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- **Table S4.** Regression analyses to determine the effects of new UA-PD on cPRA increases (0-1.000) after failed transplant and relisting, augmented to evaluate the interaction between the effects of HLA-DR and DQ. Results are stratified by receipt of an initial (A) deceased donor kidney or (B) living donor kidney and by pre-transplantation cPRA (0%, 1-50%, 51-100%). The interaction term "DR & DQ Interaction" estimates the change in cPRA given an additional HLA-DR UA-PD in the presence of an HLA-DQ UA-PD. These results are presented graphically as recipient scenarios in Figures S2-S3.
- **Figure S1.** Scenarios applying regression analyses from Table S3 to demonstrate interactions between the effects of HLA-DR and HLA-DQ mismatches (MM) on the probabilities of being relisted with new HLA-DR and HLA-DQ UA-PD (0-100.0%). Scenarios are presented for example A) deceased donor recipients and B) living donor recipients. Rightmost columns depict expected changes in probabilities of new HLA-DR and HLA-DQ UA-PD given the presence or absence of HLA-DR and/or HLA-DQ mismatches, as compared a recipient with no HLA-DR or DQ mismatches.
- **Figure S2.** Scenarios applying regression analyses from Table S4 to demonstrate interactions between the effects of HLA-DR and HLA-DQ UA-PD on cPRA increases (0-100.0%) in prior deceased donor recipients with pre-transplant cPRA of 0% (A) or 1-50% (B). The rightmost column depicts expected increases in cPRA after graft failure and relisting with new HLA-DR and/or HLA-DQ UA-PD as compared to a recipient relisted with no HLA-DR or DQ UA-PD.
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	Failed	Deceased Donor K	idney	Failed Living Donor Kidney		
Characteristics	African American Recipient	White Recipient	Hispanic Recipient	African American Recipient	White Recipient	Hispanic Recipient
Median age, yr (IQR)	50 [40,59]*	54.5 [39,64]*	44 [29,56]*	43 [33,53]*	45 [32,56]*	35 [26,46]*
Male, n (%)	951, 60.57%	647, 57.87%	290, 57.88%	165, 56.31%	512, 60.52%	117, 53.92%
Median BMI, kg/m² (IQR) Cause of ESRD, n (%)	28, [24, 33]*	28, [24, 32]*	27, [24, 31]*	28, [24, 33]*	27, [23, 31]*	26.5, [22,31]*
GN	424, 27.01%	313, 28.00%	157, 31.34%	118, 40.27%	353, 41.73%	88, 40.55%
Diabetes	304, 19.36%*	183, 16.37%*	115, 22.95%*	48, 16.38%*	86, 10.17%*	33, 15.21%*
Hypertension	557, 35.48%*	154, 13.77%*	81, 16.17%*	83, 28.33%*	86, 10.17%*	37, 17.05%*
PKD	65, 4.14%*	151, 13.51%*	27, 5.39%*	12, 4.10%*	107, 12.65%*	11, 5.07%*
Other	220, 14.01%*	317, 28.35%*	121, 24.15%*	32, 10.92%*	214, 25.30%*	48, 22.12%*
Education level, n (%)						
No college	678, 43.18%*	438, 39.18%*	326, 65.07%*	103, 35.15%*	263, 31.09%*	91, 41.94%*
Some college	468, 29.81%*	303, 27.10%*	106, 21.16%*	104, 35.49%	249, 29.43%	67, 30.88%
College graduate	265, 16.88%*	214, 19.14%*	42, 8.38%*	61, 20.82%*	222, 26.24%*	40, 18.43%*
Graduate degree Insurance status	96, 6.11%*	92, 8.23%*	13, 2.59%*	17, 5.80%*	75, 8.87%*	7, 3.23%*
Private	312, 19.87%*	345, 30.86%*	81, 16.17%*	119, 40.61%*	491, 58.04%*	83, 38.25%*
Medicare	1,100, 70.06%*	687, 61.45%*	322, 64.27%*	151, 51.54%*	314, 37.12%*	101, 46.54%*
Medicaid	117, 7.45%*	58, 5.19%*	81, 16.17%*	18, 6.14%*	27, 3.19%*	25, 11.52%*
Other/Missing	41, 2.61%	28, 2.50%	17, 3.39%	5, 1.71%	14, 1.65%	8, 3.69%
Median time on dialysis	1,687.5, *	1,162 *	1,533.5*	661.5*	452*	418.5*
pre-transplant, days (IQR)	[1,026, 2,472]	[641.5, 1,769]	[880, 2,469]	[388, 1,301.5]	[268, 846]	[270.5, 736]
Median time from initial	745*	579.5*	613*	1,062	955.5	1,096
transplant to return to	[218, 1,239]	[168, 1,265]	[192, 1,174]	[593, 1,465]	[412, 1,439]	[569, 1,464]
dialysis, days (IQR)	[220, 2,200]	[200, 2,200]	[232, 2,27.]	[555, 1, 155]	[:==, =,:==,	[503, 2, 10.]
Median time between graft	1,104*	648.5*	891*	1,440	1,376	1,411
failure/relisting, days (IQR)	[258, 1,911]	[110, 1,628]	[168, 1,802]	[749, 2,107]	[432, 2,194]	[530, 2,200]
Pre-transplant cPRA	[230, 1,311]	[110, 1,020]	[100, 1,002]	[, 13, 2,10,]	[102, 2,131]	[330, 2,200]
0%	961, 61.21%	734, 65.65%	323, 64.47%	198, 67.58%*	628, 74.23%*	165, 76.04%*
0.1-50%	318, 20.25%	221, 19.77%	105, 20.96%	65, 22.18%	160, 18.91%	35, 16.13%
50.1-80%	132, 8.41%*	57, 5.10%*	30, 5.99%*	13, 4.44%	32, 3.78%	11, 5.07%
80.1-98.0%	119, 7.58%	88, 7.87%	31, 6.19%	11, 3.75%	21, 2.48%	5, 2.30%
>98.1%	40, 2.55%	18, 1.61%	12, 2.40%	6, 2.05%	5, 0.59%	1, 0.46%
Peak post-graft loss cPRA	40, 2.3370	10, 1.01/0	12, 2.4070	0, 2.0370	3, 0.3370	1, 0.4070
0%	290, 18.47%*	289, 25.85%*	98, 19.56%*	51, 17.41%*	242, 28.61%*	55, 25.35%*
0.1-50%	127, 8.09%*	133, 11.90%*	51, 10.18%*	25, 8.53%*	121, 14.30%*	22, 10.14%*
50.1-80%	145, 9.24%	120, 10.73%	55, 10.98%	33, 11.26%	105, 12.41%	30, 13.82%
80.1-98.0%	276, 17.58%	210, 18.78%	78, 15.57%	60, 20.48%	169, 19.98%	40, 18.43%
>98.1%	732, 46.62%*	366, 32.74%*	219, 43.71%*	124, 42.32%*	209, 24.70%*	70, 32.26%*
HLA-A mismatch, n (%)	7.52, 10.0270	300, 32.7 470	_15, 15.7170	12.1, 12.32/0	_05, _ 7.7070	, 0, 02.20/0
0	93, 5.92%*	145, 12.97%*	59, 11.78%*	40, 13.65%	152, 17.97%	44, 20.28%
1	563, 35.86%	435, 38.91%	186, 37.13%	146, 49.83%	460, 54.37%	110, 50.69%
2	914, 58.22%*	538, 48.12%*	256, 51.10%*	107, 36.52%*	234, 27.66%*	63, 29.03%*
HLA-B mismatch, n (%)	31 1, 30.22/0	330, 4 0.12/0	200, 01.10/0	10., 30.32/0	20 1, 27.00/0	00, 20.00/0
0	47, 2.99%*	96, 8.59%*	30, 5.99%*	28, 9.56%	89, 10.52%	28, 12.90%
1	349, 22.23%*	313, 28.00%*	100, 19.96%*	124, 42.32%*	410, 48.46%*	116, 53.46%*
2	1,174, 74.78%*	709, 63.42%*	371, 74.05%*	141, 48.12%*	347, 41.02%*	73, 33.64%*
HLA-C mismatch, n (%)	1,174, 74.70%	703, 03.4270	3/1, /4.03%	141, 40.1270	347,41.02%	73, 33.04%
	110, 7.01%*	122 11 000/*	// O 700/*	E6 10 110/	167 10 740/	17 21 660/
0 1		123, 11.00%*	44, 8.78%*	56, 19.11%	167, 19.74%	47, 21.66%
-	645, 41.08% 815, 51.91%*	472, 42.22% 523, 46.78%*	191, 38.12% 266, 53.09*	138, 47.10%* 99, 33.79%*	423, 50.00%* 256, 30.26%*	128, 58.99%* 42, 19.35%*
2				44 33 /4%"	/ 3D 3H / D% "	4/ I4 45% [*]

0	126, 8.03%*	180, 16.10%*	58, 11.58%*	47, 16.04%	129, 15.25%	41, 18.89%
1	738, 47.01%	536, 47.94%	246, 49.10%	147, 50.17%	437, 51.65%	104, 47.93%
2	706, 44.97%*	402, 35.96%*	197, 39.32%*	99, 33.79%	280, 33.10%	72, 33.18%
HLA-DQ mismatch, n (%)						
0	257, 16.37%*	227, 20.30%*	80, 15.97%*	72, 24.57%	188, 22.22%	42, 23.96%
1	863, 54.97%	599, 53.58%	267, 53.29%	172, 58.70%	487, 57.57%	124, 57.14%
2	450, 28.66%	292, 26.12%	154, 30.75%	49, 16.72%	171, 20.21%	41, 18.89%

Table S1. Demographics and relevant characteristics stratified by deceased vs. living donor category of the initial failed graft and recipient race/ethnicity. Asterisks (*) denote statistically significant differences between the three race/ethnicity groups (p<0.05).

	Difference Between	n Initial and Maximum cP	RA After Failed Tr	ansplant and Re-Listing
A) Deceased	All Data	Initial cPRA was 0	(0,.5]	(.5,1]
DQ UA	0.235***	0.279***	0.198***	0.047***
	(0.009)	(0.011)	(0.018)	(0.009)
DR UA	0.124***	0.145***	0.097***	0.038***
	(0.010)	(0.012)	(0.020)	(0.010)
A UA	0.231***	0.283***	0.195***	0.060***
	(0.011)	(0.013)	(0.022)	(0.010)
BUA	0.205***	0.225***	0.232***	0.057***
	(0.011)	(0.013)	(0.023)	(0.010)
C UA	0.113***	0.137***	0.088***	0.023**
	(0.011)	(0.013)	(0.022)	(0.011)
A Mismatch	-0.026***	-0.030***	-0.013	-0.005
	(0.006)	(0.008)	(0.013)	(0.006)
B Mismatch	-0.022***	-0.018*	0.004	0.002
	(0.008)	(0.010)	(0.016)	(0.007)
C Mismatch	-0.012*	-0.007	-0.011	-0.010
	(0.007)	(0.008)	(0.013)	(0.006)
DO Mismatch	-0.020***	-0.013	-0.024	-0.001
	(0.007)	(0.008)	(0.015)	(0.007)
DR Mismatch	-0.010	-0.004	-0.004	-0.007
	(0.007)	(0.009)	(0.015)	(0.007)
Time between failure and	0.000***	0.000***	0.000***	0.000
relisting	(0.000)	(0.000)	(0.000)	(0.000)
Time between relisting	0.000***	0.000***	0.000	0.000
and final follow up	(0.000)	(0.000)	(0.000)	(0.000)
PRA Original	-0.600***	(0.000)	-0.663***	-0.558***
- Lu - Oligiimi	(0.013)		(0.052)	(0.024)
Constant	0.308***	0.195***	0.316***	0.499***
Constant	(0.016)	(0.021)	(0.032)	(0.023)
Observations	3,443	2,173	711	559
R-squared	0.737	0.771	0.698	0.642

Standard errors in parentheses, *** p<0.01, ** p<0.05, * p<0.1

	Difference Between	Initial and Maximum cP	RA After Failed Ti	ansplant and Re-Listing
B) Living	All Data	Initial cPRA was 0	(0,.5]	(.5,1]
DQ UA	0.290***	0.332***	0.266***	0.055***
DQUA	(0.015)	(0.018)	(0.033)	(0.020)
DR UA	0.187***	0.192***	0.190***	0.063***
DRUA	(0.016)	(0.019)	(0.036)	(0.020)
A UA	0.218***	0.248***	0.159***	-0.001
71 071	(0.018)	(0.021)	(0.037)	(0.024)
BUA	0.168***	0.177***	0.183***	0.076***
BOA	(0.018)	(0.021)	(0.037)	(0.022)
C UA	0.130***	0.144***	0.107***	-0.004
CON	(0.019)	(0.023)	(0.038)	(0.028)
A Mismatch	-0.044***	-0.037***	-0.051*	0.020
71 Wildinaten	(0.012)	(0.014)	(0.027)	(0.015)
B Mismatch	0.008	0.007	0.033	0.017
Divisinaten	(0.014)	(0.017)	(0.032)	(0.016)
C Mismatch	0.003	0.004	-0.016	0.001
Civilsinaten	(0.012)	(0.014)	(0.028)	(0.015)
DQ Mismatch	-0.028**	-0.035**	-0.027	-0.009
DQ IVIISIIIATOII	(0.013)	(0.016)	(0.029)	(0.017)
DR Mismatch	0.036***	0.041***	0.009	-0.015
Dit Mishaten	(0.013)	(0.015)	(0.029)	(0.017)
Time between failure and	0.000***	0.000***	0.000	-0.000
relisting	(0.000)	(0.000)	(0.000)	(0.000)
Time between relisting	0.000***	0.000***	0.000***	0.000*
and final follow up	(0.000)	(0.000)	(0.000)	(0.000)
PRA Original	-0.609***	(0.000)	-0.569***	-0.526***
8	(0.028)		(0.097)	(0.058)
Constant	0.180***	0.142***	0.235***	0.421***
	(0.020)	(0.023)	(0.045)	(0.052)
Observations	1,424	1,036	273	115
R-squared	0.696	0.714	0.678	0.656

Standard errors in parentheses, *** p<0.01, ** p<0.05, * p<0.1

Table S2. Regression analysis to determine the relationship between UA-PD at each HLA locus and changes in cPRA after relisting (Figure 4), stratified by failed (A) deceased and (B) living donor kidney and by pre-transplantation cPRA (0%, 1-50%, 51-100%). The relationship between new UA-PD and cPRA was largely independent of potential contributors to sensitization—overall HLA mismatches, time between graft failure and relisting, time between relisting and final follow up and pre-transplantation cPRA.

A) Deceased	A UA	B UA	C UA	DR UA	DQ UA
DR & DQ Interaction	-0.041**	-0.046**	-0.017	-0.075***	-0.113***
	(0.018)	(0.018)	(0.015)	(0.018)	(0.017)
A Mismatch	0.204***	0.034**	0.004	0.009	0.009
	(0.013)	(0.013)	(0.011)	(0.013)	(0.013)
B Mismatch	0.014	0.145***	0.021	-0.016	-0.007
	(0.017)	(0.017)	(0.014)	(0.016)	(0.016)
C Mismatch	0.001	0.027*	0.095***	0.008	-0.030**
	(0.014)	(0.014)	(0.011)	(0.014)	(0.013)
DR Mismatch	0.074***	0.083***	0.014	0.270***	0.126***
	(0.023)	(0.023)	(0.019)	(0.022)	(0.022)
DQ Mismatch	0.040	0.046	0.007	0.095***	0.402***
	(0.028)	(0.028)	(0.023)	(0.027)	(0.027)
Constant	0.087***	0.060*	0.024	0.034	0.039
	(0.032)	(0.033)	(0.027)	(0.031)	(0.031)
Observations	3,443	3,443	3,443	3,443	3,443
R-squared	0.085	0.054	0.030	0.079	0.138
	C4 1		41		

Standard errors in parentheses
*** p<0.01, ** p<0.05, * p<0.1

B) Living	A UA	BUA	C UA	DR UA	DQ UA
DR & DQ Interaction	-0.038	-0.049*	-0.037*	-0.105***	-0.133***
•	(0.026)	(0.027)	(0.020)	(0.026)	(0.026)
A Mismatch	0.262***	0.012	0.000	0.006	-0.007
	(0.021)	(0.021)	(0.016)	(0.021)	(0.021)
B Mismatch	-0.014	0.160***	0.001	-0.014	-0.029
	(0.027)	(0.028)	(0.020)	(0.026)	(0.026)
C Mismatch	0.003	0.036	0.126***	0.022	0.005
	(0.023)	(0.024)	(0.018)	(0.023)	(0.023)
DR Mismatch	0.025	0.046	0.007	0.307***	0.136***
	(0.033)	(0.034)	(0.025)	(0.032)	(0.033)
DQ Mismatch	0.035	0.055	0.037	0.077*	0.457***
	(0.041)	(0.042)	(0.032)	(0.041)	(0.041)
Constant	0.085**	0.078**	0.020	0.028	0.042
	(0.035)	(0.036)	(0.027)	(0.035)	(0.035)
Observations	1,424	1,424	1,424	1,424	1,424
R-squared	0.127	0.072	0.055	0.094	0.171

Standard errors in parentheses *** p<0.01, ** p<0.05, * p<0.1

Table S3. Regression analyses to determine the effects of HLA mismatches [0-2] on the probabilities of UA-PD (0-1.000), augmented to evaluate the interactions between the effects of HLA-DR and HLA-DQ. Results are stratified by receipt of an initial (A) deceased donor kidney or (B) living donor kidney. The interaction term "DR & DQ Interaction" estimates the change in probability of an UA-PD with each additional HLA-DR mismatch in the presence of an HLA-DQ mismatch, or with each additional HLA-DQ mismatch in the presence of an HLA-DR mismatch. These results are presented graphically as recipient scenarios in Figure S1.

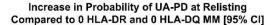
A) Deceased	All Data	Original is 0	After Failed Transplar Original is (0.0.5]	Original > 0.
A) Deceased	All Data	Original is 0	Original is (0,0.5]	Original > 0.
DR & DQ Interaction	-0.289***	-0.366***	-0.257***	-0.075***
	(0.016)	(0.019)	(0.032)	(0.016)
DQ UA	0.352***	0.424***	0.307***	0.077***
97.44	(0.011)	(0.013)	(0.022)	(0.011)
DR UA	0.284***	0.359***	0.230***	0.075***
	(0.013)	(0.016)	(0.025)	(0.012)
A UA	0.217***	0.255***	0.191***	0.058***
	(0.011)	(0.013)	(0.022)	(0.010)
B UA	0.195***	0.217***	0.217***	0.055***
	(0.011)	(0.013)	(0.022)	(0.010)
C UA	0.105***	0.120***	0.094***	0.022**
	(0.010)	(0.012)	(0.021)	(0.010)
A Mismatch	-0.024***	-0.028***	-0.008	-0.004
	(0.006)	(0.007)	(0.013)	(0.006)
B Mismatch	-0.026***	-0.022**	-0.000	0.000
	(0.008)	(0.009)	(0.015)	(0.007)
C Mismatch	-0.010	-0.004	-0.012	-0.009
	(0.006)	(0.007)	(0.012)	(0.006)
DQ Mismatch	-0.014**	-0.006	-0.020	0.001
17.0	(0.007)	(0.008)	(0.014)	(0.007)
DR Mismatch	-0.015**	-0.007	-0.012	-0.009
	(0.007)	(0.008)	(0.014)	(0.007)
Time between failure and relisting	0.000***	0.000***	0.000**	-0.000
	(0.000)	(0.000)	(0.000)	(0.000)
Time between relisting and final	0.000***	0.000***	0.000	0.000
follow up	(0.000)	(0.000)	(0.000)	(0.000)
PRA Original	-0.601***		-0.661***	-0.558***
	(0.012)		(0.049)	(0.023)
Constant	0.290***	0.167***	0.301***	0.495***
	(0.015)	(0.019)	(0.031)	(0.023)
Observations	3,443	2,173	711	559
R-squared	0.759	0.803	0.724	0.656

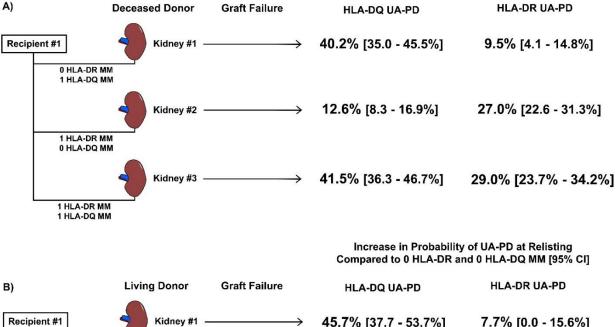
Standard errors in parentheses. *** p<0.01, ** p<0.05, * p<0.1

B) Living	Difference Between Init	tial and Maximum c	PRA After Failed Tra	nsplant and Re-Listi
CELTIFICATION TO A COLOR OF THE	All Data	Original is 0	Original is (0,0.5]	Original > 0.5
DR & DQ Interaction	-0.359***	-0.409***	-0.340***	-0.008
Dr. & DQ Interaction	(0.027)	(0.031)	(0.057)	(0.040)
DQ UA	0.417***	0.481***	0.379***	0.057**
DQOA	(0.017)	(0.020)	(0.037)	(0.023)
DR UA	0.392***	0.417***	0.397***	0.068**
DICOA	(0.022)	(0.025)	(0.048)	(0.034)
A UA	0.199***	0.220***	0.154***	-0.002
1011	(0.017)	(0.020)	(0.035)	(0.024)
B UA	0.164***	0.174***	0.181***	0.075***
	(0.017)	(0.020)	(0.035)	(0.022)
C UA	0.115***	0.126***	0.086**	-0.004
COA	(0.018)	(0.021)	(0.036)	(0.029)
A Mismatch	-0.043***	-0.034***	-0.044*	0.020
A Mismatch	(0.011)	(0.013)	(0.025)	(0.015)
3 Mismatch	0.010	0.005	0.041	0.017
J Wildington	(0.013)	(0.015)	(0.030)	(0.016)
C Mismatch	-0.002	0.002	-0.030	0.001
o minimum	(0.011)	(0.013)	(0.026)	(0.016)
OQ Mismatch	-0.017	-0.021	-0.021	-0.008
5 Q Iviisimuteii	(0.012)	(0.014)	(0.027)	(0.017)
OR Mismatch	0.024**	0.030**	-0.007	-0.016
ort minimuten	(0.012)	(0.014)	(0.027)	(0.017)
Time between failure and relisting	0.000***	0.000***	0.000	-0.000
-	(0.000)	(0.000)	(0.000)	(0.000)
Fime between relisting and final follow up	0.000***	0.000***	0.000**	0.000*
	(0.000)	(0.000)	(0.000)	(0.000)
PRA Original	-0.609***		-0.569***	-0.525***
	(0.026)		(0.091)	(0.059)
Constant	0.153***	0.103***	0.219***	0.420***
	(0.019)	(0.022)	(0.042)	(0.053)
Observations	1,424	1,036	273	115
R-squared	0.731	0.756	0.716	0.656

Standard errors in parentheses. *** p<0.01, ** p<0.05, * p<0.1

Table S4. Regression analyses to determine the effects of new UA-PD on cPRA increases (0-1.000) after failed transplant and relisting, augmented to evaluate the interaction between the effects of HLA-DR and DQ. Results are stratified by receipt of an initial (A) deceased donor kidney or (B) living donor kidney and by pre-transplantation cPRA (0%, 1-50%, 51-100%). The interaction term "DR & DQ Interaction" estimates the change in cPRA given an additional HLA-DR UA-PD in the presence of an HLA-DQ UA-PD. These results are presented graphically as recipient scenarios in Figures S2-S3.





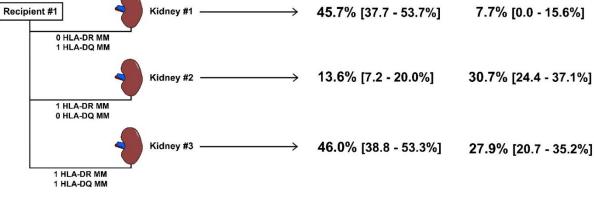


Figure S1. Scenarios applying regression analyses from Table S3 to demonstrate interactions between the effects of HLA-DR and HLA-DQ mismatches (MM) on the probabilities of being relisted with new HLA-DR and HLA-DQ UA-PD (0-100.0%). Scenarios are presented for example A) deceased donor recipients and B) living donor recipients. Rightmost columns depict expected changes in probabilities of new HLA-DR and HLA-DQ UA-PD given the presence or absence of HLA-DR and/or HLA-DQ mismatches, as compared a recipient with no HLA-DR or DQ mismatches.

Deceased Donor Recipients

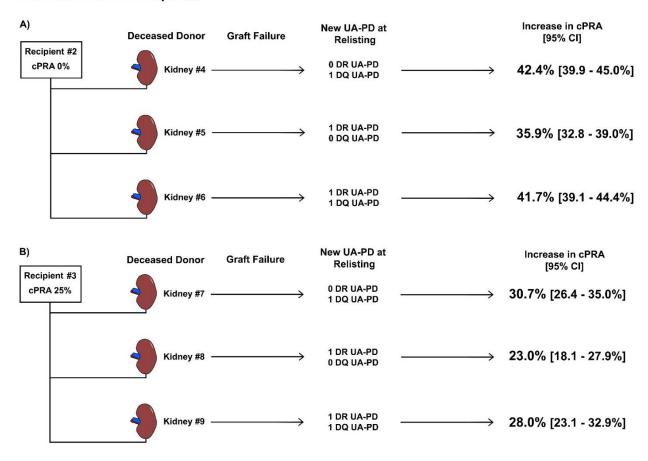


Figure S2. Scenarios applying regression analyses from Table S4 to demonstrate interactions between the effects of HLA-DR and HLA-DQ UA-PD on cPRA increases (0-100.0%) in prior deceased donor recipients with pre-transplant cPRA of 0% (A) or 1-50% (B). The rightmost column depicts expected increases in cPRA after graft failure and relisting with new HLA-DR and/or HLA-DQ UA-PD as compared to a recipient relisted with no HLA-DR or DQ UA-PD.

Living Donors

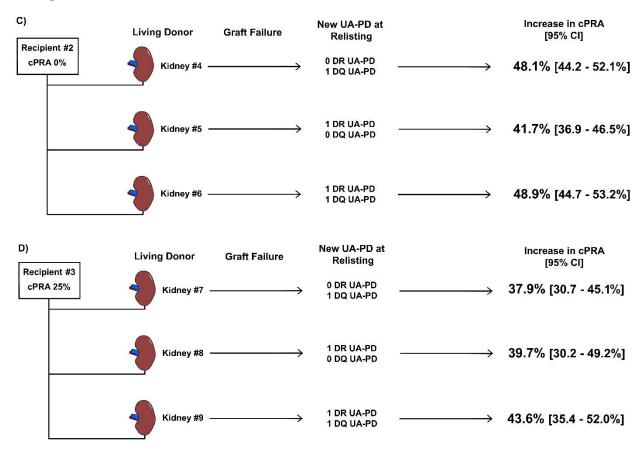


Figure S3. Scenarios applying regression analyses from Table S4 to demonstrate interactions between the effects of HLA-DR and HLA-DQ UA-PD on cPRA increases (0-100.0%) in prior living donor recipients with pre-transplant cPRA of 0% (A) or 1-50% (B). The rightmost column depicts expected increases in cPRA after graft failure and relisting with new HLA-DR and/or HLA-DQ UA-PD as compared to a recipient relisted with no HLA-DR or DQ UA-PD.