# cfDNA in Heart Transplantation: A Powerful Biomarker

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# Faculty / Presenter Disclosure

- Faculty: Kiran K. Khush, MD, MAS
- Relationships with commercial interests:
  - Consulting Fees: CareDx, Inc.

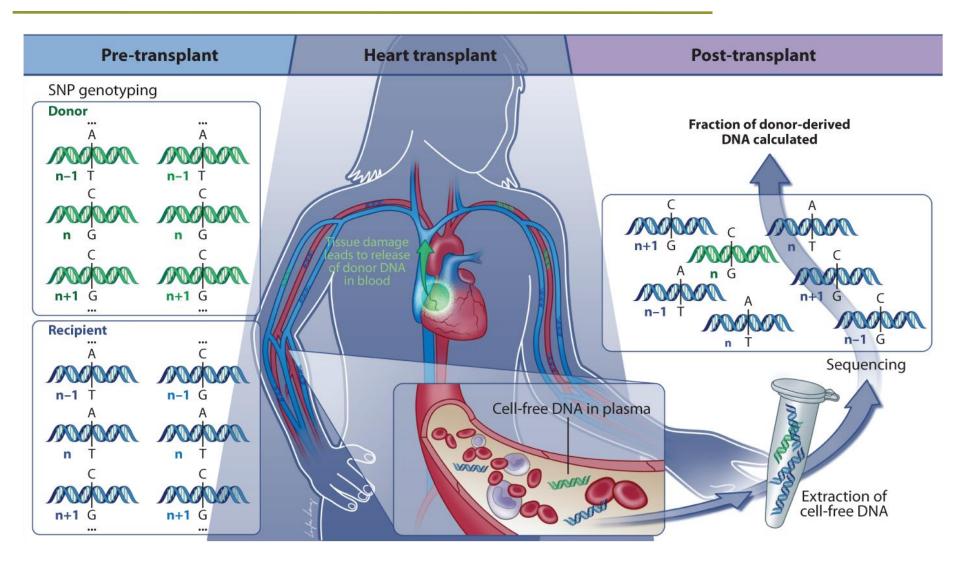


## **Mitigating Potential Bias**

 Only work that has been published and/or discussed at scientific meetings will be presented.

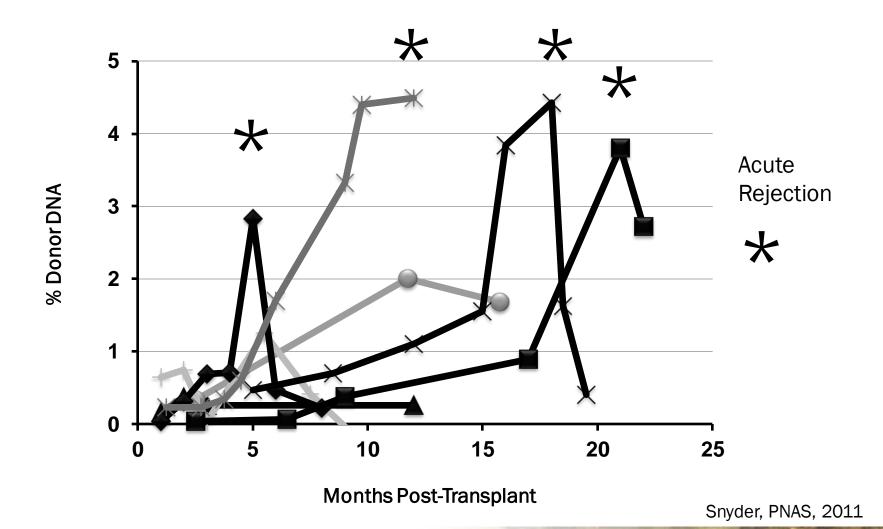


#### Cell free donor DNA as a marker of acute rejection



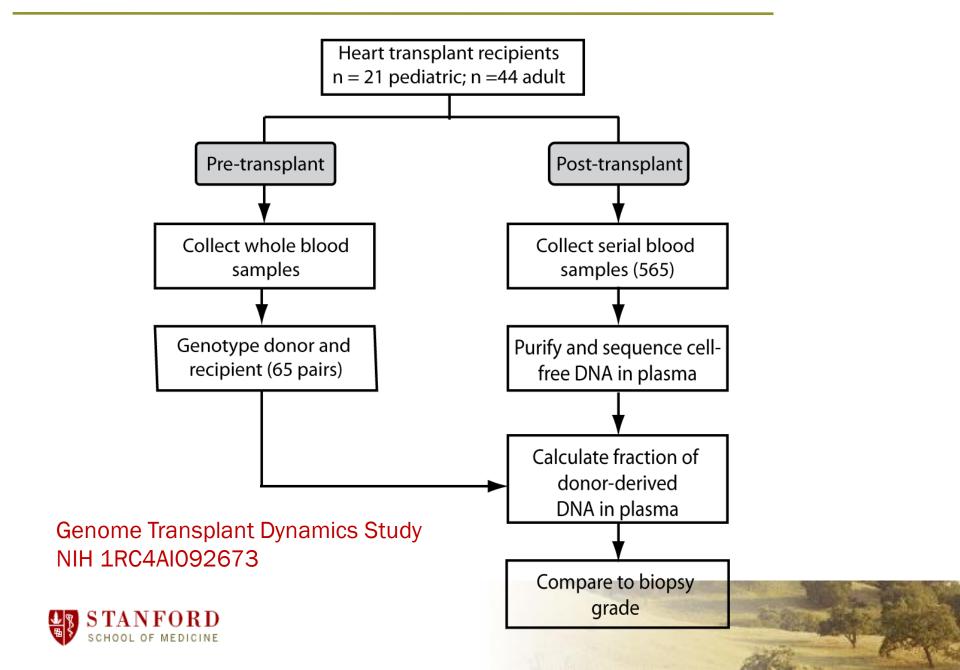


#### **Preliminary Results: Heart Transplant**

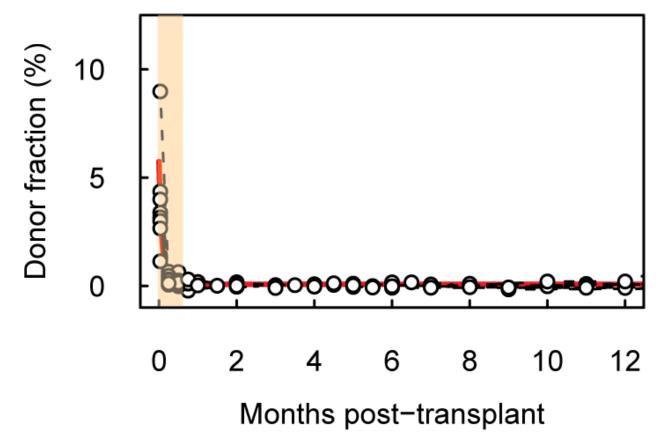




#### Heart transplant: prospective study design and numbers



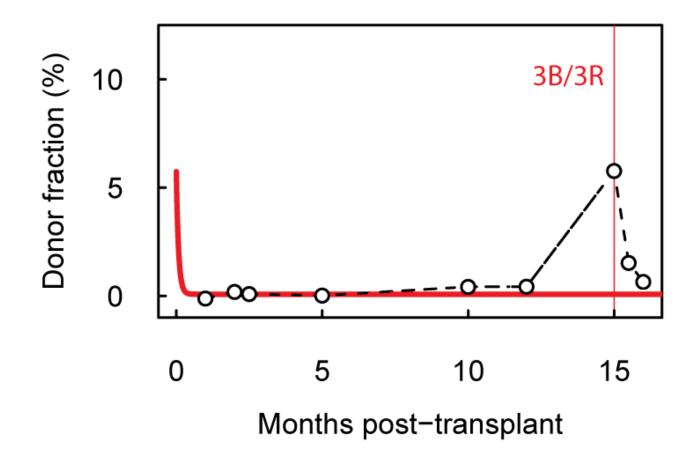
#### **Dd-cfDNA** in the absence of rejection



Elevated signal immediately post transplant followed by a quick decay (decay time 2.4 days) to a low baseline level

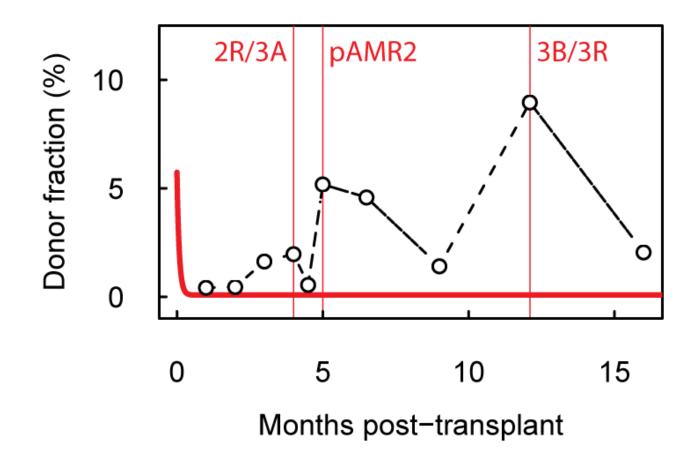


#### **Dd-cfDNA** at the time of acute cellular rejection



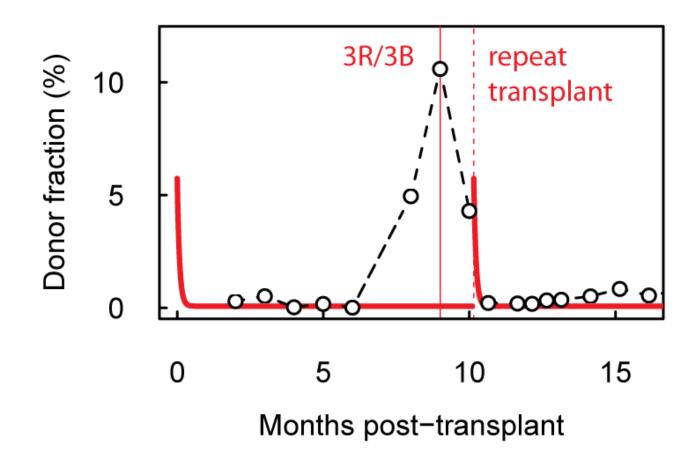


## Dd-cfDNA at the time of acute cellular and antibodymediated rejection



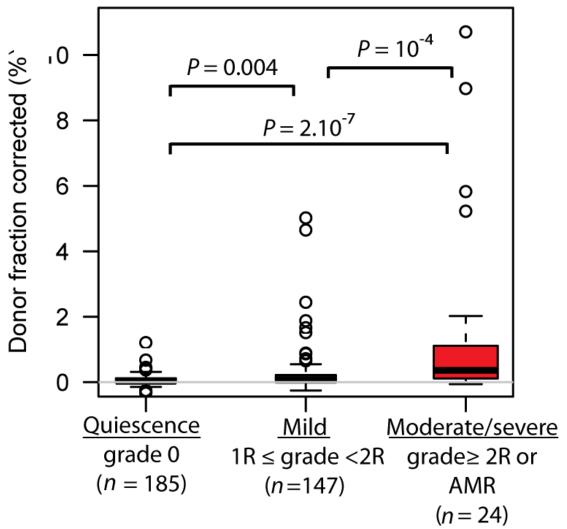


#### **Dd-cfDNA** in a case of graft loss and re-transplantation



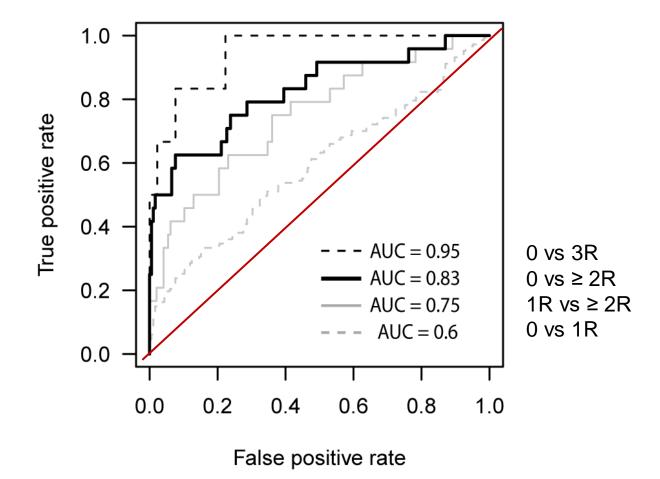


#### **Dd-cfDNA**: analysis of diagnostic performance



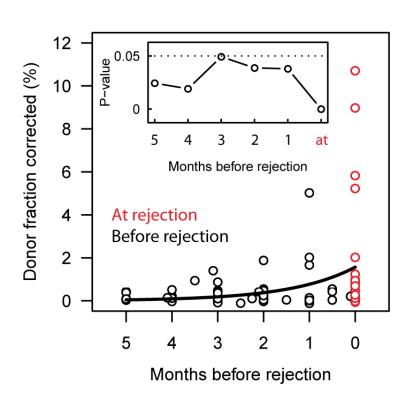


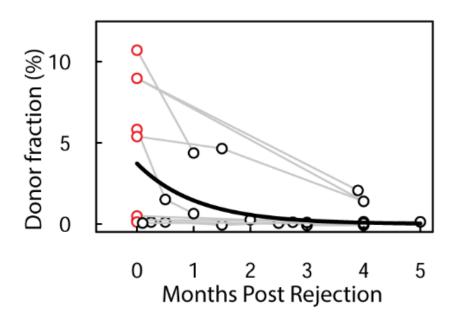
#### **Analysis of diagnostic performance**





#### Early diagnosis and monitoring of therapeutic response



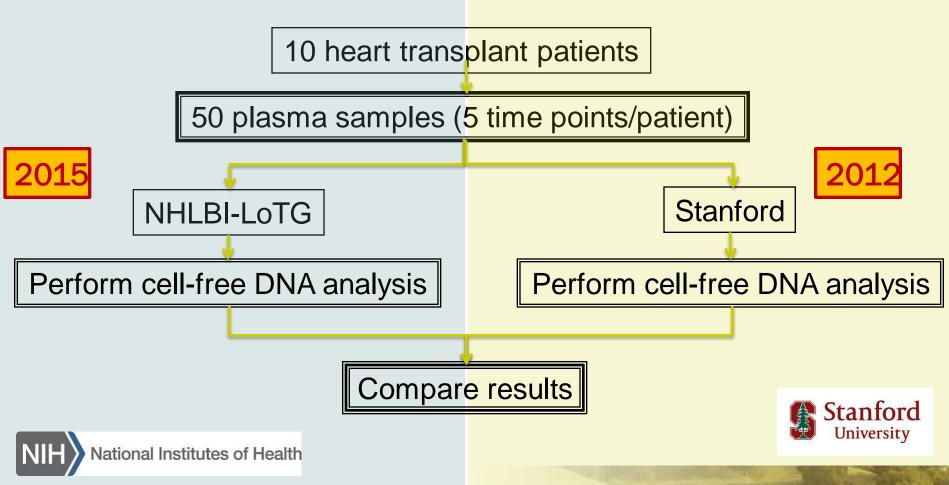




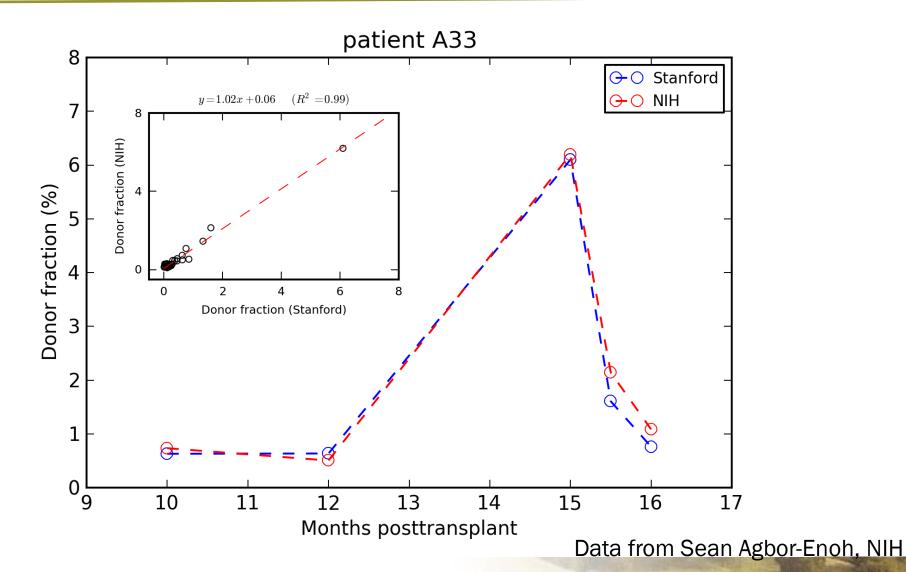
#### Replication/Reproducibility Studies

NHLBI - Genomic Research Alliance for Transplantation (GRAfT)





## Replication/Reproducibility Studies





#### dd-cfDNA: A Rapidly Evolving Technology

- Technology used in previous studies to measure SNP alleles:
  - Shotgun sequencing methods (Stanford) (1)
  - Targeted amplification (Wisconsin, Chronix) (2)
  - Both requiring recipient AND donor genotypes
- A new approach has been developed (CareDx, Inc: AlloSure™) with targeted amplification of SNPs (n=266) that DOES NOT require genotyping of the donor or recipient (3)
- "One genome" informatics algorithm (4)

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(1) Snyder et al., PNAS 108(15):6229, 2011
De Vlaminck et al., Sci Transl Med. 6(241):241, 2014
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(2) Beck et al., Clin Chem 59:12, 2013 Hidestrand et al., JACC 63:1224, 2014

nidestrand et al., JACC 63.1224, 2012

- (3) Grskovic et al, Jol Mol Diagnostics, Nov 2016
  - (4) Sharon et al. Submitted for publication





# CARGO II: Retrospective Analysis of dd-cfDNA (AlloSure™) in acute heart transplant rejection

CARGO II observational study: Heart transplant recipients from 17 centers; 737 patients, 7977 samples

Clinical status, including endomyocardial biopsy grades (graded by four independent pathologists) and blood were collected at routine surveillance visits for up to two years.

#### Rejection (R) cohort

-2/4 pathologists graded sample as 2R or 3R

N=58 patients

#### Selection for cfDNA Analysis

- -blood drawn prior to biopsy
- at least one preceding sample available

N=28 patients

#### **Study of Treatment Effect**

-3 visits per patient (two subsequent to rejection within 60 days)

N=17 patients



#### Quiescent (Q) cohort

-4/4 pathologists graded sample as OR

N=249 patients

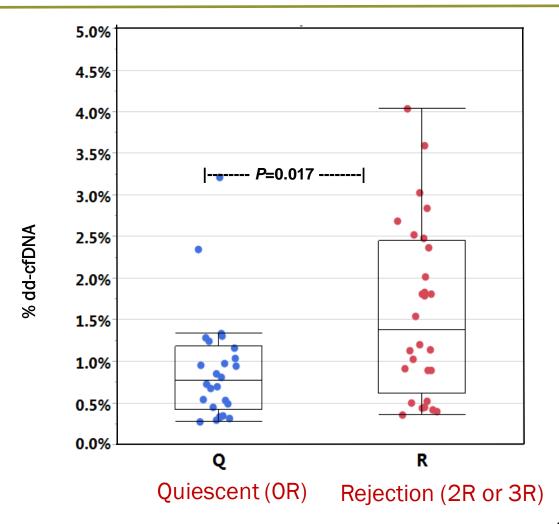
#### Selection for cfDNA Analysis

- blood drawn prior to biopsy
- no rejection treatment
- steroid dose < 20 mg
- at least 2 preceding samples available
- patients matched with the R set for race, age

N=26 patients



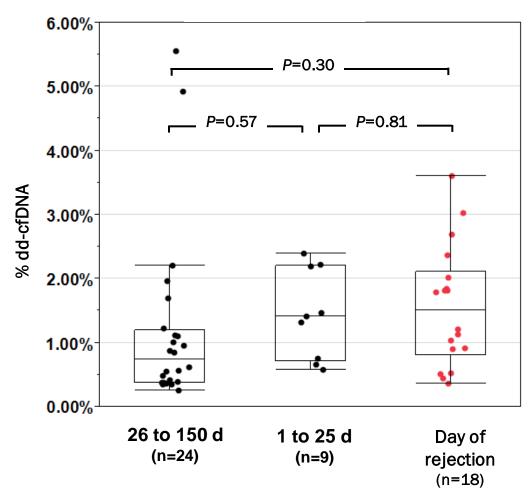
# **Increased dd-cfDNA Levels Correlate with Acute Rejection in Heart Transplant Recipients**



Crespo-Leiro, ISHLT 2015



# dd-cfDNA tends to increase within one month prior to acute rejection

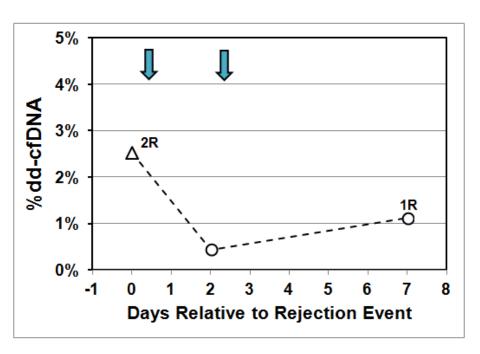


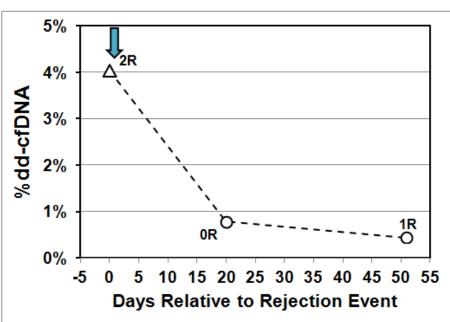
Days prior to rejection



#### dd-cfDNA Levels Decrease Following Rejection Treatment

 $\mathbf{J}$  = Rejection treatment







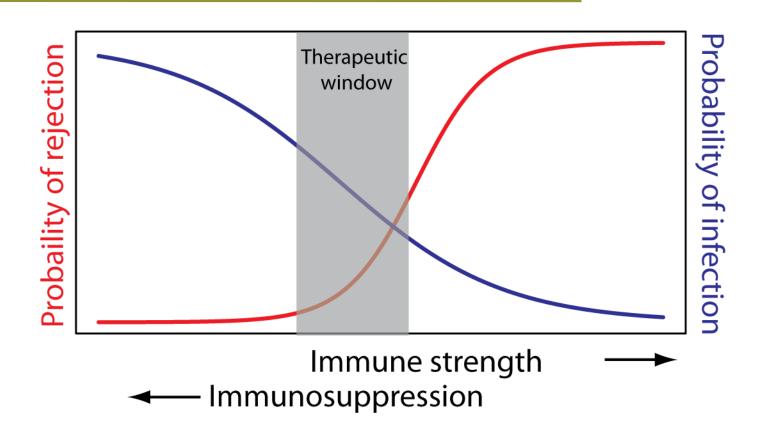


#### **Conclusions**

- Donor-derived cell-free DNA is an informative pan-organ noninvasive marker of acute rejection after solid organ transplantation.
- Dd-cfDNA may enable clinicians to non-invasively distinguish acute rejection from other post-transplant complications
- Measurement of serial dd-cfDNA levels may permit earlier detection of acute rejection, before graft damage/dysfunction occurs.
- dd-cfDNA levels reliably fall after treatment of acute rejection
- Early measurements of dd-cfDNA may identify transplant recipients at risk of chronic graft injury



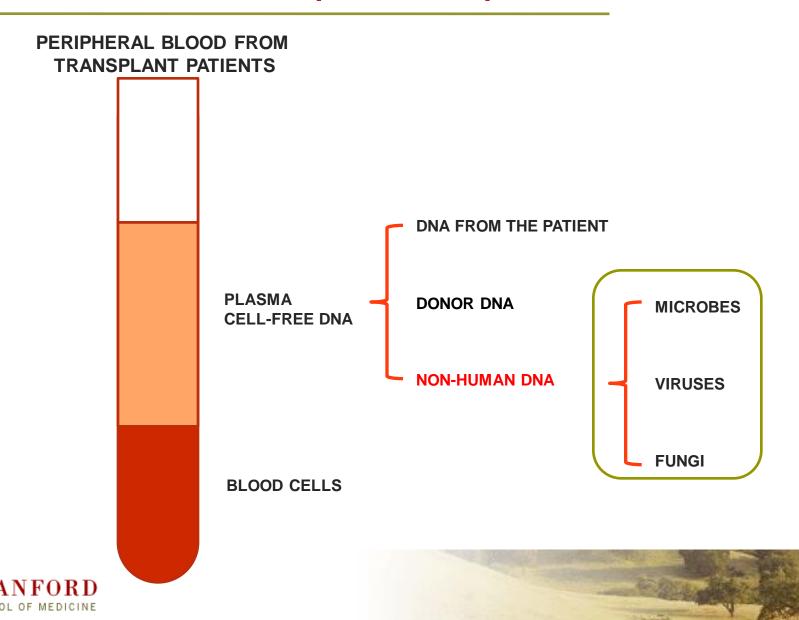
# The challenge of post-transplant therapy



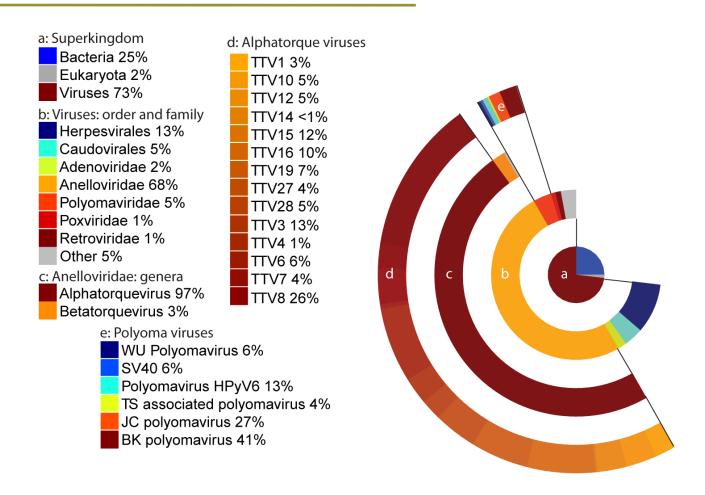
The therapeutic window is narrow, and can vary between patients. Sometimes rejection and infection can present in similar ways.



## Non-human DNA is also present in plasma



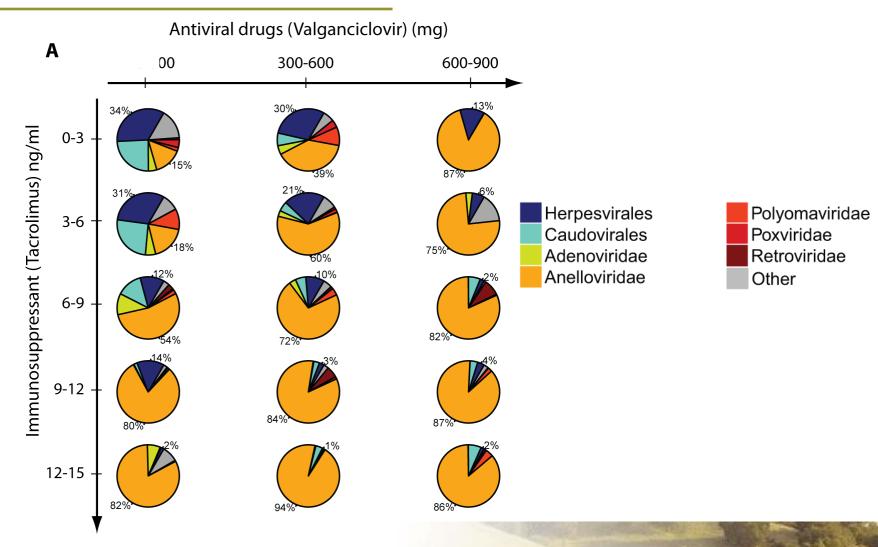
### Relative genomic abundance



The anelloviridae fraction is primarily composed of viruses from the alphatorque genus.

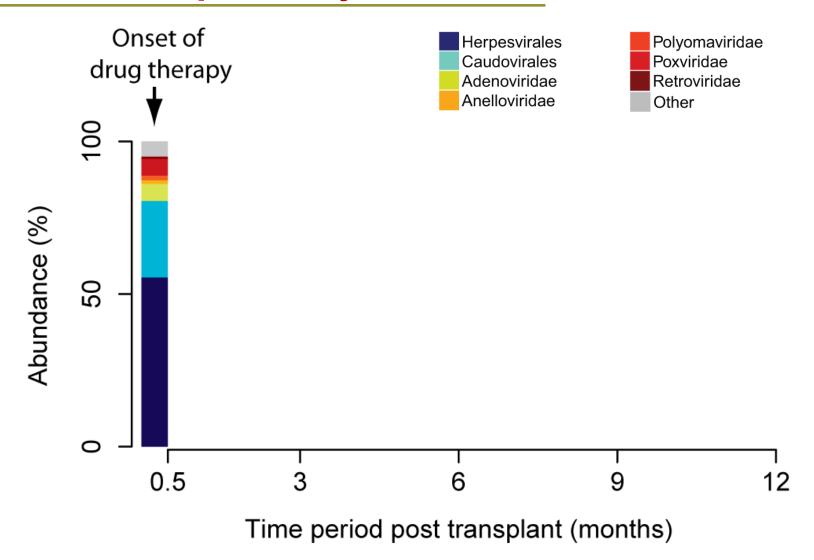


# Immunosuppressants and antivirals alter structure of the virome



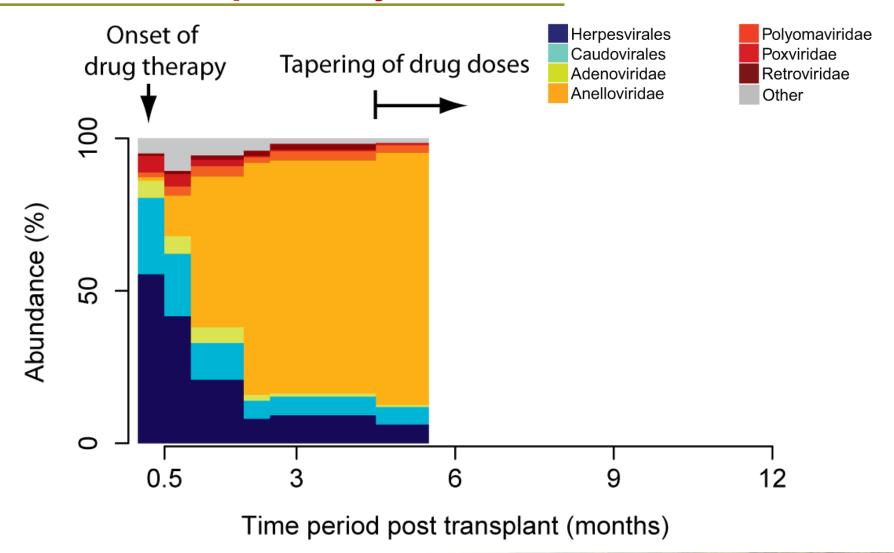


# Virome temporal dynamics



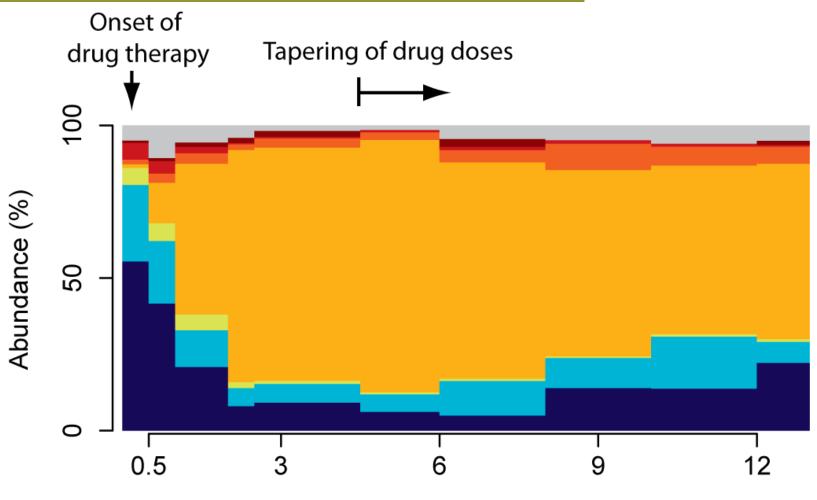


# Virome temporal dynamics





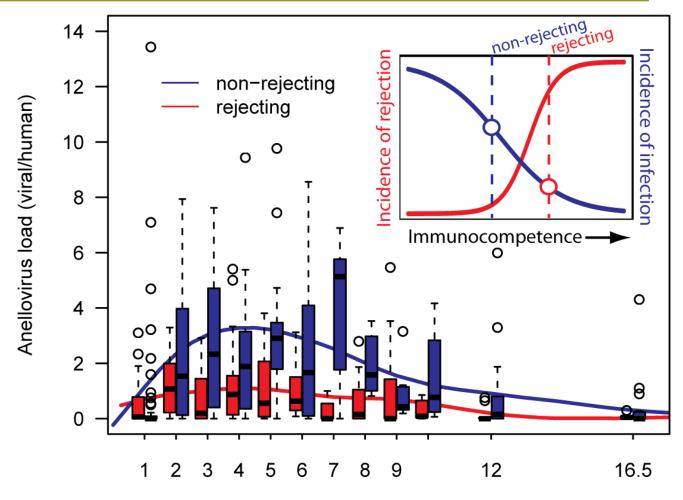
# Virome temporal dynamics

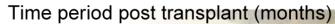






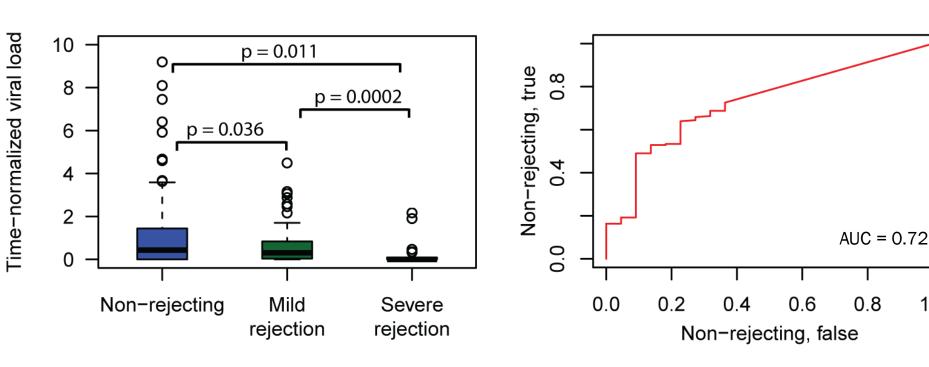
# Anellovirus load for rejecting vs non-rejecting recipients







# Anellovirus load for rejecting vs nonrejecting recipients

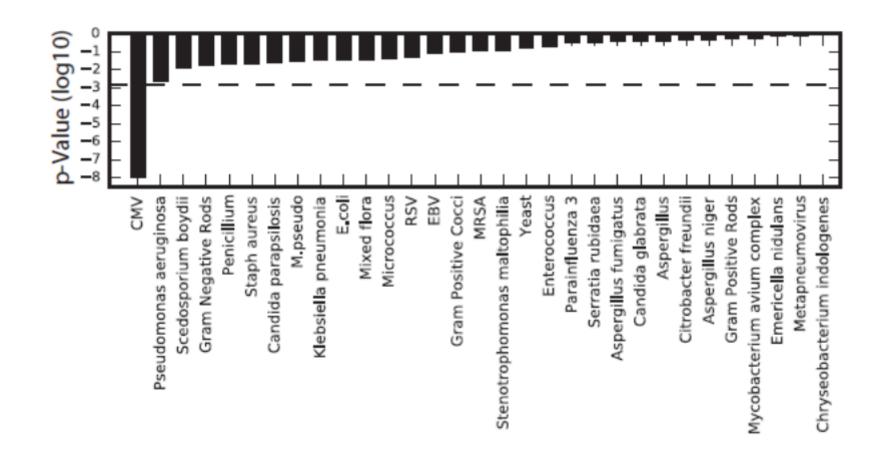


Can anellovirus load be used as a marker of a patient's net state of immunosuppression?

1.0

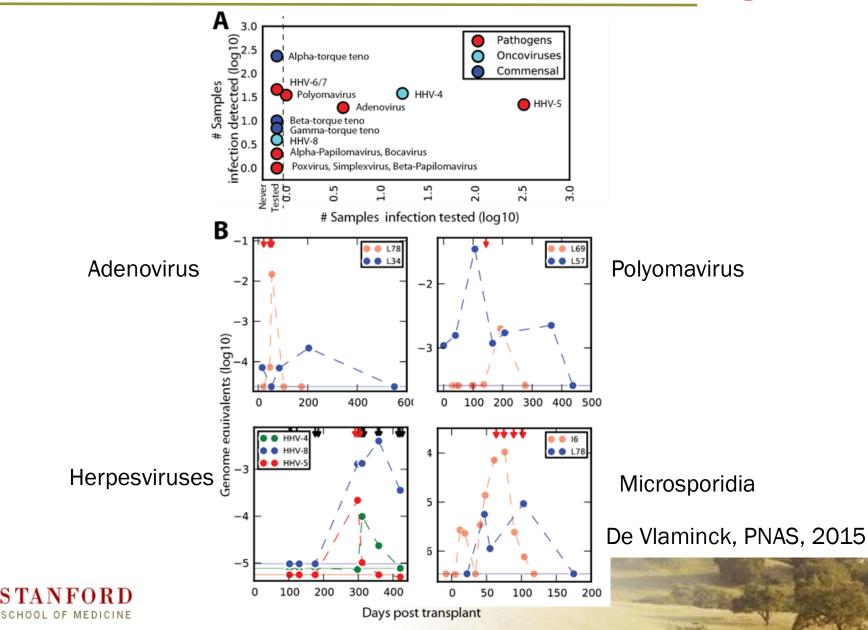


# Correlation between shotgun sequencing and clinical lab results





# Non-Biased Detection of Specific Pathogens



## **Conclusions**

- cfDNA sequencing can be used to study the microbiome, and changes over time
- Structure of the virome is strongly affected by immune modulation and antivirals.
- The total viral load increases markedly at the onset of immunosuppressive therapy.
- Anellovirus load allows stratification of rejecting and non-rejecting recipients.
- Non-biased sequencing of the virome may enable diagnosis of infectious complications



#### **Future Directions**

- D-OAR: Prospective registry study of AlloSure™ assay to study test performance
  - 23 heart transplant centers
  - ~700 study subjects and ~2500 samples collected as of this week
  - Transition from research-grade to clinical-grade testing will facilitate adoption for patient management





#### **Future Directions**

- Mitochondrial cfDNA as a marker of acute rejection after transplantation (DeVlamick, Cornell)
- Other sample types
  - Fingerprick blood samples point of care diagnostics
  - cfDNA in urine to monitor infections and rejection
- Identifying the tissues of origin of cell-free DNA
  - Genome-wide methylation patterns
     K. Sun, D. Lo, PNAS, 2015
  - Patterns of nucleosome and transcription factor occupancy
     M. Snyder, J. Shendure, Cell, 2016





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