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## Cardiovascular Psychiatry and Neurology

Cardiovascular Psychiatry and Neurology  
Volume 2009 (2009), Article ID 409562, 4 pages  
doi:10.1155/2009/409562

### Hypothesis

## Chromatin from Peripheral Blood Mononuclear Cells as Biomarkers for Epigenetic Abnormalities in Schizophrenia

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Received 15 April 2009; Accepted 1 June 2009

Academic Editor: Hari Manev

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### Abstract

**Background.** Studies have implicated abnormalities in epigenetic regulation in schizophrenia. We hypothesize that identifying abnormalities in chromatin structure in peripheral blood mononuclear cells (PBMC) from schizophrenia patients could (a) help identify genetic and (b) lead to targeted pharmacological interventions. **Testing.** We will examine the relationship between symptoms, demographics, hormonal fluctuations, substance abuse, comorbid illnesses, and epigenetic parameters in PBMC. In addition, we will examine the effects of mood stabilizers, as well as experimental agents both as clinically prescribed and as novel agents, on the effects of these agents on chromatin. **Implications.** If PBMC could be used to identify genetic mechanisms then this could lead to a “biomarker” approach to the diagnosis of schizophrenia. This approach may provide an informed method for the treatment of psychiatric disorders.

### 1. Background

Chromatin conformation regulates the access and attachment (repressors) to gene promoters. Deacetylation of histones, catalyzed by histone deacetylase (HDAC) enzymes, dimethylation of lysine 9 of histone 3 (H3K9me2), and DNA methyltransferase (DNMT) family of enzymes, are several examples of epigenetic marks that define a restrictive chromatin state both site specifically and globally [1], and play a key role in gene regulation. Because chromatin “plasticity” can be modified, the use of epigenetic marks to the regulation of gene expression in clinical populations is possible.

Increasing evidence indicates the existence of epigenetic gene regulation in schizophrenia. Supporting this hypothesis includes studies in which mice administered with antipsychotics show endophenotypes of schizophrenia [2], as well as studies using antipsychotics to modulate schizophrenia candidate genes through epigenetic mechanisms [3 - 5]. Several postmortem studies have found increased repressive enzymes including DNMT1 and HDAC1 in schizophrenia brains, suggesting abnormalities in the coordination of repressive processes [9]. Finally, studies of PBMC from human subjects have found reduced levels of the “open” chromatin mark H3K9me2 in schizophrenia patients compared to nonpsychiatric controls or to controls treated with “closed” chromatin mark H3K9me2 compared to controls [12] using HDAC inhibitors both as clinically administered [13] as well as

## 2. Presentation of the Hypothesis

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### 2.1. Understanding Epigenetic Gene Regulatory Abnormalities in PBMC

Unlike many illnesses in which investigators and clinicians are able to study the afflicted patient is still alive, psychiatry is limited by the inability to study the brain representing the higher order cognitive dysfunctions present in schizophrenia, making it impossible. Therefore, there exists a long history of searching for biomarkers of pathology within the brain. There are several factors which make epigenetic gene regulation in the brain.

Firstly, previous studies have demonstrated that PBMC can provide a window into environment/life experiences on chromatin structure and DNA methylation. Epigenetic parameters in lymphocytes from monozygotic twins at birth are virtually indistinguishable, while 3-year-old twins are virtually indistinguishable in terms of DNA methylation of histone 3, and acetylated histone 4, 50-year-old twins had significant differences. Older twins had greater differences in terms of epigenetic parameters. To note that these differences were consistent in subjects across all ages, the study of epigenetic parameters in lymphocytes is a reliable method for assessing the impact of environment on that chromatin from lymphocytes may provide a “molecular window” into environment, life experience, and stochastic factors which would not be possible to study in the brain.

Secondly, the analysis of gene regulation in nucleated blood cells provides a window into their disorder, including response to pharmacological, metabolic, and environmental factors. This approach for prospective longitudinal clinical research, and applicable to postmortem brain studies. This is because PBMC share much of the same environment as such as neurohormones, neuropeptides, chemo/cytokines, metal ions, and other epigenetically relevant substances present in the blood that are not present in the brain. Methionine, which both worsens psychosis in schizophrenia and is a precursor to homocysteine, which has been found to be elevated in the plasma of schizophrenia patients [17 - 20], and valproic acid, which significantly alter global chromatin structure in a dose dependent manner [18 - 20], are

Finally, PBMC contain the full complement of epigenetic enzymes : neurons and peripheral nucleated cells [21, 22]. Previous studies overall abnormalities in epigenetic mechanisms also thought to be disease, a disorder known to be associated with dysfunctions transcriptional repression across several chromosomes was found shown that peripheral markers are able to discern differences in illness [24 - 26], as well as show similarities in epigenetic parameters [24, 25].

Although, PBMC may be capable of reflecting overall changes in epigenetics for studying brain-specific processes such as synaptic plasticity a one gene at a particular time in a PBMC may not be reflective of the expression of a particular gene in a pyramidal cell at a given time or glial cell. In addition, there are some disorders in which abnormalities, for example, cancer cells have particular abnormalities that one would expect schizophrenia is characterized by brain-specific chromatin alterations in PBMC. Also, global DNA methylation tissue patterns have been found to be similar [28]. However, the differences between individuals or diagnostic groups are found to be present across tissues [23].

We now hypothesize that PBMC may be capable of reflecting changes in chromatin structure or DNA methylation. It may also help understand the effects of drugs of abuse on chromatin. Finally, it could provide a tool to both monitor and alter chromatin as well as identify patients most likely to benefit from these treatments.

There is now a growing literature to support the idea that PBMC reflect epigenetic machinery within an individual. In a previous study we showed that for four weeks with the only HDAC inhibitor approved for psychiatry, valproic acid (VPA) was shown to increase GAD67 mRNA expression in the brain [2]. In the serum levels of VPA there was a significant increase in GAD67 mRNA at the time that chromatin in PBMC from schizophrenia subjects was altered. This was confirmed when it was noted that there was less of an increase in histones 3 and 4 in schizophrenia subjects [13]. However, unexposed subjects had significantly lower levels of acetylated histone 3 compared to schizophrenia subjects. These associations with these measures of chromatin state and symptoms were stronger in schizophrenia subjects with higher baseline Young Mania Rating Scale (YMRS) scores. These subjects show greater increases in acetylated histone 3 after four weeks of VPA treatment. Baseline acetylated 3 levels showed greater improvement on the PANSS Syndrome Scale (PANSS). In other words, those subjects who had higher baseline levels alter their chromatin in response to VPA, and those subjects with higher baseline levels to respond favorably to VPA treatment [11].

As a consequence of these studies it was decided to attempt to use PBMC as a model. PBMC are held several advantages relative to clinical treatment using PBMC are protected from unforeseeable or undisclosed changes such as noncompliance, substance use, environmental exposures such as differences in metabolism of HDAC inhibitors. Further, the in vivo use of a known chromatin altering drug approved for use in psychiatric patients without altering medications to nonpsychiatric controls would not be ethical.

Similar to the previous studies in which systemic administration of VPA altered gene expression in human PBMC and in brains from mice, we found correlations between

cultured with equivalent doses of VPA [11]. Also in keeping with t to have an abnormally restrictive chromatin state as indicated b levels of the “closed’ ’ chromatin mark, H3K9me2 [10, 12]. Mo demonstrating the lack of “plasticity’ ’ in schizophrenia chroma we found less change in GAD67 mRNA expression, H3K9me2 level with an HDAC inhibitor compared to nonpsychiatric controls [10, 1 of onset had higher levels of H3K9me2 [12].

### 3. Testing the Hypothesis

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One means of testing the impact of particular environmental fac substances such as hormones, medications, or drugs of abuse anc structure. These findings could then be evaluated in light of find example, recent epidemiological evidence indicates that women mi data from PBMC cultures reveal sex differences in the ability of Therefore, it is a reasonable hypothesis that female sex horr schizophrenia risk through their ability to alter chromatin. The im be tested by culturing PBMC with estrogen, progesterone, or testos be compared to chromatin from animal and postmortem studies to tissues.

In addition, extensive studies could be conducted both in vivo and schizophrenia or characteristics of schizophrenia are associated w An examination of baseline levels of epigenetic parameters, suc epigenetic enzyme expression could be conducted using samp abnormalities exist. Previous studies indicate abnormalities i schizophrenia [9 - 11, 13]. Chromatin altering medications could b cultured with their PBMC as molecular probes into this system. altering agent clinically approved in psychiatry, this greatly limits t of determining those schizophrenia patients with abnormalities in e from the culture could be used to select subjects most likely established using in vitro methods, clinical trials could be implem epigenetic mechanisms could be treated with agents aimed at norr

### 4. Implications of the Hypothesis

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If PBMC, whether exposed to chromatin altering medications as c could serve as a reliable model of overall epigenetic mechanisms, revealing pathological chromatin state in schizophrenia. This app reaction of a subject’ s chromatin to medications prior to clinical method for selecting psychiatric medications for certain disorders. with symptoms and disease characteristics could lead to a better u potential for improved diagnostic validity more informed by a patie

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