

1 **SUPPLEMENTARY DATA**

Supplementary Table 1 Primers and probes used in this study

Name	Target	Sequence ^a
HSV-1 US4_Fw	Human herpesvirus 1 <i>US4</i>	TCCTSGTTCCTMACKGCCTCCC
HSV-1 US4_Rv	Human herpesvirus 1 <i>US4</i>	CGTCTGGACCAACCGCCACACAGGT
HSV-1 US4_probe	Human herpesvirus 1 <i>US4</i>	<i>FAM</i> -GCAGICAYACGTAACGCACGCT- <i>BHQ1</i>
HSV-1 LAT_Fw	Human herpesvirus 1 <i>LAT</i>	CCCACGTA CTCCAAGAAGGC
HSV-1 LAT_Rv	Human herpesvirus 1 <i>LAT</i>	AGACCCAAGCATAGAGAGCCAG
HSV-1 LAT_probe	Human herpesvirus 1 <i>LAT</i>	<i>FAM</i> -CCCACCCCGCCTGTGTTTTTGTG- <i>BHQ1</i>
VZV ORF38_Fw	Human herpesvirus 3 <i>ORF38</i>	AAGTTC CCCCCGTTTCGC
VZV ORF38_Rv	Human herpesvirus 3 <i>ORF38</i>	CCGCAACA ACTGCAGTATATATCGTCTCA
VZV ORF38_probe	Human herpesvirus 3 <i>ORF38</i>	<i>FAM</i> -TGGACTTGAAGATGAACTTAATGAAGC- <i>BHQ1</i>
HMBS_Fw	Homo sapiens <i>HMBS</i>	GCCTGCAGTTTGAAATCAGTG
HMBS_Rv	Homo sapiens <i>HMBS</i>	CGGGACGGGCTTTAGCTA
HMBS_probe	Homo sapiens <i>HMBS</i>	<i>FAM</i> -CGCAGGCACTCGTACTGCTCGCT- <i>BHQ1</i>
ApoE2_Fw	Homo sapiens <i>APOE2</i>	GCGGACATGGAGGACGTGT
ApoE2_Rv	Homo sapiens <i>APOE2</i>	CCTGGTACTG CCGAGGCA
ApoE3_Fw	Homo sapiens <i>APOE3</i>	CGGACATGGAGGACGTGT
ApoE3_Rv	Homo sapiens <i>APOE3</i>	CTGGTACTG CCGAGGCG
ApoE4_Fw	Homo sapiens <i>APOE4</i>	CGGACATGGAGGACGTGC
ApoE4_Rv	Homo sapiens <i>APOE4</i>	CTGGTACTG CCGAGGCG
ApoE_Probe	Homo sapiens <i>APOE</i>	<i>FAM</i> -CAGCTCCTCGGTGCTCTGGC- <i>BHQ1</i>
ApoE_seq1	Homo sapiens <i>APOE</i>	AGCCCTTCTCCCCGCCTCCCACTGT
ApoE_seq2	Homo sapiens <i>APOE</i>	CTCCGCCACCTGCTCCTTCACCTCG

^a5' and 3'-modifications of TaqMan probes are indicated. *FAM* 6-Carboxyfluorescein, *BHQ1* black hole quencher 1

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Supplementary Table 2 FFPE TG samples used in this study

Subject	Age ^a	Gender ^b	PMI ^c	Cause of death	Neurological disease	HSV-1 status ^d
1	81	F	3:35	Cachexia and dehydration	Alzheimer's disease	negative
2	82	F	5:55	Cardiac arrest	Alzheimer's disease	negative
3	89	F	4:30	Peritonitis	Alzheimer's disease	positive
4	87	F	8:10	Uremia by dehydration	Alzheimer's disease	positive
5	86	F	5:15	Cachexia and dehydration	Alzheimer's disease	positive
6	81	M	8:21	Euthanasia	Parkinson's disease	positive
7	62	F	12:35	Cachexia with pulmonary insufficiency	Multiple sclerosis	positive
8	80	F	9:30	Cardiac arrest	Bipolar disorder	positive
9	103	F	6:35	Dehydration and pneumonia	Tauopathy	positive
10	89	M	6:50	Urosepsis	non-demented control	positive

^aAge in years; ^bF, female; M, male; ^cPMI, post-mortem interval; ^dHSV-1 infection status based on serology and HSV-1 LAT ISH

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Supplementary Table 3 TBI patients used in this study

Subject	Age ^a	Gender ^b	Interval TBI – death
1	84	M	0 days (acute)
2	48	M	7 days
3	13	F	11

^aAge in years; ^bF, female; M, male

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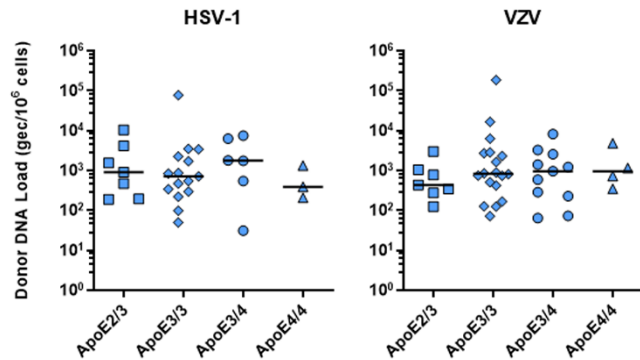
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Supplementary Table 4 FFPE AD samples used in this study

Subject	Age ^a	Gender ^b	PMI ^c	Cause of death	Neurological disease	HSV status ^d
1	88	M	4:40	Cachexia and dehydration	Alzheimer's disease	positive
2	74	M	8:00	Dehydration by possible cystitis	Alzheimer's disease	positive
3	86	M	05:10	Cachexia and dehydration	Alzheimer's disease	positive

^aAge in years; ^bF, female; M, male; ^cPMI, post-mortem interval; ^dHSV infection status based on HSV-specific IgG serology data

11 **SUPPLEMENTARY FIGURE 1**



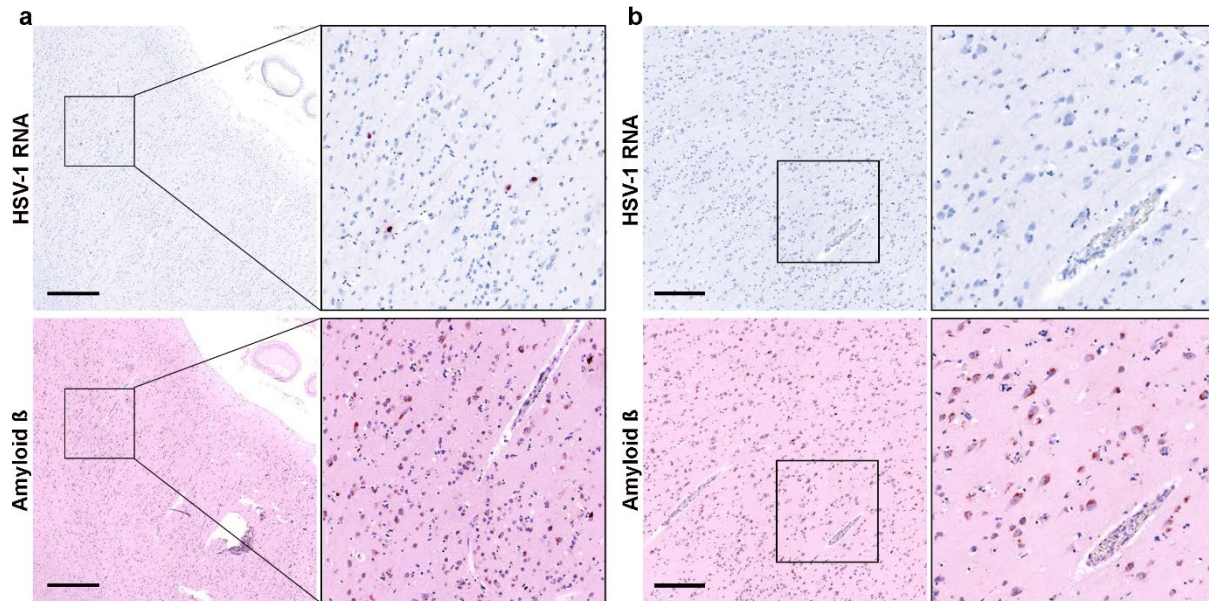
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14 **Supplementary Fig. 1** Quantification of latent HSV-1 and VZV DNA load in human TG
15 stratified on *APOE* genotype. HSV-1- and VZV-specific qPCR and *APOE* genotyping was
16 performed on DNA extracted from the trigeminal ganglia (TG) of AD patients and controls.

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18 **SUPPLEMENTARY FIGURE 2**



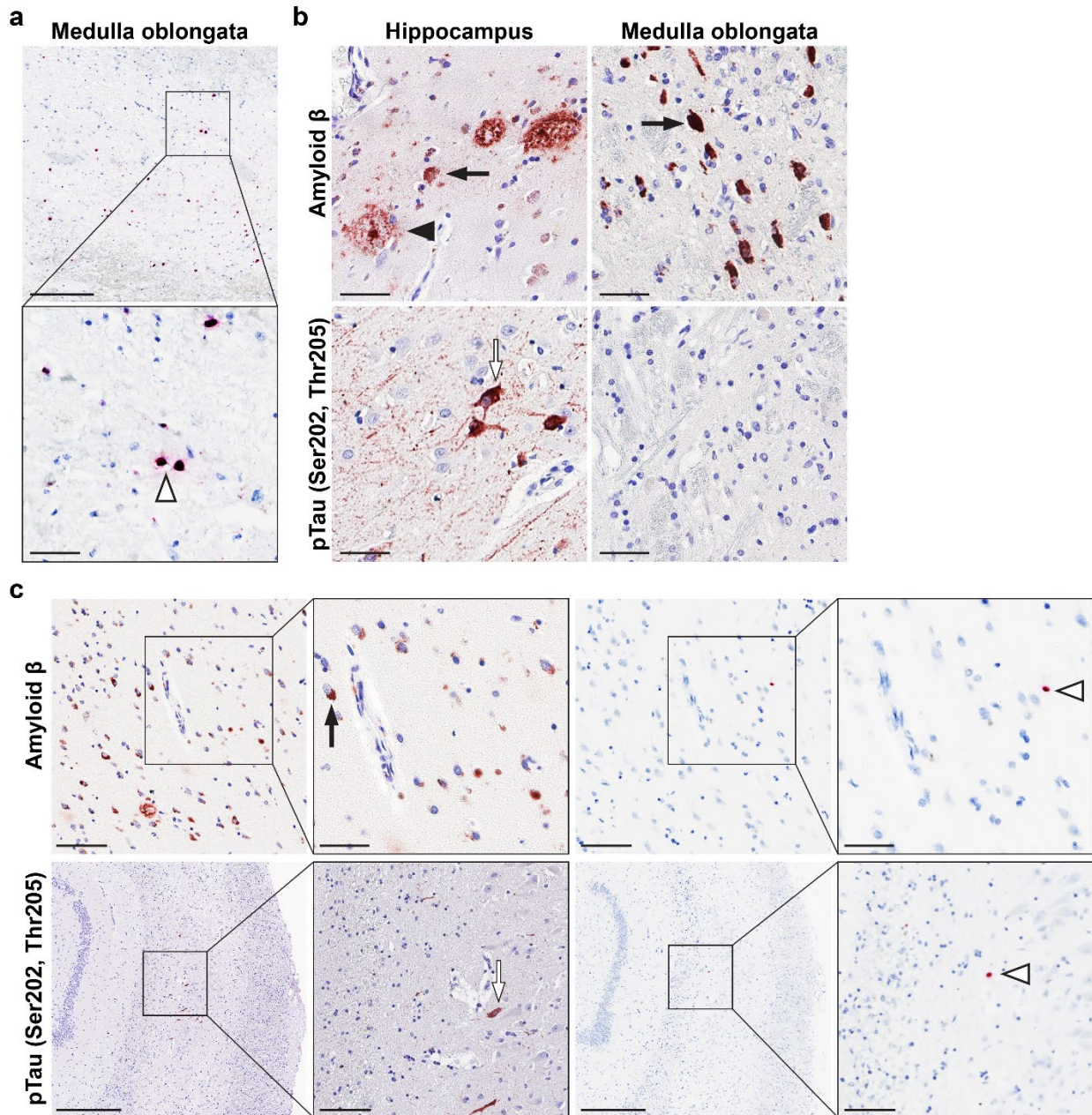
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20 **Supplementary Fig. 2** Intra-neuronal accumulation of A β protein is not restricted to
21 areas of HSV-1 infection in brains of HSE patients. Consecutive slides were stained for
22 HSV-1 RNA by ISH and A β by IHC. A β can be seen in areas with HSV-1 RNA (**a**) and in
23 areas without HSV-1 RNA (**b**). Data shown for HSE donor 5. Scale bar: 500 μ m (**a**) and
24 250 μ m (**b**).

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26 SUPPLEMENTARY FIGURE 3

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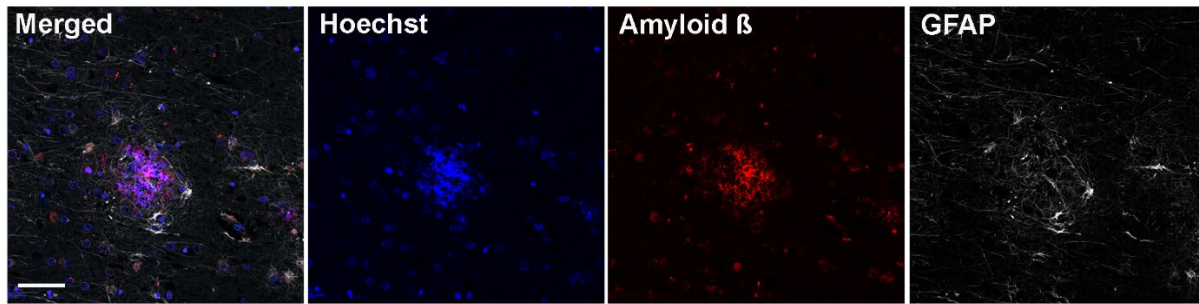
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29 **Supplementary Fig. 3** Lytic VZV infection is not associated with A β plaques or NFT in
30 brain of a VZV encephalitis patient. **a** Brain section stained for VZV RNA by ISH. Box
31 indicates area shown at higher magnification. Open arrowhead indicates examples of
32 VZV RNA-expressing cells. Scale bars indicate 250 μ m (top) and 50 μ m (bottom). **b** Brain

33 sections IHC stained for A β and pTau (Ser²⁰²/Thr²⁰⁵). Scale bars indicate 50 μ m. **c**
34 Consecutive brain sections were IHC stained for A β and pTau (Ser²⁰²/Thr²⁰⁵) or stained
35 for HSV-1 RNA by ISH. Boxes indicate areas shown at higher magnification. Filled arrow
36 indicates intracellular A β protein, open arrow indicates NFT and open arrowhead
37 indicates VZV RNA-expressing cells. Scale bars indicate 50 μ m (A β , high magnification),
38 100 μ m (A β , low magnification; pTau high magnification) and 500 μ m (pTau, low
39 magnification).

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41 **SUPPLEMENTARY FIGURE 4**

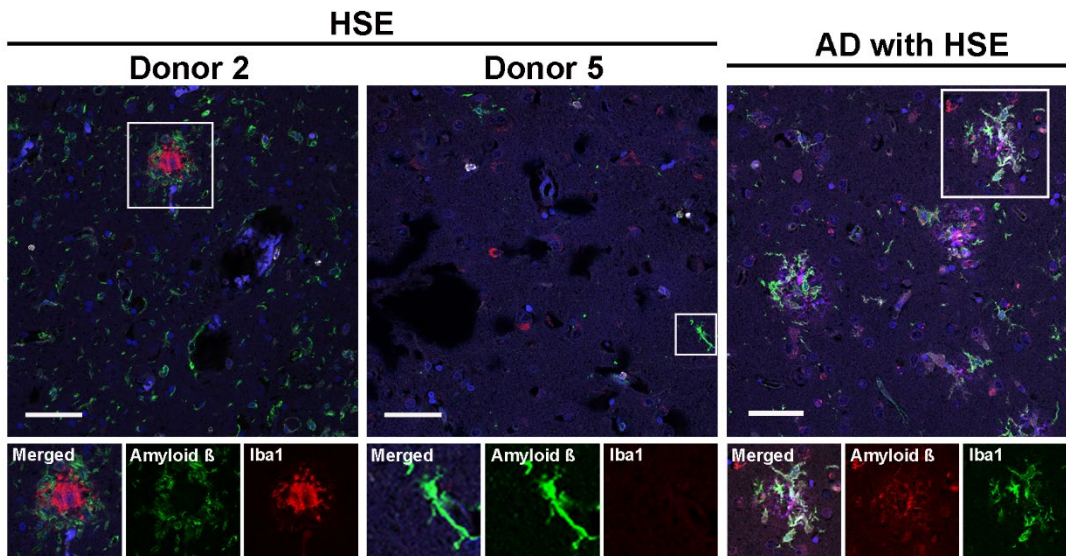


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43 **Supplementary Fig. 4** Prominent GFAP staining of astrocytes interacting with A β
44 plaques brain of an AD patient with concurrent HSE. IF staining for A β (red), GFAP (white)
45 and nuclei (Hoechst-33342; blue). Scale bar: 50 μ m.

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47 **SUPPLEMENTARY FIGURE 5**



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49 **Supplementary Fig. 5** Microglia morphology and density in the brain of HSE patients
50 and AD patient with HSE. Brain tissue sections were IF stained for A β (red), Iba1 (green)
51 and nuclei (Hoechst-33342; blue). Boxes indicate areas shown at higher magnification.
52 Scale bar: 50 μ m.

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