

Jonathan Z. Li<sup>1</sup>, Autumn Kittilson<sup>1</sup>, Meghan Melberg<sup>1</sup>, Elizabeth Wonderlich<sup>2</sup>, Jennifer Kinslow<sup>3</sup>, Evgenia Aga<sup>4</sup>, Ronald Bosch<sup>4</sup>, Yijia Li<sup>1</sup>, Mark Pilkinton<sup>5</sup>, Lynsay MacLaren<sup>6</sup>, Michael Keefer<sup>7</sup>, Ed Acosta<sup>8</sup>, Lawrence Fox<sup>9</sup>, Liz Bar<sup>10</sup>, John Mellors<sup>11</sup>, Robert Coombs<sup>12</sup>, Steven Deeks<sup>13</sup>, Rajesh Gandhi<sup>14</sup>, Michael Busch<sup>15</sup>, Alan Landay<sup>3</sup>, Bernard Macatangay<sup>11</sup>, and Davey M. Smith<sup>16</sup> for the A5345 Study Team

<sup>1</sup>Brigham and Women's Hospital, Harvard Medical School, <sup>2</sup>Southern Research, <sup>3</sup>Rush University, <sup>4</sup>Harvard T.H. Chan School of Public Health, <sup>5</sup>Vanderbilt University Medical Center, <sup>6</sup>Whitman Walker Health, <sup>7</sup>University of Rochester, <sup>8</sup>University of Alabama, <sup>9</sup>National Institute of Allergy and Infectious Diseases, <sup>10</sup>ACTG Community Scientific Subcommittee, <sup>11</sup>University of Pittsburgh, <sup>12</sup>University of Washington, <sup>13</sup>University of California, San Francisco, <sup>14</sup>Massachusetts General Hospital, Harvard Medical School, <sup>15</sup>Vitalant Research Institute, <sup>16</sup>University of California, San Diego

## BACKGROUND

- Strategies for ART-free HIV remission ultimately require validation through treatment interruption (TI) studies to confirm a delay in HIV rebound
- Identification of biomarkers predictive of time to rebound is essential for developing and evaluating promising treatment strategies
- AIDS Clinical Trials Group (ACTG) trial A5345 is a prospective TI study to determine associations between virologic, immunologic and host biomarkers and time to rebound
- We compared the timing of HIV rebound for early-treated and chronic-treated cohorts

## A5345 STUDY DESIGN

### Key Inclusion Criteria

- ART initiation:
  - Initiated ART >6 months after date of infection (Cohort A, chronic-treated) OR
  - Initiated ART during Fiebig stages III-V (Cohort B, early-treated)
- ≥2 years of suppressive ART

### ART Restart Criteria

- HIV-1 RNA at ≥1,000 copies/mL on two consecutive measurements

**Table 1: Participant Characteristics**

	Overall (N=45)	Chronic (N=33)	Early (N=12)
Age (median years)	45	46	38
Sex			
Male	91%	88%	100%
Female	9%	12%	0%
Race/ethnicity			
White	56%	73%	8%
Black	9%	12%	0%
Hispanic	18%	12%	33%
Asian, Pacific Islander	13%	0%	50%
More than one race	4%	3%	7%
Country			
USA	87%	100%	50%
Thailand	13%	0%	50%
Baseline CD4 (cells/mm <sup>3</sup> )	771	783	742
Pre-ART Viral Load (log <sub>10</sub> copies/mL)	4.6 (N=35)	4.5 (N=23)	5.7 (N=12)

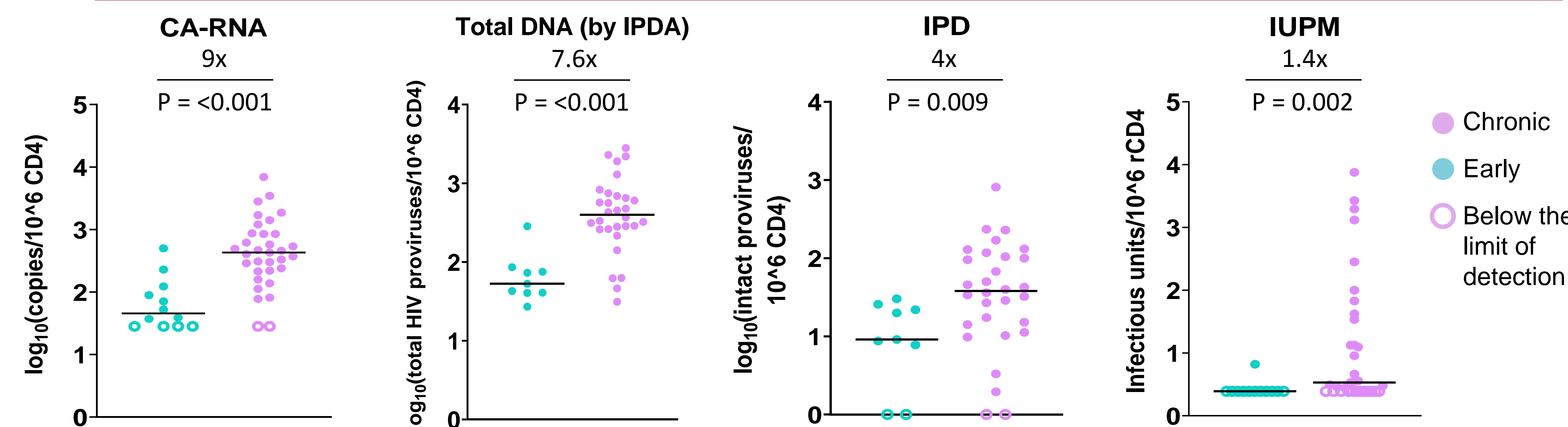
## Laboratory Assays

- Pre-TI viral reservoir markers were quantified by the following assays:
  - Unspliced cell-associated RNA (CA-RNA)
  - Total HIV DNA (total-DNA)
  - Intact proviral DNA (IPD) by the IPDA
  - Residual viremia by the integrase single-copy assay (iSCA)
  - Infectious units/million rCD4 cells (IUPM) by the differentiation quantitative viral outgrowth assay (dQVOA)
  - T cell phenotyping and HIV-specific immune responses to Gag, Env, Pol (CD107a, TNFα, IL-2, IFNγ)
  - Antibody levels (HIV-1+2 Ab, HIV Combo Ab, LAg-Avidity)

## RESULTS

- Early-treated participants had smaller (lower HIV DNA, IUPM, IPD) and less transcriptionally active HIV reservoirs (lower CA-RNA) (**Figure 1**)
- Lower reservoir activity was modestly associated with a delay in HIV rebound including CA-RNA (Spearman's  $r = -0.26$ ,  $p = 0.08$ ), the pre-specified primary measure (**Table 2**)
- The strongest predictor of time to HIV rebound was IPD in chronic-treated participants and iSCA in early-treated participants (**Table 2**)
- Pre-treatment interruption, significant correlations were detected across multiple reservoir markers and CA-RNA strongly correlated with total-DNA but only modestly correlated with intact proviral DNA (**Figure 2**)
- A higher percentage of HIV gag-specific CD8+ T cell cytotoxic response (CD107a) was associated with a lower IPD (Spearman's  $r = -0.37$ ,  $P=0.02$ )

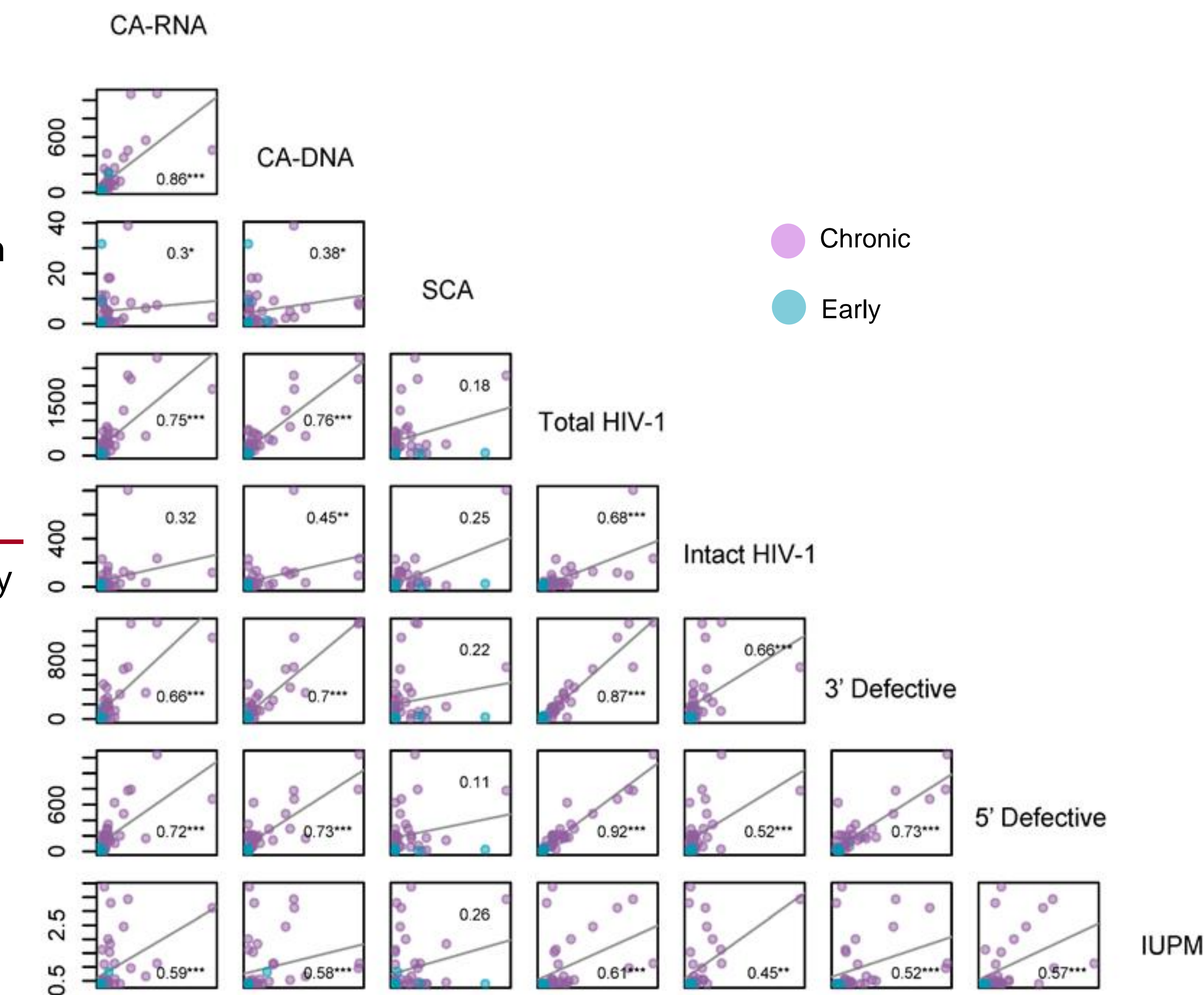
**Figure 1: Early-treated participant reservoirs are smaller and less active**



**Table 2: Reservoir predictors of HIV rebound timing**

	Overall N	Spearman		Overall N	Spearman		Overall N	Spearman	
		Correlation	p-value		Correlation	p-value		Correlation	p-value
CA-RNA (copies/10 <sup>6</sup> CD4)	45	-0.26	0.08	33	-0.27	0.13	12	-0.31	0.33
Total DNA (copies/10 <sup>6</sup> CD4)	39	-0.18	0.26	30	-0.26	0.17	9	-0.50	0.17
IPD (intact proviruses/10 <sup>6</sup> CD4 cells)	39	-0.28	0.09	30	-0.37	0.044	9	-0.27	0.48
iSCA (copies/mL)	45	-0.28	0.06	33	-0.11	0.55	12	-0.68	0.015
IUPM (infectious units/10 <sup>6</sup> rCD4 cells)	45	-0.18	0.25	33	-0.17	0.35	12	-0.18	0.58

**Figure 2: Reservoir marker correlations**



## CONCLUSIONS

- The strongest predictors of time to HIV rebound are the size of the intact reservoir for chronic-treated individuals and HIV residual viremia for early-treated individuals
- Smaller reservoir size and lower transcriptional activity in early-treated participants suggests early ART may lower the barrier to HIV remission
- We identified key HIV-specific CD8 immune responses associated with a smaller intact HIV reservoir size

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