

# Distinct genomic landscapes characterize mismatch-repair deficiency(dMMR)/microsatellite instability-high (MSI-H) gastrointestinal (GI) cancers stratified by tumor mutation burden (TMB)

Jingyuan Wang<sup>1,2,3</sup>, Joanne Xiu<sup>4</sup>, Alex P. Farrell<sup>4</sup>, Francesca Battaglin<sup>1</sup>, Hiroyuki Arai<sup>1</sup>, Joshua Millstein<sup>1</sup>, Shivani Soni<sup>1</sup>, Wu Zhang<sup>1</sup>, Anthony F. Shields<sup>5</sup>, Axel Grothey<sup>6</sup>, Benjamin A. Weinberg<sup>7</sup>, John L. Marshall<sup>7</sup>, Emil Lou<sup>8</sup>, Moh'd Khushman<sup>9</sup>, Davendra P.S. Sohal<sup>10</sup>, Michael J. Hall<sup>11</sup>, Matthew Oberley<sup>4</sup>, David Spetzler<sup>4</sup>, and Heinz-Josef Lenz<sup>1</sup>.

<sup>1</sup>Division of Medical Oncology, Norris Comprehensive Cancer Center, Keck School of Medicine, University of Southern California, Los Angeles, California, USA. <sup>2</sup>Department of Medical Oncology, Cancer Center, Zhongshan Hospital, Fudan University, Shanghai, China. <sup>3</sup>Department of Gastrointestinal Oncology, Peking University Cancer Hospital and Institute, Beijing, China. <sup>4</sup>Caris Life Sciences, Phoenix, Arizona, USA. <sup>5</sup>Department of Oncology, Karmanos Cancer Institute, Wayne State University, Detroit, Michigan, USA. <sup>6</sup>West Cancer Center and Research Institute, Germantown, TN, USA. <sup>7</sup>Ruesch Center for The Cure of Gastrointestinal Cancers, Lombardi Comprehensive Cancer Center, Georgetown University Medical Center, Washington, D.C., USA. <sup>8</sup>University of Minnesota, Minneapolis, Minnesota, US. <sup>9</sup> Division of hematology and oncology, St. Louis/Siteman Cancer Center, Washington University. <sup>10</sup> University of Cincinnati, Ohio, USA. <sup>11</sup>Department of Clinical Genetics, Fox Chase Cancer Center, 333 Cottman Avenue, Philadelphia, PA, USA.

Abstract ID: 2618

## Background

- TMB-H was reported to be predictive of response to immune checkpoint inhibitors[1-2].
- However, genomic signatures contributing to TMB-H independent from dMMR/MSI-H status are not well-studied.
- We aimed to characterize specific molecular features of a large cohort of MSS GI tumors with TMB-H.

## Methods

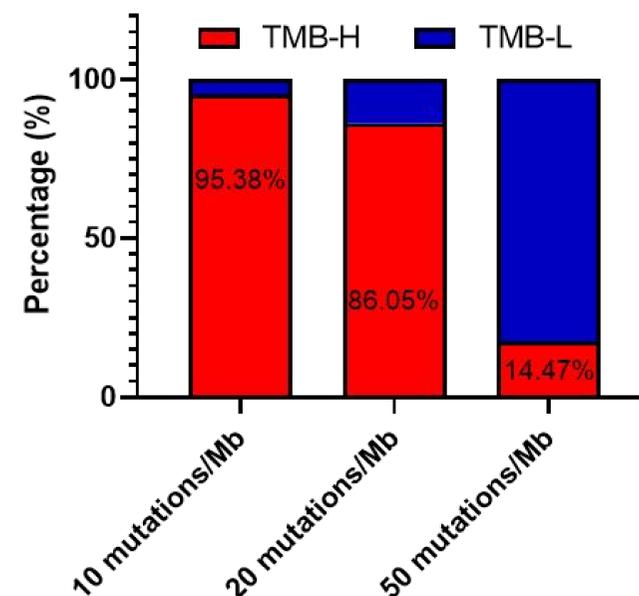
- NGS was performed on genomic DNA isolated from formalin-fixed paraffin-embedded tumor samples using the NextSeq or NovaSeq 6000 platforms (Illumina, Inc., San Diego, CA). All variants were detected with greater than 99% confidence based on allele frequency and amplicon coverage, with an average sequencing depth of coverage of greater than 500 and an analytic sensitivity of 5%[3].
- Microsatellite instability (MSI)/ MMR status was determined by a combination of NGS (>=46 loci), IHC and fragment analysis.
- TMB-H were defined using differing TMB cutoffs (10, 20, 50 mutations/Mb), according to the standard algorithm by Friends of Cancer Research TMB Harmonization Project[4].
- Molecular features were compared in four groups (TMB<10 vs 10-20 vs 20-50 vs ≥50mutations/Mb) using Fisher-Exact or Chi-square and adjusted for multiple comparison by Benjamini-Hochberg.
- Significance was determined by  $q < .05$ .

### Reference:

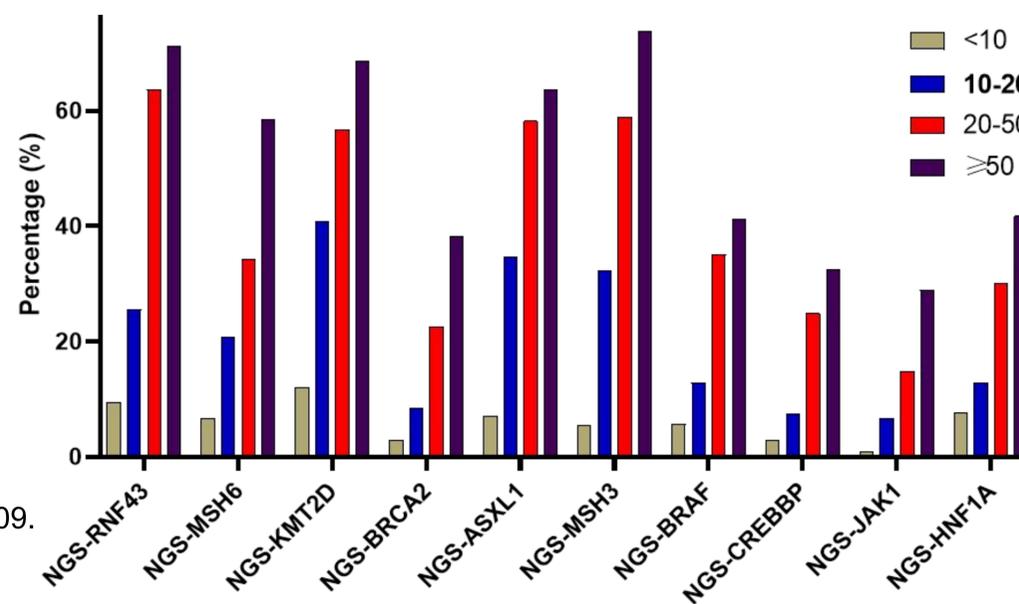
- Ann Oncol. 2019 Jul 1;30(7):1096-1103.
- Oncologist. 2020 Sep;25(9):803-809.
- Lancet Oncol. 2023 Feb;24(2):151-161.
- J Immunother Cancer 2020; 8.

## Results

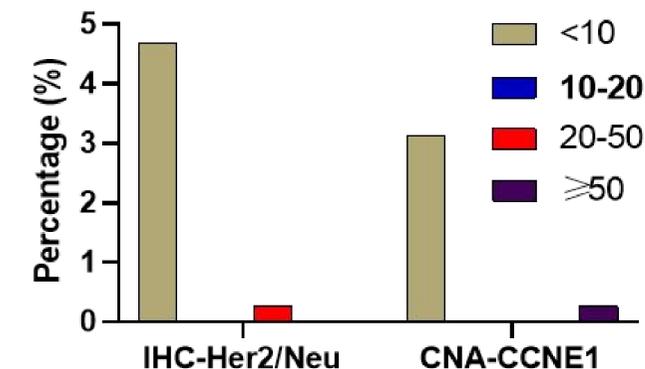
**Fig 1.** Tumors with TMB over 10, 20, 50 mutations/Mb were observed in 95.38%, 86.05% and 14.47% respectively in the dMMR/MSI-H GI cohort (n=2272).



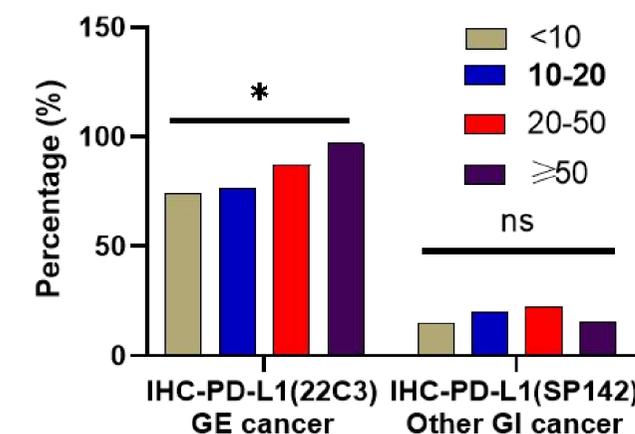
**Fig 2.** Distinct mutational landscapes according to different TMB levels ( all  $P_{adj} < 0.0001$ ).



**Fig 3.** The rates of CNNE1 amplification and HER2 overexpression were the highest in tumors with TMB below 10 mutations/Mb (all  $P_{adj} < 0.05$ ).



**Fig 4.** The association between PD-L1 positivity and TMB levels was only observed in gastroesophageal cancers.



## Conclusion

This is the largest study to investigate the distinct molecular landscape of dMMR/MSI-H GI cancers with different degrees of TMB. These data may inform our understanding of the efficacy of ICB in dMMR/MSI-H GI tumors.