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TRANSPLANT SUMMIT 2019

NO SIZE FITS ALL: Uncovering the Potential of Personalized Transplantation

Disclosure

CareDx, Inc: scientific advisor, speakers' bureau, research grant recipient, steering committee



Learning Objectives

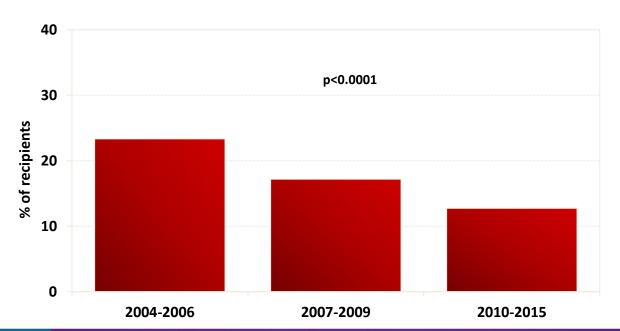
- To appreciate the need for non-invasive biomarkers for surveillance of graft health
- To understand the principles of cell-free DNA testing
- To review major studies to-date of donor-derived cell free DNA testing for acute rejection monitoring after heart transplantation
 - Stanford shotgun sequencing (GTD- Genome Transplant Dynamics)
 - Targeted sequencing
 - 。 AlloSure®
 - 。 myTAI_{HEART}
 - One genome method





Need for acute rejection surveillance

Adult Heart Transplants
% of Recipients Experiencing <u>Treated</u> Rejection Between Transplant Discharge
and 1-Year Follow-Up by Era



Treated rejection = Recipient was reported to (1) have at least one acute rejection episode that was treated with an anti-rejection agent; or (2) have been hospitalized for rejection.





Non-Invasive Markers of Rejection

- Electrocardiogram
 - Surface
 - Intramyocardial
- Cardiac Imaging
 - Echocardiography
 - Magnetic Resonance Imaging
 - Nuclear Imaging

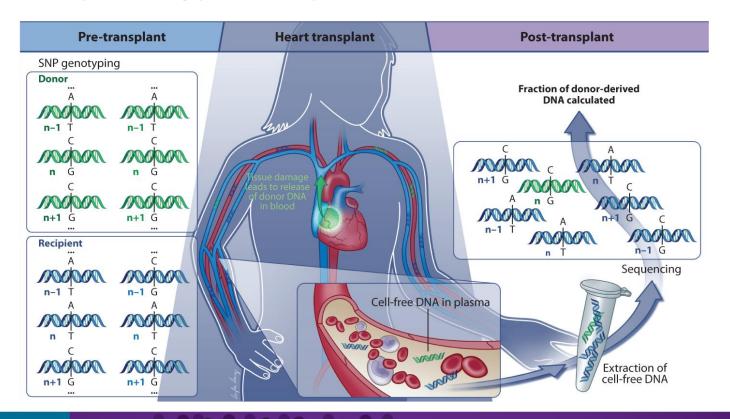
Biomarkers

- B-type Natriuretic Peptide (BNP)
- Troponin (Tnl, TnT)
- High-sensitivity CRP

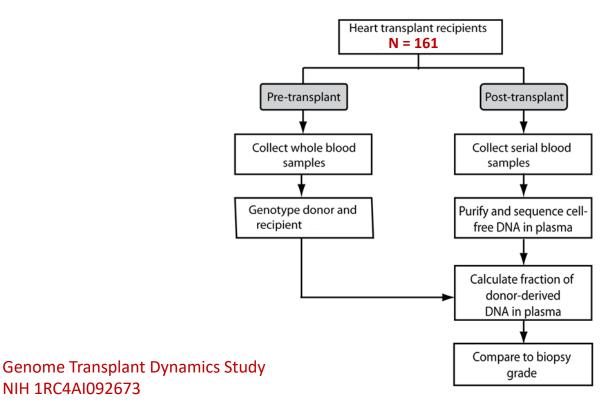
- Genomic Markers
 - Gene Expression Profiling
 - Cell-free DNA
 - mRNA/miRNA
 - Proteomics



Cell-Free DNA: a promising post-transplant biomarker

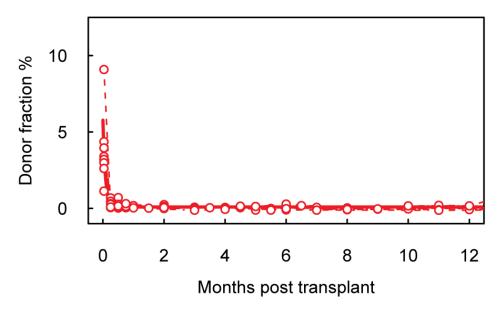


cfDNA prospective study design



NIH 1RC4AI092673

dd-cfDNA signal in absence of rejection

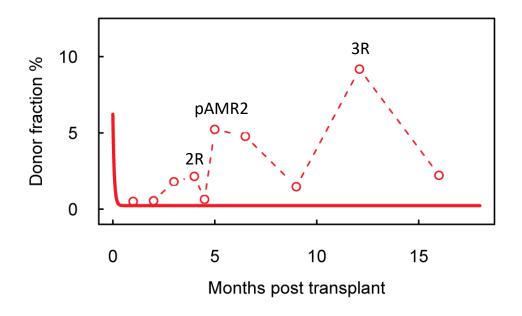


Elevated signal immediately post transplant followed by a quick decay (2.4 days) to a low baseline level





dd-cfDNA signal at time of acute rejection

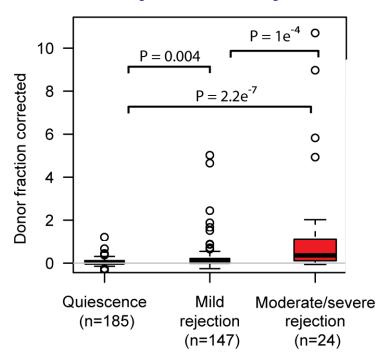


Elevated donor DNA at time of rejection





Comparison of dd-cfDNA levels: no rejection vs rejection

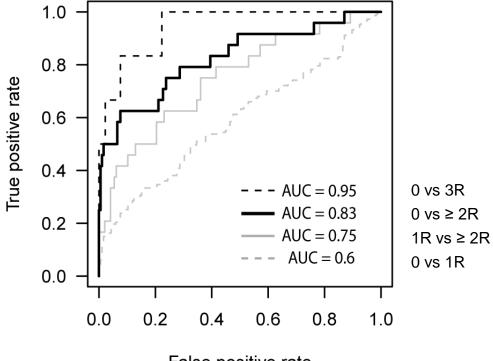


Significant increase in fraction of donor-derived DNA at rejection





Test performance: Detection of acute rejection



False positive rate





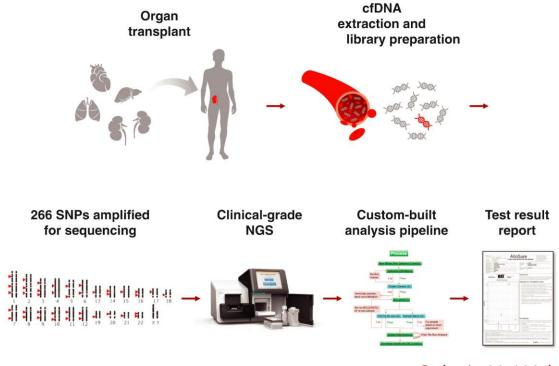
dd-cfDNA: A Rapidly Evolving Technology

- Technology used in previous studies to measure SNP alleles:
 - Shotgun sequencing methods (Stanford) (1)
 - Targeted amplification (Wisconsin) (2)
 - Both requiring recipient AND donor genotypes
- New approaches (AlloSure, myTAI_{HEART}, one genome method)
 have been developed to discriminate donor from recipient after
 sequencing cfDNA from a recipient blood sample only (3, 4)
 - (1) Snyder et al., PNAS, 2011 De Vlaminck et al., Sci Transl Med, 2014
 - (2) Beck et al., Clin Chem, 2013 Hidestrand et al., JACC, 2014
 - (3) Sharon E et al. PLoS Comput Biol, 2017
 - (4) Grskovic et al, J Mol Diagnostics, 2016





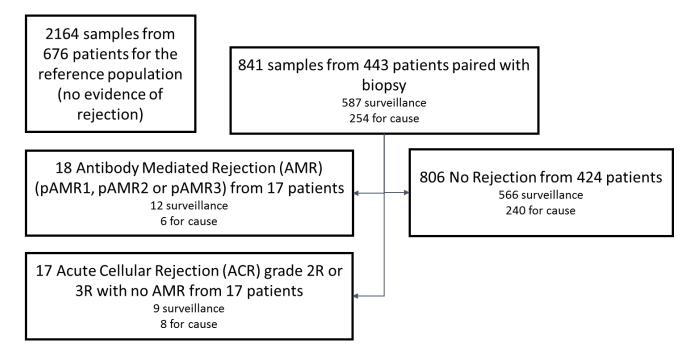
AlloSure® dd-cfDNA assay



Grskovic, M. J Molecular Diagnostics, 2016



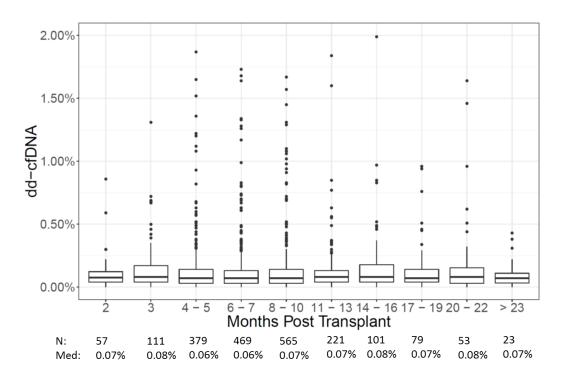
Prospective AlloSure® study in heart transplantation: D-OAR Registry







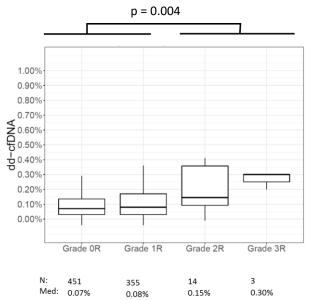
D-OAR Study: dd-cfDNA levels post-transplant (no rejection)

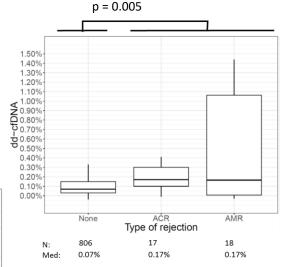


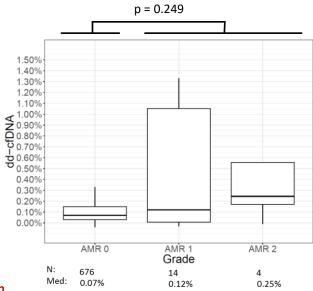




D-OAR Study Results

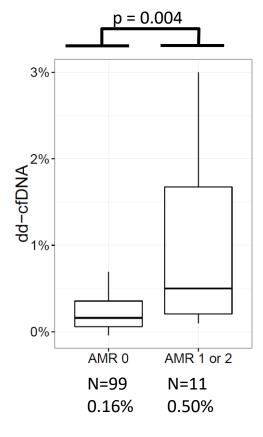








D-OAR Study Results: Cedars Sinai cohort

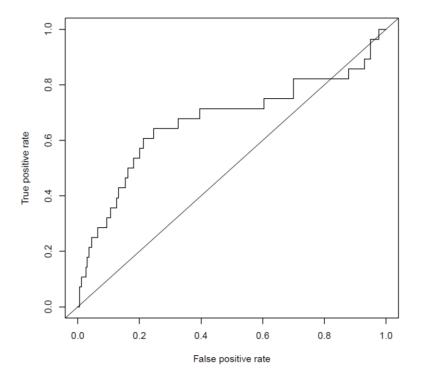


33 patients, 110 samples





AlloSure test performance



Threshold 0.2%

AUC 0.64

Sensitivity 44%

Specificity 80%

PPV 9%

NPV 97%





Now Offering... my TA

- First non-invasive, cell-free DNA based test for increased risk of moderate or higher acute cellular rejection in heart transplant patients
- Rapid, cost-effective, clinically validated laboratory developed test
- Can be used in patients as young as 2 months of age and as early as
 7 days post-transplant
- Results available next business day after receipt
- CLIA Certified / CAP Accredited



myTAI_{HEART}

- Targeted sequencing "relies on selected highly-informative genomic regions" [Digital Analysis of Selected Regions (DANSR)]
- Method 1: donor and recipient genotyped
- Method 2: recipient genotyping only
- 88 pediatric heart transplant patients, mean age 13 years (0.1-30 years)
- 158 blood samples paired with biopsy

Hidestrand M, et al. JACC, 2014 Ragalie W, et al. JACC, 2018





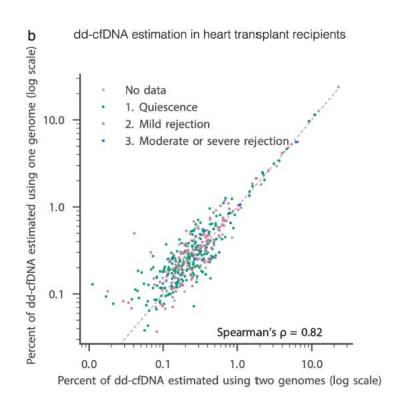
$\mathbf{myTAI}_{\mathsf{HEART}}$

| Biopsy Grade | N | % dd-cfDNA (threshold 0.2%) | |
|--------------|-----|-----------------------------|----------|
| | | Method 1 | Method 2 |
| OR | 134 | 0.11% | 0.25% |
| 1R | 21 | 0.37% | 0.89% |
| 2R | 3 | 0.97% | 1.22% |
| 3R | 0 | - | - |

| Method | Comparison | P-value | AUC |
|--------|-------------|---------|------|
| 1 | OR vs 1R/2R | 0.02 | 0.78 |
| 2 | OR vs 1R/2R | <0.001 | 0.84 |

"One genome" method

- Shotgun sequencing of total cfDNA
- Computational approach to estimate dd-cfDNA levels in the absence of a donor genotype



Sharon E, et al. PLOS Computational Biology, 2017





SHORE Study: Surveillance HeartCare Outcomes Registry

HeartCare = AlloMap + AlloSure

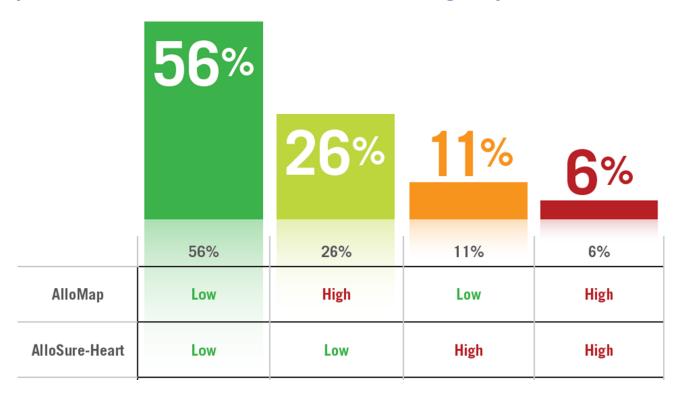


Hypothesis:

- AlloMap (gene expression test) detects immune activation and is clinically validated for ACR monitoring
- AlloSure (cfDNA assay) detects graft injury and is validated for ACR and AMR monitoring
- Perhaps a COMBINED AlloMap + AlloSure monitoring approach will be better than either test alone



SHORE Study: Surveillance HeartCare Outcomes Registry



SHORE study design

- 5 year study
- 35 sites in US
- 1,600 patients
- **Primary objective**: To assess the clinical utility of surveillance using HeartCare testing, in association with the clinical care of heart transplant recipients
- **Endpoints**: deaths, number of biopsies, number of rejection events, measures of graft function



Heart Allograft Routine Testing Schedule (HARTS)

A blood test can be administered more frequently and conveniently than an invasive procedure, providing for more continuous rejection surveillance.











(Months Post-Transplant)

Courtesy of CareDx, Inc.





Conclusions: dd-cfDNA for heart transplant monitoring

- We have come a long way in the quest for non-invasive ways to assess graft health after transplant
- Fewer biopsies are being performed than ever before
- Fewer procedural complications, more satisfied patients
- AlloSure has been tested in prospective registry study (D-OAR)
 - Detects both ACR and AMR at a threshold of 0.2%
- Upcoming SHORE study (CareDx): Combined AlloMap + AlloSure for heart transplant monitoring
- MyTAI_{HEART} is available for use (pediatric patients)
- "One genome method" is being studied for clinical use





THANK YOU



