Australian and New Zealand Paired Kidney Exchange Program

Protocol 4: ANZKX Tissue Typing Laboratory Guidelines



ANZKX Tissue Typing Laboratory (TTL) Guidelines

This document outlines the Tissue Typing Laboratory requirements to enable implementation of the ANZKX Program.

Acronyms referenced in these Guidelines

ATTC = ANZKX Tissue Typing Coordinator CP = Compatible pairs DSA = Donor Specific Antibody MFI = Mean Fluorescence Intensity OLI = One lambda Inc. (antibody assay) OM = OrganMatch SAB = Single Antigen Bead TTL = Tissue Typing Laboratory TWL = Transplant Waiting List Tepnel = Lifecodes Immucor Inc. (antibody assay) UA = Unacceptable Antigens

Introduction

The ANZKX Program uses an algorithm to find matches among the pool of donor-recipient pairs to create 2 way or larger exchanges, or non-directed donor chains of transplants.

First, matched pairs are found by comparing the ABO blood groups of each donor and each recipient, then by checking the Unacceptable Antigens (UA) of recipients and comparing these in turn with the HLA-typing of each ABO compatible donor.

UA include primarily HLA alleles to which a recipient has an antibody. They may also include mismatched antigens from previous transplants or pregnancies or those associated with a high eplet load as per each transplant unit's requirements.

These guidelines outline the requirements and processes necessary to achieve the goal of the ANZKX Program. The TTL are encouraged to complete testing requirements promptly to enable pairs who are ready to be matched to participate in the ANZKX program. This is due to the change to continuous matching throughout the year instead of having match runs as done previously.

1 HLA typing requirements for donors and recipients

1.1 Registration requirements for donors and recipients

Registration for donors and recipients is managed by their enrolments in Organ Match. It is important to create their OrganMatch enrolments as soon as the donor-recipient pair have been deemed suitable for the program, and document when the pair is likely to be ready for matching via notes in the organ match records.



1.2 HLA typing requirements for donors

For donors all loci are required to be at 2 field level. For entry onto the ANZKX register, donors must have an authorised HLA typing recorded into OrganMatch (OM) for each of the following mandatory HLA loci:

HLA-A*, HLA-B*, HLA-C*, HLA-DRB1*, HLA-DPB1*, HLA-DPA1*, HLA-DQA1*, HLA-DQB1* and HLA-DRB3/4/5*.

1.2.1 Overseas Donors

Australian or New Zealand recipients are able to be co-registered with a donor who resides overseas if the following conditions are met:

- The donor is reviewed by the Australian or New Zealand unit prior to registration.
- HLA typing is performed in Australia or New Zealand to ANZKX specifications.
- Adequate blood is obtained at time of sampling to enable freezing of cells for later use e.g. crossmatch.
- TTL to verify that a sufficient number of cells are isolated from this sample to enable multiple FLOW crossmatches to be performed. If necessary recall donor for additional specimen for cells to be frozen by the TTL in case the donor returns to their home country and is not available for crossmatching samples to be collected.
- The donor is able to accommodate any potential surgery timeline.

1.3 HLA typing requirements for recipients

For recipients (new and existing) all loci typing at 2 field level is required. For entry onto the ANZKX register, patients must have an authorised HLA typing recorded into OrganMatch for each of the mandatory HLA loci:

HLA-A*, HLA-B*, HLA-C*, HLA-DRB1*, HLA-DPB1*, HLA-DPA1*, HLA-DQA1*, HLA-DQB1* and HLA-DRB3/4/5*.

Full typing of recipients is required for the purposes of epitope matching and to exclude the possibility of self-reactive antibodies, particularly for DP locus.

1.4 Donors that express an HLA allele not included in one of the single antigen bead assays

Currently Organ Match is unable to exclude from matching recipients with donor-specific antibody (DSA) against a donor expressing a 2 field HLA allele that is not covered by the Single Antigen Bead (SAB) assays. For this reason if a donor expresses a 2 field HLA allele that is not covered by one of the SAB assays, special management procedures are required (e.g. C*07:01 is not currently included in the One Lambda Inc. (OLI) kit).

The local TTL will be responsible for entering the following comment into the OrganMatch donor record for each donor expressing an HLA allele that is not covered by the SAB assay.

"Following alleles are not represented in the current OLI/Tepnel SAB lot: (e.g.) B*35:02, DRB1*08:03".



1.5 HLA nomenclature

Whenever possible the 2 field molecular nomenclature (describing a specific allele of the antigen) will be used for HLA antigens and HLA-antibodies used in the OrganMatch database for the purpose of the ANZKX Program.

In exceptional circumstances information may only be available based on historical data obtained by serological typing. This could be the case if a patient had a previous transplant and HLA specificities for the previous donor are only available at the 1 field resolution to describe an antigen. In rare instances a patient may have a record of a historical HLA antibody demonstrated by serological methods, but the historical serum is no longer available to determine 2 field specificity by Luminex SAB (see 5.1). In this instance the antigens based on the specificity of the serological antibody should be listed in the UA table in OM (see 3.1).

As the 1 field antigens will generally be identical to the first field of the 2 field alleles, the few exceptions to this rule will need to be listed in the OrganMatch HLA antigen/allele relationships table exclusively using 1 field molecular nomenclature. All the required antigen and allele code updates must be validated by the TTL.

Examples:	2 field molecular	1 field molecular	Serological	
	A*11:01	A*11	A11	
	C*03:04	C*03	Cw10	
	DRB1*03:01	DRB1*03	DR17	

2 HLA antibody screening of patients

For entry into the ANZKX program, patients must have an authorised antibody record tested by one of the Luminex Single Antigen Bead test providers (OLI or Life Technologies) for both Class I and Class II HLA antibodies.

Confirmatory SAB testing must be performed using a sample collected at least one month earlier or later than the primary sample. EDTA serum treatment is recommended for OLI SAB testing.

Supplementary bead sets are available if using the one lambda platform and should be used for testing at least one sample when there are Class I and or 2 antibodies detected using the standard bead sets.

Individual antibody strengths expressed as mean fluorescence intensity (MFI) must be entered into the record for all Luminex-detected Class I and Class II HLA antibodies.

Authorised antibodies to be used in defining UA are assigned by each local Australian or New Zealand TTL in consultation with their clinicians. These should, in general, be the level which is likely to give a positive crossmatch, ie >2000MFI for OLI and >1500MFI Tepnel or the level that is acceptable to the Transplant unit of the patient.

Antibody results at the 2 field level are required to be stored in OrganMatch along with MFI values for all excluded and non-excluded antibodies detected by SAB assay.

2.1 Testing frequency

For patients with ongoing registration in ANZKX single antigen bead antibody testing should be repeated:

- every 6 months
- after any sensitisation event
- if matched in a chain and not tested within the previous 3 months
- and again within 1 month of the booked transplant date



2.2 HLA antibody strength

Published data indicates that in the presence of DSA, values of <2000MFI (by OLI kit) are unlikely to yield a positive **CDC-crossmatch**, whereas >8000MFI are extremely likely to have a positive CDC crossmatch. The cut-off for a positive **flow cross match** is somewhere around 1000 – 2000MFI and certainly a value below this is unlikely to have a positive flow cross match.

A review of the Asia Pacific Histocompatibility and Immunogenetics Association (APHIA) QC data has suggested that in general Tepnel has lower MFI and that a value of 1500MFI is the equivalent of a 2000MFI for OLI and using such cut -offs will result in very similar calculated panel reactive antibody (cPRA) amongst all TTL.

When listing antibody sourced UA, the following is suggested to determine was is an acceptable match:

Cut-offs for One Lambda Inc. (OLI) Luminex assay are as follows:

To be entered in OrganMatch by default as unauthorised, i.e. do not exclude from matching.

• Weak 500 - 2000

To be entered in OrganMatch by default as authorised, i.e. will exclude from matching.

• Moderate 2000 to 8000 or Strong >8000

Cut-offs for Lifecodes Immucor Inc. ('Tepnel') Luminex test are as follows:

To be entered in OrganMatch by default as unauthorised, i.e. do not exclude from matching.

• Weak 300 - 1500

To be entered in OrganMatch by default as authorised, i.e. will exclude from matching.

• Moderate 1500 to 4000 or Strong >4000

Occasionally matches may be offered where there may be a DSA outside of the above parameters where a higher UA antibody threshold has been used, i.e. highly sensitised patients to allow more potential matches. Prior discussion with the clinical unit has usually occurred to ensure this is an acceptable offer prior to a chain being created (see 6.3).

3 Defining UA

3.1 Serum selection for use in antibody sourced UA

Evidence suggests that the current serum may underestimate true sensitisation and post-transplant immunological risk. The <u>default</u> serum authorised for match run is the patient's 'peak serum', unless otherwise advised by the recipient's clinical unit.

It is acknowledged that the amount of one antibody may be high in one serum sample and another antibody may be higher in a different serum sample. In Organ Match an Antibody consolidation can be used to select all relevant antibody results.

Ultimately, the selected panel of antibodies authorised as UA for the match run is at the discretion of the patient physician in combination with the local Australian or New Zealand TTL.

See section 2.2 for suggested cut-offs for antibody sourced UA depending on the testing technology used.

3.2 Exclude previous Mismatches

Previous transplant data must be entered into the OrganMatch database.

Previously transplanted mismatched antigens will be considered as unacceptable in the organ matching process unless otherwise agreed between clinicians and the local TTL.



The HLA typing of a previous donor must be promoted to the HLA Typing Profile and authorised to enable the mismatched antigens to be visible in the UA table of the recipient. If in a serological HLA typing format, a molecular conversion can be performed for use in the donor HLA typing profile and the recipient UA. Where possible this must be in a minimum of 1 field but ideally 2 field molecular typing format. From this table they can then be selected and authorised as unacceptable mismatches for matching.

If a previous transplant donor record does not exist in OM, the record should be created and a historical transplant registered. If retrospective transplant registration of historical transplants is not possible in OM, the previous donor mismatches will need to be entered into the recipient UA table using the "add antigen" button.

Type= Unacceptable Antigen. Antigen = 1 or 2 field molecular results entered as a string separated by comma and space.

E.g. Historic typing = A2, 24; B62, B60, Cw10, Cw9 would be entered as a string

A*02, A*24, B*15:01, B*40:01, C*03:03, C*03:04

Reason: "previous transplant mismatch"

Clinicians, following discussion with the TTL, may deem it safe to remove a previously transplanted mismatched antigen from the authorised mismatches (because of current and historical absence of a specific antibody to the previous mismatch).

3.3 Additional UA

Clinicians, in consultation with their TTL, may decide to enter any other unwanted antigens into UA list.

Examples for this strategy are:

- incompatible spousal live donor pairs, where the donor is the husband/male partner and one or more specific antigens of the husband need to be excluded (for instance because of presence of DSA <2000MFI to husband antigens).
- Exclusion of antigens where the predicted eplet load would be very high in case of a match, even if no antibodies to that antigen are detectable

These additional antigens are categorised as Other UA in OrganMatch and will need to be entered into the Recipient UA Table using the "add antigen" button.

Type= unacceptable Antigen. Antigen = 1 or 2 field molecular results entered as a string separated by comma and space.

A*02:01, DRB4*01, DQB1*03:01

Reason: "high eplet load" or "spouse mismatches".



4 Willing to accept ABO-incompatible transplant

An additional strategy to increase options for sensitised patients is to accept kidneys from ABO-incompatible, but HLA acceptable donors within the ANZKX pool. The acceptance of ABO-incompatible donors for some of the highly sensitised recipients in the ANZKX Program results in a "virtual" expansion of the donor pool

The TTL is required to enter acceptance of ABOi in the 'Willing to accept' tab of the KPD enrolment and also indicate which ABOi blood groups they are willing to accept and also enter a consent date. It is possible to select only A2 or both A1 and A2. OrganMatch requires the consent date to be entered.

Willing to accept					
Accept Hep B Core Yes <u>No</u>	Date of Consent dd/mm/yyyy				
Accept ABOi	Date of Consent * dd/mm/yyyy	ABO Groups	ABO Groups		
		A	AB	В	0
		A1 A2	A1B A2B		
		Plance calent an ARO			

The referring transplant unit is responsible to determine whether acceptance of an ABO-incompatible donor is a feasible option for their recipients and to enter this information in the KPD enrolment in OrganMatch.

5 Rules for inclusion of compatible pairs in the ANZKX Program

Participation of compatible pairs (CP) in ANZKX can be attractive to CP who have a high degree of HLAmismatch, if the KPD allocation algorithm provides a better HLA match for the CP recipient.

Because the ANZKX program is not designed to help CP, it is important to define allocation metrics that enable the CP to receive a better-matched kidney, without disadvantage to incompatible pairs (ICP). The following pathway for inclusion of compatible pairs in ANZKX was approved by RTAC.

<u>CP can be included in ANZKX in order to gain a better HLA-matched donor, provided the HLA mismatch to</u> the own donor has a high level of eplet mismatch (EpMM) (for example combined class I and II Eplet mismatch more than approximately 65).

- The majority of CP mismatched at HLA-A, B, DR will meet these criteria.
- The majority of these CP will be able to find a match in a relatively short period of time dependant on the blood group of the donor being entered. The ANZKX team will regularly liaise with the recipient transplanting unit regarding the likelihood of the pairs matchability.
- CP who do not meet the 65 EpMM threshold can also be considered, if they wish to avoid a specific mismatch that would impact a future chance of a repeat transplant.

Better immunological matching can be achieved by excluding high eplet mismatched antigens in the UA profile of the CP recipient (see 3.3).

- The list of UA should translate into a virtual cPRA of approximately 70-80%:
- Assigning a virtual cPRA up to 80% will:
 - not excessively affect CP match probability;
 - not disadvantage ICP in favour of CP recipients;
 - not reduce match probability for highly sensitised ICP.
- The virtual cPRA of 70-80% may be adjusted if no matches are found.



6 Method of matching by OrganMatch for ANZKX

The OrganMatch PKE program will match:

- 1 recipients UA against
- 2 donors HLA antigens

Whilst aiming to:

- maximise the number of suitable donor-recipient pairs
- maximise number of blood group ABO identical pairs (ABO identical > ABO compatible)

The Match Probability (MP) is an indicator of how easy or difficult it will be for a pair to be matched in a chain.

The Match Probability (MP) will be calculated as: **MP** = (**a**/**b**), where:

a = the number of acceptable donors in the run (i.e. donors having no HLA antigens to be excluded because of patient DSA or previously transplanted antigens).

b = total number of ABO compatible/acceptable donors in the run.

MP range is 0 to 1: 0 implies no compatible donors and 1 implies all ABO compatible donors are suitable matches.

6.1 Method of matching recipient UA against donor HLA

The default method of matching will be to match 2 field patient UA to 2 field donor HLA:

- A 2 field patient UA specificity will result in exclusion of any donor with that precise allele authorised in the HLA typing. However donors with any authorised other allele of the antigen will not be excluded.
- These will be picked up in the match review process and the match blocked if necessary.
- Recipient UA can be antibody sourced, previous donor mismatches or other as described in section 3.

6.2 Individualised assignment of unacceptable HLA antigens

In the ANZKX Program a high proportion of the incompatible pairs are enrolled as a result of HLAincompatibility. Some highly-sensitised recipients with broad sensitisation have only a limited number of rare donor HLA genotypes that they can be matched with and this leads to the accumulations of patients with broad sensitization and high antibody strength against common antigens in the ANZKX pool. Alternative allocation strategies are required to assist these patients and one possible solution is the individualised assignment of unacceptable HLA antigens.

Therefore, there may be instances where a transplant team decides to remove some HLA antibody specificities with >2000 MFI from the authorised unacceptable antigen list for a particular patient to increase the chance of them being matched. This means that donors with HLA alleles to which a particular recipient has donor-specific antibody (DSA) with a reactivity of >2000 MFI will not be excluded from matching with this recipient.

For these cases a recipient evaluation form outlining HLA antibody to be ignored (see appendix) should be made available to the ATTC and ANZKX Coordination Centre.

If DSA >2000 MFI are present in non-authorized sera the patient's physician will be notified after the potential match is identified, before a chain is offered for consideration to all centres involved.

The individualised assignment of unacceptable mismatches should consider the sensitisation status of the patient:



- Patients with a cPRA <75% are <u>very likely</u> to find a compatible donor in the ANZKX pool with no DSA to the matched donor.
- Patients with cPRA >75% (highly-sensitised) and in particular those with cPRA >95% are <u>less likely</u> to find a compatible donor in the ANZKX pool with no DSA to the matched donor.

Individualised assignment of UA should be reserved for highly-sensitised recipients.

The selection of the best available compatible donor with the longest projected graft lifespan should be considered for those likely to require re-transplantation. Therefore, it is also possible to consider excluding 'acceptable' mismatches (where there would be no DSA to a specific donor HLA antigen) for patients with minimal sensitisation, in particular paediatric patients and young adults. In this instance one or more specific alleles can be added to the UA list as "others".

7 Further testing after a match run

7.1 Flow cytometry crossmatching

Flow crossmatching is performed routinely for ANZKX matched pairs.

The Flow crossmatch is to be performed by the Tissue Typing lab of the donor state. The ATTC will coordinate Flow Crossmatching by advising the state tissue typing coordinators where to send recipient serum samples for testing and which donor crossmatches are required.

Flow Crossmatch results are to be entered into OM and verified by the State TTL. The ATTC should be notified when the results have been verified. Once all of the results have been entered into OM for all of the match pairs in a particular chain the ATTC will generate the Match Event reports and advise the national directors they are ready for reporting to the transplant units of the match pairs involved.

7.2 Allocation of a non-directed donor chain kidney (last in chain) to the transplant waitlist recipient

When a non-directed donor (NDAD, also known as altruistic donor) is referred to ANZKX and starts a chain of transplants, the last recipient in the chain will have a living donor who has not donated during the match cycle. This donor will donate to the deceased donor list (and close the chain) of either New Zealand or the relevant Australian state that entered the NDAD donor to ANZKX. Allocation of the last donor in the chain to the state-of-origin waitlist will follow the standard OrganMatch allocation rules for deceased donors in Australia, or NZKAS rules in New Zealand.

The timing of the allocation will be as close as possible to the scheduled date of surgery planned for the NDAD chain. However, the timing must take into consideration the need for crossmatching to be performed and reported to the recipient's team.

The unit performing the transplant of the waitlist recipient will also need sufficient time to assess/ review intended recipient medical suitability and for the laboratory to repeat a Luminex SAB test on a current serum, as there could be patients on the Transplant Waiting List (TWL) who have screening results up to 12 months previous as these are generally reviewed annually unless the labs have been notified of a sensitising event.

It is therefore suggested that allocation should take place no later than 3 weeks prior to the scheduled date for the chain surgeries. Possibly a shortlist of 3 possible recipients should be made in case a contraindication comes to light clinically or immunologically within that 3 weeks' time frame. The TWL enrolment of the recipient, once identified, must be put "on hold" pending transplant.

7.3 HLA antibody testing after matching and prior to transplantation

Following a successful match ideally transplantation should occur as quickly as possible, preferably within 4 to 6 weeks. In the event of surgical delay HLA antibody testing should be performed within 4 weeks prior to the scheduled surgical date. If the SAB profile has not altered significantly then a repeat FLOW crossmatch



is not required. If a repeat FLOW crossmatch is required due to an altered SAB profile which would influence the decision to proceed or not, then it should be performed in consultation with the transplant unit.

8 Coordination of tissue typing work

The ANZKX Tissue Typing Coordinator (ATTC) is responsible to track all of the steps of the tissue typing process for enrolled pairs and provide updates to the ANZKX Coordination Centre, including:

- confirmation of completion of tissue typing of donors and recipients referred to ANZKX;
- post-run review of recipient's antibody record against matched donor to ensure no authorised DSA have been missed;
- coordination of any serum requiring transfer to another lab for the crossmatch
- coordination of crossmatch test dates;
- review of crossmatch results;
- generation of crossmatch reports (donor de-identified);
- distribution of reminders to local Australian and New Zealand TTL's for SAB retesting of matched patients, as required.

Following a match run the ATTC is responsible along with the Australian and New Zealand TTL's for ensuring that patients in matched pairs are placed "On Hold" for their TWL enrolment. This will usually occur on the day a chain has been generated, reviewed and deemed to be suitable for all of the pairs involved.

Transplant candidates who have their TWL enrolment placed on hold will be reactivated in the OrganMatch waitlist promptly on the advice of the ATTC in consultation with the ANZKX Coordination Centre when:

- The referring team refuses a match offered to one of their recipients (e.g. the team would accept ABOi donor, but no DSA or specific DSA, but only if ABOc donor).
- A patient has final positive cell-based crossmatch result to the matched donor (in this case the other patients in the same chain will be reactivated on the waitlist).
- A chain breaks down because of recipient or donor unsuitability reasons.

9 Protocol for non-directed altruistic donors (NDAD)

9.1. Overarching principles

- Australian NDADs must meet the usual donor criteria to be entered into ANZKX.
- NDADs can be either entered directly into the ANZKX program or some Australian/New Zealand Transplant Centre's might choose to first check whether they match a highly sensitised recipient on the TWL. The NDADs preference should be taken into account when deciding how to allocate their kidney.
- It was decided by RTAC that if the latter option is chosen then the NDAD should only be offered to a TWL recipient if matched at Level 1-3 match of the current National Matching Algorithm:
- HLA A+B+DR = 0 mismatches (level 1) or
- Class 1 PRA>80% and HLA A+B+DR ≤2 mismatches (level 2+3).
- If a match is not found in the National pool at level 1-3 then these donors will then be allocated through the ANZKX program.
- The last donor in chain will be allocated **<u>nationally</u>** to an Australian TWL recipient fulfilling Level 1-3 match, or a New Zealand equivalent as per the NKAS algorithm.



 New Zealand NDADs entered into ANZKX will be allocated to the ANZKX, with the last donor in the chain allocated in accordance with the NZKAS to a New Zealand recipient on the deceased donor waiting list.

9.2. Step-by-step pathway (Australian NDADs only)

9.2.1 NDAD-Assessment and tissue typing:

- 1 Any NDAD should be referred to a local Australian or New Zealand TTL for typing only once they are deemed suitable to donate (i.e. passed medical, surgical and psychological suitability investigations).
- 2 Registration for NDAD is managed by their enrolments in Organ Match. It is important to create their Organ match enrolments as soon as the donor has been deemed suitable for the program, and document when the NDAD is likely to be ready for matching via a note in the NDADs organ match record.
- 3 Any suitable NDAD referred to the Australian or New Zealand TTL will be tested to the level required for ANZKX (i.e. including confirmatory typing and 2 field for all alleles)
- 4 The participating Australian or New Zealand TTL will have the donor "ready" on OrganMatch as soon as required testing is complete.
- 5 Matching of NDADs is performed at the time they are entered into ANZKX and testing completed. If there are particular timing requirements these need to be communicated to the NTTC via email and a note added to the NDAD Organ Match record.

9.2.2 End-of chain donor allocation:

- 6 The end-of-chain donor will be allocated against the Australian or New Zealand TWL approximately 3 weeks prior to the planned surgery date, according to the following rules:
 - a. Allocation to any recipient on the Australian TWL fulfilling level 1-3 match or New Zealand equivalent

if no suitable recipient:

b. Allocation to any recipient on the <u>TWL of the Australian state or New Zealand equivalent from which</u> <u>the NDAD originated.</u>

if still no suitable recipient:

c. Follow national Australian and/or New Zealand allocation override rules

if still no suitable recipient:

- d. Repeat allocation closer to the date of surgery of NDAD chain as per rules a-c above.
- 7 The ATTC will arrange a blood specimen* from the end-of-chain donor to be sent to their local Australian or New Zealand TTL for crossmatch against the TWL tray.
- 8 If the crossmatch against the end-of-chain donor is negative the matched recipient will also be temporarily off-listed.

*The TTL will freeze donor cells in the event there is no TWL recipient so they can be used for crossmatch several weeks later when an alternative recipient has been identified.

Comment: In the case that the kidney of the last donor in chain is directed to a recipient in an Australian state or New Zealand centre other than the NDAD origin state/centre, the standard payback rule will apply to compensate the state and/or centre of origin.



10 Protocol for orphaned kidneys and recipients

10.1 Definitions

Orphaned kidney: refers to a kidney removed from an ANZKX donor that cannot be transplanted into the matched recipient.

Orphaned recipient: refers to an ANZKX recipient whose co-registered donor has donated, but who has been unable to receive a kidney from the matched donor.

The protocol for orphaned kidneys and orphaned recipients was developed by the National Paired Kidney Exchange Program Advisory Group and has been revised by ANZKX, RACOS and RTAC (in Australia). In NZ, the NRTLT is responsible for the oversight of the NZKAS, including dealing with orphaned kidneys and recipients.

In the rare event where an exchange cannot proceed due to unforeseen clinical or logistical circumstances, the following is recommended:

10.2 Orphaned kidney

On the day of transplant surgery a recipient may suffer an acute event immediately prior to going to theatre, during induction or during their operation such that the procedure needs to be abandoned. Because donor surgery always occurs before recipient surgery, the donor has already had their kidney removed. This will result in an 'orphaned kidney'.

Donors are asked in advance to consent to their kidney being allocated to someone suitable on the deceased donor TWL if this circumstance arises. It is for this unlikely, but possible, contingency that a specimen of donor's whole blood (40ml) is taken at anaesthetic induction and transported with the kidney.

10.2.1 Process for determination and allocation of an orphaned kidney

Recipient Centre: must immediately notify the ANZKX Coordination Centre if the recipient has become acutely ill and is unable to undergo or continue with transplant surgery.

Steps for allocation of an orphan kidney:

- Depending on the logistics, an allocation to a highly sensitized recipient on the deceased donor waiting list (Level 1-3 on the National Allocation formula) should be sought. A decision regarding whether further transport of the kidney is possible for such an allocation will be made by the ANZKX Coordination Centre although advice from RACOS can be requested if required.
- If this is not possible or there is no recipient matched at level 1-3 on the National Allocation formula then:
 - If the kidney is in transit, the kidney will be allocated to a recipient on the transplant waiting list in the state/country of destination.
 - If the kidney is still in the state/country of origin, it will be allocated to a TWL recipient within this state/country of origin.
 - In New Zealand the kidney would be allocated to the top ranked recipient in the country or transplant service as per NKAS algorithm.
- The ANZKX Coordination Centre will alert the ATTC to perform an urgent OrganMatch search/virtual crossmatch to identify a suitable recipient.
- The ATTC will generate an allocation list via OrganMatch or NKAS system.



- Virtual crossmatch will be performed by the ATTC using the ANZKX matching criteria and the top 5 potential recipients.
- The ATTC will notify the ANZKX Coordination Centre about the identified potential match (state/country of origin or state/country of destination) and the transplanting centre under which the identified recipient is listed.
- The ANZKX Coordination Centre will contact the recipient's transplant centre to alert them of the probable allocation. The immunological and clinical information regarding the match will be available to be viewed by the transplant unit in OrganMatch.
- The ATTC will alert relevant TTL to prepare for an urgent cross-match test.
- The state TTL will perform CDC crossmatching against the top 5 recipients. This information may be provided to the physician retrospectively if necessary.
- The TTL performing the crossmatch will send a report to the recipient's centre, with copy to the ANZKX Coordination Centre and ATTC.
- The ANZKX Coordination Centre will ensure the TWL recipient centre has been duly informed.
- The ANZKX Coordination Centre will report the critical incident to the OTA/ANZKX Oversight committee and monitor outcomes.
- The ANZKX Coordination Centre will facilitate communication of the resultant issues and outcomes between donor and recipient centres as required.

10.3 Orphaned recipient

The definition and approach to Orphaned recipients is outlined in ANZKX Protocol 1, section 5.2.

10.3.1 Priority listing of orphaned recipients

RTAC agreed that orphaned recipients should receive priority listing for a suitable kidney from the National deceased donor organ pool. This is because the recipient's co-registered donor has already donated his/her kidney and thus the recipient no longer has recourse to an ANZKX exchange.

The process of prioritisation for Australian patients is that the 'orphaned recipient' will receive OrganMatch priority listing (Level 4 interstate exchange) for a suitable kidney from the National deceased donor organ pool.

Orphaned recipients in Australia are assigned a base score of 57,500,000 before other modifiers are applied. This base score prioritises an orphaned recipient after a Rank 3 (58,000,000; 4/6 matched and cPRA >80) and ahead of a Rank 5 (57,000,000; 6/6 matched and cPRA <50).

In New Zealand the recipient would be allocated in Rank 1 as per the NKAS algorithm, available on the NRTLT website.

Pre-emptive recipients are not listed in OrganMatch, as activation on the deceased donor waitlist starts with the first day on dialysis. In these cases, exception will be made after notification and approval by RTAC that the pre-emptive recipient can be listed for priority allocation on the deceased donor waiting list.



VERSION CONTROL					
Version	Date	Author	Comments		
V1.0	July 2019	ANZKX Team	AKX transitioned ANZKX		
V2.0	Feb 2021	ANZKX Team	Match run structure changed to continuous matching		
V3.0	Nov 2021	ANZKX Team	MMEx transition to OrganMatch		

