**Abstract:**

The multiplex endocrine neoplasia 2A (MEN2A) syndrome is a monogenic disease caused by mutation of the RET protooncogene leading to medullary thyroid cancer (MTC), pheochromocytoma and primary hyperparathyroidism. The specific mutations determine the timing of the preventive thyroidectomy. In 2007, in a 26 year old female patient, after detailed endocrine investigation, total thyroidectomy with both sided lateral neck lymph node dissection was performed with suspicion of medullary thyroid cancer. The final histology proved the diagnosis. Three years later hyperventricular peaks occurred, pheochromocytoma was found and removed by laparoscopic adrenalectomy. The family was screened for MEN2A. The mutation TGC634TTC(Cys634Phe) of the codon 634 was found in the patient, in her father and one brother. After spontaneous conception and normal pregnancy in October 2012 she gave birth to a boy. The son was screened for MTC with calcitonin at the age of 11 and 13 months (15.6-16 pg/ml) and was proven to be RET positive. Preventive total thyroidectomy and partial parathyroidectomy was done at the age of 18 months. The final histology found no sign of medullary thyroid cancer or primary hyperparathyroidism. As the patient did not want to have another child affected by the disease, application of assisted reproductive technologies were recommended. After ovarian stimulation, 4 embryos were biopsied in trophectoderm stage. Karyomapping (preimplantational genetic screening for chromosomal abnormalities and preimplantational genetic diagnostics for the specific mutation) was performed revealing one healthy and three MEN2A affected embryos, two of them with maternal aneuploidy as well.

**Conclusion:**

For the monogenic disease affected families, it is important to offer the possibility of preimplantational genetic diagnosis ensuring that the affected embryos are not transferred. Healthy children don’t require lifelong endocrine follow-up and operations, and no psychological support.

**Multiplex endocrine neoplasias (MEN):** 2 or more endocrine tumors occur as a part of a defined MEN syndromes, in a single patient and there is evidence for either a causative mutation or hereditary transmission. Inheritance: autosomal dominant.

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**The family (his)story**

The first patient

The 1st Hungarian karyomapping for MEN2A

**Techniques for preimplantational/gene:**

Compared to other genetic anomaly (CGA) methods for analyzing copy number variations (CNVs) relative to ploidy level in the DNA of a test sample compared to a reference sample. It can identify two germline DNA samples arising from two sources suspected to contain differences in terms of either gains or losses of either whole chromosomes or subchromosomal regions.

**Karyomapping:** uses genome wide linkage-based analysis to determine the inheritance of single-gene defects from as little as a single embryonic cell. It enables all chromosomes to be screened, optimizes the likelihood that embryos with no identifiable genetic abnormalities are transferred during an IVF cycle.

**Results:**

Short GnRH ovarian stimulation protocol was used during which the patient produced 4 eggs. All were fertilized with ICSI (intracytoplasmatic sperm injection). All four eggs reached the trophectoderm stage and were biopsied. Karyomapping was performed revealing three embryos affected by the ret 634 codon mutation and two aneuploid ones. Only one embryo was healthy, which was transferred. Unfortunately, the pregnancy terminated on the 12th gestational week by a spontaneous abortion.

**Conclusions:**

Prenatal diagnostics is only applicable for determination of existing birth defect or inherited disease of the foetus in the early phase of pregnancy.

Preimplantational genetic screening (PGS) and diagnostics (PGD) is used for optimizing the likelihood that embryos with no identifiable genetic abnormalities (chromosomal abnormalities, germline mutations) are transferred during an in vitro fertilization (IVF) cycle.

In our case karyomapping revealed 75% genetically affected embryos, from which 75% were positive for ret 634 codon mutation and 50% were aneuploidy.

Karyomapping allowed us to avoid genetically affected (MEN2A positive 3/ aneuploid 2) embryos to be transferred.

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