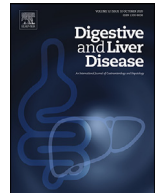




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Acquired microsatellite instability status and loss of HER2 positivity during treatment of gastro-esophageal junction adenocarcinoma [☆]

Dear Editor,

Gastro-oesophageal adenocarcinomas (GOAs) are a common cause of cancer mortality worldwide. Nowadays, beyond histological classification, the management of GOAs depends on molecular classification. The Cancer Genome Atlas consortium identified four different molecular subtypes of GOAs potentially allowing to tailor the treatment strategies [1]. About 20% of patients show an overexpression or amplification of the Human Epidermal Growth Factor Receptor-2 (HER2). These tumors are generally microsatellite stable (MSS) and present chromosomal instability [1]. The anti-HER2 monoclonal antibody trastuzumab combined with platinum and fluoropyrimidine chemotherapy is the standard first line regimen for these patients [2]. Microsatellite unstable (MSI) and DNA mismatch repair deficient (dMMR) cancers are a distinct subtype of GOAs with a prevalence of up to ~20% and being more frequent in early stage [3]. These tumors are often associated with a hypermutated phenotype resulting in extensive formation of cancer-specific neoantigens triggering robust anti-cancer immune responses. They benefit extensively from anti-PD-1 immune checkpoint blockade [2,4]. Here, we describe a rare case of acquisition of MSI and the concomitant loss of HER2 positivity potentially due to selective therapeutic pressure in a metastatic MSS GOA treated with platinum-based chemotherapy and trastuzumab.

A 61-year-old man, without family history of cancer, was diagnosed in January 2018 with a moderately (G2) to poorly (G3) differentiated HER2-positive (IHC 2+, mean HER2 copy number of 5.75 and HER2/CEP17 ratio of 2.09 as assessed by *in situ* hybridization) and MSS gastro-esophageal junction adenocarcinoma with peritumoral and mediastinal lymph nodes and multiple liver metastases. The histology appeared homogeneous across the 2 biopsies (samples) performed in the primary tumor. Combined treatment with trastuzumab and platinum-based chemotherapy was initiated: capecitabine (1000 mg/m² BID) for 14 days, oxaliplatin (130 mg/m²) and trastuzumab (8 mg/kg loading dose followed by 6 mg/kg) every three weeks. Partial response of all metastases was observed on imaging three months after treatment start. Due to poor tolerance and significant neuropathy, chemotherapy was stopped and trastuzumab monotherapy continued for additional 4 months. Seven months after treatment start (August 2018), the patient experienced symptomatic dysphagia. Imaging described a progression of the primary tumor and metastatic

lymph nodes without changes in liver metastases. Local progression of the tumor was confirmed by endoscopy. Rebiopsy of the same primary tumor site demonstrated the persistence of a (G2)-G3 adenocarcinoma, however HER2-negative (IHC 1+, mean HER2 copy number of 5.05 and HER2/CEP17 ratio of 1.38) and Mismatch Repair deficiency (dMMR, characterized by loss of expression of *MLH1* and *PMS2* proteins at immunohistochemistry staining) (Fig. 1). Like the diagnostic biopsy, no mixed tumor histology was observed. The MSI status was confirmed by the polymerase chain reaction (PCR) amplification and analysis of seven MSI biomarkers (IdyllaTM MSI assay: *ACVR2A*, *BTBD7*, *DIDO1*, *MRE11*, *RYR3*, *SEC31A*, *SULF2*). *MLH1* promoter was not hypermethylated (SALSA-MLPA ME011-D1 method). Considering the MSI status, a treatment with anti-PD1 immunotherapy (nivolumab, 3 mg/kg every two weeks) was initiated leading to partial response for 12 months. At progression (September 2019), the patient was sequentially treated with other types of immunotherapies (association of anti-PD-L1 and anti-TIM3 antibodies for 3 months, anti-CD25 antibody during two months into early-phase clinical studies) and anti-angiogenic treatment (ramucirumab) with and without chemotherapy (paclitaxel). Rebiopsy of the same primary tumor site was performed at each disease progression ($n = 4$) before each treatment modification and revealing the same histology pattern, as well as the persistence of HER2 negativity and MSI/dMMR status. Molecular characteristics of biopsies from the diagnosis (Jan 2018), the first (Aug 2018) and the last (Aug 2020) disease progression before treatment modification are summarized in the Fig. 1. Best supportive care was initiated in December 2020 and the patient eventually died in January 2021.

Comprehensive DNA sequencing (FoundationOne CDx assay) was performed on the initial diagnostic biopsy specimen of 2018 and the latest biopsy performed at progression in August 2020 (Figs. 1 and 2). Sequencing of the initial biopsy specimen confirmed the MSS status with a tumor mutational burden (TMB) of 5.0 Muts/Mb and found a clonal *TP53* and a sub-clonal *ERBB3* alteration (Table 1). Sequencing of the progressive lesion showed a genomic signature of MSI with a TMB of 23.9 Muts/Mb. Accordingly, several new variants were detected, mainly consisting of frameshift mutations (Table 1). Interestingly, the same *TP53*, *ERBB3*, and *STK11* alterations were still present, and among others two *MSH6* frameshift mutations had appeared. We observed no base substitution, insertion, deletion or copy number alteration in *MLH1* or *PMS2*.

HER2 and MSI molecular alterations are usually not coexisting or overlapping in time. Each distinguished profile predicts efficacy of a specific treatment. The addition of trastuzumab to platinum-based chemotherapy provided a significant survival benefit for patients with HER2-positive GOAs [2]. Loss of HER2 positivity after trastuzumab-based chemotherapy has been reported in ~30% of HER2 IHC3+ score and/or FISH amplified GOAs [5]. Several

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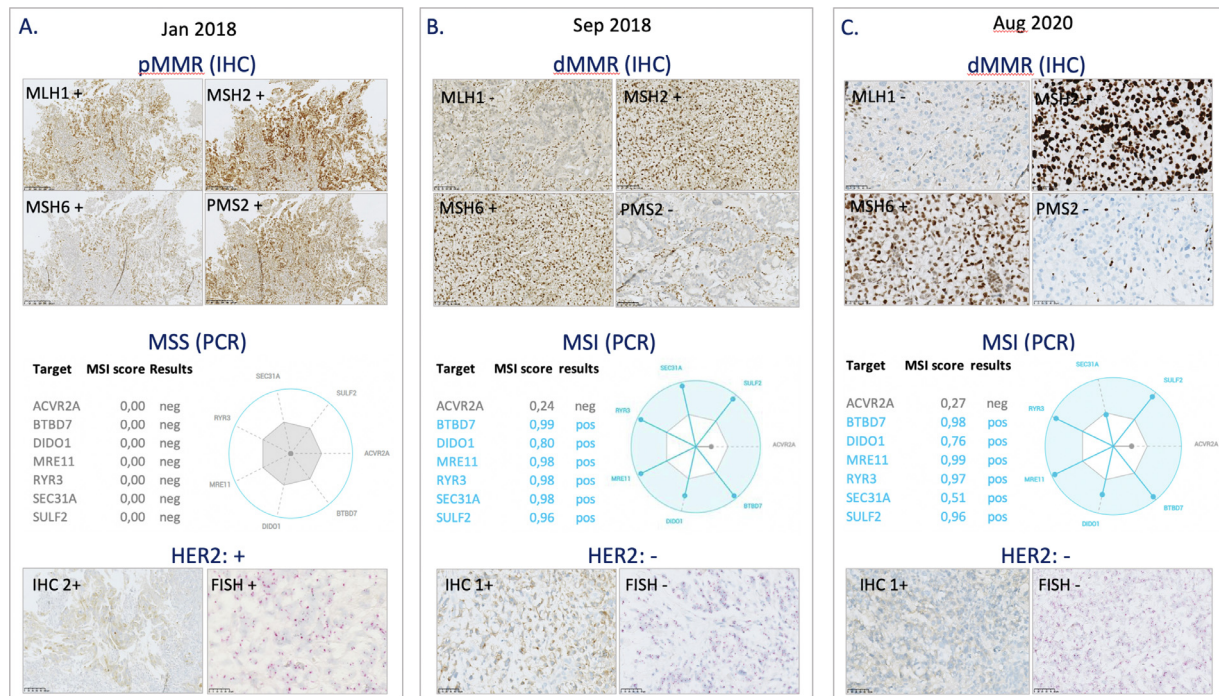


Fig. 1. Overtime molecular characterization for MSI and HER2 status. A: pMMR adenocarcinoma was characterized by IHC expression of *MLH1*, *MSH2*, *MSH6* and *PMS2* proteins. MSS was determined by PCR amplification and subsequent fragment analysis of seven monomorphic microsatellite biomarkers (*ACVR2A*, *BTBD7*, *DIDO1*, *MRE11*, *RYR3*, *SEC31A*, *SULF2*). No mutation has been found in the seven biomarkers (MSI score <0.5). HER2 had a weak to moderate membranous expression (IHC 2+) and was amplified (FISH: mean HER2 copy number of 5.75 and HER2/CEP17 ratio of 2.09). B and C: dMMR adenocarcinoma was characterized by loss of expression of *MLH1* and *PMS2* proteins at IHC staining. MSI was determined by PCR amplification and subsequent fragment analysis of seven monomorphic microsatellite biomarkers (*ACVR2A*, *BTBD7*, *DIDO1*, *MRE11*, *RYR3*, *SEC31A*, *SULF2*). The result is considered MSI-High because 6/7 biomarkers have a mutation detected (MSI score >0.5). HER2 had faint/barely perceptible membranous reactivity (IHC 1+) and was not amplified (FISH: mean HER2 copy number of 5.05 and HER2/CEP17 ratio of 1.38). dMMR: Mismatch Repair deficiency; FISH: fluorescence in situ hybridization; IHC: immunohistochemistry; MSI: microsatellite instability; MSS: microsatellite stability; neg: negative; PCR: polymerase chain reaction; pMMR: Mismatch Repair proficiency; pos: positive.



Fig. 2. Nucleotide and amino acid sequence (IGV browser) of the *MSH6* region carrying the two frameshift mutations acquired during cancer progression. The red lines delineate the nucleotidic repetitive region in which they are located.

studies including MSI cancers (GOAs and other tumor types) have reported higher treatment efficacy and survival benefit of anti-PD-1 immunotherapy, regardless of tumor origin [6].

MSI status has been correlated with older age, female gender, middle and lower gastric tumor localization, and poorly differentiated adenocarcinoma. MSI in GOAs is mainly related to germline mutations in a mismatch repair gene (Lynch Syndrome) or somatic methylation of a CpG island in the promoter region of *MLH1* and less frequently with somatic mutations of the mismatch repair machinery [1,2,6]. MSI status is generally considered an early genetic event rather than a status acquired through selective pressure. In our case, the evolution from MSS to MSI after platinum-based chemotherapy was demonstrated both at DNA and protein level. Nevertheless, we could not demonstrate the mechanism behind this evolution: *MLH1* promoter was not hypermethylated and DNA sequencing did not reveal a genetic alteration leading to the loss of expression of *MLH1* and *PMS2*, whereas the *MSH6* mutations did not lead to the loss of the corresponding protein expression. A post-translational mechanism is plausible, and we could hypothesize the treatment (platinum chemotherapy) led to reduced *MLH1* expression, as already reported [7,8]. The exhaustion of the biopsies however precludes us from performing RNA sequencing analyses. The somatic *MSH6* variants must be considered as a consequence of acquired microsatellite instability, rather than the cause. Indeed, these genomic positions are located in a mononucleotidic

or trinucleotidic repetitive region (Fig. 2). Heterogeneous loss of *MSH6* after chemotherapy has been described in literature in colorectal and endometrial cancer, whereas tumors with loss of *MLH1* and *PMS2* have been shown to acquire secondary *MSH6* alterations inside a microsatellite locus [9]. DNA sequencing furthermore confirmed the progressive lesion was phylogenetically related to the diagnostic cancer sample, given the persistence of the early *TP53* and *ERBB3* alterations.

Another hypothesis related to our observation could be the possibility of intra-tumoral heterogeneity of MMR status rather than acquired MSI tumor, as previously reported [10]. Heterogeneous staining patterns of MMR proteins (restricted spatial loss restricted to some tumor area) associated with confirmed MSI status has been exceptionally reported in colorectal cancer [10,11]. In our case, we never observed a mixed histology or a spatially heterogeneous MMR protein expression in the multiple biopsies performed overtime (5 time-points) at the same primary tumor site. Nevertheless, a mixed tumor compound (HER2 and MSI) could still have been missed by the initial biopsy. Clonal selection during exposure to trastuzumab is likely to occur in GOAs as reflected by poor efficacy of this treatment beyond progression [12]. Shared *TP53*, *ERBB3*, and *STK11* clonal and subclonal alterations suggested at least a common phylogenetic process in the tumor and exclude the presence of two different cancer types. The hypermutator-phenotype of MSI GOAs enables high evolvability producing ex-

Table 1

FoundationOne CDx assay findings.

January 2018					August 2020				
TMB	MS	Gene	Alteration	VAF (%)	TMB	MS	Gene	Alteration	VAF (%)
5.0 muts/Mb	stable	ERBB3	p.(V104M)	3,4	23.9 muts/Mb	unstable	ERBB3	p.(V104M)	10,1
		TP53	p.(G279E)	28,2			TP53	p.(G279E)	12
		/					TP53	p.(S362Afs*8)	11,4
		INPP4	Exon 21–22 deletion				INPP4	Exon 21–22 deletion	
		PTEN	p.(T319*)	24,2					
		STK11	P281fs*6	4,4			STK11	P281fs*6	11,1
							MSH6	G1070fs*9	
							MSH6	F1088fs*5	
							PTCH1	N97fs*20	
							PTCH1	S1203fs*52	
							MLL2	Q2932fs*10	
							QK1	K134fs*14	
							CREBBP	S976fs*22	
							ARID1A	P146fs*86	
							CDK12	3761–2A>C	
			PTEN	loss					
			ROS1	amplification					

VAF: Variant Allele Frequency, MS: Microsatellite, TMB: Tumor Mutational Burden.

treme intratumor heterogeneity [13]. Nevertheless, it seems highly unlikely that we faced a MSI GOA since the diagnosis, in which a subclone characterized by a driver alteration (HER2+) and a concomitant loss of MSI status was identified in the initial biopsy, definitively eliminated by trastuzumab and chemotherapy and not found in sequential biopsies overtime. Like MSI status, HER2 overexpression in premalignant gastric lesions suggests its potential involvement in the early steps of gastric carcinogenesis [1].

In conclusion, we described a rare case of MSI acquisition (and concomitant loss of HER2 positivity) in a patient with metastatic MSS gastro-esophageal junction cancer treated with platinum-based chemotherapy and trastuzumab. Despite our efforts, we were unable to identify the mechanism that led to MSI status. Biopsies performed at each stage of disease progression supported our decision on the best treatment option that allowed the patient to subsequently benefit from immunotherapy. This case also reinforces the need to rebiopsy progressive lesions to ensure optimal treatment selection.

Compliance with Ethical Standards

All the authors of this manuscript declare no conflict of interest.

The study protocol conforms to the ethical guidelines of the 1975 Declaration of Helsinki (6th revision, 2008) as reflected in a priori approval by the institution's human research committee.

The authors obtained informed consent to publish the report from patient's legal representative (patient deceased).

Conflict of interest

None declared.

References

- [1] Cancer Genome Atlas Research Network/Analysis Working Group: asan University, BC Cancer Agency, et al. Integrated genomic characterization of oesophageal carcinoma. *Nature* 2017;541(7636):169–75.
- [2] Lordick F, Carneiro F, Cascinu S, et al. Gastric cancer: ESMO Clinical practice guideline for diagnosis, treatment and follow-up. *Ann Oncol* 2022;33:1005–20.
- [3] Puliga E, Corso S, Pietrantonio F, et al. Microsatellite instability in gastric cancer: between lights and shadows. *Cancer Treat Rev* 2021;95:102175.
- [4] Chao J, Fuchs CS, Shitara K, et al. Assessment of pembrolizumab therapy for the treatment of microsatellite instability-high gastric or gastroesophageal junction cancer among patients in the KEYNOTE-059, KEYNOTE-061, and KEYNOTE-062 clinical trials. *JAMA Oncol* 2021;7:895–902.
- [5] Seo S, Ryu MH, Park YS, et al. Loss of HER2 positivity after anti-HER2 chemotherapy in HER2-positive gastric cancer patients: results of the GASTRIC cancer HER2 reassessment study 3 (GASTHER3). *Gastric Cancer* 2019;22:527–35.
- [6] Le DT, Durham J, Smith K, et al. Mismatch repair deficiency predicts response of solid tumors to PD-1 blockade. *Science* 2017;357:409–13.
- [7] Kishi K, Doki Y, Yano M, et al. Reduced MLH1 expression after chemotherapy is an indicator for poor prognosis in esophageal cancers. *Clin Cancer Res* 2003;9:4368–75.
- [8] Sood S, Patel FD, Srinivasan R, et al. Chemoradiation therapy induces in vivo changes in gene promoter methylation & gene transcript expression in patients with invasive cervical cancer. *Indian J Med Res* 2018;147:151–7.
- [9] Graham RP, Kerr SE, Butz ML, et al. Heterogenous MSH6 loss is a result of microsatellite instability within MSH6 and occurs in sporadic and hereditary colorectal and endometrial carcinomas. *Am J Surg Pathol* 2015;39:1370–6.
- [10] Tachon G, Frouin E, Karayan-Tapon L, et al. Heterogeneity of mismatch repair defect in colorectal cancer and its implications in clinical practice. *Eur J Cancer* 2018;95:112–16.
- [11] Greenberg A, Kariv R, Solar I, et al. Geographic heterogeneity for mismatch repair proteins is associated with defects in DNA repair. *ISR Med Assoc J* 2020;22:32–6.
- [12] Makiyama A, Sukawa Y, Kashiwada T, et al. Randomized, Phase II study of trastuzumab beyond progression in patients with HER2-positive advanced gastric or gastroesophageal junction cancer: WJOG7112G (T-ACT study). *J Clin Oncol* 2020;38:1919–27.
- [13] von Loga K, Woolston A, Punta M, et al. Extreme intratumour heterogeneity and driver evolution in mismatch repair deficient gastro- oesophageal cancer. *Nat Commun* 2020;11:675.

Lynn Gabrielle Alexis

Department of Medical Oncology, Institut Roi Albert II, Cliniques Universitaires Saint-Luc, Université Catholique de Louvain (UCLouvain), Brussels, Belgium

Hélène Dano

Department of Pathology, Institut Roi Albert II, Cliniques Universitaires Saint-Luc, Université Catholique de Louvain (UCLouvain), Brussels, Belgium

Anne-France Dekairelle

Department of Genetics, Institut Roi Albert II, Cliniques Universitaires Saint-Luc, Université Catholique de Louvain (UCLouvain), Brussels, Belgium

Cédric Van Marcke*

Department of Medical Oncology, Institut Roi Albert II, Cliniques Universitaires Saint-Luc, Université Catholique de Louvain (UCLouvain), Brussels, Belgium

Marc Van den Eynde**

Department of Medical Oncology and Hepato-Gastroenterology, Institut Roi Albert II, Cliniques Universitaires Saint-Luc, Université Catholique de Louvain (UCLouvain), Brussels, Belgium

*Corresponding author: Cliniques universitaires Saint-Luc, Avenue Hippocrate, 10 - 1200 Brussels - Belgium.
E-mail address: marc.vandeneynde@uclouvain.be (M. Van den Eynde)

* These authors contributed equally to this work.