## **Supplemental Material**

Cytologic Scoring of Equine Exercise-Induced Pulmonary Hemorrhage: Performance of Human Experts and a Deep Learning-Based Algorithm

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**Supplemental Table S1.** Confusion matrix of the hemosiderin grades of alveolar macrophages assigned to the same cells by the ten annotators and the ground truth annotations.

|                           |   | Ground tru | Ground truth's cell grade |        |       |       |  |
|---------------------------|---|------------|---------------------------|--------|-------|-------|--|
|                           |   | 0          | 1                         | 2      | 3     | 4     |  |
| Annotators'<br>cell grade | 0 | 28,646     | 6 2,448                   | 87     | 16    | 0     |  |
|                           | 1 | 11,668     | 3 25,385                  | 2,821  | 53    | 8     |  |
|                           | 2 | 486        | 6 15,215                  | 15,213 | 1,784 | 5     |  |
|                           | 3 | ç          | ) 1,147                   | 7,102  | 4,503 | 160   |  |
|                           | 4 | Z          | . 71                      | 1,051  | 2,275 | 1,060 |  |

For the 158,143 annotations made by the annotators, a match was found in the ground truth dataset (Euclidean distance between the annotators' and ground truth annotation of  $\leq$  50 pixels) in 121,217 (76.7%) instances. Of the 121,217 alveolar macrophages, 74,807 (61.71%) were assigned the same hemosiderin grade by the annotators and the ground truth. Of the 46,410 macrophages with divergent hemosiderin grade, 43,473 (93.67%) had a divergence by one grade level, 2,829 (6.10%) had a divergence by two grade levels, 104 (0.22%) had a divergence by three grade levels, and 4 (<0.01%) had a divergence by four grade levels.

**Supplemental Table S2.** Confusion matrix of the hemosiderin grades of alveolar macrophages assigned to the same cells by the ground truth annotator and the deep learning-based algorithm.

|                        |   | Ground truth's cell grade |        |   |        |   |        |        |       |
|------------------------|---|---------------------------|--------|---|--------|---|--------|--------|-------|
|                        |   | 0                         |        | 1 |        | 2 |        | 3      | 4     |
| Algorithmic cell grade | 0 |                           | 58,341 |   | 2,934  |   | 3      | 1      | 0     |
|                        | 1 |                           | 2,413  |   | 62,627 |   | 2,867  | 0      | 0     |
|                        | 2 |                           | 10     |   | 4,347  |   | 36,516 | 1,971  | 0     |
|                        | 3 |                           | 3      |   | 4      |   | 1,326  | 11,615 | 263   |
|                        | 4 |                           | 1      |   | 0      |   | 1      | 184    | 1,223 |

For the 218,003 algorithmic predictions, a match (Euclidean distance between the pathologist's and ground truth annotation of  $\leq$  50 pixels) was found in the ground truth dataset in 186,650 (85.62%) instances. Of the 186,650 alveolar macrophages, 170,322 (91.25%) were assigned the same hemosiderin score by the ground truth annotator and the algorithm. Of the 16,328 macrophages with divergent hemosiderin grade, 16,305 (99.86%) had a divergence by one grade level, 18 (0.11%) had a divergence by two grade levels, 4 (0.02%) had a divergence by three grade levels, and 1 (<0.01%) had a divergence by four grade levels.

| based algoritr            | ım. |      |                        |        |        |       |     |  |  |
|---------------------------|-----|------|------------------------|--------|--------|-------|-----|--|--|
|                           |     | Algo | Algorithmic cell grade |        |        |       |     |  |  |
|                           |     | 0    |                        | 1      | 2      | 3     | 4   |  |  |
| Annotators'<br>cell grade | 0   |      | 30,228                 | 2,301  | 73     | 18    | 0   |  |  |
|                           | 1   |      | 11,595                 | 24,986 | 2,911  | 44    | 5   |  |  |
|                           | 2   |      | 426                    | 14,378 | 15,603 | 1,530 | 5   |  |  |
|                           | 3   |      | 10                     | 918    | 7,279  | 4,237 | 168 |  |  |

**Supplemental Table S3.** Confusion matrix of the hemosiderin grades of alveolar macrophages assigned to the same cells by the ten annotators and the deep learning-based algorithm.

For the 158,143 annotations made by the annotators, a match (Euclidean distance between the annotators' and ground truth annotation of  $\leq$  50 pixels) was found in the algorithmic predictions in 121,025 (76.53%) instances. Of the 121,025 alveolar macrophages, 76,051 (62.84%) were assigned the same hemosiderin grade by the annotators and the algorithm. Of the 44,974 macrophages with divergent hemosiderin grade, 42,411 (94.30%) had a divergence by one grade level, 2,477 (5.51%) had a divergence by two grade levels, 84 (0.19%) had a divergence by three score levels, and 2 (<0.01%) had a divergence by four grade levels.

51

1,011

2,249

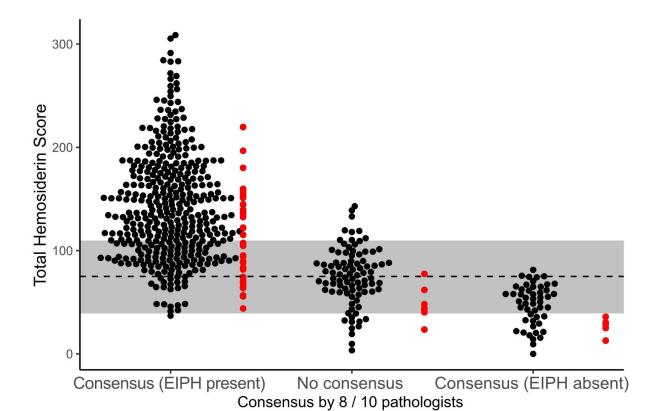
997

2

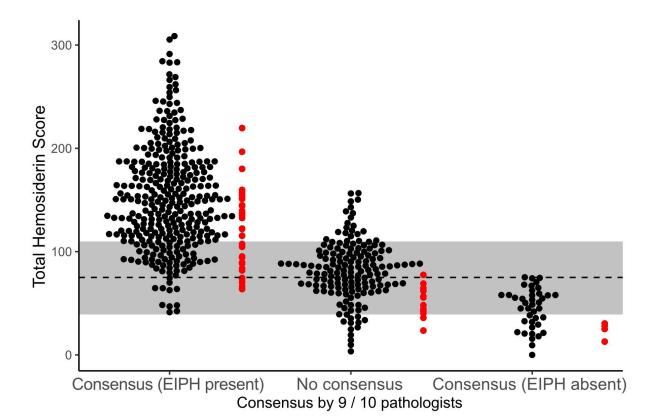
4

| Consensus definition        | Number of cases with consensus |                            |  |  |  |
|-----------------------------|--------------------------------|----------------------------|--|--|--|
|                             | Annotator's THS                | grade-<br>standardized THS |  |  |  |
| By all of the 10 annotators | 24 / 52 (46%)                  | 40 / 52 (77%)              |  |  |  |
| By at least 9/10 annotators | 36 / 52 (69%)                  | 47 / 52 (90%)              |  |  |  |
| By at least 8/10 annotators | 43 / 52 (83%)                  | 50 / 52 (96%)              |  |  |  |
| By at least 7/10 annotators | 49 / 52 (94%)                  | 52 / 52 (100%)             |  |  |  |
| By at least 6/10 annotators | 50 / 52 (96%)                  | 52 / 52 (100%)             |  |  |  |
| By at least 5/10 annotators | 52 / 52 (100%)                 | 52 / 52 (100%)             |  |  |  |

**Supplemental Table S4.** Degree of consensus by the ten annotators on exercise-induced pulmonary hemorrhage diagnosis (THS score above or below 75).



**Supplemental Fig. S1.** Scatter plots for total hemosiderin scores (THSs) determined by the ten annotators (black dots) and deep learning-based algorithm (red dots). The 52 cases are separated based on their consensus of the exercise-induced pulmonary hemorrhage (EIPH) diagnosis by 8 out of 10 annotators. The left scatter plot represents the THS values for cases with a consensus on THS values above the cutoff of 75 (N = 32), the middle scatter plot represents the THS values for cases with no consensus (N = 16) and the right scatter plot represents the THS values for cases with a consensus the THS values for cases with a consensus the THS values for cases with no consensus on THS values below the cut-off of 75 (N = 4). The broken line indicates the diagnostic cut-off at a THS of 75 and the grey bar around the broken line is the reference range determined in this study.



**Supplemental Fig. S2.** Scatter plots for total hemosiderin scores (THSs) determined by the ten annotators (black dots) and deep learning-based algorithm (red dots). The 52 cases are separated based on their consensus of the exercise-induced pulmonary hemorrhage (EIPH) diagnosis by 9 out of 10 annotators. The left scatter plot represents the THS values for cases with a consensus on THS values above the cut-off of 75 (N = 32), the middle scatter plot represents the THS values for cases with no consensus (N = 16) and the right scatter plot represents the THS values for cases with a consensus the THS values for cases with a consensus the THS values for cases with no consensus on THS values below the cut-off of 75 (N = 4). The broken line indicates the diagnostic cut-off at a THS of 75 and the grey bar around the broken line is the reference range determined in this study.