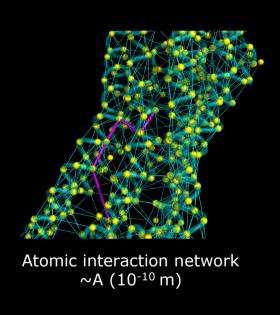
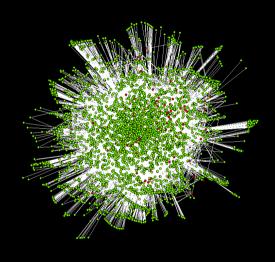
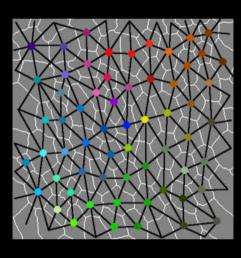
Uncovering organizing principles in biological systems by applying network theory at different scales





Molecular interaction network ~nm (10⁻⁹ m)



Cellular interaction network ~um (10⁻⁶ m)

CNN, Cambridge, 27 Sep 2011

M. Madan Babu

Senior Investigator
MRC Laboratory of Molecular Biology, Cambridge, UK

Regulatory Genomics and Systems Biology

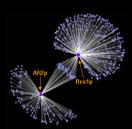
Molecular Level

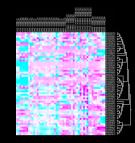
Discover new features of regulatory proteins $YP \times X \times P$ LK KAKYPAESPDYFVDF VKTKAKYPVEPPD<mark>C</mark>VVDF LKTNAKYPTEPPDCVVDF LKTNAKYPYEAPDCSVDF IQ SSQH<mark>P</mark>TSAPSCVTEL PD PTNFPYSTPTVNSNL IQ NRDYPNSPPSVSADV hpL..pbP.psPph.sph Molecular Cell (2008) Cell Stem Cell (2010) EMBO J (2011)

Molecules & regulation

Defining features of regulatory proteins

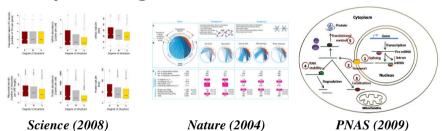






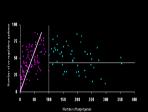
Systems Level

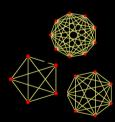
Principles of regulation for cellular homeostasis

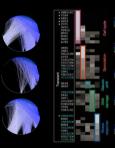


Systems & regulation

Principles of regulation in cellular systems

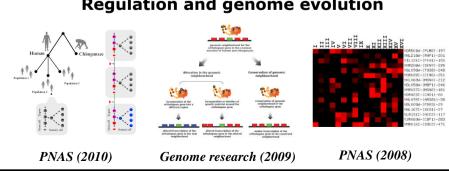






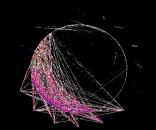
Genome Level

Regulation and genome evolution



Genomes & regulation

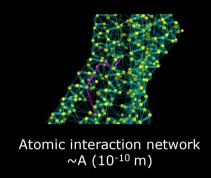
Interplay between regulation and genome evolution



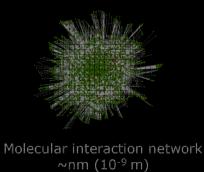




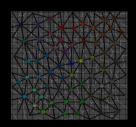
Outline



Application of network approach to understand conformation changes in proteins



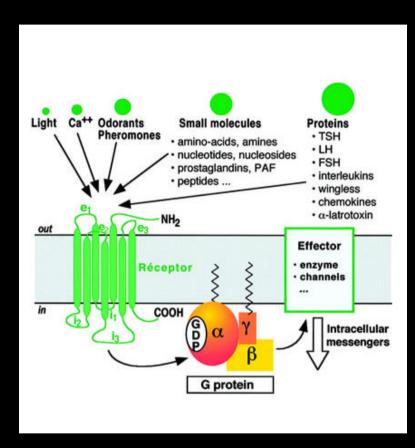
Application of network approach to understand the dynamics of molecular interaction networks



Cellular interaction network ~um (10⁻⁶ m)

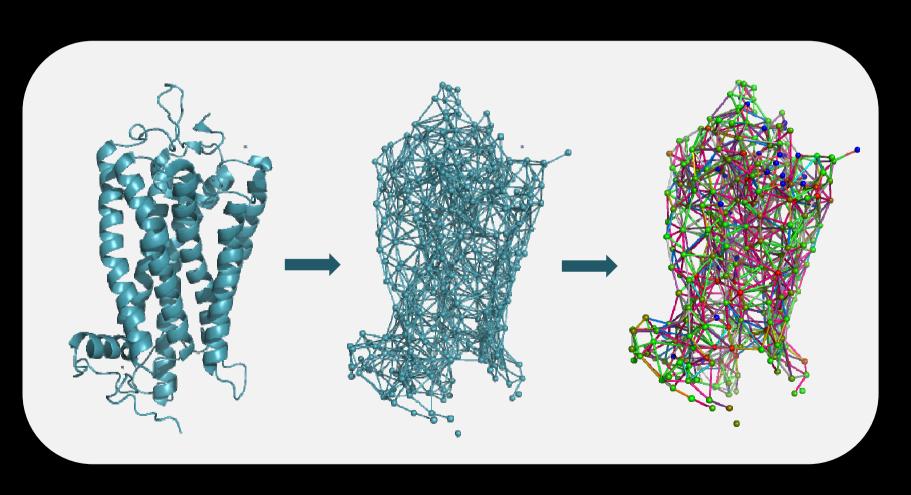
Application of network approach to understand patterns in cell-to-cell contacts during development

Conformational changes in proteins

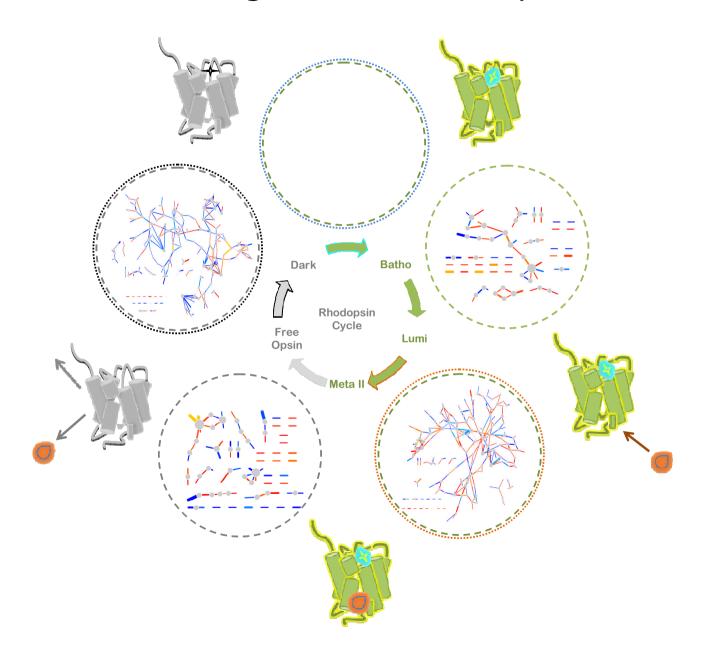


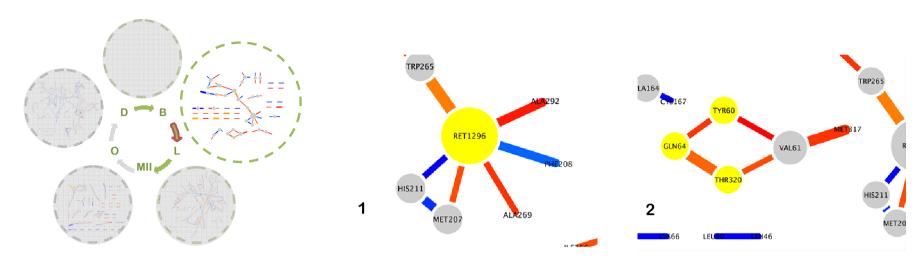
GPCR – an evolutionary success

To understand the conformational cycle in GPCR activation by investigating how the atomic interaction network changes in the different structures

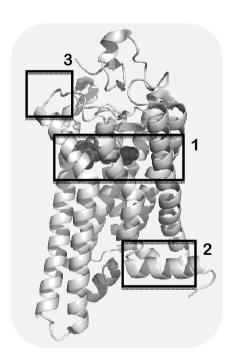


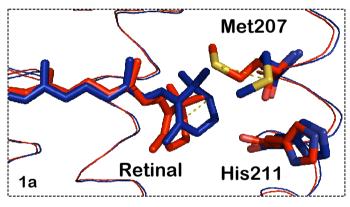
Network alignment and comparison

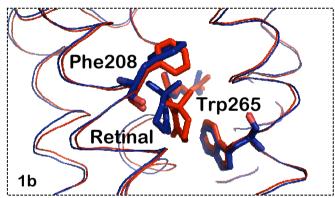




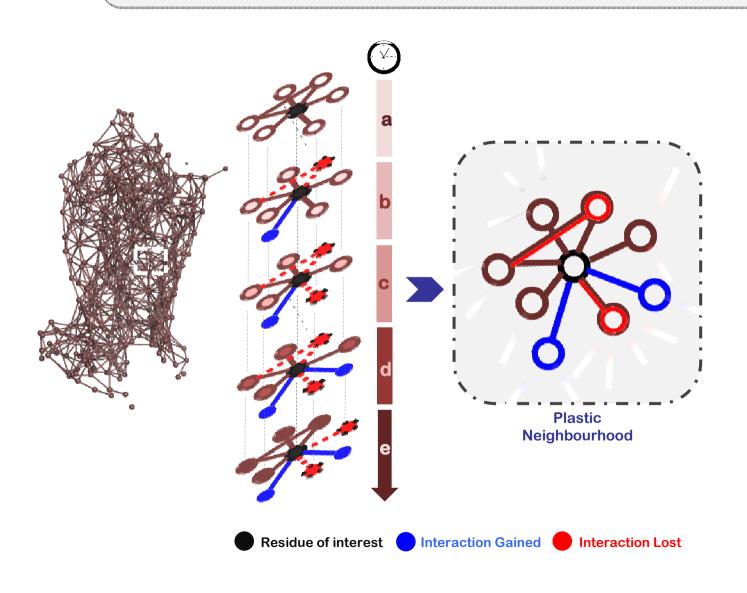
Removed: 44, Added: 24



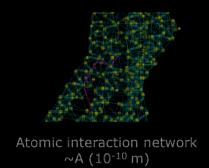




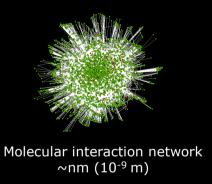
Which are the <u>key residues</u> participating in the activation processes?



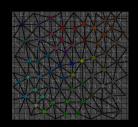
Outline



Application of network approach to understand conformation changes in proteins



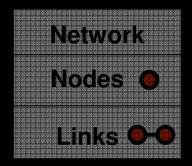
Application of network approach to understand the dynamics of molecular interaction networks

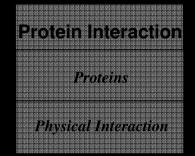


Cellular interaction network ~um (10⁻⁶ m)

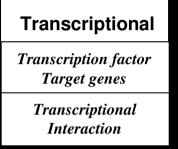
Application of network approach to understand patterns in cell-to-cell contacts during development

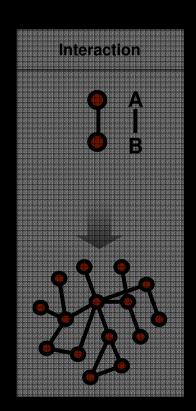
Molecular interaction networks in biology

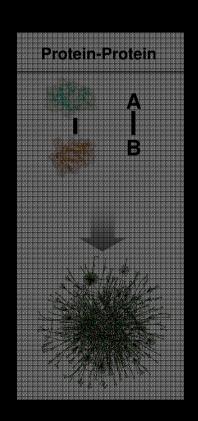


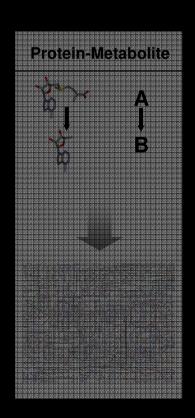


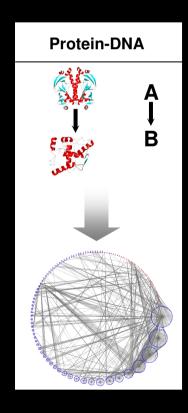




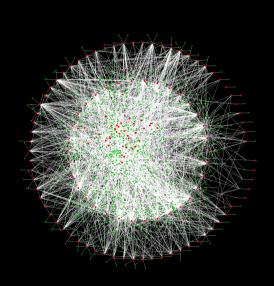








To understand the dynamics of the gene regulatory network in different time scales by investigating how network structure change in different time scales



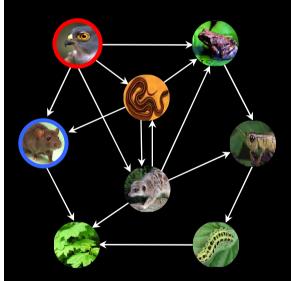
Transcriptional network

- Time scale of the evolution of species (~million years) sequence information; Nat Gen 2004
- Time scale of the life of an organism (~hours or days) expression information; Nature 2004
- Time scale of the life of molecules (~minutes)
 Kinetic information; Mol Sys Biol 2009

Methods to infer hierarchical organization in networks

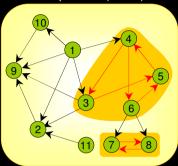
Food web network

 $predator \longrightarrow prey$



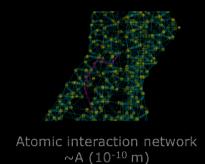
Topological Sort: An approach to infer hierarchical organization in networks

1. Identify strongly connected components (SCCs)

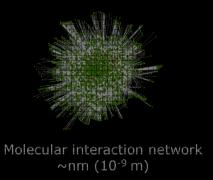


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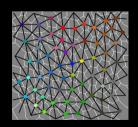
Outline



Application of network approach to understand conformation changes in proteins



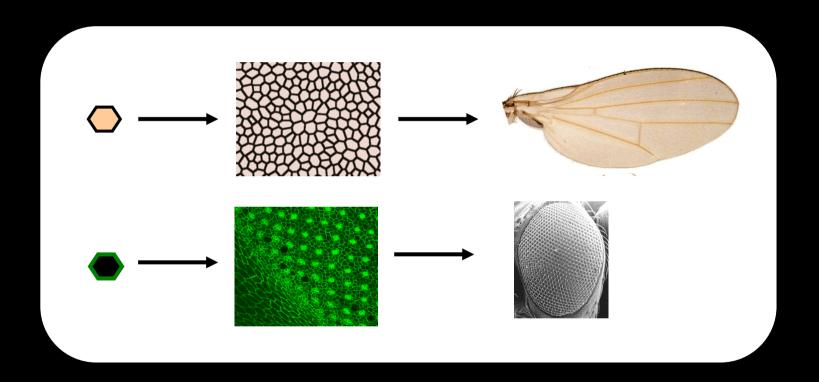
Application of network approach to understand the dynamics of molecular interaction networks



Cellular interaction network ~um (10⁻⁶ m)

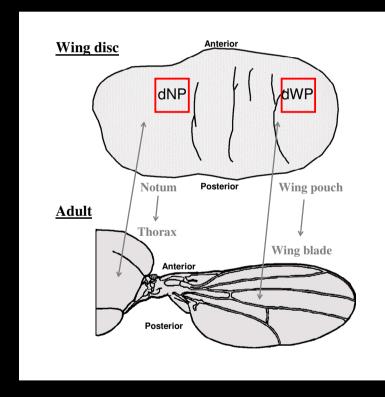
Application of network approach to understand patterns in cell-to-cell contacts during development

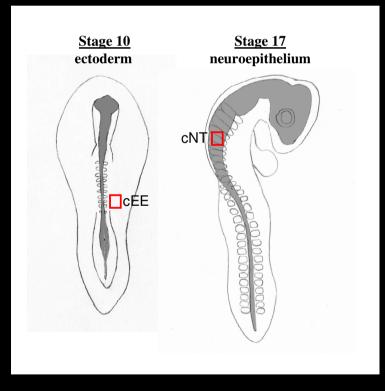
Patterns in the organisation of cells forming an epithelia



The emergence of differences in arrangement of cells is the first step towards establishment of many organs. Understanding this process is limited by the lack of systematic characterization of epithelial organisation.

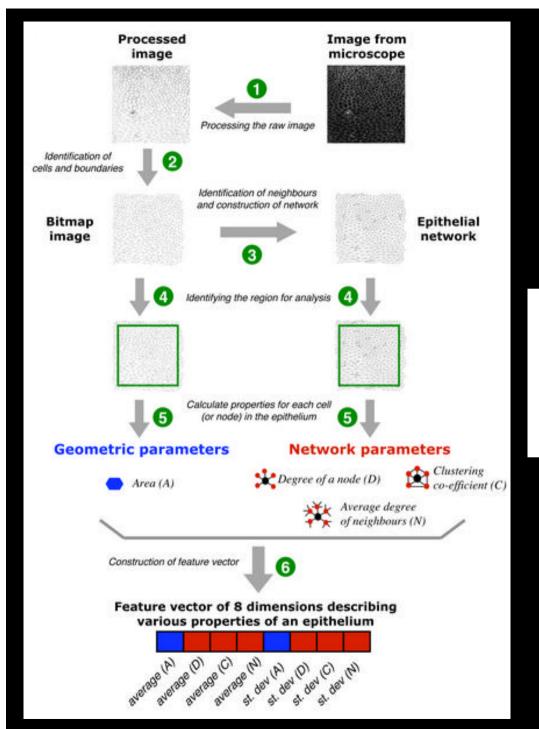
To understand organization of cells in epithelia of different types that are separated spatially, temporally and genetically perturbed by investigating topological patterns in the network of cellular contacts from different organisms



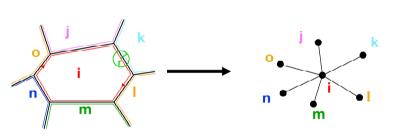


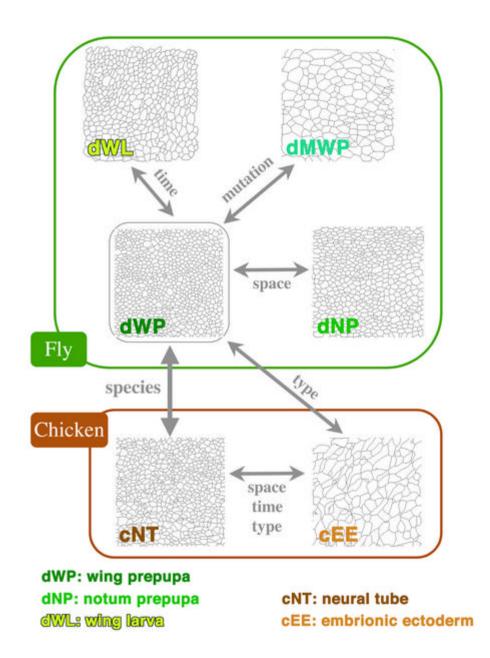
Drosophila

Chicken



Cell contact network

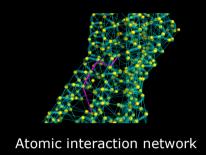




The features of individual cells, together with attributes of the cellular network produce a defining signature that distinguishes epithelia from different organs, species, developmental stages and genetic conditions.

The approach permits characterization, quantification and classification of normal and perturbed epithelia and establishes a framework for understanding molecular mechanisms that underpin the architecture of complex tissues.

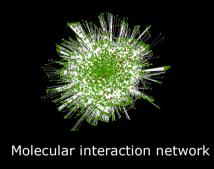
Acknowledgements





A J Venkatakrishnan Gebhard Schertler Jorg Standfuss

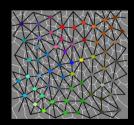
A J Venkatakrishnan et al, unpublished





Raja Jothi Balaji Santhanam Teresa Prytzycka Arthur Wuster Joerg Gsponer L Aravind

Jothi et al, Mol Sys Biol, 2009



Cellular interaction network



Luisma Escudero Luciano Costa Anna Kicheva James Briscoe Matthew Freeman

Escudero et al, Nature Communications, in press

Regulatory Genomics and Systems Biology Group



MADAN BABU MOHAN

Regulatory Genomics & Systems Biology

Past and present group members & associate members



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Natalia de Groot Postdoctoral Scientist 2010 2013



Marija Bulian Postdoctoral Scientist 2011 2014



Kai Kruse PhD Student 2010 2013





Venkat PhD Student 2009 2012



van der Lee Masters Student 2010 2011

Robin



Weber Joint PhD Student 2007 2010

Katie



Sven Sewitz Post-doc



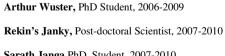
Guilhem Chalancon Masters Student 2009 2011



Charles Ravarani Masters Student 2010 2010



Marie Schrynemackers, Masters Student, 2010 Subhajyoti De, Associate post-doc, 2008-2010 **Joerg Gsponer**, Associate post-doc, 2008-2009 Nitish Mittal, Visiting PhD student, 2008-2009 Matthias Futschik, Visiting scientist, 2007 Henning Claussen, Visiting student, 2007





Sarath Janga PhD Student, 2007-2010 Pradeep Kota, Visiting student, 2007 ...and members of the Theoretical and Computational Biology group at the LMB for helpful discussions













Fondation Philippe Wiener - Maurice Anspach























Human Frontier Science Program