

The Swine Leukocyte Antigen (SLA) Nomenclature System: Current Status after 10 Years of its Establishment

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[ABSTRACT]

The SLA Nomenclature Committee was established in 2002 at the 28th ISAG Annual Conference in Göttingen, Germany. It is now also affiliated to the Veterinary Immunology Committee of the International Union of Immunological Societies (IUIS-VIC), which is the umbrella organization for immunology societies worldwide. To date, there are 131 class I (SLA-1, SLA-2, SLA-3, SLA-6) and 174 class II (DRA, DRB1, DQA, DQB1, DMA) alleles officially designated. There are also 31 class I (SLA-1-3-2-6) and 26 class II (DRA-DRB1-DQA-DQB1) high-resolution (allele level) haplotypes designated, while designation of low-resolution (allele group level) haplotypes is in progress. In conclusion, the SLA system is among the most well characterized MHC systems in non-primate species. Continuous efforts on characterizing new SLA alleles and studying of SLA diversity in various pig populations will further our understanding of the allelic architecture and polymorphism of the SLA system, which may ultimately facilitate the research on swine immunology, vaccine development, and the use of swine as a large biomedical animal model.

2012 ISAG SLA Nomenclature Committee

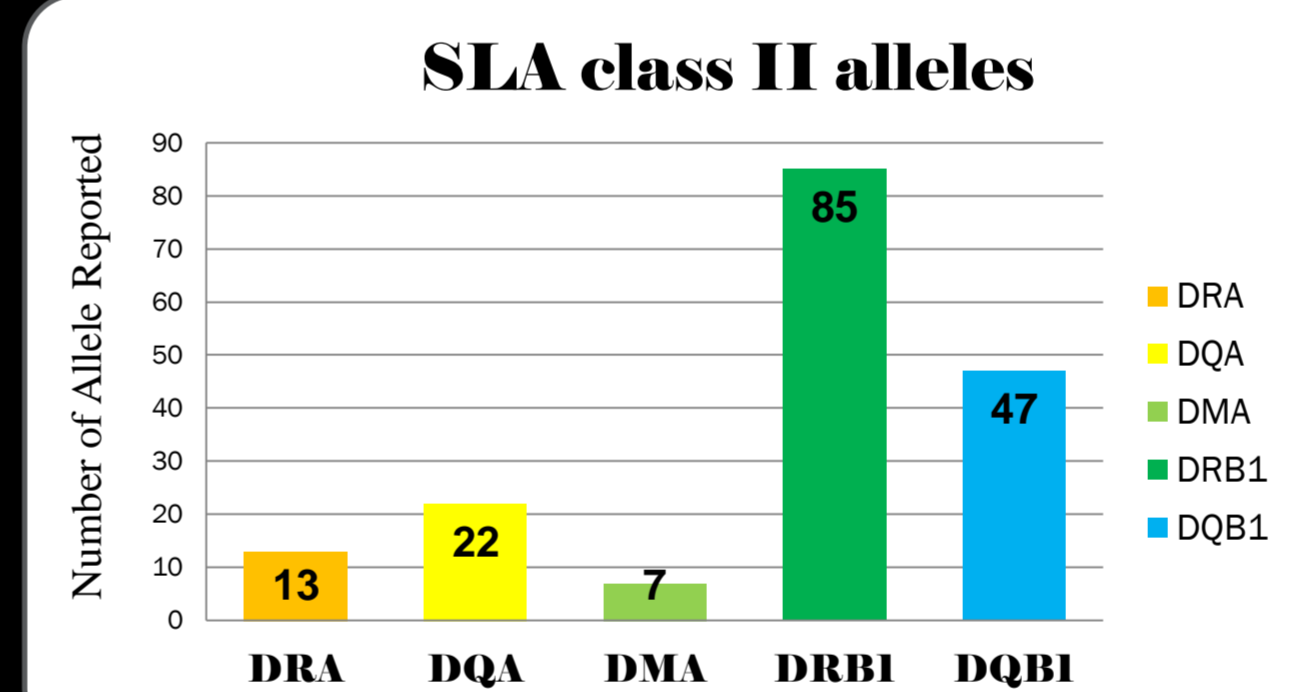
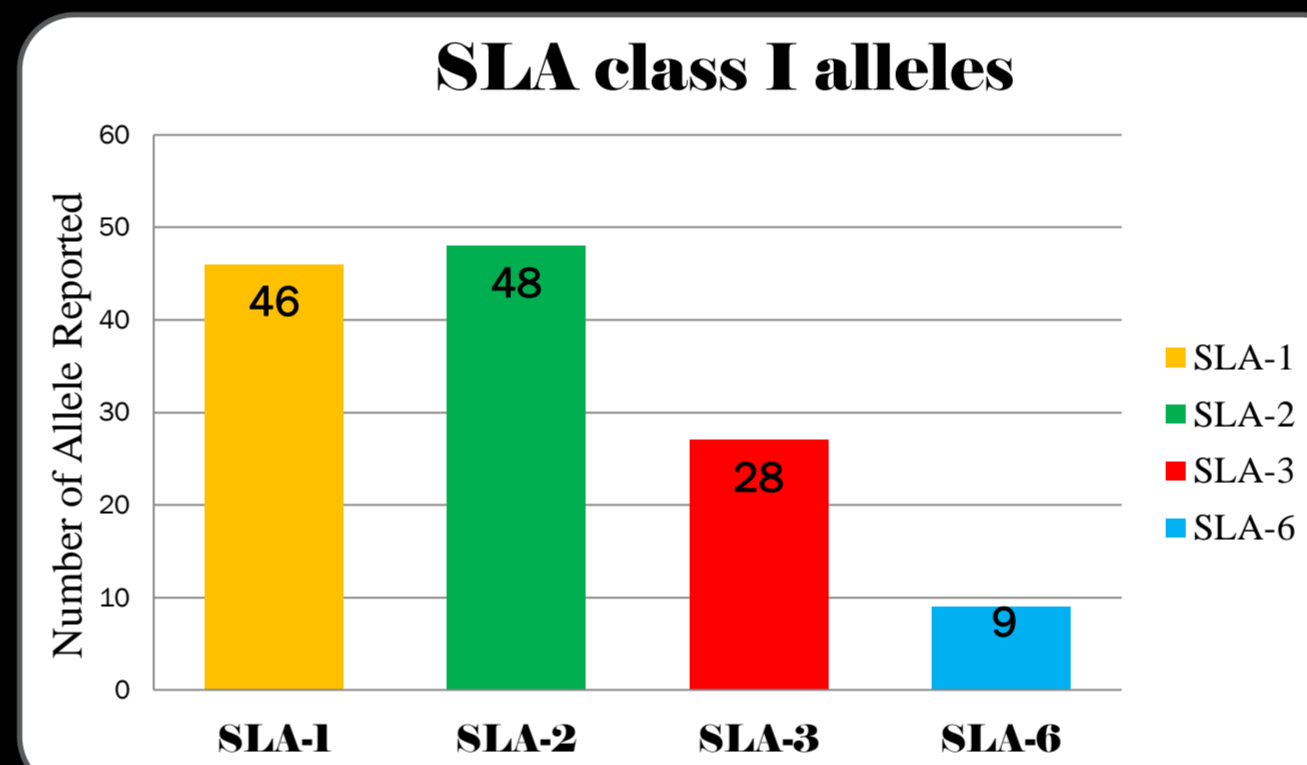
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[Statistics of Current SLA Sequence Database]

	Identified alleles	
2005	SLA class I	88
	SLA class II	127

	Previously identified Alleles (2005)	Newly identified alleles	Total
2009	SLA class I	88 + 37	125
	SLA class II	127 + 37	164

	Previously identified Alleles (2009)	Newly identified alleles	Total
Currently	SLA class I	125 + 6	131
	SLA class II	164 + 10	174



IPD-MHC SLA Sequence Database Website http://www.ebi.ac.uk/ipd/mhc/sla

IPD-MHC Database

Swine (SLA) Sequences - Release 1.2.0 16/05/2008

Welcome to the IPD-MHC Swine Leukocyte Antigen (SLA) website. The site is intended as a resource for information on the nomenclature and DNA sequence data for the genes of the swine MHC complex. The data presented represents work published or submitted to public databases by many authors and has been compiled and edited by the members of the SLA Nomenclature Committee of the International Society for Animal Genetics (ISAG).

The information presented here is based on the reports of the SLA Class I Nomenclature Workshops:

- Smith DM, Lunney JK, Martens GW, Ando A, Lee JH, Ho CS, Schook LB, Renard C, Chardon P. Nomenclature for factors of the SLA class-I system, 2004. *Tissue Antigens* (2005), 65:136-9
- Smith DM, Lunney JK, Ho CS, Martens GW, Ando A, Lee JH, Schook LB, Renard C, Chardon P. Nomenclature for factors of the swine leukocyte antigen class-II system, 2005. *Tissue Antigens* (2005), 66:623-9
- Ho CS, Lunney JK, Ando A, Rogel-Gaillard C, Lee JH, Schook LB, Smith DM. Nomenclature for factors of the SLA system, update 2008. *Tissue Antigens* (2009), 73:307-15

Both articles are freely available from [BioRxiv](#).

The following additional information on the SLA region is also available:

- Conditions for Acceptance of New Allele Sequences
- Map of the SLA Class I Region
- Map of the SLA Class II Region
- Phylogeny of SLA-1, SLA-3 and SLA-6 (pdf)
- Phylogeny of SLA-2 (pdf)
- Phylogeny of SLA-DRB1 (pdf)
- Phylogeny of SLA-DQA (pdf)
- Phylogeny of SLA-DQB1 (pdf)

[References]

- Ho CS, Lunney JK, Ando A, Rogel-Gaillard C, Lee JH, Schook LB, Smith DM. Nomenclature for factors of the SLA system, update 2008. *Tissue Antigens* 2009;73:307-15.
- Ho CS, Lunney JK, Franzo-Romain MH, Martens GW, Lee YJ, Lee JH, Wysocki M, Rowland RRR, Smith DM. Molecular characterization of swine leukocyte antigen (SLA) class I genes in outbred pig populations. *Animal Genetics* 2009;40:468-78.
- Ho CS, Lunney JK, Lee JH, Franzo-Romain MH, Martens GW, Rowland RRR, Smith DM. Molecular characterization of swine leukocyte antigen (SLA) class II genes in outbred pig populations. *Animal Genetics* 2010;41:428-32.
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- Renard C, Hart E, Schra H, Beasley H, Coghill P, Howe K, Harrow J, Gilbert J, Sims S, Rogers J, Ando A, Shigenari A, Shiina T, Inoko H, Chardon P, Beck S. The genomic sequence and analysis of the swine major histocompatibility complex. *Genomics* 2006;88:96-110.
- Tanaka-Matsuda M, Ando A, Rogel-Gaillard C, Chardon P, Uenishi H. Difference in number of loci of swine leukocyte antigen classical class I genes among haplotypes. *Genomics* 2009;93:216-73.

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[SLA class I and II haplotypes]

Table 1. High-resolution (Hp) 31 SLA class I and 26 SLA class II haplotypes

Hp ^a	Breed ^b	Previous designation	SLA-1	SLA-3	SLA-2	SLA-6	Hp ^a	Breed ^b	Previous designation	DRB1	DQB1	DQA	DRA
1a.0	Large White	H01	0101	0101	0101	0101	0.1	Large White, Korean native pig	H01	0101	0101	0101	010101
1b.0	Large White	H28	01rh28	01rh28	0101	ND ^c	0.2	NIH, Sinclair, Hanford	a, b	0201	0201	0201	010101
2.0	NIH, Sinclair, Hanford	a, b, H10	0201, 0701	Null ^d	0201	w02sa01	0.3	NIH	c	0301	0301	0102	0201
3.0	NIH	e, H59	Null	0301	0301	0103	0.4	NIH	d	0201	040101	020201	010102
4a.0	NIH, Duroc	d, H04	0401	0401	0401	0102	0.5	Yucatan	x	0501	0201	020202	020301
4b.0	Yucatan	x	0401	0401	040201	0104	0.6	Yucatan	w	0501	0801	0103	020203
4c.0	Meishan	k	0401	0401	0401	0104	0.7	Yucatan	y	0601	0601	01my01	0203my01
5.0	Yucatan	w	0401	05sw01	w08sw01	Null	0.8	Yucatan	z	0801	0202	0203	010101
6.0	Yucatan	y	08sy01	0601	05sy01	03sy01	0.9	Westran	None	0201	0402we01	03we01	0101we01
7.0	Yucatan	z	0801	0701	0502	0101	0.10	Sinclair, Hanford	a	0401	0801	ND ^c	ND
8.0	Westran	None	02we02, 04we01	0302	07we01	01we01	0.11	Sinclair	c	0901	0402	ND	020202
9.0	Sinclair, Hanford	a	0601	0501	0601	ND	0.12	Sinclair	d	0602	0701	0301	020201
10.0	Sinclair	e	0501	hm22	0302	ND	0.13	Hanford, Duroc	e, d2	0403	0303	ND	ND
11.0	Sinclair	d	0101, w09sm09	0701sm19	0501	ND	0.14	Meishan	M, K	0901	0801	0301	010103
12.0	Hanford	e	08sm08, w09sm09	0502	10sm01	ND	0.15a	Meishan	N	0401	0201	0203	0201
13.0	Hanford	f, d2	w10sm21	0401	w13sm20	ND	0.15b	Banna	None	0402	0202	020202	020301
14.0	Large White	H12	0102	01rh12	07rh12	ND	0.16	Clawn	e1	Hae21	0601	ND	ND
15.0	Large White	H34	0102	07rh34	05rh34	ND	0.17	Clawn	e2	0801	0501	ND	ND
16.0	Clawn	e1	0401	0602	w09an02	ND	0.18	Meishan	L	1401	040102	02es01	010103
17.0	Clawn	e2	ND	03an02	06an03	ND	0.19	SNU, Austrian Pietrain		0404	0701	ND	ND
18.0	Meishan	M	0401	0304	06me01	0102	0.20	Austrian Pietrain (n = 2)		0602	0303	ND	ND
19.0	Meishan	N	08sm05, 13sm21	0602	w09sn01	0105	0.23	Korean native pig, Landrace, Austrian Pietrain		1001	0601	ND	ND
20.0	Meishan	L	w10es01, es02	0101	H10102	0103	0.24	Austrian Pietrain (n = 1)		0701	0201	ND	ND
21.0	Commercial breeds	H03	rh03	0601	05rh03	ND	0.25	Hampshire ^e , Austrian Pietrain	None	1301	0901	ND	ND
25.0	Hampshire ^e	None	H101	0302	0701	ND	0.30	Korean native pig, Duroc	d1	1101	0503	02jh01	020202
27.0	Duroc	d1	06an04, 08an03	0101	0102	ND	0.38	Landrace		0101	0901	ND	ND
28.0	Landrace	H11	0901 (a), 1501 (b)	070102	0503	ND							
56.0	Korean native pig	None	Hjh01	0303	jh01	w04jh01							
59.0	Korean native pig	None	Hjh02	0503	jh02	0102							
60.0	Duroc	d2	an02	0502	1002	ND							
62.0	Landrace	H12	1401	hb06	0602	ND							

^a SLA class I and II haplotype assignments based on Smith *et al.*, 2005.

^b Breed in which the haplotype was sequence-based typed; haplotype may be found in other breeds.

^c ND, not determined specific allele at this locus.

^d Null, no expression of this locus.

^e Haplotype Hp-25.0 and Hp-0.25 were found in the LLC-PK1 pig cell line (ATCC) derived from a Hampshire pig.