

I polimorfismi dei geni *MICA* e *NKG2D* influenzano l'efficacia della terapia immunosoppressiva modificando il rischio di Rigetto Cronico Anticorpo-mediato nel Trapianto di Rene

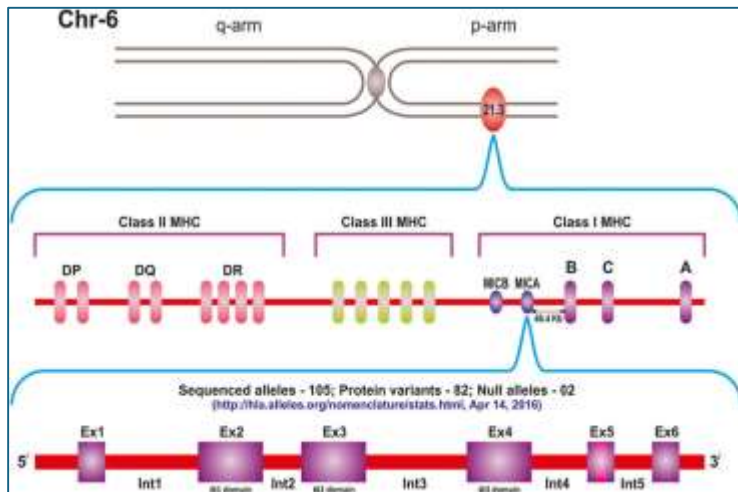


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MICA Gene - MHC Class I

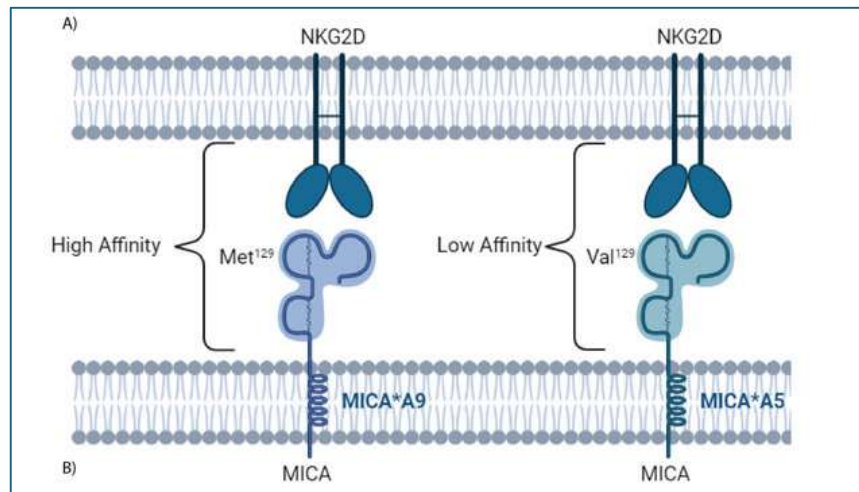
Receptor - NKG2D

MICA Location

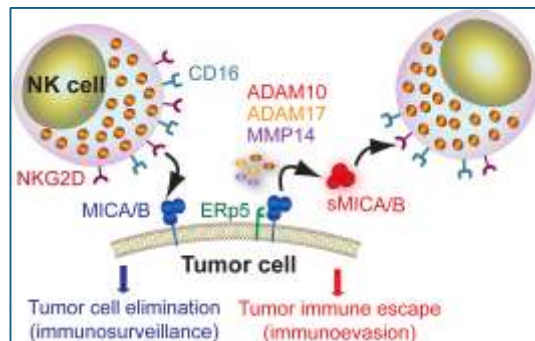


Baranwal AK, Mehra NK. Major Histocompatibility Complex Class I Chain-Related A (MICA) Molecules: Relevance in Solid Organ Transplantation. Front Immunol. 2017

MICA-129 MET/VAL Dimorphism



Major Histocompatibility Complex Class I Chain-Related α (MICA) STR Polymorphisms in COVID-19 Patients –Bautista et al.



Leveraging NKG2D Ligands in Immuno-Oncology

- MICA*008: rilascio esosomi
- MICA*002: taglio proteolitico
- MICA*009: taglio proteolitico
- MICA*010:01: non espresso

Aploblocco -1

Attività
citotossica

dbSNP

Alt
a

Bassa

Aploblocco -2

Attività
citotossica

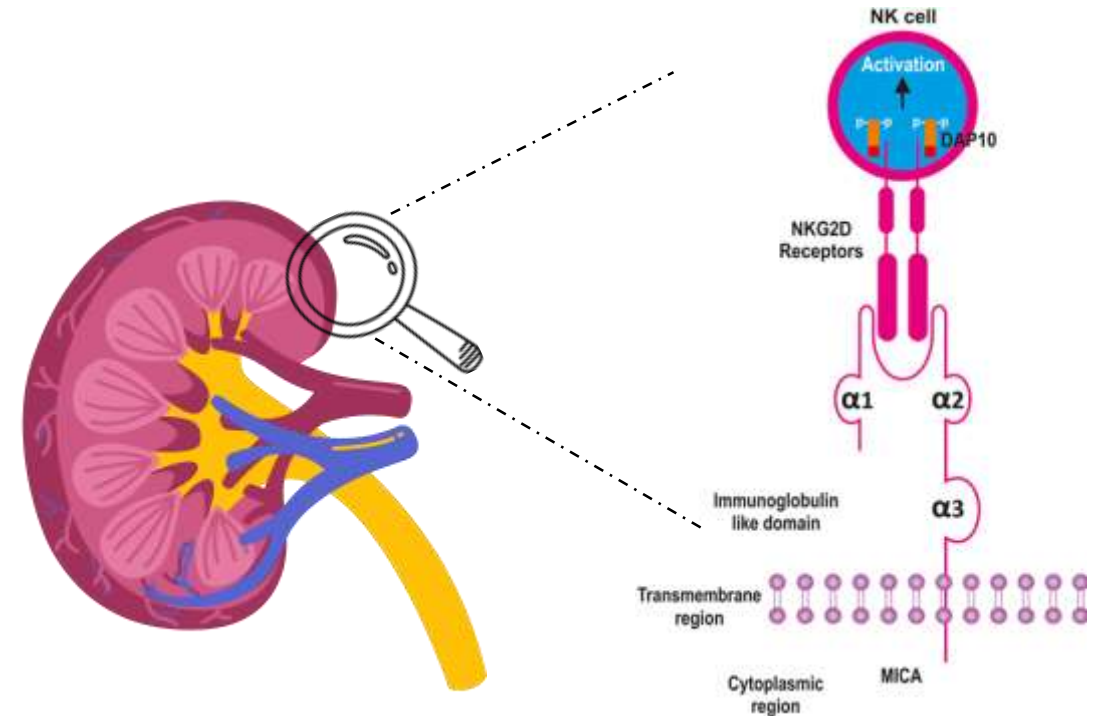
dbSNP

Alt
a

Bassa

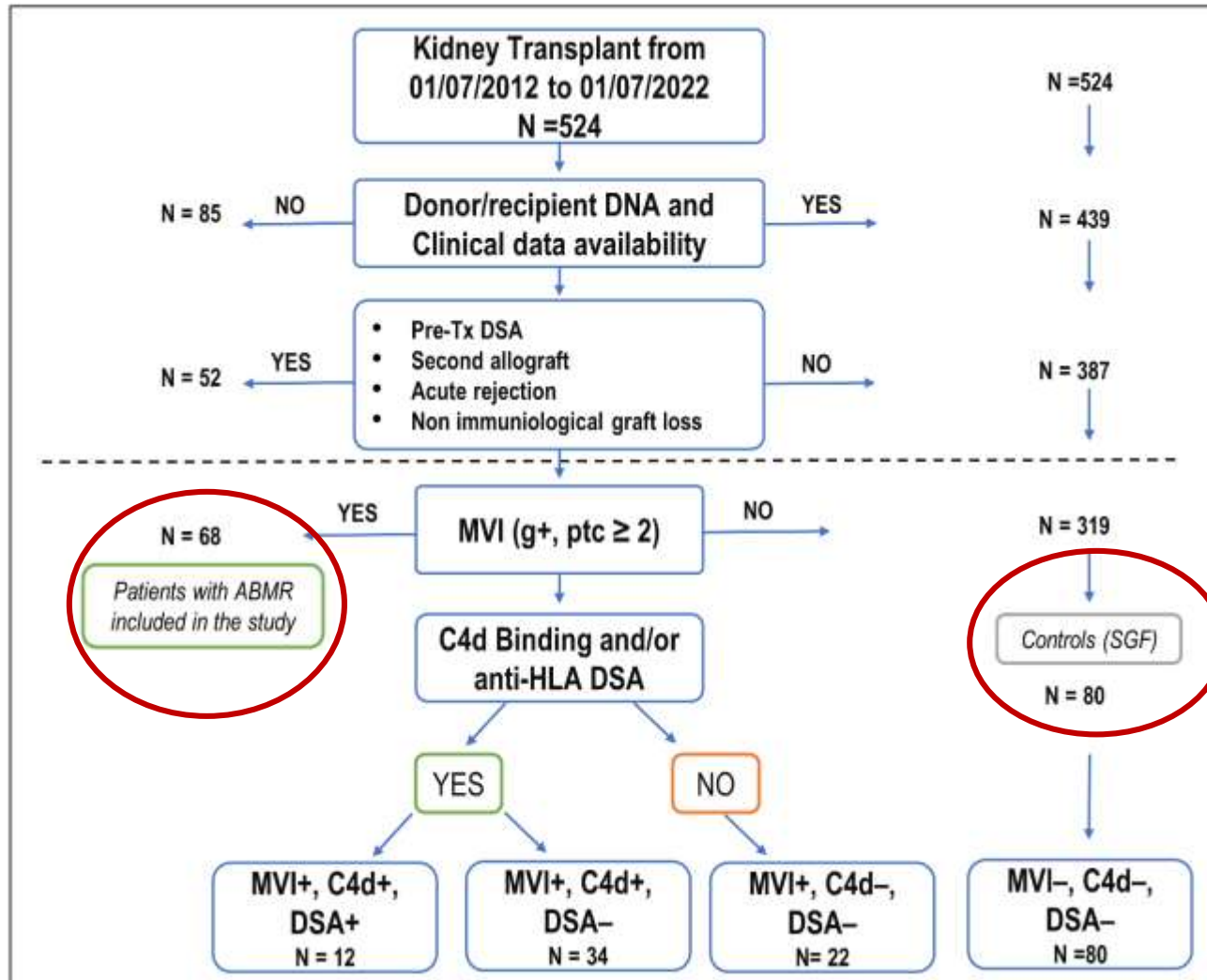
SCOPO DEL LAVORO

Indagare la correlazione tra la compatibilità allelica **MICA** tra ricevente e donatore e il genotipo NKG2D (**KLRK1**) con il rischio di AMR (rigetto anticorpo-mediato) e i loro potenziali effetti clinici e implicazioni per la terapia di mantenimento dell'organo.



Baranwal AK, Mehra NK. Major Histocompatibility Complex Class I Chain-Related A (MICA) Molecules: Relevance in Solid Organ Transplantation. Front Immunol. 2017

METODI



Dati clinici (e.g. Glomerular filtration, Serum creatinine levels, maintenance therapy..)

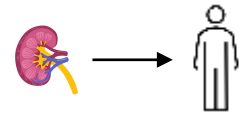
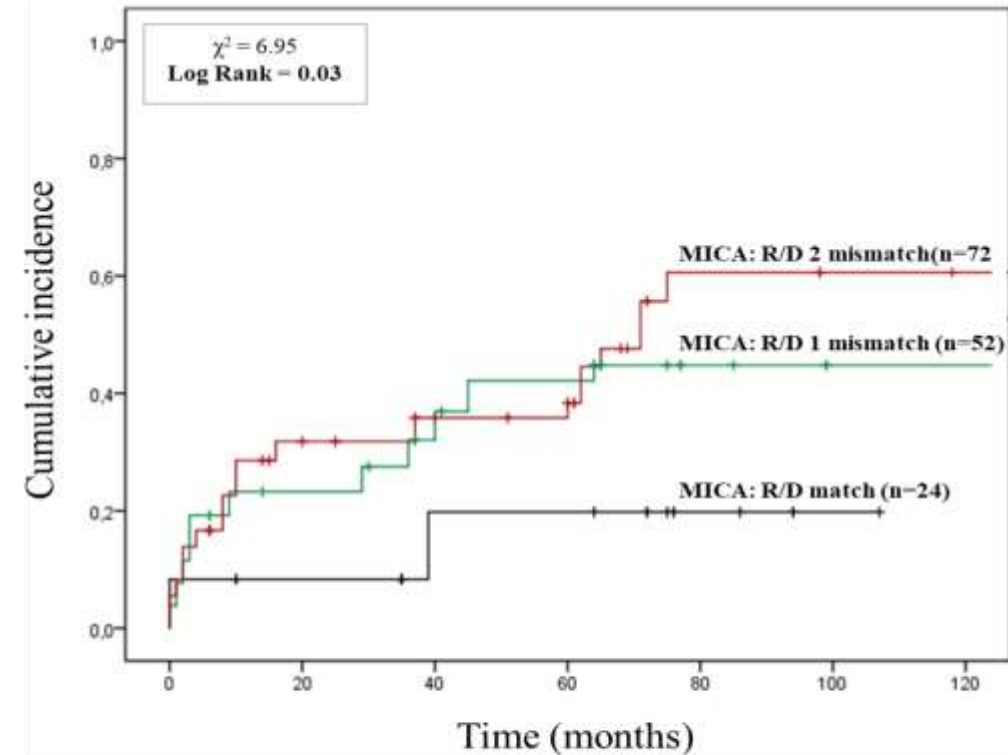
Analisi Genetica

1. HLA loci:

HLA-A, HLA-B, HLA-C, HLA-E, HLA-F, HLA-G, HLA-H, HLA-DRB1, HLA-DRB3, HLA-DRB4, HLA-DRB5, HLA-DQA1, HLA-DQB1, HLA-DPA1, HLA-DPB1, MICA, and MICB.

2. NKG2D Ricevente (KLRK1): *rs1049174, rs2255336 (Aplotipo-1 e 2)*

MICA R/D Mismatch



**MICA R/D
Match**

2MM
53.8%

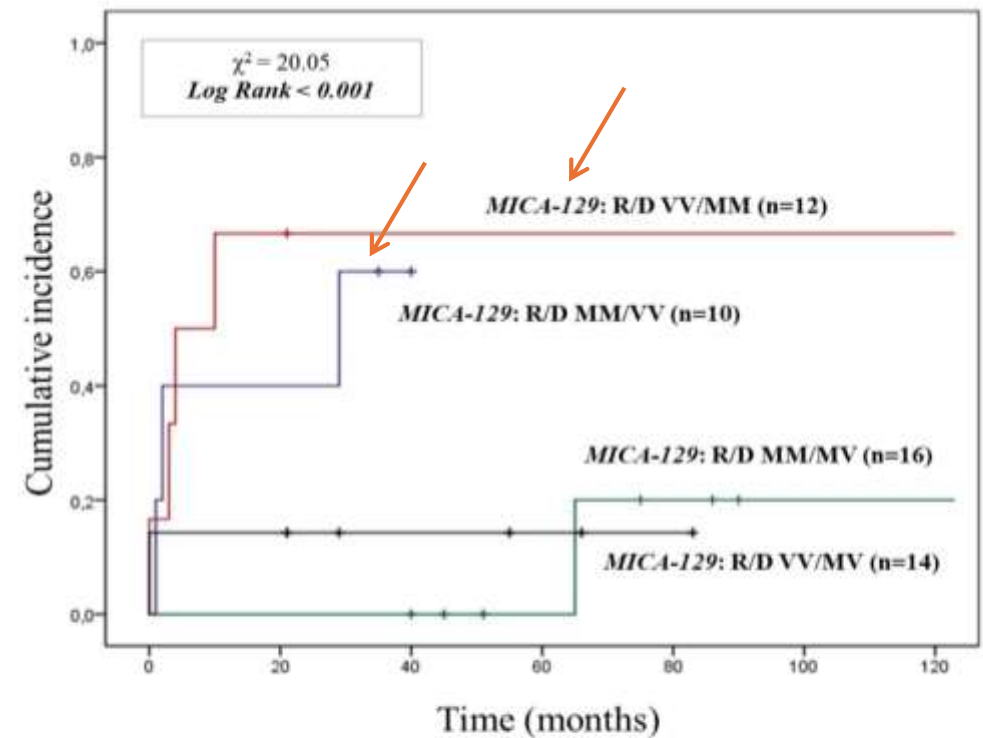
1MM:
50%

0MM
16.7%

INCIDENZ

Time (Months)	0	20	40	60	80	100	120	Number at risk
MICA 2 MM	72	40	27	22	8	6	5	
MICA 1 MM	52	36	26	22	11	7	7	
MICA 0 MM	24	20	14	14	6	2	0	

MICA R/D Mismatch Met/Val 129



**MICA R/D
Met/Val 129**

VV/MM
66,7%

MM/VV
60%

MM/MV
20%

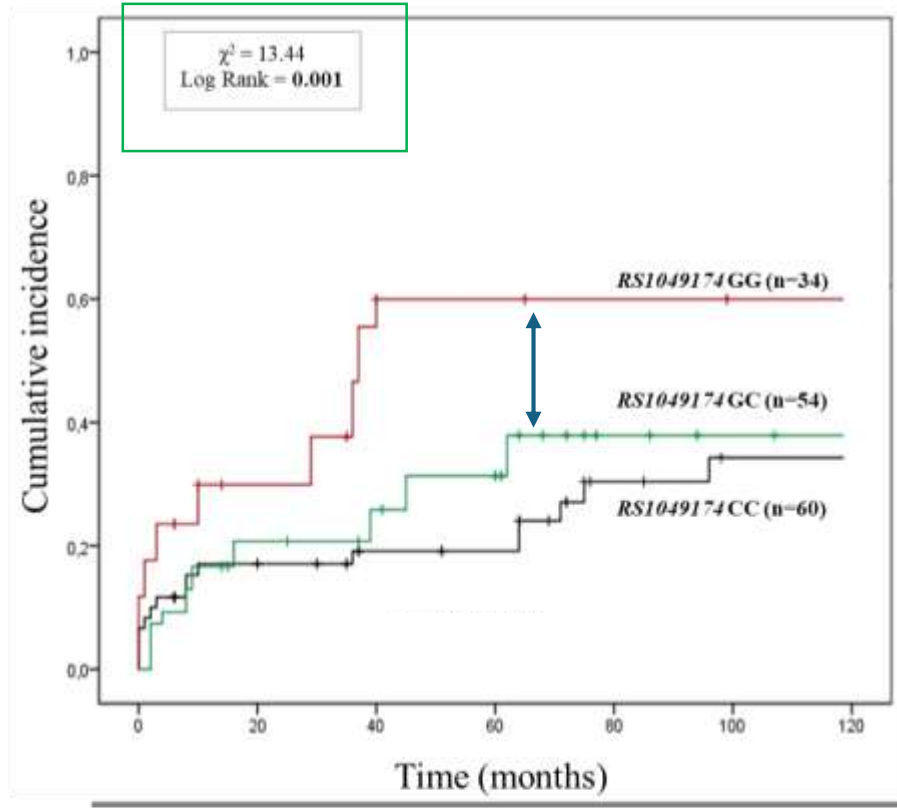
VV/MV
14,3%

INCIDENZ

Time (Months)	0	20	40	60	80	100	120	Number at risk
MICA-129: R/D VV/MM	12	4	3	3	3	3	1	
MICA-129: R/D MM/VV	10	6	2	-	-	-	-	
MICA-129: R/D MM/MV	16	16	14	10	6	2	2	
MICA-129: R/D VV/MV	14	12	6	6	2	-	-	

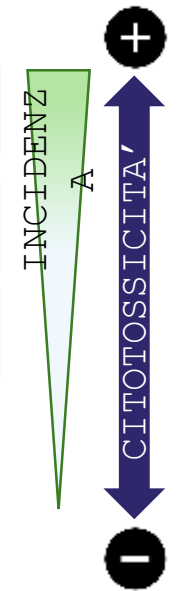
rs1049174
rs2255336
C G
Low High
A

Recipient *rs1049174* NKG2D (Aplotipo-1)



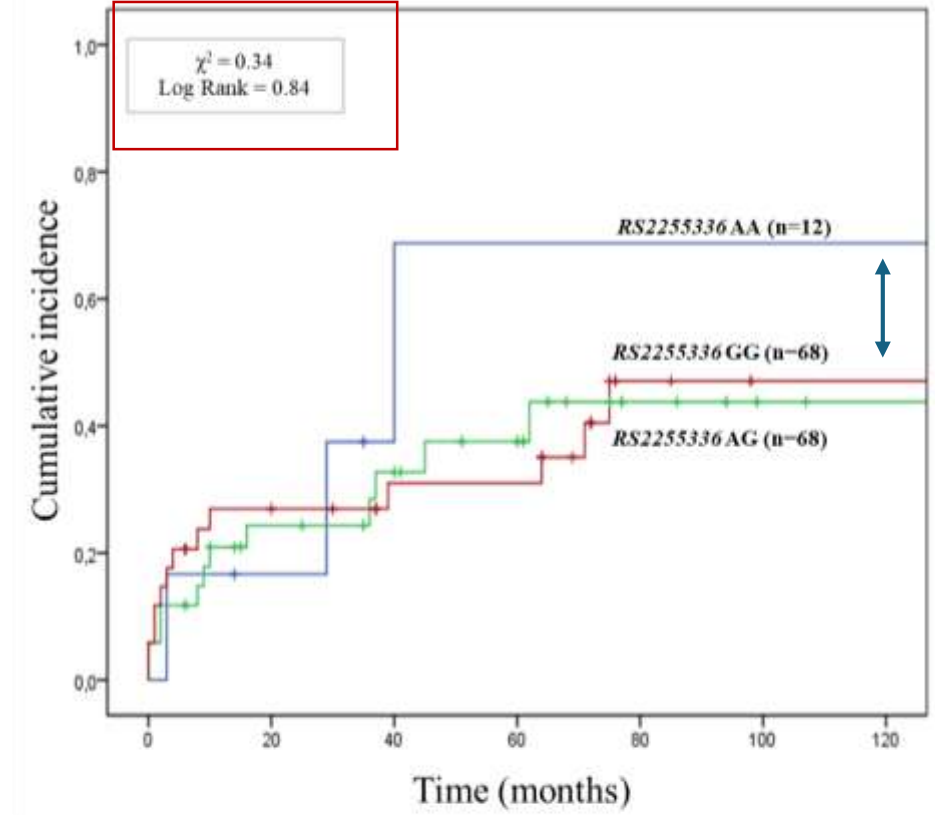
rs1049174 NKG2D

GG: 58.8%
GC 46,6%
CC 37%



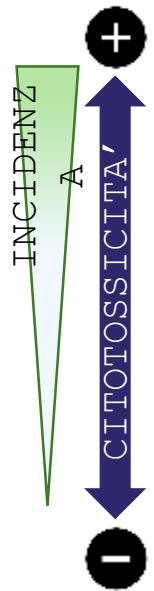
Time (Months)	0	20	40	60	80	100	120	
NKG2D [CC]	60	46	35	33	19	16	16	Number at risk
NKG2D [GC]	54	39	29	25	9	5	3	
NKG2D [GG]	34	18	10	7	5	3	3	

Recipient *rs2255336* NKG2D (Aplotipo-2)



rs2255336 NKG2D

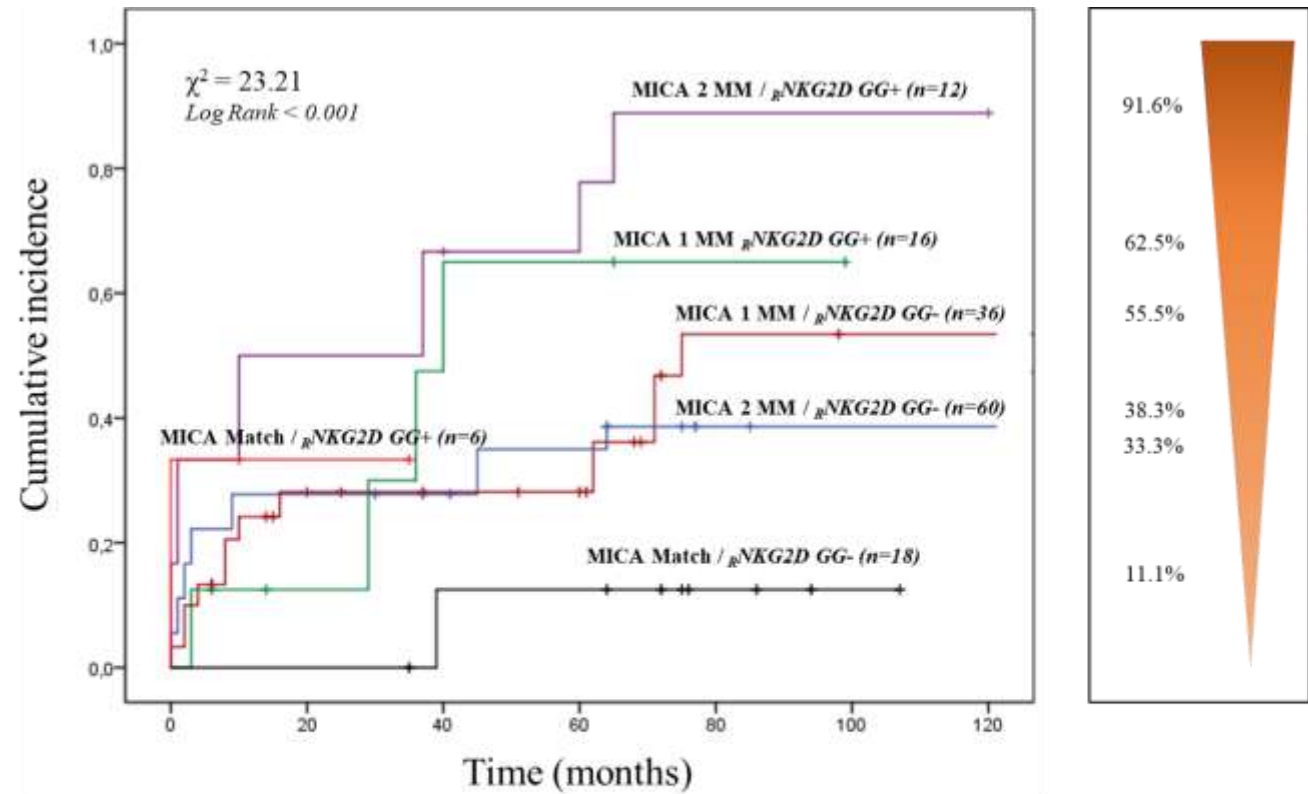
AA: 66.7%
GG 50%
AG 38.2%



Time (Months)	0	20	40	60	80	
NKG2D [AA]	12	8	4	2	2	Number at risk
NKG2D [AG]	68	44	32	24	10	
NKG2D [GG]	68	46	34	34	12	

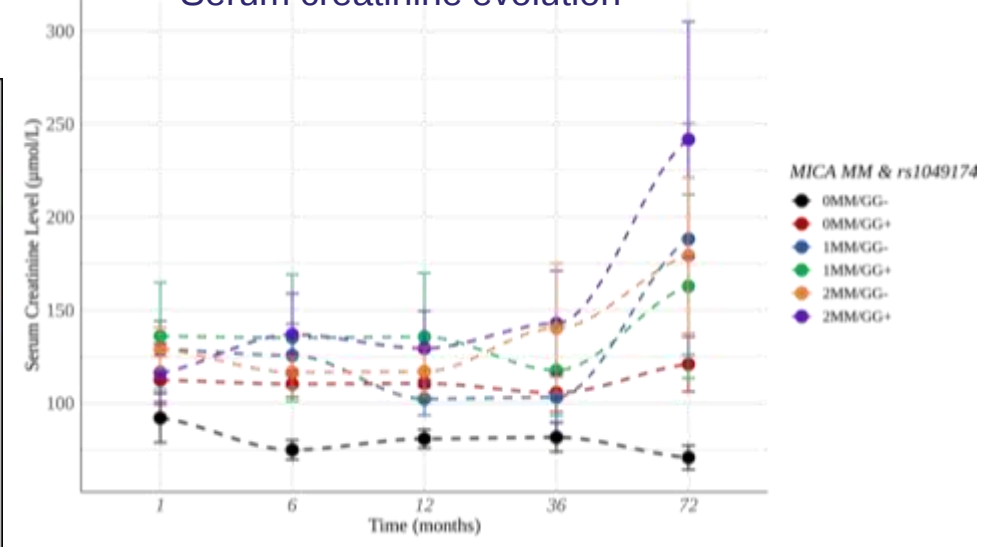
rs1049174 C G
rs2255336 G A
Low High

MICA mismatch + rs1049174 NKG2D

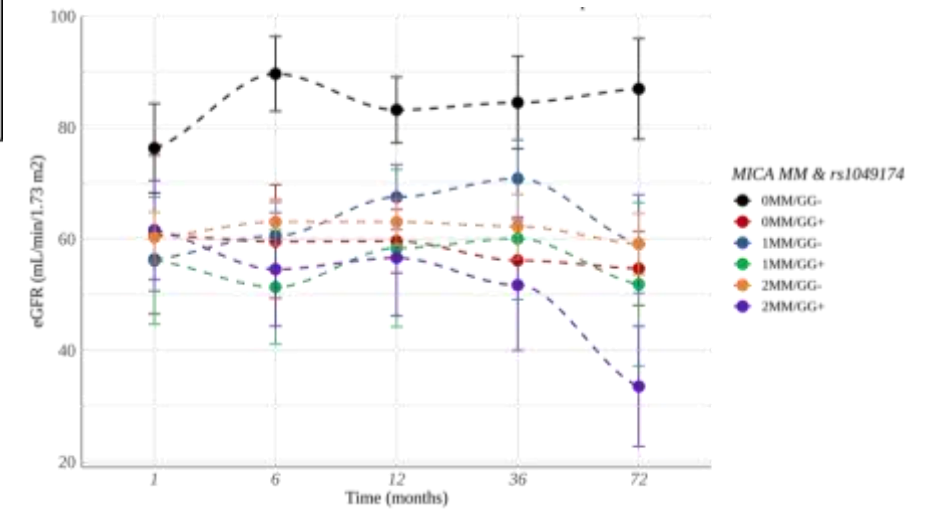


Time (Months)	0	20	40	60	80	100	120	Number at risk
2MM/GG+	12	6	4	2	0	0	0	
1MM/GG+	16	10	6	4	2	0	0	
1MM/GG-	36	26	22	18	9	7	7	
2MM/GG-	60	36	24	22	7	5	5	
0MM/GG+	6	4	2	0	0	0	0	
0MM/GG-	18	18	14	14	12	3	0	

Serum creatinine evolution



Glomerular filtration evolution



Correlation between maintenance therapy, antibody-mediated rejection and genetic profiles

Gene profiles of transplanted patients	Maintenance therapy	SGF (N = 80)	AMR (N = 68)	Comparisons of ABMR vs SGF		
				P value	OR	(95% CI)
MICA alleles 2MM	Calcineurin inhibitors	18	14	0.477	1.6	(0.6 – 4.4)
	mTOR inhibitors	18	22			
Donor MICA Met129/ recipient G	Calcineurin inhibitors	12	2	0.012	0.139	(0.03 – 0.7)
	MTOR inhibitors	20	24			
rs1049174 GG	Calcineurin inhibitors	9	5	0.015	0.2	(0.0 – 1.0)
	mTOR inhibitors	5	15			
rs2255336 AA	Calcineurin inhibitors	4	0	0.002	0	(0.0 – 0.3)
	mTOR inhibitors	0	8			
MICA 2MM + rs1049174 GG	Calcineurin inhibitors	2	0	0.025	0	(0.0 – 0.7)
	mTOR inhibitors	0	10			

*Calcineurin inhibitors: Cyclosporine, tacrolimus

*mTOR inhibitors: Everolimus, Sirolimus

Take-home message

Mismatch R/D nel gene MICA

All'aumentare del numero di mismatches aumenta la probabilità di rigetto anticorpo mediato.

Dimorfismo Met/Val 129 MICA

La presenza di due mismatches influenza in modo comparabile il rischio di rigetto.

- Donor MetMet/RecipientValVal
- Recipient MetMet/Donor ValVal

Polimorfismi recettore NKG2D nel ricevente

-Aplotipo-1: **rs1049174** 3'UTR sito di legame del microRNA-1245 a regolazione negativa. La variante G maggiore espressione di NKG2D in vitro.

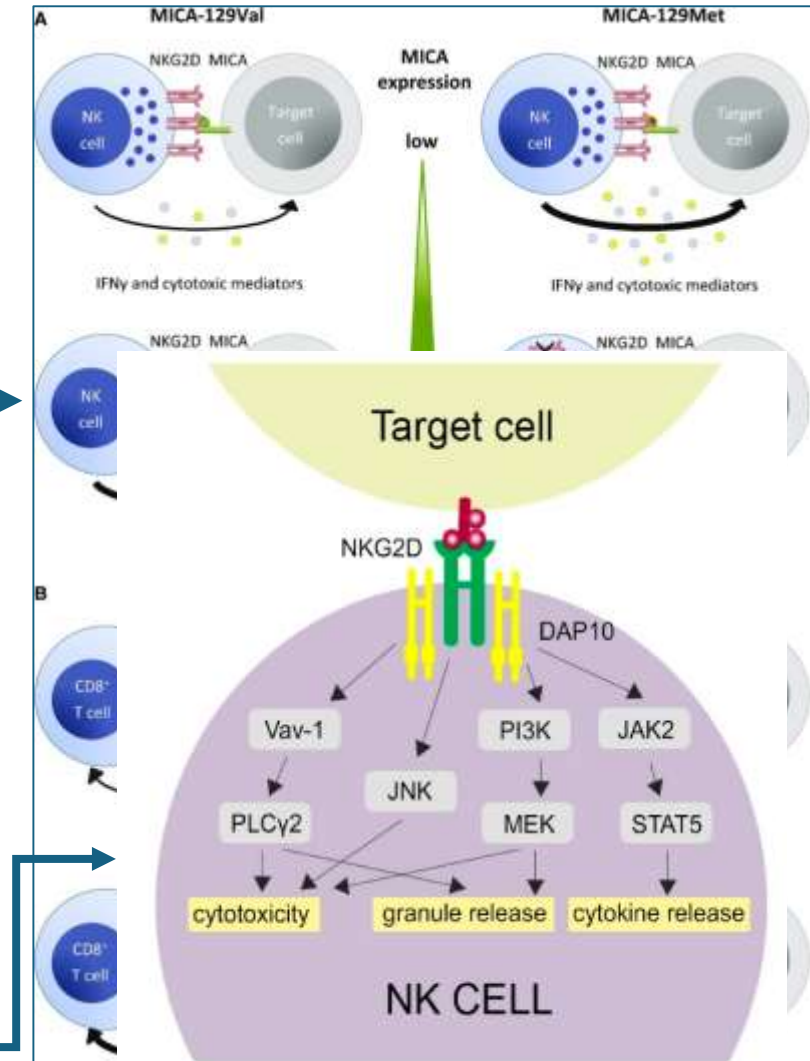
-Aplotipo-2: **rs2255336** sostituzione amminoacidica vicino al sito di legame DAP10 maggiore affinità di legame.

Polimorfismo recettore NKG2D nel ricevente + il doppio mismatches R/D nel gene MICA

La presenza del polimorfismo rs1049174 [GG] nel ricevente e la presenza del doppio mismatches determinano il grado massimo di probabilità di rigetto.

Polimorfismi recettore NKG2D

In presenza del polimorfismo rs1049174 [GG] la risposta alla



Impact of Siemaszko, J, NKG2D Natural Killer Cell Receptor—A Short
Disease Risk Description and Potential Clinical Applications. *Cells* 2021

Grazie per l'attenzione



Università degli studi di Cagliari

Prof. Giglio Sabrina
Dr. Lorrai Michela
Dr. Mereu Caterina
Dr. Murgia Michela
Dr. Sanna Celeste

ASL Cagliari



Dr. Giuressi Erika
Dr. Lai Sara
Dr. Littera Roberto
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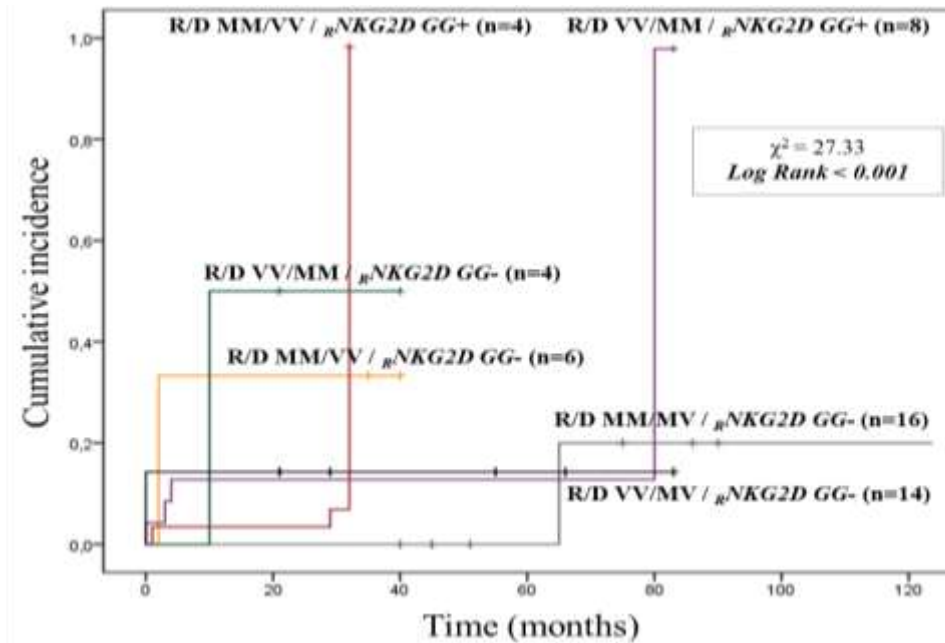
ARNAS Brotzu

Prof. Pani Antonello
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***Tutto il personale
dell'Ospedale Binaghi**

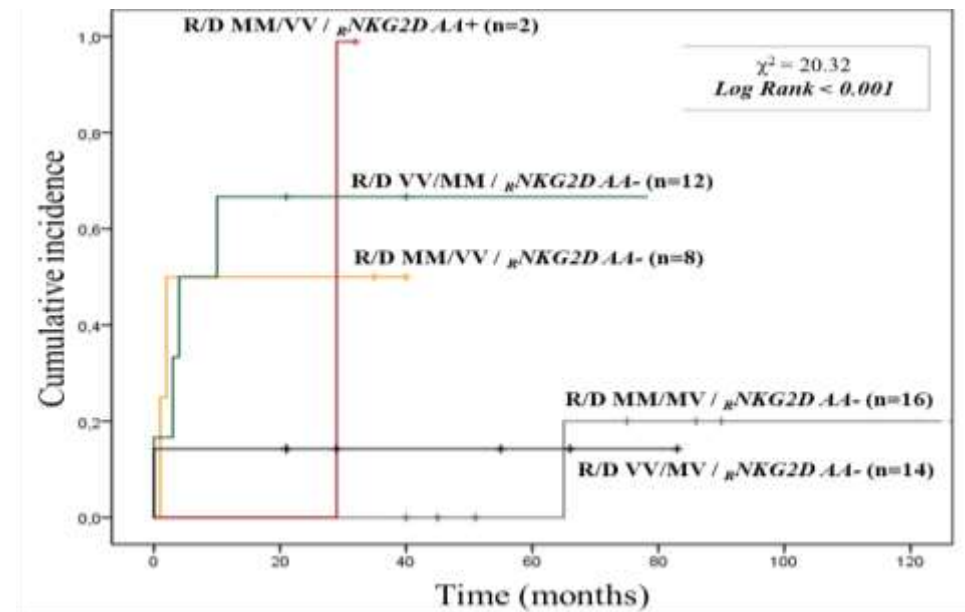


MICA Met/Val 129 & *rs1049174* NKG2D



Time (Months)	0	20	40	60	80	100	120	Number at risk
<i>R/D MM/VV / _RNKG2D GG+</i>	4	2	0	0	0	0	0	
<i>P/D VV/MM / _RNKG2D GG+</i>	8	2	2	2	0	0	0	
<i>R/D VV/MM / _RNKG2D GG-</i>	4	2	0	0	0	0	0	
<i>R/D MM/VV / _RNKG2D GG-</i>	6	4	3	1	0	0	0	
<i>R/D MM/MV / _RNKG2D GG-</i>	16	16	15	12	8	4	2	
<i>R/D VV/MV / _RNKG2D GG-</i>	14	12	6	4	2	0	0	
<i>R/D MM/MV / _RNKG2D GG+</i>	0	-	-	-	-	-	-	
<i>R/D VV/MV / _RNKG2D GG+</i>	0	-	-	-	-	-	-	

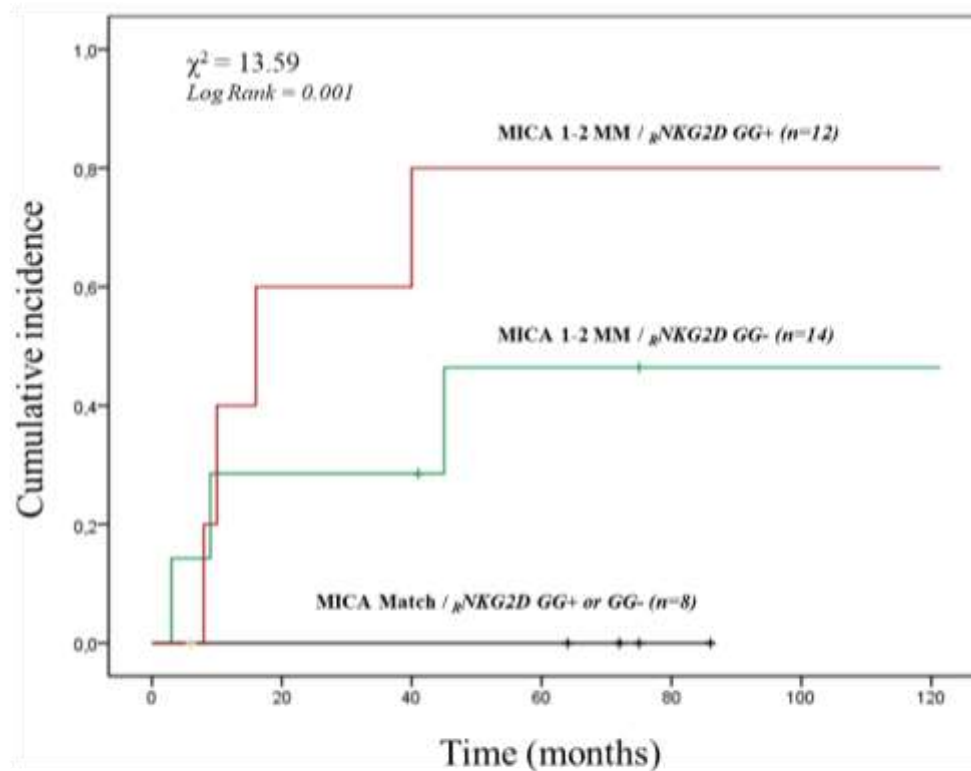
MICA Met/Val 129 & *rs2255336* NKG2D



Time (Months)	0	20	40	60	80	100	120	Number at risk
<i>R/D MM/VV / _RNKG2D AA+</i>	2	2	0	0	0	0	0	
<i>R/D VV/MM / _RNKG2D AA-</i>	12	4	1	1	1	1	0	
<i>R/D MM/VV / _RNKG2D AA-</i>	8	4	1	0	0	0	0	
<i>R/D MM/MV / _RNKG2D AA-</i>	16	16	14	10	6	2	2	
<i>R/D VV/MV / _RNKG2D AA-</i>	14	12	6	4	2	0	0	
<i>R/D VV/MM / _RNKG2D AA+</i>	0	-	-	-	-	-	-	
<i>R/D MM/MV / _RNKG2D AA+</i>	0	-	-	-	-	-	-	
<i>R/D VV/MV / _RNKG2D AA+</i>	0	-	-	-	-	-	-	

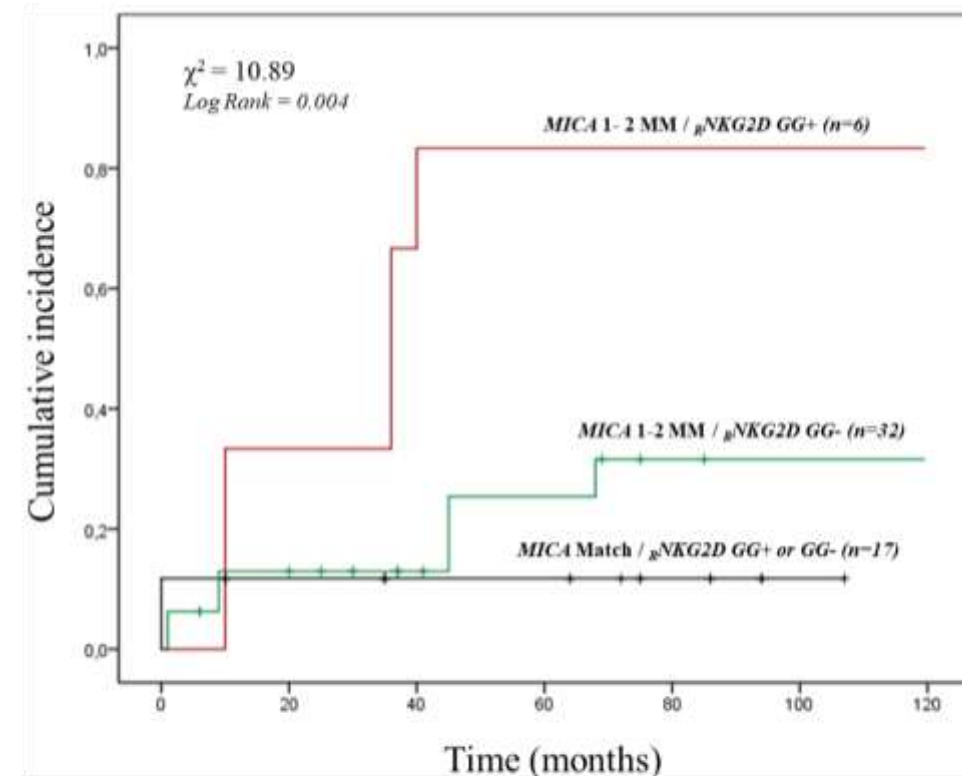
rs1049174 C G
rs2255336 G A
Low High

Kidney Antibody-mediated rejection: *R/D HLA-DRB1* match



Time (Months)	0	20	40	60	80	100	120	Number at risk
MICA 1-2 MM / <i>NKG2D</i> GG+	12	5	5	1	1	1	1	
MICA 1-2 MM / <i>NKG2D</i> GG-	14	10	10	6	5	5	5	
MICA Match / <i>NKG2D</i> GG+ or GG-	7	7	7	7	1	0	0	

Kidney Antibody-mediated rejection: *R/D HLA-A,-B,-C* ≤3 mismatches



Time (Months)	0	20	40	60	80	100	120	Number at risk
MICA 1-2 MM / <i>NKG2D</i> GG+	6	4	2	1	1	1	1	
MICA 1-2 MM / <i>NKG2D</i> GG-	32	26	16	12	7	5	5	
MICA Match / <i>NKG2D</i> GG+ or GG-	17	13	10	10	5	1	0	