

References

Paper:

[MetaEuk—sensitive, high-throughput gene discovery, and annotation for large-scale eukaryotic metagenomics](#)

Two papers for the use of metagenomics in protein prediction from Journal “Science”:

1. [Protein structure determination using metagenome sequence data.](#)
2. [Big-data approaches to protein structure prediction](#)

Pictures for the Video:

1. [MetaEuk logo](#)
2. [Ocean organisms](#)
3. [laboratory cultivation](#)
4. [Ecology](#)
5. [Pharmacy](#)
6. [MMseqs2 logo](#)
7. [AlphaFold logo](#)