MSH2 and MLH1 testing

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DNA testing is recommended in families fulfilling at least "suspected HNPCC" criteria. After exclusion of FAP (characteristic FAP features include polyposis, congenital hypertrophy of the retinal pigment epithelium, cysts and osteomata of bones of the maxilla and mandible, desmoid tumours), immunohistochemical analyses (IHC) of MLH1, MSH2 and MSH6 expression in malignant tissues should be performed (absence of the protein may indicate the mutated gene).

The results of several studies performed in our centre characterised the frequencies and spectrum of MSH2 and MLH1 mutations in Poland [1]. Similarly to other populations, the most frequent causes of HNPCC in Poland are MLH1 and MSH2 mutations, constituting 90% of all mutations associated with this syndrome. MLPA detects 10% of these mutations. In over 60% of all HNPCC families recurrent mutations can be found. Thus after IHC, MLPA for MSH2 and MLH1 should be performed. Next, with MLPA negative, DNA tests searching for recurrent mutations, characteristic for the Polish population, should be applied. The last step should include DHPLC [2] and sequencing of the cases indicated by DHPLC results.

References

- Kurzawski G, Suchy J, Lener M, Kłujszo-Grabowska E, Kładny J, Safranow K, Jakubowska K, Jakubowska A, Huzarski T, Byrski T, Debniak T, Cybulski C, Gronwald J, Oszurek O, Oszutowska D, Kowalska E, Góźdź S, Niepsuj S, Słomski R, Pławski A, Łacka-Wojciechowska A, Rozmiarek A, Fiszer-Maliszewska Ł, Bebenek M, Sorokin D, Sasiadek MM, Stembalska A, Grzebieniak Z, Kilar E, Stawicka M, Godlewski D, Richter P, Brozek I, Wysocka B, Limon J, Jawień A, Banaszkiewicz Z, Janiszewska H, Kowalczyk J, Czudowska D, Scott RJ, Lubiński J. Germline MSH2 and MLH1 mutational spectrum including large rearrangements in HNPCC families from Poland (update study). Clin Genet 2006; 69: 40-47.
- Kurzawski G, Safranow K, Suchy J, Chlubek D, Scott RJ, Lubiński J. Mutation analysis of MLH1 and MSH2 genes performed by denaturing high-performance liquid chromatography. J Biochem Biophys Methods 2002; 51: 89-100.