

5 success

5 accuracy

3 accuracy

Bulk accuracy

DRB345 accuracy

Time point accuracy

Composite accuracy

## 5 success

	Field 1					Field 2					Field 3		
	scHLAcount	arcasHLA	HLAminer	PHLAT	OptiType	scHLAcount	arcasHLA	HLAminer	PHLAT	OptiType	scHLAcount	arcasHLA	PHLAT
A	$0.68 \pm 0.04$	$0.86 \pm 0.03$	$0.71 \pm 0.03$	$0.89 \pm 0.03$	$0.92 \pm 0.02$	$0.44 \pm 0.04$	$0.64 \pm 0.03$	$0.41 \pm 0.03$	$0.81 \pm 0.03$	$0.87 \pm 0.03$	$0.06 \pm 0.02$	$0.79 \pm 0.03$	$0.8 \pm 0.03$
B	$0.44 \pm 0.04$	$0.82 \pm 0.02$	$0.56 \pm 0.03$	$0.88 \pm 0.02$	$0.92 \pm 0.02$	$0.21 \pm 0.03$	$0.63 \pm 0.03$	$0.2 \pm 0.03$	$0.81 \pm 0.03$	$0.9 \pm 0.02$	$0.2 \pm 0.03$	$0.8 \pm 0.03$	$0.83 \pm 0.03$
C	$0.82 \pm 0.03$	$0.82 \pm 0.02$	$0.66 \pm 0.02$	$0.91 \pm 0.02$	$0.94 \pm 0.01$	$0.16 \pm 0.03$	$0.55 \pm 0.03$	$0.35 \pm 0.03$	$0.81 \pm 0.03$	$0.9 \pm 0.02$	$0.02 \pm 0.01$	$0.64 \pm 0.03$	$0.82 \pm 0.03$
DPA1	$0.71 \pm 0.02$	$0.95 \pm 0.02$	$0.66 \pm 0.02$	—	—	$0.69 \pm 0.02$	$0.94 \pm 0.02$	$0.54 \pm 0.03$	—	—	$0.65 \pm 0.03$	$0.94 \pm 0.02$	—
DPB1	$0.25 \pm 0.03$	$0.85 \pm 0.03$	$0.43 \pm 0.02$	—	—	$0.25 \pm 0.03$	$0.84 \pm 0.03$	$0.42 \pm 0.02$	—	—	$0.34 \pm 0.04$	$0.84 \pm 0.03$	—
DQA1	$0.81 \pm 0.03$	$0.92 \pm 0.02$	$0.85 \pm 0.02$	$0.94 \pm 0.01$	—	$0.55 \pm 0.03$	$0.88 \pm 0.02$	$0.55 \pm 0.03$	$0.86 \pm 0.03$	—	$0.71 \pm 0.04$	$0.88 \pm 0.02$	$0.86 \pm 0.03$
DQB1	$0.83 \pm 0.03$	$0.91 \pm 0.02$	$0.88 \pm 0.02$	$0.91 \pm 0.02$	—	$0.54 \pm 0.04$	$0.76 \pm 0.03$	$0.67 \pm 0.03$	$0.89 \pm 0.02$	—	$0.59 \pm 0.04$	$0.78 \pm 0.03$	$0.87 \pm 0.02$
DRB1	$0.7 \pm 0.04$	$0.9 \pm 0.02$	$0.73 \pm 0.03$	$0.89 \pm 0.03$	—	$0.48 \pm 0.04$	$0.88 \pm 0.03$	$0.34 \pm 0.03$	$0.85 \pm 0.03$	—	$0.49 \pm 0.04$	$0.88 \pm 0.03$	$0.78 \pm 0.03$

## 5 accuracy

	Field 1					Field 2					Field 3		
	scHLAcount	arcasHLA	HLAminer	PHLAT	OptiType	scHLAcount	arcasHLA	HLAminer	PHLAT	OptiType	scHLAcount	arcasHLA	PHLAT
A	$0.41 \pm 0.03$	$0.86 \pm 0.03$	$0.7 \pm 0.03$	$0.88 \pm 0.03$	$0.81 \pm 0.03$	$0.24 \pm 0.02$	$0.63 \pm 0.03$	$0.4 \pm 0.03$	$0.8 \pm 0.03$	$0.76 \pm 0.04$	$0.02 \pm 0.01$	$0.59 \pm 0.03$	$0.44 \pm 0.04$
B	$0.26 \pm 0.02$	$0.82 \pm 0.03$	$0.56 \pm 0.03$	$0.88 \pm 0.02$	$0.81 \pm 0.03$	$0.13 \pm 0.02$	$0.62 \pm 0.03$	$0.2 \pm 0.03$	$0.81 \pm 0.03$	$0.78 \pm 0.03$	$0.07 \pm 0.02$	$0.6 \pm 0.03$	$0.58 \pm 0.04$
C	$0.48 \pm 0.02$	$0.81 \pm 0.02$	$0.66 \pm 0.02$	$0.9 \pm 0.02$	$0.82 \pm 0.03$	$0.1 \pm 0.02$	$0.55 \pm 0.03$	$0.34 \pm 0.03$	$0.81 \pm 0.03$	$0.78 \pm 0.03$	$0.01 \pm 0.01$	$0.47 \pm 0.03$	$0.44 \pm 0.04$
DPA1	$0.62 \pm 0.02$	$0.94 \pm 0.02$	$0.66 \pm 0.02$	—	—	$0.6 \pm 0.02$	$0.93 \pm 0.02$	$0.53 \pm 0.03$	—	—	$0.56 \pm 0.03$	$0.92 \pm 0.02$	—
DPB1	$0.21 \pm 0.03$	$0.84 \pm 0.03$	$0.42 \pm 0.02$	—	—	$0.21 \pm 0.03$	$0.83 \pm 0.03$	$0.41 \pm 0.02$	—	—	$0.21 \pm 0.03$	$0.83 \pm 0.03$	—
DQA1	$0.72 \pm 0.03$	$0.91 \pm 0.02$	$0.74 \pm 0.03$	$0.93 \pm 0.02$	—	$0.48 \pm 0.03$	$0.87 \pm 0.02$	$0.47 \pm 0.03$	$0.85 \pm 0.03$	—	$0.45 \pm 0.03$	$0.87 \pm 0.02$	$0.72 \pm 0.03$
DQB1	$0.53 \pm 0.02$	$0.9 \pm 0.02$	$0.75 \pm 0.03$	$0.9 \pm 0.02$	—	$0.35 \pm 0.03$	$0.75 \pm 0.03$	$0.56 \pm 0.03$	$0.88 \pm 0.02$	—	$0.3 \pm 0.03$	$0.66 \pm 0.03$	$0.7 \pm 0.03$
DRB1	$0.4 \pm 0.02$	$0.9 \pm 0.03$	$0.72 \pm 0.03$	$0.88 \pm 0.03$	—	$0.28 \pm 0.02$	$0.87 \pm 0.03$	$0.33 \pm 0.03$	$0.84 \pm 0.03$	—	$0.29 \pm 0.03$	$0.88 \pm 0.03$	$0.67 \pm 0.04$

## DRB345 accuracy

	Field 1		Field 2		Field 3	
	arcasHLA	HLAminer	arcasHLA	HLAminer	arcasHLA	HLAminer
	Unfiltered Filtered	Unfiltered Filtered	Unfiltered Filtered	Unfiltered Filtered	Unfiltered Filtered	Unfiltered Filtered
DRB3	0.46 ± 0.03 0.79 ± 0.04	0.44 ± 0.03 0.78 ± 0.03	0.46 ± 0.03 0.79 ± 0.04	0.35 ± 0.03 0.65 ± 0.04	0.46 ± 0.03 0.79 ± 0.04	— —
DRB4	0.44 ± 0.02 0.91 ± 0.02	0.77 ± 0.04 0.88 ± 0.03	0.4 ± 0.02 0.69 ± 0.04	0.14 ± 0.03 0.17 ± 0.03	0.39 ± 0.03 0.66 ± 0.04	— —
DRB5	0.35 ± 0.03 0.9 ± 0.03	0.33 ± 0.03 0.83 ± 0.03	0.35 ± 0.03 0.88 ± 0.03	0.15 ± 0.03 0.33 ± 0.04	0.33 ± 0.03 0.86 ± 0.03	— —

### Supplemental Table X - *HLA-DRB345* accuracy

Accuracy of *HLA-DRB345* genotypes filtered or unfiltered by KNN copy number classifier. Values represent mean accuracy +/- SEM