

## **Research achievements: Sanjeev Galande (2015 to 2022)**

### **Extent of the contributions:**

Out of these, the two red colored entries are the most significant recent publications from my laboratory in 2023 and have a direct translational/public health impact. The blue colored entries are the next set of significant recent publications from my laboratory that report fundamental discoveries (Basic science). I have conceived and supervised both these studies along with the collaborator, hence we are co-corresponding authors. The work described in the paper published in *Frontiers in Neuroscience* (# 2 Mahajan et al., 2023) is done as a collaboration with Dr. Nixon Abraham of IISER Pune. Here, the entire project was conceived by me during the pandemic. Since none of the labs could order any strains of mice from overseas, I thought of generating the line in-house. The entire planning and execution of the knockout strategy was done by me. Dr. Nixon's lab performed the phenotyping of the lab. I supervised the entire project. This is one of the most significant contributions for research on COVID. Another significant paper of this year is the paper published in *Journal of Developmental Origins of Health and Disease* ((# 6 Khare et al., 2023). This is a result of the long-standing collaboration with the diabetes research unit of KEM hospital, Pune. They built the cohort and collected the samples while my laboratory performed the cellular and molecular work as well as data analysis. I supervised the entire project and this was part of the Centre of Excellence Program of DBT. The results of this study have profound implications on the strategies towards dietary interventions for the malnourished population in India.

I all other publications where I am the corresponding author, the ideas and projects were conceived by me along with the PhD students or postdoctoral fellows. I provided the supervision, contributed to data analysis, wrote/edited the manuscripts and also dealt with the revision process. The funding to support this work was also obtained by me from various agencies.

**Certified that I have not received any award for the research work (2015-2023) listed above.**



Sanjeev Galande

## Research Contributions in past 8 years:

All the awards that I have received are based on the work done prior to 2015. The list of papers published in the past 8 years is given below (in reverse chronological order):

**Publications:**            \*Equal contribution            #Corresponding author

1. Suresh V. \*, Muralidharan, B. \*, Pradhan, S. J. \*, Bose, M., D'Souza, L., Parichha, A., Reddy, P. C., **Galande, S. #** and Tole, S. # 2023. Regulation of chromatin accessibility and gene expression in the developing hippocampal primordium by LIM-HD transcription factor LHX2. *PLoS Genetics* 19(8):e1010874.doi: 10.1371/journal.pgen.1010874.
2. Mahajan S., Sen S., Sunil A., Srikanth P., Marathe S.D, Shaw K., Sahare M., **Galande S. #**, and Abraham N.A. # 2023. Knockout of ACE2 receptors lead to morphological aberrations in rodent olfactory centers and dysfunctions associated with sense of smell. *Frontiers in Neuroscience*, 17:1180868. doi: 10.3389/fnins.2023.1180868
3. Gungi, A., Saha, S., Pal, M. and **Galande, S. #**, 2023. H4K20me1 plays a dual role in transcriptional regulation of regeneration and axis patterning in Hydra. *Life Science Alliance*, 6(5).
4. Chee, J.M., Lanoue, L., Clary, D., Higgins, K., Bower, L., Flenniken, A., Guo, R., Adams, D.J., Bosch, F., Braun, R.E., Brown, S.D., H.-J. Genie Chin, Dickinson, M.E., Hsu C.-W., Dobbie M., Gao X., **Galande S.**, Grobler A., Heaney J.D., Herault Y., de Angelis M.H., Mammano F., Nutter L.M.J., Parkinson H., Qin C., Shiroishi T., Sedlacek R., Seong J.-K., Xu Y., The International Mouse Phenotyping Consortium, Brooks B., McKerlie C., Lloyd K.C.K., Westerberg H. and Moshiri A. 2023. Genome-wide screening reveals the genetic basis of mammalian embryonic eye development. *BMC biology*, 21(1), pp.1-15.
5. O'Rourke, M.B., Januszewski, A.S., Sullivan, D.R., Lengyel, I., Stewart, A.J., Arya, S., Ma, R.C., **Galande, S.**, Hardikar, A.A., Joglekar, M.V. and Keech, A.C., Jenkins A.J., Molloy M. P., 2023. Optimised plasma sample preparation and LC-MS analysis to support large-scale proteomic analysis of clinical trial specimens: Application to the Fenofibrate Intervention and Event Lowering in Diabetes (FIELD) trial. *PROTEOMICS–Clinical Applications*, 17(3), p.2200106.
6. Khare, S.P. \*, Madhok, A. \*, Patta, I., Sukla, K.K., Wagh, V.V., Kunte, P.S., Raut, D., Bhat, D., Kumaran, K., Fall, C. and Tatu, U., Chandak G.R., Yajnik C. S. #, **Galande, S. #** 2023. Differential expression of genes influencing mitotic processes in cord blood mononuclear cells after a pre-conceptional micronutrient-based randomised controlled trial: Pune Rural Intervention in Young Adolescents (PRIYA). *Journal of Developmental Origins of Health and Disease*, 14: 437–448. doi: 10.1017/S204017442200068X
7. Higgins, K., Moore, B.A., Berberovic, Z., Adissu, H.A., Eskandarian, M., Flenniken, A.M., Shao, A., Imai, D.M., Clary, D., Lanoue, L. and Newbigging, S., Nutter L.M.J., Adams D.J.,

- Bosch F., Braun R.E, Brown S.D.M., Dickinson M.E., Dobbie M., Flicek P., Gao X., **Galande S.**, Grobler A., Heaney J.D., Herault Y., de Angelis M.H., Chin H.-J. G., Mammano F., Qin C., Shiroishi T, Sedlacek R., Seong J.-K., Xu Y., The IMPC Consortium, Lloyd K.C.K., McKerlie C., and Moshiri A. 2022. Analysis of genome-wide knockout mouse database identifies candidate ciliopathy genes. *Scientific Reports*, 12(1), p.20791.
8. Shetty A\*, Tripathi SK\*, Junttila S\*, Buchacher T\*, Biradar R, Bhosale SD, Envall T, Laiho A, Moulder R, Rasool O, **Galande S**, Elo LL<sup>#</sup>, and Lahesmaa R<sup>#</sup>. (2022) A systematic comparison of FOSL1, FOSL2 and BATF-mediated transcriptional regulation during early human Th17 differentiation *Nucleic Acids Res* 50:4938–4958, PMID: 35511484.
  9. Pradhan SJ, Reddy PC, Smutny M, Sharma A, Sako K, Oak MS, Shah R, Pal M, Deshpande O, Dsilva G, Tang Y, Mishra R, Deshpande G, Giraldez AJ, Sonawane M, Heisenberg CP<sup>#</sup>, **Galande S<sup>#</sup>**. (2021) Satb2 acts as a gatekeeper for major developmental transitions during early vertebrate embryogenesis. *Nat Commun.* 12:6094. doi: 10.1038/s41467-021-26234-7. PMID: 34667153
  10. Pandit P, **Galande S<sup>#</sup>**, Iris F. (2021) Maternal malnutrition and anaemia in India: dysregulations leading to the 'thin-fat' phenotype in newborns. *J Nutr Sci.* 10:e91. doi: 10.1017/jns.2021.83. eCollection 2021. PMID: 34733503
  11. Unni M, Reddy PC, Pal M, Sagi I, **Galande S<sup>#</sup>**. (2021) Identification of Components of the Hippo Pathway in Hydra and Potential Role of YAP in Cell Division and Differentiation. *Front Genet.* 12:676182. doi: 10.3389/fgene.2021.676182. eCollection 2021. PMID: 34691138.
  12. Shetty A, Bhosale SD, Tripathi SK, Buchacher T, Biradar R, Rasool O, Moulder R, **Galande S**, Lahesmaa R. (2021) Interactome Networks of FOSL1 and FOSL2 in Human Th17 Cells. *ACS Omega*.6:24834-24847. doi: 10.1021/acsomega.1c03681. eCollection 2021 Sep 28. PMID: 34604665
  13. Sharma A, Mir R, **Galande S<sup>#</sup>**. (2021) Epigenetic Regulation of the Wnt/ $\beta$ -Catenin Signaling Pathway in Cancer. *Front Genet.* 12:681053. doi: 10.3389/fgene.2021.681053. eCollection 2021. PMID: 34552611
  14. Madhok A, Bhat SA, Philip CS, Sureshbabu SK, Chiplunkar S<sup>#</sup>, **Galande S<sup>#</sup>**. (2021) Transcriptome Signature of V $\gamma$ 9V $\delta$ 2 T Cells Treated With Phosphoantigens and Notch Inhibitor Reveals Interplay Between TCR and Notch Signaling Pathways. *Front Immunol.* 12:660361. doi: 10.3389/fimmu.2021.660361. eCollection 2021. PMID: 34526984.
  15. Shah R, Sharma A\*, Kelkar A\*, Sengupta K, and **Galande S<sup>#</sup>**. 2021. A novel cis regulatory element regulates human XIST in CTCF-dependent manner. *Mol Cell Biol*, 41(8):e0038220. doi: 10.1128/MCB.00382-20DOI: <https://doi.org/10.1128/MCB.00382-20>.
  16. Pillai, A.\*, Gungi, A.\*, Reddy P.C.<sup>#</sup> and **Galande, S.<sup>#</sup>**, 2021. Epigenetic regulation in Hydra: conserved and divergent roles. *Frontiers in Cell and Developmental Biology*, 9, p.1155.
  17. Ramanujam, P. L. \*, Mehrotra, S. \*, Kumar, R. P., Verma, S., Deshpande, G., Mishra, R. K. #, & **Galande, S<sup>#</sup>** (2021). Global chromatin organizer SATB1 acts as a context-dependent regulator of the Wnt/Wg target genes. *Scientific Reports*, 11(1), 3385. doi: 10.1038/s41598-021-81324-2.

18. Reddy, PC\*, Gungi A\*, Ubhe, S, and **Galande S<sup>#</sup>** (2020) Epigenomic landscape of enhancer elements during *Hydra* head organizer formation. *Epigenetics and Chromatin* Oct 12;13(1):43. doi: 10.1186/s13072-020-00364-6. PMID: 33046126
19. Naik S\*, Unni M\*, Sinha D, Rajput SS, Reddy PC, Kartvelishvily E, Solomonov E, Sagi I, Chatterji A, Patil S<sup>#</sup> and **Galande S<sup>#</sup>**. (2020) Differential tissue stiffness of body column facilitates locomotion of *Hydra* on solid substrates. *J Exp. Biol.* Oct 29;223(Pt 20):jeb232702. doi: 10.1242/jeb.232702. PMID: 32958523.
20. Mehrotra S<sup>#</sup>, Bansal P, Oli N, Jayarajan S and **Galande S**. (2020) Defective Proventriculus regulates cell specification in the gastric region of *Drosophila* intestine. *Frontiers in Physiology* Jul 14;11:711. doi: 10.3389/fphys.2020.00711. eCollection 2020. PMID: 32760283.
21. Patta I, Madhok A, Khare SP, Gottimukkala KP, Verma A, Giri S, Dandewad V, Seshadri V, Lal G, Misra-Sen J and **Galande S<sup>#</sup>** (2020), Dynamic regulation of chromatin organizer SATB1 via TCR induced alternative promoter switch during T-cell development. *Nucleic Acids Res* 48:5873-5890. PMID: 32392347.
22. Lloyd KCK, Adams DJ, Baynam G, Beaudet AL, Bosch F, Boycott KM, Braun RE, Caulfield M, Cohn R, Dickinson ME, Dobbie MS, Flenniken AM, Flicek P, **Galande S**, Gao X, Grobler A, Heaney JD, Herault Y, de Angelis MH, Lupski JR, Lyonnet S, Mallon AM, Mammano F, MacRae CA, McInnes R, McKerlie C, Meehan TF, Murray SA, Nutter LMJ, Obata Y, Parkinson H, Pepper MS, Sedlacek R, Seong JK, Shiroishi T, Smedley D, Tocchini-Valentini G, Valle D, Wang CL, Wells S, White J, Wurst W, Xu Y, Brown SDM. (2020) The Deep Genome Project. *Genome Biol.* 21:18. doi: 10.1186/s13059-020-1931-9.
23. Reddy PC, Pradhan SJ, Karmodiya K, **Galande S<sup>#</sup>**. (2020) Origin of RNA Polymerase II pause in eumetazoans: Insights from *Hydra*. *J Biosci.* 45. pii: 8.
24. Narlikar G, **Galande S**. Topical collection on Chromatin Biology and Epigenetics. *J Biosci.* 45. pii: 1.
25. Reddy PC, Gungi A, Ubhe S, Pradhan SJ, Kolte A, **Galande S<sup>#</sup>**. (2019) Molecular signature of an ancient organizer regulated by Wnt/ $\beta$ -catenin signalling during primary body axis patterning in *Hydra*. *Commun Biol.* 2:434. doi: 10.1038/s42003-019-0680-3. eCollection 2019.
26. Khare SP, Shetty A, Biradar R, Patta I, Sathe A, Reddy PC, Chen Z, Lahesmaa R and **Galande S\***. (2019) NF- $\kappa$ B signaling and IL-4 signaling regulate SATB1 expression via alternative promoter usage during Th2 differentiation. *Frontiers in Immunology* 10:667. doi: 10.3389/fimmu.2019.00667. eCollection 2019.
27. Muley VY<sup>#</sup>, Akhter Y, and **Galande S<sup>#</sup>**. (2019) PDZ domains across the microbial world: molecular link to the proteases, stress response and protein synthesis. *Genome Biology and Evolution*;11:644-659. doi: 10.1093/gbe/evz023.
28. Tripathi ST\*, Valikangas T\*, Shetty A\*, Khan MM, Bhosale SD, Moulder R, Komsı E, Salo V, Albuquerque RSD, Rasool O, **Galande S**, Elo LE<sup>#</sup>, and Lahesmaa R<sup>#</sup>. (2019) Quantitative mass spectrometry-based proteomics reveals the dynamic protein landscape during initiation of human Th17 cell polarization. *iScience*;11:334-355. doi: 10.1016/j.isci.2018.12.020. Epub 2018 Dec 26.

29. Balyan R, Gund R, Chawla AS, Khare SP, Pradhan SJ, Rane S, **Galande S**, Durdik JM, George A, Bal A and Rath S. (2019) Correlation of Cell-Surface CD8 Levels with Function, Phenotype and Transcriptome of Naïve CD8 T Cells. *Immunology*;156:384-401. doi: 10.1111/imm.13036. Epub 2019 Jan 13.
30. Naik R and **Galande S**<sup>#</sup>. (2018) SATB family chromatin organizers as master regulators of tumor progression. *Oncogene* 38:1989-2004.
31. Mir R, Sharma A, Pradhan SJ, **Galande S**<sup>#</sup>. (2018) Regulation of transcription factor SP1 by  $\beta$ -catenin destruction complex modulates Wnt response. *Mol Cell Biol* 38:pii: e00188-18. doi: 10.1128/MCB.00188-18. Print 2018 Nov 15.
32. Jangid RK, Kelkar A, Muley VY, **Galande S**<sup>#</sup>. (2018) Bidirectional promoters exhibit characteristic chromatin modification signature associated with transcription elongation in both sense and antisense directions. *BMC Genomics*. 19:313.
33. Manjunath GP\*, Ramanujam PL\*, and **Galande S**<sup>#</sup>. (2018) Structure function relations in PDZ domain containing proteins: implications for protein networks in cellular signaling. *J. Biosci.* 43:155-171.
34. Sawant AA, **Galande S**, Srivatsan SG. (2018) Imaging Newly Transcribed RNA in Cells by Using a Clickable Azide-Modified UTP Analog. *Methods Mol Biol*. 1649:359-371.
35. Muralidharan B, Keruzore M, Pradhan S, Roy B, Shetty A, Kinare A, D'souza L, Maheshwari U, Karmodiya K, Suresh A, **Galande S**, Bellefroid E, and Tole S. (2017) Dmrt5, a novel neurogenic factor, reciprocally regulates Lhx2 to control the neuron-glia cell fate switch in the developing hippocampus. *J Neurosci*. 37:11245-11254.
36. Reddy PC, Ubhe S, Sirwani N, Lohokare R, and **Galande S**<sup>#</sup>. (2017) Rapid divergence of histones in Hydrozoa (Cnidaria) and evolution of a novel histone involved in DNA damage response in hydra. *Zoology (Jena)* 123:53-63.
37. Muralidharan B, Khatri Z, Maheshwari U, Gupta R, Roy B, Pradhan S, Karmodiya K, Padmanabhan H, Shetty A, Balaji C, Kolthur-Seetharam U, Macklis J, **Galande S**, and Tole S. (2017) Lhx2 interacts with the NuRD complex and regulates cortical neuron subtype determinants Fezf2 and Sox11. *J Neurosci*. 37:194-203.
38. Gottimukkala KP, Jangid R, Patta I, Sultana DA, Sharma A, Misra-Sen J, and **Galande S**<sup>#</sup>. (2016) Regulation of SATB1 during thymocyte development by TCR signaling. *Mol. Immunol.* 2016 Jul 22;77:34-43.
39. Keisuke S, Pradhan SJ, Barone V, Inglés-Prieto Á, Rogers KW, Müller P, Ruprecht V, **Galande S**, Janovjak H, and Heisenberg CP. (2016) A temporal pattern of Nodal signaling regulates cell fate specification. *Cell Reports*. 16:866-877.
40. Pusalkar M, Ghosh S, Jaggar M, Husain BF, **Galande S**, Vaidya VA. (2016) Acute and Chronic Electroconvulsive Seizures (ECS) Differentially Regulate the Expression of Epigenetic Machinery in the Adult Rat Hippocampus. 2016; *Int J Neuropsychopharmacol*. 2016 May 17. pii: pyw040. doi: 10.1093/ijnp/pyw040. [Epub ahead of print].

41. Sawant AA, Mukherjee PP, Jangid RK, **Galande S<sup>#</sup>**, Srivatsan SG<sup>#</sup>. (2016) A clickable UTP analog for the posttranscriptional chemical labeling and imaging of RNA. *Org Biomol Chem*. 14:5832-5842.
42. Reddy PC, Sinha I, Kelkar A, Habib F, Pradhan SJ, Sukumar R, and **Galande S<sup>#</sup>**. (2015) Comparative sequence analyses of genome and transcriptome reveal novel transcripts and variants in the Asian elephant *Elephas maximus*. *J Biosci*. 40:891-907.
43. Pusalkar M, Suri D, Kelkar A, Bhattacharya A, **Galande S**, and Vaidya VA. (2015) Early stress evokes dysregulation of histone modifiers in the medial prefrontal cortex across the life span. *Dev. Psychobiol*. 58:198-210.
44. Karmodiya K<sup>#</sup>, Pradhan SJ, Joshi B, Jangid R, Reddy PC, **Galande S**. (2015) A comprehensive epigenome map of *Plasmodium falciparum* reveals unique mechanisms of transcriptional regulation and identifies H3K36me2 as a global mark of gene suppression. *Epigenetics Chromatin*. 2015 Sep 17;8:32. doi: 10.1186/s13072-015-0029-1. eCollection 2015.
45. Sawant AA, Tanpure AA, Mukherjee PP, Athavale S, Kelkar A, **Galande S<sup>#</sup>** and Srivatsan SG<sup>#</sup>. (2015) A versatile toolbox for posttranscriptional chemical labeling and Imaging of RNA. *Nucleic Acids Res*. 44(2):e16. doi: 10.1093/nar/gkv903. Epub 2015 Sep 17.
46. Reddy PC, Unni MK, Gungi A, Agarwal P, and **Galande S<sup>#</sup>**. (2015) Evolution of Hox-like genes in Cnidaria: Study of Hydra Hox repertoire reveals tailor-made Hox-code for Cnidarians. *Mech Dev*. 138 Pt 2:87-96.
47. Hardikar AA<sup>#</sup>, Satoor SN, Karandikar MS, Joglekar MV, Puranik AS, Wong W, Kumar S, Limaye A, Bhat DS, Januszewski A, Umrani MR, Ranjan AK, Apte K, Yajnik P, Bhonde RR, **Galande S**, Keech AC, Jenkins AJ, and Yajnik CS. (2015) Susceptibility to obesity and diabetes following multigeneration undernutrition is not reversed by two generations of nutrient availability. *Cell Metab*. 22:312-319.
48. Mir R, Pradhan SJ, Patil P, Mulherkar R and **Galande S<sup>#</sup>**. (2015) Wnt/ $\beta$ -catenin signaling regulated SATB1 promotes colorectal cancer tumorigenesis and progression. *Oncogene*. 35:1679-1691.



Sanjeev Galande