

RESEARCH ARTICLE

Investigation of canine extracellular vesicles in diffuse large B-cell lymphomas

Marek Kulka^{1*}, Kieran Brennan², Margaret Mc Gee²

1 Department of Pathology and Veterinary Diagnostics, Institute of Veterinary Medicine, Warsaw University of Life Sciences, Warsaw, Poland, **2** UCD School of Biomolecular and Biomedical Science, Conway Institute, University College Dublin, Dublin, Ireland

* marek_kulka@sggw.edu.pl

Abstract

Diffuse large B-cell lymphomas (DLBCLs) are the most common lymphoproliferative diseases in dogs. DLBCL diagnosis to date has relied on histopathological analysis; however liquid biopsies have gained attention in recent years as a source of diagnostic and prognostic information. Liquid biopsies can be a source of circulating DNA, miRNA, circulating tumour cells or extracellular vesicles (EVs). In this study EVs were isolated from the plasma of healthy dogs, and dogs with lymphoma, and adenocarcinoma by iodixanol density gradient centrifugation. These EVs were positive for the EV markers CD63 and TSG101 and the pan-B cell markers CD79a, CD21, CD45, CD20. NTA analysis revealed that the DLBCL and adenocarcinoma dogs had elevated plasma EVs relative to the healthy dogs. Furthermore, the modal size of lymphoma EVs had decreased relative to healthy dogs while adenocarcinoma EVs were unchanged. This study demonstrates that the plasma EV population is altered in canine lymphoma patients in a manner similar to previous studies on human lymphomas. The similar changes to the EV population in dogs, together with the similar pathological features and treatment protocols in canine and human non-Hodgkin lymphomas would make dogs a good comparative model for studying the role of EVs in DLBCL development and progression.

OPEN ACCESS

Citation: Kulka M, Brennan K, Mc Gee M (2022) Investigation of canine extracellular vesicles in diffuse large B-cell lymphomas. PLoS ONE 17(9): e0274261. <https://doi.org/10.1371/journal.pone.0274261>

Editor: Girijesh Kumar Patel, Texas Tech University Health Science, Lubbock, UNITED STATES

Received: January 11, 2022

Accepted: August 24, 2022

Published: September 20, 2022

Copyright: © 2022 Kulka et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Data Availability Statement: All relevant data are within the paper and its [Supporting information files](#).

Funding: This work has been funded by National Science Centre (grant number MINIATURA 2019/03/X/NZ5/01168). The funder had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing interests: The authors have declared that no competing interests exist.

Introduction

Lymphoproliferative diseases are the most common neoplastic diseases in dogs [1]. The majority of canine lymphomas are diffuse large B-cell lymphomas (DLBCLs), characterized by a diffuse proliferation of large neoplastic lymphoid cells with a nuclear size more than twice the size of a red blood cell. There are two common types [2] the immunoblast and centroblastic variants, where the centroblastic variant is the most frequent and described as proliferation of large lymphoid cells similar to proliferating cells of the germinal center with oval to round vesicular nuclei (fine chromatin, multiple nucleoli) and small amount of basophilic cytoplasm. The immunoblastic variant [2], is more aggressive and described as round cells with centrally located nucleoli and a broad rim of basophilic cytoplasm [3].

Lymphomas initially present with enlargement of peripheral lymph nodes. When the internal lymph nodes are affected by the disease, then symptoms such as dyspnea, cough, and

fatigue may occur. Further development of DLBCL is associated with organ infiltration (liver, spleen, bone marrow). Patients may show non-regenerative or regenerative anemias and lymphocytosis or lymphopenia [2, 4]. The 'gold standard' treatment for canine lymphoma consists of a multi-agent, CHOP-based chemotherapy protocol (cyclophosphamide, doxorubicin, vincristine, and prednisone) [5, 6]. The similar pathological features and treatment protocols in canine and human non-Hodgkin lymphomas make dogs a good large comparative model in studying the DLBCLs [7]. Whereas the pathology of lymphomas is well recognized, the majority of current research is focused on early cancer diagnosis. Liquid biopsies are starting to play an important role in defining the first signs of lymphoproliferation. This minimally invasive approach can be used to monitor different components in blood such as circulating tumour DNA, miRNA and more recently extracellular vesicles (EVs) [8].

EVs are a heterogeneous group of nanometer-sized, membrane-bounded vesicles that play an important role in intercellular communications [9]. They originate from the inward budding of multivesicular endosomes (40–150 nm exosomes) or outward budding from the plasma membrane (>100 nm microvesicles or apoptotic bodies) [10, 11]. EVs are secreted by presumably all cell types and are found in most biofluids, including blood, serum, urine, and cerebral spinal fluid [12, 13]. EVs can transfer a wide range of nucleic acids, lipids, and soluble and membrane-associated proteins, and have been shown to play a pivotal role in normal physiology and disease, including maintenance of cellular homeostasis, regulation of gene transcription, activation and modulation of immune response and cancer progression [12–14].

Tumour-derived EVs have been reported to enhance cell proliferation, invasion, migration, and angiogenesis [15] and can influence drug resistance by transferring multidrug-resistant proteins, miRNAs, and exporting chemotherapeutic drugs [16, 17]. Several studies have examined exosome-derived RNA as a potential biomarker in DLBCL [18–25]. DLBCL-derived EVs can regulate several functions of natural killer cells [25] as well as macrophage polarization [26].

Currently, DLBCL diagnosis relies on tissue specimen examination, which is invasive and expensive. EV encapsulation can improve protein and miRNA stability in biofluids making EVs an attractive source of biomarkers for the development of non-invasive tests for the early diagnosis or follow-up and the prediction of treatment response [27–29]. Liquid biopsies could improve lymphoma management due to the non-invasive nature, and the ability to reflect spatial inter- and intra-tumour heterogeneity, and the possibility of repeated measurements through longitudinal profiling without the need for a tissue biopsy.

The aim of this preliminary study is to examine changes in the EV population in patients with DLBCL and how these changes in the EV population can be applied in the diagnosis of DLBCL.

Materials and methods

Animals, material collection

Blood was collected by certified veterinary surgeons during medical checkups or before planned surgery after obtaining verbal consent from the owner. Each patient was clinically evaluated. One group consisted of 14 healthy dogs with an age ranging from 1 year and 2 months to 7 years 3 months old. A second group consisted of 13 patients with an age ranging from 4 years to 15 years and 4 months that were diagnosed with diffuse large B-cell lymphoma (DLBCL) based on the lymph nodes fine needle biopsy results (cytological analysis and 12 patients by immunohistochemistry and 1 with PARR). During clinical examination patients presented with lymphadenopathy, 8 had cough and dyspnea. The third group consisted of two

patients aged 7–8 years that were diagnosed with adenocarcinoma, which presented with mammary tumours in one of their milk strips.

Blood samples were taken from the cephalic vein and collected into appropriate collection test tubes with EDTA-K2 and into clotting test tubes. Serum was aspirated after centrifugation of clotted samples. The whole blood was tested for CBC and serum for basic biochemical panels: alanine aminotransferase (ALT), aspartate aminotransferase (AST), alkaline phosphatase (AP), urea, creatinine, total protein (TP), albumins, globulins, glucose, lipase also blood smears were prepared stained with May-Grunwald Giemsa stain (according to the manufacturer's recommendations) and assessed. The plasma samples were separated from blood cells using Lymphoprep™ (Stem Cell Technologies). The plasma fraction from the Lymphoprep™ was centrifuged at 600×g at 4°C, for 10 min and then the supernatant was centrifuged at 2,000×g at 4°C, for 20 min. The platelet free plasma was stored at -80°C for the EV isolations.

Ultracentrifugation

All ultracentrifugations were performed using a Beckman Coulter Optima MAX-UP ultracentrifuge (stopping without break), with centrifugation durations based on a “50 nm cut-off size”, as described in Livshits et al., with an additional 5 min added to allow the rotor to come up to speed. Plasma samples were defrosted and transferred to a 13.5 ml Beckman Coulter ultracentrifuge tube (Prod. No. 355630) [30]. The plasma was diluted to 8.6 ml with particle-free PBS for ultracentrifugation. The tubes were centrifuged at 120000 g (RCF avg, 40800 rpm) for 2 hours at 20°C, using a Beckman Coulter MLA-55 rotor. The supernatant was removed, and the EV pellet was resuspended in 100 µl residual PBS.

Iodixanol density gradient centrifugation

Density gradient centrifugation was performed using a modified protocol from [31, 32]. A 54% iodixanol-PBS working solution was prepared by diluting a stock solution of OptiPrep™ (60% (w/v) aqueous iodixanol from Axis-Shield PoC, Norway) with 10x particle-free PBS (Gibco, Waltham, MA, USA). Iodixanol solutions (41% (w/v), 35% (w/v)), were prepared by diluting the 54% iodixanol-PBS working solution in 1x particle-free PBS (Gibco, Waltham, MA, USA). To form the gradient, firstly a homogenous 41% (w/v) base layer of the gradient (estimated density ~1.223 g/ml) was produced by adding 336 µl of the 54% iodixanol-PBS working solution to a 16 × 76 mm thinwall, ultra-Clear™ tube (Beckman Coulter), together with resuspended pellet. Next, 1.7 ml 35% (w/v) iodixanol (estimated density ~1.192 g/ml), and 1.7 ml 14% iodixanol (estimated density ~1.08 g/ml) were layered successively on top of the vesicle suspension. Centrifugation was performed at 219,373g (RCF avg, 55000 rpm) for 16 h at 20°C in Beckman Coulter Optima MAX-UP ultracentrifuge, using Beckman Coulter MLA-55 rotor (stopping without break). Fractions (~200 µl) were collected from the top of the tube and then 50 µl of each fraction was pipetted into a 96 well plate and the absorbance of the fractions was measured at 340 against an iodixanol standard curve to determine the fraction density. The fractions with densities between 1.08–1.19 g/ml were combined and diluted to a density <1.03 g/ml with particle-free PBS and the diluted fractions were centrifuged at 120000 g (RCF avg, 40800 rpm using Beckman Coulter MLA-55 rotor) for 2 hours at 4°C in Beckman Coulter Optima MAX-UP ultracentrifuge (stopping without break). The supernatant was removed, and the EV pellets were resuspended in 100 µl residual PBS.

NTA analysis

Particle number and size distribution in plasma samples was determined by nanoparticle tracking analysis (NTA) using a NanoSight LM10 (Malvern Panalytical, UK) configured with

a 405 nm laser and a high sensitivity scientific CMOS camera. Samples were diluted (plasma 20:400) in particle-free PBS (Gibco, Waltham, MA, USA). Samples were analyzed at 24°C and 15×60s videos were captured with a camera level of 15. Data was analyzed using NTA 3.2 Dev Build 3.2.16 software with a detection threshold of 5.

Western blot analysis

15 µg protein isolated from the serum EV fraction was separated by SDS-PAGE (10% polyacrylamide) according to a standard protocol. The polypeptides were electroblotted onto nitrocellulose membrane (Immobilon[®]-FL PVDF membrane) and probed with a primary antibody overnight at 4°C. Alix (1:200; Santa Cruz sc-53538), Tsg101 (1:500; Abcam ab83), CD63 (1:5000; ABIN 1440014), CD79a (1:500; ABIN 2472431), CD20 (1:2000; ABIN 2717455), CD21 (1:5000; ABIN 94032), CD45 (1 µg/ml; ABIN 6940460) antibodies were used as primary antibodies. Anti-goat (for CD63) or anti-mouse (for the rest) antibodies conjugated to horseradish peroxidase (Bio-Rad, Hercules, CA, USA) were used as secondary antibodies at a dilution of 1/5000 or 1/10000, respectively. Visualization was performed with a chemiluminescent reagent system (Clarity[™]Western ECL, Bio-Rad, Hercules, CA, USA). Blots were quantified by densitometry using Quantity One 4.6.2 software (Bio-Rad Ltd.).

Statistical analysis

All groups were checked for normal distribution and for comparisons an unpaired T-test was used. *P* values were considered significant when < 0.05.

Results

All blood morphological and biochemical parameters were within reference values [33] with no significant difference between healthy control dogs and dogs with adenocarcinoma. Within the DLBCL group 2 patients displayed normocytic normochromic non regenerative anemia, and 3 displayed mild lymphopenia. Biochemical parameters were within normal limits, with the one exception where two dogs displayed a mild increase in ALT.

Plasma was prepared by depletion of cells and platelets by centrifugation. Plasma EVs were isolated by ultracentrifugation and iodixanol density gradient ultracentrifugation. The EVs were resuspended in PBS and particle size and size distribution of the canine EVs were examined by NTA analysis (Fig 1A). DLBCL patients had higher EV concentrations than healthy controls, with DLBCL patient plasma having $5.19 \pm 3 \times 10^8$ particles / ml and healthy control plasma having $2.86 \pm 1.8 \times 10^8$ particles / ml (*P* = 0.019) (Fig 1B). DLBCLs EVs also had a smaller modal size (129.7 ± 15.6 nm) when compared to healthy controls (159.9 ± 22.6 nm) (*P* = 0.0013) (Fig 1C). Adenocarcinoma patients had higher EV concentration ($6.08 \pm 5.6 \times 10^8$ particles / ml) than healthy controls (*P* = 0.027), however, no change in modal size was observed.

Western blot analysis was performed using an equal amount of protein from each serum EV sample. The EVs markers; Tsg101 and CD63, and the lymphoma markers; CD45, CD79a, CD21, were detected in all EV samples. However, only one DLBCL patient was positive for CD20 in EVs (Fig 2).

15 µg protein from 6 lymphoma (L1-6) and 6 control (C1-6) serum EV samples were analyzed alongside a molecular weight marker (Bio-Rad # 161-0375) by SDS-PAGE and western blot analysis. The membranes were probed with anti-TSG101 (49 kDa, 1:500; Abcam ab83), anti-CD63 (40–60 kDa, 1:5000; ABIN 1440014), anti-CD79a (44–50 kDa, 1:500; ABIN 2472431), anti-CD20 (30–50 kDa, 1:2000; ABIN 2717455), anti-CD21 (145 kDa, ABIN 94032), anti-CD45 (180–220 kDa, 1 µg/ml; ABIN 6940460). The EVs markers TSG 101, CD63 and

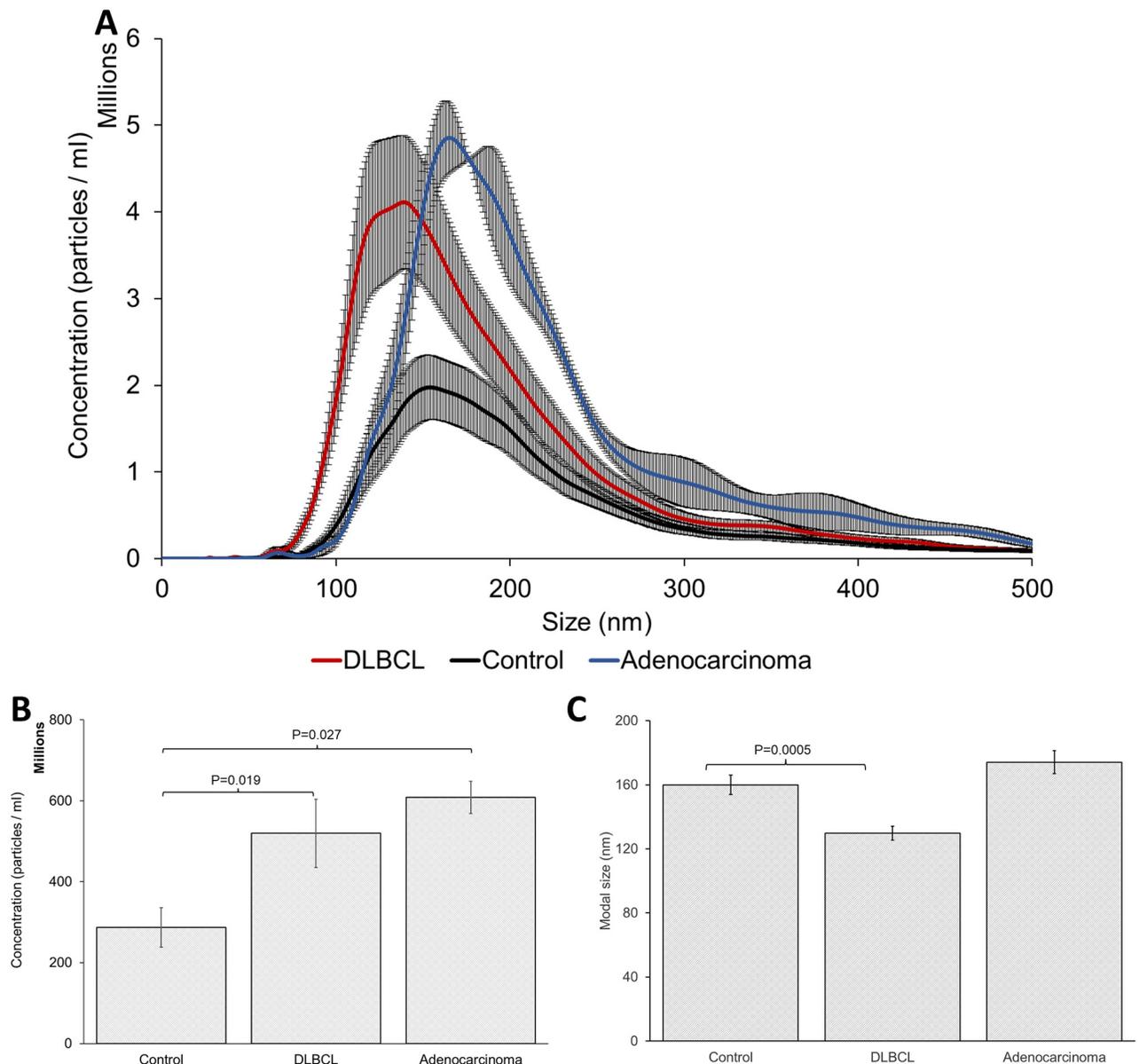


Fig 1. NanoSight analysis of particle size distribution of EVs isolated from canine plasma. (A) Line graph of healthy control, DLBCL and adenocarcinoma patient particle size distribution. (B) Bar graph of total EV concentration of particles / ml and (C) modal size of healthy control, DLBCL and adenocarcinoma patient EV samples.

<https://doi.org/10.1371/journal.pone.0274261.g001>

lymphoma markers CD45, CD79a, CD21 were detected in patients with DLBCLs (L) and healthy dogs (C), CD20 was detected in one patient with DLBCL.

Discussion

EVs play a role in cell-to-cell communication, promotion of metastasis and immune suppression. EVs have gained a lot of interest as a source of diagnostic and prognostic biomarkers for human cancer research and are becoming more important in veterinary medicine. EVs are a promising option for lymphoma monitoring as the EV content can reflect the cell from which they originate and they are readily accessible in biofluids without the need for tissue biopsy.

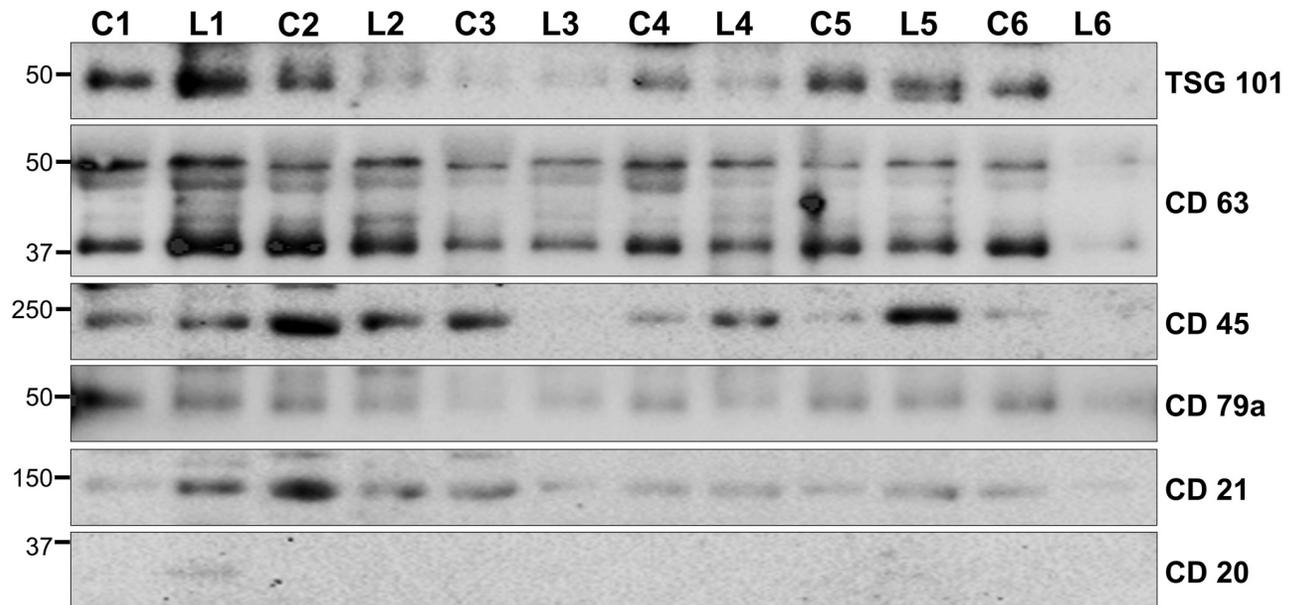


Fig 2. Western blot analysis of protein lysates from isolated from canine plasma.

<https://doi.org/10.1371/journal.pone.0274261.g002>

Canine DLBCL patients were found to have an elevated concentration of serum EVs relative to healthy patients. A similar observation has been made in human hematological neoplastic disorders, with elevated plasma EV levels observed in multiple myeloma, Hodgkin's lymphoma and to a lesser degree in non-Hodgkin's lymphoma and chronic lymphocytic leukemia [34]. A higher concentration of EVs was also observed in dogs with adenocarcinoma, which confirmed studies made on canine and feline mammary tumour cell lines [35]. In addition to increased EV concentration, DLBCL patients had a smaller EV modal size relative to healthy dogs, which was comparable to the decreased EV modal size observed in human hematological cancers [34]. It is not possible to distinguish whether this increase in smaller EVs is due to an upregulation of the exosome biogenesis pathways or whether it is due to increased small microvesicle budding from the plasma membrane [36] as specific markers to distinguish these two pathways do not currently exist. All samples were positive for the EV membrane marker CD63 and the intra-EV marker TSG101 [37]. DLBCL has no specific immunophenotype for either canine or human cancers; however, they express pan-B markers including CD79a, CD21, CD45, CD20 [1]. In this study, the DLBCL patient EVs were positive for CD79a, CD21, and CD45, however, only one EV sample was positive for CD20, suggesting some diversity within the DLBCL group. This heterogeneity in B-cell expression profile may help diagnostic approaches for DLBCL patients in the future however, a larger study would be needed to assess the diagnostic significance of CD20 in DLBCL plasma EVs. The presence of these lymphoma markers in control patients indicates that EVs from normal immune cells are also present in the plasma EV population. Furthermore, the lack of a strong correlation between the NTA data and the western blot results suggests that alternative protein markers will need to be examined to determine if it is possible to identify the population of EVs that is increasing in lymphoma patients.

In conclusion, we have found that canine DLBCL patients have elevated levels of plasma EVs, that express lymphocyte markers. These EVs had a smaller modal size relative to healthy controls suggesting that not all EV sizes were increased, but instead the production of small EVs was upregulated. This observation is similar to previous studies on human lymphomas [7,

8, 18]. The fact that canine and human DLBCL patients share similar pathological features and treatment protocols suggests they may also share the same EV signaling pathways, making dogs a good large comparative model to study DLBCL.

Supporting information

S1 File. Supplemental western blot figure.

(PPTX)

S2 File. Supplementary NTA figure.

(XLS)

Author Contributions

Conceptualization: Marek Kulka, Kieran Brennan, Margaret Mc Gee.

Data curation: Marek Kulka.

Formal analysis: Marek Kulka, Kieran Brennan, Margaret Mc Gee.

Funding acquisition: Marek Kulka, Margaret Mc Gee.

Investigation: Marek Kulka, Kieran Brennan.

Methodology: Marek Kulka, Kieran Brennan.

Project administration: Marek Kulka.

Resources: Marek Kulka.

Software: Marek Kulka.

Supervision: Marek Kulka, Margaret Mc Gee.

Validation: Marek Kulka.

Visualization: Marek Kulka.

Writing – original draft: Marek Kulka, Kieran Brennan.

Writing – review & editing: Marek Kulka, Kieran Brennan, Margaret Mc Gee.

References

1. Seelig DM, Avery AC, Ehrhart EJ, Linden MA. The Comparative Diagnostic Features of Canine and Human Lymphoma. *Vet Sci*. 2016; 3(2): 11. <https://doi.org/10.3390/vetsci3020011> PMID: 28435836
2. Aresu L. Canine Lymphoma, More Than a Morphological Diagnosis: What We Have Learned about Diffuse Large B-Cell Lymphoma. *Front Vet Sci*. 2016; 3: 77. <https://doi.org/10.3389/fvets.2016.00077> PMID: 27630997
3. Xie Y, Pittaluga S, Jaffe ES. The histological classification of diffuse large B-cell lymphomas. *Semin Hematol*. 2015; 52(2): 57–66. <https://doi.org/10.1053/j.seminhematol.2015.01.006> PMID: 25805585
4. Henriques J, Felisberto R, Constantino-Casas F, Cabeçadas J, Dobson J. Peripheral blood cell ratios as prognostic factors in canine diffuse large B-cell lymphoma treated with CHOP protocol. *Vet Comp Oncol*. 2021; 19(2): 242–252. <https://doi.org/10.1111/vco.12668> PMID: 33247533
5. Comazzi S, Marconato L, Argyle DJ, Aresu L, Stirn M, Grant IA, et al. European Canine Lymphoma Network. The European canine lymphoma network: a joining initiative to generate consensus guidelines for the diagnosis and therapy in canine lymphoma and research partnership. *Vet Comp Oncol*. 2015; 13(4): 494–7. <https://doi.org/10.1111/vco.12128> PMID: 26463403
6. Benjamin SE, Sorenmo KU, Krick EL, Salah P, Walsh KA, Weinstein NM, et al. Response-based modification of CHOP chemotherapy for canine B-cell lymphoma. *Vet Comp Oncol*. 2021; 19(3): 541–550. <https://doi.org/10.1111/vco.12693> PMID: 33729654

7. Ito D, Frantz AM, Modiano JF. Canine lymphoma as a comparative model for human non-Hodgkin lymphoma: recent progress and applications. *Vet Immunol Immunopathol.* 2014; 159(3–4): 192–201. <https://doi.org/10.1016/j.vetimm.2014.02.016> PMID: 24642290
8. Garnica TK, Lesbon JCC, Ávila ACFCM, Rochetti AL, Matiz ORS, Ribeiro RCS, et al. Liquid biopsy based on small extracellular vesicles predicts chemotherapy response of canine multicentric lymphomas. *Sci Rep.* 2020; 10(1): 20371. <https://doi.org/10.1038/s41598-020-77366-7> PMID: 33230132
9. Cappariello A, Loftus A, Muraca M, Maurizi A, Rucci N, Teti A. Osteoblast-Derived Extracellular Vesicles Are Biological Tools for the Delivery of Active Molecules to Bone. *J Bone Miner Res.* 2018; 33(3): 517–533. <https://doi.org/10.1002/jbmr.3332> PMID: 29091316
10. Théry C, Amigorena S, Raposo G, Clayton A. Isolation and characterization of exosomes from cell culture supernatants and biological fluids. *Curr Protoc Cell Biol.* 2006; Chapter 3:Unit 3.22. <https://doi.org/10.1002/0471143030.cb0322s30> PMID: 18228490
11. Nishimura T, Oyama T, Hu HT, Fujioka T, Hanawa-Suetsugu K, Ikeda K, et al. Filopodium-derived vesicles produced by MIM enhance the migration of recipient cells. *Dev Cell.* 2021; 56(6): 842–859.e8. <https://doi.org/10.1016/j.devcel.2021.02.029> PMID: 33756122
12. Schorey JS, Bhatnagar S. Exosome function: from tumor immunology to pathogen biology. *Traffic.* 2008; 9(6):871–81. <https://doi.org/10.1111/j.1600-0854.2008.00734.x> PMID: 18331451
13. Rajagopal C, Harikumar KB. The Origin and Functions of Exosomes in Cancer. *Front Oncol.* 2018; 8:66. <https://doi.org/10.3389/fonc.2018.00066> PMID: 29616188
14. Kalluri R, LeBleu VS. The biology, function, and biomedical applications of exosomes. *Science.* 2020; 367(6478):eaau6977. <https://doi.org/10.1126/science.aau6977> PMID: 32029601
15. Chen X, Yang F, Zhang T, Wang W, Xi W, Li Y, et al. MiR-9 promotes tumorigenesis and angiogenesis and is activated by MYC and OCT4 in human glioma. *J Exp Clin Cancer Res.* 2019; 38(1): 99. <https://doi.org/10.1186/s13046-019-1078-2> PMID: 30795814
16. Zhang X, Yuan X, Shi H, Wu L, Qian H, Xu W. Exosomes in cancer: small particle, big player. *J Hematol Oncol.* 2015; 8: 83. <https://doi.org/10.1186/s13045-015-0181-x> PMID: 26156517
17. Litwińska Z, Łuczkowska K, Machaliński B. Extracellular vesicles in hematological malignancies. *Leuk Lymphoma.* 2019; 60(1): 29–36. <https://doi.org/10.1080/10428194.2018.1459606> PMID: 29745272
18. Caivano A, La Rocca F, Simeon V, Girasole M, Dinarelli S, Laurenzana I, et al. MicroRNA-155 in serum-derived extracellular vesicles as a potential biomarker for hematologic malignancies—a short report. *Cell Oncol (Dordr).* 2017; 40(1):97–103. <https://doi.org/10.1007/s13402-016-0300-x> PMID: 27761889
19. Provencio M, Rodríguez M, Cantos B, Sabín P, Quero C, García-Arroyo FR, et al. mRNA in exosomes as a liquid biopsy in non-Hodgkin Lymphoma: a multicentric study by the Spanish Lymphoma Oncology Group. *Oncotarget.* 2017; 8(31):50949–50957. <https://doi.org/10.18632/oncotarget.16435> PMID: 28881619
20. Di C, Zhang Q, Wang Y, Wang F, Chen Y, Gan L, et al. Exosomes as drug carriers for clinical application. *Artif Cells Nanomed Biotechnol.* 2018; 46(sup3):S564–S570. <https://doi.org/10.1080/21691401.2018.1501381> PMID: 30431368
21. Rutherford SC, Fachel AA, Li S, Sawh S, Muley A, Ishii J, et al. Extracellular vesicles in DLBCL provide abundant clues to aberrant transcriptional programming and genomic alterations. *Blood.* 2018; 132(7): e13–e23. <https://doi.org/10.1182/blood-2017-12-821843> PMID: 29967128
22. Feng Y, Zhong M, Zeng S, Wang L, Liu P, Xiao X, et al. Exosome-derived miRNAs as predictive biomarkers for diffuse large B-cell lymphoma chemotherapy resistance. *Epigenomics.* 2019; 11(1):35–51. <https://doi.org/10.2217/epi-2018-0123> PMID: 30211623
23. Ting CY, Liew SM, Price A, Gan GG, Bee-Lan Ong D, Tan SY, et al. Clinical significance of aberrant microRNAs expression in predicting disease relapse/refractoriness to treatment in diffuse large B-cell lymphoma: A meta-analysis. *Crit Rev Oncol Hematol.* 2019; 144: 102818. <https://doi.org/10.1016/j.critrevonc.2019.102818> PMID: 31733445
24. Xiao XB, Gu Y, Sun DL, Ding LY, Yuan XG, Jiang HW, et al. Effect of rituximab combined with chemotherapy on the expression of serum exosome miR-451a in patients with diffuse large b-cell lymphoma. *Eur Rev Med Pharmacol Sci.* 2019; 23(4): 1620–1625. https://doi.org/10.26355/eurev_201902_17121 PMID: 30840285
25. Zare N, Haghjooy Javanmard S, Mehrzad V, Eskandari N, Kefayat A. Evaluation of exosomal miR-155, let-7g and let-7i levels as a potential noninvasive biomarker among refractory/relapsed patients, responsive patients and patients receiving R-CHOP. *Leuk Lymphoma.* 2019; 60(8): 1877–1889. <https://doi.org/10.1080/10428194.2018.1563692> PMID: 30714442
26. Liu W, Zhu M, Wang H, Wang W, Lu Y. Diffuse large B cell lymphoma-derived extracellular vesicles educate macrophages to promote tumours progression by increasing PGC-1 β . *Scand J Immunol.* 2020; 91(2): e12841. <https://doi.org/10.1111/sji.12841> PMID: 31833575

27. Mitchell PJ, Welton J, Staffurth J, Court J, Mason MD, Tabi Z, et al. Can urinary exosomes act as treatment response markers in prostate cancer? *J Transl Med.* 2009; 7: 4. <https://doi.org/10.1186/1479-5876-7-4> PMID: 19138409
28. Valadi H, Ekström K, Bossios A, Sjöstrand M, Lee JJ, Lötvall JO. Exosome-mediated transfer of mRNAs and microRNAs is a novel mechanism of genetic exchange between cells. *Nat Cell Biol.* 2007; 9(6): 654–9. <https://doi.org/10.1038/ncb1596> PMID: 17486113
29. Kosaka N, Iguchi H, Ochiya T. Circulating microRNA in body fluid: a new potential biomarker for cancer diagnosis and prognosis. *Cancer Sci.* 2010; 101(10): 2087–92. <https://doi.org/10.1111/j.1349-7006.2010.01650.x> PMID: 20624164
30. Livshits MA, Khomyakova E, Evtushenko EG, Lazarev VN, Kulemin NA, Semina SE, et al. Isolation of exosomes by differential centrifugation: Theoretical analysis of a commonly used protocol. *Sci Rep.* 2015; 5: 17319. <https://doi.org/10.1038/srep17319> PMID: 26616523
31. Parolini I, Federici C, Raggi C, Lugini L, Palleschi S, De Milito A, et al. Microenvironmental pH is a key factor for exosome traffic in tumor cells. *J Biol Chem.* 2009; 284(49): 34211–22. <https://doi.org/10.1074/jbc.M109.041152> PMID: 19801663
32. Brennan K, Martin K, FitzGerald SP, O'Sullivan J, Wu Y, Blanco A, et al. A comparison of methods for the isolation and separation of extracellular vesicles from protein and lipid particles in human serum. *Sci Rep.* 2020; 10(1): 1039. <https://doi.org/10.1038/s41598-020-57497-7> PMID: 31974468
33. Harvey J. *Veterinary Hematology: A Diagnostic Guide and Color Atlas.* 1st ed. Elsevier Saunders; 2012.
34. Caivano A, Laurenzana I, De Luca L, La Rocca F, Simeon V, Trino S et al. High serum levels of extracellular vesicles expressing malignancy-related markers are released in patients with various types of hematological neoplastic disorders. *Tumour Biol.* 2015; 36(12): 9739–52. <https://doi.org/10.1007/s13277-015-3741-3> PMID: 26156801
35. Sammarco A, Finesso G, Cavicchioli L, Ferro S, Caicci F, Zanetti R, et al. Preliminary investigation of extracellular vesicles in mammary cancer of dogs and cats: Identification and characterization. *Vet Comp Oncol.* 2018; 16(4): 489–496. <https://doi.org/10.1111/vco.12405> PMID: 29851284
36. Paulaitis M, Agarwal K, Nana-Sinkam P. Dynamic Scaling of Exosome Sizes. *Langmuir.* 2018; 34(32): 9387–9393. <https://doi.org/10.1021/acs.langmuir.7b04080> PMID: 29542322
37. Théry C, Witwer KW, Aikawa E, Alcaraz MJ, Anderson JD, Andriantsitohaina R, et al. Minimal information for studies of extracellular vesicles 2018 (MISEV2018): a position statement of the International Society for Extracellular Vesicles and update of the MISEV2014 guidelines. *J Extracell Vesicles.* 2018; 7(1): 1535750. <https://doi.org/10.1080/20013078.2018.1535750> PMID: 30637094