



HHS Public Access

Author manuscript

Clin Neurophysiol. Author manuscript; available in PMC 2022 July 01.

Published in final edited form as:

Clin Neurophysiol. 2021 July ; 132(7): 1647–1662. doi:10.1016/j.clinph.2021.03.021.

Modulation of motor cortical excitability by continuous theta-burst stimulation in adults with autism spectrum disorder

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Abstract

Objective.—To test whether change in motor evoked potential (MEP) induced by continuous theta-burst stimulation (cTBS) of motor cortex (M1) distinguishes adults with autism spectrum disorder (ASD) from neurotypicals, and to explore the contribution of two common polymorphisms related to neuroplasticity.

Methods.—44 adult neurotypical (NT) participants (age 21–65, 34 males) and 19 adults with ASD (age 21–58, 17 males) prospectively underwent M1 cTBS. Their data were combined with previously obtained results from 35 NT and 35 ASD adults.

Results.—MEP at 15 minutes post-cTBS (T15) was a significant predictor of diagnosis ($p=0.04$) in the present sample ($n=63$). T15 remained a significant predictor in a larger sample

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Conflicts of Interest Statement

A.P.-L. is a co-founder of Linus Health and TI Solutions AG; serves on the scientific advisory boards for Starlab Neuroscience, Magstim Inc., and MedRhythms; and is listed as an inventor on several issued and pending patents on the real-time integration of noninvasive brain stimulation with electroencephalography and magnetic resonance imaging. A.R. is a founder and advisor for Neuromotion and PrevEp, and serves on the scientific advisory board or has consulted for Cavion, Epihunter, Gamify, Neural Dynamics, NeuroRex, Praxis, Roche, Otsuka, and is listed as an inventor on a patent related to integration of TMS and EEG. The remaining authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

(n=91) and when partially imputed based on T10–T20 from a yet-greater sample (N=133). T15 also remained a significant predictor of diagnosis among brain-derived neurotrophic factor (*BDNF*) Met+ and apolipoprotein E (*APOE*) ε4– subjects (p 's<0.05), but not among Met– or ε4+ subjects (p 's>0.19).

Conclusions.— MEP at T15 post-cTBS is a significant biomarker for adults with ASD, and its utility is modulated by *BDNF* and *APOE* polymorphisms.

Significance.—M1 cTBS response is a physiologic biomarker for adults with ASD in large samples, and controlling for *BDNF* and *APOE* polymorphisms can improve its diagnostic utility.

Keywords

transcranial magnetic stimulation; continuous theta-burst stimulation; biomarker; autism spectrum disorder; *BDNF*; *APOE*

1. Introduction

Autism spectrum disorder (ASD) refers to a group of complex neurodevelopmental disorders characterized by: (1) persistent deficiencies in social communication and social interaction, and (2) limited interests and repetitive behavior (American Psychiatric Association, 2013). ASD has an estimated prevalence of 14.5 per 1,000 in the U.S. (Christensen et al., 2018) and often results in significant impairments in activities of daily living, both in children and adults (Haertl et al., 2013). Because of the large heterogeneity of the clinical endophenotype in ASD and symptom manifestation over a range of ages and to different degrees, the clinical diagnosis of ASD can be challenging and is typically based on behavioral interviews and subjective clinical impression. Thus, an objective neurophysiologic biomarker that can facilitate ASD diagnosis is highly desirable, especially to improve diagnostic accuracy and to enable metrics of target engagement and clinical/behavioral outcomes in therapeutic interventions.

Data from rodent ASD models and studies on genetic syndromes with high prevalence of ASD symptoms in humans indicate aberrant mechanisms of synaptic plasticity in ASD pathophysiology, including use-dependent changes in synaptic strength (Bhakar et al., 2012; Bourgeron, 2009; Gipson and Johnston, 2012; Krueger and Bear, 2011; Peça et al., 2011; Percy, 2011) and abnormalities in long-term potentiation (LTP) and long-term depression (LTD) of excitatory synaptic strength (Dani et al., 2005; Gogolla et al., 2009; Huber et al., 2002; Narita et al., 2002; Rinaldi et al., 2008a, 2008b, 2007; Tordjman et al., 2007).

Plasticity mechanisms similar to LTP and LTD can be studied noninvasively in humans using transcranial magnetic stimulation (TMS) (Barker, 2017; Barker et al., 1985; Thickbroom, 2007; Ziemann, 2004). TMS is a neurophysiological technique based on the principle of electromagnetic induction that enables triggering or modulation of neural activity in the brain (Hallett, 2007) and is considered safe when applied following the recommended guidelines (Rossi et al., 2009; Rossini et al., 2015). Delivering a single TMS pulse (spTMS) to primary motor cortex (M1) can induce a motor evoked potential (MEP) in the target muscle. TMS has been used in various forms including spTMS, paired-pulse TMS (ppTMS), and repetitive TMS (rTMS) at specific intensities, frequencies, and patterns of

stimulation to study, modify, or restore activity in corticospinal pathways, as well as various brain regions and networks (see Valero-Cabré et al. 2017 for a review).

Several spTMS studies have found no significant difference in baseline M1 excitability in ASD (Enticott et al., 2012; Enticott et al., 2013; Minio-Paluello et al., 2009; Théoret et al., 2005). PpTMS studies have found no alteration in short-interval intracortical inhibition (SICI) (Jung et al., 2013; Théoret et al., 2005) or intracortical facilitation (ICF) in ASD (Enticott et al., 2010; Peter G Enticott et al., 2013; Théoret et al., 2005). The findings of long-interval intracortical inhibition (LICI) in ASD have been mixed, with some ASD individuals exhibiting abnormal intracortical inhibition while others have responses similar to those of neurotypical (NT) individuals (Enticott et al., 2010; Enticott et al., 2013; Oberman et al., 2010).

A form of rTMS referred to as theta-burst stimulation (TBS) of M1 (Huang et al., 2005) consists of 50Hz bursts of triplet TMS pulses repeated at 5 Hz for a total of 600 pulses, in one of two protocols: (1) intermittent theta-burst stimulation (iTBS) with a 2-sec on, 8-sec off pattern for 190 sec that typically induces MEP facilitation by ~35% for up to 60 min; (2) continuous theta-burst stimulation (cTBS) for 40 sec that typically induces MEP suppression by ~25% for up to 50 min (Wischniewski and Schutter, 2015). MEP facilitation and suppression by iTBS and cTBS protocols are considered to involve mechanisms similar to LTP and LTD, respectively (Huang et al. 2005; Huerta and Volpe 2009). The return of post-TBS MEP amplitudes to their baseline levels is considered a neurophysiologic index of the efficacy of the mechanisms of cortical plasticity (Oberman et al., 2010, 2016, 2014, 2012; Pascual-Leone et al., 2011, 2005; Suppa et al., 2016; Tremblay et al., 2015). TBS aftereffects involve mechanisms of gamma-aminobutyric acid- (GABA-)ergic and glutamatergic plasticity (Benali et al., 2011; Huang et al., 2008, 2007; Stagg et al., 2009; Trippe et al., 2009).

Studies by Oberman and colleagues (Oberman et al., 2016, 2012) found greater and longer-lasting TBS-induced changes in MEP amplitude in adults with ASD compared to NT adults, indicating an exaggerated, *hyperplastic* response to TBS in ASD. Recently, we found that children and adolescents with high-functioning ASD (HF-ASD) had abnormally greater facilitatory responses to cTBS relative to typically developing children (Jannati et al., 2020). Moreover, cTBS measures of plasticity showed a maturational trajectory in children and adolescents with HF-ASD, in which the extent of, or the maximum, cTBS-induced suppression of MEPs increased linearly with age (Jannati et al., 2020; Oberman et al., 2014).

These results collectively support the utility of cTBS cortical plasticity measures as biomarkers for individuals with ASD across the lifespan (Jannati et al. 2020). In recent years, however, several studies have documented large inter- and intra-individual variability in M1 cTBS responses among healthy adults (Corp et al., 2020; Goldsworthy et al., 2014; Hamada et al., 2013; Hordacre et al., 2017; Jannati et al., 2019, 2017; Nettekoven et al., 2015; Vallence et al., 2015). Such variability may limit the biomarker utility of cTBS for differentiating individuals with ASD from their NT counterparts. Careful consideration of possible sources of within- and across-individual variability is thus important.

Here we address these issues by comparing M1 cTBS aftereffects in a relatively large sample of NT participants (n=44) and a group of ASD participants (n=19), and then validating the obtained results by combining individual participant data from the present sample with the corresponding values from 70 additional subjects from two separate datasets reported previously (Oberman et al., 2012). The total sample (N=133) included 79 NT and 54 ASD subjects across three datasets and two centers. Moreover, we stratified the participants with available DNA data in the present sample for two single-nucleotide polymorphism (SNPs) identified as important contributors to variability of rTMS and other measures of neuroplasticity: the Val66Met SNP in the brain-derived neurotrophic factor (*BDNF*) gene (Antal et al., 2010; Chang et al., 2014; Cheeran et al., 2008; Di Lazzaro et al., 2015; Fried et al., 2017; Jannati et al., 2019, 2017; Lee et al., 2013) and the presence of the $\epsilon 4$ allele in the apolipoprotein E (*APOE*) gene (Jannati et al., 2019; Mahley and Rall Jr, 2000; Nichol et al., 2009; Peña-Gomez et al., 2012; White et al., 2001; Wolk et al., 2010). We then calculated the standard measures of biomarker utility for each comparison of cTBS responses.

2. Methods

2.1. Participants

63 adults (age range 21–65, 12 females) participated in this study. The local Institutional Review Board approved the study in accordance with the Declaration of Helsinki. All participants provided written informed consent prior to enrollment and received monetary compensation for their participation. All participants were screened for TMS contraindications (Rossi et al., 2009), and all had a normal neurological examination. The present sample (*Dataset 1*) consisted of two groups: (1) high-functioning adults with non-syndromic ASD (n = 19); (2) neurotypical age- and gender-matched control participants (n = 44). Local community advertisements and autism associations and clinics were used for participant recruitment. Participants in the ASD group were required to provide documentation of a clinical diagnosis made by a psychiatrist or clinical psychologist, met the ASD criteria in the Diagnostic and Statistical Manual of Mental Disorders, 5th edition (DSM-5) (American Psychiatric Association, 2013), and were independently assessed with the Autism Diagnostic Observation Schedule (ADOS; mean score = 8.68; SD = 4.35). Participants in the NT group had no neurological or psychological disorder. Screening for TMS contraindications was based on the safety recommendations of the International Federation of Clinical Neurophysiology (IFCN) (Rossi et al., 2009). Detailed demographic characteristics of the participants are presented in Table 1 and comparisons of those characteristics between ASD and NT groups are presented in Table 2. Racial categories were defined according to the National Institutes of Health guidelines for inclusion of minorities as subjects in clinical research (NIH Office of Extramural Research, 2001). Handedness in the present sample was determined by asking participants their hand preference. Individual handedness data were not available for the other two datasets.

The two other datasets, reported previously (Oberman et al., 2012), included *Dataset 2* collected in Boston, Massachusetts (n=40; 20 ASD subjects; age 18–64, 4 females) and *Dataset 3* collected in Barcelona, Spain (n=30; 15 ASD subjects; age 29–52, 2 females).

2.2. Neuropsychological testing

The Abbreviated Battery of Stanford–Binet IV intelligence scale (Thorndike et al., 1986) and the Autism-Spectrum Quotient (AQ) (Baron-Cohen et al., 2001) were completed for both groups in Dataset 1. All participants in the ASD group had an Abbreviated IQ > 70 (Table 1). AQ scores were used to quantify the situation of each participant on the continuum from normality to autism and to rule out clinically significant levels of autistic traits in the NT group (Baron-Cohen et al., 2001). The Autism Diagnostic Observation Schedule (ADOS) Module 4 was conducted by a research-reliable investigator (L.O.) to assess the current social-communicative behavior in the ASD group (Lord et al., 2000). Corresponding neuropsychological details of Datasets 2 and 3 were reported previously (Oberman et al., 2012). Individual age data were available for all three datasets, whereas individual IQ scores were only available for Datasets 1 and 2. ASD and NT groups in each of the three datasets were age-, gender-, and IQ-matched.

2.3. Genetic testing

Saliva samples from 56 participants in Dataset 1, including 17 participants (89.5%) in the ASD group and 39 participants (88.6%) in the NT group, were assessed for single-nucleotide polymorphisms (SNPs) in the brain-derived neurotrophic factor (*BDNF*), Val66Met, and *APOE* genes. Four NT and two ASD participants did not consent to providing DNA samples, and one NT sample was deemed unusable.

The Oragene Discover OGR-250 Kit (DNA Genotek Inc., Ottawa, ON, Canada) was used to extract genomic DNA from saliva samples using standard methodology and the prepIT•L2P reagent (DNA Genotek Inc., 2015). Each sample's quality was assessed using PicoGreen fluorometry for double-stranded DNA quantification, Nanodrop spectrophotometry as to estimate purity using A260/A280 ratios, and agarose gel electrophoresis to visualize DNA integrity. A TaqMan single tube genotyping assay, using polymerase chain reaction (PCR) amplification and a pair of fluorescent dye detectors targeting the SNP, was used to analyze the rs6265 SNP of the *BDNF* gene, and the rs429358 and rs7412 SNPs of the *APOE* gene.

2.4. Transcranial magnetic stimulation

Participants sat in a comfortable chair with the right arm and hand in a natural pronated $\sim 90^\circ$ angle on a pillow, were instructed to keep their right hand relaxed and to keep their eyes open during the session. Participants were also monitored for drowsiness during TMS. Live electromyogram (EMG) was monitored to ensure hand relaxation throughout the session.

TMS procedures were conducted according to the IFCN-recommended guidelines (Rossi et al., 2009; Rossini et al., 2015). All TMS pulses were applied with a MagPro X100 stimulator, using the 'normal' settings, attached to a MC-B70 Butterfly Coil (outer diameter: 97mm; MagVenture A/S, Farum, Denmark), generating biphasic pulses with induced current in the antero-posterior–postero-anterior (AP-PA) direction in the brain. The intensity of single pulses was set at 120% of individual resting motor threshold (RMT) whereas cTBS was applied at 80% of individual active motor threshold (AMT). The coil was held tangential to the scalp, with the handle pointing posteriorly and at 45° angle relative to the mid-sagittal line. To achieve consistent targeting, neuronavigation was performed with

Polaris infrared-optical tracking (Northern Digital Inc., Waterloo, ON, Canada) and Brainsight software for frameless stereotaxy (Rogue Research Inc., Montreal, QC, Canada). To ensure the consistency of the coil position and orientation, the participant's brain MRI or an MRI template was used to register the participant's head using pre-defined cranial landmarks and 12 additional samples from the scalp (Ruohonen and Karhu, 2010).

A PowerLab 4/25 data-acquisition device and LabChart software (AD Instruments, Colorado Springs, CO, USA) were used to record surface EMG from the right FDI, using the belly-tendon electrode montage. The TMS and EMG systems were synchronized via triggered pulses issued by the TMS device. EMG signal was digitized at 1 kHz, epoched from 100 ms pre-trigger to 500 ms post-trigger, amplified within ± 10 mV range, and band-pass filtered (0.3–1000 Hz).

At the beginning of each TMS session, the motor hotspot, defined as the optimal spot for eliciting maximal motor evoked potentials (MEPs) from the right FDI was localized, and the RMT, defined as the lowest stimulation intensity that elicited MEPs ≥ 50 μ V on at least five of ten trials, was measured. To assess baseline cortico-motor reactivity, three blocks of 30 single TMS pulses were applied to M1, with a 5-min inter-block interval and at a random 4–6 s interval between successive pulses. Individual MEPs within each block that were > 2.5 SD from the mean were excluded from further analyses. There were never more than three MEPs excluded from each block. It has been estimated that at least 20 pulses are required to obtain a reliable estimate of the MEP amplitude at a given point in time (Chang et al., 2016; Goldsworthy et al., 2016).

The peak-to-peak amplitudes of MEPs in the three pre-cTBS blocks were averaged to calculate the baseline MEP amplitude. The AMT was then measured as the lowest stimulation intensity that evoked MEPs ≥ 200 μ V on at least five of ten trials while the participant slightly contracted the FDI. To control the effects of voluntary hand movements on cTBS aftereffects (Iezzi et al., 2008), there was a 5-minute break between the AMT measurement and the delivery of cTBS, during which participants were asked to maintain hand relaxation. cTBS was applied as triplet pulses at 50 Hz, delivered as 200 bursts repeated every 200 ms for 40 s (for a total of 600 pulses). To assess post-cTBS cortico-motor reactivity, 30 single TMS pulses were applied at 5, 10, 15, 20, 30, 40, 50, and 60 min following the cTBS (*T5–T60*). Each time point of interest was in the middle of each post-cTBS block.

Datasets 2 and 3, reported previously (Oberman et al., 2012), were collected with Magstim Super Rapid stimulators and monophasic pulses inducing posterior-anterior (PA) currents in the brain. TMS procedures for Datasets 2 and 3 were identical to those employed in the present study, except T15 data were not collected in Dataset 2 and individual T5, T10, T20 and T50 data were not collected in Dataset 3.

2.5. Statistical analyses

Research Electronic Data Capture (REDCap) electronic data capture tools hosted at Beth Israel Deaconess Medical Center (BIDMC) (Harris et al., 2019, 2009) were used for study data collection and management. Statistical analyses were conducted with Stata 16.1

(StataCorp, College Station, TX, USA) and MATLAB R2016b (The MathWorks, Natick, MA, USA) software.

RMT and AMT were expressed as percentage of maximum stimulator output (MSO), and the baseline MEP amplitude was calculated as the average of baseline MEP amplitude in 3 blocks of 30 single TMS pulses. The aftereffects of cTBS were calculated as the percent change from baseline (%) by calculating the average amplitude of 30 MEPs at T5–T60 for each participant. Significant deviations from normal distribution in MEP values were found with the Shapiro–Wilk test. Thus, each post-cTBS MEP amplitude was first baseline-corrected in each participant. A natural log-transformation was then applied to the baseline-corrected MEP amplitudes at each post-cTBS time point (MEP) (Nielsen, 1996a, 1996b; Pasqualetti and Ferreri, 2011). The same transformations were done on the raw MEP values in Datasets 2 and 3.

Grand-average MEPs were calculated separately for each time-point in each group. In Dataset 1, MEP at T5–T20 for one NT subject and MEP at T15 in another NT subject were not obtained due to technical difficulties. The missing MEPs in Dataset 1 were imputed using multiple regression on *Age* and *IQ* score by sampling from conditional distribution with bootstrap (Gelman et al., 2014; Royston, 2004). The uncollected MEPs at T15 in Dataset 2 were imputed as the average of MEPs at T10 and T20 for each subject.

For the purpose of investigating the utility of cTBS aftereffects for differentiating ASD and NT individuals, assessing the differential responses of the two groups (relative to each other) is more important than assessing whether cTBS induced a significant effect in either group. For this reason, and to avoid multiple comparisons that would reduce the power of our contrasting analyses, we focused our analyses on the separability of cTBS responses between the two groups rather than the significance of cTBS-induced modulation within each group.

Pairwise comparisons were conducted to assess whether demographic, neuropsychological, genetic, and neurophysiological measures (RMT, AMT, and baseline MEP amplitude) were significantly different between ASD and NT groups (Table 2). Continuous variables were compared using Student's *t*-tests, while proportions were compared using Fisher's Exact tests. All analyses were two-tailed, and α level was set to 0.05.

To compare cTBS aftereffects between ASD and NT groups, MEPs were entered into a 2 (*Diagnosis*) \times 8 (*Time*) repeated-measures analysis of variance (ANOVA). However, as the maximum modulation of MEP amplitudes typically occurs within the first 20 minutes after cTBS (Table 4 in Wischniewski and Schutter 2015), planned pairwise comparisons between MEPs at T5–T20 in the two groups were conducted using Student's *t*-tests. When indicated, false discovery rate (FDR) was controlled by adjusting the *p*-values for multiple comparisons using the Benjamini-Hochberg method (Benjamini and Hochberg, 1995). T15 was selected *post hoc* (see Results) as the time-point at which cTBS plasticity measures were most altered in ASD relative to NT controls. To assess the predictive utility of cTBS aftereffects, logistic-regression analyses were conducted with *Diagnosis* (ASD vs. NT) as dependent variable (DV) and *MEP* at T15 as the main independent variable (IV).

Studies have found TMS measures can be influenced by demographic variables such as age (Corp et al., 2020; Freitas et al., 2011), sex (Cahn et al., 2003; De Gennaro et al., 2004; Huber et al., 2003), genetic polymorphisms (Antal et al., 2010; Chang et al., 2014; Cheeran et al., 2008; Di Lazzaro et al., 2015; Peña-Gomez et al., 2012), and baseline neurophysiological measures including rMT, aMT, and baseline MEP amplitude (Corp et al., 2020, In Press; Fried et al., 2017; Jannati et al., 2017). Race and education can sometimes act as, albeit imperfect, proxy variables for socioeconomic status and possibly other/unknown covariates in studies with health-related outcomes (Geronimus et al., 1996; Krieger, 1992, 1990; Krieger and Gordon, 1999; Soobader et al., 2001). To control for potential confounders, demographic (age, sex, race, and education), neuropsychological (IQ), genetic (*BDNF* and *APOE* SNPs, when available), and baseline neurophysiological measures (RMT, AMT, baseline MEP amplitude) were added, one at a time, as covariates to the logistic-regression model. For each model, area under the receiver operating characteristic curve (AUROC), positive and negative predictive values (PPV and NPV) and percentage correctly classified (Dx) were calculated.

To validate the utility of T15 MEP in a larger sample, we conducted an overall logistic regression with *Diagnosis* as DV and MEP at T15 as IV. *Dataset* was included as a covariate to control for differences in TMS equipment and pulse waveform and other potential differences in subject composition across the datasets. The logistic-regression analyses involving T15 were conducted both with and without including the imputed T15 values in Dataset 2. We also controlled for *Age* and *IQ* (when available) as additional covariates. As exploratory analyses, we checked whether post-cTBS time points other than T15 were also significant predictors of *Diagnosis* in the overall regression models. The analyses referring to $n = 91$ consisted of participants in Datasets 1 and 3, for whom T15 MEP was obtained.

Based on previous studies that found a developmental trajectory in cTBS response in children and adolescents with ASD (Jannati et al., 2020; Oberman et al., 2014), we assessed the changes in cTBS response across the adult lifespan (age 21–65) by evaluating the correlation between age and MEP at T15 separately for each group across the two datasets in which the age data were available (Datasets 1 and 3; $n=91$).

To control for influences of *BDNF* and *APOE* genotypes on cTBS aftereffects, MEPs of ASD and NT groups in Dataset 1 were compared separately for *BDNFMet-*, *BDNFMet+*, *APOE e4-*, and *APOE e4+* participants. For *APOE e4+* participants, T30 was selected *post hoc* (see Results) as the time-point at which cTBS aftereffects were most altered in the ASD group relative to NT controls. Thus, the logistic-regression analyses to predict *Diagnosis* among *APOE e4+* participants were conducted with MEP at T30 as the IV.

To assess whether *BDNF* or *APOE* genotype influenced the type of cTBS response, we classified participants in Dataset 1 as *inhibitor*, *facilitator*, or *non-responder* if their MEP at T15 was equivalent to 90% (-0.105), 110% (0.095), or between 90 and 110%, ($-0.105, 0.095$), respectively (Nettekoven et al., 2015). We then compared the proportion of participants with different types of cTBS response between the ASD and NT groups and their corresponding *BDNF* and *APOE* subgroups. Unless explicitly specified, all three types

of responses were included in the comparisons of proportions between different groups/subgroups.

3. Results

Table 1 details demographics, *BDNF* and *APOE* SNPs, single-nucleotide polymorphisms, and neuropsychological measures for individual participants. Group means \pm SD for those measures and their comparisons in the ASD and NT groups as well as in their *BDNF* and *APOE* subgroups are presented in Table 2. Education data were available for 16 (84.2%) and 42 (95.4%) participants in the ASD and NT groups, respectively. *BDNF* data were available for 17 (89.5%) and 40 (90.9%) participants in the ASD and NT groups, respectively. *APOE* data were available for 16 (84.2%) and 40 participants (90.9%) in the ASD and NT groups, respectively. Considering the high percentage of participants in the two groups for whom education and genetic data were available, we decided it would be beneficial to control for those covariates even in the presence of incomplete data.

3.1. Demographics and neuropsychological testing

There was no significant group difference in age, sex, handedness, or education (p 's $>$ 0.1), but the proportion of White participants was significantly higher in the ASD group than in the NT group ($p = 0.050$). Therefore, to control for potential confounding effect of Race, we repeated the main-group logistic regression analysis among White participants.

In the neuropsychological measures, the IQ scores were comparable between the two groups ($p = 0.41$), indicating that ASD participants did not significantly differ from NT controls in terms of overall cognitive function. As expected, participants with ASD had significantly higher AQ scores than their NT counterparts in the whole sample and in all *BDNF* and *APOE* subgroups (p 's $<$ 0.001).

3.2. Genetic analyses

Among 56 participants with available *BDNF* results, the proportions of *BDNF* Val/Val and Val/Met genotypes were 55.4% and 44.6%, whereas among 55 participants with available *APOE* results, the proportions of *APOE* $\epsilon 4^-$ and $\epsilon 4^+$ genotypes were 65.5% and 34.6%, respectively. The two groups were comparable in *BDNF* Met $^-$: Met $^+$ and *APOE* $\epsilon 4^-$: $\epsilon 4^+$ ratios (p 's $>$ 0.5).

ASD and NT participants in each *BDNF* and *APOE* subgroup were comparable in terms of demographics and IQ scores (p 's $>$ 0.05; Table 2). *APOE* $\epsilon 4^-$ participants were more educated in the NT group than in the ASD group ($p = 0.015$).

3.3. Measures of corticospinal excitability and plasticity

All participants tolerated TMS procedures with no complications or unexpected side effects. As detailed in Table 2, RMT, AMT, and baseline MEP amplitudes were comparable between the ASD and NT groups (p 's $>$ 0.3), and their *BDNF* and *APOE* subgroups (p 's $>$ 0.2). These results indicate the ASD participants did not differ significantly from NT controls in baseline corticospinal excitability.

The repeated-measures ANOVA indicated the MEPs at T5–T60 did not vary significantly by *Diagnosis*, $F(1,61) = 0.64$, $p = 0.428$, $\eta^2_p = 0.01$, *Time*, $F(7,422) = 1.82$, $p = 0.082$, $\eta^2_p = 0.03$, or their interaction, $F(7,422) = 1.37$, $p = 0.218$, $\eta^2_p = 0.02$. Planned *t*-tests at T5–T20 found ASD subjects had significantly greater inhibition of MEPs at T15 ($p = 0.029$), which did not survive FDR adjustment ($P_{FDR} = 0.116$). In the NT group, 29.5%, 13.6%, and 56.8% of participants were inhibitors, non-responders, and facilitators, respectively, whereas the corresponding proportions in the ASD group were 57.9%, 10.5%, and 31.6%. There was no significant overall difference in the type of cTBS response between the two groups ($p = 0.103$). When the 8 non-responders were excluded from the two groups, the proportion of inhibitors was significantly greater in the ASD group than in the NT group ($p = 0.044$).

MEPs at T5–T10 were not significantly different between the two groups (p 's > 0.2). Figure 1 shows the cTBS aftereffects in ASD and NT groups and the ROC curve associated with the logistic regression.

Logistic regression analysis found that T15 MEP was a significant predictor of *Diagnosis*, $\beta = -1.55$, $p = 0.038$, indicating a more negative MEP at T15 was predictive of ASD diagnosis, AUROC = 0.64. As detailed in Table 3, follow-up analyses found that T15 MEP remained a significant predictor of *Diagnosis* after controlling for *IQ*, *Handedness*, *BDNF*, *APOE*, *RMT*, *AMT*, or *Baseline MEP* (p 's < 0.046) but not after controlling for *Age*, *Sex*, or *Education* (p 's > 0.052). However, none of the added covariates was a significant predictor (p 's > 0.08), and the predictive powers of all the logistic-regression models were comparable (AUROC range 0.64–0.70). Controlling for *Education* or *Race* resulted in the most predictive models (Dx > 80%), whereas controlling for other covariates resulted in 72–78% correct classification (Table 3). T15 MEP remained a significant predictor of *Diagnosis* among White participants ($p = 0.039$; AUROC = 0.70).

Across all three datasets (N = 133), while controlling for *Dataset* and *Age*, T15 MEP (partially imputed based on T10 and T20) was a significant predictor of *Diagnosis* ($\beta = -1.25$, $p = 0.004$; AUROC = 0.70). Across Datasets 1 and 3 in which T15 MEP values were obtained (n = 91), while controlling for *Dataset* and *Age*, T15 MEP was also a significant predictor of *Diagnosis* ($\beta = -1.54$, $p = 0.025$; AUROC = 0.71). Among subjects in Datasets 1 and 2 for whom individual IQ scores were available (n = 89), while controlling for *Dataset*, *Age*, and *IQ*, T15 MEP remained a significant predictor of *Diagnosis* ($\beta = -1.42$, $p = 0.016$, AUROC = 0.76). The effects of *Dataset*, *Age*, or *IQ* were not significant in any of the models above (p 's > 0.09).

Across all three datasets (N = 133), among post-cTBS time points other than T15 and while controlling for *Dataset* and *Age*, MEPs at T30 ($\beta = -1.28$, $p = 0.002$; AUROC = 0.73) and T40 ($\beta = -1.17$, $p = 0.002$; AUROC = 0.72) were significant predictors of *Diagnosis* but MEPs at other time points were not (p 's > 0.07). Among subjects in Datasets 1 and 2 for whom individual IQ scores were available (n = 89), while controlling for *Dataset*, *Age*, and *IQ*, none of the post-cTBS time points other than T15 was a significant predictor of *Diagnosis* (p 's > 0.053).

MEP at T15 was negatively correlated with *Age* among NT subjects across datasets 1 and 3 ($n = 57$; $r = -0.32$, $p = 0.016$) but not among ASD subjects in the same datasets ($n = 34$; $r = -0.23$, $p = 0.189$).

3.4. BDNF and APOE influences in cTBS measures of plasticity

Among 53 participants in Dataset 1, after controlling for both *BDNF* and *APOE* SNPs as covariates, T15 MEP remained a significant predictor of *Diagnosis* ($p = 0.018$; AUROC = 0.69; Dx = 79.3%), whereas neither *BDNF* nor *APOE* status was a significant predictor (p 's > 0.2).

Table 4 presents results of the logistic regression analyses predicting *Diagnosis* with T15 MEP as the IV within *BDNF* and *APOE* subgroups. Because controlling for *Education* or *Race* in the overall logistic regression resulted in the models with the highest Dx values, *Education* and *Race* were added, one at a time, as covariates to each model within genetic subgroups. For *APOE* $\epsilon 4+$ participants, T30 MEP was chosen *post hoc* as the time point at which cTBS aftereffects in ASD participants showed the largest alteration compared to NT participants, and similar logistic regression analyses as above were conducted with T30 MEP as the IV. The analyses found T15 MEP was a significant predictor of *Diagnosis* among 25 *BDNFMet+* participants (16 NT, 9 ASD; $p = 0.048$; AUROC = 0.69; Dx = 83.3%) and among 36 *APOE* $\epsilon 4-$ participants (26 NT, 10 ASD; $p = 0.037$; AUROC = 0.69; Dx = 79.4%), but not among 31 *BDNFMet-* participants (23 NT, 8 ASD; $p = 0.490$; AUROC = 0.57; Dx = 73.3%). Figures 2–5 show the comparisons of cTBS aftereffects in *BDNF* and *APOE* subgroups of ASD and NT participants and the ROC curves associated with the significant logistic-regression models.

In the analysis of *APOE* $\epsilon 4-$ participants, when *Education* was added as a covariate, it was a significant predictor ($p = 0.041$), whereas T15 MEP only showed a trend ($p = 0.064$), even though the model itself remained significant, $p = 0.009$, AUROC = 0.83; Dx = 83.9%. It is possible that the loss of significant effect of T15 MEP was due to the listwise deletion of a few participants for whom *Education* data were not available (Table 2).

Among *APOE* $\epsilon 4+$ participants, T15 MEP was not a significant predictor of *Diagnosis*, with or without controlling for *Education* or *Race* (p 's > 0.2). Choosing T30 MEP as the IV and controlling for *Race* resulted in a significant model, $p = 0.017$, AUROC = 0.86; Dx = 84.2%, and both T30 MEP and *Race* had non-significant effects (p 's < 0.1) (Table 4).

Among either *APOE* $\epsilon 4-$ or $\epsilon 4+$ participants, there was no significant difference between ASD and NT groups in the type of cTBS response ($p = 0.780$), whereas among *BDNFMet+* participants, the proportion of inhibitors was significantly greater in the ASD group than in the NT group ($p = 0.006$). Among either *APOE* $\epsilon 4-$ participants or *APOE* $\epsilon 4+$ participants, there was no significant difference between ASD and NT groups in the type of cTBS response ($p = 0.257$ and $p = 0.185$, respectively).

4. Discussion

The present study compared cTBS measures of brain plasticity in adults with HF-ASD and their age-, gender-, and IQ-matched NT controls. We found, among 63 participants including 19 participants with ASD, cTBS-induced MEP suppression at 15 minutes post-stimulation (T15) was a significant predictor of diagnosis with moderate (72–80%) classification accuracy, depending on the covariate included in the model. The discriminatory power of T15 MEP was not driven by demographic, neuropsychological, or genetic differences between ASD and NT groups. Moreover, we found T15 MEP remained a significant predictor of ASD diagnosis when the present data were combined with the results of two previously reported cohorts, for a total of 133 subjects including 54 ASD subjects. In exploratory analyses, MEPs at T30 and T40 were the only other post-cTBS time point that were significant predictors of ASD diagnosis when controlling for Dataset and Age. In combined analyses, T15 MEP had the highest discriminatory power among all post-cTBS time points.

A novel contribution of the present study was to control for influences of *BDNF* and *APOE* SNPs on cTBS aftereffects, and to assess the discriminatory power of cTBS responses between ASD and NT individuals within the more-homogenous SNP subgroups of either *BDNF* or *APOE* genes. Stratifying the study sample by either *BDNF* or *APOE* SNP status made a noticeable difference in the discriminatory power of cTBS aftereffects. While T15 MEP was not a significant predictor of diagnosis among *BDNFMet-* participants, it was a stronger predictor among *BDNFMet+* and *APOE e4-* participants, with a classification accuracy of up to 88% and 84%, respectively. The overall pattern of cTBS aftereffects among *APOE e4+* was distinct from that among the other genetic subgroups, with MEP at T30 showing the greatest alteration in cTBS response in ASD participants, predicting diagnosis with an accuracy of up to 84%.

4.1. Overall cTBS measures of plasticity in ASD and NT adults

The main finding of the present study is that, despite large inter-individual variability in both groups, M1 cTBS responses were still able to differentiate individuals with ASD from their NT counterparts in a sample that was larger than most other TBS studies (Wischniewski and Schutter, 2015). The lack of a significant overall cTBS-induced suppression of MEPs in the NT group is consistent with the results of several recent studies in healthy adults (Goldsworthy et al., 2014; Hamada et al., 2013; Hordacre et al., 2017; Jannati et al., 2019, 2017; Nettekoven et al., 2015; Vallence et al., 2015). The present overall cTBS results in the NT group further confirm the large interindividual variability in TBS aftereffects (Corp et al., 2020).

As in previous studies investigating brain plasticity mechanisms with rTMS (Fried et al., 2017, 2016; Jannati et al., 2020, 2019, 2017; Oberman et al., 2010, 2016, 2014, 2012), we focused on the primary motor cortex in the left hemisphere. It is unlikely, however, that the observed differences in cTBS metrics of cortical plasticity between the two groups result from an ASD-related pathophysiological process specific to motor cortex. A structured neurological exam or medical history review found no evidence of gross or fine motor abnormalities in our ASD participants. Moreover, consistent with previous studies (Enticott

et al., 2010; Peter G Enticott et al., 2013; Peter G. Enticott et al., 2013; Minio-Paluello et al., 2009; Oberman et al., 2016, 2012; Théoret et al., 2005), baseline neurophysiological measures including RMT, AMT, and baseline MEP amplitude were comparable in the two groups, indicating that baseline M1 excitability, cortico-motor reactivity, and the function of the corticospinal pathway are not necessarily affected in ASD.

The greater and longer-lasting inhibitory response to cTBS is likely due to ASD-related abnormalities in the efficacy of plasticity mechanisms in the cortex. These include both classic LTD-like synaptic plasticity as well as nonsynaptic plasticity mechanisms, including biochemical and genetic changes (Tang et al., 2017). Although abnormalities in motor domain are not among the core symptoms of ASD, motor deficits among individuals with ASD have been reported, including alterations in motor learning (Sharer et al., 2016). Moreover, there is evidence to suggest abnormalities in motor function may precede higher-level social and communication impairments, including inferring others' intentions, gesturing, and imitation (Casartelli et al., 2016; Mostofsky and Ewen, 2011). It remains unclear whether these aberrant TMS-derived physiological responses are causal or a consequence of ASD pathology. It also remains to be investigated whether and to what extent these findings translate to non-motor cortical regions. Such investigations would require combining TMS and electroencephalography (EEG) or other neuroimaging techniques (Pascual-Leone et al., 2011; Thut et al., 2005; Thut and Pascual-Leone, 2010a, 2010b; Tremblay et al., 2019).

The present results build upon and expand the main finding of previous cTBS studies in adults with ASD (Oberman et al., 2016, 2012) that identified abnormal plastic regulation in the form of prolonged cTBS aftereffects in adults with ASD compared to NT adults. Maximum sensitivity and selectivity of cTBS measures of plasticity found in the present study were comparable with those found previously, 0.75 and 0.85, respectively (Oberman et al., 2012). One noticeable difference was the substantially earlier return to baseline of cTBS-modulated MEP amplitudes in the present study (by T20) compared to those found previously, i.e., T40–T60 in the ASD group (Oberman et al., 2012). This difference in pattern of cTBS aftereffects may be attributed to interindividual variability of cTBS responses, differences across datasets in proportions of *BDNF*, *APOE*, and other relevant SNPs, demographics, neuropsychological characteristics, or neuroactive medications (i.e., any drug that can influence neural activity in the central nervous system, whether as a stimulant or suppressant) in the ASD group. Moreover, the inter-study variability in the combined analyses likely resulted in the lower diagnostic utility of later time points compared to T15.

The present results can also be considered in the context of studies on aberrant TBS responses in children and adolescents with HF-ASD (Jannati et al., 2020; Oberman et al., 2014). Those studies showed a paradoxical, facilitatory response to cTBS in approximately one-third of children and adolescents with HF-ASD (Oberman et al., 2014) or at the group level relative to typically developing children (Jannati et al., 2020). Moreover, the extent of (Oberman et al., 2014) or the maximum inhibitory response to cTBS (Jannati et al., 2020) showed a maturational trajectory, increasing linearly up to the age of 16. Among the adult subjects in the present study (aged 21–65), the NT group showed a maturational trajectory,

with increasingly greater inhibitory cTBS response across the lifespan, whereas the ASD group did not show such a relationship. Assuming that the lack of significant correlations is not due to low power, these findings considered together with the greater and/or longer-lasting cTBS aftereffects observed in ASD adults (Oberman et al. 2012, 2016; present study) suggest a gradual change in the pattern of alteration in cTBS-derived plasticity measures in ASD across the lifespan relative to NT individuals. Individuals with ASD seem more likely to show a facilitatory response to cTBS in childhood and early adolescence. As ASD individuals grow older, they become more likely to show a greater inhibitory response to cTBS that may plateau by the age of 21. NT individuals show a more-inhibitory, but stable, response in childhood and adolescence, relative to individuals with ASD. In adulthood, NT individuals continue to show an increasingly inhibitory response to cTBS, perhaps due to the effects of normal aging on excitatory:inhibitory ratio in the cortex, which may be aberrant in ASD (Ben-Ari et al., 2012).

Two points are worth mentioning in comparing the present results to our previous cTBS results among healthy adults (Jannati et al., 2017). First, in that study (which included 21 of the 44 NT participants in the present study), we found T10 was the time point with the greatest explanatory power of interindividual variability in cTBS responses in healthy adults, whereas in the present study T15 was the strongest predictor of diagnosis, with little discriminability between ASD and NT groups at T10 (Figures 1–5). These contrasting results indicate that not only the most suitable post-TBS time point(s) to differentiate a given clinical population such as those with ASD from healthy individuals can differ from those in other clinical populations (e.g., McClintock et al. 2011; Tremblay et al. 2015; Fried et al. 2016), they can also differ from the time points that best characterize the gamut of TBS responses in healthy adults (Jannati et al., 2017) and across the adulthood (Freitas et al., 2011).

Second, we previously found AMT was a significant predictor of cTBS response in healthy adults (Jannati et al., 2017), whereas in the present study neither AMT nor RMT or baseline MEP amplitude were significant predictors of ASD diagnosis. These results suggest, while baseline neurophysiological measures can influence TBS responses among healthy individuals (Corp et al., 2020), they may not necessarily be a good differentiator of ASD and NT populations.

Interestingly, the finding that cTBS aftereffects retained their diagnostic utility across datasets that were obtained with two different types of TMS device (Magstim vs. MagPro) and pulse waveforms (monophasic vs. biphasic) suggest the utility of cTBS response as a physiologic biomarker is not necessarily susceptible to differences in TMS equipment and pulse-waveform characteristics (Davila-Pérez et al., 2018; Di Lazzaro and Rothwell, 2014).

4.2. The roles of BDNF and APOE polymorphisms in cTBS aftereffects

Factors contributing to inter- and intra-individual variability in TBS responses include the activation state of intracortical networks (Hamada et al., 2013), functional connectivity in the motor system (Nettekoven et al., 2015, 2014), state-dependent factors (Suppa et al., 2016), and SNPs that influence neuroplasticity, including *BDNF* (Antal et al., 2010; Chang et al., 2014; Cheeran et al., 2008; Di Lazzaro et al., 2015; Fried et al., 2017; Jannati et al.,

2019, 2017; Lee et al., 2013) and *APOE* (Jannati et al., 2019; Mahley and Rall Jr, 2000; Nichol et al., 2009; Peña-Gomez et al., 2012; White et al., 2001; Wolk et al., 2010).

The prevalence of *BDNFMet+* in the American population (AMR) population is estimated at ~28.2 % (1000 Genomes Project Consortium et al., 2015), whereas the prevalence of *APOE ε4+* is estimated at ~14.5% (Eisenberg et al., 2010), indicating approximate prevalence of 61.4%, 10.4%, 24.1%, and 4.1% for *Met-/ε4-*, *Met-/ε4+*, *Met+/ε4-*, and *Met+/ε4+* subgroups, respectively. The current results suggest, after adjusting for the relevant demographics (Table 4), predictiveness of cTBS aftereffects may be greater among *Met+/ε4-* and *Met+/ε4+* subgroups, followed by the *Met-* subgroups.

Compared to the logistic regressions based on continuous measures of cTBS response, subdividing the proportion of participants based on their type of cTBS response (inhibitor, non-responder, or facilitator) resulted in a less-sensitive measure, as it was only able to differentiate ASD and NT participants among *BDNFMet+* participants but not in the other subgroups.

The present results on the influence of *BDNF* and *APOE* SNPs on cTBS aftereffects have two main implications: (1) The existence and pattern of abnormality in cTBS responses in ASD adults are substantially modulated by both *BDNF* and *APOE* SNPs (Figures 2–5); (2) Limiting the analyses to each SNP subgroup of *BDNF* or *APOE* noticeably improves the predictive power of cTBS aftereffects in most, but not all, SNP subgroups. These two findings indicate that using the same logistic-regression model for differentiating all individuals with ASD from their NT counterparts is not optimal. Instead, we propose a hierarchical decision tree based on *BDNF* and *APOE* SNPs and perhaps other characteristics of a given individual with ASD (Tables 3 and 4). For example, if an individual with ASD is *BDNFMet-*, the next best step may be to compare his/her cTBS response with the cTBS responses of NT individuals who have a similar *APOE ε4* status, while also controlling for potentially important covariates. The post-cTBS time-point of interest in each step of analysis and estimations of the PPV and NPV of the cTBS biomarker may need to be chosen and adjusted accordingly, i.e., based on the specific SNP subgroup to which that individual belongs and perhaps demographic covariates such as race and/or level of education (Tables 3 and 4).

4.3. Additional considerations

A number of factors may limit the generalizability of the present findings. First, because our ASD sample was relatively small (n=19), it is likely there were heterogeneities among individuals with HF-ASD in demographics, clinical endophenotype, symptom severity, behavioral interventions and neuroactive medications, as well as underlying structural and functional brain differences that were not adequately represented in our sample. Such sampling errors could have reduced the representativeness of cTBS response in our ASD sample. We attempted to address these limitations by validating the present results in a relatively large, multi-site cohort including two previously obtained datasets. Such efforts are necessary for better assessment of the biomarker utility of cTBS, as well as enabling more-robust indices of target engagement and therapeutic response to experimental pharmacotherapy (e.g., Lemonnier et al. 2017) and potential rTMS treatments for ASD

(Cole et al., 2019). Larger samples of TMS data from healthy individuals (e.g., Corp et al. 2020) and clinical populations may also allow for evaluating multivariate models that take into account potentially important covariates such as demographic, neuropsychological, and neurophysiological measures as well as multiple relevant SNPs at the same time, enabling more-encompassing predictive models for individuals with ASD and those with other neuropsychiatric disorders.

Second, the representativeness of our ASD sample was limited in two ways: (1) because of lack of established feasibility and tolerability of rTMS in low-functioning individuals with ASD, all of our ASD participants were high-functioning; (2) because of the potential risk of rTMS-induced seizure, however small (Lerner et al., 2019), we excluded ASD participants with history of epilepsy. The prevalence of history of epilepsy in ASD can be as high as 26% (Viscidi et al., 2013). Moreover, a history of epilepsy is associated with more severe ASD symptoms, history of developmental regression, and poorer adaptive and language functioning (Viscidi et al., 2013). Thus, it is possible that if our ASD group included low-functioning individuals and/or those with a history of epilepsy, the overall pattern of cTBS responses in the ASD group would have been substantially different from present results. One option to study TMS measures of plasticity in individuals with ASD and current or history of epilepsy, while further reducing the potential risk of TMS-induced seizure, is to use TMS protocols such as paired associative stimulation (PAS) (Stefan et al. 2000; Ziemann 2004; see Suppa et al. 2017 for a recent review) that involve applying spTMS to the brain but still enable measuring TMS indices of plasticity in ASD (Jung et al., 2013).

Third, we were only able to stratify our sample by either *BDNF* or *APOE* SNP, as our sample size did not allow for robust assessment of the biomarker utility of cTBS aftereffects while stratifying simultaneously for both SNPs. Due to potential interactions between the effects of *BDNF* and *APOE* SNPs on rTMS plasticity metrics, the discriminatory power of cTBS aftereffects may differ substantially among those four SNP subgroups, even though, to our knowledge, such interactions have not been reported.

Fourth, a potentially important limitation of the present study was the absence of a sham control. Even though including a sham-TMS condition in purely neurophysiological studies such as the present work may seem not as critical as in studies with behavioral or psychological outcomes, it may nevertheless help assess possible natural fluctuations in corticospinal excitability that are not related to the rTMS intervention. For example, studies have found increased corticospinal excitability over time (Julkunen et al., 2012) or as a result of receiving successive single TMS pulses over the duration of a study visit (Pellicciari et al., 2016). Assuming these potential effects are similar between the ASD and NT groups, they would be canceled out when comparing their cTBS aftereffects, but still the inclusion of a sham-TMS condition would enable a more-robust assessment of post-cTBS changes in MEP amplitude that are not associated with the cTBS itself.

Fifth, another factor that could have influenced the difference in cTBS responses between ASD and NT groups is the potential effects of the various neuroactive medications received by all but three of the ASD participants. Due to the relatively small sample size and the disparate mechanisms of those neuroactive medications, controlling for them as a covariate

when comparing the cTBS responses between the two groups was not be statistically feasible. Future studies with larger cohorts may benefit from dividing ASD participants based on the type of neuroactive medications they receive and controlling for those medications as a covariate in multivariate models. A further valuable control would be to assess the same cTBS measures among non-ASD participants who receive the same or similar medications for other reasons.

Lastly, another limitation of our study was the lack of data on other SNPs that influence rTMS measures of brain plasticity, e.g., the catechol-O-methyltransferase (*COMT*) Val158Met SNP (Lee et al., 2014) that can interact with the effect of *BDNF* polymorphism on rTMS responses (Witte et al., 2012).

For cTBS measures of plasticity to ultimately become a valid and reliable physiologic biomarker for ASD, more consistent results are needed. Beyond controlling for relevant demographics, medications, and genetic variations known to influence neuroplasticity, possible future directions in this regard include assessing the effect of a range of stimulation intensities, including higher intensities than what has been used conventionally for TBS, conducting repeated stimulation sessions, and controlling for response to realistic sham-TMS, which itself may vary across individuals. Other future directions include combining TMS-EMG plasticity metrics with electroencephalography (EEG) and functional magnetic resonance imaging (fMRI) measures of TBS-induced plasticity, including modulation of TMS-evoked potentials and changes in resting-state functional connectivity (Eldaief et al., 2011; Farzan et al., 2016; Halko et al., 2014; Pascual-Leone et al., 2011).

4.4. Conclusions

The results of the present study show the utility of cTBS measures of M1 plasticity as a biomarker for adults with ASD, and the importance of controlling for *BDNF* and *APOE* SNPs in comparing those measures between ASD and NT individuals. The validation of the main findings from the present cohort in a larger multi-site cohort indicates the promise of TBS measures of plasticity as physiologic biomarkers for individuals with ASD.

Acknowledgement

We thank Stephanie Changeau, Aaron Boes, and Simon Laganier (BIDMC) for assistance with neurological examinations, and Ann Connor and Joanna Macone (BIDMC) for regulatory oversight and compliance, and for assistance with evaluation of participants' health and medical history.

This study was primarily funded by the National Institutes of Health (NIH R01 MH100186). A.P.-L. is further partly supported by the National Institutes of Health (R24AG06142, and P01 AG031720), the National Science Foundation, and the Barcelona Brain Health Initiative (La Caixa and Institute Guttmann), as well as sponsored research agreements with Neuroelectrics for home-based transcranial current stimulation and EGI for high-density EEG in cognitive disability. A.J. was further supported by postdoctoral fellowships from the Natural Sciences and Engineering Research Council of Canada (NSERC 454617) and the Canadian Institutes of Health Research (CIHR 41791). L.O. was further supported by the Simons Foundation Autism Research Initiative (SFARI) and the Nancy Lurie Marks Family Foundation. A.R. was further supported by the NIH (R01 NS088583), The Boston Children's Hospital Translational Research Program, Autism Speaks, Massachusetts Life Sciences, The Assimon Family, Brainsway, CRE Medical, Eisai, Neuroelectrics, Roche, Sage Therapeutics, and Takeda Medical. The content is solely the responsibility of the authors and does not necessarily represent the official views of the involved institutions or granting agencies.

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Highlights

- cTBS aftereffects are different in a large sample of individuals with ASD compared to neurotypical controls.
- T15 cTBS aftereffect allows to distinguish individuals with ASD diagnosis from neurotypical controls.
- Diagnostic utility of cTBS aftereffects for ASD is modulated by *BDNF* and *APOE* SNPs.

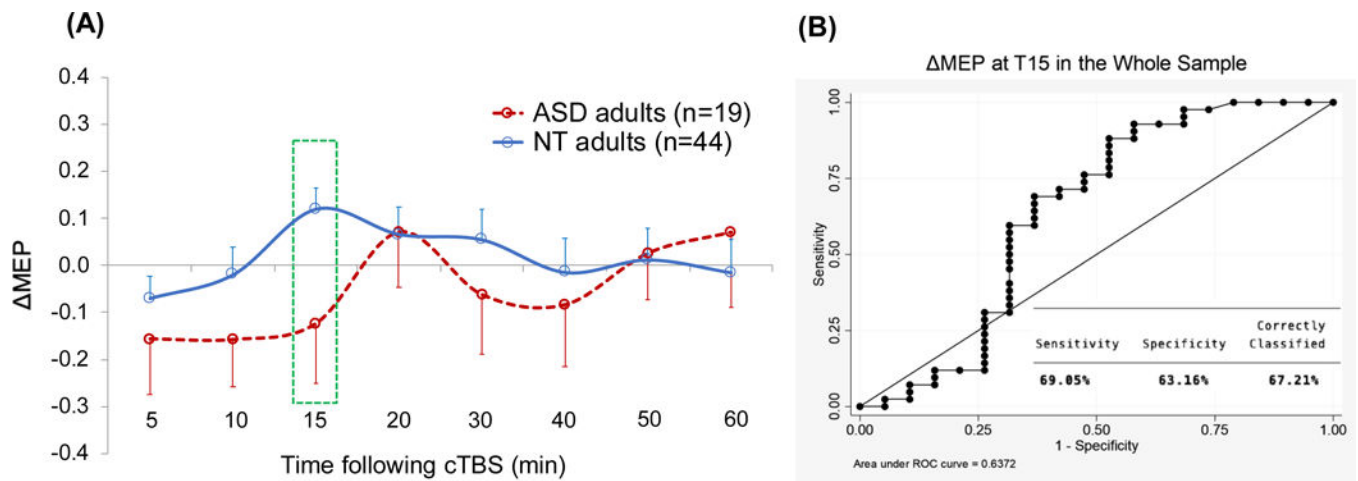


Figure 1.

(A) Grand-average change from baseline in MEP amplitude recorded from the right FDI muscle at 5–60 min following cTBS of the left motor cortex in ASD and NT groups. Error bars represent standard error of the mean. (B) The receiver operating characteristic curve associated with the logistic regression analysis with *Diagnosis* (ASD vs. NT) as IV and MEP at T15 as DV. ASD, autism spectrum disorder; cTBS, continuous theta-burst stimulation; MEP, natural log-transformed, baseline-corrected amplitude of motor evoked potential; DV, dependent variable; FDI, first dorsal interosseous; IV, independent variable; NT, neurotypical; Tn, *n* minutes post-cTBS.

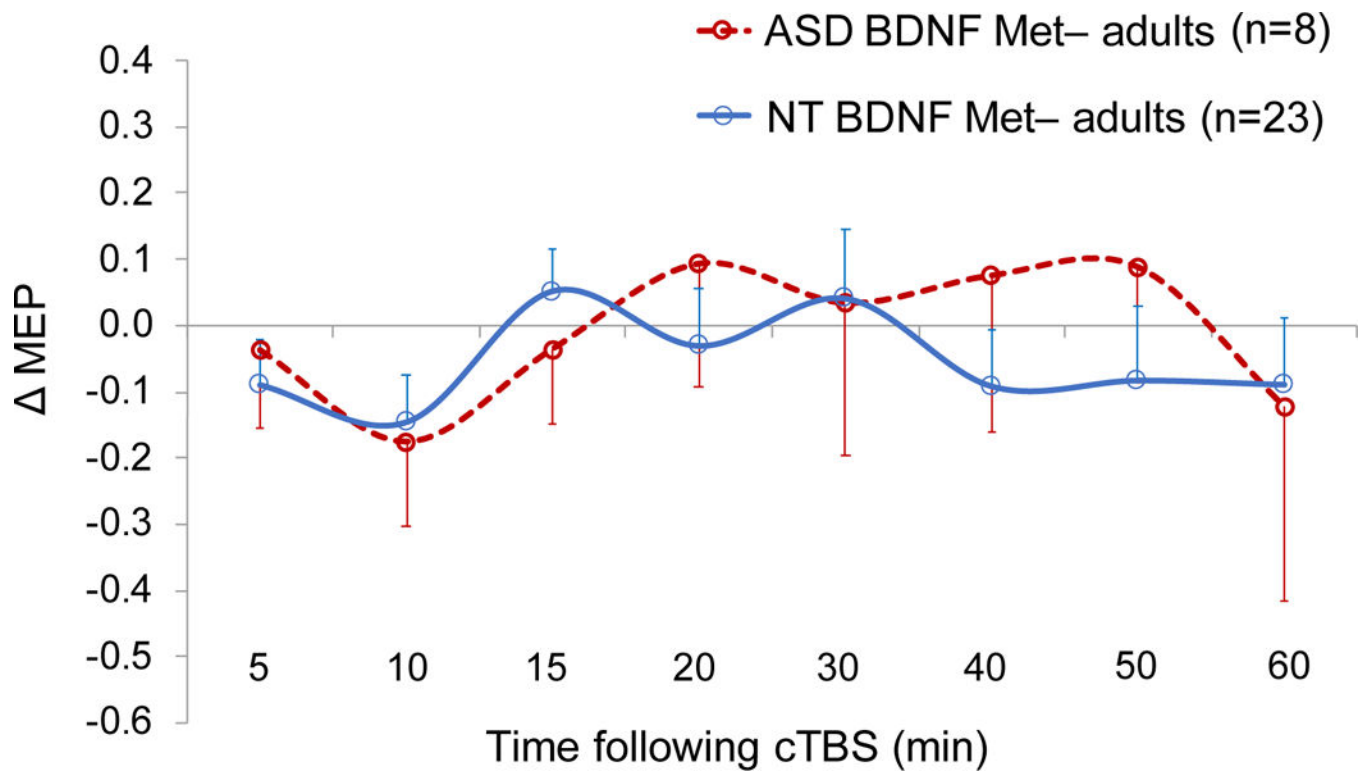


Figure 2. Grand-average change from baseline in MEP amplitude recorded from the right FDI muscle at 5–60 min following cTBS of the left motor cortex in ASD and NT subjects with *BDNF* Met– genotype. Error bars represent standard error of the mean. ASD, autism spectrum disorder; *BDNF*, brain-derived neurotrophic factor; cTBS, continuous theta-burst stimulation; MEP, natural log-transformed, baseline-corrected amplitude of motor evoked potential; FDI, first dorsal interosseous; Met, methionine; NT, neurotypical.

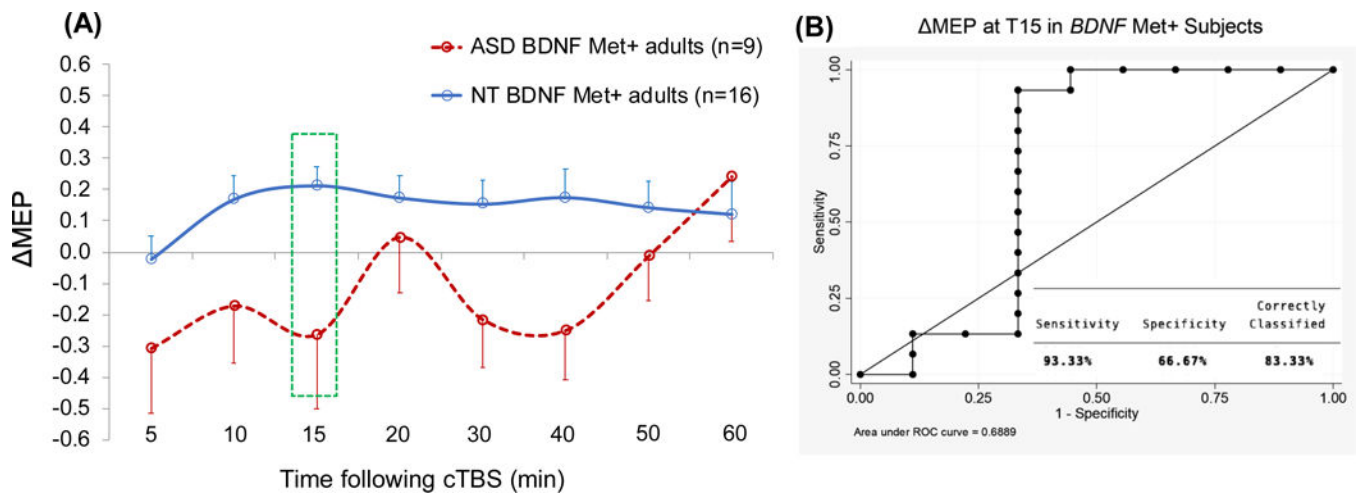


Figure 3.

(A) Grand-average change from baseline in MEP amplitude recorded from the right FDI muscle at 5–60 min following cTBS of the left motor cortex in ASD and NT subjects with *BDNF*Met+ genotype. Error bars represent standard error of the mean. (B) The receiver operating characteristic curve associated with the logistic regression analysis with *Diagnosis* (ASD vs. NT) as IV and MEP at T15 as DV. ASD, autism spectrum disorder; *BDNF*, brain-derived neurotrophic factor; cTBS, continuous theta-burst stimulation; MEP, natural log-transformed, baseline-corrected amplitude of motor evoked potential; DV, dependent variable; FDI, first dorsal interosseous; IV, independent variable; Met, methionine; NT, neurotypical; Tn, *n* minutes post-cTBS.

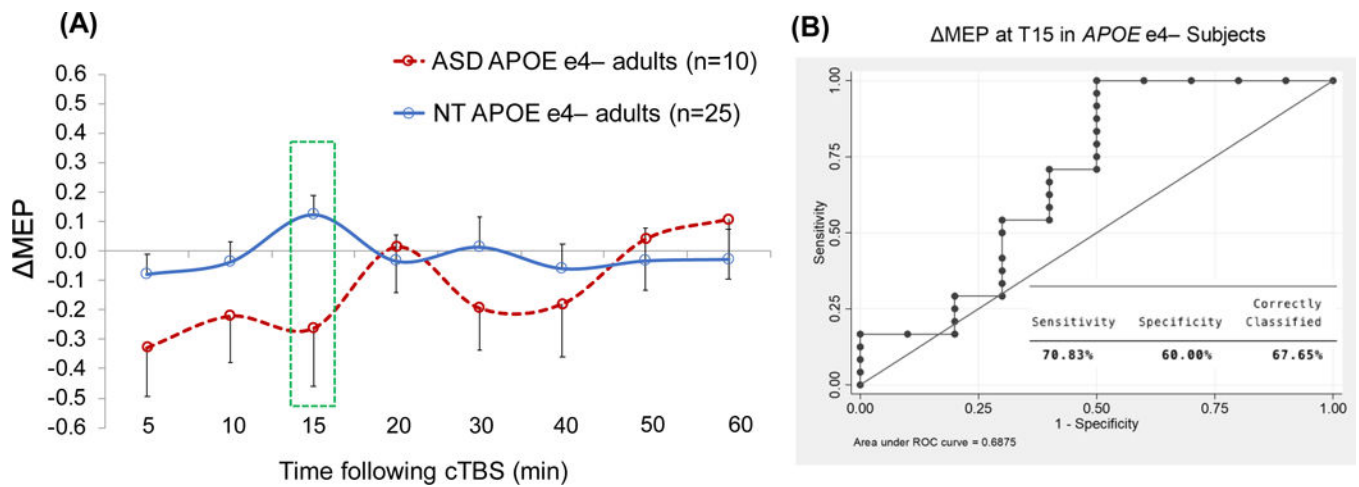


Figure 4.

(A) Grand-average change from baseline in MEP amplitude recorded from the right FDI muscle at 5–60 min following cTBS of the left motor cortex in ASD and NT subjects with *APOE* $\epsilon 4-$ genotype. Error bars represent standard error of the mean. (B) The receiver operating characteristic curve associated with the logistic regression analysis with *Diagnosis* (ASD vs. NT) as IV and Δ MEP at T15 as DV. *APOE*, apolipoprotein E; ASD, autism spectrum disorder; cTBS, continuous theta-burst stimulation; Δ MEP, natural log-transformed, baseline-corrected amplitude of motor evoked potential; DV, dependent variable; FDI, first dorsal interosseous; IV, independent variable; NT, neurotypical; Tn, n minutes post-cTBS.

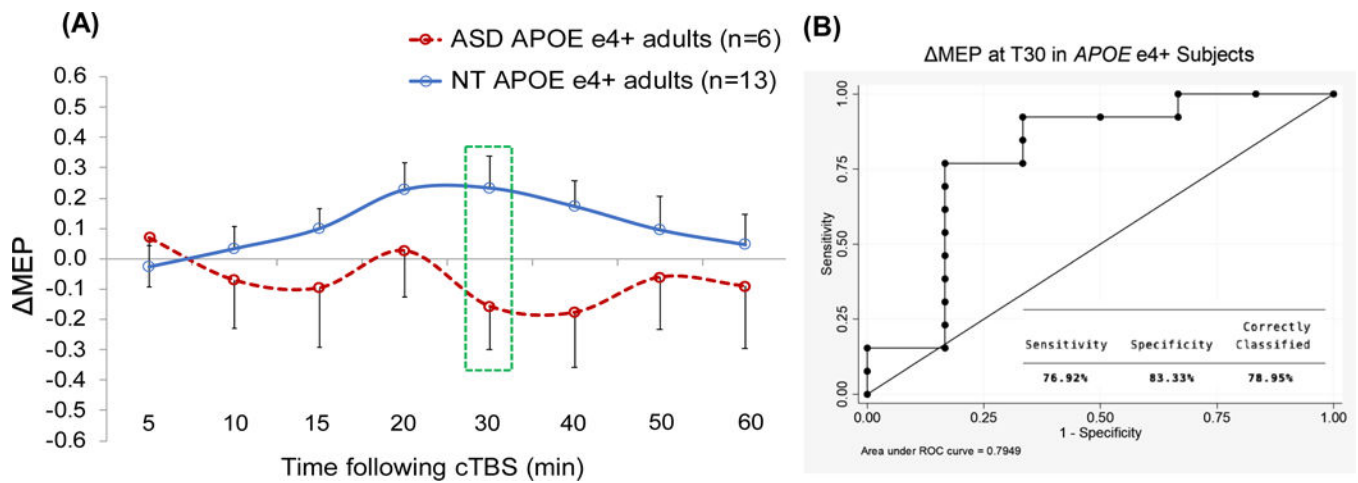


Figure 5.

(A) Grand-average change from baseline in MEP amplitude recorded from the right FDI muscle at 5–60 min following cTBS of the left motor cortex in ASD and NT subjects with *APOE* e4+ genotype. Error bars represent standard error of the mean. (B) The receiver operating characteristic curve associated with the logistic regression analysis with *Diagnosis* (ASD vs. NT) as IV and Δ MEP at T30 as DV. *APOE*, apolipoprotein E; ASD, autism spectrum disorder; cTBS, continuous theta-burst stimulation; MEP, natural log-transformed, baseline-corrected amplitude of motor evoked potential; DV, dependent variable; FDI, first dorsal interosseous; IV, independent variable; NT, neurotypical; Tn, n minutes post-cTBS.

Table 1.

Participants' demographics, neuropsychological measures, and single-nucleotide polymorphisms.

Group	Age	Sex	Race	Education* (yr)	Handedness	BDNF [†]	APOE [‡]	IQ	Verbal KN	Nonverbal FR	ADOS	AQ	Neuroactive medication(s)	Neurolog Psychia Comorb
<i>ASD (n = 19)</i>														
	21	M	Other	–	L	Val/Val	ε3/ε4	94	9	9	7	–	Risperidone, Methylphenidate	–
	21	M	Asian	13	R	Val/Met	ε3/ε3	124	16	12	15	26	Escitalopram	Anxiety, Depressio
	25	M	White	12	R	Val/Val	ε3/ε3	79	8	5	8	–	Amphetamine/ Dextroamphetamine	ADHD
	27	M	White	15	R	Val/Met	ε3/ε3	94	8	6	10	33	Risperidone, Bupropion	Anxiety, Depressio
	28	M	White	20+	R	Val/Met	ε3/ε4	121	16	11	7	20	–	Dysthymia
	31	M	White	–	R	Val/Val	ε3/ε3	79	9	4	17	–	Buspirone, Escitalopram	Anxiety, Depressio
	32	M	White	13	R	–	–	70	9	1	4	16	Aripiprazole, Hydroxyzine, Fluvoxamine, Methimazole	Anxiety, Depressio
	33	F	White	20	R	Val/Val	ε3/ε4	118	17	9	5	35	Sumatriptan	Depressio Migraine
	34	M	White	19	R	Val/Met	ε3/ε4	118	13	13	4	35	–	–
	37	M	Asian	18	R	Val/Met	ε3/ε3	115	15	10	16	21	–	ADHD, Depressio
	41	M	White	17	R	Val/Met	ε3/ε3	109	10	13	13	26	Acetyl-L-carnitine	Depressio Essential tremor
	41	M	White	16	R	Val/Met	ε3/ε3	88	4	12	8	17	–	–
	42	M	White	15	R	Val/Met	ε3/ε3	124	14	14	3	28	–	Anxiety, Depressio
	49	M	White	19	R	Val/Val	ε3/ε4	112	17	7	12	–	Aripiprazole, Zolpidem, Melatonin, Lisdexamfetamine	Anxiety, Depressio
	50	M	White	12	R	Val/Met	ε3/ε3	103	12	9	8	34	Citalopram	Anxiety, Depressio Fainting/ spells
	50	M	–	16	R	Val/Val	ε3/ε3	112	12	12	10	32	Sertraline	Social An Disorder
	52	M	White	–	L	Val/Val	ε3/ε4	118	19	7	3	43	–	ADHD, Anxiety
	52	M	White	11	R	–	–	100	9	11	5	36	–	Anxiety, Migraine
	58	F	White	16	R	Val/Val	–	121	18	9	10	36	Desipramine, Frovatriptan, Modafinil	Depressio Migraine
<i>NT (n = 44)</i>														

Group	Age	Sex	Race	Education* (yr)	Handedness	BDNF [†]	APOE [‡]	IQ	Verbal KN	Nonverbal FR	ADOS	AQ	Neuroactive medication(s)	Neurolog Psychia Comorb
	21	M	White	15	R	Val/Val	e3/e4	109	14	9	-	12	-	-
	21	M	Multiracial	16	R	-	-	112	11	13	-	7	-	-
	21	M	Asian	16	R	-	-	94	8	10	-	17	-	-
	22	M	Asian	17	R	Val/Met	e3/e4	103	9	12	-	16	-	-
	22	M	White	17	R	Val/Val	e2/e3	112	10	14	-	7	-	-
	22	F	White	16	R	Val/Val	e3/e3	100	10	10	-	6	-	-
	22	M	White	16	R	Val/Val	e3/e3	118	17	9	-	18	-	-
	23	M	Multiracial	16	R	Val/Val	e3/e3	106	11	11	-	20	-	-
	23	M	White	17	R	Val/Val	e2/e3	127	16	13	-	11	-	-
	23	M	White	17	R	Val/Met	e3/e3	118	13	13	-	22	-	-
	23	M	Black	14	R	Val/Val	e2/e3	103	9	12	-	8	-	-
	23	M	Asian	17	R	Val/Val	e3/e3	115	11	14	-	19	-	-
	23	F	Other	16	R	Val/Met	e3/e3	118	14	12	-	10	Sumatriptan	Migraine
	23	F	Asian	16	L	Met/Me t	e4/e4	103	14	7	-	19	-	-
	24	M	Asian	16	R	Val/Met	e3/e3	100	10	10	-	14	-	-
	24	M	Multiracial	17	R	Val/Met	e2/e3	115	12	13	-	12	-	-
	24	M	White	13	R	Val/Met	e3/e4	118	12	14	-	17	Cannabis	Anxiety, Depressio
	24	F	White	17	R	Val/Met	e3/e3	106	15	7	-	14	-	Fainting/ spells
	25	M	White	16	R	Val/Met	e3/e3	130	17	13	-	16	-	-
	25	M	Asian	18	R	Val/Val	e3/e3	91	7	10	-	17	-	-
	25	F	Asian	17	R	Val/Val	e3/e4	94	8	10	-	28	-	-
	25	F	Other	20	R	-	-	106	12	10	-	10	-	-
	26	M	Other	19	R	Val/Val	e3/e4	121	17	10	-	23	-	-
	28	F	White	-	R	Val/Val	e3/e3	100	10	10	-	16	-	-
	28	M	White	-	R	Val/Val	e3/e3	115	13	12	-	19	-	Headache Fainting/ spells
	28	M	Multiracial	18	R	Val/Val	e3/e3	100	9	11	-	13	-	-
	29	F	White	18	R	-	-	118	12	14	-	9	-	-
	30	M	White	20	L	Val/Met	e3/e4	97	10	9	-	13	-	-
	30	M	White	18	R	Val/Met	e3/e3	109	10	13	-	11	-	-
	32	M	White	19	R	Val/Val	e3/e3	100	9	11	-	14	-	Anxiety, Depressio
	32	M	Asian	20+	R	Val/Met	e3/e4	103	8	13	-	11	-	-
	45	M	White	16	R	Val/Val	e2/e3	94	9	9	-	14	-	-
	46	M	White	13	R	Val/Val	e3/e4	88	9	7	-	21	-	-
	47	M	Black	18	R	Val/Met	e3/e3	88	9	7	-	15	-	-
	47	M	White	20+	R	Val/Val	e3/e3	133	17	14	-	11	-	Migraine
	47	M	Black	18	R	Val/Val	e3/e4	118	16	10	-	11	-	-
	49	M	Asian	16	R	Val/Met	e3/e3	112	12	12	-	22	-	-

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Group	Age	Sex	Race	Education* (yr)	Handedness	<i>BDNF</i> [†]	<i>APOE</i> [‡]	IQ	Verbal KN	Nonverbal FR	ADOS	AQ	Neuroactive medication(s)	Neurolog Psychia Comorb
	50	F	Black	12	R	–	–	97	10	9	–	10	–	–
	51	M	White	16	R	Val/Val	e3/e3	115	14	11	–	21	–	–
	53	M	White	16	R	Val/Val	e3/e4	106	13	9	–	11	–	–
	56	F	Black	14	R	Val/Val	e3/e4	97	9	10	–	9	–	–
	56	M	White	16	R	Val/Met	e3/e3	109	13	10	–	16	–	–
	62	M	White	16	R	Val/Met	e3/e4	118	12	14	–	25	–	–
	65	M	Asian	20+	R	Val/Val	e3/e4	124	15	13	–	14	–	–

Participants in each group are sorted by age. Racial categories were defined according to the National Institutes of Health policy and guidelines on the inclusion of minorities as subjects in clinical research (NIH Office of Extramural Research, 2001). IQ scores were estimated using the Abbreviated Battery of Stanford-Binet IV intelligence scale. *APOE*, apolipoprotein E; AQ, autism-spectrum quotient; ASD, autism spectrum disorder; *BDNF*, brain-derived neurotrophic factor; IQ, intelligence quotient; Met, methionine; NT, neurotypical; Val, Valine. Values marked by “–” were either not reported by the participant or not available.

* Education data were available for 16 and 42 participants in the ASD and NT groups, respectively.

[†] *BDNF* data were available for 17 and 40 participants in the ASD and NT groups, respectively.

[‡] *APOE* data were available for 16 and 40 participants in the ASD and NT groups, respectively.

Table 2.

Participants' demographics, single-nucleotide polymorphisms, neuropsychological scores, and baseline neurophysiological measures.

	All (N = 63)	NT (n = 44)	ASD (n = 19)	<i>p</i>	NT Met- (n = 23) [‡]	ASD Met- (n = 8) [‡]	<i>p</i>	NT Met+ (n = 16) [‡]	ASD Met+ (n = 9) [‡]	<i>p</i>	NT e4- (n = 26) [§]	ASD e4- (n = 10) [§]	<i>p</i>	NT e4+ (n = 13) [§]	ASD e4+ (n = 6) [§]	<i>p</i>
Age (yr, mean ± <i>SD</i>)	34.57 ± 12.77	33.05 ± 13.15	38.11 ± 11.38	0.150	33.96 ± 13.69	39.88 ± 13.96	0.303	32.94 ± 3.31	35.67 ± 3.02	0.589	31.74 ± 2.14	36.50 ± 3.21	0.246	37.69 ± 4.57	36.17 ± 4.92	0.843
Sex (M : F)	51 : 12	34 : 10	17 : 2	0.318	19 : 4	6 : 2	0.634	13 : 3	9 : 0	0.280	22 : 4	10 : 0	0.559	10 : 3	5 : 1	1.000
Race (White : non-White)	37 : 26	22 : 22	15 : 4	0.050	13 : 10	6 : 2	0.433	8 : 8	7 : 2	0.229	16 : 10	7 : 3	0.716	5 : 8	5 : 1	0.141
Education (yr, mean ± <i>SD</i>) [‡]	16.52 ± 2.1	16.79 ± 2.09	15.81 ± 3.06	0.171	16.76 ± 2.07	16.60 ± 3.13	0.888	16.94 ± 1.95	16.22 ± 2.86	0.465	16.63 ± 1.56	14.89 ± 2.15	0.015	17.23 ± 2.65	19.75 ± 0.96	0.087
<i>BDNF</i> (Met- : Met+) [‡]	31 : 25	23 : 16	8 : 9	0.560	23 : 0	8 : 0	N/A	16 : 0	9 : 0	N/A	16 : 10	3 : 7	0.139	7 : 6	4 : 2	1.000
<i>APOE</i> (e4- : e4+) [§]	36 : 19	26 : 13	10 : 6	0.765	16 : 7	3 : 4	0.372	10 : 6	7 : 2	0.661	26 : 0	10 : 0	N/A	13 : 0	6 : 0	N/A
Handedness (Right : Left)	59 : 4	42 : 2	17 : 2	0.578	24 : 0	6 : 2	0.056	14 : 2	9 : 0	0.520	27 : 0	10 : 0	1.000	11 : 2	4 : 2	0.557
IQ (mean ± <i>SD</i>)	107.29 ± 13.03	108.18 ± 11.12	105.21 ± 16.82	0.411	107.63 ± 12.13	104.13 ± 17.56	0.533	109.19 ± 10.30	110.67 ± 13.17	0.758	108.11 ± 12.02	102.70 ± 17.00	0.286	108.54 ± 10.16	113.50 ± 9.99	0.334
AQ (mean ± <i>SD</i>) [¶]	18.42 ± 8.67	14.75 ± 5.13	29.20 ± 7.98	0.001	14.91 ± 5.58	36.50 ± 4.65	0.001	15.81 ± 4.34	26.67 ± 6.48	0.001	14.88 ± 4.59	27.13 ± 5.96	0.001	16.08 ± 6.03	33.25 ± 9.60	0.001
RMT (% MSO, mean ± <i>SD</i>)	35.65 ± 7.42	36.00 ± 8.29	34.84 ± 4.97	0.574	35.04 ± 7.28	34.88 ± 6.49	0.954	36.69 ± 10.21	34.22 ± 3.99	0.497	34.88 ± 8.89	33.40 ± 5.19	0.625	37.38 ± 7.77	34.67 ± 3.44	0.429
AMT (% MSO, mean ± <i>SD</i>)	26.52 ± 5.32	26.09 ± 5.61	27.52 ± 4.55	0.329	25.35 ± 4.60	28.00 ± 6.04	0.206	26.56 ± 6.81	26.33 ± 3.12	0.925	26.15 ± 5.94	26.40 ± 4.40	0.906	25.23 ± 4.90	26.67 ± 3.67	0.533
Baseline MEP amplitude (mV, mean ± <i>SD</i>)	1.34 ± 1.33	1.27 ± 1.31	1.51 ± 1.39	0.523	1.45 ± 1.58	1.54 ± 1.71	0.885	1.19 ± 1.03	1.57 ± 1.29	0.422	1.37 ± 1.58	1.86 ± 1.69	0.418	1.28 ± 0.88	0.83 ± 0.68	0.287

Racial categories were defined according to the National Institutes of Health guidelines (NIH Office of Extramural Research, 2001). Comparisons of proportions were conducted with Fisher's exact test. Education and single-nucleotide polymorphisms were not statistically compared between the subgroups because the data were not available for the total sample. N/A indicates that a statistical comparison between the two subgroups was not applicable because the measure of interest itself was the basis for subdividing the sample. Significant *p*-values are in bold font. The *p*-values were not adjusted for multiple comparisons. AMT, active motor threshold; *APOE*, apolipoprotein E; *APOE* e4+, e2/e4 or e3/e4 genotype; *APOE* e4-, e2/e3 or e3/e3; AQ, autism-spectrum quotient; *BDNF*, brain-derived neurotrophic factor; *BDNF*Met-, Val66Val; *BDNF*Met+, Val66Met; IQ, intelligence quotient; MEP, motor evoked potential; Met, methionine; MSO, maximum stimulator output; N/A, not applicable; RMT, resting motor threshold; *SD*, standard deviation; Val, valine.

[‡] Education data were available for 16 and 42 participants in the ASD and NT groups, respectively.

[‡]*BDNF* data were available for 17 and 39 participants in the ASD and NT groups, respectively.

[§]*APOE* data were available for 16 and 39 participants in the ASD and NT groups, respectively.

[¶]AQ scores were available for 15 and 44 participants in the ASD and NT groups, respectively.

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Table 3.

Changes in the logistic regression model predicting diagnosis after adding covariates.

	LR χ^2_{model}	P_{model}	β_{IV}	z_{IV}	P_{IV}	$\beta_{\text{covariate}}$	$z_{\text{covariate}}$	$P_{\text{covariate}}$	AUROC	PPV	NPV	% Dx
<i>T15 MEP</i>	4.79	0.029	– 1.55	–2.07	0.038	–	–	–	0.637	100.00%	72.41%	73.77%
<i>T15 MEP plus Education[†]</i>	4.94	0.084	– 1.32	–1.75	0.081	–0.18	–1.32	0.187	0.644	100.00%	78.43%	80.36%
<i>T15 MEP plus Race</i>	8.13	0.017	– 1.31	–1.74	0.081	1.15	1.75	0.080	0.704	88.89%	78.85%	80.33%
<i>T15 MEP plus BDNF[‡]</i>	6.04	0.049	– 1.76	–2.11	0.034	0.53	0.85	0.397	0.635	100.00%	75.51%	77.78%
<i>T15 MEP plus APOE[*]</i>	7.16	0.028	– 2.13	–2.41	0.016	0.20	0.31	0.756	0.699	100.00%	75.51%	77.36%
<i>T15 MEP plus Handedness</i>	5.54	0.063	– 1.58	–2.09	0.036	–0.94	–0.88	0.379	0.647	100.00%	75.00%	77.05%
<i>T15 MEP plus IQ</i>	5.85	0.054	– 1.63	–2.12	0.034	–0.02	–1.03	0.304	0.691	83.33%	74.55%	75.41%
<i>T15 MEP plus Age</i>	6.02	0.049	– 1.45	–1.93	0.053	0.02	1.11	0.266	0.668	71.43%	74.07%	73.77%
<i>T15 MEP plus Sex</i>	5.41	0.067	– 1.42	–1.86	0.063	–0.65	–0.76	0.447	0.664	100.00%	72.41%	73.77%
<i>T15 MEP plus Baseline MEP</i>	4.80	0.091	– 1.54	–2.01	0.044	0.02	0.11	0.915	0.639	100.00%	72.41%	73.77%
<i>T15 MEP plus RMT</i>	4.79	0.091	– 1.54	–2.00	0.045	–0.001	–0.03	0.976	0.639	100.00%	72.41%	73.77%
<i>T15 MEP plus AMT</i>	6.65	0.036	– 1.78	–2.27	0.023	0.08	1.35	0.177	0.677	66.67%	72.73%	72.13%

Models including a covariate are ranked based on their % Dx. *Diagnosis* was 0 for NT and 1 for ASD. IV refers to T15 MEP. *Sex* was 0 for males and 1 for females. *Race* was 0 for White and 1 for non-White. Racial categories were defined according to the National Institutes of Health guidelines (NIH Office of Extramural Research, 2001). *Handedness* was 0 for right-handed and 1 for left-handed. *BDNF* was 0 for Met– and 1 for Met+. *APOE* was 0 for $\epsilon 4-$ and 1 for $\epsilon 4+$. Significant *p*-values are in bold font. % Dx, percentage correctly classified; AMT, active motor threshold; *APOE*, apolipoprotein E; ASD, autism spectrum disorder; AUROC, area under the receiver operating characteristic curve; *BDNF*, brain-derived neurotrophic factor; IQ, intelligence quotient; IV, independent variable; LR, likelihood ratio; MEP, baseline-corrected, natural log-transformed amplitude of the motor evoked potential; Met, methionine; NPV, negative predictive value; NT, neurotypical; PPV, positive predictive value; RMT, resting motor threshold; T15, 15 minutes post-cTBS.

* *APOE* data were available for 16 and 39 participants in the ASD and NT groups, respectively.

[†] Education data were available for 16 and 42 participants in the ASD and NT groups, respectively.

[‡] *BDNF* data were available for 17 and 39 participants in the ASD and NT groups, respectively.

Table 4.

The logistic regression models within *BDNF* and *APOE* subgroups.

	LR χ^2_{model}	P_{model}	β_{IV}	z_{IV}	P_{IV}	$\beta_{\text{covariate}}$	$z_{\text{covariate}}$	$P_{\text{covariate}}$	AUROC	PPV	NPV	% Dx
<i>BDNF</i> Met – (n = 31; 23 NT, 8 ASD)												
<i>MEP</i> ^{T15}	0.50	0.480	-1.0	-0.69	0.490	–	–	–	0.571	–	73.33%	73.33%
<i>MEP</i> ^{T15} plus <i>Education</i> [†]	0.11	0.950	0.52	0.32	0.750	-0.002	-0.01	0.992	0.570	–	80.00%	80.00%
<i>MEP</i> ^{T15} plus <i>Race</i>	1.26	0.533	-1.09	-0.75	0.452	0.79	0.84	0.399	0.622	–	73.33%	73.33%
<i>BDNF</i> Met + (n = 25; 16 NT, 9 ASD)												
<i>MEP</i> ^{T15}	5.20	0.023	-2.12	-1.98	0.048	-0.50	-1.04	0.296	0.689	100.00%	78.95%	83.33%
<i>MEP</i> ^{T15} plus <i>Education</i> [†]	5.25	0.073	-2.09	-1.92	0.055	-0.05	-0.21	0.833	0.711	100.00%	83.33%	87.50%
<i>MEP</i> ^{T15} plus <i>Race</i>	5.90	0.052	-1.84	-1.68	0.093	0.86	0.83	0.408	0.682	85.71%	82.35%	83.33%
<i>APOE</i> e– (n = 36; 26 NT, 10 ASD)												
<i>MEP</i> ^{T15}	5.46	0.019	-2.11	-2.09	0.037	–	–	–	0.688	100.00%	77.42%	79.41%
<i>MEP</i> ^{T15} plus <i>Education</i> [†]	9.48	0.009	-2.03	-1.85	0.064	-0.63	-2.04	0.041	0.828	83.33%	84.00%	83.87%
<i>MEP</i> ^{T15} plus <i>Race</i>	5.48	0.065	-2.15	-2.06	0.040	-0.12	-0.14	0.892	0.671	100.00%	77.42%	79.41%
<i>APOE</i> e+ (n = 19; 13 NT, 6 ASD)												
<i>MEP</i> ^{T15}	1.67	0.197	-2.18	-1.21	0.228	–	–	–	0.705	100.00%	72.22%	73.68%
<i>MEP</i> ^{T15} plus <i>Education</i> [†]	4.42	0.110	-1.68	-0.86	0.388	0.63	1.51	0.131	0.789	–	73.33%	64.71%
<i>MEP</i> ^{T15} plus <i>Race</i>	4.81	0.091	-1.85	-1.06	0.288	2.02	1.61	0.108	0.808	66.67%	84.62%	78.95%
<i>APOE</i> e+ (n = 19; 13 NT, 6 ASD)												
<i>MEP</i> ^{T30}	3.69	0.055	-2.63	-1.62	0.106	–	–	–	0.795	100.00%	76.47%	78.95%

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	LR χ^2_{model}	P_{model}	β_{IV}	z_{IV}	P_{IV}	$\beta_{\text{covariate}}$	$z_{\text{covariate}}$	$P_{\text{covariate}}$	AUROC	PPV	NPV	% Dx
<i>T30</i> <i>MEP</i> plus <i>Education</i> [†]	5.63	0.060	−1.92	−1.30	0.193	0.63	1.40	0.163	0.827	50.00%	84.62%	76.47%
<i>T30</i> <i>MEP</i> plus <i>Race</i>	8.19	0.017	−2.93	−1.73	0.083	2.60	1.87	0.062	0.859	80.00%	1.20%	84.21%

Models in each genetic subgroup are ranked based on their % Dx. *Diagnosis* was 0 for NT and 1 for ASD. IV refers to MEP at T15 or T30, as specified. *Race* was 0 for White and 1 for non-White. Racial categories were defined according to the National Institutes of Health guidelines (NIH Office of Extramural Research, 2001). Significant *p*-values are in bold font. In each subgroup, the highest % Dx value of a significant model is in bold font. % Dx, percentage correctly classified; *APOE*, apolipoprotein E; ASD, autism spectrum disorder; AUROC, area under the receiver operating characteristic curve; *BDNF*, brain-derived neurotrophic factor; IV, independent variable; LR, likelihood ratio; MEP, baseline-corrected, natural log-transformed amplitude of the motor evoked potential; Met, methionine; NPV, negative predictive value; NT, neurotypical; PPV, positive predictive value; *Tn*, *n* minutes post-cTBS.

* *APOE* data were available for 16 and 39 participants in the ASD and NT groups, respectively.

† Education data were available for 16 and 42 participants in the ASD and NT groups, respectively.

‡ *BDNF* data were available for 17 and 39 participants in the ASD and NT groups, respectively.

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