



An innovative genetic test to  
identify predisposition to  
producing embryos with  
aneuploidies

[www.fertiladvance.it](http://www.fertiladvance.it)



**FERTILADVANCE**

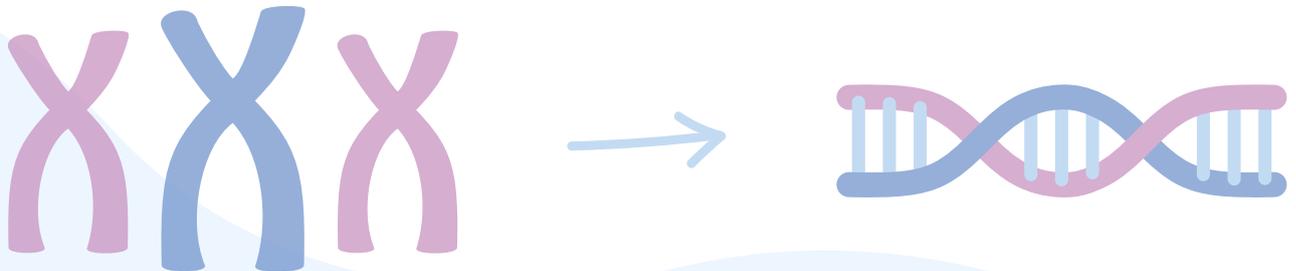
ANEUPLOIDY RISK



## WHY TALK ABOUT ANEUPLOIDIES

**Aneuploidies** are numerical chromosome abnormalities: a cell may carry one extra chromosome (**trisomy**) or one missing chromosome (**monosomy**) compared with the normal chromosomal complement. In reproductive medicine, these abnormalities are among the most frequent causes of **implantation failure**, **early miscarriage**, and, more rarely, **ongoing pregnancies affected by a chromosomal abnormality** up to birth.

A key aspect—often relevant for clinical assessment and for interpreting laboratory findings (including **PGT-A**)—is that aneuploidy can have **different biological origins**, with different implications for pathophysiology and risk.





## ANEUPLOIDIES: MEIOTIC ORIGIN VS MITOTIC ORIGIN

### Aneuploidies of meiotic origin

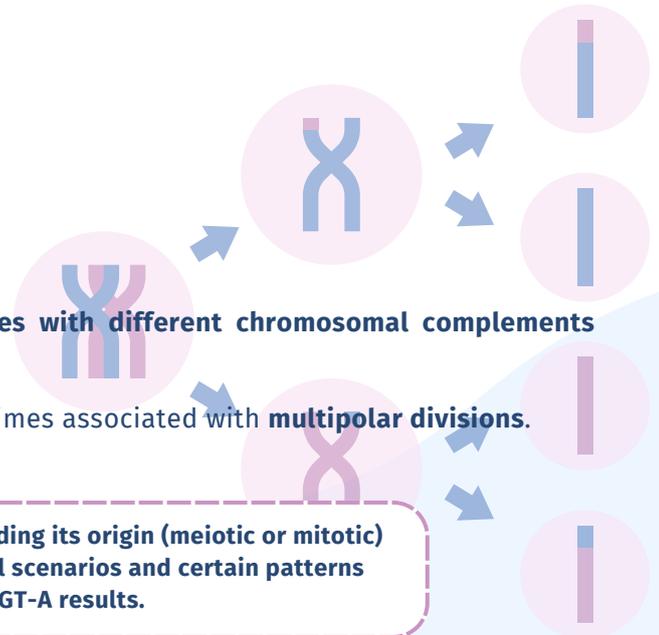
These arise during **meiosis**, the process that leads to the formation of gametes (in particular, the **oocyte**). In humans, most meiotic aneuploidies are of maternal origin and show a well-known relationship with **age**; however, in addition to age, **individual genetic factors** may also contribute.

### Aneuploidies of mitotic (post-zygotic) origin

These occur **after fertilization**, during the very first **mitotic divisions** of the embryo, due to chromosome segregation errors. Such errors may lead to:

- ▶ **embryonic mosaicism (the presence of cell lineages with different chromosomal complements within the same embryo);**
- ▶ **complex patterns** with multiple aneuploidies, sometimes associated with **multipolar divisions**.

Aneuploidy is not a “single” phenomenon. Understanding its origin (meiotic or mitotic) can help specialists better interpret specific clinical scenarios and certain patterns observed in ART pathways and PGT-A results.





# FERTILADVANCE

ANEUPLOIDY RISK

## WHAT IS FERTILADVANCE ANEUPLOIDY RISK

**FERTILADVANCE ANEUPLOIDY RISK** is an advanced genetic test designed to assess predisposition to producing embryos with aneuploidies by analyzing a selected set of **13 genes** associated with an increased likelihood of **meiotic and/or mitotic** aneuploidies.

### Main aims

The purpose of the test is not to “diagnose” an embryonic aneuploidy, but to provide specialists with an **individual risk-stratification tool** useful to:

- complement **reproductive genetic counseling**;
- enable a more informed **interpretation of complex clinical scenarios** (e.g., repeated implantation failures, recurrent pregnancy loss, a higher-than-expected proportion of aneuploid embryos);
- contextualize **results from previous investigations** (e.g., PGT-A patterns suggestive of meiotic or mitotic errors).



## ANALYZED GENES AND BIOLOGICAL RATIONALE

Genes have been identified in which **pathogenic** variants are associated with an increased likelihood of **meiotic** and/or **mitotic** aneuploidies.

The test includes the analysis of the following genes



SMC1B, RNF212, CCNB1IP1  
(HEI10), C14orf39, CCDC66,  
CEP120, AURKB, AURKC, KIF18A,  
KIF20A, TLE6, TP73, PLK4

Overall, the analyzed genes fall into three major functional areas:

- 1 **meiotic cohesion and recombination** (key determinants of non-disjunction risk);
- 2 **segregation and checkpoint control** (relevant in both meiosis and mitosis);
- 3 **centrosome function and spindle dynamics** (relevant to early mitotic errors and complex/mosaic patterns).

1. Carioscia et al. 2026 Common variation in meiosis genes shapes human recombination and aneuploidy Nature 2026 Jan 21. doi: 10.1038/s41586-025-09964-2.  
2. Sun et al. 2023 Identifying risk variants for embryo aneuploidy using ultra-low coverage whole-genome sequencing from preimplantation genetic testing. Am. J. Hum Genet 2023; 110:2092-2102  
3. Tyc et al, 2020 Exome sequencing links CEP120 mutation to maternally derived aneuploid conception risk. Hum Reprod. 2020 9;35:2134-2148.  
4. Nguyen et al. 2017 Identification and characterization of Aurora kinase B and C variants associated with maternal aneuploidy. Mol Hum Reprod 2017 13;23:406-416.  
5. Biswas, et al. 2024 Maternal genetic variants in kinesin motor domains prematurely increase egg aneuploidy, Proc. Natl. Acad. Sci. U.S.A.(2024) 121 (45) e2414963121.  
6. Lledo et al. 2023 Identification of novel candidate genes associated with meiotic aneuploidy in human embryos by whole-exome sequencing. J Assist Reprod Genet . 2023;40:1755-1763.  
7. McCoy et al. 2015 Common variants spanning PLK4 are associated with mitotic-origin aneuploidy in human embryos. Science. 2015 10;348:235-238.



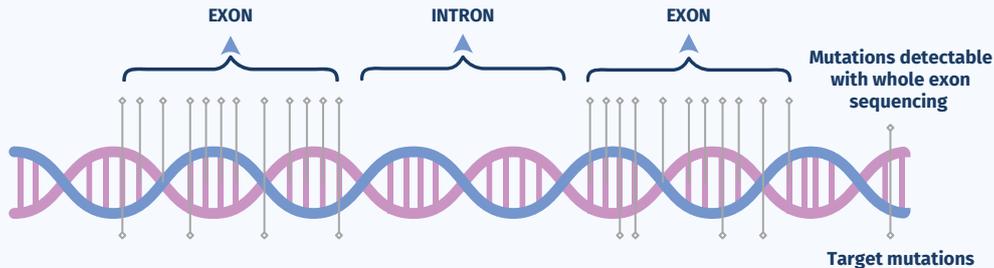
# FERTILADVANCE ANEUPLOIDY RISK

## TECHNOLOGY: HIGH RESOLUTION AND ADVANCED BIOINFORMATICS



**FERTILADVANCE ANEUPLOIDY RISK** uses next-generation sequencing (NGS) **technology** with sequencing of the entire coding region (exons) of the investigated genes and an advanced bioinformatics pipeline.

**Advantage over “target” tests:** it does not search only for selected mutations, but enables identification of **any variant** within the sequenced regions.





## TEST RESULTS



### Positive: pathogenic variant detected

The presence of rare variants with plausible functional impact in genes involved in meiosis/mitosis may indicate a **predisposition to producing embryos with aneuploidies**. Interpretation should always be integrated with clinical data and, when appropriate, with genetic counseling.



### Negative: no pathogenic variant detected

No variants associated with an increased risk of producing embryos with chromosomal aneuploidies are identified. A negative result **does not exclude** aneuploidy risk, as the phenomenon is multifactorial and depends also on maternal age and other biological variables not investigated by the test.



## TEST INDICATIONS

The test may be considered when investigating a possible genetic contribution to an increased aneuploidy risk in scenarios such as:

- ▶ **repeated ART/IVF cycles** with a high proportion of aneuploid embryos, especially if unexpected for maternal age;
- ▶ **recurrent implantation failure** or **recurrent pregnancy loss**, particularly when there are data suggestive of a chromosomal origin;
- ▶ a previous conception/pregnancy affected by **chromosomal aneuploidy**;
- ▶ the need for a **preconception** or **pre-IVF** assessment to clarify possible genetic contributors;
- ▶ the presence of patterns suggestive of **mitotic errors** (e.g., “chaotic” patterns, suspected tripolarity) or **meiotic errors** (recurrent aneuploidies consistent with non-disjunction).



# FERTILADVANCE

ANEUPLOIDY RISK

A SIMPLE, STEP-BY-STEP WORKFLOW



**Kit  
request**



**Completion of kit  
documentation**



**Sample  
collection**



**Sample  
shipment**



**Report  
delivery**

**Turnaround time**



**15 days**

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## LABORATORIES

Rome: Via Arduino 38 - 00162 Tel.: 06.21115020  
E-mail: [info@genomicalab.it](mailto:info@genomicalab.it)  
[www.genomicalab.it](http://www.genomicalab.it)

## REGISTERED OFFICE

Rome: Via Arduino 38 - 00162  
PEC: [info@pec.genomicalab.it](mailto:info@pec.genomicalab.it)  
VAT No.: 14554101007 - REA: RM - 1530210

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