#### Time to DNA-PCR Positivity in Non-Breastfed HIV-Infected Infants (Primarily B HIV Subtype)

An analysis of data from the Perinatal AIDS Collaborative Transmission Study (PACTS) and the Women and Infants Transmission Study (WITS)

November 16, 2020







- Estimate the time to DNA PCR test positivity in non-breastfed, HIV infected infants
- Evaluate differences in DNA PCR test positivity according to exposure to maternal/infant HIV regimen
- Combine data from the Perinatal AIDS Collaborative Transmission Study (PACTS) and the Women and Infants Transmission Study (WITS)





## Background

- Accurate diagnostic tests to detect HIV infection in infants are critical to ensure early treatment.
- HIV DNA-PCR tests have imperfect sensitivity when given soon after birth.
- Our previous work in infants infected with non-B subtype HIV showed that time to HIV test positivity may be longer in infants exposed to combination therapy, but data are limited.
  - Balasubramanian, R., Fowler, M.G., Dominguez, K., Lockman, S., Tookey, P. A., Huong, N. N. G., Nesheim, S., Hughes, M. D., Lallemant, M., Toswill, J., Shaffer, N., Sherman, G., Palumbo, P., Shapiro, D. E. (2017): Time To First Positive HIV-1 DNA PCR May Differ With Antiretroviral Regimen In Infants Infected With Non-B Subtype HIV-1, AIDS, 31 (18), pp. 2465-2474. PMCID: PMC5710822.





### Cohorts

- Pediatric AIDS Collaborative Transmission Study (PACTS)
  - A multicenter, prospective cohort study (1986-1999) of HIV-infected pregnant women and their newborns conducted in 4 US cities to monitor the incidence of mother-to-child HIV transmission and to describe the natural course of pediatric HIV disease progression.
- Women and Infants Transmission Study (WITS)
  - WITS is a prospective epidemiologic study (1999-2007) of the natural history of HIV infection in pregnant women and their infants carried out at obstetric/gynecologic and pediatric clinics in Boston, Chicago, Manhattan, Brooklyn, San Juan, and Houston.





## Inclusion/exclusion criteria

- Inclusion Criteria:
  - HIV infection status: Infant HIV-infected or indeterminate; AND
  - Infant has at least one DNA-PCR test before age 3 months
- Exclusion Criteria:
  - DNA PCR tests with missing result or missing age at time of blood draw
  - Excluded infants whose mother's ARV exposure was unknown





## Summary of participant characteristics

#### WITS:

- Number of HIV positive infants: 129
- Number of HIV positive mothers: 126

#### PACTS:

- Number of HIV positive infants: 299
- Number of HIV positive mothers: 298





# Classifying Maternal and Infant ARV Regimen

Maternal ARV :

 Most complex ARV received in the 3<sup>rd</sup> trimester and at the time of labor/delivery.

Infant ARV :

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Prophylactic regimen with a start date prior to 45 days after birth.





## Maternal ARV by Infant ARV

	Infant ARV				
iviaternal ARV	None	ZDV	Other		
A: no ARV	193	5	0		
B: Single NRTI	54	35	0		
C: 2-3 NRTIs	9	2	0		
D: 2-3 NRTIs with sdNVP	6	2	0		
E/F: E (3 or more ARVs, with NNRTIs, no PI) + F (3 or more ARVs, with PI)	80	23	3		
G: sdNVP only	5	1	0		
H: ZDV + sdNVP	8	2	0		





# Timing of DNA PCR tests







### Participant characteristics

		Matornal viral		Delivery Type		
Maternal ARV	Maternal CD4 closest to delivery	load closest to delivery	Gestational age	Vaginal	CS no labor/RO M	CS with labor/RO M
A: no ARV	530 [282 - 775]	11250 [1,125 - 41,250]	39 [35 - 40]	81	3	16
B: Single NRTI	400 [287 - 617]	8,000 [9 - 42,000]	37 [35 - 40]	82	7	11
C: 2-3 NRTIs	249 [65 - 536]	9 [0 - 37,714]	38 [36 - 40]	78	11	11
D: 2-3 NRTIs with sdNVP	455 [300 - 552]	1,183 [906 - 2,830]	38 [36 - 40]	67	0	33
E/F: 3 or more ARVs, with NNRTIs or PI)	459 [298 - 738]	0 [0 - 2,380]	38 [36 - 39]	44	38	18
G: sdNVP only	300 [97 - 473]	52,873 [10,441 - 106,879]	37 [31 - 38]	67	33	0
H: ZDV + sdNVP	471 [364 - 704]	3,553 [0 - 66,666]	37 [35 - 38]	50	30	20





## Statistical methods

- Used methods for interval-censored data
  - Time to DNA PCR test positivity is uncertain; in interval between last negative and first positive DNA-PCR test
- Regression modeling to adjust for potential confounders
  - Parametric Weibull proportional hazards models





## Cumulative Probability of Positive DNA-PCR

	Maternal ARV	Number of Infants	Birth - 1 day	≤ 14 days	≤ 30 days	≤ 90 days
	A [No ARV]	198	0.29 [0.22-0.38]	0.58 [0.5-0.66]	0.68 [0.58-0.77]	0.81 [0.68-0.91]
	B [Single NRTI]	89	0.25 [0.17-0.35]	0.51 [0.39-0.65]	0.60 [0.46-0.76]	0.74 [0.56-0.89]
	C [2-3 NRTIs]	11	0.04 [0.01-0.26]	0.1 [0.01-0.52]	0.13 [0.02-0.62]	0.18 [0.03-0.76]
	D [2-3 NRTI + sdNVP]	8	0.09 [0.02-0.34]	0.21 [0.06-0.62]	0.27 [0.08-0.72]	0.37 [0.11-0.84]
	E/F [3+ ARVs]	106	0.05 [0.02-0.09]	0.11 [0.06-0.18]	0.14 [0.08-0.23]	0.2 [0.12-0.32]
	G [sdNVP only]	6	0.28 [0.1-0.66]	0.57 [0.23-0.93]	0.66 [0.29-0.97]	0.8 [0.39-0.99]
	H [ZDV + sdNPV]	10	0.11 [0.03-0.32]	0.25 [0.09-0.59]	0.31 [0.11-0.69]	0.42 [0.16-0.82]
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# Effect of Maternal ARV on time to test positivity

(Weibull PH model; HR>1 Means Earlier Detection of HIV)

Maternal ARV Groups	Unadjusted Hazard Ratio [95% CI]	Adjusted Hazard Ratio [95% CI] 5.8 [2.6-12.8]	
A [No ARV]	7.5 [4.06-13.8]		
B	6.2	4.7	
[Single NRTI]	[3.1-12.4]	[2.0-11.1]	
C	0.9	1.8	
[2-3 NRTIs]	[0.1-6.9]	[0.2-14.7]	
D	2.1	2.0	
[2-3 NRTI + sdNVP]	[0.5-9.3]	[0.4-10.5]	
E/F [3+ ARVs]	1	1	
G	7.2	3.5	
[sdNVP only]	[2.0-25.6]	[0.8-14.5]	
H	2.5	2.5	
[ZDV + sdNPV]	[0.7-8.7]	[0.7-9.3]	
P value for ARV Group	1.28e-12	5.95e-4	





## Time to test positivity by infant ARV

(Weibull PH model; HR>1 Means Earlier Positive)

Infant ARV	Unadjusted Hazard Ratio [95% CI]	Adjusted Hazard Ratio [95% CI]
None	1.0 [0.6-1.5]	0.68 [0.39-1.19]
ZDV	1.0	1.0
Other	0.65 [0.09-4.87]	0.51 [0.06-4.31]
P value for infant ARV	0.90	0.38





### Summary

- Time to DNA-PCR test positivity was later with receipt of 
   <u>></u>3 ARVs than with
   no ARV or single NRTI
- Differences in test positivity rates were observed at birth and remained at 3 months after birth.
- Adjusting for maternal/infant characteristics did not attenuate the findings.





# Study team/Acknowledgements

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