



Deep SHAP to revealed nucleotide contributions (Wang *et al.*, 2019). Deep SHAP is a compositional approximation of SHAP values since it is challenged to compute SHAP values exactly, especially for a complex deep neural networks (Lundberg and Lee, 2017). In our understanding, the method based on input perturbation often requires better generalization ability of the model (even for artificial ridiculous noise data). Moreover, recent work indicates that model explain techniques, which based post hoc explanations techniques and input perturbations, could be fooled to generate meaningless explanations instead of reflecting the underlying biases (Slack *et al.*, 2019), in other word, they could be unreliable and misleading, even on model with excellent performance. In light of the above, we believe it is essential to develop a model which can not only match deep learning based model in performance, but also be comparable to conventional machine learning algorithms in interpretability.

## 2 Approach

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## Acknowledgements

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