

Ly6a

4

3

2

1

locus	csq terms	alleles	BALB/c	CB6J	NONBL/BL	CASTLEJ	PWH/PhU	AMUJ	COLLJ	DMB/J	FVB/J	LPJ	MOL/REU	NONBL/BL
15:74900018	downstream gene variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900020	downstream gene variant	["T","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900041	downstream gene variant	["G","G/C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900005	downstream gene variant	["T","TTT/C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900009	downstream gene variant	["C","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900144	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900164	downstream gene variant	["A","AT/CCT"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900173	downstream gene variant	["C","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900190	downstream gene variant	["A","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900289	downstream gene variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900290	downstream gene variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900299	downstream gene variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900425	downstream gene variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900438	downstream gene variant	["G/C","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900469	downstream gene variant	["C","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900492	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900505	downstream gene variant	["C","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900642	downstream gene variant	["CT","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900643	downstream gene variant	["T","TGC/C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900698	downstream gene variant	["A","ATCTGCTC"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900698	downstream gene variant	["ATCTGCTCTC","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900698	downstream gene variant	["CTCTGCTCTGCTCTC","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900847	downstream gene variant	["T","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900852	downstream gene variant	["T","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900859	downstream gene variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74900984	downstream gene variant	["T","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901340	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901607	downstream gene variant	["GGA","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901607	downstream gene variant	["G","GGAAGAGAGA"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901607	downstream gene variant	["GGAAGAGAGAGA","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901674	downstream gene variant	["C","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901693	downstream gene variant	["A","AAG"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901693	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901725	downstream gene variant	["AG","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901774	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901776	downstream gene variant	["A","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901791	downstream gene variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74901807	downstream gene variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902145	downstream gene variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902209	downstream gene variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902261	downstream gene variant	["T","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902357	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902491	downstream gene variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902498	downstream gene variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902505	downstream gene variant	["T","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902542	downstream gene variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902569	downstream gene variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902570	downstream gene variant	["T","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902586	downstream gene variant	["C","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902592	downstream gene variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902625	downstream gene variant	["C","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902642	downstream gene variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902675	downstream gene variant	["T","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902677	downstream gene variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74902697	downstream gene variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903197	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903238	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903239	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903298	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903317	downstream gene variant	["T","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903489	downstream gene variant	["G","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903509	downstream gene variant	["A","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903519	downstream gene variant	["A","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903598	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903792	downstream gene variant	["A","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903799	downstream gene variant	["A","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903839	downstream gene variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903851	downstream gene variant	["T","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903855	downstream gene variant	["A","AT/C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74903977	downstream gene variant	["A","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74904138	downstream gene variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74904155	downstream gene variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74904209	downstream gene variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74904289	downstream gene variant	["G","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74904322	downstream gene variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74904500	downstream gene variant	["AGG","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74904729	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74904750	downstream gene variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74905070	3' prime UTR variant	["C","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74905074	3' prime UTR variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74905098	3' prime UTR variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74905271	synonymous variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74905330	missense variant	["A","G"], V105A	0	0	0	0	0	0	0	0	0	0	0	0
15:74905370	synonymous variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74905388	synonymous variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74905479	missense variant, splice region variant	["T","C"], D63G	0	0	0	0	0	0	0	0	0	0	0	0
15:74905537	intron variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74905674	intron variant	["C","CAGCATCCTGGTTT"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74905676	intron variant	["C","CAGCATCCTGGTTTAT"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906010	intron variant	["T","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906100	intron variant	["T","TTCCTGCTGAGCTAAATC"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906153	intron variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906362	intron variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906365	intron variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906688	intron variant	["G","C"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906783	intron variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906807	intron variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906836	intron variant	["C","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906872	intron variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906826	intron variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906931	intron variant	["C","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906938	intron variant	["G","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906944	intron variant	["A","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906953	intron variant	["A","G"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74906959	intron variant	["A","T"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74907128	intron variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74907169	intron variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74907170	intron variant	["G","A"]	0	0	0	0	0	0	0	0	0	0	0	0
15:74907220	intron variant	["T","T"]	0	0										