

Exploring Rust-Bio: A Fast and Safe Bio-informatics Library + E Theorem Prover [ETP] Combination as Promising Software Tools w.r.t Designing COVID-19 Next Generation DNA/RNA Sequencing Algorithms → A Simple Technical Communication Using Smart Devices + IoT + HPC Heterogeneous Computing R&D Environment/s.

Nirmal – Informatics R&D – 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 :

[a] <https://www.rust-lang.org/> [b] <https://rust-bio.github.io/>

[c] The E homepage can be found at <<http://www.e prover.org>>

[II] R&D Informatics Framework w.r.t RUST + RUSTBIO + E Theorem Prover :

{ It is very EASY to DERIVE an Advanced Bio-informatics R&D Framework Using the above mentioned Software Tools. }

[III] Important References Using RUST (((via))) our github :

[a] <https://github.com/tejdnk-2019-ShortNotes> - Many useful short notes.

[b] <https://medium.com/swlh/machine-learning-in-rust-smartcore-2f472d1ce83> -
[Machine Learning in Rust, Smartcore | by Vlad Orlov | The Startup | Medium]

[c] <https://github.com/tejdnk-2019-ShortNotes/tejdnk-Space-Medicine-Informatics-github.io/blob/master/Rust-C-GeneChip-Nir-2021.pdf>

[IV] Acknowledgment/s :

Sincere Thanks to all WHO made this happen in my LIFE.Non-Profit R&D.
Inspiring Others is Always GOOD.

[V] Conclusion/s + Future Perspectives :

One of the pioneering R&D Efforts in this HIGHLY CHALLENGING Domain using RUST + RUSTBIO + ETP.

[THE END]