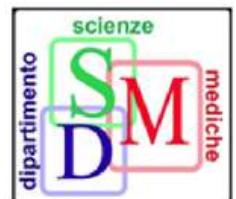


Il DNA libero circolante di origine del donatore: uno strumento diagnostico nel trapianto di organo solido



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*Il sottoscritto Silvia Deaglio
in qualità di relatore al*

**XXX CONGRESSO NAZIONALE AIBT
NAPOLI, 10/12 OTTOBRE 2024**

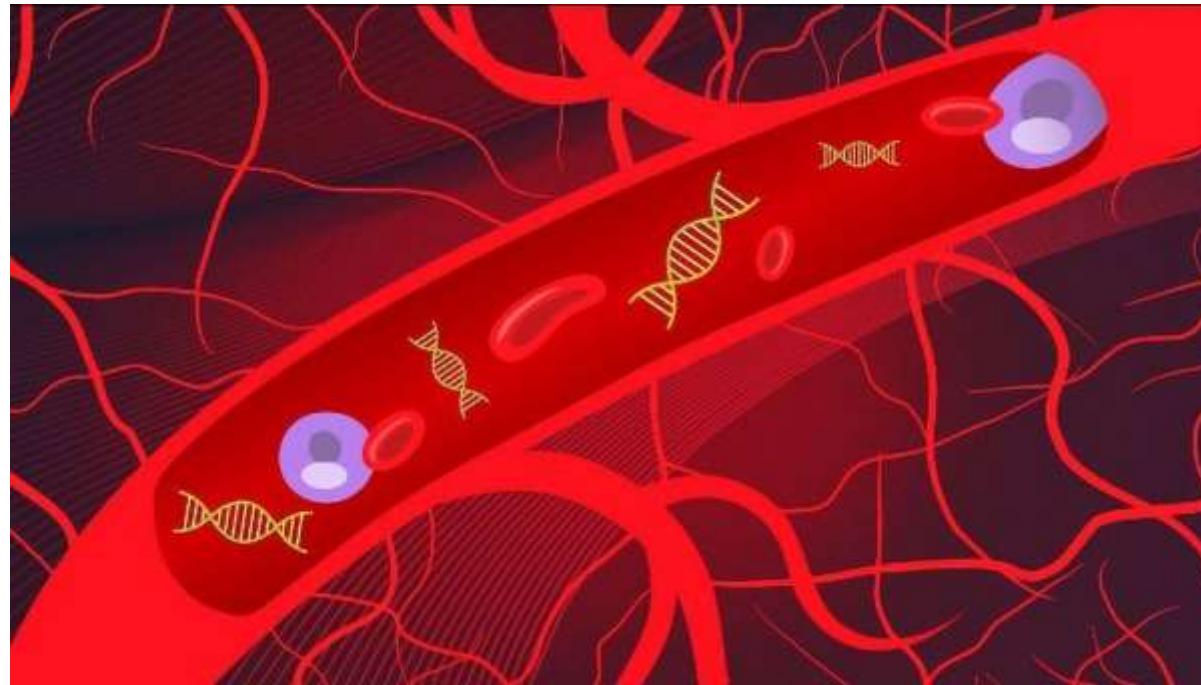
*ai sensi dell'art. 3.3 sul Conflitto di Interessi, pag. 18,19
dell'Accordo Stato-Regione del 19 aprile 2012, per conto di
Planning Congressi srl*

dichiara

*che negli ultimi due anni ha avuto rapporti diretti di
finanziamento con i seguenti soggetti portatori di interessi
commerciali in campo sanitario:*

- Astra Zeneca (sponsored research agreement)
- Heidelberg Pharma (sponsored research agreement)
- Solve Therapeutics (sponsored research agreement)

Cos'è il cfDNA?



- ✓ Piccoli frammenti di DNA che si trovano nel plasma di ciascuno di noi
- ✓ Tre ambiti di applicazione in medicina:
 - ambito oncologico
 - ambito prenatale
 - ambito trapiantologico

Come si usa oggi il

cfDNA?

✓ In ambito oncologico il ctDNA serve per:

- diagnosi precoce
- monitoraggio effetto terapia
- monitoraggio comparsa resistenza alla malattia
- MRD

LIMITARE LE BIOPSIE TISSUTALI / ASPIRATI MIDOLLARI

✓ In ambito di diagnosi prenatale il cfDNA serve per:

- stimare il rischio di alterazioni citogenetiche con un buon livello di accuratezza
- determinare il sesso del feto

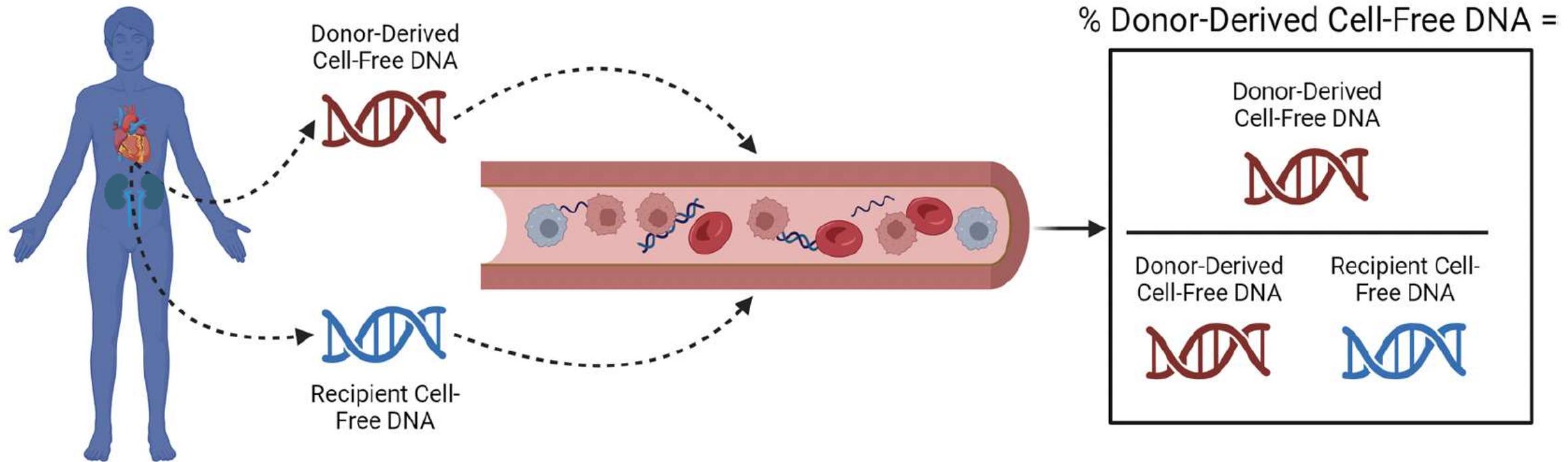
LIMITARE LA DIAGNOSI PRENATALE INVASIVA

✓ In ambito trapiantologico il dd-cfDNA serve per:

- diagnosticare il rigetto/danno d'organo
- monitorare la risposta alla terapia degli episodi di rigetto

LIMITARE LE BIOPSIE DELL'ORGANO TRAPIANTATO

Il cfDNA nel contesto del trapianto



Le pubblicazioni sul tema dd-cfDNA



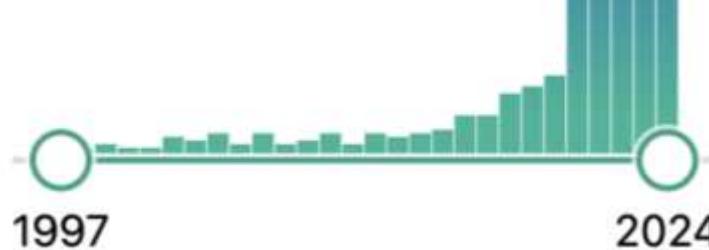
donor derived cell free dna AND transplantation

Advanced Create alert Create RSS

RESULTS BY YEAR



417



RESULTS BY YEAR



205 (49%)

1999

2024

RESULTS BY YEAR



96 (23%)

2013

2024

RESULTS BY YEAR



71 (17%)

2004

2024

RESULTS BY YEAR

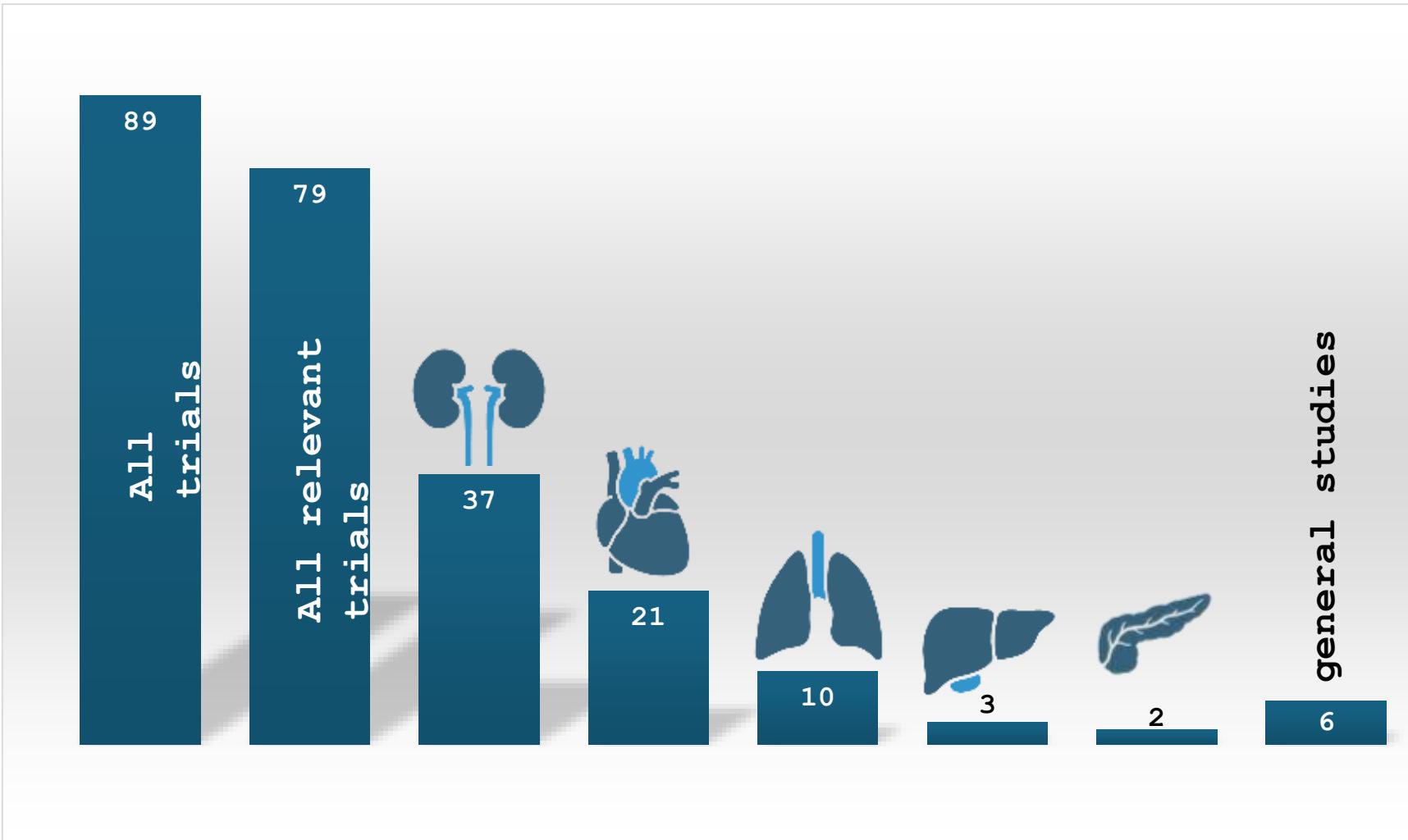


38 (9%)

2004

2024

Sperimentazioni cliniche sul dd-cfDNA



33 studi stanno reclutando o cominceranno a reclutare

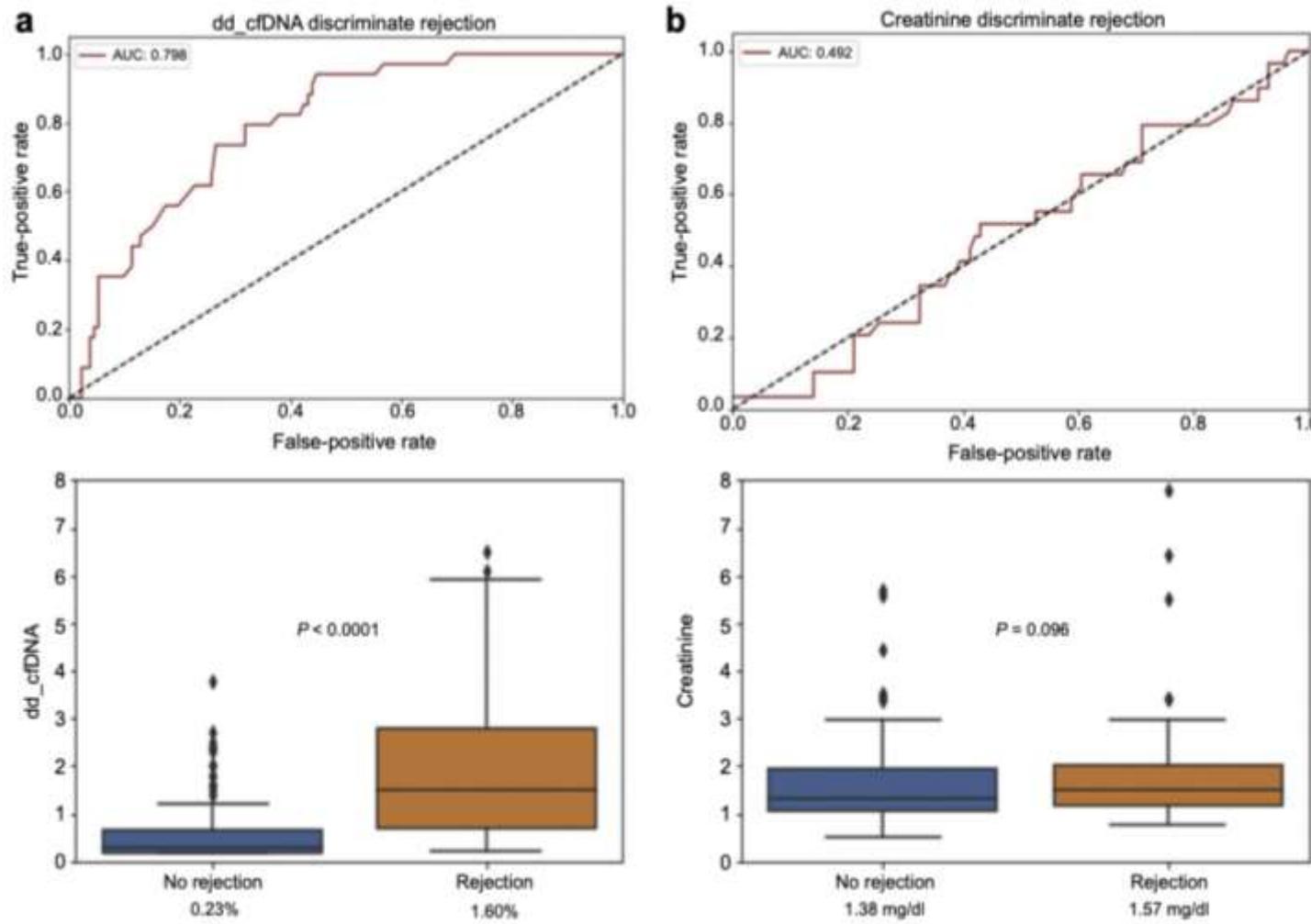
Il dd-cfDNA nel monitoraggio del rigetto acuto nel trapianto di rene

TABLE 1 | Summary of the evidence reviewed on donor-derived cell-free DNA for clinical acute rejection.

Authors	Study design	Number of samples	Results
[9]	Prospective observational, Multicenter	107 biopsies 27 rejections 80 no rejection	For any rejection (AUROC 0.74, PPV 0.61, NPV 0.84)
[10]	Subgroup analysis of prospective observational, Multicenter	87 patients 16 ABMR 53 no rejection	For ABMR with + DSA and dd-cfDNA >1%, (AUROC 0.86, PPV 0.81, NPV 0.83)
[2]	Retrospective analysis of biorepository samples, Single center	217 biopsies 38 rejections 179 no rejection	For any rejection for cause + SubAR (AUROC 0.87, PPV 0.52, NPV 0.95)
[11]	Prospective observational, Single center	63 biopsies 34 rejections 29 no rejection	For any rejection (AUROC 0.71, PPV 0.77, NPV 0.75)
[12]	Prospective observational, Single center	189 patients 22 rejections 395 stable samples	For any rejection (Absolute concentration of dd-cfDNA (AUROC 0.83) is better than dd-cfDNA (%) (AUROC 0.73)
[13]	Prospective cross-sectional, Multicenter ($n = 2$)	61 biopsies 13 ABMR 48 no rejection	For ABMR (absolute concentration AUROC 0.91 vs. dd-cfDNA (%) 0.89)
[14]	Subgroup analysis of prospective observational, Multicenter	79 patients with TCMR 1A/borderline changes	Subjects with TCMR (1A and borderline) with high dd-cfDNA had worse clinical outcomes compared to those with low dd-cfDNA
[15]	Cross-sectional for DSA screening/Retrospective testing of dd-cfDNA on bio-banked samples, Single center	From 2 independent cohort 45/30 biopsies 25/17 ABMR 20/13 no ABMR	For ABMR with +DSA AUROC for dd-cfDNA alone 0.89/0.69 AUROC for DSA alone 0.88/0.77
[3]	Prospective observational, multicenter (ADMIRAL)	219 biopsies 113 rejections 106 no rejection	For any rejection dd-cfDNA (AUROC 0.8, PPV 0.5, NPV 0.9)
[16]	Prospective observational, Single center	208 biopsies 162 rejections by histology 46 no rejection by histology	For any rejection dd-cfDNA and MMDx (AUROC 0.80), dd-cfDNA and histology (AUROC 0.75)
[17]	Prospective observational, multicenter (TRIFECTA)	300 biopsies 120 rejections 180 no rejection	dd-cfDNA levels are strongly associated with the active molecular rejection phenotype (MMDx), particularly with AMR, mixed, and active TCMR
[18]	Prospective observational, multicenter (TRIFECTA)	367 biopsies 83 (histology test set) rejection 71 (MMDx test set) rejection	Any rejection prediction AUROC in test set by logistic regression model using both dd-cfDNA (%) and absolute concentration <ul style="list-style-type: none">• 0.88 for MMDx• 0.82 for histologic rejection

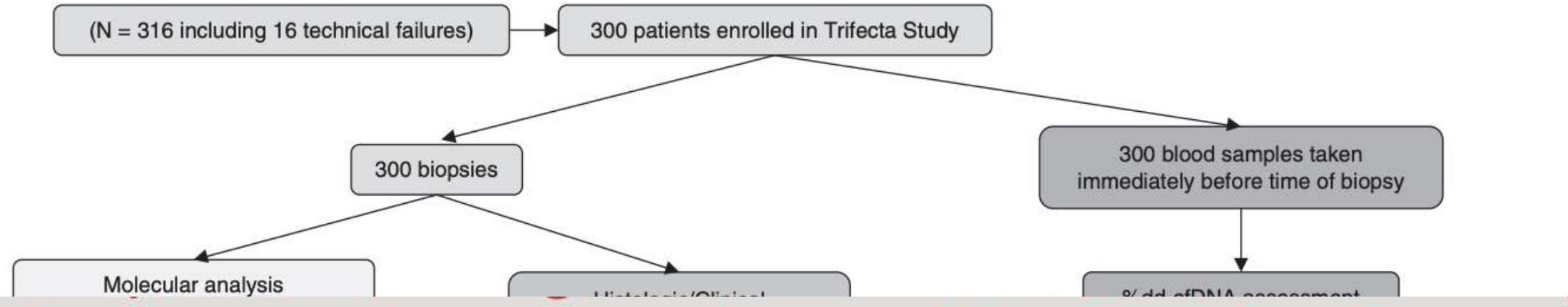
AUROC, area under the receiver operating characteristic curve; DSA, donor-specific antibody; MMDx, the molecular microscope diagnostic system; NPV, negative predictive value; PPV, positive predictive value.

Il ruolo del dd-cfDNA nel trapianto di rene: lo studio ADMIRAL

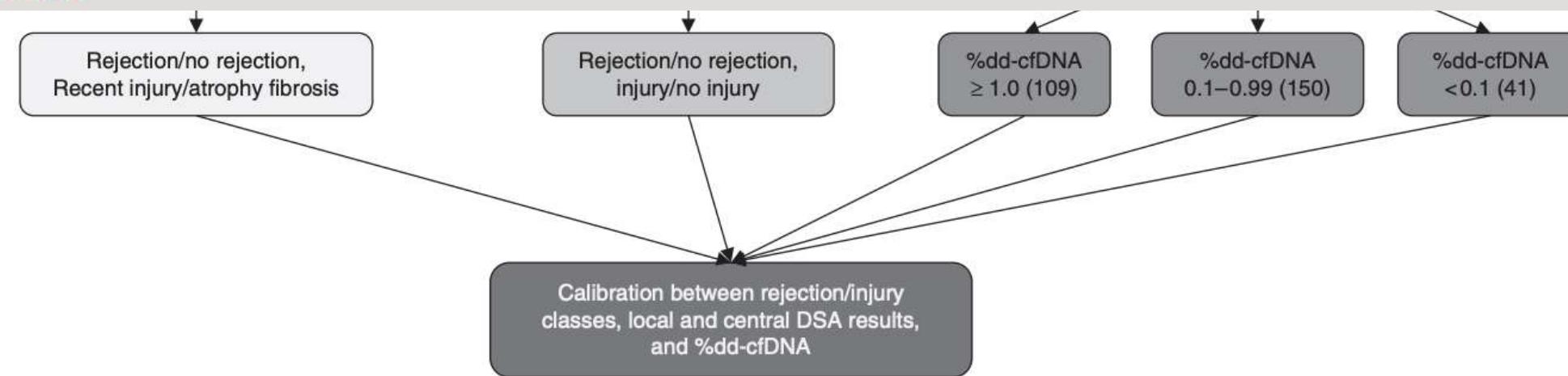


- ✓ 1092 kidney transplant recipients monitored for dd-cfDNA over a three-year period
- ✓ association of dd-cfDNA with histologic evidence of allograft rejection
- ✓ **persistently low dd-cfDNA levels may accurately identify allograft quiescence or absence of injury**

Il ruolo del dd-cfDNA nel trapianto di rene: lo studio TRIFECTA



These findings indicate that plasma dd-cfDNA levels are strongly related to the active molecular rejection processes in indication biopsies.





Cell-free DNA for the detection of kidney allograft rejection

- ✓ To assess the association of dd-cfDNA with rejection, disease activity and severity
- ✓ To assess its independent and added value to detect rejection beyond standard of care monitoring

Il disegno dello studio

Cohort: 3,733 kidney transplant evaluations between 2013 and 2022

French development cohort : 1,415 evaluations

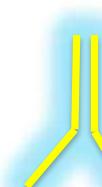
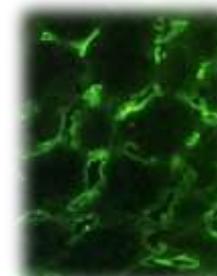
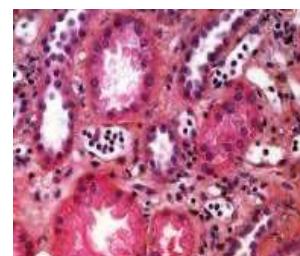
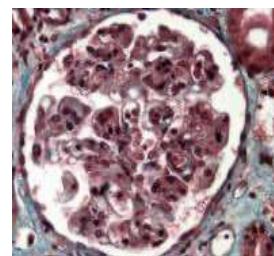
Validation cohort : 2,318 evaluations in the US (11 centers) and Belgium (1 center)

Evaluations
(n=3,733)



- Baseline characteristics
- Renal function + Proteinuria (P/C)
- HLA DSA at the time of TX
- HLA DSA at the time of biopsy
- Histology :
 - Banff 2019 scores
 - Diagnoses
- **dd-cfDNA (AlloSeq and AlloSure)**

Rejection
Banff scores



Aubert et al, Nature Medicine. 2024

Il dd-cfDNA mostra significativa associazione con il rigetto

Derivation cohort

		Number of biopsies	Number of events	OR	95% CI	p	OR 95% OR bootstrap BCA
eGFR (mL/min/1.73 m ²)		1,395	226	0.989	(0.979-0.999)	0.033	(0.979-1.000)
Proteinuria (g/g) (log transformation)		1,395	226	1.180	(1.031-1.350)	0.016	(1.029-1.350)
Kidney graft instability*	No Yes	1,170 225	159 67	1 1.768	- (1.144-2.711)	0.010	- (1.102-2.894)
Previous episode of Rejection	No Yes	1,300 95	178 48	1 4.756	- (2.852-7.920)	4.754e-09	- (2.781-7.827)
Anti-HLA DSA MFI	<500 ≥500 – 3,000 ≥3000 – 6,000 ≥6,000	859 424 54 58	76 93 20 37	1 3.081 3.310 6.338	- (2.152-4.429) (1.618-6.597) (3.186-12.791)		- (2.062-4.496) (1.726-6.607) (3.410-12.258)
dd-cfDNA (log transformation)		1,395	226	2.275	(1.902-2.739)	<2.2e-16	(1.805-2.801)

Il dd-cfDNA mostra significativa associazione con il rigetto

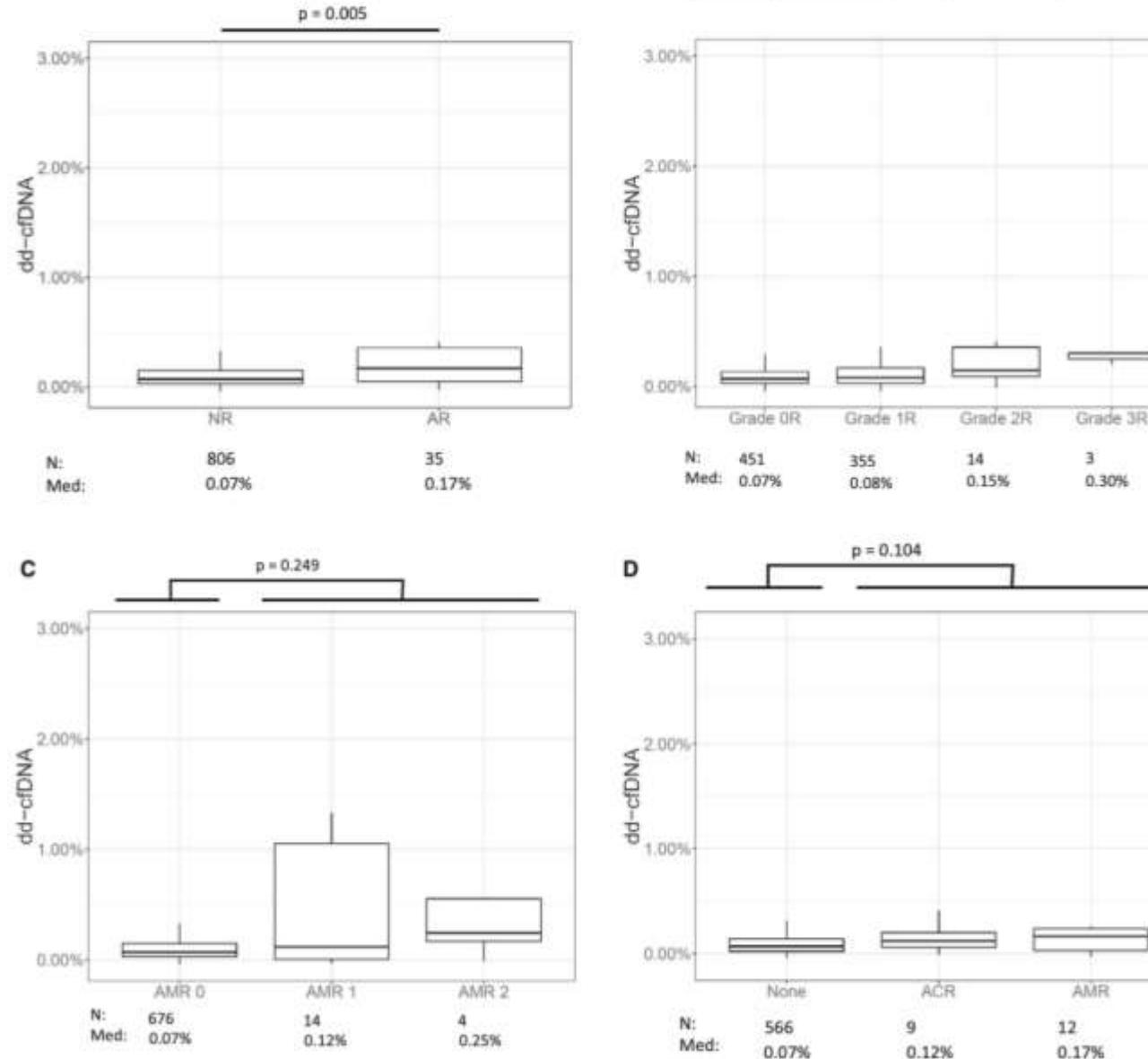
Validation cohort

		Number of biopsies	OR	95% CI	p
eGFR (mL/min/1.73 m ²)		1,911	0.990	(0.984-0.996)	0.001
Proteinuria	Absence Presence	1,341 570	1 1.468	- (1.134-1.900)	0.016
Kidney graft instability	No Yes	1,505 406	1 1.397	- (1.028 – 1.896)	0.004
Previous episode of rejection	No Yes	1,538 373	1 2.661	- (2.016-3.516)	5.17e-12
Anti-HLA DSA	Absence Presence	1,498 413	1 3.134	- (2.402-4.096)	< 2e-16
dd-cfDNA (log transformation)		1,911	2.317	(2.083-2.588)	< 2e-16

Il ruolo del dd-cfDNA nel trapianto di cuore

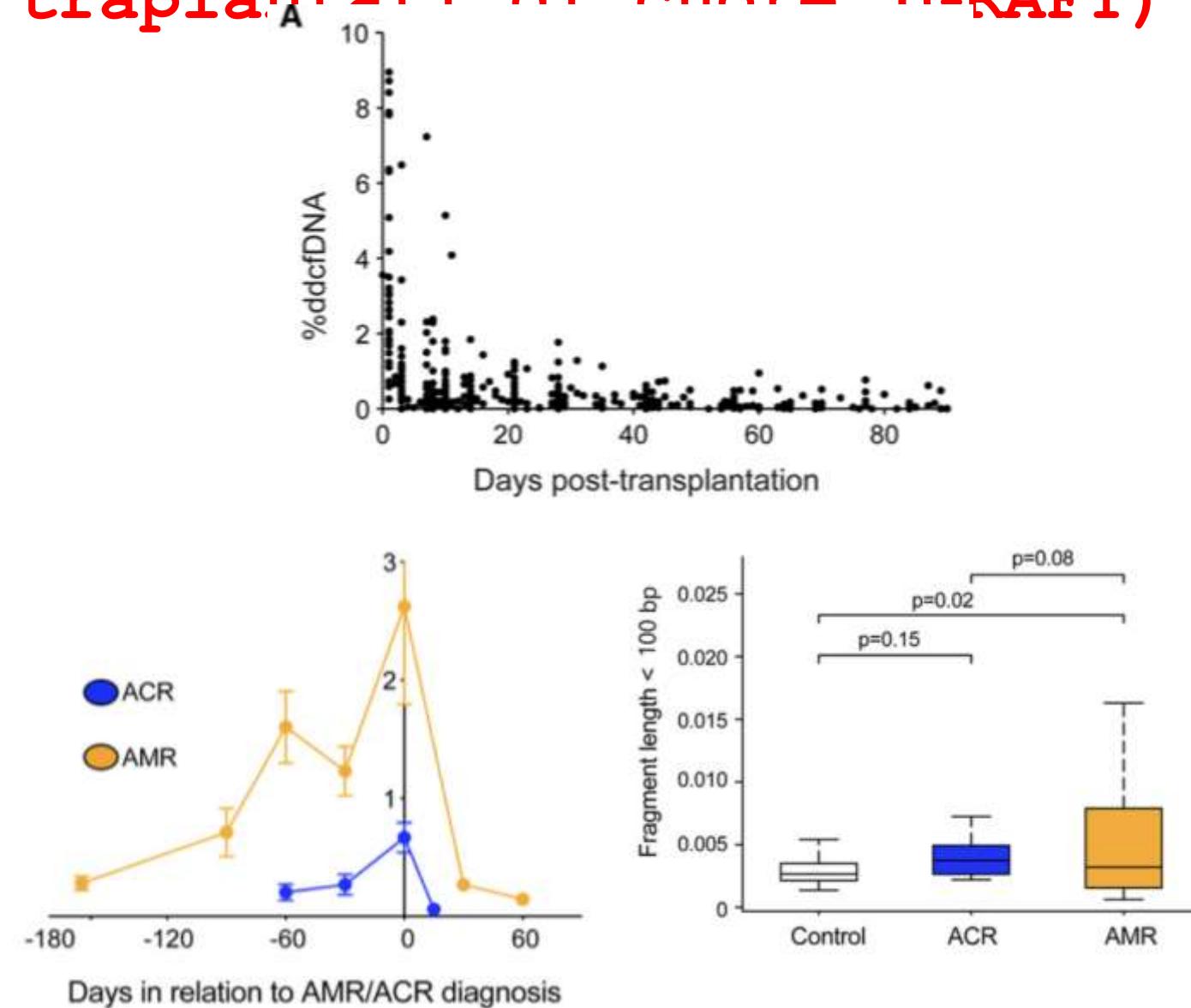
Study Name, Year of Publication	Study Design	Key Study Findings
dd-cfDNA studies		
Circulating Cell-Free DNA Enables Noninvasive Diagnosis of Heart Transplant Rejection, 2014 ¹³	<ul style="list-style-type: none"> Prospective study on Genome Transplant Dynamics (GTD) using SNV genotyping 65 adult and pediatric HT recipients with 565 longitudinally obtained plasma samples compared with EMB 	<ul style="list-style-type: none"> Established rapid clearance of dd-cfDNA after HT and a stable baseline allowing for implementation of a time-independent threshold for the diagnosis of AR when collected more than 2 wks after HT dd-cfDNA detected both ACR and AMR, and levels correlated with severity ROC analysis for detection of ACR (≥ 2R/3A or AMR) by dd-cfDNA revealed AUC of 0.83 (sens = 0.58, spec = 0.93 at a dd-cfDNA threshold level of 0.25%) Findings indicate that dd-cfDNA could replace EMB
Donor-Derived Cell-Free DNA- Outcomes AlloMap Registry (D-OAR), 2019 ¹⁴	<ul style="list-style-type: none"> Prospective observational cohort of 740 HT recipients from 26 U.S. centers aged ≥ 15 y and > 55 d post-HT who were undergoing AlloMap GEP and AlloSure dd-cfDNA testing for surveillance, plus a single-center cohort of 33 patients at high risk for AMR. Study period: 2014-2018 	<ul style="list-style-type: none"> Median dd-cfDNA was 0.07% in reference HT recipients (2,164 samples) and 0.17% with acute rejection (35 samples; $P = 0.005$) At a 0.2% threshold, dd-cfDNA had a 44% sensitivity to detect rejection and a 97% NPV In the cohort at risk for AMR (110 samples from 33 patients), dd-cfDNA levels were elevated 3-fold in AMR compared with patients without AMR (99 samples; $P = 0.004$)
Genomic Research Alliance for Transplantation (GRAFT), 2021 ¹⁵	<ul style="list-style-type: none"> Prospective observational cohort study of HT recipients aged ≥ 18 y from 5 U.S. centers Aimed to validate the test characteristics of dd-cfDNA for acute rejection and to determine the ability to predict long-term outcomes including CAV, graft failure, and mortality dd-cfDNA measured by shotgun sequencing and included donor-recipient-paired genotyping to identify SNVs (research grade assay - not available clinically) Study period: 2015-ongoing 	<ul style="list-style-type: none"> Median dd-cfDNA levels were 0.34% vs 0.04%; $P < 0.006$ for ACR grade ≥ 2 vs ACR grade 1 Median levels were 0.63% vs 0.02%; $P < 0.001$ for AMR 1 versus grade 0 rejection and 1.68% vs 0.63%; $P = 0.039$ for AMR ≥ 2 vs AMR 1 dd-cfDNA levels rose as early as 120 d before acute rejection, and fragment length > 100 bp was associated with AMR diagnosis
A novel donor-derived cell-free DNA assay for the detection of acute rejection in heart transplantation (DEDUCE), 2022 ¹⁶	<ul style="list-style-type: none"> Observational 2 center study with retrospective and prospective component Using the clinically available Prospera dd-cfDNA test (Natera) Study period: 2017-2022 	<ul style="list-style-type: none"> 811 samples from 223 patients with dd-cfDNA testing and contemporaneous EMB with 49 EMBs showing AR in 35 patients dd-cfDNA was significantly higher in AR (median 0.58%; IQR: 0.13%-1.68%) compared with non-AR (median 0.04%, IQR: 0.01%-0.11%, P corrected for multiple comparisons < 0.001) AUC-ROC of 0.86 (95% CI: 0.77-0.96) For dd-cfDNA $\geq 0.15\%$ 78.5% sens (95% CI: 60.7%-96.3%) and 76.9% spec (95% CI: 71.1%-82.7%) for AR PPV 25.1% (95% CI: 18.8%-31.5%) and NPV 97.3% (95% CI: 95.1%-99.5%) NA
Surveillance HeartCare Outcomes Registry (SHORE) [NCT03695601]	<ul style="list-style-type: none"> Prospective observational registry study, HT patients with HeartCare (AlloMap and Allo-Sure) monitoring initiated within 3 mo of HT compared with historical control group not monitored with HeartCare Planned enrollment of 2,300 patients completed (plan to match with 1,150 historical controls) Primary outcome: Percentage of patients alive at 1, 2, and 3 y post-HT Study period: 2018-2024 	

Il dd-cfDNA come marcatore di rigetto nei trapiantati di cuore (D-OAR)



- ✓ 740 pazienti da 26 centri con biopsie seriali
- ✓ La misurazione standardizzata di dd-cfDNA discrimina rigetto acuto da non-rigetto

Il dd-cfDNA come marcatore di rigetto nei trapiantati di cuore (C-RAPT)



- ✓ 171 pazienti
- ✓ Il test è attendibile 4 settimane dopo il trapianto
- ✓ Il dd-cfDNA si alza circa 4 mesi prima dell'evidenza istologica di rigetto
- ✓ La lunghezza dei frammenti di dd-cfDNA è suggestiva del tipo di rigetto
- ✓ Il test può evitare un numero significativo di biopsie protocollari

Come monitoriamo il dd-cfDNA?

NGS-based methods				Non-NGS methods				
Targeted Gene Panels	Selection of informative genes/loci to be sequenced	Oncologic field [Zhang et al. 2020, Tie et al. 2015, Dawson et al. 2013, Gandara et al. 2018]	Prenatal field [Grskovic et al. 2016, Sigdel et al. 2018, Levitsky et al. 2022]	Transplant field	qPCR	Oncologic field [Dauber et al. 2020, García-Fernández et al. 2022, Fernández-Galán et al. 2022]	Prenatal field	Transplant field
NGS sequencing of >200 SNPs vs 50 in/del markers				Highly quantitative PCR conditions, may be multiplexed				

Cosa è disponibile in Europa?

TABLE 2 Commercially-Available GEP and dd-cfDNA Assays

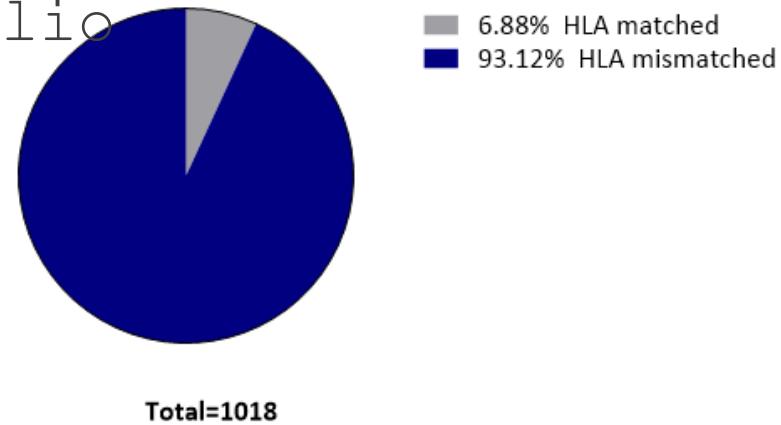
GEP/dd-cfDNA Test Name	Vendor	Availability/Approval	Key Research Studies	Practical Considerations	Notable Features
AlloMap	CareDx, Inc	Available in the U.S. FDA approved since 2008 for use ≥55 d post-HT in recipients ≥15 y CMS approved in the U.S.	CARGO ¹³ IMAGE ¹⁴ eIMAGE ¹⁵ CARGO II ¹⁶ QAR ¹⁷	Minimum needle size for phlebotomy is 22G and collect blood in CPT tubes Avoid transporting sample in pneumatic tubing systems GEP samples need to be processed in a specialized laboratory within 3 h of blood draw Centrifuge and dry ice needed to maintain 6°–37°C	Remote phlebotomy offered by vendor Class IIa recommendation in ISHLT 2010 guidelines for the care of HT recipients ¹⁸ AlloMap results within 50–72 h
AlloSure heart (dd-cfDNA) HeartCare (GEP+dd-cfDNA)	CareDx, Inc	Available in the U.S. CMS approved as HeartCare (combined with GEP AlloMap test) ≥55 d post-HT in patients ≥15 y AlloSure alone is not FDA approved; classified as CAP/CLIA test	D-OAR ¹⁹ SHORE [NCT03695601]	Minimum needle size for phlebotomy is 22G and collect blood in Streck tubes dd-cfDNA must be drawn before EMB or >24–48 h after to avoid false elevations from biopsy-induced myocardial trauma	Remote phlebotomy offered by vendor Plasma dd-cfDNA test performed at a single CareDx CLIA laboratory Analyzes 405 SNVs via NGS HeartCare results within 50–72 h
AlloSeq (dd-cfDNA)	CareDx, Inc	Available outside the U.S., in Europe and additional countries Registered and certified with CE IVD Not paired with GEP testing	NA	dd-cfDNA must be drawn before EMB or >24–48 h after to avoid false elevations from biopsy-induced myocardial trauma	Analyzes 202 SNVs via NGS at local/in-house laboratory (no shipment to CareDx required) Interpretation performed locally Plasma levels of dd-cfDNA and interpretation of results are identical to AlloSure
Prospera Heart (dd-cfDNA)	Natera, Inc	Available in the U.S. Not FDA or CMS approved, classified as CAP/CLIA test	DEDUCE ²⁰ ProTECT [Prospera Test Evaluation in Cardiac Transplant; NCT05205551, enrolling] DETECT [Donor-Derived Cell-free DNA to DETect REjection in Cardiac Transplantation; NCT05081739, planning] Trifecta-Heart [Trifecta-Heart cfDNA-MMDx Study; NCT04707872, enrolling] DEFINE [Development of Non-invasive Cell-free DNA to Supplant Invasive Biopsy in Heart Transplantation; NCT05309382, planning]	dd-cfDNA must be drawn before EMB or >24–48 h after to avoid false elevations from biopsy-induced myocardial trauma Use ≥21G needle and collect blood in Streck tubes	Analyzes 13,292 SNVs Prospera results in 48–72 h
Viracor TRAC (dd-cfDNA)	Eurofins Viracor, Inc	Available in the U.S. only, but the U.S.-based CAP/CLIA lab accepts samples from outside the U.S. Not FDA or CMS approved, classified as CAP/CLIA test	Studied in kidney ²¹ and liver ²² transplant recipients	dd-cfDNA must be drawn before EMB or >24–48 h after to avoid false elevations from biopsy-induced myocardial trauma Use 21G or 22G needle and collect blood in Streck tubes Ambient shipping temperature	Genotyping of recipient only, after which donor genotype is inferred using computational approaches TRAC results within 4–6 business days

CAP = Certified Authorization Professional; CE = Conformité Européenne; CLIA = Clinical Laboratory Improvement Amendment; CPT = Cell Preparation Tubes; CMS = Centers for Medicare and Medicaid Services; FDA = Federal Drug Administration; IVD = in vitro diagnostic; NGS = next-generation sequencing; other abbreviations as in Table 1.

La nostra esperienza sfruttando gli *HLA DRB1* in ddPCR

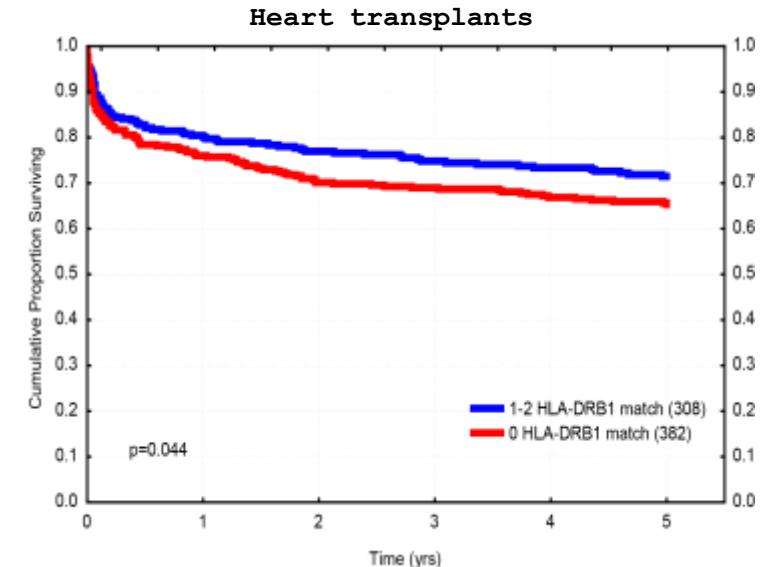
Perché il gene *HLA-*

- ✓ *DRB1*? Non è troppo polimorfico nella popolazione
- ✓ Sonde e primers erano commercialmente disponibili
- ✓ I trapianti "matched" per HLA-DR vanno in generale meglio

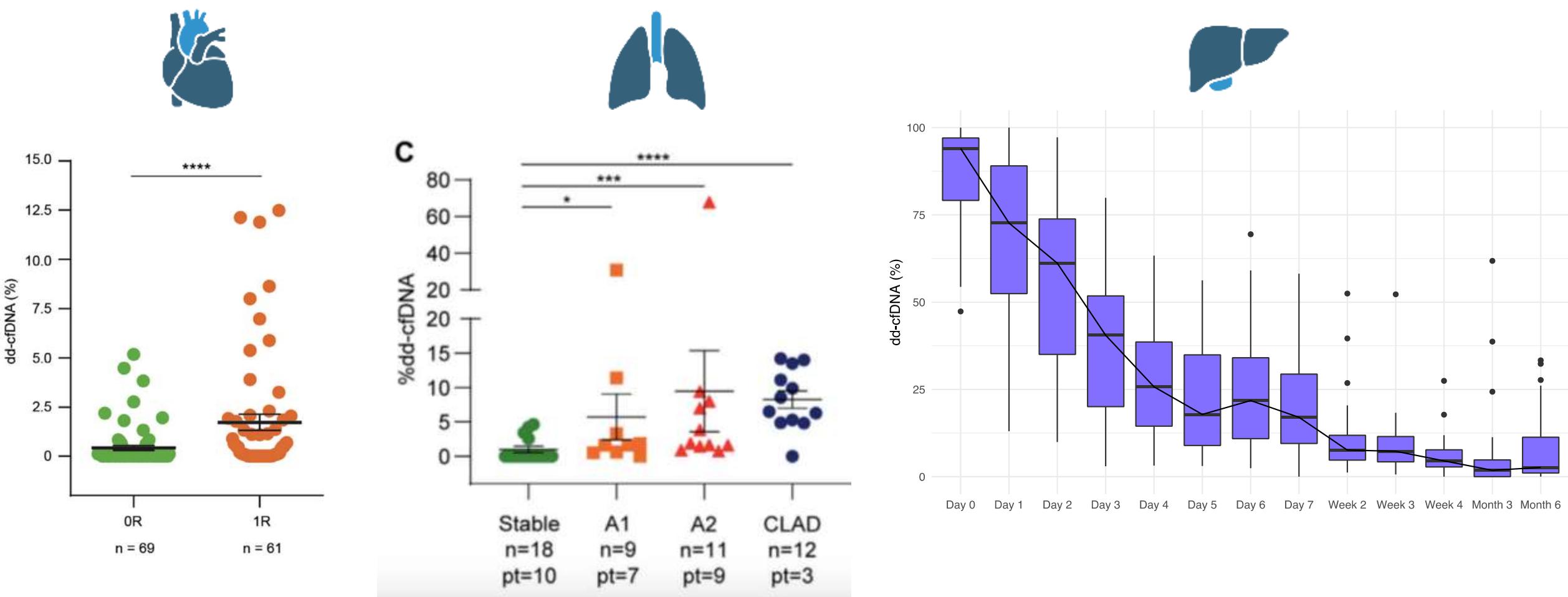


Probe ID	Target allele
dHsaEXD29156242	HLA-DRB1*01
dHsaEXD93426015	HLA-DRB1*03
dHsaEXD67695788	HLA-DRB1*04
dHsaEXD41965561	HLA-DRB1*07
dHsaEXD16235334	HLA-DRB1*08
dHsaEXD80505107	HLA-DRB1*11
dHsaEXD54774880	HLA-DRB1*13
dHsaEXD29044653	HLA-DRB1*15/16

Zou et al, Hum Immunol 2017



Validazione di un test “home-made”



Quando e come usare questo test?

- ✓ Poiché ha un valore predittivo negative lo potremmo usare per limitare le biopsie di sorveglianza
- ✓ Abbiamo offerto a una popolazione di pazienti pediatrici trapiantati di cuore e in follow-up presso il Centro di Torino il monitoraggio del dd-cfDNA come parte dei loro esami di controllo a ogni appuntamento per 12 mesi consecutive
- ✓ Al termine dello studio abbiamo correlato i dati di dd-cfDNA con le informazioni cliniche, di laboratorio, di imaging e di biopsia per capire che cosa aggiunge questo monitoraggio

La coorte pediatrica di trapiantati di cuore

TABLE 1.

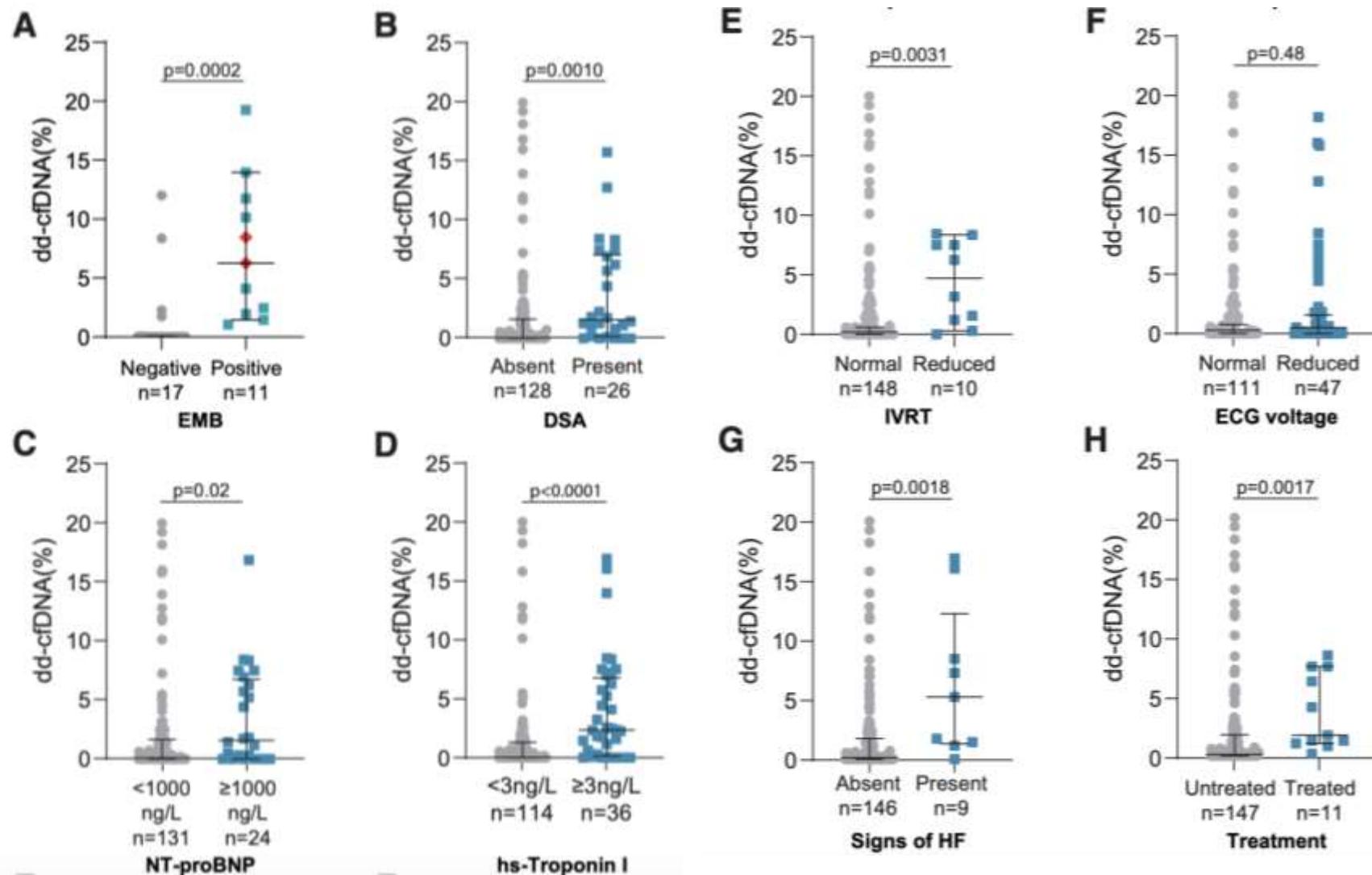
Characteristics of the enrolled cohort

ID	Sex	Age, y	Diagnosis	Transplant date	Enrollment date	Time since transplant, y	No. of EMB	dd-cfDNA determinations
1	F	18.6	CHD	01/10/06	02/22/22	16.13	1	4
2	F	15.3	DCM	07/03/08	04/26/22	13.82	0	6
3	F	17.9	DCM	02/09/10	03/08/22	12.08	1	4
4	F	15.1	CHD	08/27/10	03/29/22	11.59	1	1
5	M	14.1	DCM	12/31/10	10/04/22	11.77	1	1
6	M	16.6	CHD	04/25/11	04/08/22	10.96	1	8
7	F	12.6	HCM	07/02/11	03/21/22	10.73	1	4
8	F	13.9	DCM	03/07/13	04/05/22	9.08	1	6
9	M	11.4	DCM	04/17/13	02/24/22	8.86	1	9
10	F	10.3	DCM	01/09/14	03/01/22	8.15	1	6
11	M	10.2	CHD	09/18/14	03/14/22	7.49	0	3
12	F	8.4	DCM	06/20/15	02/24/22	6.69	1	5
13	F	18.5	CHD	01/08/16	03/09/22	6.17	1	8
14	F	12.7	CHD	07/16/16	05/09/22	5.82	1	5
15	F	7.0	DCM	10/21/16	04/19/22	5.50	1	5
16	F	8.3	DCM	04/26/17	05/23/22	5.08	1	4
17	F	7.2	DCM	08/01/17	03/03/22	4.59	1	3
18	F	12.0	DCM	12/05/17	02/28/22	4.24	1	5
19	M	5.6	DCM	11/04/18	02/23/22	3.31	4	11
20	M	7.0	CHD	06/16/19	05/02/22	2.88	1	5
21	F	3.5	DCM	11/12/20	04/11/22	1.41	1	4
22	F	9.9	CHD	12/22/20	03/15/22	1.23	1	7
23	M	14.6	RCM	07/01/21	02/25/22	0.65	2	13
24	M	16.5	CHD	07/17/21	03/10/22	0.65	1	5
25	F	2.4	DCM	08/25/21	03/14/22	0.55	1	7
26	M	14.7	CHD	03/17/22	03/24/22	0.02	1	11
27	M	0.8	CHD	07/30/22	10/20/22	0.22	0	2
28	F	7.1	DCM	01/08/23	01/20/23	0.03	0	3
29	M	11.3	CHD	01/11/23	02/13/23	0.09	0	3

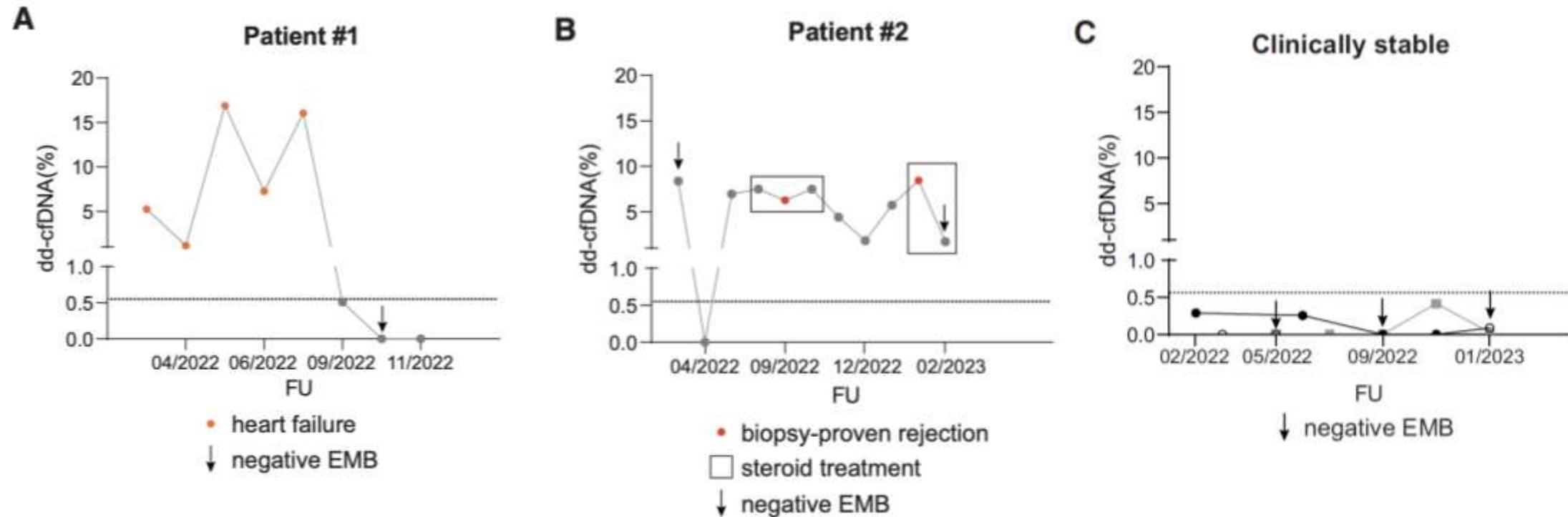
For each patient, sex, age, diagnosis, date of transplant and enrollment, and number of EMBs, and dd-cfDNA samples are listed.

CHD, congenital heart defect; DCM, dilated cardiomyopathy; dd-cfDNA, donor-derived cell-free DNA; EMB, endomyocardial biopsy; F, female; HCM, hypertrophic cardiomyopathy; M, male; RCM, restrictive cardiomyopathy.

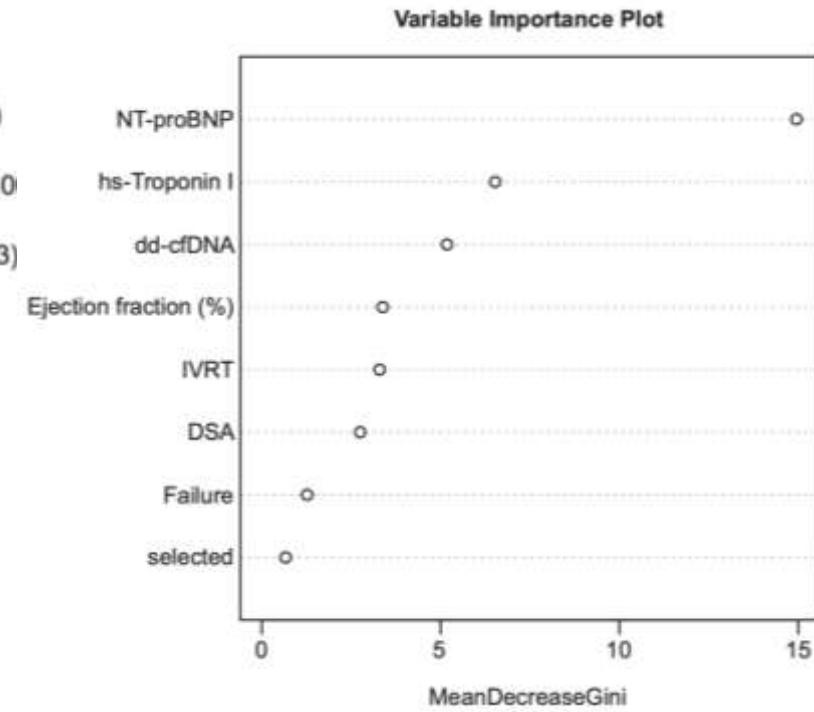
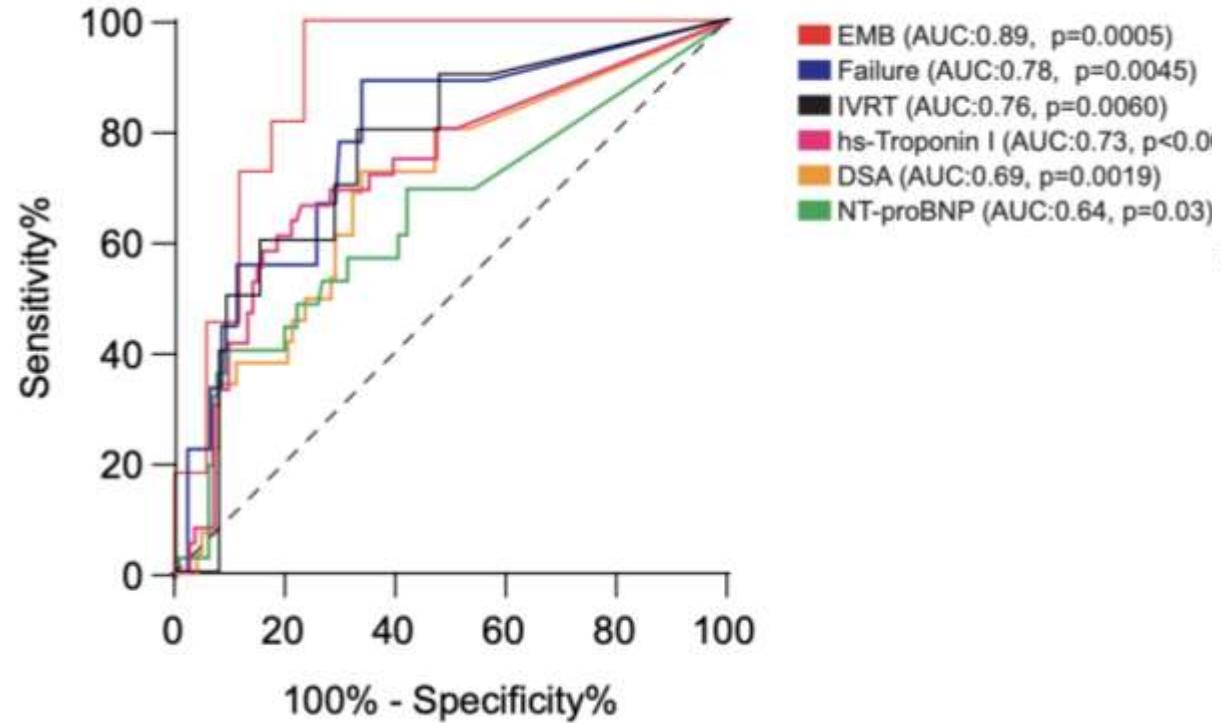
La coorte pediatrica di trapiantati di cuore



The pediatric heart transplant cohort

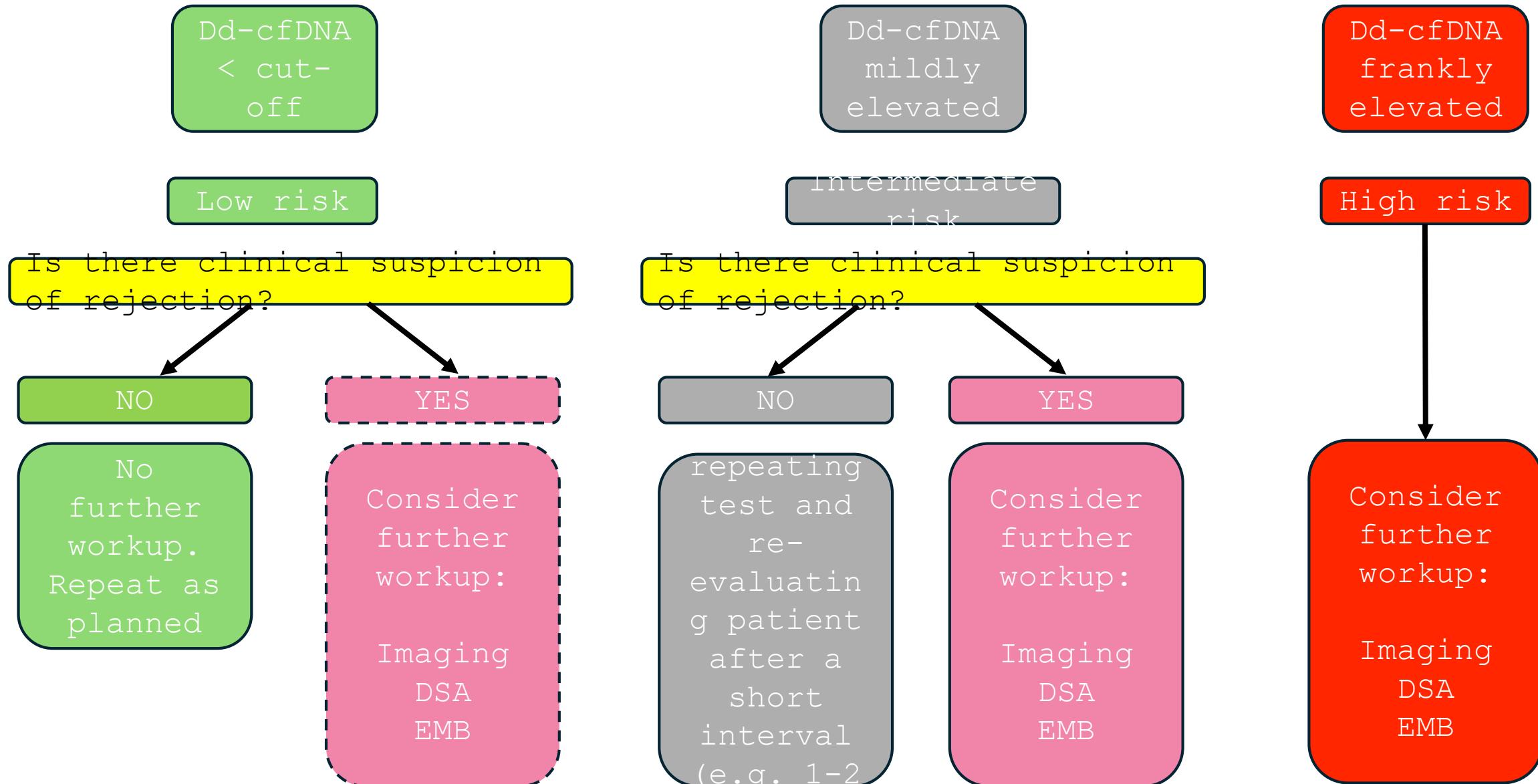


The pediatric heart transplant cohort



- ✓ Con il cut-off di 0.55%, il test aveva il 100% di sensibilità
- ✓ Pazienti con livelli di dd-cfDNA levels sotto il cut-off potrebbero posporre la loro biopsia di sorveglianza
- ✓ Con questi criteri avremmo evitato una biopsia a 7 pazienti, tutte con esito OR

Un modello operativo per usare il dd-cfDNA nei trapiantati di cuore



E nel trapianto di rene come lo usiamo? Le raccomandazioni ESOT

Question 1. In kidney transplant patients with stable graft function, is plasma dd-cfDNA measurement a reliable diagnostic tool for subclinical acute rejection monitoring when compared with standard of care (eGFR/creatinine monitoring or surveillance biopsy)?

Recommendation 1.1 - We suggest that clinicians consider measuring serial plasma dd-cfDNA in patients with stable graft function to exclude the presence of subclinical antibody-mediated rejection. Quality of Evidence - Moderate Strength of Recommendation - Weak in Favor

QUANDO? Insieme ai DSA (4/5 volte nei primi 24 mesi) e/o in concomitanza di una biopsia nei centri che eseguono biopsie protocolari
Question 2. In kidney transplant patients with acute allograft dysfunction, is plasma dd-cfDNA measurement a reliable diagnostic tool for acute rejection monitoring when compared with standard of care (eGFR/creatinine monitoring or for cause biopsy)?

Recommendation 2.1 - We recommend that clinicians measure plasma dd-cfDNA in patients with acute graft dysfunction to exclude the presence of rejection, particularly antibody-mediated rejection. Quality of Evidence - Moderate. Strength of Recommendation - Moderate in Favor

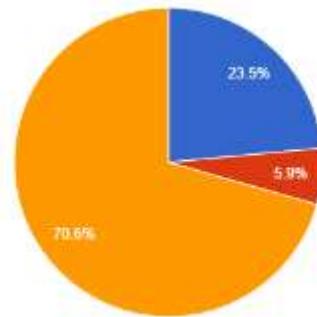
QUANDO? In concomitanza della biopsia. Ma cosa aggiunge?

SURVEY cfDNA: domande

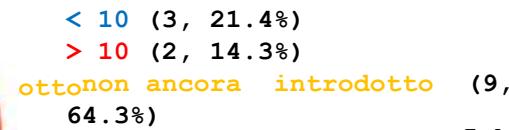
- Quanti analisi eseguite in un anno (se recentemente introdotto quante in un mese) ?
- Per quali pazienti eseguite il test?
- Per quali indicazioni cliniche eseguite il test?
- Quanti e quali plasmi sono analizzati per ciascun paziente?
- Quali provette sono utilizzate per la raccolta del plasma
- Come viene estratto il DNA?
- Tipo di kit estrazione automatica
- Viene valutata la quantità del DNA estratto prima del test
- Se sì, con quale metodica?
- Quale metodica è utilizzata per il test
- Con quale software analizzate i risultati?
- Quale cut off (%) viene utilizzato come soglia?

SURVEY cfDNA: risposte (17)

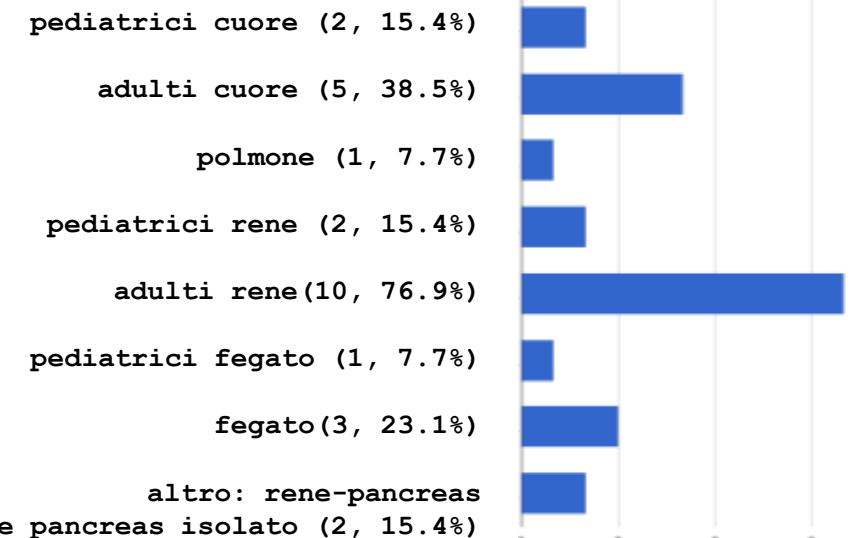
Quanti test eseguiti in un anno ?



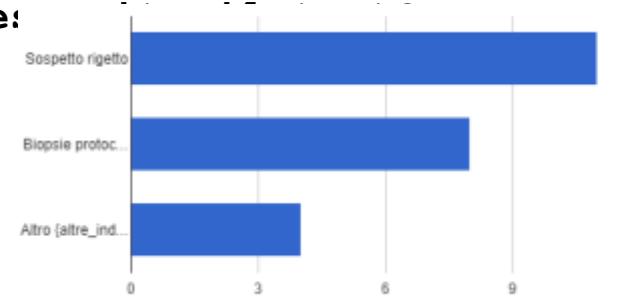
Quanti test eseguiti in un centri)



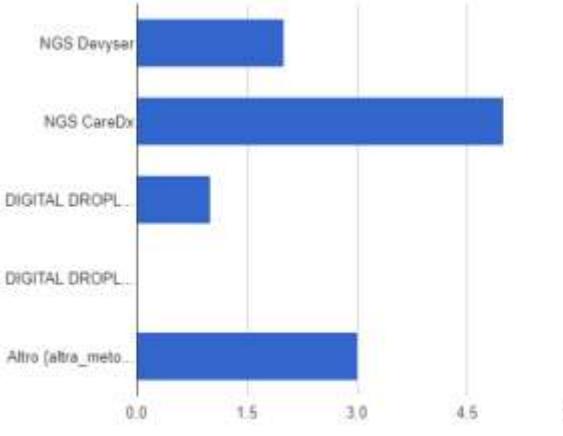
Per quali pazienti eseguite il test?



Per quali indicazioni cliniche es:



Quale metodica è utilizzata per



SURVEY cfDNA: next

- Cut-off definito per ciascun organo/kit utilizzato
- Controlli di qualità
- Identificare una prestazione con dicitura/costo compatibile con il test

Per concludere

- ✓ Il monitoraggio del dd-cfDNA nel trapianto di rene e di cuore sta acquistando importanza
- ✓ I sistemi più consolidati per il suo monitoraggio sono basati su NGS (in Europa) e su NGS e dd-PCR (negli Stati Uniti)
- ✓ Il test ha un valore predittivo negativo elevato
- ✓ Occorre disegnare delle sperimentazioni cliniche multicentriche e coordinate per potere analizzare numeri alti di pazienti

Lavoro di squadra!

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