Agenesis of the Corpus Callosum: Genetic, Developmental and Functional Aspects of Connectivity

Lynn K. Paul^{1,2}, Warren S. Brown², Ralph Adolphs¹, J. Michael Tyszka¹, Linda J. Richards³, Pratik Mukherjee⁴, and Elliott H. Sherr⁴.

¹ California Institute of Technology
 ² Travis Research Institute
 ³University of Queensland
 and ⁴University of California, San Francisco

Address correspondence to:

Lynn K Paul HSS 228-77 Caltech, Pasadena, CA 91125

Website: http://www.emotion.caltech.edu/AgCC

And to:

Elliott H. Sherr Department of Neurology, UCSF 350 Parnassus Ave, Suite 609 San Francisco, CA 94143-0137

Website: http://www.ucsf.edu/brain/callosum/callosum.htm

Agenesis of the corpus callosum (AgCC), a failure to develop the large bundle of fibers that connect the cerebral hemispheres, occurs in 1:4000 individuals. Genetics, animal models and detailed structural neuroimaging are now providing insights into the developmental and molecular bases of AgCC. Neuropsychological, EEG, and fMRI approaches are examining the resulting impairments in emotional and social functioning, and have begun to explore the functional neuroanatomy underlying impaired higher-order cognition. The study of AgCC could provide insight into the integrated cerebral functioning of healthy brains, and may offer a model for understanding certain psychiatric illnesses, such as schizophrenia and autism.

Introduction

The brain's complexity arises from its connectivity – this is highlighted by the disproportionate increase in white matter volume through primate evolution ¹. The largest connective structure in the brain is the corpus callosum, consisting of over 190 million axons that transfer information between the two cerebral hemispheres (**Figure 1**) ². The corpus callosum contains homotopic and heterotopic interhemispheric connections. Although there has been debate about whether the connections are primarily excitatory (integrating information across hemispheres) or inhibitory (allowing the hemispheres to inhibit each other to maximize independent function) ³ it appears to be primarily excitatory and this is the focus of most research on interhemispheric transfer (IHT).

Corpus callosum function in humans was first investigated in classic studies of "split brain" patients, whose callosum is severed surgically for the treatment of epilepsy (**Box 1**) ^{4,5}. However, there is another population that provides valuable insight about the functions of the corpus callosum and the role of altered connectivity in neurodevelopmental disorders: individuals with developmental absence (agenesis) of the corpus callosum (AgCC) (**Box 2**).

AgCC encompasses complete absence as well as hypogenesis (partial absence) of the corpus callosum (**Box 3**). This review covers a broad range of findings from research into AgCC, including animal models of callosal development, genetic and environmental contributions to AgCC, neuroimaging in acallosal humans, and neuropsychological outcomes in individuals with Primary AgCC. Therefore, the interdisciplinary nature of this review provides a framework for bridging once largely non-overlapping domains of neuroscience: genetics and neuropsychology.

AgCC is a complex condition, which may result from disruption in any one of the multiple steps of callosal development, such as cellular proliferation and migration, axonal growth, or glial patterning at the midline. We review the molecular mechanisms underlying these processes. Later sections address behavioral and neuropsychological aspects of AgCC. We briefly examine research on IHT and alternative hypotheses regarding behavioral symptoms. Although the contribution of AgCC to our understanding of callosal function is complicated by concomitant anatomic changes (Box 2), we suggest that AgCC may be a powerful model for understanding cortico-cortical plasticity in other neurological and psychiatric populations.

Development of the Corpus Callosum

Corpus callosum formation involves multiple steps, including correct midline patterning, formation of telencephalic hemispheres, birth and specification of commissural neurons and axon guidance across the midline to their final target in the contralateral hemisphere. Much of

what we know about the stages of callosal development comes from animal models ^{6,7}. Several principal mechanisms have been hypothesized to regulate callosal formation.

Guidance by pre-exisiting axon tracts

The first axons to cross the midline arise from neurons in the cingulate cortex. In mice, these pioneer axons cross the rostral midline at embryonic day (E)15.5 8, providing a path for the fasciculation of later-arriving neocortical axons. In humans, pioneer axons express the guidance receptor Neuropilin -1 9, which may guide these axons themselves or the later-arriving callosal neurons from the neocortex. Cingulate cortex neurons also project axons into the rostrolateral cortex, perhaps to guide neocortical axons toward the midline initially. In more caudal regions of the corpus callosum, the hippocampal commissure, which in mice is formred a day earlier than the corpus callosum, may provide a growth substrate 10, 11.

Midline glial structures

Multiple glial structures including the glial wedge, midline zipper glia, and indusium griseum (**Figure 3**), are present at the developing midline and are likely required for corpus callosum formation ¹²⁻¹⁵. The glial wedge is a bilaterally symmetrical structure composed of radial glial cells that reside ventral to the corpus callosum at the corticoseptal boundary. It prevents callosal axons from entering the ventrally located septum, and once callosal axons have crossed the midline, repels the axons away from the midline into the contralateral hemisphere ^{13, 16}. Guidance by the glial wedge occurs through both Slit-Robo and Wnt-Ryk signaling ^{13, 16-18}. Midline zipper glia are found ventral to the developing corpus callosum at the septal midline. Their fusion at the midline has been hypothesized to be necessary for subsequent callosal axons to grow across the midline ¹⁹.

The indusium griseum glia (IGG), which are dorsal to the developing corpus callosum, also express Slit-2 and may help in guiding commissural axons toward their site of midline crossing ¹³. Recent work in conditional FGFR1/GFAP Cre knockout mice has shown the importance of these glia in corpus callosum formation ¹⁴. When FGFR1 is selectively eliminated from glia (and not neurons), the corpus callosum fails to form. Further analysis showed that FGFR1 is required for the migration of the IGG and selectively for the development of this midline glial structure ¹⁴. However, when FGFR1 is knocked out earlier in development, all midline glial structures at the corticoseptal boundary fail to develop, suggesting that FGFR1 plays a signaling role at multiple stages of callosal development ¹⁵. Similarly, Nfia and Nfib knockout mice ^{20, 21}, whose IGG and glial wedge are absent or significantly reduced in size, do not form a corpus callosum. However, midline glia are not the only guidance mechanisms at the midline. FGFR1 heterozygote (KO/WT) ¹⁵ and GAP43 knockout mice ²² do form midline glial structures (and express Slit 2), but callosal axons fail to cross, suggesting that multiple mechanisms regulate callosal development.

The subcallosal sling lies dorsal to the glial wedge and ventral to the developing corpus callosum. When the sling is severed experimentally, the corpus callosum fails to form ¹⁹, suggesting a role for this structure in callosal guidance, although the first callosal axons cross prior to the formation of the subcallosal sling. In mice, the majority of cells that make up the sling prenatally are neurons ²³, but in humans, whose subcallosal sling contains a large number of glia, the cellular origins of this structure are more complex ⁹ (Figure 3). Finally, additional neurons that have been identified within the corpus callosum ²⁴ may play a role in axon guidance.

Target recognition and selective pruning in the contralateral hemisphere

After crossing the midline, callosal axons grow into the contralateral hemisphere toward their designated target region, usually homotopic to their region of origin, and then innervate the appropriate cortical layer. Such processes probably involve both molecular recognition of the appropriate target region and activity-dependent mechanisms that regulate axon targeting to the correct layer and the subsequent refinement of the projection ²⁵. In cats and ferrets, refinement of callosal visual projections occurs through the selective pruning of axons after eye opening ²⁵⁻²⁷. Correct pruning and stabilization at the border of areas 17 and 18 (but to a lesser extent in other areas) requires visual input ²⁸. A similar refinement of developmentally exuberant projections occurs in the somatosensory cortex ²⁵. It is not yet clear whether defects in axonal pruning may affect corpus callosum size and contribute to callosal hypoplasia (See Box 3) in humans.

Animal Models of AgCC

Animal models of AgCC provide a basis for identifying genes that may be involved in human AgCC. The inactivation of genes that cause AgCC in mice often also trigger neurological deficits in other large fiber-tracts such as the internal capsule, and consequently leads to death at birth in many cases. These phenotypes, mostly resulting from gene deletions, may be too severe to model human AgCC, and such gene deletions may also result in embryonic or perinatal death in humans. However, a number of mouse models exist in which AgCC is partially or fully penetrant, but the animals have normal lifespans. Strains such as 129 and BTBR have been used to map quantitative trait loci that affect corpus callosum size ²⁹. Recent studies have shown that the gene DISC1 is homozygously inactivated in all 129 strains ³⁰ and this genetic mutation may be causally linked to AgCC in these mice. Interestingly, this gene has also been implicated in schizophrenia, and may be an important mechanistic link between the two disorders ³¹.

Finally, recent studies have used novel tools for labeling and isolating functional subsets of neurons to identify markers that are unique to callosal projection neurons ³². These studies have identified the gene Lmo4 as a candidate transcription factor for specifying callosal "identity" to projection neurons. This approach will potentially lead to a greater understanding of how neurons acquire their functional identities.

Causes of AgCC in humans

Genetic causes

The genetics of AgCC in humans are quite variable and reflect the underlying complexity of callosal development. Current evidence suggests that a combination of genetic mechanisms, including single-gene Mendelian, single-gene sporadic mutations and complex genetics (which may have a mixture of inherited and sporadic mutations) may have a role in the aetiology of AgCC. Retrospective chart reviews and cross-sectional cohort studies report that 30-45% of individuals with AgCC have identifiable causes for AgCC. Approximately 10% have chromosomal anomalies and the remaining 20-35% have recognizable genetic syndromes (Table 1) ³³. However, if we only consider individuals with complete AgCC, then the percentage of patients with identifiable syndromes drops to 10-15%, and thus 75% of patients with complete AgCC do not have an identified cause.

One example of AgCC associated with a Mendelian disorder is X-linked lissencephaly with AgCC and ambiguous genitalia (XLAG), which results from a mutation in the Aristaless related homeobox gene (ARX). The first description of this disorder included only male patients. However, females with mutations in ARX, who, because of X-inactivation, have clinical symptoms that range from normal to those with spasticity, mental retardation and seizures. MRI scans of these female patients are either normal or show Isolated AgCC with Probst bundles (Box 3) ³⁴. Male ARX knockout mice also have AgCC and mimic many of the other clinical and anatomic findings in XLAG ³⁵, including a significant reduction in cortical interneurons, which likely explains the severe and uncontrollable seizures in this condition ³⁶.

Another syndrome caused by a single-gene mutation with considerable overlap between the human and animal phenotype is CRASH syndrome (corpus callosum agenesis, retardation, adducted thumbs, spastic paraplegia and hydrocephalus), which is accompanied by diminutive cortico-spinal tracts within the brainstem. CRASH is caused by mutations in the L1CAM gene that codes for a transmembrane cell adhesion protein broadly expressed in the central nervous system. Mice with L1CAM gene inactivation show complete or partial AgCC, hydrocephalus, small corticospinal tracts, reduced neuron numbers, and additional abnormalities in the elaboration of apical dendrites from cortical pyramidal neurons ³⁷. Recent work suggests that inhibition of homophilic binding of the L1CAM protein can cause hydrocephalus, but that preventing corpus callosum formation also requires disruption of heterophilic interactions with other proteins including integrins ³⁸. Gene dosage effects have also been observed in mouse knockout models for the genes DCC and GAP-43. Here, heterozygote mice show partial AgCC whereas homozygote knockout mice have complete AgCC with additional anomalies ²².

Andermann syndrome, an autosomal recessive condition prevalent in the Saguenay-Lac-St-Jean region of Quebec, presents with callosal hypoplasia or AgCC, cognitive impairment, episodes of psychosis and a progressive central and peripheral neuropathy. It is caused by mutation of the potassium-chloride cotransporter, KCC3 ³⁹. KCC3 knockout mice display neurodegeneration, and also have hearing loss and progressive neuropathy ⁴⁰. However, in contrast to ARX and L1 mouse mutants, they have normally formed corpora callosa. Interestingly, some patients with KCC3 mutations also have a fully formed corpus callosum, and there is even phenotypic variability within families, suggesting that additional genetic influences are at play.

The variable effects of gene inactivation on callosal development in mice and humans is also evident in Meckel Gruber syndrome. In humans, mutation of Meckelin (the gene in MKS3) causes occipital encephaloceles, hepatic ductal cysts and polydactyly. In mice, mutation of the same gene causes AgCC, hydrocephalus and cysts within the kidney ⁴¹. Table 1 shows other disorders associated with callosal agenesis that show a clear recessive pattern of inheritance but for which the causative gene has not been identified, (Table 1).

In spite of the progress of research into single-gene Mendelian causes of AgCC, most individuals with AgCC do not have a clearly inherited cause or a recognized genetic syndrome, suggesting that AgCC can be caused by sporadic (*de novo*) genetic events. One salient example of this is Mowat-Wilson syndrome (MWS), which, in addition to AgCC, causes Hirschsprung disease, congenital heart disease, genitourinary anomalies, microcephaly, epilepsy and severe cognitive impairment ⁴². MWS is caused by heterozygous inactivating mutations in the gene ZFHX1B on chromosome 2q22, which codes for SIP1 (Smad interacting protein 1) ⁴³. AgCC is not observed in all MWS cases, suggesting that haploinsufficiency or gene dosage of SIP1 interacts with other genetic polymorphisms to alter callosal development ⁴².

Aicardi syndrome is another disorder likely caused by sporadic mutations, in this case on the X chromosome. Only observed in females and XXY males with Klinefelter syndrome, this

disorder consists of AgCC, infantile spasms and chorioretinal lacunae. Patients with Aicardi syndrome show a constellation of additional cerebral and ophthalmologic abnormalities, so it is likely that the mutation participates widely and early in prosencephalic development. By inference, it is likely that other cases of AgCC are caused by haploinsufficiency at other genetic loci. This is exemplified by many reports of patients with AgCC who have sporadic chromosomal mutations, with particular loci identified repeatedly ⁴⁴. Recent data using microarray-based comparative genomic hybridization demonstrate that patients with AgCC have chromosomal deletions or duplications that are smaller than those that can be detected using conventional cytogenetics ⁴⁵. Indeed, in collaboration with the California Birth Defects Monitoring Program, we have demonstrated that the risk of having a child with AgCC is nearly three-fold higher for mothers aged 40 and above, consistent with causal sporadic chromosomal changes (E. Sherr, unpublished observations).

As noted for the single-gene disorders discussed above, not every patient displays AgCC, suggesting that many cases of AgCC are caused by polygenic and other complex interactions. Moreover, the abundance of case reports of AgCC associated with specific diseases (partially listed in Table I) probably also reflects complex underlying mechanisms. This is exemplified by a recent report examining the MRI findings of individuals with Sotos syndrome, which is caused by haploinsufficiency of the NSD1 gene. In this study, only one patient had complete AgCC and the rest (35/36) had either diffuse hypoplasia or thinning of the posterior body of the corpus callosum ⁴⁶. These findings suggest that some genes that are often referred to as modifier genes only partially contribute to callosal formation. Common disorders that affect behavior and for which the influence of many modifier genes is the likely mode of inheritance include autism and schizophrenia ^{47, 48}. Moreover, as there are many reports of AgCC or abnormally formed corpora callosa in patients with autism and schizophrenia ^{49, 50}, it is possible that the modifier genes that affect callosal development overlap significantly with those that cause these neuropsychiatric disorders.

Environmental factors

Finally, it is important to note that environmental factors may contribute to AgCC as well. While much less is known about these than the genetic factors we have reviewed above, one clear example of environmental influences on callosal development is provided by fetal alcohol syndrome (FAS). Both clinical and experimental evidence indicates that alcohol exposure *in utero* decreases gliogenesis ⁵¹ and glial-neuronal interactions ⁵², processes that are critical to corpus callosum development ⁵³. Additionally, a growing body of literature suggests that ethanol disrupts the transcription and biochemical function of L1CAM ⁵⁴⁻⁵⁶, implicating an interplay of environment and genetics in AgCC. The incidence of AgCC in FAS is approximately 6.8% ⁵⁷, with an even higher incidence of corpus callosum malformations short of complete AgCC. In many FAS cases, the corpus callosum is hypoplastic; this may result not only from disruption of early events in callosal formation, but also from later dysregulation of axonal pruning. Such mechanisms may also cause callosal hypoplasia in other disorders such as schizophrenia and autism ^{25, 31}. Other environmental factors may also influence post-natal and prenatal callosal development, including musical training ⁵⁸⁻⁶⁰, hypothyroidism ^{61, 62}, and enrichment or deprivation of experience ⁶³.

Behavioral Impairment in AgCC

Consistent with the broad range of genetic factors involved in AgCC, the cognitive, behavioral and neurological consequences of AgCC are wide-ranging. One approach to defining clinical subsets of the AgCC patient population is to categorize individuals according to specific

neuroanatomical findings, and subsequently relate these to the behavioral symptoms within these groups. For example, a number of studies have suggested that the presence of polymicrogyria, pachygyria (abnormally broad gyri), heterotopia, detected using MRI, correlates with moderate to severe developmental delay ^{64, 65}. However, creating accurate anatomical groupings is very difficult given the diversity of symptoms in patients with a particular anatomical abnormality.

The comparison between complete and partial AgCC has revealed conflicting data, with multiple studies showing no difference in behavioral and medical outcomes between the two conditions, whereas one earlier study reported a worse outcome for individuals with complete AgCC ⁶⁶. One hospital-based study reported that just under a third of AgCC patients were developmentally "normal" or only mildly delayed ⁶⁵. A longitudinal study of 17 children prenatally diagnosed with isolated AgCC, showed that nearly all patients had at least mild behavioral problems ⁶⁷. This suggests that even isolated AgCC when not ascertained clinically, still causes behavioural and cognitive impairment. Parents often report that when their child was diagnosed with AgCC, they were told that the outcome was unclear, ranging from severely delayed to "perfectly normal". However, as more individuals with Primary AgCC are identified and assessed with sensitive standardized neuropsychological measures, a pattern of deficits in higher-order cognition and social skills has become apparent even in the so-called "normal" individuals with AgCC.

Connectivity deficits

Historically, most research in subjects with AgCC focused on the consequences of callosal absence, with the expectation that AgCC patients would exhibit a "disconnection syndrome" similar to that seen in commissurotomy patients⁴. The classic disconnection syndrome (Box 1) involves complete lack of IHT and interhemispheric integration of sensory and motor information presented independently to each of the hemispheres ⁴, with surprisingly subtle behavioral consequences in everyday life.

Studies using tachistoscopic presentation of visual stimuli and studies of evoked potentials provide the most compelling information about functional connections and IHT in AgCC at various stages of sensory processing. Figure 4 illustrates the visual cortical disconnection effects in AgCC, as well as the limits of these disconnection symptoms. First, as illustrated by visual evoked potentials, there is a complete lack of IHT at the level of early visual processing in AgCC 68. The hemispheric disconnection of the primary visual system in AgCC patients results in a unique pattern of deficits in laboratory tasks that involve comparisons across the two visual fields: intact comparisons of simple stimuli and impaired comparisons of complex stimuli. First, despite the lack of transfer of early visual information, individuals with AgCC exhibit a normal ability to make comparisons of simple and easily encoded stimuli, implying intact interhemispheric transfer of simple or familiar information. For example, they exhibit an intact interhemispheric Stroop interference effect 69 and the typical bilateral field advantage for comparison of familiar and easily encoded visual information across hemifields ⁶⁸. These findings confirm that information can be transferred between the hemispheres in AgCC. One theory to explain the preserved capacity for IHT of simple stimuli in AgCC patients is that simple information can be transferred via other connecting pathways, such as the anterior commissure. Structural and functional exploration of these alternate pathways for IHT is a critical frontier in AgCC research.

In contrast, the capacity for IHT may be limited by task complexity. For example, the performance of patients with AgCC when comparing tachistoscopically presented visual information is significantly less accurate for more visually complex, less familiar, and less easily

verbalized stimuli ⁶⁸ (Figure 4). Similar limitations in IHT in AgCC patients were evident on other tasks that required transfer or integration of the products of more complex cognitive operations, required more rapid processing and relied on less prior experience ^{68, 70-72}. Taken together, these studies indicate that there is greater reliance on the corpus callosum for rapid and effective interhemispheric interactions as task requirements increase (e.g., when stimuli become more complex or response criteria become more constrained).

The question remains, however, about what causes the behavioral disturbances evident in Primary AgCC. Studies using dichotic listening ^{73,74}, PET ⁷⁵, and fMRI ⁷⁶ have revealed that language lateralization is normal in Primary AgCC patients and is in some cases even exaggerated. Although there is no published evidence for normal or abnormal localization of other higher cognitive functions in this population, we may hypothesize that if localized functioning in the cortex in Primary AgCC patients is normal, a lack of information transfer between localized processing regions in opposite hemispheres could contribute to behavioral difficulties. This leaves callosal disconnection as a viable explanation for the behavioural disturbances in Primary AgCC patients.

Neuropsychological impairment

The major anatomic feature of Primary AgCC is the absence of the corpus callosum, and it is presumed to be the cause of the cognitive and behavioral changes in Primary AgCC patients. However, colpocephaly (Box 3) and Probst bundles are common in people with Primary AgCC (and never without AgCC), and together with other more subtle anatomical changes, likely also affect behavior. Functional and anatomic imaging approaches, coupled with incisive neuropsychological assessments, may in the future be able to map the neural processes and neuropathology associated with AgCC onto the diverse behavioral anomalies. For now, we begin by describing the symptom profile.

Primary AgCC has surprisingly limited impact on general cognitive ability. Although the full-scale IQ may be lower than expected based on family history, scores frequently remain within the average range ⁷⁷. Interestingly, in an unexpectedly large number of persons with Primary AgCC (as many as 60%), performance IQ and verbal IQ are significantly different ^{77, 78}. However, there is no consistency with respect to which of the two is more affected. Impairments in abstract reasoning ^{79, 80}, problem solving ⁸¹⁻⁸³, generalization (the ability to extrapolate from one case to others) ⁸⁴, and category fluency (the ability to list multiple items that belong to a semantic category, e.g., names of animals)⁸⁰ have all been consistently observed in patients with Primary AgCC. Data from large sample sizes suggest that problem solving abilities become more impaired as task complexity increases (Brown and Paul, unpublished observations). While neuropsychological research into domains such as memory, attention, and spatial skills is under way in large samples of patients with Primary AgCC, currently published results in these domains are limited to a few case-studies that do not yet provide consistent findings.

The most comprehensively examined higher cognitive domain in AgCC patients is language. Overall, individuals with Primary AgCC have intact general naming (e.g. naming objects from line drawings ^{85,86}), receptive language (e.g. comprehension of sentences ^{78,86}), and lexical reading skills ⁸⁷. However, impairments have been reported in comprehension of syntax and linguistic pragmatics ^{88,89}, and in phonological processing and rhyming ^{86,88-90}. With respect to linguistic pragmatics, persons with Primary AgCC are impaired in comprehension of idioms, proverbs, vocal prosody ^{91,92}, and narrative humor ⁹³. Within humor, they exhibit difficulty in over-riding literal interpretation bias and are poor at using context to infer meaning ⁹²⁻⁹⁴. Primary AgCC patients also show marked difficulties with expressive language, for

example in the verbal expression of emotional experience, which is consistent with a diagnosis of alexithymia ⁹⁵. In a study of a large sample of AgCC patients with adequate expressive language skills, parents consistently described "meaningless" or "out-of-place" conversation as particularly common ⁹⁶. Interestingly, recent studies of language support the dynamic dual pathway model according to which syntax and semantics are lateralized to the left hemisphere and prosody to the right hemisphere ⁹⁷⁻¹⁰². In this model, the corpus callosum is the main path for coordination of this lateralized information, particularly for coordinating syntactic and prosodic information ⁹⁷⁻⁹⁹, the very areas of linguistic weakness evident in AgCC.

Parents of individuals with Primary AgCC consistently describe impaired social skills and poor personal insight as the features that interfere most prominently with the daily lives of their children ^{96, 103-105}. Specific traits include emotional immaturity, lack of introspection, impaired social competence, general deficits in social judgment and planning, and poor communication of emotions (e.g. prefer much younger friends, marked difficulty generating and sustaining conversation, take all conversation literally, do not take perspective of others, inability to effectively plan and execute daily activities such as homework, showering, or paying bills ^{96, 105}). Consequently, Primary AgCC patients often have impoverished and superficial relationships, suffer social isolation, and have interpersonal conflict both at home and at work due to misinterpretation of social cues.

Responses of adults with Primary AgCC on self-report measures also suggest diminished self-awareness ¹⁰⁴. The patients' self-reports are often in direct conflict with observations from friends and family. One potential factor contributing to poor self-awareness may be a more general impairment in comprehension and description of social situations. For instance, when asked to generate stories about social pictures, adults with Primary AgCC provided stories lacking in logic, narrative content, and social understanding ¹⁰⁶. It appeared that they had difficulty recognizing the implications of pictures depicting social scenes, imagining a sequence of events, and organizing relevant ideas in order to present an appropriate narrative. Similarly, when presented with highly provocative social pictures (e.g., photos of mutilations), adults with AgCC tended to under-estimate the emotional valence and intensity of the pictures, particularly for negatively valenced pictures ¹⁰⁷. Taken together, the neuropsychological findings in Primary AgCC highlight a pattern of deficits in problem solving and in social pragmatics of language and communication.

AgCC and neuropsychiatric disorders

The deficits in social communication and social interaction in Primary AgCC patients overlap with the diagnostic criteria for autism (DSM-IV)¹⁰³. Furthermore, people with Primary AgCC may display a variety of other social, attentional, and behavioral symptoms that may resemble those of certain psychiatric disorders. Psychiatric diagnoses are based on clusters of behaviors, which are very complex and likely involve multiple neural mechanisms ¹⁰⁸. Examination of symptom overlap between psychiatric disorders and AgCC may help isolate the symptoms that are directly caused by a dysfunction in cortico-cortical connectivity.

There are also structural similarities between AgCC and some psychiatric disorders. Indeed, structural correlates of abnormal brain connectivity are evident in essentially every psychiatric disorder that has been examined. For instance, several studies have found altered morphology of the corpus callosum in schizophrenia patients, including changes in size and shape, as well as microstructural changes in callosal regions that are reflected in diffusion MRI (dMRI) ³¹. There are also a number of reports of complete AgCC in schizophrenia patients ^{50,80,109}, underscoring a direct connection between AgCC and schizophrenia and countering claims that the smaller anatomic changes in the corpora callosa in schizophrenic patients are not causally related to the

condition. Corpus callosum size is also decreased in some cases of autism, especially its anterior sectors ¹¹⁰ ¹¹¹. Moreover, in one study, 8.5% of AgCC individuals had a diagnosis of autism, compared to only 1% of their siblings¹¹². Microstructural changes in the corpus callosum have also been found in patients with Tourette syndrome ¹¹³ and attention deficit hyperactivity disorder ¹¹⁴, ¹¹⁵. One recent study provides evidence linking genetic changes in KCC3 (the gene mutated in Andermann syndromed **this is defined in the genetics section**) with bipolar disorder ¹¹⁶, even though these patients did not have evident changes in callosal anatomy. As more causes of AgCC are identified, we anticipate further genetic correlations between AgCC and neuropsychiatric disorders will be found.

The functional consequences of structural changes in brain connectivity, which may be revealed by dMRI analysis¹¹⁷ including diffusion tensor imaging (DTI), contribute to cognitive impairment. Functional connectivity studies show that the strength of the correlations between brain activation in different regions and anatomical abnormalities is strikingly task-dependent. For example, children with attention deficit hyperactivity disorder show a disturbed transcallosally mediated motor inhibition ¹¹⁸. Functional connectivity studies in AgCC patients may reveal the means by which these highly atypical brains attain such apparently "typical" interhemispheric interaction. In turn, understanding the functional limits of such connectivity may contribute to knowledge about psychopathological conditions with apparent corpus callosum involvement.

AgCC, like many psychiatric disorders, but unlike callosotomy, results from abnormal *development* of connectivity. It may therefore be able to shed light on the behavioral and cognitive consequences of abnormal connectivity during development in general, as well as on potential compensation due to early intervention—a topic that is now receiving much interest, especially in studies of autism ¹¹⁹. Of course, most people with autism, schizophrenia or ADHD do not typically have gross absence of the corpus callosum. Nonetheless, insofar as AgCC models one specific component (namely, altered connectivity) of what is likely to contribute to the cognitive symptoms of these psychiatric diseases, *it may allow us to isolate a subset of symptoms that arise primarily from altered connectivity.* As disorders as complex as autism are not likely to have a single correct explanation ¹⁰⁸, finding clear genetic and neuroanatomical models that can dissect particular aspects of such disorders may be invaluable. Taken together, AgCC might be a powerful model for studying behavioural and cognitive aspects of a number of psychiatric disorders.

Integrating Findings Across Disciplines

We have emphasized the genetic and developmental nature of AgCC, and described its cognitive neuropsychology. How can data from these different domains best be synthesized? In linking genes and development to behavior and cognition, one approach is to postulate multiple intermediate traits or "endophenotypes", a compelling concept that has been developed to dissect the causes of complex psychiatric disorders ¹²⁰. One type of endophenotype category is the anatomy. We propose that in AgCC, the principal anatomic endophenotype is the absence of the corpus callosum. Regardless of the diverse genetic and developmental factors that result in AgCC, callosal absence in itself may directly lead to the behavioral and cognitive symptoms we have described in this review. However, in AgCC additional neuroanatomical factors such as Probst Bundles, colpocephaly, abnormal ipsilateral connections and abnormal cortical folding may well contribute separately to the clinical outcome of AgCC patients, functioning as independent endophenotypes within the AgCC clinical complex, as well as contributing to other disorders like schizophrenia and autism.

Thus, the endophenotype concept as applied to AgCC proposes that the abnormal neuroanatomy is generated by genetic and environmental factors operating on development, and that the neuroanatomy in turn generates the behavioral phenotype seen in AgCC. Such a picture offers intriguing possibilities for drawing parallels with psychiatric illness. Are there sets of genetic mutations or environmental factors that might contribute both to AgCC and to psychiatric illnesses such as schizophrenia and autism? Are there sets of cognitive and behavioral impairments that are common both AgCC and those psychiatric illnesses? Commonalities at either the genetic/environmental, anatomic or the behavioral level would provide preliminary support for hypotheses that callosal and other cortico-cortical white matter tract impairments may be central to these disorders. We thus suggest that geneticists, anatomists, cognitive neuroscientists, and psychiatrists need to collaborate closely in order to take full advantage of the insights that AgCC can offer to understanding psychiatric illness.

Conclusions and Future Directions

Research on AgCC holds great promise for multiple scientific disciplines. In the field of genetics, much needs to be learned about the mode of inheritance. Most current data point to sporadic and polygenic inheritance. Identification of additional fully penetrant genetic causes will provide important insight into callosal development and function. As these genetic causes are illuminated, understanding the range of behavioral phenotypes that correlate with the genetics may be particularly useful for informing family planning decisions, for understanding related psychiatric conditions and for developing early intervention strategies for children whose developmental trajectories may be more accurately predicted.

The biological basis of AgCC is complex; this is reflected by the large number of human congenital syndromes associated with AgCC. It is perhaps one of the most complicated neurological birth defects simply because so many developmental processes are involved in the final readout of a fully formed corpus callosum. It is this observation that makes AgCC a plausible model for many other neurological and psychiatric illnesses with neurodevelopmental components. Callosal development can be affected by defects in cellular proliferation and migration, axon growth and guidance, glial development and patterning at the midline. Understanding the basis of the many disorders associated with AgCC, such as schizophrenia and autism, requires not only the identification of genes that regulate each of these processes but also a deep understanding of the function of each of the genes and how they work together in separate and overlapping molecular pathways to produce a corpus callosum.

AgCC is also particularly interesting to those studying network plasticity and compensation, as it does not result in the classic disconnection syndrome seen following surgical disconnection in adulthood. Careful integration of imaging and electrophysiology methods may provide important information about the intra and inter-hemispheric connections in AgCC, similar to current work with split-brain patients ⁵. In turn, AgCC provides a powerful test-bed for the integration of methods such as dMRI, fMRI, and EEG to examine effective connectivity: given the demonstrable gross absence of specific structural connectivity, how does this translate to the functional deficits? Generating a functional map of AgCC brains will inform critical questions about cortical and subcortical re-organization: where are particular functional regions located (e.g., specific visual areas, areas involved in language)? To what extent do their locations differ from those in healthy brains? Are there some functional regions whose anatomical location remains relatively invariant, whereas others can shift location more

variably? Such questions have been much investigated in studies of plasticity in animal brains; next to nothing is known about this in the human brain.

Of course, the people most invested in AgCC research are the individuals and families who deal with this condition. Neuropsychological and behavioral characterization of AgCC may help clarify distinctions between it and various behavioral diagnoses (for example autism, Tourette syndrome, and ADHD). Methods from cognitive neuroscience will be the most fruitful route to understand the mechanisms underlying the cognitive and psychosocial characteristics that are common in Primary AgCC. In turn utilizing this information, clinicians can develop more nuanced interventions for key deficits in AgCC such as social skills, problem solving, and planning, with the goal of enhancing the everyday lives of these individuals.

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Box 1. AgCC and the classic "split-brain"

Surgical commissurotomies ("split-brain") are typically conducted in adulthood for the treatment of intractable epilepsy, while AgCC is a brain abnormality present at birth. In commissurotomy patients all cerebral commissures including the anterior commissure, are severed, whereas the anterior commissure is intact in almost all Primary AgCC patients. In callosotomy patients the anterior commissure is not surgically severed ⁵.

Individuals with commissurotomy manifest a "disconnection syndrome" that includes (1) absence of callosal transfer of sensory information (e.g. Refs ^{121, 122}), and (2) deficiency in bimanually coordinated motor activity (e.g. Ref ¹²³).

The pre-surgical existence of a seizure disorder complicates interpretation of higher cognitive functions in split-brain cases. However, Roger Sperry comments that "speech, verbal intelligence, calculation, motor coordination, verbal reasoning and recall, personality and temperament are all preserved to a surprising degree in the absence of hemispheric interconnection" ⁴. Nevertheless, deficits have been noted in cognitive processing time, arithmetic, abstract reasoning ⁴, and short-term memory ¹²⁴. Commissurotomized patients also may exhibit alexithymia ¹²⁵.

Overall, AgCC patients are have better, though limited, interhemispheric integration than commissurotomy patients on many forms of visual and tactile information (e.g., ^{68, 126}). The relative importance of age at onset of AgCC versus commissurotomy for interhemispheric transfer (IHT) is illustrated by the finding that patients with early callosotomy and children with AgCC show little evidence of a disconnection syndrome on IHT tests with simple tactile information, while adolescent and adult callosotomy patients show marked transfer deficits ¹²⁷. This suggests that neural plasticity in children may allow for reinforcement of alternative neural pathways and that presence of the anterior commissure alone may not be sufficient to explain residual IHT in AgCC. The extent to which this compensatory plasticity involves unique recruitment of anterior commissure fibers remains unclear.

Despite the difference in functional interhemispheric connectivity, commissurotomy and AgCC both result in impairments of reasoning in complex novel situations¹²⁸ Social situations require extremely rapid processing of very complex information that is typically handled within lateralized regions (i.e. lexical and affective processes) and therefore may be particularly sensitive to corpus callosum abnormality.

Box 2. Prevalence and Features of AgCC

Agenesis of the corpus callosum (AgCC) encompasses a broad range of diagnoses. A synthesis of recent neonatal and prenatal imaging studies suggested that AgCC occurs in at least 1:4,000 live births^{129, 130}, and other imaging studies ^{131, 132} demonstrated that 3-5% of individuals assessed for neurodevelopmental disorders have AgCC.

Complete and partial AgCC likely result from disruption of the early stages callosal development that could result from genetic, infectious, vascular, or toxic causes ^{65, 133-135}. Further heterogeneity in AgCC can arise from concomitant abnormalities in the anterior commissure. A recent study reported that the anterior commissure was small or absent in 60%, yet enlarged in 10% of AgCC cases ¹³⁶. The latter may provide insight into brain plasticity, as it has been hypothesized that interhemispheric connections in AgCC could be re-routed through the anterior commissure ^{68, 137}. This idea is indirectly supported by better clinical outcomes in individuals with a normal or large anterior commissure (E. Sherr, unpublished observations).

The contribution of AgCC to our understanding of callosal function is complicated by concomitant anatomic changes including *colpocephaly and Probst bundles*. It is possible that cognitive and behavioral differences between AgCC and split-brain patients arise from these other anatomic differences. Colpocephaly refers to the dilatation of the posterior aspect of the lateral ventricles, frequently including the temporal horns. This does not represent hydrocephalus ¹³⁸ but may represent reduction of ipsilateral cortical association tracts ¹³⁹. Probst bundles are the misrouted callosal axons that run parallel to the interhemispheric fissure and can also be observed in cases of partial AgCC. Apparently within the Probst bundles, a structure called the *sigmoid bundle* has been recently identified in several cases of partial AgCC (Figure 2, ¹⁴⁰). This long, heterotopic commissural tract appears to connect the left frontal lobe with the right occipitoparietal cortex.

Other brain malformations can also be associated with AgCC ¹³⁶. One AgCC autopsy study documented a lack of pyramidal tract decussation, suggesting a more global disorder of midline crossing ¹⁴¹. This pattern is also observed in many animal models of callosal agenesis ^{7, 142, 143}. All concomitant anatomical abnormalities, including changes in commissural fibers outside the corpus callosum, may be relevant to clinical outcome.

Box 3. Key Diagnostic Definitions in AgCC

Complete AgCC

A congenital condition characterized by total absence of the corpus callosum.

Hypogenesis of the Corpus Callosum

(Also known as Partial AgCC) A congenital condition characterized by partial absence of the corpus callosum. The absence must be evident from birth and not be representation of a degenerative condition.

Hypoplasia of the Corpus Callosum

Condition in which the corpus callosum is fully formed, but is thinner than expected for age and sex of the individual.

Isolated AgCC

Neuroanatomic description which includes complete absence of the corpus callosum, without other confounding brain abnormalities such as polymicrogyria, heterotopia or schizencephaly. Individuals with isolated AgCC frequently have colpocephaly and Probst Bundles.

Primary AgCC

Primary AgCC refers to a symptom profile which includes isolated AgCC and generally intact intellectual functioning as indicated by full-scale IQ >/= 80.

Anterior commissure

Small band of approximately 50,000 axons that connect the cerebral hemispheres. The anterior commissure connects the temporal lobes and is located at the base of the fornix.

Probst Bundles

Misrouted callosal axons that run parallel to the interhemispheric fissure and can also be observed in cases of partial AgCC.

Colpocephaly

Dilatation of the posterior aspect of the lateral ventricles, frequently including the temporal horns. This does not represent hydrocephalus but may represent reduction of ipsilateral cortical association tracts.

Sigmoid Bundle

A long heterotopic commissural tract found in some cases of partial AgCC. It appears to connect the left frontal lobe with the right occipitoparietal cortex.

Figure 1. Neuroanatomy of the corpus callosum

The human corpus callosum contains approximately 190 million axons. **a.** Organization of a human corpus callosum based on histological and neuroimaging findings. **b.** Diffusion magnetic resonance imaging (dMRI) and tractography modeling provide important provides imaging (dMRI) and tractography modeling provides provides imaging (dMRI) and tract

b. Diffusion magnetic resonance imaging (dMRI) and tractography modeling provide important information about the corpus callosum fiber tracts and the cortical regions they connect. These dMRI data¹⁴⁴ of transcallosal fiber tracts in normal brains resulted in a new organizational scheme that describes corpus callosum organization, and suggested that much more of the corpus callosum is involved in premotor and supplementary motor coordination than previously thought. Fibers are colored according to their projection areas: prefrontal lobe (green), premotor and supplementary motor areas (light blue), primary motor areas (dark blue), primary sensory cortex (red), parietal lobe (orange), occipital lobe (yellow), temporal love (violet). **c.** In monkeys, researchers have been able to use chemical tracers to map the organization of cortical fibers passing through the corpus callosum, providing a level of detail currently unavailable in humans.

Figure 2. Examples of Neuroanatomical Findings in AgCC

Neuroanatomic features of callosal agenesis and callosal hypogenesis revealed by magnetic resonance imaging (MRI) and diffusion tensor imaging (dMRI). Structural T1-weighted MRI (top 3 rows) and directionally-encoded color anisotropy dMRI (bottom row) are shown from a normal young adult male volunteer (left column), a young adult male with callosal agenesis (middle column), and a young adult male with callosal hypogenesis (right column). The dMRI images encode fiber orientation in white matter tracts using a 3-color scheme such that fiber pathways with predominantly left-right orientation are displayed as red, anteroposterior orientation as green, and craniocaudal orientation as blue. PB, Probst bundle; CB, cingulum bundle; AC, anterior commissure.

Figure 3. Corpus Callosum Development

Midline structures support the development of the corpus callosum in the developing human brain. Panels **a-c** depict coronal sections of human fetal brains at 17 weeks gestation. Panel **a** is labeled with an anti-GFAP antibody, panel **b** with an anti-Npn-1 antibody and panel **c** with an anti-nuclear factor 1a (Nfia) antibody.

Several midline glial structures are present at the cortical midline including the glial wedge (GW, \mathbf{a}) , the indusium griseum glia (IGG, \mathbf{a}) and the midline zipper glia $(MZG, \mathbf{d}, \mathbf{e})$. Pioneer axons, which form an additional potential guidance mechanism, express the guidance receptor Npn-1 $(\mathbf{b}, \mathbf{d}, \mathbf{e})$ and arise from the cingulate gyrus (\mathbf{b}) . In addition, the developing human brain contains subcallosal sling neurons, stained here with an antibody to Nfia (\mathbf{c}) .

Developing human and mouse brains differ in two significant ways at the midline: 1) In humans, differentiating astrocytes are found across the entire width of the midline (\mathbf{a} , see also \mathbf{d} , \mathbf{e}). These cells may either be part of the subcallosal sling or an extension of the MZG; 2) A population of NfiA/NeuN/Calretinin positive cells is present above the corpus callosum in humans (\mathbf{c}), but not mice. It is unclear whether these cells are similar to the subcallosal sling neurons or whether they may form neurons in the IGG (\mathbf{e}). Scale bar in B = 3mm in \mathbf{a} and \mathbf{b} and 400mm in \mathbf{c} . (\mathbf{a} and \mathbf{b} were previously published in 9 .)

Figure 4. Interhemispheric transfer in AgCC

Illustration of interhemispheric transfer (IHT) limitations in individuals with AgCC. A & B illustrate absence of interhemispheric conduction of the early visual evoked potential components that index sensory activity in the extrastriate visual cortex (i.e., P1 and N1 components) in both commissurotomy patients and individuals with AgCC ⁶⁸.

a. Visual evoked potential recording paradigm. Right visual field stimuli (top, solid lines) first result in evoked responses from locations within the left hemisphere (bottom left), and then following interhemispheric transfer (middle, solid arrow) evoked responses in right hemisphere locations (bottom right). The bottom panels show samples of typical evoked potentials from the left and right hemisphere recording locations within a healthy brain in response to right (solid lines) and left (dashed lines) visual fields. b. Comparison of left hemisphere evoked responses to stimuli in the right (solid lines) and left (dashed lines) visual fields. In the normal brain (b, top), the delay created by IHT is indicated by the later and smaller P1 and N1 to left visual field stimuli (dashed lines) compared to right field responses (solid lines). P1 and N1 components for left visual field stimulation are absent in left hemisphere recordings of both the person with AgCC (b, middle) and the commissurotomy patient (b, bottom), indicating that the corpus callosum is necessary for IHT of visual information. c, d. b, Experimental conditions that reveal limitations in ability to compare visual information from right and left visual fields 68. Each square is an example of a stimulus used in a letter (c) and dot-pattern (d) matching task. While participants looked at a central fixation point (solid diamond), two stimuli to be matched were flashed tachistoscopically in various configurations (bilateral or unilateral) in each trial. AgCC patients could make bilateral letter matches as well as control participants (presumably using extra-callosal pathways). However, AgCC patients could not successfully match bilaterally presented dot-patterns, which is a more complex task that cannot use semantic simplification, suggesting a limit on information transfer via non-callosal pathways.

Table 1. Syndromes associated with AgCC#

Α.	Syndro a.	omes with identified genes Andermann syndrome neuropathy and dementia	Gene symbol KCC3	Salient Features AgCC, progressive
	b.	XLAG AgCC, intractable epilepsy	ARX	Lissencephaly,
	c.	3.5	ZFHX1B	Hirschsprung
	d.	AgCC with fatal lactic acidosis deficiency, AgCC, brain malformations	MRPS16	Complex I & IV
	e.	770 1 0 12 7 1 0 1	L1CAM	Hydrocephalus,
B. Syndromes with AgCC seen consistently, no gene yet identified				
	a.		,	AgCC, polydactyly,
	b.	craniofacial changes, MR Aicardi syndrome		AgCC, chorioretinal
	c.	lacunae, infantile spasms, MR Chudley McCullough syndrome hydrocephalus, AgCC, colpocephaly		Hearing loss,
	d.	Donnai Barrow syndrome hernia, exomphalos, AgCC, deafness		Diaphragmatic
	e.	FG syndrome		MR, AgCC,
		craniofacial changes, macrocephaly		
	f.	Genitopatellar syndrome urogenital malformations, AgCC		Absent patellae,
	g.	Temtamy syndrome coloboma, craniofacial changes, MR		AgCC, optic
	h.	Toriello Carey syndrome changes, cardiac defects, MR		AgCC, craniofacial
	i.	Vici syndrome Recurrent infections, MR		AgCC, Albinism,
C.	Syndro a.		SPG11	Progressive
	b.	spasticity and neuropathy, Thin CC Craniofrontonasal syndrome craniosynostosis, facial asymmetry, bific	d nose	Coronal
	c.	Fryns syndrome	a nobe	CDH, pulmonary
	d.	hypoplasia, craniofacial changes Marden Walker syndrome		Blepharophimosis,
	e.	micrognathia, contractures, AgCC Meckel Gruber syndrome		Encephalocele,
		polydactyly, and polycystic kidneys		
	f.	Microphthalmia with linear skin defects	s Microopthalmia, linear skin	
	g.	Opitz G syndrome craniofacial changes, AgCC, MR		Pharyngeal cleft,
	h.	Orofaciodigital syndrome		Tongue hamartoma,
		microretrognathia, clinodactyly		T 1 .
	i.	Pyruvate decarboxylase deficiency seizures, severe MR and spasticity		Lactic acidosis,

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j. Rubinstein-Taybi syndrome Broad thumbs and

great toes, MR, microcephaly

k. Septo-optic dysplasia (DeMorsier syndrome) Hypoplasia of

septum pellucidum and optic chiasm

1. Sotos syndrome Physical overgrowth, MR, craniofacial changes

m. Warburg Micro syndrome Microcephaly, microopthalmia, microgenitalia, MR

n. Wolf-Hirschhorn syndrome Microcephaly, seizures, cardiac defects, 4p-

#reliable incidence data is unavailable for these very rare syndromes

*Many of these may also consistently have a thin or dysplastic corpus callosum, such as Sotos syndrome or AgCC with spastic paraparesis (SPG11). The overlap between AgCC and these conditions is still under investigation. Other gene symbols are omitted from section C.

Glossary

hypoplasia See Box 3

interhemispheric transfer (IHT)

Transmission of information between the cerebral hemispheres, typically assessed with laterally presented stimuli (see Box 1 and Figure 4).

pioneer axons

These are axons that innervate targets early in development and form a substrate for the guidance of later developing axons.

retrospective chart reviews

A retrospective analysis of medical records of a group of individuals with a particular condition or disease, typically used to study rare diseases for which prospective identification and follow up are difficult.

Mendelian disorder

A disorder controlled by a single genetic locus which is inherited according to Mendel's Laws.

X-inactivation

Early embryonic inactivation of the genes in each cell on one of a female's X chromosomes (may also occur in males with Klinefelter syndrome who have more than one X-chromosome). X-inactivation results in dosages of the proteins from genes on the X-chromosomes being equivalent to those in typical males (who only have one X chromosome to begin with).

isolated AgCC

Neuroanatomic description which includes complete absence of the corpus callosum, without other confounding brain abnormalities such as polymicrogyria, heterotopia or schizencephaly. Individuals with isolated AgCC frequently have colpocephaly and Probst Bundles.

MKS3

encephaloceles

A neural tube defect (NTD) that results in a sac-like protrusion of brain tissue and overlying meninges. These NTD's are frequently associated with other brain or craniofacial malformations and clinically with broad-ranging neurologic problems.

polydactyly

The anatomic variant of having more than the normal number of digits on the hands or feet. This is observed in approximately 1:500 births and is usually inherited as an autosomal dominant trait with variable penetrance.

Hirschsprung disease

A developmental disorder of the enteric nervous system resulting in absence of the neuronal ganglion cells in the distal colon, resulting in a functional obstruction of the colon. Can present

with a dramatically distended colon (megacolon) or with bowel perforation. It is a cardinal feature of Mowat-Wilson syndrome.

haploinsufficiency

A clinically evident symptom arising when one of the two copies of a gene is mutated, leaving a single functional copy and a presumed reduction in the level of the encoded protein.

Klinefelter syndrome

A genetic syndrome defined as a 47, XXY karyotype in a phenotypic male. Patients frequently have small testes, minimal sperm production, breast enlargement in puberty and psychosocial problems.

chorioretinal lacunae

Punched out lesions in the pigmented layer of the retina that cluster around the optic disc that are pathognomonic for Aicardi syndrome.

microarray-based comparative genomic hybridization (CGH)

A method that compares the quantity of DNA across the whole genome between two individuals. Two DNA samples are labeled red and green respectively and both hybridized to a slide that has arrayed many thousands of spots containing DNA from unique places in the genome. The color ratio at each spot determines the relative amount of DNA present between the two samples.

Sotos syndrome

Also known as cerebral gigantism, is a genetic disorder that results in early physical overgrowth and cognitive impairment. Most cases are caused by haploinsufficiency of the gene NSD-1, which is a coregulator for steroid receptors.

heterotopia

In general, this terms refers to the displacement of neuronal cell bodies into the white matter.

commissurotomy

Surgical procedure that involves severing the corpus callosum as well as the anterior commissures (may also include severing of posterior and hippocampal commissures). This is the original "split-brain" procedure as reported by Roger Sperry ⁴.

tachistoscopic

Presentation of visual stimuli more rapidly than the eyes can move. Tachistoscopic presentation thus results in a visual stimulus being perceived in only one hemisphere and representation of the image in the opposite hemisphere will require interhemispheric transfer of information.

Stroop interference effect

A reaction time measure when identifying one feature of a stimulus, while inhibiting a dominant tendency to identify it according to an interfering feature (for example the normally increased reaction time when naming the ink color of the word Red printed in green ink).

bilateral field advantage

The normal decrease in reaction time when comparing two stimuli presented in opposite visual hemifields, compared to presentation of both within one hemifield. The reason for this advantage is dual processing, i.e. each hemisphere only has to process on stimulus. Without efficient IHT, there cannot be such an advantage.

anterior commissure

Small band of approximately 50,000 axons that connect the cerebral hemispheres. The anterior commissure connects the temporal lobes and is located at the base of the fornix.

dichotic listening

An research method testing language lateralization by simultaneously presenting different auditory input to each ear. The degree to which subjects preferentially recall information from one ear or the other is an indication of which hemisphere is dominant in language processing.

syntax

Grammatical arrangement of words and phrases in a sentence, which impacts relationships of meaning. For example changing the placement of a word or phrase can change the meaning.

linguistic pragmatics

The processes that allows one to go beyond literal meaning of language and actually interpret the speaker's intended meaning. This may involve utilizing second-order meanings, body language, vocal inflection, context, and other factors.

alexithymia

Impairment in expression of one's feelings and mood-states. A dominant hypothesis is that alexithymia arises from compromised connection between the language processing in the left hemisphere and affect processing in the right.

valence

A continuous scale from pleasant to aversive.

dMRI: diffusion magnetic resonance imaging

This broad term covers both diffusion weighted MRI data acquisition and image analysis of this data ¹¹⁷, including diffusion tensor imaging (DTI:TERM). The MRI signal is weighted by the amount of water diffusion within tissues. The weighting can vary with direction allowing diffusion anisotropy arising from microscopic restrictions in biological tissues to be observed.

Andermann syndrome **This is defined in the genetics section**

DTI: diffusion tensor imaging

Anisotropic diffusion within tissues is modeled as a second-rank tensor which can be calculated from diffusion-weighted MRI acquired in six or more non-collinear directions. The tensor at each point in the image can be visualized as an oriented and scaled ellipsoid. More simply, quantities such as the mean diffusivity and fractional anisotropy can be calculated from the tensor and visualized as conventional images. The tensor contains information about likely axonal fiber direction and can be used to create virtual fiber tracts through the DTI, reflecting structural connectivity in white matter.

callosotomy

Surgical procedure that involves severing only the corpus callosum, either in part or in its entirety, leaving other commissures intact. This has also been described by some as a "splitbrain" procedure ⁵.

Endophenotypes

Hereditary characteristics that are associated with a condition, but are not a direct symptom of that condition

Author Biographies

Lynn K. Paul – Dr. Paul began research on agenesis of the corpus callosum with Dr. Warren Brown in 1993, as part of her doctoral work in clinical psychology at Fuller Theological Seminary. Her research has focused on the cognitive and psychosocial characteristics in individuals with Primary AgCC. In 2004, she began the Corpus Callosum Research Program within Dr. Adolph's laboratory and is now a Senior Research Fellow at Caltech. This program involves histological, neuroimaging, and social-cognition studies of AgCC. Dr. Paul was also the founding president of the National Organization for Disorders of the Corpus Callosum, a non-profit that gathers and disseminates information about these conditions (www.nodcc.org).

Elliott H. Sherr - Dr. Sherr is an Assistant Professor in Neurology and Pediatrics at UCSF. He directs the Brain Development Research Program, studying the genetics and biology of AgCC and other developmental brain malformations. He obtained his undergraduate degree at Stanford University and his M.D. and Ph.D. at Columbia University.

Ralph Adolphs - Dr. Adolphs received his undergraduate degree from Stanford University, and his graduate training in neurobiology from the California Institute of Technology. Since 1993 he has been investigating the neural basis of emotion and social behavior in humans. Much of this work studied neurological patients with focal brain lesions, research he conducted first as a postdoc and subsequently as faculty at the University of Iowa. Since 2004, Dr. Adolphs has been professor of psychology and neuroscience at the California Institute of Technology, while maintaining an adjunct appointment also at the University of Iowa. His long-term goals are to understand how emotion modulates perception, attention, memory, and social cognition. Dr. Adolphs is the recipient of awards from the Sloan Foundation, the Klingenstein Fund, the EJLB Foundation, and currently holds a 21st Century Science Award from the James S. McDonnell Foundation.

Warren Brown - Warren S. Brown (Ph.D., University of Southern California) is currently Professor of Psychology at the Fuller Graduate School of Psychology, where he is Director of the Lee Travis Research Institute. Prior to Fuller, Brown was at the UCLA Department of Psychiatry, where he was involved in research on childhood schizophrenia, Alzheimer's disease, and autism. He continues to be a member of the UCLA Brain Research Institute. His primary research interest is the role of the corpus callosum in human higher cognitive and social processes. For the last 15 years, Brown's laboratory has been studying the cognitive and psychosocial consequences of agenesis of the corpus callosum in individuals who have intelligence within the normal range.

Pratik Mukherjee - Dr. Mukherjee is a neuroradiologist and assistant professor of radiology and bioengineering at the University of California, San Francisco. His research over the past 8 years has focused on applications of diffusion tensor imaging (DTI) to the noninvasive microstructural analysis of biological tissues with special emphasis on the developing brain. He published some of the first DTI studies in the areas of stroke, hypertensive encephalopathy, and childhood brain development and is currently conducting prospective studies of DTI in

traumatic brain injury and callosal dysgenesis. Dr. Mukherjee continues to develop diffusion MRI methods that surpass DTI for non-invasive mapping of human white matter.

Linda Richards – Dr. Richards completed her PhD in 1994 at The Walter and Eliza Hall Institute, Melbourne, Australia in the laboratory of Prof. Perry Bartlett and postdoctoral training at The Salk Institute for Biological Studies, California, with Prof. Dennis O'Leary. She began her laboratory at The University of Maryland in 1997, where she remained until 2005 when she was appointed as an Associate Professor at The University of Queensland, Australia. Her laboratory investigates the development of the cerebral cortex and midline commissure formation in the brain. The focus is on the development and patterning of the cortical midline and aspects of glial and neuronal development that impact the formation of midline structures such as the corpus callosum.

J. Michael Tyszka – Dr. Tyszka is a graduate of Cambridge University where he developed a long-standing interest in magnetic resonance imaging under Laurie Hall. He began his collaboration with Scott Fraser's Biological Imaging Center at Caltech in the mid 1990s before joining the group full-time in 2001. He has published widely in the magnetic resonance imaging technical literature and his research interests range from engineering in MR microscopy, through MR diffusion histology, to the integration of structural and effective connectivity models of human brain function. He is currently Director of Magnetic Resonance Physics at the Caltech Brain Imaging Center in Pasadena, California.