

Zinc and Alzheimer's Disease

Ashley I. Bush *et al.* present data from an *in vitro* study of an interaction of aqueous Zn ions with β -amyloid protein (A β) (1), the protein that accumulates in neuritic plaques in patients with Alzheimer's disease (AD). Zinc was shown to bind to the protein, inducing aggregates that precipitated from solution (tinctorial amyloid formation). One important question is the specificity of this interaction, and Bush *et al.* partly address this by assaying 14 other metal ions (3 μ M) on an equimolar basis.

Although Bush *et al.* state that none of these other metals induced significant aggregation, the data presented show that Cu and Hg effected an approximate 12% decline in filterability of the A β solution; the decline with Zn was about 25%, although other data in the report suggest that 3 μ M Zn may induce up to 40% reduction. However, almost all of the other experiments described in this report used Zn at 25 μ M, presumably to maximize effects [this was the uppermost value in the dose-response graph, figure 2B in (1)] and to be operating with concentrations of Zn normally found in plasma. What would the A β aggregate-inducing ability of Cu and Hg ions (and indeed the other metals) be if tested at 25 μ M, or, more pertinently, if tested at a normal range of concentrations plasma? A review of such concentrations reveals that 3 μ M in plasma is unlikely; for Mg and Ca it is about three or four orders of magnitude too low; for Fe and Cu, it is just below their minimums of about 10 μ M; for Pb, Al, Hg, Cd, Mn, and Co, it is, respectively, 3, 5, 10, 30, 65, and 100 times the recommended maximum in non-occupationally exposed groups. This information therefore most likely discounts these latter metals as possible effectors of A β aggregation, but clearly Mg and Ca (and perhaps Fe and Cu) ions should be tested at concentrations at which they normally exist in plasma.

Bush *et al.* state that elevated (80%) Zn concentrations in cerebrospinal fluid (CSF) have been found in AD cases. However, in the reference they cite, it is evident that these data apply not to confirmed AD cases but to patients with Alzheimer-type dementia (ATD) (2). Furthermore, although the average concentrations CSF-Zn in these ATD cases was 80% higher than in age-matched controls, only 21% of these cases (7/33) had the arbitrarily-set, elevated concentration of CSF-Zn. Of a group of six AD cases, verified by autopsy, one had elevated concentrations CSF-Zn as compared with one of eight controls. Hershey *et al.* (2) also provide data on Cu, Fe, and Si (an element not tested by Bush *et al.*). In the CSF of the

33 ATD patients, concentrations of Si were elevated in eight patients. The average concentration over all cases was 130% higher than it was in the control group, while analyses of the autopsy-verified AD cases and controls revealed CSF-Si elevated in 5/7 cases versus 0/8 controls. Also, average concentrations of ATD-CSF-Cu and Fe in patients were 120% and 125% above those in the control group, respectively.

These data may further encourage extended A β aggregation studies with Cu and Fe, and suggest that Si (normally present as a complex anion or silicate polymer) should be included. Normal serum concentrations of Si are around 5 μ M, but in renal dialysis, concentrations of Si in the plasma of patients can range from 25 to 100 μ M, and there are data showing that an elevated concentration of Si has a protective role in limiting Al bioavailability (3).

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Bush *et al.* (1) report that the aggregation of human AD amyloid peptide, A β ₁₋₄₀, in tris saline is significantly enhanced by Zn cations. Their qualitative conclusion (that zinc induces A β aggregation) confirms our earlier work (2) that in either tris buffers or human cerebrospinal fluid at 37°C, Zn was more effective than a variety of other metals tested. Two points must be addressed in studies of this kind: the accuracy and reproducibility of the measurements, and the extrapolation of results obtained at higher concentrations of peptide and metal to the much lower ones found physiologically. Neither of these points has been adequately addressed by Bush *et al.* (1). First, their measurements of peptide concentration and aggregation are incorrect because the method used is inappropriate. Second, their extrapolation from data gathered at 1 to 25 μ M peptide concentrations to the physiological range (<0.005 μ M, a thousandfold lower) is unsubstantiated. Significant aggregation of A β is induced by Zn²⁺, but at concentrations a hundredfold higher than those reported by Bush *et al.* (1).

Bush *et al.* (1) used ultraviolet spectroscopy at low wavelength, that is, optical

density at 214 nm (OD₂₁₄), to measure concentrations and aggregation of human A β (mostly at 1.6 μ M). On the basis of this method, they report that Zn²⁺ concentrations above 0.3 μ M cause aggregation of A β into an insoluble form that can be removed from solution by centrifugation or filtration. At a concentration of about 3 μ M Zn²⁺, half of the A β aggregated. In contrast to normal human A β , rat A β and radiolabeled human A β were aggregated at significantly higher (>30 μ M) concentration of Zn²⁺. These A β concentrations are at the limits of the sensitivity of the OD₂₁₄ method (1, 6).

OD₂₁₄ is rarely used for peptide quantitation (8, 9). Essentially all organic molecules absorb at 214 nm, producing a high background that is sensitive to artifact. At the concentrations used by Bush *et al.* (1), tris buffer has a much greater OD₂₁₄ than 1.6 μ M A β . Furthermore, the buffer absorbance is highly sensitive to wavelength, pH, temperature, dissolved oxygen, and Zn²⁺. The optical density of peptides at 214 nm is mostly provided by the peptide bond (8, 9) and as such is dependent on the conformation of the peptide (8-11) as well as on its concentration. For example, the OD₂₁₄ of polylysine at constant concentration changes by more than 50% as its conformation changes from α helix to β sheet (11). The addition of metals such as Zn to peptides and proteins affects their conformation and therefore their optical density. Bush *et al.* have concluded (6) that Zn affects the conformation of A β and then assume (1) that a change in OD₂₁₄ reflects only a change in A β concentration, with no effect of the conformational change.

Because the behavior of A β is highly dependent on its concentration, extrapolation to physiological (10⁻⁹ M) concentrations of data gathered at >10⁻⁶ M (1) is not valid (3, 4) without experimental support. We used unlabeled A β (tracked by immunoassay), or radiolabeled A β (tracked by counting), or mixtures of the two (tracked independently by each method) in various ratios to examine the effect of Zn on A β aggregation at A β concentrations varying from 10⁻¹⁰ to 10⁻⁶ M. Bush *et al.* (1) report that the labeled and unlabeled peptides behave differently, but we found that in all cases ¹²⁵I-A β tracked unlabeled A β accurately, validating the use of the radiolabeled tracer in these experiments.

With the use of filtration (1) or centrifugation (1, 2) to remove insoluble aggregates, we found that approximately 200 μ M Zn²⁺ is required to aggregate half of 1 μ M A β from solution at pH 7.4 and 37°C; this EC₅₀ is about a hundred times higher than that determined by Bush *et al.* (1) from optical density measurements. The EC₅₀ for Zn varies with A β concentration, with

somewhat higher Zn concentrations required to aggregate the same fraction of peptide at physiological concentrations. The deposition of A β at physiological concentrations onto plaques in AD tissue sections or homogenates is also enhanced by Zn, but only at Zn²⁺ concentrations above 50 μ M (unpublished results). Although the rat and human A β peptides are certainly different in their behavior in some systems (7), we disagree with Bush *et al.* (1) that the peptides differ significantly in the concentration of Zn required to clear them from aqueous solution. In contrast, similar concentrations of Zn were required to aggregate half of either human or rat A β at peptide concentrations between 10⁻¹⁰ and 10⁻⁶ M.

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Response: We agree with the interpretation of Maggio *et al.* that Zn appears to induce a major conformational effect on A β ₁₋₄₀ in vitro, as indirectly indicated by our findings that Zn(II), at physiologically relevant concentrations, provokes the rapid aggregation of human A β ₁₋₄₀ into particles that are retained by filters of pore sizes 0.1 to 0.65 μ m (1). In contrast, the rat and mouse form of the peptide is relatively unaffected by Zn, perhaps explaining why these species are spared cerebral amyloid pathology with age. To quantify the retention of A β ₁₋₄₀, we assayed the amounts of peptide entering the filtrate by measuring optical density at 214 nm (OD₂₁₄). We validated this technique by preparing standard curves of OD₂₁₄ ver-

sus peptide concentration for each of the experimental buffers used (2). These curves were linear over the range of concentrations used in these experiments (even after incubation with Zn), which indicated that the changes in absorbance observed in our studies most likely reflect changes in peptide concentration (3).

Our aggregation assay compares the OD₂₁₄ of an incubated peptide solution to the OD₂₁₄ of the same solution that has been filtered after the same incubation. Maggio *et al.* maintain that the changes in OD₂₁₄ that we have reported in comparing a filtered to an unfiltered solution may be reflecting changes not only in concentration, but also in the peptide consequent to the act of filtration. Their comment does not take into account our other findings that support the interpretation that Zn(II) at ≤ 25 μ M induces the rapid and abundant aggregation of A β . These findings (1) include (i) retention by filtration of zinc:A β complexes (with 1:1 stoichiometry) after the incubation of the peptide with 25 μ M Zn(II), confirmed by the lack of retention of the tracer ⁶⁵Zn in the absence of peptide; and (ii) the ability of Zn(II) at 25 μ M (and not EDTA) to rapidly induce the production of tinctorial amyloid from soluble A β ₁₋₄₀. The observation of visible amyloid after treatment of A β ₁₋₄₀ with Zn is independent of assays that depend on measurements of OD₂₁₄.

In this rebuttal, we present results of other experiments corroborating those obtained (1) with the use of OD₂₁₄ measurements. To confirm that the reduction in OD₂₁₄ after the filtration of Zn-incubated A β solutions is a result of retention of the aggregated peptide by the filtration membrane, we assayed both the filtrate and the retentate for A β by immunoassay. To mea-

Table 1. Confirmation by immunoassay of zinc-induced precipitation of A β . A β ₁₋₄₀ (1 μ M, 400 μ l) was brought to the metal concentrations indicated in tris-buffered saline (100 mM NaCl, 20 mM tris-HCl, pH 7.4), incubated, and filtered through 0.2 μ m pore-size filters as previously described (7). The filtrate was diluted 1:100 in TBS and then an ELISA was performed for A β immunoreactivity against a standard curve (4). The data indicate the amount of peptide assayed in the filtrate as a fraction of peptide in the starting material ($n = 3$). Zn(II) chloride induced loss of A β ₁₋₄₀ whereas Fe(III) citrate had no effect, confirming the results of the OD₂₁₄ measurements.

Filtrate (μ M)	Filtered fraction	
	Mean	SD
No metal	0.90	0.01
Zn(II) 8	0.39	0.08
Fe(III) 8	0.91	0.02
Zn(II) 25	0.20	0.13
Fe(III) 25	0.91	0.03

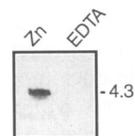
sure the amount of A β that enters the filtrate after the filtration of a peptide solution that has been incubated with Zn, we developed an ELISA that uses immobilized Zn or Cu to capture A β peptide from solutions or suspensions (4). The results (Table 1) confirm our earlier observations that >60% of a 1 μ M solution of A β ₁₋₄₀ incubated with Zn(II) (≤ 8 μ M) is lost after filtration. When we applied heated SDS sample buffer to the filter following the peptide filtration, and then assayed the eluted peptide by protein immunoblot (5), we confirmed (Fig. 1) that the peptide was retained by the filter in the presence of Zn, but not in the presence of EDTA.

Maggio *et al.* state that they do not detect precipitation of A β at peptide concentrations of ≤ 1 μ M unless they incubate the peptide with Zn concentrations approximately an order of magnitude (>200 μ M) greater than those that we used. Thus, the techniques used by these investigators appear to be unable to demonstrate precipitation of a 1 μ M A β solution by 25 μ M Zn(II), a strongly positive result that we have now confirmed by three different assay techniques (OD₂₁₄ measurement, ELISA, and western blot). Since Maggio *et al.* are unable to reproduce a result that is, for us, a positive control, it is unlikely that they will be able to detect precipitation of A β at lower peptide concentrations.

We are presently examining the effects of physiological concentrations of zinc upon A β at peptide concentrations below the limit of detectability of 214 nm spectrophotometry. Because iodinating the peptide at a critical tyrosine residue (position 10) (6) abolishes the zinc-induced precipitation of A β (1), we do not believe that ¹²⁵I-A β is a suitable tracer for such studies (7). Therefore, we are exploring alternative means of labeling the peptide without damaging its physicochemical properties.

Fitzgerald questions the specificity of metal-induced A β ₁₋₄₀ aggregation, point-

Fig. 1. Confirmation of zinc-induced retention of A β ₁₋₄₀. A β ₁₋₄₀ was brought to 1 μ M (800 μ l) in TBS \pm 50 μ M EDTA or 25 μ M ZnCl₂, incubated, and filtered through a 0.2 μ m pore-size filters as previously described (7). The membrane was washed twice with the incubation buffer (500 μ l). Sample buffer (8% SDS, 68 mM tris-HCl, pH 6.8, heated to 95°C, 100 μ l) was applied to the membrane and the filter centrifuged (10,000g \times 5 min). Filtered sample buffer was collected in a clean vessel, 10 μ l was loaded onto a tricine gel and a protein immunoblot was performed with 6E10 (gift of K. S. Kim and H. Wisniewski). Migration of the band relative to the molecular size markers (low M.W., Amersham, Arlington Heights, Illinois) is indicated. Figure illustrates $n = 7$ experiments.



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