[OCaml-LLVM/Polly-Owl-Satallax(Theorem Prover in OCaml) - CoqTP/OCaml - Q*cert/OCaml/Eigen] in the Context of IoT/HPC-High Performance Computing - Heterogeneous Environments to [Design+Test+Implement DNA based Theoretical Gene Therapy Informatics] R&D Framework - A Simple Suggestion & Novel Approach.

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[I] Inspiration + Introduction :

"The brain is one of the richest green fields of science. There's so much yet to be discovered." - Paul Allen.

"In my experience, each failure contains the seeds of your next success—if you are willing to learn from it." -Paul Allen. [Microsoft Co-founder]

"A multi-disciplinary team approach to drug discovery – Revolutionize The Way Drugs Are Discovered & Developed – Currently, it takes 12 years and \$2.6 billion to get a single drug to market, with the drug discovery and development process costing \$1.4 billion.Most drugs only target one gene at a time – Neurological diseases are caused by complex interactions between many genes. Many drug discoveries fail because researchers target only one gene at a time."

[Source - <u>https://www.vergegenomics.com/</u> && <u>https://www.vergegenomics.com/#current-approach</u>]

OCaml – Functional Programming Language – <u>https://ocaml.org/</u> – OCaml is an industrial strength programming language supporting functional, imperative and object-oriented styles. <u>Install OCaml</u>.

Owl – Scientific Library/Machine Learning Concepts – <u>https://devmesh.intel.com/projects/owl-an-ocaml-numerical-library/https://github.com/owlbarn/owl/opam.ocaml.org/packages/owl/</u> – Owl is an OCaml library for scientific computing and machine learning using functional programming.Owl is an OCaml library for scientific computing. It enables academic researchers to fast prototype machine learning algorithms and construct deep neural networks with very concise code. It also facilitates industry programmers to develop robust analytical applications using functional language at a large scale.

Q*cert – Query Compiler – Compilation and verification of data languages <u>https://querycert.github.io/</u> Q*cert, a framework for the development and verification of query compilers. It supports a rich data model and includes an extensive compilation pipeline 'out of the box'.Q*cert is built using the Coq proof assistant (<u>https://coq.inria.fr</u>). A significant subset of the provided compilation pipeline has been mechanically checked for correctness.

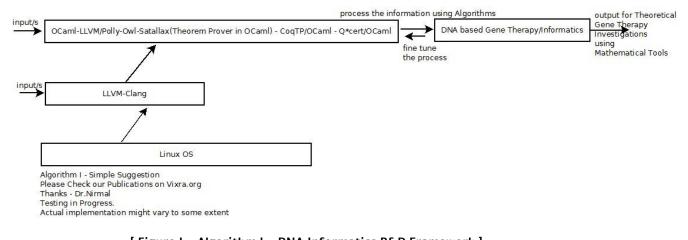
Satallax – Theorem Prover – *satallaxprover.org* – / *www.ps.uni-saarland.de*/~*cebrown*/*satallax* – *https://en.wikipedia.org*/*wiki*/*Automated_theorem_proving* –/ *https://link.springer.com/chapter/10.1007/978-3-642-31365-3_11* – Satallax is an automated theorem prover for higher-order logic. The particular form of higher-order logic supported by Satallax is Church's simple type theory with extensionality and choice operators.

Coq – **Theorem Prover** – *https://coq.inria.fr* – Coq is a formal proof management system. It provides a formal language to write mathematical definitions, executable algorithms and theorems together with an environment for semi-interactive development of machine-checked proofs.

LLVM – **http://llvm.org/** – The LLVM Project is a collection of modular and reusable compiler and tool chain technologies. LLVM began as a <u>research project</u> at the <u>University of Illinois</u>.

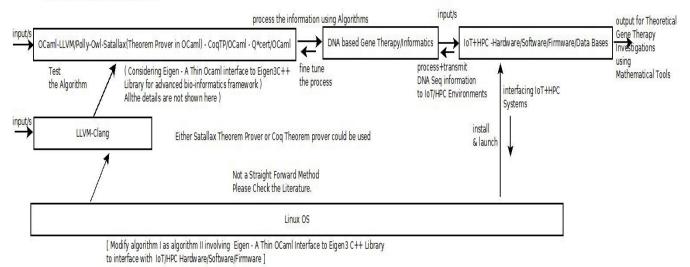
[II] R&D of DNA based Informatics Framework Implementation :

ALGORITHM I & INFORMATICS FRAMEWORK BASED ON OCAML/LLVM-POLLY/SATALLAX-THEOREM PROVER COQ THEOREM PROVER/Q*CERT



[Figure I – Algorithm I – DNA Informatics R&D Framework] Approximate Algorithm Only – Actual Implementation Might Vary to Some Extent – Please Note. Testing in Progress – Please Check some of our publications on Gene Therapy. We are interested in Targeting Neurosciences.

ALGORITHM II & INFORMATICS FRAMEWORK BASED ON OCAML/LLVM-POLLY/SATALLAX THEOREM PROVER COQ THEOREM PROVER/Q*CERT/OWL/EIGEN-INTERFACING WITH IoT/HPC HARDWARE/SOFTWARE/FIRMWARE TO PROBE GENE THERAPY USING MATHEMATICAL CONCEPTS/ALGORITHMS



Please Note the difference between Algorithm I & Algorithm II.

Algorithm II - Modified Algorithm I

[Figure II – Algorithm II – DNA Informatics R&D Framework] Approximate Algorithm Only – Actual Implementation Might Vary to Some Extent – Please Note. We are interested in Targeting Neurosciences.

[III] Related R&D Information on Mathematics+Software Used :

https://github.com/colinbenner/ocaml-llvm

https://ocaml.org/

http://caml.inria.fr/

https://www.tutorialspoint.com/compile_ocaml_online.php

https://github.com/ocaml/ocaml

https://try.ocamlpro.com

https://github.com/ocaml

https://discuss.ocaml.org

https://ocaml.xyz

https://reasonml.github.io/docs/en/comparison-to-ocaml.html

https://fdopen.github.io/opam-repository-mingw/installation

https://github.com/querycert/qcert

https://en.wikipedia.org/wiki/Coq

http://www.ps.uni-saarland.de/~cebrown/satallax/

https://polly.llvm.org -/ https://polly.llvm.org/get_started.html / https://github.com/llvm-mirror/polly -/ polly.llvm.org/projects.html / <u>Polly - Polyhedral optimizations for LLVM</u> - Polly is a high-level loop and data-locality optimizer and optimization infrastructure for LLVM. It uses an abstract mathematical representation based on integer polyhedra to analyze and optimize the memory access pattern of a program.

https://github.com/owlbarn/owl - Owl - OCaml Scientific and Engineering Computing @ http://ocaml.xyz

https://github.com/owlbarn/eigen – Eigen – A Thin OCaml Interface to Eigen₃ C++ Library – Simply put, Eigen is a very thin OCaml interface to Eigen₃ C++ template library. This library is used by another OCaml numerical library –– Owl to provide basic support for both dense and sparse matrix operations.

[Owl's OCaml Interface to Eigen3 C++ Library - http://eigen.tuxfamily.org/]

https://bioengineer.org/new-gene-therapy-research-at-neuroscience-2019/

https://www.elsevier.com/books/gene-therapy-in-neurological-disorders/...

https://www.novartis.com/our-focus/cell-and-gene-therapy

https://en.wikipedia.org/wiki/Gene_therapy

[IV] Conclusion/s With Future Perspectives :

Transforming drug discovery using AI/ML(artificial intelligence/machine learning) is very much challenging & promising ,hence, "Leveraging Machine Learning " to drastically reduce the time and cost of drug discovery is the need of the hour. To the best of our knowledge we have used advanced Mathematical Concepts/Theorem Provers/Functional Programming/Ocaml Language/LLVM/Polly to "Leverage" the Health Care Industry R&D domains.We, sincerely hope and are quite sure that in the near future, many researchers would certainly find this approach useful in their R&D Works.

*** Algorithms I & II gave us an opportunity to TEST the performances of Satallax Theorem Prover/Coq Theorem Prover – by studying their underlying logic & mechanisms in OcamI based HPC Environments.

[V] Acknowledgment/s :

Non-Profit R&D. Special Thanks to all WHO made this happen – My Friends+Mentors.

[VI] References :

- [a] <u>https://www.vergegenomics.com/publications</u>
- [b] <u>http://www.vixra.org/author/nirmal_tej_kumar</u>
- [c] <u>http://www.vixra.org/author/n_t_kumar</u>
- [d] <u>http://www.vixra.org/author/d_n_t_kumar</u>
- [e] <u>http://www.vixra.org/author/nirmal</u>
- [f] <u>https://www.semanticscholar.org/author/Nirmal-Tej-Kumar/12354503/suggest</u> [Semanticsscholar Indexing]

[From - Allen Institute/AI Tool]

[g] https://allenai.org/ - Allen Institute Website -Useful in Searching R&D Information Using Semanticscholar.

*** A free tool for uncovering supplement-drug interactions: supp.ai | Powered by Semantic Scholar .

[THE END]