

Selective pressures for the high prevalence of MEFV variants induced by smallpox infection in the “Old World”: A hypothesis

Sirs,

High frequency of the heterozygous carriers of MEFV mutations, which are associated with familial Mediterranean fever, in certain ethnic groups poses an important question of selective advantage against yet unknown fatal infections by a tendency to develop stronger inflammatory response (Fig. 1). Cattan recently discussed the possibility of a lower rate of mortality from tuberculosis conferred by MEFV variants based on mortality records of Tunis in the first half of the 20th century (1). I herein propose that smallpox may be a more likely selective pressure for the MEFV variations. Smallpox had been one of the most dreadful infections in human life, with a case fatality rate of 20% (2). Historical sources indicate that smallpox was already endemic in Egypt and Mesopotamia by the second century AD. It spread to Northern Europe in the 11th and 12th centuries, and to America much later (2). These records may suggest that smallpox may have acted as a selective pressure longer and increased the MEFV allele frequencies in geographic regions where FMF is prevalent.

Recently, Galvani and Slatkin evaluated the bubonic plague and smallpox as the selective pressures for the chemokine receptor CCR5-Δ32 allele using an age-structured model (3). They suggested that smallpox can explain better than plague the selective rise of CCR5-Δ32 allele to current frequencies with a relatively high case fatality rate, the persistence of the infection more continuously since the origin of the allele, the affection of mainly younger people with reproductive potential, and a higher cumulative death toll (3). Their argument about the timing of the appearance of the resistance allele and the number of generations that the disease can drive the allele to its current frequencies can be applied to the MEFV variants. Age of MEFV mutations reaching up to 2500 years can be explained by considering an incomplete dominance model, in which a heterozygous allele confers less protection against disease mortality compared to the carriers of two copies of the resistance alleles and also the presence of more than one variant with varying degrees of protection (3, 4).

The MEFV variants can cause a reduced expression of pyrin, which results in increased activation caspase 1 (also known as interleukin-1β (IL-1β) converting enzyme (ICE)) and IL-1β. Poxviruses developed many defence strategies to suppress host immunity, which included serpin protease inhibitors (serpins) interfering with proteolytic activity of ICE, thus inhibiting the IL-1β activation (5). Inactivation of the ser-

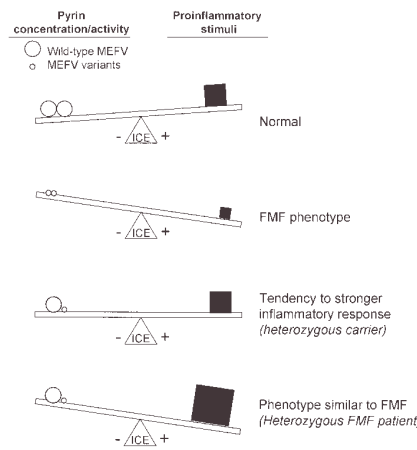


Fig. 1. Proposed model of pyrin activity in fine-tuning of inflammation through controlling the interleukin-1β converting enzyme (ICE) activity in normal individuals and in carriers of 1 or 2 copies of MEFV variants.

pin genes in the vaccinia virus resulted in an increased vaccine safety with attenuated activity without compromising the strength of the immune responses (5). Thus, inherent increased ICE activity resulting from MEFV mutations can be advantageous by conferring decreased smallpox mortality, if not inhibiting the contraction of infection itself.

On the other hand, selective increase of heterozygous carriers of the MEFV mutations due to smallpox or other infections may also constitute a population health problem. FMF has been known as an autosomal recessively inherited disorder. However, depending on the type or location of the mutations (i.e. ΔM694, complex E148Q/M694V or H478Y), occasional cases with true autosomal dominant inheritance can be seen (6, 7). Other genetic polymorphisms or accompanying inflammatory disorders can also affect the expression of FMF in heterozygous carriers (8, 9). It can be assumed that depending on the site, intensity, duration or other characteristics of accompanying inflammation, FMF phenotype can be developed in individuals carrying a single MEFV mutation, at least temporarily (Fig. 1). MEFV mutations can also affect the severity of the accompanying inflammatory condition, as observed in patients with multiple sclerosis and rheumatoid arthritis (10, 11).

Pyrin seems to act as negative regulator in the fine-tuning of inflammation through affecting the ICE activity, and MEFV-variants-related disadvantages, especially for populations with a high heterozygous carrier rate, need to be studied further.

AHMET GÜL, *Professor of Medicine Istanbul University, Istanbul Faculty of Medicine, Department of Internal Medicine, Division of Rheumatology, 34390 Capa, Istanbul.*
E-mail: agul@istanbul.edu.tr

References

- CATTAN D: Familial Mediterranean fever: Is low mortality from tuberculosis a specific advantage for MEFV mutation carriers? Mortality from tuberculosis among Muslims, Jewish, French, Italian and Maltese patients in Tunis (Tunisia) in the first half of the 20th century. *Clin Exp Rheumatol* 2003; 21 (Suppl. 30): S53-4.
- EYLER JM: Smallpox in history: the birth, death and impact of a dread disease. *J Lab Clin Med* 2003; 142: 216-20.
- GALVANI AP, SLATKIN M: Evaluating plague and smallpox as historical selective pressures for the CCR5-D32 HIV resistance allele. *Proc Natl Acad Sci USA* 2003; 100: 15276-9.
- INTERNATIONAL FMF CONSORTIUM: Ancient missense mutations in a new member of the RoRet gene family are likely to cause familial Mediterranean fever. *Cell* 1997; 90: 797-807.
- LEGRAND FA, VERARDI PH, JONES LA, CHAN KS, PENG Y, YILMA TD: Induction of potent humoral and cell-mediated immune responses by attenuated vaccinia virus vectors with deleted serpin genes. *J Virol* 2004; 78: 2770-9.
- BOOTH DR, GILLMORE JD, LACHMANN HJ *et al.*: The genetic basis of autosomal dominant familial Mediterranean fever. *Q J Med* 2000; 93: 217-21.
- ALDEA A, CAMPISTOL JM, AROSTEGUI JI *et al.*: A severe autosomal-dominant periodic inflammatory disorder with renal AA amyloidosis and colchicine resistance associated to the MEFV H478Y variant in a Spanish kindred: An unusual familial Mediterranean fever phenotype or another MEFV-associated periodic inflammatory disorder? *Am J Med Genet* 2004; 124A: 67-73.
- HOLMES AH, BOOTH DR, HAWKINS PN: Familial Mediterranean fever gene. *N Engl J Med* 1998; 338: 992-3.
- LIVNEH A, AKSENTIEVICH I, LANGEVITZ P *et al.*: A single mutated MEFV allele in patients suffering from familial Mediterranean fever and Behçet's disease (FMF-BD). *Eur J Hum Genet* 2001; 9: 191-6.
- SHINAR Y, LIVNEH A, VILLA Y *et al.*: Common mutations in the familial Mediterranean fever gene associate with rapid progression to disability in non-Ashkenazi Jewish multiple sclerosis patients. *Genes Immun* 2003; 4: 197-203.
- RABINOVICH E, LIVNEH A, LANGEVITZ P *et al.*: Severe disease in patients with rheumatoid arthritis carrying a mutation in the Mediterranean fever gene. *Ann Rheum Dis* 2005; 64: 1009-14.

What effect do dietary antioxidants have on the symptoms and structural progression of knee osteoarthritis over two years?

Sirs,

There has been evidence that antioxidant intake may be associated with reduced progression of radiographic knee osteoarthritis (OA). We performed a prospective cohort study to examine the effect of dietary antioxidants on symptoms, cartilage volume and their change over 2 years in subjects with knee OA.

One hundred and thirty-six subjects who fulfilled American College of Rheumatology clinical and radiographic criteria for knee OA (1) entered the study. General health, pain, stiffness, and function were assessed using 36-Item Short-Form Health Survey (SF-36) (2) and Western Ontario