local temporal changes in the visual scene. Global motion integration processes seem to be quite specific (3, 7) and may localize to separate independent functional subdivisions within the dorsal stream (3, 8–10). Accordingly, it is essential to define

Furthermore, it has recently been shown that *GTF2IRD1* (a member of the *GTF2I* transcription factor family) plays a pivotal role in the craniofacial and cognitive phenotype (15), and recent analysis in patients with small deletions has shown that haploinsufficiency of *GTF2IRD1* and/or *GTF2I* may also contribute to the visual phe-

and understand the correlation of hierarchical structural/func-

tional abnormalities at distinct levels of processing, including the

Given the expression of several critical genes involved in WBS

in the neural retina, it is surprising that a functional and structural characterization of this region has so far, to our knowledge,

not been described (reviewed in ref. 6). For example, GTF2I (11)

and LIMK1 (12, 13) are expressed in the retina (14) and may play a

notype in WBS (16, 17).

modulatory role in the visual phenotype.

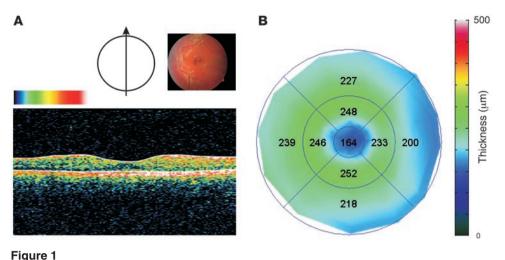
retina and the cortex.

During early retina development in mice, *Gtf2ird1* is involved in cell proliferation, thereby influencing eye size as well as the relationship between all neural layers (18). Later in development, the localization of its protein to the outer nuclear layer suggests that it may influence the differentiation of this particular region (18). Further evidence for an important role of that protein in retinal development and the likely modulatory role of other genes is suggested by the fact that the mammalian GTF2IRD1 protein binds to the retinoblastoma (Rb) protein, thereby serving as a positive transcription regulator of development (19). Since disruption of Rb function might result in deficiencies during the developmental process, it is important to verify whether

Nonstandard abbreviations used: cd, candela; CS, contrast sensitivity; HRT, Heidelberg retinal confocal tomography; mfERG, multifocal electroretinography; OCT, optical coherence tomography; RNFL, retinal nerve fiber layer; RT, retinal thickness; VA, visual acuity; WBS, Williams-Beuren syndrome.

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Central retinal phenotype in a representative WBS subject. (A) Optical biopsy using OCT of the central macular region of the retina reveals a normal layering of retinal structures in a WBS (WBS1) participant (color-coded log reflectivity map: red, high; black, low). An intact foveomacular depression is visible. However, quantitative morphometry revealed a decrease in RT. The bottom hyperreflective layer corresponds to the pigment epithelium and the top one to the ganglion cell RNFL. The hyporeflective region represents the photoreceptor layer. Top right inset represents a fundus photograph of the subject's retina. Arrow in left inset shows axis of depicted OCT image. (B) Color-coded thickness map of the central 20 degrees of the same retina depicted in A (for details, see Methods). A generalized loss is visible (for statistical details, see text). Numbers indicate regional RT.

there are functional consequences stemming from an impact on the integrity of multiple retinal cell layers in WBS.

We therefore aimed to identify a retinal phenotype by examining low-level visual processing. This would also help explain our previous findings suggesting separable early (magnocellular) and high-level (dorsal stream) visual deficits in WBS patients well characterized at the molecular level (3). We provide objective anatomical and physiological data, which we believe are novel, by means of noninvasive optical coherence tomography (OCT), confocal scanning laser tomographic imaging (Heidelberg retinal confocal tomography [HRT]), and multifocal electrophysiological (multifocal electroretinography [mfERG]) techniques that can explain an independent visual phenotype in WBS. Accordingly, our results demonstrate that the objectively identified neural phenotype at the level of the retina predicts low-level visual deficits (3) and is independent of higher-level visual cortical-dorsal stream damage.

Results

Molecular characterization. Molecular studies showed that all 13 patients had the most common 1.55 Mb deletion mediated by nonallelic homologous recombination between the blocks of segmental duplications that flank the WBS locus (20). Therefore, they all are hemizygous for at least 26 genes located in the commonly deleted interval, including GTF2IRD1 and GTF2I at the telomeric edge (see Supplemental Figure 1; supplemental material available online with this article; doi:10.1172/JCI32556DS1). Deletion was on the maternal chromosome in 8 patients and on the paternal chromosome in 5. We precisely localized the breakpoints on the blocks of the segmental duplications of the recombinant chromosome of each patient between 2 paralogous sequence variants. Thus, we could determine whether the deletion affected the functional copies of 2 genes in the breakpoint regions, NCF1 and GTF2IRD2 (see Supplemental Figure 1). The finding of a recombinant chromosome with a gain of telomeric-type paralogous sequence variants permitted us to infer that the transmitting progenitor was a carrier of an inverted chromosome in just 1 case. The telomeric and functional copy of GTF2IRD2 was affected by the deletion in 3 cases; NCF1 was also deleted in those cases as well as in 2 additional patients. The remaining 8 patients had deletions including GTF2I but not affecting the distal 2 genes NCF1 and GTF2IRD2.

In summary, all cases had similar deletion size and were hemizygous for GTF2IRD1 and GTF2I. The parental origin of the hemideletion and the precise location of the breakpoints did not seem to play a major explanatory role in the observed phenotype (see below), which is consistent with the similarity of deletion size.

Baseline craniofacial and ocular phenotypic profile in WBS patients: relative preservation of low-level measurements of visual performance, such as visual acuity. We performed complete eye examinations on our panel of 13 WBS patients. All patients displayed the common facial features of flat midface, periorbital fullness, slight telecanthus, and variable degrees of epicanthal folds. The characteristic full-blown stellate pattern of the anterior iris leaflet was present in all but 2 subjects (whose 4 eyes showed another consistent pattern of abnormality; for details, see Methods). Our WBS subjects had overall preserved best-corrected distance visual acuity (VA) $(-8.9/10 \pm 1.5 \text{ SD})$, which is consistent with previous reports (6). Stereopsis, which is known to be consistently reduced or absent in WBS, was partially preserved in our sample (for details on these measurements and refractive correction, see Methods). Examination of the fundus of the eye, documented with fundus photography after papillary dilation, was performed in all cases. We did not observe significant changes in the macula and periphery of the retina. The optic discs showed subtle deviations from their normal appearance, namely a hint of a semitranslucent, membrane-like structure covering the disc surface in 5 cases and slight deviations in cupping patterns. However, the normal appearance of the central retina and the subtle abnormalities in the optic nerve region did not predict the profound changes that we were able to document with high-resolution imaging techniques (OCT and HRT).

Noninvasive assessment of retinal layers: evidence for loss of inner layers and the ganglion cell retinal nerve fiber layer. We obtained cross-sectional images of ocular microstructures using OCT. This optical assessment of the central macular region of the retina revealed an apparently normal pattern of organization of the different retinal layers in all WBS participants. Figure 1A depicts a representative example in a WBS subject, showing an apparently intact foveomacular depression. However, quantitative morphometry revealed a decrease in retinal thickness (RT) in comparison with



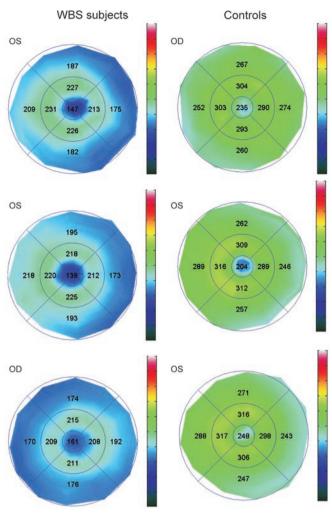


Figure 2 RT maps (in μ m) of the central 20 degrees of the retina depicted in 3 control subjects and 3 WBS participants (WBS3, WBS7, and WBS8).

an age-matched control, healthy population (Figures 1 and 2). The OCT maps depicted in Figure 2 further document the generalized loss in thickness across the outer retina in 3 other WBS subjects compared with 3 representative controls. Figure 3 illustrates the overall reduction in all 9 OCT-defined regions explored in a region covering 20 degrees of the WBS retina in comparison with healthy subjects. In spite of the preserved morphology of the foveomacular pit, RT maps show a significant reduction when compared with age-matched controls (P < 0.0001 for all 9 OCT-defined regions; Mann-Whitney U test).

Figure 3 shows that the ganglion cell retinal nerve fiber layer (RNFL) was more subtly damaged in comparison with the inner retinal (photoreceptor and bipolar cell) layers. Regions 6, 7, and 9 (outer contiguous regions with nasal bias) showed significant reduction of the ganglion cell RNFL (P < 0.01; Mann-Whitney U test).

Noninvasive morphometry of the optic nerve region using OCT and confocal scanning laser imaging (HRT): evidence for structural reorganization and global loss of ganglion cell nerve fibers. We also analyzed the optic disc region with OCT. We generally observed an abnormal pattern of anatomical organization of this region in WBS patients, in contrast with the normal macular pattern. Abnormalities included

loss of normal surface curvature and greater surface irregularities. Figure 4 illustrates orthogonal OCT slices in 3 WBS subjects, which show a striking loss of the normal concavity that defines the disc cup. A partial or almost complete obliteration is evident in these cases. Subject WBS3 in Figure 4 shows an intriguing cup occlusion by a thin membrane-like formation, which in fact transforms the concavity into a cyst-like structure. It is worth noting that a similar membrane was present in the contralateral eye of the same subject (not shown), which suggests that both eyes underwent very similar developmental trajectories. The cup reorganization did not necessarily lead to full obliteration. In fact, in 2 of our subjects (corresponding to 4 eyes), we observed localized and abnormal steepening of the cup walls, concomitant with widening of the "neck" region and deepening of the cup (see below).

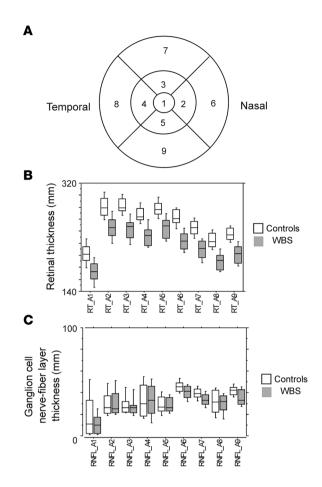
We confirmed the surprising differences and bimodal pattern of reorganization in surface geometry in most WBS subjects, which were documented on OCT images of the optic nerve region by means of confocal scanning laser imaging. Measurements such as mean cup depth, cup/disc area ratio, and cup volume below surface and rim area were often outside the 95% confidence intervals, confirming the findings observed by means of OCT. We also verified that some WBS subjects showed a tendency for the cup structure to be obliterated (compare Figures 4 and 5 for OCT and HRT measurements). Other subjects' data showed the opposite phenomenon, e.g., an increased depth of the depression (e.g., partially in the case of WBS6 and almost completely in the case of WBS5), with abnormally steep walls (see Figure 5B). Interestingly, within-subject patterns were similar across eyes, suggesting common developmental trajectories of the optic nerve and correlated ontogenesis (Figures 4 and 5). Accordingly, either bilateral obliteration or increased depth patterns could be observed (Figures 4 and 5). This between-subjects bimodal pattern suggests the occurrence of hysteresis phenomena that may reflect space-time shifts in morphogenetic processes that fail to reach a final structural stability.

This pattern of results was confirmed when we attempted to independently provide a mathematical fit to the raw image data with "cylindrical fits" of the "neck region" of the optic disc depression (for details, see Methods). This procedure yielded simple estimates of depression depth in normal control subjects and uncovered a bimodal profile in WBS subjects, similar to the one found by above mentioned classical OCT and HRT based methods: we noted instances in which no depression was found by the fitting algorithm and, in cases in which it was found, it could then reach large values. Similar results were obtained with a least-square fitting procedure that modeled the optic nerve head by a smooth 2D surface described by up to 10 free parameters (21). The WBS subjects in whom the blind fitting procedure did not find a depression were the same ones in which OCT measurements documented partial or almost complete obliteration of the normal cup structure. The outcome of the fitting procedure (failure vs. no failure) was significantly different in the WBS population (Fisher's exact test; P value = 0.0033), further confirming the OCT evidence for anatomical abnormalities in WBS subjects.

Finally, we analyzed mean RNFL thickness in the optic disk region. We found a significant decrease in our WBS sample (P = 0.0013; Mann-Whitney U test).

mfERG recordings show patterns of localized impairment of retinal responses in WBS. The mfERG is a technique that provides spatial information about physiological responses in the retina not readily available in the classical ERG. Figure 5C depicts the group analysis





(percentile box plots) of multifocal electrophysiological responses in WBS subjects as compared with age-matched controls (for statistical details, see below).

Figure 6A shows the typical pattern of responses observed in a control subject. Central waveforms were in general proportionately larger than in the visual periphery due to the higher cell density, as further documented by the color-coded 3D plot, which shows responses normalized by the area of stimulation. Figure 6B shows responses from the 2 simultaneously recorded eyes of a WBS subject. Both show regions of preserved waveforms and areas with clearly abnormal responses. The retinal (intraocular) pattern of regional asymmetry in response integrity also generalizes to the interocular comparison.

Group analysis (see also Figure 5C) was sorted by rings of increasing eccentricity (R1–R5). It was evident that the P1 component was more significantly reduced in WBS than the N1 component. Non-parametric statistics confirmed this observation, with a significant involvement of all rings concerning the P1 response (Mann-Whitney U test: ring 1, P = 0.0001; ring 2, P = 0.0001; ring 3, P = 0.0001; ring 4, P = 0.0001; and ring 5, P = 0.0001). A less dramatic but still significant effect was found for the N1 component. In this case, we found an overall significant effect that was, however, relatively more modest across rings (Mann-Whitney U test: ring 1, ns; ring 2, P = 0.014; ring 3, P = 0.0001; ring 4, P = 0.0001; and ring 5, P = 0.031).

Correlations between structural and functional markers: evidence for independent sources of retinal damage that predict local psychophysical deficits independently of visual integration performance measurements. Figure 7 illustrates the basic dichotomy identified by our study: a retinal

Figure 3

Analysis of RT and RNFL thickness across groups. (A) Scheme of regions where RT and RNFL were analyzed. Numbers code different regions. (B) Overall thickness reduction in all 9 regions (see A for respective local labels) explored in the central 20 degrees of the WBS retina in comparison with age-matched control subjects, as illustrated by percentile plots. Box boundaries correspond to upper and lower twenty-fifth percentiles, outer bars to the tenth percentiles, and middle bars to the median. In spite of the preserved morphology of the foveomacular pit, we observed a significant thickness reduction in WBS subjects when compared with age-matched controls (P < 0.0001 for all 9 regions, Mann-Whitney U test). (C) The ganglion cell RNFL thickness in the central part of the retina was less dissimilar between groups, differences only being significant in outer and nasal sectors. RNFL, ganglion cell RNFL.

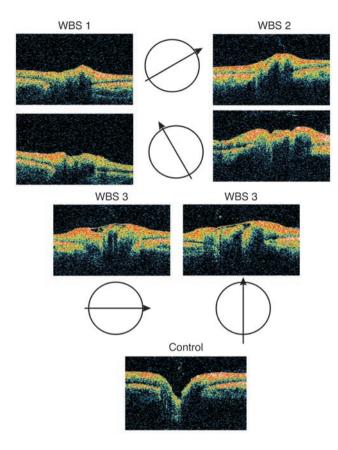
phenotype can predict local magnocellular deficits but not highlevel cognitive function (see below). RT measurements proved to be the best predictive retinal biomarkers and also explained part of the variance associated with the disruption in their efferent ganglion cell outputs, as revealed by the significant correlation with the reduced thickness of the ganglion cell RNFL (Rho = 0.344; P = 0.001; Spearman's correlation analysis). On the other hand, no correlations between RNFL thickness and functional measurements were observed, indicating a significant involvement of the outer retina in the correlations found.

The sensitivity of RT measurements for explaining low-level (not requiring motion integration) visual performance in WBS subjects is probably due to the more local sampling properties of outer retinal neurons (less convergence). Low-level local speed discrimination tasks showed significant correlations with RT measurements for both temporally constrained and unconstrained tasks (Rho = -0.451, P = 0.038; and Rho = -0.509, P = 0.019, respectively; see Figure 7). Note that correlations were negative because low-speed discrimination corresponded to high-threshold velocities that match a standard speed and low outer RT (fewer sampling retinal neurons). The same held true for tasks requiring local motion direction discrimination, regardless of whether subjects had unlimited stimulus presentation or stimulation was confined to approximately 1 second (Rho = -0.79, P = 0.0003; and Rho = -0.77, P = 0.0004, respectively; see Figure 7).

Interestingly, and unlike all local tasks, we found that global motion tasks using exactly the same type of local stimuli yielded patterns of performance that were found to be in general independent of retinal measurements (see Figure 7). Indeed, global motion integration 2D coherence tasks (requiring detection of dot-defined 2D surfaces immersed in noise) did not show any correlation with retinal structural measurements for short presentations (which implies relatively limited processing times of up to 1 second for motion integration). These conditions enabled us to isolate a predominant cortical mechanism, unlike the local discrimination tasks. For unlimited stimulus presentations, where local sampling could be optimized, cumulative bottom-up retinocortical processing was probably helpful, since a beneficial effect could then be found (Rho = -0.69, P = 0.0012; see Figure 7).

Concerning 3D motion coherence tasks (requiring detection of dot-defined 3D spheres immersed in noise), extremely time-constrained (200 ms) stimulus presentations may have led to suboptimal sampling and a retinal bottleneck for such short time frames. Indeed, in a few cases with low RT, stimulus visibility was probably compromised because performance then dropped





to near chance levels. In the remaining subset, no correlation was found with retinal measurements (see Figure 7).

In summary, we also have evidence for an independent cortical component both for 2D and 3D motion integration tasks. Figure 7 depicts the basic dichotomy, suggesting the outer retinal layers as the main cause of the low-level functional phenotype. It also highlights the absence of correlation of RT with measurements of dorsal stream function, such as visuospatial ability, as measured by Benton's Judgement of Line Orientation Test, which requires angular comparisons, and is known to assess parietal function. RT was also not correlated with measurements of ventral stream function, as assessed by a standard test of facial discrimination. Our measurements of dorsal stream and ventral stream function were, as expected, uncorrelated. Motion integration measurements were uncorrelated with magnocellular contrast-sensitivity (CS) measurements, further suggesting that a magnocellularbased CS deficit cannot be directly related to higher-level steps in motion processing.

Concerning the correlational structure of motion tests, they share, as expected, some common variance, but it is interesting to point out that our analysis also suggests that our assessment techniques are probably tapping distinct mechanisms within the dorsal stream. Accordingly, 3D structure from motion performance measurements was shown to dissociate from 2D motion coherence and visuospatial performance (using either a nonparametric or partial-correlation approach), suggesting that they require distinct cortical networks for adequate performance.

Given that the characterization of the deletions showed that they were all identical in size, variation was only present in the number of functional copies of 2 genes in the breakpoint regions,

Figure 4

Optical biopsy (OCT) of the optic disc region of the retina depicting abnormal cupping and layering of retinal structures in 3 WBS participants (WBS1, WBS2, WBS3). Arrows depict orthogonal axes of optical section. Although we observed that there is a bimodal pattern of deviation in our WBS group (see text), the cup region most often loses concavity, either by obliteration or partial covering by a thin membrane (subject WBS3).

NCF1 and *GTF2IRD2*, and the parental origin of the hemideletion. These variables, however, did not show predictive value regarding phenotype severity.

Discussion

Our study addresses structure-function correlations that unrayel a hitherto unsuspected neural retinal phenotype in Williams-Beuren syndrome that explains patterns of visual deficits that are independent of previously described cortex-based phenotypes (6). This suggests a new framework for hierarchical phenotyping and identification of independent mechanisms contributing to complex and multifactorial diseases of the nervous system. We show, we believe for the first time, patterns of dysfunction in early retinal circuits that are consistent with the previous demonstration that several genes, including *LIMK1* (reviewed in ref. 6) and GTF2IRD1, are pivotal in the development of retinal neural layers (18) and contribute to visual deficits in WBS (16, 17). Furthermore, GTF2I, which is also included in the deletion region in WBS, is also involved in neuronal maturation and is exclusively present in neurons (11). Converging evidence for the involvement of these genes in the WBS visual phenotype is provided by recent studies in patients with small deletions, confirming that the presence of GTF2IRD1 in humans leads to a less severe visual phenotype, and animal studies showing that loss of Gtf2ird1 in mice also results in a phenotype that is consistent with the expression pattern of Gtf2ird1 in the developing brain and craniofacial areas, including the eye and the retina (15, 18). However, one should not neglect the role of other genes expressed in the retina, such as LIMK1 and GTF2I (Gene Expression Omnibus profiles data: http://www.ncbi.nlm.nih.gov/geo/).

Moreover, spatiotemporal patterns of expression of genes such as *GTF2IRD1* and *GTF2I* can be related to the observed phenotypical features. During early retina development, *Gtf2ird1* is involved in proliferation across all neuronal layers, thus controlling final cell number (18). Later on in development, it instructs differentiation of the outer layer with an impact on cellular function (19). Finally, Proschel et al. showed that *Limk1* is expressed in the inner nuclear layer of the retina during embryogenesis and also in adulthood (14).

Given these facts, we have explicitly looked for anatomical and physiological evidence for the involvement of the neural retina in the visual phenotype of WBS. Accordingly, we have searched for objective measurements of anatomical and physiological impairment of the neuronal retina in this disorder by means of multifocal electrophysiology, which allows comparison of focal responses from the retina, and by imaging of retinal structures using OCT and confocal scanning laser retinal imaging. The integrity of outer neural retinal layers, as measured by OCT, was the best predictor of the low-level retinal phenotype (Figure 7). Further functional evidence for retinal impairment in neuroretinal retinal layers was provided by the observed pattern of elec-



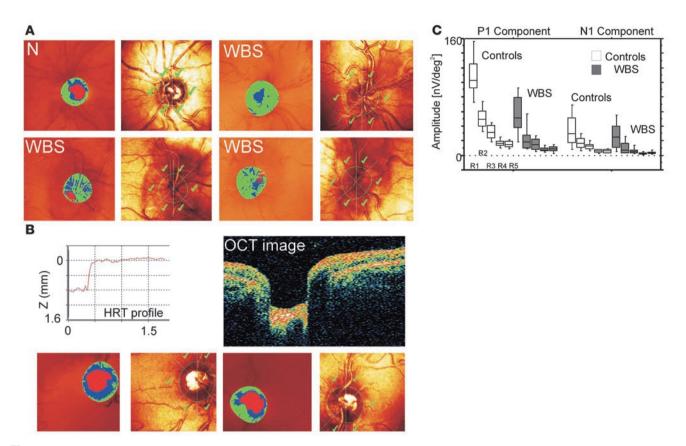


Figure 5

Confocal scanning ophthalmoscopy confirms anatomical abnormalities suggested by OCT results, and multifocal electrophysiology shows functional impairment. (A) The central deep segmented red region is often obliterated or reduced and displaced in WBS subjects as compared with normal subjects (N). The red area marks the cup. The rest of the disc area is divided into a sloping (blue) and stable (green) neuroretinal rim. Right-side images in each panel mark the optic nerve region in average confocal images. Loss of concavity is even more evident in these images. (WBS subjects: lower left, WBS3, right eye; lower right, WBS3, left eye; upper right, WBS1). (B) Abnormal deviation of the normal cup pattern by further deepening of that region in a WBS subject (WBS5), suggesting that the pattern of deviation is bimodal. The walls of the cup region are abnormally steep, as shown in the measurements in the HRT images (see top left, height profile) and confirmed by the OCT biopsy (top right; for comparison, see normal profile in Figure 4, bottom). Bottom row of images also suggest that the "neck" of the cup opening is abnormally large in this subject. (C) The P1 component of the multifocal first order kernel response was most significantly affected in WBS, as illustrated by percentile plots. Box boundaries correspond to upper and lower twenty-fifth percentiles, outer bars to the tenth percentiles, and middle bar to the median. The earlier N1 component is less significantly affected. R1–R5, rings of increasing eccentricity.

trophysiological responses, which showed local abnormalities in N1 and P1 wave components (22, 23).

Interestingly, we found evidence that this phenotype is independent of additional visual cortical–dorsal stream deficits, given that correlation of low-level magnocellular impairment with high-level motion coherence measurements (requiring motion integration of 2D and 3D dot-defined objects immersed in noise) was absent. In the present study, we directly addressed this issue by establishing structure-function correlations.

The unsuspected (in spite of the gene expression data) neural phenotype at the level of the retina may have escaped rigorous scrutiny given the fact that most WBS subjects have relatively normal VA, which was also the case in our sample. Our suspicion of a possible neuroretinal phenotype was also raised by our own previous work, which suggested separable low-level magnocellular deficits in peripheral vision and high-level structure from motion deficits (3). Our correlation analysis confirmed that the latter deficits can be explained by the previously described visual cortical-dorsal stream abnormalities (6) whereas peripheral low-level

deficits can be correlated with objective functional and structural impairment in the retina.

Optic coherence tomography and confocal scanning laser retinal imaging data were also critical in revealing abnormal optic nerve development. In this regard, we found 2 distinct morphogenetic trajectories. It remains to be clarified which developmental processes are critical in deciding between these trajectories. Interestingly, the same pattern was often observed to be similar across eyes within the same subject, suggesting that this bimodal decision process is constrained by a given genetic background. Most importantly, these abnormalities were accompanied by significant ganglion cell RNFL loss.

In spite of the previously discovered role for *GTF2IRD1*, *GTF2I*, and *LIMK1* as potential genetic determinants of mammalian craniofacial and neural development, including development of the retina, it is clear that mechanisms of cumulative dosage effects of these genes will vary across and within neural tissues. Other factors, such as parental origin of the hemideletion or the precise location of the breakpoints, did not seem to play a major



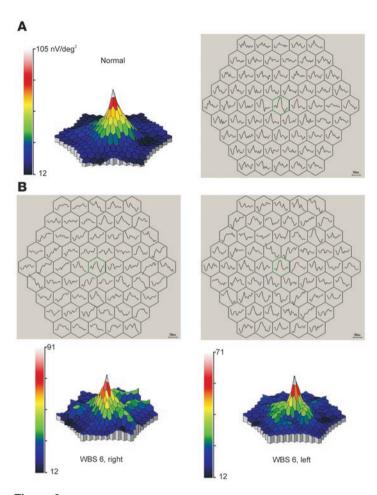


Figure 6

Objective functional evidence for retinal impairment in WBS. Multifocal electrophysiological maps (average curves extracted by means of reverse correlation depicted for contiguous stimulus hexagons and respective color-coded amplitude maps) in a normal control subject (A) and the 2 retinas of a WBS subject (B, WBS6). The central peak is preserved, but many areas are clearly impaired, with a large heterogeneity within and across eyes in WBS. 3D Plots are normalized for area of stimulation. For each stimulus hexagon, the peak amplitude, defined as the difference between N1 and P1 (peak negative and positive deflections, marked as cyan and yellow in the high-resolution image version), was calculated. Hexagon with green outline represents the central fixation region.

explanatory role in the observed phenotype, which is consistent with the similarity of deletion size. The variations in focal physiological impairment that we have found may well reflect regional variations in gene dosage (15, 24).

In conclusion, the structural and functional data of this multimodal study show clear evidence of what we believe is a novel visual phenotype at the level of the retina. The fact that patterns of impairment are dissociable from cortical–dorsal stream function (Figure 7) shows that the discovery of a novel mechanism of disease can also bring new insights into brain function. Indeed, our findings seem to seriously challenge theories that explain global coherence deficits based on magnocellular impairment (e.g., dyslexia) (25–29).

Future imaging and electrophysiological studies of individuals with partial deletions (11–13, 15, 30) would be of special interest in separating the role of the genes involved in different aspects of

visual processing. It is likely that some but not all higher-level cortical processes may overcome peripheral impairment by taking advantage of functional redundancy (3, 9). Finally, our work poses constraints on theoretical models explaining the necessity/sufficiency of cortical and/or retinal abnormalities in explaining the global visual phenotype in this and other developmental conditions.

In summary, our study provides a framework to explain low-level (magnocellular) visual deficits in genetic neurodevelopmental conditions and their relation to defective cortical (dorsal stream) mechanisms.

Methods

Clinical assessment

Participants. The patient group comprised 13 genetically confirmed WBS subjects, 6 females and 7 males, with a chronological mean age of 18.62 ± 6.38 SD, ranging from 10 to 31 years.

Ophthalmologic examination and assessment of visual function. A complete eye examination was performed on all WBS individuals. This exam included best-corrected VA (Snellen optotypes), stereopsis evaluation using Randot, intraocular pressure measurement (Goldmann applanation tonometer), slit lamp examination of anterior chamber structures, fundus examination, and photography after pupil dilation.

VA was relatively well preserved (8.9/10 \pm 1.5 SD), and ocular pressure ranged also within normal values (13.3 \pm 0.99 SD). Refractive errors ranged from moderate myopia to moderate hyperopia (average spherical correction: 0.56 \pm 2.69) with variable astigmatism but with no common pattern among our study group. Of 13 patients, 4 (~30%) had measurable strabismus (1 case of alternating esotropia and 3 cases of exotropia). The 9 remaining patients had preserved binocular function, and in 3 of these we could actually demonstrate a high-depth discrimination score of 20 seconds of arc using the Randot Stereo test. Overall, the cases with remaining stereopsis had relatively good scores (mean depth discrimination score: 70 arc s \pm 20.6 SD). Eye movement examination was unremarkable in all cases.

All but 2 patients presented the characteristic stellate pattern of the anterior iris leaflet. In the remaining 2 WBS subjects (4 eyes), we observed a pattern of sectorial atrophy of the anterior iris leaflet, mostly peripheral, which was very consistent across eyes, within subjects.

All control subjects were volunteers without evidence of either optic disc or retinal disease and with normal or corrected-to-normal VA. All were not previously exposed to the tests performed. Normal subjects were individuals with no visual field defect, no structural optic disc abnormalities, and no history of intraocular pressure greater than 21 mmHg. Subjects were excluded if they had a history of any retinal disease, surgery or laser procedures, optic disc anomalies such as coloboma or optic disc drusen, or any neurological diseases associated with visual field defects.

The control group for OCT and HRT analysis was composed of n=22 eyes of normal subjects who had undergone an eye examination at our center. Concerning the electrophysiological recordings, we used control recordings from 20 normal age-matched eyes. Chronological age-matching is the most adequate for the correct analysis of the obtained objective measurements. These measurements are critically sensitive to the level of maturation of retinal pathways.

Psychophysical assessment of visual function. Psychophysical assessment of visual function was performed using procedures described in detail previously in the context of psychophysical characterization of WBS visual phenotypes



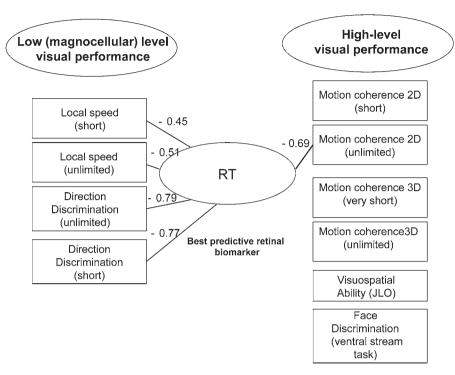


Figure 7

Retinal measurements can predict low-level motion performance deficits but not high-level motion integration and dorsal stream function measurements. Correlations with high-level ventral stream (face discrimination) performance are, as expected, also absent. Connecting lines depict significant correlations. Rho values next to lines depict correlation strength (they are in general negative because lower RTs correspond to less neurons in the outer retina, and lower performance implies higher thresholds). JLO, Benton's Judgment of Line Orientation.

(in addition to the classical dissociation of performance concerning Benton's Facial Recognition Test and Benton's Judgement of Line Orientation Test) in relation to mental and chronological age–matched controls (3). In brief, we applied CS strategies. The stimuli used were patches of 0.25 cycle/deg of vertically oriented sinusoidal gratings, undergoing 25 Hz counterphase flicker. Standard voltage–luminance curves were measured for each phosphor with software and hardware (including a Minolta colorimeter) provided by CRS, which ensured γ correction. Mean background luminance was 61.7 candela (cd)/m². Subject reliability scores were as described previously, with only 2.3% fixation errors. All participants did CS perimetry under monocular conditions, and the first tested eye was chosen in a random manner.

Concerning motion stimuli, we used random dot kinematograms presented within a circular spatial window of 6-degree visual angle (except when otherwise stated) generated using Vision-Works for Windows (Vision Research Graphics) in a Trinitron GDM-F520 monitor. Viewing distance was 56 cm. Pixel size was 0.056 deg² and dot size was 3×2 pixels. Dot density was 3 dots/deg². For all tests, the background luminance was approximately 0 cd/m², and 1 pixel had approximately 18 cd/m². In all experiments, a 2-alternative forced-choice staircase method (temporal or spatial) was used (with 12 reversals, 6 practice and 6 experimental). Steps were 0.01 log units in size unless otherwise stated. Durations of fixed stimulus presentations were of 1.133 s or 200 ms, in the case of motion coherence of spherical surfaces task, after which a gray background appeared and was present until the subject responded and the next trial commenced.

Visual thresholds were measured for the following tasks: (a) 2D motion coherence: 100% noise dots alternated randomly within 1 aperture with coherent motion in a particular direction that varied pseudorandomly from

trial to trial. Dots moved at 3 deg/s. Subjects had to report the presence or absence of coherent motion. (b) Direction discrimination: the stimulus was presented within a single circular aperture in the middle of the screen and consisted of dot-defined surfaces moving coherently in a horizontal manner to the right (0 deg) or vertically downwards (270 deg) at 2.5 deg/s. Subjects had to indicate whether dots were moving horizontally (0 deg.). Staircase implementation was such that correct answers caused the vertical stimulus to change its direction of motion to approximate the horizontally moving stimulus (from 270 toward 0 degrees). Step size was 0.02 log units. (c) Local speed discrimination (2 distinct tasks, requiring spatial 2 alternative forced choice (2AFC), either with 1 second or unrestricted temporal presentation): 2 windows were positioned with 6 degrees horizontal separation. Dots moved horizontally, to the right. The initial speed of the test stimulus was 50 deg/s, and the standard stimulus was moving at 15 deg/s. Subjects had to indicate which aperture contained the fastest moving dots. (d) 3D Motion coherence: the stimulus consisted of dots placed on the surface of a rotating sphere 3 degrees in diameter revolving around an axis, whose angle varied in a pseudorandom way. Speed of revolution was 20 rpm. The sphere alternated within 1 aperture with a

stimulus consisting of 100% noise dots that moved at 2 deg/s. Subjects had to report the presence or absence of a rotating sphere. All participants gave their informed consent, and the study followed the guidelines of the ethics committee of the Faculty of Medicine of Coimbra.

Molecular characterization of deletions. The diagnosis of WBS was first confirmed by FISH. Hemizygosity for the critical region of 7q11.23 involving the elastin locus was tested using a commercially available probe specific for the WBS critical region (WBSCR) (Qbiogene Inc.).

PCR analyses of single- and multiple-copy microsatellites were performed in order to define the size and parental origin of the deletion. Genotyping of paralogous sequence variants was done thereafter to map the exact breakpoints of the deletion within the specific block of segmental duplications at the WBS locus. The locations of all markers used in these analyses are shown in Supplemental Figure 1. PCR conditions and oligonucleotides used for the analyses were as previously described in detail (19, 31). In brief, site differences corresponding to single-nucleotide changes in each amplicon were detected using restriction enzymes (New England Biolabs), size fractionation on 1%–3% agarose gels, and relative quantification of digital images with the Quantity One software package (Bio-Rad).

Electrophysiological recordings. We recorded mfERGs using Dawson-Trick-Litzkow fiber electrodes after pupil dilation (for details see below) prior to fundus photography with a RETIscan System (Roland Consult) (32). The stimulus used in the mfERG consisted of 61 hexagons, covering in total a visual eccentricity of up to 30 degrees, and were presented on a 20-inch monitor at a viewing distance of 33 cm. Maximum luminance was approximately 120 cdm⁻². The hexagon areas increased with eccentricity in order to compensate for local differences in signal amplitude due to differences

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in cone density across the retina (leading to a 4-fold change in hexagon area size). Each hexagon was temporally modulated between light and dark according to a binary m-sequence (frame rate: 60 Hz).

Observers fixated a small cross in the center of the stimulus during recording sessions of approximately 8 seconds. Fixation was continuously checked by means of online video monitoring. To improve fixation stability, the sessions were broken into 47-second segments with brief rest periods between segments. Eight trials were recorded for each mfERG session, and the average recording time for each trial was 47 seconds. Signals were amplified with a gain of 100,000 and band pass filtered (5–300 Hz).

The reference and ground electrodes were attached to the ipsilateral outer canthus and forehead, respectively. The surface electrode impedance was less than 10 k ohm. First-order kernels were used for mfERG evaluation. For each hexagon, the peak amplitude, defined as the difference between N1 and P1, was calculated and the implicit time of P1 determined.

Multimodal imaging of retinal structures

Subject preparation. Each eye of each subject was dilated with 1% tropicamide prior to image acquisition. OCT, HRT, and fundus photography were then performed on both eyes. Ophthalmic photographers who were familiar with and trained in the use of the OCT/HRT systems obtained all scans.

Standard operational procedures for fundus photography followed the guidelines of the Fundus Photograph Reading Center (Department of Ophthalmology and Visual Sciences, University of Wisconsin, Madison, Wisconsin, USA).

OCT. OCT is a biomedical optical imaging technique that can provide noninvasive, high-resolution cross-sectional images of ocular microstructures in situ and in real time (33, 34). Low-coherence near-infrared light (820 nm wavelength and 20 nm bandwidth) from a superluminescent diode source is coupled into a fiber-optic Michelson interferometer and split into reference and sample paths: one directed at the retina and one to a reference mirror. Light retroreflected from the variable distance reference mirror is recombined with light backscattered from the eye (35). An A-scan, an axial profile of optical reflectivity versus distance, is obtained by moving the reference mirror while recording the magnitude of the resulting interference signal. This process can be repeated many times while scanning the beam through the eye fundus, thus allowing one to obtain an optical cross-section image, a B-scan (36, 37).

We used an OCT device (Stratus OCT; Carl Zeiss Meditec) to obtain real-time, objective, cross-sectional measurements of various layers of the retina, including the RNFL. Stratus OCT A-scans consisted of 1024 data points for 2-mm depth while B-scans were composed of 128, 256, or 512 adjacent A-scans, therefore generating cross-sectional images of retinal anatomy (optical biopsies) with axial resolution less than or equal to 10 μm , transversal resolution of 20 μm , and 2 mm of longitudinal scan range.

Using the Stratus OCT Fast Macular Protocol, 6 radial 6-mm (length) line scans, 30 degrees apart, and of 128 A-scans each were scanned in 1.92 seconds. Two data sets were obtained from the eye fundus at 2 different locations corresponding to 2 different acquisitions: one centered in the macula and one centered in the optic disc.

From the single acquisition centered in the macula, 2 different measurements can be obtained, the RNFL thickness information and the RT information, thus corresponding to 2 different thickness maps. This is achieved by segmenting the retina from other layers with an algorithm detecting the edge of the retinal pigment epithelium and the photoreceptor layer.

Macular RT was calculated by computing the distance between the signal from the vitreoretinal interface and the signal from the anterior boundary of the retinal pigment epithelium while the RNFL in the macular and peripapillary region was determined by obtaining the distance between the vitreoretinal interface and its adjacent highly reflective layer, with the

posterior border determined based on threshold reflectivities that are a function of the local maximum reflectance adjusted to variations in optical alignment. The RNFL, visible as a scattering layer originating from the optic disc, became thinner as it approached the fovea. An interpolation algorithm was used to correct for any missing boundaries caused by shadowing due to blood vessels or exudates (38).

Both the RNFL and RT are presented as a 9-region thickness map, showing the interpolated thickness for each area, with a central circle of 500 μm radius and 2 outer circles with radii of 1500 μm and 3000 μm . The interpolated thickness is displayed using a false color scale, in which bright colors (red and white) correspond to thickened areas and darker colors (blue and black) are assigned to thinner areas.

In the evaluation protocol of the optic nerve head, the software automatically identified the anterior and innermost borders of the second echo-rich structure (e.g., the retinal pigment epithelium/choriocapillaries layer) at each side of the optic disc. The detected position of the end of the retinal pigment epithelium/choriocapillaries reflection was corrected manually when necessary.

To improve OCT thickness mapping accuracy, we used proprietary software to correct for the location of fovea (brought into the center of the thickness map) as well as to perform interscan registration, thus correcting for saccades. This was achieved by registering all the 6 radial line scans from the Fast Macular Protocol into an atlas of the human retina (RT-Atlas) built using RT analyzer II (RTA; Talia Technology) to scan a set of 32 healthy volunteers. Principal component analysis was used to capture RT shape and define RT-Atlas, i.e., the one associated with the major spread of data. As for OCT registration, scans were brought into the RT-Atlas space and registered by estimating a total of 24 parameters defining the global position and rotation as well as individual scan fit.

Confocal scanning laser imaging. We used an HRT II device (version 2.01, Heidelberg Engineering), which is a confocal scanning laser ophthalmoscope that provides topographical images derived from optical sections at consecutive equidistant focal depth planes, yielding equidistant 2D optical section images (with 256×256 image resolution) in a 15-degree field of view, enabling the construction of a topographic image. To acquire the digital confocal images, a laser beam with a wavelength of 670 nm was focused in the retina and periodically deflected by oscillating mirrors. The amount of reflected light at each point was measured using a light-sensitive detector.

Two 115-degree field-of-view scans judged to be of acceptable quality were obtained for each eye. Images with significant movement artifacts were rejected. A mean topographic image of these scans was created using HRT software, version 2.01. Mean image topography was generated, and a trained technician outlined the optic disc margin using a mouse-drawn contour line on the mean topographic image while viewing stereoscopic photographs of the optic disc.

All initial analysis on the HRT was performed using HRT software, version 2.01. Typical optic disc parameters measured with HRT included rim area (area of neuroretinal rim located above the reference plane), mean cup depth, maximum cup depth, height variation in contour, mean height contour, cup shape, disc area, cup area, cup/disc area ratio, cup volume below surface, rim volume above reference plane, rim/disc ratio, RNFL thickness, and RNFL cross section (21, 39, 40).

Statistics. Distribution-free nonparametric statistics (Mann-Whitney U tests, Spearman's rho correlations) were carried out for all tasks to avoid biases due to deviations from normality or lack of data homoscedascity. Each subject's score was entered as a data point in correlation analyses for binocular tests (when more than one measure was available, an average score was used, yielding up to 13 data points). Concerning monocular tests, data from each eye were entered as data points (yielding up to 26 data points,



although typically 23/24 because cases in which performance fell to chance levels were excluded). Heterogeneity in performance between the 2 groups (normal and WBS) was analyzed by the F test for homogeneity of variance. All statistical analyses were performed with STATVIEW (SAS) or SPSS (version 12 for Windows; SPSS LEADS Technologies Inc.) software packages.

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