

Table 1 : Basic statistics about MADaM and AmpliSAS processing

MADaM	Locus	DRB1	DQA1	DQB1	DPB1
	N. samples	3'011	3'186	2'769	3'080
	N. TV	2.75+/-0.81	1.65+/-0.49	1.68+/-0.47	1.79+/-0.41
	Min/Max TV	1-5	1-3	1-3	1-3
AmpliSAS	Locus	DRB1	DQA1	DQB1	DPB1
	N. samples	3'011	3'186	2'769	3'080
	N. TV	2.91+/-0.82	1.98+/-0.74	1.92+/-0.6	1.95+/-0.68
	Min/Max TV	1-8	1-9	1-5	1-6

Table 1 : For each locus this table gives the number of processed samples (N. samples), the number of true variants found (N. TV) and the minimum and maximum number of true variants found per samples (Min/Max TV), according to the two algorithm used : MADaM (top) and AmpliSAS (bottom).