

Genetic Background Monitoring Report

No.: 101206-15

BRC No.: RBRC01544

Strain Name (Official Nomenclature): B6.129P2-Sirpa<tm1Nog>

Strain Type: Mutant Strain

Strain of Origin: 129P2/OlaHsd

Donor Strain: E14.1 (ES cell)

Background Strain: C57BL/6NJcl

Date: 2010/12/6

Locus	Type	ID of animals surveyed (Generation ¹) / Sex		Male	a-type	b-type (B6 type)
		Location ²				
1	D1Mit212	SSLP	Chr1:40049840-40049985	S00080306 (N10F1+N3)	C3,FVB	B6,D2,129,BALB,CBA
2	D1Mit306	SSLP	Chr1:98827535-98827643		D2,129	B6,BALB,C3,CBA,FVB
3	D2Mit433	SSLP	Chr2:57149522-57149694		D2,C3,129,CBA,FVB	B6,BALB
4	D3Mit333	SSLP	Chr3:44511652-44511774		C3,CBA,FVB	B6,D2,BALB,129
5	D3Mit106	SSLP	Chr3:111841219-111841382		D2,BALB,C1,129,CBA,FVB	B6
6	D4Mit149	SSLP	Chr4:3584271-3584381		D2,BALB,C3,129,CBA	B6,FVB
7	D4Mit108	SSLP	Chr4:38319122-38319254		BALB,C3,129	B6,D2,CBA,FVB
8	D5Mit148	SSLP	Chr5:32278263-32278411		BALB,C3,129,CBA,FVB	B6,D2
9	D5Mit168	SSLP	Chr5:137534245-137534394		D2,BALB,C3,129,CBA,FVB	B6
10	D6Mit9	SSLP	Chr6:87375892-87376028		D2,C3,129,CBA,FVB	B6,BALB
11	D7Mit230	SSLP	Chr7:56764849-56764953		D2,C3,129,FVB	B6,BALB,CBA
12	D8Mit293	SSLP	Chr8:37226054-37226173		D2,BALB,C3,129,CBA,FVB	B6
13	D9Mit191	SSLP	Chr9:46494456-46494601		D2,BALB,C3,CBA,FVB	B6,129
14	D10Mit282	SSLP	Chr10:20654078-20654202		BALB,129,FVB	B6,D2,C3,CBA
15	D11Mit219	SSLP	Chr11:72132472-72132603		D2,129	B6,BALB,C3,CBA,FVB
16	D12Mit190	SSLP	Chr12:60829453-60829579		D2,FVB	B6,BALB,C3,129,CBA
17	D13Mit179	SSLP	Chr13:44860362-44860453		BALB,C3,129,CBA,FVB	B6,D2
18	D14Mit50	SSLP	Chr14:21831604-21831793		D2,BALB,129,CBA,FVB	B6,C3
19	D15Mit138	SSLP	Chr15:39861164-39861310		C3	B6,D2,BALB,129,CBA,FVB
20	D16Mit50	SSLP	Chr16:78953071-78953200		D2,129,CBA,FVB	B6,BALB,C3
21	D17Mit152	SSLP	Chr17:65689824-65689949		D2,BALB,FVB	B6,C3,129,CBA
22	D18Mit68	SSLP	Chr18:21594126-21594238		D2,BALB,CBA,FVB	B6,C3,129
23	D19Mit16	SSLP	Chr19:20420342-20420474		D2,C3,CBA	B6,BALB,FVB
24	DXMit117	SSLP	Chr X:127822562-127822687		D2,129,FVB	B6,BALB,C3,CBA

¹Generation definitions: N_Number of backcross generations; F_Filial or inbreeding generations; M_Cross-intercross breeding generations; ?_ Use when the prior breeding history is not known; +_Generation numbers before the "+" took place in the donating investigator, after the "+", at RIKEN BRC.

² based on UniSTS annotation of NCBI Build 37

Haplotype combination:

a/a_a-type homozygous allele, b/b_b-type homozygous allele, a/b_heterozygous allele