

# IGOR MELNYK

Capital One ♦ AI Foundations ♦ New York, NY

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## SUMMARY

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Distinguished Applied Researcher at Capital One with 9+ years of experience working in the areas of Machine Learning and AI with the focus on Generative models in Natural Language Processing, Computer Vision and Biology

## EDUCATION

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**PhD in Computer Science and Engineering**, University of Minnesota, Minneapolis, MN June, 2016

**MS in Computer Science**, University of Colorado, Boulder, CO May, 2009

**BS in Computer Science**, Dnipro National University, Dnipro, Ukraine June, 2004

## RESEARCH INTERESTS / SKILLS

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<b>Machine Learning / AI</b>	Generative Models, Knowledge Graph, Protein Design, Anomaly Detection Style Transfer, Information Theory, Image Captioning
<b>Technical Skills</b>	Python/C++, Pytorch/Tensoflow

## EXPERIENCE

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<b>Distinguished Applied Researcher</b> Capital One	Aug 2024 - Present New York, NY
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<b>Research Staff Member</b> IBM Research AI	Sep 2016 - Aug 2024 Yorktown Heights, NY
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<b>Research Assistant</b> University of Minnesota	Sep 2009 - June 2016 Minneapolis, MN
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<b>Data Scientist</b> NASA Ames Research Center	May 2014 - Aug 2014 Mountain View, CA
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<b>Software Developer</b> Cisco Systems	May 2008 - Aug 2008 Boulder, CO
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## SELECTED PUBLICATIONS

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- EpMAN: Episodic Memory AttentionN for Generalizing to Longer Contexts, *ACL*, 2025
- AlphaFold Distillation for Protein Design, *Scientific Reports*, 2025
- Distributional Preference Alignment of LLMs via Optimal Transport (AOT) *NeurIPS*, 2024
- Larimar: Large Language Models with Episodic Memory Control, *ICML*, 2024
- Risk Aware Benchmarking of Large Language Models, *ICML*, 2024
- Reprogramming Pretrained Language Models for Antibody Sequence Infilling, *ICML*, 2023
- Knowledge Graph Generation From Text, *EMNLP*, 2022

## SELECTED CODE

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- [github.com/huggingface/trl](https://github.com/huggingface/trl): Alignmnet via Optimal Transport (AOT) is part of HuggingFace TRL library
- [github.com/IBM/AFDistill](https://github.com/IBM/AFDistill): AlphaFold Distillation for Inverse Protein Folding
- [github.com/IBM/ReprogBERT](https://github.com/IBM/ReprogBERT): Model Reprogramming for Antibody Design
- [github.com/imelnyk/ArxivPapers](https://github.com/imelnyk/ArxivPapers): Arxiv Papers Digest