

Table 5. Molecular genetic characteristics of Ig L chain V region segments and CDR3 of TCL1 mice

Animal no.	Source	Isotype	V _L family	V _L gene	% difference from germ line	LCDR3 length	J _L gene	LCDR3			pI
TCL1-012	Spleen	κ	V _κ 4	V4-91	0.3%	13	J _κ 5	QQ	GSSILT	FGAG	5.5
TCL1-018	Spleen	κ	V _κ 3	V3-2	0%	12	J _κ 4	QH	NHGSLP	FT	7.17
			V _κ 8	V8-19	12.2%	10	J _κ 2	QQ	SF E VP	YT	3.62
TCL1-017	LN	κ	V _κ 8	V8-34	2.1%	10	J _κ 5	QQ	SYSA	PLT	5.5
			V _κ 5	V5-43	0.0%	10	J _κ 1	QQ	SNSWP	WT	5.5
			V _κ 3	V3-2	0.0%	10	J _κ 2	QQ	S K EVPY	T	6.13
			V _κ 3	V3-10	0.0%	10	J _κ 2	QQ	NN E DY	YT	3.32
			V _κ 14	V14-111	8.9%	11	J _κ 4	LQ	H G E H SP	FT	6.0
TCL1-015	Spleen	κ	V _κ 4	V4-59	0.6%	9	J _κ 1	HQ	WSSY	R T	9.05
			V _κ 3	V3-2	2.6%	10	J _κ 1	QQ	S K EVP	WT	6.13
			V _κ 6	V6-15	0.0%	10	J _κ 2	QQ	YNSYP	YT	5.5
			V _κ 14	V14-111	0.0%	10	J _κ 2	LQ	H G E S P	YT	5.14
			V _κ 4	V4-59	0.6%	9	J _κ 1	HQ	WSSY	R T	9.05
TCL1-016	Spleen	κ	V _κ 1	V1-110	1.4%	10	J _κ 1	SQ	ST H VP	QT	7.0
			V _κ 6	V6-15	0.0%	10	J _κ 1	QQ	YNSYP	LT	5.5
			V _κ 14	V14-111	0.0%	10	J _κ 4	LQ	H G E S P	FT	5.14

Animal no.	Source	Isotype	V _L family	V _L gene	% difference from germ line	LCDR3 length	J _L gene	LCDR3	pI
			V κ 19	V19-93	0.7%	10	J κ 2	LQ YD NLL YT	3.43
			V κ 4	V4-91	0.0%	11	J κ 4	QQ GSSIPN FT	5.5
			V κ 6	V6-23	4.1%	10	J κ 4	QQ YSNYP FT	5.5
TCL1-007	Spl & LN	κ	V κ 9	V9-124	0.0%	10	J κ 1	LQ YASYP VD	3.43
TCL1-018	Spl & LN	κ	V κ 4	V4-57	0.0%	10	J κ 5	QQ RSS YP LT	9.05
			V κ 6	V6-32	0.0%	10	J κ 2	QQ DYSSP LT	3.43
			V κ 3	V3-9	0.0%	10	J κ 1	MQ SM D P WT	3.32
			V κ 19	V19-93	0.7%	10	J κ 4	LQ YD NLL FT	3.43

Red letters indicate basic, positively charged amino acids; green letters indicate acidic, negatively charged amino acids.

Spl, spleen; LN, lymph node.