

Decay Rate and HIV-1 DNA Reservoir Size Following Early Infant Antiretroviral Therapy

Department of Molecular Microbiology and Immunology, Johns Hopkins School of Public Health, Baltimore, MD; ²Pediatrics Infectious Disease, Northwestern University School of Medicine, Chicago, IL; ³Pediatric Infectious Disease, Johns opkins Medical Institutes, Baltimore, MD; ⁴School of Medicine; University of California, San Diego; ⁵Department of Pediatrics, University of Massachusetts Medical School, Worcester, MA; ⁶Department of Biostatistics, Johns Hopkins School

INTRODUCTION

STUDY POPULATION

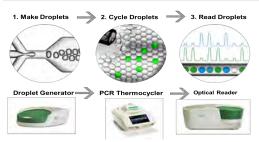
- Cure among HIV-1-infected individuals is precluded due to early establishment of latent HIV-1 reservoirs, primarily in resting memory CD4+ T cells1-3.
- · Early combination antiretroviral therapy (cART) restricts the size of the latent reservoir, and affords long-term virologic control in the absence of antiretroviral therapy (HIV-1 remission) 4.
- Rapid institution of cART at 30 hours of age may have contributed to prolonged HIV-1 remission (27 months instead of ~2 weeks) in the "Mississippi Child" by restricting reservoir size 5,6.
- The rapidity with which HIV-1 reservoirs are formed in perinatal infection, and the effects of early cART on the dynamics of decay of HIV-1-infected cells and their transcriptional states are not defined.

OBJECTIVE

Characterize decay of biomarkers of HIV-1 infection under Lopinavir/ritonavir-based ART in perinatally-infected infants during the first two years of life.

- · Cellular HIV-1 DNA : Total HIV-1 DNA (Pol); Episomal HIV-1 DNA (2-LTR circle)
- <u>Cellular HIV-1 RNA</u>: multiply spliced RNA (Tat/Rev); unspliced RNA (Gag)
- · Viral Outgrowth Assay: Replication competent HIV-1 DNA from resting CD4+ T cells 7.

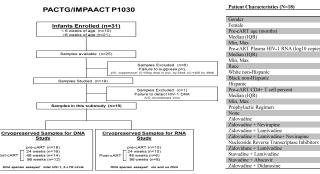
METHODS



o HIV-1 DNA- normalized to RPP30 (LOD: 2.6 copies/106 cells) o HIV-1 RNA- normalized to total input RNA (LOD- msRNA- 0.18 copies/ ugRNA; usRNA- 0.35 copies/ugRNA)

Adapted from Bio-Rad

Statistical Analysis: Slopes for HIV-1 DNA estimated using linear mixed effects model. Relationships between HIV-1 DNA, RNA, and viral load are summarized by follow-up week using either Pearson (for log-transformed variables) or Spearman correlation coefficients ***p≤0.0001; *p≤0.05 indicated statistical significance.

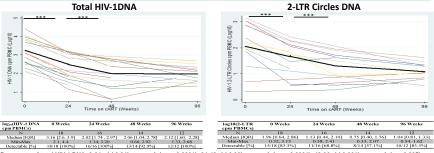


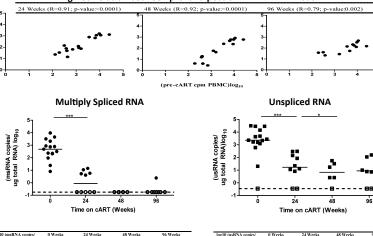
RESULTS

BM

DNA

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8 -0.75 [-0.75, -0.75] -0.75, 0.38

24 We

2] -0.07 [-0.75, 0.83] -0.75 [-0.75, -0.7 -0.75, 1.15 -0.75, -0.75

High Correlation between pre- and post-cART HIV-1 DNA load

96 Week

0.94 [=0.13, 1.78] =0.46, 2.19

.23 [0.57, 2.23] -0.46, 2.48

2.04 (1.35,3.78) 0.99, 5.39

5.77 (5.40,6.27) 3.72, 6.88

34.50% (29.0, 41) 11.0, 49.0

0 (0%)

12 (66 7%)

4 (28.6%)

10 (55.6%) 3 (16.7)

0 (0%) 1 (5.6%)

10 (55.6%)

7 (38.9%)

1 (5.6%)

CONCLUSIONS

msRNA usRNA

msRNA

usRNA msRNA

usRNA usRNA

RESULTS

HIV-1 DNA

and Q-VOA

Spearmans (p)

Min-Max

p-value (p)

usRNA/total HIV-1DNA

Median [IQR]

 Rapid establishment of a large pool of HIV-1 infected cells within two months of life in perinatal infection

Correlation of Total HIV-1 DNA and Viral Outgrowth Assay (VOA)

DNA/IUPM 66.0 [14.81, 240.15] 169.79 [52.19, 375.08] 148.33 [66.36, 273.61]

Ratio of HIV-1 RNA(us) and Total DNA

Correlation of HIV-1 RNA with pVL and Total DNA

Correlation with Plasma Viral Load (pVL

48 Weeks

0.39

0.339

8 65 751 55

18.24 [6.25, 36.60] 2.26 [0.37, 3.57] 0.68 [0.15, 1.07] 2.46 [0.73, 4.55] 0.05, 70.14 0.06, 15,44 0.03, 1.30 0.59, 81

0.72

24 Weeks

96 Weeks

0.44

53 59 933 3

0.02

0.002

0.02

0.46

A JOHNS HOPKINS

0.381

24 Weeks

0.66

0.028

7.06 641 60

0 Week

Correlation with HIV-1 DNA

- Pre-ART infected cell frequencies influence time to undetectable viral load and eventual reservoir size.
- · HIV-1 DNA was in excess of replication competent genomes at all time points tested.
- HIV-1 infected cells decrease in a biphasic manner in the first year of life with cART but stabilizes for an additional year.
- Preferential clearance of transcriptionally active cells with cART, though unspliced RNA detected through 96-weeks.
- · Findings have implications for refining cART for infants, including with immunotherapeutic approaches, towards long-term remission



NICHD

REFFERENCES

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Decay rate of total HIV-1 DNA :0-24 wk(-0.04log10 cpm/wk; p:<0.0001); 24-48wk(-0.02log10 cpm/wk; p:<0.0001); 48-96wk(-0.001log10 cpm/wk; p:0.295)