

# Disrupted Activity in the Hippocampal–Accumbens Circuit of Type III Neuregulin I Mutant Mice

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Neuregulin I (*Nrg1*), a schizophrenia susceptibility gene, is involved in fundamental aspects of neurodevelopment. Mice lacking any one of the several isoforms of *Nrg1* have a variety of schizophrenia-related phenotypes, including deficits in working memory and sensorimotor gating, loss of spines in pyramidal neurons in the ventral subiculum, loss of dendrites in cortical pyramidal cells, loss of parvalbumin-positive interneurons in the prefrontal cortex, and altered plasticity in corticolimbic synapses. Mice heterozygous for a disruption in exon 7 of the *Nrg1* gene lack Type III (cysteine-rich-domain-containing) isoforms and have sensorimotor gating deficits that may involve changes in the activity of a circuit involving projections from the ventral hippocampus (vHPC) to medium spiny neurons in the nucleus accumbens (nACC). To explore the neural basis of these deficits, we examined electrophysiological activity in the nACC and vHPC of these mice. Under urethane anesthesia, bursts of spontaneous activity propagated from the vHPC to the nACC in both wild-type and mutant mice. However, these bursts were weaker in mutant nACC, with reduced local field potential amplitude and spiking activity. Single units in mutant nACC fired less frequently within the bursts, and more frequently outside of the bursts. Moreover, within-burst nACC spiking was less modulated by vHPC activity, as determined by phase-locking to the low-frequency oscillatory components of the bursts. These data suggest that the efficacy of vHPC input to the nACC is reduced in the Type III *Nrg1* heterozygotes, supporting a role for *Nrg1* in the functional profile of hippocampal–accumbens synapses.

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## INTRODUCTION

The nucleus accumbens (nACC) is a critical mediator of limbic motor computations. The nACC receives inputs from the ventral hippocampus (vHPC), prefrontal cortex, ventral tegmental area, and amygdala, and is thus in a privileged position to have a role in context-informed volitional behaviors (Groenewegen *et al*, 1987; Wright and Groenewegen, 1995; Mulder *et al*, 1998; Groenewegen *et al*, 1999; O'Donnell *et al*, 1999; Grace, 2000; French and Totterdell, 2002, 2003; Groenewegen and Trimble, 2007; Sesack and Grace, 2010). The interaction of these numerous afferents within the nACC has been implicated in circuits controlling sensory–motor gating and working memory, behaviors

that are typically disrupted in patients with schizophrenia (Braff and Geyer, 1990; O'Donnell and Grace, 1998; Freedman *et al*, 2003; Barch, 2005; Simpson *et al*, 2010).

Although the causal pathophysiology of schizophrenia is not well understood, it is clear that genetic susceptibility factors have an important role. One such factor is the gene for neuregulin 1 (*Nrg1*), which has been linked to schizophrenia in multiple, independent linkage studies (Stefansson *et al*, 2002; Harrison and Weinberger, 2005; Harrison and Law, 2006; Li *et al*, 2006; Munafo *et al*, 2006; Nicodemus *et al*, 2009). A complex, variably spliced molecule, *Nrg1* has been implicated in the regulation of diverse processes, including neuronal survival, neuronal migration, axonal pathfinding, and peripheral myelination (Michailov *et al*, 2004; Taveggia *et al*, 2005; Lopez-Bendito *et al*, 2006; Bjarnadottir *et al*, 2007; Krivosheya *et al*, 2008; Chen *et al*, 2010). Of the various classes of *Nrg1* isoforms, the Type III *Nrg1* variety is of particular interest. Type III *Nrg1* affects synaptic function by acting as both a ligand of and a receptor for ErbB receptor tyrosine kinases, inducing bidirectional signaling in a juxtacrine manner (Bao *et al*, 2003; Hancock *et al*, 2008). Mice with specific disruptions in exon 7 of the *Nrg1* gene (*Nrg1<sup>tm1Lwr</sup> +/–* or Type III *Nrg1<sup>+/-</sup>*)

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have decreased expression of Type III *Nrg1* isoforms, and are deficient in sensory–motor gating and spatial working memory behaviors (Wolpowitz *et al*, 2000; Chen *et al*, 2008), similarly to patients with schizophrenia (Thaker, 2000). Analyses of corticolimbic (eg, cortex to basolateral amygdala and vHPC to nACC) circuits *in vitro* reveal altered synaptic plasticity in Type III *Nrg1*<sup>+/-</sup> compared with control (Zhong *et al*, 2008). Given these *in vitro* findings and the behavioral impairments in sensorimotor gating and memory in Type III *Nrg1*<sup>+/-</sup> mice (Chen *et al*, 2008), we tested the hypothesis that disruption of Type III *Nrg1* signaling would alter the profile of vHPC–nACC circuits *in vivo*. Here we examine the effect of altered Type III *Nrg1* expression in vHPC–nACC circuits *in vivo* by recording simultaneously from the nACC and vHPC in urethane-anesthetized wild-type (WT) and Type III *Nrg1*<sup>+/-</sup> mice. Consistent with previous findings (Goto and O'Donnell, 2001a, b) we describe bursts of spontaneous neural activity in the nACC, which follow similar bursts in the vHPC in mice of both genotypes. Characterizing nACC unit and local field potential (LFP) data in and around these bursts, we show that decreased expression of Type III *Nrg1* disrupts the temporal organization of nACC activity, consistent with diminished efficacy of inputs from the vHPC to the nACC. These results support the notion that changes in functional connectivity in the vHPC–nACC circuit may underlie the deficits in sensory–motor gating and spatial working memory behaviors displayed by Type III *Nrg1*<sup>+/-</sup> mice.

## MATERIALS AND METHODS

### Animals

Type III *Nrg1* heterozygous mutant mice (*Nrg1*<sup>tm1Lwr</sup> +/–, referred to here as Type III *Nrg1*<sup>+/-</sup>) were back crossed to >99.9% C57Bl6 background. Mutant and WT littermates were generated by crossing Type III *Nrg1*<sup>+/-</sup> × Type III *Nrg1*<sup>+/-</sup> or Type III *Nrg1*<sup>+/-</sup> × C57/Bl6 WT as described (Wolpowitz *et al*, 2000). Animals were maintained on a 12-h light/dark cycle and provided with food and water *ad libitum*. Animal testing was conducted in accordance with the Principles of Laboratory Animal Care and the guidelines of the National Institutes of Health and was approved by the State University of New York at Stony Brook, Columbia University and New York State Psychiatric Institute Institutional Animal Care and Use Committees.

### Surgical Procedure

WT and Type III *Nrg1*<sup>+/-</sup> mice (30–40 g; *n* = 6 WT, 13 mutants) were anesthetized with 200 mg/kg urethane (Sigma, St Louis, MO), and a tracheostomy was performed for insertion of custom-made glass tubes (1.2 mm OD, WPI, Sarasota, FL) to maintain a patent airway. Animals were positioned in a stereotaxic device (David Kopf Instruments, Tujunga, CA) and the skull leveled. Burr holes were drilled (Foredom, Bethel, CT) over the nACC (+1 A/P, 1–1.5 M/L) and the vHPC (–3.9 A/P, 2.2–2.5 M/L). Core temperature was maintained at 37°C by a feedback-controlled device (Warner Instruments, Hamden, CT) and 100% humidified oxygen was blown past the tracheotomy tube. Animals were

allowed to equilibrate for ≥1 h before cell search began. nACC was searched by differential recording through two tungsten recording electrodes (5 MΩ, 0.01 inch; A-M Systems, Carlsborg, WA) glued together with cyanoacrylate. The electrode tips were configured in an offset manner in both the vertical and horizontal directions. One tip was ~400 μm lower, and ~400 μm lateral to the other, with the lower/medial tip placed in the nACC and the upper/lateral tip used for a reference. A similar offset recording/reference pair was positioned in the vHPC, except that the recording electrode was placed laterally and ventrally relative to the reference electrode. The use of different local references in each brain region to record the field potentials in the nACC and vHPC reduces the likelihood that the results can be accounted for by volume conduction or contamination of common noise from the reference electrode.

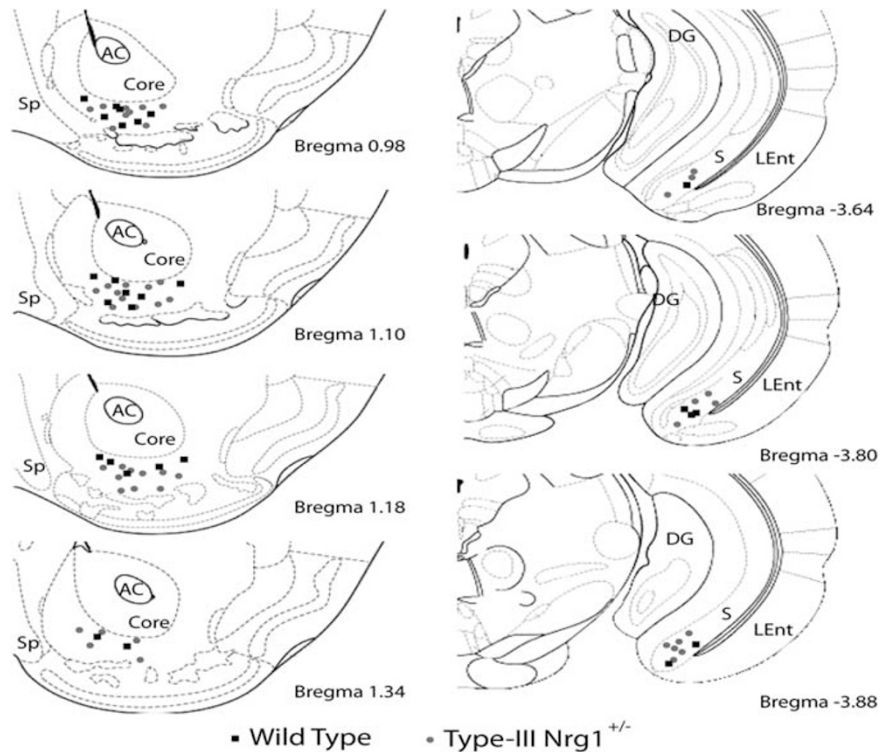
### Cell Recording Protocol

Electrodes were lowered into the target regions until spontaneously active units were identified. Successfully isolated units were recorded for a minimum of 15 min. For LFP recordings, data were filtered (0.5–200 Hz), and digitized at 2 kHz using Spike2 acquisition software (CED, Cambridge, UK). Cell data were acquired at 40 kHz using Spike2; spikes were threshold-detected with a minimum signal-to-noise ratio of 4:1. Waveforms were digitized (36 points per waveform). Individual waveforms were discriminated off-line using template-matching software (Spike2), which was then confirmed by Cluster Analysis (Spike2). LFPs were recorded from the same electrodes, Bessel filtered (0–100 Hz bandpass), and acquired at 2 kHz. After recording, anodal current was passed (20 μA for 1 min), marking the site of the recording. Only one recording site was utilized per animal. Animals were overdosed with urethane, and perfused transcardially with cold PBS followed by 10% formalin. After 10 min of perfusion fixation, the brain was removed and stored in 10% formalin overnight. Brains were then immersed in 30% sucrose in 0.1 M PBS until equilibration. Samples were sectioned (50 μm) in the coronal plane, and marked recording sites were visualized and plotted on standard sections. All cells included in this study were from the shell region of the nACC (Figure 1).

### Data Analysis

Cellular burst analysis was completed with a built-in Spike2 'Burst' algorithm. Parameters were set so that bursts comprised a minimum of three spikes, were initiated when instantaneous firing rate reached 10 Hz, and were terminated when firing rate decreased below 3 Hz. Bursts had to be separated by a minimum of 50 ms of silence. These parameters were selected by qualitative analysis of a representative selection of data and were determined to be non-restrictive, including most high-frequency events (Cooper and White, 2000) in the Type III *Nrg1*<sup>+/-</sup> mice.

nACC LFP recordings highlighted low-frequency, high-amplitude oscillations concurrent with spike bursts (Figure 2). These were termed 'LFP bursts'. LFP bursts were identified by a Spike2 thresholding algorithm that detected changes in the LFP that were biphasic, at least two

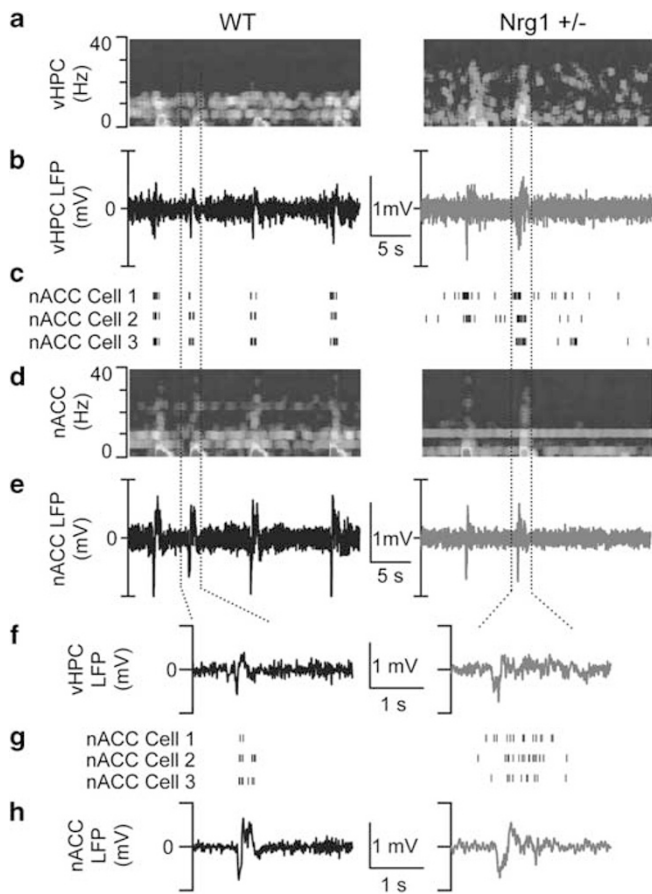


**Figure 1** Recording locations for all experiments. Recording locations from nACC (left panels) and vHPC (right panels) from WT (black squares) and Type III *Nrg1*<sup>+/-</sup> (red circles) mice superimposed onto standard atlas schematics. AP distance from the bregma in millimeter is indicated at lower right of each panel. Positive numbers are anterior to the bregma. AC, nACC; DG, dentate gyrus; LEnt, lateral entorhinal cortex; S, ventral subiculum; Sp, piriform cortex.

times the standard deviation of absolute LFP power, and began with a negative-going voltage component. After finding these LFP waves, the algorithm set the start time of the LFP burst as the zero-crossing point before the negative-going portion of the biphasic LFP waveform. Algorithm precision was cross-checked by direct inspection of all LFP burst start points by the investigator. Start points judged incorrect were reset by filtering the LFP signal from 0.5 to 4 Hz, re-running the thresholding algorithm and marking the start points as the time the filtered LFP crossed 0 mV. Peak and trough amplitudes of each LFP burst were detected and averaged to compare LFP burst sizes across genotypes. Peri-burst time histograms were constructed by using the start times of nACC LFP bursts to compile nACC cellular spike time histograms. LFP peri-burst power ratios compared the oscillatory power immediately previous to the start of the LFP burst with the power during the LFP burst. The 1-s time windows were constructed based upon the start times (see above) of each LFP burst yielding a 1-s ‘pre-burst’ period and a 1-s ‘burst’ period. Raw LFP waves were filtered for  $\delta$  (0.1–4 Hz),  $\theta$  (4–12 Hz), and  $\gamma$  (30–90 Hz). Power in the  $\theta$  and  $\gamma$  bandwidths was computed by a multitaper method (MATLAB (Mathworks, Natick, MA); seven non-overlapping windows) and  $\delta$  power was quantified as the size of the area under the filtered LFP for each 1 s period of time (pre and burst) for every LFP burst. Each peri-burst measurement was computed as a ratio (burst/pre-burst) to normalize for LFP amplitude differences. Results are stated as the average ratio by animal.

Spike phase-locking and LFP lag analysis were conducted using custom scripts written in MATLAB (Mathworks). The strength of single-unit phase-locking to  $\theta$  oscillations was assessed using Rayleigh’s *Z*-statistic of circular uniformity. To determine whether spikes were phase-locked to  $\theta$ , instantaneous,  $\theta$  phases of LFPs were determined through the Hilbert transform on the  $\theta$ -filtered vHPC LFP, and a phase was assigned to each spike based on the time of the spike’s occurrence. A phase of 0 refers to the trough of the  $\theta$  cycle as recorded, although the variability in placement of the electrodes precludes assigning absolute phase relative to known landmarks. Rayleigh’s *Z*-statistic values were obtained by the analysis of phase-locking of 100 randomly chosen spikes from a minimum of 900 s of experimental time. To ensure a random sample of spikes, this process was repeated 1000 times per experiment. The final *Z*-value reported is the average *Z*-statistic of these 1000 phase-locking iterations.

To determine whether vHPC leads nACC activity or vice versa, two different methods were used. First, to determine the temporal relationship between single-unit activity and vHPC  $\theta$  oscillations, phase-locking was calculated for 80 different temporal offsets (in 2.5 ms increments) for each unit recording. Units with significant Bonferroni-corrected phase-locking in at least 1 of the 80 shifts were used for the analysis in Figure 6c. No mutant units were significantly phase-locked using this stringent criterion. Negative lags indicate vHPC leading the nACC. Second, to determine the temporal relationship between the low-frequency



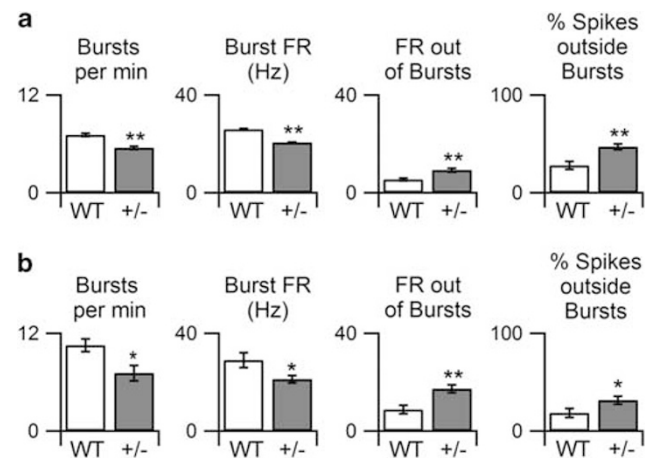
**Figure 2** nACC activity is disorganized in Type III *Nrg1*<sup>+/-</sup> mice. Spectrograms and LFP recordings from the vHPC (a and b) and nACC (d, e) reveal synchronous bursts of low-frequency power in both brain regions. (c) Raster plots of nACC unit activity. Note the staccato, bursting discharge pattern of nACC cells in the WT (left panels) and the less precise organization of spikes from nACC cells in the Type III *Nrg1*<sup>+/-</sup> mouse (right panels). (f–h) These simultaneous events are seen in expanded views showing the structure of the cellular discharge (g) and the LFP bursts recorded in the vHPC (f) and nACC (h).

components of the LFP, we filtered each LFP from 0.1 to 50 Hz and obtained the instantaneous amplitude using the Hilbert transform. We then cross-correlated the instantaneous amplitudes of vHPC and nACC and measured the lag to the peak of the cross-correlation (Adhikari *et al*, 2010). Negative lags indicate that vHPC leads nACC.

## RESULTS

### Disorganized Patterns of Neural Activity in Type III *Nrg1*<sup>+/-</sup> nACC

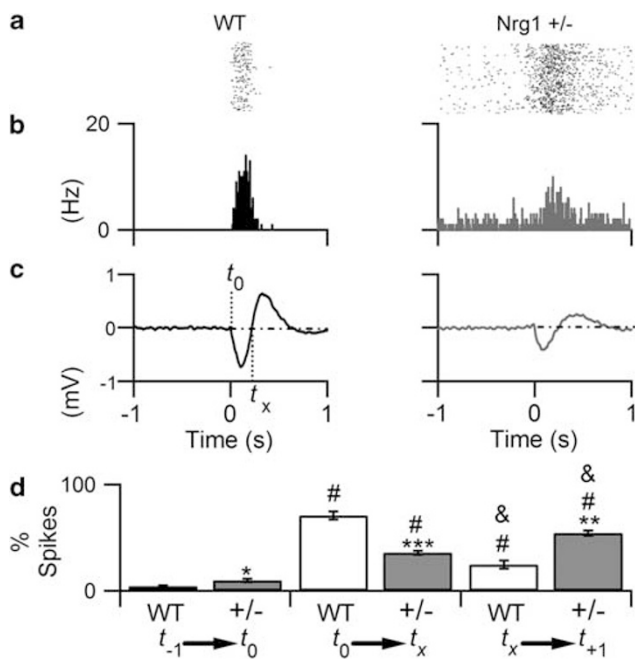
Recordings of neural activity were obtained from the nACC and vHPC of 6 WT and 13 Type III *Nrg1*<sup>+/-</sup> mice under urethane anesthesia. High-impedance electrodes and standard spike sorting techniques were used to record spontaneous activity from 21 WT and 35 mutant nACC neurons. Simultaneous LFP recordings were obtained from the CA1 pyramidal layer of the vHPC. Mean nACC neuron firing rates ( $19.8 \pm 2.2$  Hz for WT vs  $17.4 \pm 1.4$  for Type III *Nrg1*<sup>+/-</sup>,  $p = 0.5$ ) did not differ by genotype.



**Figure 3** nACC neuronal firing patterns are altered in Type III *Nrg1*<sup>+/-</sup> mice. (a and b) Left to right: Mean  $\pm$  SEM burst rate, intraburst spike rate, extraburst spike rate, and % of spikes occurring outside of bursts in WT (white bars) and Type III *Nrg1*<sup>+/-</sup> (gray bars) mice, for cell bursts (a), defined by short interspike intervals in the spike record, and LFP bursts (b), defined by the occurrence of increased low-frequency power in the LFPs. \* $p < 0.01$ , \*\* $p < 0.001$ .

Recordings of nACC and vHPC activity in WT animals suggested a high degree of temporal coordination across areas. The discharge pattern of nACC cells in WT animals was characterized by discrete staccato bursts of spikes interspersed with long, interburst pauses (Figure 2c and g; left-side panels). nACC spike bursts were accompanied by large amplitude, low-frequency LFP events (Figure 2d, e and h). Similar large-amplitude, low-frequency LFP events were recorded in the vHPC (Figure 2a, b and f), which coincided with LFP waves in the nACC. The LFP events in both locations consisted of a stereotyped bi- or triphasic waveform with high  $\delta$ - (1–4 Hz) and  $\theta$ - (4–12 Hz) range power (Figure 2a and d).

Although nACC and vHPC activity in Type III *Nrg1*<sup>+/-</sup> mice displayed a pattern of bursts of cellular activity accompanied by large, low-frequency LFP events that was similar to that of WT mice, the frequency and coordination of this activity was changed considerably in the Type III *Nrg1*<sup>+/-</sup> recordings compared with WT (Figure 2a–h; right-side panels). Activity bursts in the Type III *Nrg1*<sup>+/-</sup> mice appeared to occur less often, and spiking activity within the bursts appeared to be less tightly correlated with the LFP events in either area. To quantify these differences, two independent methods were used to identify activity bursts. First, spike bursts of nACC single units were categorized by an interspike interval algorithm classifying the start of a burst as the point where unit firing frequency rose above 10 Hz, and the end of the burst as the point where firing frequency fell below 3 Hz. Second, nACC LFP bursts were identified by an amplitude-thresholding algorithm (see Materials and Methods). Results yielded by both methods were similar. Bursts of activity in the nACC, whether distinguished by cell discharge frequency or LFP events, occurred less frequently in Type III *Nrg1*<sup>+/-</sup> mutants than in WT animals (Figure 3a and b). During these bursts, nACC single units from Type III *Nrg1*<sup>+/-</sup> mice discharged at a lower rate and produced fewer spikes per burst ( $6.9 \pm 0.1$  vs  $5.8 \pm 0.05$ ;  $p < 0.001$ ) relative to WT mice. In contrast, outside of these bursts, firing rates of nACC



**Figure 4** nACC neuronal burst firing patterns are altered in Type III *Nrg1*<sup>+/-</sup> mice. (a–c) Peri-burst firing patterns from representative WT (left panels) and Type III *Nrg1*<sup>+/-</sup> (right panels) units. (a) Raster plots of unit firing during 31 and 33 bursts for WT and Type III *Nrg1*<sup>+/-</sup> examples, respectively. (b) Peri-burst time histograms averaged over these bursts. (c) Average burst signal in the LFP, triggered at the initiation of the burst in the LFP and averaged over these bursts.  $t_0$ , burst start;  $t_x$ , first zero crossing after  $t_x$ . (d) Within-burst firing patterns in WT and Type III *Nrg1*<sup>+/-</sup> mice averaged over all units. Percentage of peri-burst spikes that occur within the 1 s preceding the LFP burst ( $t_{-1}$  to  $t_0$ ); between the beginning of the LFP burst and the first zero-crossing ( $t_0$  to  $t_x$ ), and from the first zero crossing to 1 s after the LFP burst initiation ( $t_x$  to  $t_{+1}$ ). \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ ; WT vs +/- Type III *Nrg1*<sup>+/-</sup>. # $p < 0.01$ ; vs ' $t_{-1}$  to  $t_0$ ' bin; and & $p < 0.01$ ; ' $t_0$  to  $t_x$ ' bin vs ' $t_x$  to  $t_{+1}$ ' bin.

neurons were significantly higher in Type III *Nrg1*<sup>+/-</sup> mice than in WT mice (Figure 3a and b). Relatively few spikes from WT cells were found to occur outside of bursts, whereas almost half of all discharge events in Type III *Nrg1*<sup>+/-</sup> units occur outside of bursts (Figure 3a and b). To ensure the reliability of the unit analysis, cells were pooled by animal and the above measures re-tested on a per animal basis. All results remained significantly different. Hence, the temporal organization of nACC spike bursting is disrupted in Type III *Nrg1*<sup>+/-</sup> animals as compared with WT mice.

To further explore the organization of nACC neuron firing, we examined the dynamics of unit discharge throughout the course of LFP events. Rasters and peri-burst time histograms time-locked to the start of LFP events (Figure 4a and b) revealed a temporally precise onset and rapid offset of firing in WT nACC units. nACC neurons from Type III *Nrg1*<sup>+/-</sup> mice, however, showed altered cell-LFP synchronization. Considerable firing outside of bursts occurred in the mutants, and the neurons took longer to reach peak firing rate and maintained elevated firing rates throughout and beyond the LFP event. To quantify this effect, we compared the relative firing rates of nACC neurons in WT vs Type III *Nrg1*<sup>+/-</sup> mice across the peri-burst period. For each individual LFP event, unit discharge was compared between three distinct periods describing the

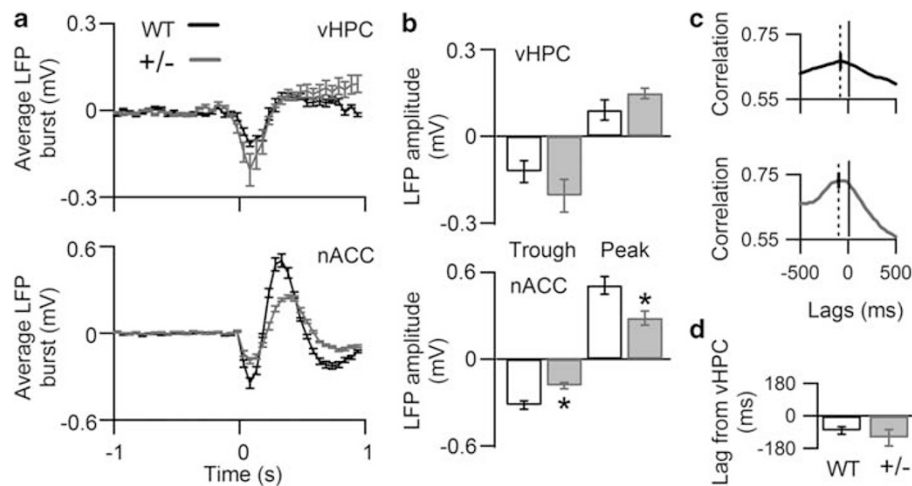
shape of the LFP event: the 1 s before the start of the LFP event ( $t_{-1}$  to  $t_0$ , period 1), from the start of the event to the subsequent zero mV crossing ( $t_0$  to  $t_x$ , period 2), and from the zero crossing to 1 s after the start ( $t_x$  to  $t_{+1}$ , period 3). Spikes in each period were counted and normalized to total spikes over the entire 2 s event period (Figure 4c and d). nACC units from Type III *Nrg1*<sup>+/-</sup> mice had higher activity before the start of the LFP events, yet had fewer spikes before the zero crossing, and more spikes after the zero crossing than neurons from WT animals. This template of activity is consistent with (1) increased extraburst firing, (2) slower onset, and (3) slower offset of spiking activity in *Nrg1* heterozygous mice compared with WT. Therefore, the composition of burst discharge patterns of nACC neurons is disorganized in Type III *Nrg1*<sup>+/-</sup> mice.

In addition to the evidence from nACC single-unit firing patterns, comparison of the simultaneously recorded LFP activity in WT and Type III *Nrg1*<sup>+/-</sup> mice also revealed significant differences in the mutants. As noted above, spiking activity was accompanied by stereotyped large amplitude LFP bursts in WT and Type III *Nrg1*<sup>+/-</sup> mice. However, nACC LFP bursts in Type III *Nrg1*<sup>+/-</sup> mice were considerably smaller than those in WT mice. The peak and the trough amplitude of the waves recorded in nACC of Type III *Nrg1*<sup>+/-</sup> mice were significantly reduced compared with those from WT mice (Figure 5a and b). In contrast, there was a nonsignificant trend toward an increase in the amplitude of LFP bursts recorded from the vHPC between genotypes (Figure 5a and b), indicating that the mutant animals appear to have decreased nACC activity that is not due to decreases in the strength of vHPC bursts. Similar results were obtained in the frequency domain, comparing LFP power during and immediately preceding the bursts (Supplementary Figure S1).

#### Decreased Functional Connectivity between the vHPC to the nACC in Type III *Nrg1*<sup>+/-</sup> Mice

The nACC receives direct monosynaptic input from the vHPC (Pennartz and Kitai, 1991; Mulder *et al*, 1998). We wondered whether the decreased nACC LFP burst amplitude and disorganization of spiking activity in the Type III *Nrg1*<sup>+/-</sup> mice might be the result of a deficit in the functional connectivity of vHPC inputs to the nACC. To explore this possibility, we examined the relationship between neural activity in the nACC and vHPC. We first explored the timing of the LFP bursts in the two regions. LFPs from the two regions were filtered for low frequencies (1–50 Hz) and instantaneous amplitude was calculated using the Hilbert transform. In both Type III *Nrg1*<sup>+/-</sup> and WT mice, cross-correlations of low-frequency amplitude envelopes (Adhikari *et al*, 2010) from nACC and vHPC showed a peak at negative lags, indicating that changes in amplitude in the vHPC precede those in the nACC (Figure 5c). Peak lags in the two genotypes were not different by *t*-test ( $p = 0.4$ ) (Figure 5d). This result suggests that under these recording conditions, the vHPC leads the nACC.

To further study the possible directionality of the circuit, we examined the relationship between nACC unit discharge and the  $\theta$ -frequency components of the vHPC low-frequency bursts. The  $\theta$ -frequency oscillations have been



**Figure 5** nACC LFP bursts reflect vHPC input and are weaker in Type III *Nrg1*<sup>+/-</sup> mice. (a) Grand average LFP bursts ( $\pm$  SEM) from the vHPC (top panel) and nACC (bottom panel) from WT (black lines) and Type III *Nrg1*<sup>+/-</sup> (gray lines) mice. (b) Mean  $\pm$  SEM amplitude of LFP bursts recorded from the vHPC (top panel) and nACC (bottom panel) in WT (white bars) and Type III *Nrg1*<sup>+/-</sup> (gray bars) mice at both the trough and peak of the burst. (c) Example cross-correlations of low-frequency (0–50 Hz) amplitude between the nACC and vHPC in WT (top) and Type III *Nrg1*<sup>+/-</sup> (bottom). Peak lags (dotted vertical lines) occur at negative values, indicating that changes in vHPC power precede changes in nACC power. (d) Mean  $\pm$  SEM lag between vHPC and nACC, calculated from the peak of the cross-correlations of low-frequency power in the nACC and vHPC. Difference is nonsignificant (see text). \* $p < 0.05$ , WT vs *Nrg1*<sup>+/-</sup>.

shown to influence neuronal firing in a variety of hippocampal downstream target regions in awake animals (Seidenbecher *et al*, 2003; Jones and Wilson, 2005; DeCoteau *et al*, 2007), although the relevance of the LFP bursts reported here to awake  $\theta$  oscillations is unclear. In both genotypes, nACC neuron firing was significantly modulated by the  $\theta$ -frequency component of the vHPC LFP bursts. Figure 6a shows the effect of this modulation: spikes from individual nACC cells that occur during LFP bursts are more likely to fire at specific phases of the  $\theta$ -frequency components of the vHPC bursts, that is, they tend to be ‘phase-locked’ to the vHPC LFP. The strength of this phase-locking can be quantified by computing the Rayleigh’s *Z*-statistic (see Materials and Methods); strong phase-locking results in higher *Z*-values. On average, nACC cells from Type III *Nrg1*<sup>+/-</sup> mice were more weakly phase-locked than those from WT mice (Figure 6b), consistent with the hypothesis of decreased functional connectivity between these two brain regions.

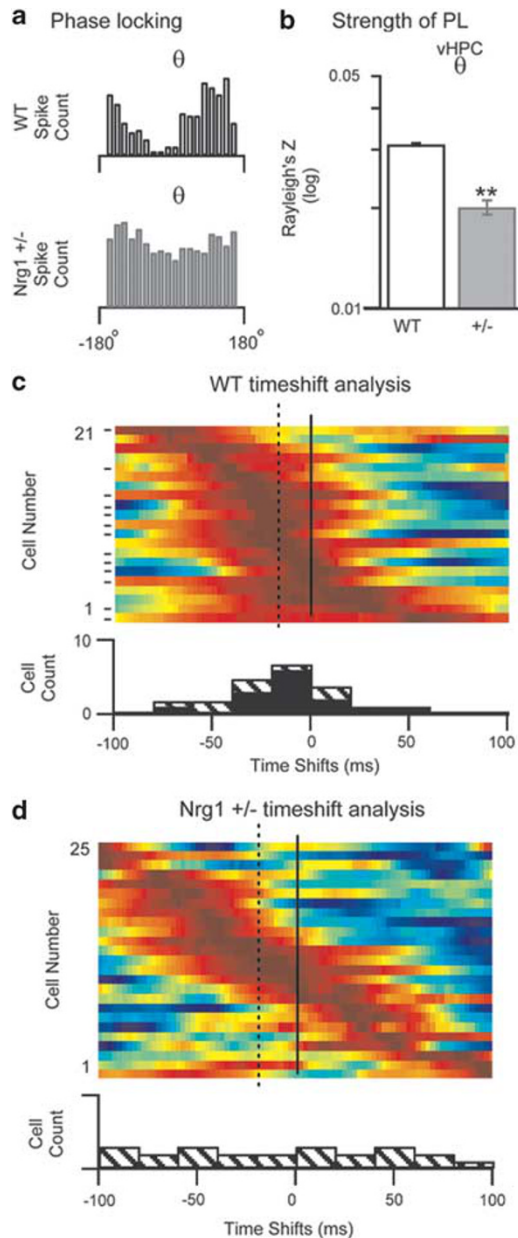
The cross-correlation of LFP amplitude envelopes suggests a predominance of vHPC to nACC directionality; the data are consistent with the notion that nACC activity follows activity in the vHPC. If this directionality were indeed accurate, then nACC cell discharge should be most strongly phase-locked to vHPC activity of the recent past (Jones and Wilson, 2005; Siapas *et al*, 2005). To examine this, phase-locking strength was computed after shifting the spike trains of nACC units in time, both backwards and forwards, relative to the vHPC LFP (Figure 6c and d). Spike trains were shifted in 2.5 ms steps over lags ranging from  $-100$  to  $+100$  ms. In WT animals, 13 of 21 nACC neurons were significantly phase-locked at any lag using conservative criteria ( $p < 0.05$  after Bonferroni correction for multiple comparisons). As predicted, the majority of cells phase-locked most strongly to vHPC LFP of the past, with a mean time shift at maximal Rayleigh’s *Z* of  $-17$  ms (Figure 6c). In the Type III *Nrg1*<sup>+/-</sup> mice, consistent with

the weaker phase-locking noted above, none of the neurons were significantly phase-locked after Bonferroni correction. Nonetheless, nACC units from Type III *Nrg1*<sup>+/-</sup> mice phase-locked more strongly to the vHPC LFP of the past (mean time shift  $-20$  ms, Figure 6d). Therefore, in animals of either genotype, nACC unit activity follows vHPC LFP activity. These results confirm the directionality within the vHPC/nACC circuit as suggested by the LFP results. Taken together with the evidence of decreased strength of phase-locking in the Type III *Nrg1*<sup>+/-</sup> mice, these results are consistent with a deficit in functional connectivity between the vHPC and nACC *in vivo*.

## DISCUSSION

Pursuant to the hypothesized role of circuits involving the nACC in complex behaviors disrupted in schizophrenia, we examined activity in the nACC and vHPC regions *in vivo* in mice carrying a heterozygous mutation in a gene that has been associated with increased susceptibility to schizophrenia, *Nrg1*. Both WT and Type III *Nrg1*<sup>+/-</sup> mice display bursts of synchronized activity in these regions, and the temporal pattern of this synchronized activity suggests that nACC activity followed that in the vHPC. In the Type III *Nrg1*<sup>+/-</sup> mice, the temporal structure of the nACC discharge was significantly different from that recorded in WT animals, and the strength of the modulation of nACC spiking by vHPC activity was significantly reduced. These findings demonstrate a role for Type-III *Nrg1* in regulating function within the vHPC/nACC circuit, and raise the possibility that decreased functional connectivity between the vHPC and nACC may underlie the sensory-motor gating deficits seen in Type III *Nrg1*<sup>+/-</sup> mice (Chen *et al*, 2008).

Although the precise nature of the spontaneous bursts of activity may be unclear, several lines of evidence suggest



**Figure 6** nACC unit response to vHPC input is weaker in *Nrg1*<sup>+/-</sup> mice. (a) Spiking of representative single units from WT (top) and Type III *Nrg1*<sup>+/-</sup> (bottom) neurons as a function of the phase of the  $\theta$ -frequency component of the LFP bursts recorded in vHPC. (b) Strength of phase-locking (PL) in WT (open bar) and *Nrg1*<sup>+/-</sup> (gray bar) nACC units. Mean  $\pm$  SEM of the Rayleigh's Z-statistic from of 21 and 25 units in WT and Type III *Nrg1*<sup>+/-</sup> mice, respectively. (c and d) Normalized phase-locking strength of each recorded nACC neuron as a function of lag. Each row represents an individual neuron ordered by the lag at which phase-locking is maximal. Note that nACC units are most strongly modulated by the vHPC  $\theta$  phase of the recent past in both WT (c) and Type III *Nrg1*<sup>+/-</sup> (d) mice. Horizontal lines on the left axis indicate units that demonstrate stronger phase-locking than expected by chance ( $p < 0.05$  after Bonferroni correction). Note that 13/21 WT and 0/21 Type III *Nrg1*<sup>+/-</sup> units are significantly phase-locked by this test. Bottom: Distribution of lags at which each cell was maximally phase-locked. Filled bars represent units with Bonferroni-corrected significant phase-locking. Mean lags at maximal phase-locking are  $-17.3$  and  $-19.1$  ms for WT and Type III *Nrg1*<sup>+/-</sup> units, respectively. Negative lags indicate phase-locking to the vHPC of the past.  $**p < 0.001$ , Type III *Nrg1*<sup>+/-</sup> vs WT.

that the disorganization of activity seen in the mutant nACC is the result of diminished functional input from the vHPC to the nACC. First, although the size of the vHPC LFP bursts are the same or larger in Type III *Nrg1*<sup>+/-</sup> mice when compared with WT, nACC bursts are dramatically smaller, whether measured as absolute amplitude or as the relative increase in low-frequency power. Second, the synchrony between the two areas is decreased in the Type III *Nrg1*<sup>+/-</sup> mice, as measured by the strength of phase-locking of nACC unit spikes to vHPC low-frequency activity. Third, in both genotypes, nACC activity follows that in the vHPC, as demonstrated using both phase-locking and amplitude cross-correlation methods. There are two possible mechanisms by which nACC activity would follow vHPC activity. Either the bursts of activity seen in the nACC are driven directly by input from the vHPC, or both areas are driven by a third area but with different latencies. Given the presence of monosynaptic connections from the vHPC to the nACC (DeFrance *et al*, 1985), the former explanation is the most parsimonious.

The known biology of Type III *Nrg1* is consistent with an important role in the regulation of neural activity in the nACC. Type III *Nrg1* is a membrane-bound protein expressed along axons and at presynaptic terminals of specific subsets of neurons (Michailov *et al*, 2004; Taveggia *et al*, 2005; Hancock *et al*, 2008; Zhong *et al*, 2008). Type III *Nrg1* has been shown to induce short- and long-term changes in cellular signaling cascades and activate downstream gene targets acting as both a ligand and a receptor with the ErbB family of tyrosine kinases. Thus, Type III *Nrg1*-ErbB interactions constitute a prime example of bidirectional signaling in the nervous system (Bao *et al*, 2003; Hancock *et al*, 2008; Mei and Xiong, 2008). Type III *Nrg1* is expressed in the vHPC, medial prefrontal cortex, and amygdala, all of which send converging inputs onto medium spiny neurons in the nACC (Groenewegen *et al*, 1999; French and Totterdell, 2002, 2003; Chen *et al*, 2008) ErbB4 is expressed postsynaptically on nACC medium spiny neurons (Steiner *et al*, 1999). *Nrg1*-ErbB4 signaling interactions are important in synapse formation, stabilization, and plasticity (Wolpowitz *et al*, 2000; Li *et al*, 2007; Woo *et al*, 2007), and the expression of Type III *Nrg1* in the vHPC is essential to vHPC-nACC synapses in particular. *In vitro* studies of vHPC slices and nACC neurons, in which vHPC axons from Type III *Nrg1*<sup>-/-</sup> (null), <sup>+/-</sup>, or WT mice innervate WT nACC neurons, demonstrate that functional synapses are formed regardless of presynaptic genotype. However, examination of nascent synapses formed by vHPC slices from Type III *Nrg1*<sup>+/-</sup> mice and postsynaptic WT nACC neurons *in vitro* reveals differences in the synaptic profile compared with their WT vHPC/ WT nACC counterparts (lower release probability, altered profile of postsynaptic glutamate receptor channels; C Du and LW Role, unpublished data). These data are consistent with the hypothesis that decreased *Nrg1* signaling at vHPC-nACC synapses may underlie the deficits in *in vivo* functional connectivity described here.

We find that vHPC activity is relatively preserved but that the nACC response to this activity is diminished in *Nrg1*<sup>+/-</sup> mice. The locus of the deficit in the heterozygote animals is unclear. This could be because of (1) a presynaptic deficit, reducing release of neurotransmitter onto nACC neurons,

(2) a postsynaptic deficit, reducing the response of nACC neurons to vHPC release of neurotransmitter, and/or (3) a nACC local circuit deficit, reducing the network response to vHPC input. Higher-resolution recording and analyses of synaptic function are required to address this important issue. However, we do find substantive deficits in vHPC/nACC circuits, whether pre- and/or postsynaptic in origin. Indeed, ongoing work using *in vitro* preparations supports the hypothesis that both pre- and postsynaptic are involved in altered synaptic profile of Type III Nrg1<sup>+/-</sup> vHPC/WT nACC connections (Zhong *et al*, 2008; and manuscripts in preparation by C Zhong, D Talmage, L Jiang, M Mertz, and L Role).

Although its behavioral relevance is not addressed directly by the current experiments, disorganized nACC activity may have a role in the sensorimotor gating deficits seen in Type III Nrg1<sup>+/-</sup> mice in which pre-pulse inhibition of acoustic startle is strongly disrupted (Chen *et al*, 2008). Findings suggestive of decreased hippocampal-accumbens connectivity have also been described in a neurodevelopmental model of schizophrenia neuropathology characterized by impaired pre-pulse inhibition (Moore *et al*, 2006). Indeed, functional amygdala, prefrontal cortical, and vHPC connections into the striatum are implicated in the regulation of striatal gating to auditory stimuli (Rosen *et al*, 1991; Swerdlow *et al*, 2000).

Understanding the impact of schizophrenia risk alleles on the function of neural systems is critical if the causal pathophysiology of schizophrenia is to be fully elucidated. Here we describe the impact of a heterozygous deletion of the cysteine-rich transmembrane domain-containing Type-III family of isoforms of Nrg1 on the function of the vHPC-nACC circuit. We found a disorganization of network activity in the nACC in mice carrying this deletion, and relate this disorganization to decreased functional connectivity between the vHPC and the nACC. The disruption in this circuit may underlie the behavioral deficits in sensorimotor gating, providing an important mechanistic inroad into future investigations into pathophysiological mechanisms known to operate in schizophrenia patients.

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## DISCLOSURE

The authors declare no conflict of interest.

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