

Molecular Characterization of Bladder Cancer in Smokers versus Nonsmokers

Monika Joshi¹, MD, MRCP, Sherri Z. Millis², MS, PhD, Donald Lamm³, MD, Sandeep Reddy², MD, Sheldon L. Holder¹, MD, PhD, Nicholas J. Vogelzang⁴, MD, Joseph J. Drabick¹, MD ¹ Penn State Hershey Cancer Institute, Hershey, PA; ²Caris Life Sciences, Phoenix, AZ; ³BCG Oncology, Phoenix, AZ; ⁴Comprehensive Cancer Centers of Nevada, Las Vegas, NV

Abstract #4528 Background

- Bladder cancer (BC) is one of the most common malignancies of the urinary tract and is the 4th most common cancer among men.
- It is estimated that by the end of 2015, the US will have approximately 74,000 new BC cases, accounting for 16,000 cancerrelated deaths.
- Smoking is considered an important risk factor for BC. Recent data demonstrate an increase in BC incidence in nonsmokers as well.
- Molecular characterization of BC in nonsmokers has not been well studied.

Methods

- 676 consecutive BC profiled at a CLIA-certified laboratory from 2006 through 2014 were evaluated for differences in molecular characterization between smokers and nonsmokers.
- Smoking status (nonsmokers [NS]; current or reformed smokers [R/S]), patient characteristics, age, sex and survival data were collected on each subgroup.
- Formalin fixed paraffin-embedded (FFPE) samples were analyzed. Tumors were verified by a board-certified pathologist to confirm diagnosis.
- Protein expression was determined by IHC analysis, using commercially available detection kits and automated staining techniques. In-situ hybridization, fluorescent (FISH) or chromogenic (CISH), was used for evaluation of *HER-2/neu* and *EGFR*. *ALK* rearrangements, either inversion or translocation of the ALK gene at 2p23, were identified by separation of the fusion signals.
- Gene sequencing was performed by next-generation sequencing (NGS). Full details are available at www.carislifesciences.com.

Results

- Identified trends included differences in the PI3 kinase, Wnt and EGFR pathways.
 - •Percentage of *PIK3CA* mutations was higher in NS (43%) than R/S (11%).
 - •Wnt pathway aberrations (e.g. *CTNNB1* and *APC* mutations) occurred more frequently in R/S.
 - •*EGFR* amplification occurred in 22% NS but only 11% in R/S. •HER2 was amplified only in R/S (23% vs. 0%, p=0.05).
- Three of eight (37.5%) R/S had an ALK 2p23 rearrangement. For comparison, ALK is found in ~5% non-small cell lung cancers.
- *TP53* did not differ between the populations.
- Survival data from 31 patients (14 NS, 17 R/S) showed overall average survival in the NS cohort was 175 days longer than in the R/S cohort.

Patient Demographics



Lif

Figure 1. Distribution of 69 cases with known smoking status. Data were provided by reformed smokers (55%).

Results, Immunohistochemistry (IHC)

EGFR (anti-EGFR) TOP2A (anthracyclines) SPARC (nab-paclitaxel) PDGFRA (TKI's) AR (hormonal therapies) cKIT (CKIT inhibitors) low MGMT (temozolomide) Low PTEN (mTor inhibitors) low RRM1 (gemcitabine) ERCC1 (cisplati

TOPO1 (topotecan, irinotecan) Her2 (Her2-Targeted therapies) low TS (capecitabine, fluorouracil

Table 1. Categorization of cases and gender/age breakdown.

ategory; Total Cases	Gender; Age	
ll bladder cases, n=676	M: 75%; F: 25%	
	Median age: 66 (range 20-95)	
etime nonsmoker, n=30	M: 21%; F: 9%	
	Median age: 66	
Current smoker, n=8	M: 6%; F: 2%	
	Median age: 65	
eformed smoker, > 5 yr,	M: 22%; F: 9%	
n=31	Median age: 69	
Grouped, total current or reformed smoker (R/S), n=39		



participating physicians. Most were current or

Results, Gene Sequencing

Figure 3. Mutation frequency. Genes are shown from left to right by decreasing percent mutation. Genes with no alterations identified included ALK, CSF1R, FGFR1, FLT3, GNA11, GNAQ, GNAS, JAK2, JAK3, MPL, NOTCH1, NPM1, PTPN11, VHL. Genes seen in just one case included ABL, AKT1, ATM, BRAF, CDH1, cKIT, cMET, HNF1A, KDR, MLH1, NRAS, PDGFRA, RET, SMAD4, SMARCB1. Additionally, while no cases in the subgroup were tested for *BRCA1/2*, the overall incidence in bladder cancer was 6.5% for BRCA1 and 13% for BRCA2.



Figure 2. Levels of protein expression. Overexpression is reported as percent positive of total cases tested; loss is reported as percent negative.



Results, Fluorescence or Chromogenic *in situ* Hybridization (ISH) Figure 4. Amplification or rearrangement. HER2 amplification was not



seen in documented lifetime nonsmokers (p value = 0.05).

Results, Overall Survival

Figure 5. Kaplan		
Meier Curve.		
Overall survival		
curves of NS (red)		
versus		
R/S (blue) from		
the date of		
diagnosis. Statistical		
significance was not		
reached (p=0.21)		
due to small sample		
size.		



175 days longer in the NS cohort.

	Avg Days Survived	Range	Alive
NS, n=14	843	185 – 2966	6/14
R/S, n=17	668	112 – 1479	3/17

Conclusions

- treatment options.
- More studies need to be conducted to identify other nonsmokers.
- clinical trial design in this subgroup of BC patients.
- subset for PARP inhibitor clinical trials.
- associated outcomes.

References

- bladder carcinoma. Nature 507:315-22, 2014.
- bladder cancer among men and women. JAMA 306:737-45, 2011.
- Nonbladder Origin. *Clinical Genitourinary Cancer*, 2014.



Table 2. Average survival, range, and vital status. Average survival was

• The difference in molecular biology between R/S and NS with BC suggests a different oncogenesis, with potentially different

mutational abnormalities between smokers vs. lifetime

• Increased incidence of *PIK3CA* mutations in NS may inform

• *BRCA1* and *BRCA2* testing in bladder cancer might identify a

• Follow-up on the ALK translocation patients is recommended to determine whether crizotinib was utilized and, if so, review

Millis et al. Molecular Profiling of Infiltrating Urothelial Carcinoma of Bladder and

^{..} Cancer Genome Atlas Research N: Comprehensive molecular characterization of urothelial 2. Freedman ND, Silverman DT, Hollenbeck AR, et al: Association between smoking and risk of