ALZHEIMER'S DISEASE AND RELATED DISORDERS - ORIGINAL ARTICLE

Effect of copper intake on CSF parameters in patients with mild Alzheimer's disease: a pilot phase 2 clinical trial

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Abstract A plethora of reports suggest that copper (Cu) homeostasis is disturbed in Alzheimer's disease (AD). In the present report we evaluated the efficacy of oral Cu supplementation on CSF biomarkers for AD. In a prospective, randomized, double-blind, placebo-controlled phase 2 clinical trial (12 months long) patients with mild AD received either Cu-(II)-orotate-dihydrate (verum group; 8 mg Cu daily) or placebo (placebo group). The primary outcome measures in CSF were A β 42, Tau and Phospho-Tau. The clinical trial demonstrates that long-term oral intake of 8 mg Cu can be excluded as a risk factor for AD based on CSF biomarker analysis. Cu intake had no effect on the progression of Tau and Phospho-Tau levels in CSF. While A β 42 levels declined by 30% in the placebo group (P = 0.001), they decreased

only by 10% (P = 0.04) in the verum group. Since decreased CSF A β 42 is a diagnostic marker for AD, this observation may indicate that Cu treatment had a positive effect on a relevant AD biomarker. Using mini-mental state examination (MMSE) and Alzheimer disease assessment scale-cognitive subscale (ADAS-cog) we have previously demonstrated that there are no Cu treatment effects on cognitive performance, however. Finally, CSF A β 42 levels declined significantly in both groups within 12 months supporting the notion that CSF A β 42 may be valid not only for diagnostic but also for prognostic purposes in AD.

Keywords Cu · Alzheimer · Clinical trial · Tau · CSF, plasma · Abeta

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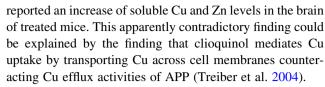
Introduction

One major pathological hallmark of Alzheimer's disease (AD) is the deposition of β -amyloid (A β) plaques in the brain. A β peptides result from enzymatic processing of the β -amyloid precursor protein (APP) by β - and γ -secretases. Since cerebrospinal fluid (CSF) is in direct contact with the central nervous system (CNS), measurement of the CSF concentration of $A\beta$ peptides has shown to represent one biomarker, which—especially in combination with increased Tau protein concentration in CSF-can improve the safety of clinical AD diagnosis. The normal level of $A\beta$ 1-42 is regularly found decreased to about 50% in patients with AD (Lewczuk et al. 2004; Wiltfang et al. 2001, 2002). Among other possible factors which require a confirmation the most accepted explanation is the accumulation of A β in plaques leading to decreased A β 42 levels in the CSF of AD patients.

Potentially toxic $A\beta$ peptides are generated from the copper-binding APP by two independent proteolytic events (Bayer et al. 2001; Glenner and Wong 1984; Hesse et al. 1994; Kang et al. 1987). APP is actively involved in balancing Cu concentrations in cells. In APP-knock-out mice, Cu levels were found increased in cerebral cortex and liver (White et al. 1999), whereas overexpression of APP was reported to result in significantly reduced Cu levels in brain tissue of different APP transgenic mouse strains (Bayer et al. 2003; Phinney et al. 2003) and in mice overexpressing the C-terminal fragment of APP (and enhanced $A\beta$ secretion) (Maynard et al. 2002).

The N-terminal Cu binding domain (CuBD-I) of APP shows structural homology to the Cu binding domain of Cu chaperons (Barnham et al. 2003) binding Cu with nanomolar affinity (Hesse et al. 1994). A secondary CuBD-II appears in $A\beta$ after its release from APP (Atwood et al. 2000), and Cu application was reported to increase $A\beta$ aggregation in vitro [reviewed in (Bush 2003)]. APP reduces Cu(II) to Cu(I), leading to oxidative modification of APP (Multhaup et al. 1996), which is facilitated through the protein surface localization of the binding site thus resembling so called cytoplasmic Cu chaperones (Barnham et al. 2003).

In cell culture systems, Cu supplementation was found to stimulate the non-amyloidogenic APP pathway thereby suppressing the formation of β amyloid (Borchardt et al. 1999). More recently, APP was shown in yeast cells to have a Cu efflux activity thereby explaining why APP over-expressing mice have a reduced Cu level in their brains (Bayer et al. 2003; Phinney et al. 2003; Treiber et al. 2004). Earlier studies in animals have reported that elevated Cu is a risk factor for developing the AD related pathology. Cherny et al. (2001) showed that clioquinol, a Cu and Zn chelating agent, can remove β -amyloid plaque pathology. However, it was unclear how this effect worked, since the authors



The observation that dietary Cu supplementation in a transgenic mouse model for AD does not only increase bioavailable brain Cu levels and restores superoxide dismutase-1 (SOD-1) activity but also lowers $A\beta$ levels in the brain and prevents premature death, supports the notion that a disturbed Cu homeostasis may be associated with the pathological process in AD (Bayer et al. 2003). Furthermore, increasing Cu levels by genetic means reduced $A\beta$ plaque load and rescued the premature death of APP transgenic mice (Phinney et al. 2003).

Normally, Cu contained in the food is taken up in the stomach and then absorbed in the small intestine. About 30-50% of the Cu is absorbed. Cu is distributed from the liver throughout the body and transported in the bloodstream bound to ceruloplasmin. The liver is the most important organ for Cu distribution and storage. Cu is excreted via the biliary system. Usually, 2 mg of Cu per day are taken with food. Ingestion of as much as 10 mg of Cu per day is considered to be safe. The clinical reference value for physiological Cu plasma concentrations is 65-165 μg/dl [reviewed in (Kessler et al. 2005)]. In an earlier report we could show a negative correlation between plasma Cu and ADAS-cog score in AD patients, i.e. cognitive performance in patients with Cu levels within the lower third of the physiological range was worse than in patients with higher Cu plasma levels (Pajonk et al. 2005). In addition, AD patients fulfilling the criteria of CSF diagnosis for AD (defined as at least two of three markers [A β 42, Tau, Phospho-Tau] being out of the reference range) had significantly lower blood Cu levels than AD patients who fulfilled none or only one of these CSF criteria (Kessler et al. 2006).

Taken together, this brought us to hypothesize that restoring brain Cu homeostasis might have a beneficial influence on the progression of AD. We speculated that oral Cu supplementation may stabilize CSF biomarker parameters in AD patients. Thus, we monitored the progression of the disease in AD patients by recording the changes in CSF and plasma within 12 months in a prospective, randomized, double-blind pilot phase 2 clinical trial.

Material and methods

Study population

Criteria for participation in the study included written informed consent as well as caregiver consent, a diagnosis



of probable AD by means of NINCDS-ADRDA criteria (McKhann et al. 1984), a MMSE score of 20–25. All patients received 5–10 mg donepezil daily. Patients with severe and unstable somatic diseases and patients with present or known history of alcohol, drug or medication abuse were excluded. Patients taking drugs for coexistent diseases were included except those taking psychotropic drugs, "nootropics" or health food supplements. The trial has been approved by the appropriate ethics committee and has therefore been performed in accordance with the ethical standards laid down in the 1964 Declaration of Helsinki.

Study design

The study had a monocentric, prospective, double-blind, placebo-controlled, parallel-group randomized design. In total 68 patients were recruited. Screening procedure consisted of a medical history, physical examination, psychometric tests, blood tests, MRT scan and diagnostic lumbar puncture. After 1 year a second lumbar puncture was obtained to monitor possible CSF changes. Patients were randomly allocated either to placebo or to verum (Cu orotate). The oral dosage of Cu orotate was 51.62 mg (corresponding to 8 mg Cu) once daily. To maintain blinding, capsules containing Cu orotate and placebo had identical shape and colour. The duration of the study was 12 months. The study was registered at http://www.clinicaltrials.gov with the identification number NCT00608946.

Outcome measures

The primary efficacy variables were the changes from the baseline score in the CSF A β 42, Tau and Phospho-Tau levels. Efficacy evaluations were performed at baseline and after 12 months.

CSF sampling

CSF was drawn from patients by lumbar puncture in the afternoon, sampled in polypropylene vials and centrifuged (1,000g, 10 min, 4°C). Aliquots of 250 μ l were stored at -80°C. Freezing of samples was conducted by directly cooling 250 μ l of CSF in polypropylene tubes down to -80°C without an intermediate temperature stage. The samples did not undergo additional freeze and thaw cycles.

 $A\beta 42$, Tau and Phospho-Tau quantification by ELISA

CSF amyloid beta 1–42 (A β 42), CSF total Tau, and CSF Tau phosphorylated at threonine-181 (Phospho-Tau) were quantified using ELISA as recommended by the

manufacturer (Innogenetics, Gent, Belgium; carried out at the Medizinisches Labor Bremen, Germany). CSF samples for ELISA were analysed at different time points.

Urea gel electrophoresis for A β 1–37, 1–38, 1–39, 1–40 and 1–42 quantification by Western blotting

CSF proteins were separated on 10% Bicine/Tris gels containing 8 M urea (22), and $A\beta$ peptide species were revealed by western blotting using antibody 1E8. Synthetic $A\beta$ peptides of different size were run in parallel in the same gel system and under the same conditions for the identification and quantification of $A\beta$ peptides by densitometry. Immunoreactive band intensities were quantified with the Quantity One v4.1 software (Bio-Rad). All samples were run as duplicates and each gel carried a five-step dilution series of a synthetic $A\beta$ peptide mix. Bands were quantified relative to this dilution series. The interand intra-assay coefficients were below 10%. Mean values were used for subsequent calculations. The CSF samples were analysed altogether at one time point.

Cu, Zn and ceruloplasmin in blood

Blood was collected to determine the level of Cu and, Zn by using atomic absorption spectroscopy (AAS) and to analyze ceruloplasmin by the nephelometric method (see below). Blood samples were collected into metal-free tubes that contained lithium heparin as an anti-coagulant. Concentrations of Cu were measured in lithium-heparin plasma samples, utilizing flame AAS (PerkinElmer, AAnalyst 800). Samples were diluted with deionised water, and the analysis was performed using standards prepared in glycerol to approximate the viscosity characteristics of the diluted samples. Standard atomic absorption conditions were utilized for Cu (air acetylene, wave length 324.8 nm). The between-day coefficient of variation (CV%) for Cu assay was 4.24% (at 88 µg/dl), and the between-day coefficient of variation (CV%) for Zn assay was 7.21% (at 317 µg/dl). Cu, Zn and ceruloplasmin samples were analysed immediately after collection.

Ceruloplasmin has been determined by immunochemical reaction utilizing the nephelometric method (Behring Nephelometer BN II). Reagents and quality control sera were provided by Dade Behring Company (Germany). The results were evaluated by means of logit-log function. Samples containing particles were centrifuged prior to testing. Plasma specimens were automatically diluted 1:20 with N Diluent. The analytical imprecision has been determined as follows: intra-assay CV 1.4%, interassay CV 1.8%. The expected reference range utilising this method was between 0.2 and 0.6 g/l (P_{2.5}–P_{97.5} percentile).



Cu levels in CSF

Inductively coupled plasma mass spectrometry (ICP-MS): for analysis of metal concentrations (Cu and Zn) in CSF, samples were prepared by HNO₃ closed-vessel microwave digestion and diluted in Milli-Q water to a final concentration of 6.5% HNO₃ for analysis by ICP-MS. ICP-MS was performed by using a HP4500 Series 300 ShieldTorch system instrument (Agilent, Waldbronn, Germany) in peak-hopping mode with spacing at 0.05 atomic mass units, three points per peak, three scans per replicate, and an integration time of 300 ms per point. The rate of plasma flow was 15 l/min with an auxiliary flow of 1.0 l/min. The replicative form power was 1.2 kW. The sample was introduced by using a crossflow nebulizer at a flow rate of 1.02 l/min. The CSF samples were analysed at one time point.

Safety measures

Standard adverse event reporting was conducted. Before baseline, the Cu content in drinking water at the patients' home was analyzed under standard conditions; concentration of Cu in drinking water must not exceed 2 mg/l. For safety reasons, blood analyses (including especially plasma Cu levels and liver enzymes) were performed at week 1, 2, 3 and 4 after baseline and at month 3, 6, 9 and 12.

Data analysis

For statistical analyses SPSS 15 for Windows was used. Significance level was $\alpha = 0.05$. All tests were two-tailed. Demographic variables and data on the course of the disease at screening time (t_1) were compared between the verum and the placebo group with one-way analysis of variance

Table 1 Demographic and clinical data of patients at screening time (t1) that have completed the clinical trial

		Placebo		Verum	χ^2	df	P
Gender (no. male; no. female)		m.:12; f.: 21		m.: 17; f: 18	1.04	1	0.31
APOE: number of E4 alleles (0, 1, 2)		0: 9; 1: 13; 2: 2	0: 7; 1: 16; 2: 2		0.54	2	0.76
	n	Mean ± SEM	n	Mean ± SEM	F	df	P
Age (years)	33	69.48 ± 1.39	35	70.37 ± 1.12	0.25	1, 66	0.62
Disease Duration (months)	33	24.73 ± 2.69	34	33.26 ± 4.97	2.25	1, 65	0.14
Age at onset (years)	33	67.48 ± 1.34	34	67.50 ± 1.09	0.00	1, 65	0.99
Education (years)	32	10.31 ± 0.35	34	11.41 ± 0.47	3.53	1, 64	0.065
CDT (score)	33	2.82 ± 0.23	34	2.88 ± 0.21	0.04	1, 65	0.84
MMSE (total score screening)	33	23.52 ± 0.52	35	24.17 ± 0.61	0.66	1, 66	0.42

The placebo and verum group did not differ significantly in any of the variables

m, Male; f, female; n, number of cases; m, mean; SEM, standard error of the mean; χ^2 , χ^2 statistics; df, degrees of freedom; F, F statistics; CDT, clock drawing test; MMSE, mini mental state examination

Table 2 Demographic and clinical data of patients with CSF collections at screening time (t_1) at the beginning and at the end of the study

		Placebo	Ve	erum	χ^2	df	P	
Gender (no. male; no. female)		m.:4; f.: 13	m.	m.: 11; f: 7		1	0.025	
APOE: number of E4 alleles (0, 1, 2)		0: 6; 1: 8; 2: 1		2; 1: 12; 2: 2	3.10	2	0.21	
	n	Mean ± SEM	n	Mean ± SEM	F	df	Р	
Age (years)	17	67.76 ± 1.72	18	68.94 ± 1.67	0.24	1, 33	0.63	
Disease Duration (months)	17	23.12 ± 3.79	18	31.50 ± 8.52	0.78	1, 33	0.38	
Age at onset (years)	17	65.94 ± 1.72	18	66.44 ± 1.50	0.05	1, 33	0.83	
Education (years)	17	10.59 ± 0.54	18	12.06 ± 0.67	2.86	1, 33	0.10	
CDT (score)	17	2.76 ± 0.36	18	2.94 ± 0.27	0.16	1, 33	0.69	
MMSE (total score screening)	17	23.71 ± 0.83	18	23.67 ± 0.95	0.00	1, 33	0.98	

m, Male; f, female; n, number of cases; m, mean; SEM, standard error of the mean; χ^2 , χ^2 statistics; df, degrees of freedom; F, F statistics; CDT, clock drawing test; MMSE, mini mental state examination



(ANOVA). χ^2 test on independence was used to analyze, if the gender distribution was different between the two groups and if the number of APOE E4 alleles differed between the groups (Tables 1, 2). Primary dependent variables were ELISA A β levels of A β 42, Tau and Phospho-Tau. They were expressed as absolute numbers at the starting point (t_1) and 12 months later at the end of the study (t_{12}). Secondary dependent variables were: Western blot A β levels, expressed as % of total A β : 1–42, 1–40, 1–39, 1–38, 1–37%, or as A β ratios 1–42/1–40, 1–42/1–38, 1–38/1–40,

such as cholesterol, plasma Zn, ceruloplasmin, plasma Cu and CSF Cu values, expressed as absolute numbers at t_1 and t_{12} . Independent between-subject factor was group (placebo, verum), within-subject factor was time of measurement. Kolmogorov–Smirnov tests were used to test, if there were significant deviations of the dependent variables from the normality assumption. At t_1 (Table 3) and t_{12} (Table 4) one-way ANOVA with fixed factor group was performed for the dependent variables and found to be normal distributed. For the variables, where the normality

Table 3 Descriptive Statistics at screening time (t1), ANOVA/Mann-Whitney U test

	Placebo		Verum		% Diff. verum versus placebo	Factor group		
	n	Mean ± SEM	n	Mean ± SEM		df	F	P
WB: Aβ 1–42%	11	4.6 ± 0.49	11	4.0 ± 0.39	-13.8	1, 20	1.0	0.3
WB: Aβ 1–40%	11	66.0 ± 1.37	11	67.5 ± 0.80	2.1	1, 20	0.8	0.4
WB: Aβ 1–39%	11	9.4 ± 0.60	11	9.4 ± 0.26	0.1	1, 20	0.0	1.0
WB: Aβ 1–38%	11	13.1 ± 0.65	11	12.5 ± 0.44	-4.4	1, 20	0.6	0.5
WB: Aβ 1–37%	11	6.8 ± 0.60	11	6.6 ± 0.32	-2.8	1, 20	0.1	0.8
WB: Aβ 42/40	11	0.069 ± 0.01	11	0.059 ± 0.01	-15.0	1, 20	1.3	0.3
WB: Aβ 42/38	11	0.375 ± 0.05	11	0.328 ± 0.03	-12.4	1, 20	0.5	0.5
WB: Aβ 38/40	11	0.200 ± 0.01	11	0.186 ± 0.01	-7.1	1, 20	0.9	0.4
Cholesterol plasma (mg/dl)	20	211.0 ± 8.98	17	211.1 ± 9.52	0.1	1, 35	0.0	1.0
Zn plasma (µg/ml)	24	67.3 ± 1.61	26	65.9 ± 1.47	-2.0	1, 48	0.4	0.5
Ceruloplasmin plasma (mg/dl)	25	27.8 ± 1.07	29	27.0 ± 0.98	-2.8	1, 52	0.3	0.6
Cu plasma (µg/dl)	25	109.0 ± 4.68	27	98.5 ± 3.41	-9.6	1, 50	3.3	0.1

ANOVA and Mann–Whitney U test were used where appropriate and show no significant difference between the verum and placebo group n, Sample size; m, mean; SEM, standard error of the mean; % Diff., difference verum versus placebo group in per cent terms; df, degrees of freedom; F, F statistics; P, error probability for falsely rejecting the null hypothesis, that there are no mean differences between the two treatment groups; WB, Western blot: % of total $A\beta$

Table 4 Descriptive statistics at the end (t_{12}) , ANOVA/Mann–Whitney U test

	Placebo		Verum		% Diff. verum versus placebo	Factor group		
	n	Mean ± SEM	n	Mean ± SEM		df	F	P
WB: Aβ 1–42%	11	4.38 ± 0.46	11	3.8 ± 0.36	-14.3	1, 20	1.2	0.3
WB: Aβ 1–40%	11	65.9 ± 1.25	11	68.1 ± 1.02	3.2	1, 20	1.7	0.2
WB: Aβ 1–39%	11	9.5 ± 0.56	11	9.1 ± 0.36	-4.3	1, 20	0.4	0.5
WB: Aβ 1–38%	11	12.9 ± 0.58	11	12.5 ± 0.37	-3.6	1, 20	0.5	0.5
WB: Aβ 1–37%	11	7.2 ± 0.65	11	6.6 ± 0.37	-8.5	1, 20	0.7	0.4
WB: Aβ 42/40	11	0.066 ± 0.01	11	0.056 ± 0.01	-15.7	1, 20	1.4	0.3
WB: Aβ 42/38	11	0.359 ± 0.05	11	0.301 ± 0.03	-16.1	1, 20	0.9	0.4
WB: Aβ 38/40	11	0.198 ± 0.04	11	0.184 ± 0.03	-7.0	1, 20	1.0	0.3
Cholesterol plasma (mg/dl)	20	207.8 ± 7.46	17	211.3 ± 8.38	1.7	1, 35	0.1	0.8
Zn plasma (µg/ml)	24	67.2 ± 2.22	26	70.1 ± 2.19	4.4	1, 48	0.9	0.5
Ceruloplasmin plasma (mg/dl)	25	29.2 ± 0.97	29	27.8 ± 0.89	-4.6	1, 52	1.0	0.3
Cu plasma (µg/dl)	25	100.8 ± 4.32	27	100.7 ± 2.94	-0.1	1, 65	0.0	1.0

ANOVA and Mann–Whitney U test were used where appropriate and show no significant difference between the verum and placebo group n, Sample size; m, mean; SEM, standard error of the mean; % Diff., difference verum versus placebo group in per cent terms; df, degrees of freedom; F, F statistics; P, error probability for falsely rejecting the null hypothesis, that there are no mean differences between the two treatment groups; WB, Western blot: % of total $A\beta$



assumption was rejected, non-parametric Mann-Whitney U test with fixed factor group was computed. These analyses were based on the patients that completed the examinations. For Western blot and ELISA variables and CSF Cu it was tested if the change over time $(t_{12}-t_1)$ was gender related [ANOVA (gender \times group) or Mann–Whitney U test: males versus females separately for placebo and verum patients, respectively]. As main analysis the general linear model procedure (GLM) was used to perform a multivariate analysis of variance (MANOVA) with a repeated measure design using the within-subject factor time of measurement $(t_{12} \text{ vs. } t_1)$ and the between-subject factor group if the dependent variables were tested to be normal distributed. If the normality assumption was rejected, the non-parametric Wilcoxon test with the within-subject factor time $(t_{12} \text{ vs. } t_1)$ was done separately for the placebo and the verum group. The change over time $(t_{12}-t_1)$ of CSF copper values were correlated with cholesterol and A β 42 differences (t_{12} – t_1) using Spearman correlation coefficients.

Power calculation

The power calculation was based on the success criterion of the medication having an effect on the primary efficacy variables for CSF ELISA A β 42 levels applying a 5% significance level for each analysis. Assuming a large effect size of $[(m_{\text{verum}} - m_{\text{placebo}})/\sigma] = 1$ (notations: m_{verum} mean change over time $(t_{12}-t_1)$ in the verum group, m_{placebo} mean change over time $(t_{12}-t_1)$ in the placebo group, σ pooled standard deviation) and a sample size of 35 patients (CSF A β 42: 18 in the verum group, 17 in the placebo group), the study had a power of 83% to detect significant differences between the groups.

Results

Subject recruitment and demographics

At t_1 , between the placebo and the verum group there were no significant mean differences for age, disease duration, age at onset of the disease, clock drawing test and mini mental state examination at screening time. There was a trend for a longer duration of education in the verum group (F = 3.50; df = 1, 65; P = 0.066). The gender distribution was not significantly different between the verum and the placebo group. The number of APOE E4 allele carriers was not significantly different between the two groups (Table 1). Table 2 shows the demographic data of those patients with CSF samples at the beginning and the end of the study. There were also no significant mean differences for age, disease duration, age at onset of the disease, clock drawing test and mini mental state examination at

screening time. However, the gender distribution was significantly different with the number of male patients was significantly higher in the verum group.

Proof of concept

Kolmogorov–Smirnov test showed that in the placebo group and in the verum group as well, there were significant deviations from the normal distribution for ELISA A β 42, Tau and CSF Cu values. Consequently, for these variables, non-parametric tests were used. For all other dependent variables, the normality assumption could be maintained, and thus, for further analysis, parametric tests were performed. The change over time of all Western blot and ELISA variables and of CSF Cu was not significantly different between male versus female patients.

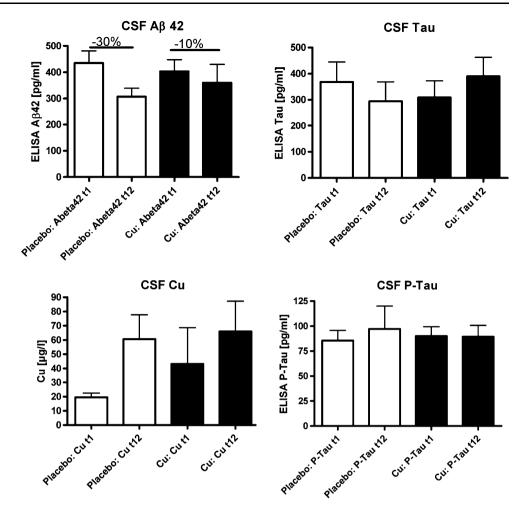
For most data there were no significant mean differences between the placebo and the verum group at t_1 (Table 3) and at t_{12} (Table 4). There were significant effects of within-subject factor time for A β concentration detected by Western blot 1–42% (t_{12} vs. t_1 –5.9%; F = 5.7; df = 1, 20; P = 0.03), A β ratio 1–42/1–40% (t_{12} vs. t_1 –6.0%; F = 5.1; df = 1, 20; P = 0.04), and ceruloplasmin % (t_{12} vs. $t_1 + 3.9\%$; F = 6.5; df = 1, 52; P = 0.01), however, these changes over time were not group related, i.e., there was no treatment effect using MANOVA with repeated measures design. Furthermore, there was a significant interaction of time and group for Cu in plasma (t_{12} vs. t_1 : placebo -7.5%; verum +2.3%; F = 5.2; df = 1, 50; P = 0.03). For Cu in CSF there was a significant increase in the placebo group (t_{12} vs. t_1 : +211.6%; Z = -2.6; P = 0.01), whereas there was only a trend for an increase over time in the verum group (t_{12} vs. t_1 : +52.9%; Z = -1.9; P = 0.06).

There were no significant correlations for the change over time $(t_{12}-t_1)$ between cholesterol and Cu values in plasma (placebo: $\rho=0.02$; df=17; P=0.94; verum: $\rho=0.21$; df=15; P=0.42) as well as in CSF (placebo: $\rho=0.33$; df=12; P=0.25, verum: $\rho=-0.15$; df=10; P=0.64).

In contrast to the other values CSF-A β 42, CSF-Tau and CSF-Cu levels did not show a normal distribution, therefore non-parametric Wilcoxon tests were used to compare time point t_1 at the beginning and t_{12} at the end of study. CSF A β 42 was significantly decreased in both Cu-treated (P=0.04) and placebo (P=0.001) groups after 12 months. Of interest A β 42 levels deceased by 30% in the placebo group, whereas only 10% in the Cu treated group, which may be regarded as a mild treatment effect. No significant effects were seen in Tau, P-Tau and Cu levels. CSF Cu levels in the placebo group increased significantly (P=0.01) due to lower levels at the beginning, however, there was no significant group effect. P-Tau levels were



Fig. 1 CSF parameters before and at the end of the clinical study. A\beta42, Tau and Cu CSF levels did not elicit normal distribution, therefore nonparametric Wilcoxon tests were used to compare time point t_1 at the beginning and t_{12} at the end of study. a CSF A β 42 was significantly decreased in both Cu-treated (P = 0.04) and placebo (P = 0.001) groups after 12 months. Of interest A β 42 levels deceased by 30% in the placebo group, whereas only by 10% in the Cu treated group. **b** No significant effect was seen in Tau levels. c CSF Cu levels in the placebo group increased significantly (P = 0.01) due to lower levels at the beginning, however, there was no significant Cu-treatment effect. **d** P-Tau levels showed no significant differences. All values represent mean and SEM. Number of samples for $A\beta42$, Tau and P-Tau was 17 in placebo and 18 in the verum group. Number of samples for Cu analysis in CSF was 13 in the placebo and 14 in the verum group



normal distributed therefore MANOVA analysis with repeated measures design was used. Again, P-Tau levels showed no significant differences (Fig. 1). There were no significant correlations for the change over time $(t_{12}-t_1)$ between Cu values in CSF and cholesterol (placebo: $\rho = 0.33$; df = 12; P = 0.25, verum: $\rho = -0.15$; df = 10; P = 0.64) as well as between CSF-Cu and CSF-A β 42 (placebo: $\rho = 0.32$; df = 10; P = 0.31, verum: $\rho = -0.39$; df = 12; P = 0.16).

Discussion

APP and $A\beta$ are both metalloproteins which bind Cu in vitro and were suggested to be involved in brain Cu homeostasis. Maynard et al. (2002) have shown that overexpression of the $A\beta$ sequence contained within a carboxy-terminal fragment of APP elicits significantly reduced Cu levels in transgenic mouse brain. This effect might be further enhanced by the drastically reduced bioavailability of Cu possibly due to the binding to plaque $A\beta$ (Lovell et al. 1998).

The APP mediated Cu efflux activity observed in APP overexpressing living cells best explained the intracellular Cu deficiency and a subsequently reduced SOD-1 activity (Bayer et al. 2003; Treiber et al. 2004). In addition, a genetically 1.5-fold up-regulated Cu level was associated with an increased survival of APP transgenic mice and lowered endogenous murine $A\beta$ levels prior to detectable $A\beta$ plaques formed by the human APP transgene (Phinney et al. 2003). These observations led us conclude that restoring brain Cu homeostasis might have a beneficial influence on the progression of AD biomarkers. Thus, in the present clinical trial we investigated potential beneficial effects of oral intake of Cu-(II)-orotate-dihydrate (8 mg Cu daily) in AD patients and CSF was collected at beginning and at the end of the study after 12 months.

The treatment was generally well tolerated although there was no clear-cut beneficial effect on CSF biomarker levels in AD patients. One may speculate that Cu treatment normalized Cu levels in plasma in the verum group by enhanced uptake and transport and improved tissue homeostasis. The plasma Cu levels declined only in the placebo group during the 12-months period, however the



placebo group had higher Cu levels at the beginning of the study by accident. Cu levels in the verum group were unchanged, which seems to be paradoxical. Previously, we have reported that significantly lower levels of Cu in plasma were found in those AD patients, who fulfilled the criteria of CSF diagnosis for AD (Kessler et al. 2006). In addition we demonstrated reduced Cu levels in plasma in patients with higher ADAScog scores (making more mistakes in this neuropsychological test) (Pajonk et al. 2005).

Reduced CSF levels of the $A\beta42$ in AD patients have been found in numerous studies, with high sensitivity and are commonly used for diagnostic evaluation [reviewed in (Andreasen et al. 2003)]. We also observed that $A\beta42$ levels were reduced using Western blot and ELISA techniques after a 12 month observation period corroborating these earlier studies. However we did not find a change in Tau or Phospho-Tau levels over time. On the other side FDG-PET, PIB-PET and fMRI may be more suitable as in vivo pathological markers.

Recently, it has been demonstrated that CSF Cu levels negatively correlate with A β 42 levels in AD patients (elevated Cu and decreased A β 42) (Strozyk et al. 2007). In good agreement with this report, we found that Cu treatment apparently had a small effect on A β 42 levels. A β 42 declined in both the verum and placebo groups over time however much more dramatically with 30% in the placebo group (P=0.001). The A β 42 levels declined only by 10% in the verum group (P=0.04). In contrast to this observation Cu treatment had no beneficial effect on cognitive abilities tested by MMSE and ADAS-cog in AD patients of the present clinical phase II pilot study as already shown (Kessler et al. 2008). No effect on cognitive performance and a stabilizing effect on CSF A β 42 levels is to be contradictory and therefore needs further and independent evaluations.

One animal study showed that treatment of 21-monthsold Tg2576 mice with clioquinol, a Cu, Zn chelator, inhibited plaque formation and concomitantly increased soluble brain Cu and Zn levels (Cherny et al. 2001). Moreover, lowered insoluble A β levels (by 49%) and increased soluble A β levels (by 50%) were accompanied with elevated Cu levels (and Zn). This increase of Cu- and Zn-ions might either be attributed to an inefficiency of the chelator with its known weak affinities for Zn ($K_1 = 7.0$) and for Cu ($K_1 = 8.9$) or even more likely, due to a facilitated uptake in brain of clioquinol–Cu complexes. The latter hypothesis has been experimentally confirmed in a yeast system in vitro (Treiber et al. 2004).

When AD patients were treated with clioquinol, the placebo group deteriorated faster than the clioquinol group suggesting a beneficial effect upon clioquinol treatment (Ritchie et al. 2003). In contrast, clioquinol treatment of APP transgenic mice was associated with premature death, which could be rescued by Cu supplementation (Schäfer

et al. 2007). At present there is no simple explanation for these apparently contradictory results.

Sparks and Schreurs (2003) challenged rabbits with a high-cholesterol diet and reported that the intake of minor concentrations of Cu (0.12 mg/l) in the drinking water impaired behavior and induced plaque-like structures in the hippocampus and temporal lobe. These surprising findings implied that tap water Cu concentration might influence AD. In the present study we did not find any correlation between plasma or CSF Cu and cholesterol levels.

Postmortem Cu levels in CNS of AD patients were found decreased (Deibel et al. 1996) or unchanged (Loeffler et al. 1996). Controversial results have also been published on the level of Cu in plasma and brain in AD patients. Plasma Cu levels being within the normal range (65–165 μg/dl) in AD patients are in good agreement with an earlier study (Jeandel et al. 1989). However, other studies suggest that elevated Cu levels might be a risk factor for AD without providing a rationale (Squitti et al. 2002, 2003, 2004). These results are in contrast to the present study, which might be explained by the fact that we studied Cu levels in the same patients within a 12 months period and can therefore better control for inter-individual differences. Short-term high Cu intake has been reported not to affect Cu status or functions related to Cu status, only long-term high Cu intake can result in increases in some parameters in young men (Turnlund et al. 2004).

The present clinical trial demonstrates that long-term oral intake of 8 mg Cu [Cu-(II)-orotate-dihydrate] can be excluded as a risk factor for AD based on CSF biomarker analysis. In addition, A β 42 levels declined significantly within 12 months in CSF indicating its value as a prognostic biomarker in addition to its common use for diagnosing AD.

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