

An odyssey and a journey through digestive oncology: Immunotherapy in digestive cancers

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Disclosure information

- Participation to advisory boards for: Array, Astrazeneca, Bayer, Biocartis, Bristol-Myers Squibb, Celgene, Daiichi, Pierre-Fabre, Incyte, Ipsen, Lilly, Merck Sharp & Dohme, Merck KGaA, Novartis, Roche, Servier, Sirtex, Taiho
- Research grants from: Bayer, Boehringer Ingelheim, Celgene, Ipsen, Lilly, Roche, Merck Sharp & Dohme, Merck KGaA, Novartis, Roche, Servier paid to the institution.
- Receives an honorarium for speaking as this symposium is supported from Bristol Myers Squibb
- New data are discussed on drugs, including data for which an approval is not yet granted



Colorectal Cancer Subtypes Microenviroment targeting in mCRC

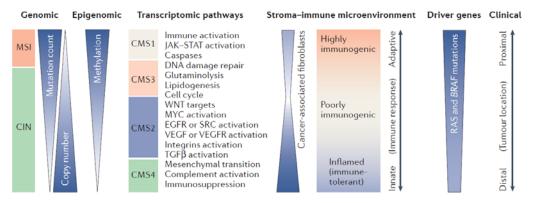
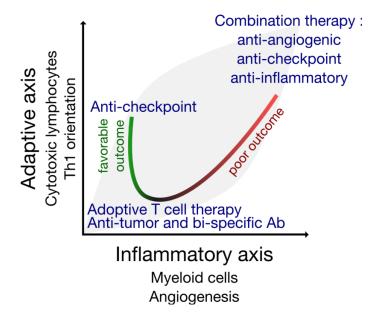


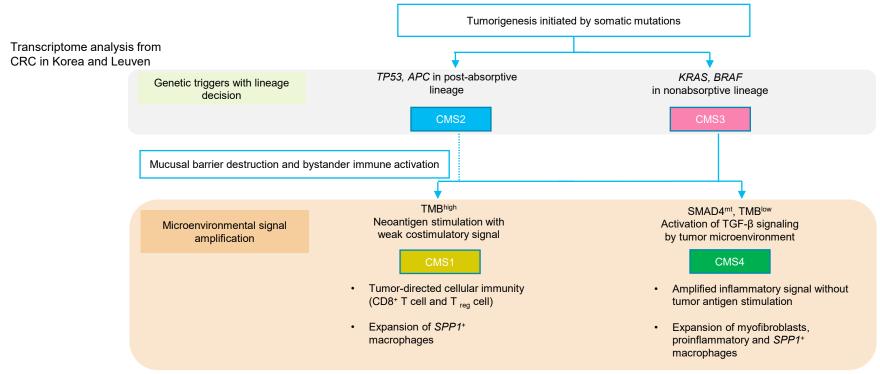
Figure 1 | Schematic representation of CRC subtypes. Microsatellite instability (MSI) is linked to hypermutation, hypermethylation, immune infiltration, activation of RAS, *BRAF* mutations, and locations in the proximal colon. Tumours with chromosomal instability (CIN) are more heterogeneous at the gene-expression level, showing a spectrum of pathway activation ranging from epithelial canonical (consensus molecular subtype 2 (CMS2)) to mesenchymal (CMS4). Tumours with CIN are mainly diagnosed in left colon or rectum, and their microenvironment is either poorly immunogenic or inflamed, with marked stromal infiltration. A subset of CRC tumours enriched for RAS mutations has strong metabolic adaptation (CMS3) and intermediate levels of mutation, methylation and copy number events. ECFR, epidermal growth factor receptor; JAK, Janus kinase; STAT, signal transducer and activator of transcription; TGFβ, transforming growth factor-β; VEGF, vascular endothelial growth factor; VEGFR, VEGF receptor.





1. Dienstmann R et al. Nat Rev Cancer 2017;17(2):79-92. 2. Becht et al. Advances Immunol 2016;130:95-190.

Lineage-dependent gene expression programs influence the immune landscape of colorectal cancer





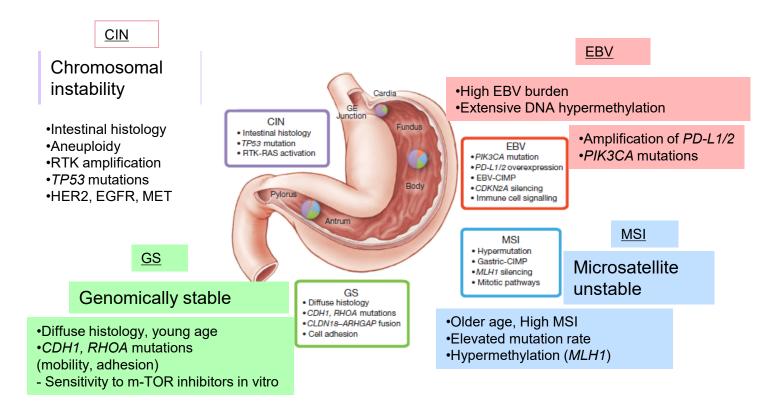
Adapted from Lee HO et al. Nat Genet 2020;52(6):594-603.

Colorectal Cancer: Two Different Diseases

CIN+ (85%) Chromosomal Instability	MSI-H (15%) Genetic (Microsatellite) Instability
Aneuploidy, loss of heterozygosity/loss of genetic material	Diploidy, no loss of heterozygosity
Proficient Mismatch Repair system Microsatellite stable (MSS)	Deficient Mismatch Repair system Microsatellite instability (MSI)
Sporadic or Familial Adenomatous Polyposis (FAP)	Sporadic or Lynch syndrome
95% of metastatic colorectal cancer	5% of metastatic colorectal cancer. Prognosis and chemosensitivity of MSI seems worse vs MSS
More prevalent in distal location	More prevalent in proximal location
Frequent mutation of KRAS	Frequent mutation of BRAFV600E
Tumor mutation burden low	Tumor mutation burden high Increased immune infiltration, higher tumour neo-antigens
No clear efficacy of immune check point inhibitor	Efficacy of immune check point inhibitor in phase I and II

MSI-H/MMRd tumors exhibit high mutational load and increased immune inflammation

Comprehensive Molecular Characterization in Gastric Cancer

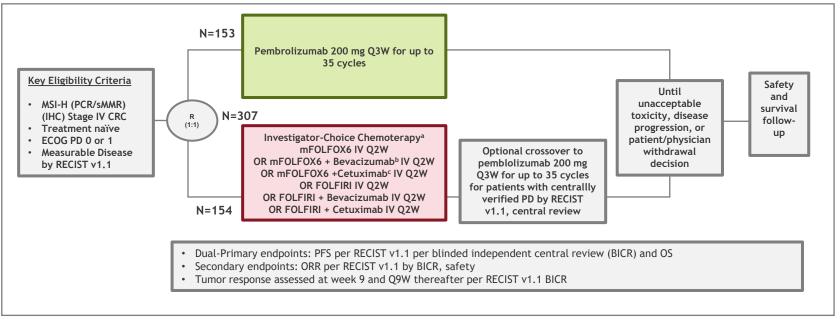




1. The cancer Genome Atlas Research Network. Nature 2014;513:202-9. 2. Lei et al. Gastroenterology 2013;145:554-565.

Keynote-177: Pembrolizumab vs chemotherapy in first line MSI-H mCRC

Study design (NCT02563002)



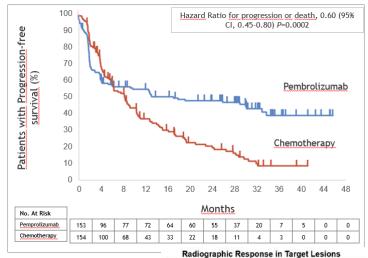
^aChosen before randomization; ^bBevacizumab 5 mg/kg IV; ^cCetuximab 400mg/m2 over 2 hours then 250 mg/m2 IV over 1 hour weekly.

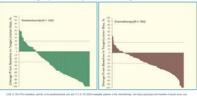
IHC: immunohistochemistry with hMLH1, hMSH2, hMSH6; PCR polymerase chain reaction; PFS, progression-free survival; OS: overall survival; ORR: overall response rate; Q9W: every 9 weeks.

Checkpoint inhibition in first line MSI-H mCRC

Keynote 177

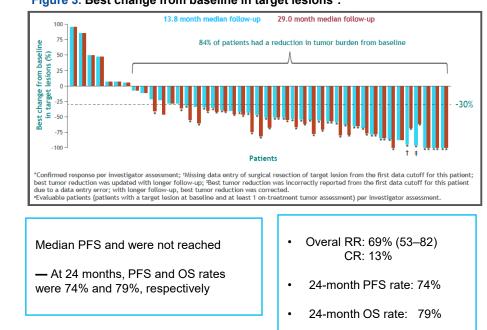
Figure 1. Progression-free Survival in Patients with MSI-H–dMMR Metastatic Colorectal Cancer ¹





Non-randomized phase 2 cohort in first line: checkmate 142 study, N=45

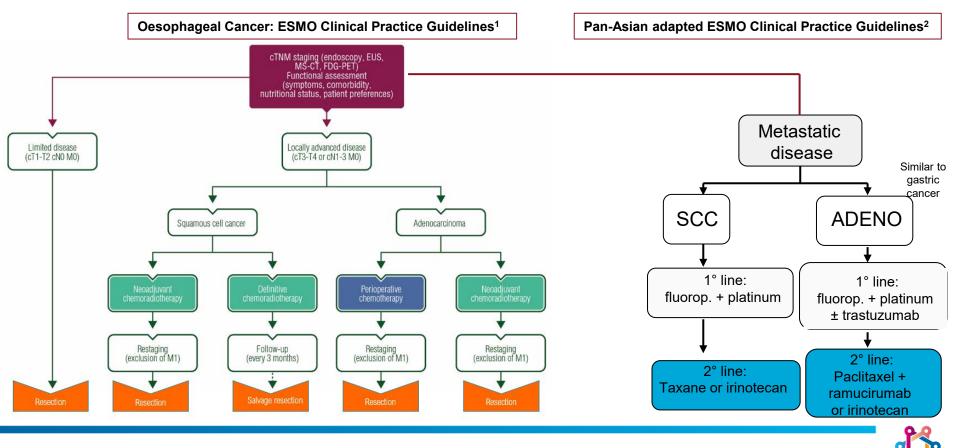
* CHMP positive opinion – EMA approval is not yet granted³ Figure 3. Best change from baseline in target lesions².





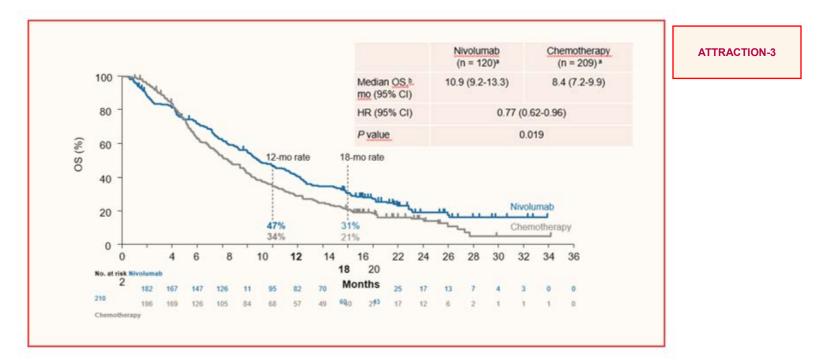
1. Andre T, Van Cutsem E et al. *N Engl J Med* 2020;383(23):2207-2218. 2. Adapted from Lenz H, Van Cutsem E et al. *J Clin Oncol* 2020;38:15_suppl 4040. 3. https://www.ema.europa.eu/en/news/meeting-highlights-committee-medicinal-products-human-use-chmp-17-20-may-2021

Oesophageal Cancer ESMO guidelines and JSMO/ESMO guidelines



1. Lordick F et al. Ann Oncol 2016;27(suppl 5):v50-v57. 2. Adapted from Muro K, Van Cutsem E et al. Published in 2018 – Ann Oncol 2019; 30:19–33.

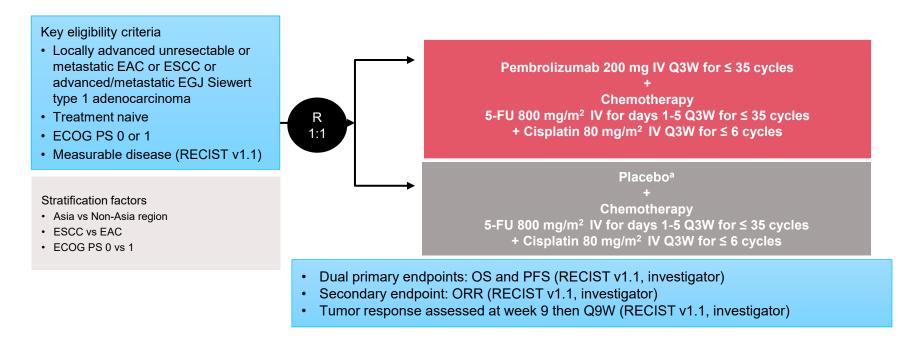
First approved checkpoint inhibitor in Oesophageal Cancer nivolumab versus docetaxel or paclitaxel in patients with ESCC



Although the benefit in CPS<1 (OS HR = 0.84 (95% CI 0.62, 1.14)) patients was lower in magnitude than in PD-L1 CPS≥1 patients (OS HR = 0.69 (95% CI: 0.51, 0.94)), nivolumab was approved in the EU for 2L esophageal cancer.



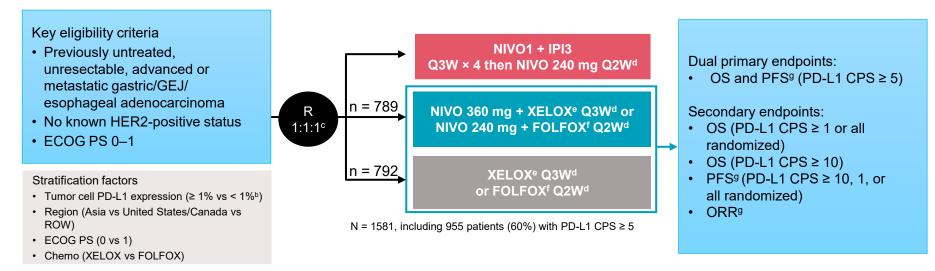
KN590: randomized, open-label, phase 3 study in 1st line esophageal cancer



^aSaline IV Q3W for ≤35 cycles. All treatments were continued for the specified number of cycles or until disease progression, intolerable toxicity, withdraw al of consent, or physician decision; EAC, esophageal adenocarcinoma; EGJ, esophagogastric junction, ESCC, esophageal squamous cell carcinoma.



CheckMate 649: randomized, open-label, phase 3 study in 1st line gastric cancer, gastroesophageal junction cancer, and esophageal adenocarcinoma



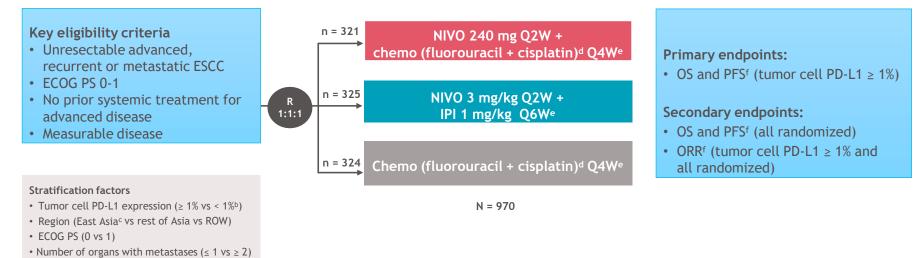
At data cutoff (May 27, 2020), the minimum follow-up was 12.1 months^h

^aClinicalTrials.gov number, NCT02872116; ^b< 1% includes indeterminate tumor cell PD-L1 expression; determined by PD-L1 IHC 28-8 pharmDx assay (Dako); ^cAfter NIVO + chemo arm was added and before new patient enrollment in the NIVO1+IPI3 group was closed; ^dUntil documented disease progression (unless consented to treatment beyond progression for NIVO + chemo), discontinuation due to toxicity, withdrawal of consent, or study end. NIVO is given for a maximum of 2 years; ^eOxaliplatin 130 mg/m² IV (day 1) and capecitabine 1000 mg/m² orally twice daily (days 1–14); ^fOxaliplatin 85 mg/m², leucovorin 400 mg/m², and FU 400 mg/m² IV (day 1) and FU 1200 mg/m² IV daily (days 1–2); ^gBICR assessed; ^hTime from concurrent randomization of the last patient to NIVO + chemo vs chemo to data cutoff.



Adapted from Moehler M et al. Oral presentation at ASCO 2021. Abstract 4002.

CheckMate 648: randomized, open-label, phase 3 study in 1st line for advanced esophageal squamous cell carcinoma



At data cutoff (January 18, 2021), the minimum follow-up was 12.9 months^g

^aClinicalTrials.gov. NCT03143153; ^b< 1% includes indeterminate tumor cell PD-L1 expression; determined by PD-L1 IHC 28-8 pharmDx assay (Dako); ^cEast Asia includes patients from Japan, Korea, and Taiwan; ^dFluorouracil 800 mg/m² IV daily (days 1-5) and cisplatin 80 mg/m² IV (day 1); ^eUntil documented disease progression (unless consented to treatment beyond progression for NIVO + IPI or NIVO + chemo), discontinuation due to toxicity, withdrawal of consent, or study end. NIVO is given alone or in combination with IPI for a maximum of 2 years; ^fPer blinded independent central review (BICR); ^gTime from last patient randomized to clinical data cutoff.



Adapted from Chau I et al. Oral presentation at ASCO 2021. Abstract LBA4001.

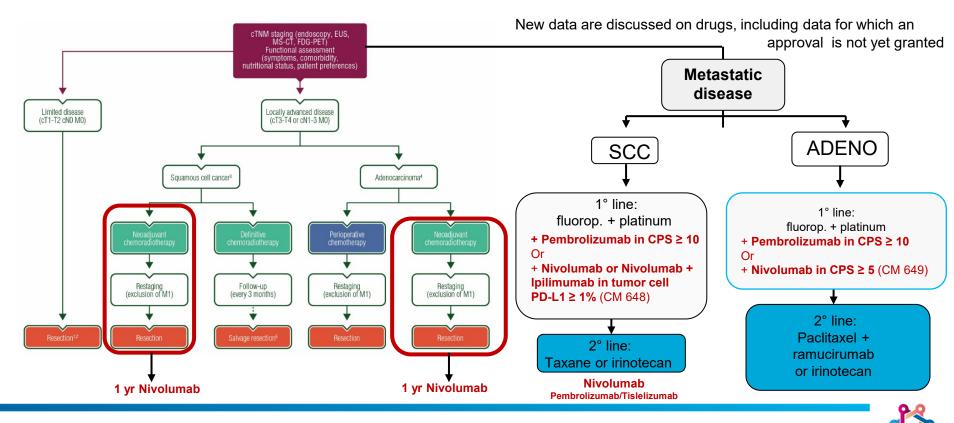
ESMO 2020 & ASCO 2021 in Upper GI Cancer: Practice Changing Data in Unmet Need Populations

POSITIVE TRIALS

- □ Metastatic (1° line) oesophageal cancer (SCC & adenocarcinoma):
 - ★ KEYNOTE 590¹: Pembrolizumab + chemotherapy vs chemotherapy in PD-L1 CPS ≥ 10
- □ Metastatic (1° line) oesophageal cancer (SCC):
 - CheckMate 648²: Nivolumab + chemotherapy or Nivolumab + Ipililumab vs chemotherapy in PDL ≥ 1% (and all comers)
 - **ESCORT-1**³: Camrelizumab + chemotherapy vs chemotherapy
- D Metastatic (1° line) gastric cancer, gastroesophageal junction cancer, and esophageal adenocarcinoma
 - CheckMate 649⁴: Nivolumab + chemotherapy vs chemotherapy in PD-L1 CPS ≥ 5
- Resected esophageal or gastroesophageal junction cancer (SCC & adenocarcinoma) following neoadjuvant chemoradiotherapy:
 - CheckMate 577⁵: nivolumab vs placebo

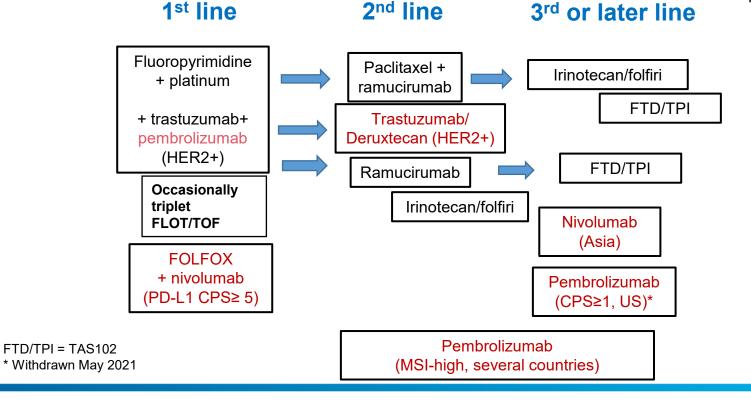
Kato et al. Oral presentation at the ESMO 2020. LBA8. 2. Chau I et al. Oral presentation at ASCO 2021. Abstract LBA4001. 3. Rui-hua X et al. Oral presentation at ASCO 2021. Abstract 4000. 4. Janjigian Y et al. *Lancet* 2021. Published Online June 5, 2021, https://doi.org/10.1016/S0140-6736(21)00797-2. 5. Kelly R, Van Cutsem E et al. *N Engl J Med* 2021;384:1191-203.

Esophageal Cancer ESMO guidelines and JSMO/ESMO guidelines updates in 2021 (personal opinion EVC based on evidence)



 Lordick F et al. Ann Oncol. 16;27(suppl 5):v50-v57. 2 MODIFIED by Eric Van Cutsem from Muro K, Van Cutsem E et al. Published in 2018 – Ann Oncol 2019;30:19–33. Updated algorithm for metastatic gastric cancer in 2021 (personal opinion EVC based on evidence)

> New data are discussed on drugs, including data for which an approval is not yet granted





MODIFIED by Eric Van Cutsem from Muro K, Van Cutsem E et al. JSMO-ESMO guidelines. Ann Onc 2019;30(1):19-33.

IMbrave150: Atezolizumab + Bevacizumab vs Sorafenib for First-line Treatment of Advanced HCC

Multicenter, randomized, open-label pase III trial

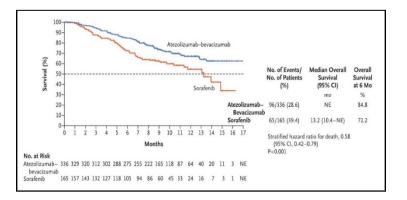
GO30140: randomized phase 1b study showed potential benefit of atezolizumab + bevacizumab for patients with advanced HCC (ORR 36%)

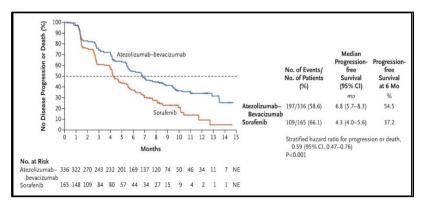


*Trial included subgroups of high-risk patients excluded from other contemporary phase III trials: \approx 40% had macrovascular invasion; specifically included patients with 50% hepatic involvement or main portal vein invasion or invasion of the portal vein branch contralateral to the primarily involved lobe.

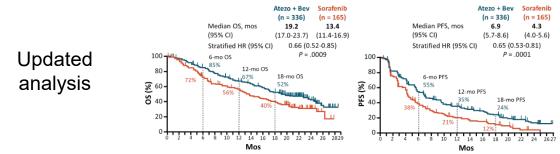


IMbrave150: Atezolizumab + Bevacizumab vs Sorafenib for First-line Treatment of Advanced HCC





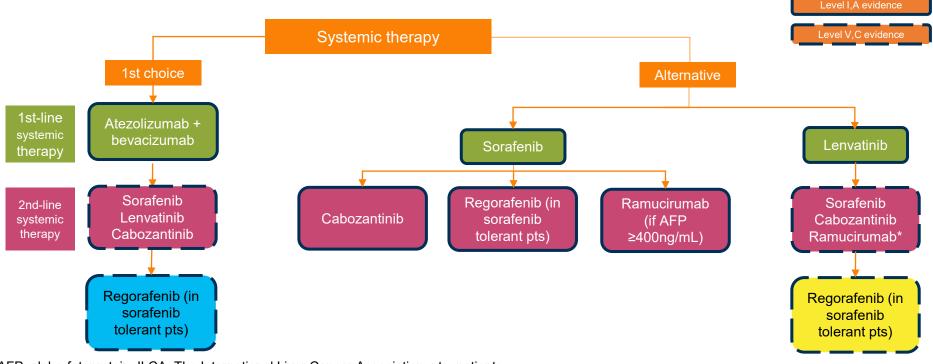
Primary analysis OS/PFS HR: 0.58/0.59 (median f/u 8.6 mos)





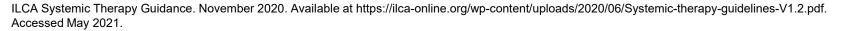
1. Finn RS et al. N Engl J Med 2020;382(20):1894-1905. 2. Finn RS et al. Abstract presented at the ASCO GI; Abstract 267.

Current HCC treatment guidelines ILCA 2020



AFP, alpha-fetoprotein; ILCA, The International Liver Cancer Association; pts, patients.

*If AFP ≥400ng/mL





Summary

- Very important practice changing data in 2020/2021 with checkpoint inhibitors in:
 - Esophageal cancer
 - Gastric cancer
 - Hepatocellular cancer

Leading to new treatment algorithms

- ► This is a fast-moving field with new data expected
- Optimal selection of patients is even more crucial





Importance of biomarkers in GI tumors

P. Pauwels (UZA, UA)



Programmed death ligand IHC scoring



Tumor proportion score (TPS): algorithm for assessment of PD-L1 expression on tumor cells

$$TPS (\%) = \frac{Positive \ tumour \ cells}{Total \ number \ of \ tumour \ cells} \times 100$$

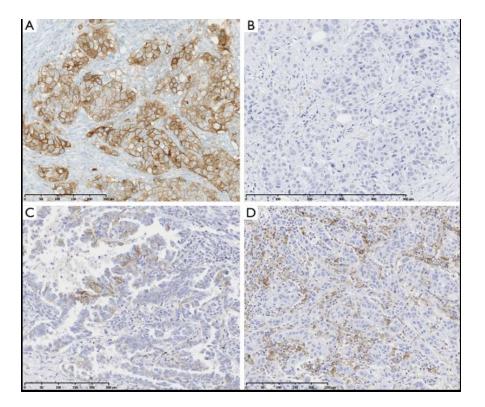


Combined positive score (CPS): algorithm for assessment of PD-L1 expression on both tumor and immune cells

 $CPS = \frac{Positive \ intratumour \ cells +}{Total \ number \ of \ tumour \ cells} \times 100$



IHC to assess staining pattern of PD-L1 in tumor and immune cells



IHC: Immuno Histochemical Staining



Lantuejoul S et al. J Thorac Dis 2019;(Suppl 1):S89-S101

Key messages

- Across multiple tumor types, response to immunotherapy is seen in patients with high expression of PD-L1, especially when PD-L1 is expressed in the immune cells¹
- PD-L1 expression varies across tumor types, and thresholds and scoring systems to determine PD-L1 positivity can vary between antibody assays²⁻³
- PD-L1 expression may be more prevalent on immune cells than tumor cells in certain cancers, such as melanoma, gastric, and colon¹
- Clinical utility of PD-L1 have been evaluated in several GI clinical trials, PD-L1 scoring method varies between the clinical trials of different immunotherapies⁴⁻⁷



1. Herbst RS et al. Nature 2014;515:563–567. 2. Krigsfeld G et al. Poster presentation at the 108th American Association for Cancer Research Annual Meeting; April 1–5, 2017; Washington, DC, USA. Abstract CT143. 3. Udall M et al. Diagn Pathol 2018;13:12.2. 4. Moehler M et al. Oral presentation at the European Society for Medical Oncology Annual Meeting (Virtual); September 13–21, 2020. Abstract 3047; 5. Lei M et al. Oral presentation at the 110th American Association for Cancer Research Annual Meeting; March 29–April 3, 2019; Atlanta, GA, USA. Abstract 2673. 6. Kelly RJ et al. Oral presentation at the European Society for Medical Oncology Annual Meeting (Virtual); September 19–21, 2020. Abstract 2968. 7 Boku N et al. Oral presentation at the European Society for Medical Oncology Annual Meeting (Virtual); September 19–21, 2020. Abstract 2968. 7

MSI/MMR testing

MSI/MMR: Microsatellite instability/mismatch repair

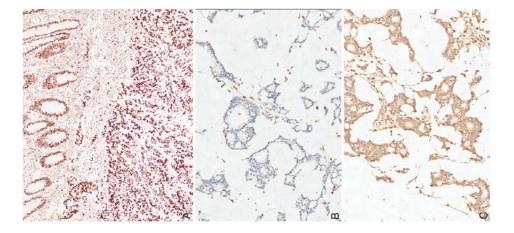


MMR: DNA repair pathway that identifies and corrects DNA mismatches





dMMR results from loss of expression of 1 or more of the proteins MLH1, MSH2, MSH6, or PMS2 involved in the MMR process









Presence or not of mutant markers to determine MSI-H or microsatellite stable (MSS)

ELSEVIER



Clinical Performance of the Idylla MSI Test for a Rapid Assessment of the DNA Microsatellite Status in Human Colorectal Cancer

Karen Zweenepoel,*¹ Julie Holmgaard Daeland,¹ Koen De Winne,* Vincent Raes,* Christine Beyn,*¹ Sazan Lambin,* Robina Dendooven,* Glenn Brieckx,* Torben Steliniche,¹ and Patrick Pauwels*¹

From the Laboratory of Pathninginal Austany, ¹ Autorey University Respiral (U2B), Edgern, Relgion; the Center for Oscological Research Autorey (CORE),¹ University of Autorety, Wiley, Belgium; and the Department of Pathning,¹ Autora University Research, Department

в Α Sample MSI Status MSS ACVR2A No mutation detected MSI Score 0.00 BTBD7 No mutation detected MSI Score 0.02 No mutation detected DID01 MSI Score 0.00 MRE11 No mutation detected MSI Score 0.04 RYR3 No mutation detected MSI Score 0.00 SEC31A No mutation detected MSI Score 0.00 SULF2 No mutation detected MSI Score 0.00 **Quality Status** 7 MSI biomarkers have been properly amplified and therefore the assay result is VALID. -NR21 BAT26 BAT25 NR24 MONO27 D MSI-H Sample MSI Status Mutation detected ACVR2A MSI Score 1.00 BTBD7 Mutation detected MSI Score 1.00 Mutation detected MSI Score 0.98 MRE11 Mutation detected MSI Score 1.00 RYR3 No mutation detected MSI Score 0.38 SEC31A No mutation detected MSI Score 0.04 SULF2 Mutation detected MSI Score 1.00 **Ouality Status** 7 MSI biomarkers have been properly amplified and therefore the assay result is -VALID. - 2 + + + +

Figure 1 Readout from the Idylla MSI test and Promega MSI analysis for a microsatellite stable (MSS) result (A and B) and a microsatellite instability—high (MSI-H) result (C and D).

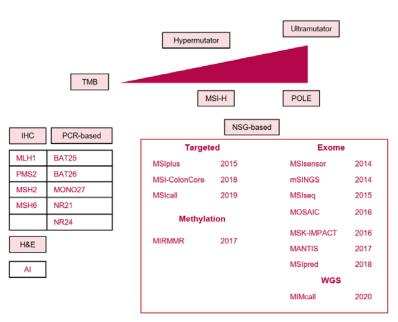
NR21

BAT26

BAT25 NR24

MONO27

MSI status is evaluated by IHC, PCR-based assays, and NGS-based techniques





Yamamoto H eta al. Arch Toxicol 2020;94(10):3349-3357.

Key messages

- MMR system is a DNA repair pathway that identifies and corrects DNA mismatches dMMR results from loss of expression of 1 or more of the proteins involved in the MMR process. dMMR can cause MSI^{1,2}
- MSI-H, indicative of dMMR, is found across several tumor types^{3,4}
- MSI-H/dMMR tumors are highly immunogenic, which may make these tumors susceptible to immune checkpoint inhibitors⁵
- MSI-H is reported to be around 15%⁶
- Pathologic staging is a key determinant of CRC prognosis and treatment. However, stageindependent outcome variability in patients with CRC supports the implementation of robust prognostic and predictive markers, such as MSI-H and dMMR⁷
- dMMR/MSI can be asssed by IHC, PCR and NGS⁸⁻¹⁰
- Clinical utility of MSI-H have been evaluated in several GI clinical trials¹¹⁻¹⁴

