conquer breast

1st GBO Meeting

1st European Course for MDs in training Going Beyond in Oncology

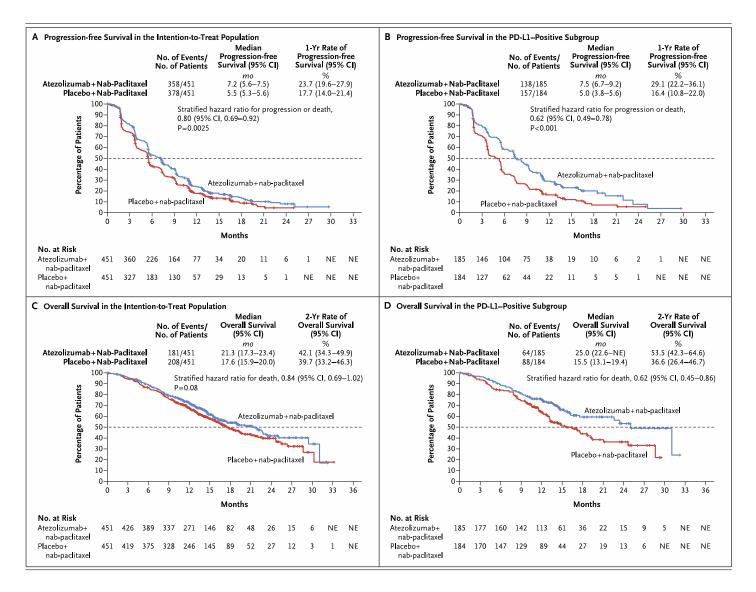
NOVEMBER, 28th 2018 MILANO

ROSA GRAND HOTEL MILANO Piazza Fontana. 3

Tissue biomarkers for immunotherapy

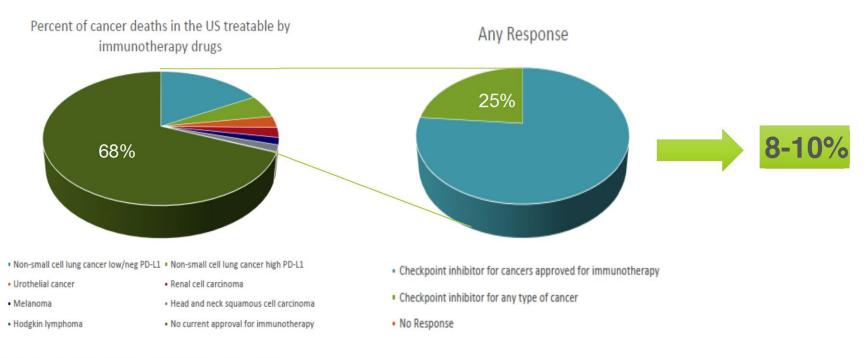
Giancarlo Pruneri, MD
University of Milan
National Cancer Institute INT, Milan

Breast cancer: immunotherapy goes on stage



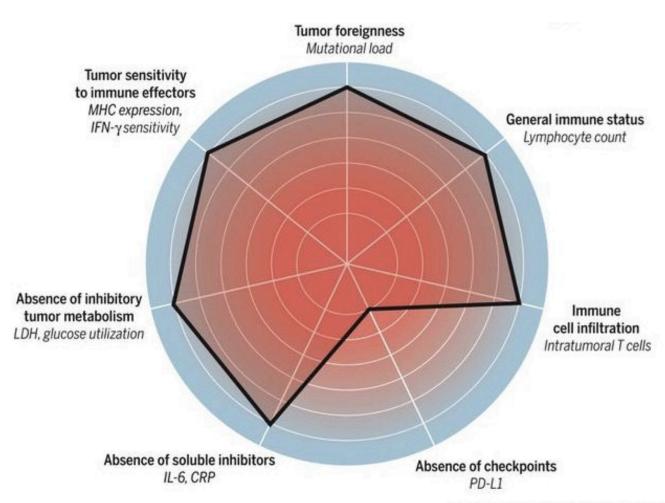
Immuno-Oncology's Substantial Clinical Impact

Few Patients Actually Benefit from Currently Approved IO Drugs



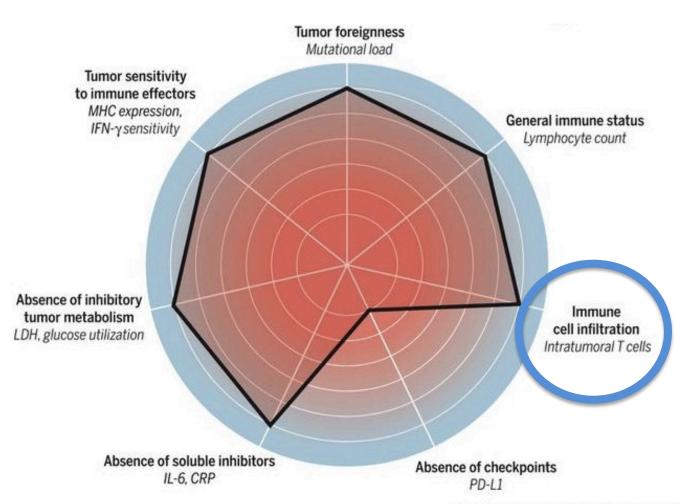
Source: Adapted from Abola and Prasad, JAMA, 2016

The cancer immunogram



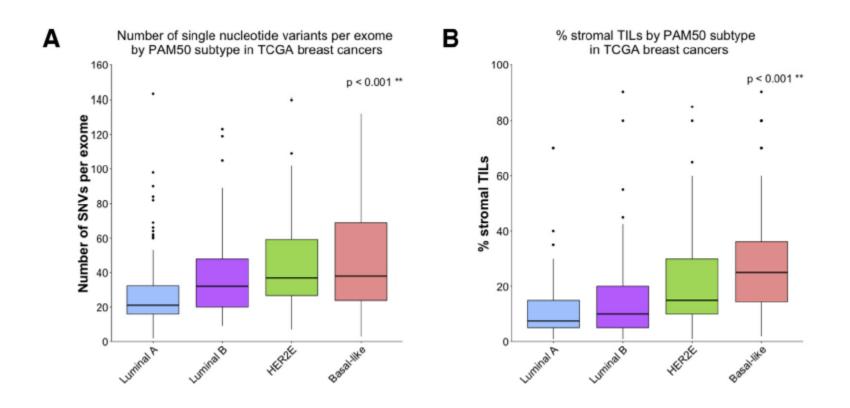
Christian U. Blank et al. Science 2016;352:658-660

The cancer immunogram

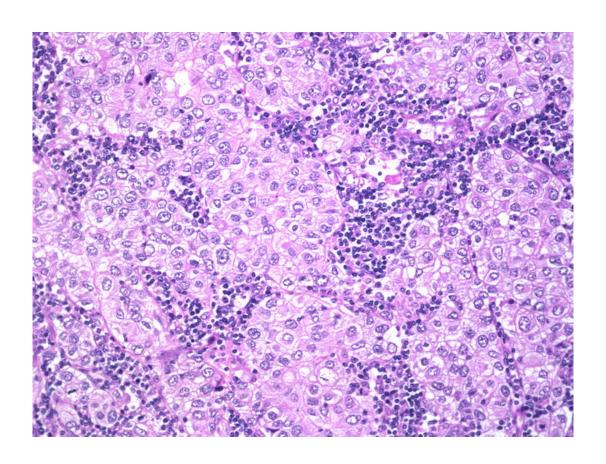


Christian U. Blank et al. Science 2016;352:658-660

TILs and mutational load in BC subtypes



TILs: what if...

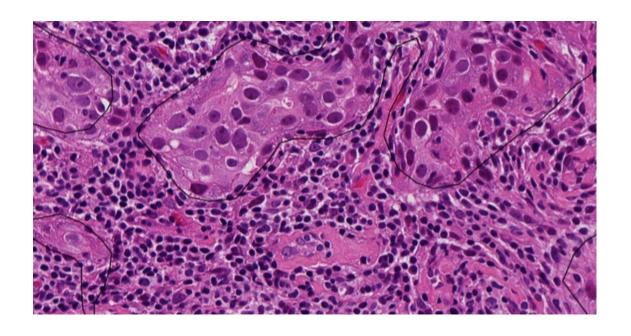


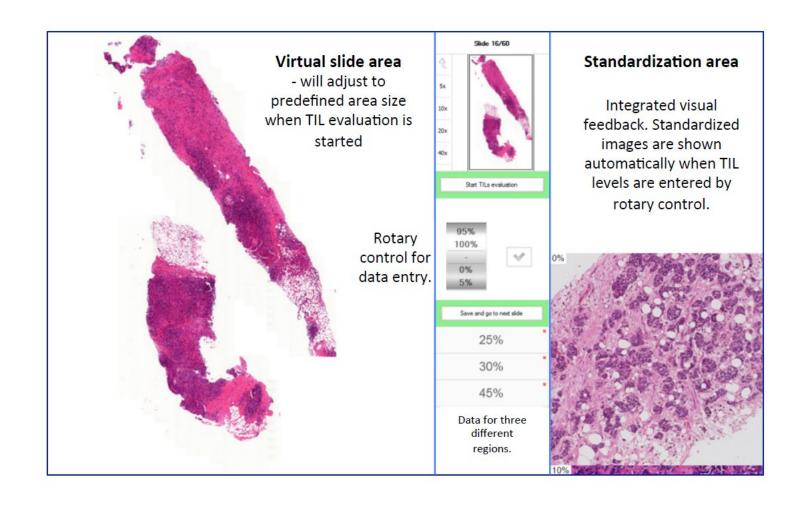
Do TILs have validity?

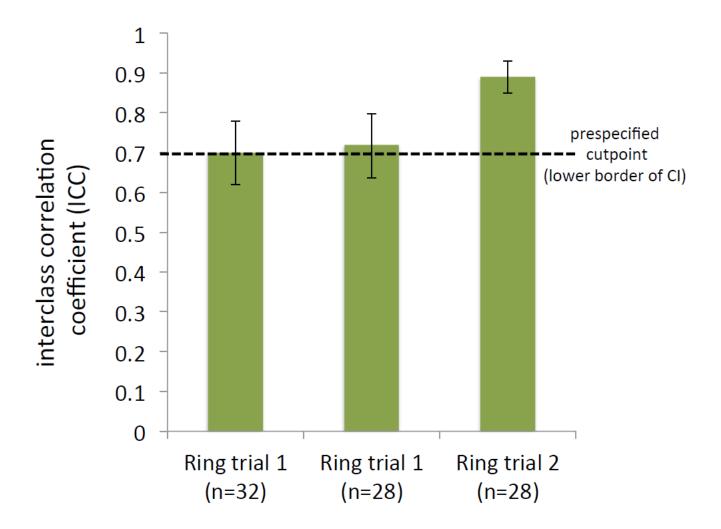
- Analytical validity refers to the accuracy, reliability, and reproducibility of the assay as demonstrated by preanalytic, technical, and scoring or interpretation methods
- Clinical validity refers to the ability of a tumor biomarker test to divide one population into two or more groups that differ either biologically or clinically. CV alone is insufficient to recommend that the test be used to guide treatment decisions

Pre-defined parameters for TILs evaluation in breast cancer

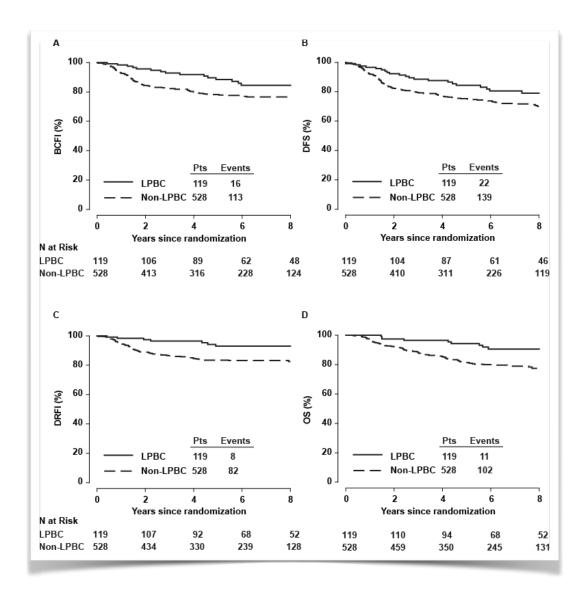
Stromal TILsBetween the tumor cells but within tumor stroma



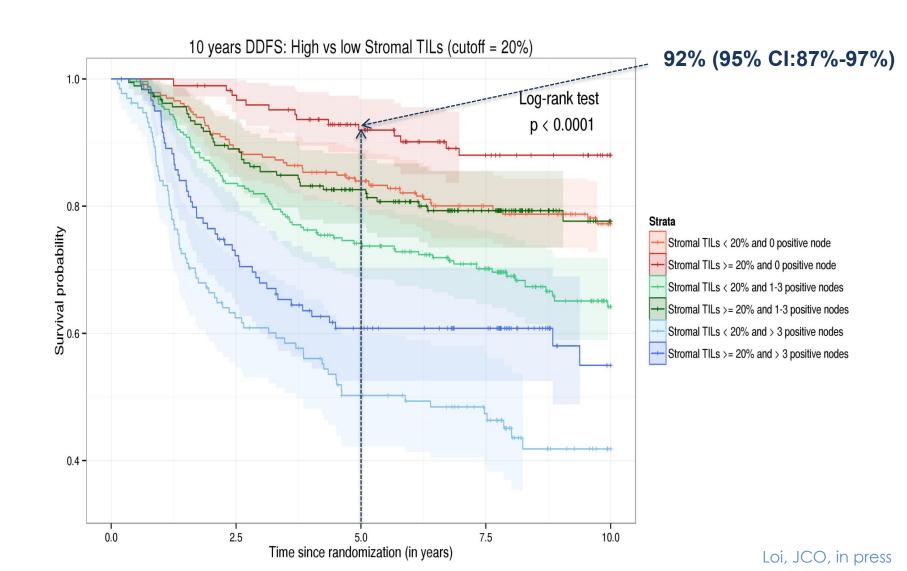




TILs guidelines in prospective trials: IBCSG 22-00



Stromal TILs ≥20% in node negative TNBC patients have excellent 5 yr D-DFS estimated by Kaplan-Meier



Clinical utility

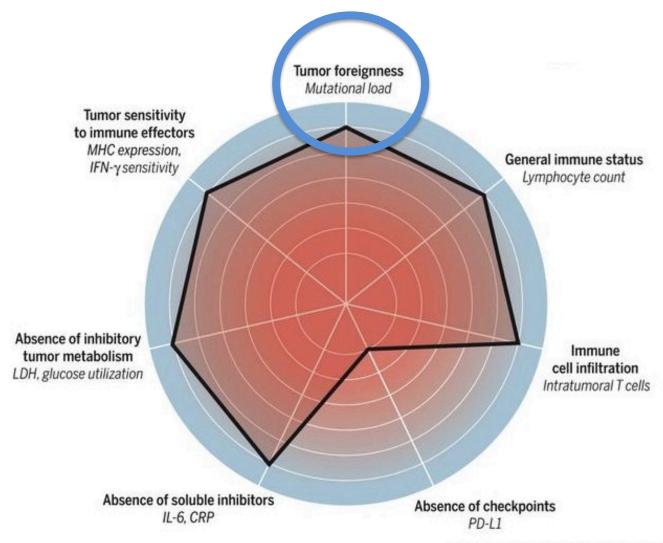
- A biomarker-based test has clinical utility if its use is associated with a favorable balance of benefits to harms compared with treatment of the patient in the absence of the biomarker test results
- Benefits may include improvement in survival end points (EFS, DFS, PFS, OS)
- A biomarker must contribute clinically useful information beyond that already provided by clinical or pathologic indicators in standard use, unless the new test can provide equivalent information at lower cost, less invasively, or with less inconvenience or risk. The magnitude of the benefit must be clinically meaningful and outweigh risks, costs, and/or inconvenience associated with use of the test

Use of Biomarkers to Guide Decisions on Adjuvant Systemic Therapy for Women With Early-Stage Invasive Breast Cancer: American Society of Clinical Oncology Clinical Practice Guideline

Lyndsay N. Harris, Nofisat Ismaila, Lisa M. McShane, Fabrice Andre, Deborah E. Collyar, Ana M. Gonzalez-Angulo, Elizabeth H. Hammond, Nicole M. Kuderer, Minetta C. Liu, Robert G. Mennel, Cathy van Poznak, Robert C. Bast, and Daniel F. Hayes

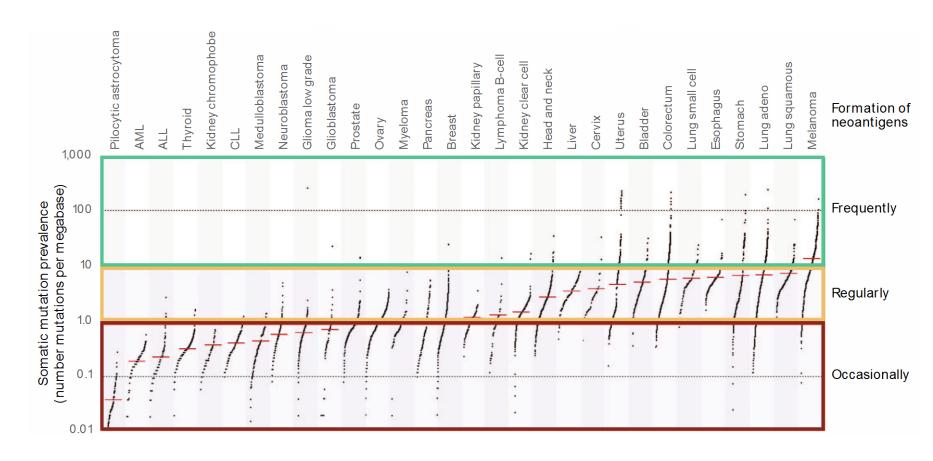
- If a patient has ER/PgR-positive, HER2-negative (node-positive or node-negative) breast cancer, the clinician should not use TILs lymphocytes to guide decisions on adjuvant systemic therapy. Type: informal consensus. Evidence quality: insufficient. Strength of recommendation: strong.
- If a patient has HER2-positive breast cancer or TN breast cancer, the clinician should not use TILs to guide decisions on adjuvant systemic therapy. Type: evidence based. Evidence quality: intermediate. Strength of recommendation: strong.

The cancer immunogram

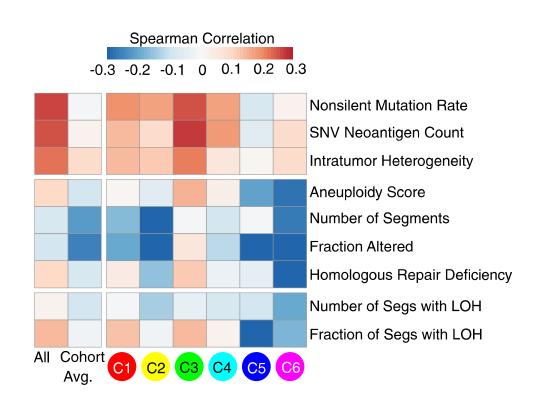


Christian U. Blank et al. Science 2016;352:658-660

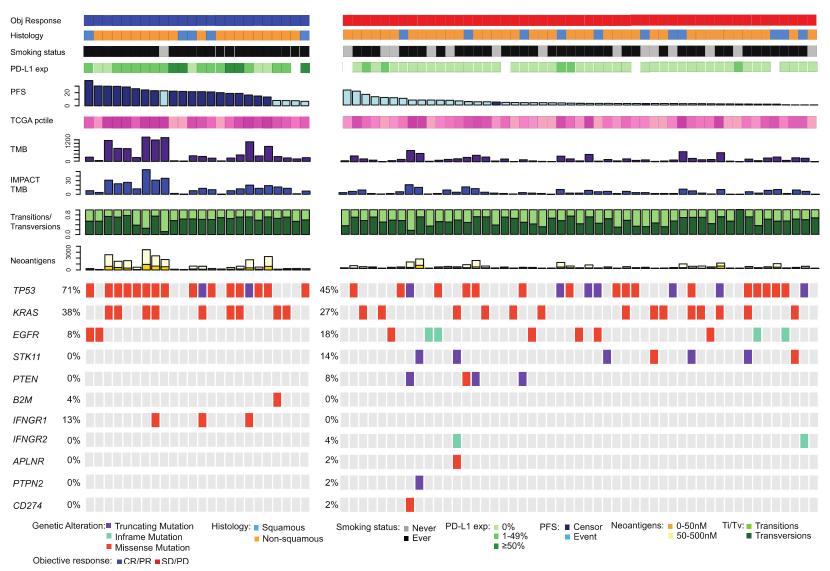
Tumors with >150 non-syn mut (10 mut/mb DNA) form neoantigens



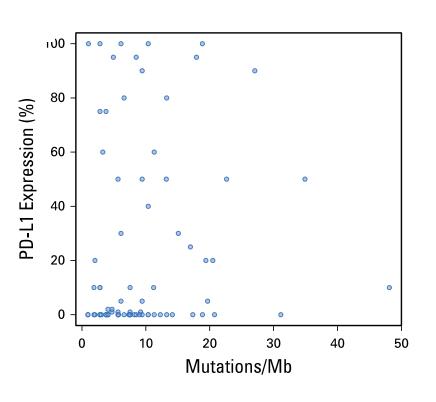
Immune infiltration correlates with tumor somatic variation

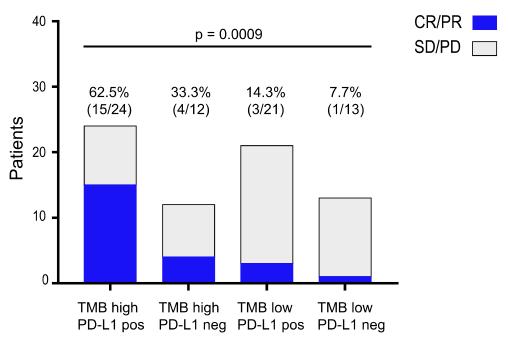


TMB in clinical trials CheckMate-012 Nivo+Ipi NSCLC



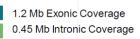
PDL-1 IHC and TMB: towards an integrate approach?

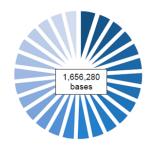




TMB in the real world INT experience on 35 early NSCLC patients

The Oncomine Tumor Mutation Load Assay covers 1.7Mb across 409 cancer-driver genes, relevant across major cancer types





- · signaling cascades
- apoptosis genes
- DNA repair genes
- transcription regulators
- inflammatory response genesgrowth factor genes

- Set up
- TAT
- Data interpretation
- Reproducibility

Signature Pattern of Somatic Mutations - Background

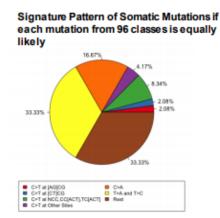
Carcinogens/Biological Processes have unique Patterns

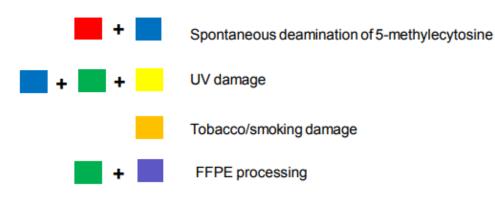
High C>T at CpG is consistent with Spontaneous deamination of 5-methylcytosine¹

High C>T at CpC, CpC, TpC, T>A, and T>C is consistent with UV damage²

High C>A is consistent with smoking damage³

High C>T (site independent) is consistent with FFPE processing⁴







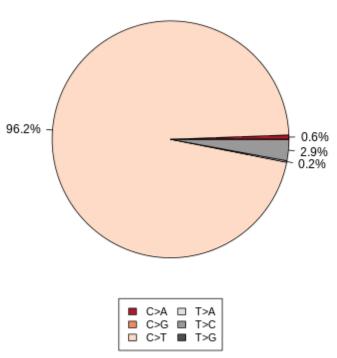
¹ Alexandrov LB et al. Nature. 2013; ² Hayward NK et al. Nature. 2017; ³ Alexandrov LB et al. Cancer Etiology. 2016; ⁴ Wong SQ et al. BMC Medical Genomics. 2014;

Real world hurts

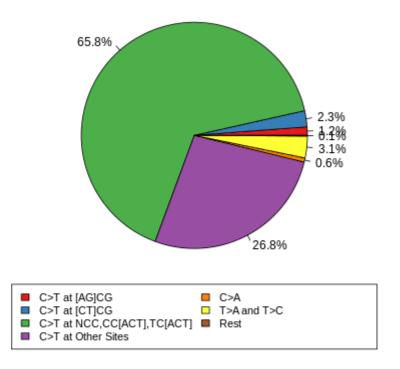
Mutation Load (Mutations/Mb): 1902.52



Somatic Mutations across Substitution Type



Signature Pattern of Somatic Mutations



Additional Information:

- High C>T at CpG is consistent with Spontaneous deamination of 5-methylcytosine¹
- High C>T at CpC, CpC, TpC, T>A, and T>C is consistent with UV damage²
- High C>A is consistent with smoking damage³
- High C>T (site independent) is consistent with FFPE processing⁴

¹Alexandrov LB et al. Nature. 2013; ²Hayward NK et al. Nature. 2017; ³Alexandrov LB et al. Cancer Etiology. 2016; ⁴Wong SQ et al. BMC Medical Genomics. 2014;

S07-9455 8-4 v1 c1476 2018-06-15-16-51-43-489 Mutation Load (Mutations/Mb): 11.04 Sample Results Sample QC Views: Substitution Type and Signature Pattern of Somatic Mutations ▼ Signature Pattern of Somatic Mutations Somatic Mutations across Substitution Type 55.6% 5.6% 77.8% 5.6% 5.6% 11.1% 16.7% 11.1% 7-9455 8-4_v1_c3271_2018-09-24-18-57-17-958 lutation Load (Mutations/Mb): 3.07 Sample QC ample Results Substitution Type and Signature Pattern of Somatic Mutations 💌 Somatic Mutations across Substitution Type Signature Pattern of Somatic Mutations 40% 40% 20% 20%

20%

40%

Workflow 10% standard

+ Spontaneous deamination of 5-methylecytosine

+ UV damage

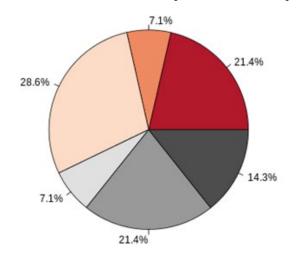
Tobacco/smoking damage

+ FFPE processing

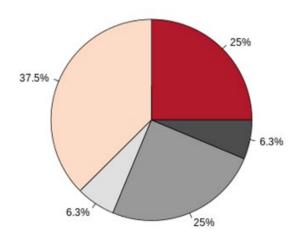
Workflow 10% modified

Formalin fixation

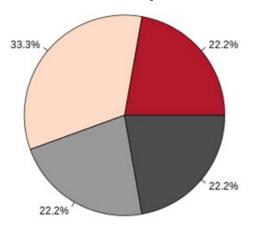
8 hrs MUTATION LOAD (Mutations/Mb): 8.54



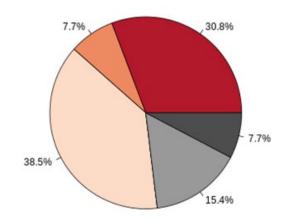
72 hrs MUTATION LOAD (Mutations/Mb): 9.78



24 hrs MUTATION LOAD (Mutations/Mb): 5.63



1 week MUTATION LOAD (Mutations/Mb): 7.94



Immunotherapy in breast cancer, take-home

- Immunotherapy is entering the clinical arena for BC patients
- TILs will be likely included among predictive/prognostic biomarkers
- Integrating biomarkers is warranted (GEP, TILs, TMB)
- Clinical practice: cancer centers with a NGS lab and large volumes vs, biotech companies
- Tissue workflow critical (IHC for diagnosis HRs, PDL-1, ISH, TMB)
- Multidisciplinary team mandatory (surgeons, oncologists, radiologists, pathologists, molecular biologists)