

Not another statistics/bioinformatics talk

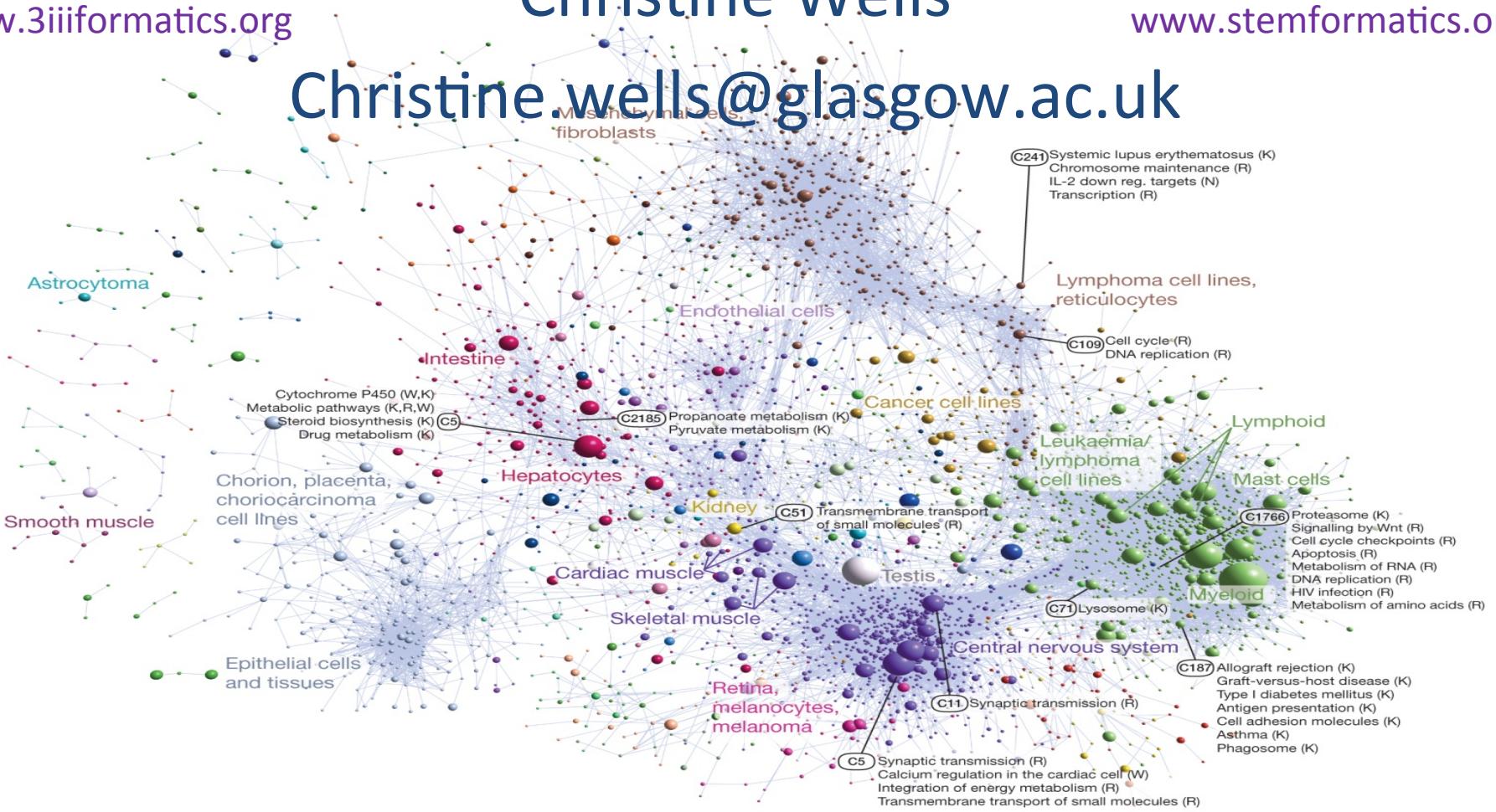
3IIIIFORMATICS
A COLLABORATION WITH
STEMFORMATICS

www.3iiiformatics.org

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Urgent need

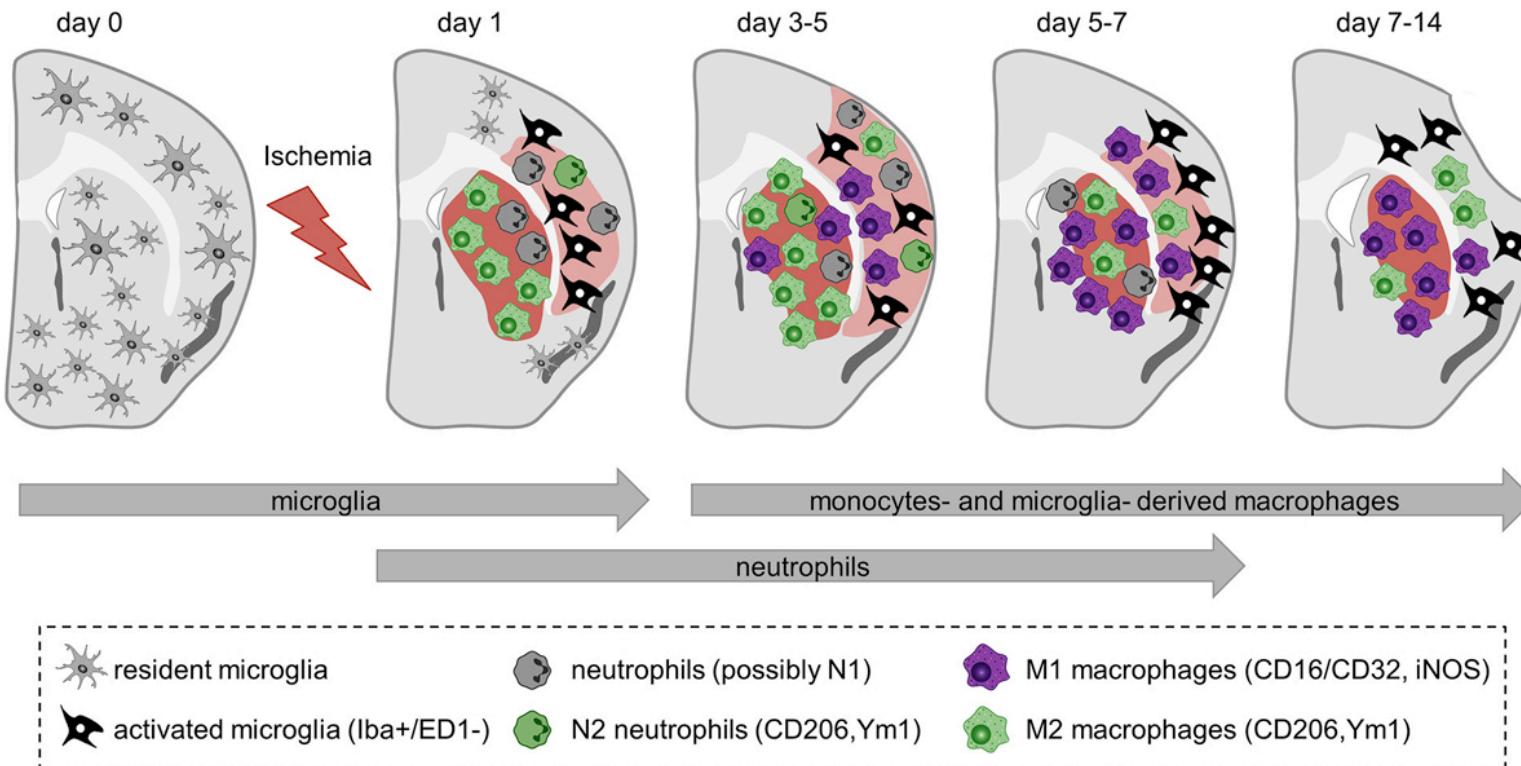
- Clinical material
- Cell models
 - > to identify the major Clec4e+/Mincle cells
 - > resolve why Mincle KO animals have better recovery from stroke.
 - > understand regulation of peripheral inflammatory cell recruitment to injured brain.

The role of microglia and myeloid immune cells in acute cerebral ischemia

Corinne Benakis, Lidia Garcia-Bonilla, Costantino Iadecola and Josef Anrather*

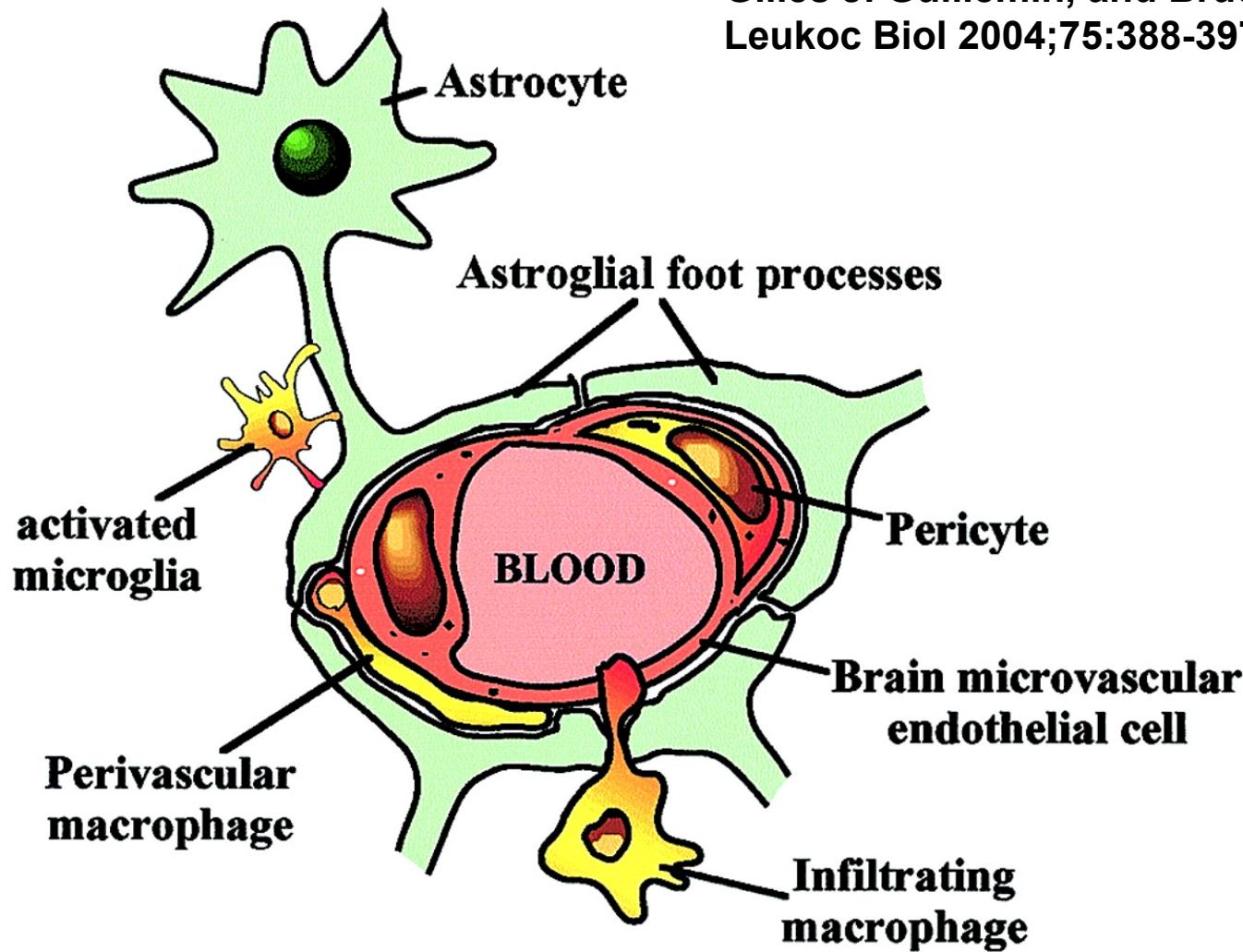
Feil Family Brain and Mind Research Institute, Weill Cornell Medical College, New York, NY, USA

Spatio-temporal profile of microglia and myeloid immune cells and their polarization state in transient brain ischemia



Simplified schematic drawing of the myeloid cells in the brain perivascular area.

Gilles J. Guillemin, and Bruce J. Brew J
Leukoc Biol 2004;75:388-397



Crazy collaborations?

- Map the myeloid cells in the brain that regulate vascular permeability
- Cell trafficking
- Injury
- Repair.
 - >capacity to identify subsets unambiguously
 - >capacity to model interactions/behaviours



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1 2 3 4 5 6

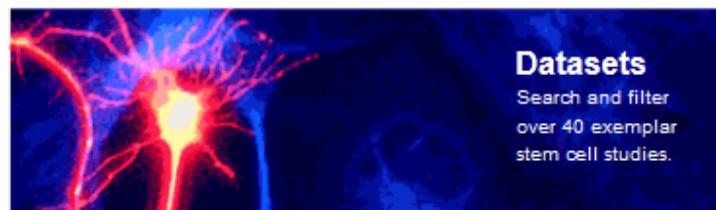
Read (9,63)

Welcome to Stemformatics

This is a portal to a series of public experiments describing mouse and human stem cells and how they differentiate to become mature cells, tissues and organs.

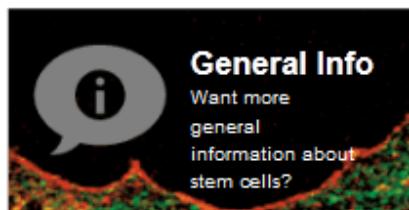
You'll find data from leading stem cell laboratories in a format that is easy to search, easy to visualise and easy to export. Log in to the Workbench to run and save your own analyses.

[START EXPLORING](#)



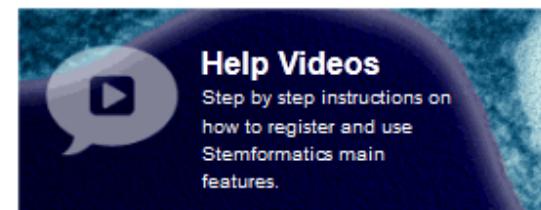
Datasets

Search and filter over 40 exemplar stem cell studies.



General Info

Want more general information about stem cells?



Help Videos

Step by step instructions on how to register and use Stemformatics main features.

PARTNERS

RECENT NEWS

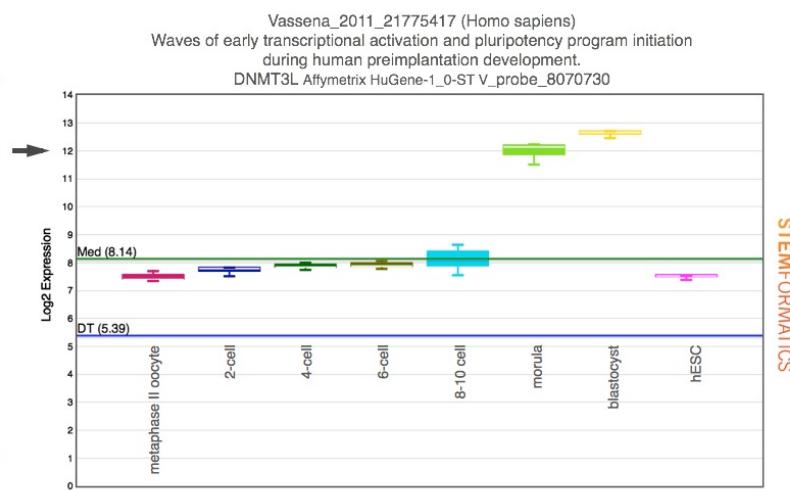
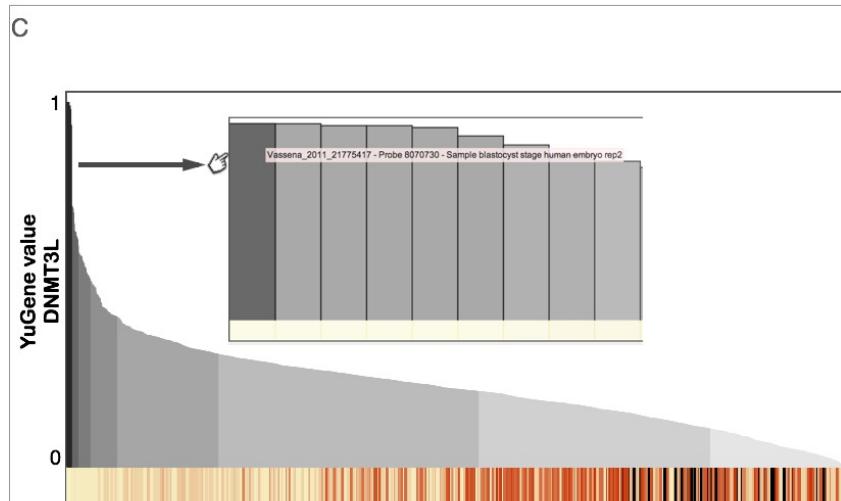
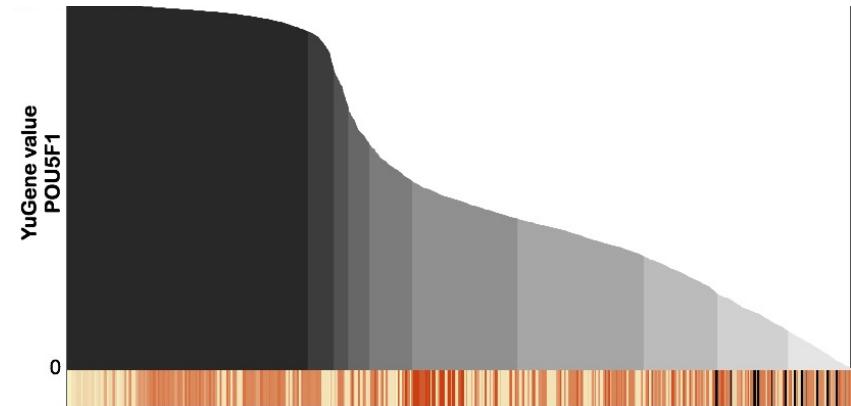
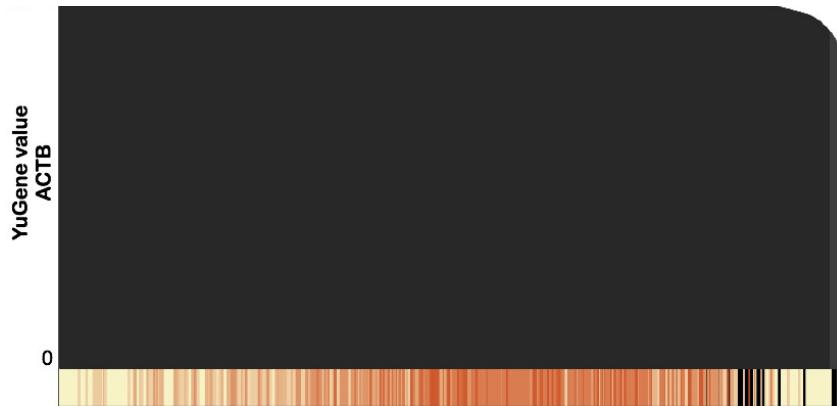
WORKBENCH

Stemformatics.org

Collaboration platform for 'omics data

YuGene

Compare expression across many different datasets



Emergent properties = data integration and visualisation

Intuitive visualization of gene expression.

STEMFORMATICS

Login | Register | Forgot pass phras

GENES > DATASETS > **GRAPHS >** ANALYSES > MY JOBS > ABOUT US >

Gene Search >> CLEC4E >> CLEC4E Summary >> Dataset Browser >> Watkins_2009_19228925 Summary >> Expression Results

CHANGE DATASETS CLEC4E SEARCH

Export & Share Graph Options Analysis Genome Browser Information Help

Gene Expression Graph for Gene CLEC4E grouped by Sample Type
A HaemAtlas: characterizing gene expression in differentiated human blood cells

Log₂ Expression

Med (8.79)

DT (5.00)

CLEC4E Probe

CLEC4E Probe ILMN_1771664

Legend

- megakaryocyte (MK)
- erythroblast (EB)
- granulocyte
- monocyte
- B lymphocyte
- helper T lymphocyte
- cytotoxic T lymphocyte
- natural killer cell (NK)

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Move quickly to additional views/analyses

Customize legend

Hover-over detail



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Genes > Manage Gene Lists > [Gene List View](#)

Gene List View - biomarkers

Show help information

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Hierarchical Cluster

Multi-Gene Expression Graph

Gene List Annotation

Innate DB Pathway Analysis

Innate DB Gene Ontology Analysis

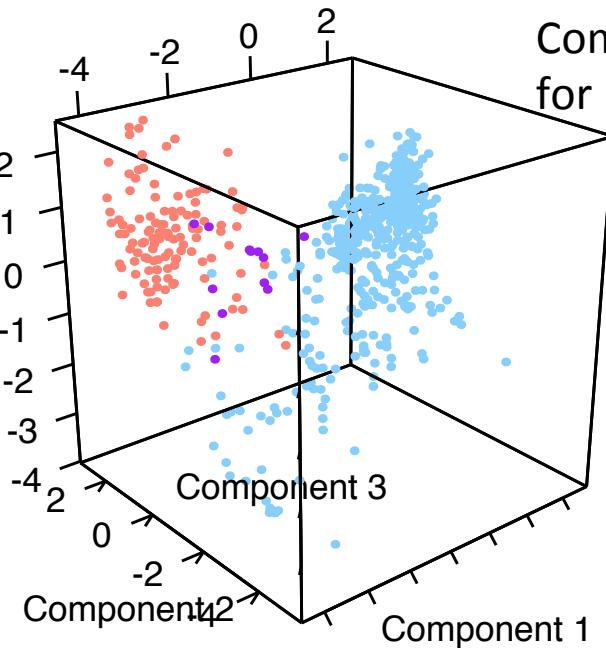
Innate DB Network Analysis

Innate DB Interactor Analysis

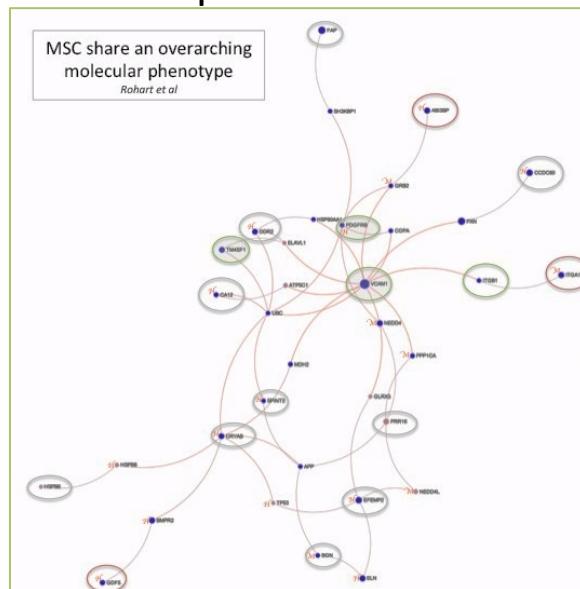
Innate DB Transcription Factor BS Analysis

Gene Name	Ensembl ID	Description	Action
SPATA20	ENSG00000006282	spermatogenesis associated 20	Delete
PROM1	ENSG00000007062	prominin 1	Delete
ERCC1	ENSG00000012061	excision repair cross-complementing rodent repair deficiency, complementation group 1 (includes overlapping antisense sequence)	Delete
SNAI2	ENSG00000019549	snail homolog 2 (Drosophila)	Delete
	ECGF ECGF1		

Combine multiple datasets
for meta-analyses



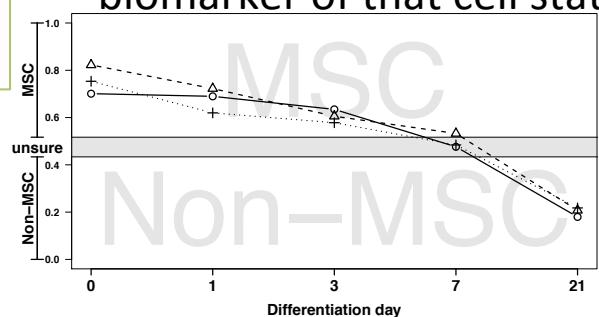
Find signatures of
particular cell states



Florian Rohart



Apply the signature as a
biomarker of that cell state



AIBN, UQ – Wells lab

Rowland Mosbergen

Silvia Manzanero

Othmar Korn

Elizabeth Mason

Florian Rohart

Nick Matigian

Edward Huang

Suzy Butcher

Barbara Rolfe

(Past members)

Anthony Beckhouse

Kelly Hitchens

Dipti Vijayan

Sandra Lo

Jill Shepherd

Ale Vitale

Ashley Waardenberg

Isaac Englert

Celena Heazlewood

UoG Wells lab

Heather Mackie

Shane Kelly

Acknowledgements

StemCore

Ernst Wolvetang

Lars Nielsen

UQ

Mark Walker

Garrie Arumugam

Nicholas Fisk

Sean Grimmond

Robert Ashman

Biostatistics

Kim Anh LeCao – **UQ**

Jess Mar – **AECM, NY**

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Winston Hide

Emmanuel Dimmont

John Quackenbush – **Dana Farber**

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RIKEN Genome Sciences group

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SCA

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Martin Pera (U. Mel)

Nadia Rosenthal

(AARMI)

NHMRC, ARC,

QLD Government

JEM research foundation

Wellcome Trust

ARUK

GLAZGO