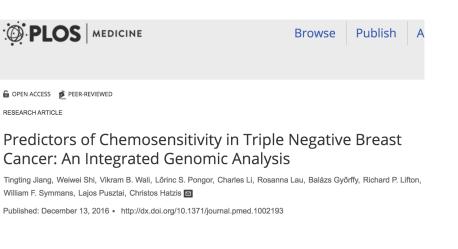
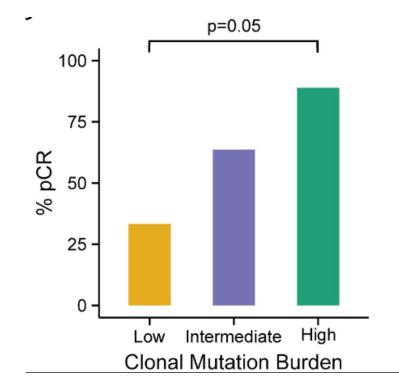
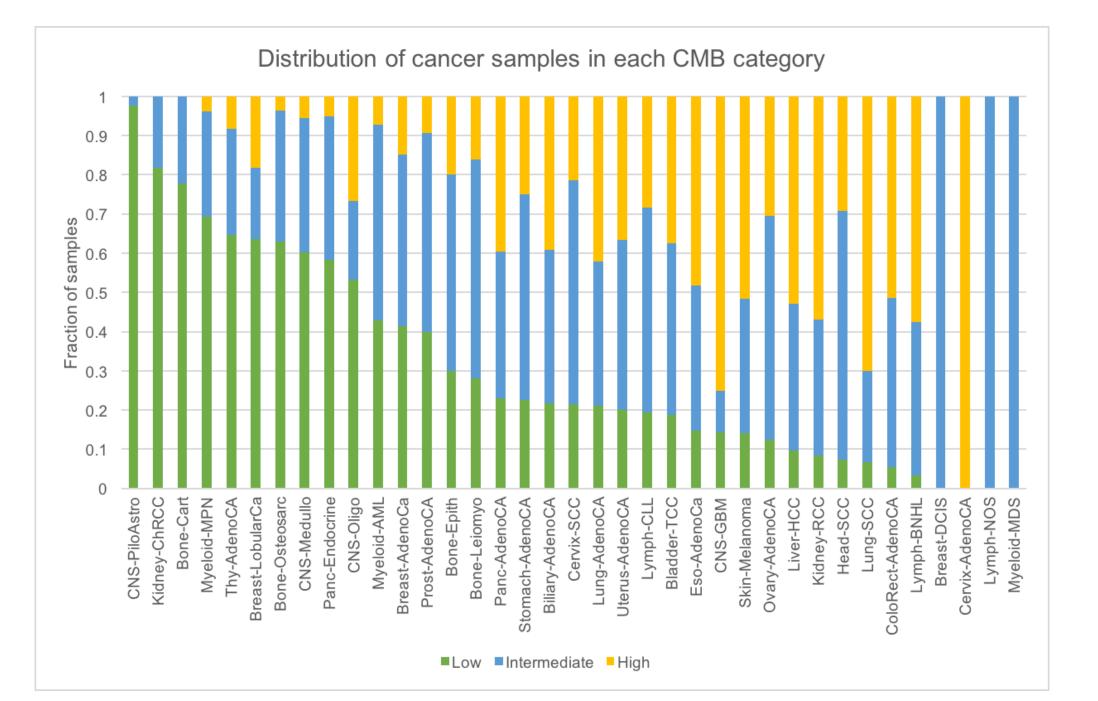
Clonal Mutation Burden Analysis for PCAWG Somatic SNVs

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.



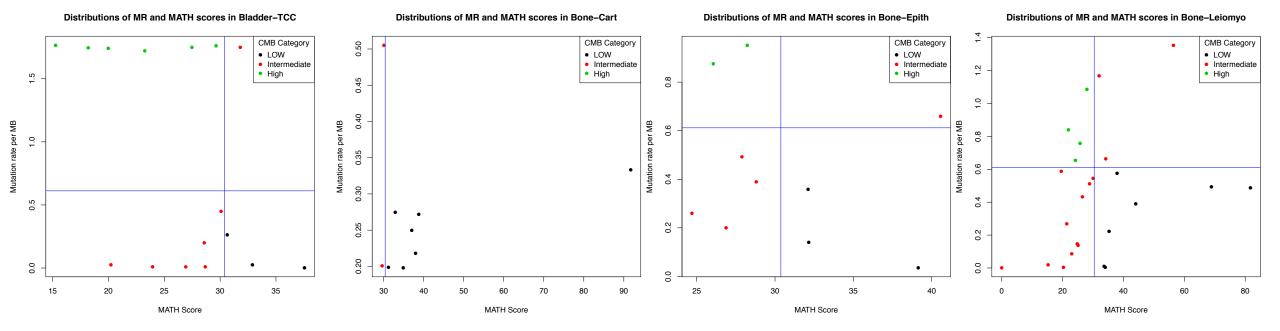




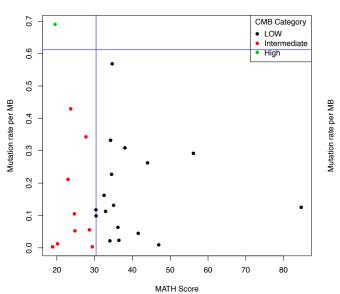
CMB Category • LOW • • • Intermediate • High ۲ 1.5 • Mutation rate per MB • • 1.0 ۲ 0.5 0.0 • 20 30 50 40 60

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

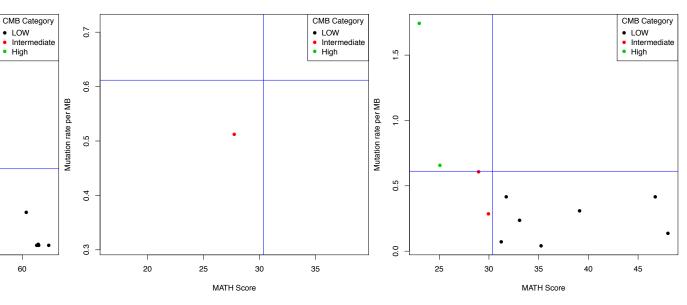
MATH Score

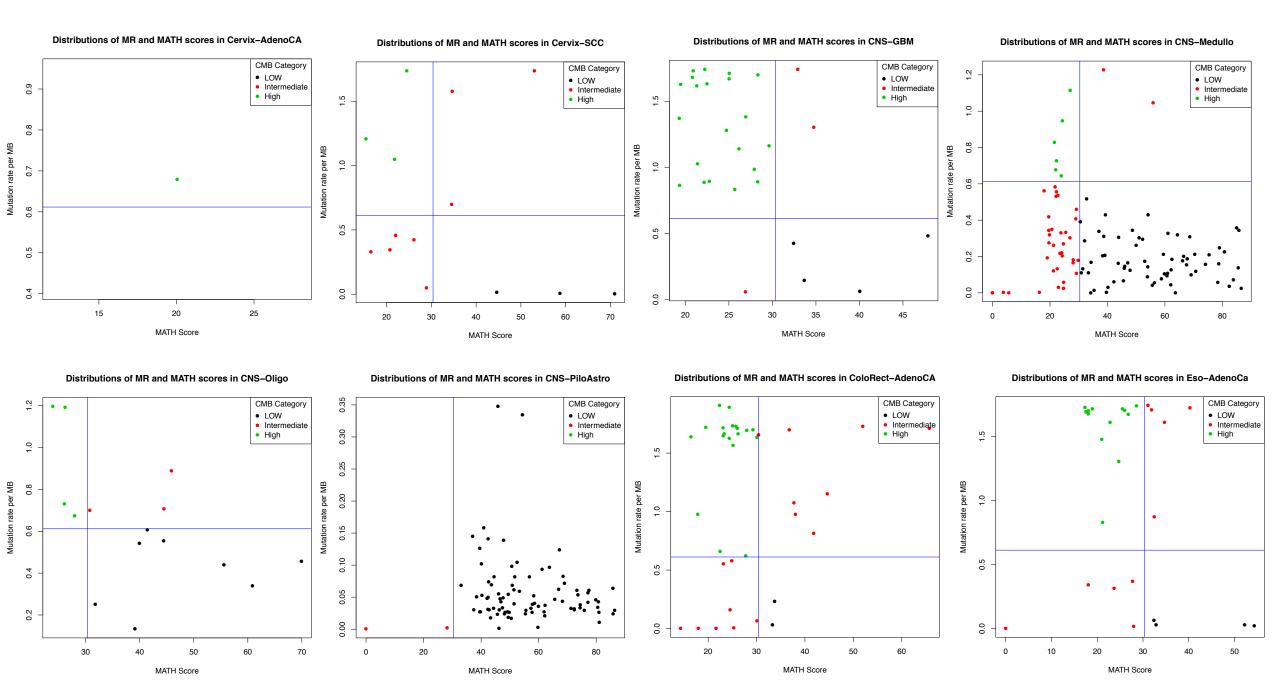
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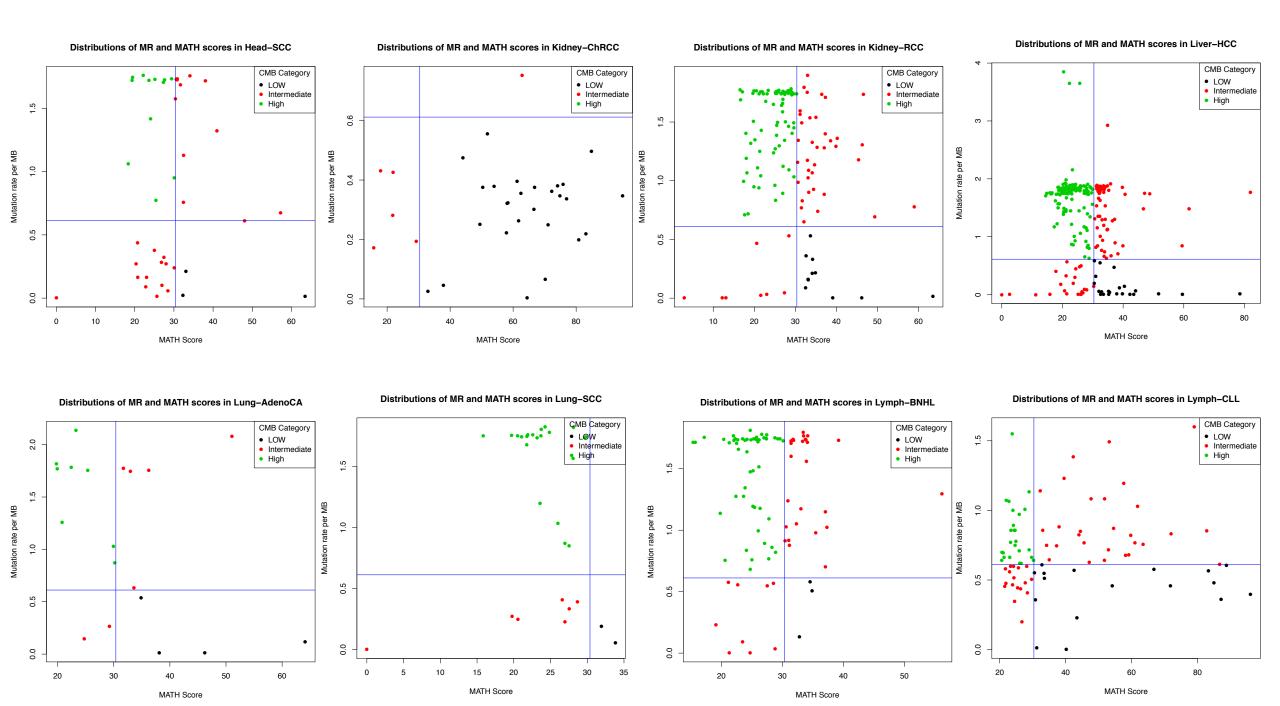
50

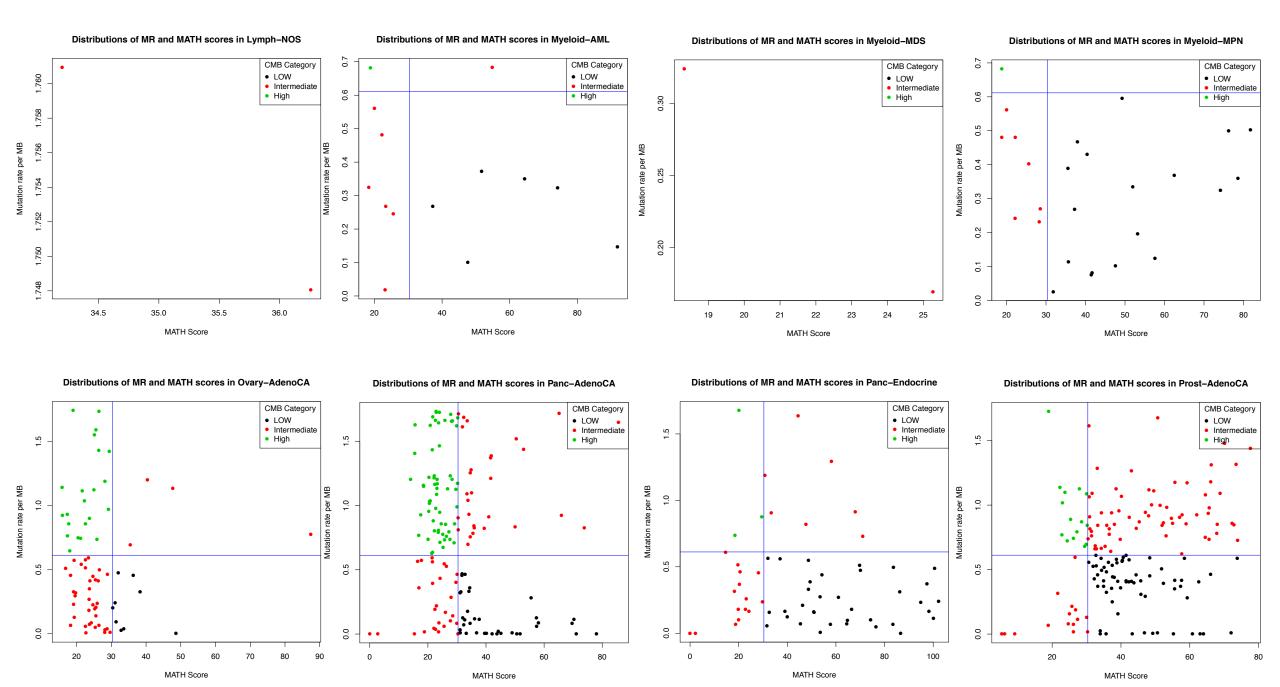
Distributions of MR and MATH scores in Breast-DCIS

Distributions of MR and MATH scores in Breast-LobularCa

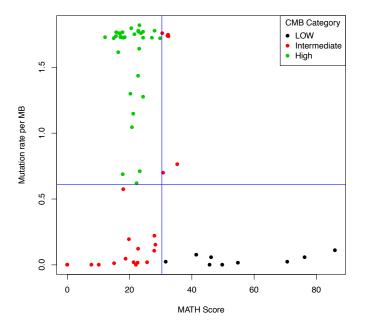




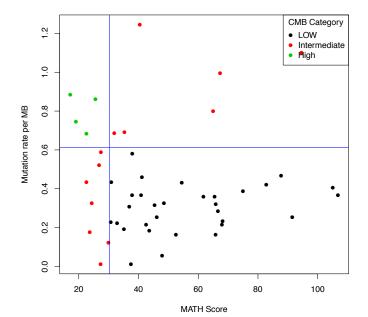




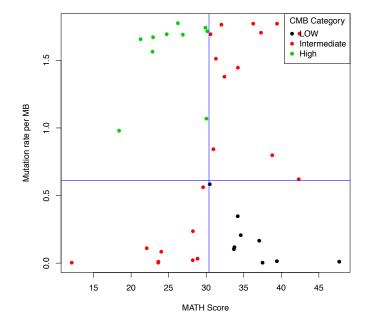
Distributions of MR and MATH scores in Skin–Melanoma



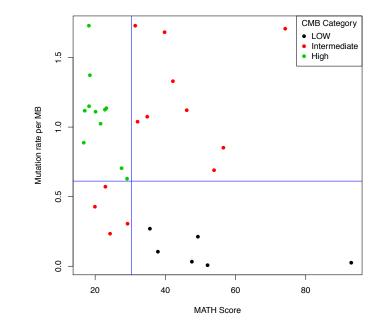
Distributions of MR and MATH scores in Thy-AdenoCA



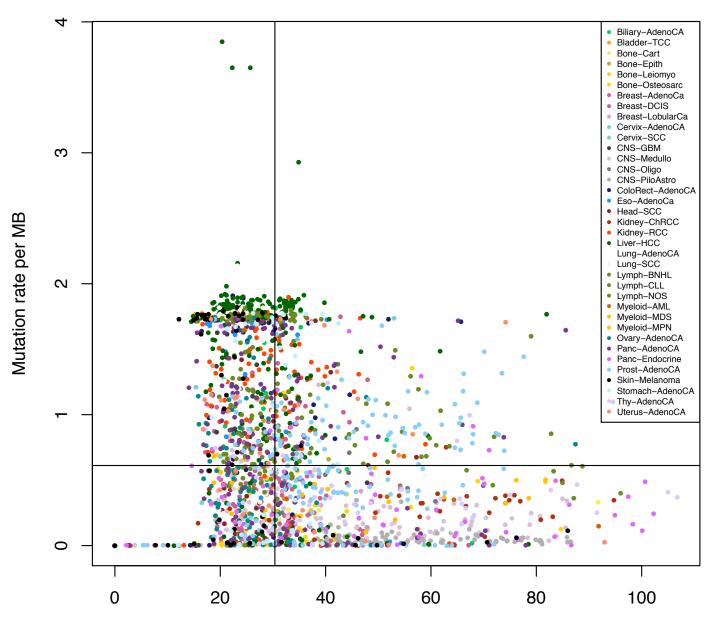
Distributions of MR and MATH scores in Stomach-AdenoCA



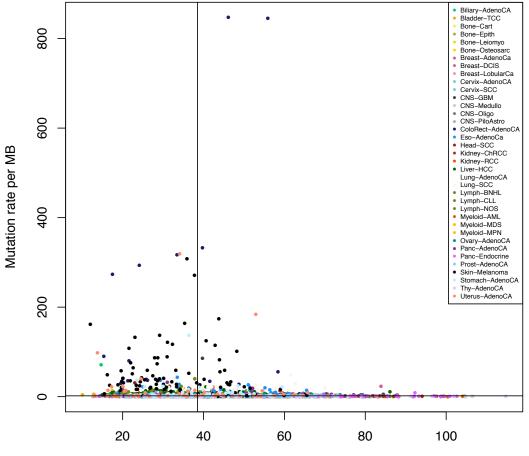
Distributions of MR and MATH scores in Uterus-AdenoCA



Distributions of MR and MATH scores in all samples

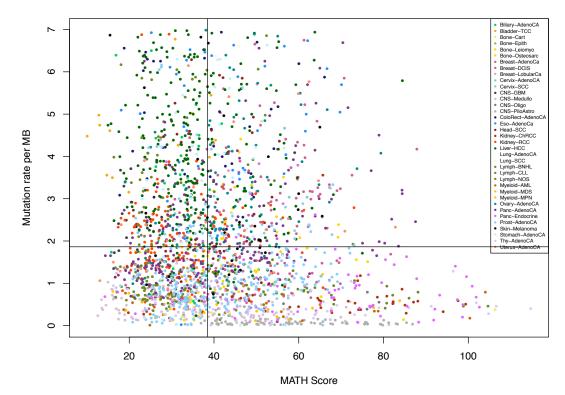


MATH Score



Distributions of MR and MATH scores in all samples

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



MATH Score