

RESEARCH

Open Access



Time trends in herpesvirus seroepidemiology among Swedish adults

Jan Olsson^{1*}, Sema Nourmohammadi¹, Emma Honkala¹, Anders Johansson², Göran Hallmans³, Bodil Weidung⁴, Hugo Lövheim⁵ and Fredrik Elgh¹

Abstract

Background Human herpesviruses are widespread among the human population. The infections often occur unnoticed, but severe disease as well as long-term sequelae are part of the symptom spectrum. The prevalence varies among subpopulations and with time. The aim of this study was to describe the seroprevalence of Immunoglobulin G against *Herpes simplex 1*, *Herpes simplex 2*, Epstein-Barr virus and Cytomegalovirus in the adult Swedish population over a time period of several decades.

Methods Serum samples ($n = 892$) from biobanks, originating from 30-year-old women, 50-year-old men and 50-year-old women sampled between 1975 and 2018, were analyzed for presence of anti-herpesvirus antibodies. Linear regression analysis was used to test for a correlation between birth year and seroprevalence. Multiple linear regression analysis was used to differentiate between other factors such as age and gender.

Results Birth year correlated negatively with the prevalence of immunoglobulin G against *Herpes simplex 1* and Epstein-Barr virus ($p = 0.004$ and 0.033), and positively with Immunoglobulin G against Cytomegalovirus ($p = 0.039$). When participant categories were analyzed separately, birth year correlated negatively with the prevalence of Immunoglobulin G against *Herpes simplex 1* and *Herpes simplex 2* ($p = 0.032$ and 0.028) in 30-year-old women, and with the prevalence of Immunoglobulin G against Cytomegalovirus in 50-year-old men ($p = 0.011$).

Conclusions The prevalence of Immunoglobulin G against *Herpes simplex 1*, *Herpes simplex 2* and Epstein-Barr virus decreases in later birth cohorts. This indicates a trend of declining risk of getting infected with these viruses as a child and adolescent.

Keywords Herpes, Herpes simplex, Epstein-Barr virus, Cytomegalovirus, Seroprevalence, Epidemiology, Time trends, Immunoglobulin G

Background

Human herpesvirus (HHV1-8) are globally distributed among the human population, causing a variety of diseases. Albeit phylogenetically closely related, the primary infections caused by the different HHVs are diverse both in symptoms spectra and the interplay in a wider sense between virus and human host. HHV-1&2 (HSV-1&2) cause oral and genital herpes with typical lesions, HHV-3 (VZV) cause chickenpox and shingles, HHV-4 (EBV) cause infectious mononucleosis, HHV-6&7 are the causative agents for roseola. Congenital infections of HHV-2 and HHV-5 (CMV)

*Correspondence:

Jan Olsson
jan.l.olsson@umu.se

¹ Department of Clinical Microbiology, Umeå University, Umeå, Sweden

² Department of Odontology, Umeå University, 901 97 Umeå, Sweden

³ Department of Public Health and Clinical Medicine, Umeå University, Umeå, Sweden

⁴ Department of Public Health and Caring Sciences, Clinical Geriatrics, Uppsala University, Uppsala, Sweden

⁵ Department of Community Medicine and Rehabilitation, Geriatric Medicine, Umeå University, Umeå, Sweden



© The Author(s) 2024. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>. The Creative Commons Public Domain Dedication waiver (<http://creativecommons.org/publicdomain/zero/1.0/>) applies to the data made available in this article, unless otherwise stated in a credit line to the data.

poses a high risk for the developing fetus [1]. Typical for infections by HHV are the lifelong carriership once infected together with episodes of reactivated infection [1]. Several HHVs are linked to cancer; carriership of EBV drastically increase the risk for lymphoma and for carcinoma in the nasopharynx region [2], and HHV-8 is strongly associated with Kaposi's sarcoma [3]. In addition to the well-characterized symptoms associated with primary infections and reactivations, several HHVs have been attributed roles in the development of chronic neurological disorders, most prominently represented by HSV-1 for Alzheimer's disease and EBV for Multiple Sclerosis [4–7]. Several HHVs may also act in concert to alter the disease trajectory, especially CMV and HHV-6 have been suggested such adjuvant roles [4, 8, 9]. When studied from an epidemiological perspective, factors such as geographic location, socio-economic status and age influence the rate of acquisition of HHV infection [1]. In addition, host factors such as comorbidities, coinfections and immunosuppression influence both acquisition, reactivation and severity of symptoms [10–12]. As most herpesvirus infections never end up in medical records, the detection of anti-HHV Immunoglobulin G (IgG) is currently the gold standard when screening for a history of HHV infection. An individual might be carrier of anti-HHV IgG (positive) or not (negative) and the proportion of positive individuals in a population is denoted seroprevalence. Well-grounded estimates of seroprevalence are helpful not only when assessing impact from herpesvirus infections on present and future neurological disease burden, but also to motivate vaccine development, and to guide decision-making regarding screening programs and treatment recommendations for a population. Among the HHVs, HHV-3 (VZV), HHV-6 A&B, and HHV-7 exhibit a seroprevalence close to 100% [1]. For VZV does a proportion of seropositive cases stem from vaccination. We performed the present study to estimate the seroprevalence of HHV-1 (HSV-1), HHV-2 (HSV-2), HHV-4 (EBV) and HHV-5 (CMV) over several decades in healthy adults from northern Sweden.

Methods

With the aim to estimate time trends in seroprevalence of HHV-1 (HSV-1), HHV-2 (HSV-2), HHV-4 (EBV) and HHV-5 (CMV), serum from biobanks were analysed for presence of anti-HHV IgG with ELISA. By including participants of uniform age born between 1937 and 1988, seroprevalence in adults as a function of year of birth was monitored. Linear regression analyses were used to test for correlation.

Participants

Samples were retrieved from two separate biobanks; The Northern Sweden Maternity cohort and The Västerbotten Intervention Programme / Northern Sweden Health and Disease Study Cohort (VIP/NSHDS) cohort. The Northern Sweden Maternity cohort consists of serum samples collected in conjunction with screening for infectious agents among pregnant women in their first trimester. The biobank includes samples from four counties (Västernorrland, Jämtland, Västerbotten and Norrbotten) in northern Sweden from 1975 and onward. Approximately 2400 samples are added each year, and at the time of sample retrieval for this study the maternity biobank comprised of approximately 102,000 individuals [13]. The VIP/NSHDS cohort consists of samples from residents in Västerbotten County. People at the age of 40, 50 and 60 are since 1985 invited to contribute a plasma sample in connection with health surveys for risk factor screening performed by the healthcare region of Västerbotten, Sweden [14]. At the time of sample retrieval consisted the biobank of 110,663 individuals.

The samples that were used from the maternity cohort are dated from 1975 to 2018 and from VIP/NSHDS 1987 to 2017. The range of year-of birth thus spans between 1937 and 1988, allowing for representation from birth cohorts that have entered a geriatric stage in life, as well as representation from relatively young individuals. Previous studies have indicated that span to cover a significant shift in seroprevalence, at least for anti-HSV-1 IgG [15]. Both cohorts are administered by The Biobank Research Unit at Umeå University, and are by agreement with the healthcare region- made accessible for such research that serves the public health of the population. Integrity of the cohorts regarding IgG analyses have been proven earlier [16, 17]. For each sampling year, 12 serum samples from 30 year old women from the maternity cohort and 6 plasma samples each from 50 year old men and 50 year old women from the VIP/NSHDS cohort were retrieved in a randomized manner. From the first year of the maternity cohort (1975) were only four samples possible to retrieve. A total of 892 samples were included.

Serum and plasma analyses

Enzyme-Linked Immunosorbent Assay (ELISA) was used to detect anti-HSV-1 IgG, anti-HSV-2 IgG, anti-EBV IgG, and anti-CMV IgG.

ELISA assays deployed were, for anti-HSV-1 IgG and anti-HSV-2 IgG: Herpesselect®-assays (Focus Diagnostics), for anti-EBV IgG: VCA IgG (VIROTECH Diagnostics), and for anti-CMV IgG: an in-house ELISA based on total tissue culture antigen, as previously

described in [18]. Thresholds for a sample to be considered positive is for the HerpeSelect® (anti-HSV-1 IgG, anti-HSV-2 IgG): 0.9X index value, for anti-EBV IgG more than 9.0 VE and for anti-CMV IgG more than 5U. Samples in the greyzone ($n = 5-9$) were considered positive, the rationale being that this is biobank samples stored for a long time, and the chance is higher that some IgG is lost than the opposite have happened. The HerpeSelect®-assays are claimed by the manufacturer to exhibit a sensitivity for HSV-1 / HSV-2 of 91/96% and a specificity of 92/97% (HerpeSelect® 1 ELISA IgG REF EL0910G Rev. K product package insert). Independent assessments have reported 70/92% sensitivity and 92/57% specificity [19], and 99/97% sensitivity, 77/89% specificity [20] when compared to western blot. The anti-EBV IgG assay has, in an early assessment, been reported to generate 2.7% false positives and 7.9% false negatives [21]. The performance of the in-house method was assured through external quality programs managed by Equalis (www.equalis.se/en/), UK Neqas (www.ukneqas.org.uk) and Lab-Quality (www.labquality.com). In addition, internal controls consisting of previously run sera were included in every test.

Statistics

The seroprevalence was calculated by dividing the number of positive samples by all samples for each cohort. A linear regression analysis was used to test for a correlation between birth year and seroprevalence for the three groups separately: 30 year old women, 50 year old women and 50 year old men. A multiple linear regression analysis was used to differentiate between other factors and included overall prevalence in relation to birth year, age and gender.

$P < 0.05$ was regarded as statistically significant. The IBM SPSS 25 software for Mac was used for statistical calculations.

Results

Three groups were studied separately: 30 years old women, 50 years old women and 50 years old men. Table 1 shows the background characteristics and seropositivity for each group.

The relationship between birth year and seroprevalence was investigated for each of the three groups (Table 2). Statistically significant correlations were found between birth year and decreasing prevalence of anti-HSV-1

Table 1 Background characteristics

	Females age 30 y	Males age 50 y	Females age 50 y
<i>n</i>	520	186	186
Birth years	1945 – 1988	1937 – 1967	1937 – 1967
Sampling years	1975 – 2018	1987 – 2017	1987 – 2017
Anti-HSV-1 IgG positive <i>n</i> (%)	350 (67.3)	141 (75.8)	137 (73.7)
Anti-HSV-2 IgG positive <i>n</i> (%)	77 (14.8)	22 (11.8)	35 (18.8)
Anti-EBV IgG positive <i>n</i> (%)	507 (97.5)	180 (96.8)	185 (99.5)
Anti-CMV IgG positive <i>n</i> (%)	382 (73.5)	143 (76.9)	153 (82.3)

Abbreviations: *n* number, *y* years, HSV-1 Herpes simplex 1, IgG Immunoglobulin G, HSV-2 Herpes simplex 2, EBV Epstein-Barr virus, CMV Cytomegalovirus

Table 2 Herpes virus seroprevalence by birth year – linear regression analysis

		Female 30 y	Male 50 y	Female 50 y
Anti-HSV-1 IgG positive	Birth year beta	-0.004	-0.003	-0.007
	Birth year <i>p</i> -value	0.032	0.413	0.053
	Constant	7.603	6.399	14.380
Anti-HSV-2 IgG positive	Birth year beta	-0.003	-0.002	0.004
	Birth year <i>p</i> -value	0.028	0.480	0.234
	Constant	5.537	3.791	-7.289
Anti-EBV IgG positive	Birth year beta	-0.001	-0.002	-0.0003
	Birth year <i>p</i> -value	0.115	0.166	0.578
	Constant	2.672	4.903	1.651
Anti-CMV IgG positive	Birth year beta	0.002	-0.009	-0.004
	Birth year <i>p</i> -value	0.195	0.011	0.240
	Constant	-3.220	17.954	8.038

IgG ($p=0.032$) and anti-HSV-2 IgG ($p=0.028$) among 30-year-old women, and for anti-CMV IgG among 50-year-old men ($p=0.011$).

In a multiple linear regression analysis including overall seroprevalence in relation to birth year, age and female sex, birth year correlated negatively with the seroprevalence of anti-HSV-1 IgG ($p=0.004$) and anti-EBV IgG ($p=0.033$), while age correlated positively with anti-CMV IgG ($p=0.039$) (Table 3).

In the Female age 50 y group, there was a positive correlation between being anti-HSV-1 IgG positive and anti-CMV IgG positive (0.233, $p=0.001$) and a negative correlation between being anti-HSV-1 IgG positive and anti-HSV-2 IgG positive (-0.149, $p=0.042$). No other correlations were seen between antibody presence in any group.

Discussion

Seroprevalence trends of anti-HSV-1 IgG, anti-HSV-2 IgG, anti-EBV IgG and anti-CMV IgG were studied in a population of 30-year-old pregnant women and 50-year-old men and women from northern Sweden. Although the vast majority of seroconversions for the studied HHVs take place in childhood and early adolescence, some increase in seroprevalence, as a consequence of de novo infection of adults, is predicted as a population is ageing [15]. Correct prediction of seroconversion rates requires longitudinal samples, a research asset not commonly available. In a multivariate regression analysis, we could confirm a significant increase of anti-CMV IgG seropositivity as a function of age, likely due to seroconversion in adult years. This is in line with previous reports [15, 22, 23].

For groups combined, later birth year correlated with lower seroprevalence for anti-HSV-1 IgG and anti-EBV IgG. For groups studied separately, later birth year correlated with lower seroprevalence for anti-HSV-1 IgG and anti-HSV-2 IgG among 30-year-old women, and with lower seroprevalence for anti-CMV IgG among 50-year-old men. The observed decrease in anti-HSV-1 IgG seropositivity for 30-year-old women is dramatic, from around 88% among women born in 1945–1948 to 69% among women born 1985–1988. This is in line with figures reported by other studies from Sweden [15], Finland [24], England [25] and the United States [26].

The diagram depicting a moving average over 10 years (Fig. 1a) suggests that the main part of the drop in anti-HSV-1 IgG seropositivity occurred between birth years 1950 and 1960, a time when the studied population proportionally changed from a rural to a more urban lifestyle. Interestingly, after that shift, the trend does not seem to continue. Speculatively, in the 1970s, daycare service and pre-schools with large groups of children were implemented on a large scale, a milieu notorious for transmission of infectious agents. The moving average curve for anti-HSV-2 IgG suggests a peak in seroprevalence for birth year around 1960, followed by a significant decrease (Fig. 1b). Other cohorts exhibit a similar major difference between birth years born around 1960 and those born a few years later [27]. The decline for anti-HSV-2 IgG follows a trend shared by several sexually transmitted diseases (STDs) [28]. This effect has been attributed changes in sexual behavior, not to the least necessitated by the HIV pandemic [28]. The overall declining trend is in line with results from other studies [24, 26, 29]. The introduction of acyclovir as a suppressive treatment option for HSV infections in the mid-1980s is presumed to have reduced the rate of reactivation and changed the pattern of spread [30], as have the fact that HSV-1, instead of HSV-2, by now is the primary cause of genital HSV infections at least in some populations [31]. Public health implications of the decrease in seroprevalence include that a larger proportion of women enter pregnancy seronegative, and thus run a risk of contracting the infection and transferring it to the fetus.

EBV is a very common infection in our population, where over 96% were seropositive in this study. The declining trend in relation to birth year is statistically significant, but so small that implications for populational health are questionable (Fig. 1c). In a study from Finland, no declining trend in seroprevalence of EBV could be detected [24].

Comparing the moving average curves for anti-CMV IgG (Fig. 1d) and anti-HSV-1 IgG (Fig. 1a) gives an illustration of the similarities in spread between the two agents, which is further supported by the correlation between having these two agents among 50-year-old females. Interestingly, both anti-HSV-1 IgG and anti-CMV IgG seem to have their minimum around birth year 1970–1980. Further studies should investigate if the

Table 3 Herpes virus seroprevalence by birth year, age and sex – multiple linear regression analysis

	Anti-HSV-1 IgG positive	Anti-HSV-2 IgG positive	Anti-EBV IgG positive	Anti-CMV IgG positive
Birth year beta (p -value)	-0.004 (0.004)	-0.002 (0.103)	-0.001 (0.033)	-0.0002 (0.875)
Age beta (p -value)	0.0003 (0.898)	0.001 (0.679)	0.0003 (0.695)	0.004 (0.039)
Female sex beta (p -value)	-0.022 (0.648)	0.070 (0.059)	0.027 (0.080)	0.054 (0.225)

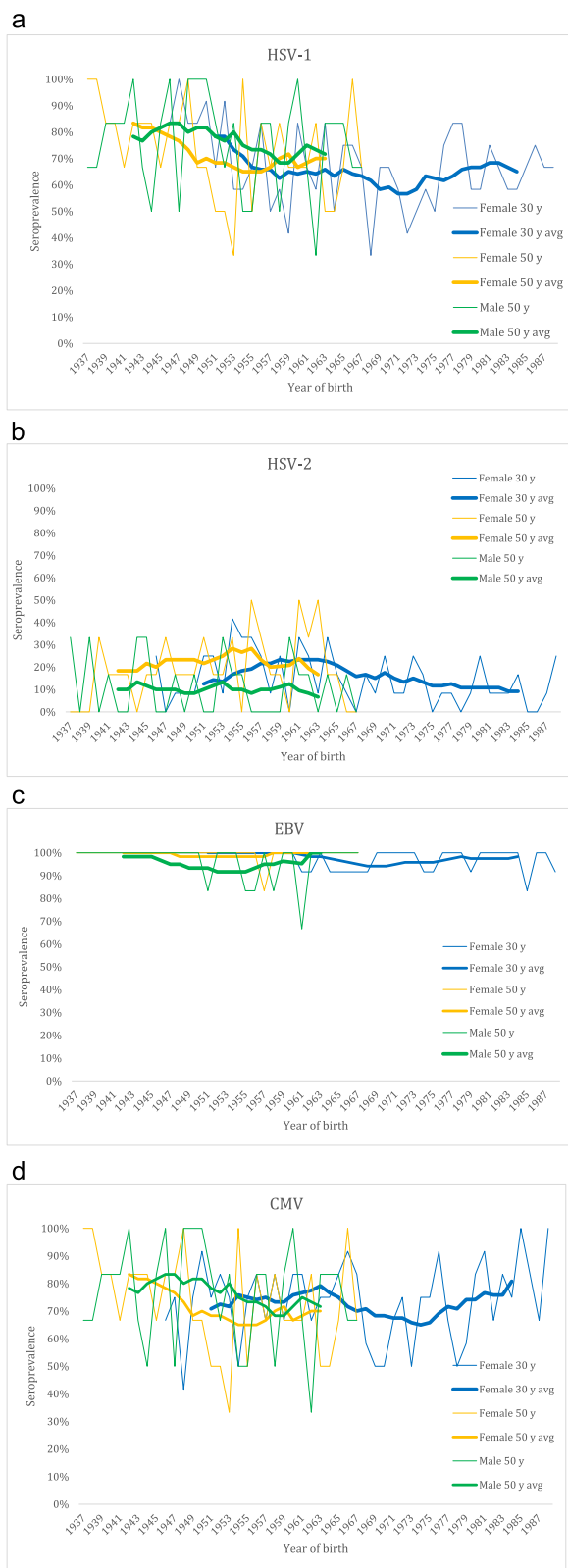


Fig. 1 Seroprevalence of anti-HHV IgG among subjects born between 1937 and 1988. **a** represents anti-HSV-1 IgG, **b** represents anti-HSV-2 IgG, **c** represents anti-EBV IgG, **d** represents anti-CMV IgG. Subjects are divided into females aged 30 years (blue), females aged 50 years (yellow), and males aged 50 years (green). Thick lines represent the moving average over 10 years

trend of increasing seroprevalence in later birth years is statistically significant.

Strengths and weaknesses of the study

The limited information on each sample does not permit for detailed suggestions of underlying causes and correlations of the reported trends such as comorbidities, rates of reactivations, disease outcomes and other clinical data.. Also, the small number of samples from each year make yearly prevalence estimates as well as trends for sub-timespans uncertain. Therefore, to equalize random year-to-year variation we also present the 10-year moving average. A weakness of the study is that data for females originates from two separate cohorts, while data for males originates from one single cohort. This must be considered when comparisons are made. In addition, the 30-year old women are all pregnant, a fact that probably skews the selection, Strengths of the study include the employment of population-based biobanks [13], that allows for estimation of 50-year long time trends in age- and gender specific anti-HHV IgG prevalence among persons born between 1937 and 1988. The population-based biobanks represent an unselected sample of the underlying general population, with minimal risk of further selection bias apart from the above mentioned. This makes it likely that the observed trends do reflect the actual population prevalence of these viruses.

Conclusion

The seroprevalence of anti-HSV-1 IgG, anti-HSV-2 IgG and anti-EBV IgG decreases in later birth cohorts. This indicates an overall trend of declining risk of getting infected as a child and adolescent. Healthcare implications include a higher proportion of the population being susceptible to primary infection at adult age.

Abbreviations

- HHV Human herpesvirus
- IgG Immunoglobulin G
- HSV *Herpes simplex virus*
- EBV Epstein-Barr virus
- CMV Cytomegalovirus
- ELISA Enzyme Linked Immunosorbent Assay
- VZV *Varicella zoster virus*
- STD Sexually transmitted disease

Acknowledgements

The authors wish to acknowledge Per Juto, MD, PhD, for his work to establish the CMV IgG method used in this and many other studies as well as in clinical diagnostics. The authors wish to acknowledge Göran Wadell, MD, PhD for his work to oversee the maternity cohort for several decades and to pioneer the work to make the cohort accessible for research.

Authors' contributions

JO, SN, and EH performed the serological analyses. AJ and GH assembled the study cohorts. BW and HL performed data analysis and interpretation. HL and FE conceptualized the study. SN and JO drafted the manuscript. All authors have contributed to and approved the final version of the manuscript.

Funding

Open access funding provided by Umeå University. This study was supported financially by grants from Region Västerbotten, Wallenberg Centre for Molecular Medicine (WCMM) at Umeå University, the Swedish Dementia Association, the Swedish Alzheimer Fund and the Umeå University Foundation for Medical Research. The funding bodies had no role in the design of the study, in the collection, analysis, and interpretation of data, nor in writing the manuscript.

Availability of data and materials

The datasets used and analyzed during the current study are available from the corresponding author on reasonable request.

Declarations

Ethics approval and consent to participate

The study is approved by a regional ethical board (Dnr 2017/384–31). The research subjects have provided informed consent for the serum- and plasma samples to be used for research purposes at the time of the sampling.

Consent for publication

All participants who have provided consent for their serum/plasma to be used in research have also given consent for publication of research based on stored serum/plasma.

Competing interests

The authors declare no competing interests.

Received: 18 December 2023 Accepted: 19 February 2024

Published online: 02 March 2024

References

- Arvin, A., et al., Human Herpesviruses Biology, Therapy, and Immunoprophylaxis. 2007, Cambridge University Press, Cambridge. p. 1 online resource (1408 p).
- Khan G, Hashim MJ. Global burden of deaths from Epstein-Barr virus attributable malignancies 1990–2010. *Infect Agent Cancer*. 2014;9(1):38.
- Mesri EA, Cesarman E, Boshoff C. Kaposi's sarcoma and its associated herpesvirus. *Nat Rev Cancer*. 2010;10(10):707–19.
- Olsson T, Barcellos LF, Alfredsson L. Interactions between genetic, lifestyle and environmental risk factors for multiple sclerosis. *Nat Rev Neurol*. 2017;13(1):25–36.
- Lovheim H, et al. Reactivated herpes simplex infection increases the risk of Alzheimer's disease. *Alzheimers Dement*. 2015;11(6):593–9.
- Lovheim H, et al. Herpes Simplex Virus, APOEε4, and Cognitive Decline in Old Age: Results from the Betula Cohort Study. *J Alzheimers Dis*. 2019;67(1):211–20.
- Lopatko Lindman K, et al. A genetic signature including apolipoprotein Eε4 potentiates the risk of herpes simplex-associated Alzheimer's disease. *Alzheimer's & Dementia: Translational Research & Clinical Interventions*. 2019;5:697–704.
- Lovheim H, et al. Interaction between Cytomegalovirus and Herpes Simplex Virus Type 1 Associated with the Risk of Alzheimer's Disease Development. *J Alzheimers Dis*. 2018;61(3):939–45.
- Bistrom M, et al. Epstein-Barr virus infection after adolescence and human herpesvirus 6A as risk factors for multiple sclerosis. *Eur J Neurol*. 2021;28(2):579–86.
- Asha K, Sharma-Walia N. Targeting Host Cellular Factors as a Strategy of Therapeutic Intervention for Herpesvirus Infections. *Front Cell Infect Microbiol*. 2021;11: 603309.
- Busnadiago I, et al. Critically ill COVID-19 patients with neutralizing autoantibodies against type I interferons have increased risk of herpesvirus disease. *PLoS Biol*. 2022;20(7): e3001709.
- Henze L, et al. Management of herpesvirus reactivations in patients with solid tumours and hematologic malignancies: update of the Guidelines of the Infectious Diseases Working Party (AGIHO) of the German Society for Hematology and Medical Oncology (DGHO) on herpes simplex virus type 1, herpes simplex virus type 2, and varicella zoster virus. *Ann Hematol*. 2022;101(3):491–511.
- Pukkala E, et al. Nordic biological specimen banks as basis for studies of cancer causes and control—more than 2 million sample donors, 25 million person years and 100,000 prospective cancers. *Acta Oncol*. 2007;46(3):286–307.
- Hallmans G, et al. Cardiovascular disease and diabetes in the Northern Sweden Health and Disease Study Cohort – evaluation of risk factors and their interactions. *Scand J Public Health Suppl*. 2003;61:18–24.
- Olsson J, et al. Herpes virus seroepidemiology in the adult Swedish population. *Immun Ageing*. 2017;14:10.
- Esberg A, et al. 43-Year Temporal Trends in Immune Response to Oral Bacteria in a Swedish Population. *Pathogens*. 2020;9(7):544.
- Lovheim H, et al. Herpes simplex infection and the risk of Alzheimer's disease: A nested case-control study. *Alzheimers Dement*. 2015;11(6):587–92.
- Sjostrom S, et al. Human immunoglobulin G levels of viruses and associated glioma risk. *Cancer Causes Control*. 2011;22(9):1259–66.
- Agyemang E, et al. Performance of Commercial Enzyme-Linked Immunoassays for Diagnosis of Herpes Simplex Virus-1 and Herpes Simplex Virus-2 Infection in a Clinical Setting. *Sex Transm Dis*. 2017;44(12):763–7.
- Ashley-Morrow R, et al. Performance of focus ELISA tests for herpes simplex virus type 1 (HSV-1) and HSV-2 antibodies among women in ten diverse geographical locations. *Clin Microbiol Infect*. 2004;10(6):530–6.
- Gartner BC, et al. Evaluation of four commercially available Epstein-Barr virus enzyme immunoassays with an immunofluorescence assay as the reference method. *Clin Diagn Lab Immunol*. 2003;10(1):78–82.
- Hecker M, et al. Continuous cytomegalovirus seroconversion in a large group of healthy blood donors. *Vox Sang*. 2004;86(1):41–4.
- Lachmann R, et al. Cytomegalovirus (CMV) seroprevalence in the adult population of Germany. *PLoS ONE*. 2018;13(7): e0200267.
- Puhakka L, et al. Decrease in seroprevalence for herpesviruses among pregnant women in Finland: cross-sectional study of three time points 1992, 2002 and 2012. *Infect Dis (Lond)*. 2016;48(5):406–10.
- Forbes H, et al. Risk factors for herpes simplex virus type-1 infection and reactivation: Cross-sectional studies among EPIC-Norfolk participants. *PLoS ONE*. 2019;14(5): e0215553.
- Xu F, et al. Trends in herpes simplex virus type 1 and type 2 seroprevalence in the United States. *JAMA*. 2006;296(8):964–73.
- Lowhagen GB, et al. Epidemiology of genital herpes infections in Sweden. *Acta Derm Venereol*. 1990;70(4):330–4.
- Edgardh K. Adolescent sexual health in Sweden. *Sex Transm Infect*. 2002;78(5):352–6.
- Berntsson M, et al. Decreasing prevalence of herpes simplex virus-2 antibodies in selected groups of women in Sweden. *Acta Derm Venereol*. 2009;89(6):623–6.
- Lopatko Lindman K, et al. Long-term time trends in reactivated herpes simplex infections and treatment in Sweden. *BMC Infect Dis*. 2022;22(1):547.
- Johnston C, et al. Viral Shedding 1 Year Following First-Episode Genital HSV-1 Infection. *JAMA*. 2022;328(17):1730–9.

Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.