Pharmahacks 2025 Error 404 Team Name Not Found

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Multiple Sequence Alignment

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Reinforcemen t Learning

Agent: Model **Action**: Step **Environment**:

Sequence Alignment

State: Score calculation for current alignment

Reward:Positive

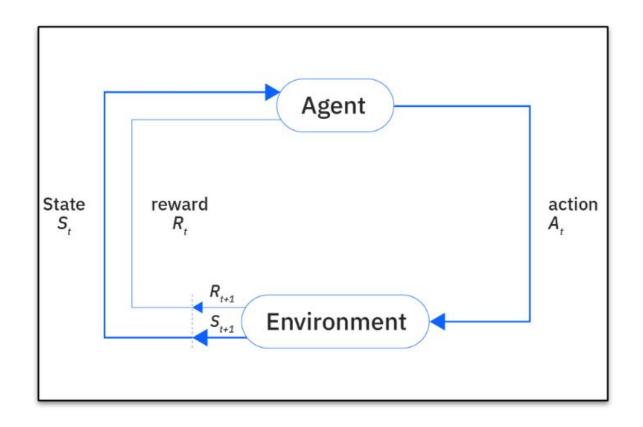
Score(New) > Score(Old)

- Step is made

Negative

Score(New) < Score(Old)

- Step is not made
- Model is penalized

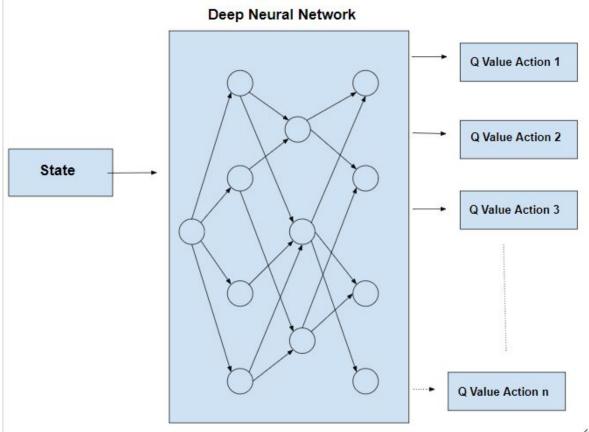


Challenge: Exponential number of possible steps

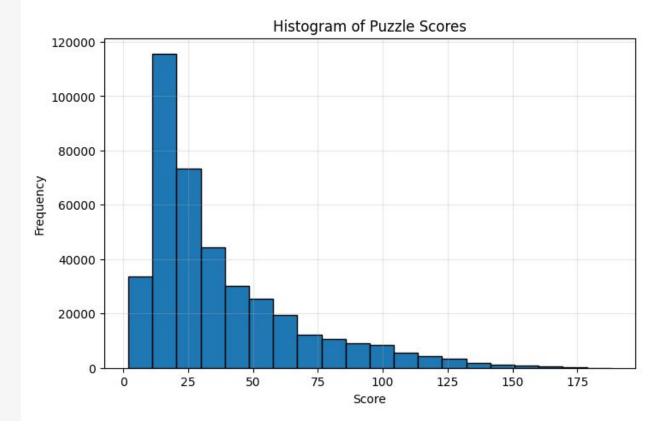
Model: Deep-Q Learning

Mathematical approximation of all the possible steps given a state

Neural Network chooses the best action (step)



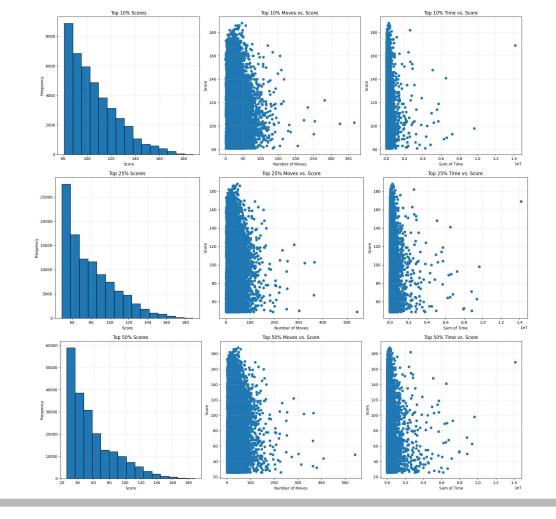
Visualize Data Spread



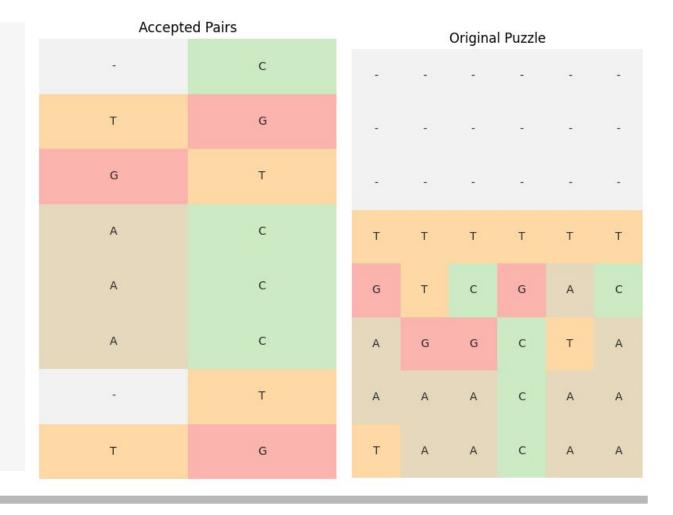
Trend's from Top Scorers

No strong trends between number of moves or gameplay time and high scores

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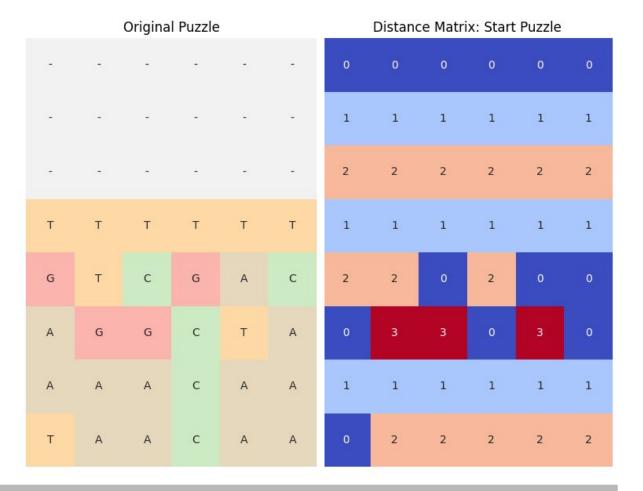
Model Theory



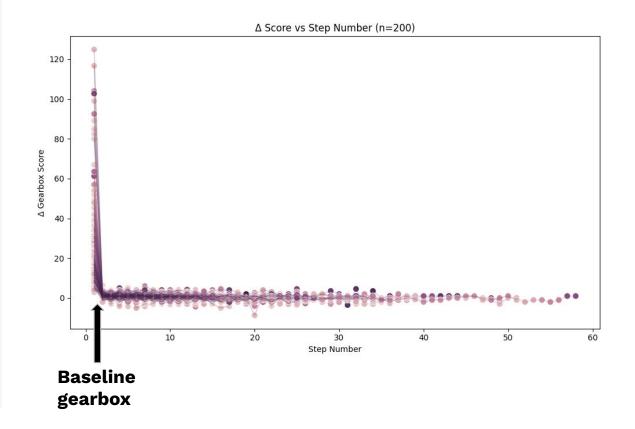
Distance Algorithm

Calculate how far each base pair is from alignment

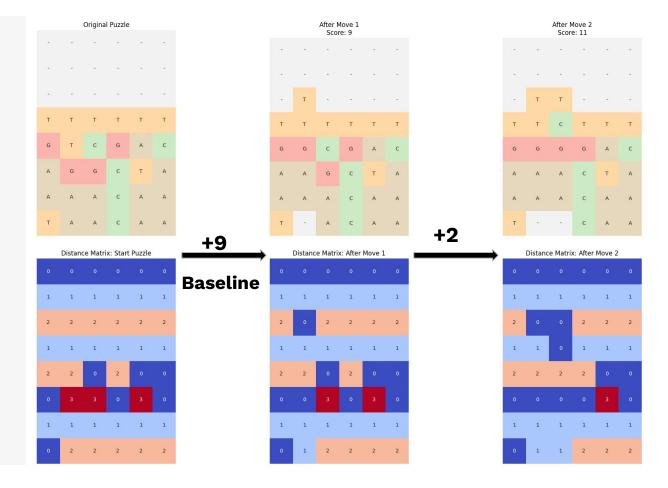
Spatial awareness task component enhances model accuracy and efficiency



Importance of first steps



Algorithm input: Correlate changes in distance matrix to gearbox score



Model + Task

Adding the distance algorithm task as a parameter in the neural network helps refine learning to make it more MSA-specific

Features:

- Steps
- Accepted Pairs
- Start
- Distance Matrix

Target Variable:

- Score

Goal:

Model solution score should beat test solution score

Moves variable omitted for simplicity

Model

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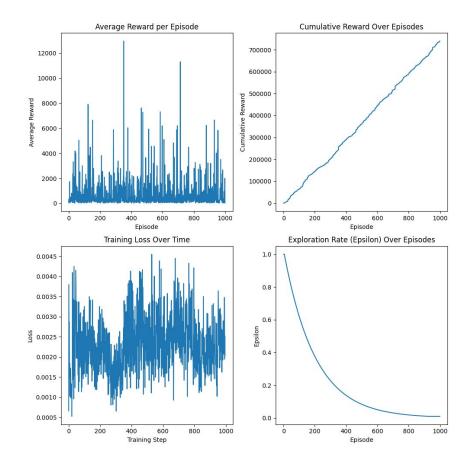
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O2 Results

Evaluation Metrics



Evaluation Metrics

Model Metrics currently in queue as of submission deadline

Conclusions

Biological Significanc e

- This algorithm is a starting point for small sequences
- Much more complicated to apply to large sequences
- Could be refined to be more biology-specific with the integration of the spatial task algorithm

Future Directions

- Increasing models layers and episodes
- Evaluate carbon footprint
- Use Leave-One-Out cross validation to refine algorithm
- Expand on distance matrix calculations to refine efficiency