

Table S1. Performance characteristics for different dd-cfDNA cut-offs.

	dd-cfDNA Cut-offs					
	0.6%	0.8%	1.0%	1.2%	1.4%	1.6%
Sensitivity (95% CI)	95.2% (87.8–100)	93.5% (85.0–100)	88.7% (77.7–99.8)	85.5% (73.2–97.8)	80.7% (66.9–94.5)	69.4% (53.2–85.6)
Specificity (95% CI)	59.6% (51.7–67.6)	66.3% (58.7–74.0)	72.6% (65.4–79.8)	79.5% (73.0–76.0)	82.8% (76.7–88.9)	86.9% (81.4–92.3)
PPV (95% CI)	44.0% (38.8–49.2)	48.1% (42.0–54.2)	51.9% (44.7–59.2)	58.2% (49.7–66.7)	61.0% (51.6–70.3)	63.8% (52.8–74.8)
NPV (95% CI)	97.4% (93.4–100)	96.9% (92.8–100)	95.1% (90.5–99.7)	94.3% (89.7–98.9)	92.8% (88.0–97.6)	89.5% (84.5–94.5)

Table S2. Multiple groups comparisons.

dd-cfDNA			eGFR		
Comparison	z	Padi	Comparison	z	Padi
AR - BL	5.711116	0	AR - BL	-1.17381	0.460946
AR - OI	4.569421	2.00E-05	AR - OI	-1.95735	0.15092
BL - OI	0.13599	0.891829	BL - OI	-1.15746	0.247085
AR - STA	6.746807	0	AR - STA	-8.65195	0
BL - STA	1.107485	0.804253	BL - STA	-9.08226	0
OI - STA	0.644728	1	OI - STA	-5.19251	1.00E-06

Significant (adj. $p < 0.0001$) results are bolded. The test to compares dd-cfDNA and eGFR for AR vs. Non-AR was a Kruskal-Wallis test.

Table S3. Cohort breakdown into for-cause and protocol biopsy for AR versus Non-AR.

Status	Biopsy_Reason	Total	Median	Low	High	Mean	SD
AR	For cause	25	2.04	0.09	23.9	3.85	4.81
	Protocol	13	3.56	0.12	23.4	6.16	6.44
Non-AR	For cause	78	0.645	0.02	6.54	1.08	1.27
	Protocol	101	0.27	0.03	6.78	0.80	1.28

Table S4. Descriptive statistics dd-cfDNA as a function of ABMR/TCMR

	Count	Median	Low	High	Mean	SD
ABMR	16	2.22	0.12	23.9	5.55	7.64
Both	12	2.56	0.09	8.8	4.00	3.21
TCMR	10	2.69	1.01	9.77	3.96	3.11

Table S5. Histology report for all active rejection cases.

Histology Report	dd-cfDNA Assay Result	
	<1% dd-cfDNA	>1% dd-cfDNA
ABMR	0	1
ABMR + acute bTCMR	0	1
ABMR + acute TCMR IA	0	4
ABMR + acute TCMR IIB	0	2
ABMR + bTCMR	3	2
ABMR + chronic active bTCMR	0	5
ABMR + chronic active TCMR IA	0	1
ABMR + chronic active TCMR IB	1	0
ABMR + chronic active TCMR II	0	2
ABMR + chronic active TCMR IIA	0	1
acute ABMR	0	1
acute TCMR IA	0	2
bABMR + acute TCMR IA	0	2
bC4d-positive ABMR + TCMR-IA	1	0
chronic active ABMR + bTCMR	0	2
chronic active TCMR IA	0	1
chronic active TCMR IB	0	3
TCMR IA	0	2
TG + cABMR	0	1

Table S6. Descriptive statistics for dd-cfDNA as a function of donor type.

Group	Donor_type	Total	Median	Low	High	Mean	SD
AR	Living Related	1	0.97	0.97	0.97	0.97	N/A
	Living Non-related	2	4.88	1.39	8.37	4.88	4.94
	Deceased Non-related	35	2.41	0.09	23.9	4.74	5.59
Non-AR	Living Related	14	0.265	0.02	3.89	0.67	1.09
	Living Non-related	75	0.65	0.05	6.78	1.17	1.54
	Deceased Non-related	90	0.405	0.06	6.3	0.75	1.02

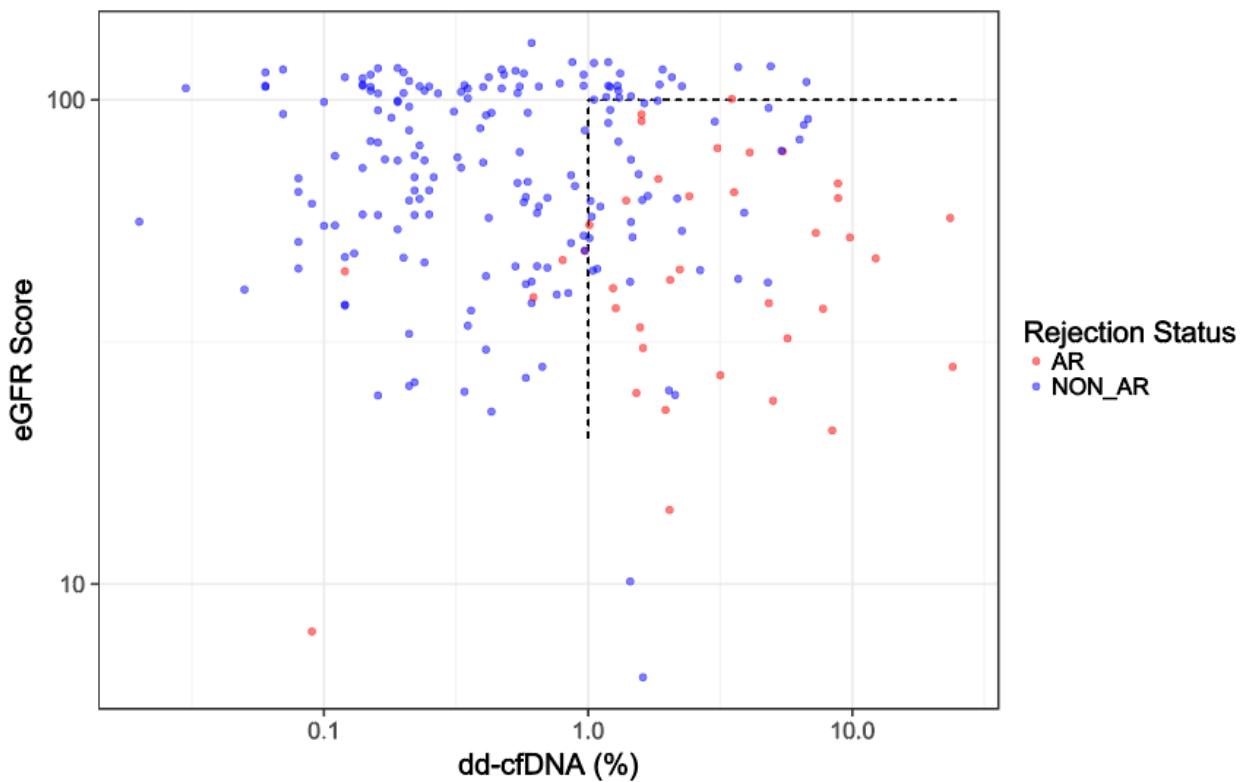


Figure S1. dd-cfDNA and eGFR in biopsy matched samples.

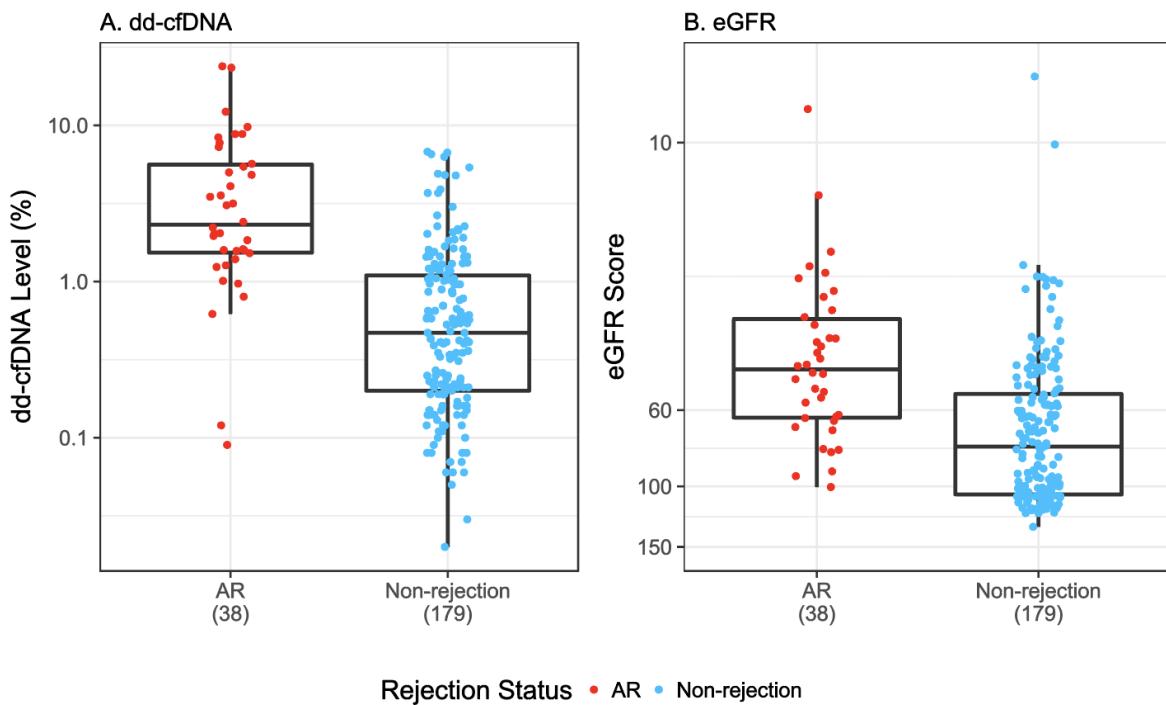


Figure S2. Discrimination of active rejection by dd-cfDNA versus eGFR in AR versus non-AR.

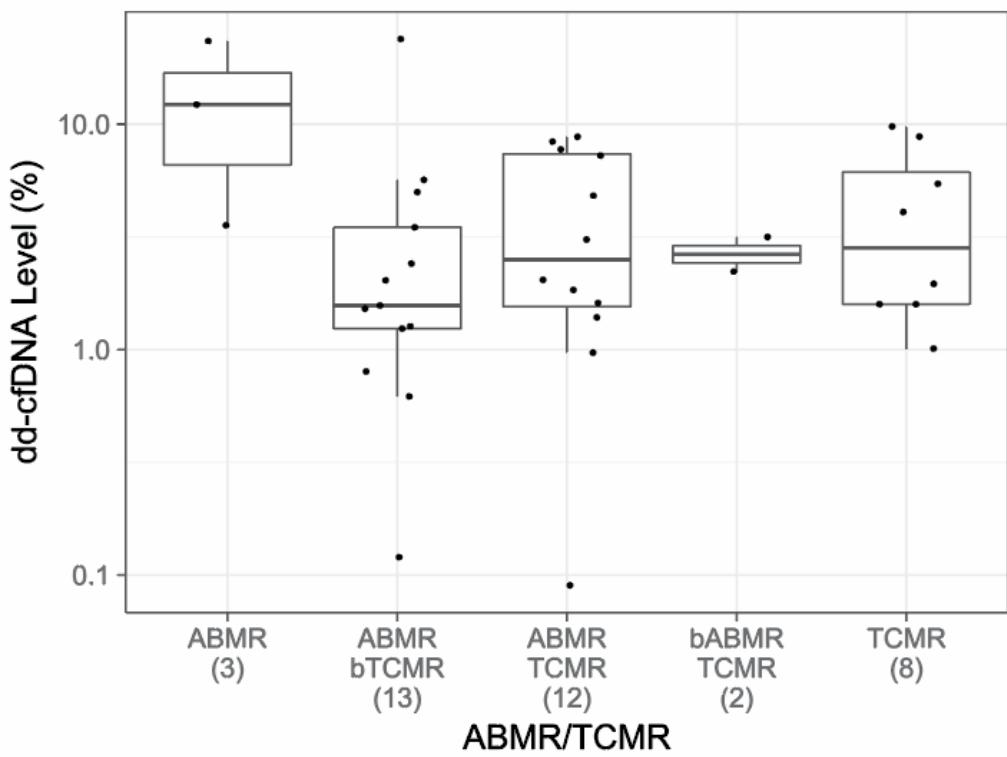


Figure S3. Percent dd-cfDNA by ABMR/TCMR status.