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Random subsampling techniques for sea bass mortality prediction

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Agenda

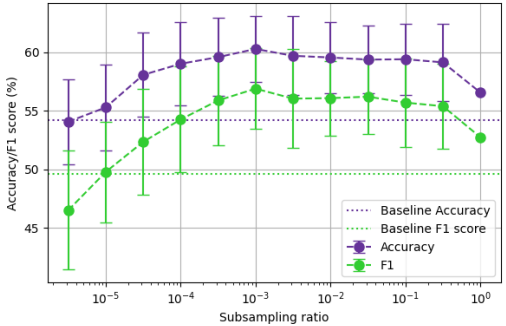
1. Introduction
2. Results
3. Conclusions

Introduction

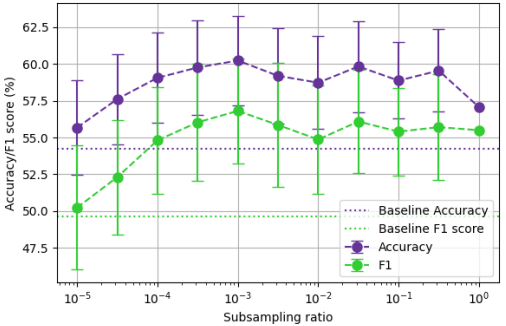
Methodology: subsampling

Results

Results: uniform subsampling



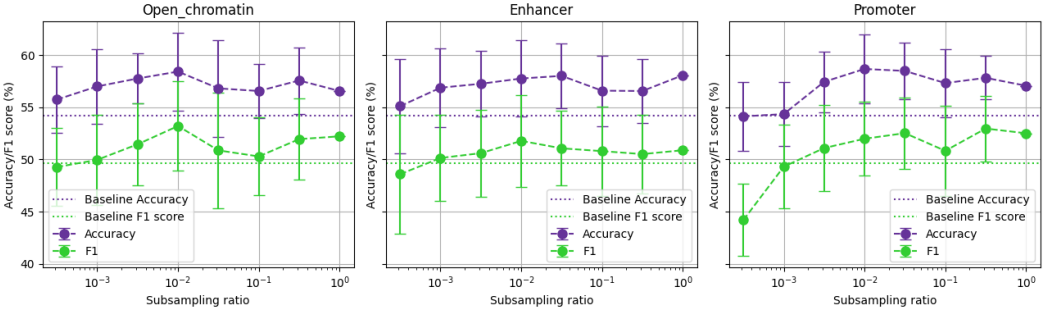
(a) Plot of the scores when subsampling uniformly on the whole genome.



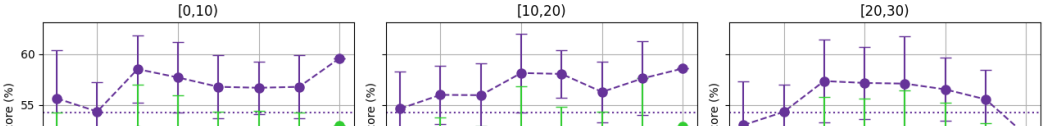
(b) Plot of the scores when subsampling uniformly on each chromosome.

Figure: Plots of accuracy and F1 scores for different subsampling rates on the whole genome.

Results: annotated subsampling



(a) Plots of the scores while subsampling genes annotated in different categories.



Conclusions