

## Erratum to: Microsatellite instability in colorectal cancer: from molecular oncogenic mechanisms to clinical implications

Aziz Zaanan · Katy Meunier · Fatiha Sangar ·  
Jean-François Fléjou · Françoise Praz

Published online: 31 May 2011  
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**Erratum to: Cell Oncol.**  
**DOI 10.1007/s13402-011-0024-x**

In the above mentioned review two errors in the legend of Fig. 1 has escaped the author's attention during proofreading. The correct legend of Fig. 1 should read:

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The online version of the original article can be found at <http://dx.doi.org/10.1007/s13402-011-0024-x>.

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A. Zaanan · K. Meunier · F. Sangar · J.-F. Fléjou · F. Praz  
INSERM, UMR\_S 938, Centre de Recherche Saint-Antoine,  
75012 Paris, France

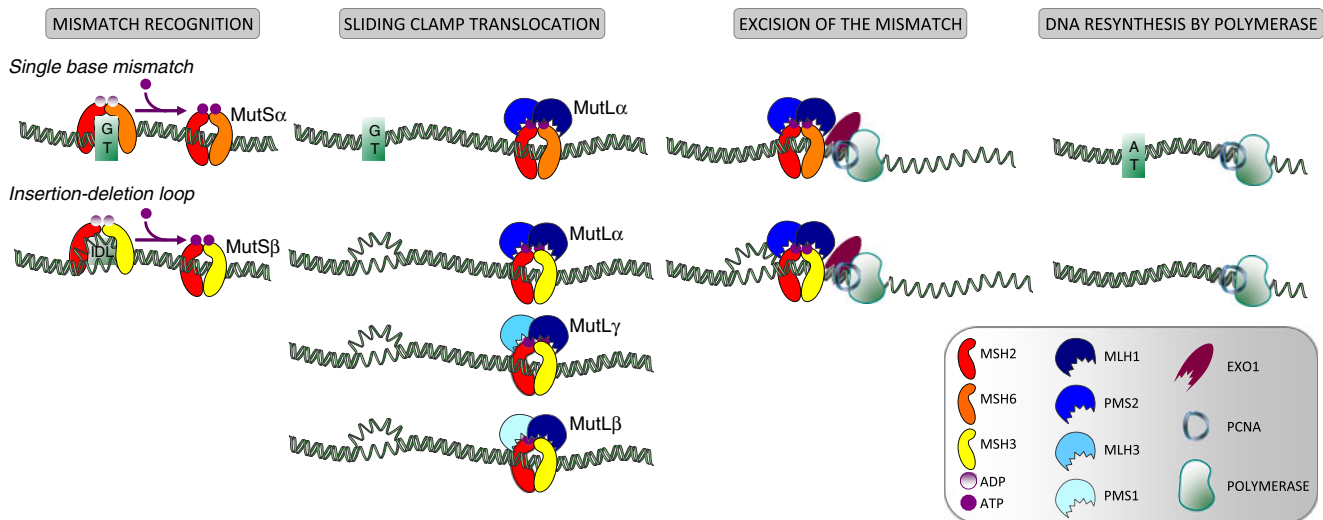
A. Zaanan · K. Meunier · F. Sangar · J.-F. Fléjou · F. Praz  
UPMC Univ Paris 06, UMR\_S 938,  
Centre de Recherche Saint-Antoine,  
75012 Paris, France

A. Zaanan  
Service d'Oncologie Médicale, Hôpital Saint-Antoine,  
Assistance Publique–Hôpitaux de Paris,  
75012 Paris, France

K. Meunier  
Service de Chirurgie Oncologique, Centre Paul Papin,  
Angers, France

J.-F. Fléjou  
Service d'Anatomie et Cytologie Pathologique,  
Hôpital Saint-Antoine, Assistance Publique–Hôpitaux de Paris,  
75012 Paris, France

F. Praz (✉)  
Centre de Recherche Saint-Antoine,  
INSERM UMRS\_938-UPMC,  
184 rue du Faubourg Saint-Antoine,  
75571 Paris, Cedex 12, France  
e-mail: [Francoise.Praz@inserm.fr](mailto:Francoise.Praz@inserm.fr)



**Fig. 1** Eukaryotic DNA mismatch repair. Single nucleotide mismatches, as well as 1 nucleotide loops typically occurring at microsatellite mononucleotide repeated sequences, are recognized by the MutS $\alpha$  heterodimer formed by MSH2 and MSH6; this complex binds to DNA as a sliding clamp after MSH2 ADP has been exchanged for ATP (upper panel). The heterodimer MutL $\alpha$ , formed by MLH1 and PMS2, then binds to MutS $\alpha$ . The MutS $\alpha$ -MutL $\alpha$  complex moves along the DNA until it meets the DNA polymerase complex, PCNA and Exo I. The exonuclease then removes up to several hundred bases from the newly synthesized DNA strand, to allow correct resynthesis

of DNA by the replicative polymerase. IDL of two or more nucleotides are preferentially recognized by the MutS $\beta$  complex, formed by MSH2 and MSH3 (lower panel). After binding to DNA, MutS $\beta$  exchanges ADP for ATP and recruits the MutL $\alpha$  complex, a heterodimer of MLH1 and PMS2 heterodimer. Subsequently, excision and resynthesis are performed as described for the repair of single base mismatches. MLH1 may also form heterodimers with MLH3 and PMS1 to form respectively the MutL $\gamma$  and MutL $\beta$  complexes, but their roles in human MMR function are not entirely clear