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Fagerlund A(1), Ween O(1), Lund T(1), Hardy SP(2), Granum PE(1).

Author information:

(1)Department of Food Safety and Infection Biology, The Norwegian School of Veterinary Science, PO Box 8146 Dep., N-0033 Oslo, Norway.

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doi: 10.1007/s00122-0  
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protein-protein interaction networks (Li et al.2012a), as well as DNA regulatory machinery elements (Barabasi et al.2011).

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domain that has been proposed in the MLH gene family on the basis of genetic and crystallographic analysis of the N-

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terminal fragment of  
MutL (Ban et al. 1999).  
Mutator phenotype  
exhibited by mlh1ts  
alleles at permissive  
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temperatures: To  
determine whether  
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