

Translational Research on the Way to Effective Therapy for Alzheimer Disease

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Context: Alzheimer disease (AD) is a major public health issue with a prediction of 12 million Americans being affected by 2025 from the present 4 million. Molecular and genetic findings have provided significant insights into the roles that amyloid, tau, and apolipoprotein E isoforms have in the causation of AD. A central issue in AD pathogenesis is the amyloid cascade hypothesis. It states that abnormal amyloid processing and accumulation is the primary causative factor of AD and other associated neuropathologic abnormalities are of secondary consequence. It is presented to provide the rationale for novel drug and vaccination therapeutic strategies. Future research directed at prediction and prevention of AD through a genomic and proteomic analysis with identification of multiple polymorphic genes that interact, resulting in increased risk for late-onset AD, are the realistic and ultimate goals. A new approach for drug development is required, one that will emphasize a genomic and proteomic analysis to identify at-risk gene sets whose genetic expression is sufficient to cause late onset, sporadic AD. Prediction and prevention of disease prior to clinical signs and symptoms are the goals.

Objective: A review and analysis from electronic literature databases and subsequent reference searches of

the molecular genetic data, including pertinent genetic mutations and abnormal biochemical findings causal of AD, are cited. The amyloid cascade hypothesis, the contributions of apolipoprotein E, and hyperphosphorylated tau are discussed as to their roles in pathogenesis. Molecular targets for potential drug and vaccination therapies are cited from a critical assessment of the molecular and biomedical data. These data form the basis for rational, target-specific drug and vaccination therapies currently employed and planned for the near future. Phase 2 and 3 clinical trial results of drug and vaccination therapies are cited.

Conclusions: A new approach is needed as current pharmacologic therapy directed at symptomatic relief has proved to be marginally effective. The genomic and proteomic basis of AD will be defined in the near future, and corresponding molecular therapeutic targets will be identified. Genomic neurology has arrived and its application to resolving AD is our best hope.

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THE MOLECULAR AND GENETIC data related to early-onset Alzheimer disease (AD) causation are in strong support of the amyloid cascade hypothesis. In support of the amyloid cascade hypothesis is that amyloid vaccination therapy in selected patients may have shown slowing of cognitive loss and resolution of amyloid burden in autopsied brain. However, current pharmacologic therapy has been of limited benefit to slow effectively the cognitive loss in patients with AD. It is proposed that a genomic and proteomic analysis may be of value to identify at-risk gene sets whose genetic expression is sufficient to cause late-onset, sporadic AD. This information would provide essential data for the prediction of disease prior to the appearance of clinical signs and symptoms and

serve to direct genomic and proteomic research strategies to prevent it.

PATHOGENESIS OF DISEASE

Research in AD is proceeding at a rapid pace. Clinicians now have drugs that marginally ameliorate the cognitive and behavioral symptoms of AD. More effective therapies directed at the biological basis of disease pathogenesis are needed. Advances in knowledge of the molecular and genetic aspects of AD are providing therapeutic targets to attack more directly the molecular processes of disease.¹ Much of this knowledge comes from study of familial AD. Although the early-onset autosomal-dominant form of AD clearly results from specific genetic mutations, late-onset sporadic AD, representing 90% of patients, appears to result from multifac-

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Table. Genetic Aspects of Alzheimer Disease³⁻¹¹

Chromosome	Gene Effects	Functional Effects
1: autosomal dominant, early onset	Missense mutations of presenilin 2 gene	Increased synthesis and release of A β 42
14: autosomal dominant, early onset	Missense mutations of presenilin 1 gene	Increased synthesis and release of A β 42
19: risk factor, late onset	Inheritance of <i>APOE</i> ϵ 4 allele	Increased β amyloid aggregation
21: Down syndrome	Reduplication of <i>APP</i> gene	Increased APP processing; plaques and tangles by age 40 years
21: autosomal dominant, early onset	Missense mutations of <i>APP</i> gene	Increased synthesis and release of A β 40, A β 42

Abbreviations: A β 40, 40 amino acid length β amyloid protein; A β 42, 42 amino acid length β -amyloid protein; APOE, apolipoprotein E; APP, amyloid precursor protein.

torial environmental events and genetic influences, as inheritance of the apolipoprotein E ϵ 4 allele and other late-onset acting polymorphic genes.¹ The importance of nongenetic factors in AD is underscored by the fact that only one third of identical twins are concordant of the disease.² The environmental risk factors have remained elusive, but progress in finding genes causal of AD as well as increasing risk for it has been steady and impressive. Mutations in the amyloid precursor protein (*APP*) gene on chromosome 21q and of the presenilin 1 (*PS1*) and presenilin 2 (*PS2*) genes on chromosomes 14q and 1q, respectively, account for approximately one half of early-onset forms of autosomal-dominant inherited disease and about 3% to 5% of AD overall (**Table**). Additional gene loci that may increase the risk for AD have been identified on chromosomes 9, 10, 12, and 19.

Alzheimer disease may be considered as a form of amyloidosis resulting from the abnormal processing and intramembranous proteolysis of APP, a transmembrane protein whose function is unknown. The primary role of altered amyloidogenesis in the causation of AD has been convincingly supported.^{1,2,12-24} An increased synthesis of β amyloid peptide (A β) from APP in the early-onset forms of AD due to *APP* and *PS1* and *PS2* mutations is a central point in support of the amyloid hypothesis, which states that increased amyloidogenesis and/or decreased amyloid clearance with increased amyloid fibrillization are primarily causal of the pathogenesis of AD.²⁵⁻²⁷ Other molecular pathologic abnormalities, such as tangles composed of hyperphosphorylated tau, are of secondary importance. A more cautious view would be to say that amyloid deposition is necessary but not sufficient to cause the dementia of AD. Increased tau deposition facilitates A β toxicity.

Amyloid precursor protein processing involves 3 classes of enzymes: α -, β -, and γ -secretase. As shown in **Figure 1**, APP is first cut enzymatically by α - or β -secretase. The products of these first cleavages are cut again by γ -secretase, yielding a soluble fragment from the portion of the molecule produced by an α - γ cleavage and a self-aggregating fragment (β amyloid 40-42) from the portion of the molecule produced by the β - γ cleavage.

Environmental or other nongenetic factors can activate α secretase in sporadic AD, reducing levels of A β .²⁸ The insulin-like degrading enzyme insulin, neprilysin, and other enzymes as cathepsins are important for β amyloid degradation and clearance.²⁹ A balance between rates of A β synthesis and its degradation and removal from brain is a central research issue in late-onset

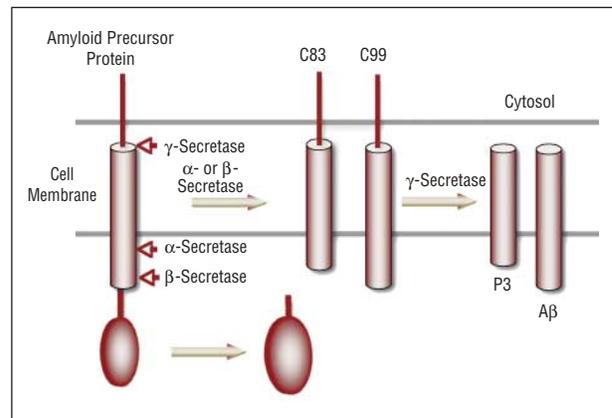


Figure 1. Amyloid precursor protein processing. Amyloid precursor protein is processed by α - or β -secretase to products C83 or C99, respectively. γ -Secretase processes C83 and C99 to P3 and A β , respectively. β Amyloid then can polymerize to form oligomers (A β)_n in the amyloid plaque.

AD. β amyloid increases selectively in brain with AD and not in other organs relates to the fact that β -secretase 2, which produces nonamyloidogenic fragments from APP, is active in nonneural tissue, thus reducing APP as a substrate for β -secretase in nonneural tissue to form A β .³⁰

Mutations in the *APP*, *PS1*, and *PS2* genes causal of AD (**Table**) have been studied in transgenic mouse lines and have provided important insights into the processes of amyloid deposition.³¹ The development of amyloid plaques in the APP 717 transgenic mouse depends on the expression of apolipoprotein E.³²

AMYLOID PLATELET BIOMARKERS

Amyloid precursor protein processing by platelets in patients with AD is different from that of control subjects.³³ There was partial normalization of the mean ratio of the 120- to 130-kD APP isoforms to the 110-kD isoforms in patients treated for 6 weeks with a statin drug, suggesting possible use of this biomarker in clinical trials in evaluating statins and other drugs.³⁴

PROTECTIVE GENES

Just as the apolipoprotein E ϵ 2 allele appears to delay the onset of AD, there may be other genetically determined factors that protect against AD. As the genetic degree of Cherokee Indian ancestry increases, the representation of AD decreases.¹ Choctaw Native American people have a later age at onset of AD with Mini-Mental Status Ex-

1. Amyloid containing (A β) extracellular neuritic plaques and intraneuronal neurofibrillary tangles (hyperphosphorylated tau) accumulate in the Alzheimer disease brain associated with progressive clinical dementia.
2. The degree of dementia correlates with the degree of synapse loss and neurofibrillary tangles and also with A β burden, although to a lesser degree.
3. In the autosomal-dominant forms of early-onset Alzheimer disease with mutations in the *APP*, *PS1*, or *PS2* genes, there is increased production of A β 40 and/or A β 42.
4. Aged patients with Down syndrome have increased A β synthesis and formation of typical Alzheimer disease plaques and tangles and may develop a dementia of Alzheimer disease type.
5. In sporadic late-onset Alzheimer disease, which represents the majority of patients, there is a decreased clearance of A β from brain and CSF with increased accumulation in brain as amyloid containing plaques and also as intracellular toxic oligomers. In early or prodromal Alzheimer disease, plasma A β levels decrease due to the decreased clearance of A β from brain and CSF. In late-onset Alzheimer disease, A β synthesis is not increased, as in the early-onset autosomal-dominant genetic forms of disease.
6. A β deposition in brain is a very early and toxic event in Alzheimer disease neuropathologic abnormalities and precedes clinical symptoms of Alzheimer disease, suggesting it is the prime pathogenic cause of Alzheimer disease. It is a central feature of the amyloid cascade hypothesis of Alzheimer disease pathogenesis.
7. Prevention and reversal of amyloid plaque formation with A β vaccination has been shown to correlate with behavioral improvement in transgenic mouse models of Alzheimer disease. Active A β vaccination in patients with Alzheimer disease resulted in slowing of cognitive loss and reduced brain amyloid burden.
8. Apolipoprotein E ϵ 4 is a risk factor for developing Alzheimer disease. It acts as a pathological chaperone aiding in the polymerization of A β into β -pleated sheets of amyloid in plaques. Apolipoprotein E with astrocytes also mobilizes and removes A β , which is impaired in transgenic mice carrying a mutant gene causal of Alzheimer disease and is a factor in increasing amyloid plaques in this experimental model.
9. Cell surface membrane receptor signaling involving APOE-R2, VLDLR, and LRP, and their ligands, including APOE, A β , APP, α 2-macroglobulin, and reelin, may be directly involved in Alzheimer disease pathogenesis due to altered signaling pathways misdirecting APP processing and/or A β degradation and clearance and the degree of tau hyperphosphorylation. Single nucleotide polymorphisms in the genes for proteins involved in neuronal cell surface membrane receptor signaling are genetic factors that may be linked to Alzheimer disease.
10. Late-onset Alzheimer disease results from the specific expression of genes that increase the risk for induction of the neuropathologic abnormalities causal of disease. Induction of the expression of this subset of genes is in part due to the effect of environmental vectors.

Figure 2. Standard model of Alzheimer disease: a synthesis of pathogenic mechanisms in support of the amyloid cascade hypothesis.^{36,37} A β indicates β amyloid; APOE, apolipoprotein E; APP, amyloid precursor protein; CSF, cerebrospinal fluid; LRP, lipoprotein receptor-related protein; VLDLR, very low-density lipoprotein receptor.

amination scores equivalent to those of white people, suggesting again a protective genetic influence. Increased vascular disease is a major factor for adding to the disease burden in the Choctaw patient and the emergence of AD.³⁵ These data indicate that genetic protective and vascular risk factors are involved in a variable manner resulting in AD in the Native American patient.

From the molecular and genetic data reviewed emerges a standard model for AD that includes its central pathogenic features (**Figure 2**) (modified from another article¹).

SYMPTOMATIC THERAPY

CHOLINESTERASE INHIBITORS

Cholinergic basal forebrain neurons that provide cholinergic innervation diffusely throughout the neocortex are especially vulnerable to the neuropathologic process of AD. Loss of cholinergic input due to neuronal forebrain degeneration is thought to be an important con-

tribution to cognitive loss in patients with AD. These observations led to the development of cholinesterase inhibitors to increase acetylcholine levels in brain by inhibiting the enzymes that metabolize it. Four cholinesterase inhibitors have been approved as therapy: tacrine hydrochloride, donepezil hydrochloride, rivastigmine tartrate, and galantamine hydrobromide.

The regimens of the 3 widely used cholinesterase inhibitors follow: donepezil hydrochloride, 5 mg daily at first and then 10 mg daily; rivastigmine tartrate, 1.5 mg twice daily at first and then 6 mg twice daily; galantamine hydrobromide, 4 mg twice daily and then 12 mg twice daily. Cognitive assessments of all 3 cholinesterase inhibitors have shown similar efficacy. There is a statistically significant slowing of cognitive loss using the Alzheimer's Disease Assessment Scale–Cognitive Subscale, and the 3 cholinesterase inhibitors have similar levels of response.³⁸

However, in a comprehensive assessment of donepezil by the AD 2000 consortium in 2004, it was found that donepezil produced no measurable reduction in rate of institutionalization or progress of disability. It was not found to be cost-effective because donepezil did not delay institutionalization sufficiently to offset the cost of the medication. The AD 2000 consortium found no evidence that costs of caring for patients with AD in the community are reduced by donepezil.³⁹

Presented at the Ninth International Conference on Alzheimer Disease and Related Diseases (July 2004), a study reported on the effects of donepezil compared with placebo to delay the time that patients with mild cognitive impairment developed AD. Seven hundred ninety patients were randomized, and 769 patients had initial, baseline evaluations. Patients were followed up for 3 years and about 30% dropped out prior to the end of the study. The important finding was that donepezil appeared to delay the onset of AD for about 6 months after which there were no significant differences between donepezil and placebo groups. Clearly, donepezil had positive effects on overall cognition, memory, and language tests related to those of the placebo group.⁴⁰

These studies, in general, indicate that donepezil reduces memory and cognitive loss during the prodromal or early phase of AD for several months.³⁸⁻⁴⁰ Behavioral and psychological symptoms associated with mild to moderate AD improved by 6.2 points on the Neuropsychiatric Inventory for patients receiving 10 mg of donepezil daily for 12 weeks compared with patients receiving placebo in a recently completed study in 2004.⁴¹ Cholinesterase inhibitor therapy provides clinical benefit to patients in the early phases of AD by slowing the rate of cognitive and memory abilities, and it is recommended for appropriate patients.

MEMANTINE

The Food and Drug Administration has approved memantine, an N-methyl-D-aspartate antagonist, for treatment of moderately advanced AD. It is proposed that its mechanism of action is to reduce potential glutamatergic excitotoxicity.

A study of memantine in moderate to severe AD

showed its effectiveness in the Activities of Daily Living Inventory and the Severe Impairment Battery but not the Global Deterioration Scale as compared with placebo.⁴²

It is prescribed at a dose of 5 mg daily and increased to 5 mg twice daily and then to 10 mg twice daily. In patients with moderate to severe AD receiving stable doses of donepezil, memantine resulted in significantly better outcomes than placebo on measures of cognition, activities of daily living, global outcome, and behavior and was well tolerated. It is suggested by these positive data that memantine offers a new approach for therapy of patients with more advanced disease⁴³ and may be administered together with cholinesterase inhibitor therapy in appropriate patients.

HORMONE REPLACEMENT THERAPY

The Women's Health Initiative Study of estrogen and medroxyprogesterone demonstrated an increased occurrence of dementia in postmenopausal women.⁴⁴ Estrogen therapy alone in postmenopausal women did not reduce dementia or incidence of mild cognitive impairment and increased risk for both conditions combined.⁴⁵ Further for women aged 65 years or older, estrogen therapy had an adverse effect on cognition.⁴⁶ It is now clear that hormone replacement therapy has adverse effects, increasing the risk for dementia and impaired cognition. Use of hormone therapy to prevent dementia or cognitive decline in women 65 years of age or older is not recommended.⁴⁴⁻⁴⁶

ANTIOXIDANTS

One significant study is a randomized, placebo-controlled clinical trial using selegiline (N,2-dimethyl-N-2 propynyl phenethylamine hydrochloride, an irreversible inhibitor of monoamine oxidase B), vitamin E, and both agents together compared with placebo in patients with AD. The one positive result was that the time to nursing home placement, the time to death, and the time to severe dementia were extended in selegiline, in vitamin E, and in combined groups compared with patients receiving placebo.⁴⁷ Vitamin E at 2000 international units a day was recommended based as a modest means to slow the progress of the disease.⁴⁷

However, the Cochrane Dementia Group Register of Clinical Trials was searched for effectiveness of vitamin E in AD in 2000. It was concluded that there is insufficient evidence of efficacy of vitamin E in the treatment of AD.⁴⁸

Use of vitamin E and vitamin C supplements in combination in the Cache County Study was associated with reduced prevalence and incidence of AD and thus merits further study for the primary prevention of AD.⁴⁹

ANTI-INFLAMMATORY DRUGS

Increased acute phase reactants including cytokines and interleukins and minor signs of cellular inflammation in patients with AD have resulted in clinical trials to study the effectiveness of anti-inflammatory drugs. Naproxen (a cyclooxygenase-1 and -2 inhibitor) and rofecoxib (a

cyclooxygenase-2 inhibitor) have been studied. A multicenter, randomized double-blind, placebo-controlled parallel group trial with 1-year exposure to rofecoxib or naproxen was conducted with patients with mild to moderate AD. The primary outcome measure was the 1-year change in the Alzheimer Disease Assessment Scale-Cognitive Subscale score. The results of this study indicated that rofecoxib or low-dose naproxen did not slow cognitive decline.⁵⁰ These drugs do not reduce the rate of cognitive loss nor functional decline in patients with AD.⁵⁰ A prospective, placebo-controlled, double-blind prevention trial with an anti-inflammatory agent has not been completed.

CHOLESTEROL LOWERING THERAPY

Statins, 3-hydroxy-3-methylglutaryl coenzyme A (HMG CoA) reductase inhibitors, may prevent AD. Evidence for a protective effect of HMG CoA reductase inhibitors has been provided by comparing the prevalence of the diagnosis of AD in 3 groups of individuals from Veterans Administration hospital records. It was found that the cohort of individuals receiving HMG CoA reductase inhibiting drugs had a prevalence 60% to 73% lower than that of either the total patient population or patients taking other medications typically used to treat hypertension or cardiovascular disease.⁵¹ In another retrospective study, persons aged 50 years or older who were prescribed HMG CoA reductase inhibitors had substantially lower risk of developing dementia, independent of the presence or absence of untreated hyperlipidemia or exposure to non-HMG CoA reductase lipid-lowering agents. The adjusted risk ratio for those receiving HMG CoA reductase inhibitors was 0.29 (confidence interval, 0.13-0.63; $P = .002$).⁵² These data from retrospective medical-record reviews need to be validated with prospective, double-blind, placebo-controlled trials with statins.

Recently, a prospective, randomized, dose-finding, 36-week treatment trial with statins (simvastatin or atorvastatin) was conducted with 39 patients with hypercholesterolemia. Plasma levels of A β 40 and A β 42 were measured. Both statins reduced total plasma cholesterol levels by 56%, but levels of A β 40, A β 42, and total A β were unchanged. Thus, this study does not support the effect of statins on altering the processing of APP in humans.⁵³

The Heart Protection Study Collaborative Group⁵⁴ and the Prospective Study of Pravastatin in the Elderly at Risk (PROSPER)⁵⁵ have both recently reported that neither simvastatin nor pravastatin appeared to slow cognitive decline in the elderly during 5 years of treatment in the Heart Protection Study and 3.2 years in the PROSPER. These findings suggest that lower rates of dementia found in the medical record reviews^{51,52} among persons taking statins may have been due to other factors that were actually responsible for the lower risks.⁵⁶

VACCINATION

Monthly immunization for 11 months with injections of A β 42 was found to prevent the development of A β plaque

formation, neuritic dystrophy, and astrogliosis in mice transfected with the V717F mutant human *APP* gene.⁵⁷ Partial regression and clearance of plaques were seen in animals that had already developed them. Substantial reduction of reactive gliosis also occurred.

Based on the impressive reduction in brain amyloid burden in the immunized transgenic mice, safety and efficacy trials of the injectable vaccine were initiated in patients with AD. It was shown that patients immunized with a primary injection of preaggregate A β 42 followed by 1 booster injection in a placebo-controlled study did generate antibodies that recognized A β plaques, diffuse A β deposits, and vascular A β in brain blood vessels. Thus, A β 42 vaccination in patients with AD induces antibodies that have a high degree of selectivity for the pathogenic target structures.⁵⁸

Some patients who generated such antibodies showed slower rates of decline of cognitive functions and activities of daily living.^{59,60} The A β 42 immunization was apparently having a positive therapeutic effect, but unfortunately, the clinical trial had to be terminated because 6% of immunized patients developed autoimmune meningoencephalitis.⁶¹ The brains of 4 people with mild to moderate AD who had been vaccinated and died from unrelated causes were examined. Each brain showed an almost complete lack of A β with maintenance of neurofibrillary tangles.^{62,63} The vaccine targeted normal myelin in addition to A β as some patients developed acute demyelinating lesions.^{61,62,64} The specificity of the A β peptide vaccine must be reviewed because the role of microglial nonspecific activation by any vaccine reducing the accumulation of A β remains an unresolved issue.

A program using passive immunization with preformed A β antibodies is currently being developed to prevent secondary encephalitis. Genetic vaccination using the complementary DNA for A β in transgenic mice generated antibodies against A β without activating cytotoxic T cells causal of autoimmune meningoencephalitis and thus might be a form of active immunization to develop for future clinical trials.⁶⁵ Immunization therapy for AD should be viewed optimistically because it has shown a positive response in lowering the A β burden in selected patients with AD. The significant adverse effects need to be reviewed carefully and new immune strategies designed, including active immunization with the A β gene, which shows some promise.^{64,65}

SECRETASE INHIBITORS

Of great interest are the reports¹ of the cloning, isolation, and characterization of the membrane-bound aspartyl protease that cleaves full-length APP at the β -secretase cleavage site and finds it to be the predominant β -cleavage activity in human brain. Overexpression of the β -secretase gene increased the amount of β -secretase cleavage products. Demonstration of the specific β site APP-cleaving enzyme provides a direct approach to develop pharmacologic agents that can specifically inhibit this gene's expression. γ -secretase inhibition by a specific inhibitor is also a potential target to inhibit A β synthesis. Inhibition of β - or γ -secretase would prevent A β synthesis and affect the rate of plaque production.¹

Efforts to achieve a clinically useful secretase inhibitor form an area of active research.

CLIOQUINOL

Metal chelation using clioquinol has been reported in a pilot study with 36 patients with AD to reduce the rate of cognitive loss in a double-blind, placebo-controlled, phase 2 clinical trial. Clioquinol's effect in this preliminary study is due to its ability to chelate zinc and copper associated with amyloid plaques. The mobilization and removal of brain amyloid is believed to be basis of its therapeutic effect.⁶⁶ Clioquinol increased serum zinc and copper levels, which is explained only in part by the chelation model.

INSULYSIN

It has been recently demonstrated that the peptidase insulysin has a quantitatively significant and rate-limiting role in degrading brain A β peptides in mice in vivo. Because insulysin also has a prominent role in insulin degradation, decreased or aberrant insulysin expression is not only a likely risk factor for AD but also a reasonable mechanism to explain a high incidence of abnormalities in glucose and insulin metabolism in AD. Insulysin activation is a possible therapeutic approach to reduce brain levels of A β .²⁹

CONCLUSIONS

It is clear from the molecular and genetic data that AD is a clinical syndrome with a common set of clinical features due to several genotypes causal of early-onset autosomal-dominant disease (*APP*, *PS1*, and *PS2* mutations) or, more significantly, in the vast majority of patients with late-onset disease, due to the complex interaction of polygenetic influences and environmental risk factors. The present view is that a subset of risk-producing polymorphic genes are expressed and result in the neuropathologic abnormalities and clinical dementia of AD. The future resolution of its molecular pathogenesis and subsequent pharmacologic therapies will depend on a genomic and proteomic analysis of patients, their nonaffected family members, and control subjects. Genomic and proteomic profiling as shown on DNA and protein microarrays, it is believed, will show a pattern of expression that correlates with a high-risk state for subsequent AD. Genomic and proteomic analyses for prediction and prevention of AD may supplant current clinical diagnosis and symptomatic treatment.

Pharmacogenomic therapy designed to prevent progression of preclinical to overt clinical disease with dementia based on the genomic or proteomic profile of the individual patient is the intent and hope. It is premature to predict exactly what therapeutic effect will result from a genomic/proteomic approach. Gene linkage and sequencing studies for finding at-risk genes for AD have found some promising leads, including insulysin (chromosome 10) and possibly α 2-macroglobulin (chromosome 12). A recent genomic analysis suggested that pa-

tients with AD show about 3 to 5 times higher genetic variation than a control population.⁶⁷ There are data indicating that the therapeutic response in patients with AD to cholinesterase inhibitors is genotype specific.⁶⁸ A growing view is that genomics provide the potential to offer insights into the molecular and genetic basis of pathogenesis of complex diseases like AD.⁶⁸⁻⁷⁰ This approach may eventually prove to be productive in finding a cohort of genes and expressed products, relatively small in number, whose polymorphic profile will identify high-risk states for AD. In this regard, a proteomic analysis of AD brains has shown quantitative differences in the expression of proteins in 6 areas of brain. The molecular identity of 37 proteins with significantly altered expression was determined.⁷⁰ Identification of altered expression of genes and proteins primarily causal of AD from those showing secondary and reactive change is the challenge. When this goal is achieved, it will be possible to consider pharmacogenomic therapy at an early point in the disease process with careful clinical diagnostic criteria for AD before irreversible neuropathologic changes result.⁷¹

A new approach is needed as current pharmacologic therapy directed at symptomatic relief has proved to be marginally effective. The genomic and proteomic basis of AD will be defined in the near future, and corresponding molecular therapeutic targets will be identified. Genomic neurology has arrived and its application to resolving AD is our best hope.³⁰

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