
Concept Paper

A Molecular Framework for Autistic Experiences: Mitochondrial Allostatic Load as A Mediator between Autism and Psychopathology

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Abstract: Molecular research into Autism Spectrum Conditions (ASC) is undergoing an essential shift towards a biopsychosocial framework that is informed by autistic experiences. In this context, research aims are moving away from correcting external autistic behaviors and towards alleviating internal distress. ASC is associated with high rates of depression, suicidality and comorbid psychopathologies, but the underlying mechanisms that mediate this relationship are poorly understood. Here, we integrate emerging psychosocial characterizations of internal autistic experiences within a molecular framework to yield insight into the prevalence of psychopathology in ASC. We demonstrate that recent conceptualizations of social camouflaging and autistic burnout resonate closely with the accepted definitions for early life stress (ELS) and chronic adolescent stress (CAS). We propose that social camouflaging could be considered a distinct form of ELS that contributes to allostatic load, culminating in a pathophysiological state that is experienced as autistic burnout. Autistic burnout is thought to contribute to psychopathology via both psychosocial and neurophysiological mechanisms, but these remain largely unexplored by molecular researchers. Considering recent insights from converging bodies of work in molecular neuroscience, we discuss the substantial evidence implicating mitochondrial dysfunction in ASC to propose a novel role for mitochondrial allostatic load in the relationship between autism and psychopathology. An interplay between mitochondrial metabolism, neuroimmune and neuroendocrine signaling is increasingly implicated in stress-related psychopathologies and these molecular players are also associated with neurodevelopmental, neurophysiological and neurochemical aspects of ASC etiology. Together, this suggests an increased exposure to, and an underlying molecular susceptibility to, ELS that increases the risk of psychopathology in ASC. This article describes an integrative framework shaped by autistic experiences that highlights novel avenues for future research into molecular mechanisms that affect the quality of life and well-being of autistic individuals. Moreover, this framework emphasizes the need for increased access to diagnosis, accommodations, and resources to improve mental health outcomes in autism.

Keywords: autism; autistic burnout; social camouflaging; early life stress; suicidality; psychopathology; mitochondrial allostatic load

1. Introduction

Research into autism spectrum conditions (ASC) operates at the intersection of disparate psychosocial and biomedical disciplines that traditionally work in isolation rather than within cohesive interdisciplinary frameworks. However, the complex and rapidly evolving context of autism research calls for a novel integrative approach. The current conceptualization of ASC, used in both academic and clinical contexts, is based solely on how autism is perceived by external neurotypical observers rather than an understanding of internal autistic experiences (1). Therefore, much of the molecular research into ASC has been rooted in the assumption that effective therapeutic strategies would aim to

decrease autistic behaviors. However, as autism research is increasingly informed by inclusive participatory research practices, the field is moving towards a more comprehensive integration of the biopsychosocial disability model (2,3). This integrative model recognizes that while the biological (4–7) and behavioral (8–11) differences associated with ASC present challenges to independent functioning, the inability to function within a societal framework that accommodates only one neurotype is not indicative of an underlying biological deficit (12,13). Congruent with this, psychosocial autism research is shifting its focus towards improving quality of life and minimizing internal distress rather than eradicating or correcting autistic behaviors (14).

This paradigm shift is yet to be fully integrated into the molecular and medical fields of autism research, as these disciplines require unambiguous definitions of disease, reductionist animal and cell culture models, and highly specific questions with limited scope. The challenges of biomedical research are already compounded by the etiological heterogeneity of ASC (15,16) and the limitations of its conceptualization in diagnostic texts (1). These limitations are increasingly well-recognized in clinical settings (1), especially with respect to the reliability of autism assessments in females and people of color across varied sociocultural contexts (17–24). Yet, molecular studies require deeply-phenotyped cohorts characterized according to standardized diagnostic criteria in order to accurately distinguish between complex molecular signatures (25). Thus, it has been difficult for molecular researchers to move away from the deficits-based medical model of disability to integrate evolving understandings of autistic experiences into their research questions. In addition, much of molecular research uses limited measurements of behavior and flawed indicators of functioning. The efficacy of an intervention is often measured by a decrease in ASC diagnostic criteria but it is becoming clear that this is not a good indicator of health, well-being or quality of life for autistic individuals. In fact, the chronic suppression of autistic traits contributes to anxiety, depression, burnout, poor quality of life and poor mental health in ASC (26–29). It is now recognized that any therapeutic intervention should focus on improving physical and psychological well-being rather than decreasing ASC-associated behaviors, allowing for the fact that health may look different in autism. Importantly, the relationship between internal distress and the external presentation of autistic behaviors will reflect the variability that is inherent to all facets of ASC etiology. Molecular interventions that target biological or behavioral differences may certainly be able to improve quality of life. Still, it should be acknowledged that much of molecular research has aimed to decrease the severity of the autism phenotype without understanding that this is not synonymous with an improved quality of life.

This highlights a weakness in molecular autism research in that it is generally not informed by autistic experiences. Yet, molecular research is essential to provide accurate information about ASC to autistic individuals, their families and peers, clinicians, and the general public. Knowledge about the origins and manifestations of ASC can be used to inform the implementation of effective accommodations in schools and workplaces, as well as to combat the widespread misinformation and stigmatization surrounding autism. Additionally, clinicians and researchers are still grappling with the current diagnostic label that does not easily distinguish between ASC and co-morbidities, or reflect the heterogeneity in autism presentations, internal lived experiences or underlying biological processes. In conjunction with other psychosocial models, molecular research could be a useful tool to improve the resolution, specificity and utility of diagnostic labels. Ultimately, this could lead to a better understanding of individual autism presentations, their specific challenges, and relevant accommodations. Therefore, it is essential that molecular researchers grapple with the task of integrating the rigor of scientific experimental design with evolving understandings of autism in a biopsychosocial framework.

However, an interdisciplinary framework is not only needed drive autism research that is relevant, useful and transformative for the autistic community (3,14). An integrative approach also has the potential to yield novel insights into molecular ASC etiology. It is widely recognized that many promising findings from *in vitro* and *in vivo* molecular studies have failed to translate into reproducible results in human clinical trials (16,30–

34). This is largely attributed to the complex interplay between genetic, epigenetic and environmental mechanisms that contribute to ASC etiology. Furthermore, molecular research is inherently reductionist and grounded in a one-dimensional understanding of ASC that is removed from the internal experience of autism. This could be another prevailing limitation that hinders the translatability of molecular research in a clinical and sociological context (1). As such, an integrative approach is needed to highlight the convergence between different scientific disciplines involved in autism research. Moreover, this approach could provide new insight into the cellular mechanisms at the intersection of environmental stressors and biological processes that directly impact the quality of life and well-being of autistic individuals. In particular, integrating molecular research with recent characterizations of internal autistic experiences could be crucial to understand the high rates of psychopathology and suicidality in ASC.

Here, we incorporate recent conceptualizations of social camouflaging and autistic burnout within a scientific framework based on molecular neuroscience to suggest an increased risk of exposure to, and an underlying molecular susceptibility to, early life stress (ELS) and Chronic Adolescent Stress (CAS) in ASC. Interactions between psychosocial stress, underlying physiology and allostatic load have previously been proposed to influence clinical outcomes in autism (35,36). However, this model remains understudied by molecular researchers and emerging definitions for distinct autistic experiences have not yet been assimilated with recent discoveries pertaining to the biological mechanisms that mediate the response to ELS. This article proposes an integrative framework shaped by autistic experiences that demonstrates how psychosocial and biological factors converge to increase the prevalence of psychopathology in ASC. This framework emphasizes the need to improve access to diagnoses, accommodations, and resources to mitigate poor mental health outcomes in autism. Moreover, this highlights novel avenues for future research into molecular mechanisms that directly affect the quality of life and well-being of autistic individuals.

2. Psychopathology and Suicidality in ASC: the role of social camouflaging and autistic burnout

ASC is associated with a high prevalence of comorbid psychopathologies that substantially impair functioning and decrease quality of life (37–39). A growing body of literature demonstrates significantly increased rates of generalized anxiety disorder (GAD), major depressive disorder (MDD), post-traumatic stress disorder (PTSD), self-harm, and suicidal ideation in ASC (29,40,41). Between 63% (40) and 66% (42) of autistic adults are reported to be at risk of suicide; where suicide attempts are 3–4 times more frequent in ASC (43) and mortality by self-harm and suicide is significantly increased (39). Moreover, the onset of MDD occurs earlier in ASC; it was recently reported that 39% of 10–13 year-olds with autism reached clinically significant thresholds for measures of depression compared to only 6.5% of typically developing children (44). Current research suggests that psychopathology in ASC is associated with social difficulties (41), alexithymia (40), cognitive rigidity (45), loneliness and lack of social support (46) or increased exposure to traumatic experiences (47). Compounding this, there are unique challenges associated with the assessment of suicidal ideation and suicide risk given that alexithymia and communication difficulties are core components of ASC etiology (48). Moreover, it has been suggested that underlying biological mechanisms in ASC etiology could mediate a susceptibility to depression and other mood disorders (44,49). Thus, it is critical to understand the internal experiences and physiological mechanisms involved in this relationship in order to identify prevention strategies. Emerging research has begun to characterize a distinct phenomenon known as autistic burnout according to how it is experienced and defined by autistic individuals. This is an understudied, yet integral aspect of the internal experience of autism that could play a central role in distinguishing psychological health from psychopathology.

Recent studies have generated complementary definitions for autistic burnout, which is described as a “*a state of physiological and psychological incapacitation, exhaustion, and distress in every area of life*” (50). This is a recurring condition marked by loss of function, reduced tolerance to stimulus and an increased manifestation of autistic traits (51) that directly impacts “*functioning, mental health, quality of life and well-being*” (52). Autistic burnout is conceptually similar to non-autistic burnout in that both result from chronically operating beyond personal capacity. However, current research emphasizes that autistic burnout is distinct from both clinical depression and non-autistic burnout (50,51) with respect to its causes, manifestations and functional implications (53). Autistic burnout results from “*chronic life stress and a mismatch of expectations and abilities without adequate supports*” (50). The main contributor to autistic burnout is social camouflaging, defined as “*the need to suppress autistic traits or disability... in order to meet family, social, vocational, or other mainstream expectations*” (50). There are three core components of social camouflaging; i) deliberately adopting new social behaviors to make up for social difficulties (compensation) (26,54); ii) consciously suppressing autistic traits in order to appear non-autistic (masking); and iii) implementing strategies to manage internal discomfort invisibly (assimilation) (21). Chronic social camouflaging leads to internal distress, an eroded sense of self, and physical and mental exhaustion (26,55,56) and is associated with psychopathology (28), self-injury (57–59), and a higher risk of lifetime suicidality (29). However, camouflaging also facilitates access to employment, relationships and social acceptance and is an unavoidable prerequisite for independent functioning (27,57).

In fact, autistic individuals are required to exert substantial effort to perform typically “autonomic” aspects of everyday life. Consequently, autistic burnout is characterized by an inability to maintain basic levels of independent functioning, which is further exacerbated by the intensified presentation of autistic traits (51). Moreover, the inescapable demand to meet the neurotypical threshold for functionality makes recovery from autistic burnout much more difficult. Autistic burnout is described as “pervasive” (50), “chronic” (52), and “recurring” (51) which most closely aligns with a concept known as habitual burnout, when functional impairments become embedded into daily life (60). Notably, the symptoms of habitual burnout include insomnia, anxiety, behavioral dysregulation, depression, self-doubt, emptiness and isolation (60). The manifestations and functional implications of autistic burnout are thus highly specific to its context (53) and present a significant risk for the development of psychopathology. Both MDD and autistic burnout are associated with chronic fatigue, cognitive incapacitation, and suicidal ideation; however, the anhedonia and existential hopelessness that characterizes MDD is not always a defining feature of autistic burnout (51). While depression is described as a lack of *motivation* to participate in life, autistic burnout is experienced as a lack of *capacity* to do so (50). Nevertheless, autistic burnout can lead to the subsequent development of MDD and vice versa. Moreover, the struggle to survive in a constant state of burnout can seem objectively irrational, while the fundamental logistics of living become a physiological impossibility. Thus, autistic burnout represents a distinct, independent and fundamentally different path to suicidal ideation even in the absence of clinical psychopathology (50).

Notably, autistic individuals who remain undiagnosed throughout childhood are particularly susceptible to both autistic burnout and psychopathology due to an absence of adequate supports and the chronic suppression of autistic traits. Furthermore, undiagnosed autistic individuals lack an explanation for their differences and challenges and tend to internalize these as personal failings (56,61). A pervasive experience among late-diagnosed autistic individuals is a lifelong sense of “inherent wrongness”, alienation, and a deep sense of shame about who they are (62). This culminates in low self-esteem, a lack of self-worth, and destructive core beliefs that are all risk factors for the development of psychopathology (63). Altogether, this demonstrates how the pervasive lack of awareness and stigma surrounding ASC contributes to autistic burnout (52) and how early diagnosis could play an important role in preventing psychopathology. While the link between ASC and psychopathology is well-established, there is a critical gap in the literature on what

physiological factors lead to autistic burnout, how they differ from, or contribute to the etiology of MDD and crucially, how these could be prevented or treated. Moreover, the molecular mechanisms that mediate the relationship between ASC etiology, autistic burnout and psychopathology are unknown and remain largely unexplored by molecular researchers. However, we propose that converging bodies of work in molecular neuroscience highlight a potentially important connection between early life adversity, allostatic load and psychopathology in ASC.

3. Early life stress, chronic adolescent stress and allostatic load in ASC

There is substantial evidence from both human epidemiological studies and animal models that ELS and CAS increase the risk of psychopathology and suicidality in adulthood (64–70). Notably, the established definitions for ELS and CAS closely mirror those emerging to characterize autistic experiences of childhood and adolescence, which points to the relevance of molecular ELS research in the context of ASC. While it has historically been challenging to reach a consensus for the definition of ELS or CAS, current models agree that stress refers to *“environmental events or chronic conditions that objectively threaten the physical and/or psychological well-being of an individual”* (65), undermine a child’s *“sense of safety, stability, and bonding”*(71) and for which *“adequate coping resources are unavailable”* (66). While much of the literature focuses on adverse childhood experiences, adolescence is also a critical period for brain development that is highly sensitive to stress, and CAS is known to contribute to the development of depression, anxiety, and other stress-related psychopathologies (65). Researchers have previously distinguished between good, tolerable stress and toxic stress; these differentially induce adaptation, resilience or pathology respectively. Here, the defining feature of toxic stress is *“a lack of internal resources or external support systems, resulting in chronic physiological dysregulation”*(72).

Importantly, cumulative evidence suggests that autism is associated with a significantly increased exposure to toxic stress in childhood and adolescence. Children with ASC are statistically more likely to experience early adversity (73), social vulnerability and isolation (74), bullying, discrimination, and ostracism (75), and other forms of victimization (76). Differences with respect to social communication, development, and integration can make it difficult for autistic children to feel a sense of a sense of *“safety, stability or bonding”* with their families or peers. Moreover, situations that are *“tolerable”* for neurotypical children are experienced as *“toxic”* stressors due to distinct neurophysiological differences in autism. Exposure to every-day sensory stimuli is painful, uncontrollable, and thus *“personally threatening”* due to difficulties with sensory processing. Personal accounts of social camouflaging describe a need *“to exceed what nature has given”* (26) in order to manage both the *“physically assaultive”, “uncertain, exhausting nature of the social environment”* (77), and the persistent fear of stress, rejection and being misunderstood by peers (78). In this context, autistic children are pushed beyond their capacity just to achieve a degree of social participation; even further resources are expended to camouflage sufficiently to achieve social acceptance. Accordingly, social camouflaging has been conceptualized as the discrepancy between the internal experience of autism and the external presentation of autistic traits (21,79,80), while autistic burnout has been described as *“having all of your internal resources exhausted beyond measure”* (50). Thus, the literature consistently points to a mismatch between external demands and internal capacity, and clearly demonstrates an absence of *“adequate coping resources”* which is consistent with the accepted definitions for toxic stress.

Since camouflaging strategies become increasingly complex and intentional from middle childhood, it could be argued that social camouflaging acts as a distinct form of CAS in autism. Middle childhood is a critical period for socialization that is marked by the development of self-concept and an increasing capacity for self-regulation (81). The self-concept tasks of middle childhood culminate in the ability to *“take the self as an object and view it as others do”*, internalize expectations for one’s own behavior and develop strategies to meet these expectations (81). This enables an autistic child to both identify the

need to suppress autistic traits and develop enough self-regulation to implement camouflaging as a replacement for natural behaviors. The extent to which these tasks are completed will vary widely across the autism spectrum depending on differences in both underlying biology and sociocultural factors that contribute to socialization. Equally, there are significant differences with respect to an individuals' ability and motivation to mask their autistic traits. This has important implications for the relationship between ASC etiology, chronic stress and psychopathology. What has previously been conceptualized as a more severe autism phenotype – presenting with extreme sensory sensitivities, greater difficulties with self-regulation and social communication or atypical developmental profiles - might be associated with a lower tendency to mask but an increased exposure to early childhood adversity. On the other hand, a greater tendency to mask may be marked by a subclinical phenotype but a greater susceptibility to CAS induced by the chronic suppression of autistic traits, which may explain the increased rates of depression and suicidality in high-masking autistic individuals. This emphasizes that different external presentations of autism are not always reflective of the profound physiological and psychological dysregulation experienced internally. In fact, despite the etiological heterogeneity of ASC, an underlying susceptibility to stress-related pathologies could be a common mechanism responsible for negative clinical outcomes in autism.

Furthermore, the resonance between descriptions of social camouflaging, autistic burnout, and toxic stress serves as a point of convergence between psychosocial and molecular research into ASC. From a molecular standpoint, ELS and CAS are thought to contribute to allostatic load, which describes the pathophysiological consequences of chronic stress on the metabolic, endocrine and immune systems required for homeostasis (82). Allostatic load provides an integrative model with which to quantify the cumulative impact of biological, psychological and social factors that contribute to pathology (82). In this context, social camouflaging could act as a distinct form of toxic stress that contributes to allostatic load in autism. Autistic burnout may represent a distinct pathophysiological state resulting from allostatic overload, which increases the risk of psychopathology via both physiological and psychosocial mechanisms. In fact, Singletary (2015) previously recognized that autistic children are exposed to significant psychosocial and biological stress during early life and proposed an interaction between ELS, underlying physiology and allostatic overload contributing to ASC etiology (35). Scarpa *et al.* (2021) further proposed that underlying features of ASC etiology also predispose autistic individuals to pathological responses to stress and trauma (36). The recent characterization of unique autistic experiences that closely overlap with the accepted definitions for ELS and CAS points to the need for a more integrative framework to characterize interactions between neurobiology, environmental stressors and allostatic overload in the context of ASC. Notably, the molecular mechanisms that mediate the association between ELS, CAS and psychopathology are also known to contribute to ASC etiology. This could signify an underlying molecular susceptibility to ELS, that may yield mechanistic insight into the relationship between autistic burnout, psychopathology and suicidality.

4. Mitochondrial allostatic load and convergent molecular mechanisms in ELS, ASC, and psychopathology

There are well-characterized neurobiological and psychological aspects of ASC etiology that predispose autistic individuals to stress-related pathologies (36). However, the molecular mechanisms that mediate this susceptibility are complex and poorly understood. While the concept of allostatic load is well-established(83–85), the field has not yet described specific molecular mechanisms that mediate pathological stress responses, nor identified reproducible molecular biomarkers to measure allostatic load(86). The Hypothalamic-Pituitary-Adrenal (HPA) axis has been the focus of stress pathophysiology research in ASC, but the current evidence for HPA axis dysregulation in autism is inconsistent(87). Recently, mitochondrial metabolism has been recognized as a central regulator of the major neuroendocrine and neuroimmune systems involved in allostasis (82). In

particular, emerging research highlights the relevance of mitochondrial allostatic load (MAL) in stress pathophysiology (85). MAL refers to adaptive changes to mitochondrial morphology, dynamics, and function that are observed in response to chronic stress. Notably, mitochondrial dysfunction is a biological signature that is not only consistently observed in idiopathic ASC but also independently implicated in ELS and psychopathology. Thus, MAL could serve as a central molecular mechanism involved in the development of psychological disorders in autistic individuals.

4.1. Mitochondrial allostatic load

A growing body of research suggests that mitochondrial dysfunction is involved in the relationship between ELS and psychopathology. Clinical studies show that ELS is associated with indicators of oxidative stress, including decreased levels of glutathione peroxidase and an increase in superoxide dismutase, protein carbonylation, and total reactive antioxidant potential (88). In line with this, several preclinical models for ELS show an increase in protein carbonylation and a decrease in superoxide dismutase and catalase activity in brain tissue (89) and increased reactive oxygen species (ROS), mitochondrial glutathione, and cytochrome c release in cardiac tissue (90). Ruigrok et al., (2021) recently showed that ELS led to impaired ETC activity in the hypothalamus and altered the expression of genes involved in mitochondrial fission and antioxidant defense in the hippocampus, leading to cognitive impairments(91). Moreover, a recent systematic review presented consistent evidence for reduced mitochondrial energy production capacity and electron chain (ETC) complex activity, altered mitochondrial morphology and changes in mitochondrial DNA (mtDNA) copy number in animal models for chronic stress (92). Notably, the duration and type of stress, as well as underlying genetic differences significantly affect how mitochondrial function is altered in response to stress(92). Conversely, experimentally disrupting mitochondrial function has been shown to alter the physiological and behavioral consequences of psychological stress(82).

Similarly, extensive evidence suggests a role for mitochondrial dysfunction in the etiology of MDD, which has recently been comprehensively reviewed (85,93–98). Mood disorders are frequently reported in patients with mtDNA mutations and up to half of patients with primary mitochondrial diseases present with MDD; conversely, depression is associated with elevated mtDNA deletions, aberrant mtDNA copy number and oxidative mtDNA damage (95,99). Numerous independent studies have reported increased oxidative stress and altered ETC activity (100) and decreased levels of antioxidants and antioxidant enzymes (101) in the brain tissue of patients with MDD. Of note, Karabatsiakos *et al.* (2014) reported significantly impaired mitochondrial respiration in patients with MDD which correlated significantly with the severity of depressive symptoms (102). Pre-clinical studies also consistently demonstrate an increase in ROS and a decrease in ATP and glutathione in models for depression (95). Importantly, many pre-clinical models for MDD are induced by exposure to forms of ELS, and such models consistently implicate mitochondrial mechanisms in the subsequent development of depressive behaviors (98). *In vivo* models have shown that mitochondrial ETC complex activity is disrupted in brain tissue following maternal deprivation (103), chronic restraint stress (104), acute restraint stress(105), social isolation (106), chronic mild stress (107) and chronic unpredictable stress (108). Moreover, several transcriptomic profiling studies have reported significant dysregulation of mitochondrial genes involved in lipid metabolism, oxidative phosphorylation (OXPHOS) and glucose homeostasis in animal models for chronic and acute stress (96,109–111). Thus, both human epidemiological studies and animal models suggest a role for mitochondrial dysfunction in the development of MDD after ELS. In fact, Allen, Caruncho and Kalynchuk, (2021) recently reviewed the central role of MAL in the development of psychopathology following ELS, highlighting how mitochondrial dysfunction contributes to the symptoms of MDD and emphasizing the utility of targeting mitochondrial function to develop novel pharmacological treatments (85).

The role played by MAL in the response to ELS is particularly relevant in the context of autism, given that mitochondrial dysfunction is also a well-established component of ASC etiology. A 1998 review of ASC clinical data first demonstrated decreased glucose uptake and ATP synthesis in the cortex, as well as lactic acidosis, carnitine deficiency and elevated urine levels of Krebs cycle metabolites (112). Since then, multiple lines of evidence have emerged to support a role for mitochondrial dysfunction in ASC, which have been well described in recent reviews (113–119). Of note, clinical studies have reported altered mtDNA copy numbers and deletions (116,120–123), as well as mtDNA mutations (124,125), mitochondrial ETC defects (120,122,126) and altered plasma levels of lactate, pyruvate, alanine, creatine kinase, glutathione-S-transferase and caspase 7 (124,127). Recent studies in ASC-derived lymphoblastoid cell lines (LCLs) showed significantly increased mitochondrial respiration and mitochondrial membrane potential as well as elevated activities of ETC complex 1,3 and 4 (128), while disruptions to mitochondrial bioenergetics, dynamics and morphology were observed in ASC-derived fibroblasts (129,130). Moreover, multiple independent transcriptomic studies have reported down-regulated expression of genes involved in mitochondrial respiration in ASC brain tissue (131–134), while ASC transcriptomic, proteomic and DNA methylation functional enrichment signatures consistently converge on mitochondrial OXPHOS (135). Furthermore, increased oxidative stress is a well-documented aspect of ASC etiology, and several recent reviews have described the cumulative evidence for altered glutathione metabolism, lipid peroxidation, protein oxidation, DNA oxidation and antioxidant enzyme activity in ASC (136–141). Therefore, there is substantial evidence for altered mitochondrial metabolism in ASC, which may serve as a molecular susceptibility to the pathophysiological consequences of ELS. In line with this, emerging research highlights mitochondria as essential modulators of both the neuroendocrine and neuroimmune components of allostasis.

4.2. MAL and the HPA axis: interdependent mechanisms in stress-related psychopathology

The HPA axis is the major neuroendocrine system that initiates the stress response and plays a central role in coordinating molecular components of allostatic load (65). Children are particularly vulnerable to HPA disruption during periods of higher HPA axis plasticity. These periods occur between 0-5yrs old and during adolescence; thus, HPA signaling is a key mediator of pathological responses to ELS (66,142) and CAS (65). Recent systematic reviews have demonstrated that HPA axis dysregulation plays an important role in mood disorders, suicidal behavior(143), and in mediating the relationship between ELS and major depression(144). Transcriptomic studies in brain tissue indicate that HPA axis signaling is involved in the development of MDD following ELS exposure (64,145). This is reflected by altered cortisol and catecholamine responses in individuals with a history of ELS who develop PTSD, depression and suicidality(146). Finally, recent work in animal models has demonstrated distinct alterations to HPA axis signaling, stress reactivity, and cognitive and emotional functioning in response to both ELS (64) and CAS (147).

Importantly, there is a growing body of literature demonstrating a bidirectional interaction between the HPA axis and mitochondrial metabolism (65). Mitochondria are directly responsible for the synthesis of stress-responsive progestogens, mineralocorticoids, androgens, estrogens and glucocorticoids (GCs) like cortisol and corticosterone (148). Moreover, mitochondrial metabolism not only regulates, but is also modulated by, GC signaling(82,149); this is comprehensively reviewed by Kokkinopoulou and Moutsatsou (2021) (150). GCs are released in response to HPA axis activation and can cross the blood-brain barrier to act on GC receptors (GRs) and regulate genes involved in neurogenesis, neuroplasticity, and neurotransmission (151). *In vitro* studies have demonstrated that GCs improve mitochondrial oxidation, membrane potential, and calcium buffering capacity at physiological levels, while supraphysiological levels impair mitochondrial function (152). More recent work shows that chronic GC treatment reduces the activity of specific mitochondrial ETC complexes and increases mitochondrial ROS production and mitochondrial fragmentation (153–155). GCs can influence mitochondrial metabolism by regulating

the expression of relevant genes via GC response elements (GREs) in the nucleus and potentially via the three known GREs in mtDNA to influence the expression of mitochondrial genes (150,156,157). In particular, GCs have been shown to increase the expression of peroxisome proliferator-activated receptor gamma coactivator-1 alpha, mitochondrial transcription factor A, and nuclear respiratory factors 1 and 2 which are central transcriptional regulators of mitochondrial biogenesis (149,157,158). Notably, animal models have shown that both chronic stress and corticosterone treatment increase mtDNA copy number (159), and conversely, that mtDNA genetic variants can alter corticosterone production and HPA signaling (160,161). Moreover, human epidemiological studies have shown that mtDNA copy number is elevated in people with a history of ELS (162), and in major depression (159) and recent work has demonstrated that the positive relationship between ELS and mtDNA copy number was mediated by differential methylation of the GR (NR3C1) (162). Altogether, this highlights a mechanistic relationship between HPA axis activation and mitochondrial biogenesis, dynamics and function that points an intersection between MAL and well-established stress-responsive processes involved in psychopathology.

Thus, both mitochondrial function and GC signaling are independently implicated in the response to ELS and the development of MDD. Moreover, recent data indicate that these mechanisms are interdependent, which may reveal a novel molecular susceptibility to ELS that has been overlooked in the context of ASC (87). Several candidate autism risk genes are known to increase the risk of developing mental illness after exposure to ELS (82,163–166) but the associated mechanisms are unclear. Although much research has investigated a role for stress-responsive HPA axis signaling in the relationship between autism and psychopathology, this has yielded inconsistent results. Several studies have reported HPA hyperactivation (167–170) and increased stress reactivity (171) in ASC, but these findings lack intra- and inter-individual reproducibility (167). Many studies assess HPA function using single end-point measurements of salivary cortisol, but these cannot reflect complex interactions between biological signaling processes and genetic architecture with sufficient resolution. More targeted molecular studies have shown disruptions to GR expression in ASC brain tissue (172); while genetic polymorphisms in the GR co-chaperone FKBP Prolyl Isomerase 5 (FKBP5) were associated with serum cortisol levels in ASC (173) and moderated the relationship between autistic traits and social anxiety (174). However, a recent review reported conflicting evidence for HPA dysregulation, and its association with psychopathology, in ASC (44). Thus, the relationship between HPA axis signaling and psychopathology in autism is complex and likely mediated by pleiotropic biological factors that contribute to an increased susceptibility to HPA axis dysregulation (175). In this context, the emerging relationship between mitochondrial metabolism and HPA axis stress-response signaling highlights mitochondrial dysfunction as a common biological signature that is consistently observed in idiopathic ASC and could act as a novel susceptibility to stress-related psychopathology.

4.3. The neuroimmune system: a molecular mediator between MAL and the HPA stress response

The neuroimmune system is the second established component of allostatic load that works in concert with HPA neuroendocrine signaling to modulate the stress response. A dysregulation of neuroimmune homeostasis is well-documented in human clinical studies of neuropathology and is observed in response to ELS (66,176,180) and in several psychopathologies (181–183). In fact, it has been proposed that the neuroendocrine-immune network is a central molecular mediator between the psyche, psychosocial stress and psychopathology (185). Accordingly, preclinical models have demonstrated that chronic stress consistently induces microglial activation and proinflammatory cytokine signaling in key brain regions involved in psychiatric disorders (176). This disrupts microglial proliferation and phagocytic capacity, leading to impaired neuronal development and functional deficits in cognitive performance, memory, reward processing, and processing of social stimuli (177). Furthermore, recent animal studies showed that ELS-induced

depressive behaviors were associated with neuroinflammation (89) and aberrant microglial activity (178), while CAS was also shown to increase hippocampal immune reactivity (179). Notably, there is also extensive evidence for neuroinflammation and gliosis in ASC (140,184–189), including reports of elevated inflammatory cytokines in ASC blood and post-mortem brain tissue, as well as immunohistochemistry, positron emission tomography and morphological data that demonstrates glial over-proliferation and over-activation in ASC (190–195). A seminal transcriptomic study in 2011 reported significant differential expression of genes involved in inflammation, astrocyte function and microglia activation in ASC brain tissue (196) and the glial fibrillary acidic protein (GFAP) was shown to be overexpressed at both the mRNA and protein levels in ASD post-mortem brain tissue (197). Post-mortem studies have also reported a significant upregulation of microglial-specific genes (194), increased microglial density (192,198) and excessive microglial activation (193,199) in ASC brain tissue and two recent systematic reviews have found consistent evidence for dysregulated microglial morphology, organization and activation in autism (188,200). Neuroimmune signaling therefore denotes another mechanistic overlap between ASC neurophysiology and the pathophysiological response to ELS.

In fact, the neuroimmune system also serves as an intermediary between MAL and the HPA axis on a molecular level. Mitochondrial metabolism is a central modulator of glial cell phenotypes and neuroinflammatory state (82) while the bidirectional relationship between inflammatory and GC signaling has been well-documented in the context of MDD and other psychopathologies (201,202). Glial functions are controlled by tightly regulated interactions between microglia, astrocytes, oligodendrocytes and neurons, which are closely coupled to glial metabolism(203). Importantly, glial phenotypes are dynamic, shift in response to different stimuli and can be either neuroprotective or neurotoxic (204). The transition to an inflammatory phenotype under pathogenic conditions is known as reactive gliosis, which is characterized by transcriptional, biochemical, metabolic and morphological remodeling (205). This has distinct implications for the metabolism, physiology and function of different glial cell types (206–211), each of which are implicated in specific aspects of ASD etiology (184,188,212–215). In particular, the tight coupling of mitochondrial metabolism and microglial activation is highly relevant in the context of ELS and ASC. Microglia are the only glial cells present in the early embryonic brain (216) that play an essential role in the regulation of phagocytosis, synaptic pruning, neurogenesis, neuronal excitability, synaptic activity and axonal myelination (211,217–219), all of which are thought to be dysregulated in autism (184,220,221).

Mitochondrial dysfunction can induce microglial activation via both extrinsic and intrinsic mechanisms. Mitochondrial dysfunction in surrounding tissue leads to the release of damage-associated molecular patterns (DAMPs) that bind to microglial receptors (117). Moreover, microglial activation is marked by intrinsic metabolic reprogramming, and different microglial phenotypes are associated with distinct metabolic pathways (218). While resting microglia rely primarily on OXPHOS (222), the shift towards the inflammatory M1 phenotype is driven by a decrease in mitochondrial respiration and an upregulation of glycolysis that enhances flux through the pentose phosphate pathway and increases lactate production to meet increased cellular energy demands (204). Recent work has demonstrated that mitochondrial metabolism directly regulates microglial activation (223,224) and that disrupting mitochondrial function has profound implications for microglial inflammatory state (217,218,224,225). *In vitro* models have shown that inhibiting mitochondrial ETC activity (226) or increasing mitochondrial fragmentation (218) induces microglial activation and pro-inflammatory cytokine signaling which could be mitigated by inhibiting mitochondrial fission (227) or normalizing mitochondrial membrane potential and ROS (218). On the other hand, mitochondrial stress impaired the transition to the anti-inflammatory M2 phenotype (228). Thus, mitochondrial dysfunction exacerbates the proinflammatory M1 phenotype leading to the release of neurotoxic inflammatory cytokines and oxidative stress (229). Conversely, exposure to changing cytokine stimuli has been shown to alter microglial metabolic state. Inflammatory stimuli were found to induce a metabolic switch from OXPHOS to glycolysis in cultured microglia (230),

marked by increased lactate production and decreased OXPHOS, mitochondrial oxygen consumption and ATP synthesis (231,232). This metabolic shift correlated with increased production of proinflammatory cytokines (233) and nitric oxide (230), highlighting the close coupling of mitochondrial metabolism, microglial phenotypes and neuroinflammation.

Stress-sensitive neuroendocrine mediators including both catecholamines and GCs are also known to modulate immune system responses (82) by suppressing leukocyte trafficking and activation, as well as inflammatory cytokine production (202). Conversely, inflammatory cytokines modulate GC signaling by decreasing GR expression, altering GR phosphorylation or disrupting GR nuclear translocation and DNA binding (234). Cytokine-stimulated immune cells can also mediate HPA axis activation by inducing CRH production in the hypothalamus, leading to the secretion of adrenocorticotrophic hormone (ACTH) and cortisol by the pituitary gland and adrenal cortex respectively (158). Of particular relevance, interleukin 6 (IL6) has been shown to activate HPA axis signaling and stimulate GC secretion in animal models and conversely, cortisol is known to inhibit the peripheral production of IL6 (158). IL6 is one of the most consistently upregulated inflammatory markers associated with ELS (180), MDD (183) and ASC (235) and recent pre-clinical studies have demonstrated that neuroimmune and HPA neuroendocrine interactions may mediate the neuroinflammatory response to ELS. ELS was shown to increase GR promoter methylation and decrease GR expression, which was associated with upregulated proinflammatory cytokine signaling (158). Similarly, chronic stress was shown to disrupt GR signaling on a transcriptomic level, while increasing microglial activation (236). Considering that GR signaling regulates mtDNA gene expression and mitochondrial functions - and that the mitochondria are central to immune modulation - it has been proposed that HPA-immune signaling could be mediated by mitochondrial mechanisms (82). Notably, several recent studies demonstrated that ELS induces sex-specific changes in both mitochondrial metabolism and markers of neuroinflammation (237–239), which supports a role for MAL in mediating the neuroinflammatory stress responses that lead to psychopathology. Altogether, this points to a tightly coupled interplay between mitochondrial metabolism, neuroimmune and neuroendocrine signaling that underlies the established relationship between ELS, MAL, and psychopathology. This discussion has highlighted that both mitochondrial dysfunction and neuroinflammation are well-established hallmarks of ASC etiology that each play central roles in modulating the HPA axis stress response. Thus, these molecular signatures could serve as an underlying susceptibility to ELS that may contribute to the high rates of psychopathology in autism.

4.4. From MAL to neurophysiology, neurochemistry, and behavior: a mechanistic perspective

A role for MAL in the relationship between autism and psychopathology is supported by the mounting evidence that points to mitochondrial metabolism as a key regulator of neurodevelopment, neurophysiology and neurochemistry. In fact, mitochondria modulate distinct neurological processes that are disrupted in response to ELS and involved in the etiology of both ASC and psychopathology. ELS has been shown to dysregulate the proliferation, differentiation and survival of neuronal stem cells (NSCs) (64), culminating in disruptions to synaptogenesis, synaptic pruning, long-term potentiation, and the balance between excitatory and inhibitory synapses (146). ELS-induced disruptions to neurodevelopment, synaptogenesis, synaptic plasticity and myelination are thought to contribute to the cognitive and behavioral differences observed later in life (240). Extensive synaptic remodeling also occurs during adolescence and CAS has been shown to alter neuroplasticity with implications for psychology and behavior throughout adulthood (65). These neurophysiological differences have previously been attributed to HPA axis signaling, as GCs are known to modulate NSC fate and neurogenesis via interactions with the brain-derived neurotrophic factor (BDNF), phosphoinositide 3-kinase (PI3K)/Akt, hedgehog, and Wnt pathways (241,242). However, just as the concept of MAL has become

increasingly well-established in stress research, mitochondria are also emerging as central players in the regulation of neurogenesis and NSC self-renewal (243–245).

Neuronal differentiation is closely tied to metabolic state, which is coupled to, and regulated by, mTOR signaling and this relationship plays an important role in mediating the metabolic shift that is required to facilitate neurogenesis (246–248). Notably, both mTOR signaling (246,249–251) and mitochondrial function (252–257) are essential for synaptic signaling, function and plasticity. Wang et al., (2020) recently showed that ELS impaired synaptic plasticity by inhibiting mTOR signaling in the mouse hippocampus, resulting in increased anxiety-like behavior and cognitive impairments (258) while Sanchez et al., (2022) also demonstrated that ELS induced hyperactivation of the mTOR complex and reduced oxidative metabolic capacity in the amygdala (259). This suggests that the mTOR-dependent modulation of NSC metabolic state contributes to the neurophysiological, cognitive and behavioral changes induced by ELS. As an important regulator of microglial activation, mitochondrial metabolism also modulates both neuronal apoptosis and synaptic pruning during neurodevelopment; moreover microglial metabolism is essential to regulate myelin processing in the adult brain(207). Mitochondrial dysfunction is known to disrupt myelination by decreasing the myelinating potential of oligodendrocytes, increasing oxidative damage and disrupting neuronal fatty acid metabolism (260). Myelin is used as an alternative energy source if glycolysis is inhibited (207) and the physiological turnover of myelin is dependent on microglial phagocytic efficiency. In fact, microglial dysfunction has been shown to precede major myelin breakdown in hereditary demyelinating disorders(211). These data are highly relevant in the context of ASC which is also characterized by disruptions to mTOR signaling (251,261,262), synaptic pruning(200,263) and myelination (214) leading to cortical hyperconnectivity and the resultant behavioral phenotypes (219,264). Thus, the mitochondrial modulation of neurodevelopment, synaptogenesis and myelination manifests in neurophysiological features that are consistently associated with ELS and are well-established components of ASC etiology.

In addition, both ELS and ASC are marked by disruptions to neurotransmission and neuroendocrinology. Preclinical models have demonstrated that both ELS and CAS disrupt glutamatergic and GABAergic neurotransmission, leading to impaired social recognition, decreased social interest, reduced cognitive function, cognitive inflexibility and increased anxiety-like behaviors (265–268). ELS has also been shown to disrupt both serotonin and dopamine signaling in animal models (238). Several preclinical models have shown that ELS alters the mRNA expression of dopamine receptors, the density of dopamine transporters and the rate of dopamine metabolism and turnover in brain tissue (269). Moreover, a meta-analysis of rodent studies found robust associations between ELS exposure and specific dopamine metabolites in the striatum (270). Similarly, Adjimann, Argañaraz and Soiza-Reilly, (2021) recently summarized the current evidence from animal models suggesting that serotonin plays a central role in the development of psychopathology in response to ELS (271). These data are supported by human cohort studies which have shown that ELS alters glutamate/glutamine cycling (272), dopaminergic neurotransmitter responses(273,274) and serotonin transporter binding (275) in patients who presented with MDD, while serotonin receptor genetic polymorphisms have been shown to mediate the effect of ELS on the subsequent development of depression (276). CAS also alters neuroendocrinology on several fronts and has been shown to induce glutamatergic excitotoxicity and decrease levels of serotonin, basal dopamine and norepinephrine (65). Notably, an unbalanced excitatory-to-inhibitory synaptic signaling ratio (163,166,277) and disruptions to glutamatergic (166,215,278–282), serotonergic (283–287) and dopaminergic (163,165,288) neurotransmission are well-established aspects of ASC neurochemistry that are associated with distinct aspects of ASC etiology. Glutamatergic excitotoxicity is thought to contribute to anxiety, perseveration, restlessness, migraines, tics and motor stereotypies, social interaction and social memory, and language and cognitive impairments (282). Disruptions to serotonin signaling are linked to altered social cognition, facial recognition, emotion processing and communication (286) while dopamine signaling is associated with disruptions to sleep, mood and attention (163,165,288).

This underlying neuroendocrinological dysregulation is not only associated with core facets of ASC etiology, but also has direct functional implications for stress-related psychopathology. A dysregulation of glutamatergic and GABAergic signaling contributes to the etiology of anxiety disorders (328–331), PTSD (332), eating disorders (333–335), OCD (336) and substance use disorders (SUDs) (337,338). The serotonergic, dopaminergic and noradrenergic systems are implicated in MDD, anxiety and SUDs (339–341) while serotonin is also associated with mood disorders, aggression, anti-social conduct, OCD, stress disorders and ADHD (342–344) and dopamine is thought to play a role in eating disorders (345,346), executive dysfunction and behavioral inhibition (347,348). Moreover, well-documented interactions between neurotransmitter, neuroimmune and neuroendocrine pathways play a central role in modulating suicide risk after ELS exposure (143). Glutamate is involved in both driving HPA responses and limiting HPA overactivation (318–320) and conversely, GC signaling is known to modulate glutamatergic synapse plasticity and excitability (321). Serotonergic neurotransmission excites CRH neurons in the amygdala (183) and acts as an important regulator of GR signaling in the PFC after exposure to acute stress (322). On the other hand, animal models show that corticosteroids decrease serotonin receptor binding densities (323,324) and this relationship is implicated as a mediator of suicide risk following ELS (276). Finally, GCs also activate the mesolimbic dopamine pathway (325,326) by upregulating the rate limiting step of dopamine synthesis, downregulating dopamine degradation, clearance and synaptic uptake, or by acting directly on GRs in dopamine-receptive neurons (327).

Crucially, MAL functions as a mechanistic link between the molecular, neurochemical and behavioral aspects of ASC that have been highlighted here as potential contributors to psychopathology. Separate reviews have comprehensively described the close coupling of mitochondrial function with glutamatergic (289–291), serotonergic (292–294) and dopaminergic (295–299) neurotransmission, although this remains understudied in the context of MAL and psychopathology. Firstly, glutamatergic neurotransmission is tightly coupled to neuronal and astrocytic metabolism (300,301). Glutamate functions as a substrate for both the mitochondrial TCA cycle and the transsulfuration pathway, thereby fueling OXPHOS and antioxidant synthesis to promote mitochondrial function and mitigate against oxidative stress (290,301–304). Thus, mitochondrial metabolism is essential for glutamate-glutamine cycling, and cytoplasmic glutamate levels are regulated by mitochondrial TCA cycle flux as well as intracellular redox homeostasis (305). Dopamine autooxidation is also regulated by intracellular redox state, and an accumulation of oxidative dopamine metabolites can lead to mitochondrial membrane depolarization (306) and decreased ETC activity and ATP synthesis (298). Moreover, dopamine can also be directly taken up by mitochondria where it reversibly inhibits the first ETC complex, leading to oxidative stress and impaired mitochondrial energy production (297). Conversely, mitochondrial dysfunction can disrupt both glutamatergic and dopaminergic neurotransmitter signaling; TCA cycle anaplerosis diverts glutamate away from glutamine recycling while oxidative stress promotes the generation of oxidative dopamine derivatives. Notably, the consequent shift towards glycolysis in glial cells upregulates the production of lactate and serine, which potentiates glutamatergic excitotoxicity and neuroinflammation (215,307).

Similarly, a reciprocal relationship between mitochondrial function and the two branches of tryptophan metabolism modulates the synthesis of serotonin, melatonin, and kynurenine (KYN) metabolites (308). Serotonin positively regulates mitochondrial biogenesis, oxidative capacity and ATP synthesis (294,309) while serotonin deficiency disrupts amino acid and lipid metabolism, TCA cycling, OXPHOS and antioxidant activity (310,311). Serotonin also functions as a precursor to melatonin, which regulates mitochondrial OXPHOS, redox homeostasis and inflammatory responses (312). Moreover, melatonin regulates the expression of the rate limiting enzymes involved in serotonin synthesis and catabolism (313), thereby modulating the balance between the serotonin and KYN branches of the tryptophan catabolic pathway. Importantly, the KYN pathway functions as a key mediator between neuroendocrinology and neuroinflammation that is tightly

regulated by interactions between glial cells and neurons. The KYN metabolites kynurenic acid (KYNA) and quinolinic acid (QUIN) inhibit and activate glutamate receptors respectively (112) and flux in the QUIN branch of the KYN pathway that alters *de novo* NAD⁺ synthesis also modulates metabolic and oxidative state (314). Mitochondrial dysfunction shifts the KYN pathway towards QUIN production by increasing the demand for NAD⁺ (315); conversely, QUIN contributes to mitochondrial dysfunction by increasing cytoplasmic calcium concentration and superoxide production in microglia (316). A pathogenic shift towards a reactive glial phenotype under inflammatory conditions upregulates microglial tryptophan import, which both decreases local tryptophan availability for serotonin synthesis and further increases the production of QUIN in microglia (316). Moreover, QUIN upregulates glutamatergic neurotransmission and inhibits glutamate uptake from the synaptic cleft, thereby promoting excitotoxicity (317). Altogether, this demonstrates that a disruption of the relationship between mitochondrial function and the glutamatergic, serotonergic and dopaminergic neurotransmitter cycles increases oxidative stress and shifts glial metabolism towards glycolysis, thereby contributing to gliosis, excitotoxicity and neuroinflammation (215,297,317).

Thus, these neurotransmitter systems are intrinsically coupled to, and mediators of, the relationship between mitochondrial dysfunction, neuroinflammation and HPA axis signaling that is implicated in both ELS and ASC. This discussion highlights that MAL plays a central role in the link between molecular stress signatures and neurophysiological and neurochemical mechanisms that lead to psychopathology. Moreover, these mechanisms are known to contribute to distinct neurological and behavioral aspects of ASC etiology and could function as an underlying susceptibility to allostatic load. This provides a conceptual framework for a molecular signature that may underly autistic burnout, mediate the relationship between social camouflaging and suicidality, or predispose autistic individuals to psychopathology; and thus represents a crucial target for future molecular research that aims to improve the quality of life of autistic individuals.

5. Conclusion: an integrative biopsychosocial framework for psychopathology in ASC

Molecular research into ELS is increasingly integrating various paradigms from disparate scientific disciplines in order to develop a more cohesive understanding of pathological stress responses (349). While ELS research is distinctly and fundamentally separate from autism research, this discussion has highlighted the utility of such an integrative framework to shape molecular research into ASC. The argument put forward in this article builds on separate bodies of work that are well-documented in their respective fields to propose a framework for the development of psychopathology in ASC (Figure 1). This framework integrates emerging literature that is foregrounding autistic experiences with the molecular model for allostatic load that has been developed to quantify the cumulative impact of biological and psychosocial stress (82). This framework proposes that ASC is associated with both an increased exposure to, and an underlying molecular susceptibility to, ELS that contributes to allostatic overload and the subsequent development of psychopathology.

Autistic children are more likely to be exposed to early childhood adversity, psychosocial stress and experiences that threaten their safety or well-being, while also having inadequate internal resources to cope with these stressors. Moreover, the recent characterization of concepts like social camouflaging and autistic burnout in academic literature has highlighted that autistic children are exposed to chronic and distinct forms of stress throughout adolescence that often go unrecognized. During middle childhood, children with ASC are increasingly required to suppress autistic traits in order to meet social and academic expectations, although the extent to which each individual is motivated or able to meet these expectations is mediated by a complex interplay between underlying physiology and sociocultural factors that contribute to socialization. Chronic social camouflaging creates an increasing mismatch between internal capacity and external expectations, culminating in autistic burnout. Importantly, both social camouflaging and autistic

burnout are known to contribute to the development of psychopathology and suicidality in ASC but the physiological mechanisms that are involved in these phenomena, or their relationship to each other, remain unknown and largely unexplored by molecular researchers.

The molecular mechanisms that are known to mediate the relationship between ELS, CAS and psychopathology revolve around interdependent signaling between mitochondrial metabolism, inflammatory immune responses and stress-responsive GC signaling. MAL is emerging as a central modulator of the neurophysiological processes that are disrupted by ELS and contribute to psychopathology. Notably, mitochondrial dysfunction is an underlying component of ASC physiology that is sensitive to GC signaling and plays a central role in regulating HPA axis and innate immune responses. Mitochondria are known to play an essential role in the regulation of neurodevelopment by facilitating the metabolic shift towards OXPHOS that is required for neuronal differentiation. Mitochondria also modulate microglial activation, which is essential for synaptogenesis and synaptic pruning during neurodevelopment. Moreover, the relationship between mitochondrial metabolism and microglial activation regulates myelin turnover, neuronal function, redox homeostasis and glial inflammatory state in the adult brain. The stress-responsive HPA axis is intrinsically coupled to both mitochondrial metabolism and inflammatory signaling, and this relationship is consistently shown to be disrupted by ELS and altered in psychopathology. Notably, the serotonergic, dopaminergic and glutamatergic neurotransmitter systems that are directly involved in many psychological disorders are closely coupled to mitochondrial metabolism, modulated by the HPA axis and represent independent mechanisms that contribute to gliosis and excitotoxicity. This review has highlighted how these molecular mechanisms are implicated in the neurophysiology and neurochemistry of autism and contribute to the development of psychopathology following ELS.

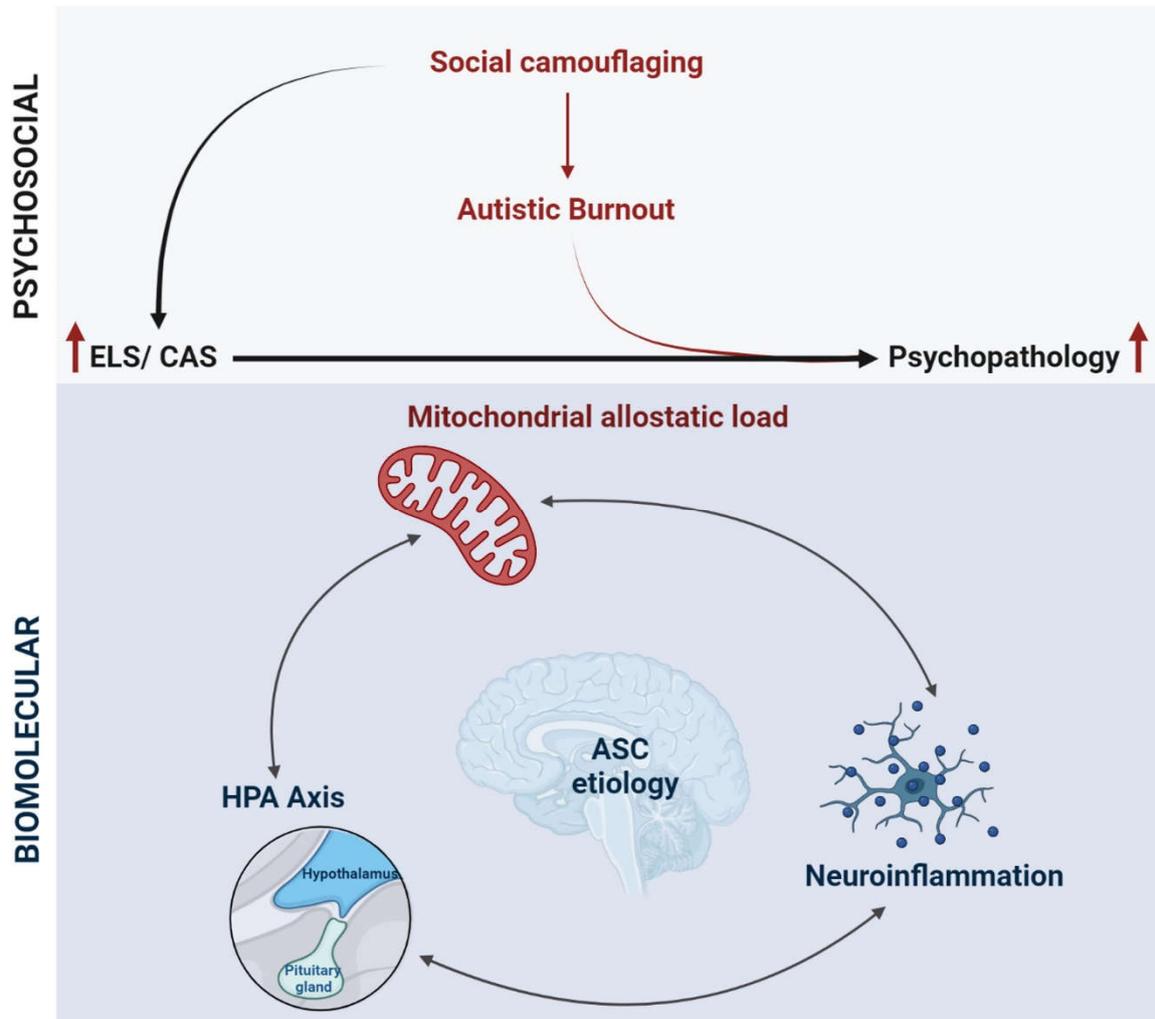


Figure 1. An integrative biopsychosocial vulnerability framework for the development of psychopathology in ASC. There is a well-established relationship between exposure to early life stress (ELS) or chronic adolescent stress (CAS) and the development of psychopathology (68–70,350). Autism is associated with both a significantly increased risk of exposure to ELS (35,36) and high rates of psychopathology and suicidality (29,39,40,43). Emerging characterizations of internal autistic experiences including social camouflaging (21,55,56) and autistic burnout (50–52) overlap closely with established definitions for ELS (65,66) and lead to increased levels of anxiety, depression, self-injury and suicidality in ASC (28,29,57,58). Thus, the development of psychopathology in autism may be mediated by an increased exposure to toxic stressors culminating in allostatic overload. Molecular stress research has demonstrated that mitochondrial allostatic load (MAL) (85) plays a central role in the development of psychopathology following ELS by disrupting a tightly coupled signaling network between neuroimmune (66,176,180,181,183) and neuroendocrine signaling (65,66,142). Separate reviews have comprehensively described the tightly regulated biochemical relationships between mitochondrial dysfunction and gliosis (204,218,223,351); mitochondrial metabolism and HPA axis signaling (82,149,150); and the HPA axis and inflammatory immune response pathways (201,202). ASC etiology is characterized by mitochondrial dysfunction (117–119), gliosis and neuroinflammation (184–189,200) and a complex signature of HPA axis dysregulation (44,175). Moreover, this three-way molecular interplay converges on the regulation of key neurodevelopmental processes implicated in the response to ELS and psychopathology that are also associated with distinct neurophysiological, neurochemical and behavioral aspects of ASC etiology (211,219,243,245,289,294,297). This could point to a molecular vulnerability to the development of psychopathology in ASC, highlighting how psychosocial and biological factors converge to increase the risk of psychopathology and suicidality.

Considering both the increased risk of exposure to, and underlying vulnerability to, ELS in autism emphasizes how psychosocial and biological factors converge to increase

the risk of psychopathology. This interplay could act as a potential mechanism that contributes to the high rates of depression and suicidality in autistic adults and may reveal a biological signature that underlies autistic burnout. Autistic burnout, depression and suicidality are factors that directly impair health, well-being and function for people with autism and much of the literature highlights the importance of improved access to accommodations, earlier diagnosis and decreased stigmatization of autistic traits in mitigating psychopathology in ASC. In addition, further investigating an underlying susceptibility to ELS could inform research into novel molecular interventions. Targeted therapeutic strategies might be able to protect children with ASC from the development of psychopathology, facilitate recovery from autistic burnout, or identify diagnostic tools to differentiate autistic burnout from clinical depression. Ultimately, this highlights the utility of molecular research that considers autistic experiences in order to progress towards findings that could tangibly improve the quality of life for autistic individuals.

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