

Prevalence of enzootic simian viruses among urban performance monkeys in Indonesia

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Summary

Animal reservoirs are the most important sources of emerging infectious diseases that threaten human populations. Global travel and tourism bring ever-increasing numbers of humans into contact with animals, increasing the likelihood of cross species transmission of infectious agents. Non-human primates come into contact with humans in a variety of contexts and may harbor infectious agents with zoonotic potential. We investigated the prevalence of infection with enzootic simian viruses among 20 urban performance monkeys (*Macaca fascicularis*) in Jakarta, Indonesia. This report documents for the first time evidence of infection with four simian viruses in urban performance monkeys. Simian foamy virus was detected by PCR in 52.9% of the macaques. Antibodies to simian retrovirus were detected in 10.5% of the macaques. Antibodies to *Cercopithecine Herpesvirus 1*, were detected in 5.3% of the macaques. Similarly, antibodies to simian T-cell lymphotropic virus were detected in 5.3% of the macaques. No evidence of infection with simian immunodeficiency virus was detected in these macaques. These results suggest that urban performance monkeys are a reservoir for enzootic simian viruses known to be capable of infecting humans.

keywords simian retrovirus, simian T-cell lymphotropic virus, *Cercopithecine Herpesvirus 1*, simian foamy virus, macaque, primate zoonoses, Asia, performing monkeys

Introduction

Recent epidemics of zoonotic diseases such as HIV, SARS, avian flu (influenza H5N1), and hantavirus illustrate the importance of animal reservoirs as sources of emerging infectious diseases that threaten human populations globally. Animal/human contact in a variety of contexts can lead to zoonotic transmission. Some of these contexts, such as those associated with animal husbandry and pet ownership, zoological parks and petting zoos, laboratory research involving captive animals, monkey temples and the bushmeat and exotic animal trade are recognized (e.g., Wolfe *et al.* 1998, 2004; Engelthaler *et al.* 1999; Shortridge *et al.* 2000; Engel *et al.* 2002; Peeters 2002; CDC, MMWR 2003a, 2003b, 2003c; Bell *et al.* 2004; Reed *et al.* 2004; Jones-Engel *et al.* 2005a). The risk of zoonotic exposure associated with other contexts of animal/human contact such as urban and eco-tourism, however, is just now being explored.

By virtue of their genetic, physiological, and behavioural similarities with humans, non-human primates (NHPs) are

thought to be likely sources of pathogens that can pose a significant threat to human populations. The HIV pandemic is a cogent example of this threat. Recent research suggests that cross-species transmission of SIV from chimpanzees (*Pan spp.*) and sooty mangabeys (*Cercocebus atys*) to humans has occurred likely as a result of butchering practices associated with the bushmeat trade (Hahn *et al.* 2000). Subsequent human-to-human transmission eventually resulted in the world-wide spread of HIV in human populations.

Research on NHP-to-human transmission has yielded evidence that several enzootic simian viruses have the capacity to infect humans. Serological studies have suggested NHP-to-human transmission of CHV-1, SIV, SRV and SFV in laboratory and zoo workers exposed to captive primates (Khabbaz *et al.* 1994; Sandstrom *et al.* 2000; Lerche *et al.* 2001; Brooks *et al.* 2002; Huff & Barry 2003; Switzer *et al.* 2004), as well as in individuals involved in the bushmeat trade in Africa (Wolfe *et al.* 2004). The simian T-cell lymphotropic virus (STLV), a retrovirus

enzootic to Old World Primates, is hypothesized to be the progenitor of human T-cell lymphotropic virus (HTLV), through multiple cross-species transmissions (Franchini 1994; Vandamme *et al.* 1998; Gessain & Mahieux 2000). HTLV is implicated as an etiologic agent of adult T-cell leukemia and tropical spastic paresis (Gessain *et al.* 1985). *Cercopithecine herpesvirus 1* [CHV-1 (Herpes B)] is an alphaherpesvirus endemic to Asian macaques (Genus *Macaca*) (Weigler 1992; Huff & Barry 2003). Similar to herpes simplex in humans, CHV-1 in macaques causes mild symptoms consisting primarily of oral and perioral vesicular lesions. In humans, however, CHV-1 produces a fulminating meningoencephalitis with a mortality rate near 70% (Hummeler *et al.* 1959; CDC, MMWR 1989, 1998; Holmes *et al.* 1990, 1995).

Urban performance monkeys in Asia

Although performing NHPs are encountered throughout the world, Asian cultures have perhaps the longest and most vibrant tradition of using NHPs for entertainment. In Japan, a thousand-year history of training performance monkeys (*Macaca fuscata*) continues today through the Suo-Sarumawashi (Japanese Monkey Performance) (Ohnuki-Tierney 1987). Similar training schools can be found throughout South and Southeast Asia. At the Punjab Institute of Mental Health in Lahore, Pakistan, for example, performance monkeys (*Macaca mulatta*) are used to entertain psychiatric patients.

Performance Monkey 'niche'

The ecology of performance monkeys is distinct from that of primates that come into contact with humans in other contexts, such as temple monkeys, laboratory NHPs, and NHPs consumed as bushmeat. The acquisition of performance monkeys, the care they receive from their owners, the characteristics of the urban environments they typically inhabit, and the unique circumstances of the performances themselves are all components of the performance monkey 'niche' that influence the amount and kinds of contact these primates have with humans and, in turn, influence the likelihood of zoonotic transmission occurring in this context.

Regarding the risk of NHP-to-human transmission of infectious agents, it is important to consider the origin of performance macaques. Although no systematic research has been published describing how owners in other parts of Indonesia and in other countries acquire their animals, the majority of the macaques in the present study were obtained from animal markets, most likely from Pramuka, the large animal market in Jakarta. This is significant because animal

markets bring together a variety of species, primate and non-primate, from diverse geographic origins, each with its potentially unique burden of infectious agents (Karesh *et al.* 2005). Animals in these markets are typically maintained at high density and often in poor conditions, which may compromise immunity and facilitate disease transmission (Malone *et al.* 2002). Monkeys included in the present study were typically purchased from animal markets while still quite young. Intensive training with the owner lasted anywhere from 5 months to a year or more depending on how quickly the monkey learned his/her routines.

Performance monkeys usually live with their owner's family. Most of the owners we spoke with owned more than one monkey. As with pet NHPs, close physical contact including the sharing of food and water resources is common in these settings. Previous research suggests that a variety of infectious agents may be transmitted from pet NHPs to humans as well as from humans to pet NHPs (Mack & Noble 1970; de Arruda *et al.* 1989; Ostrowski *et al.* 1998; Jones-Engel *et al.* 2001, 2004, 2005b; Huemer *et al.* 2002; Peeters *et al.* 2002).

Performances can be quite elaborate and usually include music and commentary from the owners. The performing monkeys are often dressed in costume and perform acrobatics (Figure 1). Performances are typically held in



Figure 1 A juvenile performance monkey from Jakarta included in the present study.

areas such as popular tourist destinations where they are likely to draw large audiences. Owners of performance monkeys may encourage monkeys to climb onto spectators' shoulders and head for photos and monkeys may also come into physical contact with spectators as they collect money for their performance.

The urban environment, which performance monkeys typically inhabit, is in itself a significant aspect of the performance monkey 'niche'. The mobility of urban populations as well as large urban centres as hubs for international travel may bring together NHPs with people who otherwise might never have contact with NHPs.

Our purpose was to investigate the prevalence of select simian pathogens in a sample of urban performance monkeys from Jakarta, Indonesia, in order to determine whether these NHPs harbor enzootic infectious agents. Specifically, we investigated the prevalence of four simian-born retroviruses known to infect humans: simian immunodeficiency virus, simian retrovirus, simian foamy virus, simian T-cell lymphotropic virus, and an alphaherpes virus, *Cercopithecine herpesvirus 1*. Identifying the prevalence of these simian infectious agents in the urban performance monkey population is the first step in estimating the risk of NHP-to-human transmission to a variety of people who may come into contact with these monkeys. These include the monkeys' owners/trainers and their families, people who live in the community where the monkeys are kept, and tourists who observe performances.

Materials and methods

Sampling

In July 2002, 20 performance monkeys, all long-tailed macaques (*Macaca fascicularis*), were identified in Kampung Dukuh, East Jakarta, a village known to be a community where owners of performance monkeys are concentrated. After explaining our research protocol to each monkey owner, we sedated the macaques with 3 mg/kg of Telazol® (tiletamine HCl/zolazepam HCl). Using universal precautions, approximately 10 ml of blood were drawn from the femoral vein of each monkey. Macaques were monitored closely during anaesthesia and recovery. Approximately 6 ml of blood were placed in a separator tube and centrifuged in the field to extract the serum. The remaining whole blood was placed in a tube containing EDTA. Both the blood and serum samples were frozen and stored at -80 °C. Each macaque's weight and dental formula were collected and recorded for age assessment. Age was estimated based on the observed pattern of dental

eruption using a published dental aging protocol for *M. fascicularis* (Richtsmeier *et al.* 1993). Study and data protocols were reviewed and approved by the University of Washington's Institutional Animal Care and Use Committee (3143-03).

Laboratory analyses for SRV, STLV, SIV, SFV and CHV-1

Enzyme-linked immunoabsorbent assays (ELISAs) were used to detect antibodies to SRV, STLV, SIV and CHV-1. Immunoblot assays were performed to confirm all positive SIV, STLV and SRV ELISA results.

SRV antigens for EIA

To produce large volumes of SRV-2 preparations for EIA, one vial of infected A549 cells (ATCC CCL-185) was thawed and added to a T-25 culture flask containing Dulbecco's Modified Eagle Medium (Gibco) supplemented with 10% FBS (Hyclone) and penicillin/streptomycin solution. After 2 days at 37°C, the cells were trypsinized and transferred to a T-175 culture flask containing 100 ml of medium. The A549 cell culture supernatant was collected and cells were passed every 3–4 days at 1×10^5 cells/ml until a total of approximately 3–6 l of culture supernatant was harvested. The supernatant was clarified first by low speed centrifugation at 1000 rpm (Beckman GH-38 rotor) for 15 min at 4 °C then by filtration through 0.45 µm PES filters. The clarified and pooled supernatant was concentrated approx. tenfold by tangential flow filtration using a Pellicon^R-2, PLCHK cassette filter (Millipore) with a 100 000 NMWL cutoff. The concentrate was partially purified by ultracentrifugation on a 20% glycerol cushion at 19 000 rpm (Beckman Type 19 rotor) for 3 h at 4 °C. The pellets were dispersed in Dulbecco's Phosphate Buffered Saline (Gibco), layered on a 30% and 45% (wt/wt) sucrose step gradient, and centrifuged at 45 000 rpm (Beckman SW55Ti rotor) for 1.5 h at 4 °C. The SRV band at the interface of the two sucrose concentrations was aspirated and mixed with PBS, then applied to a 20% glycerol cushion. The virus was pelleted by ultracentrifugation at 45 000 rpm (Beckman SW55Ti rotor) for 1 h at 4 °C. Viral antigen was quantified using BCA colorimetric detection (Pierce) then frozen at -80 °C in aliquots of 1 mg/ml in PBS. Preparation lots of antigen have resulted in yields of approximately 0.5 mg of protein per liter of A549 cell culture supernatant. Quality control of each lot of antigen consists of titration and comparison using SDS PAGE followed by commassie blue staining and Immunoblot analysis.

STLV, SIV and CHV-1, purified virus for EIA

For STLV assays we relied on the considerable cross-reactivity of HTLV-1 (Zeptomatrix, Buffalo, NY) (Franchini & Reitz 1994) For SIV assays we purchased SIVmac251 (Zeptomatrix). For CHV-1 assays we used the considerable cross reactivity of herpes simplex-1 (HSV-1) purified viral lysate (Advanced Biotechnologies, Columbia, MD) (Hilliard *et al.* 1989).

ELISA for SRV, STLV, SIV and CHV-1

ELISA plates (Polysorp Nunc) were coated with 100 µl of SDS-treated (0.1% SDS at 56 °C for 30 min) viral protein at 70 ng/well in carbonate–bicarbonate buffer pH 9.6 (Sigma) and incubated overnight at 4 °C in a moist chamber. The viral lysates were removed and plates were incubated with blocking buffer, 5% non-fat dry milk (BioRad) in PBS-0.1% tween, for 1 h at 37 °C. The plates were washed in PBS-tween four times at this step and at all subsequent wash steps. Macaque plasma samples were heat inactivated at 56 °C for 30 min prior to application to ELISA plate. Plasma diluted 1:100 in blocking buffer was applied in duplicate to each well in 100 µl volumes and incubated at 37 °C for 1 h. Positive control plasma from macaques was a pool of historical samples confirmed to be positive by producing bands to all viral proteins on Immunoblot at three or more sequential time points. Negative controls were pooled negative plasma from the Washington National Primate Research Center (WaNPRC) specific pathogen-free colonies that have tested negative on four or more consecutive time points over two or more years confirmed to be negative on Immunoblots. After the plate was washed, 100 µl of secondary antibody (goat anti-monkey IgG-peroxidase, Sigma, 1:15 000) was added and plates were incubated at 37 °C for 1 h. The colour reaction was run using OPD (20 mg tablets, Sigma) in phosphate-citrate buffer containing urea hydrogen peroxide (Sigma) for 20 min. The colour reaction was stopped with 1N sulphuric acid and plates were read at 492 nm. Plate cut-off values were determined for each plate on each run using the equation: Cut-off value = average of eight or more neg. controls + (SD × 2.010) with a 95% confidence level (Frey *et al.* 1998).

SRV STLV and SIV Immunoblots

Sucrose gradient-purified SRV was loaded onto preparative SDS-PAGE 4–15% minigels (BioRad) at 60–75 ng/mm and blots were prepared on nitrocellulose as previously described (Benveniste *et al.* 1993) HTLV, cross-reactive with STLV, and SIV Immunoblot strips were purchased at

Zeptomatrix, Buffalo, NY. SRV strips were incubated; rocking overnight at room temperature, with heat-inactivated plasma diluted 1:100 in blocking buffer. STLV and SIV strips (Zeptomatrix) were incubated and tested following the manufacturer's instructions. Positive controls (1:250) consisted of pooled historical positive samples that produce bands to all viral structural proteins [SRV- (p10/p12, p14, p27, pp18, gp20 and gp70), (STLV – p19, p24/26, GD22, gp44, rgp44-HTLV-1), (SIV – p27, gp120, gp160)]. Negative controls were the same as those used for ELISA. Strips were washed in PBS-tween three times and incubated for 1 h at room temperature with goat anti-human IgG alkaline phosphatase (1:12 000, Sigma) diluted in blocking buffer. Strips were washed three times and reacted with BCIP/NBT solution (Sigma) in a dark location for 10–15 min. Reaction was stopped by washing in water, changed several times over 10 min. Reactions were deemed positive if core and envelop bands were present, indeterminate if only core or only envelop were present and negative if bands did not appear or were not darker than negative control plasma.

Purification of DNA for PCR

DNA was purified from whole blood using the QIAamp DNA Blood Mini Kit following the manufacturer's protocols (Qiagen) with modifications. Briefly 20 µl protease (2× concentrate) and 200 µl buffer AL were combined with 200 µl whole blood and incubated at 56 °C for 30 min. After incubation 200 µl ethanol was added and the entire mixture applied to a QIAamp spin column. The purified DNA was eluted from the column with 70 µl of 10 mM Tris-HCL pH 8.0 and the concentration and purity was determined spectrophotometrically at OD260 and 280. If the 260/280 ratio did not fall between 1.70 and 2.05 DNA was rejected for PCR analysis. Rejected DNA was reapplied to a new column and purified a second time.

Real-time PCR for SRV

PCR was performed using prepared master mix from the manufacturer (BioRad) containing sybrgreen or no dye. PCR reactions (50 µl volume containing 0.5 µg DNA + 20 pmol primer + water in BioRad master mix) and gp70 and gp20 primers were set up for each DNA sample. Each sample was run with both gp70 and gp20 primers in duplicate so all samples were tested in quadruplicate. The primers used for detection were derived from consensus sequences for SRV serotypes 1–5. The primers located in the gp70 region are conserved in serotypes 1–3 (5737, CYAGATGGCTACCAGAACG; 5943, ARGGCTTACCGTGTGTTG; and 5974

CCAGCACAGTCACAAGGCTTA) and primers located in the gp20 region are conserved in all known serotypes 1–5 (7585, CTGGWCAGCCAATGACGGG and 7695, CGCCTGTCTTAGGTTGGAGTG with probe (7621 Fam-TCACTAACCTAAGACAGGAGGGYCGTCA-BHQ1). These primers are used routinely at both WaN-PRC (R. Grant unpublished data) and California NPRC (N. Lerche personal communication) for SRV diagnostic testing. Standard controls consist of known copy number plasmids, containing SRV-2 envelop, serially diluted in negative macaque genomic DNA. In addition, positive controls were used in each PCR that consisted of 0.5 µg DNA extracted from Raji cells infected with SRV serotypes 1–5. Negative control DNA consisted of genomic DNA pooled from negative macaques in the WaNPRC specific pathogen-free colonies that have tested negative on four or more consecutive time points over two or more years. Thermal cycles were 94 °C – 2 min, followed by 41 cycles of 94 °C – 20 s, 65 °C – 30 s with data collection after the 65 °C step of each cycle. A final melt curve was run on the samples by heating from 60 to 95 °C in increments of 0.5 °C. Samples were judged positive or negative based on the threshold cycle (T_c) of the PCR product and the melting temperature of the product. To be considered positive, a sample was required to be reactive to both gp70 and gp20 primers or to be proven positive by DNA sequencing of gp70 or gp20 region. Samples with a T_c less than the negative control were deemed presumptive positive. Presumptive positive samples were further analysed by melting temperature profile of the PCR product. Samples were confirmed positive if the product had a melting temperature peak of 84–86 °C, within the range of all known serotypes of SRV-1 through SRV-5. A presumptive positive with an incorrect melting temperature peak was deemed negative after DNA sequencing of PCR product and confirmation of non-viral sequences. Based on routine use of this assay for screening more than 2800 samples per year from the macaque colony at the WaNPRC since 2002 and cell culture confirmation of more than 200 samples per year, the PCR assay has a specificity of >99% and sensitivity of >98% (R. Grant unpublished data). The limit of sensitivity for the assay is 50 copies/reaction based on the optimized plasmid standard controls.

Nested PCR for detection of STLV

A protocol to amplify a wide range of primate T-lymphotropic viruses (PTLV) using outer generic primers followed by more specific nested primers in the tax gene region of STLV was performed on DNA from peripheral blood cells as previously described (Vandamme *et al.* 1997). Results were analyzed on 2% Nusieve (Cambrex, Inc.) agarose gels

with ethidium bromide staining. The sensitivity and specificity of this assay has not been independently determined in our laboratory but plasmid copy number controls indicate a limit of sensitivity of 10–50 copies/reaction.

PCR detection of SFV

The presence of SFV DNA was determined using a nested PCR. Five hundred nanograms of purified DNA were combined with a PCR reaction mixture with a final concentration of 10 mM Tris (pH9.0), 50 mM KCl, 0.1% Triton X-100, 2 mM MgCl₂ 200 µM each dNTP, 0.15 mg/m BSA, 1u Taq polymerase, and 400 nM of each primer in a total volume of 50 µl. The primer pairs used were: first round, forward, 5' CAG T GA ATT CCAGAATCTCTTC 3', reverse, 5' CACTTATCCCCTAGATGGTTC 3' and second round, forward, 5' CCAGAATCTCTTCATCCCTACTACTA 3', reverse, 5' GATGGTTCCTAAGCAAGGC 3' (Khan *et al.* 1999). Touchdown PCR was used for both rounds with reaction conditions of: initial denaturation at 94 °C for 2 min, followed by seven cycles of 94 °C for 30 s, 60 °C for 30 s, and 72 °C for 45 s with a 2 °C decrease in annealing temperature per cycle to 48 °C, followed by 33 cycles of 94 °C for 30 s, 48 °C for 30 s, and 72 °C for 45 s with a final extension at 72 °C for 2 min. Second round conditions were the same except 19 cycles were used instead of 33.

Results

Biological samples were collected from 20 macaques. Twelve were female. Dental eruption sequences classified one macaque as an infant, 10 as juveniles, 6 as subadults and three as adults at the time of sampling. Unfortunately, serum from one of the macaques, 02TK39, was unavailable for testing and whole blood samples from three macaques, 02TK16, 02TK20 and 02TK36, yielded insufficient DNA for SFV PCR.

Serological tests

Serologic results are summarized in Table 1. Although 8 of 19 serum samples were above cutoff on SRV ELISA only two (10.5%) were confirmed positive by immunoblot tests, suggesting non-specific cross-reactivity on five of the SRV ELISAs. One SRV ELISA reactive serum sample was indeterminate on immunoblot. The result of STLV testing revealed 2 of 19 plasma samples were reactive on ELISA but only one (5.3%) was confirmed positive by immunoblots and the second was indeterminate. One (5.3%) of the 19 macaques tested was ELISA positive for CHV-1. None of the 19 macaque sera tested were reactive on SIV ELISA.

Table 1 Sample information and results for serological and molecular analyses

ID	Sex	Age Class*	Weight†	SRV			STLV			SIV	SFV	CHV-1
				ELISA	Blot	PCR	ELISA	Blot	PCR	ELISA	PCR	ELISA
02TK16	F	2	1.5	React	Neg	Neg	Neg		Neg	Neg		Neg
02TK17	F	2	1.2	Neg		Neg	Neg		Neg	Neg	Neg	Neg
02TK18	F	4	3.0	Neg		Neg	Neg		Neg	Neg	POS	Neg
02TK19	M	2	1.8	React	IND	Neg	Neg		Neg	Neg	POS	Neg
02TK20	F	3	2.5	Neg		Neg	Neg		Neg	Neg		Neg
02TK21	F	3	2.2	Neg		Neg	Neg		Neg	Neg	Neg	Neg
02TK22	M	2	1.0	React	Neg	Neg	Neg		Neg	Neg	POS	Neg
02TK23	M	2	1.7	React	POS	Neg	Neg		Neg	Neg	Neg	Neg
02TK25	F	3	2.0	Neg		Neg	Neg		Neg	Neg	POS	Neg
02TK26	F	2	1.0	React	Neg	Neg	Neg		Neg	Neg	Neg	Neg
02TK28	F	2	2.0	React	Neg	Neg	Neg		Neg	Neg	POS	Neg
02TK29	M	2	1.2	Neg		Neg	React	POS	POS	Neg	POS	Neg
02TK31	M	3	3.5	Neg		Neg	Neg		Neg	Neg	Neg	Neg
02TK32	F	1	1.0	Neg		Neg	Neg		Neg	Neg	POS	Neg
02TK33	F	3	3.0	React	Neg	Neg	Neg		Neg	Neg	POS	POS
02TK35	M	2	3.0	React	POS	Neg	React	Neg	Neg	Neg	Neg	Neg
02TK36	F	2	1.8	Neg		Neg	Neg		Neg	Neg		Neg
02TK37	M	4	4.5	Neg		Neg	Neg		Neg	Neg	Neg	Neg
02TK38	M	4	3.8	Neg		Neg	Neg		Neg	Neg	POS	Neg
02TK39	F	3	2.7								Neg	
Total % Positive					10.5%			5.3%		0.00%	52.9%	5.3%

POS, positive; Neg, negative; IND, indeterminate.

* Age in years based on observed dental eruption pattern. 1, infant; 2, juvenile; 3, subadult; 4, adult.

† Weight in kilograms.

PCR tests

The majority of the animals, 9 out of 17, (52.9%) tested for SFV by PCR were positive. Repeated attempts to amplify SRV from all samples, including those positive by immunoblot, using PCR primers in two different regions of the genome were not successful. STLV was amplified from the single seropositive macaque (5.3%) but the seroindeterminate animal was PCR negative. STLV PCR specific for common isolates was negative, however, PCR to amplify generic PTLV DNA was strongly positive, indicating a divergent STLV present in the positive macaque. DNA was of sufficient quality and control betaglobin primers amplified the correct size product (data not shown).

The results of the serological and PCR assays were analyzed quantitatively to test associations by sex and age category. The results from these tests indicate there is no statistically significant association between seropositivity (for antibodies to SRV, STLV or CHV-1) and sex (OR = 3.073, 95%CI = 19.513–0.484, $P = 0.352$), or age ($\chi^2 = 0.04$, $P = 0.843$). Similarly, there is no statistically significant association between a positive SFV PCR

result and sex (OR = 0.818, 95%CI = 4.976–0.135, $P = 1.00$), or age ($\chi^2 = 0.08$, $P = 0.778$).

Discussion

This report documents the presence of antibodies to SRV, STLV and CHV-1, as well as the presence of simian foamy virus DNA in a sample of urban performance monkeys from Jakarta, Indonesia. Two of 20 urban performance monkeys had evidence of past or present infection with more than one enzootic simian virus.

Although two of the performance monkeys were found to be seropositive for SRV, viral DNA could not be amplified from any of the 20 animals surveyed through repeated attempts using primers in conserved and non-conserved locations of the genome, gp20 and gp70, respectively. The finding of SRV seropositive macaques with no detectable virus in circulating cells has been reported in SRV infections (Lerche *et al.* 1994). Routine surveys of macaques typically find those animals with antibody responses to the neutralizing epitopes of gp70 are negative or express very low copy numbers of SRV in blood cells (R. Grant unpublished data).

Although it is not unusual for animals to have long-term antibodies to SRV, it is not always possible to detect virus in these animals months or years after infection (Nick Lerche, personal communication, R. Grant unpublished data). As this study of performance monkeys used whole blood frozen at the field collection site, it is possible that some of the viral DNA was destroyed in the process of freezing and thawing many months later in the laboratory. Therefore, animals expressing low copy numbers of SRV may be undetectable in our assays and may explain the lack of any PCR positives.

The single STLV seropositive animal was positive by nested PCR. Only when using primers that amplify generic PTLV (Vandamme 1997), were we able to show a positive PCR, indicating infection with a more divergent isolate. Primers that amplify the most common HTLV/STLV genotypes did not work for this animal. Previous reports of STLV from Indonesian macaques have described divergent isolates with variable genotypes (Richards *et al.* 1998) and animals from other islands of Indonesia have reacted to the generic primers but not specific primers (L. Jones-Engel, unpublished data). Our current results are consistent with previous findings that STLV is present in Indonesian macaques but genetic diversity is unknown.

Seroprevalence patterns of the enzootic viruses studied here generally paralleled those seen in previously studied pet macaque (*Macaca nigra*, *Macaca nigrescens*, *Macaca maura*, *Macaca tonkeana*, *M. fascicularis*, *Macaca nemestrina*) populations on Sulawesi, Indonesia, but were lower than seroprevalence rates observed among a population of free-ranging macaques (*M. fascicularis*) at monkey temples in Bali, Indonesia. The percentage of performance monkeys seropositive for CHV-1 was consistent with the seropositivity observed among pet macaques in Sulawesi (15.4%; $n = 104$) but lower than that observed among temple macaques in Bali (82.0%; $n = 38$) (Engel *et al.* 2002; Jones-Engel 2002). SFV seroprevalence among the performance monkeys was also similar to the Sulawesi pets (53.7%; $n = 95$) and lower than that observed among Bali temple monkeys (89.5%; $n = 38$) (Jones-Engel *et al.* 2005a). Although the relatively small numbers included in the present study warrant caution in our interpretations, the seroprevalence patterns observed here may be explained by the similarity of this population to pet macaques in general, e.g. pet and performance monkeys are typically acquired when they are very young, limiting their duration of exposure to conspecifics especially during sexual maturation (Jones-Engel *et al.* 2005b). The higher seroprevalence of SFV and CHV-1 observed among free-ranging temple monkeys in Bali likely reflects intra-group transmission in more 'natural' contexts.

Risk of transmission

Urban performance monkeys have ongoing contact with their owners over a period of time, putting owners at risk for exposure to infectious agents that their animals may harbor. In many respects these performance monkey owners are probably like NHP pet owners (Jones-Engel *et al.* 2005b). Contact between NHP pets and their owners is often intimate, as pets climb about and cling to their owners, especially around head and shoulders. Food sharing between pets and owners is common. Pets often groom their owners, focusing on any wounds or irregularities in the skin. These activities all have the potential to bring macaque body fluids, especially oral secretions, into contact with their owner's mucus membranes, a potential portal of entry for infectious agents. Of the viruses examined in this research, owners may be at greatest risk for infection with SFV, which is prevalent among these macaques and which has been shown to be transmitted from macaques to humans in other contexts (Sandstrom *et al.* 2000; Brooks *et al.* 2002; Switzer *et al.* 2004; Wolfe *et al.* 2004; Jones-Engel *et al.* 2005a). High levels of SFV RNA are often found in normal (non-immunocompromised) macaque buccal swabs (M. Linial and S. Murray unpublished data) as such SFV can be thought of as having opportunity of transmission given everyday activities – which have the potential to bring saliva into contact with owners' mucous membranes, thought to be a likely route of transmission.

Transmission of the other viruses examined in this study may be relatively infrequent given their low seroprevalence among the performance monkeys. Certainly, the risk of exposure to these viruses among audience members is far less than that of owners, given their relatively fleeting contact with performance monkeys. However, monkeys that climb on audience members' head and shoulders could potentially transmit viruses to audience members. The likely small risk to audience members must be considered multiplied by the potentially very large number of individuals a monkey will come into contact with during his/her performance 'career'.

None of the monkeys tested exhibited serological evidence of exposure to simian immunodeficiency virus. While SIV is typically found only among African NHPs there exists a potential for the introduction of this virus into Asian NHP populations via the global trade in animals. This is an issue for performance monkeys in Java because most of them are acquired from animal markets, which are transit points for imported animals. It is therefore vital to regularly monitor Asian NHP populations for SIV as the virus has significant pathogenic potential in many of the Asian NHP species (Hirsch *et al.* 1989; Daniel *et al.* 1987).

Conclusion

Given the emphasis placed on HIV/SIV, which has its origins in Africa, the issue of NHP to human transmission of infectious agents in Asia and South America has been largely ignored. The above research provides preliminary insight into an underappreciated context of human-NHP interaction and points the way to further study. More work in this field is needed to characterize the number and species of performing monkeys and their owners in countries throughout Asia. It is also important to know about the epidemiology of audience members as well as characterizing the types and extent of contact between performance monkeys and audience members. Finally, serostudies of performance monkey owners and their families will elucidate the rate at which viral transmission occurs in this context.

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M. A. Schillaci *et al.* **Simian Viruses in Urban Performance Monkeys**

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