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Biological Correlates of Sexual Transmission of HIV

Practical Consequences and Potential Targets for Public Health

Abstract

The probability of sexual transmission of HIV depends on the infectiousness of the index case and the susceptibility of the sexual contact. The risk of HIV transmission is heterogeneous and may be greatest during the initial sexual contacts in a steady partnership. Several factors, including systemic and mucosal acquired protective immune-response might be responsible for the apparent decrease of persex-act risk of transmission in a given partnership over time. Biological studies can be used to better understand the complex information obtained by epidemiological surveys. The infectiousness of HIV depends on the inoculum, and virologic factors. The genital tract viral load of the index case is likely the most important determinant of transmission. At the population level, interventions that reduce the genital shedding of HIV by reducing systemic blood viral load and/or local inflammatory processes are likely to have a beneficial impact on HIV incidence. Antiretroviral drugs are likely to reduce sexual transmission of HIV. However, these drugs may not all prove equally. Compartmentalized HIV replication in the male and female genital tract have been observed. Treatment with antiretroviral drugs that poorly penetrate the genital tract harbour the risk of local production and spread of resistant viruses. In addition, increased risk taking behaviour could offset the benefits of reduced probability of transmission at the population level. Biological data about HIV transmission must be used to inform public health policies and optimize HIV prevention strategies.

Keywords

HIV · Sexual Transmission · HIV Prevention Strategies · Susceptibility · Infectiousness

Worldwide, the predominant mode of HIV transmission is clearly heterosexual contact. However, the biological correlates for HIV transmission are less clear because of the heterogeneity of sexual transmission. This heterogeneity is best illustrated by individual case reports where transmission occurs in some partners after limited sexual contact, while other partners of the same individual remain uninfected [1]. The heterogeneity of sexual transmission makes the risk of a single contact much less predictable than for other modes of transmission.

The probability of sexual transmission of HIV is a function of the infectiousness of the index case, the mode of the sexual contact, and the susceptibility of the person exposed to the virus. We have recently reviewed factors contributing to the infectiousness of the infected individual [2] and a review of the susceptibility factors has been published by Buchacz [3]. Here we summarize recent findings about susceptibility and infectiousness, with a focus on infectiousness factors that can be influenced by public health prevention strategies.

Estimating the transmission risk: Where biology meets epidemiology

Estimates for the risk of sexual transmission of HIV have been generated from epidemiological surveys. In most instances, partner studies have been used to calculate the per-partnership transmission risk or the per-sexual-contact risk of transmission. The calculation of the per-sexact risk is based on the assumption that the risk remains stable over a long period (stage) of the disease [4, 5]. However, studies that report a high transmission risk after a single exposure to an HIV-infected prostitute question the validity of this assumption [6, 7]. The data from partner studies is best predicted by mathematical models that account for a large heterogeneity in HIV infectiousness [8]. Increasing evidence is evolving that indicates a more complex temporal performance for both HIV infectiousness and susceptibility to HIV. Biological studies have therefore been used to complement the information obtained from epidemiological studies; the two combined ap-

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Biologische Korrelate der sexuellen Transmission von HIV

Zusammenfassung

Die Wahrscheinlichkeit einer Übertragung von HIV auf sexuellem Wege hängt ab von der Infektiosität der Index-Person und der Suszeptibilität der Person, mit der die Index-Person sexuellen Kontakt hat. Das Risiko einer HIV-Übertragung ist sehr heterogen und dürfte während der ersten sexuellen Kontakte in einer sich entwickelnden Beziehung am größten sein. Verschiedene Faktoren, u.a. auch erworbene schützende systemische und Schleimhaut-assoziierte Immunantworten, dürften für den offensichtlichen Rückgang der Übertragungswahrscheinlichkeit pro Kontakt in einer gegebenen Partnerschaft im Zeitverlauf verantwortlich sein. Biologische Untersuchungen können weiteres Licht auf die komplexen Informationen werfen, die aus epidemiologischen Erhebungen resultieren. Die Infektiosität von HIV hängt von der inokulierten Menge und von virologischen Faktoren ab. Die Viruslast im Genitaltrakt der Index-Person ist vermutlich die wichtigste Determinante für eine Übertragung. In einer gesamten Population führen Interventionen, die die Viruskonzentration im Genitaltrakt senken, sei es durch Verminderung der Viruslast im Blut oder durch Behandlung entzündlicher Prozesse im Genitaltrakt, zu einer Senkung der HIV-Inzidenz. Antiretrovirale Substanzen vermindern wahrscheinlich die sexuelle Übertragungsrate von HIV. Allerdings dürfte dies nicht für alle Substanzen im selben Ausmaß gelten. Sowohl der männliche als auch der weibliche Genitaltrakt bilden ein Kompartiment, in welchem sich HIV unabhängig vom lymphatischen System vermehrt. Die Behandlung mit antiretroviralen Medikamenten, die schlecht in den Genitaltrakt penetrieren, birgt die Gefahr einer lokalen Produktion und Ausbreitung resistenter Viren. Zusätzlich könnte eine Zunahme von Risikoverhalten die Verminderung der Übertragungswahrscheinlichkeit durch Reduktion der Viruslast wieder zunichte machen. Die biologischen Erkenntnisse über die HIV-Übertragung müssen als Grundlage für Public-Health-Strategien und zur Optimierung der Präventionsstrategien genutzt werden.

Schlüsselwörter

HIV · Sexuelle Übertragung · Infektiosität · Empfänglichkeit · Präventionsstrategien

Originalien und Übersichtsarbeiten

proaches will undoubtedly improve our understanding of HIV transmission and facilitate interventions to reduce it.

Heterogeneity of the HIV transmission risk

Three major factors that determine HIV transmission risk contribute to the heterogeneity observed: differences in the susceptibility to HIV, variable infectiousness of infected individuals, and a wide spectrum of sexual behavior in a given population. Heterogeneity of sexual behavior has led to the concept of core groups in which rates of partner change and sexual activity are substantially higher than in the general population [9]. Behavioral issues which affect transmission have recently been reviewed [10].

Susceptibility to HIV

The susceptibility to HIV-infection depends on host immunological and genetic factors. A homozygous CCR5 (Chemokine Receptor 5) coreceptor deletion mutant has been shown to protect against HIV infection [11]. Genetic factors, such as CCR5 and SDF (Stromal Cell-Derived Factor) variants as well as certain HLA types, are also strongly associated with non-progression in HIVinfected individuals [12]. It is conceivable that these factors could also influence the susceptibility to HIV. At least one third of individuals exposed to HIV develop a cell-mediated immune response without overt seroconversion [13, 14]. Some animal models show a protective immune response after sub-infectious exposure to HIV; by analogy, one might expect that some of the HIV-sensitized exposed partners also have a reduced susceptibility to HIV [15]. Some exposed individuals also develop an HIV-specific mucosal and systemic IgA response and these antibodies neutralize primary HIV strains [16]. These data are further supported by several reports of decreased susceptibility to HIV in multiply exposed commercial sex workers (CSW) with continued high-risk behavior [17, 18]. Existence of a (partly) protective immune response after sexual exposure to HIV might explain the decrease of HIV transmission risk after a few sexual contacts in a given heterosexual partnership, as reported in the study that is discussed below.

Local genital factors are also likely to influence the susceptibility to HIV. Upregulation of the CCR5 co-receptor and increased activation of T-lymphocytes present in the normal cervix have been documented [19]. This finding supports the vulnerability of the female genital tract for HIV and suggest a mechanism for the observed predominance of sexual transmission of HIV variants with CCR5 co-receptor usage [20].

Sexually transmitted diseases (STDs) have long been shown to influence not only infectiousness (see below) but also susceptibility to infection. Local inflammatory reactions in response to STDs are likely to attract CD4+ lymphocytes and release of cytokines that enhance transmission of HIV. Changes in vaginal flora appear to affect susceptibility to HIV as well. Bacterial vaginosis (BV) is associated with a high concentration of vaginal anaerobes, reduced lactobacilli, and an increase in vaginal pH [21]. Sturm-Ramirez and colleagues found in vitro evidence for increased levels of pro-inflammatory cytokines (TNF- α and IL-1) in patients with BV, which could in part explain the association of BV with increased susceptibility to HIV [22].

"The susceptibility to HIV-infection depends on host immunological and genetic factors."

Other local factors also potentially modulate HIV susceptibility: hormones that alter the vaginal epithelium have been investigated for their influence on vaginal HIV shedding. Recently, estrogens were shown to inhibit HIV transmission in the macaque model; this effect was associated by thickening of the vaginal epithelial wall [23].

The effect of circumcision on HIV transmission is highly debated. The estimates for the odds against transmission in circumcised men are in the range of 1.5. Several mechanisms have been proposed to explain this effect, including the reduced surface area for mucosal contact [24]. However, most studies investigating the role of circumcision on HIV transmission are confounded by religion and are therefore difficult to interpret. In a mathematical analysis of per-sex-act male-tofemale transmission probabilities, Duerr et al. found similar per-act transmission risk in otherwise comparable US (mostly circumcised) and Italian (mostly uncircumcised) HIV-discordant couples [25]. This argues against a marked effect of circumcision status on HIV transmission. The role of circumcision on susceptibility to HIV needs to be further examined before recommendations for large-scale circumcision interventions are issued as recently proposed [26].

In summary, a combination of local genital factors, immune response mechanism, and genetic predisposition are all influencing the risk of transmission by affecting susceptibility to HIV. However, based on epidemiological partner studies discussed below, infectiousness of the infected partner is likely to influence transmission probability even more than susceptibility.

Infectiousness of the HIV infected partner

Viral clade, quasispecies and transmission

Based on differences in the replication kinetics in dendritic cells between HIV subtype E and B, Essex and coworkers suggested that the subtype might play a role in transmission efficacy [27]. However, these results have not been confirmed by others [28] and it has been very difficult to document any effect of subtype differences on transmission risk. Miller and coworkers could not find a difference in transmissibility based on HIV macrophage tropism but found in vivo replication capacity to be predictive for the outcome of intravaginal inoculation in the macaque model [29]. Another animal study found that some HIV quasispecies were better adapted for vaginal transmission than others, but the significance of this finding for the case of HIV transmission is not known [30]. Ping and coworkers have argued that clade C expresses biological characteristics which favor sexual transmission [31]. A careful review of the debate has been published by Hu et al. [32]. In summary, systemic and local host factors remain the most important determinants for HIV infectiousness.

Systemic factors influencing HIV infectiousness

Epidemiological studies (mostly partner studies) have previously demonstrated

an increased HIV transmission from patients with low CD4 count, advanced stages of disease, p24 antigenemia and the presence of STDs (reviewed in [2]). Subsequently, time variation of HIV infectiousness was proposed by Jacquez et al., based on mathematical models of monogamous partner studies [5]. The authors argue that HIV infectiousness is increased during primary infection (which is consistent with an increase of blood viral load during this phase). However, a key assumption of this model, a stable per-contact transmission rate, was challenged by Downs et al. [33] and further discussed by Shiboski and Padian [34]. Several factors that might explain the observed decline in transmission risk after primary infection need to be considered. Firstly, the large heterogeneity of transmission itself may explain the finding, also termed "frailty selection" [35]. Secondly, the assumption of constant transmission rates during a given partnership might influence the results. Thirdly, key variables on infectiousness and susceptibility that might influence the curve are usually missing in these partner studies. Shiboski and Padian conclude that the nature of variation in infectiousness cannot be confirmed, but that the findings indicate a declining tendency of transmission risk in monogamous heterosexual couples with increasing length of exposure [34]. It is just as likely that the declining risk of transmission in a given partnership results from an acquired protective immune response in the non-infected partner. This would also explain the substantially higher risk estimates (3-8%) for single sexual exposures to commercial sex workers [6,7].

In general, transmission of a pathogen is concentration-dependent. Vertical transmission of HIV and occupational transmission through needle-stick injuries increase in situations with high viral load in the index person [36, 37]. Several partner studies have demonstrated the association of blood viral load with sexual transmission of HIV [38, 39, 40]. In these studies, neither CD4 count or the number of sexual contacts affected the risk of transmission. In a recently published sub-analysis of discordant couples included in a large STD-intervention study in Rakai, viral load was the most powerful predictor of HIV transmission [41]. Of note, none of the 53 HIV-infected individuals with a viral load below 1500

copies/ml transmitted the virus to their partner despite the negligible use of condoms. In all of the earlier epidemiological studies on transmission risk mentioned above, measurement of HIV-RNA concentration in blood was not available. It is likely that these studies were confounded by blood viral load. Low CD4 count, advanced stage of disease, p24 antigenemia and primary HIV infection are all associated with higher levels of HIV-RNA in blood. While an independent role of all those factors on infectiousness cannot be excluded, it is apparent that blood viral load is by far the most important cofactor.

HIV load in the genital tract as a measure of infectiousness

The strong predictive power of blood viral load for HIV transmission risk is biologically plausible. In general, transmission of a pathogen is concentration-dependent. Likewise, vertical transmission of HIV and occupational transmission through needle-stick injuries increase in situations with high viral load in the index person [42].We and others have therefore studied the concentration of HIV in genital secretions and postulated that the measurement of HIV in semen can be used to estimate an individual's infectiousness (reviewed in [2, 43]). Similar results have accumulated for female genital secretions, but the variability of the assays for the detection of HIV-RNA or DNA in female secretions is considerably higher than for semen [44, 45, 46, 47, 48]. The menstrual cycle further increases the biological variability of the virus shedding in the female genital tract [49].

The combined information on HIV viral load and tropism in semen and the CCR5 receptor density on target cells in the female genital tract has been used to predict the transmission probability of HIV in the genital tract [50]. In a mathematical model we have developed, HIV transmission was unlikely (1 in 10'000 episodes of intercourse) when HIV-RNA in semen was low (i.e. <5000 copies/ml) but rose to 3 per 100 episodes of intercourse if HIV-RNA in semen was high (10⁶ copies/ml). Such high seminal RNA values have been found in semen from HIV-positive men with STDs in Malawi. Thus, seminal viral load can serve as a surrogate marker to determine infectiousness.

Local genital factors influencing HIV infectiousness

Semen studies have revealed a weak but significant correlation (r = 0.55-0.60) of blood viral load and seminal HIV-RNA concentrations. The correlation was considerably higher (r = 0.80) in a recent study of patients with primary infection when a more sensitive HIV-RNA detection assay was used [51]. Thus, in agreement with the epidemiological studies, the biological studies indicate that a major determinant of the viral inoculum size in semen is blood viral load; approximately 60% of the seminal viral load can be predicted by blood viral load alone.

"Seminal viral load can serve as a surrogate marker to determine infectiousness."

Factors that explain the discordance between blood and semen viral load must be investigated. HIV-RNA concentration in semen is approximately one log lower than the respective level in blood. However, some patients have significantly higher levels of HIV in semen than in blood and have therefore been termed as hypersecretors [52, 53]. In one study, hypersecretor status has been associated with positive cell culture from the cellular semen compartment [54], supporting the potential effect of hypersecretor status on sexual transmission risk. Hypersecretor status has been associated with asymptomatic urethritis [55] and

was an independent risk factor for the shedding of HIV-RNA in semen.

Symptomatic STDs have long been shown to increase the risk of sexual transmission of HIV [56, 57]. Similarly to the studies of blood viral load described above, biological studies have been performed to demonstrate the influence of STDs on the viral load in genital secretions. Cohen and colleagues studied STD patients in Malawi and found an almost ten-fold increased level of HIV-RNA secretion in semen in patients with gonorrhea as compared to control patients without urethritis [58]. This increase in genital viral load was not associated with an increased HIV-RNA concentration in blood. In the same study, genital ulcer disease was also associated with an increased HIV shedding in semen, indicating the existence of an indirect mechanism by which genital inflammatory diseases stimulate the replication kinetics of HIV in semen [59]. Further analysis of the viral quasispecies in semen and blood samples from this study by heteroduplex mobility assay revealed a discordance of the two compartments in 40% of these individuals [31]. A number of other studies have documented similar associations between genital inflammatory diseases and HIV detection in both semen and cervical secretions [22, 60, 61, 62]. In a recently published study conducted in Nairobi, treatment of women with cervicitis (N.gonorrhea, C.trachomatis or unspecific cervicitis) was associated with a six-fold reduction in the shedding of HIV-1 RNA in cervical secretions [63]. Taken together, the biological

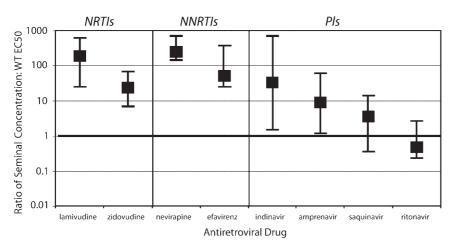


Figure 1 A Ratio of Seminal Plasma Concentrations to HIV-1 Wild-Type EC50. Ratios were calculated based on median, minimum, and maximum reported seminal drug concentrations. EC₅₀ values were obtained from Molla et al [67]

findings support the significance of STDs for HIV transmission and identify genital inflammatory diseases as another major determinant of genital shedding of HIV and HIV infectiousness.

Reducing HIV infectiousness as a preventive strategy

Efforts to reduce the spread of HIV can target sexual behavior, or the susceptibility of uninfected individuals, or the infectiousness of the HIV-infected population. Reducing HIV susceptibility by the induction of a protective immune response is the central aim of HIV vaccine development. Until a protective vaccine is developed, other strategies– such as the application of vaginal microbicides or syndromic management of STDs in high-incidence areas–are currently under development to help protect individuals from being infected.

HIV prevention efforts have only recently shifted to the infected person, who can now benefit from early detection of infection. Index cases can be encouraged to engage only in safe sex behaviors, with 100% condom usage. In addition, efforts to reduce HIV viral burden in the genital tract can be undertaken

Treatment of STDs and HIV infectiousness

STDs increase the HIV load in the genital tract. The effects of treatment of STDs on genital HIV shedding has been welldocumented in men and women [58, 60, 63]. In addition, the benefits of STD treatment on the HIV epidemic have been translated to some, but not all populations. In a community-randomized trial in Mwanza, Tanzania, improved management of STDs resulted in a 40 reduction of HIV incidence [64]. However, in a randomized study in the Rakai district of rural Uganda, mass STD-treatment did not result in a significant reduction of HIV incidence [65]. In that study, absence of a detectable effect on HIV incidence was probably a result of the mature stage of the HIV epidemic in rural Uganda [66]. In such mature, generalized, epidemic settings, a saturating effect is observed in core groups with highrisk behavior; therefore HIV incidence occurs at a lower level in the low-risk population and is less likely to be driven by STDs. Mass treatment of STDs cannot

target individuals with the highest risk of transmitting HIV. On an individual level, however, treatment of STDs is unquestionably an important intervention to reduce HIV infectiousness.

Reduction of viral load by HIV treatment

HIV-RNA concentration in maternal blood is associated with transmission risk [67, 68, 69, 70] and antiretroviral treatment of mothers prepartum results in reduction in maternal viral loads and vertical transmission rates [71]. Both systemic and local genital effects of shortcourse AZT treatment have been shown to be independently associated with reduced vertical transmission rates [72] Reduction of systemic and genital viral load is also likely to reduce the risk of sexual transmission.

A number of studies have demonstrated that highly active antiretroviral therapy (HAART) suppresses HIV replication, not only in the blood/lymphoid compartment, but also in the male and female genital tract [51, 52, 54, 73, 74, 75, 76, 77, 78, 79, 80, 81]. While some of the earlier studies were conducted among patients on suboptimal antiviral therapy, later studies have included patients with effective HAART. In our series of 114 patients with suppressed blood viral load below 400 copies/ml, only two patients had a seminal viral load slightly above the detection limit of the assay [82]. Although measurement of cell-free virus (HIV-RNA) in semen is usually suppressed in HAART-treated patients, a significant number of treated individuals harbor HIV provirus (HIV-DNA) in CD4 positive cells in their semen. In our recent study, the detection rate of HIV-DNA in semen was 17% in treated individuals compared to 38% in a drug-na control population. Zhang and colleagues have demonstrated that infectious virus can still be recovered in vitro from seminal lymphocytes after more than six months of suppressive HAART [76]. However, in a recent study on quadruple combination therapy in primary HIV infection, only 2 out of 22 patients had more than 50 HIV-DNA copies-per-ejaculate after one year of HAART [51].

In an analysis of Swiss patients with primary infection, Yerly and colleges demonstrated a significant reduction in the transmission of drug-resistant viruses after 1997 [83]. This reduction was paralleled by a gradual improvement of HAART effectiveness in the Swiss HIV Cohort Study. Currently, more than 60% of all treated patients in the Swiss HIV cohort study have HIV-RNA levels below the detection limit of 50 copies/ml in blood.

It is not known how low the blood or genital viral load must be to completely suppress the HIV-transmission risk (threshold level). In Quinn's study (cited above), a threshold level was defined at 1500 copies of HIV-RNA per ml of blood [41]. A similar observation was made for the case of vertical transmission [68]. However, any significant reduction in blood viral load is likely to result in reducing (but not eliminating) the risk of transmission.

Compartmentalization and antiretroviral drug concentrations in the male genital tract

Not all drug combinations for HAART appear to be equally effective in the genital and systemic compartment. Compartmentalization of HIV in semen has been clearly shown by the detection of different quasispecies in semen and blood in treated and untreated individuals [31, 84, 85, 86]. Recent findings of a distinct mutation pattern in virus isolates from blood and cervical samples indicate a similar compartmentalization in the case of the female genital tract [87,88]. By mathematical modeling, Kepler and Perelson have demonstrated that limited drug penetration into a small compartment is an important risk factor for the development of resistant HIV during treatment [89]. Thus, differential penetration of antiretroviral drugs into the genital tract is an important consideration if these drugs are intended to reduce the likelihood of sexual transmission of HIV-1 [90].

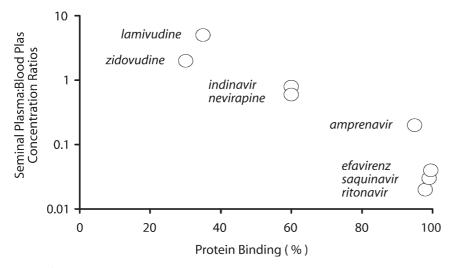
Theoretical factors that may influence antiretroviral drug distribution into the male genital tract, such as protein binding, drug transporters, and drug ionization have been reviewed in detail elsewhere [91]. Seminal concentration data are available for all three classes of antiretroviral agents currently marketed and semen to blood ratio of antiviral drug concentrations (S: B ratio) vary from undetectable to greater than one. In Figure 1, median semen concentration are shown in relation to the wild-type, protein corrected IC₅₀ of the drugs [92]. Of the nucleoside analogues, seminal concentration data are available for zidovudine (AZT), lamivudine (3TC), and stavudine (d4 T) [93, 94, 95, 96, 97]. All three appear to achieve seminal plasma concentrations greater than blood plasma [98, 99, 100]. However, data on intracellular drug phosphorylation in seminal lymphocytes is lacking.

Nonnucleoside reverse transcriptase inhibitor semen concentration data have been collected for efavirenz and nevirapine. Nevirapine concentration in semen reaches 60-100% of blood plasma levels [95], while seminal efavirenz exposure was 2.5-10% of that in blood [101]. The protease inhibitors generally achieve the lowest concentrations in the male genital tract, although S:B ratios vary for each compound (nelfinavir<saquinavir=ritonavir<amprenavir<indinavir). Detection of nelfinavir in semen was unsuccessful in one study [96] and both saquinavir and ritonavir reached less than 5% of blood plasma levels in semen during coadminstration of the two [102]. Amprenavir penetration into the seminal compartment was better than for saquinavir (S:B ratio: 0.2, 95% CI 0.05-3.0) [103]. Two groups have demonstrated high levels of indinavir in semen (median S:B ratio 0.4 and 1.2) [104] and further improvement of indinavir penetration by co-administration with low-dose ritonavir [105, 106].

Blood plasma protein binding, pglycoprotein affinity, and ionizing pH all influence drug penetration into and activity within the genital tract but protein binding appears to be a major determinant of drug concentrations in semen (Figure 2). Additionally, compounds with lower p-glycoprotein affinity (amprenavir and efavirenz) appear to achieve greater genital tract penetration than compounds with greater p-glycoprotein affinity (ritonavir and saquinavir) [107]. To date, no investigation has fully evaluated and compared combination antiretroviral therapy in the context of genital tract pharmacology and virology to determine the most potent and durable regimen for suppressing viral replication in this compartment.

Sexual behavior changes in response to HAART

The increased public awareness of the effects of HAART are likely to change





concerns about HIV transmission in the population at risk. The first study examining changes in perceived transmission risk associated with HAART was presented by Kravcik et al. in 1998 [108]. The authors described that 20% of HIV-infected individuals attending an HIV clinic thought that the risk of transmission was reduced during HAART and 19% believed that the need for safer sex practices was reduced under HAART. Increased-risk sexual behavior was also documented in a survey of serodiscordant couples [109] from California.

"The effects of HAART are likely to change concerns about HIV transmission in the population at risk."

One third of uninfected partners mentioned that they had already taken a chance with unprotected sex because of improved HIV treatment options and 40% responded that HAART had changed their transmission concerns. HIV-positive gay men studied in London in 1998 also perceived a decreased infectiousness associated with HAART, but this perception was not associated with an increased risk behavior [110]. A marked increase in risk-taking behavior (unprotected anal intercourse) - from 37% to 50% - was reported among gay men in the San Francisco Young Men's Health Study; this increase was also associated with increased rates of rectal gonorrhea

in STD clinics [111]. In a survey from the French SEROCO study group, gay or bisexual men were three times more likely to report unprotected sex with a seronegative partner after six months of HA-ART than they were prior to treatment initiation [112]. No increase in risk-taking behavior was noted among heterosexual individuals; however, the sample size was small. A decrease in sexual risk-taking behavior was documented in French IV-drug users who started HAART [113].

The increased sexual risk-taking behavior in homosexual individuals is alarming. Reduced likelihood of HIV transmission due to the widespread use of HAART could be offset by an increase in risk-taking behavior.

What happens next?

HIV prevention activities were developed in 1980 s the absence of reliable knowledge about HIV transmission. These prevention activities virtually excluded the infected index cases who would have been stigmatized with little benefit by learning their infection status. The situation in 2001 is completely different, and prevention strategies must be informed by ongoing scientific discoveries. First, vaccine and microbicide development directed at preventing infection must take advantage of our knowledge of mucosal exposure and mucosal immunity. Second, STD and behavioral interventions must embrace both HIV infected

and uninfected people, and the optimal usage for STD interventions demand intensive and emergent study. The failure of mass STD therapy to prevent incident HIV infection in a population in no way reduces the importance of treatment of STDs. Last, focus on HIV infected subjects is a strategy whose time has come. Early recognition of infection allows improved care for the infected subject as well as critical prevention opportunities. Indeed, the US Centers for Disease Prevention and Control has dedicated substantial prevention resources to this goal (Project SAFE). In addition, ART (Antiretroviral Therapy) may help to prevent transmission, and this issue has the highest global research priority. In the absence of a successful vaccine, strategies to reduce infectiousness - including use of therapy - are likely to develop in the coming years.

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Fachnachricht

Ist eine Bluttransfusion noch sicher?

Die BSE-Krise hat eine verstärkte Diskussion über die Sicherheit von Blutprodukten ausgelöst. Experten von Robert Koch- und Paul-Ehrlich-Institut haben angesichts der sich in Europa ausbreitenden Variante der Creutzfeldt-Jakob-Krankheit eine Gesamtstrategie zur Blutversorgung entwickelt. Bislang sei noch kein Fall einer vCJK-Übertragung durch Blut oder Blutprodukte bekannt geworden.

Nach "Worst-case-Szenarien" werden für Deutschland bis 2004 etwa 300-600 vCJK-Fälle erwartet, die auf Primärinfektionen über Nahrungsmittel zurückgeführt werden. Für eine Übertragung von Mensch zu Mensch gibt es bisher keine Hinweise, jedoch kann nicht ausgeschlossen werden, dass vCJK durch chirurgische Instrumente oder durch Blut oder Blutprodukte übertragbar ist. Noch gibt es keine geeigneten Nachweistests zur Erkennung der vCJK im Blut von Blutspendern. Bis ein geeigneter Test entwickelt wird, könnten als Risikovorsorge Personen von der Blutspende ausgeschlossen werden, die früher selbst Empfänger von Bluttransfusionen waren. Problem ist allerdings, dass mit dieser Maßnahme ca. 4% der Blutspender in Deutschland wegfallen würden. Bevor also Transfusionsempfänger von der Blutspende ausgeschlossen werden, sollten z.B. durch Werbeund Motivationskampagnen neue Spender gefunden werden.

Quelle: Pressestelle RKI (http://www.rki.de),

Pressestelle PEI (http://www.pei.de)