

HSC transplant: role of amino acid substitutions in antigen binding sites

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Introduction

Robust evidence supports the adverse impact of donor-recipient HLA disparity on Hematopoietic Stem Cells Transplantation (HCT) outcomes and thus it is necessary to investigate the relationship between mismatches and transplant outcome as a 10/10 match cannot be found for all patients in need of transplantation. Interesting studies conducted on amino acid substitutions (AAS) in the antigen binding sites (Pidala et al., Blood, 2013) showed the relevance of the HLA-B mismatches in position 9 (chronic GvHD increase), HLA-C in position 99 (increase in transplant-related mortality), HLA-C in position 116 (acute GvHD increase). These mismatches are classified as nonpermissive if they lead to an amino acid change.

Objectives

We studied the impact of these AAS on the transplant outcome of donor-recipient pairs from 2005 to now in collaboration with the Transplant Center (Molinette of Turin).

Methods

Patients and donors had completed high-resolution typing for HLA-A, B, C, DRB1, DQB1 (GenDx kits). In this period 931 patients underwent an HSC allografts, the study included 430 patients who received a first transplant from haploidentical sibling donors (HAPLO 106), MUD 10/10 (239), MUD 9/10 with a permissive mismatch at the loci HLA-B or HLA-C (P-MM 46), MUD 9/10 with a nonpermissive mismatch at the same loci (NP-MM 39).

Results

MUD and HAPLO sibling donors transplants were compared in the period from 2011 to 2021. We analysed patient survival related to different variables like gender, age, original disease of recipient, onset and severity of GvHD, source of HSC. **The survival at 3 years in recipient of HAPLO and, MUD.**

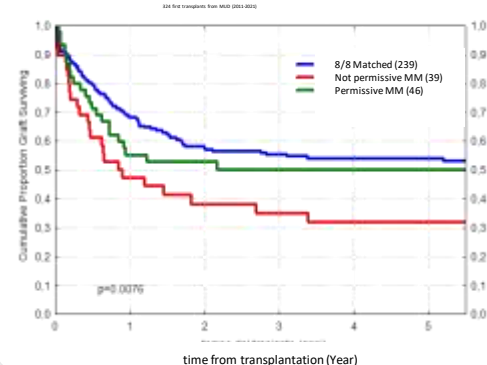
		From MUD donor	From HAPLO sibling donors	P-value
N. TX 2005-2021		305	104	
Gender (recipient (M/F))		196/109	62/42	P=0.776
Age (year) (sd)		50.5(14.9)	50.0(12.7)	P=0.748
Lymphoid disease	AML	130 (42.6%)	47 (45.2%)	P=0.659
	Erythroid	57 (38%)	11 (24%)	
	ALL	40 (12%)	11 (10.6%)	
	SALL	35 (10%)	0	
	MDS	48 (15%)	11 (10.6%)	
	Chronic myelophagocytic Disease	13 (4%)	1 (0.9%)	
Myeloid disease	Myeloma	15 (5%)	0	P=0.666
	Other	2 (0.6%)	2 (1.9%)	
	Cellular disease	Base disease	35 (10%)	
Prothrombotic blood	289 (89%)	59 (56%)		
Age donor (sd)		30/14.4	32.3(12.7)	P=0.013
Amino C-0:02		14%	16%	P=0.544

Results

We analysed patient survival related to different variables like gender, age, original disease of recipient, onset and severity of GvHD, source of HSC. **The survival at 3 years in recipient of HAPLO, MUD and P-MM transplants is similar (53%), while it is less for NP-MM (34%).**

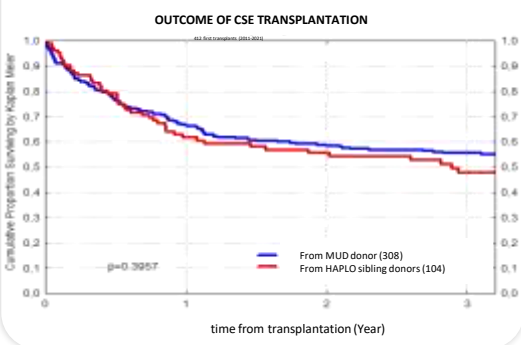
		FIRST HCT TRANSPLANTS			P-value
		8/8 MATCHED	PERMISSIVE MISMATCH	NOT PERMISSIVE MISMATCH	
N. TX 2005-2021		239	46	39	
Gender (recipient (M/F))		152/87	34/12	22/17	P=0.662
Cellular disease	Base disease	24	5	4	P=0.300
	Prothrombotic blood	215	41	35	
Age donor (sd)		29.3 (4.5)	33.0(4.5)	37.2(4.3)	P=0.004
C-0:02 status		52.7%	46%	46.6%	P=0.476
Lymphoid disease	AML	10	15	14	P=0.528
	Erythroid	0	14	9	
	ALL	10	7	3	
	SALL	4	1	1	
	MDS	38	4	4	
	Chronic myelophagocytic Disease	10	0	2	
Myeloid disease	Myeloma	15	1	4	P=0.528
	Other	0	0	0	

OUTCOME OF CSE TRANSPLANTATION
for HCT transplants from MUD (2011-2021)



Discussion & Conclusion

The survival at 5 years for P-MM is 52% significantly better than for Not Permissive-MM transplants (32%, p = 0.04). This relevant information will be provided to the Transplant Center as a tool for the selection of donors.



References

Pidala et al., Blood, 2013

