

Exploring + Testing Algorithms w.r.t Protein Folding Mechanisms & Related Bio-informatics Data Using [*Alg -I : SWI-Prolog + Bio-Java] [*Alg-II : JI Prolog + Bio-Java] [*Alg-III : JI Prolog/Jruby/Ruby + Bio-ruby] → Suggesting Novel Algorithms Design w.r.t COVID-19 Informatics R&D [Using *SD + *IoT + *HPC]

[*Alg → Algorithm /*SD – Smart Devices /*IoT – Internet of Things/*HPC – High Performance Computing]

Nirmal – Informatics R&D Collaborator – USA/UK/Israel/BRICS Group of Nations.

Current Member – ante Inst UTD Dallas TX USA.

Contact_info – hmfg2014@gmail.com

[I] Main Idea + Inspiration + Introduction :

<https://www.sciencedirect.com/science/article/pii/S0732118X20302130?via%3Dihub>

<https://doi.org/10.1016/j.newideapsych.2020.100838>

[II] SWI-Prolog + JI Prolog + Bio-Java + Bio-ruby R&D Informatics Framework :

***** Please check our Short Notes on [github/Vixra/org](https://github.com/Vixra/org). With Thanks – Dr.Nirmal.

[III] Important References Used in our R&D Work :

[a] <https://github.com/tejdkn-2019-ShortNotes> – Please Check out Notes – Thanks.

[IV] Acknowledgment/s:

Non – Profit R&D Work. Inspire Others Always. Sincere Thanks to all WHO made this happen in my LIFE.

[V] Conclusion/s + Future Perspectives : One of the Pioneering R&D Efforts in this direction.
Aim is to TEST different Software Tools Using
Prolog + Java + Ruby w.r.t COVID-19 Informatics.
Rigorous testing in progress. Keep Testing. Thanks.

[THE END]