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# ORIGINAL ARTICLE



# AJT

# Allograft and patient survival after sequential HSCT and kidney transplantation from the same donor—A multicenter analysis

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Abbreviations: aGVHD, acute graft-versus-host disease; AIC, Akaike Information Criterion; ALL, acute lymphoid leukemia; AML, acute myeloid leukemia; AZA, azathioprine; BMT-NP, bone marrow transplantation nephropathy; CML, chronic myeloid leukemia; CNI, calcineurin inhibitor; CYA, cyclosporin A; DBD, donor after brain death; DSA, donor-specific antibody; ESRD, end-stage renal disease; GN, glomerulonephritis; GVHD, graft-versus-host disease; HSCT, hematopoietic stem cell transplantation; HUS, hemolytic uremic syndrome; IS, immuno-suppression; MMF, mycophenolate mofetil; NK, not known; N, no; Nr, number; OEDTR, Austrian Dialysis and Transplant Registry; RT, renal transplantation; SCD, sickle cell disease; SMD, standardized mean difference; TAC, tacrolimus; TTP, thrombotic thrombocytopenic purpura; Yr, years; Y, yes.

Michael Eder, Christoph Schwarz, Thomas Wekerle, and Rainer Oberbauer contributed equally.

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Tolerance induction through simultaneous hematopoietic stem cell and renal transplantation has shown promising results, but it is hampered by the toxicity of preconditioning therapies and graft-versus-host disease (GVHD). Moreover, renal function has never been compared to conventionally transplanted patients, thus, whether donor-specific tolerance results in improved outcomes remains unanswered. We collected follow-up data of published cases of renal transplantations after hematopoietic stem cell transplantation from the same donor and compared patient and transplant kidney survival as well as function with caliper-matched living-donor renal transplantations from the Austrian dialysis and transplant registry. Overall, 22 tolerant and 20 control patients were included (median observation period 10 years [range 11 months to 26 years]). In the tolerant group, no renal allograft loss was reported, whereas 3 were lost in the control group. Median creatinine levels were 85 µmol/l (interquartile range [IQR] 72-99) in the tolerant cohort and 118 µmol/l (IQR 99-143) in the control group. Mixed linear-model showed around 29% lower average creatinine levels throughout follow-up in the tolerant group (P < .01). Our data clearly show stable renal graft function without long-term immunosuppression for many years, suggesting permanent donor-specific tolerance. Thus sequential transplantation might be an alternative approach for future studies targeting tolerance induction in renal allograft recipients.

#### KEYWORDS

bone marrow/hematopoietic stem cell transplantation, clinical research/practice, kidney (allograft) function/dysfunction, kidney transplantation/nephrology, tolerance: clinical

## 1 | INTRODUCTION

Renal transplantation (RT) is the treatment of choice for eligible patients with end-stage renal disease (ESRD). Whereas short- and intermediate-term outcomes of renal allografts have been improved due to fewer early rejections and eventually through better standardized immunosuppressive treatment strategies, evidence that these effects have significantly enhanced long-term transplant kidney survival is still missing.<sup>1,2</sup> Therefore, the induction of tolerance is considered as the ultimate goal in the field of transplantation. Currently, the only clinically successful way to induce donor-specific tolerance is to establish full or mixed chimerism through hematopoietic stem cell transplantation (HSCT).<sup>3</sup>

Since the concept of chimerism leading to selective donor-specific tolerance has been developed in the past century by Billingham, Medawar, and colleagues,<sup>4</sup> substantial progress has been made in terms of preconditioning and the understanding of relevant pathways involved herein.<sup>3</sup> However, despite promising results in murine studies, its widespread use has been hampered by the cytotoxic preconditioning, which is currently required for bone marrow transplantation. Clinically, the first evidence of tolerance through HSCT was delivered by Sayegh et al in 1991, who described a 30-year-old man, diagnosed with acute myelomonocytic leukemia and a 14-year-old girl with acute nonlymphocytic leukemia. Both received HSCTs from

their HLA-identical brothers. After a few years and gradual deterioration of their renal function, the original stem cell donors agreed to also donate a kidney. Except for small maintenance doses of steroids, no immunosuppression was given in the further course and transplant kidney function remained stable.<sup>5</sup> Based on this success more case reports followed, including HLA-mismatched donors, who successfully donated renal transplants after HSCT.<sup>6-21</sup>

Due to a full donor chimerism in these patients, a robust state of tolerance for the donated organ is expected. Nevertheless, most of these patients received at least short-term immunosuppression after transplantation. This was done not only to prevent renal allograft rejection but also to prevent graft-versus-host disease (GVHD), which theoretically could be triggered by solid organ transplantation.<sup>22,23</sup> However, even though the concept of tolerance induction through HSCT was established more than 2 decades ago, long-term follow-up data from patients after sequential transplantation of hematopoietic stem cells and renal allografts from the same donor are mostly monocentric and limited by small patient numbers or a short follow-up.<sup>10,18,24</sup> Moreover, due to the limited patient numbers, comparisons to conventionally transplanted patients were not feasible and so far, uncertainty remains whether tolerance truly improves renal allograft survival.

In this study we aimed at analyzing the long-term outcome of patients after sequential HSCT and RT from the same donor by

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comparing renal allograft function and patient survival with a matched cohort of conventionally transplanted live donor kidney recipients from the Austrian Dialysis and Transplant Registry (OEDTR).

## 2 | METHODS

#### 2.1 | Systematic literature research

To analyze the long-term outcome of patients after sequential HSCT and RT from the same donor, we conducted a systematic literature search in PubMed and the Cochrane database. In addition, the references of all relevant papers were manually reviewed for further reports. Studies describing combined HSCT and RT were not included because of different clinical conditions (mixed chimerism vs full chimerism, patients with ESRD but without preexisting malignant comorbidity) and different study designs.

#### 2.2 | Data retrieval

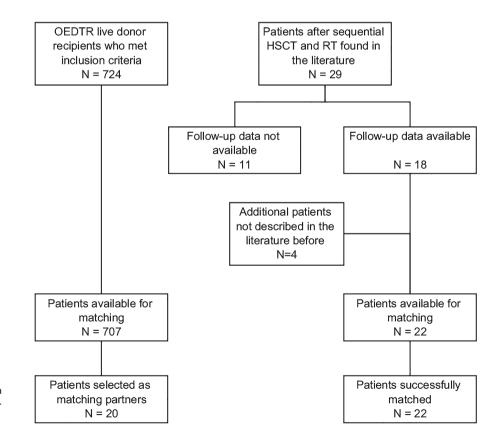
First, corresponding authors of the papers identified in the literature search were contacted and asked for follow-up data. Data were fully anonymized, or anonymization was performed immediately after reception. Patients were included in the analysis if the followup period was longer than originally described in the case report or additional information concerning patient or renal graft survival was available. Some authors reported that further, unpublished patients with previous HSCT from the same donor were transplanted. These patients were also included in our analysis. The analysis was conducted in accordance with the guidelines of good clinical practice and approved by the ethics committee of the Medical University of Vienna (EK # 1218/2017 and 1535/2012).

#### 2.3 | Requested parameters

Basic demographic and clinical characteristics were extracted from the reports: gender, relation of donor, age at each transplantation, indication for HSCT, cause of ESRD, HLA-mismatch, immunosuppressive regimen after RT, creatinine at each follow-up ( $\mu$ mol/I), episodes of GVHD, incidence of rejection of the renal allograft and date of last follow-up.

# 2.4 | Control group of conventionally transplanted living-donor recipients

To compare the long-term patient and transplant kidney-survival of sequentially with conventionally transplanted patients, data from recipients of renal transplants were extracted from the OEDTR.<sup>25</sup> The OEDTR was established by the Austrian Society of Nephrology in 1970 and has almost complete follow-up. The registry comprises data extracted from the original medical records, which were created by the responsible physician at the time of a patient's follow-up visit. All recorded ESRD patients who received their first single-organ kidney transplant from a living donor between 1990 and 2012, were eligible for this study and followed up until 2014 (N = 724). To achieve a more balanced analysis of the long-term renal allograft function, we excluded patients who lost



**FIGURE 1** Flow chart illustrating data retrieval and further patient selection for the matching procedure

		Conventional F	Conventional RT		
Variable	Tolerant patients	Matched	SMD	Pre-matching	SMD
Number of patients	22	20	-	707	-
Age at RT* (y)	33 (14)	33 (15)	4%	39 (17)	33%
Age of kidney donor* (y)	46 (13)	47 (11)	5%	49 (11)	27%
Sex match at renal transplantation* (matched)	12 (55%)	12 (60%)	10%	268 (38%)	23%
Sex (female)	10 (46%)	8 (40%)	10%	241 (34%)	35%
Overall serum creatinine after RT (µmol/I)	85 (72, 99)	118 (99, 143)	-	142 (115, 178)	-
Median total follow-up time (y)	10 (7, 22)	11 (10, 14)	-	7 (4, 13)	_
Mean number of mismatched HLA antigens at RT*	nismatched HLA		84%	2.9 (1.6)	172%
Mean number of mismatched HLA antigens at HSCT	1 (1.3)	_			
Age at HSCT (y)	23 (17, 30)	-			
Time between HSCT and RT (y)	5 (3, 9)	_			
Median IS free90 (59, 100)Patients are treated with throughout follow-upamount ofthroughout follow-upfollow-up (%)throughout follow-up			n immunosuppress	ion	

**TABLE 1** Patient characteristicscompared between sequential andconventional group

Patient characteristics compared between the tolerant and conventional groups. Standardized mean differences were computed to quantify the difference between the groups before and after matching. Variables noted by \* were used in the matching procedure. Data presented by mean (standard deviation), median (first quartile, third quartile), or absolute frequency (relative frequency).

their renal allograft within the first 3 months (Figure 1). Because this general patient population differs from the population of tolerant patients due to their medical history, we decided to match patients from each group according to immunologic and clinical key characteristics (Table 1). Caliper-matching criteria were chosen reflecting the previously described influence of those parameters on long-term transplant kidney survival and the availability of patient information common in the literature data and the OEDTR.<sup>26-28</sup> We aimed at a 1:1 matching because of the limited pool of potential matching candidates from the OEDTR once all criteria, especially optimal HLA match, and missing data were taken into account. Patients were selected from the OEDTR without replacement, such that each control patient occurred only once. Only if for a given sequential patient no other suitable matching partner was found in the registry data, we allowed that a single individual from the OEDTR acted as partner for more than one sequential patient. The rationale for this was the extraction of a subgroup from the OEDTR that is as close to the group of literature patients as possible while still retaining reasonable variability within the selected subgroup. Differences between the groups for several key characteristics are quantified as standardized mean differences

(SMDs), which express the difference of the means between the groups in units of the pooled standard deviation.

The main analyses presented here used a matching based on the variables that were available for each patient. By varying the size of the control group and allowing more matches for each tolerant patient and drawing patients with and without replacement from the OEDTR, we checked how sensitive the results from statistical analysis were to the choice of the control group. Furthermore, results were compared to analyses done on matchings based on patients with complete data only.

#### 2.5 | Statistical analysis

Metric variables are expressed as mean and standard deviation or median and range (interquartile or absolute). Categorical variables are expressed as absolute and relative frequencies. Transplant kidney and patient survival are depicted using Kaplan-Meier graphs. Due to the low number of events overall as well as crossing Kaplan-Meier curves, we used the Uno c-index to compare the matched tolerant and conventional group.<sup>29</sup> It can be interpreted as a conditional probability that for any pair of tolerant and conventional

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No.         State         Description         Description         State with a production of the state							Age (y)			
1MWistorth SyndromeMotionMotionBMotion </th <th>No.</th> <th>Sex</th> <th>Reason for HSCT</th> <th>Donor</th> <th>HLA mismatch</th> <th>Cause of ESRD</th> <th>At RT</th> <th>Donor</th> <th>IS after RT</th> <th>IS-free (y)</th>	No.	Sex	Reason for HSCT	Donor	HLA mismatch	Cause of ESRD	At RT	Donor	IS after RT	IS-free (y)
2     F     ALL     Brother     HL-identical     HO     23     25       3     F     AML     Sitter     Three mismatches     Raditon nephrits     14     NK       3     F     AML     Sitter     Three mismatches     BMT-NP     37     35       4     F     AML     Brother     HL-identical     BMT-NP     39     35       5     F     AML     Brother     HL-identical     BMT-NP     39     35       6     M     Norbus Holdpin     Brother     HL-identical     BMT-NP     39     35       17     M     Morbus Holdpin     Brother     HL-identical     BMT-NP     39     35       18     M     M     Morbus Holdpin     Brother     HL-identical     BMT-NP     30     35       19     M     Morbus Holdpin     Brother     HL-identical     BMT-NP     30     35       10     M     Morbus Holdpin     Brother     HL-identical     HL-identical     30     35       10     M     M     M     HL-identical     HL-identical     Morbus Holdpin     30     35       11     M     M     M     M     M     M     30     30	L.	Σ	Wiskott-Aldrich Syndrome	Mother	Two mismatches	Mesangial proliferative GN	18	42	Glucocorticoids, CyA	24
3         F         Application         Eather         Three mismatches         Radion nephritis         14         N           4         F         CML         Siter         Two mismatches         BMT-NP         37         35           5         F         AML         Brother         HA-identical         BMT-NP         37         35           6         AML         Brother         HA-identical         BMT-NP         37         35           1         M         Mothus Hodgin         Brother         HA-identical         BMT-NP         37         35           1         M         Mothus Hodgin         Brother         HA-identical         BMT-NP         37         35           1         M         Volume         Siter         HA-identical         Dishets type 1         37         35           1         M         Mothus         Mothus         Mothus         Mothus         37         35           1         M         Mothus         Mothus         Mothus         Mothus         37         35           1         M         Mothus         Mothus         Mothus         Mothus         36         36           1         M	Ν	ш	ALL	Brother	HLA-identical	HUS	28	25	None, later AZA and Glucocorticoids	0
4FCMLSisterWormsmatchesBMT-NP37355FAVLBruhenHU-vienticialHMT-NP40NK7MAVLBruhenHU-vienticialBMT-NP39337MMorbus HodykinBruhenBMT-NP39337MMorbus HodykinBruhenBMT-NP39339FAVLUnrelatedHU-vienticialBMT-NP393010MNorbus HodykinSisterHU-vienticialBMT-NP393011MCMLSisterHU-vienticialBMT-NP39303012FAULSisterHU-vienticialBMT-NP30303013MMorbusSisterHU-vienticialBMT-NP3030303014AULSisterHU-vienticialBMT-NP303030303015AULSisterHU-vienticialThe socialed3030303016MMorbusSisterHU-vienticialBMT-NP3030303016MMorbusSisterHU-vienticialBMT-NP3030303017MMorbusSisterHU-vienticialBMT-NP3040303018MMorbusSisterHU-vienticialBMT-NP30414119	ო	ш	Aplastic anemia	Father	Three mismatches	Radiation nephritis	14	ХX	Glucocorticoids, CyA	21
5         F         AML         Bother         HLA:dentical         HUS         40         80	4	ш	CML	Sister	Two mismatches	BMT-NP	37	35	None	19
6MAMLSisterHL-identicalBMT-NP39337MMorbus HodgkinBotherHL-identicalBMT-NP48518FAMLUnrelatedHL-identicalBMT-NP29NK9FCMLSisterHL-identicalBMT-NP29NK9NSisterHL-identicalDiabetes type 137NK10MOmMutherSisterHL-identicalNh Nowom645511MAMLSisterHL-identicalNh Nowom595712MAMLSisterHL-identical10205713MAMLSisterHL-identical10205714MAMLSisterHL-identical10205715MMutherSisterHL-identical10205716MMutherSisterHL-identical10205716MMutherSisterHL-identical10205716MMutherSisterHL-identical10202116MMutherMutherMutherMuther214617MutherMutherMutherMuther214616MutherMutherMutherMuther214616MutherMutherMutherMuther2146<	5	ш	AML	Brother	HLA-identical	HUS	40	ХX	Glucocorticoids and AZA	NK
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8       F       AML       Unrelated       BMT-NP       29       NK         9       F       CML       Sister       HLA:dentical       Diabetes type 1       37       NK         10       M       Vymphoma       Sister       HLA:dentical       Diabetes type 1       37       NK         11       M       Vymphoma       Sister       HLA:dentical       Unknown       64       56       57         12       M       AML       Muther       Sister       HLA:dentical       100       50       57         13       M       AML       Muther deficiency       Sister       HLA:dentical       100       50       57         14       M       Muther deficiency       Muther deficiency       Muther deficiency       Sister       HLA:dentical       100       50       57	7	Σ	Morbus Hodgkin	Brother	HLA-identical	BMT-NP	48	51	None	4
9       F       CML       Sister       HLA:dentical       Diabetes type 1       37       NK         10       M       CML       Sister       HLA:dentical       Unknown       64       55         11       M       Lymphona       Sibing       HLA:dentical       Unknown       64       55         12       M       AML       Father       Mother       One mismatches       CN1-associated       21       45         13       M       AML       Father       Wonforme       Sister       HLA:dentical       20       57       45         14       M       MIL       AML       Ether       Two mismatches       CN1-associated       21       45         15       M       Idiopatitic anemia       Sister       HLA:identical       Cnobi infection       23       41         16       M       Unrelated       HLA:identical       Cnobi infection       24       45         17       F       AML       Unrelated       HLA:identical       Combinication of aCMD       17       45         18       M       Unrelated       HLA:identical       Combinication of aCMD       24       46         19       M       HLA:identical<	00	ш	AML	Unrelated	HLA-identical	BMT-NP	29	NK	None	6
10MCMLSisterHLA:identicalUnknown645611MkymphomaSiblingHLA:identicalHUS/TTP596712FALLNomphomaSiblingHLA:identicalHUS/TTP296713MAMLFatherNommerchesCNI-associated214514MAMLSisterHLA:identicalITP282615MMulture-deficiencyNotherNotherLoss of first renal allograft292615MIdopathic aplastic anemiaSisterHLA:identicalLoss of first renal allograft292616MMulture-deficiencyNotherNotherLoss of first renal allograft292616MIdopathic aplastic anemiaSisterHLA:identicalComplication of aCMD172617FAMLUnrelatedInterationComplication of aCMD17214518MNoncryticaFatherTure mismatchesBMT-NP21264618MAULSisterThe mismatchesBMT-NP21454619MAULSisterThe mismatchesBMT-NP21454619MAULSisterThe mismatchesBMT-NP21454619MAULSisterThe mismatchesBMT-NP21464619MAUL	6	ш	CML	Sister	HLA-identical	Diabetes type 1	37	NK	None	17
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12       F       ALL       Mother       Onemismatches       BMT-NP       30       57         13       M       AML       Father       Two mismatches       CN1-associated       21       45         14       M       AML       Sister       HLAridentical       TTP       28       26         15       F       Comb immune-deficiency       Mother       Three mismatches       Loss of first renal allograft       22       41         16       M       Idiopathic aplastic anemia       Sister       HLAridentical       Chronic       28       26         17       F       Comb immune-deficiency       Mother       Three mismatches       Loss of first renal allograft       22       41         16       M       Unrelated       HLA-identical       Chronic       41       28       41         17       AML       Unrelated       HLA-identical       Chronic       41       42       44	11	Σ	Lymphoma	Sibling	HLA-identical	HUS/TTP	59	67	MMF, Glucocorticoids	5
13       M       AML       Eather       Two mismatches       CN-associated       21       45         14       M       AML       Sister       HLA-identical       TTP       28       26         15       F       Comb. immune-deficiency       Mother       Three mismatches       Loss of first renal allografi       22       61         16       M       Idiopathic aplastic anemia       Sister       HLA-identical       Loss of first renal allografi       23       61         17       F       AML       Unrelated       HLA-identical       Condication of aCvHD       17       43         18       M       Polymorphic       Father       Unrelated       HLA-identical       Condication of aCvHD       17       44         18       M       Polymorphic       Father       Iwo mismatches       BMT-NP       21       43         18       M       M       Bult       Rominations of anomic a	12	ш	ALL	Mother	One mismatch	BMT-NP	30	57	MMF, Glucocorticoids	5
14         ML         Siter         HLA-identical         TTP         28         26           15         F         Comb. immune-deficiency syndrome         Mother         Three mismatches         Loss of first renal allograft         22         61           16         M         Idiopathic aplastic anemia         Sister         HLA-identical         Chonic         41         52         61           17         F         ML         Unrelated         HLA-identical         Chonic         41         52         61           18         M         Polymorphic         Father         Unrelated         Complication of aGvHD         17         46           18         M         Polymorphic         Father         Two mismatches         BMT-NP         21         43           19         M         Aut         Att         24         44         24           10         M         Aut         Eather         Two mismatches         BMT-NP         21         46           10         M         Aut         Attentment         21         45         46           10         M         Aut         Attentment         21         46         46         46         46         <	13	Σ	AML	Father	Two mismatches	CNI-associated	21	45	Tac, MMF, Glucocorticoids	6
15     F     Comb. immune-deficiency syndrome     Mother     Three mismatches     Loss of first renal allograft     22     61       16     M     Idiopathic aplastic anemia     Sister     HLA-identical     Chronic     41     52     61       17     F     AML     Unrelated     HLA-identical     Chronic     61     12     46       18     M     Polymorphic     Father     Two mismatches     BMT-NP     21     43       19     M     Polymorphic     Father     Two mismatches     BMT-NP     21     43       19     M     Polymorphic     Father     Two mismatches     BMT-NP     21     43       10     M     ALL     Three mismatches     BMT-NP     21     43       20     M     B-cell non-Hodgkin lymphoma     Sister     HLA-identical     23     46       21     M     Aplastic anemia     Sister     HLA-identical     CNI toxicity     54     53       22     F     Sister     HLA-identical     CNI toxicity     54     54       23     F     Sister     HLA-identical     CNI toxicity     54     55       24     M     Ablastic anemia     Sister     HLA-identical     CNI toxicity <td>14</td> <td>Σ</td> <td>AML</td> <td>Sister</td> <td>HLA-identical</td> <td>TTP</td> <td>28</td> <td>26</td> <td>Tac, MMF</td> <td>ę</td>	14	Σ	AML	Sister	HLA-identical	TTP	28	26	Tac, MMF	ę
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17     F     AML     Unrelated     HL-identical     Complication of aGVHD     17     46       18     M     Polymorphic     Father     Two mismatches     BMT-NP     21     43       19     M     ALL     Father     Three mismatches     BMT-NP     25     46       20     M     B-cell non-Hodgkin lymphoma     Sister     HL-athentical     Acute and chronic     30     22       21     M     Aplastic anemia     Sister     HL-athentical     CN toxicity     54     63       22     F     Sister     HL-athentical     CN toxicity     54     55     55       23     F     Sister     HL-athentical     CN toxicity     54     55       23     F     Sister     HL-athentical     CN toxicity     54     55	16	Σ	Idiopathic aplastic anemia	Sister	HLA-identical	Chronic nephangiosclerosis	41	52	Glucocorticoids	6
18       M       Polymorphic       Father       Two mismatches       BMT-NP       21       43         19       M       ALL       Father       Three mismatches       BMT-NP       25       46         20       M       B-cell non-Hodgkin lymphoma       Sister       HLA-identical       Acute and chronic       30       22         20       M       B-cell non-Hodgkin lymphoma       Sister       HLA-identical       Acute and chronic       30       22         21       M       Aplastic anemia       Sister       HLA-identical       CNI toxicity       54       63         22       F       Sister       Mother       Three mismatches       Focal segmental       28       56         23       F       Sickle cell disease       Mother       Three mismatches       Focal segmental       28       56         24       F       Sickle cell disease       Mother       Three mismatches       Focal segmental       28       56         24       SiCD       F       Sickle cell disease       Mother       28       56       50         24       F       Sickle cell disease       Mother       Three mismatches       Focal segmental       28       56   <	17	ш	AML	Unrelated	HLA-identical	Complication of aGvHD treatment	17	46	Tac, Glucocorticoids	6
19     M     ALL     Father     Three mismatches     BMT-NP     25     46       20     M     B-cell non-Hodgkin lymphoma     Sister     HLA-identical     Acute and chronic     30     22       21     M     Aplastic anemia     Sister     HLA-identical     CNI toxicity     54     63       22     F     Sickle cell disease     Mother     Three mismatches     Focal segmental     28     56       23     F     Sickle cell disease     Mother     Three mismatches     Focal segmental     28     56       24     M     Sickle cell disease     Mother     Three mismatches     Focal segmental     28     56       25     F     Sickle cell disease     Mother     Three mismatches     Focal segmental     28     56       26     SCD     SCD     SCD     SCD     SCD     SCD     School Chronic ref	18	Σ	Polymorphic rhabdomyosarkoma	Father	Two mismatches	BMT-NP	21	43	Glucocorticoids	ო
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21     M     Aplastic anemia     Sister     HLA-identical     CNI toxicity     54     63       22     F     Sickle cell disease     Mother     Three mismatches     Focal segmental     28     56       22     F     Sickle cell disease     Mother     Three mismatches     Focal segmental     28     56       23     F     Sickle cell disease     Mother     Three mismatches     Focal segmental     28     56       8     SCD     SCD     SCD     SCD     SCD     50     56	20	Σ	B-cell non-Hodgkin lymphoma	Sister	HLA-identical	Acute and chronic rejection of 1. transplant kidney	30	22	Glucocorticoids	13
22     F Sickle cell disease     Mother     Three mismatches     Focal segmental     28     56       glomerulosclerosis due to       SCD       By some authors the cause of ESRD was described as bone marrow transplant nephropathy (BMT-NP). In the literature BMT-NP is defined as chronic rer	21	Σ	Aplastic anemia	Sister	HLA-identical	CNI toxicity	54	63	Single Dose Glucocorticoid	1,5
By some authors the cause of ESRD was described as bone marrow transplant nephropathy (BMT-NP). In the literature BMT-NP is defined as chronic rer	22	ш	Sickle cell disease	Mother	Three mismatches	Focal segmental glomerulosclerosis due to SCD	28	56	Glucocorticoid, MMF, Tacrolimus	0,2
includes chemo-irradiation-associated hephropathies caused by the conditioning therapies in the absences of other nephrotoxins.	By some includes	authors the chemo-irrae	e cause of ESRD was described as bo diation–associated nephropathies co	one marrow trans aused by the con	plant nephropathy (BMT- ditioning therapies in the	NP). In the literature BMT-NP i absences of other nephrotoxin	is defined a 18. <sup>14,40</sup>	is chronic renal	disease presenting within 100 da	ys after BMT. It

patients, the risk of an event (graft loss or death) is higher for the control patient. Longitudinal measurements of serum creatinine levels were investigated using a linear mixed model in which the group status is entered as main variable of interest. We used a random intercept and random slope to accommodate the patient-specific trajectories of the creatinine levels. Interaction terms with follow-up time, nonlinear effects of time using natural splines and different random effect structures were assessed by means of the Akaike information criterion (AIC) to improve model fit. Normality of residuals was checked to ensure validity of the model. Other variables such as recipient's age at RT, HLA-mismatch (set to 0 for HSCT patients), sex match between donor and recipient, and donor age were included in the model to assess whether differences in patient characteristics between the tolerant and conventional group were adequately controlled for by the matching procedure. All tests performed were

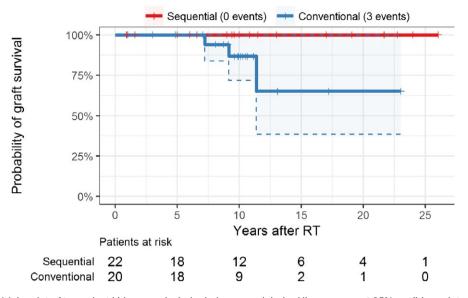
# two-sided and a P < .05 was considered statistically significant. Data management and analysis was performed by SAS software (SAS Institute Inc., Cary, USA) and the R statistical software (R Core Team, Vienna, Austria).

# 2.6 | Data retrieval and patient characteristics in sequential group

After examination of all search results, 18 relevant reports (published 1991-2017) were found, describing 29 patients with sequential renal and hematopoietic stem cell transplantation (Figure 1).<sup>5-21,24</sup> Authors of 15 reports sent follow-up data for a total of 18 patients. Despite intensive efforts, data from the other reports were not available because patients were lost to follow-up soon after the transplantation or publication. Four patients (N7, N8, N19, N21), not described in

No.	Follow-up (y)	Creatinine at last follow-up (µmol/l)	GVHD after RT	Death	Ref.	Note
1	26	133.0	Ν	Ν	(6)	
2	24	97.0	Ν	Ν	(12)	Diagnosis of radiation induced pulmonary fibrosis in 2000; lung transplantation in 2011
3	23	79.7	Ν	Ν	(13)	HCV cirrhosis (Child-Pugh A), Diabetes mellitus type 2
4	19	106.1	Ν	Ν	(14)	
5	13	NK	Ν	Υ	(14)	Died of unknown cause
6	10	106.2	Ν	Ν	(14)	
7	5	68.1	Ν	Υ	NP	Died of unknown cause, 11 y after RT
8	9	76,9	NK	Ν	NP	
9	17	80.0	Ν	Υ	(7)	Died of unknown cause, 20 y after RT
10	12	79.0	Ν	Y	(8)	Died 12 y after RT with functioning transplant kidney, cause of death presumed cardiac
11	10	131.0	Ν	Ν	(9)	
12	10	69.9	Ν	Ν	(9)	
13	9	81.0	NK	Ν	(15)	
14	6	97.2	Ν	Ν	(11)	
15	6	87.6	Ν	Ν	(16)	Second RT, successful pregnancy 4 y after second renal transplantation
16	10	80	Y	Ν	(17)	HCV cirrhosis (Child-Pugh A), manifestation of GVHD: cutaneous
17	11	48.0	Y	Ν	(18)	Manifestation of GVHD: cutaneous and musculoskeletal: scleroderma of lower and upper extremity, skin flare, avascular necrosis of knees, hips, shoulders, ankles; diagnosis of squamous cell carcinoma of the palate in 2016, currently in remission
18	4	99.1	Ν	Ν	(19)	
19	1	93.8	Ν	Ν	NP	
20	22	53.0	Y	Ν	(24)	Manifestation of GVHD: increase of liver function impairment (first seen after HSCT)
21	2	106.0	Ν	Ν	NP	
22	1	96.0	Υ	Ν	(21)	Manifestation of GVHD: oral mucosa (mild)

#### TABLE 3 Detailed clinical patient and follow-up findings



**FIGURE 2** Kaplan-Meier plot of transplant kidney survival; shaded areas and dashed lines represent 95% confidence intervals. No transplanted kidney was lost in the group of sequentially transplanted patients. RT, renal transplantation [Color figure can be viewed at wileyonlinelibrary.com]

the literature before, were included. Patient N19 was transplanted and followed-up at our clinic. The median observation period was 10 years (range 11 months to 26 years). Demographic, clinical, and transplant-associated parameters are summarized and compared to the conventional group in Table 1. A more detailed description of all tolerant patients is shown in Tables 2 and 3. Except for one patient, all indications for HSCT were of hematologic origin. Two patients received their renal transplants from unrelated but fully HLA-matched donors. Compared to the conventional group, the tolerant patients did not receive any immunosuppressive treatment during >90% of the whole follow-up period.

### 2.7 | Matching of the OEDTR control group

All 22 patients could be included into the matched comparison. We identified 707 patients from the OEDTR who met our inclusion criteria. Twenty patients were selected as matching partners for the tolerant group (Table 1). Causes of ESRD in these patients were vascular nephropathy (N = 1), glomerulonephritis (N = 5), and 14 patients were labeled as "other" cause of ESRD. Four patients had type 2 diabetes, and 2 of them developed diabetes after transplantation. The matching procedure reduced almost all SMDs into acceptable ranges. The SMD in the HLA mismatches remained high even after matching, which is due to the lack of mismatches in the group of sequentially transplanted patients.

#### 2.8 | Maintenance immunosuppression

Withdrawal of immunosuppression in the tolerant group: Seven patients did not receive any immunosuppression after renal transplantation. All other patients received at least short-term immunosuppression initially after RT (see Table 2), which was successfully withdrawn in the later course. At the time of the last follow-up, 20 of 21 patients (records not available in one patient) were free of immunosuppression. Patient N2 successful withdrew IS after RT but developed a radiation-induced pulmonary fibrosis in the later course and IS including steroids and azathioprine had to be reinitiated. In 2011, a successful lung transplantation was performed and tacrolimus was added. Maintenance IS in the control group was given as per the local center standards, that is, calcineurin inhibitor (CNI), mycophenolate mofetil (MMF), and steroids in all patients aiming at tacrolimus (TAC) troughs between 4 and 8 ng/ml after the first years.

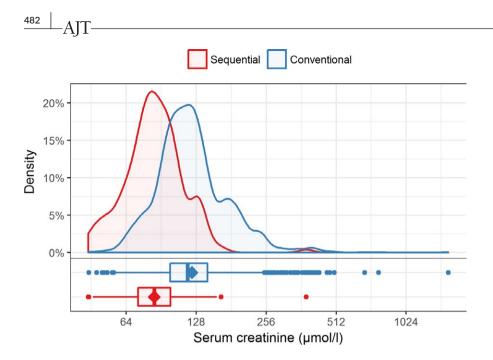
### 3 | RESULTS

# 3.1 | Overall renal allograft survival and graft function in sequentially transplanted patients compared to conventional transplanted patients

No graft loss was reported in the tolerant group, whereas 3 (15%) were lost in the control group (Kaplan-Meier analysis depicted in Figure 2). The corresponding Uno c-index comparing the 2 groups was 0.81, suggesting an improved renal graft survival in the tolerant group within the study period. The 95% confidence interval (0.67-0.96) excluded parity (0.5), providing evidence against the hypothesis that the renal allograft survival probabilities are equal in both groups (P < .001).

Serum creatinine levels over the whole study period are depicted in Figure 3, median serum creatinine levels were 85  $\mu$ mol/l (first quartile 72, third quartile 99) and 118  $\mu$ mol/l (IQR 99-143) for the tolerant and conventional group, respectively. In a crude comparison ignoring the longitudinal nature of the data, the median level in the conventional group was significantly higher than in the tolerant group (Mann-Whitney *U* test *P* < .001).

Figure 4 shows results from the mixed-model analysis of log2transformed serum creatinine levels, taking the study design into



**FIGURE 3** Comparison of serum creatinine levels (on log2 scaled axis) after RT between groups; the mean is indicated as diamond shape in the boxplots [Color figure can be viewed at wileyonlinelibrary. com]

account. No relevant difference in the change of creatinine levels over time between the 2 groups was found (interaction with follow-up time was not included in the models, either by checking significance or by AIC). The effect of time was found to be nonlinear (a model including spline terms of time using 3 knots showed improved AIC), with a decrease in creatinine levels shortly after transplantation. Group status was significant in the model (P = .006), providing evidence that serum creatinine levels in the tolerant group are on average 29% lower than in the conventional group. Estimated effects of group status remained stable regardless of modeling of time. After inclusion of other relevant key characteristics, group status remained significant (P = .045), whereas none of the other variables in the model had significant results. In an unmatched analysis, comparing all patients from the OEDTR with available creatinine measurements (N = 273) and the study group, the OEDTR patients exhibited a trend toward higher creatinine values over time (Figure S1).

In a subset of patients further immunologic assays have been performed. Both patients had no signs of humoral immune activation (measured as donor- and recipient-specific antibodies). Moreover, protocol biopsies performed one and 12 months after RT revealed no signs of allograft rejection. Notably, recipient-derived T cells measured before RT and 1 year after remained virtually unchanged (Figure 5).

# 3.2 | Overall patient survival in sequentially transplanted patients compared to conventionally transplanted patients

Of 22 patients, 4 deaths were reported. Patient N5 was lost to follow-up 13 years after RT and died in the later course. The exact time of death is not recorded and the patient is therefore treated as censored at the last follow-up visit. Patient N7 was last seen 5 years after transplantation at the treating medical center and died 11 years after renal transplantation. Patient N9 was last seen in 2012, 17

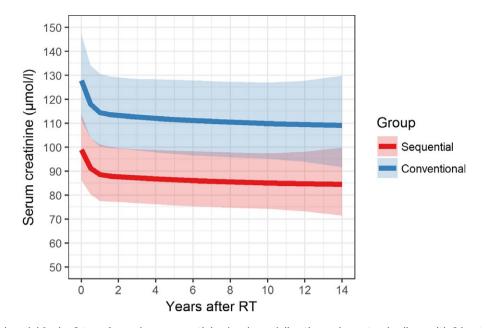
years after renal transplantation. At that time, her creatinine was in normal range. Later, no more visits to the outpatient clinic were recorded. She died in 2015. All 3 patients died of unknown causes. Patient N10 died in 2014, 12 years after RT. His presumed cause of death was of cardiac origin. Notably, the patient died with a functioning transplant kidney.

Two deaths were recorded in the matched subset of conventionally transplanted patients from the OEDTR. The Uno c-index for the corresponding Kaplan-Meier survival curves comparing the 2 groups was 0.39, suggesting a lower mortality in the conventional group within the study period (Figure 6). However, due to the low number of events, the 95% confidence interval was wide (0.08-0.71) and included parity (0.5), thus lacking evidence against the hypothesis that the overall survival probabilities are equal in both groups (P = .54).

# 3.3 | Further notable events in the patients' later course

After HSCT, patient N15 received a renal allograft from a fully matched deceased donor. Eight years later, the allograft was lost due to infection and a second renal transplantation was performed, in which the previous HSCT donor (mother) donated a kidney. Withdrawal of all immunosuppressive drugs was successful 10 weeks later. Four years later, the patient gave birth to a child without rejection. The patient developed HLA antibodies but they were not targeted against the allograft. Restart of immunosuppressive therapy was not needed in the further course.

Follow-up data (exceeding the published report) regarding graft function or patient survival from one patient described by Beitinjaneh et al were not available.<sup>20</sup> Nevertheless, the treating center reported 3 further patients not described before, who received sequential HSCT and RT. The HSCTs were performed between 1982 and 1986 (indications: acute myeloid leukemia [AML], aplastic anemia, chronic myeloid leukemia [CML]). Renal allografts were transplanted 6 to



**FIGURE 4** Mixed model for log2-transformed serum creatinine levels modeling time using natural splines with 3 knots. Estimates were transformed to original scale. The shaded areas around the model estimates constitute 95% confidence intervals. The overall difference between the 2 groups was significant. No relevant difference in the change of creatinine levels over time between the 2 groups was found. Four patients from the control group had to be excluded due to missing longitudinal serum creatinine measurements. We restricted the analysis to the first 14 years after RT due to lack of serum creatinine measurements for most OEDTR patients thereafter. RT, renal transplantation [Color figure can be viewed at wileyonlinelibrary.com]

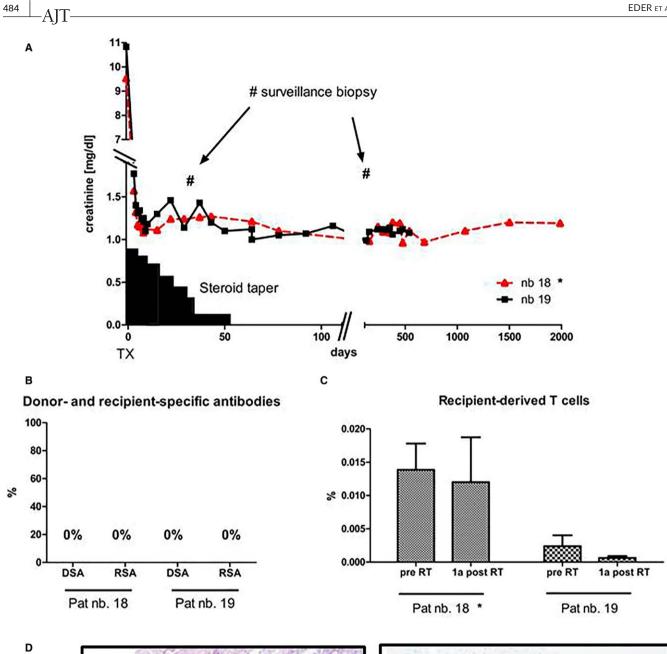
20 years later (median 16 years). All three patients died (33, 14, and 26 years after HSCT). Due to a lack of further patient data (donor age, course of creatinine, HLA mismatch, immunosuppressive regimens), we did not include these patients in our analysis.

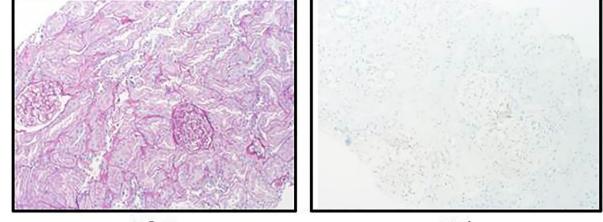
### 4 | DISCUSSION

The induction of donor-specific tolerance remains the holy grail in transplantation. Currently, the only clinically successful concept to establish donor-specific tolerance to a renal graft is via HSCT from the same donor. However, widespread use is hampered mainly by cytotoxic preconditioning before bone marrow transplantation. Thus, few patients have been reported in the literature that have received combined HSCT and RT from the same donor solely for the purpose of tolerance induction. However, several patients have been reported who had received an HSCT for a conventional indication (eg, a hematologic malignancy) and subsequently received a kidney from the same donor. The aim of our study was to contribute long-term data of such tolerant patients and to compare those findings with matched living donor transplant recipients under conventional IS.

Compared to previous studies, our analysis represents the largest and longest follow-up observation of this specific patient population and includes additional patients not described in the literature before. One of our major findings is that over a total observation period of 250 patient-years not one single allograft loss was recorded. Moreover, kidney function was significantly better compared to our matched group of conventionally living donor recipients. There are several reasons that might attribute to these results. First, in addition to receiving initial IS immediately after RT, most of the tolerant patients were successfully withdrawn from immunosuppression, thus adverse side effects and nephrotoxicity caused by conventional IS regimens were largely avoided.<sup>30</sup> Moreover, in tolerant patients without the need of ongoing immunosuppressive therapy, adherence, which is a main risk factor for chronic allograft loss, is of no concern and the likelihood of infections might be lower.<sup>31,32</sup> Furthermore it must be addressed, that this highly selected subgroup of the OEDTR represents patients with comparably promising clinical parameters (young age, low HLA mismatch) and hence may not be representative of the general population of kidney transplant recipients. On the other hand, it must also be pointed out that 4 patients from the study group died in the further follow-up period. Nevertheless, death censoring was applied within our analysis, considering that our primary aim was to evaluate the impact of tolerance on renal graft survival and taking the previous mostly severe hemato-oncologic diseases in this special group of patients into account.

Despite significantly higher serum creatinine levels in the control group and 3 renal graft losses in the follow-up, no relevant difference in the change of creatinine levels over time was found. The reason for this may also be the selection of the control group, representing patients with a very good outlook after transplantation. In an unmatched analysis, comparing all OEDTR patients with the sequential patients, the OEDTR patients exhibit a trend toward higher creatinine values over time. When interpreting the creatinine slopes





H&E

C4d

**FIGURE 5** (A) Clinical course of the 2 tolerant patients transplanted at our center (pat. N18 and N19). Both patients underwent transplantation with only a short steroid taper to reduce ischemia-reperfusion injury. No further immunosuppression was administered. (B) Humoral immune activation was assessed by measuring donor- and recipient-specific antibodies. There were no signs of humoral immune activation. (C) Chimerism was assessed via real-time PCR in separated T cells before and 1 year after RT in both patients. The proportion of recipient-derived T cells was low and virtually didn't change over the time. (D) Protocol biopsies performed 1 month and 12 months after RT showed structurally regular renal tissue with no signs of cell- or antibody-mediated rejection. Data from patient N18 have been shown previously (Schwarz 2016) with a shorter follow-up<sup>19</sup> [Color figure can be viewed at wileyonlinelibrary.com]

of the control patients, one may further conclude that, in line with recent findings from Gaston et al, those graft losses could be associated with new-onset allograft dysfunction in the later course, rather than a slow process of continuous worsening of allograft function.<sup>33</sup> Unfortunately, clinical data from the control group were limited and more detailed information regarding the cause of the renal graft losses were not available.

So far, 3 centers in the United States reported results from their prospective trials of simultaneous HSCT and RT where IS was successfully withdrawn in a considerable number of patients.<sup>34-37</sup> Nevertheless, complications such as in part severe bacterial, viral, or fungal infections were described.<sup>35,36</sup> Due to these risks, the use of combined transplantations solely for the purpose of tolerance induction, is subject to considerable controversy, especially in patients who may do well on conventional immunosuppression. Our data on the other hand indicate a sustained long term benefit regarding transplant kidney function in patients after tolerance induction via hematopoietic stem cell transplantation from the same donor.

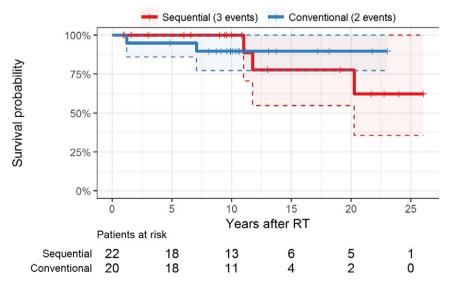
About two-thirds of the patients in our analysis received at least short-term immunosuppressive therapy after RT, which was, except for one patient (N2, lung transplantation), successfully withdrawn in the later course. Notably, none of the tolerant patients was taking long-term corticosteroid treatment.

Most patients received IS for the prevention of a potential GVHD episode. Manifestations of a GVHD were reported in 4 patients (N16, N17, N20, N22). They included skin and musculoskeletal

lesions, increased liver enzymes, or mild restrictive lung disease. Thus, considering the risk of even at least temporary IS, we think there is enough evidence to support minimalization of IS after kidney transplantation in GVHD-free and tolerant patients. Notably, a steroid taper around RT seems to be beneficial, mainly to reduce ischemia-reperfusion injury which is a well-known cause for initialtransplant kidney dysfunction.<sup>38,39</sup>

There are some limitations in our study that need to be addressed when interpreting the results. First, a publication bias cannot be ruled out. It might be that unsuccessful cases were not reported in the literature and therefore not included in this study. However, this seems to be unlikely, given the novelty of this hypothetical issue, the rarity of an available donor for both HSCT and RT when needed, and the limited number of centers with the infrastructure and experience to perform sequential transplantations.

Despite frequent communications with the authors or their successors, we were not able to gather outcome data of all 29 published patients. We tried to minimize this limitation by including further patients that have been sequentially transplanted but have not been published in the literature. Furthermore, due to the retrospective design of our study and the long period between the publication of the first cases and our analysis, the quantity of collectable data is limited and varied strongly between different patients. We therefore focused our analysis on basic parameters such as renal transplant function and patient survival. Unfortunately, this also implies that tests for donor specific antibodies (DSAs) or chimerism persistence are missing. Hence, we added supplemental material describing the



**FIGURE 6** Kaplan-Meier plot of overall patient survival; shaded areas and dashed lines represent 95% confidence intervals. Overall there were 4 deaths in the sequential group but one was treated as censored at the last follow-up visit as the exact time of death is unknown. RT, renal transplantation [Color figure can be viewed at wileyonlinelibrary.com] lack of DSAs, good transplant kidney function, as well as a virtually unchanged proportion of recipient-derived T cells in the 2 sequentially transplanted patients from our center.

Furthermore, even if publication bias is ignored, the statistical power of this study to accurately estimate differences between tolerant patients and conventionally transplanted patients is limited due to the relatively low number of available patients.

In conclusion, our data with greater numbers and lengthier follow-up than previously available indicate that tolerant patients after HSCT maintain a better function of their transplant kidney than comparable patients under conventional immunosuppression. Furthermore, overall renal graft and patient survival was at least comparable, and allowing for limitations of the analysis, perhaps superior to a matched living donor cohort. These data lend support to efforts currently underway to utilize co-transplantation of donor hematopoietic stem cells to improve outcome in larger numbers of renal transplant patients.

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#### DISCLOSURE

The authors of this manuscript have no conflicts of interest to disclose as described by the *American Journal of Transplantation*.

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#### SUPPORTING INFORMATION

Additional supporting information may be found online in the Supporting Information section at the end of the article.

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