



## **Financial Disclosures**

- Genentech Pathology Advisory Board
- UroSEEK Patent application pending

### **Overview**

- The Dual Universes of BC NMIBC vs MIBC
- The Impact of Genomics on BC Novel TAXONOMY ? Prognostics Predictive Rx Targets
  ImmunoRx Response Predictors
- UroSEEK

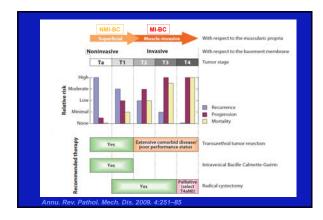
## **Bladder Cancer (BC)**

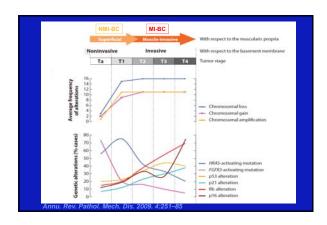
- **Disease Costs and Management Opportunities**
- Major health care cost burden:
  - Frequent cystoscopy, high rate of recurrence etc..
  - \$ 3-4 Billion per year in USA alone
  - o HIGHEST COST per patient for any type of cancer
- Unique amenability to applying molecular detection methods to urine sample (e.g. TERT)

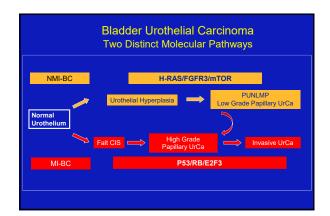
### **Urothelial Carcinoma** Two Phenotypes?

(Superficial) Non-muscle invasive BC (NMI-BC) 70-80%

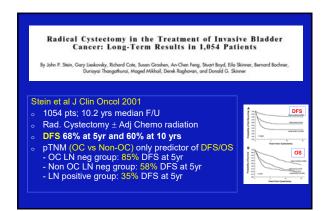
> Muscle Invasive BC (MI-BC) 20-30%

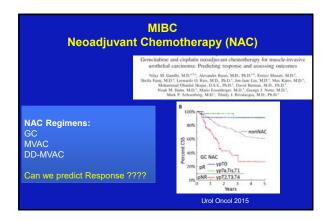


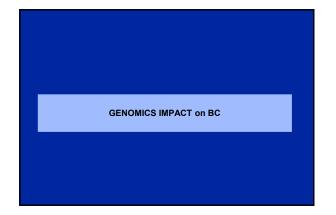


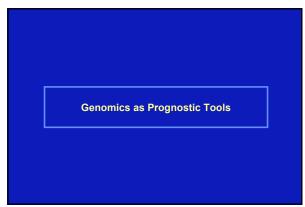












Integrated Genomic and Gene Expression Profiling Identifies Two Major Genomic Circuits in Urothelial Carcinoma

David Lindgren\*, Certified Splatals\*, Martin Laura\*, Johan Staaf\*, Gunilla Chelif\*, Kristina Lindgren\*, Statistica Sulgium\*

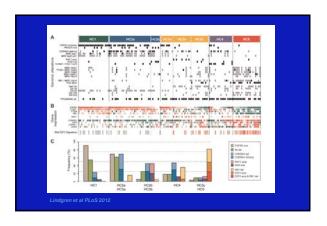
Two Genomic Circuits:

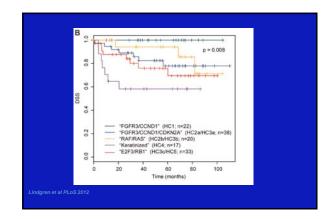
\* FGFR3 mut/Ampl; \* CCND1; PIK3CA mut; 9q (CDKN2A) deletions

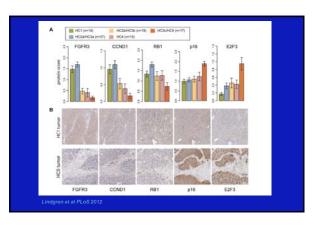
\* E2F3 ampl; RB1 del; PTEN del; \* CDKN2A; CCND1 loss, 5p gain

P53/MDM2 alterations in both circuits at advanced Dz

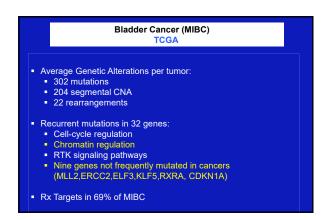
Lindgren et al. PLoS 2012





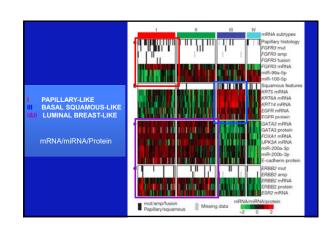


# Comprehensive molecular characterization of urothelial bladder carcinoma The Cancer Genome Atlas Research Network\* Nature: March 2014 Integrated genomic Analysis of 131 URCa

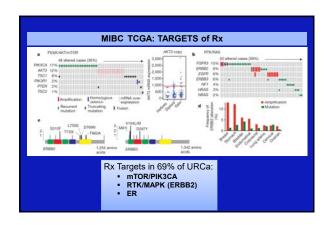


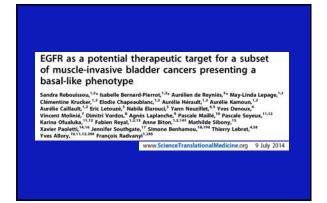
NEW GENOMIC TAXONOMY?
TCGA

INTEGARTED GENE EXPRESSION SUBTYPES
PAPILLARY-LIKE
BASAL / SQUAMOUS-LIKE
LUMINAL / BREAST-LIKE

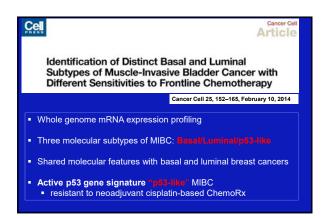


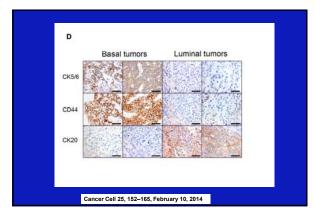
Genomic Based Novel Therapeutic Targets

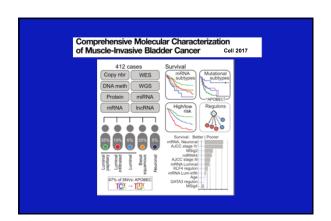


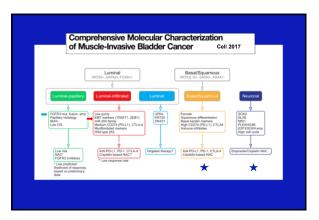


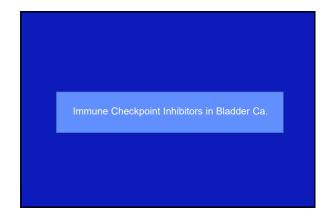
Genomic Based Predictor of Neoadjuvant ChemoRx Response

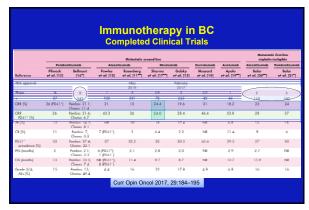


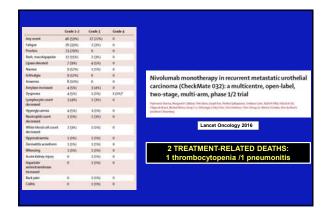


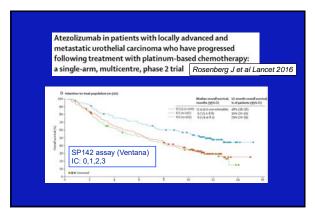


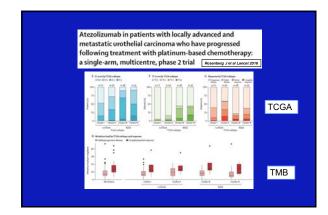


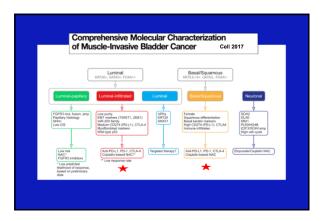












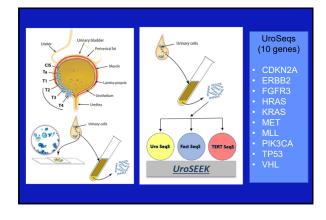
Mechanisms Driven Predictive Biomarker for Immune Checkpoint Inhibitors

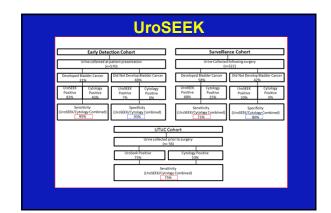
• Immunologic Biomarkers IHC/Gene Expression:
• PDL-1
• CD8+
• CD8+/Ki67
• TIM3, LAG3, IDO
• ICOS, GITR
• Chemokines Signature (Interferon 8)
• Tumor Mutational Load/Burden: NSCLC, Melanoma, BC
• Neoantigens
• dMMR/MSI: e.g. CRCa, Endometrial Ca.

• Viral Oncoproteins: HTLV-1, HPV, EBV, KSV, MCPyV, HBV, HCV

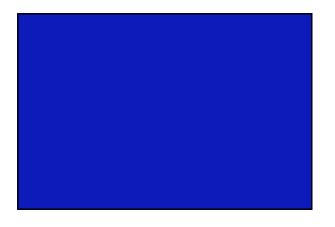
Molecular Early Detection & Surveillance Markers
The Power of Genomics

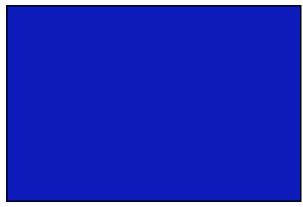






## Genomics Impact on Bladder Cancer Management: PGx Predictive Rx Targets Early Detection and Surveillance New GENOMIC TAXONOMY is being refined and awaits TRANSLATION INTO OUR LABS Immune Check Point Inhibitors are opening new doors in BC management Thank You !!!







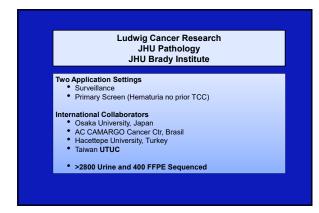
# Conclusions Genomics Impact on Bladder Cancer Management: P GX Predictive Rx Targets New GENOMIC TAXONOMY is being refined and awaits TRANSLATION INTO OUR LABS Immune Check Point Inhibitors are opening new doors in BC management Thank You !!!

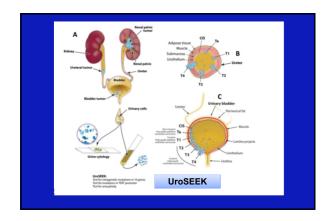
TERT promoter mutations occur frequently in gliomas and a subset of tumors derived from cells with low rates of self-renewal

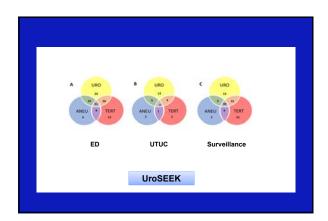
Patrick J. Killela<sup>21</sup>, Zachary J. Reitman<sup>21</sup>, Yuchen Jao<sup>21</sup>, Chetan Bettegovadh<sup>21</sup>, Mishart Agrawal<sup>21</sup>, Luis A. Diaz, Jr.<sup>2</sup>, Allan H. Friedman<sup>21</sup>, Gary L. Gallia<sup>21</sup>, Beppin C. Giovanella<sup>2</sup>, Arthur P. Grofilman<sup>2</sup>, Tong-Chuan He<sup>3</sup>, Yiping He<sup>3</sup>, Ralph H. Hruban<sup>3</sup>, George L. Jallo<sup>3</sup>, Nish Mandahi<sup>3</sup>, Alan K. Meekeh<sup>210</sup>, Fredrik Mertens<sup>3</sup>, George J. Netto<sup>3</sup>, Sehmed Rasheed<sup>3</sup>, George J. Jigigins<sup>3</sup>, Thomas A. Rosenquist<sup>3</sup>, Mark Schiffman<sup>3</sup>, Leiming Shih<sup>3</sup>, Dan Theodorescu<sup>3</sup>, Michael S. Torbenton<sup>3</sup>, Victor E. Velculescu<sup>3</sup>, Tian-Li Wang<sup>3</sup>, Nicolas Wentzenser, Laura D. Wood<sup>3</sup>, Ming Zhang<sup>3</sup>, Roger E. McLendon<sup>3</sup>, Darel D. Bigner<sup>2</sup>, Kenneth W. Kinzler<sup>3</sup>, Bert Vogelstein<sup>3,3</sup>, Nickolas Papadopoulos<sup>3</sup>, and Hal Yan<sup>3,4</sup>

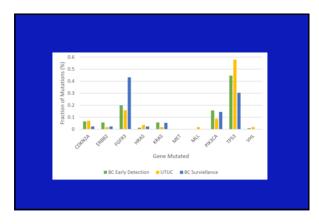
PNAS | April 9, 2013 | vol. 110 | no. 15 | 6021–6026

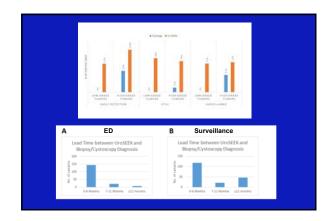


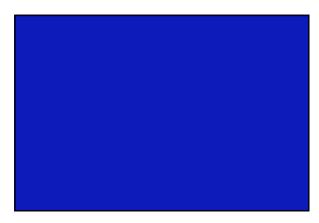


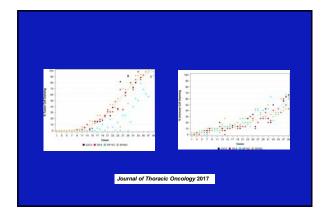


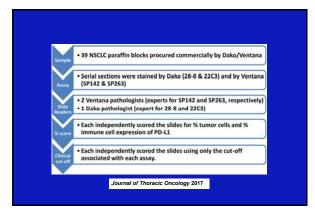


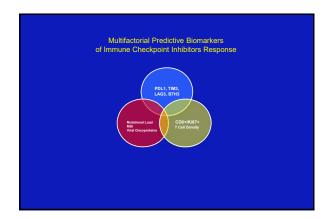


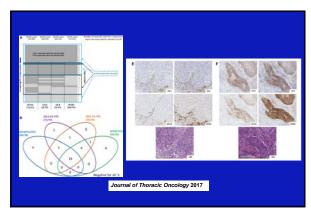












PD-L1 Immunohistochemistry Assays for Lung Cancer: Results from Phase 1 of the Blueprint PD-L1 IHC Assay Comparison Project

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