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Brain Connectivity and Neuroimaging of Social Networks in Autism

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Abstract

Impairments in social communication (SC) predominate among the core diagnostic features of autism spectrum disorders (ASDs). Neuroimaging has revealed numerous findings of atypical activity and connectivity of ‘social brain’ networks, yet no consensus view on crucial developmental causes of SC deficits has emerged. Aside from methodological challenges, the deeper problem concerns the clinical label of ASD. While genetic studies have not comprehensively explained the causes of nonsyndromic ASDs, they highlight that the clinical label encompasses many etiologically different disorders. The question of how potential causes and etiologies converge onto a comparatively narrow set of SC deficits remains. Only neuroimaging designs searching for subtypes within ASD cohorts (rather than conventional group level designs) can provide translationally informative answers.

An Ever-Increasing Public Health Challenge

ASDs represent a group of neurodevelopmental disorders with increasing prevalence, which has doubled within less than a decade (in large part, as a result of improved detection and growing awareness [1]) and is currently estimated in the USA at 1 in 59 children [2], with similar rates (1.5–2%) reported in other countries and continents [3–5]. Aside from personal suffering in affected individuals and their families, ASDs present a major public health challenge, with an estimated annual cost of ~US\$300 billion in the USA, projected in one report to approach 3–4% of the gross domestic product within the next decade [6].

Core diagnostic features of ASDs (see Glossary) are sociocommunicative impairments and restricted and repetitive behaviors and interests, with symptom onset during the first years of life [7]. ASDs are heterogeneous in presenting symptoms, developmental outcomes, and adaptive function, with symptom severity levels ranging from mild (requiring minimal support) to severe (necessitating substantial support throughout the lifespan). According to the most recent diagnostic consensus summarized in the *Diagnostic and Statistical Manual of Mental Disorders (DSM-5)* [7], core autism symptoms can occur in individuals with or without intellectual or language delays, effectively broadening the diagnosis to include

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Appendix A Supplementary data

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people with the full range of cognitive and language functioning. While delayed language acquisition is no longer among the diagnostic criteria, language skills and IQ remain the strongest predictors of outcomes in both naturalistic studies of ASDs [8,9] and in treatment response [10,11]. Males are affected about four times more frequently than females [2], although females with ASDs may be underidentified [12,13], likely due to gender differences in symptom presentation and comorbid features [14–16] (Box 1). All in all, because of the onset in early childhood, the life-long impact of ASDs on the health, economic well-being, social integration, and quality of life of affected individuals and their families is immense. On the societal level, the direct and indirect costs to the educational, healthcare, and economic sectors emphasize the need to continue the search for neurobiological causes, enhanced detection, effective and targeted interventions, and, ultimately, prevention.

Social Cognition and the ‘Social Brain’

Despite heterogeneity in behavioral manifestations across sensory, motor, attention, language, and other domains, impaired social functioning is a core feature of ASDs [17]. Deficits in what the *DMS-5* calls ‘social communication’, including diminished social responsiveness, difficulty recognizing others’ emotions and intentions, deficits in developing, maintaining, and understanding relationships, or relating to peers in a reciprocal manner, are considered the most universal and specific characteristics of the autism spectrum [18]. Social skills and competence are integral to many aspects of human behavior, laying the foundation for successful functioning in a dynamic, complex social world. The fundamental nature of social engagement and its prominence early in life is evidenced by *prenatal* sensitivity to human voices [19] and face-like stimuli [20], orientation to social cues, such as human faces and gestures within the first days of life [21,22], and heritability of socially relevant attention to facial eye and mouth regions observed in infants [23]. While it is unknown whether this primal preferential orientation to social cues is absent or reduced in fetuses and neonates who are later diagnosed with ASDs, diminished attention to social cues has been well documented in the first years of life in children with ASDs, including diminished attention to human voices [24], biological motion [25], eye gaze [23,26], and human faces [27], and impaired joint attention skills [28]. This is critical because the typically early-emerging preference for social cues is considered a precursor to the various forms of social understanding and adaptive social communication, arguably the most important attainments of early childhood development [29,30] (Box 2).

The SC functions associated with social orientation, processing of social cues, and understanding of others are supported by a distributed network of brain regions collectively referred to as the ‘social brain’ [31]. This collection of brain networks involved in processing social signals includes the dorsal and ventral medial prefrontal cortices (dMPFC and vMPFC), anterior and posterior cingulate cortex (ACC and PCC), amygdala, posterior superior temporal sulcus (pSTS), temporoparietal junction (TPJ), inferior occipital gyrus (IOG), fusiform face area (FFA), and the insula [32–34]. Extensive neuropsychological and functional neuroimaging evidence indicates that these social brain regions are related to specific domains of social cognition, such as analysis of faces and gaze by the IOG, FFA, and pSTS [35,36], emotional processing in the amygdala [37], processing of others’

intentions, feelings, and points of view (i.e., theory of mind [ToM] or mentalizing) in the dMPFC, PCC, and TPJ [38], and imitation and understanding of others' actions, which is crucial for social learning, by the regions associated with the mirror neuron system (MNS), including the premotor cortex (PMC), intraparietal sulcus (IPS), and pSTS [39]. A growing number of connectivity studies show that these social brain regions are organized in large-scale functional networks supporting specific social functions (reward-related system [40], ToM network [41], the MNS [42], and face perception network [43]; Figure 1).

Abnormalities of Social Brain Networks in ASDs

Many of the abilities supported by the social brain networks overlap with core impairments associated with ASDs, including difficulties inferring other people's emotions or intentions and recognizing facial expressions, as well as deficits in imitation, social learning, and joint attention [44–46]. While activation likelihood estimation (ALE) meta-analyses have reported consistent dysfunction (primarily hypoactivation during social tasks) of most of the individual 'social brain' regions mentioned above in ASDs [47–51], there is a growing consensus that brain abnormalities in ASDs are not localized to one or a few regions but rather implicate alterations in the connectivity of distributed brain networks. Since the early 2000s, a method of choice in the study of network connectivity has been functional connectivity MRI (fcMRI). Aberrant patterns of connectivity have been identified within and between widely distributed networks supporting core social functions [52–54]. For instance, reduced network integration, or weaker inter-regional connectivity, has been observed for the face perception [49], imitation [55], and amygdalar emotional-processing networks [56,57]. Furthermore, there is convergent evidence of reduced segregation (or differentiation) of social networks in ASDs, reflected in excessive connections with extraneous regions that are not part of neurotypical social networks. These findings appear to converge on a pattern of dual impairment, affecting both network integration and segregation, and resulting in reduced network specialization and processing efficiency of brain systems crucial for SC functions. However, not all findings to date have been consistent with this interpretation, with some reports of global overconnectivity across all social and nonsocial networks [58] or weaker connectivity between social brain networks [53,59]. One complicating aspect is that atypical maturational trajectories in ASDs may be associated with underconnectivity of a given network at one developmental stage, but overconnectivity at another [60]. We return to the question of how divergent findings may be explained in the following section.

Particularly relevant to the broad ASD phenotype, most of the regions comprising the social brain networks also support functions outside of the social domain. For instance, TPJ, considered a nexus of theory of mind and self-other distinction [38], is also associated with domain-general attentional reorienting to salient cues [61]. Similarly, while dMPFC, as part of the theory of mind network, is engaged in inferring others' mental states [62,63], it has also been implicated in abstract tasks without mentalizing content, such as categorization [64,65]. Even the FFA, considered a canonical social brain region for its crucial role in face perception, is involved in domain-general functions beyond the social context, including perceptual expertise for nonface objects [66,67].

Challenges and Perspectives

Many Methods, Uncountable Findings, and Few Answers ...

The imaging literature on brain connectivity investigating core SC symptomatology in ASDs has rapidly grown over the past decade. While a gross pattern of reduced network integration and segregation appears to emerge, as described above, the numerous findings have not added up to a fully coherent picture of brain anomalies that are crucial for impaired SC in ASDs (*cf.* [60,68]). One set of problems is methodological. First, different techniques are sensitive to different neural parameters that are directly or indirectly related to connectivity and brain network organization. *Connectivity* strictly refers to the presence of anatomical connections and, thus, to axonal and synaptic organization, which in humans can be directly examined only in postmortem studies. Diffusion-weighted imaging allows indirect inferences on anatomical connectivity based on differential water diffusion [69]. By contrast, *functional connectivity* is a derivative concept based on the often tacit assumption that distal brain regions show statistically related activity patterns only if they communicate (directly or indirectly) via anatomical connections. Within the wide field of functional connectivity, different techniques, from positron emission tomography in early studies [70] to fMRI, EEG, and magnetoencephalography, are again sensitive to different aspects of neural activity fluctuations. These differ with respect to signal source (e.g., hemodynamic versus electrophysiological), temporal frequency bands, and spatial resolution, making the integration of findings from different techniques challenging [71]. Second, even within the boundaries of a given technique, such as fMRI, different methodological approaches can result in divergent findings in the same data set [72]. The need for clearly defined best practices in data preprocessing and analysis appears plausible, but is complicated by the fact that potential incremental improvements in processing steps are proposed continuously (e.g., [73]) and any previously accepted set of practices will no longer be ‘best’ after a short time.

... but Can there be Clear Answers to an Ill-Posed Question?

Genetic investigations into ASDs, considered to promise explanatory models of causation decades ago, have failed to provide clear and simple answers as to how and why affected children fail to develop crucial SC skills. The reasons are manifold, having to do with the complexity of gene-behavior relationships, which are somewhat more tractable in syndromic forms that affect only a single locus, such as fragile X or Rett syndrome [74], but largely intractable with the current state of knowledge in polygenic disorders, such as ‘idiopathic’ ASDs. The results of 30 years of genetic ASD research show some analogy to the challenges in neuroimaging described earlier. Large-sample studies have revealed more and more risk genes, now in the hundreds [75]. Although most of these account for only a small causative factor, penetrance (the rate at which individuals with a given genetic variant develop a specific phenotype, such as ASD) varies widely [75]. Thus, causation in most individual cases of ‘idiopathic’ ASDs may be due to some permutation of ‘multiple hits’ among numerous risk genes [76], with added roles of epigenetic [77] and environmental factors [78]. In other words, both in neuroimaging and genetics, there are numerous findings that appear to apply to some cases of ASDs, but not others, and that probably need to occur in some unknown combination in order to result in ASD symptomatology.

Have neuroimaging and genetics research failed the community of families with ASDs and the funding organizations that have subsidized these fields? This question misses the point. If there has been failure that has prevented the science of ASDs to generate mechanistic models of causation and precision medicine, it is a failure to appreciate the inadequacy of a clinical label such as ‘autism spectrum disorder’ (in the singular of the *DSM-5* [7]) in the pursuit of neurobiological causes. From a developmental neurobiology perspective, the expectation that a consensus-based catalog of behavioral observations (as listed in the *DSM-5*) could tractably correspond to a small set of genetic causes or brain features is misguided. This expectation may be prompted by the existence of some single-gene disorders associated with autism symptomatology [74]. The reverse inference – that a disorder defined by a distinct set of behavioral symptoms, such as ASD, could therefore be traced back to a single cause or a small set of causes - is however unfounded.

Specificity of Sociocommunicative Symptomatology: Where is the Convergence?

The breadth of findings, both with respect to neuroimaging features found to be atypical in ASDs and with respect to the numerous genetic (and other) risk factors implicated, is remarkable given the comparatively circumscribed domain of SC core symptoms on which clinical diagnosis is based [7]. Therefore, some convergence may be expected to occur in development, from a very wide range of possible causes onto a narrower range of core symptoms. One proposal has been made in the genetics literature as to the convergence of many ASD risk genes into gene networks or modules [79,80], some of which may specifically affect synapto-genesis, synaptic function, and circuit formation [81,82]. This view may be supported by the convergence of a large number of knockout mouse models (of genes associated with ASDs) onto few distinct neuroanatomical phenotypes [83]. However, the argument could be compelling only if based on quantitative tests showing that the fraction of ASD risk genes involved in connectivity is significantly greater than would be expected from the fraction of genes somehow affecting brain connectivity within the entire genome. A corresponding argument, according to which autism is a disorder at the brain network level, has been made in the imaging literature. Although broadly accepted (*cf.* [60,68]), this argument is open to an analogous critique: Cognitive neuroscience has generally moved towards understanding all functional systems in terms of distributed networks [84], highlighting the importance of network connectivity, and this view has been *generally* applied to the study of psychopathology [85]. The view of autism as a connectivity disorder therefore conveys little *specific* insight.

However, even if one accepts a connectivity approach as specifically informative about the causation of ASDs, the convergence question remains. Since all functional systems (sensorimotor, limbic, and higher cognitive supramodal) are organized in distributed interconnected networks, why would a ‘connectivity disorder’ specifically affect SC? The question may have to be slightly softened, as it is known that ASDs are often accompanied by impaired or atypical functioning in many domains other than SC, such as sensory [86], motor [87], attention [88], and executive [89]. Although such evidence is now acknowledged by the inclusion of sensory symptomatology and greater emphasis on restricted interests and

behaviors in the *DSM-5*, deficits in social interaction and communication skills remain the defining core feature of ASDs. What can account for this domain specificity? Several scenarios may be considered (Figure 2).

Scenario 1 implies causal specificity, according to which ‘multiple hits’ ([epi]genetic and/or environmental) causing the emergence of an ASD predominantly affect neural circuits that are crucial for social cognition and communication, for example through region-specific gene expression [90]. This appears to be supported by the predominance of imaging findings implicating sociocommunicative brain networks (as described earlier). However, there is an understandable bias in study designs to preferentially target domains of core symptomatology. This alone may explain why findings for these domains are more abundant [91]. Notably, anomalous connectivity has also been reported for domains beyond core ASD symptomatology, such as motor control [92,93], salience network [94,95], and circuits connecting basal ganglia [96], thalamus [97,98], and cerebellum [99,100] with cerebral cortex. All in all, the neuroimaging literature indicates that atypical connectivity patterns in ASDs are not exclusive to sociocommunicative circuits.

A second set of scenarios relates to timing specificity. In scenario 2a, genetic and other causes have greatest impact at developmental stages crucial to the emergence of SC. Neurodevelopmental disturbances, such as gray [101] and white matter growth anomalies [102] or enlarged cerebrospinal fluid compartments [103], arise during the first 1–2 years of life, or possibly prenatally [104]. This may affect preferential social responses that have been observed prenatally and early postnatally (as described earlier), although some more complex SC abilities known to be affected in ASDs, such as ‘theory of mind’ (e.g., [105]), fully develop later in life [106]. In related scenarios, early disturbances specifically affect *precursors* crucial for subsequent SC development. In scenario 2b, these may be limbic defects affecting social motivation and reward [46]. Alternatively, in scenario 2c, these are early sensorimotor disturbances. This scenario seems supported by growing evidence of atypical or impaired development of auditory [107], visual [108], somatosensory [86], and motor functions [109] in ASDs. Scenario 2c might imply that core ASD symptomatology is sensorimotor and that SC impairments are secondary and derivative. However, there is no conclusive evidence that sensorimotor deficits are a necessary condition for the emergence of SC symptomatology in ASDs. Even if one acknowledges the plausibility of such developmental links, the specificity of resulting SC impairment would require further explanation, as other developmental disorders have also been related to early sensorimotor impairments (e.g., rapid auditory processing deficits in specific language impairment [110]).

A third scenario implies specific vulnerability and pertains to the domain of impairment itself. If SC abilities are selectively vulnerable to neurodevelopmental disturbances, causal factors that have little domain specificity may result in a phenotype with greater behavioral specificity. However, this scenario on its own is not plausible, because early neurodevelopmental disturbances can also result in other deficit profiles, such as broad impairment of overall level of functioning in intellectual disability [111], or predominant deficits in circumscribed domains, such as spoken [112] or written language [113], or attention and cognitive control [114]. It may be more plausible in combination with scenario 2, in the sense that SC abilities or their crucial building blocks could be selectively

vulnerable to specific neurodevelopmental disturbances at specific maturational stages only, possibly because of the highly distributed organization of SC networks in the brain (as described earlier). This would imply that developmental disorders achieve specificity based on the predominant timing of insult or disturbance. However, even this combination of scenarios 2 and 3 has little direct evidence in support. Indeed, based on available evidence, one would be hard pressed to pinpoint the exact timing of crucial neurodevelopmental disturbances in ASDs: Is it several years after birth when brain overgrowth peaks [101] or when, in some cases, developmental regression is observed [115]; or prenatally, when intrauterine risk factors, such as maternal inflammation, occur [116]; or at earliest stages of neuronal proliferation, as suggested by studies using induced pluripotent stem cells [117]? Since the timing of crucial insult in ASDs is undetermined (probably because it cannot actually be pinpointed to a single developmental stage), any argument that timing of insult at a stage of specific vulnerability of SC functions (or their building blocks) could account for domain specificity in outcome symptomatology appears less than compelling.

Finally, returning to questions raised earlier, we may consider whether the construct of ASD itself could be a convenient clinical fiction, serving to categorize children with developmental disorders for the purpose of treatment decisions. This would imply that convergence is an artifact of diagnostic procedures, unrelated to underlying biology. The argument has been made (e.g., [118]), and may be considered with reference to the human mind's penchant to perceive categorical boundaries even where there are none in the physical world [119], as, for example, in phonemic categorization [120]. This view is substantiated by the contemporary trend towards transdiagnostic approaches to research and clinical practice, focusing on features that cross diagnostic and categorical boundaries (e.g., [121,122]), which highlights, for instance, that SC deficits are also present in schizophrenia spectrum disorders, social anxiety, and other developmental or conduct disorders. However, while clinicians will acknowledge comorbidities and fuzzy lines of demarcation between diagnostic categories, they will likely reject the view of ASD as a convenient fiction because it is incompatible with their everyday clinical experience.

Many ASDs: Are Large Samples the Answer?

For reasons solely to do with logical transparency, we have presented the different convergence scenarios as distinct alternatives. In reality, each of them probably captures an aspect of the true complexity in the relation between causal factors and SC symptomatology, and each scenario may apply to different degrees to the varying etiologies underlying ASDs. It is likely that genetic and epigenetic risk factors affect SC abilities with *some specificity* at *maturational stages* of increased *vulnerability* and in regions with predominant effects on SC functions or its precursors. In addition, as discussed earlier, the *DSM-5* presentation of ASD as a singular disorder may indeed be misleading, because it solely reflects a relative convergence at the behavioral outcome level (onto a set of diagnostic criteria and a range of severities), of what at the biological level most likely comprises an array of diverse etiologies. The attempt to understand behavioral convergence made in the previous section is therefore the exact flipside of the need to understand diversity of causation and trajectories of neurodevelopmental disturbances.

From a purely statistical perspective, a plausible remedy for heterogeneity and expected cohort variability is the pursuit of large sample sizes. Indeed, with the increasing availability of large samples, such as the Autism Brain Imaging Data Exchange [123], some recent studies have presented findings that appear definitive solely based on ample statistical power. For one example, a recent ‘mega-analysis’ reported reduced amygdala volume in a cohort of 1508 children and adults with ASDs, compared with 1601 typically developing participants [124]. The finding is significant ($P < 0.05$, even after correction for numerous comparisons), albeit with a miniscule effect size (Cohen’s $d = 0.08$). Although this finding derives from a large sample that is probably more representative of the ASD population than smaller samples in many other studies, the effect size indicates that the finding may have little significance beyond statistics. It could be meaningful if ASD were associated with a subtle, but functionally critical decrease in amygdalar volume in every or most individuals with the disorder, but this is improbable. More likely, a minority of etiologies grouped under the current ASD diagnostic label are associated with robustly reduced amygdalar volumes, whereas no such effect is seen in most cases. Therefore, any conclusion that ASD is associated with reduced amygdala volume, while statistically accurate, would be misleading.

There is great awareness of the problem of underpowered investigations in neuroimaging [125], but a complementary problem of overpowered group-level comparisons that may lend unsuitable prominence to findings with very small effect size is not generally considered. In this age of emerging large-sample collaborative endeavors (e.g., [123,126]), a careful distinction between statistical and conceptual significance is needed (*cf.* [127]). However, while large sample studies may generate findings that provide only minimal information about the general ASD population, they are, at the same time, indispensable for the detection of variants of the disorder.

Towards Translational Goals

As discussed earlier, ASDs may encompass a large set of etiologically different disorders [128], each associated with potentially divergent trajectories of neurodevelopmental disturbances. The identification of ASD subvariants is therefore crucial, because targeted treatments may only be achievable for specific variants, rather than for the population as a whole. The approach is often characterized as ‘stratification’, with the misleading implication that subvariants are expected to form ‘layers’ differing in one dimension only. In fact, constructs such as ‘low-functioning’ versus ‘high-functioning’ autism are unlikely to capture the biological diversity of ASD subtypes.

Although it can be assumed that SC symptomatology in ASDs is associated with atypical functional and structural brain organization, etiological diversity implies that the specific features of anomaly may diverge across subvariants of the disorder. Therefore, two individuals with ASDs may show similar profiles of behavioral symptoms, but these may be linked to different brain features. Attempts to subtype ASDs based on behavioral variables have been made (e.g., [129–131]); however, variants defined by shared brain features may relate more closely to etiological variants [132]. This can be presumed because both structural and functional brain organization reflect developmental history, not simply present outcome state. For example, fundamental organizational principles of regional specialization

of cerebral cortices result from early developmental processes, such as neuronal migration along specific radial glial cells and early establishment of thalamocortical afferents [133]. Neuroimaging in children (and even in adults) is therefore ‘archeological’ because it can reveal neurodevelopmental disturbances and can unearth ‘intermediate phenotypes’ that are more directly informative of causation than behavior alone [134]. Thus, imaging has the potential to identify brain markers resulting from the convergence of differential etiologies, which may aid the development of targeted behavioral or biological interventions [135].

However, imaging measures are compounded by experiential plasticity related to environmental effects in the broadest sense (see Outstanding Questions). Therefore, imaging indices both distally reflect traces of prenatal and early postnatal neurodevelopmental disturbances and, at the same time, proximally show the neural bases of behavioral impairment in the current outcome phenotype of a single child (or adult) with ASDs. This opens two avenues for neuroimaging in ASDs that have traditionally not been clearly distinguished (Figure 3). Per its archeological ability to reveal intermediate phenotypes, neuroimaging can contribute to the identification of etiologically defined variants of ASDs, each of which may be expected to relate more tractably to genetic, epigenetic, and other causative factors. Defining such variants will be a first step towards mechanistic models and the implementation of precision medicine. On the other hand, per its ability to reveal proximal neural causation of outcome SC impairments, neuroimaging can identify targets for brain intervention techniques (e.g., transcranial magnetic or direct cortical stimulation [136,137], neurofeedback training [138]). Neither side is ready for this. The neuroimaging field needs to develop better tools to identify variants within ASD cohorts and pinpoint brain features crucially linked to SC (and other) deficits at the level of the individual; and brain stimulation techniques need to be calibrated for targeted use, in order to affect the function or connectivity of specific brain regions in a desired direction.

Concluding Remarks

Social impairments predominate among the core diagnostic criteria of ASDs. However, it is likely that these impairments result from the convergence of many different biological etiologies onto a set of phenotypes fulfilling these diagnostic criteria. Neuroimaging studies of atypical brain connectivity linked to social symptomatology must therefore move away from a focus on group-level effects towards the detection of variants within the disorder. Identifying biological subtypes and individualized targets for intervention will be the crucial challenge for neuroimaging in its quest to alleviate social impairments in children and adults with ASDs.

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Glossary

Brain network integration and segregation

the human brain is organized into large-scale networks reflecting coordinated neural activity among spatially segregated cortical areas, which takes up a large fraction of metabolism in the brain. Intrinsic networks have been observed during, or in the absence of, overt cognitive tasks or external stimuli, and can be detected during various states of consciousness, including sleep and sedation. Specialized and functionally optimized brain networks are the outcome of prolonged neurodevelopmental processes, including constructive (e.g., synaptogenesis and axonal myelination) and regressive events (e.g., synaptic pruning and axonal loss). Overabundant connectivity during early postnatal years is pruned away based on experience-driven activity patterns, resulting in distinct and functional specialized networks. These developmental principles appear to be impaired in ASDs

Core diagnostic features of ASDs

a diagnosis of an ASD is based on the presence of two core symptom domains: (i) SC impairments, including deficits in social-emotional reciprocity, in nonverbal communicative behaviors, and in developing, maintaining, and understanding relationships; and (ii) restricted and repetitive patterns of behavior and interests, such as stereotyped or repetitive speech, movements, or use of objects; excessive adherence to routines and insistence on sameness; highly restricted, fixated interests; and atypical reactivity to, or unusual interest in, sensory input. ASD symptoms must be present early during development. While often detected during toddlerhood or preschool, they may not raise concern until social demands exceed a child's capabilities (e.g., in elementary school with progressively rising academic and social demands)

Functional connectivity MRI (fcMRI)

similar to conventional functional MRI, fcMRI uses times series of brain images acquired every 2 s or faster. The contrast of interest relies on changes in the blood oxygen level-dependent (BOLD) signal associated with local neuronal activity and synaptic transmission. FcMRI detects correlations (or statistical dependence) of BOLD signals between different brain regions, interpreted as functional connectivity. In the most common type of fcMRI, called intrinsic fcMRI, images are acquired during rest (without specific task) and spontaneous synchronization of BOLD fluctuations at low frequencies (typically <0.1 Hz) is detected. However, fcMRI can also be performed to detect BOLD correlations associated with task performance or sensory stimulation

'Idiopathic' ASD

indicates cases of ASD with no known cause, as opposed to syndromic ASD with established genetic (or other) causes. Although this term (or its equivalent 'nonsyndromic' ASD) is still commonly used, it is destined to become less suitable with improving knowledge of causation in variants of the disorder

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Box 1.**Gender-Related Differences in ASDs**

Given the high male:female ratio in ASD prevalence, relatively little is known about female variants. While growing evidence indicates that profiles of SC impairments in ASDs likely differ between genders, it is unclear whether this might be due to different biological mechanisms (e.g., existence of specific factors protecting females from developing ASDs [139]), or an ascertainment bias (i.e., diagnostic criteria biased by the more prevalent male presentation [13,140]). Behaviorally, females with ASDs exhibit fewer restricted and repetitive behaviors [16], more social motivation [141], and less impaired nonverbal communication, including normative eye gaze following behavior [142], compared with males with ASDs. However, some evidence suggests that these gender differences in SC are only observed in higher-functioning samples and are absent altogether in lower-functioning samples [143]. Biologically, females may be more protected from heritable and *de novo* ASD risk variants, with sex chromosomal genes and sex hormones modulating the effects of genetic variants on the presentation of ASD symptoms [144]. Furthermore, limited evidence suggests differences in brain organization between males and females with ASDs, with the neuroanatomical features of ASDs involving different structures or growth trajectories in females than in males, including distinct patterns of early brain overgrowth (i.e., enlargement of brain volume early in life observed more prominently, or even exclusively in boys [145]), local differences in gray and white matter, and attenuation of neurotypical male > female volumetric differences [146].

Baron-Cohen's 'extreme male brain' theory of autism proposes that autism represents an amplification of specific aspects of typical sexual dimorphism in cognition, with the extreme female brain characterized as 'empathizing,' and the extreme male brain as 'systematizing' [147]. However, this 'masculinization' theory linking the ASD etiology to the masculinizing effects of fetal testosterone [148] remains controversial, with an alternative theory proposing that ASDs are associated with 'gender incoherence' or androgyny, rather than with extreme masculinization, at the neurophysiological level [149]. Notably, the two theories make the same neuroanatomical predictions for females with ASDs (i.e., a more masculinized neural signature), but not for males with autism (i.e., more feminized features according to the latter). One study that directly compared brain connectivity patterns in males and females with ASDs reported reduced gender differentiation relative to the differences observed between typically developing (TD) boys and girls, supporting neural androgyny rather than masculinization [150]. Larger studies are required to replicate these findings and to understand differences in brain connectivity that could contribute to the sex differences in SC behaviors.

Box 2.**Competing Accounts and Developmental Trajectories of Social Dysfunction in ASDs**

The neurocognitive accounts of impaired social cognition in ASDs can be largely summed up as belonging to two camps: while domain-specific explanations focus on (neural or cognitive) mechanisms specialized for social inferences and mentalizing (e.g., [151,152]), social orienting or social attention accounts propose that children with ASDs fail to preferentially orient to people, or social contingencies in general due to impairments in bottom-up, low-level attentional processes (e.g., [153]), leading to reduced opportunities for experiential learning. The latter specifically focus on social brain networks and mechanisms supporting processing of social cues, including human faces, voices, and biological motion. These bottom-up accounts emphasize that impoverished social information input (resulting from poor attention to social information) is responsible for diminished social experience and social engagement, which in turn compromise the development of higher order social cognition, such as theory of mind. Mechanistic explanations of diminished sensitivity to stimuli with social contingencies are still lacking, but the apparent indifference to social cues at early developmental stages impedes the development of social interactions, and of supporting social brain networks [17,52]. Potentially orthogonal to these models is the executive dysfunction account, which suggests that impaired executive functions, such as goal-directed initiation, inhibition, working memory, and cognitive flexibility, underlie or serve as moderators of SC impairments in ASDs [154].

Children who develop ASDs do not appear to exhibit clear impairments in SC until the end of the first year of life, with some children continuing on a more typical trajectory until later in the second or third year of life, followed by a regression in SC skills. Among the most common early behavioral signs of ASDs (observable and apparent during the second year of life) are poor eye contact, poor joint or shared attention, and diminished social or shared smiling [155], all key features of social orienting. The evidence for these aberrant social-developmental milestones comes from both retrospective studies of parental reports or home videos, and prospective studies of high-risk infants (i.e., those who have older siblings with ASDs) who are later diagnosed with ASDs. Unfortunately, domain-specific accounts of ‘mindblindness’ in ASDs [156] encounter developmental constraints, because it cannot be detected before 4–5 years of age, when mentalizing, or the ability to infer the contents of other people’s beliefs, intentions, or emotions, first emerges in typical development. Thus, while evidence suggests that mentalizing, or theory of mind, is delayed [157] and remains impaired across the lifespan in ASDs [158], this construct is not measurable during the early stages of human development, preventing any direct comparisons between the two accounts of SC dysfunction in ASDs. Likewise, the executive dysfunction accounts of SC impairment in ASDs are inherently limited by age, given the protracted development of executive functions across the human lifespan.

Highlights

With increasing prevalence, ASDs present a major public health challenge. Many neuroimaging findings indicate that the ‘social brain’ is organized atypically in ASDs.

Growing evidence shows that neural bases of ASDs cannot be pinpointed to specific regions of the brain, but that symptomatology is instead linked to atypical connectivity within and between functionally specialized brain networks (including ‘social brain’ networks). However, few neuroimaging findings have been widely replicated and no clear picture of the brain bases of sociocommunicative impairments in ASDs has emerged.

A main factor that has prevented consensus findings is etiological diversity. While diagnostic criteria focus on social deficits in ASDs, these probably result from the convergence of many different neurodevelopmental trajectories and many different causative factors.

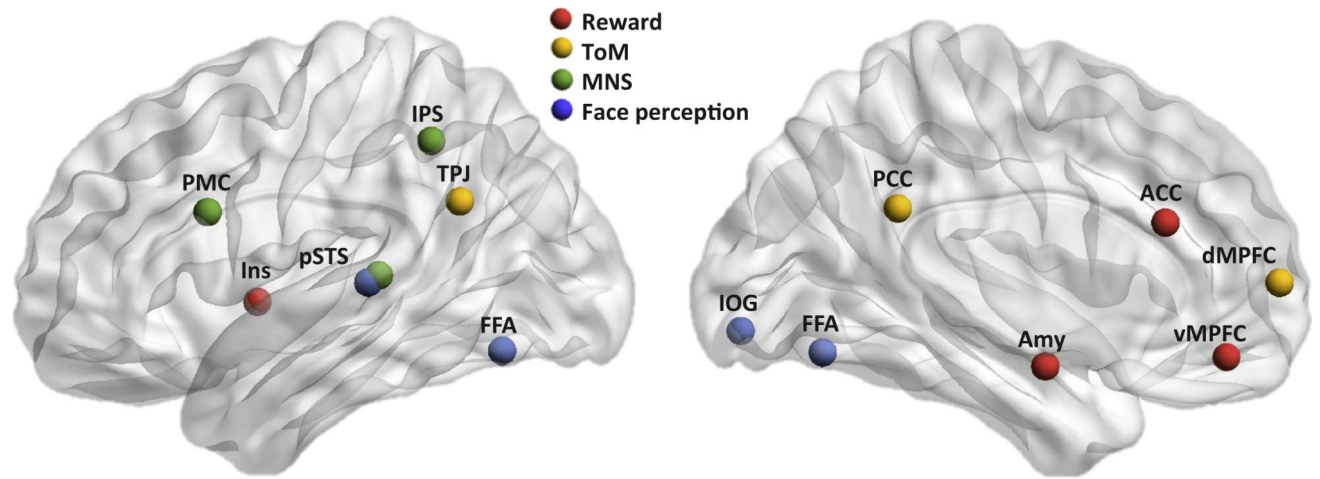
Outstanding Questions

Does atypical neurofunctional organization of social networks in ASDs reflect causation or compensation? The answer for most findings is 'probably both'. Disentangling the two is crucial for ultimate translational goals, because it will allow the distinction between atypical brain features that one may wish to ameliorate with brain intervention (or other) techniques versus features that may be promoted.

How do atypical maturational trajectories in ASDs affect comparisons with chronological age-matched TD groups? It is likely that connectivity in ASDs is atypical in ways that differ depending on maturational stage (e. g., infancy versus preteens versus adolescence); but there is little consensus on what these differences are. There may be no general pattern, but trajectories may differ across networks. In addition, heterogeneity and suspected existence of etiological subtypes are also likely to affect maturational trajectories in different ways, and there may be no single answer to this question that applies to ASDs across the board.

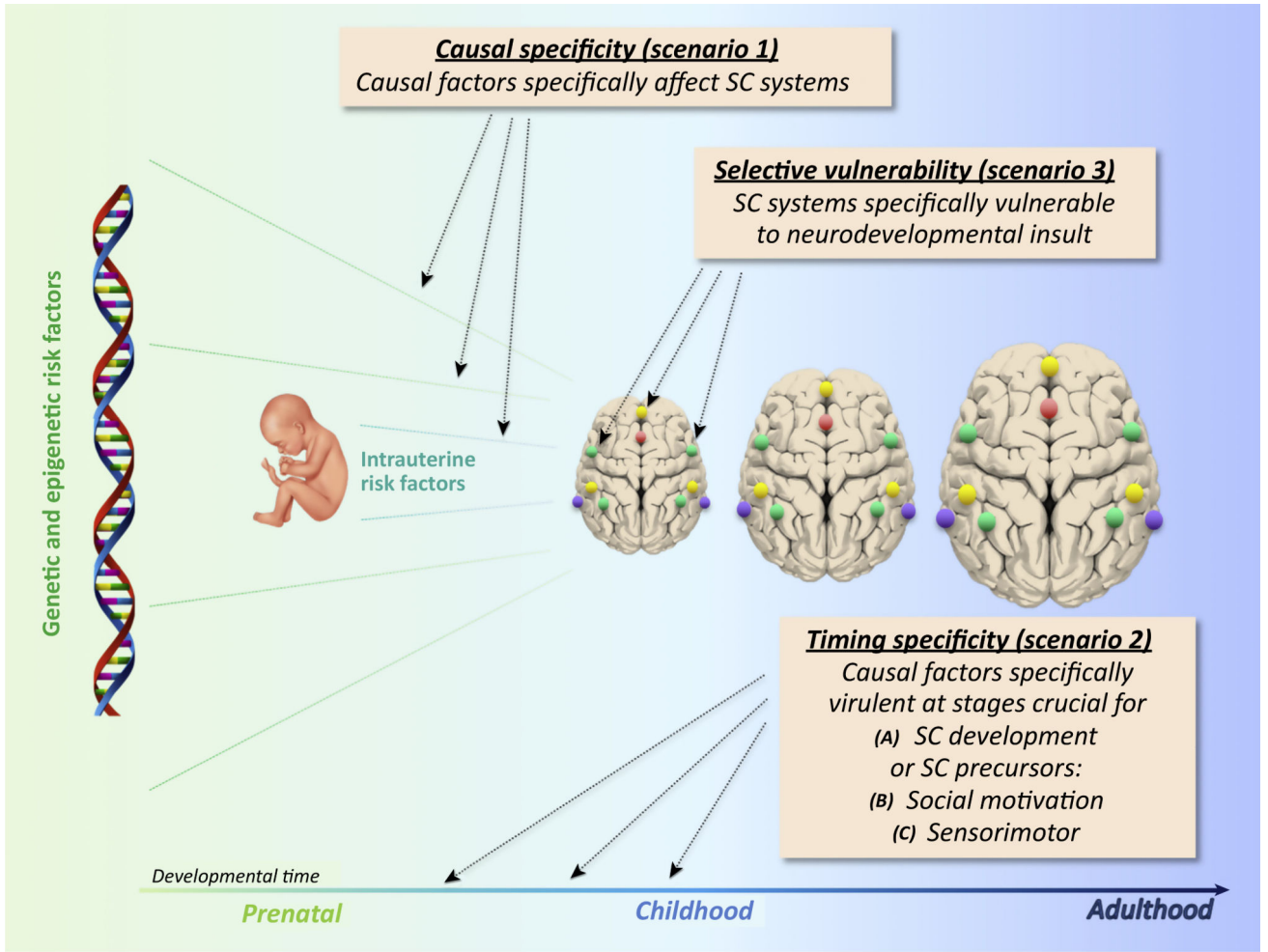
What does ASD heterogeneity imply for neuroimaging? The conventional focus on similarity within ASD groups, in comparisons with TD groups, should be redirected onto *differences*. Such focus can reveal subgroups in large samples that may reflect neurodevelopmental etiology more transparently than conventional group-level approaches.

Can machine learning improve our understanding of ASDs? Data-driven techniques are useful and can help in at least two ways: (i) They can identify brain markers that best distinguish ASD from TD participants (diagnostic prediction through supervised learning); and (ii) they can find patterns (clusterings) within large multivariate imaging data sets that may indicate brain-based variants of ASDs (through unsupervised learning).



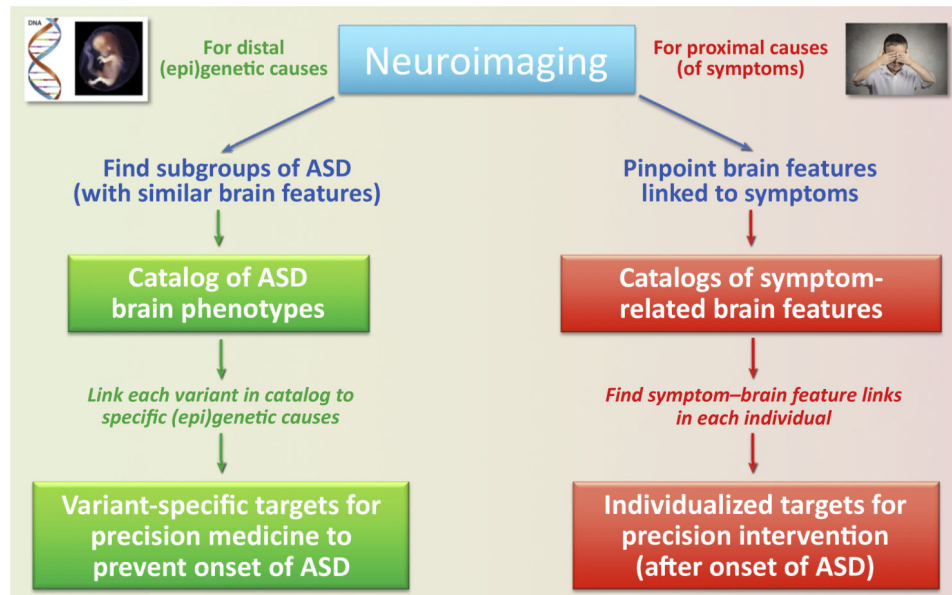
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Figure 1. Social Brain Networks. Core networks known to be involved in social cognition. Only main network regions, simplified as spheres, are shown with approximate location (excluding subcortical structures). Abbreviations: ACC, anterior cingulate cortex; Amy, amygdala; dMPFC, dorsomedial prefrontal cortex; FFA, fusiform face area; Ins, insula; IOG, inferior occipital gyrus; IPS, intraparietal sulcus; MNS, mirror neuron system; PCC, posterior cingulate cortex; PMC, premotor cortex; pSTS, posterior superior temporal sulcus; ToM, theory of mind; TPJ, temporoparietal junction; vMPFC, ventromedial prefrontal cortex.



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Figure 2. Convergence Scenarios. Cartoon of three convergence scenarios described in main text. Colored dots represent social network nodes diagrammatically. For anatomical locations see Figure 1. Abbreviation: SC, social communication.



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Figure 3.

Two Distinct Pathways towards Translational Application of Neuroimaging in Autism Spectrum Disorders (ASDs). Left: Since brain features contain information about developmental history, imaging can contribute to the search for distal (epi)genetic causes by providing a catalog of brain phenotypes, each of which may be linked to a specific etiology with specific targets for precision medicine. Right: Neuroimaging can reveal a catalog of proximal causes of ASD symptoms that require intervention. Specific links between brain features and symptoms at the level of the individual can inform targets for precision intervention. While causal neuroimaging may contribute to precision medicine that may prevent the onset of the disorder, it is also possible that such intervention would reverse critical neurobiological conditions after onset and diagnosis. Conversely, intervention targets developed from symptomatic neuroimaging may be developed for earliest signs before the onset of full symptomatology. Therefore, the timing implications of the two imaging pathways are not absolute.