SUPPLEMENTARY TABLES

Supplementary Table I. Summary of phenotyping result of canine lymphoma samples.

Sample ID	Histological Classification ^a	CD20 (6C8)	CD21	CD22	CD3	CD5	CD4	CD8	CD45
DLYM-0902	DLBCL	++	++	++	-	-	-	-	+++
DLYM-1001	DLBCL	++	++	++	-	-	-	-	+++
DLYM-1002	DLBCL	+++	+++	+++	-	-	-	-	+++
DLYM-1003	DLBCL	+++	+++	+++	-	-	-	-	+++
DLYM-1004	DLBCL	+++	+++	+++	-	-	-	-	+++
DLYM-1005	DLBCL	+++	+++	+++	-	-	-	-	+++
DLYM-1007	MZL	+++	+++	+++	-	-	-	-	+++
DLYM-1009	DLBCL	+++	+++	+++	-	-	-	-	+++
DLYM-1012	DLBCL	+++	+++	+++	-	-	-	-	+++
DLYM-1013	DLBCL ^b	+++	+++	+++	-	-	-	-	+++
DLYM-1014	DLBCL	+	+	++	-	-	-	-	+++
DLYM-1015	BL	+++	++	++	-	-	-	-	+++
DLYM-1016	DLBCL	++	+	++	-	-	-	-	+++
DLYM-1017	DLBCL	+++	+++	+++	-	-	-	-	+++
DLYM-1018	DLBCL ^b	+++	+++	+	-	-	-	-	+++
DLYM-1101	BL	+++	+++	+++	-	-	-	-	+++
DLYM-1102	DLBCL	+++	+++	+++	-	-	-	-	+++
DLYM-1103	DLBCL	++	++	++	-	-	-	-	+++
DLYM-1104	MZL	++	++	++	-	-	-	-	+++
DLYM-1106	MZL	+	+	+	-	-	-	-	+++
DLYM-1107	DLBCL	+++	+++	+++	-	-	-	-	+++
DLYM-1108	DLBCL	+	+	+	-	-	-	-	+++
DLYM-0901	TZL	-	-	-	+	+	+	-	+++
DLYM-1006	PTCL	-	-	-	+	±	+	-	+++
DLYM-1008	TZL	-	+++ ^c	-	+	+	-	±	-
DLYM-1010	PTCL	-	-	-	+++	-	±	-	+++
DLYM-1011	TZL	-	+++ ^c	-	+++	++	-	-	-
DLYM-1019	TZL	-	+++ ^c	-	++	++	-	±	+++
DLYM-1105	PTCL	-	-	-	+	±	-	±	+++

The percentage of cells that express indicated CD molecule in total cells was determined using flow cytometry. -; <30%, \pm ; 30-70%, +; >70%, ++; >80%, +++; >90%.

 $[^]a \, DLBCL; \, diffuse \, large \, B-cell \, lymphoma, \, MZL; \, marginal-zone \, lymphoma, \, BL; \, Burkitt \, lymphoma, \, TZL; \, T-zone \, lymphoma, \, PTCL; \, peripheral \, T-cell \, lymphoma \, not \, otherwise \, specified.$

^b These samples were not clearly distinguishable from late-stage MZL.

^c The expression of CD21 (complement receptor 2 or CR2) was previously reported in canine TZL. [Frantz AM, Sarver AL, Ito D, et al. Molecular profiling reveals prognostically significant subtypes of canine lymphoma. Vet Pathol 2013;50:693–703.]

Supplementary Table II. Summary of direct cytotoxicity assay using primary canine B-cell lymphoma cells.

Sample ID	IgG1/No Ab	6C8/No Ab
DLYM-1004	102.2	112.4
DLYM-1005	99.1	101.8
DLYM-1016	99.3	102.6
DLYM-1103	99.6	117.4
Mean \pm S.E.M.	100.1 ± 0.7	108.5 ± 3.8^{d}

The percent of live cells normalized to the number of cells present at the end of the culture period in the conditions with no antibody treatment is shown.

Supplementary Table III. Summary of complement-dependent cytotoxicity using primary canine B-cell lymphoma cells.

Sample ID	IgG1+RC/RC	6C8+RC/RC
DLYM-1004	87.3	78.1
DLYM-1005	101.6	100.7
DLYM-1016	84.6	90.9
Mean \pm S.E.M.	91.1 ± 5.3	89.9 ± 6.5^{e}

The percent of live cells normalized to the number of cells present at the end of the culture period in the conditions with RC only treatment sample is shown.

^dThere were no statistically significant differences in cell viability in cells treated with IgG1 and 6C8 (two-tailed t test for paired samples).

^eThere were no statistically significant differences in cell viability in cells treated with IgG1 and 6C8 (two-tailed t test for paired samples).