

ZNF804A Protein Is Widely Expressed in Human Brain Neurons: Possible Implications on Normal Brain Structure and Pathomorphologic Changes in Schizophrenia

Hans-Gert Bernstein*, Johann Steiner, Henrik Dobrowolny, and Bernhard Bogerts

Department of Psychiatry, Medical School, University of Magdeburg, Magdeburg, Germany

*To whom correspondence should be addressed; Department of Psychiatry, Medical School, University of Magdeburg, Leipziger Str. 44, D-39120 Magdeburg, Germany; tel: +493916714249, fax: +493916715223, e-mail: Hans-Gert.Bernstein@med.ovgu.de

To the editor:

We read with great interest the recently published article by Schultz and colleagues¹ about the putative link between the expression of the zinc finger protein *ZNF804A* and the cortical structure in schizophrenia and wish to comment from a “human neuroanatomical” perspective on selected aspects of this topic. A single-nucleotide polymorphism (SNP) rs1344706 in the intron of the zinc finger protein *ZNF804A* (rs1344706) is a replicated genetic risk variant associated with schizophrenia (reviewed in Hill and Bray²). In the endeavor to learn more about the role of *ZNF804A* in schizophrenia, Schultz et al¹ performed a comprehensive allelic-specific gene expression study (homozygous risk allele AA carriers vs AC/CC carriers), combining magnetic resonance imaging cortical thickness and folding computing in genotyped patients with schizophrenia and healthy controls. Furthermore, they analyzed *ZNF804A* gene expression in a prefrontal region using postmortem tissue of schizophrenia patients and controls. It was revealed that in patients, AA carriers exhibited thicker cortex in prefrontal and temporal regions and less disturbed superior temporal cortical folding than AC/CC carriers, whereas the opposite effect was found in control cases. In addition, the expression analysis showed that the risk allele is associated with lower prefrontal *ZNF804A* expression in patients. This and other studies clearly imply that the expression of *ZNF804A* (and its gene variants) significantly impacts the developmental organization of human brain, both in health and disease. Indeed, the *ZNF804A* is thought to be prominently involved in brain development by influencing neural migration, neurite outgrowth, and the formation of synapses³ and was shown to be highly expressed in fetal (peaking in the second trimester) but not adult human brain.² To better understand the role of *ZNF804A* in normal and

disease-related brain development, it would be very important to know the regional and cellular expression of this protein in developing brain and adult human brain. Amazingly enough, this aspect has not been studied so far. We therefore immunohistochemically analyzed the expression patterns of *ZNF804A* protein in 4 adult human brains from psychically healthy individuals (3 males, 1 female; aged between 43 and 55 years) and 4 fetal brains (17th, 19th, 22nd, and preterm 28th gestational weeks). All brains were obtained from the Magdeburg brain collection.⁴ Case selection procedures, the acquisition of personal data, autopsies, and the handling of autoptic material were all conducted in strict accordance with the Declaration of Helsinki and were approved by the responsible Magdeburg Ethics Committee. In adult brains, we found a wide neuronal expression in the cortex (figure 1A), thalamus, hippocampus, hypothalamus, cerebellum (figure 1B), and brain stem. In mid-gestational brains, *ZNF804A* was highly expressed in multiple neuroblasts and radial glial cells (figure 1C). Our findings strongly support, from a neuroanatomical viewpoint, the notion that *ZNF804A* might be an important player in neurodevelopmental processes.^{1–3} However, its relatively high expression in mature human brain neurons leads us to assume that *ZNF804A* is relevant for adult brain functioning as well. Especially the strong expression of this protein in normal cortical neurons (together with a proposed altered neuronal expression pattern in schizophrenia) might well contribute to differences in cortical thickness between schizophrenia and control brains as observed by Schultz et al¹ and others. Whether or not the cellular expression patterns of *ZNF804A* is really altered in schizophrenia is currently completely unknown. We therefore are about to conduct an immunohistochemical study on *ZNF804A* localization in brains of schizophrenics and healthy controls with and without the SNP

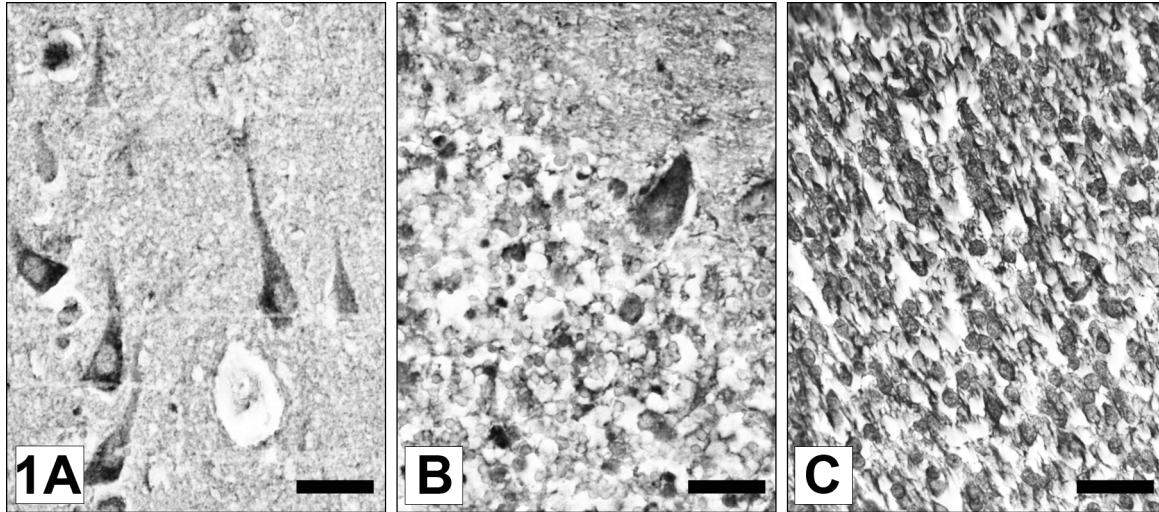


Fig. 1. Immunohistochemical demonstration of ZNF 804A in adult and developing human brain. (A) Temporal cortex of a 49 years old male. TNF 804A is expressed in layer III pyramidal cells and interneurons. (B) Cerebellum of a 55 years old female. A vast majority of granule cells and a few Purkinje cells were found to express the protein. (C) ZNF 804A expression in the cortex anlage of a fetal brain (19th gestational week). Multiple neuroblasts and radial glial cells are intensely immunostained. Scale Bars (valid for figures 1A–C) = 25 μ m.

rs1344706 to learn more about the cellular consequences of this mutation.

Acknowledgment

The authors have declared that there are no conflicts of interest in relation to the subject of this study.

References

1. Schultz CC, Nenadic I, Riley B, et al. ZNF804A and cortical structure in schizophrenia: in vivo and postmortem studies [published online ahead of print September 2008]. *Schizophr Bull.* 2013.
2. Hill MJ, Bray NJ. Evidence that schizophrenia risk variation in the ZNF804A gene exerts its effects during fetal brain development. *Am J Psychiatry.* 2012;169:1301–1308.
3. Hill MJ, Jeffries AR, Dobson RJ, Price J, Bray NJ. Knockdown of the psychosis susceptibility gene ZNF804A alters expression of genes involved in cell adhesion. *Hum Mol Genet.* 2012;21:1018–1024.
4. Bernstein HG, Stricker R, Dobrowolny H, et al. Nardilysin in human brain diseases: both friend and foe. *Amino Acids.* 2013;45:269–278.