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Abstract: The term 'whole-slide imaging (WSI)' refers to the use of robotic microscopes for computerising entire slides into digital images. This study aimed to assess the agreement between WSI and optical microscopy for evaluating canine lymphoma cytological samples. Forty-four slides were computerised using a WSI scanner. The digital and glass slides were examined by three observers with different levels of expertise. The morphology of neoplastic cells and the lymphoma grade were scored, on the basis of the updated Kiel classification, and the intraobserver agreement was assessed. Moreover, the accuracy of determining the lymphoma grade of the digital and glass slides based on the results of flow-cytometry (FC) was established.

The overall intra-observer agreement for the cytomorphological features was fair to moderate (from 0.34 to 0.52) for the three observers, whereas the intra-observer agreement for the evaluation of malignancy grade was moderate (from 0.44 to 0.53). The diagnostic agreement between FC and digital slides was slight (0.16) for the inexperienced observer, fair (0.32) for the mildly experienced observer, and moderate (0.50) for the experienced observer, whereas the diagnostic agreement between FC and glass slides was fair (0.37) for the inexperienced observer, substantial (0.63) for the mildly experienced observer, and moderate (0.50) for the experienced observer. Our findings underline the importance of observer experience in determining the malignancy grade, especially if digital slides are used. The study also identifies some technical limitations of the tested WSI scanner, mainly linked to image quality, which affected the morphological evaluation of the neoplastic cells.

## 1 Original Article

Whole-slide imaging: cytomorphological descriptive capacity and intra-observer agreement in
 canine lymphoma samples

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## 19 Abstract

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39	morphological evaluation of the neoplastic cells.	
40	Keywords: Whole-slide imaging; Telepathology; Cytology; Lymphoma; Dogs.	

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imaging (WSI)' is referred to robotic microscopes capable to computerize entire slides into digital images. The aim of this study is to assess the agreement between WSI and optical microscopy in the evaluation of canin lymphoma cytological samples. Fortyfour slides were computerize using a WSI scanner. Digital and optical slides were examined by three observers with different levels of expertise. All the morphological features of the neoplastic cells, grading, and phenotype were scored, based on the Kiel-updated classification, and the intra-observer agreement was assessed. Moreover, the accuracy in determination of grading and immunophenotype of digital and optical slides based of the results of flow-cytometry (FC) was established.

The overall intra-observer agreement for the cytomorphological features was fair for all the observers (from 0.24 to 0.37), whereas the overall intra-observer agreement in the evaluation of grading and immunophenotype was fair to moderate (from 0.31 to 0.54). The diagnostic agreement between FC and digital slides was 0.33 for observer one, 0.63 for observer two, 0.47 for observer three, whereas the diagnostic agreement between FC and optical slides was 0.24 for observer one, 0.77 for observer two, 0.73 for observer three. Our study underlines the possibility of technical limitations of the WSI scanner tested, mainly linked to the quality of the images, especially for less experienced observers.

## 42 Introduction

43	Digital pathology is a branch of pathology in which images are visualised on a computer
44	monitor rather than directly through a microscope (Weinstein, 1986). Telepathology has evolved
45	from the transmission of static images captured using microscope-mounted cameras to the use of
46	robotic microscopes controlled by pathologists at distant sites and, more recently, to whole-slide
47	imaging (WSI) (Webster and Dunstan, 2014). A WSI scanner is a robotic microscope capable of
48	digitising an entire glass slide (GS) by using a software to merge the individually captured images
49	into a composite digital image (Pantanowitz et al., 2013). The digital images can be viewed either
50	on the computer used to scan the slides, by using a specific software designed to emulate a light
51	microscope (Webster and Dunstan, 2014), or at a remote site via a high-speed internet connection
52	(Steinberg and Ali, 2001). WSI maintains the relative simplicity of static-image transfer and
53	eliminates its limitations, by making available the entire specimen for review (Wilbur, 2011) and
54	allowing the magnification of the digital slide (DS) to be changed (Wilbur et al., 2009). Moreover,
55	the most updated scanners have the capability to perform multiple line scans of the same area at
56	different fields of focus (Webster and Dunstan, 2014). This function, the so-called 'z-stack' mode,
57	is most important in acquiring images of cytological specimens, in which cells are often arranged in
58	multiple layers (El-Gabry et al., 2014).
59	
60	In human medicine, WSI is used for several reasons, mainly digital diagnostics and
61	teleconsultation (Al-Janabi et al., 2012). In veterinary medicine, despite the increased use of WSI
62	instruments in reference laboratories, few reports on digital pathology are present in the literature.
63	To the authors' knowledge, no papers focusing on the morphological descriptive capability of WSI
64	have been published. Moreover, the use of WSI has never been validated, and its capability to

65 replace the conventional microscope has not been assessed using cytological samples.

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67	One of the most commonly obtained cytological samples in canine medicine is the fine
68	needle aspirate (FNA) of lymph nodes. This technique is used to discriminate between
69	inflammatory and neoplastic disorders, among which, non-Hodgkin's lymphoma is the most
70	common primary neoplasia affecting the lymph node (Richards and Suter, 2015). The identification
71	of a clonal expansion of neoplastic lymphocytes is essential for lymphoma diagnosis. Histology,
72	immunocytochemistry, immunohistochemistry, flow cytometry (FC) and PCR to detect clonal
73	antigen receptor gene rearrangement (PARR) are fundamental tools that can be used to confirm the
74	diagnosis of lymphoma and to identify the phenotype for prognostic purposes (Burkhard and
75	Bienzle, 2013). These diagnostic procedures can be performed only in specialized laboratories.
76	Commonly, the cytological examination of lymph node samples obtained via FNA is the first step
77	in the diagnosis of lymphoproliferative diseases in dogs, because it is minimally invasive,
78	inexpensive, and fast (Amores-Fuster et al., 2015). Moreover, the cytological features are used to
79	classify the different lymphoma subtypes with prognostic significance (Ponce et al., 2004).
80	However, the morphological evaluation of neoplastic lymphocytes is complex, and the opinion of a
81	skilled clinical pathologist is often required for the correct diagnosis and classification of the
82	different lymphoma subtypes.
83	
84	The possibility to use WSI instruments could improve the quality of cytological services,
85	and could provide the possibility to share cytological samples of canine lymphomas with more
86	experienced clinical pathologists. However, before this technology can be applied in veterinary
87	cytology, the reliability of cytological evaluation and the technical aspects of the scanning process
88	must be evaluated. The aims of this study were to assess the following: 1) the intra-observer
89	agreement (IOA) between WSI (D-sight, A. Menarini Diagnostics S.r.l) and optical microscopy in
90	the evaluation of cellular morphology in canine lymphoma samples, by using the updated Kiel
91	classification (Ponce et al. 2010); 2) the IOA between WSI and optical microscopy in the
92	assessment of lymphoma grading; and 3) whether the accuracy of grading assessment varied

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93	between WSI scanner and optical microscopy, by using FC as a reference method. Moreover, the
94	influence of the level of observer expertise on their performance was evaluated.
95	
96	Materials and methods
97	Cytological samples and observers
98	The database of the Flow Cytometry Service of the Department of Veterinary Medicine
99	(University of Milan, Milan, Italy) was searched to select consecutive canine lymphoma samples
100	diagnosed on the basis of clinical, clinico-pathological, cytological, and FC data, from January 2015
101	to June 2015. Only cases with good-quality lymph node cytological smears were enrolled in the
102	study, to allow for a detailed evaluation of the morphological features of the neoplastic cells. FC
103	analysis and the criteria for lymphoma diagnosis were applied as previously described (Gelain et al.,
104	2008). The following antibodies were used: CD45 (clone YKIX716.13, Serotec), CD3 (clone
105	CA17.2A12, Serotec), CD5 (clone YKIX322.3, Serotec), CD4 (clone YKIX302.9, Serotec), CD8
106	(clone YCATE55.9, Serotec), CD21 (clone CA21D6, Serotec), CD79a (clone MCA1298F,
107	Serotec), and CD34 (clone 1H6, Pharmingen, BD Bioscience). Neoplastic cells were identified on
108	the morphological cytogram (forward scatter [FSC] versus side scatter) or on the CD45 versus FSC
109	cytogram. The percentage of neoplastic cells, the phenotype, and the mean FSC (mean cellular size)
110	were used to identify the cell type (B-cell or T-cell) and cell size. GS were stained with May-
111	Grünwald-Giemsa stain, and for each case, the slide with higher cellularity and better preservation
112	was selected.
113	
114	Three observers with different levels of expertise participated in the study, and all were
115	blinded to the FC results. The inexperienced observer, a PhD student, had the lower experience in
116	cytological evaluation; the mildly experienced observer, a postdoctoral researcher, had intermediate
117	experience in cytological evaluation; and the experienced observer, a board-certified clinical-

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118	pathologist, l	nad extensi	ve experient	ce in cyt	ological	evaluation.	None of t	he observers had

- 119 previously used the WSI technology for routine diagnostic procedures.
- 120
- 121 Digital slides

122	All the GSs were scanned using the $40 \times$ objective with the z-stack modality by using a WSI
123	scanner (D-sight, A. Menarini Diagnostics S.r.l.) to obtain the DSs. The DSs were scanned using
124	automated tissue detection and focus-point assignments with seven-line scans of the same area at
125	different fields of focus. The research case numbers assigned to the DSs were different from those
126	assigned to the corresponding GSs to minimize recall bias. DSs were subsequently uploaded to a
127	server to be evaluated by the three observers using an online software (Telepathology, Visia
128	Imaging S.r.l.) (Fig. 1). The monitors used ranged from 14 to 15.6 inches, with a screen resolution
129	of at least 1366×768 pixels, and no special monitor or setting was used. Each observer used his own
130	laptop's screen to evaluate all the DSs. Between the evaluation of GS and DS by the same observer,
131	a wash-out period of at least one month was imposed, to ensure the observers did not remember the
132	cases from the previous viewings.
133	
134	Morphological features evaluated
135	According to the updated Kiel classification (Fournel-Fleury et al., 1997; Ponce et al., 2010)
136	the following parameters were recorded for both the DSs and GSs by each observer at 40×
137	magnification: pleomorphism of the neoplastic population, cellular size, amount and colour of
138	cytoplasm, nuclear shape, nuclear chromatin pattern, and nucleoli. Specifically, the following
139	features were evaluated:
140	a. The cellular size was based on the comparison between the red blood cells and nucleus of
141	the cells, and was defined as small, medium, or large depending on whether it was smaller,
142	equal to, or larger than two erythrocytes, respectively.

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**Comment [FB84]:** Line 126 of YTVJL-D17-00838R2 Deleted – digital slides ()

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143	b. Pleomorphism of the neoplastic population was recorded if the cellular size and shape varied	_
144	among the neoplastic population.	
145	c. Cytoplasm amount, distribution and colour: the cytoplasm was described as scarce—when	/
146	only a scant amount of cytoplasm was present moderate, or abundant. The presence of	
147	unipolar distribution was also recorded. The colour of the cytoplasm was identified as clear,	$\left \right\rangle$
148	basophilic or deeply basophilic.	
149	d. Nuclear shape and nuclear chromatin pattern: round, indented or convoluted and irregular	
150	shape of the nucleus was recorded, and the chromatin was described as dense, granular or	
151	finely granular, or smooth.	
152	e. Presence of nucleoli: the presence of single or multiple prominent nucleoli was recorded.	
153	Before the beginning of the study, the observers conferred to standardize the appearance of all	
154	these morphological features.	
155	Moreover the observers were asked to identify the malignancy grade (high or low grade).	/
156	Small-cell lymphomas were classified as low-grade lymphomas while large-cell and	
157	pleomorphic mixed-small-large-cell lymphomas were classified as high-grade lymphomas.	
158	All these morphological features were scored as shown in Table 1.	
159		
160	Statistical methods	
161	In all the cases, the IOA between the DSs and GSs was assessed for each of the	
162	morphological features and the grading. Moreover, the agreement between FC and DS and between	
163	FC and GS in determining the malignancy grade was also evaluated for each observer to evaluate	
164	the accuracy of the DS and GS, respectively, by using FC as a reference method, for determining of	
165	the lymphoma grade. The IOA and the agreement in determining the malignancy grade were	
166	assessed by using linearly weighted Cohen's K. The K coefficients were interpreted as	/
167	recommended by Landis and Koch (1977): <0.00, poor; 0.00-0.20, slight; 0.21-0.40, fair; 0.41-0.60,	

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Comment [FB103]: Line 150 of YTVJL-D17-00838R2 Deleted - whereas

Comment [FB104]: Line 151 of YTVJL-D17-00838R2 Deleted - and

Comment [FB105]: Lines 153-154 c YTVJL-D17-00838R2 Deleted - based on the morphological criteria described in literature (Fournel-Fleury et al., 1997; Ponce et al., 2010),

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168	moderate; 0.61-0.80, substantial; and >0.80, almost perfect. Differences in the number of corrected		
169	classified lymphomas obtained by evaluating the DSs and GSs, compared to the results of FC, were		
170	assessed using Fisher's exact test. Statistical analysis was performed using a commercially available		
171	software program (MedCalc Statistical Software version 15.8, MedCalc Software bvba,).	_	Comment [FB122]: Line 176 of YTVJL-D17-00838R2
172			Deleted - (SPSS, version 12.0.0, SPSS Inc, Chicago, IL, USA).
173	Results		
174	In total 250 samples arrived to the Flow Cytometry Service of the Department of Veterinary		
175	Medicine, University of Milan from January to June 2015 and 163 of these (65%) also had		
176	corresponding cytological slides. In 44 out of the 163 (27%) samples, the cytological smears had		
177	good cellularity and preservation and were hence included in the present study. Based on the		
178	phenotype and size of cells assessed using FC, 24 high-grade B-cell lymphomas, four low-grade B-		Comment [FB123]: Line 179 of YTVJL-D17-00838R2
179	cell lymphomas, 10 high-grade T-cell lymphomas, and six low-grade T-cell lymphomas were		Deleted - the
			Comment [FB124]: Line 179 of
180	included in the study (Table 2).		YTVJL-D17-00838R2 Deleted - by
180 181	included in the study (1 able 2).		
	The IOA results for the cytomorphological features are listed in Table 3. For the		Deleted - by Comment [FB125]: Line 183 of
181			Deleted - by
181 182	The IOA results for the cytomorphological features are listed in Table 3. For the		Deleted - by <b>Comment [FB125]:</b> Line 183 of YTVJL-D17-00838R2
181 182 183	The IOA results for the cytomorphological features are listed in Table 3. For the inexperienced observer the agreement between the DSs and GSs was slight for the amount and		Deleted - by <b>Comment [FB125]:</b> Line 183 of YTVJL-D17-00838R2
181 182 183 184	The IOA results for the cytomorphological features are listed in Table 3. For the inexperienced observer the agreement between the DSs and GSs was slight for the amount and colour of cytoplasm, and fair for all the other morphological features. The mean IOA for		Deleted - by <b>Comment [FB125]:</b> Line 183 of YTVJL-D17-00838R2
181 182 183 184 185	The IOA results for the cytomorphological features are listed in Table 3. For the inexperienced observer the agreement between the DSs and GSs was slight for the amount and colour of cytoplasm, and fair for all the other morphological features. The mean IOA for cytomorphological features was fair. For the mildly experienced observer, the agreement between		Deleted - by <b>Comment [FB125]:</b> Line 183 of YTVJL-D17-00838R2
181 182 183 184 185 186	The IOA results for the cytomorphological features are listed in Table 3. For the inexperienced observer the agreement between the DSs and GSs was slight for the amount and colour of cytoplasm, and fair for all the other morphological features. The mean IOA for cytomorphological features was fair. For the mildly experienced observer, the agreement between the DSs and GSs was slight for cellular size, chromatin pattern, nuclear shape, and cellular		Deleted - by <b>Comment [FB125]:</b> Line 183 of YTVJL-D17-00838R2
181 182 183 184 185 186 187	The IOA results for the cytomorphological features are listed in Table 3. For the inexperienced observer the agreement between the DSs and GSs was slight for the amount and colour of cytoplasm, and fair for all the other morphological features. The mean IOA for cytomorphological features was fair. For the mildly experienced observer, the agreement between the DSs and GSs was slight for cellular size, chromatin pattern, nuclear shape, and cellular pleomorphism; fair for the amount of cytoplasm; and moderate for cytoplasm colour and the		Deleted - by <b>Comment [FB125]:</b> Line 183 of YTVJL-D17-00838R2
181 182 183 184 185 186 187 188	The IOA results for the cytomorphological features are listed in Table 3. For the inexperienced observer the agreement between the DSs and GSs was slight for the amount and colour of cytoplasm, and fair for all the other morphological features. The mean IOA for cytomorphological features was fair. For the mildly experienced observer, the agreement between the DSs and GSs was slight for cellular size, chromatin pattern, nuclear shape, and cellular pleomorphism; fair for the amount of cytoplasm; and moderate for cytoplasm colour and the presence of nucleoli. The mean IOA for cytomorphological features was fair. For the experienced		Deleted - by <b>Comment [FB125]:</b> Line 183 of YTVJL-D17-00838R2
181 182 183 184 185 186 187 188 189	The IOA results for the cytomorphological features are listed in Table 3. For the inexperienced observer the agreement between the DSs and GSs was slight for the amount and colour of cytoplasm, and fair for all the other morphological features. The mean IOA for cytomorphological features was fair. For the mildly experienced observer, the agreement between the DSs and GSs was slight for cellular size, chromatin pattern, nuclear shape, and cellular pleomorphism; fair for the amount of cytoplasm; and moderate for cytoplasm colour and the presence of nucleoli. The mean IOA for cytomorphological features was fair. For the experienced observer the agreement between the DSs and GSs was fair for cytoplasm; and moderate for cytoplasm colour and the presence of nucleoli. The mean IOA for cytomorphological features was fair. For the experienced observer the agreement between the DSs and GSs was fair for chromatin pattern, cytoplasm		Deleted - by <b>Comment [FB125]:</b> Line 183 of YTVJL-D17-00838R2

cytoplasm amount for which it was slight, and for cytoplasm colour and the presence of nucleoli forwhich it was moderate.

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195	
196	Even if no statistical differences were present, the use of WSI brought about an
197	improvement in the percentage of high-grade lymphomas correctly identified by the mildly
198	experienced observer and in the percentage of low-grade lymphomas correctly identified by the
199	inexperienced observer. Nevertheless, it affected the percentage of high-grade lymphomas
200	identified by the inexperienced and experienced observers, and the percentage of low-grade
201	lymphomas identified by the mildly experienced observer (Table 2). The agreement between the FC
202	and DS results in the assessment of grading was slight for the inexperienced observer, fair for the
203	mildly experienced observer, and moderate for the experienced observer, while the agreement
204	between the FC and GS results in the assessment of grading was fair for the inexperienced observer,
205	substantial for the mildly experienced observer, and moderate for the experienced observer (Table
206	4).
207	
208	Discussion
209	In this study, we assessed the reliability of the cytomorphological evaluation of canine
210	lymphoma samples by using a WSI scanner, and found a low IOA between the DSs and GSs for all
211	three observers for the morphological features assessed. The agreement between the DSs and FC
212	was slight for the inexperienced observer and fair to moderate for the two observers with more
213	experience. This underlines the importance of observer experience in the cytological evaluation of
214	lymphoma samples.

- 215
- 216 In recent years, researchers have shown increasing interest in both human and veterinary
- digital pathology (Maiolino et al., 2006; Kelly, 2007; Al-Janabi et al., 2012; Webster and Dunstan,
- 218 2014; Bertram et al., 2018). The three-dimensional architecture of cells in FNA samples limited the

219	application of digital pathology to cytological samples. However, with the introduction of WSI
220	scanners with the z-stack function, FNA samples have become more suitable for digitalisation and
221	visualisation on a monitor (El-Gabry et al., 2014). Nevertheless, no studies to date have validated
222	the use of WSI scanners in veterinary cytology. In our study, we chose cytological samples of
223	canine lymphoma to test our WSI scanner for many different reasons: first, lymphoma is the most
224	common haematopoietic tumour in dogs (Richards and Suter, 2015); second, the first-step in
225	diagnosis is often a FNA cytology of the lymph nodes; third, neoplastic lymphoid cells have
226	peculiar morphological features that can be appreciated by cytological analysis (Fournel-Fleury et
227	al., 2002; Ponce et al., 2003). To assess the concordance of information provided by the DSs and
228	GSs we evaluated the intra-observer variability, which is the preferred measure of performance in
229	digital pathology (Thrall et al., 2015). Moreover, to evaluate the diagnostic performance of our WSI
230	scanner, we compared the results of FC analysis to the data provided by the DS and GS analyses for
231	assessing the lymphoma grade on the basis of the fact that, nowadays, FC is considered a
232	fundamental tool in lymphoma diagnosis (Comazzi and Gelain, 2011; Comazzi et al., 2016). The
233	aim of this comparison was to evaluate whether the possible disagreement was due to the
234	descriptive capacity of the DSs or whether the challenge in the morphological identification of
235	grading was intrinsic to the cytological examination.
236	
237	In the evaluation of morphological features, the overall IOA of the three observers was fair/
238	to moderate. These IOAs are lower than those reported in the validation studies of other WSI
239	scanners used for histopathological analysis in veterinary pathology (Bertam et al., 2018) and for
240	histopathological (Thrall et al., 2015), cytological (House et al., 2013), and haematological (Gomez-
241	Gelvez et al., 2015) analyses in human pathology. The lower agreement obtained in our study is
242	likely not only because of the shortcomings of the tested WSI scanner, but also because of the
243	different design of our study compared to that of other studies in the literature. Indeed, the main
244	goal of our study was to determine the descriptive capacity of the WSI scanner rather than its

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245	'diagnostic performance'. The evaluation of morphological features of neoplastic lymphocytes
246	could also be difficult when using an optical microscope, and an unequivocal concordance on single
247	morphological features is challenging. However, our results clearly indicate that observer
248	experience is important in the cytological evaluation of lymphomas, as shown by the increasing
249	IOA from the inexperienced observer to the experienced one. Nevertheless, some technical
250	limitations of our WSI scanner contributed to the low level of agreement. In particular, the lowest
251	mean IOA was recorded in the evaluation of cytoplasm amount. The online software interface,
252	despite the z-stack scanning modality, does not have the capability to 'focus' up and down and does
253	not allows increasing the brightness of the DSs, thereby leading to a darker background than that
254	obtained for the GSs. The lack of these functions caused difficulties in the assessment of the edges
255	of the cells. Moreover, the cytoplasm was considered more frequently basophilic or deeply
256	basophilic in the DSs than in the GSs (Fig. 2). Conversely, the evaluation of the nucleoli was not
257	conditioned by the lack of these functions, and the presence of nucleoli was the microscopic detail
258	with the highest IOA. Thus, these limitations did not completely hamper our ability to evaluate
259	nuclear morphology.
260	
261	The mitotic count (MC) is defined as the number of mitoses in 10 consecutive high-power
262	fields (Meuten et al., 2016), and it is a useful parameter to define the malignancy grade of
263	lymphomas (Ponce et al., 2010). MC in cytology is somewhat controversial because of the uneven
264	distribution of cells throughout a sample, but it is considered a more reproducible and reliable tool
265	in histopathological examinations (Sapierzyński et al., 2016). Moreover, in our study, even if the
266	magnification used to analyse the DSs and GSs was the same, the area occupied by the cells in the
267	DSs viewed on the monitor was smaller than that of cells in the GSs, because a portion of the
268	monitor is occupied by the online software interface with navigation icons and commands.
269	Therefore, the number of cells/40× field of a DS was lower than that of cells/40× field of a GS, and
270	thus, the different number of cells/40× field did not allow for a comparison of the MC between the

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**Comment [FB205]:** Line 239 of YTVJL-D17-00838R2 Deleted – based on

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271	DSs and GSs. An alternative approach to comparing mitosis between the two methods is to consider
272	the mitotic index (MI), which is the ratio of the number of cells in mitosis and the number of cells
273	not in mitosis (Meuten et al., 2016). Nevertheless, this approach is time-consuming because it is
274	necessary to also count the cells not in mitosis, usually 1,000, and it is not applicable in the routine
275	diagnostic practice. For all these reasons, the MI was not included as a malignancy feature in our
276	study.

278	In addition to the possibilities to diagnose a lymphoma, the cytological examination of
279	lymph node FNAs could be useful to predict the grading of the neoplastic process. Large cells
280	and/or cellular pleomorphism could be suggestive of high grade lymphomas (e.g centroblastic B-
281	cell lymphoma or pleomorphic large T-cell lymphoma), while small cells are more frequently
282	observed in low grade lymphomas (e.g. small lymphocytic B-cell lymphoma or small clear T-cell
283	lymphoma) (Fournel-Fleury et al., 2002; Ponce et al., 2010). Despite the technical limitations of our
284	WSI scanner that led to difficulties in the evaluation of some morphological features, the IOA in the
285	determinations of grading was moderate for all the observers. These results may suggest that,
286	despite some difficulties in the evaluation of some specific characteristics, it is possible to obtain
287	acceptable results by using digital smears even with non-optimal images.
288	
289	The first aim of FC exam is the objective determination of the lymphoma's phenotype.
290	However, the information derived from FC go far beyond the simply identification of cell origin: it
291	allows the assessment of the percentage of neoplastic cells within FNA sample, their size, based on
292	the mean FSC, and the pleomorphism, based on the standard deviation of the FSC (Gelain et al.,
293	2008). Moreover, it is possible to evaluate the phenotype aberrations (both qualitative and
294	quantitative) which, in combination with morphological features (e.g size) allows to recognized
295	some specific lymphoma subtypes (e.g small clear cell lymphoma). In our study, the lymphoma's
296	grade determined by flow cytometry was compared to the lymphoma's grade obtained with DS and 12

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**Comment [FB237]:** Line 266 of YTVJL-D17-00838R2 Deleted – mitotic index

**Comment [FB238]:** Line 269 of YTVJL-D17-00838R2 Deleted - Besides

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**Comment [FB241]:** Line 272 of YTVJL-D17-00838R2 Deleted - ,

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**Comment [FB243]:** Lines 274-278 of YTVJL-D17-00838R2 Deleted - On contrary, the precise determination of the immunophenotype in canine lymphomas, essential for the prognosis, requires objective techniques, like immunohistochemistry or flow cytometry, to demonstrate lineagespecific cell antigens (Ponce et al., 2010, Comazzi and Gelain, 2011). However, an attempt at determination of phenotype could be

determination of phenotype could be made based on morphological Comment [FB244]: Line 278 of

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**Comment [FB245]:** Line 279 of YTVJL-D17-00838R2

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297	GS by the three observers. Regarding lymphoma grade, the results of the agreement with FC have
298	shown a correlation with observer experience. In fact, agreement was slight and fair for the DSs and
299	GSs, respectively, for the inexperienced observer, thus reflecting the observer's difficulty in
300	determining the lymphoma grade based only on the morphological features of the cells. In contrast,
301	the mildly experienced observer had good agreement when using the optical microscope and a
302	slightly decreased accuracy when using the WSI scanner, and the experienced observer had
303	moderate agreement when using both the DSs and GSs. These data confirm the importance of
304	observer experience in the evaluation of lymphoma samples and reflect the lower descriptive
305	capacity of the DSs than the GSs.
306	
307	From a technical perspective, the scanning time for each DS was between three and four
308	hours, depending on the amount of material on the GS. The time interval is related to the z-stack

scanning mode with 40× magnification. The use of WSI scanners with such long scanning times negatively affects their application in routine cytological diagnosis, wherein a single case often has multiple GSs. However, when challenging cases require the opinion of skilled clinical-pathologists at distant sites, this technology can be an efficient tool. To simulate a routine workflow in the use of the WSI scanner, for each case, we chose the slide with higher cellularity and better preservation,

because high cellularity and good preservation are characteristics that a regular practitioner can alsorecognize.

The limitation of the long scanning time could be solved using a more advanced WSI scanner with a higher speed of acquisition. The size of the DS (from 1 to 2 gigabytes) could also be a limitation and the use of a dedicated server is mandatory to facilitate the storage and sharing of the files (Gomez-Gelvez et al., 2015).

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321 Conclusions

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**Comment [FB250]:** Line 286 of YTVJL-D17-00838R2 Deleted - diagnostic

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Comment [FB254]: Line 287 of YTVJL-D17-00838R2 Deleted - both

**Comment [FB255]:** Line 287 of YTVJL-D17-00838R2 Deleted - one

**Comment [FB256]:** Line 288 of YTVJL-D17-00838R2 Deleted – the determination of

Comment [FB257]: Line 288 of YTVJL-D17-00838R2 Deleted – ing and phenotype of lymphoma

**Comment [FB258]:** Line 289 of YTVJL-D17-00838R2 Deleted – for unskilled observers

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322	The data from our study underline some technical limitations of the tested WSI scanner,	_	Comment [FB276]: Line 306 of YTVJL-D17-00838R2
323	mainly linked to image quality, which could limit the diagnostic power of the instrument. We have		Deleted - tested
324	also demonstrated the importance of observer experience in the correct interpretation of DSs.	$\backslash$	Comment [FB277]: Line 307 of YTVJL-D17-00838R2 Deleted - the
325	However, given the technological advances and development of new functions, the digital		Comment [FB278]: Line 307 of YTVJL-D17-00838R2 Deleted – of the images
326	cytological workflow in veterinary medicine could also be improved.	$\mathbb{N}$	Comment [FB279]: Line 307 of
327			YTVJL-D17-00838R2 Deleted - this
328	Conflict of interest statement: None of the authors of this paper have a financial or personal		<b>Comment [FB280]:</b> Line 308 of YTVJL-D17-00838R2 Deleted – 's
329	relationship with other people or organisations that could inappropriately influence or bias the		Comment [FB281]: Line 308 of YTVJL-D17-00838R2 Deleted - for
330	content of the paper.		Comment [FB282]: Line 309 of YTVJL-D17-00838R2
331			Deleted – digital slides
332	Acknowledgements		Comment [FB283]: Line 309 of YTVJL-D17-00838R2 Deleted - with
			Comment [FB284]: Line 309 of YTVJL-D17-00838R2
333	The WSI scanner (D-sight, A. Menarini Diagnostics S.r.l.) was acquired thanks to the scientific		Deleted – of technology
334	instrumentation grant (2012) by the University of Padua.		Comment [FB285]: Line 309 of YTVJL-D17-00838R2 Deleted – the release
335	The authors gratefully acknowledge the assistance of Dr. Alessandra Faggionato and of Dr.		Comment [FB286]: Line 310 of YTVJL-D17-00838R2
336	Francesco Cian in English editing.		Deleted – also in veterinary medicine
337	The preliminary results of this study were presented as an Abstract at the 18 <sup>th</sup> Congress of the		<b>Comment [FB287]:</b> Line 312 of YTVJL-D17-00838R2 Deleted – Declaration of interest
338	European Society of Veterinary Clinical Pathology, Nantes, 20-22 October 2016.		
339			
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Morphological features	1	2	3	4
Cellular size	Small	Medium	Large	
Cellular pleomorphism	Present	Absent		
Cytoplasm amount	Scarce	Moderate	Abundant	Unipolar
Cytoplasm colour	Clear	<b>B</b> asophilic	Deeply basophilic	
Nuclear shape	Round	Indented	Convoluted	Irregular
Chromatin pattern	Dense/Thickened	Granular/Coarse	Finely granular	Smooth
Nucleoli	Not present	Single	Multiple	

449 Morphological features and grading system used in the cytological evaluation.

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#### 452 Classification of the 44 lymphoma samples based on the flow cytometry results and the results of evaluation of the

453 digital and glass slides by the three observers.

	High grade lymphomas	Low grade lymphomas
Flow cytometry	34 (24 B-cells; 10 T-cells)	10 (4 B-cells; 6 T-cells)
Observer 1 - digital slides	23	9
Observer 2 - digital slides	30	5
Observer 3 – digital slides	27	7
Observer 1 – glass slides	26	8
Observer 2 – glass slides	28	8
Observer 3 – glass slides	30	7

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YTVJL-D17-00838R2 Deleted – using digital and glass slides Comment [FB291]: Table 2 of YTVJL-D17-00838R2 Deleted - High grade B-cell lymphomas . Low grade B-cell lymphomas High grade T-cell lymphomas Low grade T-cell lymphomas Flow cytometry 24 . 4 10 . 6 Observer 1 - digital slides 12,4 2,4 Observer 2 - digital slides 18 1 7 3 Observer 3 – digital slides 21 1 2 5 Observer 1 – glass slides 16 2 1 4 Observer 2 glass slides 18 1 8 6 Observer 3 –

glass slides 23 1 7 6

Comment [FB290]: Line 435 of

457 Intra-observer agreement for morphological features, grading, and phenotype. Observer one corresponded to the lower

458 level of cytological experience; observer two corresponded to the intermediate level of cytological experience; observer

459 three corresponded to the higher level of cytological expertise. The intra-observer agreement was assessed by using

460 linearly weighted Cohen's K. The K coefficients were interpreted as recommended by Landis and Koch (1977).

						Delet
Morphological features	Coefficient K observer 1 (95% CI)	Coefficient K observer 2 (95% CI)	Coefficient K observer 3 (95% CI)	Mean coefficient K (95% CI)	(	
Cellular size	0.29 (0.07-0.50)	0.00 (-0.26-0.21)	0.44 (0.14-0.72)	0.30 (0.16-0.44)		Com
Cellular pleomorphism	0.24 (-0.04-0.52)	0.17 (-0.10-0.43)	0.41 (0.14-0.68)	0.29 (0.13-0.45)		YTVJL Delet
Cytoplasm amount	0.03 (-0.16-0.21)	0.26 (0.04-0.47)	0.28 (0.04-0.51)	0.17 (0.04-0.31)	$\langle \rangle$	Com
Cytoplasm colour	0.13 (-0.10-0.36)	0.55 (0.35-0.75)	0.43 (0.21-0.64)	0.43 (0.30-0.56)	$ \$	YTVJL
Nuclear shape	0.28 (0.08-0.48)	0.14 (-0.23-0.52)	0.33 (-0.03-0.69)	0.31 (0.16-0.47)		Delet
Chromatin pattern	0.32 (0.10-0.55)	0.03 (-0.13-0.19)	0.22 (0.03-0.40)	0.22 (0.11-0.34)		Com
Nucleoli	0.33 (0.11-0.55)	0.52 (0.21-0.72)	0.50 (0.25-0.75)	0.48 (0.36-0.60)	$\neg      $	YTVJL Delet
Mean						<u> </u>
cytomorphological	0.34 (0.26-0.42)	0.40 (0.32-0.48)	0.52 (0.44-0.60)			Com
features	·					YTVJL Delet
Malignancy grade	0.53 (0.29-0.78)	0.44 (0.10-0.78)	0.46 (0.18-0.74)	0.51 (0.34-0.70)		
						Com

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Comment [FB292]: Line 442 of YTVJL-D17-00838R2 Deleted – follows as

Comment [FB293]: Table 3 of YTVJL-D17-00838R2 Deleted - 3
Comment [FB294]: Table 3 of YTVJL-D17-00838R2 Deleted - 24
Comment [FB295]: Table 3 of YTVJL-D17-00838R2 Deleted - 7
Comment [FB296]: Table 3 of YTVJL-D17-00838R2 Deleted - 9
Comment [FB297]: Table 3 of YTVJL-D17-00838R2 Deleted - 37
Comment [FB298]: Table 3 of YTVJL-D17-00838R2 Deleted - 25
<b>Comment [FB299]:</b> Table 3 of YTVJL-D17-00838R2 Deleted – 19
Comment [FB300]: Table 3 of YTVJL-D17-00838R2 Deleted - 5
Comment [FB301]: Table 3 of YTVJL-D17-00838R2 Deleted - 24
Comment [FB302]: Table 3 of YTVJL-D17-00838R2 Deleted - 28
Comment [FB303]: Table 3 of YTVJL-D17-00838R2 Deleted - 37
Comment [FB304]: Table 3 of YTVJL-D17-00838R2 Deleted – Phenotype 0.08 0.63 0. 34 0.35 Mean 0.31 0.54 0.40
Comment [FB305]: Table 3 of

468

- 463 Agreement between flow-cytometry results and the digital and glass slide assessments of lymphoma grading. Observer
- 464 one corresponded to the lower level of cytological experience; observer two corresponded to the intermediate level of
- 465 cytological experience; observer three corresponded to the higher level of cytological expertise. The diagnostic
- agreement was assessed by using linearly weighted Cohen's K. The K coefficients were interpreted as recommended by
- 467 Landis and Koch (1977).

	Coefficient K	Coefficient K	Coefficient K
	observer 1 (95% CI)	observer 2 (95% CI)	observer 3 (95% CI)
Flow cytometry results vs	0.16 (-0.13-0.50)	0.32 (0.06-0.58)	0.50 (0.23-0.77)
digital slide assessments			
Flow cytometry results vs	0.37 (0.12-0.62)	0.63 (0.39-0.87)	0.50 (0.24-0.75)
optical slide assessments			

**Comment [FB306]:** Line 446 of YTVJL-D17-00838R2 Deleted - optical

**Comment [FB307]:** Line 446 of YTVJL-D17-00838R2 Deleted – s in the

**Comment [FB308]:** Line 446 of YTVJL-D17-00838R2 Deleted – and phenotype

**Comment [FB309]:** Line 449 of YTVJL-D17-00838R2 Deleted – as follows

**Comment [FB310]:** Table 4 of YTVJL-D17-00838R2 Deleted – 0.33 0.63 0.47 0.24 0.77 0.73

#### 469 Figure legends

470

- 471 Fig. 1. Whole-slide imaging microscope imaging workstation (D-sight, A. Menarini Diagnostics
- and 40×) and a high-performance desktop computer. 1b. The main page of the online software used
- 474 for analysis (Telepathology, Visia Imaging S.r.l). The system allows to digitise, store, and preview
- 475 all digitalised slides. Thus, the user can scroll through the archive and select the slide of interest. 1c.
- 476 The online software's navigation page: using the image multi-preview, the user can select and edit
- 477 any area of interest (black asterisk). Information regarding the different magnifications available
- 478 (white asterisk), the current magnification (yellow asterisk), and the navigation map of the area
- 479 selected (red asterisk) are also present.
- 480
- 481 Fig. 2. Small-clear-cell lymphoma: 2a. Digital slide and 2b. Glass slide of the same sample ( $40 \times$
- 482 magnification). Compared to the glass slide, the digital slide, clearly shows the darker background
- and, consequently, the more basophilic cytoplasm of the cells.

Comment [FB311]: Line 455 of YTVJL-D17-00838R2 Deleted - 5

Comment [FB312]: Line 455 of YTVJL-D17-00838R2 Deleted - s

Comment [FB313]: Line 455 of YTVJL-D17-00838R2 Deleted - 4

Comment [FB314]: Line 456 of YTVJL-D17-00838R2 Deleted - desktop

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**Comment [FB316]:** Line 456 of YTVJL-D17-00838R2 Deleted - server

Comment [FB317]: Line 457 of YTVJL-D17-00838R2 Deleted - z

Comment [FB318]: Line 457 of YTVJL-D17-00838R2 Deleted - to

**Comment [FB319]:** Line 457 of YTVJL-D17-00838R2 Deleted - z

Comment [FB320]: Line 458 of YTVJL-D17-00838R2 Deleted - move

**Comment [FB321]:** Line 458 of YTVJL-D17-00838R2 Deleted – Server's

**Comment [FB322]:** Line 464 of YTVJL-D17-00838R2 Deleted – Server's Deleted - s

**Comment [FB323]:** Line 464 of YTVJL-D17-00838R2 Deleted – Server's Deleted - s

**Comment [FB324]:** Line 465 of YTVJL-D17-00838R2 Deleted – Server's Deleted - In

**Comment [FB325]:** Line 466 of YTVJL-D17-00838R2 Deleted – Server's Deleted – compared to the glass slide is evident

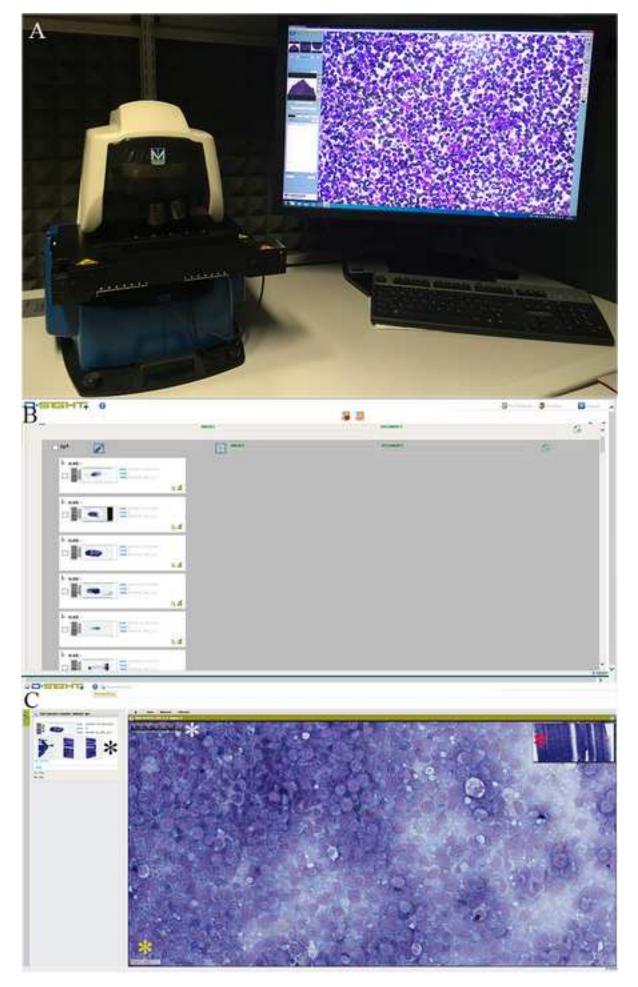
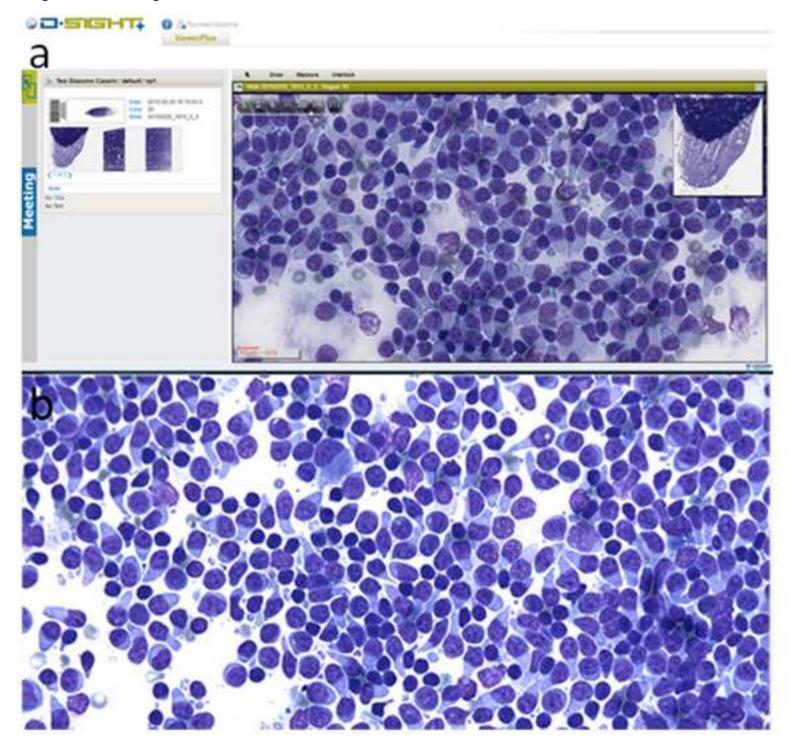


Figure2\_R1 Click here to download high resolution image



English\_editing\_certificare\_R2 Click here to download Optional e-only supplementary files: Bonsembiante et al.\_English\_editing\_certificate\_R2.pdf

# Highlights

- Whole slide imaging scanner is a robotic microscope capable to digitize a glass slide to create a composite digital image
- The cytomorphological capability of a whole slide imaging scanner is evaluated on 44 lymphoma samples
- There are some technical limitations that affected the diagnostic power of the instrument
- The release of new functions will be important to overcome the technical limitations
- This is the first time in which a whole slide imaging scanner is evaluated in Veterinary Medicine

**Revision Note** Manuscript Number: YTVJL-D-17-00838R2

Article Title: Whole-slide imaging: cytomorphological descriptive capacity and intra-observer agreement in canine lymphoma samples

**REVIEWERS' COMMENTS:** 

Reviewer #1:

**Reviewer comment** - Thanks for addressing most of the comments and concerns that were raised in the review process. I did note that the descriptive breakdown of agreement for morphological criteria were removed from the results in this version. I understand that the kappa values are listed in the tables, but I appreciated the information in the text of the previous version as well.

Author response –We added the requested information in the main manuscript (lines 182-194).

**Reviewer comment** - Regarding the impact of slide quality on whole slide imaging, I maintain that it would be valuable to have at least 1 sentence stating the percentage of samples that satisfactorily passed the QC process. According to the numbers provided in the rebuttal, only 27% of samples passed were of high enough quality for scanning. This is an important consideration for those that are interested in this technology.

**Author response** – We added this information in the main manuscript (lines 174-177). Minor comments:

**Reviewer comment** - Page 10, Line 242: I'm sorry I didn't pick up on this with the first review, but is "server" the correct word here? Is the focusing an issue with the server or the interface? **Author response** – We changed 'server' with 'online software interface' (line 251).

**Reviewer comment** - Page 12, Line 281: Wasn't the observer 2's IOA for phenotype substantial, not moderate?

**Author response** – We deleted all the information regarding the phenotype from the text, according to Reviewer 2 and Editor's comments.

Reviewer #2:

The authors have addressed many of my concerns and the paper is much improved. A few minor issues should still be addressed, in addition to some general editing for language to improve understanding.

**Reviewer comment** - Lines 73-75: Not all those tests are required to confirm lymphoma, therefore I would not say they are all fundamental. Consider rephrasing to "are fundamental tools that can be used to confirm and type lymphoma, etc." or something like that.

Author response – We added 'tool that can be used' in the main text (line 73).

**Reviewer comment** - Most instances of the word "telepathology" should be changed to "digital pathology" as no one was remotely controlling a microscope.

**Author response** – We changed 'telepathology' with 'digital pathology' in the main text (lines 43, 62, 217, 219, 229).

**Reviewer comment** - I remain in opposition to presumptively phenotyping lymphoma based on cytological morphology alone. I am concerned that this would give the impression that cytologic morphology is sufficient to type lymphoma when it is not. The publications used to justify classifying

immunophenotype based on cytological criteria did not actually use cytology as the gold standard for tumor classification. Ponce et al 2010 only examined cytology on 93/608 cases and did not use the cytology to classify the tumor types; the "cytological criteria" in their tables are the criteria used for examining histopathology slides not cytology slides. Similarly with Ponce 2004, the immunophenotype was determined by immunohistochemistry and immunocytochemistry, i.e. not based on cytological morphology. If this is kept, I would prefer to see a stronger emphasis in the discussion (and abstract) that cytological classification of lymphoma type is presumptive/preliminary and definitive methods (e.g. flow, IHC) are REQUIRED for true classification.

**Author response** – We agree with the reviewer about the needs to use immunological method to confirm the phenotype of lymphomas and we underlined it both in the text and in the revision note in the previous revision. We define the morphological evaluation only "an attempt "to determine the phenotype based on what is reported in literature (see Fournel-Fleury et al., 2002), being absolutely aware of the difficult to correctly correlate morphological features and immunophenotype in these neoplasms. However, according to the revisions of the Reviewer 2 and of the Editor, we deleted the part regarding the determination of the phenotype using cytology from the main text.

**Reviewer comment** - In my opinion, comparing morphological criteria identified on DS and GS vs FC to classify lymphomas is not so much evaluation of the ability of cytologists to identify lymphoma types per se, as it is an evaluation of the correlation between the evaluated morphological features and lymphoma immunophenotypes. Please comment.

**Author response** – Undoubtedly, the first aim of the flow cytometry exam is the objective determination of the lymphoma's phenotype. However, the information derived from FC go far beyond the simply identification of cell origin: it allows the assessment of the percentage of neoplastic cells within FNA sample, their size, based on the mean FSC, and the pleomorphism, based on the standard deviation of the FSC; it's possible to evaluate the phenotype aberration (both qualitative and quantitative) which, in combination with morphological features (e.g size) allows to recognized some specific lymphoma subtypes (e.g small clear cell lymphoma). Taken together, all these data allow to make diagnosis and to identify some lymphoma subtypes. In our study, the lymphoma's grade determined by flow cytometry was compared to the lymphoma's grade obtained with DS and GS by the three observers. We added this information in the main text (lines 289-297).

**Reviewer comment** - Throughout the paper, I would suggest replacing observer number (1, 2, or 3) with experience level, e.g. "inexperienced", "mildly experienced" and "experienced" observer. This allows someone reading the paper to understand the context without referring back to the M&M to make that determination. **Author response** – We replaced observer one with inexperienced observer, observer two with mildly experienced observer, and observer three with experienced observer in the main text.

I think the reasoning in the response to the following earlier reviewer comment is valid and should be included in the discussion:

**Reviewer comment** - Reviewer comment - Lines 106-107: "most representative slide" automatically creates bias. In a clinical situation, would a regular practitioner know how to choose the best slide? Also, based on reviewer experience, this information may not apply to a regular practitioner who often has less than the equivalent of 2 years cytology expertise.

Author response - It is our opinion that the WSI technology could be applied to cytological samples only for second opinion or for selected challenging cases in which the experience of a board-certified clinical-pathologist is required and, due to the long scanning time, not for the routine diagnostic procedures. For these reasons we included only cases with high cellularity and well preserved. The cellularity and preservation of a sample are characteristics that also a regular practitioner can recognize. We changed 'the most representative slide' with 'the slide with higher cellularity and better preservation' in the revised manuscript (lines 115-116).

Author response - We added this information in the discussion section (lines 312-315).

I also think the information below should be added to the M&M:

**Reviewer comment** - Reviewer comment - Lines 130-136: Did cytologists confer before the study to agree on the appearance of the morphological features for standardization within the study?

Author response - Yes, the cytologists conferred before the start of the study to agree on the appearance of the morphological features of the cells.

Author response – We added this information in the M&M section (line 153-154).

**Reviewer comment** - Lines 143-144: Variation in cell size is anisocytosis, not pleomorphism. Also please define the anisocytosis cut-off at which "pleomorphism" was considered present. Greater than 10% anisocytosis? Greater than 50%? Greater than 100%?

**Author response** – In the cytological evaluation of neoplastic lymph-nodes, the presence of cells with various size and shape is defined 'pleomorphism' (Fournel-Fleury et al., 2002), and the lymphomas are classified as pleomorphic (e.g. centroblastic polymorphic; small-, mixed-, or large-cell pleomorphic non-hodgkin lymphoma) when cells with different size are present (Ponce et al., 2010). In literature no cut-offs at which pleomorphism is considered present are reported.

**Reviewer comment** - Lines 145-148: Was perinuclear clearing considered in the distribution of the cytoplasm colour?

Author response – We did not consider the perinuclear halo in the distribution of the cytoplasm colour.

**Reviewer comment** - Line 152: Was variation in size or shape of nucleoli recorded? **Author response** – We considered the presence/absence and the number of the nucleoli but not the variation in shape and size

**Reviewer comment** - Lines 153-164: What is the value of comparing the classification of low vs high grade lymphoma, when they are the exact same as comparing the classification of cell size (small vs. large and mixed)? Similarly, based on the descriptions provided B vs T cell phenotype is based on potentially dichotomous criteria (clear vs. colored cytoplasm, round vs. not-round nuclei), so why not simply compare the observers' abilities to detect those?

**Author response** – The cellular size is the only objective parameter that was also included in the main text as single morphological feature, but the lymphoma's grade is determined by an overall assessment of the cytological slides.

**Reviewer comment** - It would be interesting to consider, if the data is available, how the observer's subjective assessment of cell size (small, medium, large) correlated with the FC measurement of cell size? **Author response** – Actually, this correlation is not one of the aim of our study, so we didn't record these data.

**Reviewer comment** - Lines 191-194: Are these changes statistically significant? **Author response** – The changes are not statistically significant, we added this information in the main text (line 196).

**Reviewer comment** - Line 183/Table 3: What statistical test was used to determine if the differences between observers was statistically significant or not? Alternatively, what are the confidence intervals for these Kappas?

Author response – We added the 95% confidence interval for the Kappa values in tables 3 and 4.

ADDITIONAL EDITORIAL COMMENTS:

**Reviewer comment** - We agree with Reviewer 2 that (presumptive) phenotyping/immunophenotyping of lymphoma should not done on cytomorphology alone; this section of the manuscript should be removed. **Author response** – We deleted the phenotypical assessment based on the morphological evaluation of the neoplastic cells from the main text.

**Reviewer comment** - Please change "Declaration of interest" to "Conflict of interest statement". **Author response** – Done (line 328).

In addition, one of the reviewers has made the following comments:

**Reviewer comment** - The Title is too long and should be reworded. **Author response** – The Title has been changed and reduced (lines 3-4).

**Reviewer comment** - The current Abstract is not useful and needs to be rewritten. **Author response** - We rewrote the abstract (lines 20-39).

**Reviewer comment** - Overall, the standard of English needs to be improved and requires major editing. **Author response** – The manuscript has been edited by the Elsevier Language Editing Service. The certificate is uploaded together with the other files.

**Reviewer comment** - There is a need to provide an evaluation of the statistical significance of the differences between kappa values.

**Author response** – The K value is a measure of the agreement between two observers using two different techniques or between two techniques used by the same observer. Rarely the statistical significance for Cohen's K is reported, probably because even relatively low values of K can nonethelsess be significantly different from zero but not sufficient magnitude to satisfy researchers (Bakeman and Gottman, 1997). Still, its standard error has been described (Fleiss et al., 1969). We added the 95% CI as requested by Reviewer 2.

**Reviewer comment** - The discussion is repetitive and needs to be streamlined. **Author response** – We changed the discussion (lines 209-319).



DIPARTIMENTO DI BIOMEDICINA COMPARATA E ALIMENTAZIONE



Università degli Studi di Padova

VIALE DELL'UNIVERSITÀ, 16 - 35020 LEGNARO (PD) - TEL. 0498272601 FAX. 0498272604

Legnaro (PD), 06.04.2018

Dear Editor,

Please find enclosed herewith the revised version of the manuscript titled "Whole-slide imaging: cytomorphological descriptive capacity and intra-observer agreement in canine lymphoma samples (YTVJL-D-17-00838R2)".

We answered to all the revisions of the two Reviewers and of the Editor. The title's length was reduced, the abstract was changed, and all the data regarding the determination of phenotype using cytology exam were removed from the manuscript. The manuscript was revised by the Elsevier's Language Editing Service (the certificate is attached together with the other files). To add the 95% CI of the K values we changed the statistical program (from SPSS to MedCalc). With the new statistical the K value of some parameters is slightly changed. The manuscript was revised according to the new K values.

Feel free to contact me for any questions.

Thank you in advance.

Best regards.

Federico Bonsembiante

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