Emerging Role of Immunotherapy in Gastrointestinal Cancers: Mutation and Neoantigen Burden as Predictors of Response

#### MSI mCRC

- 38 year old woman presented with 2 months of diarrhea, was also found to be anemic by her PCP
- Colonoscopy showed a right-sided mass in the colon
- R colectomy
  - 6x5x3 cm cecal poorly differentiated adenocarcinoma with lymphocytic infiltrate, T3NoMo, o/18 LN positive, MSI, RAS and BRAF WT
  - Lynch positive
- No adjuvant therapy given
- Two years later CT scan showed 2 liver masses, 5.9 x 5.4 cm and 3.3 x 3.3 cm, and abdominal lymphadenopathy

#### MSI mCRC

- Biopsy of liver mass confirmed recurrent adenocarcinoma
- Initiated nivolumab
- Partial response seen on first restaging scan (2 months)
- Continues in PR now 2 years
  - Course has been complicated by hypothyroidism

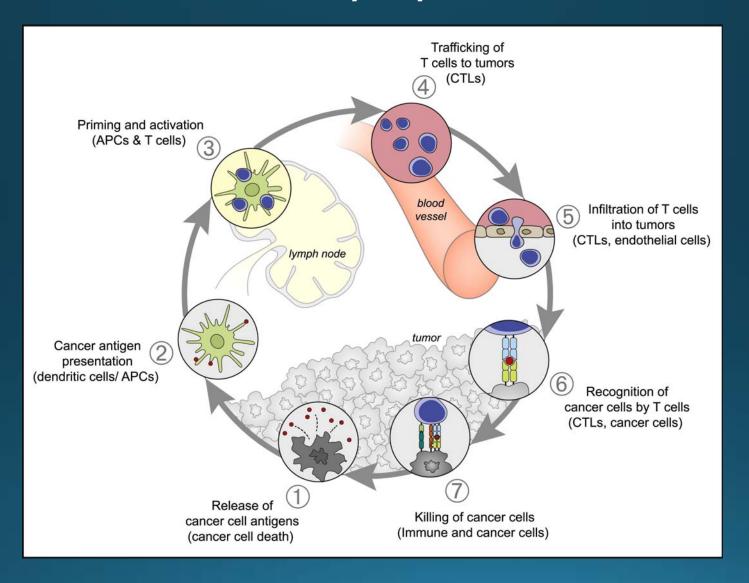
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#### Disclosures

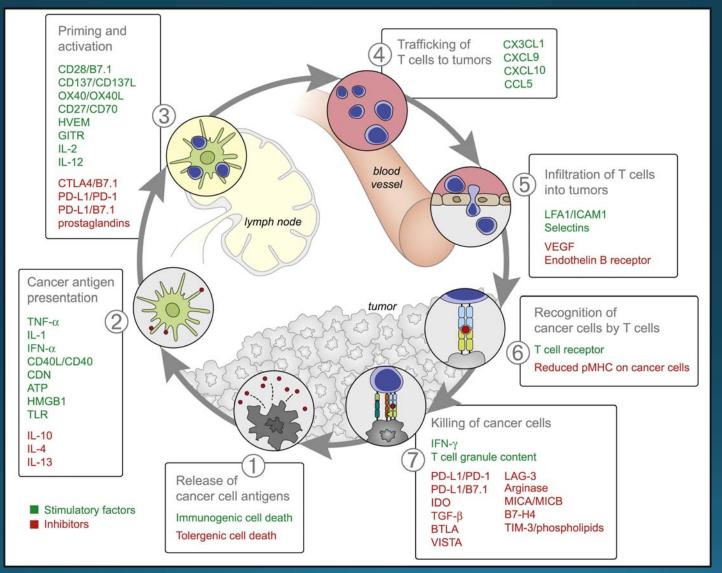
## Contracted Research

Abbott Laboratories, AbbVie Inc, Apexigen, AstraZeneca Pharmaceuticals LP, Bayer HealthCare Pharmaceuticals, Boehringer Ingelheim Pharmaceuticals Inc, Bristol-Myers Squibb Company, Celgene Corporation, Daiichi Sankyo Inc, Eisai Inc, EMD Serono Inc, Five Prime Therapeutics Inc, Forty Seven Inc, Genentech BioOncology, Gilead Sciences Inc, GlaxoSmithKline, Incyte Corporation, Kolltan Pharmaceuticals Inc, Leap Therapeutics Inc, Lilly, MacroGenics Inc, MedImmune Inc, Merck, Novartis Pharmaceuticals Corporation, OncoMed Pharmaceuticals Inc, Onyx Pharmaceuticals, an Amgen subsidiary, Pfizer Inc, Roche Laboratories Inc, Sanofi Genzyme, Stemcentrx, Taiho Oncology Inc, Takeda Oncology, TG Therapeutics Inc.

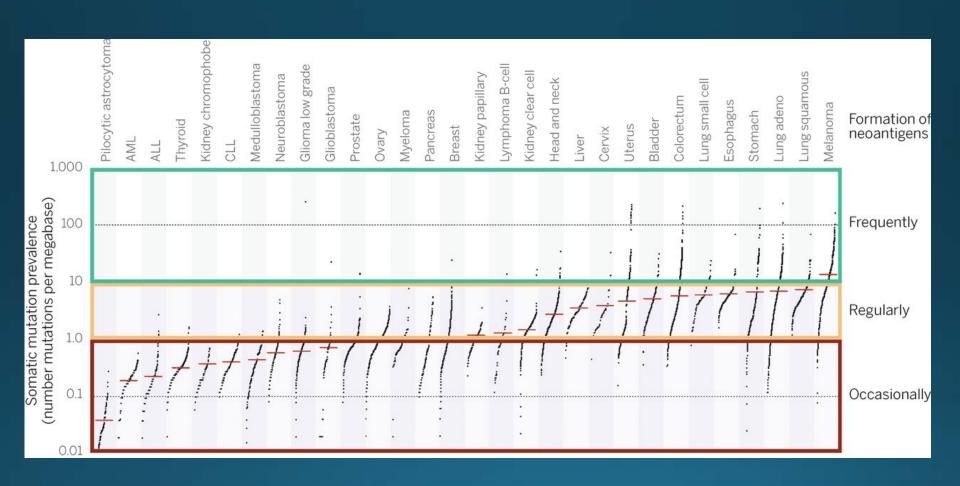
# Cancer-immunity cycle



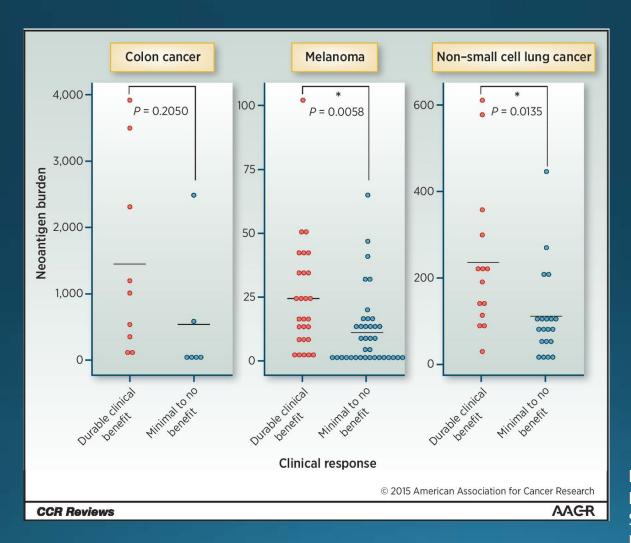
# Cancer-immunity cycle



# Mutational load/neoantigen formation

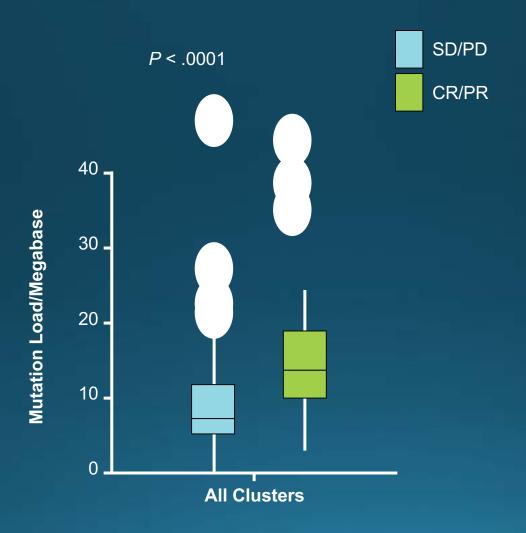


# Mutational burden and efficacy of checkpoint inhibitors

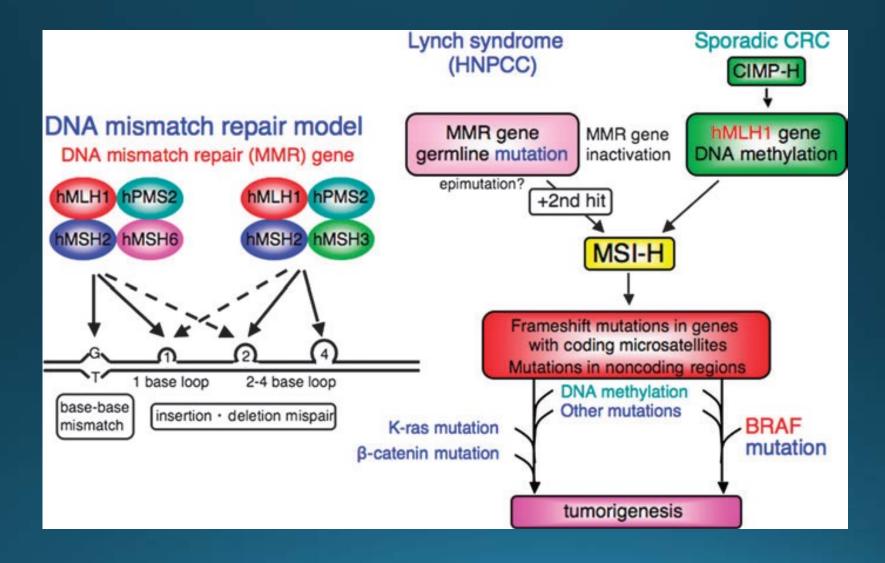


Desrichard CCR 2016 Le NEJM 2015 Snyder NEJM 2014 Rizvi Science 2015

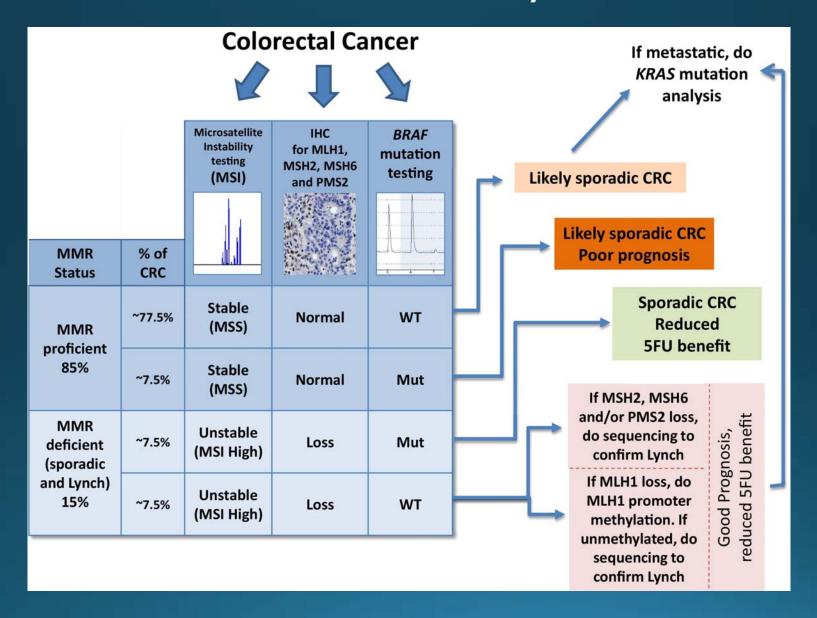
# IMvigor210: Mutation Load Associated With Objective Responses



# Microsatellite instability



# Microsatellite instability



#### IHC STAINS FOR MLH1, MSH2, MSH6 AND PMS2 PROTEINS

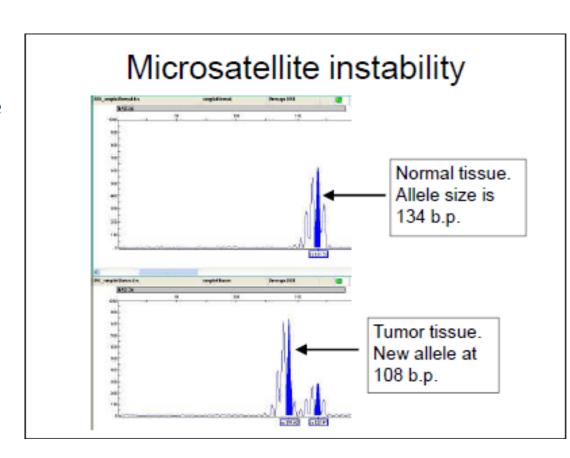
	Normal	Abnormal
MLH1		
MSH2	233	
MSH6		
PMS2		

- These proteins are located in the nucleus.
- It is normal for these proteins to be present, therefore: Loss of staining is <u>abnormal</u>.



#### **MSI TESTING BY PCR**

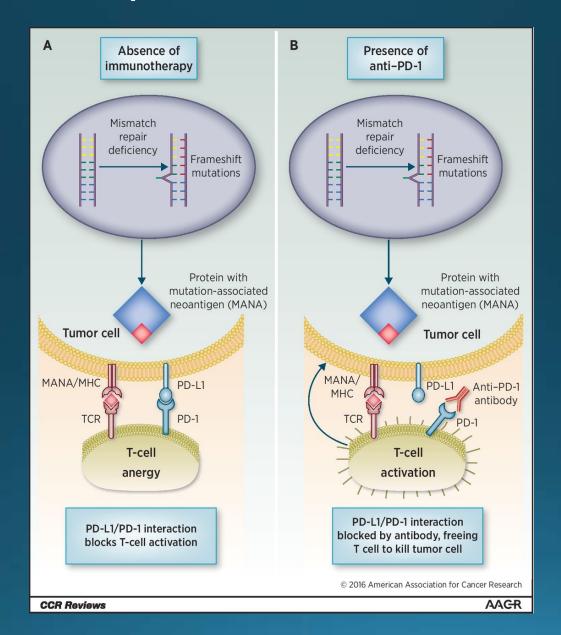
- A PCR test is used to look for a change in microsatellite size in the tumor compared to the patient's non-tumor tissue.
- Microsatellite repeats amplified from tumor and normal tissue.
- Amplified DNA separated by size using electrophoresis.
- Commercial kit available



UCSF Website: http://www.ucsfcme.com/2009/slides/MAP0901A/0GrenertMSIBasicsForPathologists%20.pdf

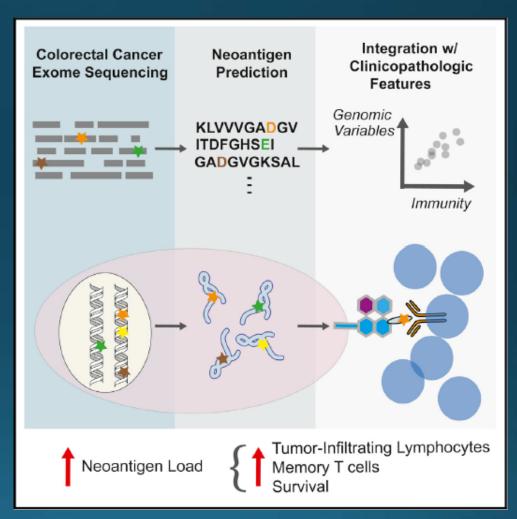


### Mismatch repair and mutations



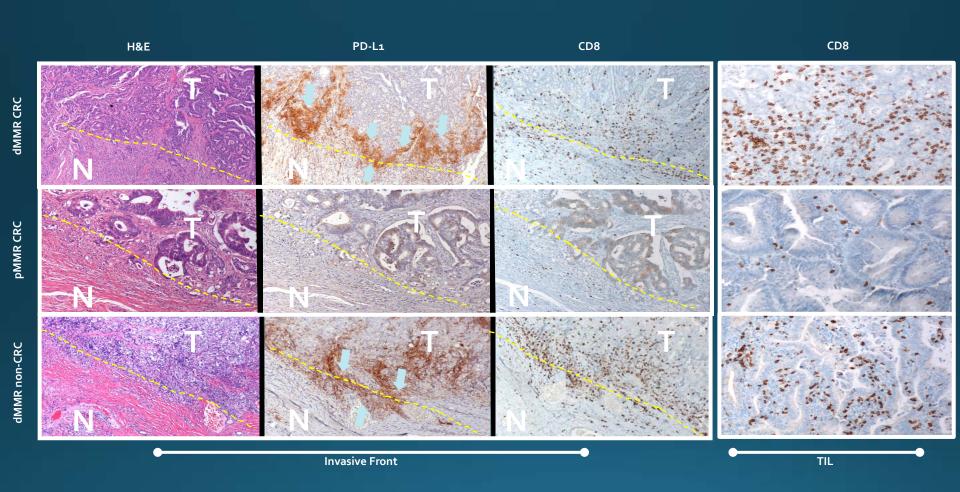
## Hypermutation and Immuno-Oncology

- In CRC, MSI-H is associated with increases in immune infiltration and expression of immune checkpoint regulators<sup>1,2</sup>
- MSI-H is also associated with increased number of mutations per tumor
- Tumor mutations produce tumor-specific neoantigens, which when expressed on the tumor cell surface, are a target for T cells
- May improve response to immunotherapy
- Elevated neoantigen load in CRC is associated with improved survival<sup>2</sup>



- 1. Llosa NJ, et al. *Cancer Discov*. 2015;5:43–51.
- 2. Giannakis M, et al. Cell Reports. 2016;15:857–865.

#### Baseline PD-L1 Expression and CD8 T Cell Infiltration



## MSI in more than CRC

Cancer Type	General population risk	Lynch syndrome (MLH1 and MSH2 heterozygotes)	
		Risk	Mean age of onset
Colon	5.5%	52-82%	44-61 years
Endometrium	2.7%	25-60%	48-62 years
Stomach	< 1%	6-13%	56 years
Ovary	1.6%	4-12%	42.5 years
Hepatobiliary tract	< 1%	1.4-4%%	Not reported
Urinary tract	< 1%	1-4%	~55 years
Small bowel	< 1%	3-6%	49 years
Brain/central nervous	< 1%	1-3%	~50 years
system Sebaceous neoplasms	< 1%	1-9%	Not reported

#### **Study Design**

#### **Colorectal Cancers**

**Non-Colorectal Cancers** 

Cohort A

Deficient in

Mismatch Repair

(n=25)

Cohort B
Proficient in
Mismatch Repair
(n=25)

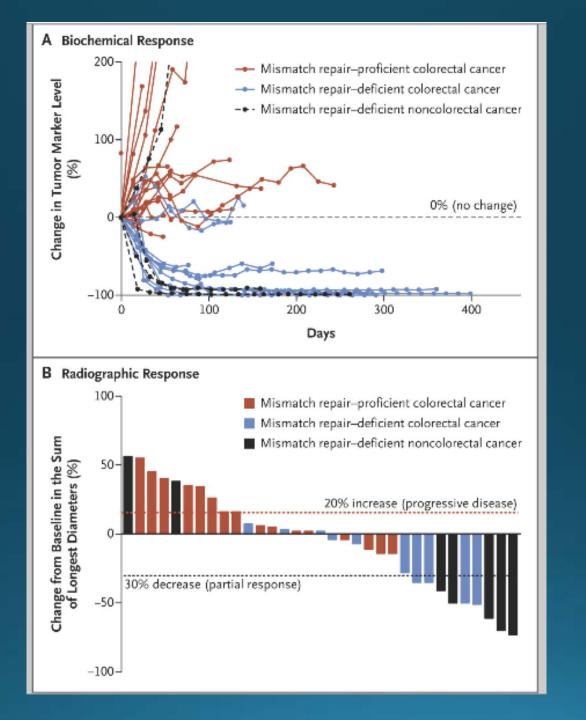
Cohort C

Deficient in

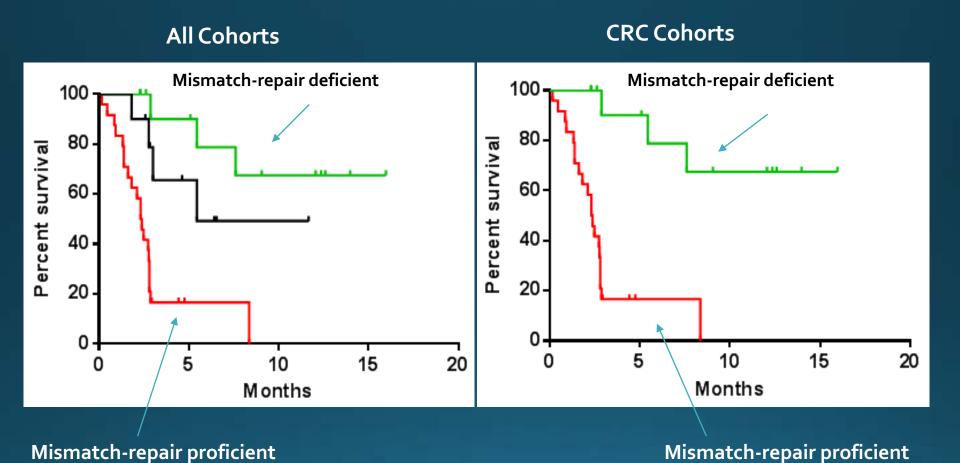
Mismatch Repair

(n=21)

- Anti-PD1 (Pembrolizumab) 10 mg/kg every 2 weeks
- Primary endpoint: immune-related 20-week PFS rate and response rate
- Mismatch repair testing using standard PCR-based test for detection of microsatellite instability



#### **Progression-Free Survival**



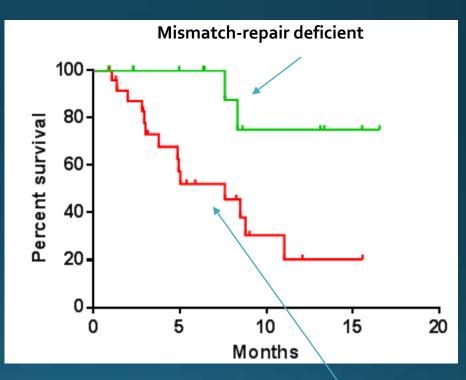
Le ASCO 2015

#### **Overall Survival**

#### **All Cohorts**

# Mismatch-repair deficient 100 80 60 40 20 Months

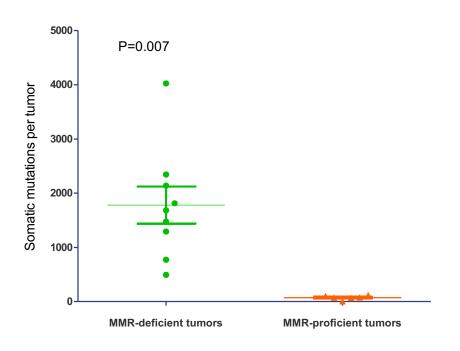
#### **CRC Cohorts**

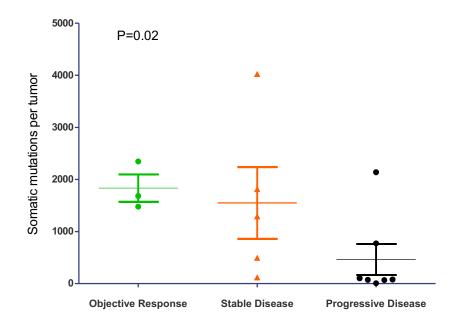


Mismatch-repair proficient

Mismatch-repair proficient

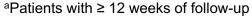
#### Mutation Burden is Associated with Efficacy





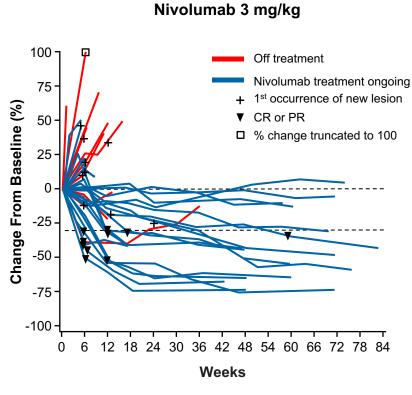
# Investigator-Assessed Best Overall Response in Patients With MSI-H Receiving Nivolumab Monotherapy

	Nivolumab 3 mg/kg (n = 47) <sup>a</sup>
ORR, n (%) (95% exact CI)	12 (25.5) (15.4, 38.1)
Complete response	0
Partial response	12 (25.5)
Stable disease	14 (29.8)
Progressive disease	17 (36.2)
Unable to determine	4 (8.5)
Median time to response, mo (range)	2.12 (1.3–13.6)
Median duration of response, mo (range)	NE (0.0 <sup>b</sup> –15.2 <sup>b</sup> )



blncludes censored observations

CR = complete response; NE = not estimable; PR = partial response



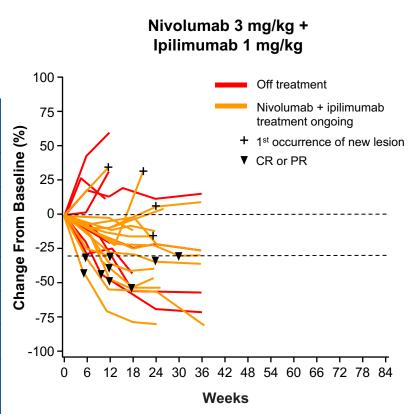
Horizontal reference line indicates 30% reduction

# Investigator-Assessed Best Overall Response in Patients With MSI-H Receiving Nivolumab + Ipilimumab

	Nivolumab 3 mg/kg + lpilimumab 1 mg/kg (n = 27) <sup>a</sup>
ORR, n (%) (95% exact CI)	9 (33.3) (18.6, 50.9)
Complete response	0
Partial response	9 (33.3)
Stable disease	14 (51.9)
Progressive disease	3 (11.1)
Unable to determine	0
Median time to response, mo (range)	2.73 (1.2–6.9)
Median duration of response, mo (range)	NE (NE-NE)

<sup>&</sup>lt;sup>a</sup>Patients with ≥ 12 weeks of follow-up

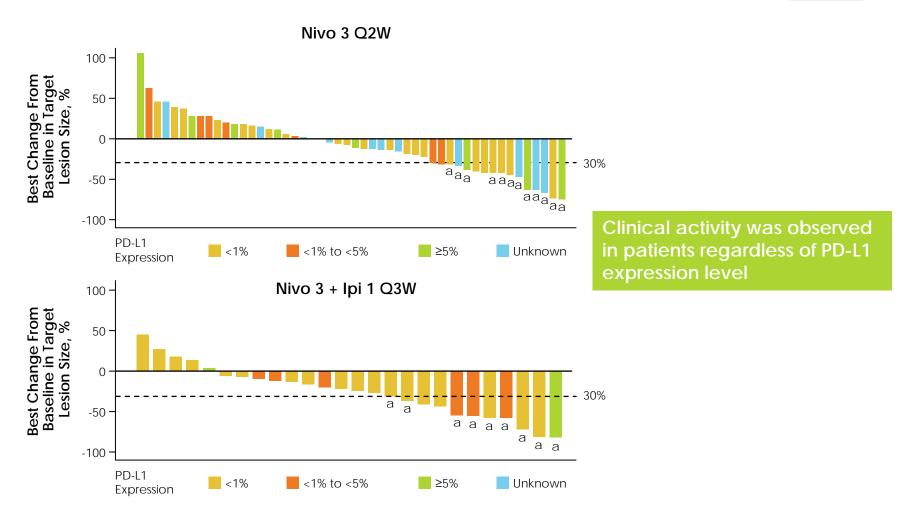
CR = complete response; NE = not estimable; PR = partial response



Horizontal reference line indicates 30% reduction

bIncludes censored observations

#### Nivolumab ± Ipilimumab: CheckMate 142 Response by pd-I1 Status



Overman M et al. Poster presentation at ESMO 2016. 479P. Database lock: April 2016.

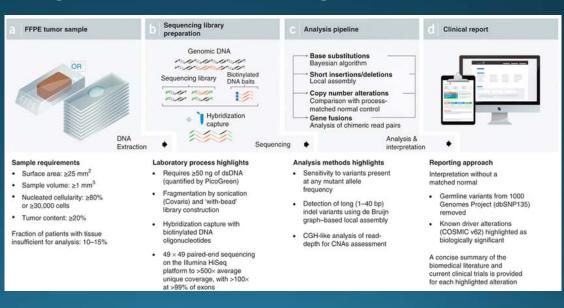
#### Testing of tumor mutational burden (TMB)

- Next generation sequencing
- Number of single nucleotide somatic mutations is counted
- This number is divided by the total number of megabases to calculate the average number of mutations/megabase (Mb) which is extrapolated to the whole genome

• In general >10 mutations/Mb designates frequent neoantigen

formation

- FMI and Caris
  - TMB high >20 mutations/Mb
  - TMB intermediate = 6-19 mutations/Mb
  - TMB low <u>< 5</u> mutations/Mb



Emerging Role of Immunotherapy in Gastrointestinal Cancers: Mutation and Neoantigen Burden as Predictors of Response

# HER2 negative gastric cancer

- 45 year old man with a history of HTN, type I DM, hypothyroidism, and allergic rhinitis presented with a 4 month history of severe reflux refractory to PPI
- Endoscopy performed showing a GE junctional mass from the distal esophagus to proximal stomach. Biopsy showed moderately differentiated adenocarcinoma, HER2 negative.
- PET scan showed hypermetabolic uptake in the gastric cardia mass and periceliac nodes
- He received neoadjuvant chemoradiation with 5FU and oxaliplatin
- He underwent resection with pathology showing minimal residual disease, but 4/21 LN were positive

# HER2 negative gastric cancer

- 4 months after surgery surveillance PET scan showed 2 new liver lesions
- He was started on docetaxel but had progressive disease in 3 months
- He was then started on clinical trial with pembrolizumab
- He had an almost partial response and remains on therapy
   14 months into treatment