Diagnosis System For Fish Diseases Trichodina and Gyrodactylus Under Microscope Based on Decision Tree and LDA.

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>Introduction

>Methodology

- **Results and Discussion**
- ≻Conclusion

>Future Recommendations



- Detection Trichodina and Gyrodactylus disease in fish, using the system.
- Aim to rapid disease diagnosis.
- Allowing for earlier treatment of infected fish to prevent the spread of disease.
- Achieve an efficient result of identification.



- **Trichodina:** is one of the parasites that you can't see without a microscope. This parasite is usually found on the gills, skin or fins of the fish.
- **Gyrodactylus:** is an interesting little parasite; species of which can be found infesting many different types of freshwater. It generally infests the skin and fins of the fish.
- Both diseases Spread in tilapia fish.

Trichodina and Gyrodactylus



Disease Diagnosis

- **Traditional:** Based on fish-farmer or fish-veterinarian using their experiences and skill during their career.
- Modern: IT technology provide the web-based guide for diagnosing fish diseases.
- On this paper: An image processing-based fish disease diagnosis system was developed.

Methodology



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Feature Extraction

• Image Feature Extraction Then Create Dataset.



| | mean | median | stander | var | target |
|-------|------------|------------|------------|-------------|------------|
| count | 262.000000 | 262.000000 | 262.000000 | 262.000000 | 262.000000 |
| mean | 133.799604 | 135.874046 | 74.407862 | 5538.005335 | 1.629771 |
| std | 6.516057 | 8.652142 | 1.216991 | 184.682151 | 0.483790 |
| min | 98.045608 | 96.000000 | 70.938491 | 5032.269528 | 1.000000 |
| 25% | 129.818424 | 130.250000 | 73.837406 | 5451.962499 | 1.000000 |
| 50% | 134.074875 | 136,000000 | 74.266483 | 5515.510532 | 2.000000 |
| 75% | 138.583188 | 142.750000 | 74,731698 | 5584.826612 | 2.000000 |
| max | 148.722055 | 158.000000 | 82.123353 | 6744.245132 | 2.000000 |

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Fisher's Linear discriminant analysis (LDA)

• Linear discriminant analysis (LDA) is one of the most popular supervised dimensionality reduction methods for the dataset.

```
# Applying Fisher's Linear Discriminant Analysis
from sklearn.discriminant_analysis import LinearDiscriminantAnalysis
lda = LinearDiscriminantAnalysis(n_components = 2)
X_train = lda.fit_transform(X_train, y_train)
X_test = lda.transform(X_test)
```



• the comparison between the classification accuracy of SURF-LDA-DT and proposed combination which is ORB-LDA-DT.

| Algorithm | Accuracy | |
|-------------|----------|--|
| ORB-LDA-DT | 87.5% | |
| SURF-LDA-DT | 85% | |



Accuracy: The algorithm recorded a final score of 87.5% As the next confusing matrix will explain.



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Classification Report Equations

- Precision = true_positive / (true_positive + false_positive)
- Recall = true_positive / (true_positive + false_negative)
- F-score = 2 * Precision * Recall / (Precision + Recall)
- The support is the number of samples of the true response that lies in that class.

Classification Report

• The precision and recall or F-score of the minor class might be the most suitable measurement.

| | precision | recall | f1-score | Support |
|--------------|-----------|--------|----------|---------|
| Trichodina | 0.83 | 0.71 | 0.77 | 7 |
| Gyrodactylus | 0.86 | 0.92 | 0.89 | 12 |
| avg/total | 0.85 | 0.85 | 0.85 | 20 |



Discussion

•Using (Decision Tree) classifier in detection of the Trichodina and Gyrodactylus diseases on fish.

•The final score of the total algorithm is **87.5%**.

•The algorithm answered 5 imaged from class one true and 2 image false and also answered 1 images from class two false and 12 images true.

•The dataset constructed and built from about 260 images from the two classes of diseases, and saved into a CSV excel file.

Conclusion

•The proposed combination of ORB-LDA-DT gives better classification accuracy than the existing combined technique which is SURF-LDA-DT.

•It is fast and efficient method to recognize and detect the Trichodina and Gyrodactylus diseases as compared to the Traditional Method.

•The Decision Tree results based on the real images of infected Fish disease dataset. The proposed combination (ORB-LDA-DT) gives 87.5% accuracy.

Future Recommendations

•Update machine learning algorithms to deep learning techniques to increase the accuracy of the system by up to 99%.

•We plan to detect the disease based on the kinds of pathogens such as parasites, bacteria, fungal, and the virus for more convenient and useful aid to the diagnosis system.

•We will make machine learning algorithms can apply to different feature Descriptors.

•We aim to identify other diseases in fish rather than only one disease.

•We aim to increase our dataset with more samples.



Thanks!

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