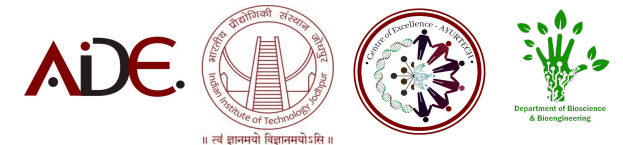


Development of frameworks for evidence-based practice of traditional medicine in integrative medicine settings



Integrating Traditional Knowledge in Evidence Based Medicine pGI January 7-10 2024

Mitali Mukerji, PhD, FASc, FNASc
Professor & Head,
Department of Bioscience & Bioengineering
Affiliate Faculty, School of Artificial Intelligence and Data
Science (AIDE)
Indian Institute of Technology Jodhpur

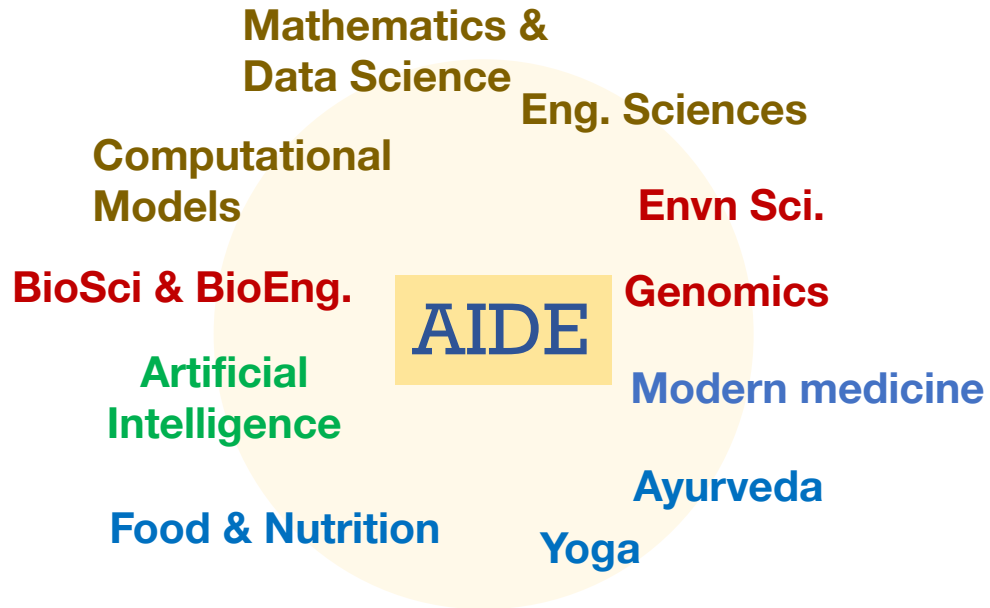


Transdisciplinary Centre of Excellence in Integrative Precision

Health

Aim

Establishment of AI driven integrative framework for population and individual risk stratification and early actionable precision health interventions



The School of Artificial Intelligence & Data Science (**AIDE**) would nucleate a transdisciplinary framework that would have application in:

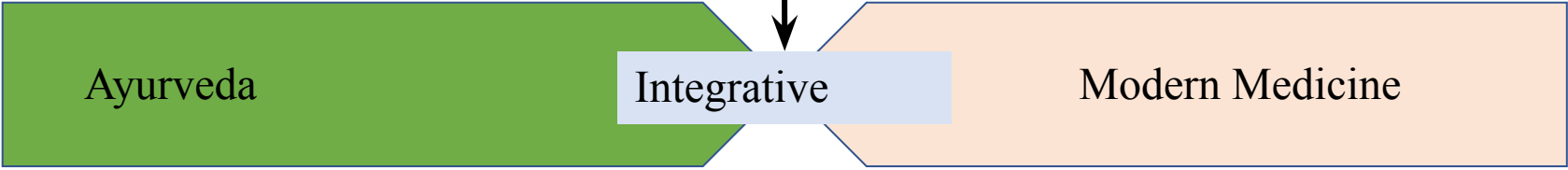
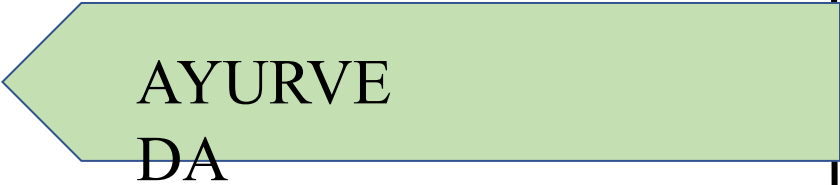
- Phenomics based non-invasive risk stratification
- Integration of principles and practice of Ayurveda with other knowledge systems
- Management of rare disease and common disease
- Development of an objective framework for Integrative medicine
- Precision health with specific focus in arid regions

Integrative medicine_ a win-win situation

*Reactive -target symptoms
Treat molecular imbalances - organ specific
Specific molecules*



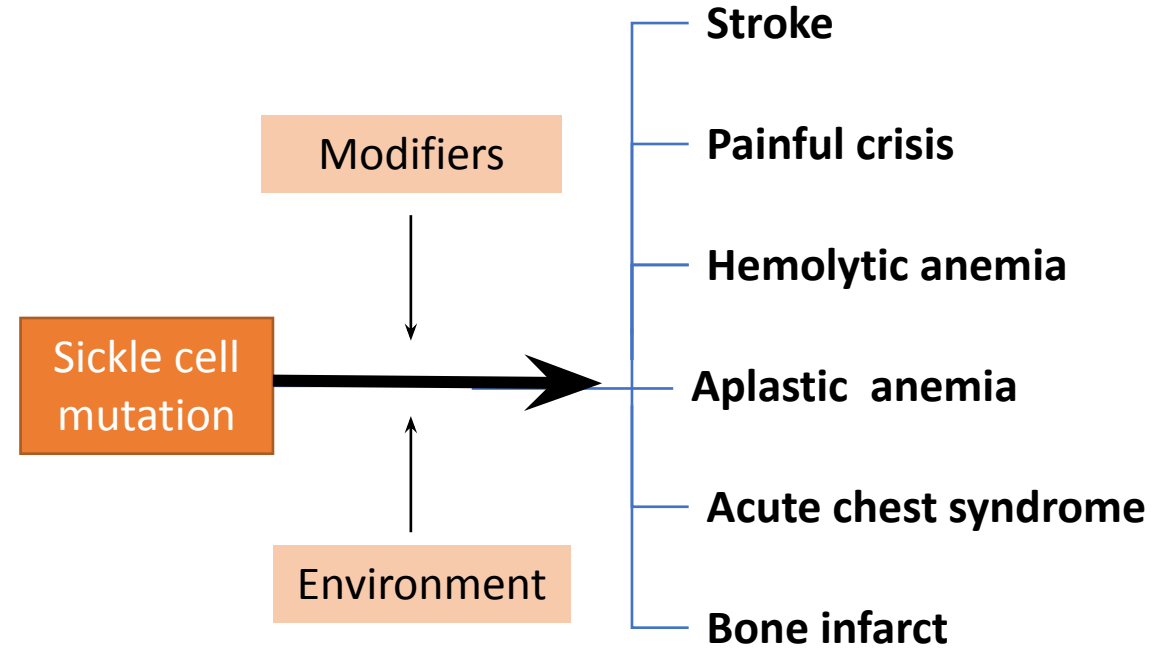
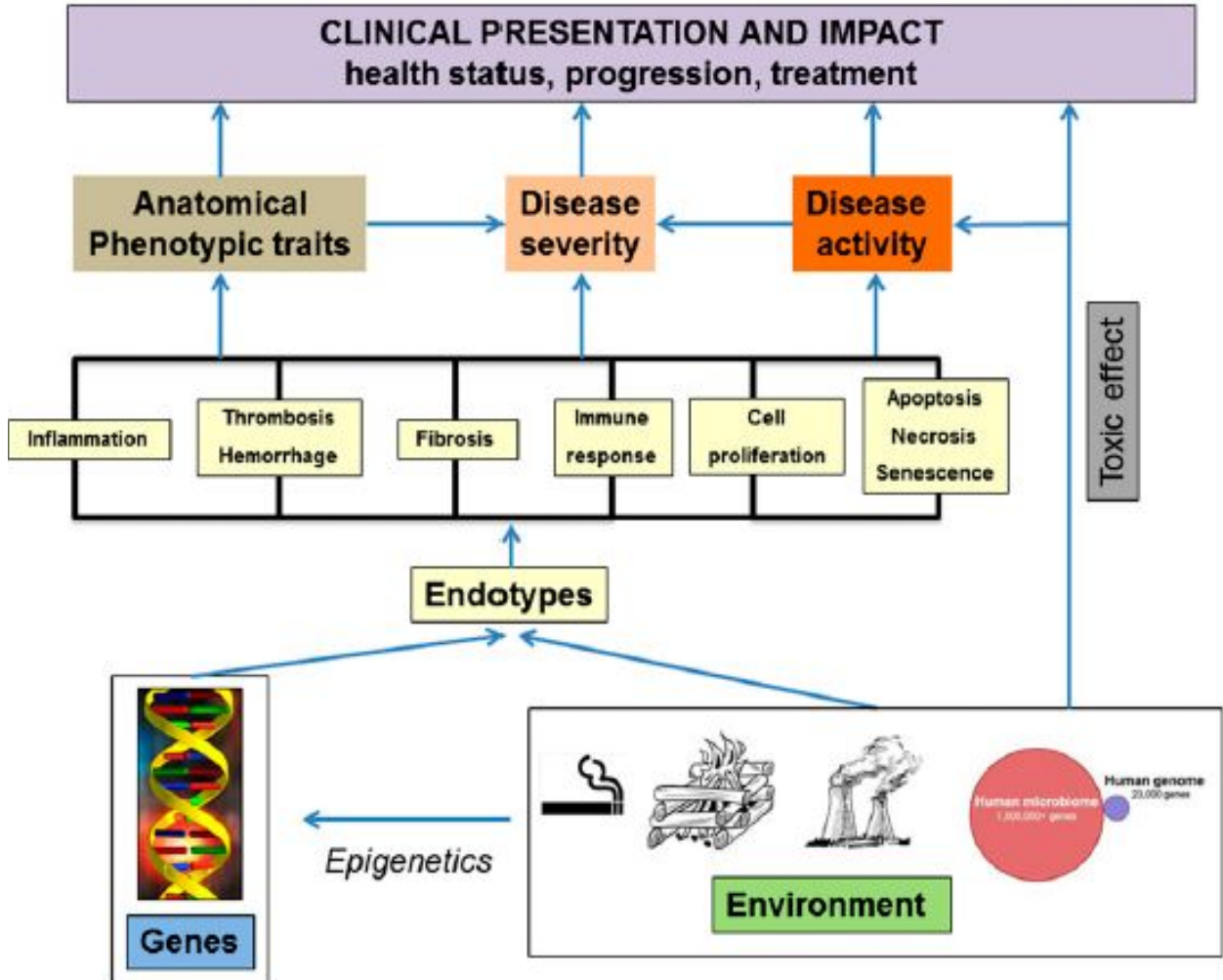
*Proactive – targets doshic imbalances
Aims at reversal - system’s perspective,
Poly-pharmacology
Holistic & individualized, based on Prakriti*



Interoperable, complementary, synergistic, personalized options

Stratification in disease management

Converging themes in diseases



Challenges

- Modifiers ?
- Endophenotypes?
- Targeted interventions ?

Agusti, Ann Am Thorac Soc Vol 10, Supplement, pp S125–S130, Dec 2013

(Shatkryikala ?)

(Samprapti ?)

A Paradigm shift from population based to individualized approach

Population response based

IMPRECISION MEDICINE

For every person they do help (blue), the ten highest-grossing drugs in the United States fail to improve the conditions of between 3 and 24 people (red).

1. **ABILIFY** (aripiprazole)
Schizophrenia



2. **NEXIUM** (esomeprazole)
Heartburn



3. **HUMIRA** (adalimumab)
Arthritis



4. **CRESTOR** (rosuvastatin)
High cholesterol



5. **CYMBALTA** (duloxetine)
Depression



6. **ADVAIR DISKUS** (fluticasone propionate)
Asthma



7. **ENBREL** (etanercept)
Psoriasis



8. **REMICADE** (infliximab)
Crohn's disease



9. **COPAXONE** (glatiramer acetate)
Multiple sclerosis



10. **NEULASTA** (pegfilgrastim)
Neutropenia



Based on published number needed to treat (NNT) figures. For a full list of references, see Supplementary Information at go.nature.com/4dr78f.

Personalised

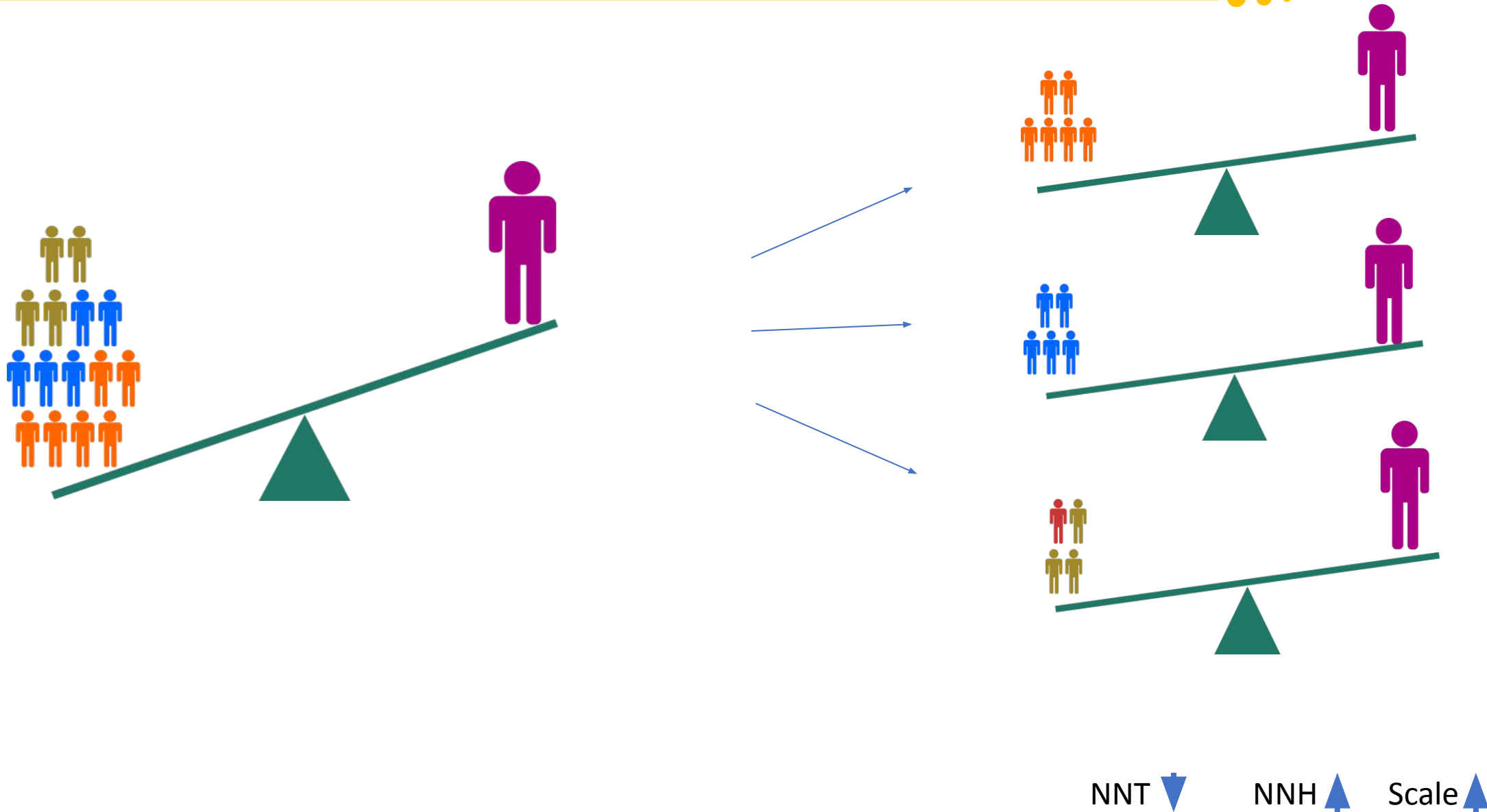


30 APRIL 2015 | VOL 520 | NATURE | 609

Time for one-person trials

Precision medicine requires a different type of clinical trial that focuses on individual, not average, responses to therapy, says **Nicholas J. Schork**.

Gap in knowledge : Health Stratification and Affordability



Definition of Genetic individuality - Modern vs

Ayurveda

Human Genome Project Surprises

There is no "The HUMAN Genome"

ESSAY

GENOME-SEQUENCING ANNIVERSARY

A Celebration of the Genome, Part III

What Does a "Normal" Human Genome

Praus et al. *Journal of Translational Medicine* (2023) 21:41
<https://doi.org/10.1186/s12967-022-03855-0>

Journal of
Translational Medicine

amag.org

RESEARCH

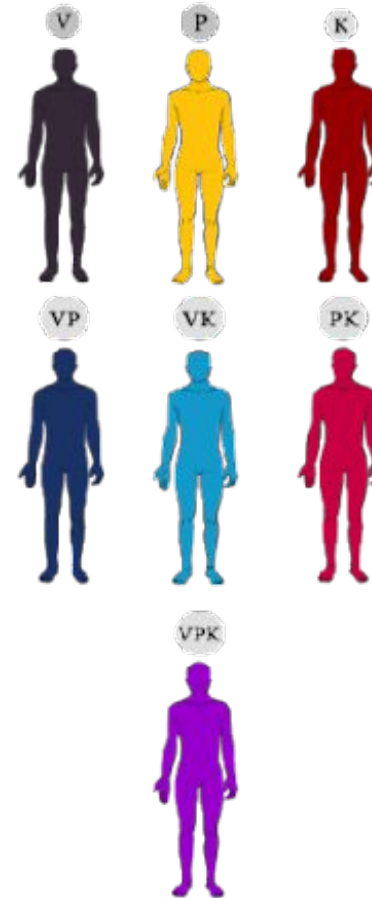
Open Access

Panomics reveals patient individuality as the major driver of colorectal cancer progression

Friederike Praus^{1†}, Axel Künstner^{2†}, Thorben Sauer¹, Michael Kohl^{1,2}, Katharina Kern¹, Steffen Deichmann³, Ákos Végvári^{4,5}, Tobias Keck³, Hauke Busch², Jens K. Habermann^{1,6} and Timo Gemoll^{1*}



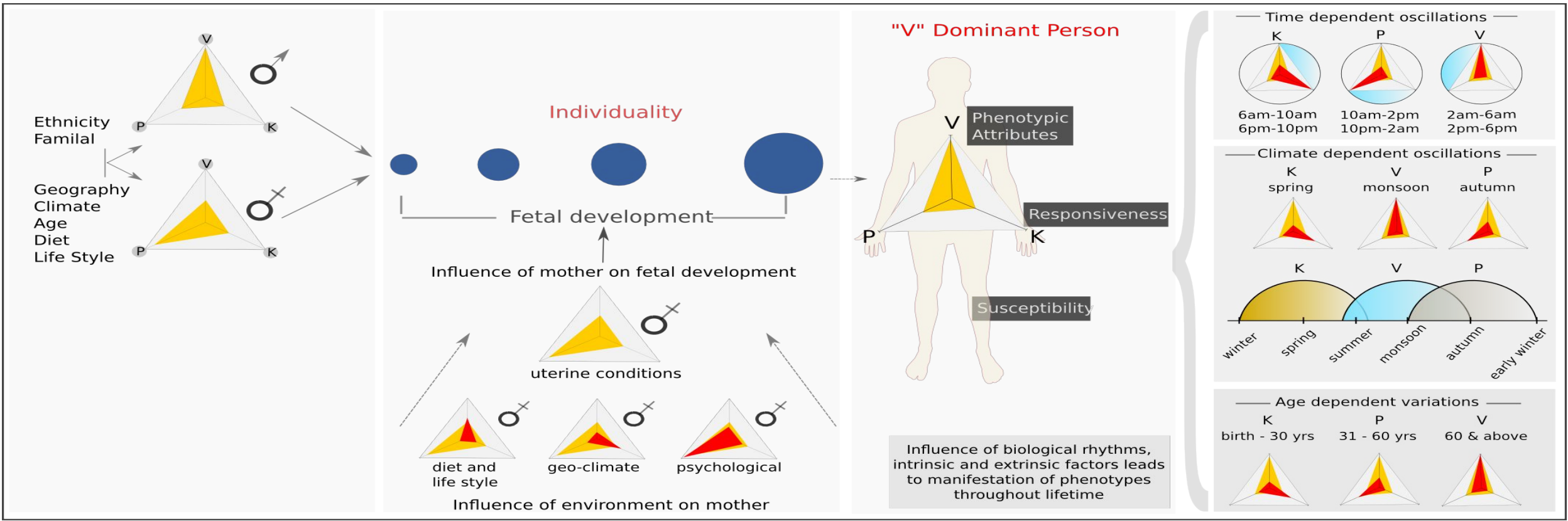
7 Prakriti types in a population



- Health
- Response
- Disease Trajectory
- Individualised management

Prasher B., Gibson G. and Mukerji M. 2016
Genomic insights into ayurvedic and western approaches to personalized medicine. J. Genet. 95, 209–228

Prakriti - GIS of an individual



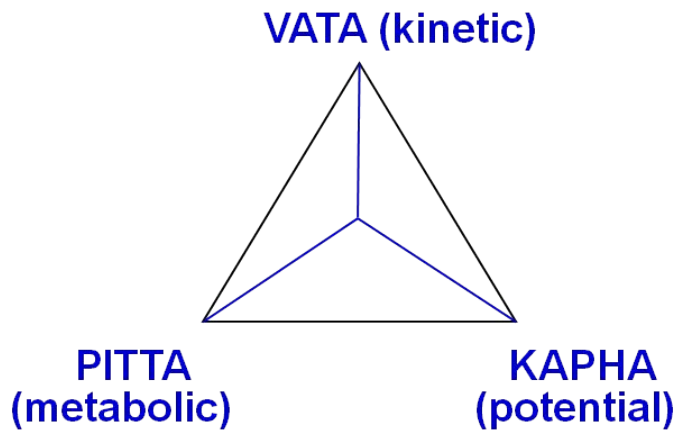
शुक्रशोणितसंयोगे यो भवेद्दोष उत्कटः ॥
 प्रकृतिर्जायते तेन तस्या मे लक्षणं शृणु ॥६३॥
genetics

तत्र प्रकृत्यादीन् भावाननुव्याख्यास्यामः । तद्यथा—शुक्रशोणितप्रकृतिं, कालगर्भाशयप्रकृतिं, आतुराहारविहारप्रकृतिं, महाभूतविकारप्रकृतिं च गर्भशरीरमपेक्षते । एतानि हि येन येन दोषेणाधिके-
epigenetics

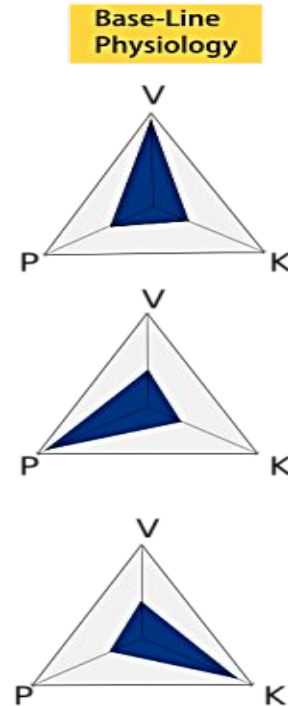
तत्र प्रकृतिर्जातिप्रसक्ता च, कुलप्रसक्ता च, देशानुपातिनी च, कालानुपातिनी च, वयोऽनुपातिनी च, प्रत्यात्मनियता चेति । जातिकुलदेशकालवयःप्रत्यात्मनियता हि तेषां तेषां पुरुषाणां ते ते भावविशेषा भवन्ति ॥ ५ ॥
ethnicity **familial** **geography** **time** **age**
Individual variation

Tridoshas: Functional attributes

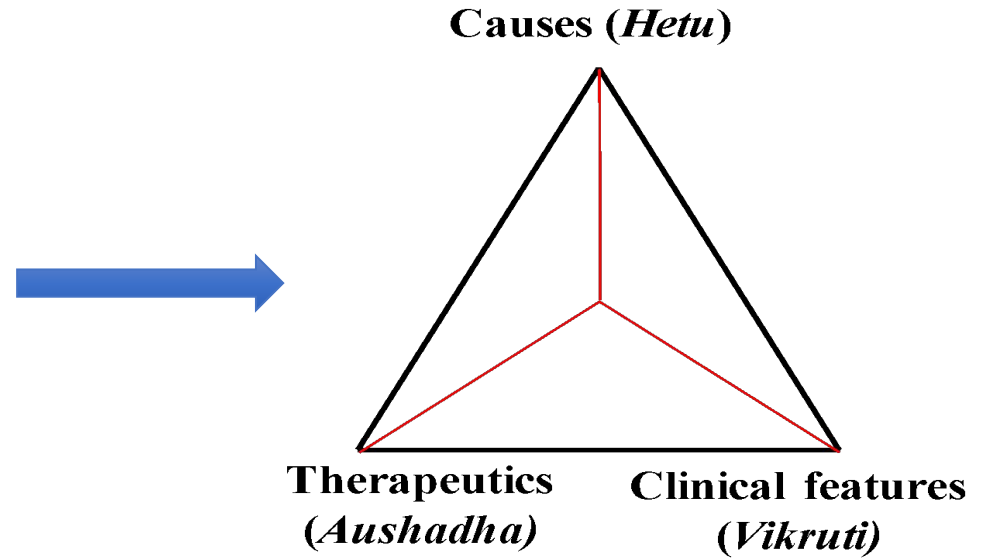
Tridosha



Doshas in Prakriti



Application of Dosha understanding in Trisutra



Basis of translation for individualised management of health & disease

Treatment evolved by “Yukti” (reasoning)

Yukti (युक्ति)-

मात्राकालाश्रया युक्तिः, सिद्धिर्युक्तौ प्रतिष्ठिता। तिष्ठत्युपरि युक्तिज्ञो द्रव्यज्ञानवतां सदा॥
चरक संहिता सूत्र स्थान २/१६

Therapeutic aptness or rational therapy depends on the matra (dose) and kala(time). Success of the treatment depends on yukti (rational therapeutics). But prior to the application of yukti (rational therapeutics) the physician should always possess complete knowledge about drugs.

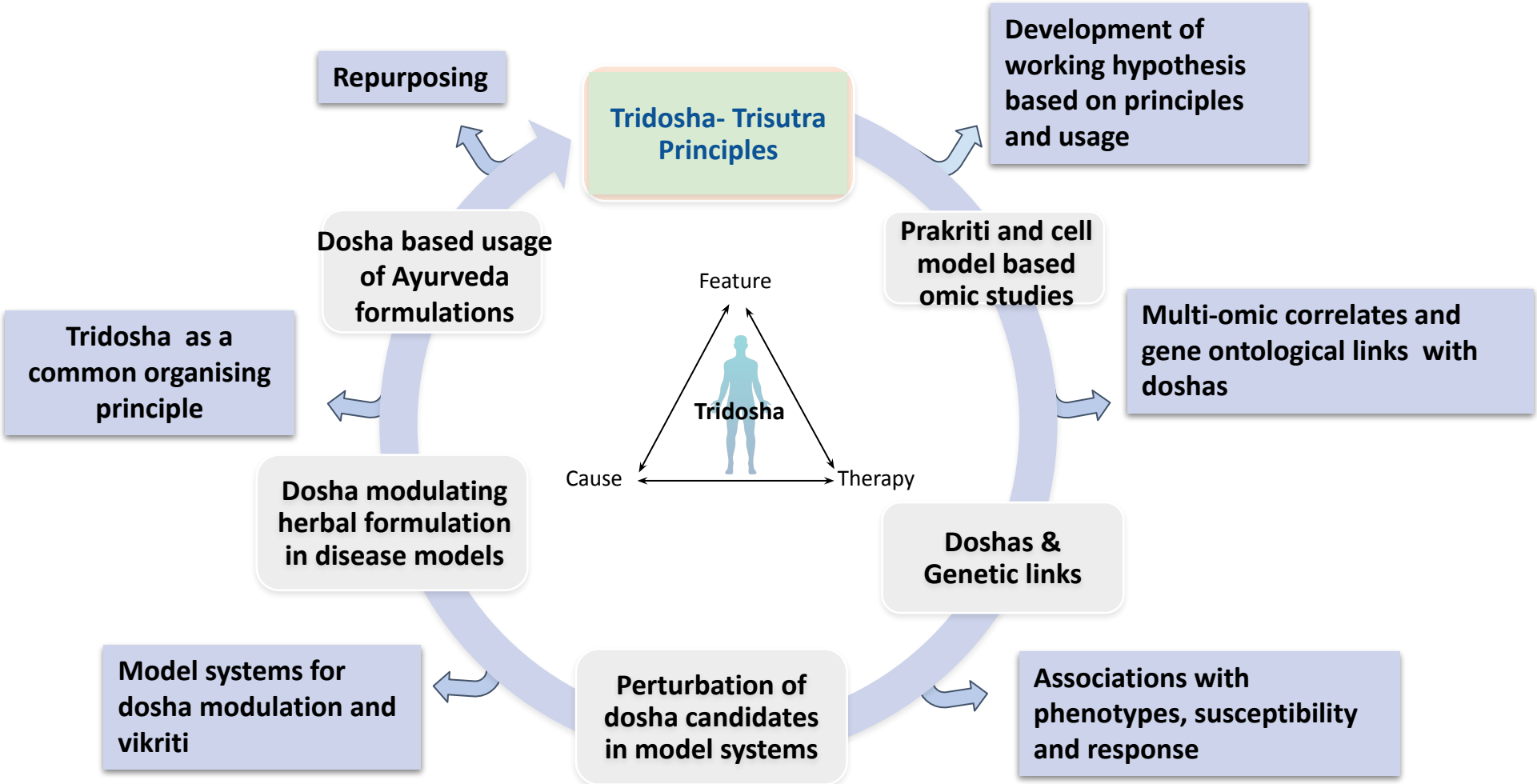
सर्वशरीरचरास्तु वातपित्तश्लेष्माणः सर्वस्मिञ्छरीरे कुपिताकुपिताः शुभाशुभानि कुर्वन्ति—

प्रकृतिभूताः शुभान्युपचयबलवर्णप्रसादादीनि, अशुभानि पुनर्विकृतिमापन्ना विकारसंज्ञकानि ॥ ९ ॥

c.su.20/9

Operational framework for exploring “trisutra” ayurveda

Summary



Molecular correlates of *Trisutra* framework : platform for translation

Differences between constitution types - Hypoxia response

Hypoxia Response in Asthma

Differential Modulation on Inflammation and Epithelial Injury

Tanveer Ahmad¹, Manish Kumar¹, Ulaganathan Mabalirajan¹, Bijay Pattnaik¹, Shilpi Aggarwal¹, Ranjana Singh¹, Suchita Singh¹, Mitali Mukerji¹, Balam Ghosh¹, and Anurag Agrawal¹

AMERICAN JOURNAL OF RESPIRATORY CELL AND MOLECULAR BIOLOGY VOL 47 2012

Cause

EGLN1 involvement in high-altitude adaptation revealed through genetic analysis of extreme constitution types defined in Ayurveda

Shilpi Aggarwal², Sapna Negi², Pankaj Jha², Prashant K. Singh², Tsering Stobdan², M. A. Qadar Pasi², Saurabh Ghosh², Anurag Agrawal², Indian Genome Variation Consortium², Bhavana Prasher^{2,4*}, and

Feature

*Genomics and Molecular Medicine, Institute of Genomics and Integrative Biology, Council of Scientific and Industrial Research (CSIR), New Delhi 110007, India; ²Human Genetics Unit, Indian Statistical Institute, Kolkata 700108, India; and ⁴Planning and Performance Division, Council of Scientific and Industrial Research (CSIR), New Delhi 110007, India

Aggarwal et al. *J Transl Med* (2015) 13:184
DOI 10.1186/s12967-015-0542-9



RESEARCH

Open Access

Combined genetic effects of *EGLN1* and *VWF* modulate thrombotic outcome in hypoxia revealed by Ayurgenomics approach

Shilpi Aggarwal^{1†}, Atish Gheware^{2,4†}, Anurag Agrawal^{1,4}, Saurabh Ghosh³, Indian Genome Variation Consortium¹, Bhavana Prasher^{2,4*} and Mitali Mukerji^{1,2,4*}

Gheware et al. *Respir Res* (2021) 22:99
https://doi.org/10.1186/s12931-021-01698-9



Respiratory Research

RESEARCH

Open Access

Adhatoda Vasica attenuates inflammatory and hypoxic responses in preclinical mouse models: potential for repurposing in COVID-19-like conditions

Atish Gheware^{1,2,3,7}, Dhvani Dholakia^{1,3,7}, Sadasivam Kannan⁵, Lipsa Panda^{4,7}, Ritu Rani^{1,2,3,7}, Bijay Ranjan Pattnaik⁴, Vaibhav Jain^{4,7}, Yash Parekh⁶, M. Ghalib Enayathullah⁶, Kiran Kumar Bokara^{6,7}, Venkatesan Subramanian^{5,7}, Mitali Mukerji^{1,2,3,7}, Anurag Agrawal^{4,7*} and Bhavana Prasher^{1,2,3,7*}

Therapy



AMERICAN JOURNAL OF PHYSIOLOGY
LUNG CELLULAR AND MOLECULAR PHYSIOLOGY

Am J Physiol Lung Cell Mol Physiol 320:L000–L000, 2021.
First published February 10, 2021; doi:10.1152/ajplung.00511.2020

RESEARCH ARTICLE

Lung Diseases in Reverse Translation: Bedside to the Bench

Adhatoda vasica rescues the hypoxia-dependent severe asthma symptoms and mitochondrial dysfunction

Prakriti, dosha and cellular outcomes

JOURNAL OF TRANSLATIONAL MEDICINE **IMPACT FACTOR 3.41** Log on/register
BioMed Central home | Journals A-Z | Feedback | Support | My details

Home | Browse articles | Supplements | Series | Search | Weblinks | Submit article | My JTM | About JTM

Top Research **Highly accessed** **Open Access** **Journal of Translational Medicine** Volume 6

Abstract **Whole genome expression and biochemical correlates of extreme constitutional types defined in Ayurveda**

Background

Methods **Bhavana Prasher¹***, **Sapna Negi¹***, **Shilpi Aggarwal¹**, **Amit K Mandal¹**, **Tav P Sethi¹**, **Shailaja R Deshmukh²**, **Sudha G Purohit²**, **Shantanu Sengupta¹**, **Sangeeta Khanna¹**, **Farhan Mohammad¹**, **Gaurav Garg¹**, **Samir K Brahmachari¹**, **Indian Genome Variation Consortium¹** and **Mitali Mukerji¹**

Results

Discussion **1** Genomics and Molecular Medicine, Functional Genomics Unit, Institute of Genomics and Integrative Biology (CSIR), Mall Road, 110007 Delhi, India

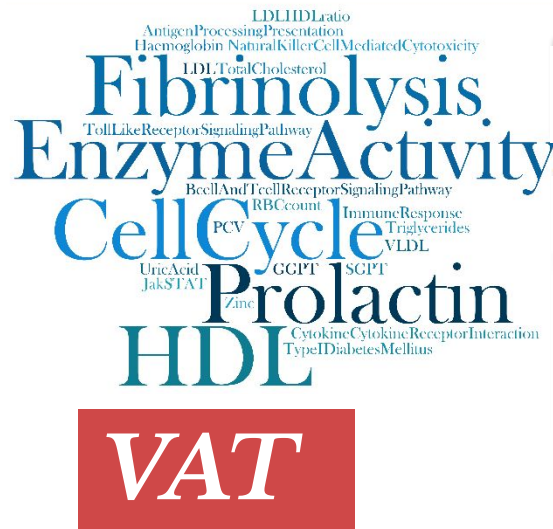
Conclusion **2** Department of Statistics, University of Pune, 411007 Pune, India

Competing interests author email corresponding author email * Contributed equally

Viewing options:
■ Abstract
■ Full text
■ PDF (652KB)
■ Additional files

Associated material:
■ Readers' comments
■ PubMed record

Related literature:
■ Articles citing this article on Google Scholar



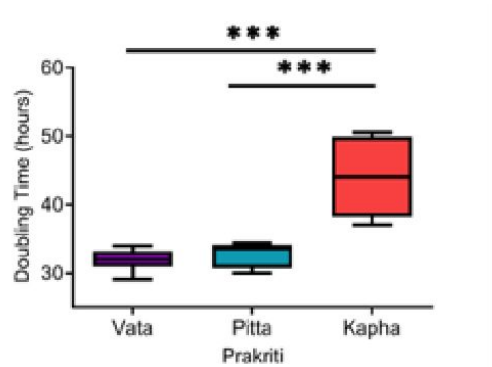
Vata regulates cell division and morphogenesis

शरीराध्यक्षास्तु परमाणुभेदेनापरिसंख्येया भवन्ति, अतिबहुत्वाद्दतिसौक्ष्म्याद्तीन्द्रियत्वाच्च । तेषां संयोगविभागे परमाणूनां कारणं वायुः कर्मस्वभावश्च ॥ १७ ॥

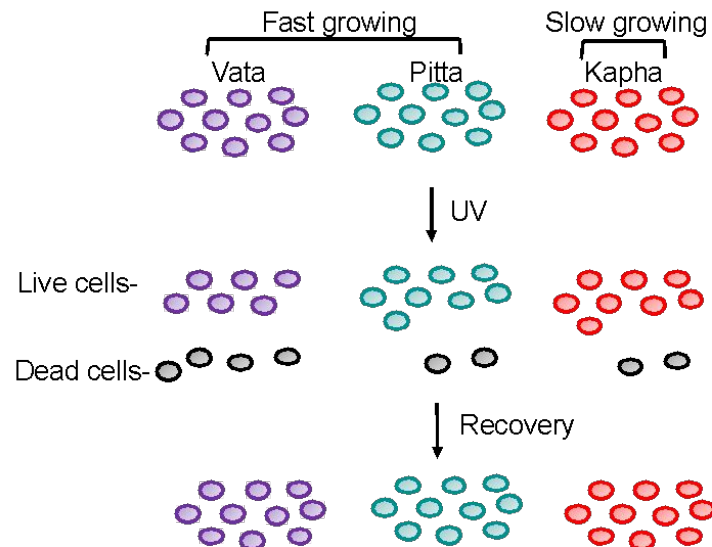
c.sa.7/17

VAT **PITT** **KAPH**

Inter-individual variability in cell proliferation rates : Who cares?



N=3 (Vata & Kapha), N=2 (Pitta); n=3, *** = P<0.0001



CELL CYCLE
<https://doi.org/10.1080/15384101.2021.1909884>

Taylor & Francis
 Taylor & Francis Group

Check for updates

RESEARCH PAPER

Baseline cell proliferation rates and response to UV differ in lymphoblastoid cell lines derived from healthy individuals of extreme constitution types

Sumita Chakraborty, Sunanda Singhmar, Dayanidhi Singh, Mahua Maulik, Rutuja Patil, Satyam Kumar Agrawal, Anushree Mishra, Madeeha Ghazi, Archana Vats, Vivek T Natarajan, Sanjay Juvekar, Bhavana Prasher, and Mitali Mukerji

¹Centre of Excellence for Applied Development of Ayurveda Prakriti and Genomics, CSIR-Institute of Genomic Delhi, India; ²CSIR Ayurgenomics Unit-TRISUTRA, CSIR-Institute of Genomics & Integrative Biology, New Delhi, Molecular Medicine, CSIR-Institute of Genomics & Integrative Biology, New Delhi, India; ³Academy of Scientific (AcSIR), Ghaziabad, India; ⁴Department of Biological Sciences, Indian Institute of Science Education & Research, West Bengal, India; ⁵Vadu Rural Health Program, KEM Hospital Research Centre, Pune, Maharashtra, India; ⁶Emerging Sciences (SPES), Baddi University of Emerging Science and Technology (BUEST), Baddi, Himachal Pradesh

SCIENTIFIC
 REPORTS
 nature research

Check for updates

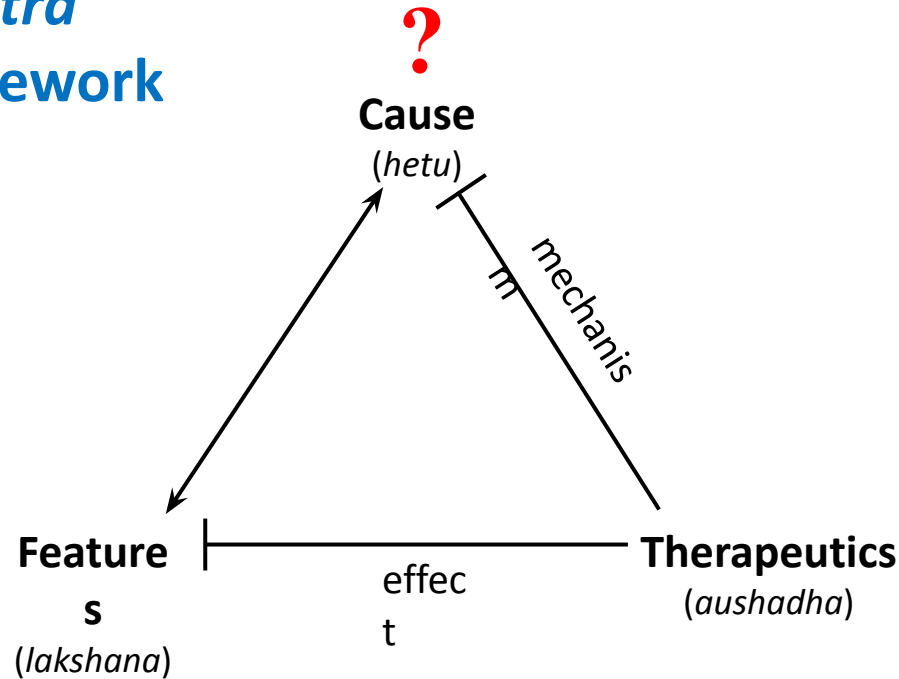
Lithium response in bipolar disorder correlates with improved cell viability of patient derived cell lines

Pradip Paul¹, Shruti Iyer², Ravi Kumar Nadella¹, Rashmitha Nayak¹, Anirudh S. Chellappa¹, Sheetal Ambardar^{2,3}, Reeteka Sud¹, Salil K. Sukumaran¹, Meera Purushottam¹, Sanjeev Jain^{1,3}, ADBS Consortium (ADBS: The Accelerator program for Discovery in Brain disorders using Stem cells)* & Biju Viswanath¹✉

Lithium is an effective, well-established treatment for bipolar disorder (BD). However, the mechanisms of its action, and reasons for variations in clinical response, are unclear. We used neural precursor cells (NPCs) and lymphoblastoid cell lines (LCLs), from BD patients characterized for clinical response to lithium (using the "Alda scale" and "NIMH Retrospective Life chart method"), to interrogate cellular phenotypes related to both disease and clinical lithium response. NPCs from two biologically related BD patients who differed in their clinical response to lithium were compared with healthy controls. RNA-Seq and analysis, mitochondrial membrane potential (MMP), cell viability, and cell proliferation parameters were assessed, with and without *in vitro* lithium. These parameters were also examined in LCLs from 25 BD patients (16 lithium responders and 9 non-responders), and 12 controls. MMP was lower in both NPCs and LCLs from BD; but it was reversed with *in vitro* lithium only in LCLs, and this was unrelated to clinical lithium response. The higher cell proliferation observed in BD was unaffected by *in vitro* lithium. Cell death was greater in BD. However, LCLs from clinical lithium responders could be rescued by addition of *in vitro* lithium. *In vitro* lithium also enhanced *BCL2* and *GSK3B* expression in these cells. Our findings indicate cellular phenotypes related to the disease (MMP, cell proliferation) in both NPCs and LCLs; and those related to clinical lithium response (cell viability, *BCL2/GSK3B* expression) in LCLs.

Exploring molecular connectivity through Ayurveda based formulations

Trisutra framework



- Repurposing of Ayurveda formulation in the norm?
- Is there a cross-talk between pathways ?
- Is there a molecular basis?
- Advantage of whole plant extract is being realized



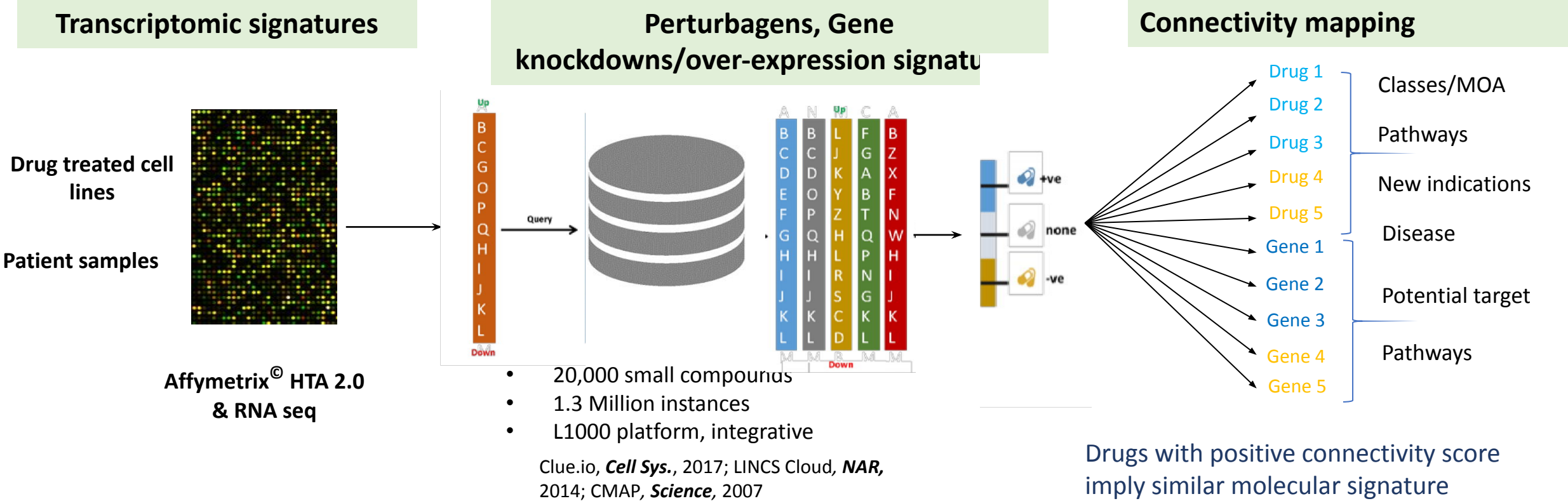
Cissampelos pareira

Ayurveda, folk-care and traditional use:

Against **snake bite**, hormonal disorders, Febrifuge, sepsis, bleeding disorders Fibroid tumours, Poor digestion, Dyspepsia and malaria etc

Estrogen & viral inhibition link???

Connectivity Map platform : For understanding/repurposing of Ayurveda formulations



Traditional use of *Cissampelos pareira* L. for hormone disorder and fever provides molecular links of ESRI modulation to viral inhibition

Madiha Haider, Dhvani Dholakia, Aleksha Panwar, Parth Garg, Vivek Anand, Atish Gheware, Khushboo Singhal, Dayanidhi Singh, Shaunak A Burse, M. Ghalib Enayathullah, Yash Parekh, Sushma Ram, Surekha Kumari, Anmol Kumar, Arjun Ray, Guruprasad R. Medigeshi, Kiran Kumar Bokara, Upendra Sharma, Bhavana Prasher, Mitali Mukerji

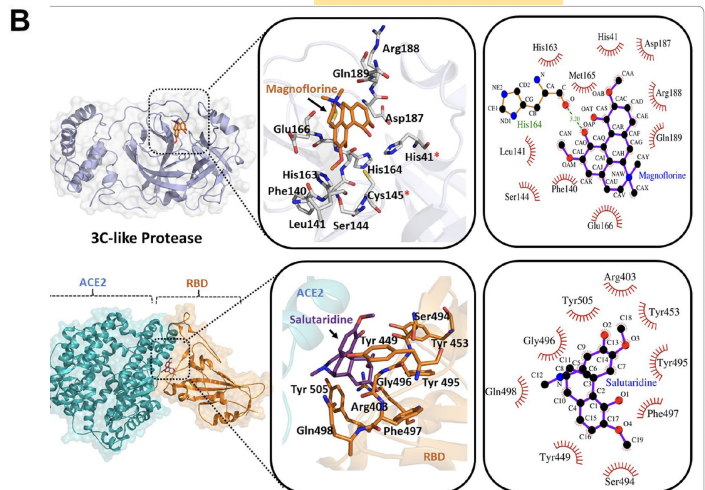
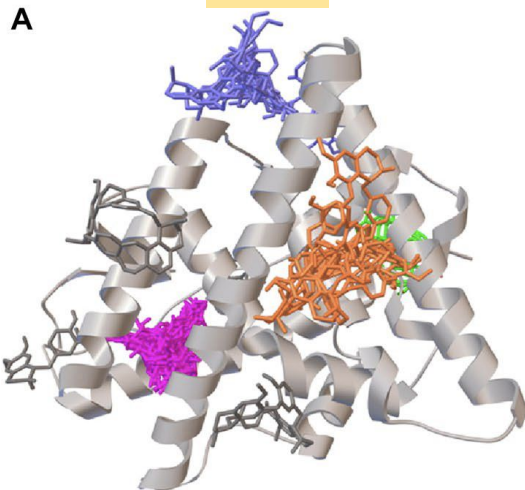
doi: <https://doi.org/10.1101/2021.02.17.431579>

This article is a preprint and has not been certified by peer review [what does this mean?].

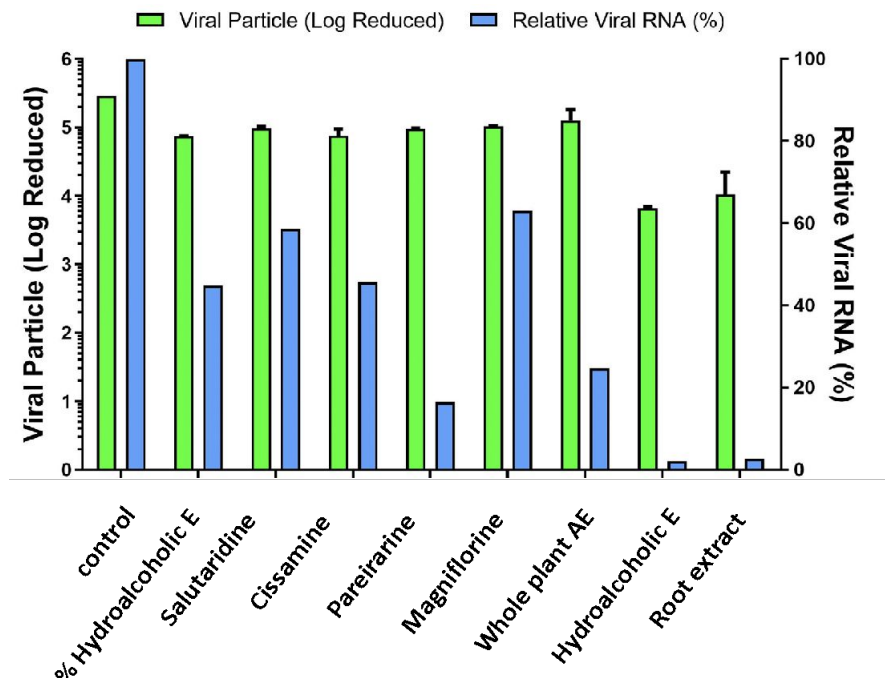
Ayurgenomics based framework : Application in exploring poly-pharmacology & repurposing

ESRI

SARS-Cov2



CIPA biomolecules binds ESRI & crucial SARS-CoV2 therapeutic targets



Whole works better than individual molecules

Traditional use of *Cissampelos pareira* L. for hormone disorder and fever provides molecular links of ESRI modulation to viral inhibition

Madiha Haider, Dhvani Dholakia, Aleksha Panwar, Parth Garg, Vivek Anand, Atish Gheware, Khushboo Singhal, Dayanidhi Singh, Shaunak A Burse, M. Ghalib Enayathullah, Yash Parekh, Sushma Ram, Surekha Kumari, Anmol Kumar, Arjun Ray, Guruprasad R. Medigeshi, Kiran Kumar Bokara, Upendra Sharma, Bhavana Prasher, Mitali Mukerji

doi: <https://doi.org/10.1101/2021.02.17.431579>

This article is a preprint and has not been certified by peer review [what does this mean?].

Haider et al. BMC Complementary Medicine and Therapies (2022) 22:114 <https://doi.org/10.1186/s12906-022-03584-3>

BMC Complementary Medicine and Therapies

RESEARCH Open Access

Anti-SARS-CoV-2 potential of *Cissampelos pareira* L. identified by connectivity map-based analysis and in vitro studies

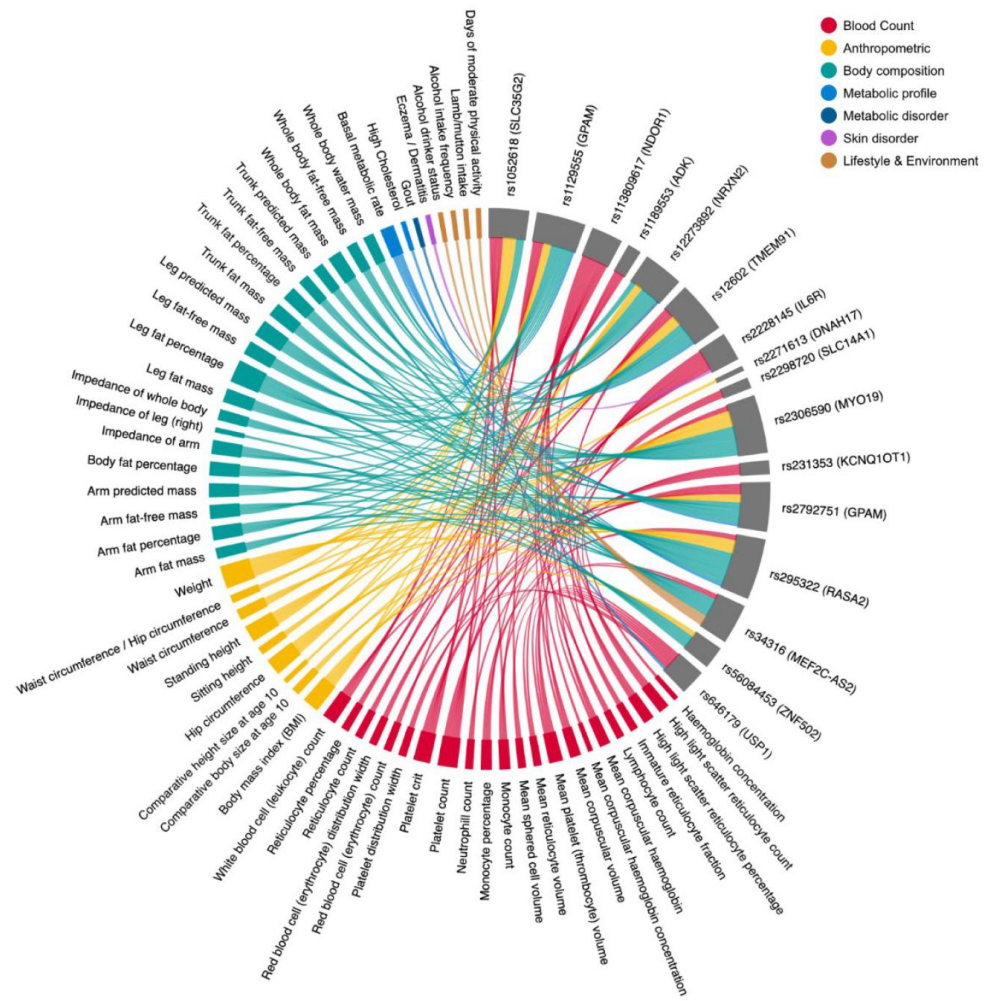
Madiha Haider^{1,2}, Vivek Anand^{1,2}, M. Ghalib Enayathullah³, Yash Parekh³, Sushma Ram³, Surekha Kumari^{4,2}, Anmol^{2,4}, Gayatri Panda⁵, Manjari Shukla⁶, Dhvani Dholakia^{1,2}, Arjun Ray⁵, Sudipta Bhattacharyya⁶, Upendra Sharma^{2,4}, Kiran Kumar Bokara³, Bhavana Prasher^{1,2,7*} and Mitali Mukerji^{1,2,6,7*}

Non-invasive phenotyping for Risk stratification : insights from exome



Article

Whole Exome Sequencing in Healthy Individuals of Extreme Constitution Types Reveals Differential Disease Risk: A Novel Approach towards Predictive Medicine



Gene	Variation details			alt allele frequency in cohorts				profile
	dbSNP	alt allele	nature	VADU		North		
				Vata	Kapha	Vata	Kapha	
ANKLE1	rs8100241	A	missense (A to T)	0.611	0.333	0.6111	0.2941	V vs K
	rs8108174	A	missense (L to Q)	0.611	0.333	0.5833	0.2778	

ARTICLE

Received 16 Jun 2015 | Accepted 20 Jul 2016 | Published 7 Sep 2016

DOI: 10.1038/ncomms12675

OPEN

Functional mechanisms underlying pleiotropic risk alleles at the 19p13.1 breast-ovarian cancer susceptibility locus

Kate Lawrenson *et al.*[#]

ARTICLE

<https://doi.org/10.1038/s42003-023-04611-w>

OPEN



ANKLE1 cleaves mitochondrial DNA and contributes to cancer risk by promoting apoptosis resistance and metabolic dysregulation

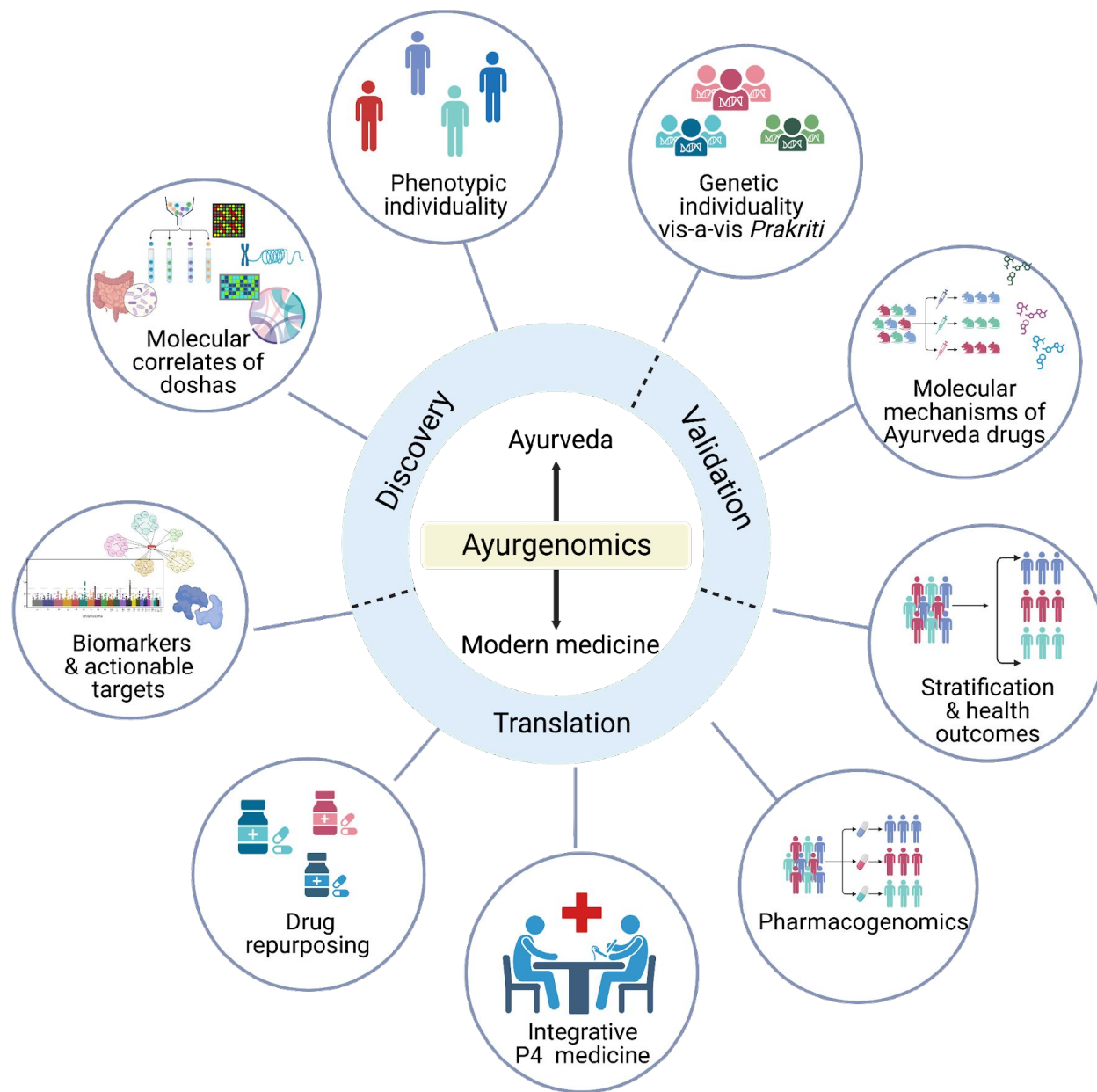
Piotr Przanowski ^{1,2}, Róża K. Przanowska ^{1,2} & Michael J. Guertin ^{3,4}

Molecular correlates of doshas through Ayurgenomics

Mukerji M (2023).

Ayurgenomics-based frameworks in precision and integrative medicine: Translational opportunities. Cambridge

Ayurgenomics – A framework for bridging Ayurveda with modern medicine



ACS
chemical
biology

IN FOCUS

pubs.acs.org/acschemicalbiology

Ayurgenomics: A New Way of Threading Molecular Variability for Stratified Medicine

Tav Pritesh Sethi,[†] Bhavana Prasher,^{*‡} and Mitali Mukerji^{*†}

ACS Chem. Biol. 2011,
6, 875–880

Ayurgenomics-based frameworks in precision and integrative medicine: Translational opportunities

Mitali Mukerji 

Department of Bioscience and Bioengineering, Indian Institute of Technology Jodhpur, Karwar, India and School of Artificial Intelligence and Data Science (AIDE), Indian Institute of Technology Jodhpur, Karwar, India

Cambridge Prisms: Precision Medicine, 1, e29, 1–14

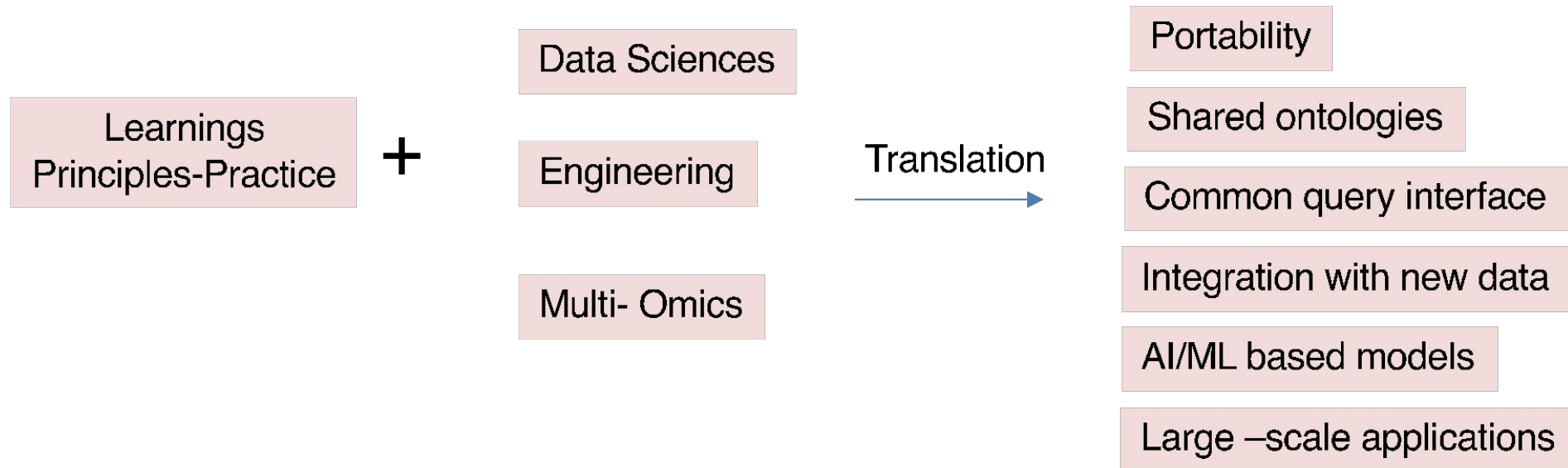
Setting up : Trans-disciplinary Centre of Excellence in integrative precision health at IIT Jodhpur

AYURTECH CoE at IITJ



Supported by Ministry of
AYUSH

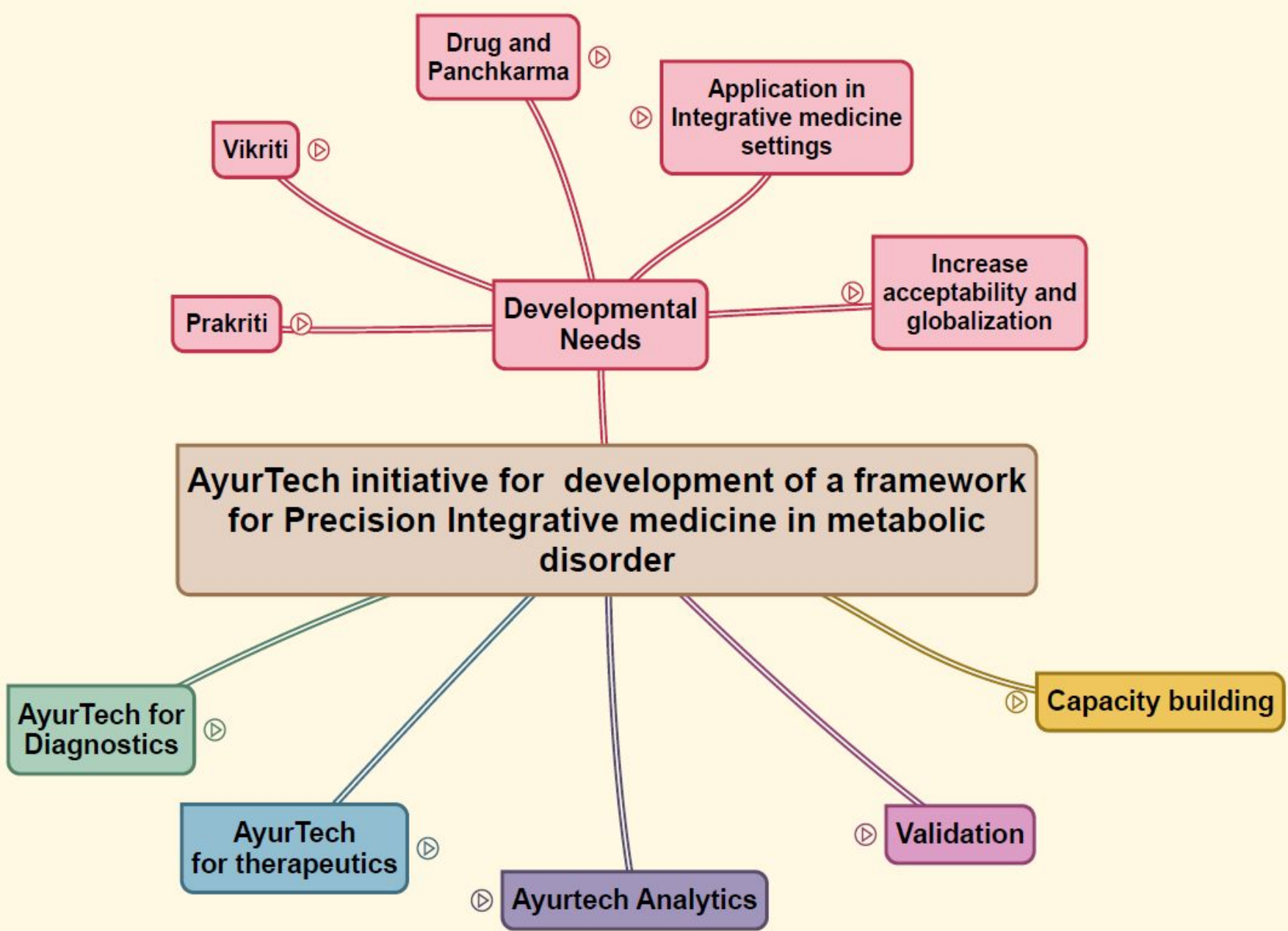
**First of it's kind unique initiative combining Electronics,
Digital health and AI for realizing “Evidence based
Ayurveda” solutions**



AYURTECH Aims to Leverage **Ayurveda** to address **GAP** areas in current **Precision Health Cohorts**

- System's understanding through common organizing principle (*tridosha*) and **network medicine** for translation (*trisuutra*)
- Individualized health baseline that inform response and trajectory in geo-spatio-temporal dimensions and **pre-emptive management** of lifestyle and nutrition
- **Disease trajectories** based on individual baselines
- Diseases treated in an individual centric manner and as moving targets
- **Unique therapeutic** modalities
 - Panchkarma
 - Multicomponent single and poly herbo-metallic formulations
 - Different preparations and mode of administrations for diverse therapies
 - Personalised dietary and yoga interventions
- Context of population, culture and habitat on **individual baselines**

Development of AyurTech Framework



First of its kind unique initiative combining Electronics, Digital health and AI for realizing “Evidence based Ayurveda” solutions

Scenario : Patient A –approaches an Ayurveda doctor with a disease called Fanconi anemia

The patient exhibits the following symptoms

HPO_TERM_NAME	CATEGORY
Deficient excision of UV-induced pyrimidine	Metabolism/Laboratory abnormality
Prolonged G2 phase of cell cycle	Metabolism/Laboratory abnormality
Hearing impairment	Ear
Strabismus	Eye
Short stature	Growth
Ectopic kidney	Genitourinary system
Male infertility	Genitourinary system
Horseshoe kidney	Genitourinary system
Duplicated collecting system	Genitourinary system
Abnormal renal morphology	Genitourinary system
Leukemia	Neoplasm
Hypergonadotropic hypogonadism	Endocrine

Abnormality of cardiovascular system morphology	Cardiovascular
Bruising susceptibility	Skin, Hair, and Nails
Abnormal heart morphology	Cardiovascular
Complete duplication of thumb phalanx	Skeletal system
Intellectual disability	Nervous System
Microcephaly	Nervous System
Abnormality of skin pigmentation	Skin, Hair, and Nails
Chromosomal breakage induced by crosslinking	Metabolism/Laboratory abnormality
Small for gestational age	Growth
Microphthalmia	Eye
Autosomal recessive inheritance	Inheritance
Absent thumb	Skeletal system
Reticulocytopenia	Blood and blood-forming tissues
Short thumb	Skeletal system
Thrombocytopenia	Blood and blood-forming tissues
Cryptorchidism	Genitourinary system
Pancytopenia	Blood and blood-forming tissues
Absent radius	Skeletal system
Renal agenesis	Genitourinary system
Anemic pallor	Skin, Hair, and Nails
Neutropenia	Blood and blood-forming tissues

What would be the Ayurveda diagnosis ?

What is the likely cause ??

Which features are informative for diagnosis??

What is the likely therapy ??

Vata regulates cell division and morphogenesis

शरीराध्यधास्तु परमाणुभेदेनापरिसंख्येया भवन्ति, अतिबहुत्वाद्दतिसौहृम्याद्दतीन्द्रियत्वाच्च । तेषां संयोगविभागे परमाणूनां कारणं वायुः कर्मस्वभावश्च ॥ १७ ॥

c.sa.7/17

Summary

Gene Location: 16q24.3

Definition

The Fanconi anemia complementation group (FANC) currently includes FANCA, FANCB, FANCC, FANCD1 (also called BRCA2), FANCD2, FANCE, FANCF, FANCG, FANCI, FANCI (also called BRIP1), FANCL, FANCM and FANCN (also called PALB2). The previously defined group FANCH is the same as FANCA. Fanconi anemia is a genetically heterogeneous recessive disorder characterized by cytogenetic instability, hypersensitivity to DNA crosslinking agents, increased chromosomal breakage, and defective DNA repair. The members of the Fanconi anemia complementation group do not share sequence similarity; they are related by their assembly into a common nuclear protein complex. This gene encodes the protein for complementation group A. Alternative splicing results in multiple transcript variants encoding different isoforms. Mutations in this gene are the most common cause of Fanconi anemia. [provided by RefSeq, Jul 2008]

No. Descendants

Hierarchy



- Abnormal cellular physiology
- Abnormality of the cell cycle
 - Prolonged G2 phase of cell cycle

Abnormality of the cell cycle HP:0011018

An abnormality of the cell cycle.

Synonyms: *Abnormality of the cell cycle*

Cross References: *UMLS:C4023594*

Export Associations

Disease Associations

Gene Associations

Disease Id	Disease Name	Associated Genes
OMIM:227650	Fanconi anemia	FANCA [2175]
OMIM:227645	Fanconi anemia, complementation group C	FANCC [2176]
OMIM:227646	Fanconi anemia, complementation group D2	FANCD2 [2177]
OMIM:600901	Fanconi anemia, complementation group E	FANCE [2178]

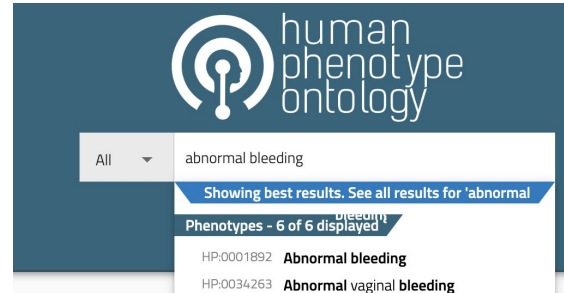
Displaying 4 out of 4.

Building conversational language between Ayurveda and Modern medicine

Bridging the ontological gap

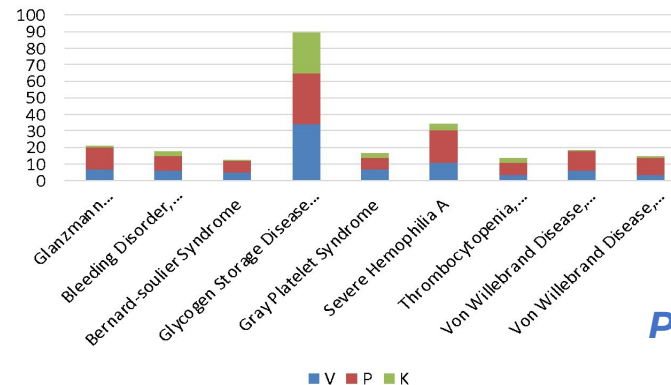
- Glanzmann Thrombasthenia
- Bleeding Disorder, Platelet-type, 24, Autosomal Dominant
- Bernard-soulier Syndrome
- Glycogen Storage Disease Due To Glucose-6-phosphatase Deficiency Type Ib
- Gray Platelet Syndrome
- Severe Hemophilia A
- Thrombocytopenia, Autosomal Dominant, 7
- Von Willebrand Disease, Type 1
- Von Willebrand Disease, Type 3

Nine syndromes



~10000 HPOs have been annotated by a team of Ayurveda physicians

300 features annotation



पित्तं विदग्धं स्वगुणैर्विदहत्याशु शोणितम् ।
 ततः प्रवर्तते रक्तमूर्ध्वं चाधो द्विधाऽपिवा ॥
 उर्ध्वं नासाक्षिकर्णास्ये मँदयोनि गुदैरधः ।
 कुपितं रोमकूपैश्च समस्तैस्तत्प्रवर्तते ॥
 -(सु.3.45)

Raktapitta

Pitta predominance in all syndromes



Prof. Santanu Chaudhury: Director IITJ & Dept of Electrical Engineering Engineering IIT Delhi

Computer Vision, AI/ML, Big Data Science

Lead Coordinators:

Prof. Mitali Mukerji: Department Bioscience and Bio Engineering, IIT Jodhpur; Human genomics, personalized medicine, Ayurgenomics

Prof. Ajay Agarwal: Department of Electrical engineering; Microelectronics professional, Nanotechnologies, MEMS, microfluidics & Micro-sensors

BioScience and Bioengineering:

Dr. Sudipta Bhattacharyya Structure-guided rational drug discovery, drug-protein interactions, biophysics, drug screening, & bioassay development

Dr. Raviraj Vankayala: nanotheranostics, biophotonics, and drug delivery

Computer Science & Engineering:

Dr. Dip Sankar Banerjee : high performance computing, data analytics, parallel computing and computer architecture.

Dr. Sumit Kalra : Software Architecture, Data Analytics, IoT and Smart Healthcare

Industry partners

Dr. Bala Pesala: Professor of Practice, IIT Jodhpur and founder and CEO of Ayur.AI & CTO, Adiuvo Diagnostics and also. Next gen deep tech companies in the digital health and diagnostics, AI/ML

Dr. Lipika Dey: TCS Research and Innovation and heads analytics and insights practices. NLP, text and data mining, information fusion, machine learning, and semantic search.

Electrical Engineering:

Dr. Rajendra Nagar: Computer Vision, Computer Graphics, 3D Geometry Processing, and Machine Learning.

Prof. Manoj Choudhary: Communications and computing, Wireless Communications & Networks (5G/6G/Next gen WLAN/UWB), AI, IOT, Connected and digital healthcare, Sensors, regulatory & standards frameworks, IPR management.

Chemistry:

Dr. Rohan Erande: synthetic organic chemistry, method development and total synthesis of bioactive natural products.

AYURTECH Lead Clinical Collaborators for framework development



Dr. S. R. Rajasthan Ayurved University

Nodal co-ordinator: Vice Chancellor,

Prof. Abhimanyu Kumar

Dr. Sanjay Srivastava: Shalya Tantra,

Dr. Prem Prakash Vyas: Bal Roga

Dr. Rakesh Kumar Sharma: Rachana Shareer

Validation studies for integrative medicine: AYUSH Centre, AIIMS Jodhpur

Nodal Coordinator : Director AIIMS Jodhpur

Dr Pankaj Bharadwaj : Additional Professor, Community Medicine & Family Medicine. Coordinator, School of Public Health. AIIMS Jodhpur.

Dr Meenakshi Sharma

S.No.	Group Name	Name of Faculty Member
1.	Prakriti	1. Dr. Rakesh Kumar Sharma
		2. Dr. Vinod Kumar Gautam
		3. Dr. Brahmanand Sharma
		4. Dr. Dinesh Chandra Sharma
2.	AYURTECH	1. Dr. Govind Prasad Gupta
		2. Dr. Gyan Prakash Sharma
		3. Dr. Sanjay Srivastav
		4. Dr. Ritu Kapoor
		5. Dr. Manisha Goel
3.	DESSERT Medicine	1. Dr. Prem Prakash Vyas
		2. Dr. Chandan Singh
		3. Dr. Manoj Adlakhia

Implementation studies: More partners have shown Expression of interest from different centres and hospitals.....

Ayurgenomics inter-disciplinary team



Dr. Bhavana Prasher MD Ayurveda
Principal Scientist TRISUTRA @IGIB & Co-PI



Prof Samir Brahmachari
(mentor & vision)

Ayurveda

Shilpi Aggarwal (PhD, BAMS)
Bharat Krishna Khuntia (MD),
Arvind Kumar (MD)

Genome informatics

Dr Ankita Narang
Dr Pradeep Tiwari
Rintu Kutum (PhD)
Dhwani Dholakia (PhD)
Tahseen Abbas (PhD)

Modern medicine, Pharma and Physiology

Tav Pritesh Sethi (PhD MBBS)
Dr Atish Gheware (PhD Mpharm)
Ritu Rani (PhD)

Genomics

Dr Sapna Negi
Dr Binuja Varma,
Dr Rajesh Pandey,
Dr Pankaj Jha
Dr Pramod Gautam
Madiha Haider (PhD)
Gaura Chaturvedi (PhD)
Dayanidhi Singh (PhD)
Sumita Chakravorty (PhD)
Sunanda (PhD)

Collaborators

Dr. Anurag Agrawal (IGIB),
Dr. Qadar Pasha (IGIB),
Dr. Debasis Dash,
Dr Saurav Ghosh (ISI Kolkata),
Dr Nar Singh Chauhan, MDU, Rohtak
Greg Gibson (Georgia Tech)
Dr Kiran Bokara CSIR-CCMB
Dr Upendra Sharma CSIR-IHBT,
Dr Guruprasad Medigeshi THSTI,
Dr Arjun Ray – IIITD

KEMHRC:

Sanjay Juvekar,
Bhushan Girase, Ankita
Shrivastava, Rutuja
Patil,
Dheeraj Aggarwal,
Bharat Choudhury

**DST , CSIR, AYUSH for
financial support**

Indian Genome Variation Consortium
TRISUTRA Ayurgenomics Consortium
CSIR-TRISUTRA Team @IGIB

Public health



Modern medicine



Ayurveda





Thank You