

GUDMAP

GenitoUrinary Development Molecular Anatomy Project

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About GUDMAP

The GenitoUrinary Development Molecular Anatomy Project (GUDMAP) is an open access online resource developed by a consortium of researchers working to provide the scientific and medical community with gene expression data, transgenic mice and tools to facilitate research and teaching. Initially GUDMAP focused on the murine urogenital system. Recently, GUDMAP has extended its efforts to include two new elements:

- **Nociceptive GUDMAP (nGUDMAP)** focuses on nociceptors and cell types associated with pain processing for the murine lower urinary tract and pelvic region.
- **Human GUDMAP (hGUDMAP)** extends the gene expression database to include data sets that annotate human bladder, urethra and kidney.

GUDMAP data includes: **Large-scale in-situ hybridisation screens, 3D OPT data, microarray gene expression data and sequencing data** of the developing mouse genitourinary (GU) system. Expression data are annotated using a **high-resolution ontology** specific to the developing GU system.

GUDMAP Data

Summary 'gene-strip' (below) provides an overview of expression data available for each gene. Clickable links connect to in-situ data & images, disease/phenotype associations and microarraydata.

Database Statistics 23-Oct-2015	
Assay Type	Entries Genes
All In Situ Hybridisation (ISH):	10766 3692
Wholemount ISH (WISH):	7352 2896
Section ISH (SISH):	3414 1437
OPT ISH:	64 32
Immunohistochemistry (IHC):	125 31
Transgenic Reporters:	115 37
Microarray:	461 -
Sequencing:	246 -

Example of GUDMAP In-Situ Entry (GUDMAP:20862), displaying images (above) and expression mapped on anatomy ontology (right).

Images: Bladder urothelium, Bladder urothelium, Bladder urothelium, Bladder urothelium, Epithelial layer of the urethra.

Expression Mappings:

- Present (unspecified strength)
- Present (strong)
- Present (moderate)
- Present (weak)
- Uncertain
- Not detected

Expression Pattern Key:

- Homogeneous
- Crystalline
- Regional
- Sporadic
- Ubiquitous
- Restricted
- Single cell

Nerve Density:

- Relative to Total
- Maximum
- Moderate
- Low

Relative to P&A:

- Increase, large
- Increase, small
- Decrease, large
- Decrease, small

Contains note

View annotated components as a list. Show annotation under groups /

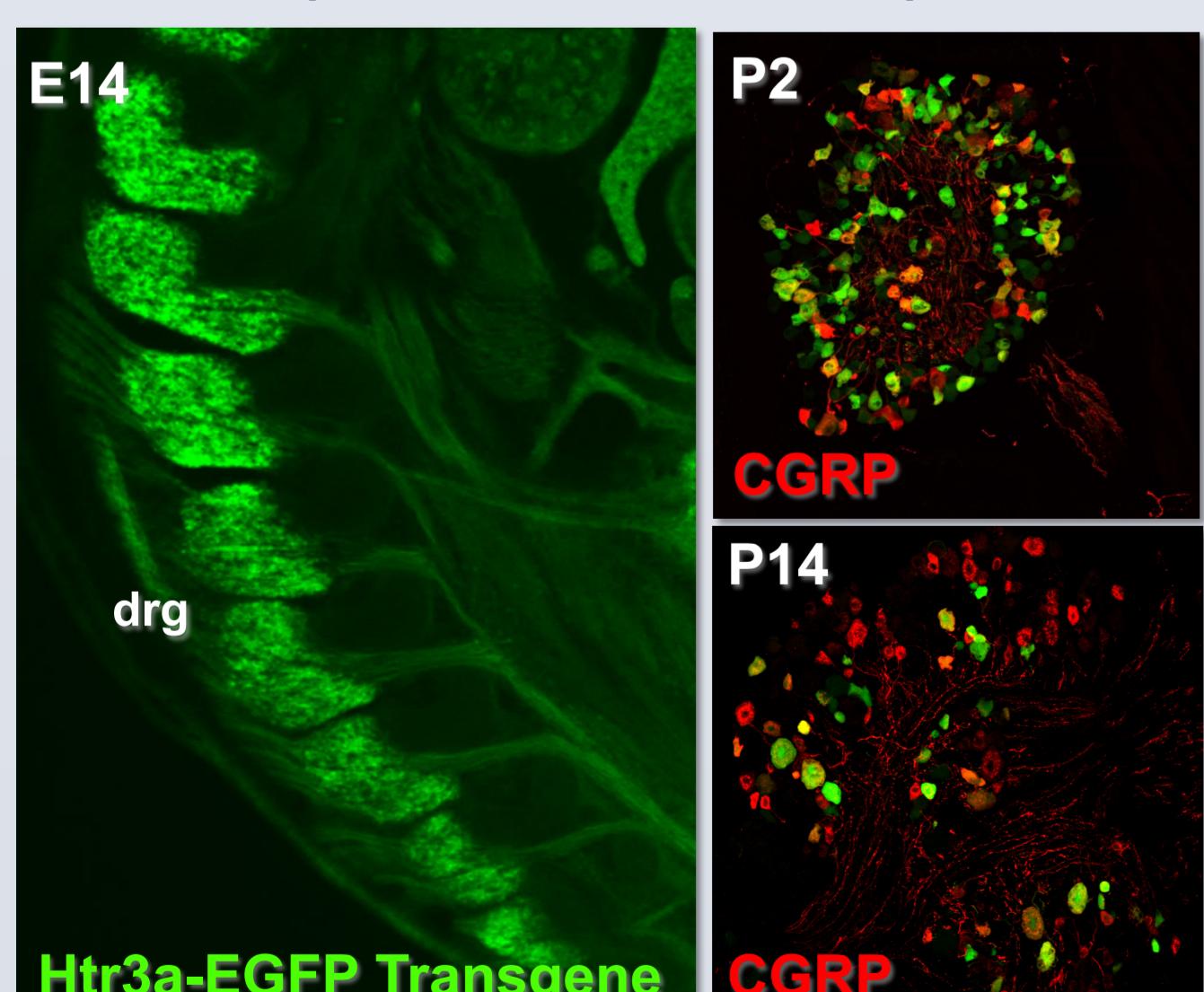
View on UCSC View on IGV

OPT 3D Atlas of Gene Expression in Developing Genital Tercle and Urethra (M. Cohn Lab)

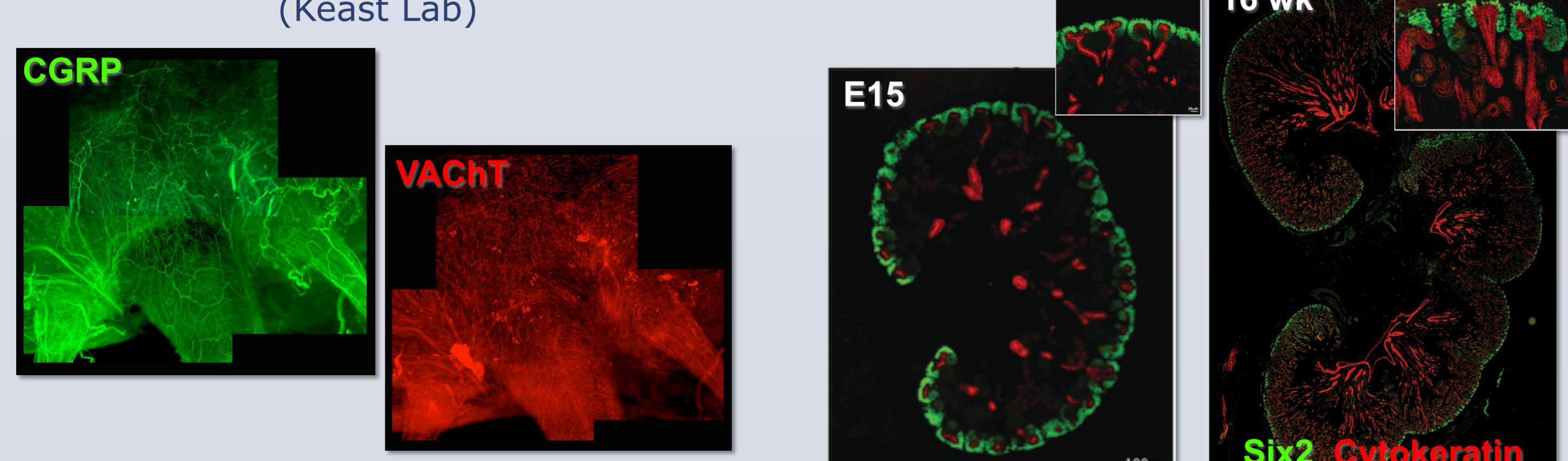


Nociceptive GUDMAP "nGUDMAP"

Developing DRG Immunohistochemistry (Southard-Smith Lab)

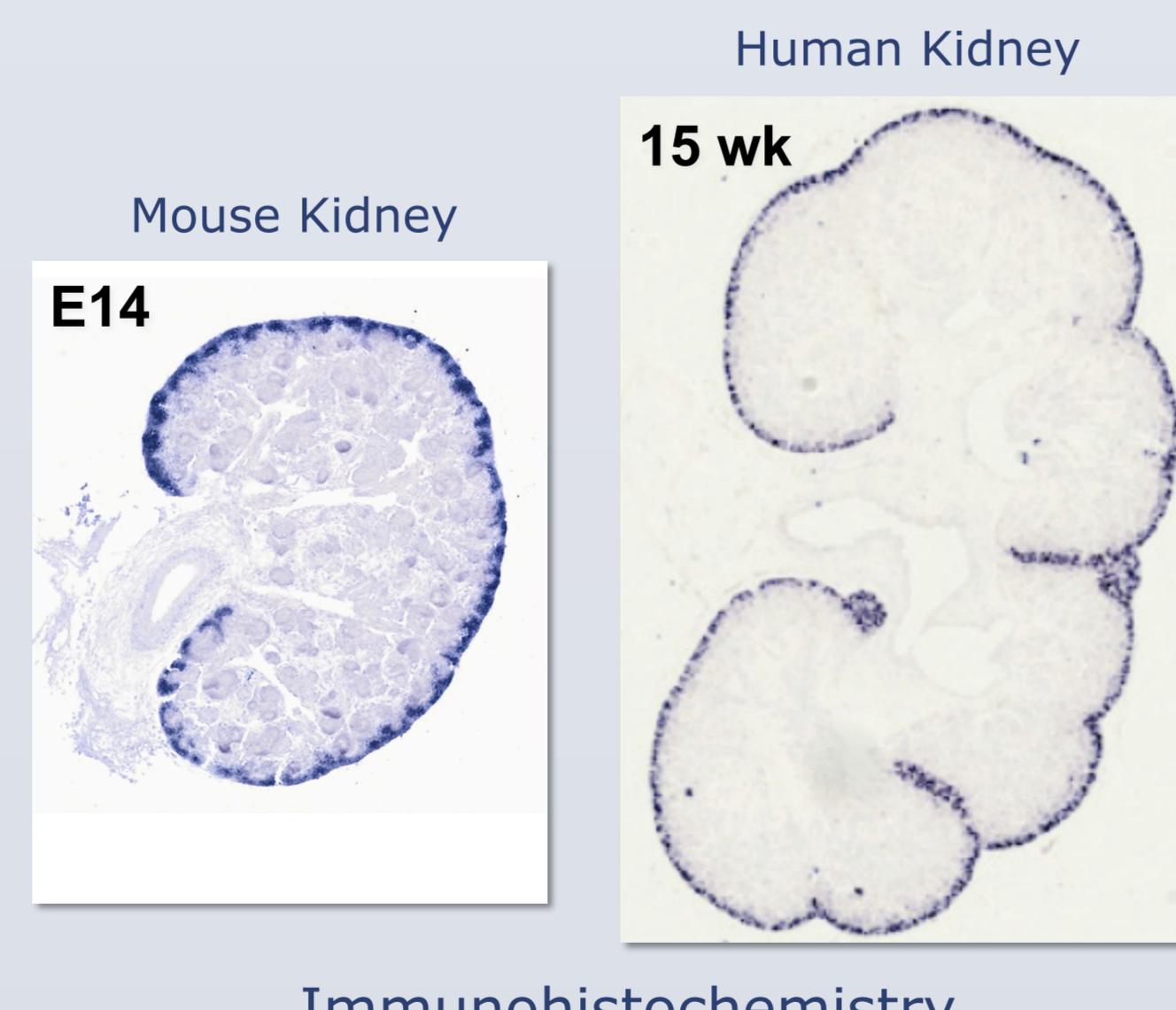


Adult LUT Immunohistochemistry (Keast Lab)

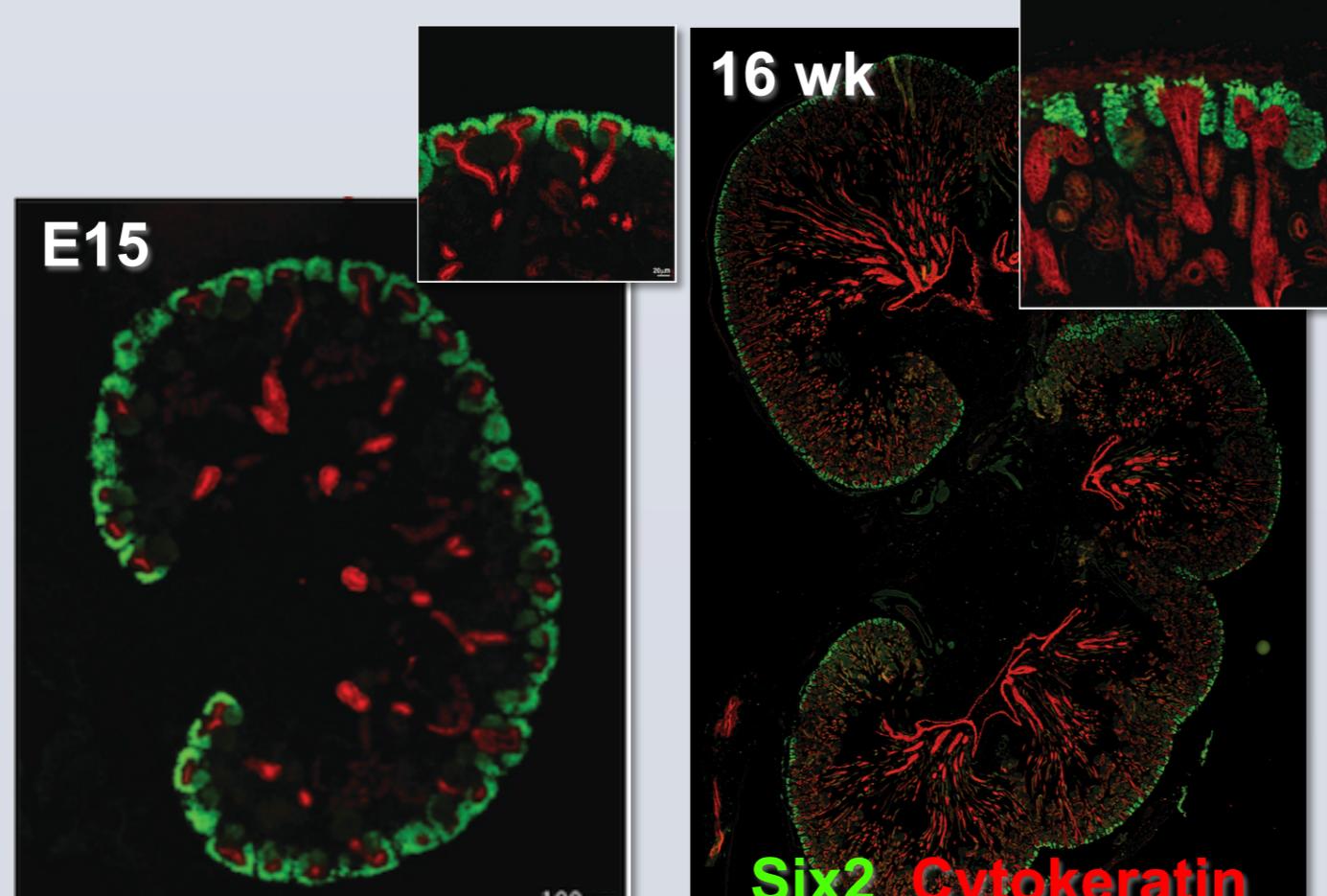


Human GUDMAP "hGUDMAP"

In Situ Hybridization (A. McMahon Lab)



Immunohistochemistry

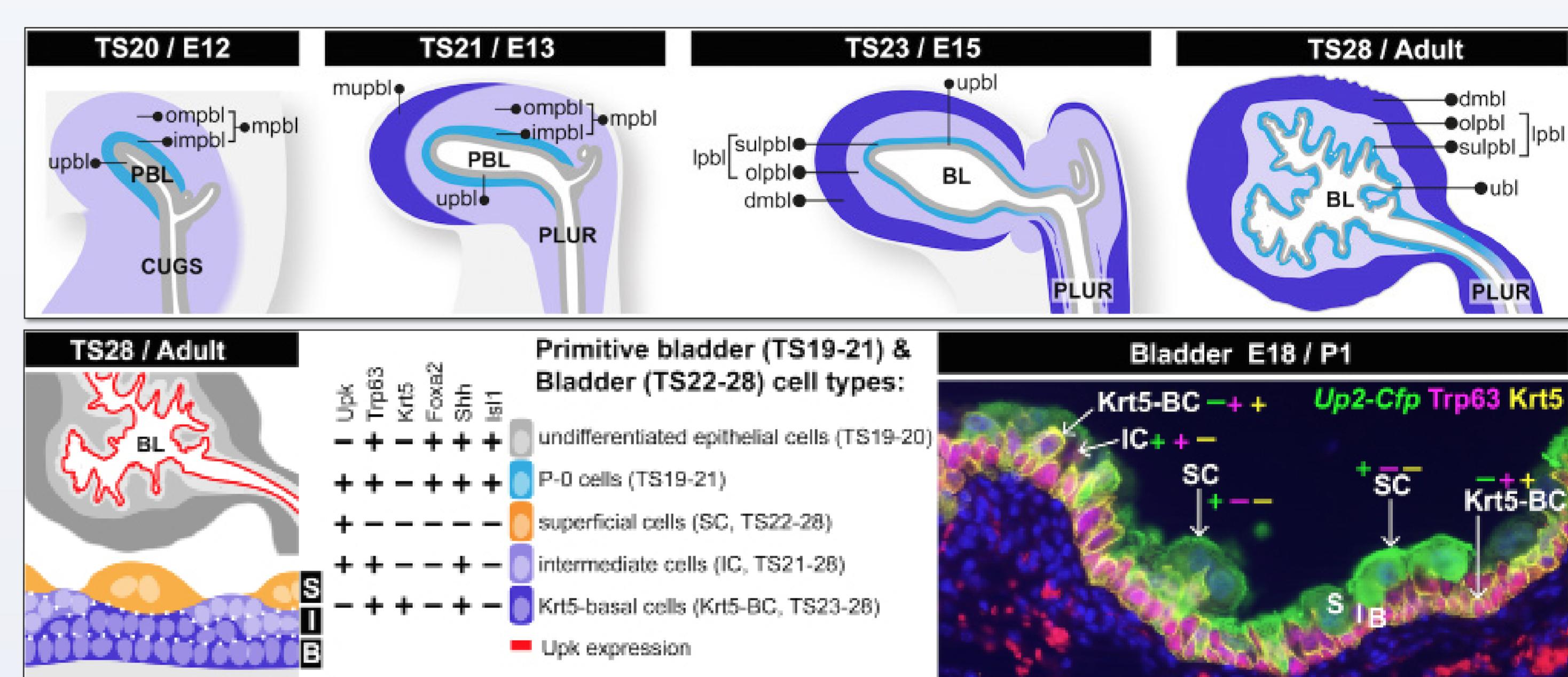


Schematics, Tutorials & Tissue Summaries

GUDMAP holds an extensive archive of high-quality schematics diagrams that illustrate different views of the developing mouse GU system

www.gudmap.org/Schematics/index.php

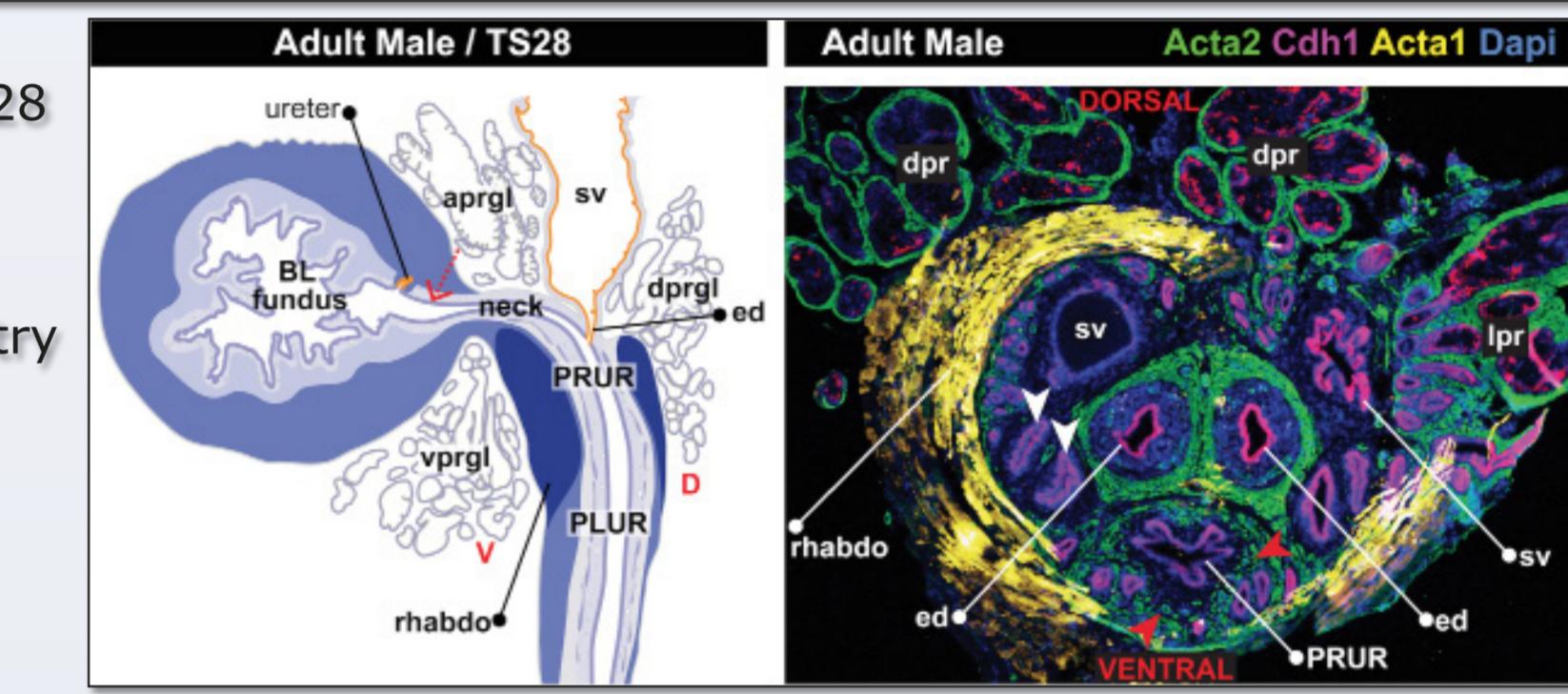
These **tutorials** supplement prior descriptions of GU organogenesis (Matt Kaufman) and enrich the GUDMAP Tissue Summary pages.



Top: Annotated sections of the bladder at TS20 (12 dpc), TS21 (13 dpc), TS23 (15 dpc) and TS28 (adult).

Middle: TS28 (adult) annotated section of the bladder with corresponding immunohistochemistry section.

Right: TS28 (adult) annotated section of the male prostate with corresponding immunohistochemistry section.

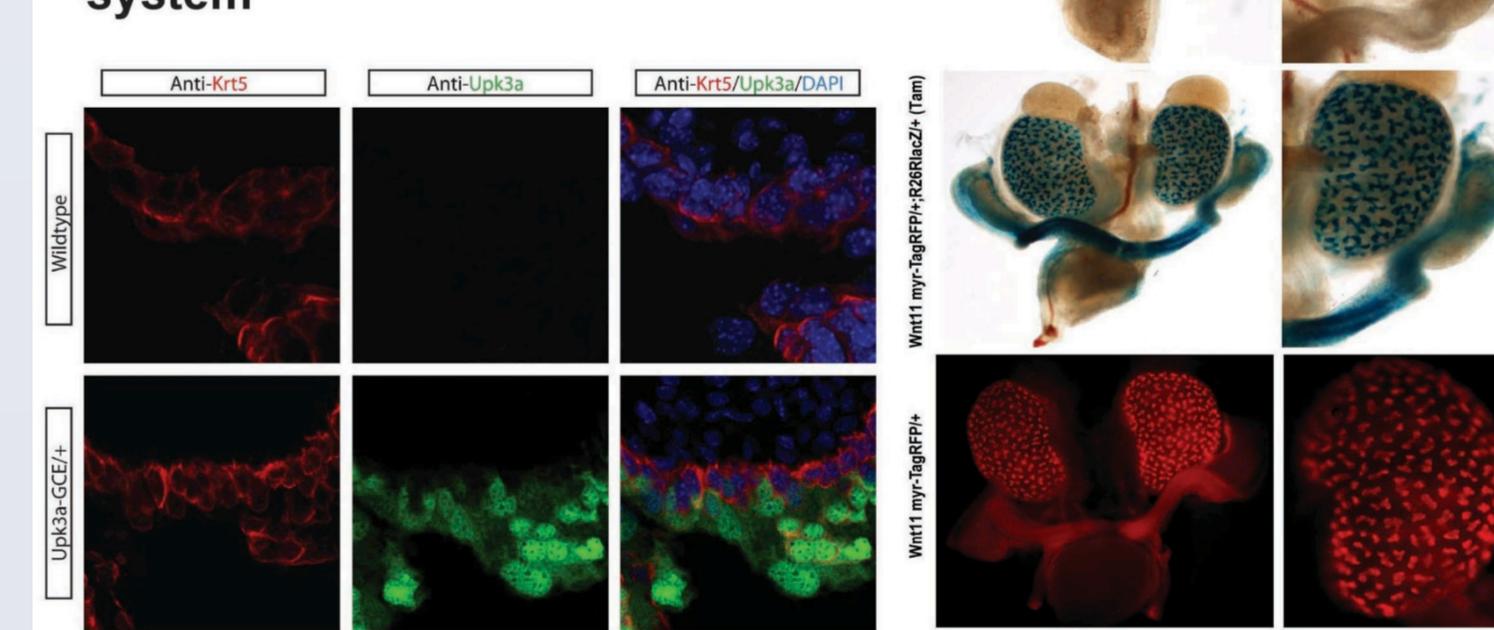


Novel Mouse Strains for Visualising, Isolating and Genetically Manipulating the GU System

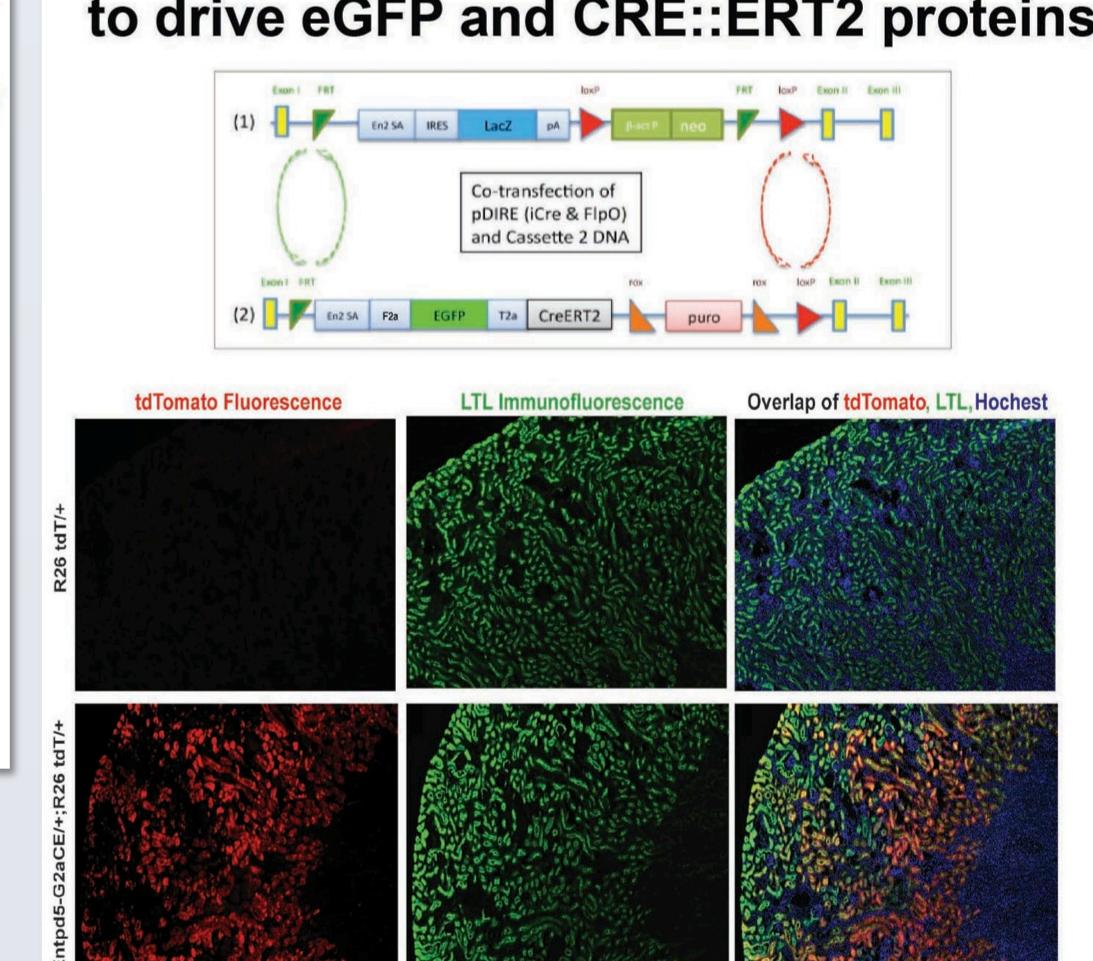
Goals:

- To mark key cell populations in order to isolate, trace or modulate gene activity
- Promote nominations of candidate loci from scientific community.
- Mice made available through the MMRRC (Jackson Labs).
- Nominate strains: www.gudmap.org/MS_GeneNoms.html

Part I: Use BAC mediated mouse transgenesis to drive eGFP and RFPT::Cre::ERT2 fusion proteins in specific cell types in the GU system

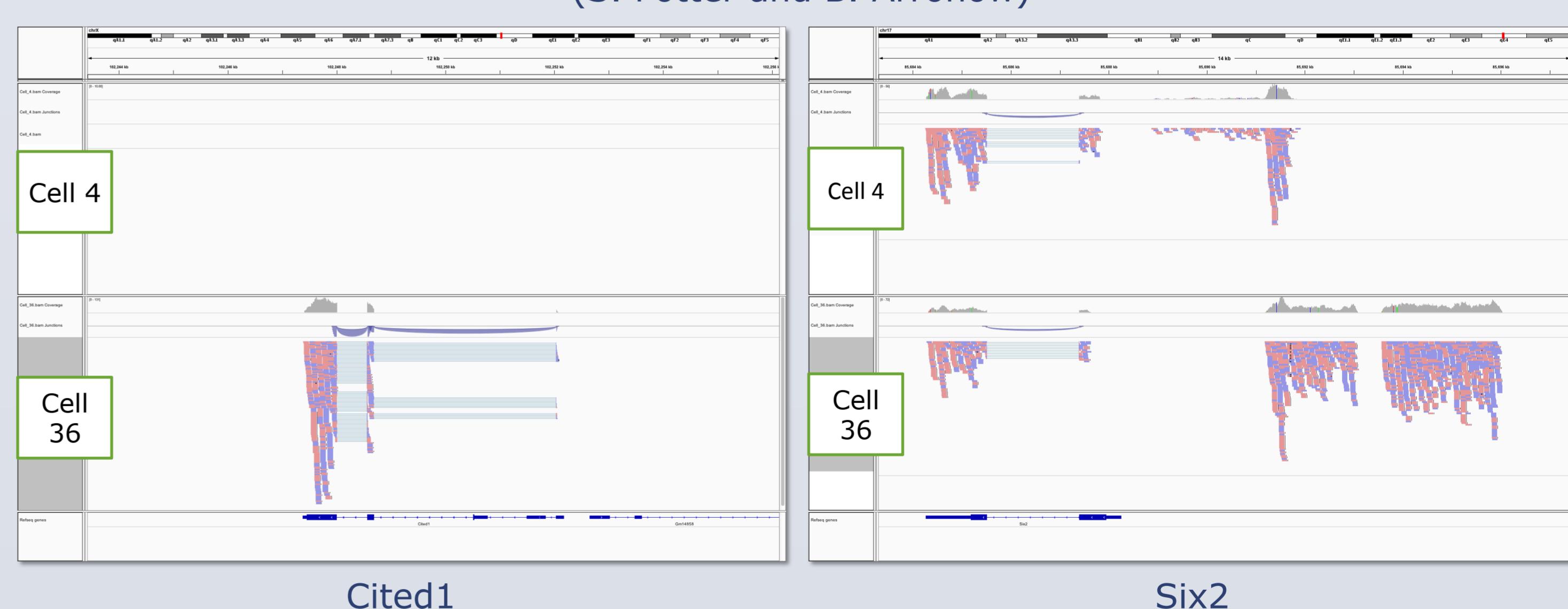


Part II: Obtain ES Cell clones through KOMP(NIH) and EUCOMM Consortia, create new alleles by dual Recombinase Mediated Cassette Exchange (dRCME) to drive eGFP and CRE::ERT2 proteins



Gene Expression Profile Analyses of GUDMAP Data

IGV Genome Browser View of Single Cell Data -- E11.5 metanephric mesenchyme (S. Potter and B. Aronow)



References

All past contributors to GUDMAP can be found at www.gudmap.org/About/Projects/

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