

The background features several stylized molecular diagrams. In the top left, a small cluster of four atoms (one blue, three orange) is connected by white lines. In the top center, a larger network of seven atoms (one blue, six orange) is shown. On the right side, a large DNA double helix is depicted with red and blue strands and white rungs. In the bottom right corner, another network of seven atoms (one blue, six orange) is shown.

# AR Protein Visualization

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# The Team

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Morristown, NJ  
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Samhitha



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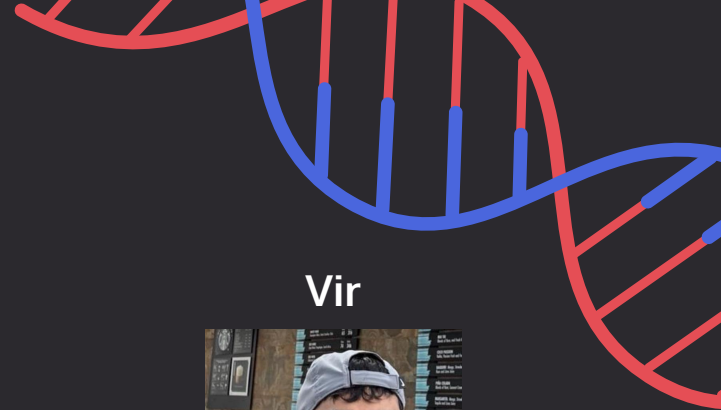
Vir



Princeton, NJ  
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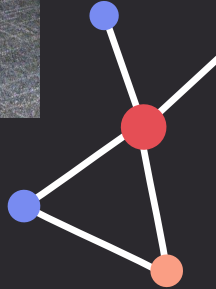
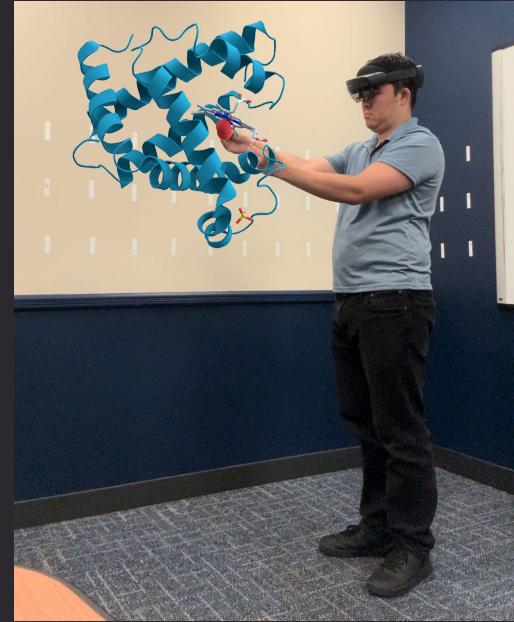


Computer  
Engineering with  
minor Computer  
Science



# Motivation

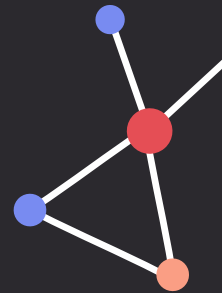
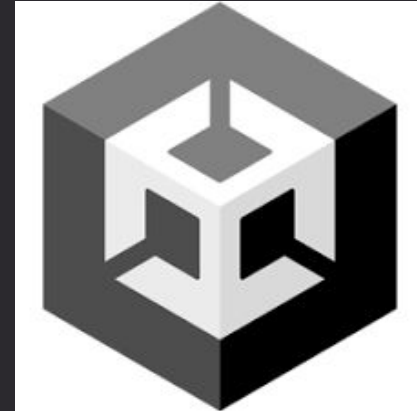
- What is Augmented Reality?
  - Computer-generated image on a user's view of the real world
- Why Augmented Reality?
  - Viewing 3D data on a 2D screen hides many details
  - Immersive and interactive exploration of complex datasets

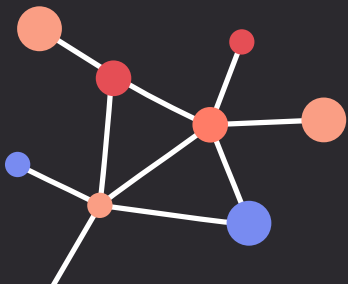


# Project and Technology Overview

Goal: Develop an interactive Augmented Reality (AR) solution to visualize and analyze molecular proteins, enhancing understanding.

- Data Source: Rutgers Protein Data Bank
- Technology: Unity Game Engine with C# scripting, Microsoft Hololens 2

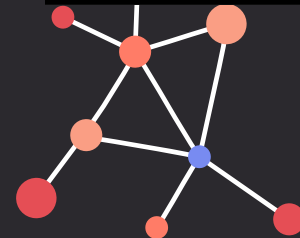
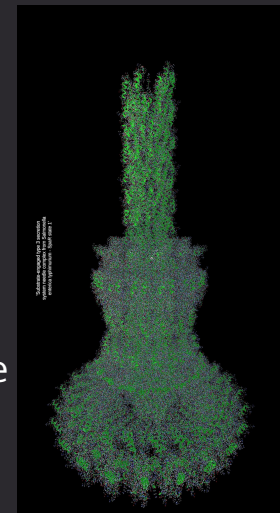
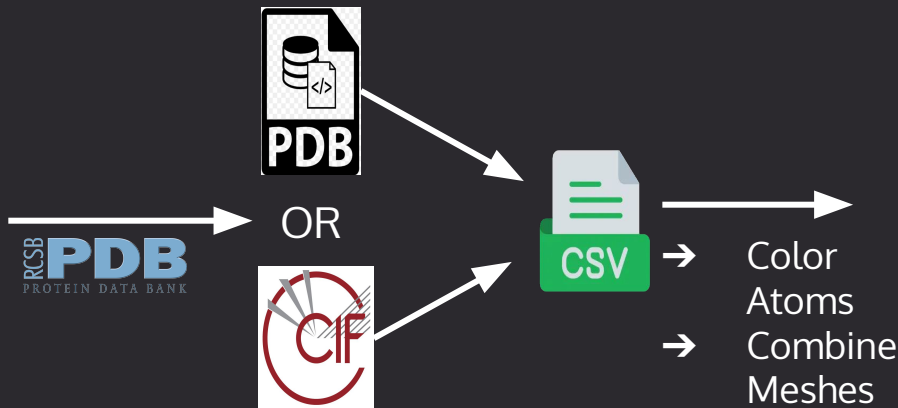




# Program Breakdown



Input code:  
"6PHQ"



# Web Scraping and CSV



- Downloads file from database
- Filters important information
- Autonomous

Original File:

```
HELX_P HELX_P1 1 VAL B 7 ? HIS B 16 ? VAL AB 7 HIS AB 16 1 ? 10
HELX_P HELX_P2 2 ASN B 25 ? TYR B 31 ? ASN AB 25 TYR AB 31 5 ? 7
HELX_P HELX_P3 3 ASP B 43 ? MET B 63 ? ASP AB 43 MET AB 63 1 ? 21
HELX_P HELX_P4 4 LYS B 74 ? GLN B 76 ? LYS AB 74 GLN AB 76 5 ? 3
HELX_P HELX_P5 5 ALA B 77 ? ARG B 87 ? ALA AB 77 ARG AB 87 1 ? 11
HELX_P HELX_P6 6 MET B 101 ? THR B 103 ? MET AB 101 THR AB 103 5 ? 3
HELX_P HELX_P7 7 ASN B 104 ? ALA B 123 ? ASN AB 104 ALA AB 123 1 ? 20
HELX_P HELX_P8 8 SER B 124 ? GLU B 129 ? SER AB 124 GLU AB 129 5 ? 6
HELX_P HELX_P9 9 PRO B 131 ? SER B 150 ? PRO AB 131 SER AB 150 1 ? 20
HELX_P HELX_P10 10 PRO B 167 ? LYS B 169 ? PRO AB 167 LYS AB 169 5 ? 3
HELX_P HELX_P11 11 GLU B 170 ? LEU B 180 ? GLU AB 170 LEU AB 180 1 ? 11
HELX_P HELX_P12 12 ALA B 207 ? ALA B 225 ? ALA AB 207 ALA AB 225 1 ? 19
HELX_P HELX_P13 13 HIS C 6 ? LEU C 12 ? HIS AC 6 LEU AC 12 1 ? 7
HELX_P HELX_P14 14 TYR C 29 ? GLU C 46 ? TYR AC 29 GLU AC 46 1 ? 18
HELX_P HELX_P15 15 LEU C 47 ? GLY C 51 ? LEU AC 47 GLY AC 51 5 ? 5
HELX_P HELX_P16 16 LYS C 72 ? VAL C 76 ? LYS AC 72 VAL AC 76 5 ? 5
HELX_P HELX_P17 17 ARG C 83 ? THR C 95 ? ARG AC 83 THR AC 95 1 ? 13
HELX_P HELX_P18 18 ASN C 108 ? LEU C 111 ? ASN AC 108 LEU AC 111 5 ? 4
HELX_P HELX_P19 19 SER C 112 ? ARG C 126 ? SER AC 112 ARG AC 126 7 ? 15
HELX_P HELX_P20 20 ALA C 129 ? GLU C 143 ? ALA AC 129 GLU AC 143 1 ? 15
HELX_P HELX_P21 21 ARG C 156 ? ALA C 160 ? ARG AC 156 ALA AC 160 5 ? 5
HELX_P HELX_P22 22 SER D 52 ? GLY D 69 ? SER AD 52 GLY AD 69 1 ? 18
HELX_P HELX_P23 23 SER D 71 ? LYS D 85 ? SER AD 71 LYS AD 85 1 ? 15
HELX_P HELX_P24 24 VAL D 88 ? SER D 99 ? VAL AD 88 SER AD 99 1 ? 12
HELX_P HELX_P25 25 ARG D 100 ? LEU D 108 ? ARG AD 100 LEU AD 108 1 ? 9
HELX_P HELX_P26 26 SER D 113 ? HIS D 123 ? SER AD 113 HIS AD 123 1 ? 11
HELX_P HELX_P27 27 LYS D 151 ? ASN D 154 ? LYS AD 151 ASN AD 154 5 ? 4
HELX_P HELX_P28 28 LEU D 155 ? LYS D 166 ? LEU AD 155 LYS AD 166 1 ? 12
HELX_P HELX_P29 29 ASP D 190 ? LEU D 194 ? ASP AD 190 LEU AD 194 5 ? 5
HELX_P HELX_P30 30 ASN D 199 ? TYR D 207 ? ASN AD 199 TYR AD 207 1 ? 9
HELX_P HELX_P31 31 GLU E 50 ? ARG E 64 ? GLU AE 50 ARG AE 64 1 ? 15
HELX_P HELX_P32 32 GLY E 103 ? GLY E 114 ? GLY AE 103 GLY AE 114 1 ? 12
HELX_P HELX_P33 33 ASN E 127 ? ARG E 140 ? ASN AE 127 ARG AE 140 1 ? 14
HELX_P HELX_P34 34 THR E 144 ? ARG E 150 ? THR AE 144 ARG AE 150 1 ? 7
HELX_P HELX_P35 35 ASP F 15 ? TYR F 33 ? ASP AF 15 TYR AF 33 1 ? 19
HELX_P HELX_P36 36 ARG F 71 ? ILE F 81 ? ARG AF 71 ILE AF 81 1 ? 11
HELX_P HELX_P37 37 ASP G 20 ? MET G 31 ? ASP AG 20 MET AG 31 1 ? 12
HELX_P HELX_P38 38 LYS G 35 ? LYS G 48 ? LYS AG 35 LYS AG 48 1 ? 14
HELX_P HELX_P39 39 GLU G 57 ? ASN G 68 ? GLU AG 57 ASN AG 68 1 ? 12
HELX_P HELX_P40 40 SER G 92 ? ARG G 111 ? SER AG 92 ARG AG 111 1 ? 20
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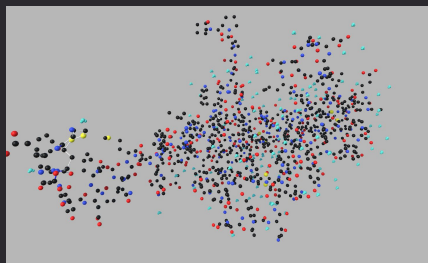


CSV File:

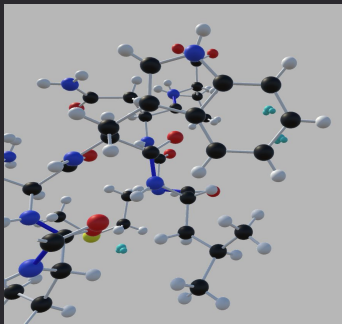
HELIX	7	16 AB			
HELIX	25	31 AB			
HELIX	43	63 AB			
HELIX	74	76 AB			
HELIX	77	87 AB			
HELIX	101	103 AB			
HELIX	104	123 AB			
HELIX	124	129 AB			
HELIX	131	150 AB			
HELIX	167	169 AB			
HELIX	170	180 AB			
HELIX	207	225 AB			
HELIX	6	12 AC			
HELIX	29	46 AC			
HELIX	47	51 AC			
HELIX	72	76 AC			
HELIX	83	95 AC			
HELIX	108	111 AC			
HELIX	112	126 AC			
HELIX	129	143 AC			
HELIX	156	160 AC			
HELIX	52	69 AD			

# Generation and Interaction

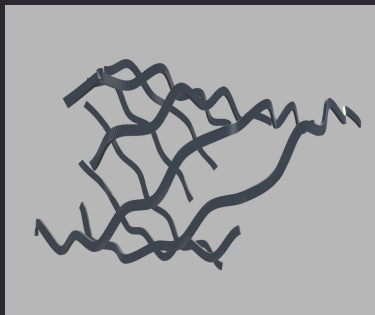
- Protein features from CSV
- Colors and Scales atoms
- Interaction Scripts attached



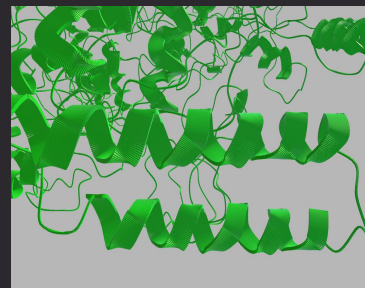
Atoms



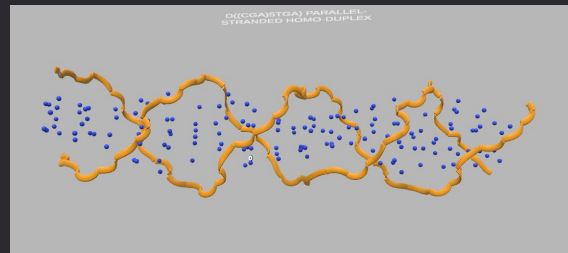
Bonds



Beta Sheets



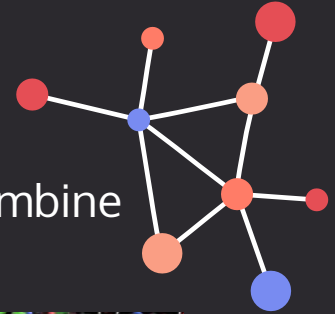
Helices



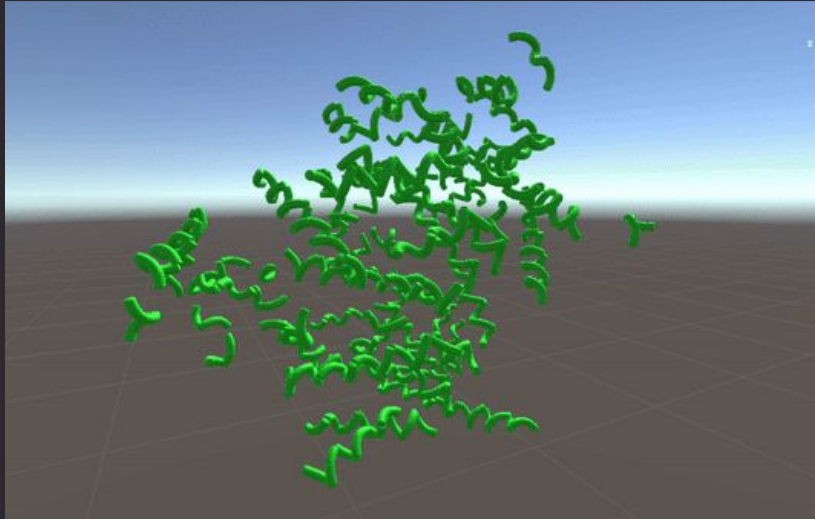
RNA



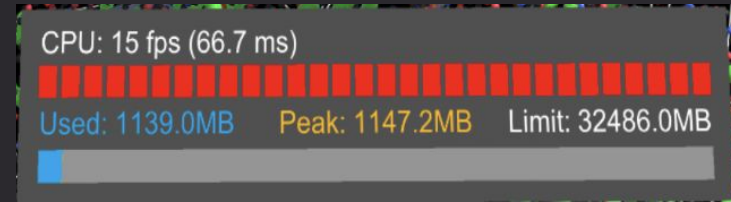
# Challenges and Optimization Efforts



Jagged curved structures → use splines to smooth out

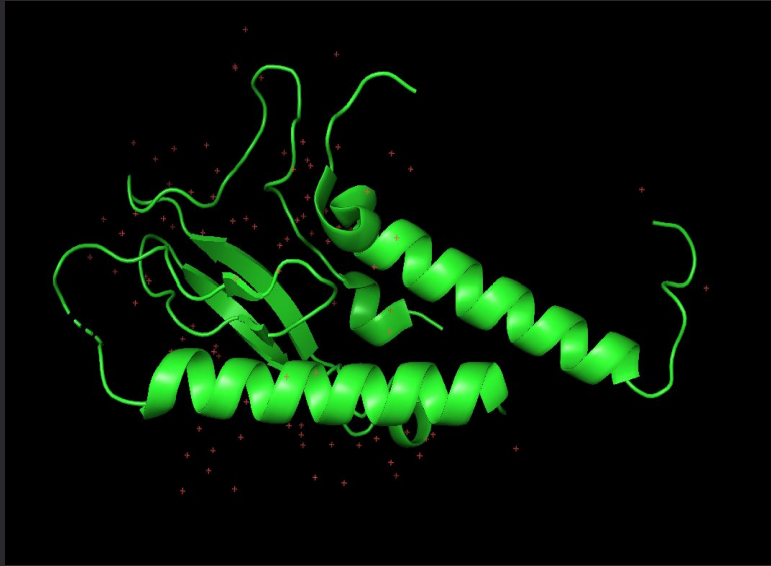


Slow model rendering → combine meshes of similar objects

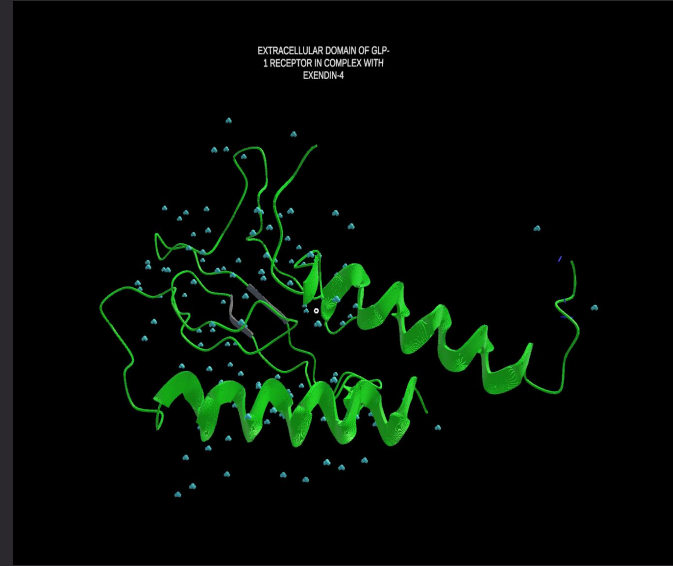




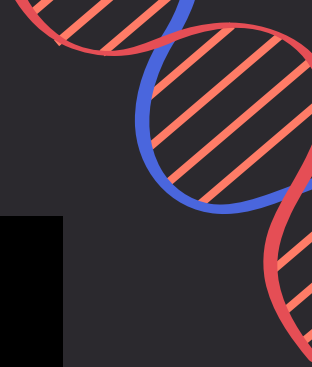
# Results



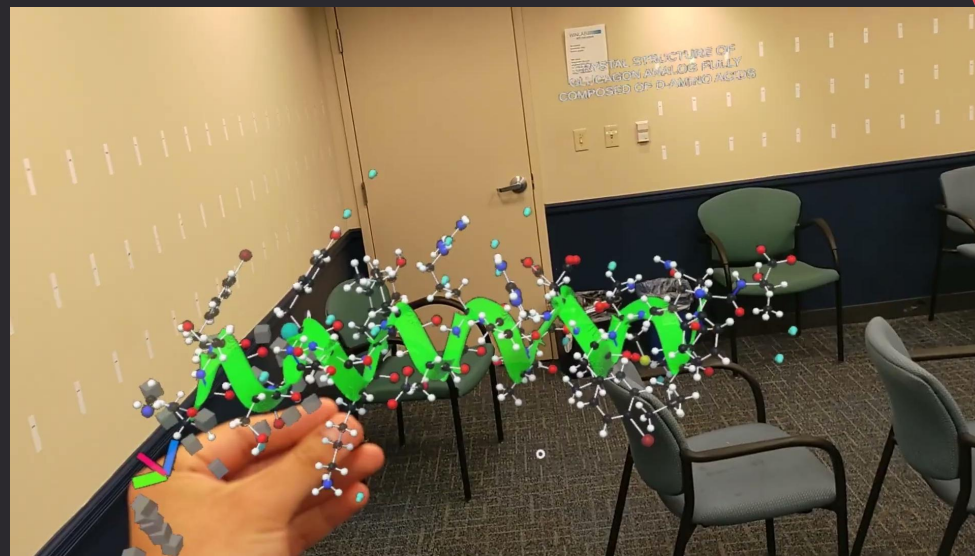
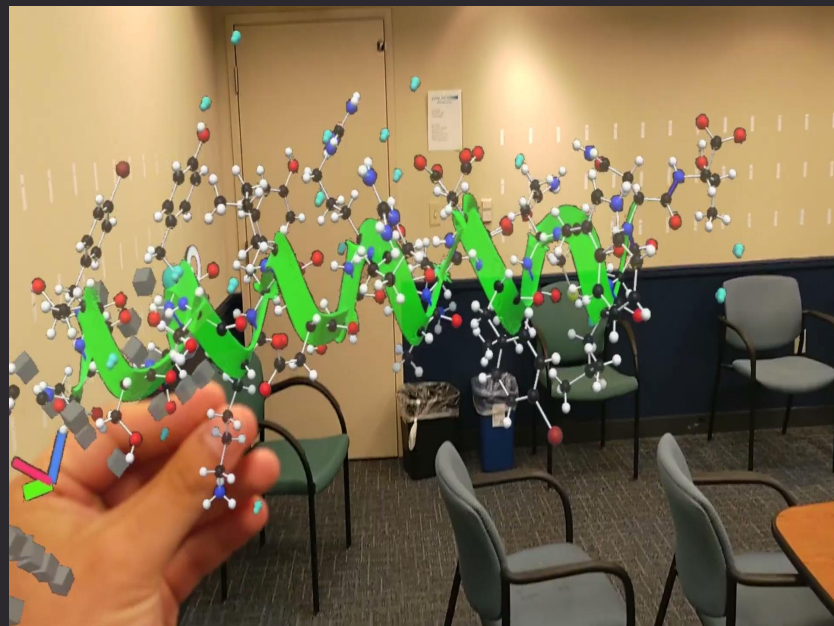
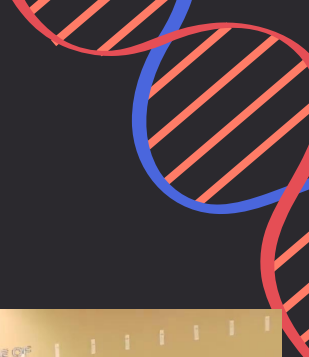
50TT PyMol Model



50TT Model in Unity

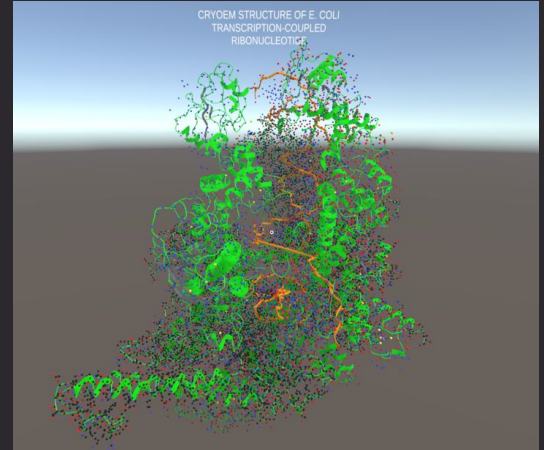


# Voice Command Feature



# Future Plans

- Greater detail in RNA and Beta-Sheets implementation
- Optimized rendering for large proteins
- Faster 3D model generation and loading time of proteins
- Deploy visualizations to a mobile application
- Feedback from professionals

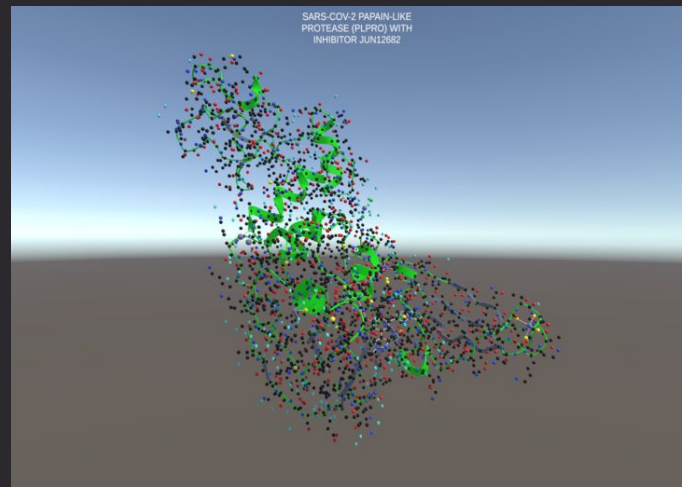


CryoEM Structure of E.Coli

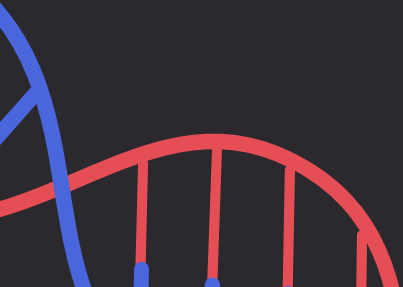
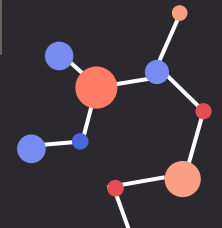
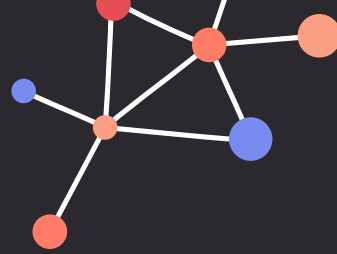


# Potential Project Applications

- Interactive Learning Experiences
- Drug Discovery and Manipulation
- Medical Training
- Public Health Awareness



SARS-CoV-2





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Wade Trappe and Ashley Guo.

Thank You!

