

**Congenital CMV and HIV Perinatal Transmission**

<sup>1</sup>Kristina Adachi, MD, <sup>2</sup>Jiahong Xu, MS, MPH, <sup>1</sup>Bonnie Ank, BA, <sup>3</sup>D. Heather Watts, MD, <sup>2</sup>Margaret Camarca, RN, MPH, <sup>4</sup>Lynne M. Mofenson, MD, <sup>5</sup>Jose Henrique Pilotto, MD, <sup>6</sup>Esau Joao, MD, <sup>7</sup>Glenda Gray, MD, <sup>8</sup>Gerhard Theron, MD, <sup>9</sup>Breno Santos, MD, <sup>10</sup>Rosana Fonseca, MD, <sup>11</sup>Regis Kreitchmann, MD, <sup>12</sup>Jorge Pinto, MD, <sup>13</sup>Marisa M. Mussi-Pinhata, MD, <sup>14</sup>Daisy Maria Machado, MD, <sup>15</sup>Mariana Ceriotto, MD, <sup>16</sup>Mariza G. Morgado, PhD, <sup>1</sup>Yvonne J. Bryson, MD, <sup>16</sup>Valdilea G. Veloso, MD, <sup>16</sup>Beatriz Grinsztejn, MD, PhD, <sup>17</sup>Mark Mirochnick, MD, <sup>4</sup>Jack Moye, MD, and <sup>1</sup>Karin Nielsen-Saines, MD, MPH for the NICHD HPTN 040 Study Team.

<sup>1</sup>David Geffen UCLA School of Medicine, Los Angeles, CA 90095-1406, U.S.; <sup>2</sup>Westat, Rockville, MD, 20850-2062, U.S.; <sup>3</sup>Office of the Global AIDS Coordinator, U.S. Department of State, Washington D.C., 20522-2920, U.S.; <sup>4</sup>*Eunice Kennedy Shriver* National Institute of Child Health and Human Development, National Institutes of Health, Bethesda, MD, 20892-7510, U.S.; <sup>5</sup>Hospital Geral de Nova Iguaçu, Nova Iguaçu, RJ, 26030-380, Brazil; <sup>6</sup>Hospital Federal dos Servidores do Estado, Rio de Janeiro, RJ, 20221-903, Brazil; <sup>7</sup>SAMRC and Perinatal HIV Research Unit, University of Witwatersrand, Cape Town, 7441, South Africa; <sup>8</sup>Stellenbosch University/Tygerberg Hospital, Cape Town, 7505, South Africa; <sup>9</sup>Hospital Conceicao, Porto Alegre, RS, 91350-200, Brazil; <sup>10</sup>Hospital Femina, Porto Alegre, RS, 90430-001, Brazil; <sup>11</sup>Irmandade da Santa Casa de Misericórdia de Porto Alegre, RS, 90020-090, Brazil; <sup>12</sup>Federal University of Minas Gerais, Belo Horizonte, MG, 30130-100, Brazil; <sup>13</sup>Ribeirão Preto Medical School, University of São Paulo, SP, 14049-900, Brazil; <sup>14</sup>Escola Paulista de Medicina-Universidade Federal de São Paulo, São Paulo, SP, 04939-002, Brazil;

<sup>15</sup>Foundation for Maternal and Infant Health (FUNDASAMIN), Buenos Aires, C1439CNU, Argentina; <sup>16</sup>Fundacao Oswaldo Cruz (FIOCRUZ), Rio de Janeiro, RJ, 21045-900, Brazil;

<sup>17</sup> Boston University School of Medicine, Boston, MA 02118.

**Corresponding Author:** Kristina Adachi, MD , David Geffen School of Medicine at UCLA, Department of Pediatrics, Division of Infectious Diseases, 10833 Le Conte Ave, MDCC 22-442, Los Angeles, CA 90095-1406, Phone: 310-825-5235 Fax: 310-206-4764 Email: [kadachi@mednet.ucla.edu](mailto:kadachi@mednet.ucla.edu)

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Change of Author Affiliations: Lynne Mofenson, MD has retired from service at the Eunice Kennedy Shriver National Institute of Child Health and Human Development, National Institutes of Health, Bethesda, MD, but is currently working with the Elizabeth Glaser Pediatric AIDS Foundation, Washington DC, U.S. D. Heather Watts was previously at the Eunice Kennedy Shriver National Institute of Child Health and Human Development, National Institutes of Health, Bethesda, MD, U.S., but is currently at the Office of the Global AIDS Coordinator, U.S. Department of State, Washington D.C., U.S.

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## ABSTRACT

**Background:** Congenital cytomegalovirus (CMV) infection (cCMV) is an important cause of hearing loss and cognitive impairment. Prior studies suggest that HIV-exposed children are at higher risk of acquiring cCMV. We assessed the presence, magnitude, and risk factors associated with cCMV among infants born to HIV-infected women, who were not receiving antiretrovirals during pregnancy.

**Methods:** cCMV and urinary CMV load were determined in a cohort of infants born to HIV-infected women not receiving antiretrovirals during pregnancy. Neonatal urines obtained at birth were tested for CMV DNA by qualitative and reflex quantitative real-time PCR.

**Results:** Urine specimens were available for 992 (58.9%) of 1684 infants; 64 (6.5%) were CMV-positive. Mean CMV load (VL) was 470,276 copies/ml (range: <200-2,000,000 copies/ml). Among 89 HIV-infected infants, 16 (18%) had cCMV versus 42 (4.9%) of 858 HIV-exposed, uninfected infants ( $p < 0.0001$ ). cCMV was present in 23.2% of infants with *in utero* and 9.1% infants with intrapartum HIV infection ( $p < 0.0001$ ). Rates of cCMV among HIV-infected infants were four-fold greater (aOR 4.4, 95% CI 2.3-8.2) and six-fold greater among HIV *in utero*-infected infants (aOR 6, 95% CI 3-12.1) compared with HIV-exposed, uninfected infants. cCMV was not associated with mode of delivery, gestational age, Apgar scores, six-month infant mortality, maternal age, race/ethnicity, HIV viral load, or CD4 count. Primary cCMV risk factors included infant HIV-infection, particularly *in utero* infection.

**Conclusion:** High rates of cCMV with high urinary CMV VL were observed in HIV-exposed infants. *In utero* HIV-infection appears to be a major risk factor for cCMV in infants whose mothers have not received combination antiretroviral therapy in pregnancy.

## INTRODUCTION

Cytomegalovirus (CMV) is a significant cause of congenital infections worldwide. In the U.S. and other industrialized countries, congenital CMV affects  $\leq 1\%$  of all newborns and accounts for over 40,000 neonatal infections per year.<sup>1,2</sup> In resource-limited countries, where maternal CMV seropositivity rates are higher and there have been limited data on congenital CMV infection (cCMV) rates, the problem of cCMV may be even more widespread.<sup>1,3,4</sup> cCMV's importance and potential clinical sequelae should not be underestimated. Approximately 10-15% of congenitally-infected infants have symptomatic disease that may progress to severe neurodevelopmental delays and sensorineural hearing loss.<sup>1,5-7</sup> Furthermore, approximately 5-17% of asymptomatic congenitally-infected infants may also develop clinical findings with later disease progression.<sup>8-10</sup>

Earlier studies have indicated that cCMV may be more common among HIV-exposed infants (2-7%) and among HIV-infected infants (4-26%).<sup>11-15</sup> These findings have particular relevance for HIV-infected infants, as they are more likely to have symptomatic cCMV, and cCMV may accelerate HIV disease progression.<sup>11, 12, 14, 16, 17</sup>

To investigate these issues further, we performed a study using data and specimens from the National Institute of Child Health and Human Development (NICHD) HIV Prevention Trials Network (HPTN) 040 perinatal clinical trial cohort of HIV-exposed infants. The primary objective of this sub-study was to determine the rate of CMV co-infection among HIV-exposed infants. Additional secondary objectives included comparing rates of cCMV among HIV-exposed, uninfected and HIV-infected infants (HIV acquired either *in utero* or *intrapartum*); determining predictors of cCMV; and evaluating mortality rates among infants with cCMV.

## METHODS

Study Design This was a sub-study of the NICHD HPTN 040 trial, also known as the International Maternal Pediatric Adolescent AIDS Clinical Trials Network (IMPAACT P1043) NICHD/HPTN 040 (or P1043), a phase 3, randomized, open-label, multi-center study that evaluated the efficacy, safety, and tolerance of three different infant antiretroviral regimens for the prevention of intrapartum HIV transmission to infants born to HIV-infected pregnant women, who had not received antiretroviral drugs during pregnancy.<sup>18</sup> The study enrolled 1684 HIV-infected pregnant women diagnosed with HIV infection at the time of labor and delivery. All women provided written informed consent. Enrollment occurred at multiple sites in Brazil, South Africa, Argentina, and the United States. Infants <32 weeks of gestational age were excluded from the study. Maternal plasma HIV RNA levels and CD4+ T-lymphocyte subsets were obtained at the time of labor and delivery. The primary endpoint of the parent study was HIV infection status at 3 months of age. Infants were followed until 6 months of age for safety and toxicity monitoring in the parent study.

HIV Diagnosis HIV DNA polymerase chain reaction (PCR) (Roche Molecular Systems Inc., Basel, Switzerland) was performed on infant blood specimens within 48 hours of birth and at 10-14 days, 4-6 weeks, 3 months, and 6 months of age. Repeat HIV DNA PCR testing was performed to confirm a positive result. Diagnosis of infant HIV infection required two positive HIV DNA PCR test results on separate specimens. Infants with a positive HIV DNA PCR test result at birth and positive results on repeat testing were classified as having *in utero* HIV infection. Infants with a negative HIV DNA PCR result at birth and a positive HIV DNA PCR



result on subsequent testing were classified as having *intrapartum* HIV infection. All HIV-exposed infants, who were enrolled in the study, were exclusively formula fed.

*Specimen Collection and CMV Testing* The presence of cCMV was evaluated in NICHD HPTN 040's population of HIV-exposed infants. cCMV and magnitude of urinary viral load was determined in HIV-uninfected, HIV-*in utero* infected, and HIV-*intrapartum* infected infants. Stored neonatal urine samples were collected within 48 hours of birth and frozen at  $-80^{\circ}\text{C}$  and stored at study sites. Infant urines were tested by qualitative Real-Time PCR for CMV DNA (FOCUS Diagnostics CMV Analyte Specific Reagent) with quantification of positive specimens. Infants with detectable CMV in the urine in the first 48 hours of life were diagnosed with cCMV.

*Statistical Analysis* Chi-square or Fisher's exact tests (when more than 25% of expected cell frequencies were less than five) was used to compare differences in proportions between cCMV and CMV-non-infected infants.

Univariate and multivariable logistic regression analysis was used to examine the relationship of cCMV and HIV infection, demographic/ geographic parameters, maternal characteristics, and infant mortality. Covariates with a p-value of less than 0.15 from univariate models were included in the initial multivariable model selection. All computations were done using SAS software v9.3 (Cary, NC, USA).

*Human Subjects* The study was approved by the institutional review boards and national ethics committees at each of the participating study sites.

## **RESULTS**

### *Rates of Congenital CMV Infection among HIV-exposed and HIV-infected Infants*

Urine specimens were available for 992 (58.9%) of the 1684 infants in the original study. Of these infants, 64 (6.5%) were found to have urines with detectable CMV, with a mean virus load

of 470,276 copies/ml (range: <200-2,000,000 copies/ml). Eleven (17.2%) infants with cCMV had results that were too elevated to quantify (>2,000,000 copies/ml), whereas approximately 8 (12.5%) had positive but low levels of detectable CMV (<200 copies/ml). The mean CMV urine virus load was higher for HIV-infected infants (697,698 copies/ml) than that of HIV-uninfected infants (448,897 copies/ml), but these differences were not significant.

Among the 992 infants, 89 (9%) were HIV-infected, 858 (86.6%) were HIV-uninfected, and 45 (4.5%) had unknown HIV status because of loss to follow-up or death before 3 months of age (the HIV diagnostic study endpoint). (Table 1) The rates of cCMV were significantly different when evaluated by infant HIV-status ( $p < 0.0001$ ). The cCMV rate among HIV-infected infants was 18% ( $n=16$  of 89 infants) compared to only 4.9% ( $n=42$  of 858 infants) among HIV-exposed, uninfected infants and 13.3% ( $n=6$  of 45) of infants with unknown HIV status. (Table 1) The difference was especially pronounced among HIV *in-utero*-infected infants, where 23.2% ( $n=13$  of 56) had cCMV as opposed to 9.1% ( $n=3$  of 33) of HIV *intrapartum*-infected infants. Compared with HIV-exposed uninfected infants, the rate of cCMV in HIV-infected infants was more than four-fold greater (OR 4.3, 95% CI 2.3-7.9) with multivariate logistic regression showing similar results aOR=4.4 (95% CI 2.3-8.2) after adjusting for study site country and maternal/race ethnicity, and was almost 6-fold greater among HIV *in-utero*-infected infants (OR 5.9, 95% CI 2.9-11.8), which was also similar in the adjusted analysis (aOR 6, 95% CI 3-12.1) after adjusting for study site country and maternal race/ethnicity. (Table 1, 2)

Deaths among infants  $\leq 6$  months of age occurred in 21 (2.1%) infants in the cohort with 2 deaths among 64 infants (3.1%) occurring in infants with cCMV. (Table 1) Infant mortality was not associated with cCMV. However, further evaluation revealed that HIV-exposed infants who died or were lost-to-follow-up before the 3-month HIV diagnostic endpoint also appeared to be

at increased risk of cCMV (OR 2.25, 95% CI 0.98-5.20), although this was marginally significant (p=0.057).

#### Other Risk Factors for Congenital CMV Infection

Apart from differences noted in rates of cCMV by infant HIV status and infant HIV mode of acquisition, cCMV rates also differed by geographical location of our study sites. (Table 1) Rates of cCMV ranged from 2-16.7% from top-enrolling sites in the Americas (primarily Brazil but also the US) but were only 2.1-2.5% in South Africa. Compared to infants in South Africa, HIV-exposed infants in the U.S. (aOR 15.6, 95% CI 1.3-193) and Brazil (aOR 5.1, 95% CI 1.4-18.8) had the greatest risk of acquiring cCMV, although it should be noted that the number of US infants was small (N= 7). In contrast, cCMV was not associated with other demographic and obstetric parameters including maternal age, race/ethnicity, mode of delivery, gestational age at delivery, or infant Apgar scores. Furthermore, maternal HIV viral load and maternal CD4 count were also not significantly associated with an increased risk of cCMV. (Table 1)

#### **DISCUSSION**

This study found high rates of congenital CMV infection (cCMV) in HIV-exposed infants, whose mothers were not receiving antiretroviral drugs in pregnancy due to late detection of HIV status resulting from delayed presentation to care. The rates of cCMV were highest among HIV-infected infants, particularly among those with *in utero* acquired HIV infection.

While CMV is among the most common etiologies of congenital infection worldwide, limited studies of cCMV have been reported from low and middle-income countries, particularly from HIV-infected pregnant women living in Latin America (Brazil) and sub-Saharan Africa (South Africa).<sup>1-4, 13, 19-23</sup> From studies of healthy pregnant women residing in industrialized nations,<sup>24</sup> we know that cCMV has been estimated to affect less than 1% of all newborn infants;<sup>1, 2</sup>

whereas, studies from Brazil and sub-Saharan Africa have generally suggested higher cCMV rates between 1.1-2.9%<sup>3,4,13</sup> and 1.4-14%,<sup>2,19-23</sup> respectively, in spite of documented high maternal CMV seroprevalence.

Our study provides further evidence demonstrating the extent to which cCMV rates may be elevated among HIV-exposed infants and HIV-infected infants. In our study, cCMV rates for HIV-exposed uninfected infants (4.9%) and HIV-infected infants (18%), particularly for infants that acquired HIV *in utero* (23%), were more than 4 to 23 times higher than cCMV rates reported among healthy infants in high-income countries (typically  $\leq 1\%$ ).<sup>1,2</sup> *In utero* HIV infection was the strongest predictor for cCMV (aOR 6, 95% CI 3-12.1).

Apart from two exceptions in smaller studies,<sup>19,25</sup> the cCMV rates among HIV-exposed, uninfected infants in our sub-study were higher than those seen in the majority of other published studies ranging from 2.2% to 4.6% among studies of HIV-exposed infants in the U.S., France, South Africa, Brazil, and Thailand.<sup>11,13-16,26-30</sup> Among HIV-infected infants, cCMV rates correlated with those reported in other studies, which typically ranged from 4.3% to 21%;<sup>11,12,14,16,29</sup> but included some studies with rates as low as 0%<sup>13</sup> and others with rates as high as 26-29%.<sup>31,32</sup>

Our study was designed to clearly discern between HIV *in utero* and *intrapartum* infections, since the primary objective of the parent study was to evaluate the use of postnatal antiretroviral prophylaxis in the prevention of *intrapartum* HIV transmission. Postpartum HIV or CMV transmission by breast milk was not of concern as formula feeding was an entry criterion for study participation. This distinction in the timing of HIV transmission in a large cohort of HIV-exposed infants allowed us to further explore potential associations between different transplacental infections, in this case HIV and CMV, which is a unique feature of our study.

Based on prior epidemiologic studies from Brazil and sub-Saharan Africa, it is presumed that the majority of women in our sub-study were CMV seropositive and demonstrated either CMV reinfection during pregnancy or re-activation given the risk factors of pregnancy-induced maternal immunosuppression, especially in the third trimester, and immunosuppression from undiagnosed and untreated HIV.<sup>26</sup> One potential explanation for the relatively high rates of cCMV in our cohort of HIV-exposed uninfected and HIV-infected infants may be that women were not diagnosed with HIV until the time of labor and delivery and were not on highly active antiretroviral treatment (HAART) during pregnancy. Some studies have suggested that the use of HAART by pregnant HIV-infected women may reduce cCMV rates among their HIV-exposed infants by improving their immune status during pregnancy.<sup>11, 26-28, 31</sup>

The findings of our study also provide additional evidence highlighting the complex inter-relationship “synergism” dynamics of HIV and CMV, suggesting that infection with one of these viruses may be a risk factor for infection with the other.<sup>12, 33, 19, 29, 34</sup> Both CMV and HIV have the ability to infect similar cells;<sup>12, 34-36</sup> *in vitro* studies have demonstrated that both viruses have the capability to stimulate gene expression and viral replication in the other.<sup>12, 14, 33-35, 37-39</sup> Other studies have also found that CMV may facilitate susceptibility to HIV infection by augmenting expression of Fc receptors, enhancing production of cytokines and other cellular products, and activating sentinel players such as T cells and monocytes.<sup>12, 33, 39, 40</sup> As with HIV and CMV, transplacental passage of one pathogen may facilitate passage of other concurrent pathogens, which has also been suggested for HIV and *Treponema pallidum* or even HIV and *Toxoplasma gondii*.<sup>34, 41-45</sup> For instance, in NICHD HPTN 040, we saw a higher rate of congenital syphilis infection overall and also among *in utero* HIV-infected infants.<sup>45</sup>

Although HIV-infection in infants was a risk factor for cCMV, other predictors such as lower maternal CD4 count (particularly  $< 200 \text{ cells/mm}^3$ )<sup>11, 27</sup> and younger maternal age were not associated with an increased risk of cCMV in our study, although there have been a reported associations in other studies.<sup>11</sup> It is possible that younger maternal age (potentially associated with a higher likelihood of CMV primary infection) may not play a significant role in settings like ours, where CMV reactivation or reinfection are likely responsible for CMV transmission. In addition, while prior studies have suggested high rates of infant death with cCMV (nearly 27%), infant mortality was low in our study and cCMV was not associated with an increased risk for infant death.<sup>12, 14, 16, 19</sup>

### Limitations

A major strength of our study was the relatively large sample size of HIV-exposed infants evaluated for cCMV compared to other studies, even though it was restricted to mother-infant pairs from our NICHD/HPTN 040 parent study<sup>18</sup> with available infant urines for specimen analysis; some sites were unable to collect urine specimens. Due to the lack of a comparison group of pregnant women without HIV, we could not evaluate cCMV rates in the general population and their risk factors for cCMV. Thus, our findings related to cCMV are only specific to infants born to HIV-infected pregnant women, who were not on HAART during pregnancy because of a late HIV diagnosis. In addition, because this was a secondary analysis, information on perinatal/postnatal CMV infection and symptomatic cCMV in infants with regards to rates of chorioretinitis, brain calcifications, microcephaly, other central nervous system anomalies, or hearing loss were not collected and is beyond the scope of this study.

## CONCLUSION

HIV-exposed infants are at significant risk for acquiring cCMV during pregnancy. HIV-infected infants, particularly those who acquired HIV *in-utero*, are at greatest risk for congenital CMV. CMV screening is an important component of a comprehensive evaluation needed for HIV-exposed infants, particularly among those born to women not on antiretrovirals during pregnancy.

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**Table 1. Demographic, Infant Delivery and HIV Related Characteristics by Infant CMV Status (Congenital CMV Infection)**

	<b>Overall</b> N=992 (column %)	<b>Infant CMV -Infected (+cCMV)</b> N=64 n (column %)	<b>Infant CMV -Uninfected (-cCMV)</b> N=928 n (column %)	<b>Unadjusted OR (95% CI)</b>	<b>p-value</b>
<b>Maternal Age, years</b>					
Mean [std. dev.]	26.5 [6.2]	26.3 [6.8]	26.5 [6.2]		
Median [min-max]	26 [13-47]	25 [14-4]	26 [13-47]		
13-24	407 (41.0)	31 (48.4)	376 (40.5)	1.14 (0.63-2.04)	0.66
25-29	289 (29.1)	13 (20.3)	276 (29.7)	0.65 (0.32-1.33)	0.24
30 and older	296 (29.8)	20 (31.3)	276 (29.7)	1.00	
<b>Maternal Race/Ethnicity</b>					
Black	419 (42.2)	26 (40.6)	393 (42.4)	1.35 (0.67-2.73)	0.40
Mixed/Mulatto	316 (31.9)	26 (40.6)	290 (31.3)	1.83 (0.90-3.70)	0.09
White/Others	257 (25.9)	12 (18.8)	245 (26.4)	1.00	
<b>Type of delivery</b>					
Cesarean after rupture/timing unknown	130 (13.1)	8 (12.5)	122 (13.2)	0.92 (0.42-2.02)	0.83
Cesarean before rupture	276 (27.8)	17 (26.6)	259 (27.9)	0.92 (0.51-1.66)	0.78
Vaginal	586 (59.1)	39 (60.9)	547 (58.9)	1.00	
<b>Infant Gestational age, weeks</b>					
36 or less	92 (9.3)	9 (14.06)	83 (8.94)	1.67 (0.79-3.49)	0.18
37 or more	900 (90.7)	55 (85.94)	845 (91.06)	1.00	
<b>Apgar score at 5 minutes</b>					
0-3	2 (0.2)	0 (0.00)	2 (0.2)	0.00 (0.00- I)	0.99
4-6	3 (0.3)	0 (0.00)	3 (0.4)	0.00 (0.00- I)	0.99
7-10	915 (99.5)	62 (100.0)	853 (99.4)	1.00	
<b>HIV Infant status</b>					
Unknown	45 (4.5)	6 (9.38)	39 (4.2)	<b>3.07 (1.23-7.66)</b>	<b>0.02</b>
<b>Positive</b>	<b>89 (9.0)</b>	<b>16 (25.0)</b>	<b>73 (7.9)</b>	<b>4.26 (2.28-7.94)</b>	<b>&lt;.0001</b>
Negative	858 (86.6)	42 (65.6)	816 (88.0)	1.00	
<b>Infant HIV Mode of Acquisition</b>					
Unknown	45 (4.5)	6 (9.4)	39 (4.2)	<b>3.07 (1.23-7.66)</b>	<b>0.02</b>

Intrapartum	33 (3.3)	3 (4.7)	30 (3.2)	1.94 (0.57-6.62)	0.29
<b>In-utero</b>	<b>56 (5.7)</b>	<b>13 (20.3)</b>	<b>43 (4.6)</b>	<b>5.87 (2.94-11.8)</b>	<b>&lt;.0001</b>
Negative	858 (86.6)	42 (65.6)	816 (88.0)	1.00	
Infant Death					
No	971 (97.9)	62 (96.9)	909 (98.0)	0.65 (0.15-2.85)	0.57
Yes	21 (2.1)	2 (3.1)	19 (2.1)	1.00	
Maternal HIV Viral load, copies/mL	----	-----	----	1.29 (0.94-1.71)	0.11
Mean [std. dev.]	63,682.2 [212,837]	59,058.3 [107,208]	64,002.1 [218,294]		
Median [min-max]	14,921 [0-3,055,766]	19,181 [195- 452,741]	14,578 [0-3,055,766]		
>100,000	117 (11.8)	8 (12.5)	109 (11.8)	2.06 (0.42-10.0)	0.37
10,000-99,999	459 (46.4)	36 (56.3)	423 (45.7)	2.38 (0.56-10.2)	0.24
1,000-9,999	310 (31.3)	18 (28.1)	292 (31.6)	1.73 (0.39-7.65)	0.47
400-999	45 (4.6)	0 (0.00)	45 (4.9)	0.00 (0.00- Inf )	0.98
0-399	58 (5.9)	2 (3.1)	56 (6.1)	1.00	
Maternal CD4 counts, cells/ $\mu$ L	----	----	----	1.02 (0.94-1.10)	0.71
Mean [std. dev.]	515.3 [303.07]	528.9 [308.01]	514.3 [302.9]		
Median [min-max]	460 [12-2160]	486 [65-1377]	459 [12-2160]		
<350	335 (34.4)	21 (33.3)	314 (34.5)	0.93 (0.52-1.66)	0.80
350 to 499	208 (21.4)	13 (20.6)	195 (21.4)	0.92 (0.47-1.82)	0.82
500+	431 (44.3)	29 (46.0)	402 (44.1)	1.00	
Study Site					
US	6 (0.6)	1 (1.6)	5 (0.5)	8.80 (0.77- 100)	0.08
<b>Brazil</b>	<b>851 (85.8)</b>	<b>60 (93.8)</b>	<b>791 (85.2)</b>	<b>3.34 (1.03-10.8)</b>	<b>0.04</b>
South Africa	135 (13.6)	3 (4.7)	132 (14.2)	1.00	

Note: All infants of HIV-infected women who had at least one available infant urine specimen were tested for CMV. N total differences for specific evaluated risk factors for congenital CMV reflect that data may not be available for every mother-infant pair.



**Table 2. Adjusted Relationship of Congenital CMV Infection (cCMV) with Potential Risk Factors**

<b>Congenital CMV Infection (cCMV)</b>		
	<b>aOR (95% CI)</b>	<b>p-value</b>
<b>Infant HIV status</b>		
Unknown	<b>5.52 (2.03-15.0)</b>	<b>0.001</b>
HIV positive	<b>4.37 (2.32-8.22)</b>	<b>&lt;.0001</b>
HIV negative	1.00	
<b>Infant HIV Mode of Acquisition</b>		
Unknown	<b>5.47 (2.01-14.9)</b>	<b>0.001</b>
Infected intrapartum	1.97 (0.56-6.91)	0.29
<b>Infected in Utero</b>	<b>5.98 (2.96-12.1)</b>	<b>&lt;.0001</b>
Negative	1.00	
<b>Study Site Country</b>		
US	<b>15.6 (1.26- 193)</b>	<b>0.03</b>
Brazil	<b>5.14 (1.40-18.8)</b>	<b>0.01</b>
South Africa	1.00	
<b>Maternal Race/ethnicity</b>		
Black	1.80 (0.86-3.76)	0.12
Mixed/mulatto	1.99 (0.96-4.09)	0.06
White/other	1.00	