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Short Report

Evaluation of Clinical, Genetic and Treatment-Related Characteristics in FMF Patients by Gender Distribution

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Individuals with familial Mediterranean fever (FMF) may be exposed to stress due to gender-related differences and consequently the frequency of attacks may be different. For example, FMF attacks can be triggered in women during menstrual periods.^{1,2} The aim of this study is to investigate the differences between males and females in clinical findings, hereditary characteristics, treatment responses and pathogen Mediterranean fever (MEFV) gene phenotype frequencies in FMF patients.

The charts of 213 patients who were followed up in the rheumatology outpatient clinic with a diagnosis of FMF were retrospectively reviewed. The data of 105 patients (70 females, 35 males) whose charts were available for all research data were evaluated. While evaluating the clinical findings; The age of attack onset, attack character (typical, atypical), dominant attack location (peritoneum, pleura, synovia, isolated fever), presence of recurrent fever, appendectomy history, family history (first degree, second degree) were evaluated. While evaluating the treatment response, the response (complete, partial, unresponsive) to colchicine treatment was questioned. The phenotype frequencies of the pathogen variations (M694V, V726A, M680I, E148Q) in the MEFV gene were determined. Findings were compared between groups.

The median age (minimum-maximum) in women and men was 37.5 (19-62) and 30.0 (19-59) years, respectively (p=0.148). Demographic clinical findings, characteristics. treatment responses of the participants are summarized in Table 1. The frequency of individuals with typical attack character was 71.4% in women and 82.9% in men. The frequency of the predominant attack localization with peritoneum was 90% in women and 88.6% in men. The frequency of recurrent fever in women and men was 67.1% and 65.7%, the frequency of appendectomy was 34.3% and 42.9%, and the presence of a family history was



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| Variables | Female (n=70) | Male (n=35) | p value |
|---------------------------------------|-----------------------|--------------|---------|
| Age (years) | 37.5 (19-62) | 30.0 (19-59) | 0.148 |
| Attack onset age (years) | 15.5 (1-50) | 16.0 (4-43) | 0.965 |
| Clinical, hereditary and treatment-re | lated findings, n (%) | | |
| Attack character | | | 0.201 |
| Typical | 50 (71.4) | 29 (82.9) | |
| Atypical | 20 (28.6) | 6 (17.1) | |
| Dominant attack localization | | | 0.557 |
| Peritoneum | 63 (90.0) | 31 (88.6) | |
| Pleura | 1 (1.4) | 0 (0) | |
| Joint | 5 (7.1) | 2 (5.7) | |
| Isolated fever | 1 (1.2) | 2 (5.7) | |
| Relapsing fever | 47 (67.1) | 23 (65.7) | 0.884 |
| Appendectomy | 24 (34.3) | 15 (42.9) | 0.392 |
| Family history | 53 (75.7) | 27 (76.1) | 0.175 |
| First degree | 40 (57.1) | 25 (71.4) | |
| Second degree | 13 (18.6) | 2 (5.7) | |
| Colchicine response | | | 0.427 |
| Good | 57 (81.4) | 25 (71.4) | |
| Limited | 8 (11.4) | 5 (14.3) | |
| No | 5 (7.2) | 5 (14.3) | |

 Table 1. Evaluation of clinical, hereditary and treatment-related characteristics of individuals with

 Familial Mediterranean fever disease by gender distribution

| Table 2. Comparison of phenotype | c frequencies of MEFV | gene mutations | according to gender |
|--|----------------------------|----------------|---------------------|
| distribution of individuals with Famil | ial Mediterranean fever di | sease | |

| Phenotype frequency (n, %) | Female (n=70) | Male (n=35) | p value | OR (CI) |
|-------------------------------|------------------|-------------|---------|----------------------|
| M694V | 42 (62.7) | 31 (88.6) | 0.006 | 4.613 (1.456-14.613) |
| V726A | 15 (22.4) | 5 (14.3) | 0.328 | 0.578 (0.191-1.749) |
| M680I | 14 (20.9) | 8 (22.9) | 0.819 | 1.122 (0.419-3.002) |
| E148Q | 10 (14.9) | 5 (14.3) | 0.931 | 0.950 (0.298-3.033) |
| | 0 0 0 1 I | di at at | | |

MEFV: Mediterranean fever, OR: Odds ratio, CI: confidence interval.

75.7% and 76.1%, respectively. The proportion of those with colchicine response was 92.8% and 85.7% in women and men, respectively. Clinical findings and colchicine response were not different between genders. The phenotype frequency of pathogen MEFV gene mutations are summarized in Table 2. The phenotype frequencies of pathogen MEFV gene mutations were 62.7% and 88.6% for M694V (p=0.006), 22.4% and 14.3% (p=0.328) for V726A, 20.9% and 22.9 for M680I (p=0.819) and 14.9% and 14.3% (p=0.931) for E148Q in women and men, respectively.

Clinical findings and treatment responses are not different in individuals with FMF disease. The frequency of the M694V mutation, which has high penetration and is associated with important complications such as amyloidosis, is higher in men. There is a need for studies to evaluate FMF activity according to gender distribution.

Conflict of Interests

Authors declare that there are none.

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