## Principles of the Kidney Matching Algorithms

## 1. Principles of Kidney TWL matching in OrganMatch

In OrganMatch, recipients listed on the kidney transplant waiting list (TWL) are matched with deceased organ donors using multi-tiered matching algorithms.

There are three parent algorithms, which run separately, in OrganMatch:

- Standard v2
- Interstate Utilisation v2
- ABOi (ABO incompatible) v2

Matching using the above parent algorithms triggers a series of functions.

The new version of the Kidney Matching Algorithm (KAv2) is effective from May 4, 2021.

#### 1.1 High-level compatibility check

The first step in matching is the high level compatibility check, which determines if recipients will progress to being matched in the next stage of the algorithm.

Check	Process	!	x	✓
HLA mismatches are identified	Compares <i>only</i> loci A, B, and DRB1. Compares at 1 field and 2 fields using the exception table.	HLA mismatches are listed	N/A	No HLA Mismatches
No Unacceptable Antigens (UA) identified	Compares donor HLA with recipient UA. Includes all HLA loci.	Potential UA for the donor are listed Will proceed to matching	UA and donor HLA match.  Does not proceed to matching.	Recipient and donor compatible. Proceeds to matching.
Extended criteria checked	Donor extended criteria is checked first and Recipient Extended Acceptance Criteria are compared.  If there is no extended criteria for the donor, proceed to matching.	N/A	Does not proceed to matching.	Proceeds to matching.
Valid ABO for program	Checks ABO compatibility.	N/A	Does not proceed to matching.	Recipient and donor compatible. Proceeds to matching.

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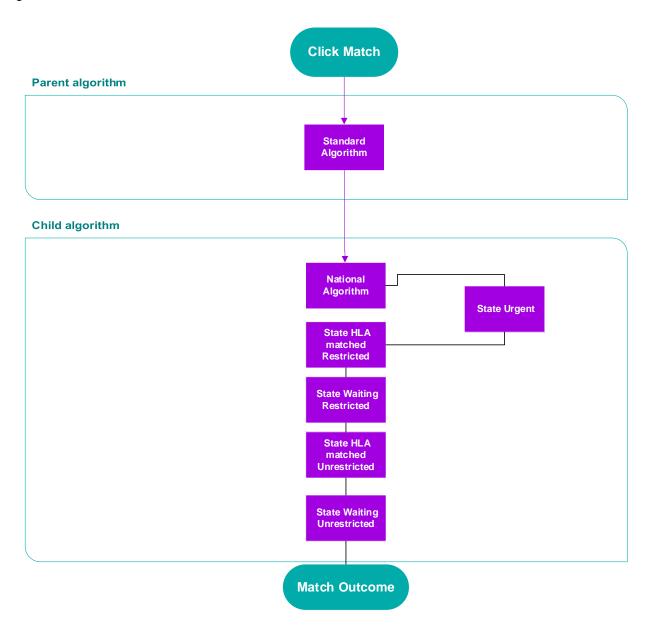
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If a recipient passes the high level compatibility check, they progress to matching, depending on which parent algorithm is selected:

## 2. Standard algorithm (KAv2)

The Standard kidney algorithm consists of a number of child algorithms. The child algorithms are executed in a set order and are dependent on the OrganMatch Lab from where the deceased organ donor is matched:



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# 2.1 National algorithm

Table 1: Criteria for National Algorithm levels.

Match level	Description	Criteria	Base score
1	Very Highly	<b>1a</b> mPRA>=99.7	99 700 000
	sensitised	<b>1b</b> mPRA>=99	99 000 000
	ABO Compatible	1c mPRA>=98	98 000 000
		<b>1d</b> mPRA>=97	97 000 000
		<b>1e</b> mPRA>=96	96 000 000
		<b>1f</b> mPRA>=95	95 000 000
National Urgent	ABO Compatible	Recipient National urgency >0	90 000 000
2	EPTS restriction	2a 0 mismatches HLA-A or HLA-B and EPTS <=25	89 000 000
	HLA matching	<b>2b</b> 1 mismatch HLA-A or HLA-B	
	Prioritises Low EPTS recipients	and EPTS<=25	88 000 000
	Matched at HLA DRB1	<b>2c</b> 2 mismatch HLA -A or HLA-B and EPTS<=25	87 000 000
	ABO Matched	2d 0 mismatches HLA -A or HLA-B	86 000 000
	KDPI max value is applied from this level down	and EPTS <=60	

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Match level	Description	Criteria	Base score
3	a/b/c HLA matching	3a 0 mismatch at HLA A or HLA B or HLA DRB1  and mPRA>80	79 000 000
	Highly Sensitised	3b 1 mismatch at HLA A or HLA B or HLA DRB1  and mPRA > 80	78 000 000
	d/e/f/g		
	HLA Matching	3c 2 mismatches at HLA A or HLA B or HLA DRB1	77 000 000
	Centre credit difference	And mPRA >80	
	difference	3d Matched at HLA DRB1	
		1 mismatch HLA A or HLA B	76 000 000
		And mPRA<= 80 And Centre credit difference <=-3	
		3e Matched at HLA DRB1	75 000 000
		2 mismatch HLA A or HLA B	
		And mPRA<= 80  Centre credit difference <=-6	
		3f mPRA > 80  Centre credit difference <=-9	74 000 000
		3g Centre credit difference <-20	73 000 000

Table 1: Criteria for national algorithm levels

See Definitions for more information on centre credit difference

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### 2.2 Calculation of kidney national algorithm score

The National Score is calculated by assigning a base score – depending on the number of HLA mismatches, Match cPRA (mpra), the patient's National Urgency Index and the centre credit difference between the donor and recipient centres (see *Table 1*) – and then add the following bonus points:

Other parameters	Bonus points added
Paediatric	250 000
Donor centre = patient centre	50
Recipient Centre credit	1000 + recipient centre credit
Recipient and Donor are HLA DRB1 homozygote	500 000 ( except level 3G)
Waiting time (on dialysis)	Number of months x 1

The lowest threshold is 73,000,000. Recipients with a scoreless than 73,000,000 then progress to the **state algorithms**.

See Definitions for more information on lab credit.

### 3. State algorithms

#### 3.1 Calculation of state HLA algorithm scores

The base score is calculated for each recipient if:

- their ABO type is compatible with donor as determined by the ABO Type Selection Rules
- they are from the same state centre as the donor.

#### 3.2 State Kidney Algorithm

EPTS-KDPI restriction occurs when EPTS-KDPI < =50. The intent of this parameter is to enable low KDPI donor kidney to be prioritised to low EPTS recipients.

If the all patients on the transplant waiting list are categorised in the "restricted group ie: EPTS-KDPI < 50, only the restricted algorithm will appear in the TWL match.

An example would be a high KDPI donor eg; KDPI = 80, and as the max EPTS is 100, all recipients would fall into the "restricted category" as EPTS-KDPI <=50 applies. There would be no patients outside of this category.

For a lower KDPI donor eg; KDPI = 20, only the patients with EPTS < 70 will fall into the restricted group to meet the restricted criteria. So this group of patients will be prioritised. The remaining patients with EPTS >70 will be in unrestricted category and they will appear under (lower rank) the restricted patients in the TWL matching.

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So you only expect both the restricted and unrestricted lists to appear with the lower KDPI donors , as the restricted list will appear above the unrestricted list

Flexibility for state to nominate at which level to move from state HLA to State waiting. The current state thresholds are

•	NSW	1E
•	QLD	1E
•	VIC	1E
•	WA	1E
•	SA	1E

ABO rules are aligned for NSW, VIC, SA, WA, QLD. QLD have an additional requirement of B donors to match O recipients.

KDPI max functionality also applies as determined by local clinical units.

The state matching algorithm sequentially flows as follows

- State HLA restricted
- State waiting restricted
- State HLA unrestricted
- State waiting unrestricted

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Table 2: Criteria for state algorithm levels

Level	Description	Details	Base Score
State Urgent	State Urgency Index >0	Urgency index added to base score	60 000 000

Level	Description	Details	Restricted base score	Unrestricted base score
State HLA	HLA mismatches A/B/DRB1	1a 000 1b 100 or 010 1c 110 1d 001 1e 200 or 020 1f 101 or 011 1g 210 or 120	49 000 000 48 000 000 47 000 000 46 000 000 45 000 000 44 000 000 43 000 000	39 000 000 38 000 000 37 000 000 36 000 000 35 000 000 34 000 000 33 000 000
State Waiting	Months on dialysis	Number of months x 1	40 000 000	30 000 000

#### Additional scores

- Paediatric bonus of 100 000 for restricted algorithms state HLA and state waiting
- Recipient and donor are HLA DRB1 homozygous bonus 500 000 to state HLA matching algorithms only.

In the event that a more than one patient has the same score, the ranking is randomised.

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# 4. Interstate Utilisation algorithm

- The Interstate Utilisation Algorithm is an additional algorithm, which is invoked by the OrganMatch user if the standard algorithm did not list enough patients to enable transplantation.
- The score is calculated by assigning a base score and bonus points, by applying the criteria described in *Table 3: Interstate Utilisation Algorithm*.
- Interstate utilisation considers all states other than the donor state, base scoring as described in *Table 3: Interstate Utilisation Algorithm*
- In the event that a patient from a OM lab which is different to the donor OM lab is matched via the national algorithm, there will be a duplication of this patient on the Interstate utilisation list.

Table 3: Interstate Utilisation Algorithm

Level	Description	Details	Restricted base score	Unrestricted base score
State HLA	HLA mismatches	1a 000	19 000 000	9 000 000
	A/B/DRB1	<b>1b</b> 100 or 010	18 000 000	8 000 000
		1c 110	17 000 000	7 000 000
		1d 001	16 000 000	6 000 000
		<b>1e</b> 200 or 020	15 000 000	5 000 000
		<b>1f</b> 101 or 011	14 000 000	4 000 000
		1g 210 or 120	13 000 000	3 000 000
State Waiting	Months on dialysis	Number of months x 1	10 000 000	0

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# 5. Other matching algorithms

#### **5.1** ABOi

- ABOi matching algorithm can only be used to match blood group AB donors with blood group A and B recipients that are registered on the TWL kidney program, and willing to accept an ABOi donor.
- The score is calculated in the same manner as the state algorithm; by assigning a base score and bonus points using the criteria described in Table 2

#### 5.2 Increased Viral Risk Donors (IVRD)

• Donors that are registered as IVRD in OrganMatch will be matched using the standard algorithm, but with patients only registered as willing to accept IVRD donors.

### 5.3 Hepatitis C (Hep C)

 Donors that are registered as Hep C in OrganMatch will be matched using the standard algorithm, but with patients only registered as willing to accept Hep C donors.

### 6. ABO selection rules

The ABO selection rules determine the acceptable organ matches, as shown:

Algorithm	Level	Donor ABO type	Patient ABO type
		А	A
		А	AB
		В	В
		В	AB
National	Level 1	AB	AB
		0	0
		0	A
		0	В
		0	AB
National		А	А

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Algorithm	Level	Donor ABO type	Patient ABO type
	Level 2 and Level 3	В	В
		AB	AB
		0	0
		А	А
	NSW	А	AB
State	WA	В	В
State	VIC	В	AB
	SA	AB	AB
		0	0
		A	A
		А	AB
0	01.5	В	В
State	QLD	В	AB
		AB	AB
		0	0
		0	В

## 7. Use of KDPI min max

Currently the default setting for all recipients in OrganMatch – Kidney TWL program, the minmax KPDI is set at 1 -100. The max value can be set by the clinical unit, which will restrict the recipient being matched at the national level (level 2 and Level 3) and also the state level.

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# **Abbreviations**

Abbreviation	Definition
ABOi	ABO incompatible
EPTS	Estimated Post transplant survival
HEP C	Hepatitis C
HLA	Human Leucocyte Antigen
IVRD	Increased viral risk donor
KAv2	Kidney Matching Algorithms version 2
KDPI	Kidney Donor Prognosis Index
ОМ	OrganMatch
TWL	Transplant waiting list

# **Definitions**

	Definition							
Centre	OrganMatch Lab							
OM lab Credit	OM Lab credit This informatio Eg : NSW lab cr	n can be edit = 48	found in	OrganMat = -4				•
	Organ exchange report							Organ <b>Match</b>
	,	euit – 31.						Organ <b>Match</b>
	Organ exchange report				Recipient OM Laboratory			
	Organ exchange report  Donor OM Laboratory	NSW	NZ	QLD	SA	VIC	WA	Total Organs Exchanged
	Organ exchange report  Donor OM Laboratory NSW	N SW 3871	<b>NZ</b> 0	373	\$A 218	477	125	Total Organs Exchanged
	Organ exchange report  Donor OM Laboratory NSW NZ	NSW 3671 0	NZ 0 0	373 0	\$A 218 0	477 0	125 0	Total Organs Exchanged 4882 0
	Organ exchange report  Donor OM Laboratory  NSW  NZ  OLD	NSW 3871 0 395	NZ 0 0 0 0	373 0 2457	\$A 216 0 80	477 0 218	125 0 47	Total Organs Exchanged 4882 0 3195
	Organ exchange report  Donor OM Laboratory NSW NZ GLD SA	NSW 3871 0 985 216	NZ 0 0 0 0 0 0	373 0 2457 119	\$A 216 0 80 1400	477 0 216 138	125 0 47 41	Total Organs Exchanged 4862 0 3196
	Organ exchange report  Donor OM Laboratory  NSW  NZ  OLD	NSW 3871 0 395	NZ 0 0 0 0	373 0 2457	\$A 216 0 80	477 0 218	125 0 47	Total Organs Exchanged 4882 0 3195

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Term	Definition
Centre	Donor state lab credit –recipient OM lab centre credit
Credit difference	Eg: If NSW donor and QLD recipient the centre credit difference
	= -43 = -1

## Referenced external documents

N/A

# **Referenced OrganMatch documents**

N/A

# **Change history**

Version number	Effective date	Summary of change
1	04/05/2021	First version
2	26/05/2021	Reformatting to align with OM documents
3	Refer to footer	Change to QLD state matching algorithm threshold. Change from 1F to 1E

# **Electronic signature**

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