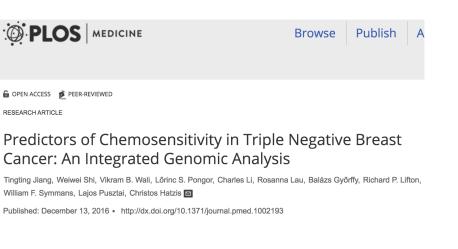
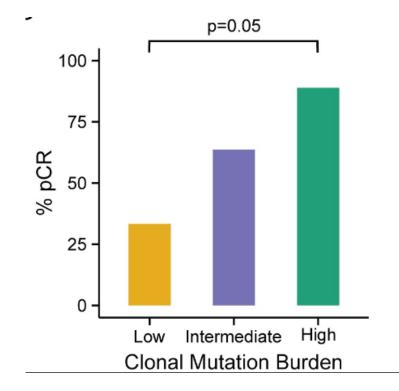
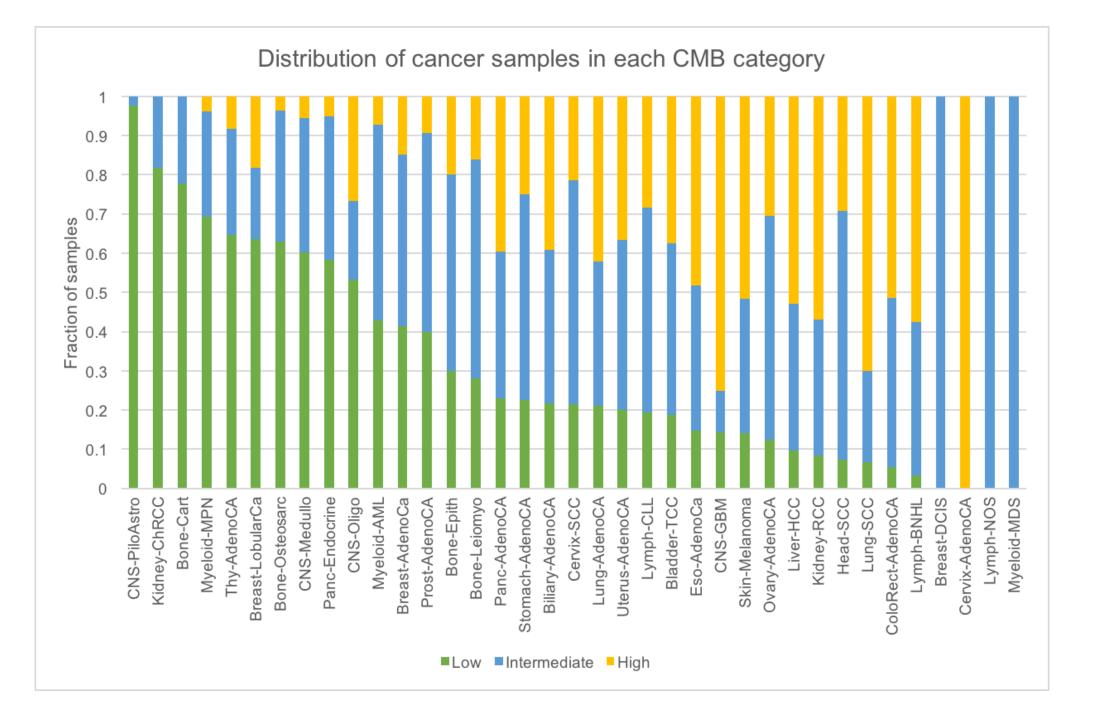
## Clonal Mutation Burden Analysis for PCAWG Somatic SNVs

Xiaotong Li

- Mutational Burden and Clonality as Predictors of Chemosensitivity in Triple Negative Breast Cancer
  - Methods
    - Combine mutation rate (MR) and mutant allele tumor heterogeneity (MATH) into a composite score, as Clonal Mutation Burden (CMB)
    - Captures both the clonality of a tumor and the number of somatic mutations per clone.
    - MR and MATH scores were mediandichotomized in the cohort, and CMB categories were defined as low (low MR, high MATH), high (high MR, low MATH), or intermediate (all others).
  - Tumors with high CMB appear to be extremely chemosensitive.



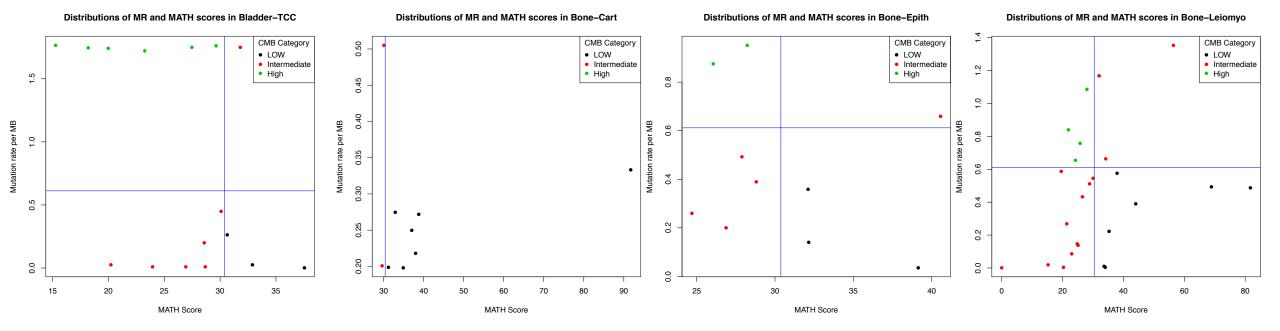




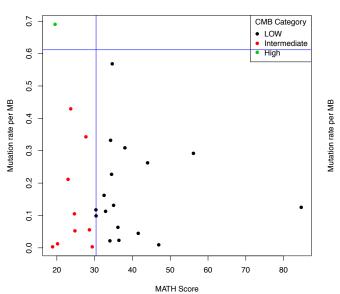
CMB Category • LOW • • • Intermediate • High ۲ 1.5 • Mutation rate per MB • • 1.0 ۲ 0.5 0.0 • 20 30 50 40 60

Distributions of MR and MATH scores in Biliary–AdenoCA

MATH Score



Distributions of MR and MATH scores in Bone-Osteosarc



Distributions of MR and MATH scores in Breast-AdenoCa

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1.5

1.0

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0.0

0

10

20

30

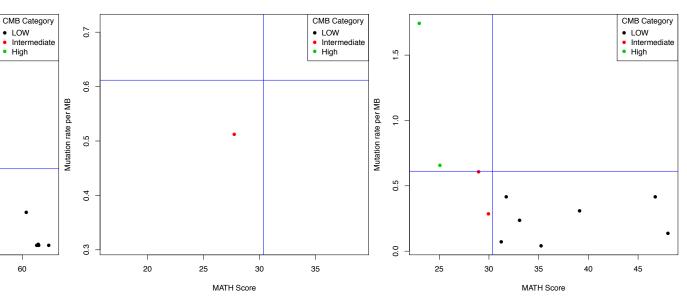
MATH Score

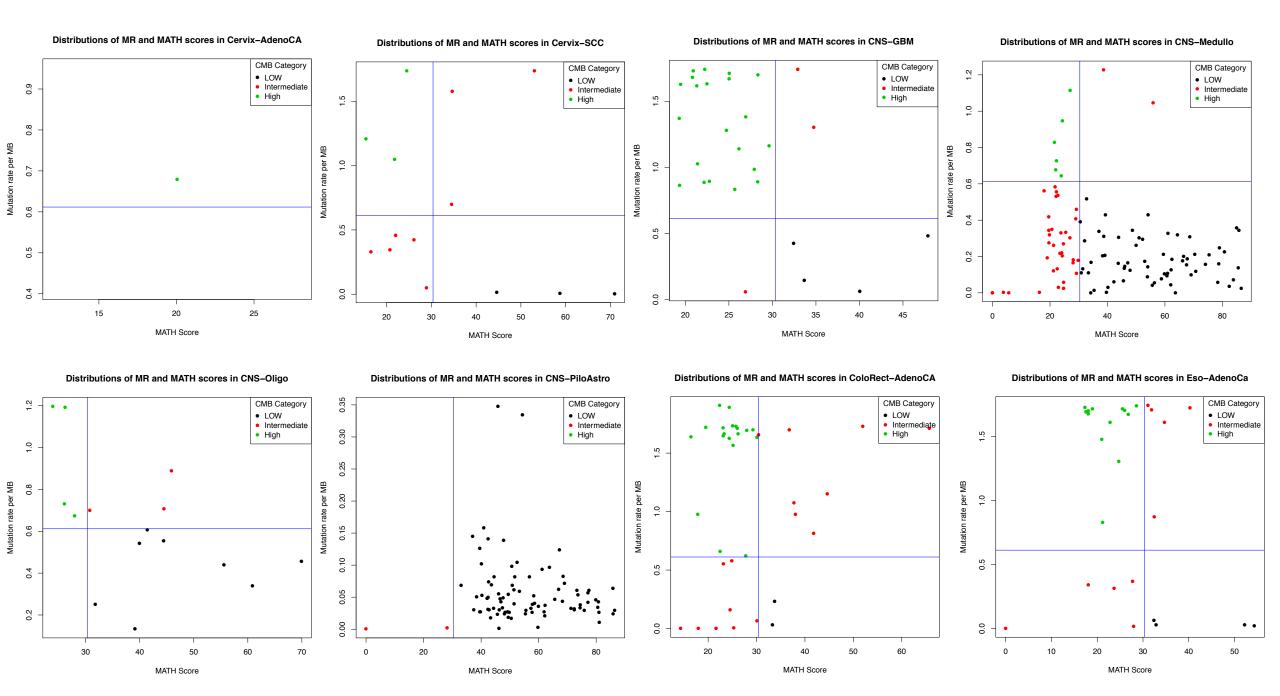
40

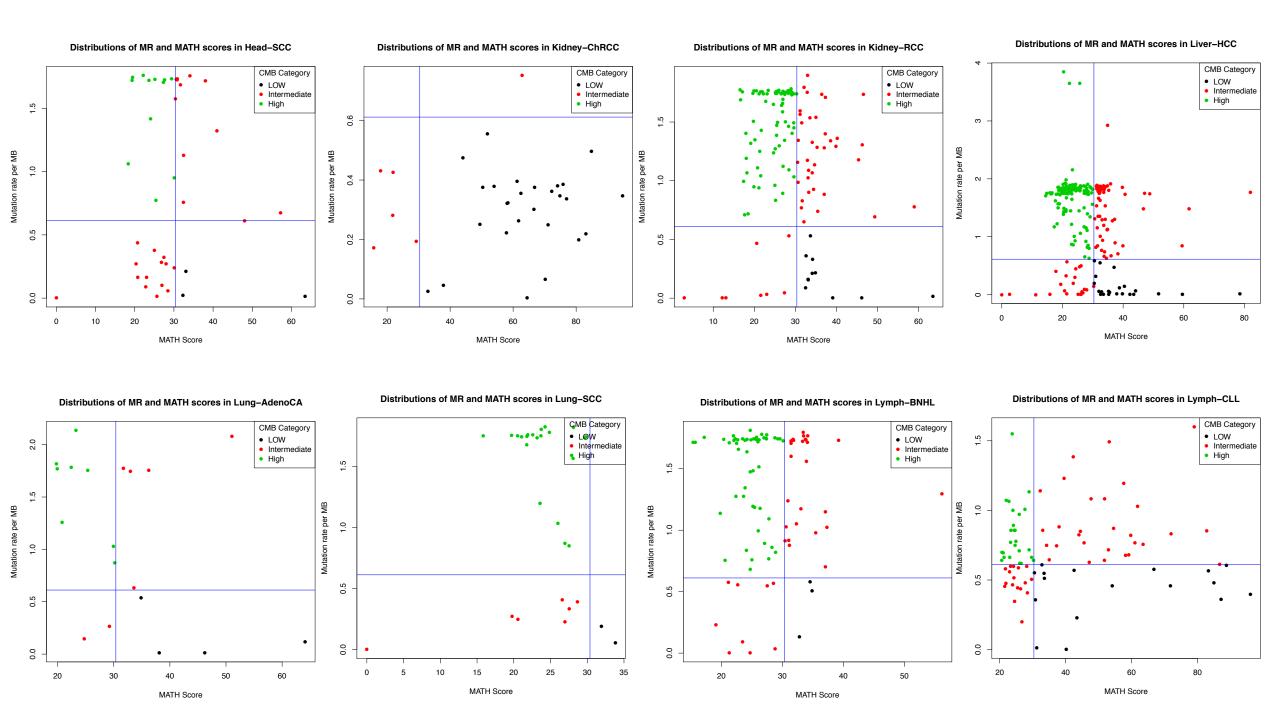
50

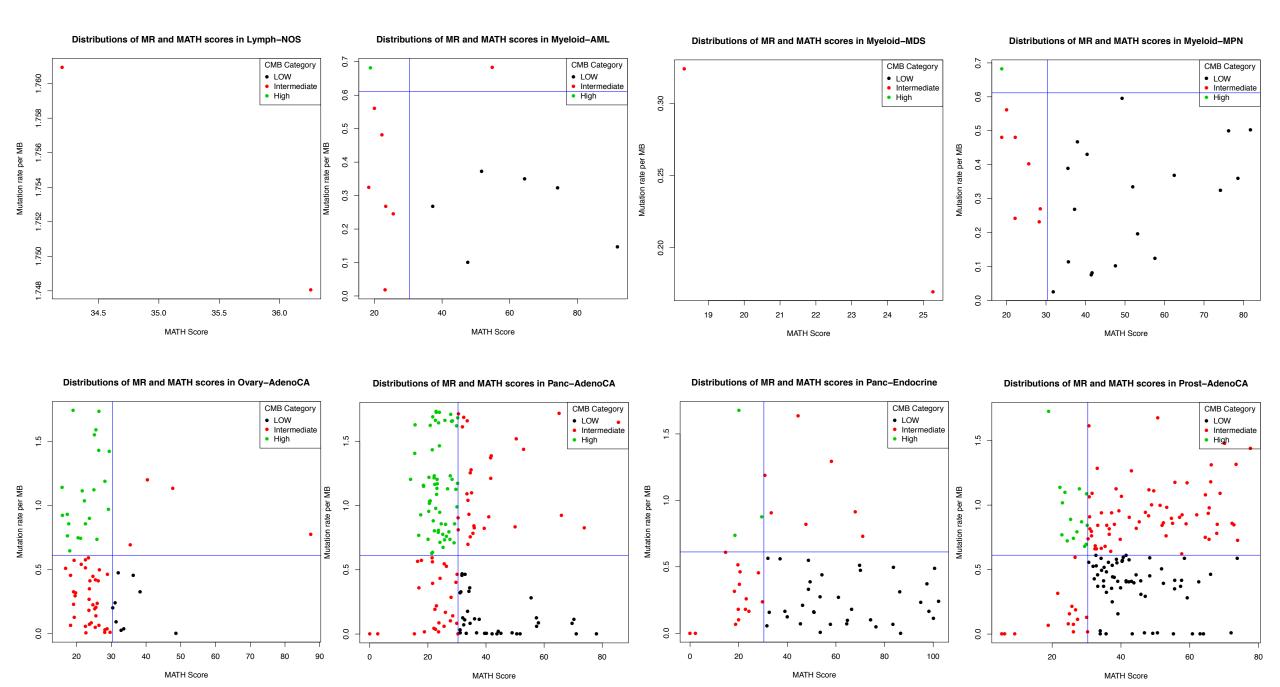
Distributions of MR and MATH scores in Breast-DCIS

## Distributions of MR and MATH scores in Breast-LobularCa

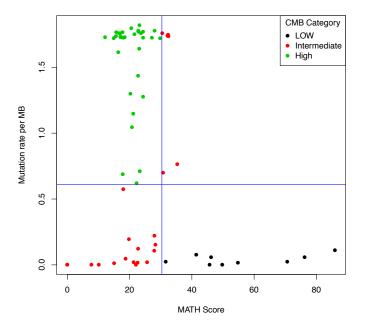




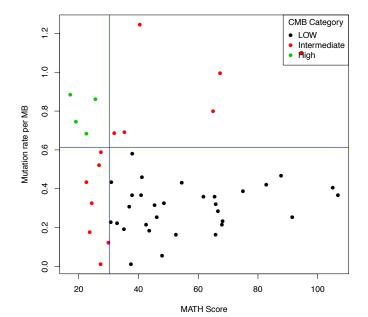




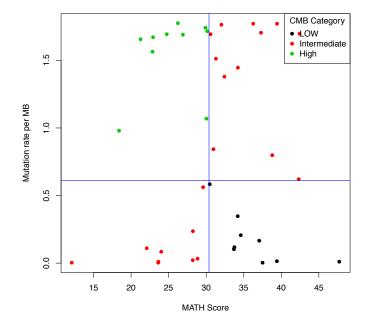
Distributions of MR and MATH scores in Skin–Melanoma



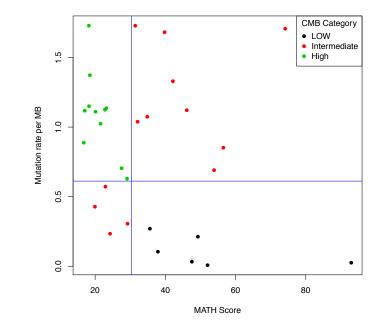
Distributions of MR and MATH scores in Thy-AdenoCA



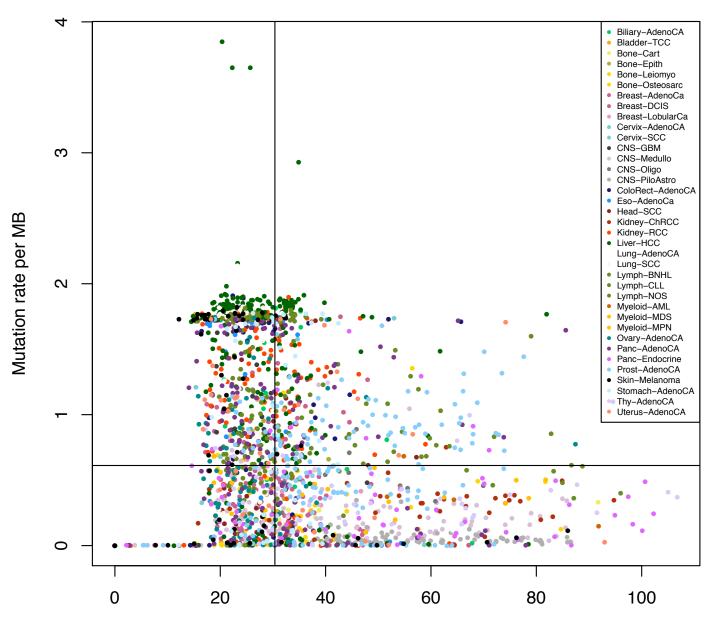
Distributions of MR and MATH scores in Stomach-AdenoCA



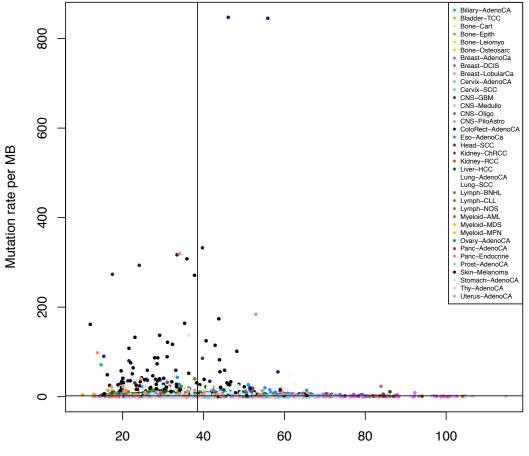
Distributions of MR and MATH scores in Uterus-AdenoCA



## **Distributions of MR and MATH scores in all samples**

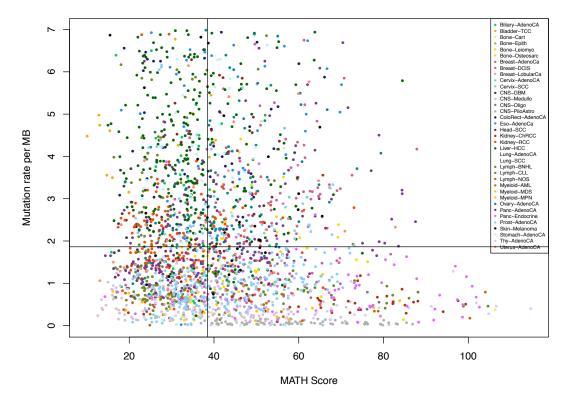


MATH Score



Distributions of MR and MATH scores in all samples

Distributions of MR and MATH scores in selected samples (MR per MB < 7)



MATH Score