



Negative Association of Cognitive Performance With Blood Serum Neurotoxicity and Its Modulation by Human Herpes Virus 5 (HHV5) Seropositivity in Healthy Women

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ABSTRACT: Identification of early influences on cognitive decline is of paramount importance in order to stem the impacts of decrements in cognitive functioning and to potentially intervene. Thus, here we focused on 132 healthy adult women (age range 26–98 years) to (a) determine whether factors circulating in serum may exert neurotoxic effects *in vitro*, (b) evaluate associations between serum neurotoxicity and cognitive performance, and (c) assess the influence of human herpes virus (HHV) seroprevalence and other factors on apoptosis and cognitive performance. The results documented that the addition of serum from healthy adult women to neural cell cultures resulted in apoptosis, indicating the presence of circulating neurotoxic factors in the serum. Furthermore, apoptosis increased with age, and was associated with decreased cognitive performance. Stepwise regression evaluating the influence of 6 HHVs on apoptosis and cognitive function revealed that only HHV5 (cytomegalovirus; CMV) seropositivity was significantly associated with apoptosis and cognitive decline, controlling for age. These findings document neurotoxic effects of serum from healthy women across the adult lifespan and suggest a unique detrimental influence associated with CMV seropositivity.

KEYWORDS: Apoptosis, human serum, neurotoxicity, human herpes viruses, cytomegalovirus, neuroblastoma N2A cells

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Introduction

Humans are exposed to myriad natural and synthetic neurotoxins, several of which have been implicated in neurodegeneration.^{1–8} Neurotoxins circulate in the blood and, therefore, are readily detected in blood and serum samples.⁹ Under healthy conditions, the brain is protected from exposure to neurotoxins via the blood-brain barrier (BBB); however, breakdown of the BBB enables entry of harmful substances into the brain, ultimately contributing to neuronal dysfunction and neurodegeneration.^{10,11} Several factors including sex, age, temperature, physical exercise, neurotransmitters, inflammatory substances, and inflammatory markers, among others, influence BBB permeability.^{12,13} In addition, infection with neurotropic viruses results in structural and functional changes in several components of the BBB, thereby affecting permeability and permitting entry of viruses and other pathogens into the brain.¹⁴

Numerous viruses are neurotropic and therefore capable of entering the nervous system and causing immediate or delayed neuropathological changes and disease.¹⁵ Human herpes

viruses (HHVs) are among the most common viruses worldwide and are associated with numerous and varied human health conditions.^{16–19} After initial infection, typically in childhood, HHVs establish latency and may be periodically reactivated by various factors including stress, fever, and UV exposure, as well as age-related immunological changes and conditions and/or treatments associated with immunosuppression.^{20,21} HHVs vary in terms of cell tropism. Some HHVs are neurotropic and establish latency within the nervous system; several others establish latency elsewhere (eg, B cells, monocytes) but are capable of entering the brain via the bloodstream or peripheral nerves. Consequently, several HHVs have been documented in brain tissue,^{15,22–26} and increasing evidence points to a role of HHVs in conditions affecting the brain.^{19,25,27–37} HHVs are ubiquitous; consequently, nearly 100% of adult humans are seropositive for at least 1 type of HHV. Five HHVs are most commonly implicated in human disease—namely, herpes simplex virus type 1 (HSV/HHV1) and 2 (HSV/HHV2), varicella zoster virus (VZV/HHV3), Epstein-Barr virus (EBV/HHV4), cytomegalovirus (CMV/



HHV5).²¹ Burgeoning evidence also links HHV6 and HHV7 to neurodegenerative diseases.^{26,38-41}

Here, we bridge these separate lines of research implicating neurotoxins and HHVs in neuropathology by assessing the effects of serum obtained from healthy adult women on neural cell cultures and with regard to HHV seropositivity. Previous research in our lab and elsewhere has documented neurotoxic effects of serum from patient populations⁴² and older adults⁴³ on neural and hippocampal progenitor cell cultures, respectively. It is not clear whether findings documenting the presence of harmful substances circulating in the blood in those populations extend to younger, healthy samples; however, since subtle neuropathological changes are evident decades before symptom onset and clinical diagnoses,⁴⁴ it is important to assess for the presence of potentially neurotoxic factors in healthy individuals across the lifespan. Thus, we first evaluated whether the addition of serum from healthy adult women to neural cell cultures resulted in apoptosis, and assessed the correspondence between apoptosis and cognitive performance. Then, in light of immense literature implicating HHVs in conditions affecting the brain, we evaluated apoptosis and cognitive performance with regard to seropositivity of neurotropic HHVs. Since apolipoprotein E and body mass index (BMI) have been shown to influence cognitive function,^{45,46} we also evaluated and controlled for their influence in this sample.

Materials and Methods

Participants

A total of 132 healthy women participated in the study as paid volunteers. This data was obtained as part of an ongoing longitudinal study involving annual data acquisition; consequently, the number of annual visits varied for participants as follows: 60 participants completed 1 annual visit, 56 completed 2 annual visits, and 16 completed 3 annual visits, for a total of 220 visits. Women were excluded from participation if they had been diagnosed at any point in their lifetime with a neurological disorder, any autoimmune disorder associated with neurocognitive dysfunction (eg, systemic lupus erythematosus, rheumatoid arthritis), any major medical condition affecting brain function (eg, brain cancer, head injury with cognitive sequelae), serious psychiatric diagnoses (eg, bipolar disorder, schizophrenia, any history of psychiatric hospitalization), or any recent/current medication or treatment known to affect brain functioning (eg, radiation, chemotherapy). Written informed consent was obtained from study participants. The institutional review board and relevant committees of the Minneapolis VA Health Care System approved the study protocol.

Cognitive assessment

The Montreal Cognitive Assessment (MoCA)⁴⁷ was administered to participants to screen for cognitive impairment. The MoCA assesses several domains of cognitive function including executive function, memory, language, and abstract reasoning, among others. Scores for each domain are added to reflect a total score ranging from 0 to 30; total scores greater than 25 are considered normal.⁴⁷ At recruitment, all participants had a total MoCA score (without the education point) of greater than 25. MoCA assessment was obtained during all but 1 visit, for a total of 219 MoCA assessments.

ApoE genotyping

DNA samples were genotyped using PCR amplification followed by restriction enzyme digestion as per recommendations.⁴⁸ Known apoE isoform standards were included in the analysis.

HHV seropositivity assays

We determined the seroprevalence of IgG antibodies against HHV1, HHV2, HHV3, HHV4, HHV5, and HHV6; we could not obtain reliable results from kits for HHV7 and HHV8. We employed standard method using commercially available indirect Enzyme-Linked Immunosorbent Assays (ELISA). The ELISA tests were performed as per the manufacturer's instructions and recommendations. The specimens were processed using a mini-automated 5-in-1 workstation (Crocodile cat. 84024-01; Berthold Technologies, Oak Ridge, TN, USA). The workstation includes the ELISA microtiter plate reader which was read at dual wavelengths for absorbance at 450 and 620 nm as reference wavelengths. Details of the virus-specific ELISA kits are as follows. (a) HHV1: Human Anti-Herpes simplex virus Type 1 IgG ELISA Kit (HSV1), Abcam Inc., Boston, MA USA, cat. ab 108737; (b) HHV2: Human Anti-Herpes simplex virus Type 2 IgG ELISA Kit (HSV2) Abcam cat. ab 108739; (c) HHV3: Human Anti-Varicella-Zoster virus IgG ELISA Kit (VZV) Abcam cat. ab 108782; (d) HHV4: Human Anti-Epstein Barr virus IgG ELISA Kit (EBV-VCA) Abcam cat. ab 108730; (e) Human Anti-Cytomegalovirus IgG ELISA Kit (CMV) Abcam cat. ab 108724; (f) Human Herpesvirus 6 IgG ELISA Kit cat. KA1457, Abnova, Taiwan. Seroprevalence for 1 or more HHVs were missing for 7 women.

Body mass index (BMI)

BMI was calculated from anthropometric measurements. Weight was not able to be obtained for a small number of acquisitions ($n=7$), precluding BMI calculations in those instances; BMI was available for 213 acquisitions and was used as a continuous measure in analyses.

Neural cell culture

Neuro-2A neuroblastoma (N2A) cells were cultured in Eagle's minimal essential medium (EMEM, ATCC, VA, USA) containing 10% fetal bovine serum (ThermoFisher Scientific, Waltham, MA) poly-D-lysine coated, 24-well plates at a concentration of 30000 to 50000/well for 48 to 72 hour. The medium was then changed to Neurobasal containing N2 supplement and L-glutamine (ThermoFisher Scientific, Waltham, MA), in the absence (medium control) or presence of human serum. For all experiments, human serum (10%) was added and the cells were cultured for 2 more days. All experiments were carried out in quintuplicate.

Cell apoptosis with terminal deoxynucleotidyl transferase mediated dUTP Nick End Labeling assay (TUNEL) assay

The extent of cell apoptosis of Neuro-2A cells incubated with women's serum was examined. All samples were examined using 4- and 8-chamber glass slides (ThermoFisher Scientific, Waltham, MA) coated with poly-D-lysine at 50 µg/ml as mentioned above. N2A cells were seeded at a concentration of 50000 to 100000 cells per chamber, in 1 ml of Neurobasal/ N2/ L glutamine medium for 2 days. In sequence, 10% of each sample of women's serum, incubated for 60 minutes at 37°C. At the end of the incubation period, the cells were examined for apoptosis. Apoptotic cells were detected using the In Situ Cell Death Detection Kit, TMR red (Terminal deoxynucleotidyl transferase (TdT) enzyme and fluorochrome labeling solution), according to the manufacturer's protocol. Briefly, the cells were fixed in ice-cold methanol for 10 minutes at room temperature, rinsed with PBS and permeabilized with 0.1% Triton X-100 in PBS for 3 minutes on ice. The cells were then incubated with 150 µl of TUNEL reaction mixture for 60 minutes at 37°C in the dark (Insitu Cell Death Detection Kit, TMR red, ThermoFisher scientific, Waltham, MA). The cells were then washed 3X with PBS and Diamond AntiFade mounting medium with 4',6-diamidino-2-phenylindole (DAPI) stain (ThermoFisher Scientific, Waltham, MA) was used for visualization of nuclei, using the EVOS FL Cell Imaging System (ThermoFisher Scientific, Waltham, MA) or an Olympus 3000 confocal microscope. The 8 to 10 images were obtained from different fields from a minimum of 2 experiments with each different experimental condition. Apoptosis was then calculated with ImageJ software by measuring the number of TUNEL-labeled cells (red nuclei) relative to the total cell number (DAPI-stained nuclei). Since experiments were run in quintuplicate, there were 5 Media and 5 Test apoptosis values per run (A_M and A_T , respectively), for a total of 220 runs \times 5/run = 1100 data values for each A_M and A_T . A_M served as the baseline for apoptosis in a particular run and, hence, was used as a covariate in all analyses on A_T .

Data Analysis

MoCA

MoCA values are counts and hence were square-root transformed, a transformation appropriate for the statistical analysis of counts⁴⁹:

$$s\text{MoCA} = \sqrt{\text{MoCA}} \quad (1)$$

All MoCA-related statistical analyses were performed using sMoCA.

ApoE

With respect to apoE genotype, 3 apoE isoforms (E4, E3, and E2) are the result of cysteine-arginine interchanges at 2 sites: there are zero interchanges in E4, 1 interchange in E3, and 2 interchanges in E2. The resulting 6 apoE genotypes (E4/4, E4/3, E4/2, E3/3, E3/2, E2/2) yield 5 groups with respect to the number of cysteine residues per mole, as follows. ApoE4/4 contains zero cysteine residues per mole (0-CysR/mole), E4/3 contains 1 (1-CysR/mole), E4/2 and E3/3 each contain 2 (0-CysR/mole), E3/2 contains 3 (0-CysR/mole), and E2/2 contains 4 (0-CysR/mole). The use of the number of cysteine residues per mole to characterize the apoE molecule converts the categorical apoE genotype scale, consisting of 6 distinct genotypes above, to a 5-point continuous scale (0-4 cysteine residues per mole), a conversion that allows the use of statistical analyses suitable for continuous variables (eg, regression) to quantify the relations between various variables and apoE.^{50,51} Therefore, we used the number of cysteine residues per mole (CysR/mole) as a covariate to evaluate the possible association of apoE with MoCA and apoptosis. In addition, as a measure complementary to CysR/mole above, we computed a binary variable, E4, based on the presence [apoE4(+)=1] or absence [apoE4(-)=0] of the apoE4 allele in the participant's apoE genotype and used it as a fixed group factor in statistical analyses below.

Implementation of statistical analyses

The IBM-SPSS Statistical package (version 29) was used to analyze the data, including Pearson correlation, partial correlation, and analysis of covariance (ANCOVA). All P-values reported are 2-tailed.

Results

Age, MoCA, apoE, BMI

The mean (\pm SEM) age of the study participants was 70.4 \pm 0.79 years, (range 25.7-97.8 years, N=220), the mean of the MoCA was 27.25 \pm 0.16 (range 17-30, N=219), and the mean BMI was 28.28 \pm 0.44 (N=213). The frequency distributions of age, MoCA, and BMI are shown in Figure 1. The frequency distribution of the apoE genotypes and the number of apoE CysR/mole are given in Tables 1 and 2, respectively;

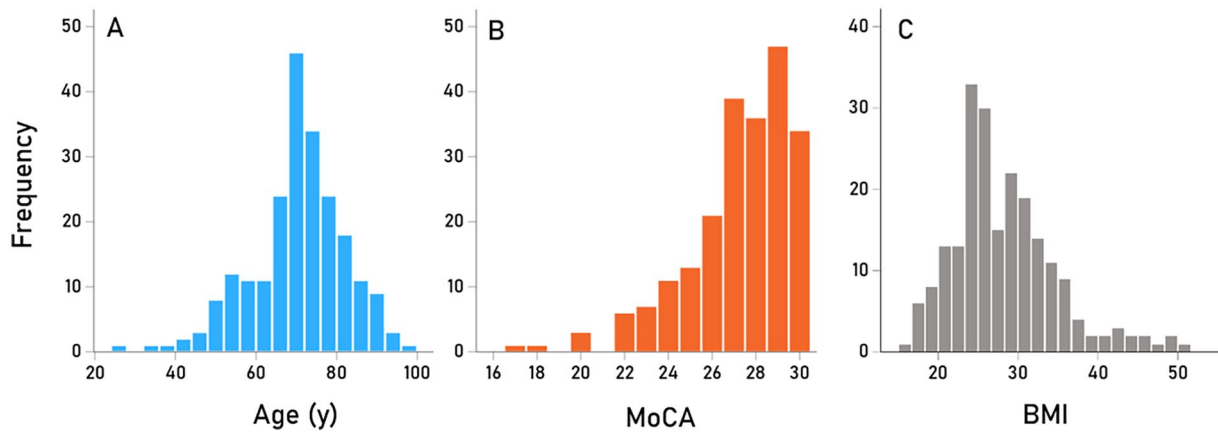


Figure 1. (A) Frequency distribution of ages at time of test. N=220 participants. (B) Frequency distribution of MoCA scores at time of test. N=220 participants. (C) Frequency distribution of BMI. N=213 participants.

Table 1. Frequency distribution of apoE genotypes.

APOE GENOTYPE	FREQUENCY (%)
23	22 (10)
24	4 (1.8)
33	131 (59.8)
34	49 (22.4)
44	13 (5.9)
Total	219 (100)

Table 2. Frequency distribution of number of apoE CysR/mole.

N OF APOE CYSR/ MOLE	APOE	FREQUENCY (%)
0	E4/E4	13 (5.9)
1	E3/E4	49 (22.4)
2	E2/E4, E3/E4	135 (61.6)
3	E2/E3	22 (10)
Total		219 (100)

there were no E2/E2 homozygotes. Analyses evaluating associations between sMoCA, apoE, BMI, and age revealed several significant associations. Age was associated with decreased sMoCA (Pearson $r = -.308$, $P < .001$, $N = 219$; Figure 2A) and decreased BMI ($r = -.373$, $P < .001$, $N = 213$). BMI was not significantly associated with sMoCA, controlling for age (partial correlation $r = -.0117$, $P = .097$, $N = 213$). Finally, sMoCA increased significantly with the number of apoE CysR/mole, controlling for age (sMoCA vs CysR/mole, partial correlation = .154, $P = .024$, $N = 218$) and was significantly higher in the apoE4(+) than the apoE4(-) group ($P = .042$, F-test in ANCOVA, where sMoCA was the dependent variable, apoE4 was a (binary) fixed factor, and age was a covariate).

Apoptosis

Next we evaluated the associations between sMoCA, apoE, BMI, and age with apoptosis. We found that apoptosis increased significantly with increasing age (Partial correlation A_T vs age = .106, controlling for A_M , $P < .001$, $N = 1220$; Figure 2B) and that sMoCA decreased significantly with apoptosis (Partial correlation sMoCA vs $A_T = -0.092$, controlling for A_M and age, $P = .002$, $N = 1095$; Figure 2C). Apoptosis was not significantly associated with BMI (partial correlation A_T vs BMI = -.002, controlling for A_M and age, $P = .941$, $N = 1065$) or apoE CysR/mole (partial correlation A_T vs number of apoE CysR/mole = .027, controlling for A_M and age, $P = .368$, $N = 1095$). Similarly, apoptosis A_T did not differ significantly between apoE4(+) and apoE4(-) groups ($P = .737$, F-test in ANCOVA, where A_T was the dependent variable, apoE4 was a binary fixed factor, and A_M and age were covariates).

HHV

The seroprevalence of HHV1-6 are given in Table 3. The effect of HHV seroprevalence on apoptosis was evaluated using a stepwise multiple linear regression, where A_T was the dependent variable, and HHV1, HHV2, HHV4, HHV5, and HHV6 seroprevalence status (coded as binary 0/1 values), age, and A_M were the independent variables. This analysis yielded only HHV5 seroprevalence as having a highly statistically significant positive effect on apoptosis ($P = .005$, t -test on the HHV5 partial regression coefficient), in addition to A_M and age (as expected). Collinearity diagnostics confirmed the absence of possible collinearities: all tolerance values were >0.9 and all variance inflation factor (VIF) <1.1 . Detailed results of the regression analysis above are given in Table 4.

The HHV5 effect was quantified by performing an ANCOVA where A_T was the dependent variable, HHV5 serostatus was a fixed factor, and A_M , age, number of apoE CysR/mole, and BMI were covariates. The effect of HHV5

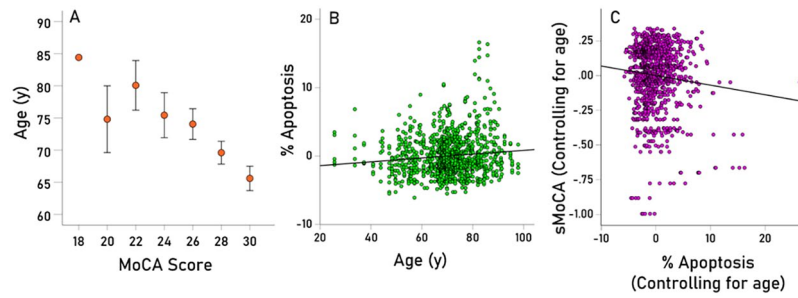


Figure 2. (A) mean (\pm SEM) MoCA scores are plotted against age. N=220. (B) % apoptosis A_T (adjusted for A_M) is plotted against age. N=1100. (C) sMoCA (adjusted for age) is plotted against % apoptosis A_T (adjusted for A_M and age). N=1100. See text for details.

Table 3. Prevalence of seropositivity for the 6 HHV strains investigated. Numbers in the column Total indicate the number of tests that gave unambiguous results in the ELISA.

HHV STRAIN		N SEROPOSITIVE	TOTAL	PERCENT SEROPOSITIVE
HHV-1	HSV1	110	209	52.6
HHV-2	HSV2	46	216	21.3
HHV-3	VZV	216	216	100.0
HHV-4	EBV	211	218	96.8
HHV-5	CMV	101	218	46.3
HHV-6		200	207	96.6

Table 4. Results of the multiple regression analysis described in the text.

	STANDARDIZED β	T-VALUE	P-VALUE	TOLERANCE	VIF ^A
HHV-1	-.023	0.719	.472	0.946	1.057
HHV-2	-.018	0.564	.573	0.930	1.075
HHV-4	-.03	0.932	.351	0.953	1.049
HHV-5	.094	2.941	.003	0.951	1.052
HHV-6	.019	0.600	.548	0.984	1.017
Age (y)	.097	3.050	.002	0.963	1.038
Media (A_M)	.215	6.790	<.001	0.979	1.021

^AVIF, variance inflation factor. Bold denotes significant effect on HHV seroprevalence on apoptosis.

seropositivity on apoptosis is illustrated in Figure 3. The mean (\pm SEM) A_T was $3.553\% \pm 0.119$ for HHV5(-) versus $4.063\% \pm 0.128$ for HHV5(+), a 14.3% increase in HHV5(+) over HHV5(-). No other HHV had a significant effect. Finally, HHV5 seropositivity had the following 2 additional MoCA-related effects. First, a significant decrease in MoCA with increasing apoptosis was observed only in the HHV5(+) group (partial correlation of A_T with sMoCA = $-.100$, $P = .025$, $N = 505$); for the HHV5(-) group this correlation was $.058$, $P = .151$, $N = 580$. And second, a positive association of MoCA with the number of apoE CysR/mole was observed in the HHV5(-) group (partial correlation, controlling for age = $.231$,

$P = .013$, $N = 116$) but was not present in the HHV5(+) group (partial correlation = $.017$, $P = .864$, $N = 101$), indicating a negative HHV5-mediated modulation of the protective effect of apoE CysR/mole on MoCA.

Discussion

In this study we evaluated the effect of serum from healthy women on neural cell cultures and with regard to cognitive performance and HHV seroprevalence. The findings documented the presence of neurotoxic factors, unrelated to BMI or apoE, that resulted in apoptosis and influenced cognitive function. Previous studies have documented serum neurotoxicity in older

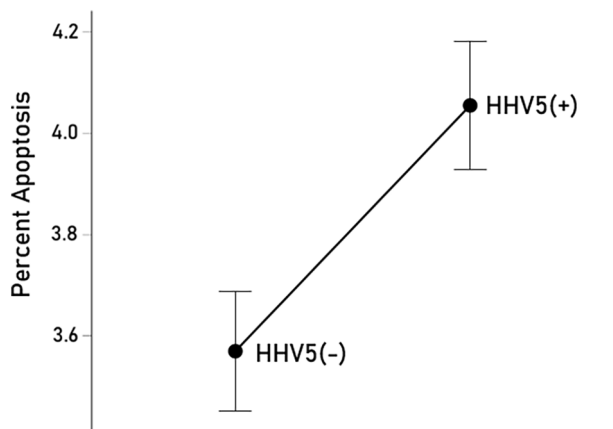


Figure 3. Mean % apoptosis $A_T \pm \text{SEM}$ (adjusted for A_M , age, apoE CysR/mole, and BMI) is plotted for HHV-5(-) and HHV-5(+) groups.

adults⁴³ and in certain patient populations⁴²; here, we show neurotoxic factors resulting in *in vitro* apoptosis and decreased cognitive performance in healthy women across the adult lifespan. This finding highlights deleterious effects of neurotoxic factors circulating in the blood. Since HHVs have been widely implicated in conditions affecting the brain, we then evaluated the influence of seroprevalence of 6 HHVs, BMI, and apoE on apoptosis and cognitive performance. The findings documented a unique influence of HHV5/CMV seropositivity on increased apoptosis and decreased cognitive performance. Notably, apoE CysR/mole was associated with protective effects on cognitive performance in CMV seronegative women but not those who were seropositive. Taken together, the findings of this study point to neurotoxic factors in serum of healthy women that may be associated with CMV exposure, adding to the literature regarding the potential for harmful downstream effects of CMV infection on brain health.

CMV is a common infection worldwide⁵² that is transmitted congenitally or through direct contact with body fluids. The seroprevalence of CMV increases with age such that approximately half of women of childbearing age and two-thirds of women age 70 to 79 are seropositive.^{53,54} Like other human herpes viruses, CMV establishes lifelong persistence characterized by periods of latency and reactivation.²⁰ Age-related increases in chronic reactivation of CMV,⁵⁵ coupled with increased CMV viral load,⁵⁶ and CMV-specific adaptive immune system dysregulation (eg, inflation of memory T cells and low or inverted CD4/CD8 ratio)^{57,58} exhaust the immune system over time.⁵⁷ These effects are thought to contribute to low-grade chronic inflammation (ie, “inflammaging”),⁵⁹ immunosenescence,⁶⁰ epigenetic aging,⁶¹ and increased mortality.^{62,63} To that end, CMV seropositivity has been associated with various chronic health conditions including atherosclerosis⁶⁴ and vascular dysfunction,⁶⁵ autoimmune disorders,⁶⁶ and cancer,⁶⁷ as well as age-related frailty⁶⁸ (cf Mathei et al⁶⁹), functional impairment,⁷⁰ and cognitive decline.^{71,72}

Here, we showed correspondence between neurotoxic effects of serum from CMV seropositive women and decrements in cognitive performance. These findings suggest that the presence of harmful agents circulating in the blood of CMV seropositive individuals may impact brain function. Previous studies have identified CMV as a risk factor for cognitive impairment and dementia⁷¹⁻⁷⁵ (cf, Mathei et al⁶⁹). Furthermore, CMV is present in a high proportion of brains from vascular dementia patients,⁷⁶ and recently, CMV has been shown to induce tau pathology, a hallmark of Alzheimer’s dementia.⁷⁷ The extent to which the neurotoxic effects of CMV seropositive serum observed in these healthy women is a predictor of future cognitive dysfunction remains to be determined and is an ongoing area of investigation in our lab.

These novel findings implicating CMV in the association between serum neurotoxicity and cognitive function must be considered within the context of study limitations. First, we used neural cell cultures as a proxy for the brain and it is not clear to what extent the deleterious *in vitro* effects correspond with *in vivo* effects in the human brain. Remarkably, the *in vitro* effects were associated with cognitive function suggesting some correspondence between *in vitro* and *in vivo* effects. Second, the present study focused exclusively on women. Since previous studies have documented sex-specific neurotoxic effects,⁷⁸ it is unknown whether the present findings generalize to men. Third, our analyses were limited to presence or absence of HHV antibodies, precluding evaluation of effects of titer concentration on apoptosis and cognitive performance. Finally, it is worth noting that the present findings do not preclude involvement of other potentially neurotoxic factors; nonetheless, the findings document serum neurotoxicity and decrements in cognitive performance in CMV seropositive women. Consequently, the development of CMV vaccines⁷⁹ may be beneficial in preventing CMV-related cognitive decline.

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Author Contributions

APG: conceived the research; LMJ, EPT, and EJW contributed to data acquisition; APG, LMJ, and EPT wrote the paper; all authors edited and approved the final version of the paper.

Data Availability

Data will be available to qualified researchers by the corresponding author on a reasonable request.

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