

---

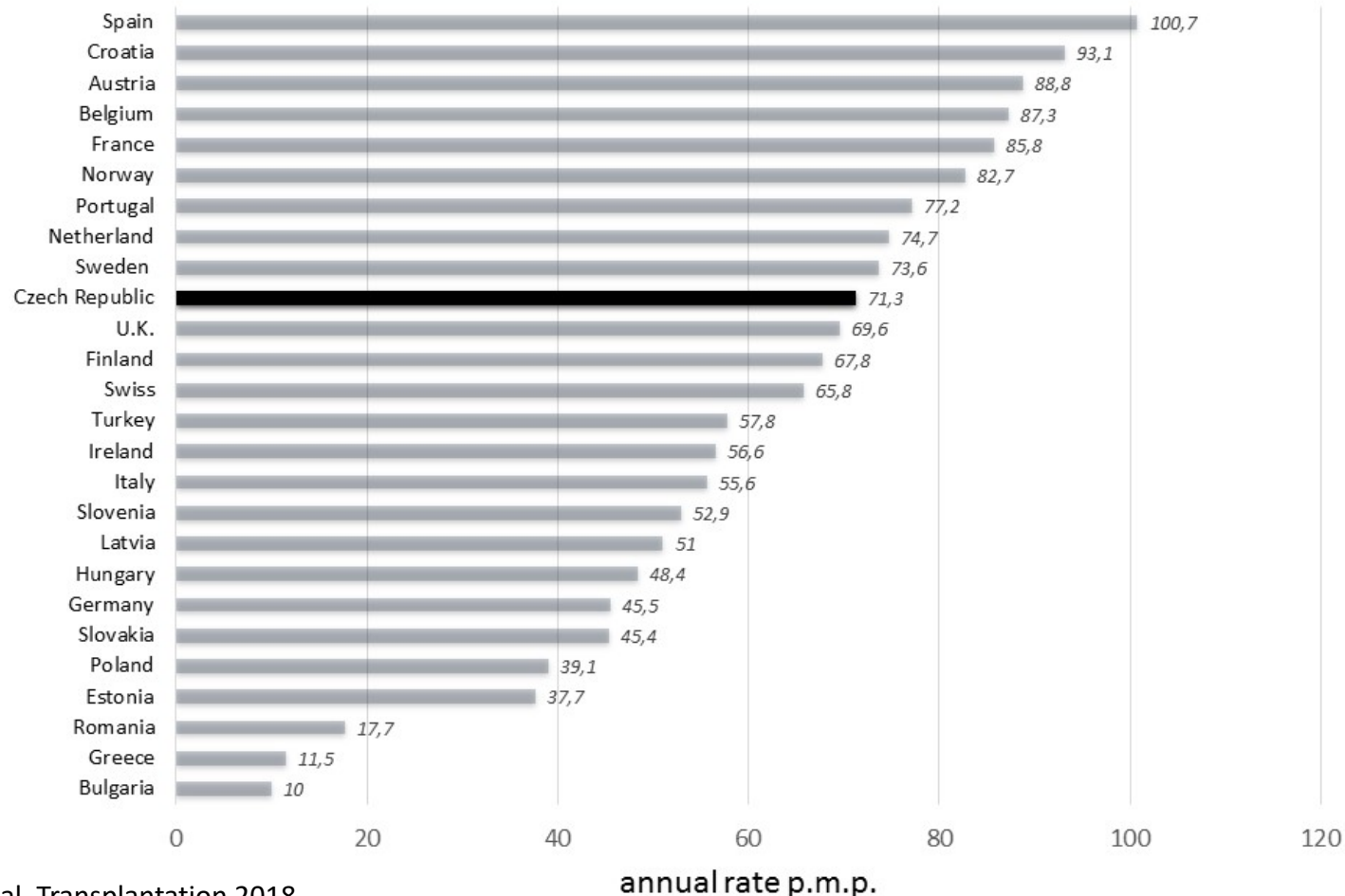
# Imunologické vyšetřování nemocných před a po transplantaci ledviny

Ondřej Viklický

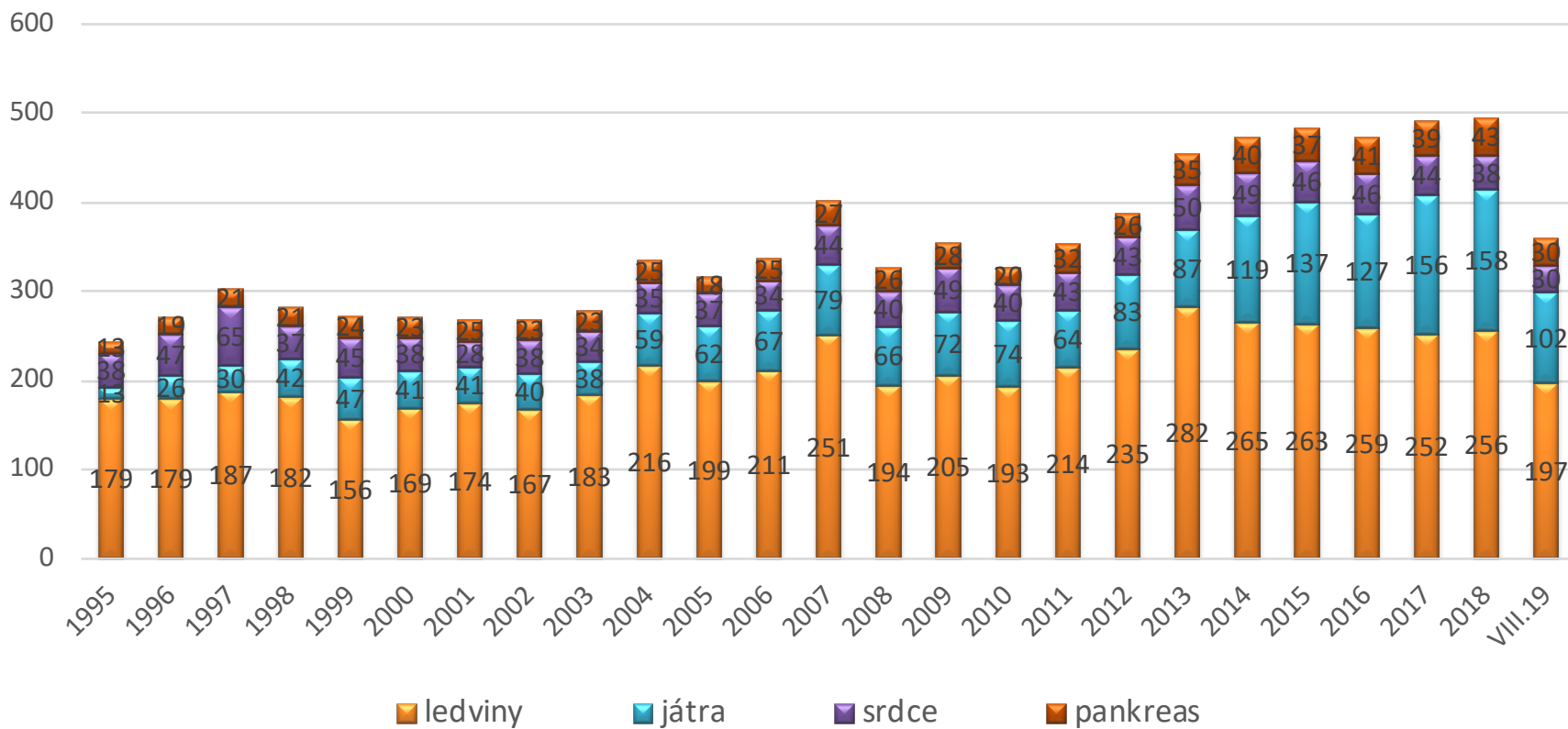
Klinika nefrologie IKEM, Praha

# Dostupnost transplantací je v ČR na vysoké úrovni

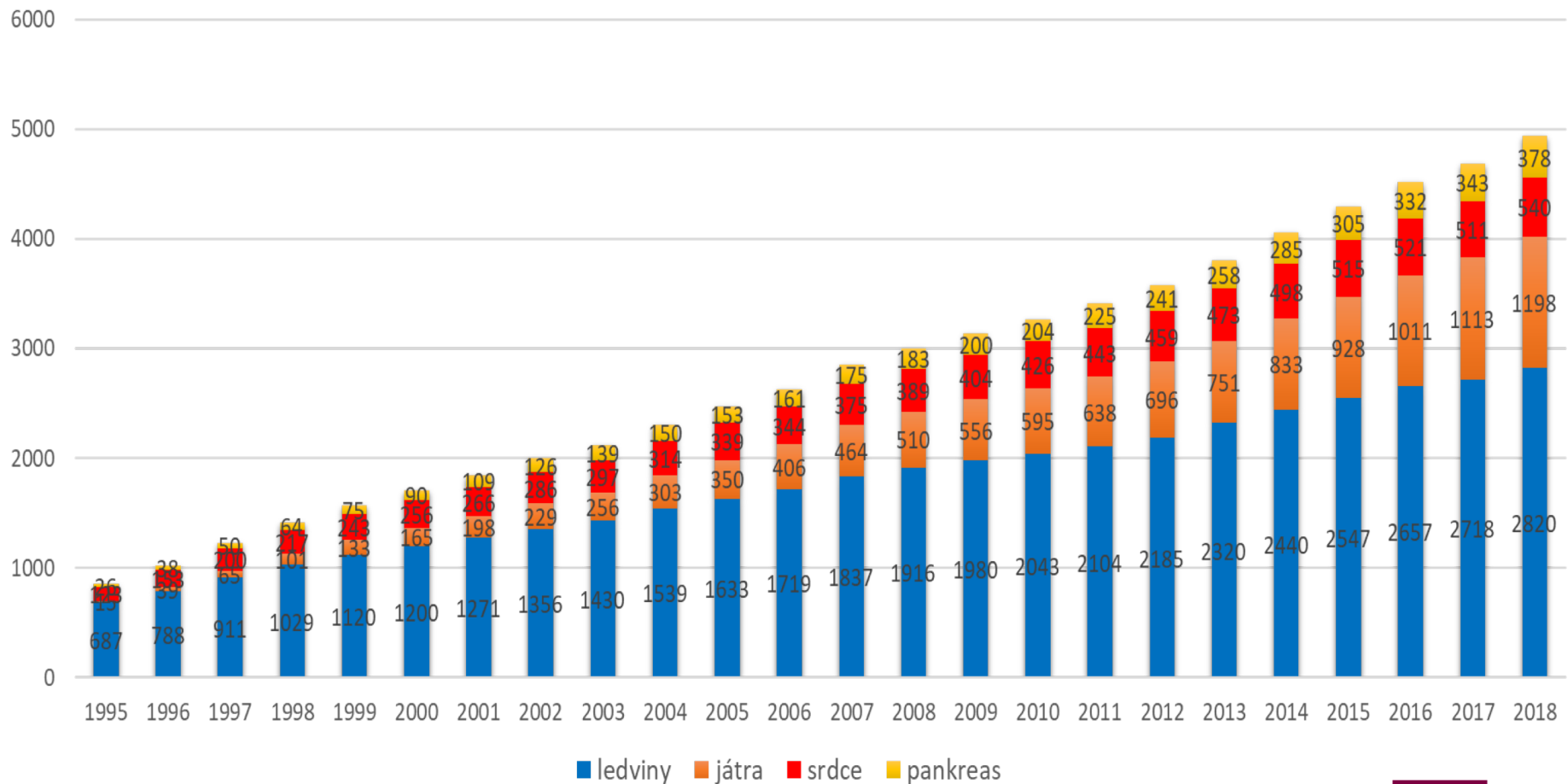
Total number of patients transplanted  
Annual rate p.m.p. 2015



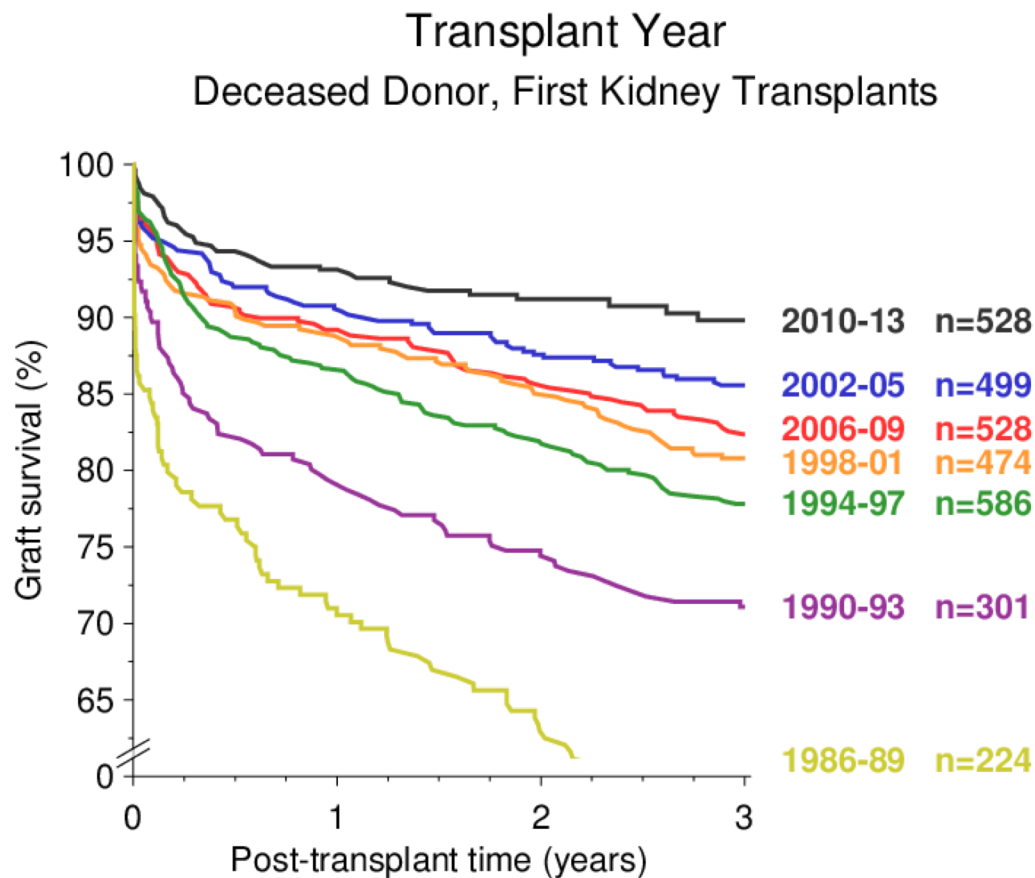
# Transplantace orgánů v TC IKEM (1995-8/2019)



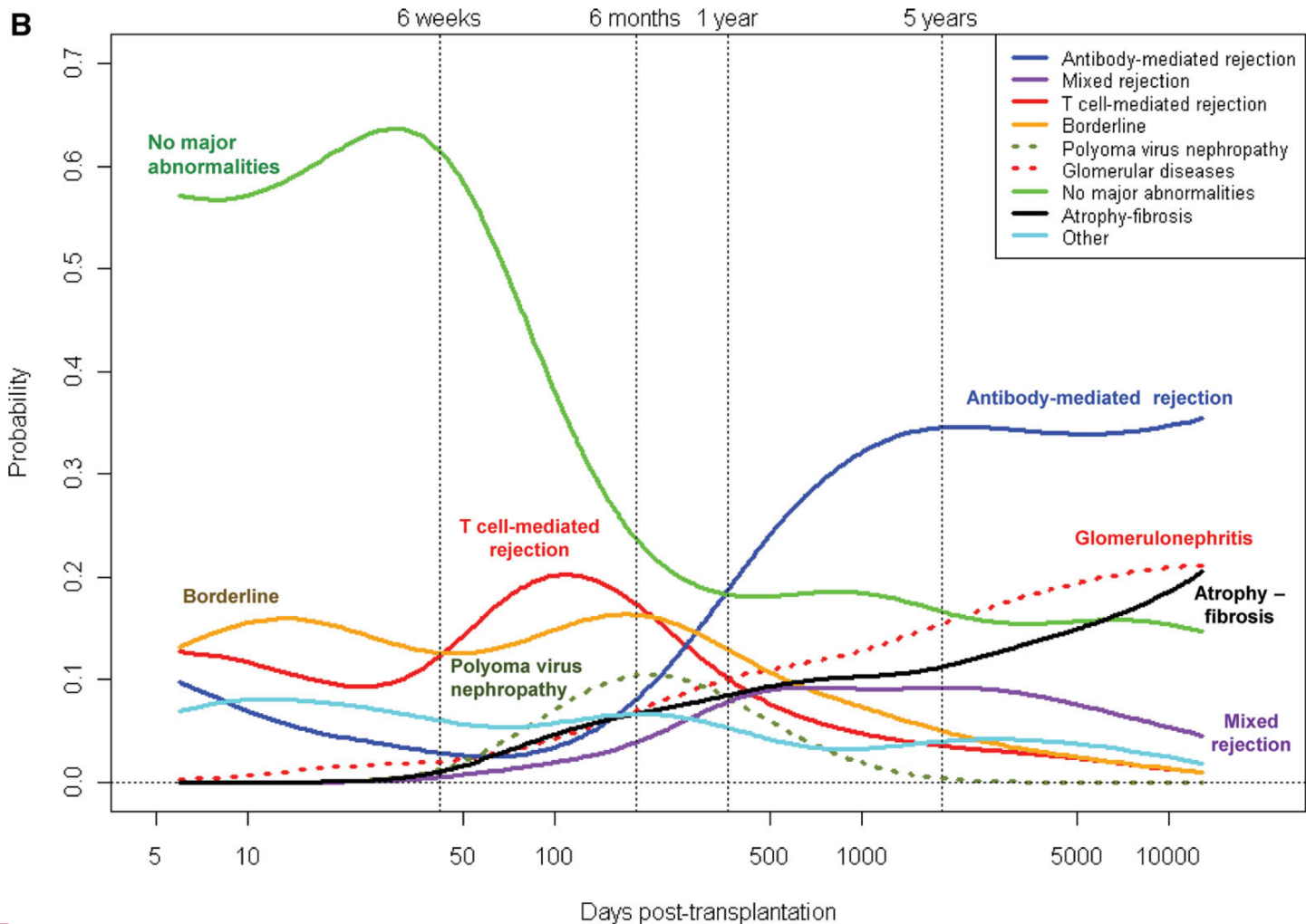
# Nemocní žijící s funkčními transplantovanými orgány v IKEM



# Zlepšení výsledků transplantací ledvin



# Výskyt patologií štěpu po transplantaci

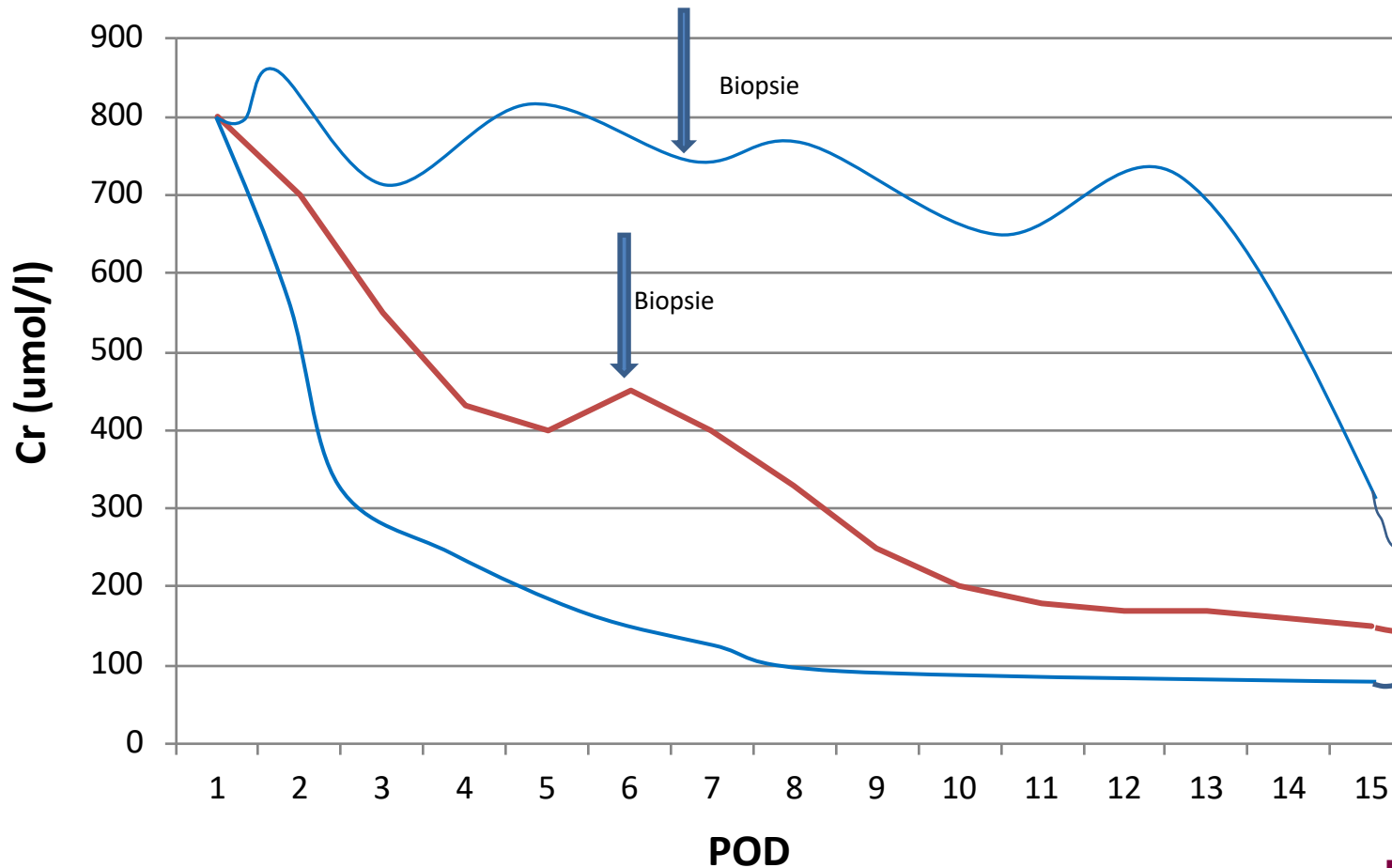


# Vyšetření imunitní odpovědi

---

- Před transplantací
- Po transplantaci
  
- Invazivní – protokolární biopsie
- Neinvazivní – z krve/moče

# Rejekce nebo ATN?

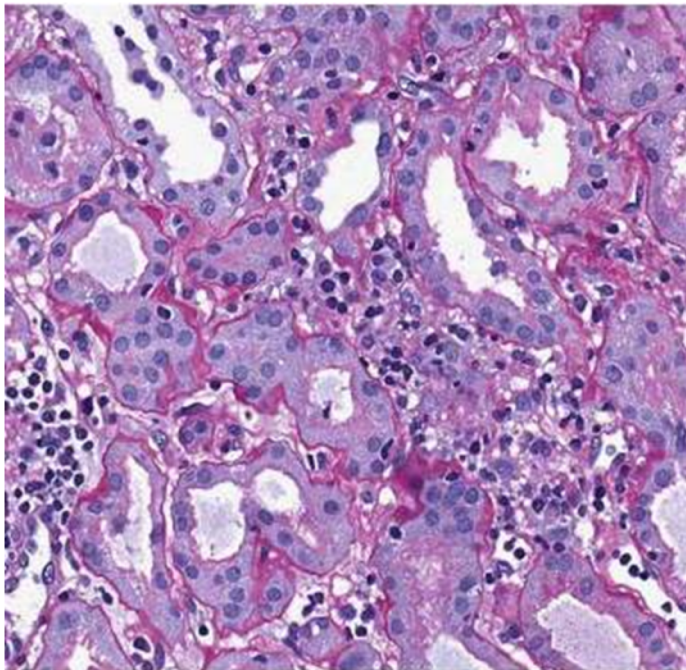


# Borderline changes

## Banff's classification

histological changes insufficient for a diagnosis of acute rejection.

Absence of intimal arteritis and the presence of tubulitis (t1,t2 or t3) with minor interstitial infiltration ((i0, i1) or interstitial infiltration(i2, i3) with mild tubulitis (t1)



## •Incidence : in case biopsies 23% of renal allografts

Meehan et al. (1999) J Am Soc Nephrol 10: 1806

Matoza et al. (2008) Transplantation Proceedings 40: 2303-2306

Viklický et al. (2013) Transplantation 95: 148-154

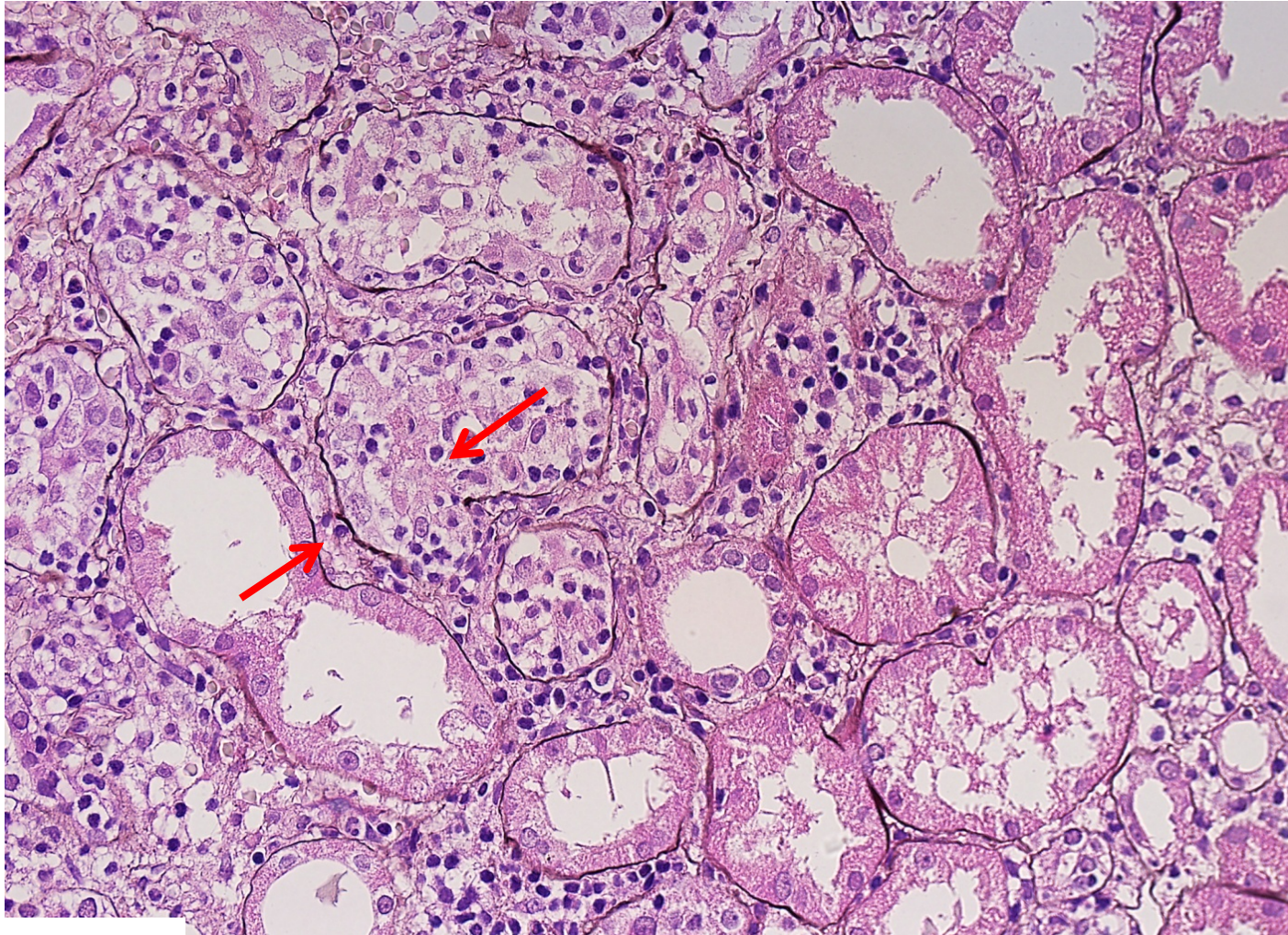
## •The outcome of borderline changes unclear

•The majority of patients with borderline changes were found to be non-rejection by molecular phenotype(67%) and only 33% were rejection-like.

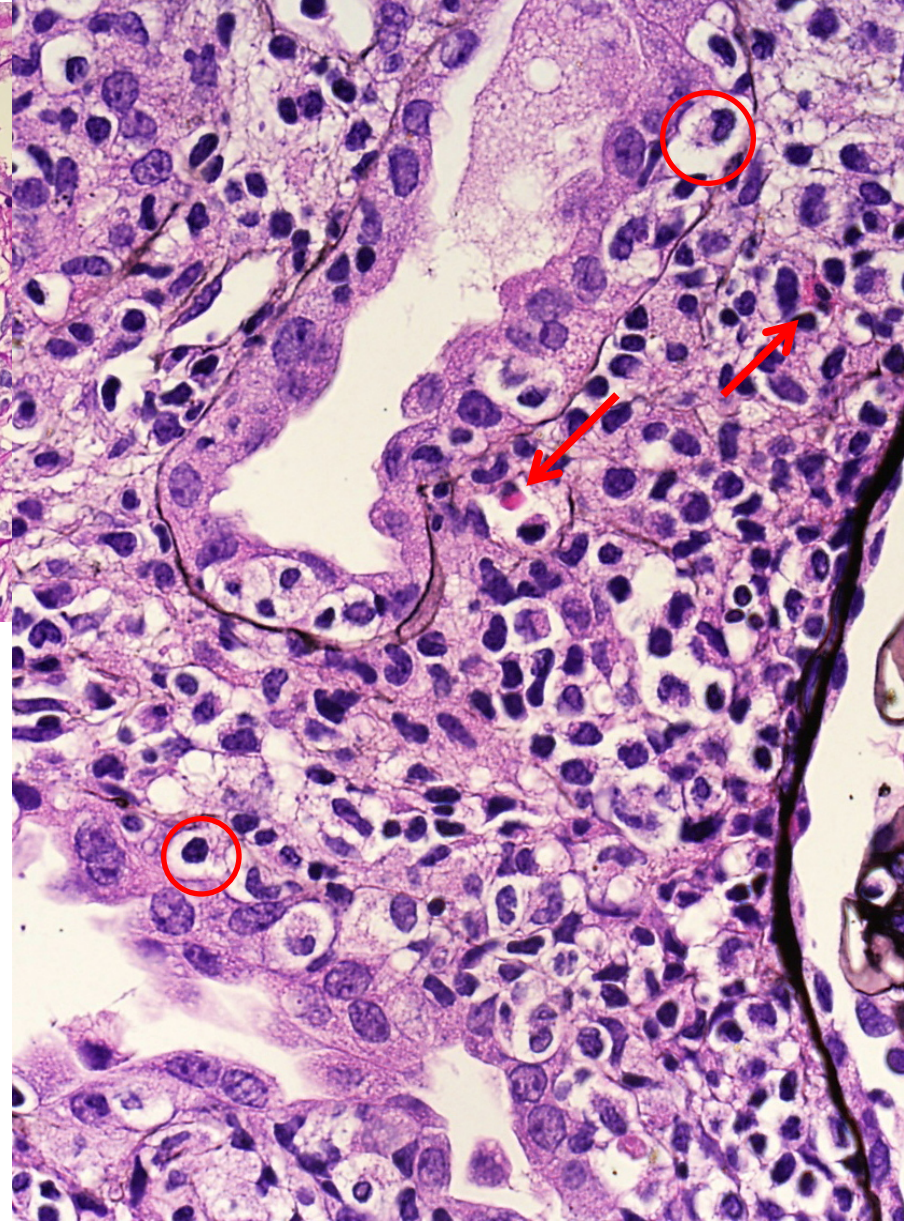
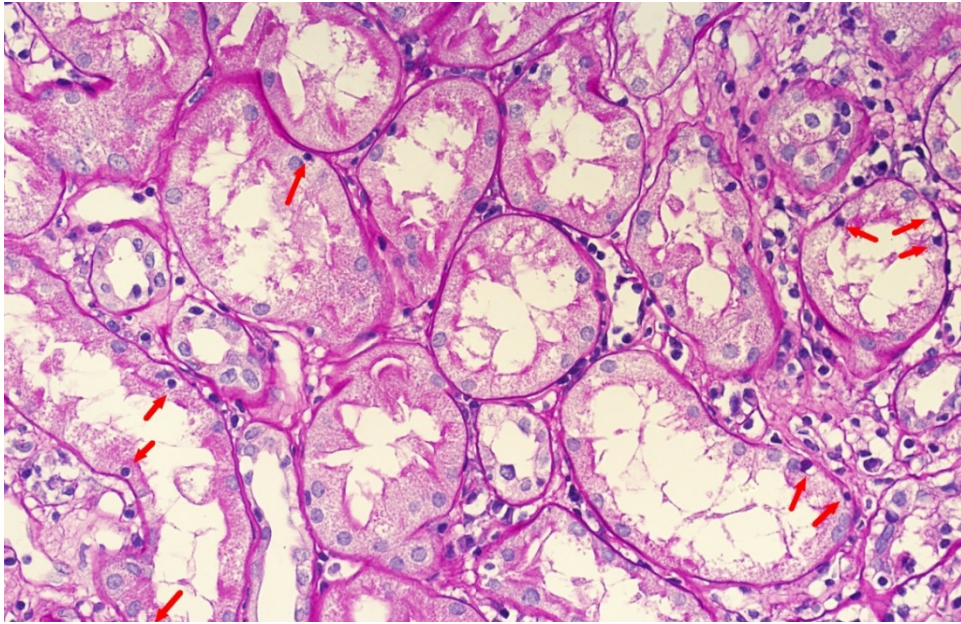
de Freitas et al. (2012) American Journal of Transplantation. 12: 191–201

# Tubulitis

---



# Acute T-cell-mediated rejection, type I *tubulitis in non-atrophic tubules*



## Banff criteria:

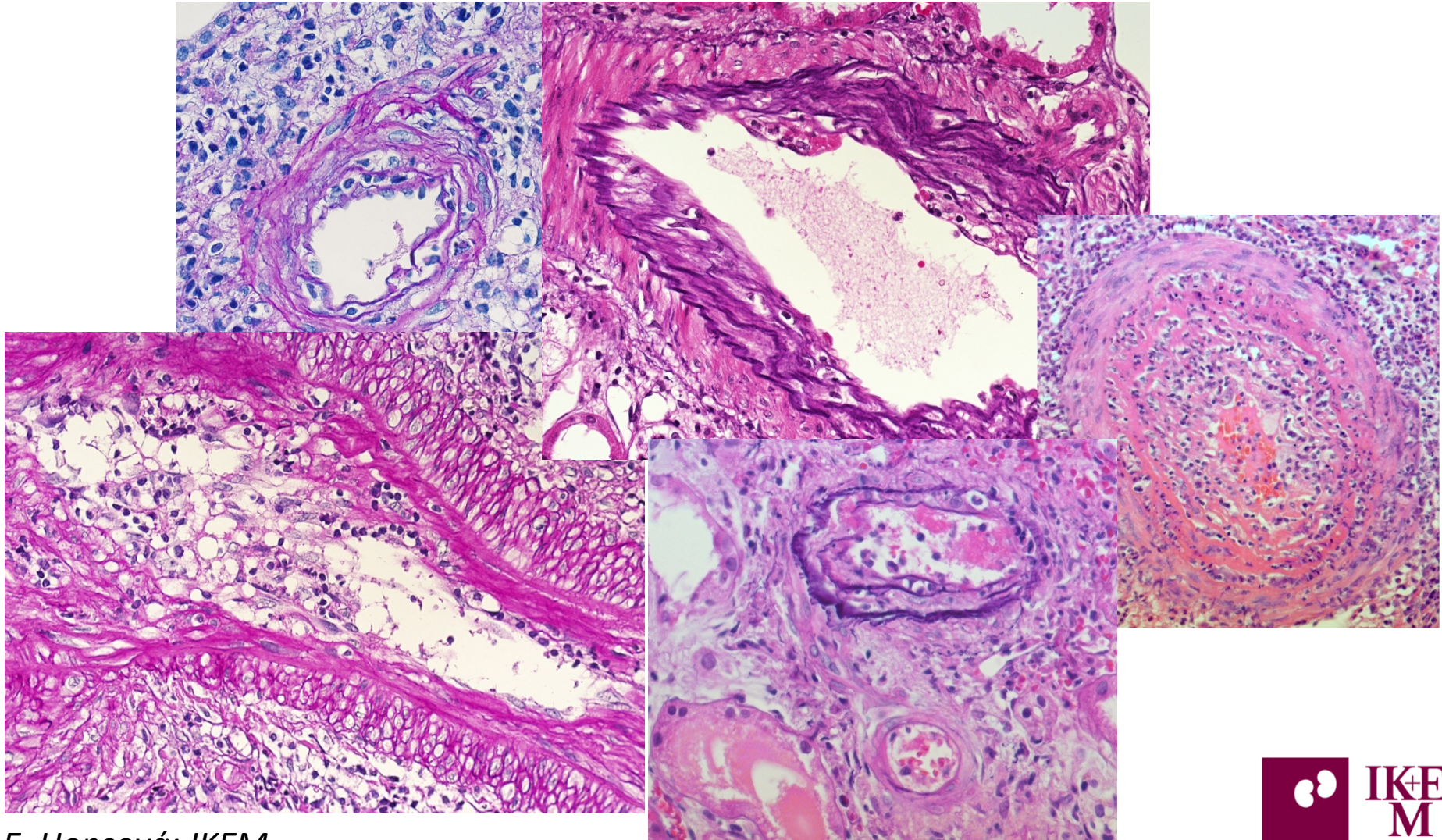
- IA.  $i > 25\%$  of parenchyma affected,  $i_2$  or  $i_3$  and  $t_2$
- IB.  $i > 25\%$  of parenchyma affected,  $i_2$  or  $i_3$  and  $t_3$

## Chapel Hill Standards

1.  $i \geq 5\%$
  2. Mild to moderate interstitial edema
  3. Tubulitis ( $t_1$  in  $\geq 3$  tubules in most inflamed area),
    - scattered eosinophils and ATI are common
    - MHC class II in tubular cells
- If criteria 1-3 are not fulfilled, but tubules strongly expressed MHC class II, then an episode of ACR is suggested

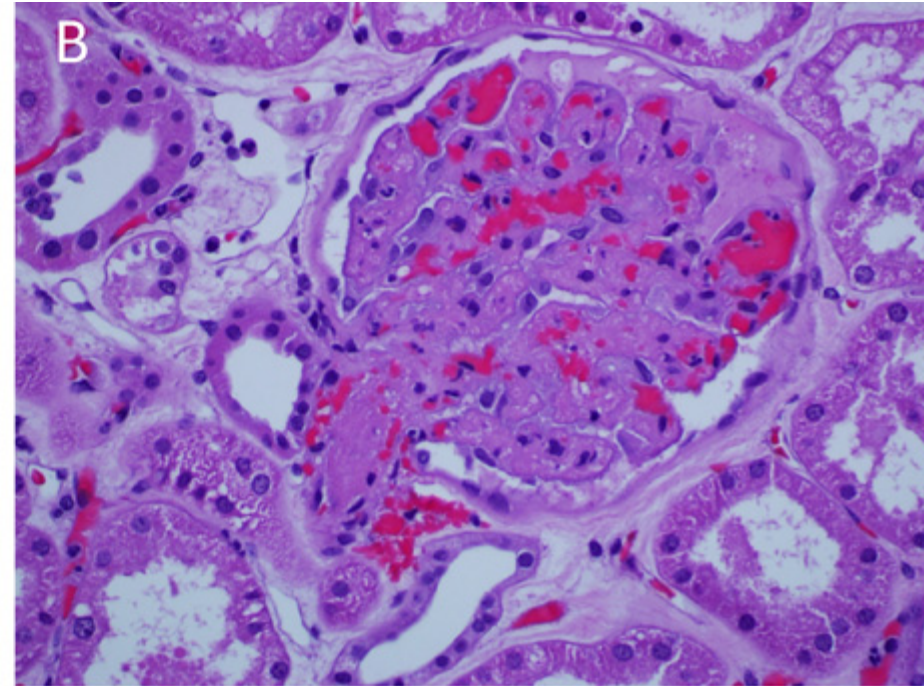
*E. Honsová; IKEM*

# Acute T-cell-mediated rejection, type II and III *arteritis or endotheliitis*



# Makroskopické a mikroskopické projevy hyperakutní rejekce

---



# Diagnostika akutní AMR

## 2002

- Diagnostika vyžaduje přítomnost 3 kritérií
  - I. Dysfunkce štěpu
  - II. Histologické známky AMR
    - C4d depozita v peritubulárních kapilárách (PTC)
    - Neutrofily v PTC a glomerulárních kapilárách
    - Tubulitis s neutrofily
    - Fibrinoidní nekrosy v arteriích a glomerulech
  - III. Detekovatelné dárcovsky specifické protilátky (DSA)

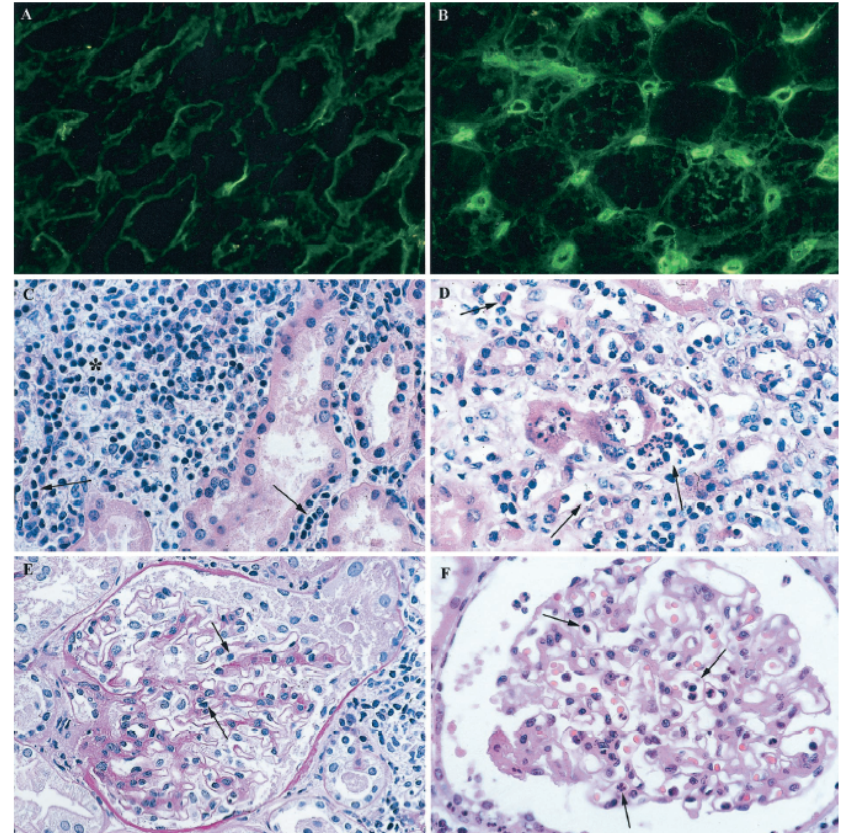
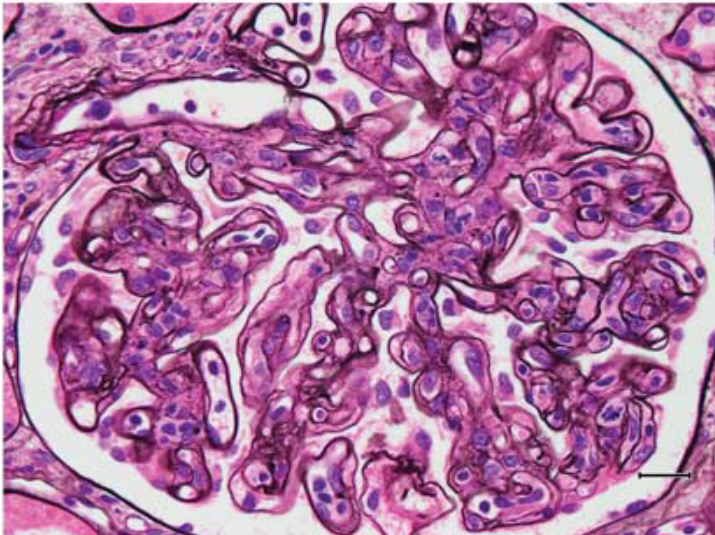
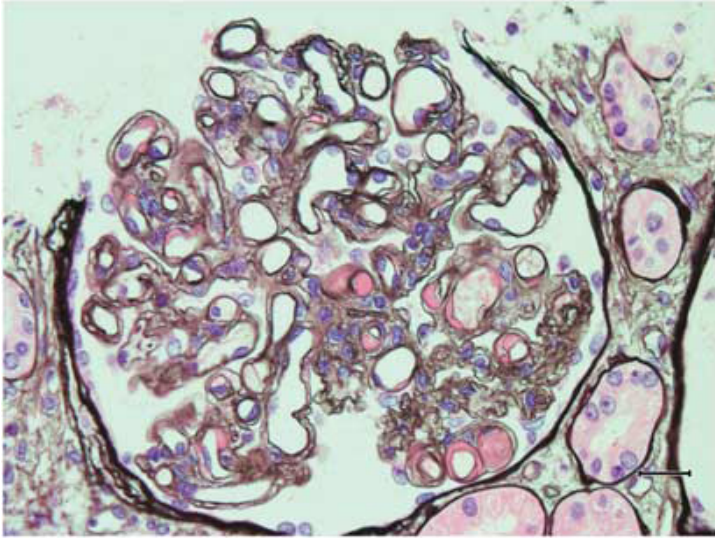


Figure 1. (A) Acute cellular rejection (ACR): no staining for C4d is seen in peritubular capillaries. (B) Acute humoral rejection (AHR): widespread and bright staining for C4d is present in the peritubular capillaries that are interspersed in between the silhouettes of tubules. (C) ACR: mononuclear cells are present in the interstitium (\*) and in peritubular capillaries (arrows). (D) AHR: abundant neutrophils are present in dilated peritubular capillaries (arrows). (E) ACR: scattered mononuclear cells are present in glomerular capillaries (arrows). (F) AHR: neutrophils are present in glomerular capillaries (arrows). Staining: C4d-FITC in A and B; Hematoxylin and eosin (H&E) in C, D, and F; and periodic acid-Schiff (PAS) in E. Magnifications:  $\times 400$  in A through D;  $\times 450$  in E and F.

# Transplantační glomerulopatie nespecifický obraz poškození?



- Protilátkami zprostředkovaná rejekce
- HCV
- Trombotická mikroangiopatie
  
- TG má nejhorší prognosu: 50% ztrát štěpů do 5 let

Haas. Kidney Int 2011; 80, 801 – 803.

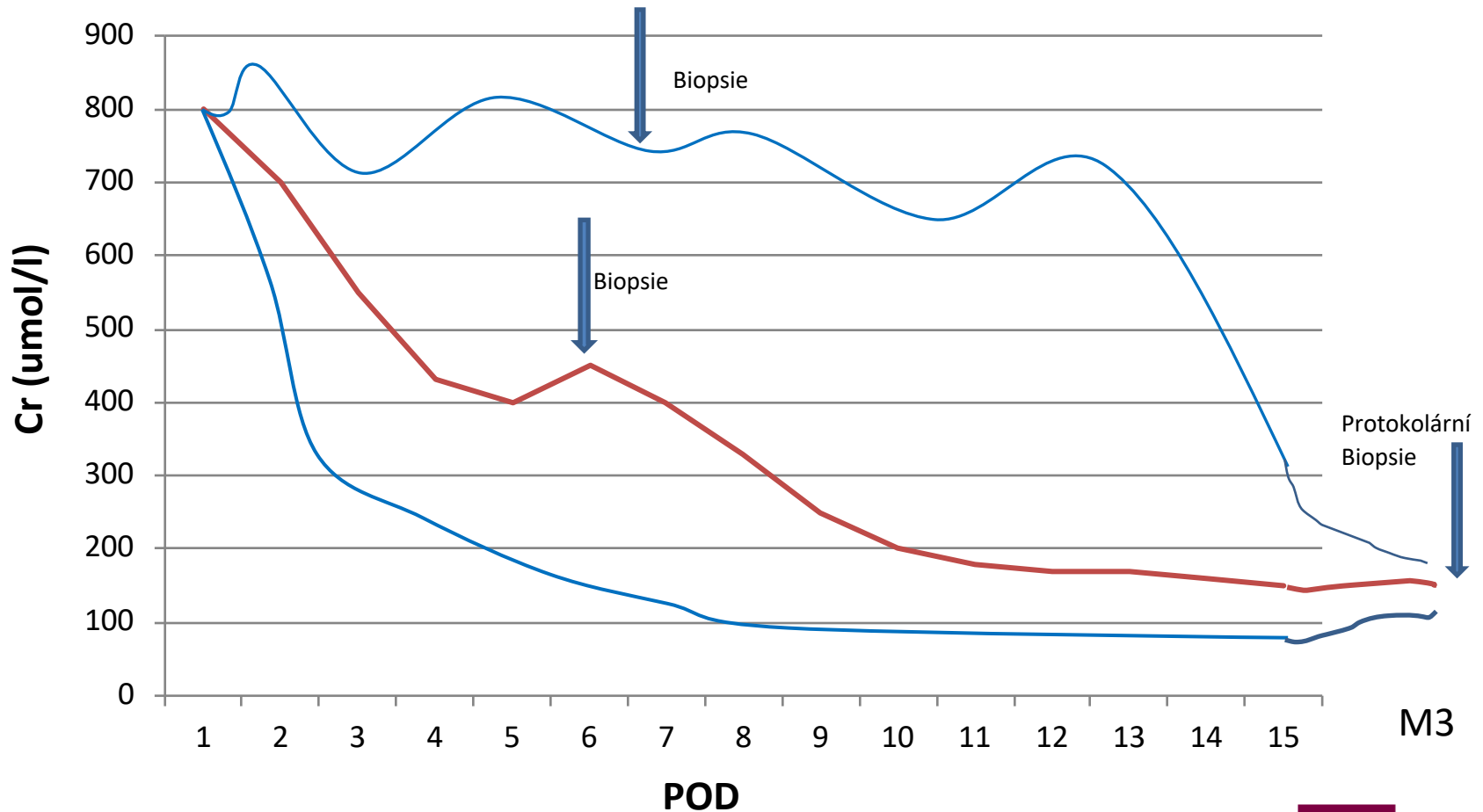
# Klasifikace histologických nálezů transplantované ledviny

---

- Category 1: Normal/nonspecific changes
- Category 2: Antibody-mediated changes
  - Acute/active ABMR
  - Chronic active ABMR
  - C4d staining without evidence of rejection
- Category 3: Borderline changes
- Category 4: TCMR
  - Acute TCMR Grade: IA, IB, IIA, IIB, III
  - Chronic active TCMR
- Category 5: Interstitial fibrosis and tubular atrophy
- Category 6: Other changes not considered to be caused by acute or chronic rejection

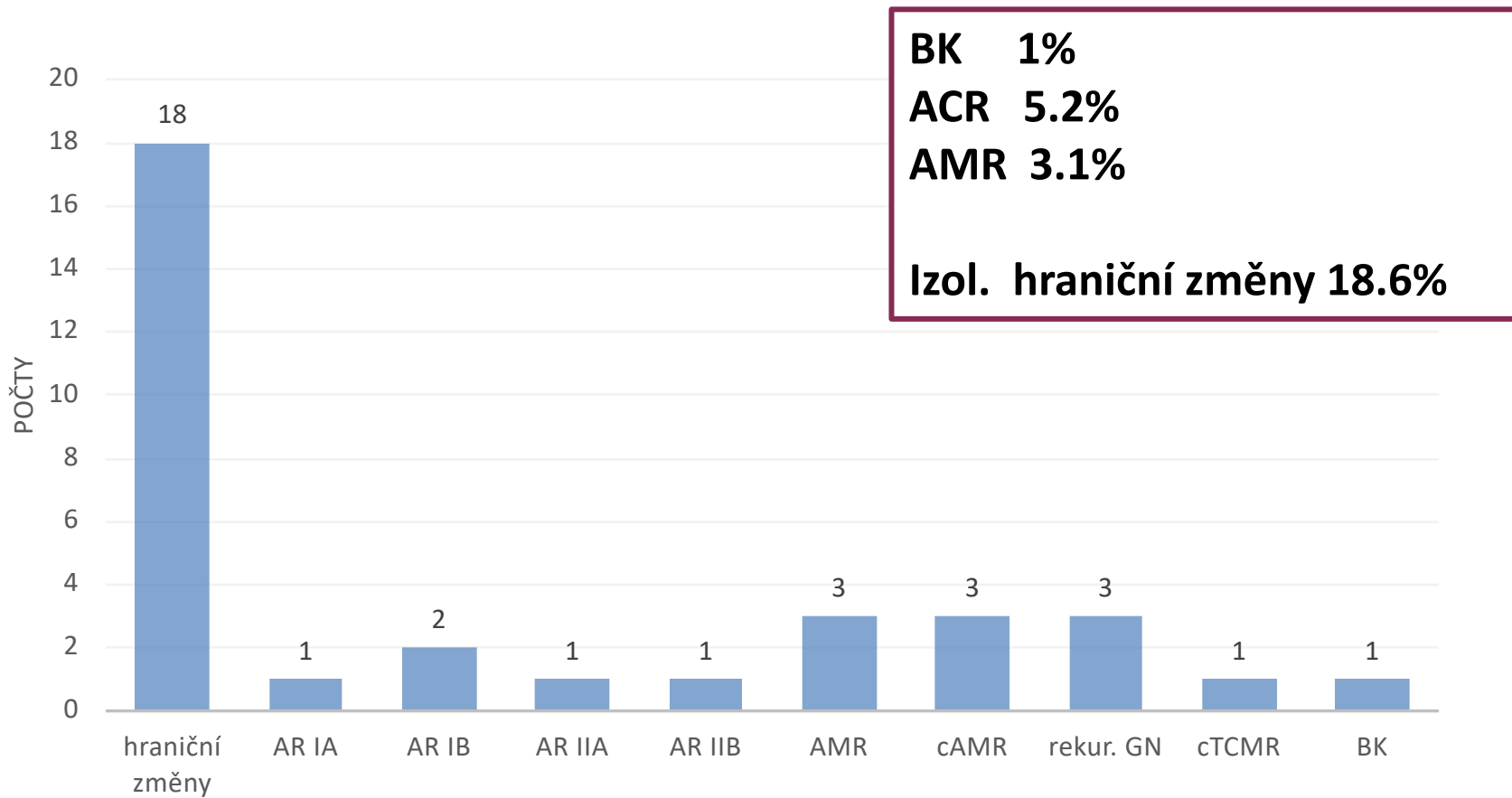
*The Banff 2015 Kidney Meeting Report, American Journal of Transplantation 2017; 17: 28–41*

# Rejekce nebo ATN?

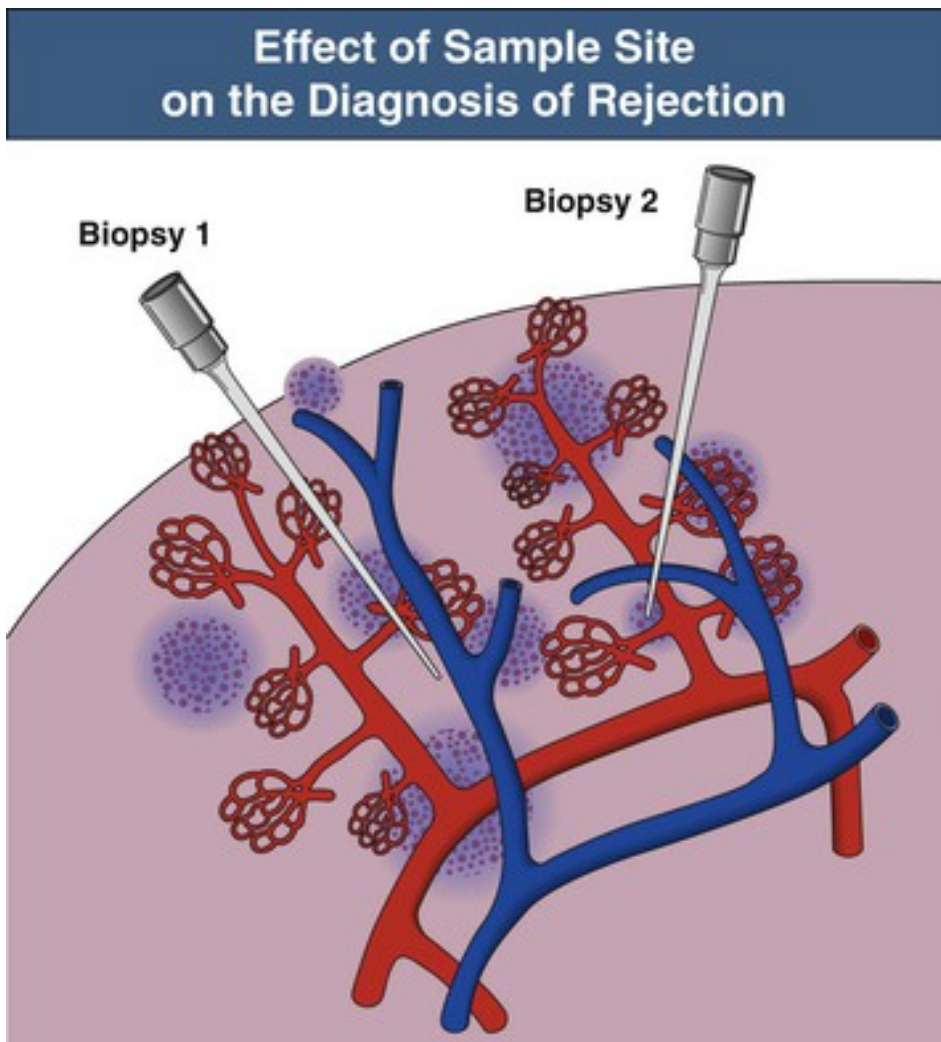


# Invazivní monitorace imunitní odpovědi: protokolární biopsie ve 3. měsíci

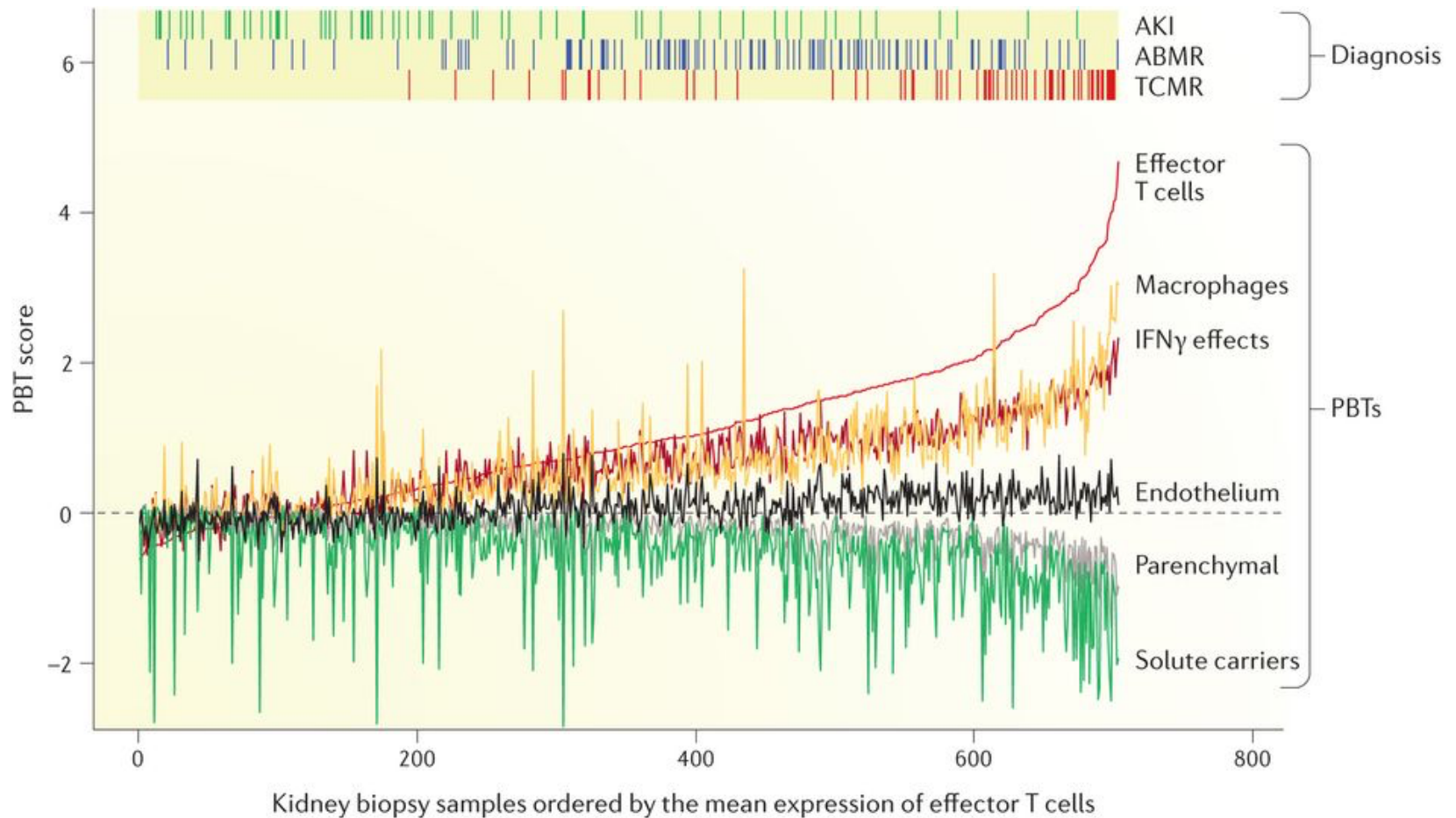
*rejekce přítomna v 8% případů*



# Possible sampling error in needle biopsies



# Molecular assesment allows diagnostics based on specific transcripts



# Explaining the Molecular Microscope® report for core kidney transplant biopsies (MMDx-Kidney)

**Patient information**  
Date of transplant, date of biopsy, etc.

**Clinical information**  
Time post-transplant, biopsy indication, DSA (if provided)

**Additional detail**  
Rejection, injury-related binary classifiers and AKI transcript set

**Comparison to normal**  
Scores of this biopsy interpreted vs. relatively normal biopsies



**Clinical interpretation**

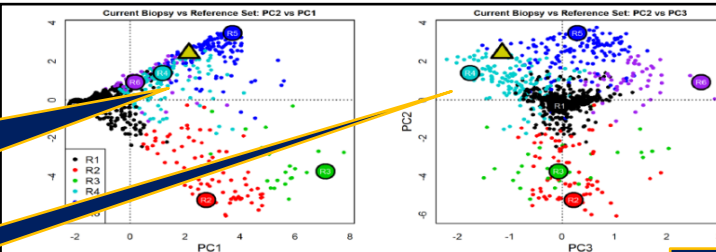
Molecular Microscope® Diagnostic Report for Kidney (MMDx-Kidney)

General Information:	
Surname	Unguran
Date of Birth	(Redacted)
Date Received (Y-M-D)	(Redacted)
Date Reported (Y-M-D)	(Redacted)
Date of Transplant (Y-M-D)	(Redacted)
Date of Biopsy (Y-M-D)	(Redacted)

**Pure molecular interpretation**  
Abnormal biopsy. Severe early-stage ABMR with g and ptc-related molecular features. No TCMR, Mild inflammation, AKI and atrophy-fibrosis. Note that MMDx cannot exclude primary renal diseases.

	Classifier/gene sets <sup>1,2</sup>	Biopsy	Range of values <sup>3</sup>	Upper limit of normal <sup>4</sup>	Interpretation
Injury Scores	Inflammation Score <sup>5</sup>	-0.32	-3.8 – 5.8	0.03	Mild
	Acute Kidney Injury (AKI) Score <sup>4</sup>	0.16	-0.6 – 1.6	0.39	Mild
	Atrophy-Fibrosis Score <sup>5</sup>	0.33	0.0 – 1.0	0.82	Mild
Rejection Scores	Rejection Score <sup>6</sup>	0.74	0.0 – 1.0	0.30	Severe
	T Cell-Mediated Rejection (TCMR) Score <sup>6,7,8</sup>	0.01	0.0 – 1.0	0.10	Normal
	Antibody-Mediated Rejection (ABMR) Score <sup>6,9,10</sup>	0.81	0.0 – 1.0	0.20	Severe

Rejection phenotype <sup>8,10</sup> (six scores, R1-R6, adding up to 1.0)	Score	Reference Set	Score
R1 Non-rejecting	0.00	All ABMR (Sum of R4, R5, and R6)	1.00
R2 TCMR	0.00	R4 Early-Stage ABMR (EABMR)	0.59
R3 Mixed Rejection	0.00	R5 Fully-Developed ABMR (FABMR)	0.41
		R6 Late-Stage ABMR (LABMR)	0.00



Survival in patients with similar biopsies in the Reference Set		Percent cortex <sup>10,11</sup>
1-year: 92%	3-years: 76%	96%

Clinical Notes

## Molecular Microscope® Diagnostic Report for Kidney (MMDx-Kidney)

	Classifier/gene sets	Biopsy score	Range of possible values <sup>3</sup>	Upper limit of normal <sup>4</sup>	Interpretation
TCMR related	TCMR-1 <sup>6</sup>	0.01	0.0 – 1.0	0.10	Normal
	TCMR-2 <sup>6</sup>	0.01	0.0 – 1.0	0.10	Normal
	Mean of 2 TCMR classifiers	0.01	0.0 – 1.0	0.10	Normal
Rejection related	Rejection <sup>6</sup>	0.74	0.0 – 1.0	0.30	Severe
Injury-scarring related	AKI score <sup>4</sup>	0.16	-0.6 – 1.6	0.39	Mild
	Atrophy-Fibrosis Score <sup>5</sup>	0.33	0.0 – 1.0	0.82	Mild
	ABMR-1 <sup>9</sup>	0.82	0.0 – 1.0	0.20	Severe
ABMR related	ABMR-2	0.77	0.0 – 1.0	0.20	Severe
	ABMR-3	0.84	0.0 – 1.0	0.20	Severe
	Mean of 3 ABMR classifiers	0.81	0.0 – 1.0	0.20	Severe
Classifiers based on histologic lesions	Glomerulitis (g) > 0 probability <sup>7</sup>	0.75	0.0 – 1.0	0.25	Severe
	Transplant glomerulopathy (tg) > 0 probability <sup>7</sup>	0.33	0.0 – 1.0	0.22	Mild
	Peritubular capillaritis (ptc) > 0 probability <sup>7</sup>	0.75	0.0 – 1.0	0.24	Severe
	DSA-positive probability	0.64	0.0 – 1.0	0.42	Moderate
	Interstitial inflammation (i) > 1 probability <sup>7</sup>	0.02	0.0 – 1.0	0.06	Normal
	Tubulitis (t) > 1 probability <sup>7</sup>	0.03	0.0 – 1.0	0.1	Normal
	Tubular atrophy (ct) > 1 probability	0.21	0.0 – 1.0	0.84	Normal
	Adherence index <sup>11</sup>	0.45	0.0 – 1.0	0.9	Normal

For classifiers: TCMR-1 = TCMR vs everything else; TCMR-2 = TCMR vs everything else, with BK/Borderline/Mixed withheld; ABMR-1 = ABMR vs everything else with TG/ABMR suspicious withheld; ABMR-2 = ABMR and Mixed vs everything else, with TG/ABMR suspicious withheld; ABMR-3 = ABMR vs everything else, with Mixed/TG/ABMR suspicious withheld.

Rank order of the most common histologic diagnoses in the 50 nearest molecular neighbors	Mean molecular scores in the 50 nearest molecular neighbors
ABMR: 54%	Rejection: 0.83
No Major Abnormalities (NOMOA): 12%	ABMR: 0.83
Transplant Glomerulopathy (TG): 8%	Atrophy-Fibrosis Score (cigt1): 0.28
Mixed Rejection: 6%	AKI Score (IRATs): 0.20
ABMR Suspicious: 6%	TCMR: 0.02

- References for the scores, classifiers, and archetypes
- Halloran PF et al. Nat Rev Nephrol 2016;12(9):534-544
  - Halloran PF et al. Kidney Int 2014;(85):258-64.
  - Mueller TF et al. Am J Transplant 2007;7(12):212-22.
  - Famulski K et al. JASN 2007;23(5):948-58.
  - Venner J et al. J Clin Invest 2016;126(11):3117-3125
  - Reeve J et al. Am J Transplant 2009 Aug;9(8):1802-10.
  - Reeve J et al. Am J Transplant 2013;13(3):645-55.
  - Sellares J et al. Am J Transplant 2013;13(4):971-83.
  - Reeve J et al. J Clin Invest 2017 Feb 22;127(8):2117-2126
  - Madill-Thomsen K et al. J Clin Invest 2017 Feb 22;127(8):2117-2126

**Summary of molecular changes**  
(Injury, rejection)

**Proportions of rejection-related AA molecular changes**  
(Normal, TCMR, ABMR)

**Visualization**  
Relationship of biopsy to others in reference set  
PC2 vs. PC1

**Visualization**  
PC2 vs. PC3

**Survival of other kidneys like this one**

**% of biopsy that is cortex**

**Histologic and molecular diagnoses in the molecular nearest neighbors of this biopsy**

**Adherence index:** Low scores in biopsies 6m-5y post-transplant correlate with possible non-adherence or under-immunosuppression

# Historical case – MMDx from biobank

---

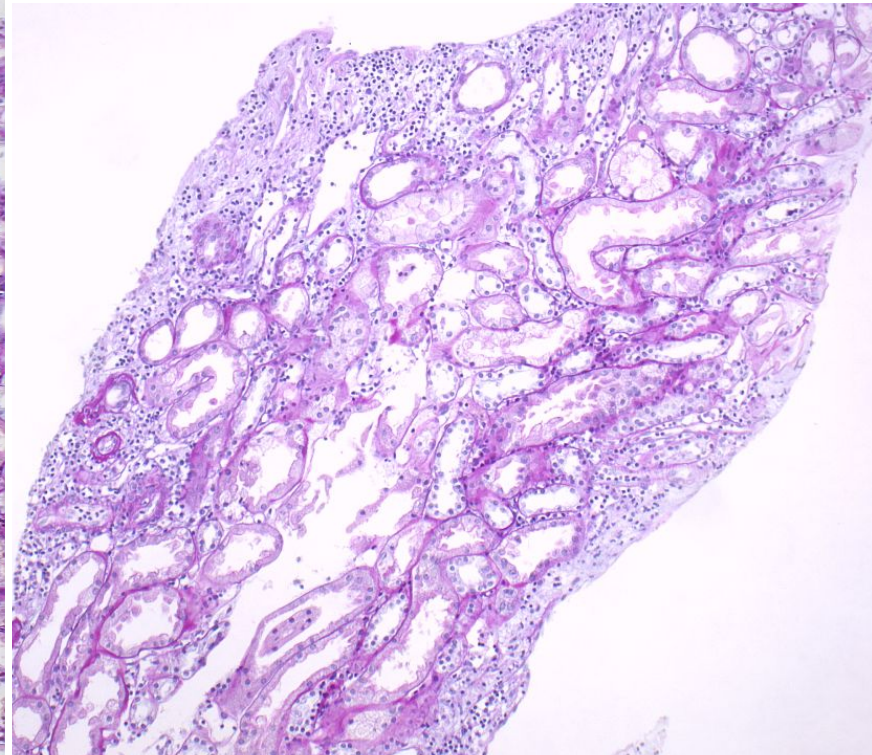
- 41 years, female
- 1st deceased donor kidney transplantation
- ECD, age 65
- DSA negative, peak PRA 4%, HLA mismatch 3
- no induction

7<sup>th</sup> POD: DGF, kidney graft biopsy:

T-cell mediated rejection, IIA with mild interstitial component (C4d negative)

Therapy: 2g of methylprednisolone, S-Cr 200 $\mu$ mol/l (2.26mg/dl)

---



Eva Honsova, IKEM

---

3M protocol biopsy

S-Cr: 156 $\mu$ mol/l (1.76mg/dl)

Interstitial inflammation with tubulitis,  
formation of fibrosis

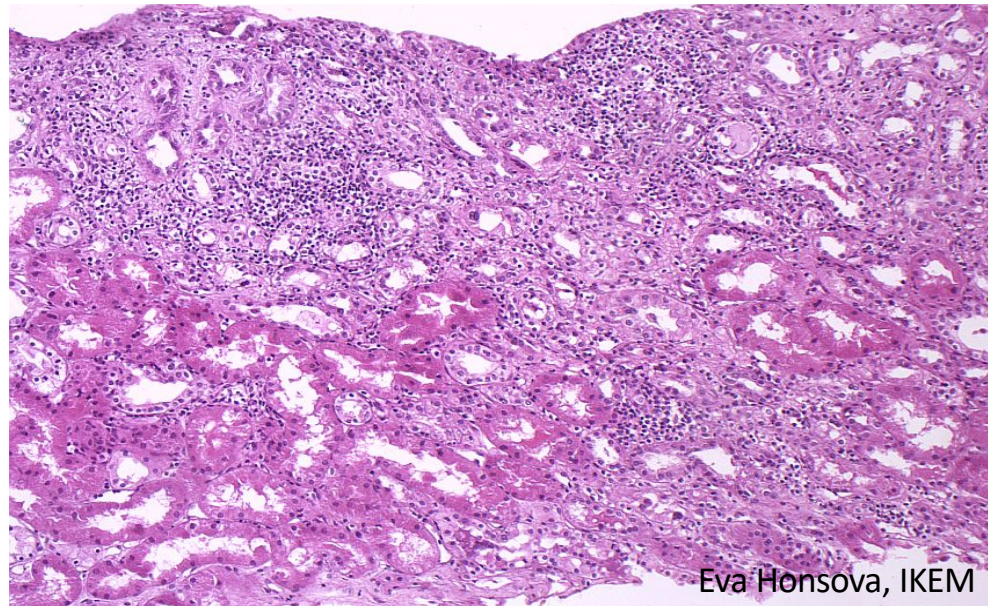
T-cell mediated rejection IA (C4d negative,  
polyoma negative)  
g1 i2 ti2 t2 cv2 ah2

Thymoglobuline 325mg (cumulative dose of  
3.7 mg/kg )

FU: 5.5 years after Tx

Stable S-Cr 178.5  $\mu$ mol/l

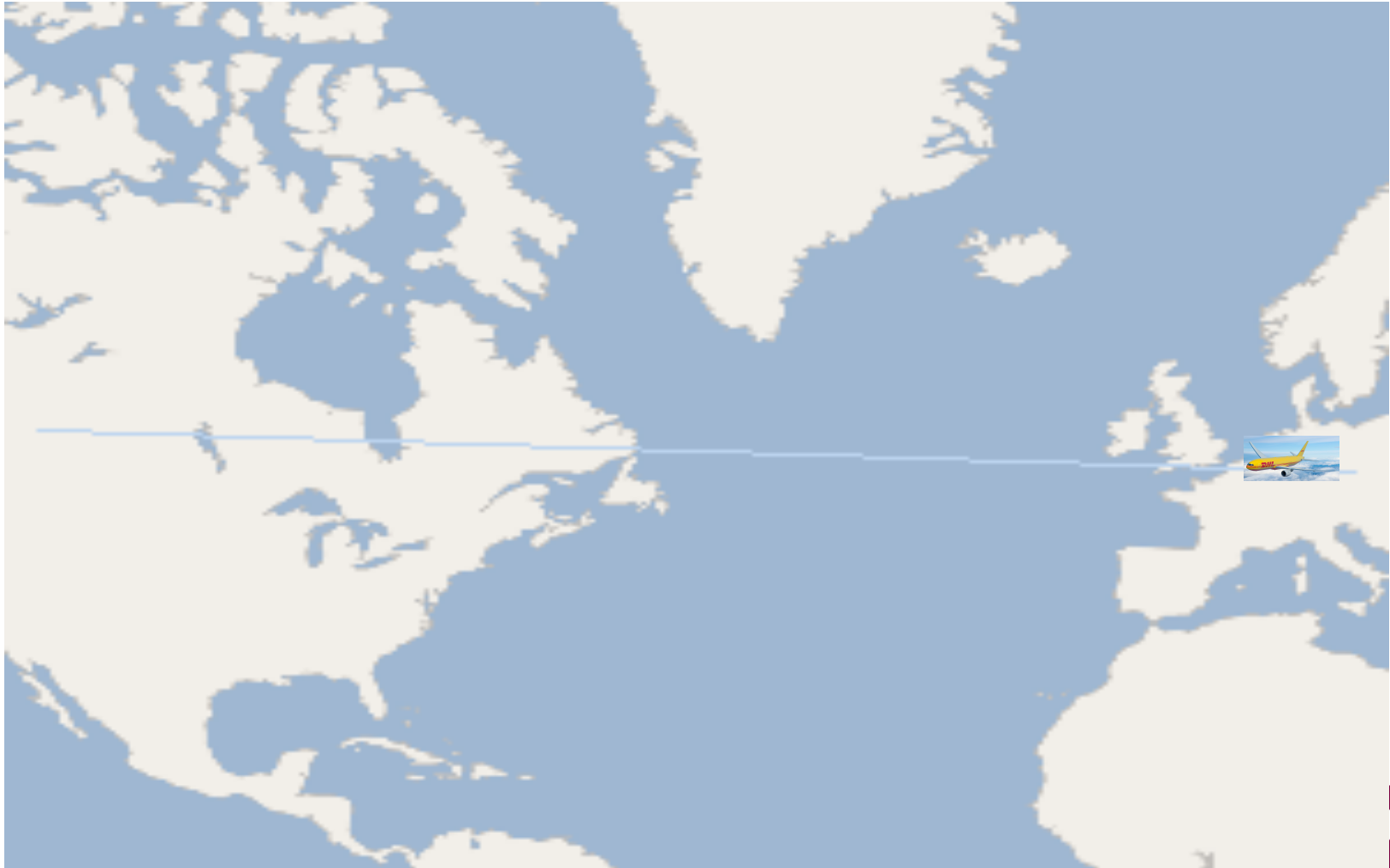
MMDx from the biobank



Eva Honsova, IKEM

# MMDx in Edmonton

---

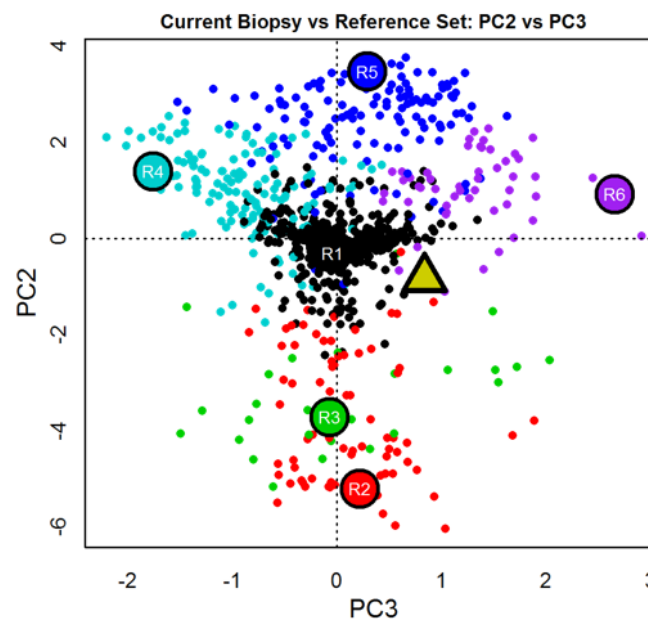
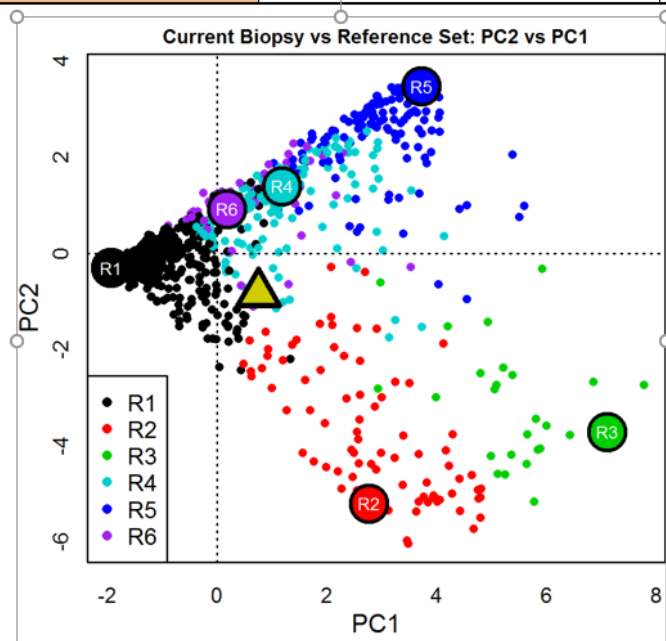


Banff histology  
g1 i2 ti2 t2 cv2 ah2

MMDx from historical  
sample:  
No rejection

	Classifier/gene sets <sup>1,2</sup>	Biopsy	Range of values <sup>A</sup>	Upper limit of normal <sup>B</sup>	Interpretation
Injury Scores	Global Disturbance Score <sup>3</sup>	9.07	-3.8 – 5.8	0.03	<b>Extensive</b>
	Acute Kidney Injury (AKI) Score <sup>4</sup>	0.88	-0.6 – 1.6	0.69	<b>Moderate</b>
	Atrophy-Fibrosis Score <sup>5</sup>	0.16	0.0 – 1.0	0.39	<b>Minimal</b>
Rejection Scores	Rejection Score <sup>6</sup>	0.06	0.0 – 1.0	0.30	<b>Normal</b>
	T Cell-Mediated Rejection (TCMR) Score <sup>7, C</sup>	0.09	0.0 – 1.0	0.10	<b>Normal</b>
	Antibody-Mediated Rejection (ABMR) Score <sup>8, C</sup>	0.07	0.0 – 1.0	0.20	<b>Normal</b>

Rejection phenotype <sup>8, D</sup> (six scores, R1-R6, adding up to 1.0)	R1 Non-rejecting	0.14	All ABMR (Sum of R4, R5, and R6)	0.59
	R2 TCMR	0.27	R4 Early-Stage ABMR (EABMR)	0.19
	R3 Mixed Rejection	0.00	R5 Fully-Developed ABMR (FABMR)	0.00
			R6 Late-Stage ABMR (LABMR)	0.40

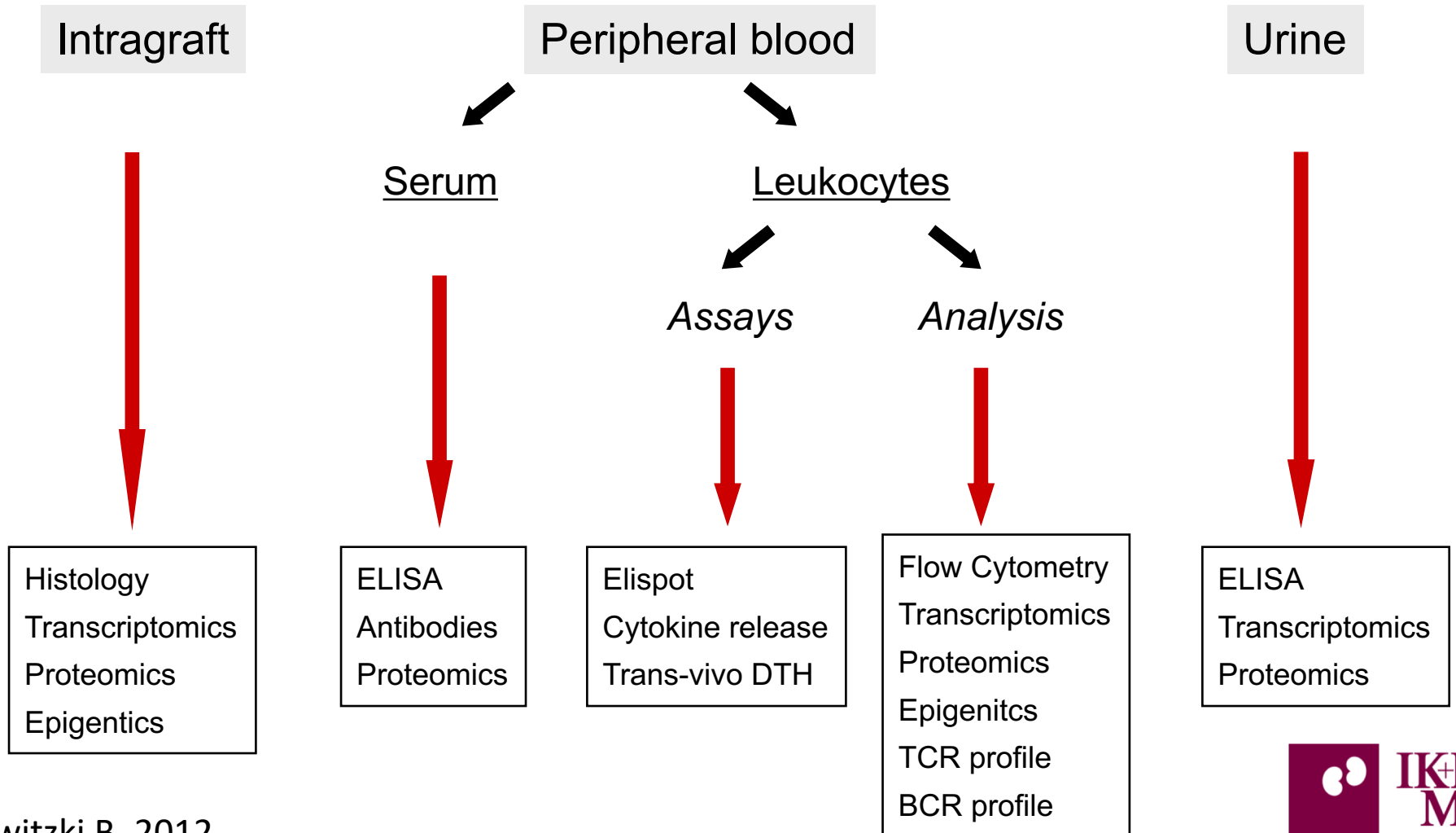


# Možnosti predikce osudu štěpu

---

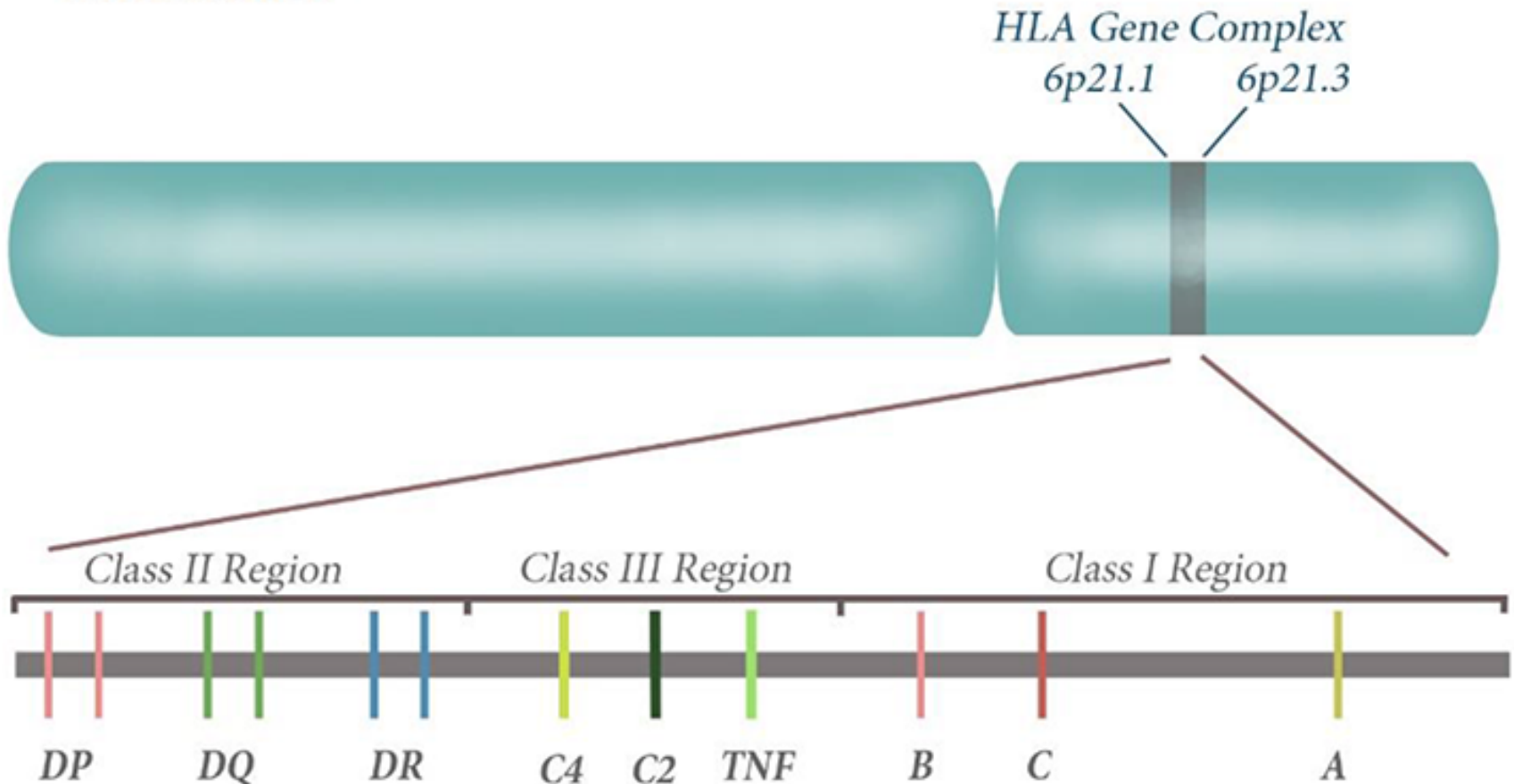


# Potenciální eseje pro před/potransplantační monitorování

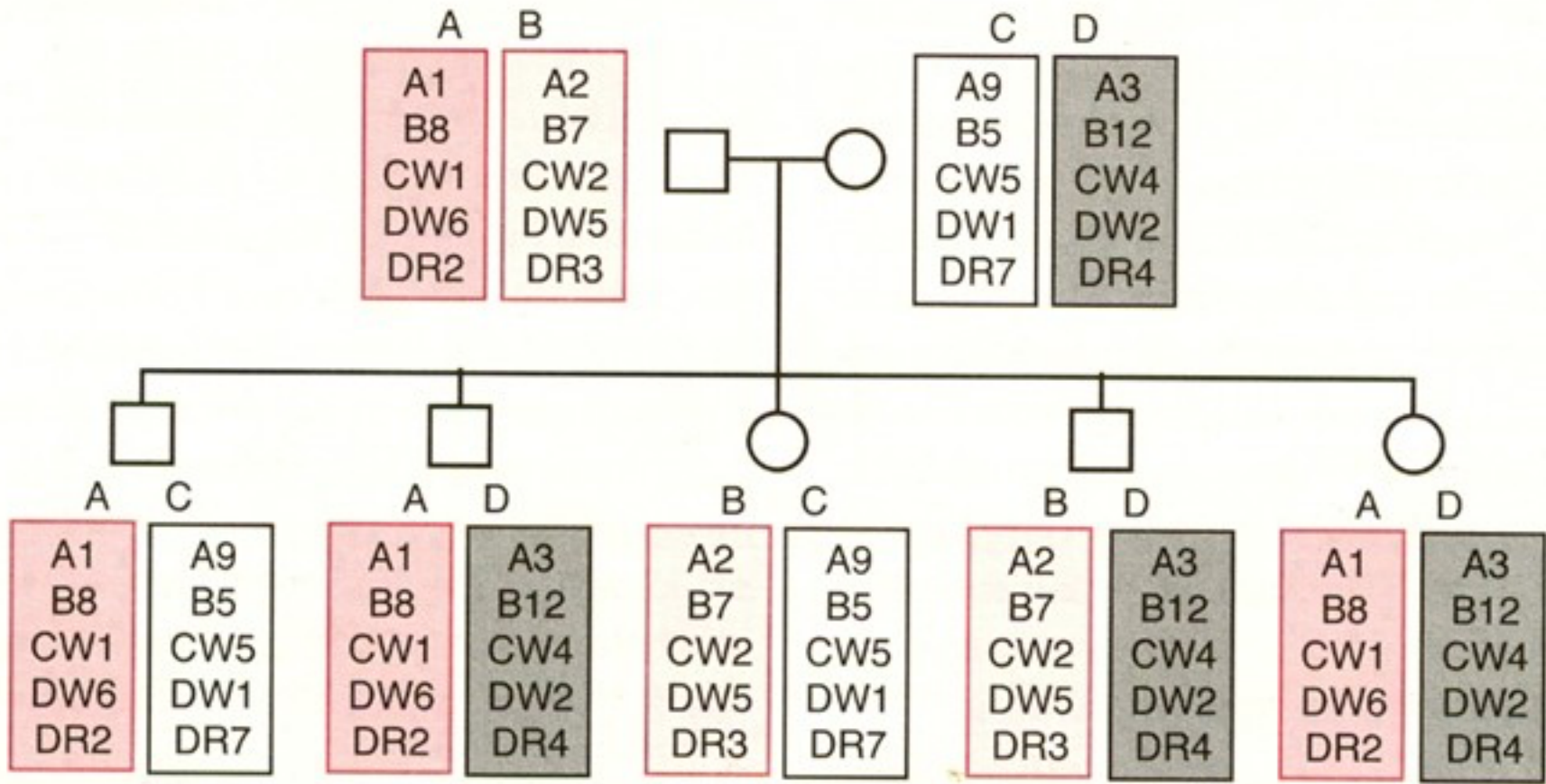


# Lidský HLA systém je kódován na 6 chromozomu

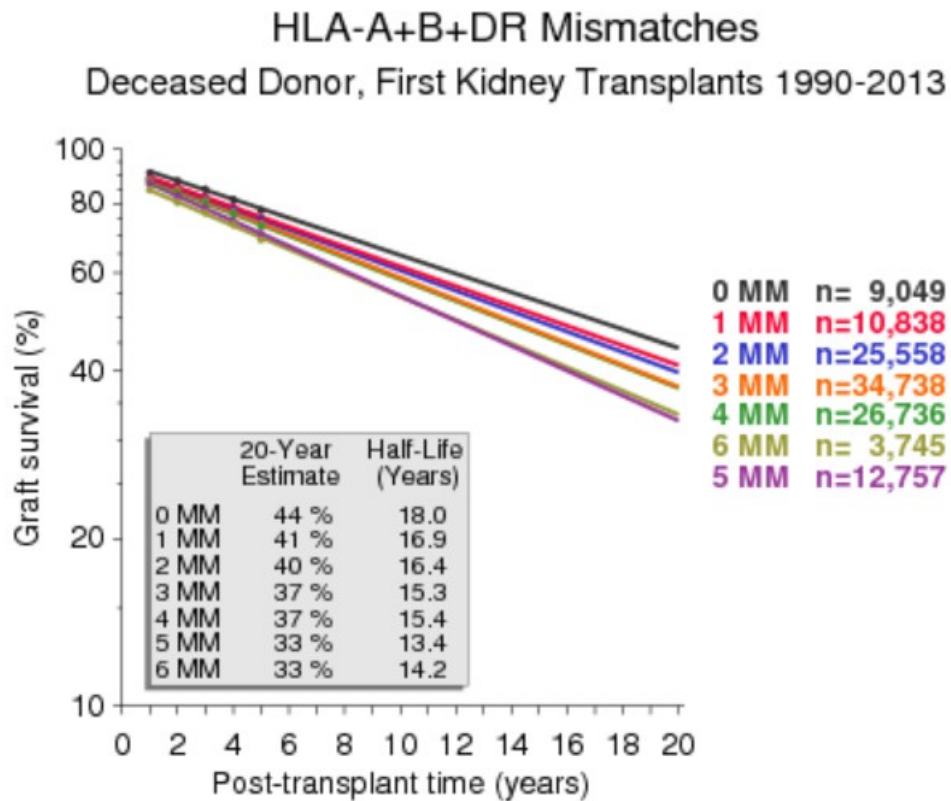
*Chromosome 6*



# Dědičnost HLA haplotypů



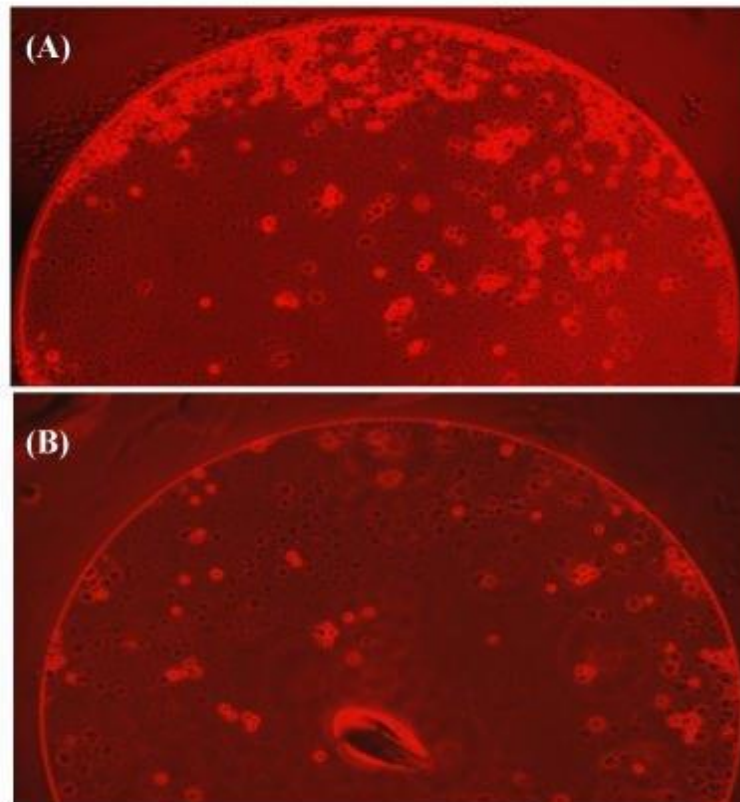
# Více je lépe...



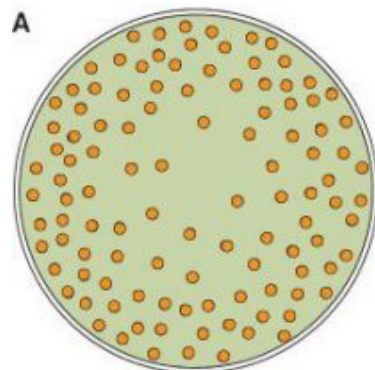
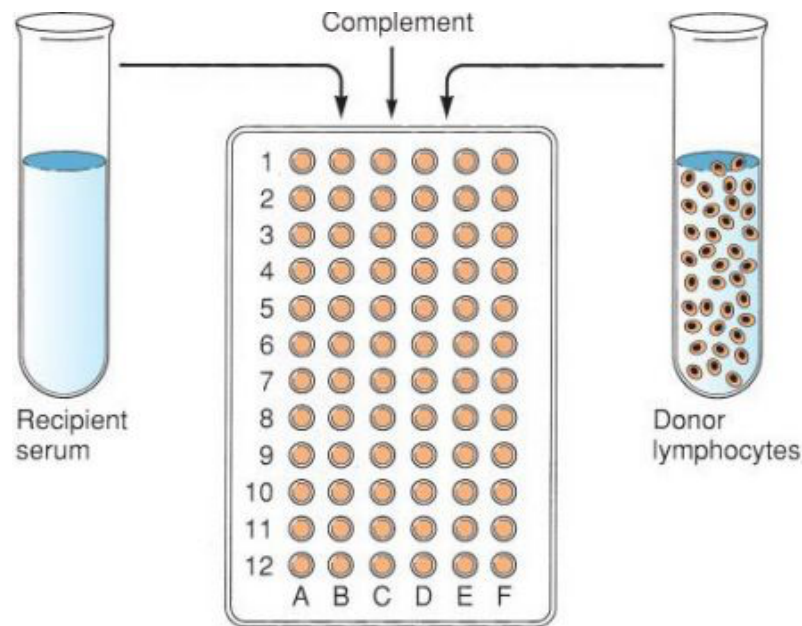
# Complement dependent cytotoxicity test (cross-match)

---

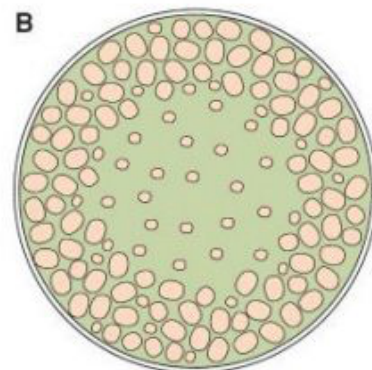
- Since 1970s prospective lymphocyte cross-match is a routine in kidney transplantation for prevention of DSA damage



# Panel reactive antibodies (PRA)



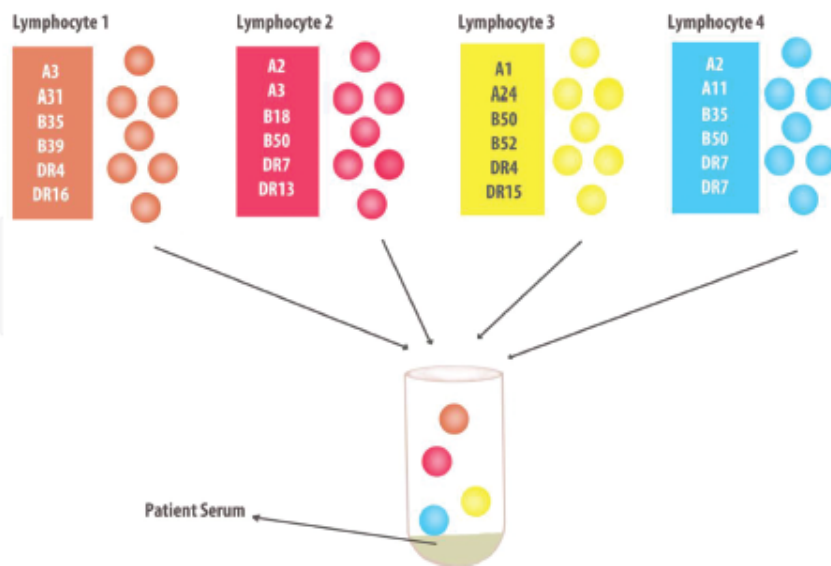
Positive crossmatch



Negative crossmatch

- Serum screening for preformed antibodies using a panel of typing cells
- Techniques identical to cross match
- Expressed as a percentage of cells with which the patient's serum reacts (0% to 100%)
- Sensitization (or increased panel reactive antibody) occurs as a consequence of prior pregnancy, blood transfusion, or transplant

# Konstrukce panelu HLA antigenů



Specificity	Number in panel	Specificity	Number in panel	Specificity	Number in panel	Specificity	Number in panel
A1	4	A80	2	B51	2	B75	1
A2	5	B7	2	B52	1	B78	1
A3	5	B8	2	B53	1	B81	2
A11	5	B13	2	B54	1	Bw4	18
A23	3	B18	2	B55	2	Bw6	26
A24	4	B27	2	B56	1	Cw1	6
A25	2	B35	2	B57	2	Cw2	6
A26	2	B37	2	B58	2	Cw4	4
A29	2	B38	1	B59	1	Cw5	1
A30	3	B39	1	B60	2	Cw6	7
A31	2	B41	2	B61	2	Cw7	9
A32	3	B42	1	B62	2	Cw8	5
A33	3	B44	2	B63	1	Cw9	2
A34	2	B45	1	B64	1	Cw10	4
A36	2	B46	1	B65	1	Cw12	1
A66	3	B47	2	B67	1	Cw14	2
A68	2	B48	2	B71	1	Cw15	3
A69	1	B49	1	B72	2	Cw16	2
A74	2	B50	1	B73	1	Cw17	3
						Cw18	3

**PRA Screening Lot 017-One Lambda**

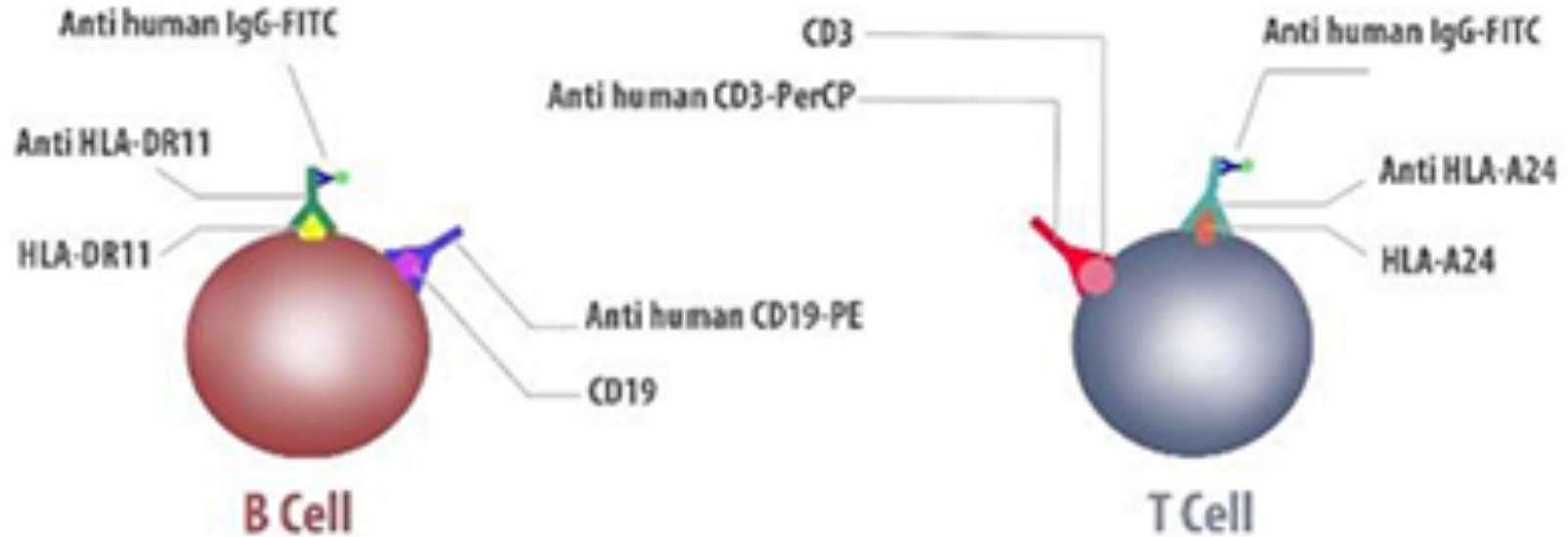
# Techniques for detection of HLA antibodies

Technology	Testing	Advantages	Limitations	Sensitivity	Specificity
Complement-dependent cytotoxic crossmatch	Donor lymphocytes are incubated with candidate serum and complement added	Highly predictive for hyperacute rejection	Subjective Cannot detect noncomplement binding or low-level antibodies	+	+++
Flow cytometric crossmatch	Donor T and B lymphocytes are exposed to candidate serum	Semiquantitative. Increased sensitivity to low-level antibody	Higher sensitivity may lead to false-positive results	++	++
Solid-phase assays: bead based	Purified HLA molecules immobilized onto solid surface	Can detect specific antigens a candidate has antibodies against	Lack of standardization Significant interlaboratory variability. Unclear significant result	++++	+

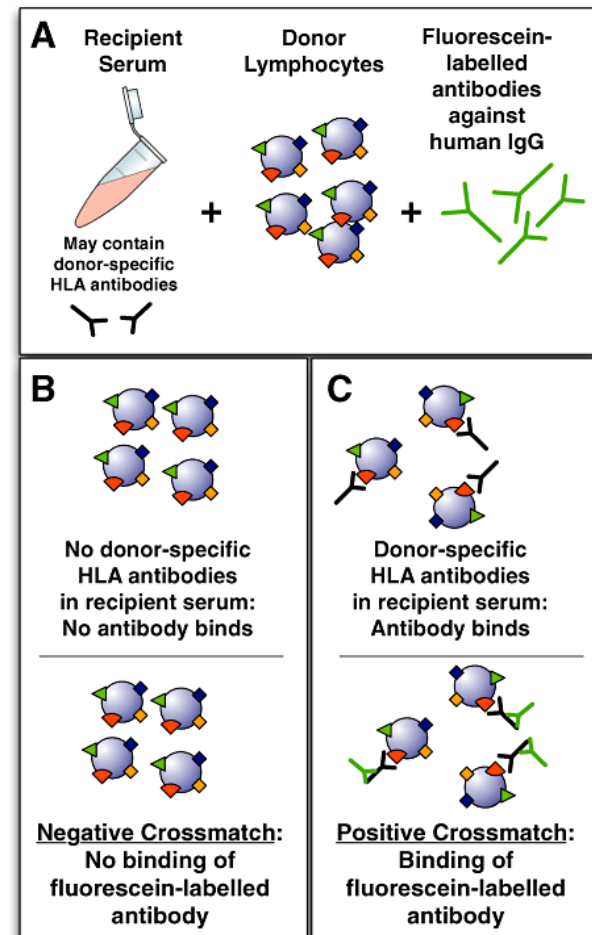
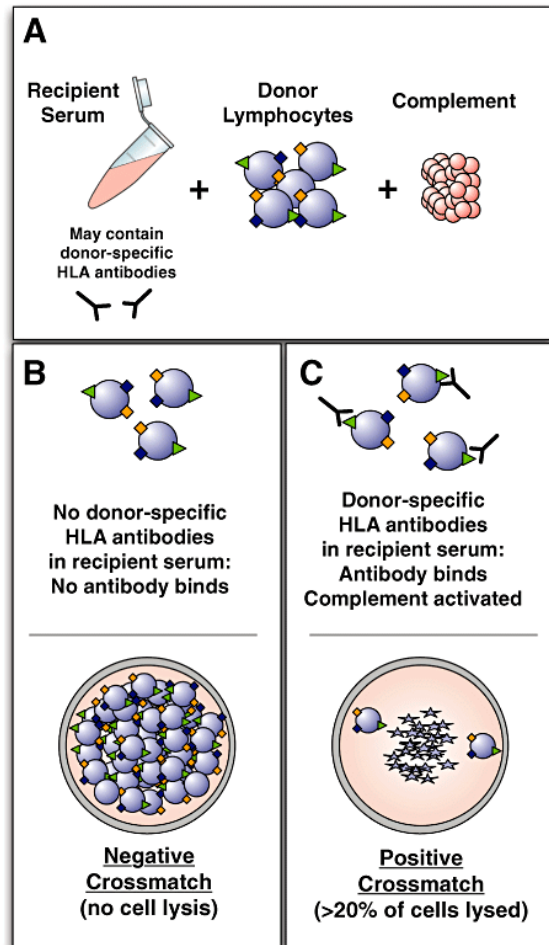


# Princip metody FCXM (flow cytometric crossmatch)

---

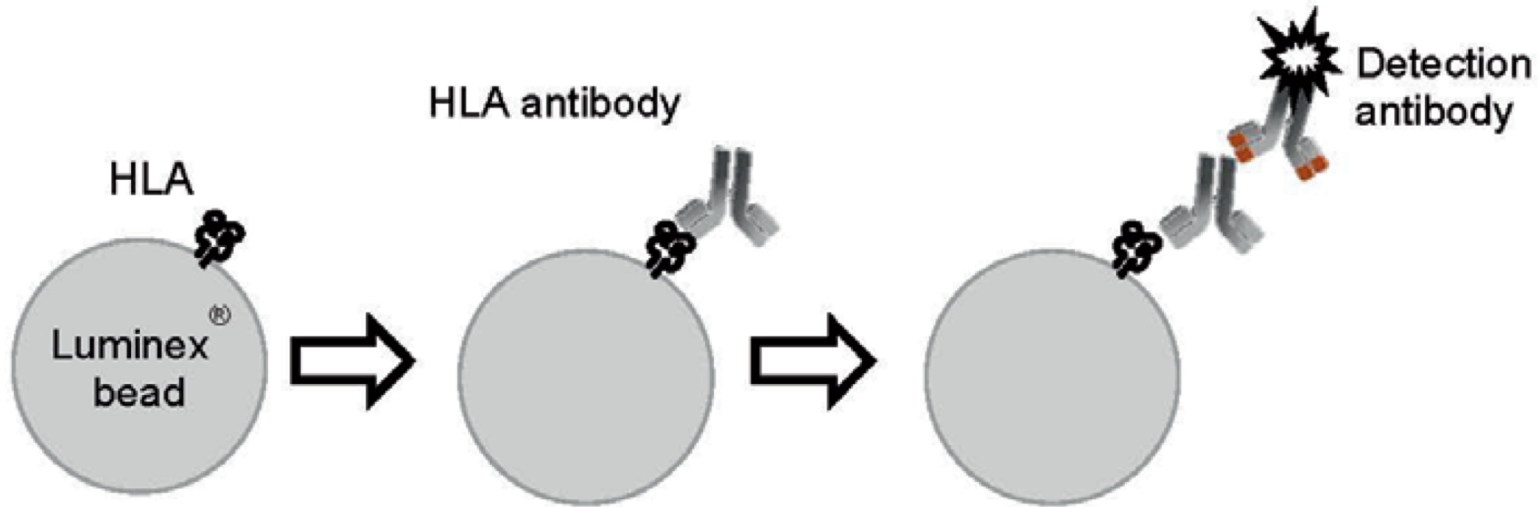


# Crossmatch testing in organ transplantation: CDC crossmatch and flow-cytometry crossmatch

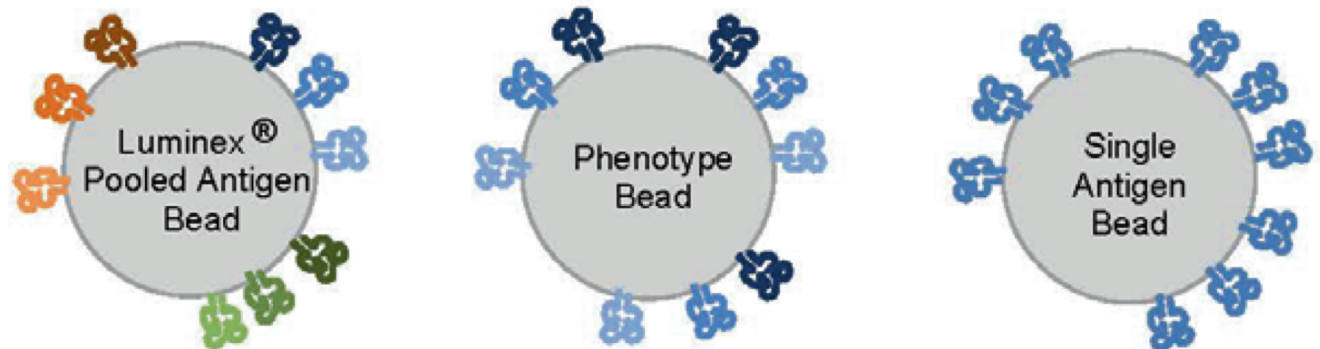


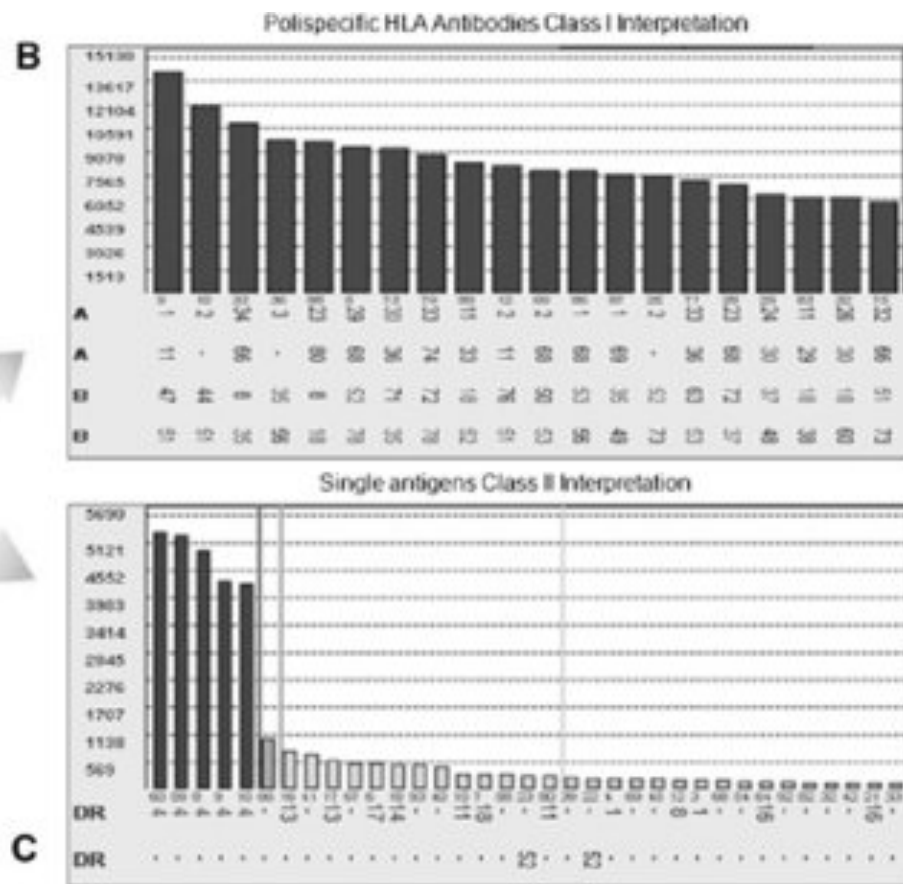
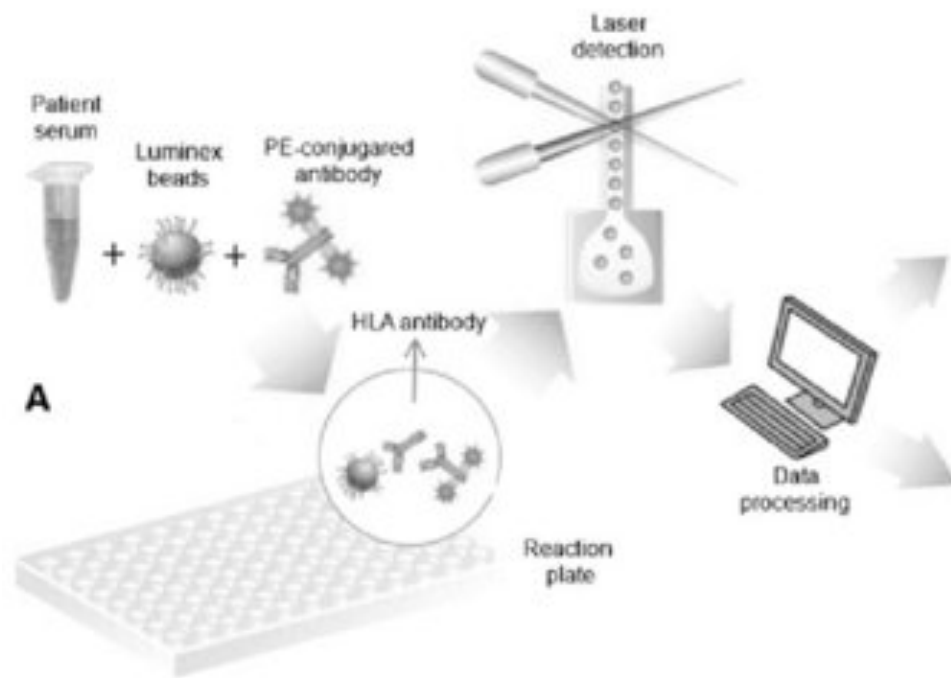
# Princip metody Luminex

A



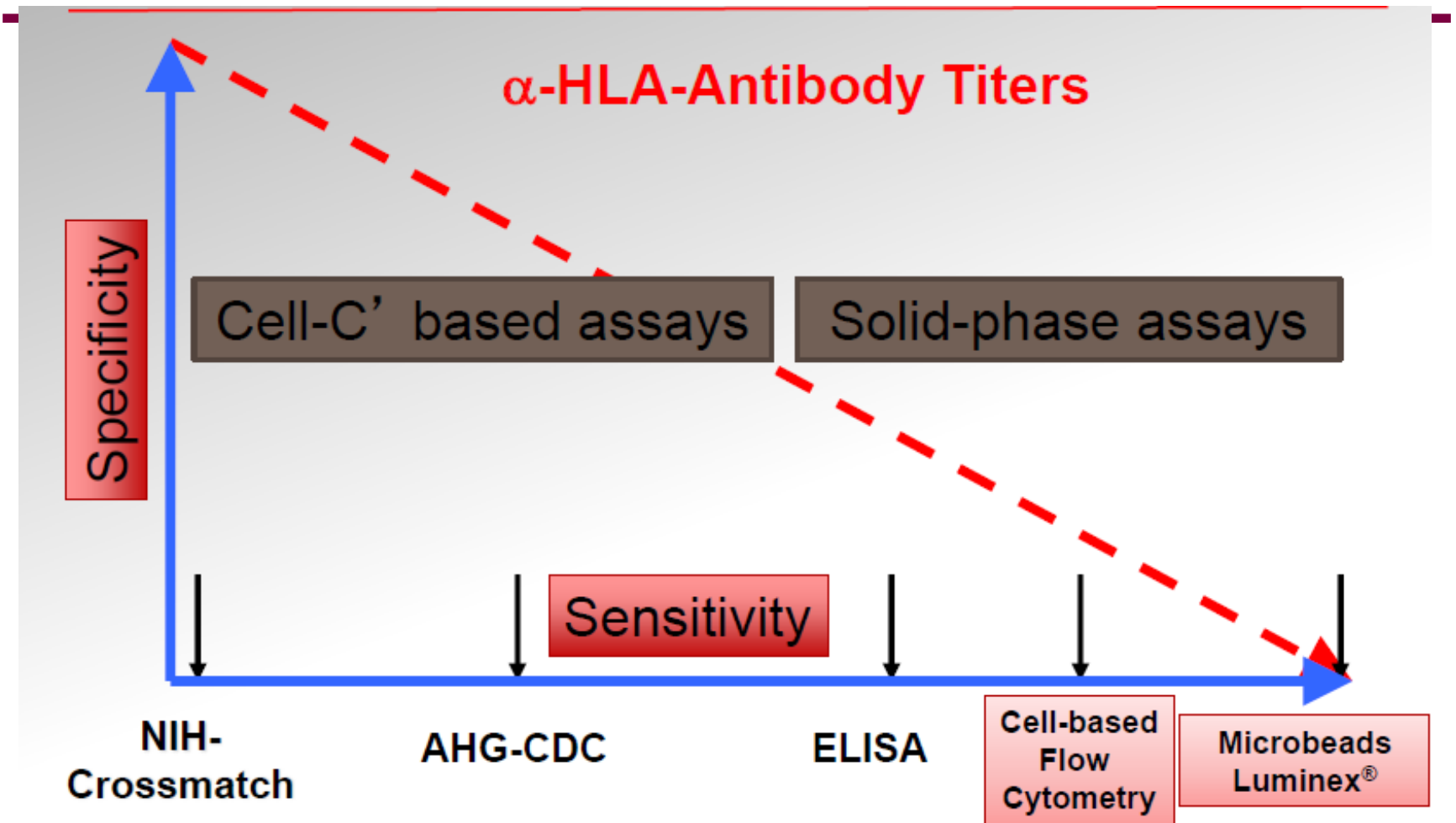
B



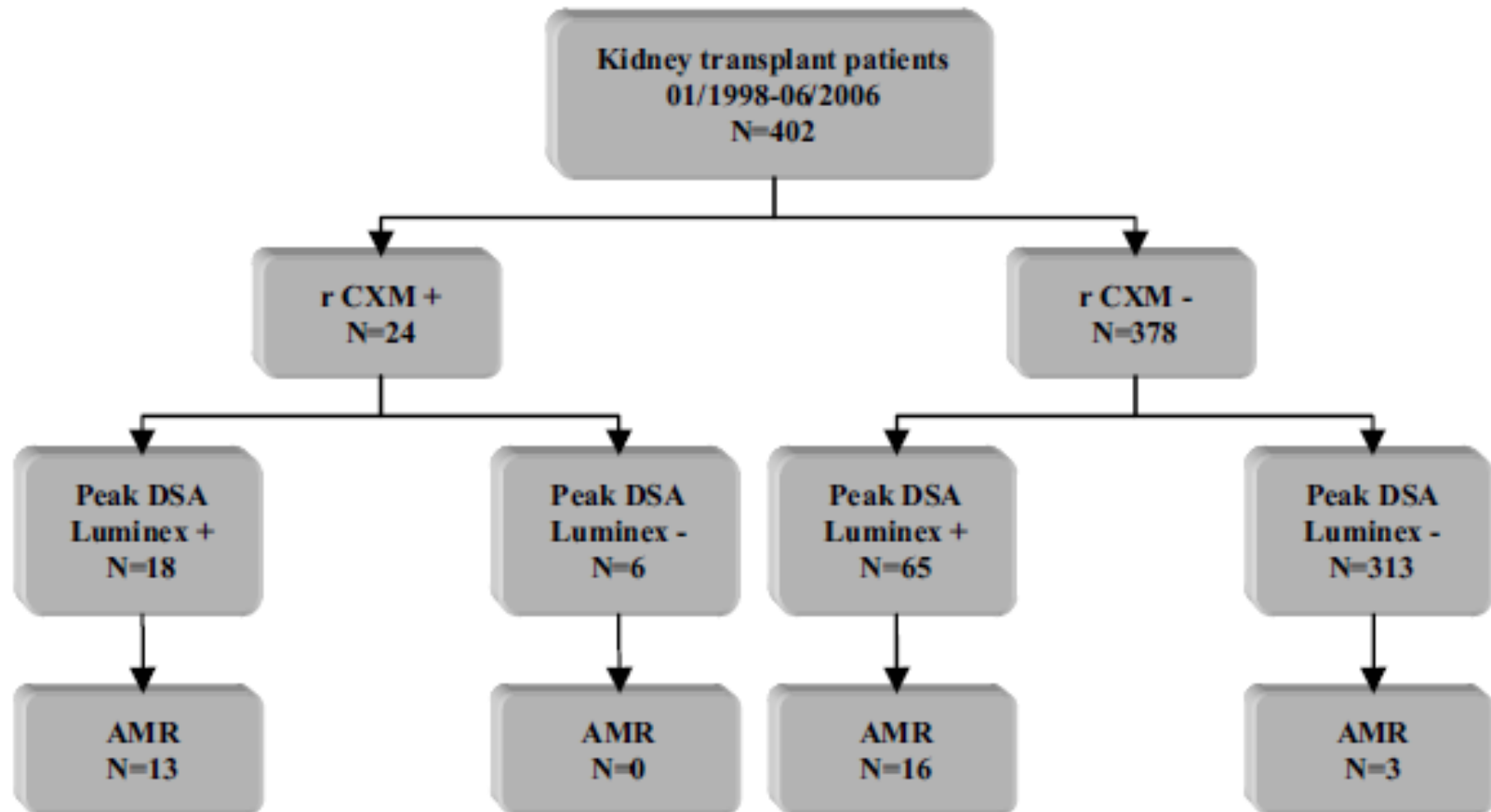




# Sensitivita testů k detekci HLA protilátek

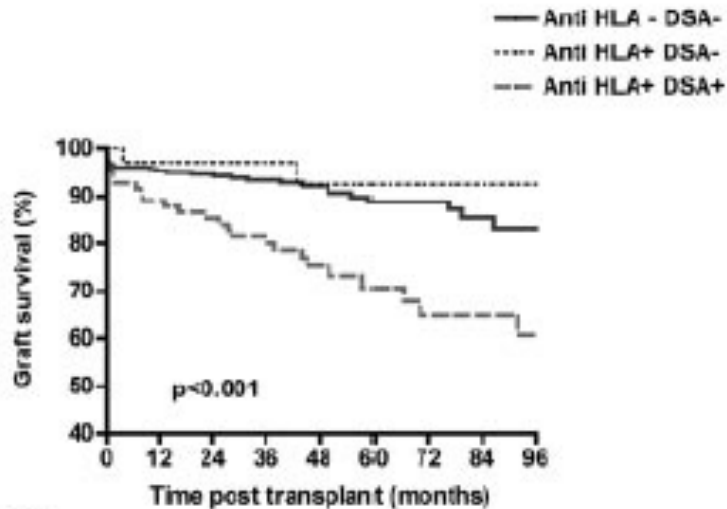


# Mnoho CDC negativních nemocných má DSA před transplantací



# Přítomnost DSA před transplantací koreluje s výsledky

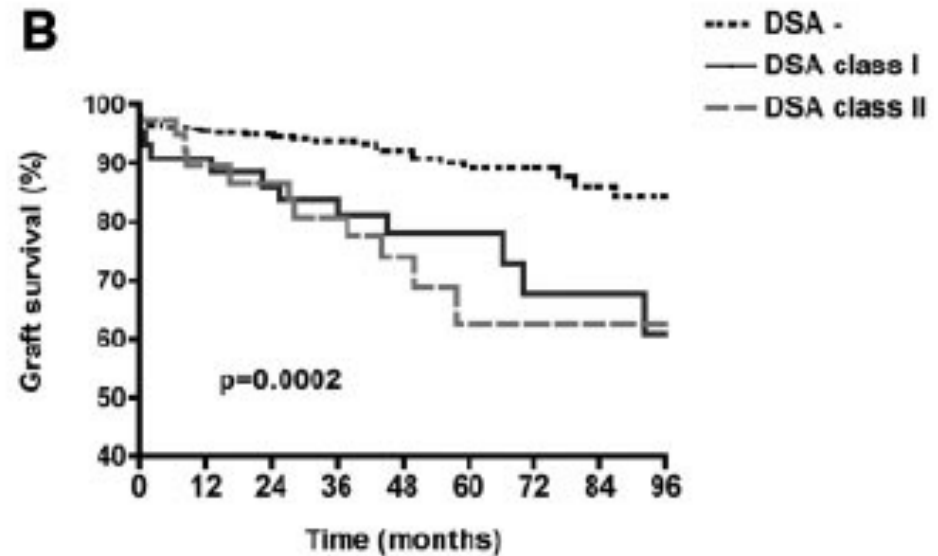
**A**



Number at risk

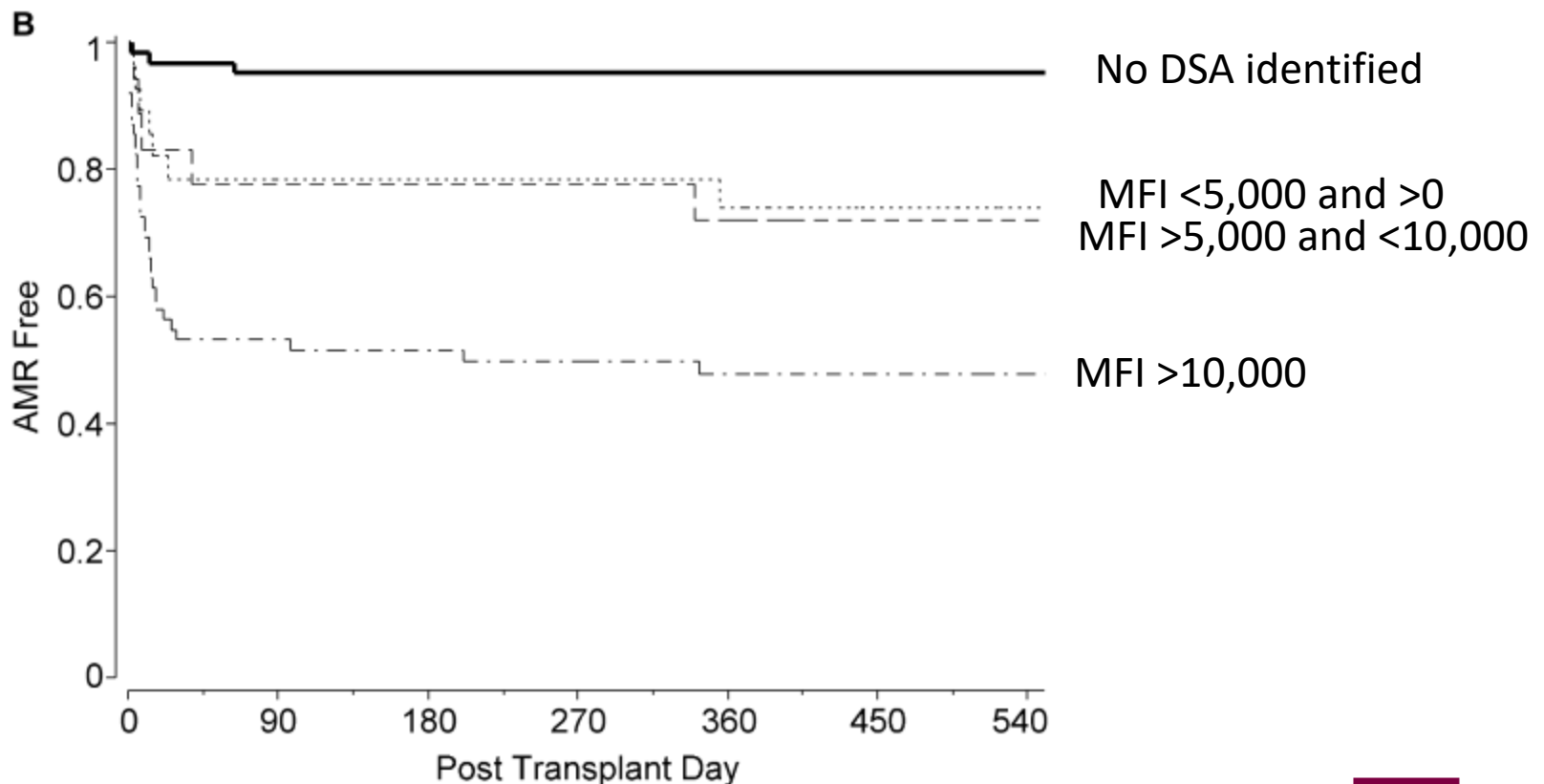
Anti HLA- DSA-	284	259	246	196	133	91	64	43	29
Anti HLA+ DSA-	35	34	33	25	16	10	10	7	6
Anti HLA+ DSA+	83	73	67	60	37	30	23	18	13

**B**



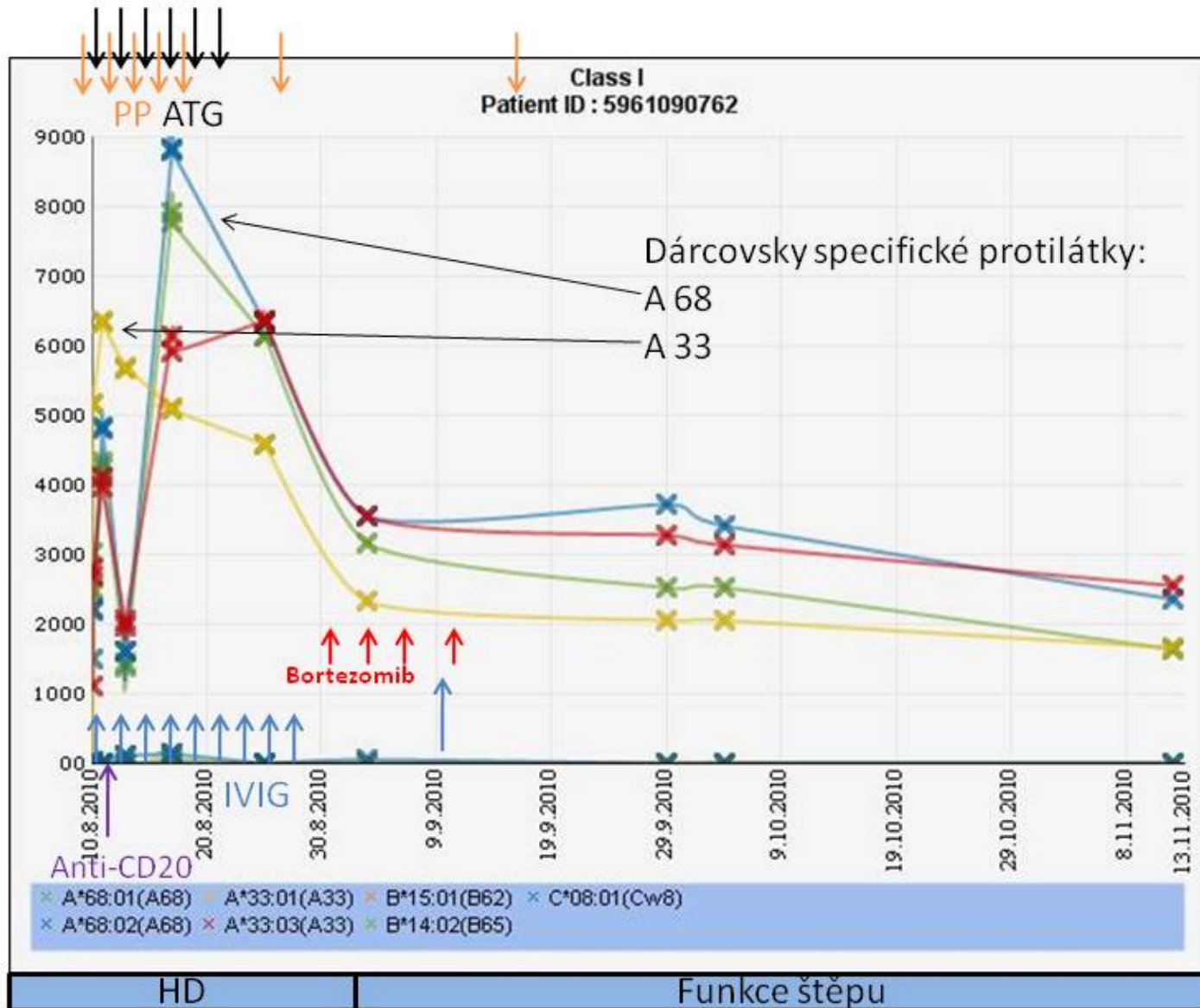
# Baseline Donor-Specific Antibody Levels and Outcomes in Positive Crossmatch Kidney Transplantation

J. M. Gloor<sup>a,\*</sup>, J. L. Winters<sup>b</sup>, L. D. Cornell<sup>b</sup>,  
L. A. Fix<sup>c</sup>, S. R. DeGoey<sup>b</sup>, R. M. Knauer<sup>b</sup>,  
F. G. Cosio<sup>a</sup>, M. J. Gandhi<sup>b</sup>, W. Kremers<sup>d</sup>  
and M. D. Stegall<sup>c</sup>

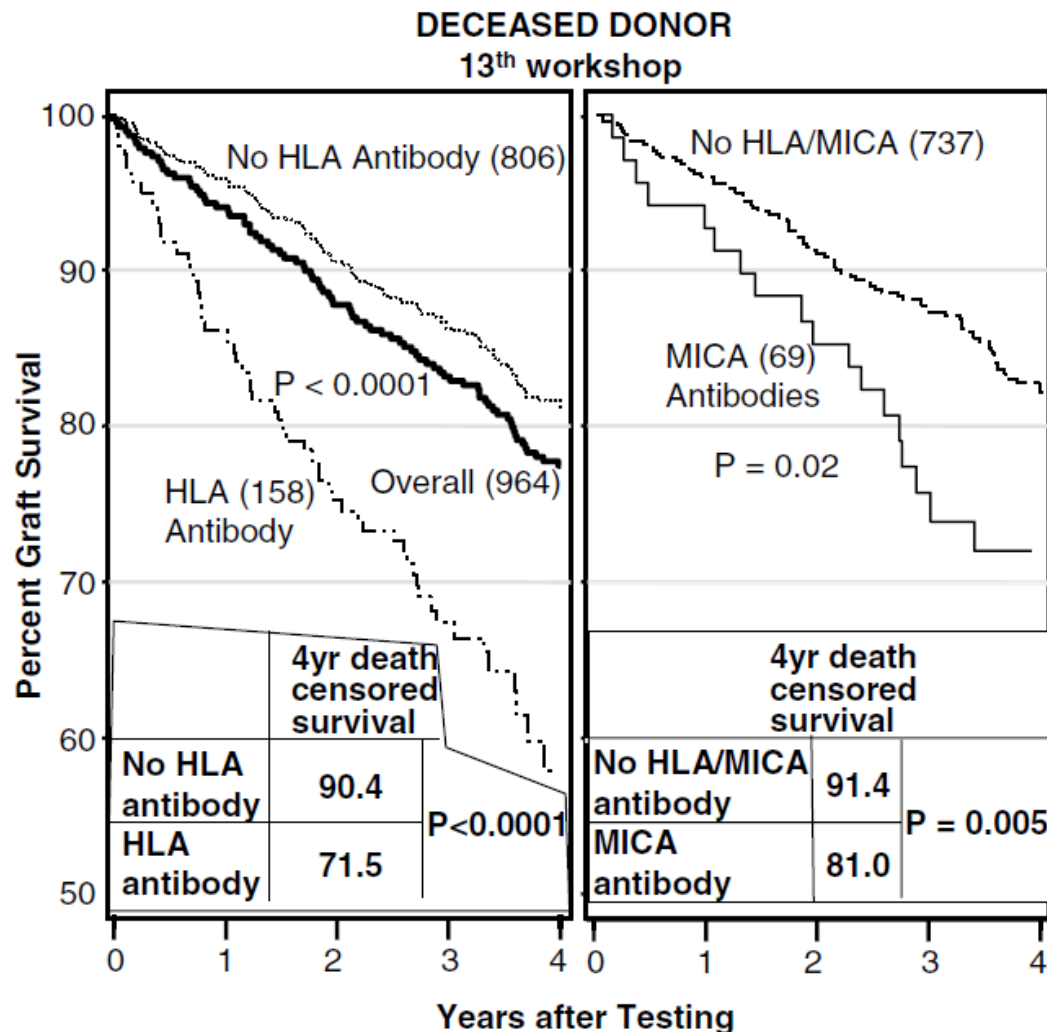


# Kidney transplantation in the presence of donor specific antibody (DSA)

IKEM 2010



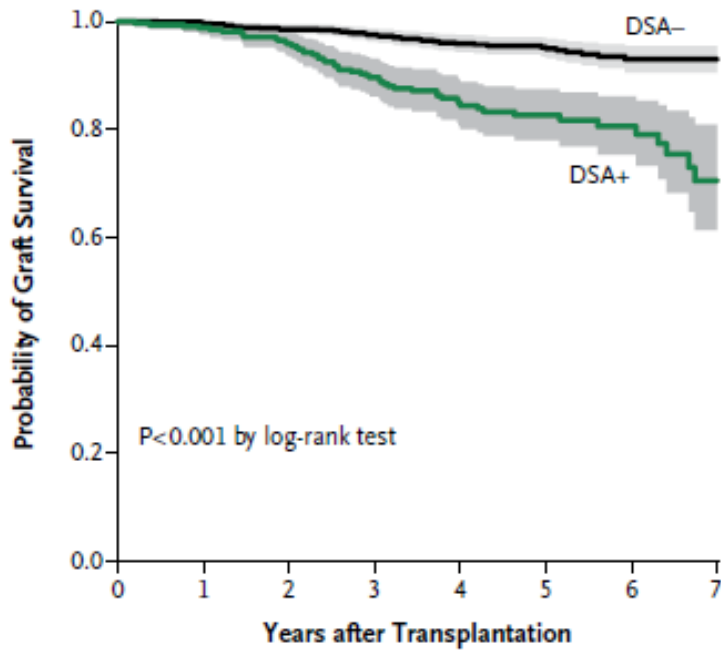
# HLA and MICA antibodies and kidney graft survival



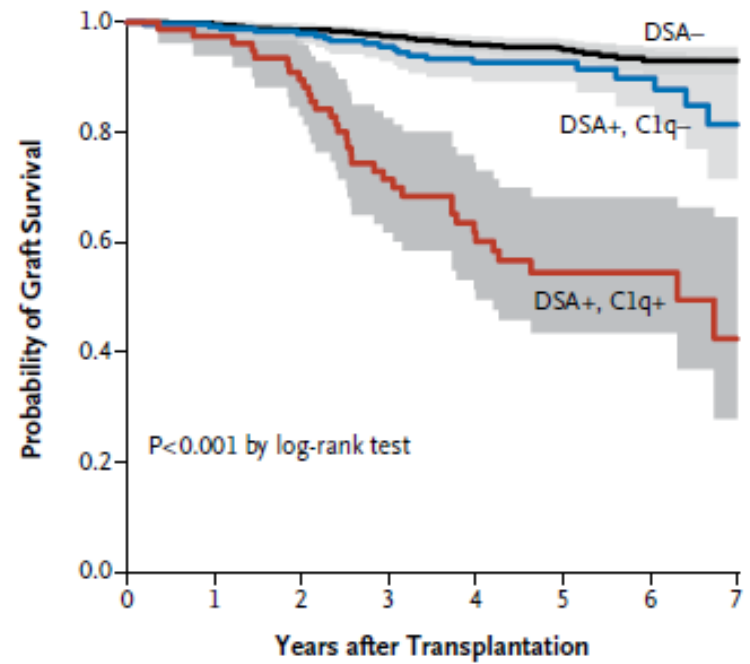
# Kidney allograft survival according to DSA and C1q status

47

**A Kidney-Allograft Survival According to DSA Status**



**B Kidney-Allograft Survival According to DSA and C1q Status**



**No. at Risk**

DSA-	700	698	667	612	504	338	164	38
DSA+	316	312	295	229	176	100	56	19

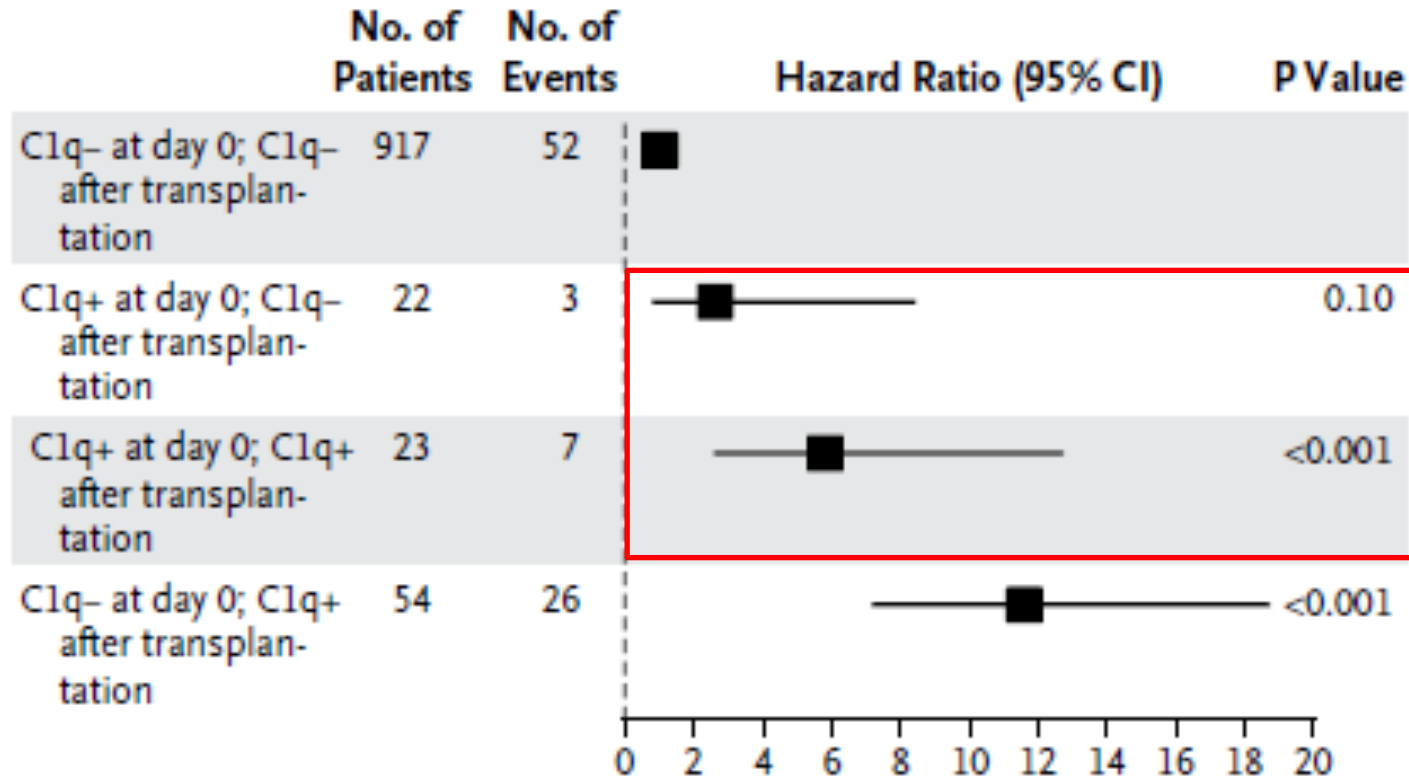
**No. at Risk**

DSA-	700	698	667	612	504	338	164	38
DSA+, C1q-	239	237	227	181	139	80	44	14
DSA+, C1q+	77	75	68	48	37	20	12	5

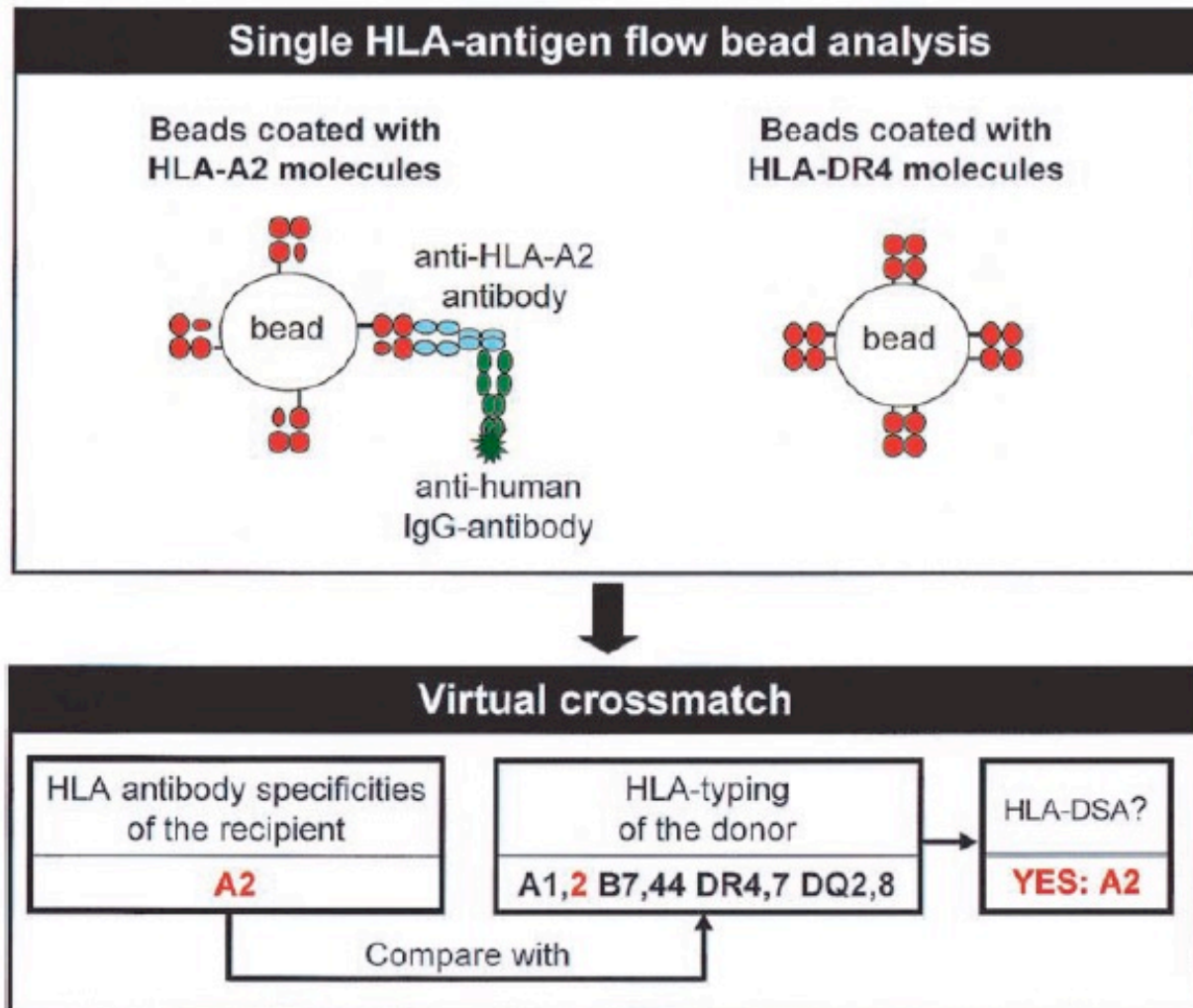
# C1q+ pretransplant does not mean risk if posttransplant status is negative

48

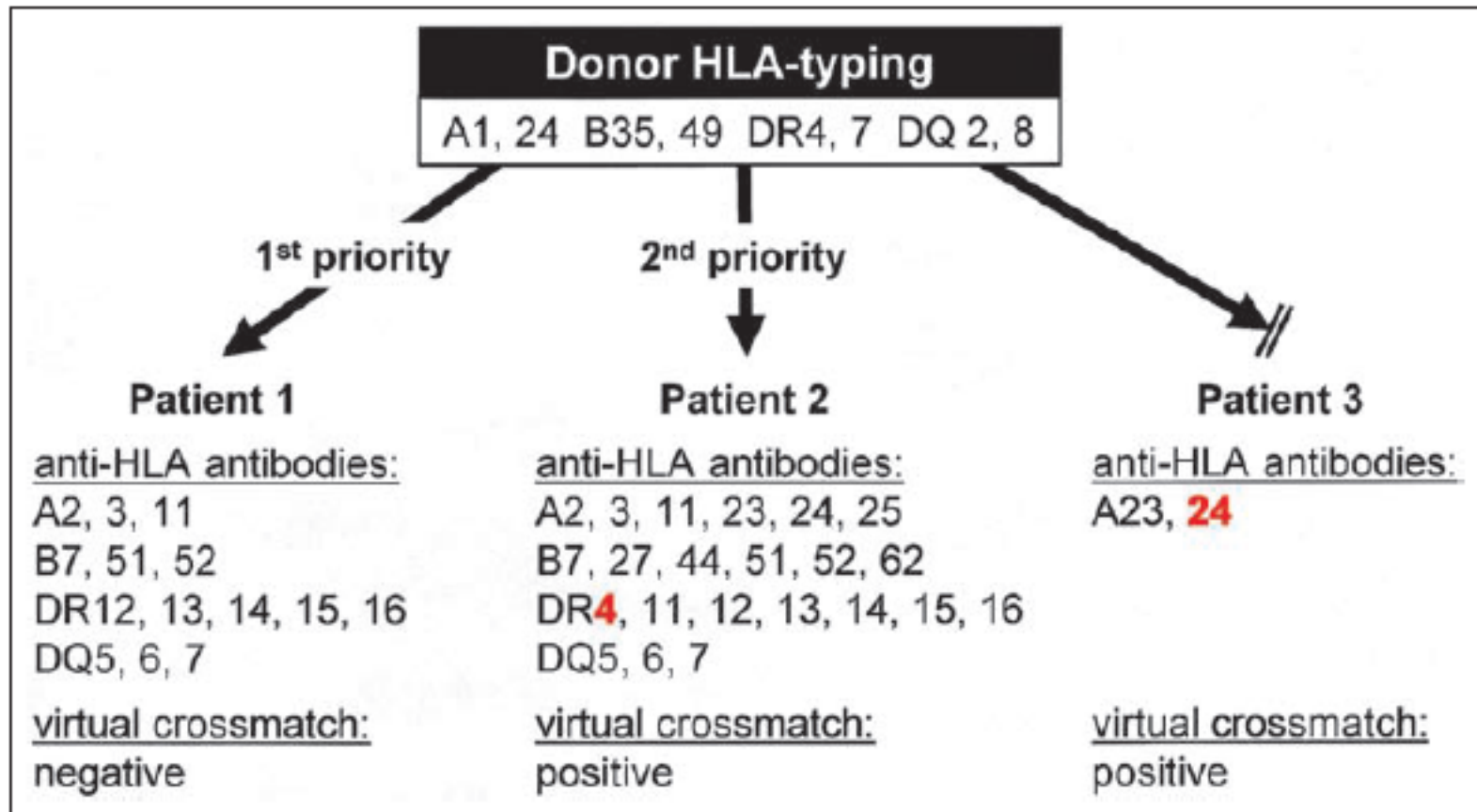
## C Risk of Kidney-Allograft Loss According to C1q Status



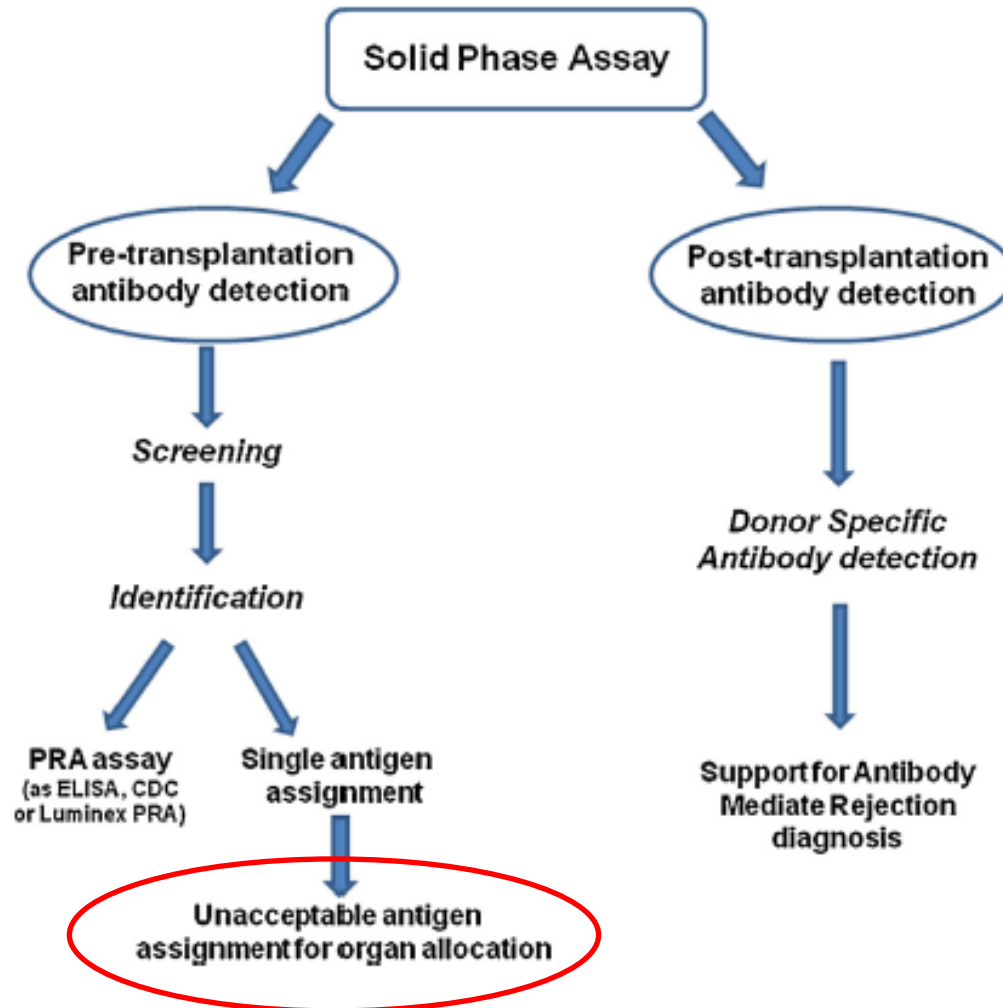
# Koncept virtuálního crossmatch



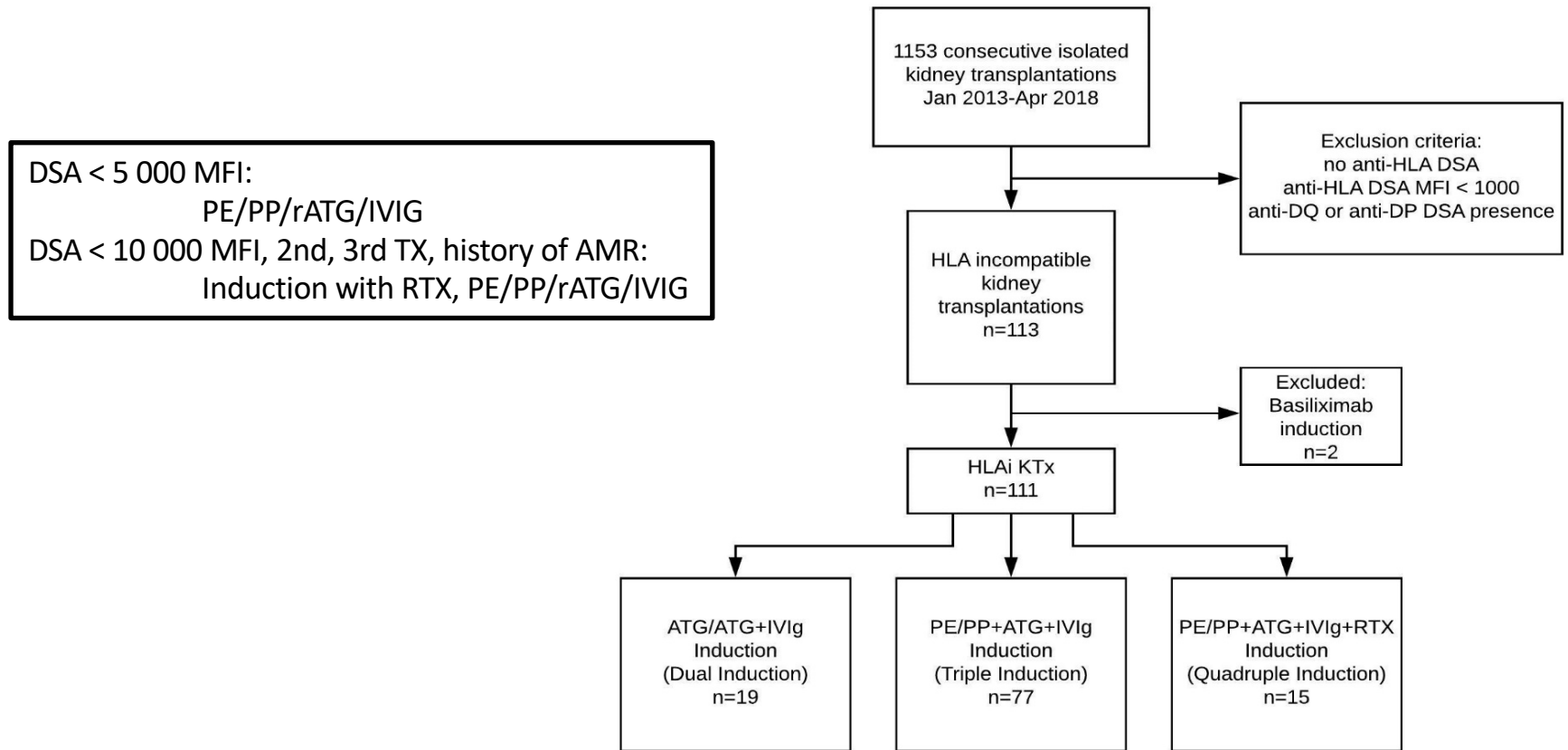
# Virtuální crossmatch a alokace ledviny



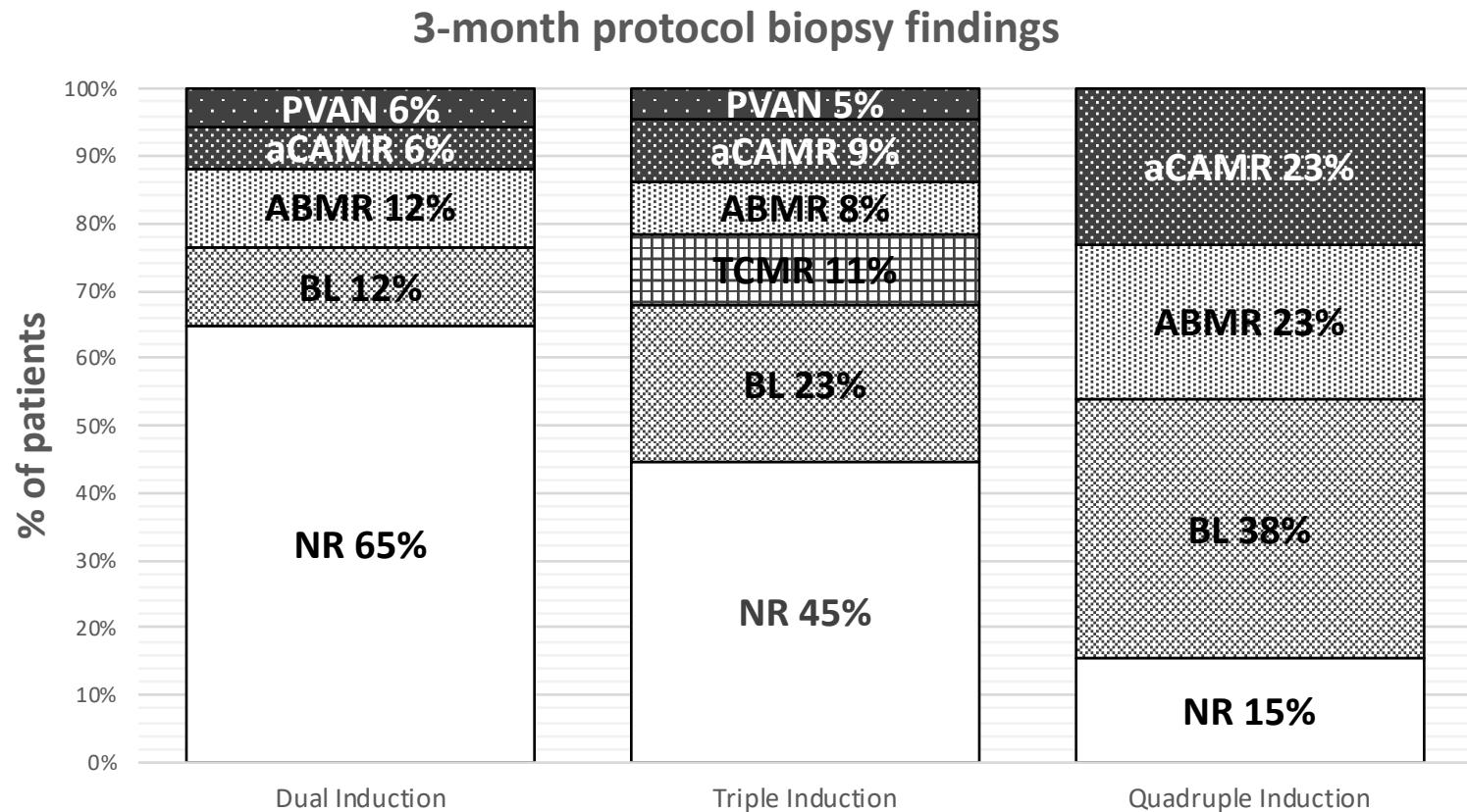
# Využití detekce anti HLA protilátek před a po transplantaci ledviny



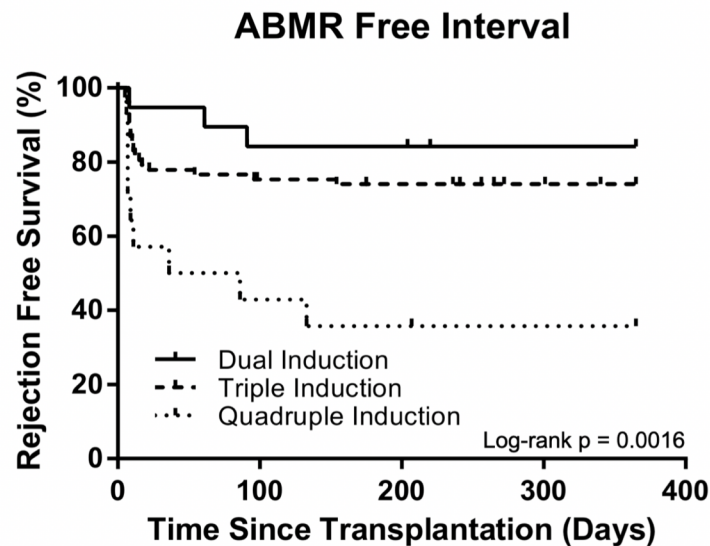
# HLAi transplantation at IKEM: induction strategy according to DSA levels



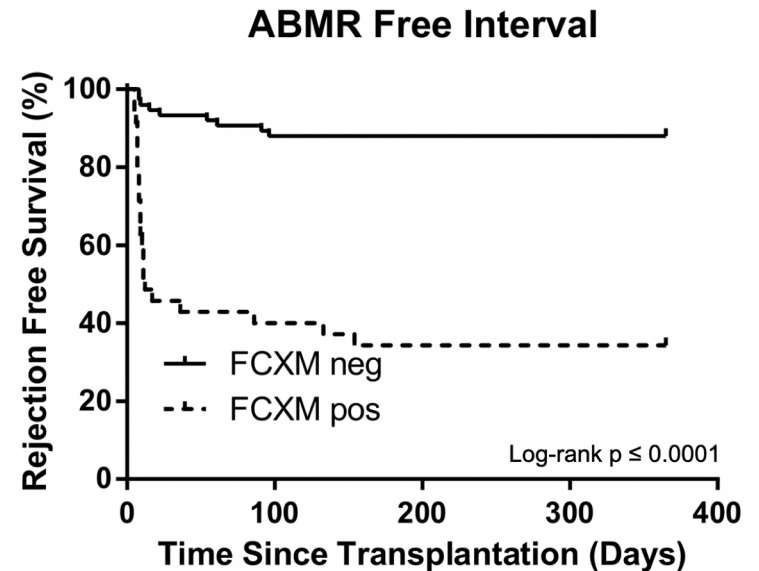
# Patients at increased immunological risk experienced more ABMR at 3M biopsies



# Pretransplant FCXM better than DSA in rejection prediction

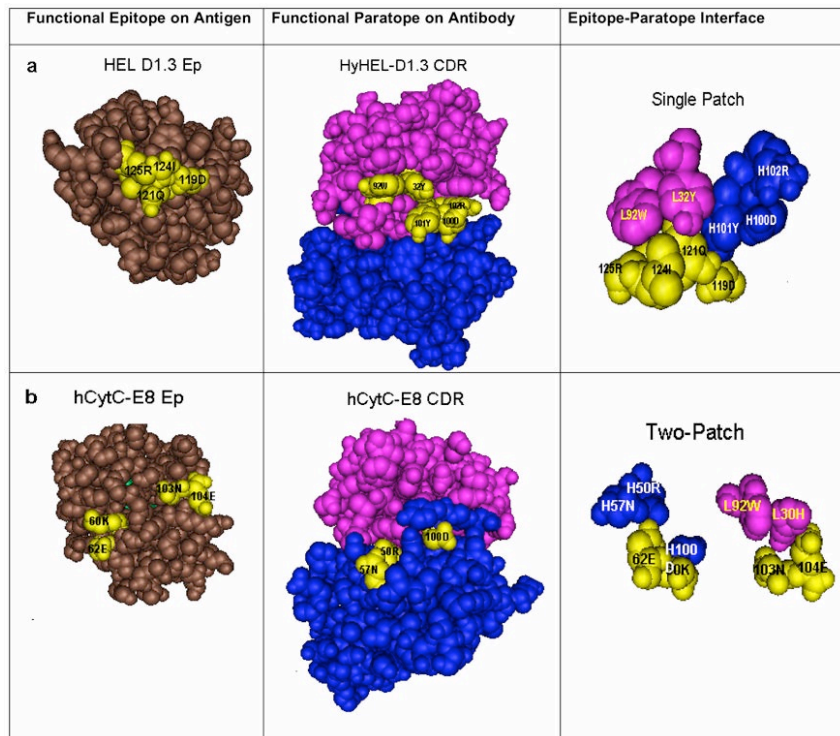


No. at Risk	0	100	200	300	400
Dual Induction	19	17	17	15	14
Triple Induction	77	58	56	51	48
Quadruple Induction	15	7	6	5	4



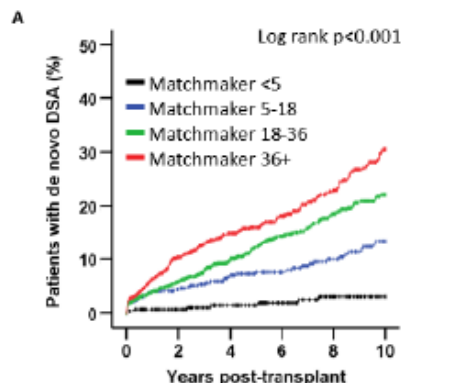
No. at Risk	0	100	200	300	400
FCXM neg	75	67	67	67	66
FCXM pos	36	15	13	13	12

# Mismatched HLA antigens without foreign epitopes for recipient do not induce antibodies similarly to fully matched grafts



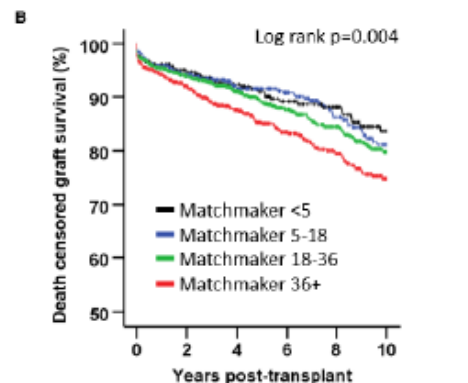
- HLAMatchmaker is a useful tool in the analysis of serum antibody reactivity of sensitized patients and the identification of potential donors with acceptable epitope mismatches
- Mismatch program to identify donors for highly sensitized patients in Eurotransplant and a similar program (Eurostam) is now being investigated for implementation in the European Union

# Donor-Recipient Matching Based on Predicted Indirectly Recognizable HLA Epitopes Independently Predicts the Incidence of *De Novo* Donor-Specific HLA Antibodies Following Renal Transplantation



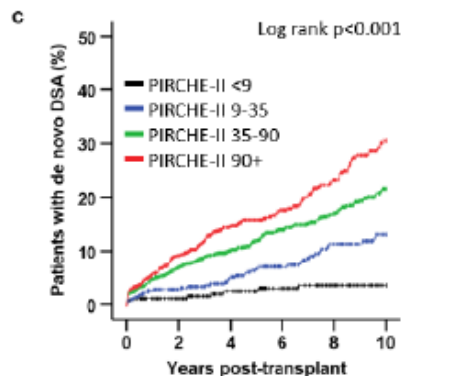
Number at risk

Years post-transplant	0	2	4	6	8	10
Matchmaker <5	323	280	236	193	145	101
Matchmaker 5-18	476	397	326	263	201	150
Matchmaker 18-36	1136	917	683	510	388	249
Matchmaker 36+	852	644	495	371	265	162



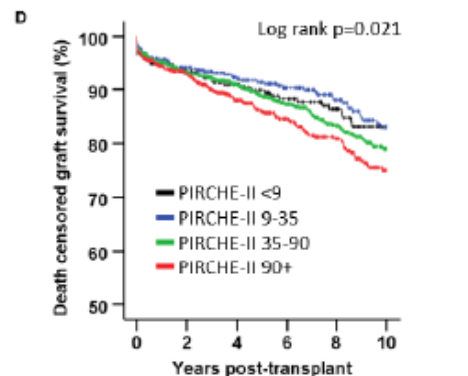
Number at risk

Years post-transplant	0	2	4	6	8	10
Matchmaker <5	323	268	222	179	136	90
Matchmaker 5-18	476	391	320	255	190	138
Matchmaker 18-36	1136	926	703	524	399	250
Matchmaker 36+	852	672	523	388	278	168



Number at risk

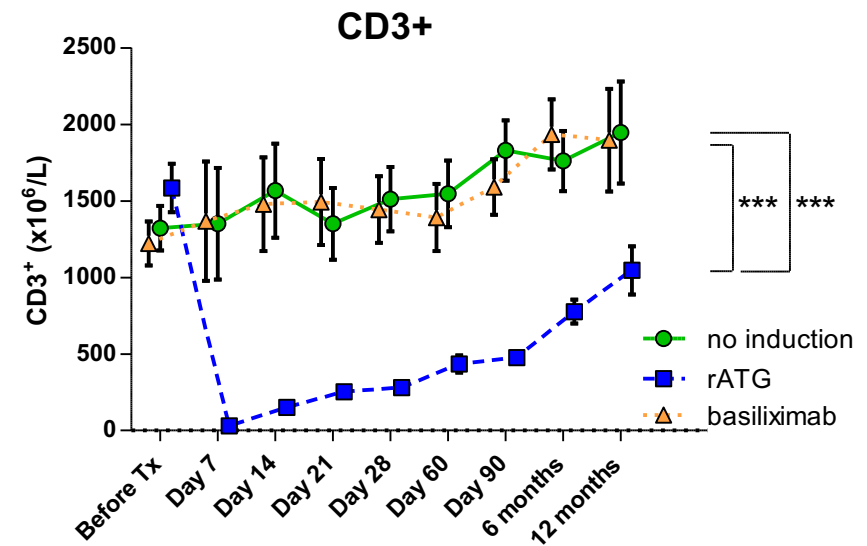
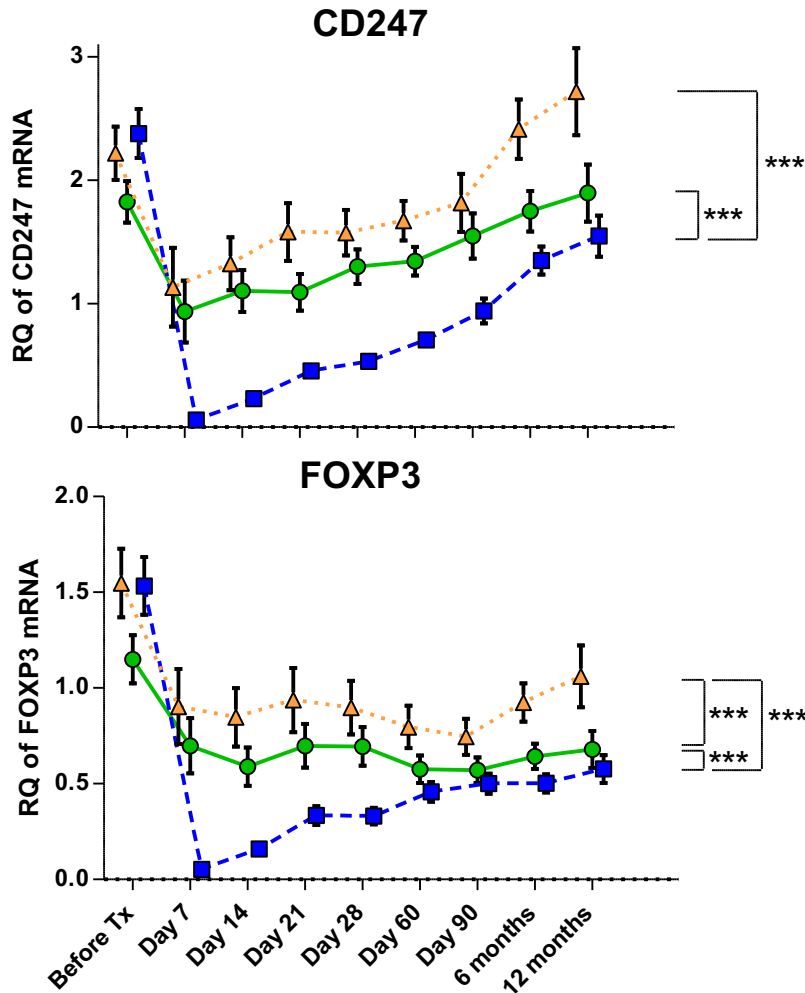
Years post-transplant	0	2	4	6	8	10
PIRCHE <9	285	246	203	168	126	87
PIRCHE 9-35	446	378	311	245	182	131
PIRCHE 35-90	1222	984	763	578	451	301
PIRCHE 90+	834	630	463	346	240	143



Number at risk

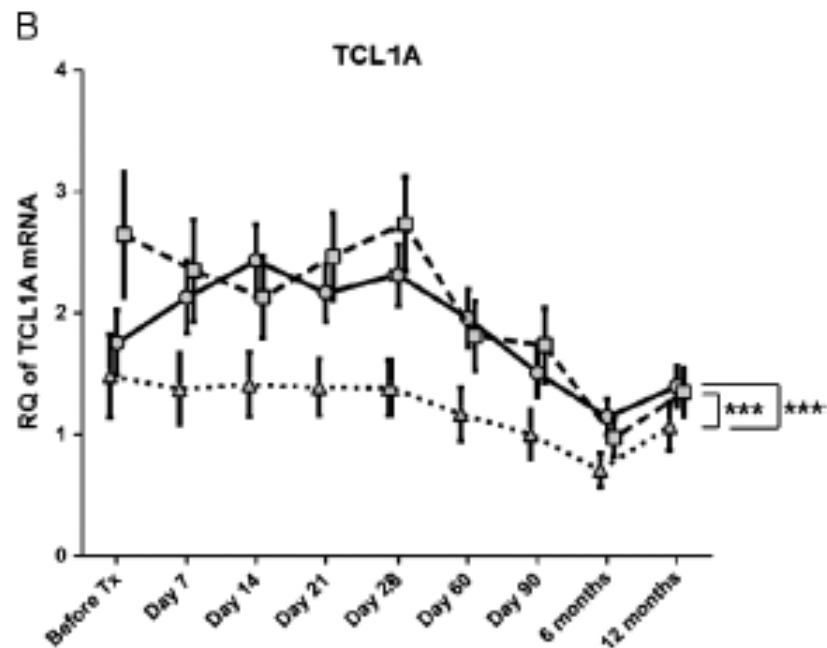
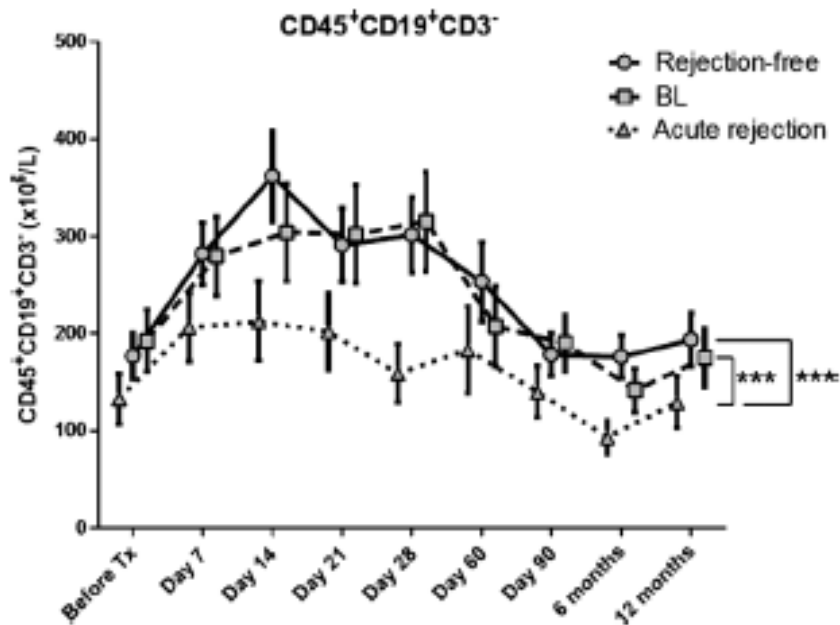
Years post-transplant	0	2	4	6	8	10
PIRCHE <9	285	234	190	155	116	77
PIRCHE 9-35	446	370	301	238	179	123
PIRCHE 35-90	1222	1001	787	590	450	298
PIRCHE 90+	834	652	490	363	258	147

# Průtoková cytometrie a periferní transkripty k monitoraci po transplantaci



# B-Cell-Related Biomarkers of Tolerance are Up-Regulated in Rejection-Free Kidney Transplant Recipients

N=69, TAC/MMF/St ± Induction



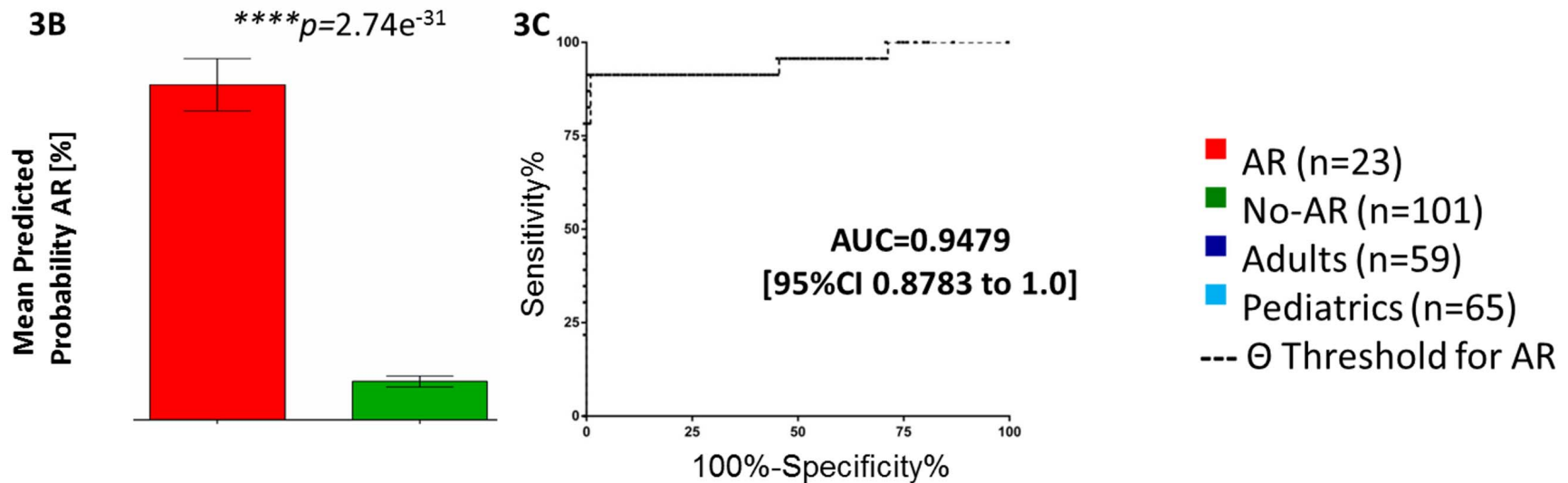
# 17-gene set—the Kidney Solid Organ Response Test (kSORT)

OPEN ACCESS Freely available online

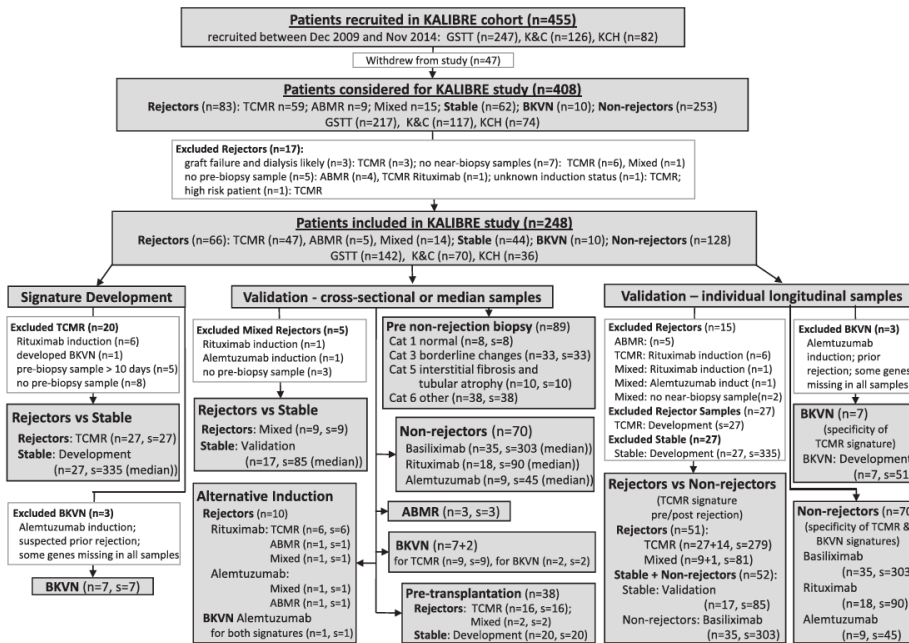
PLOS MEDICINE

## The kSORT Assay to Detect Renal Transplant Patients at High Risk for Acute Rejection: Results of the Multicenter AART Study

Silke Roedder<sup>1\*</sup>, Tara Sigdel<sup>1\*</sup>, Nathan Salomonis<sup>2\*</sup>, Sue Hsieh<sup>1</sup>, Hong Dai<sup>3,4\*</sup>, Oriol Bestard<sup>5</sup>, Diana Metes<sup>5</sup>, Andrea Zeevi<sup>5</sup>, Albin Gritsch<sup>5</sup>, Jennifer Cheeseman<sup>7</sup>, Camila Macedo<sup>5</sup>, Ram Peddy<sup>3</sup>, Mara Medeiros<sup>8</sup>, Flavio Vincenti<sup>1</sup>, Nancy Asher<sup>1</sup>, Oscar Salvatierra<sup>2</sup>, Ron Shapiro<sup>5</sup>, Allan Kirk<sup>7,8</sup>, Elaine Reed<sup>6</sup>, Minnie M. Sarwal<sup>1\*</sup>



# 7-gene peripheral signature predicts TCMR in kidney transplantation (KALIBRE study)



Contents lists available at ScienceDirect

EBioMedicine

journal homepage: [www.ebiomedicine.com](http://www.ebiomedicine.com)

**EBioMedicine**  
Published by THE LANCET

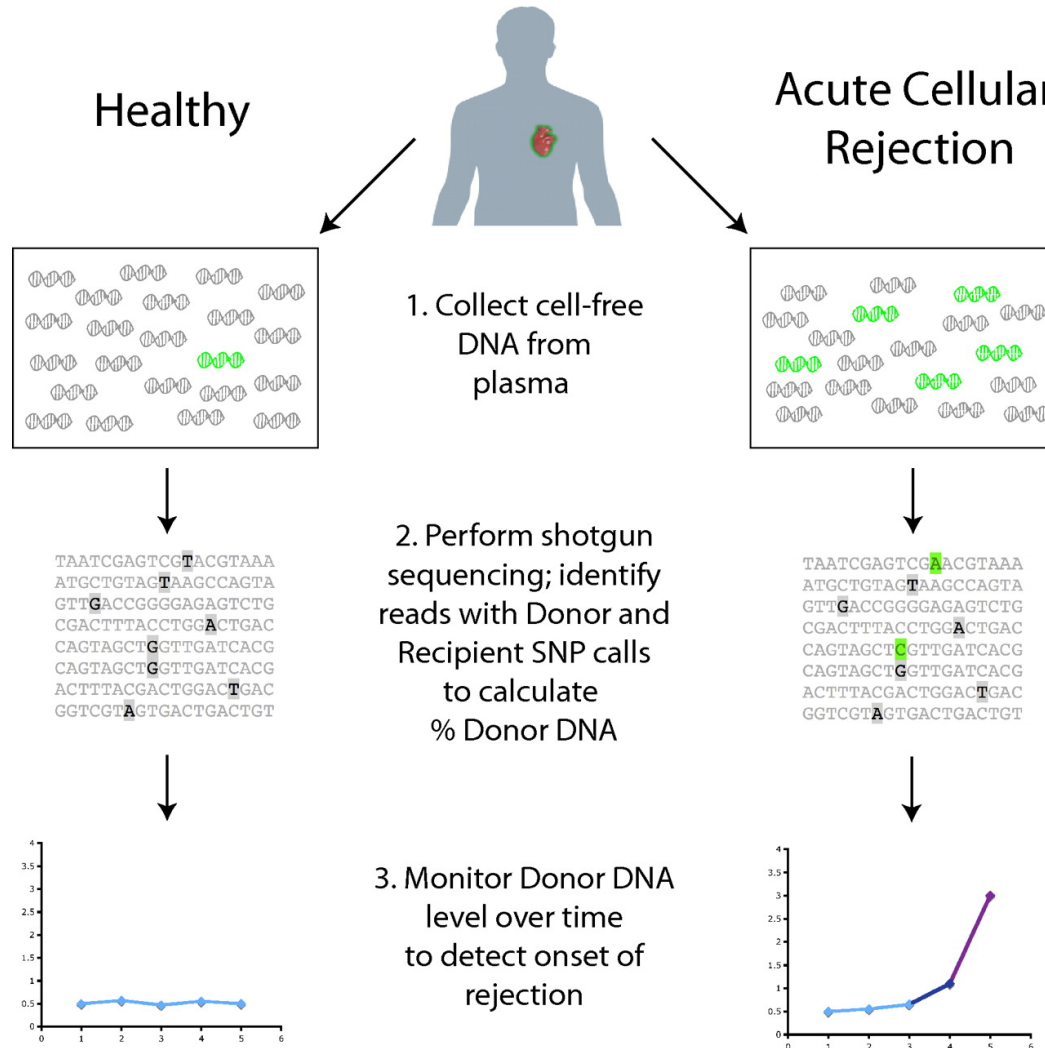
Development of a multivariable gene-expression signature targeting T-cell-mediated rejection in peripheral blood of kidney transplant recipients validated in cross-sectional and longitudinal samples

Sofia Christakoudi<sup>a,b,c,1</sup>, Manoharsingh Runglall<sup>d,1</sup>, Paula Mobillo<sup>a</sup>, Tjir-Li Tsui<sup>a,d,e</sup>, Claire Duff<sup>a,d</sup>, Clara Domingo-Vila<sup>a</sup>, Yogesh Kamra<sup>a,d</sup>, Florence Delaney<sup>d</sup>, Rosa Montero<sup>e,f</sup>, Anastasia Spiridou<sup>d,j</sup>, Theodoros Kassimatis<sup>a</sup>, Sui Phin-Kon<sup>g</sup>, Beatriz Tucker<sup>g</sup>, Christopher Farmer<sup>h</sup>, Terry B. Strom<sup>i,2</sup>, Graham M. Lord<sup>a,d,e</sup>, Irene Rebollo-Mesa<sup>a,b,k</sup>, Daniel Stahl<sup>b</sup>, Steven Sacks<sup>a</sup>, Maria P. Hernandez-Fuentes<sup>a,d,k,\*,3,4</sup>, Paramit Chowdhury<sup>e,\*,3,4</sup>

## Gene-expression signatures.

Event	Genes	Data	AUC	Sens	Spec
TCMR	All genes	Training data	0.96 (0.92–1.00)	0.85	0.89
		Cross-validation	0.80 (0.75–0.83)	0.67 (0.59–0.70)	0.81 (0.71–0.88)
TCMR	<i>IFNG, IP-10, ITGA4, MARCH8, RORc, SEMA7A, WDR40A</i>	Training data	0.95 (0.91–1.00)	0.85	0.93
		Cross-validation	0.84 (0.77–0.88)	0.67 (0.59–0.74)	0.85 (0.75–0.89)
BKVN	All genes	Training data	0.93 (0.85–1.00)	0.71	0.93
		Cross-validation	0.68 (0.64–0.73)	0 (0–0.25)	0.91 (0.85–0.94)
BKVN	<i>IL-15, IL1R2, MARCH8, PDCD1, TGFB, WDR40A</i>	Training data	0.95 (0.90–1.00)	0.71	0.91
		Cross-validation	0.73 (0.66–0.80)	0.43 (0.29–0.57)	0.89 (0.83–0.91)

# Donor derived cell-free DNA (dd-cf DNA)

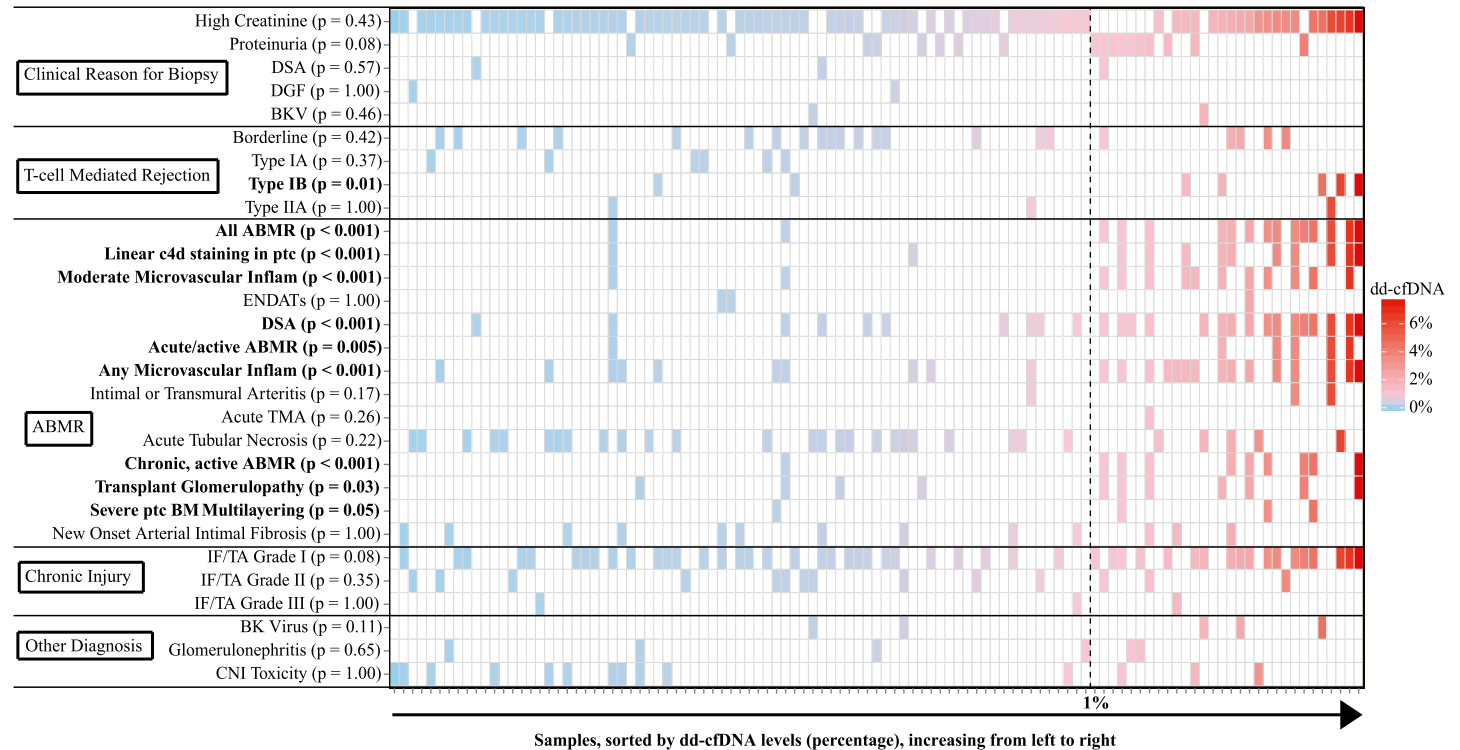


# Cell-Free DNA as a marker of active rejection?

## Cell-Free DNA and Active Rejection in Kidney Allografts

Roy D. Bloom,<sup>1</sup> Jonathan S. Bromberg,<sup>1</sup> Emilio D. Poggio,<sup>1</sup> Supharnai Bunnapradist,<sup>3</sup> Anthony J. Langone,<sup>3</sup> Puneet Sood,<sup>3</sup> Arthur J. Matas,<sup>4,5</sup> Shikha Mehta,<sup>1,7</sup> Roslyn B. Mannon,<sup>1,11</sup> Asif Sharfuddin,<sup>11</sup> Bernard Fischbach,<sup>11</sup> Mohanram Narayanan,<sup>11</sup> Stanley C. Jordan,<sup>4,11</sup> David Cohen,<sup>11</sup> Matthew R. Weir,<sup>11</sup> David Miller,<sup>11</sup> Preethi Prasad,<sup>11</sup> Robert N. Woodward,<sup>11</sup> Marica Grskovic,<sup>11</sup> John J. Srinisky,<sup>11</sup> James P. Yee,<sup>11</sup> and Daniel C. Brennan,<sup>1,11</sup> for the Circulating Donor-Derived Cell-Free DNA in Blood for Diagnosing Active Rejection in Kidney Transplant Recipients (DART) Study Investigators

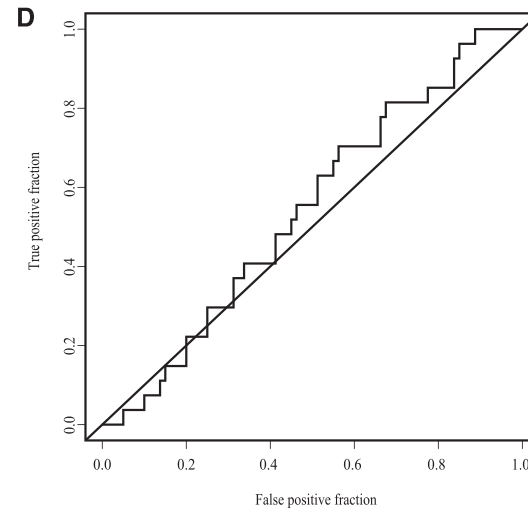
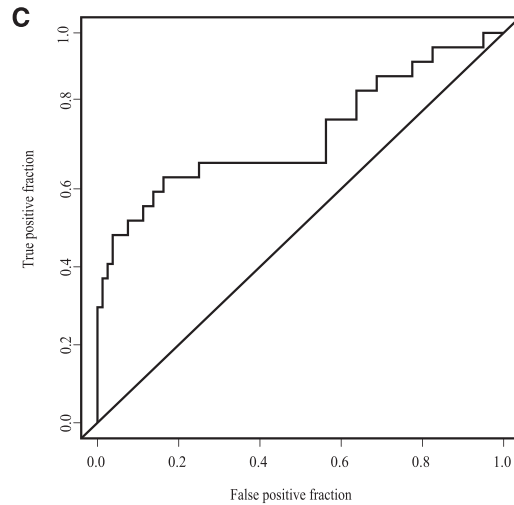
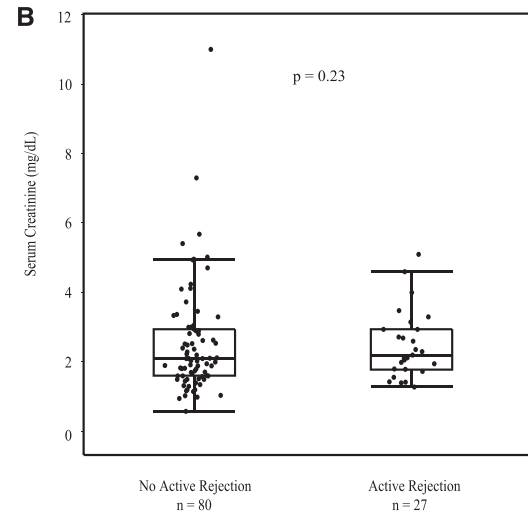
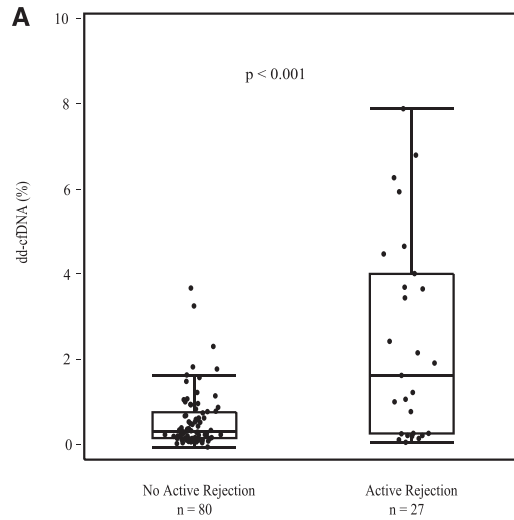
DART prospective study  
n=102  
107 biopsies



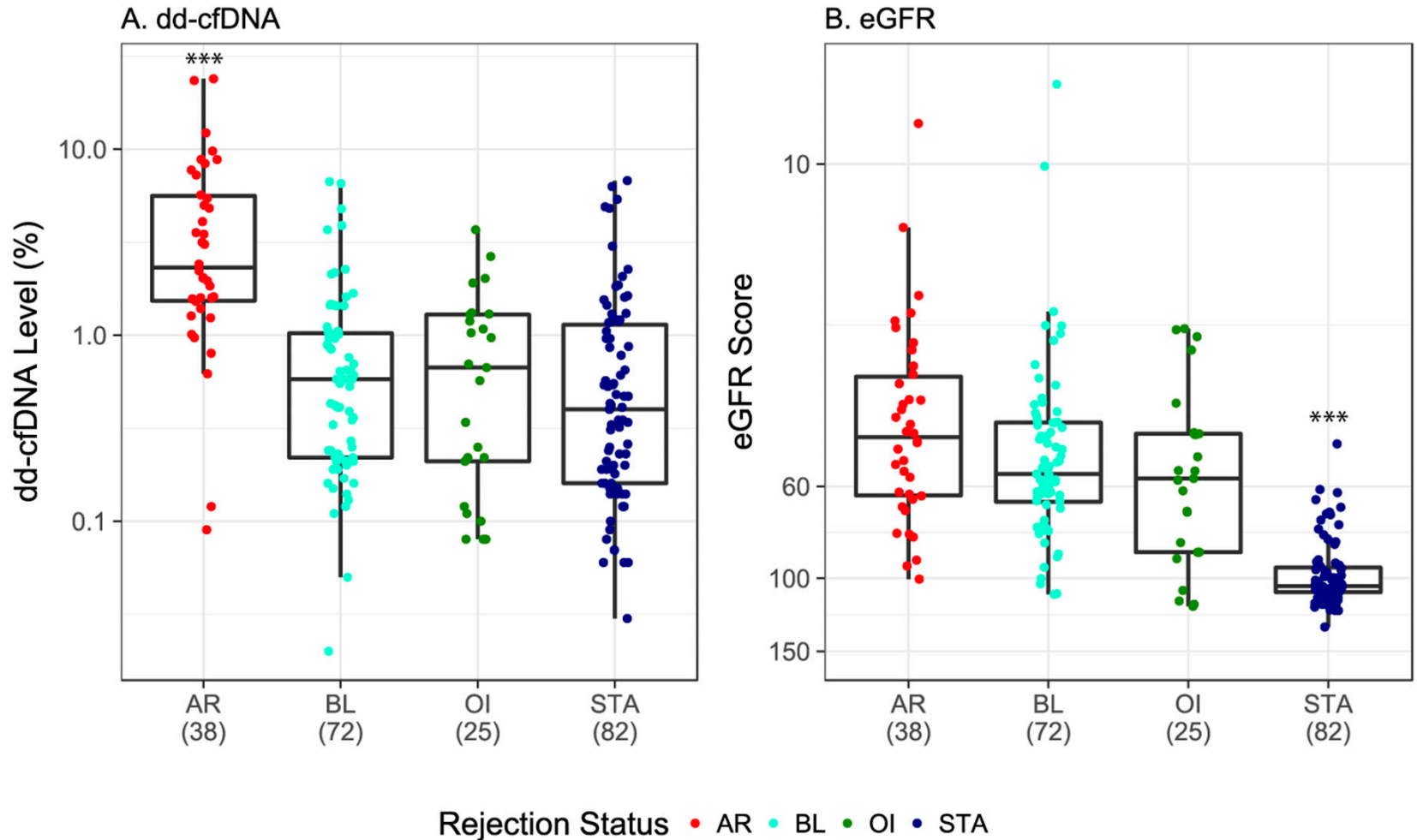
# dd-cfDNA discriminates active rejection better than renal function

dd-cf DNA

Kreatinin

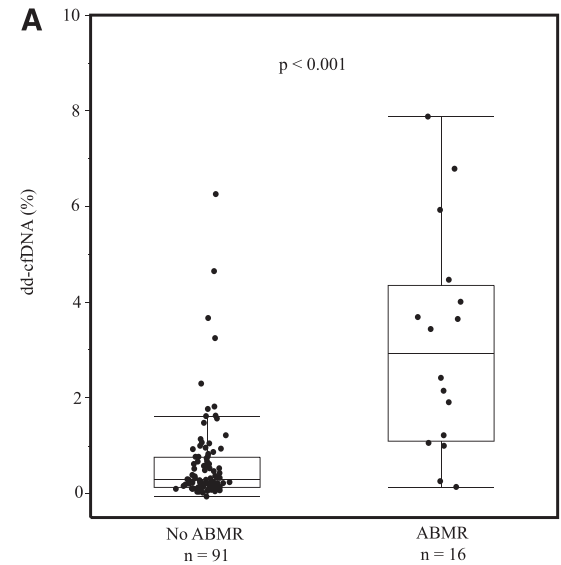
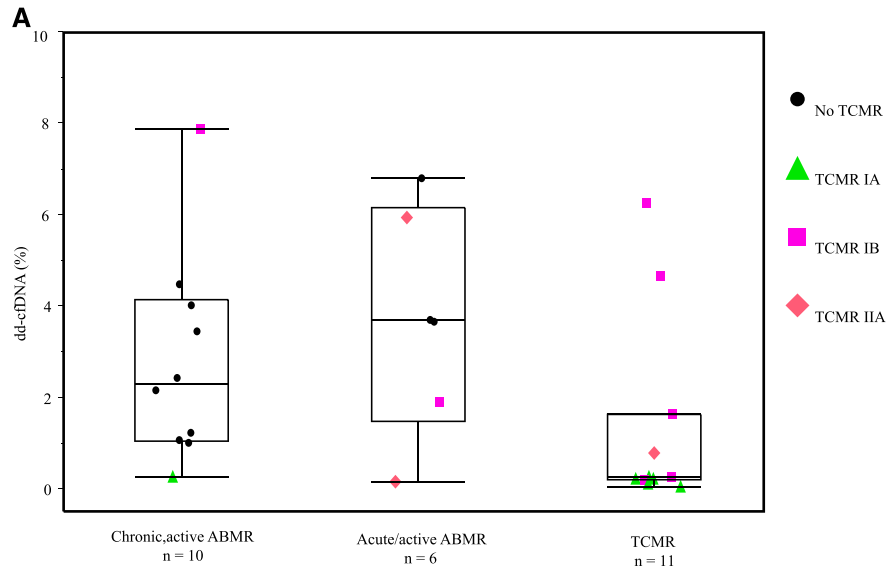


# Discrimination of active rejection by dd-cfDNA versus eGFR

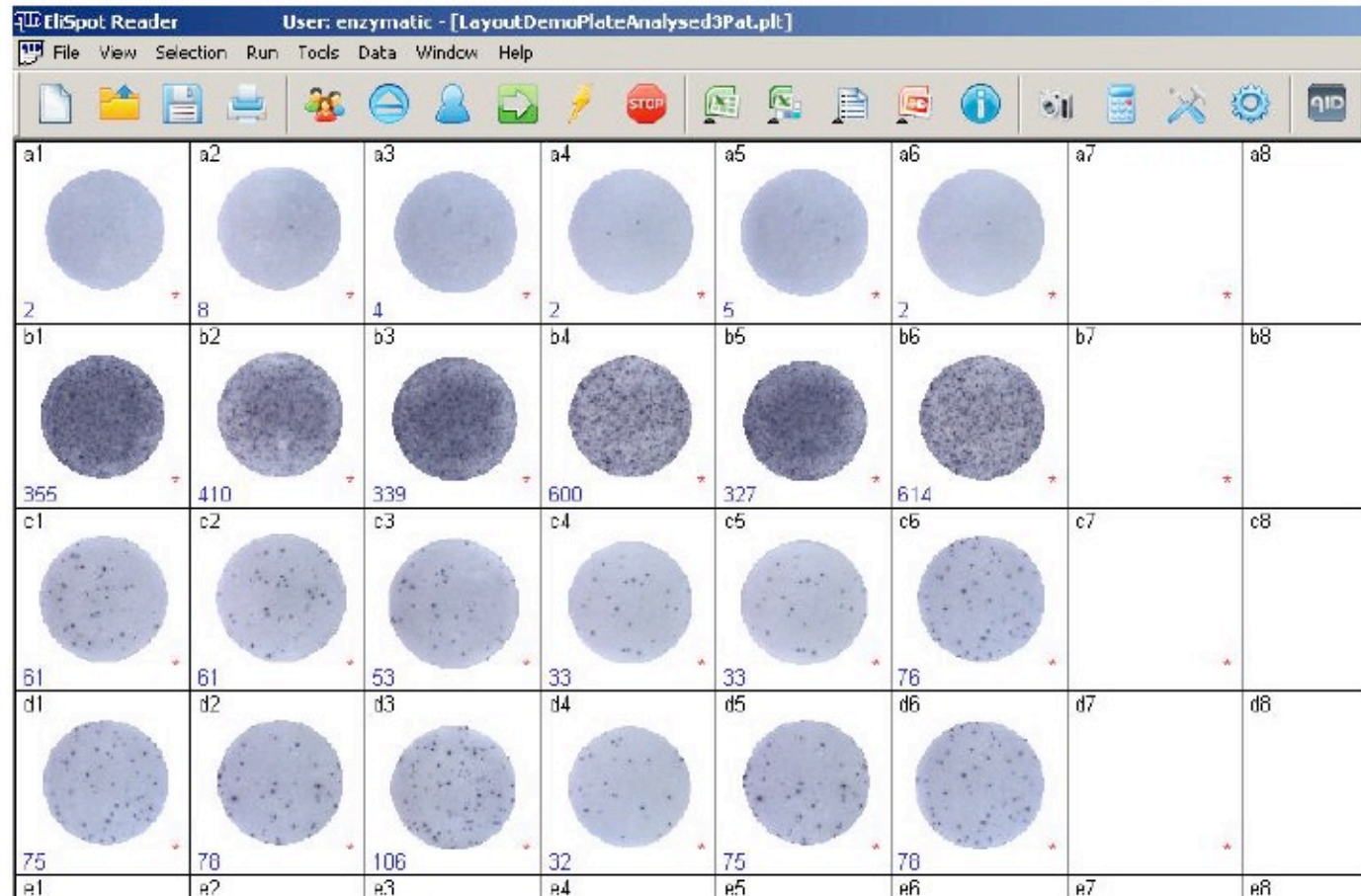
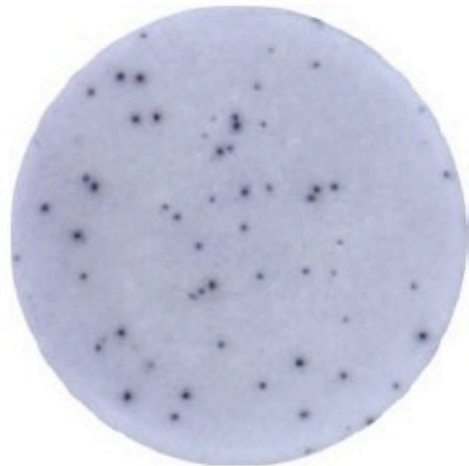


# dd-cfDNA levels are higher in ABMR

*dd-cfDNA discriminates ABMR but not TCMR*



# ELISPOT: metoda k ověření přítomnosti paměťových lymfocytů



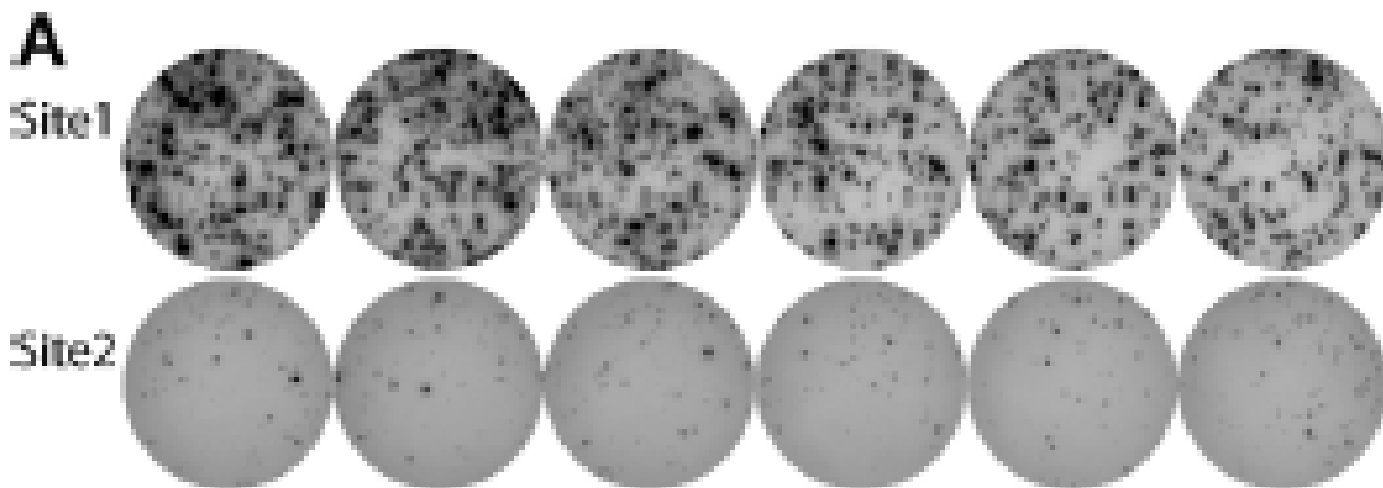
# Dárcovská aloreaktivita koreluje se subklinickou rejekcí

**Table 3 | The presence of 6-month d-s T-cell alloreactivity is associated with acute cellular subclinical rejection (TCSCR) in 6-month protocol biopsies**

TCSCR	6-Month d-s IFN- $\gamma$ Elispot		$P < 0.001$
	NEG	POS	
No	32 (91.4%)	3 (27.3%)	
Yes	3 (8.6%)	8 (72.7%)	
Total	35 (100%)	11 (100%)	46

# IFN $\gamma$ ELISPOT vyšetření bez SOP

Previous ELISPOT data suffers from not standardized methodology: impossible to compare results from different centers



*American Journal of Transplantation* 2013; 13: 1871–1879  
Wiley Periodicals Inc.

© Copyright 2013 The American Society of Transplantation  
and the American Society of Transplant Surgeons

doi: 10.1002/ajt.12286

Brief Communication

**Standardization and Cross Validation of Alloreactive  
IFN $\gamma$  ELISPOT Assays Within the Clinical Trials in  
Organ Transplantation Consortium**

---

ORIGINAL ARTICLE

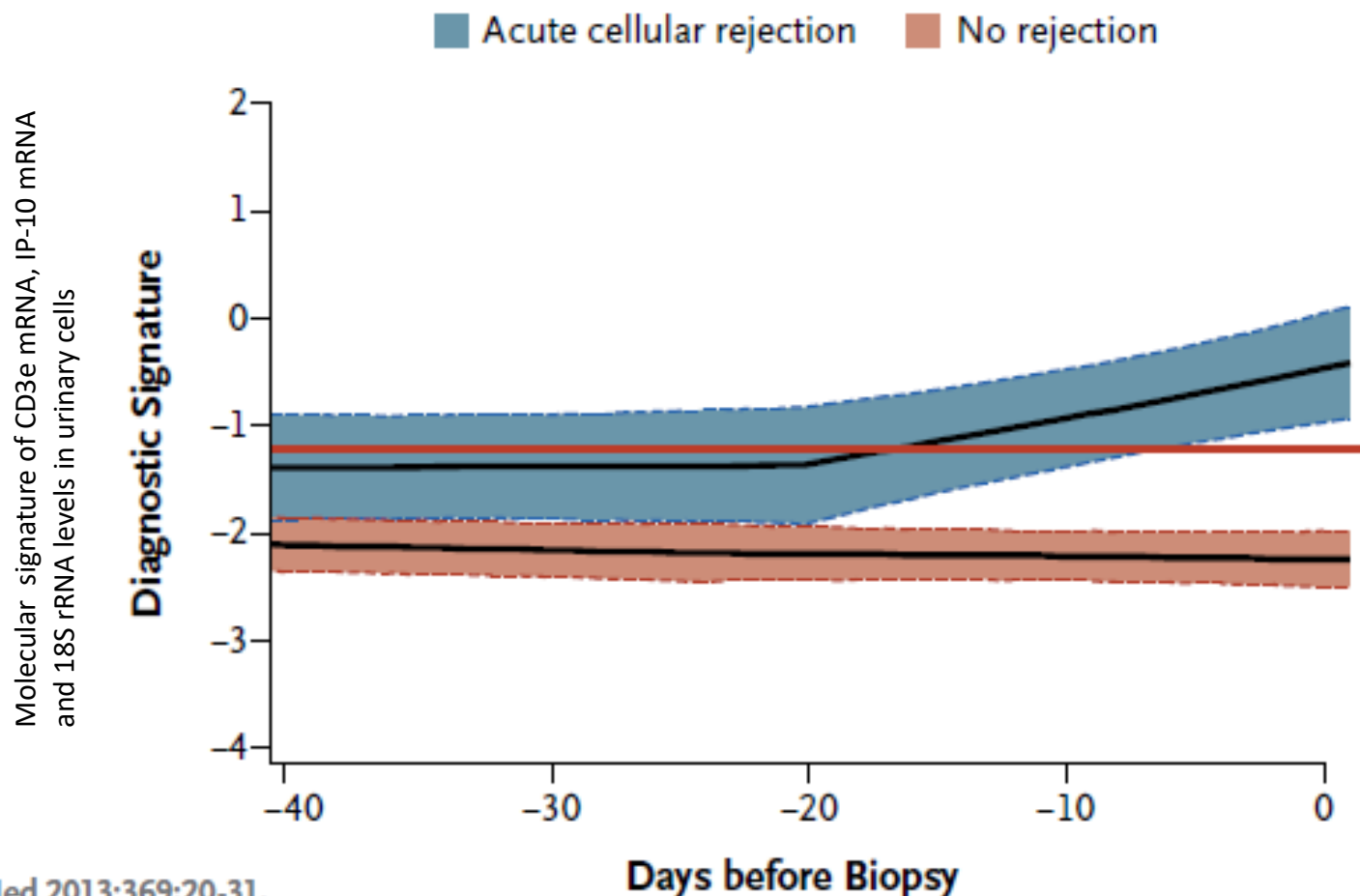
# Urinary-Cell mRNA Profile and Acute Cellular Rejection in Kidney Allografts

Manikkam Suthanthiran, M.D., Joseph E. Schwartz, Ph.D., Ruchuang Ding, M.D., Michael Abecassis, M.D., Darshana Dadhania, M.D., Benjamin Samstein, M.D., Stuart J. Knechtle, M.D., John Friedewald, M.D., Yolanda T. Becker, M.D., Vijay K. Sharma, Ph.D., Nikki M. Williams, B.S., Christina S. Chang, B.S., Christine Hoang, B.S., Thangamani Muthukumar, M.D., Phyllis August, M.D., M.P.H., Karen S. Keslar, M.S., Robert L. Fairchild, Ph.D., Donald E. Hricik, M.D., Peter S. Heeger, M.D., Leiya Han, M.D., M.P.H., Jun Liu, Ph.D., Michael Riggs, Ph.D., M.P.H., David N. Ikle, Ph.D., Nancy D. Bridges, M.D., and Abraham Shaked, M.D., Ph.D., for the Clinical Trials in Organ Transplantation 04 (CTOT-04) Study Investigators

# Vyšetření transkriptů z moče predikuje rejekci

Prospective study of 430 patients monitored, 4000 urinary samples, validation cohort

## D Both Groups, 40 Days before Biopsy



# Imunologické metody použitelné po transplantaci

---

- LUMINEX (anti HLA, DSA)
- FACS (subpopulace T a B buněk, FCXM)
- RT-PCR (mRNAs exprese – transkripty, dd-cfDNA)
- Microarray (MMDx – molekulární patologie)
- ELISA (IP-10, KIM.. nemají prediktivní roli)
- IFN-gamma ELISPOT (zřejmě slepá cesta)

# Možnosti predikce imunologické odpovědi po transplantaci

---

- Výskyt akutní rejekce se v poslední době pohybuje do 12%, v polovině případů jde o humorální rejekci
- Hlavním důvodem ztráty štěpu v dlouhodobém sledování je chronická humorální rejekce
- Předtransplantační predikce humorální odpovědi je stále založena na „starých“ testech PRA, CDC, FACS CM ale rovněž na virtuálním crossmatch (anti HLA skrínig v době zařazení do čekací listiny). Definice neakceptovatelných antigenů respektive neakceptovatelné síly protilátek zůstává nejasně definována, zdá se, že budoucnost má definice akceptovatelných epitopů!
- Z nových testů má největší šanci na klinické uplatnění test prokazující přítomnost dd cf DNA – zatím jen k monitoraci humorální odpověď