Theoretical Biology & Mathematical Modeling → w.r.t Java & its Advanced Bio-informatics Tools → A Simple Suggestion + Novel Bio-informatics R&D Framework → To Probe the Frontiers of COVID-19 Vaccines Research.

[Exploring The concepts of Protein Folding Mechanisms Using ANTLR]

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[I] Main Idea + Inspiration :

ANTLR → Probing Protein Folding Grammars → Advanced Algorithms

→ to develop COVID-19/Vaccine R&D → Design/Testing/Implementing Vaccine Research

→ by Using Java/Bio-Java/Spin Glass Theory Concepts

→ towards better understanding of Nano-Bio Material Systems w.r.t. COVID-19 Informatics.

Readers could easily come up with a RAPID PROTOTYPE of a flexible Bio-informatics Framework to suit the present requirement/s.

[II] Important References :

[a] https://github.com/tejdnk-2019-ShortNotes

[b] https://biojava.org/ && https://www.antlr.org/

[c] http://dx.doi.org/10.1590/S0100-879X2001000400001 –
Protein Folding Information.

[III] Acknowledgment/s :

[THE END]