

The Madness of Genetics: On “DNA Contamination” & the Bizarre Claim That “Specific” DNA Can Precisely Change Human Genetics and Cause Cancer

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[DNA Contamination: The Amusing Irony and Theoretical Trap](#)

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It is ironic that if the assumptions of genetic theory were correct, critics would be right. This irony shows that pharmaceuticals are paradoxically finding themselves caught in the crossfire due to theses held by both critics and the mainstream.

What does that mean?

The idea that DNA is the unchanging blueprint of life has been promoted for over 40 years. But in 2006, leading researchers confirmed: This idea was naive. DNA is dynamic and constantly changing.

The human genome

Complete decoding of the human genome (in reality just reading rows of letters that are not understood) is impossible. No two people are “genetically” alike, not even one cell is another. When you consider that the individual genomes of less than 0.0003% of the 8 billion people have been sequenced (and incompletely), one thing becomes clear: assignments of genetic letter orders for cancer, eye color, height, etc. are impossible.

An example: the color of the eyes

It was believed that it was simply genetically determined. This idea turned out to be a forced interpretation of the current data, which was refuted by more recent data. So parents with blue eyes can have a child with brown eyes. This makes it clear that the idea of a fixed genetic section has been scientifically refuted.

Plain language: If every genome is unique, we only know a tiny fraction of all genomes (0.0003%) and they are constantly changing themselves, then the attempt to define fixed gene sections for certain characteristics such as eye color or cancer is worth it , like trying to catch a river with a sieve

– it's not only inaccurate, it's simply unattainable.

The madness of genetics

When even simple examples such as eye color cannot be assigned, it seems bizarre when parties claim that foreign “specific” DNA can precisely change human genetics and cause cancer. Dr. Kary Mullis summed it up well: With PCR you can find everything in every person if you do it right. This means that every DNA or RNA sequence can be found in every person, whether healthy, sick, vaccinated or unvaccinated.

The strategy of fighting like with like may seem promising at first glance. But on closer inspection it turns out to be a theoretical trap. Instead of focusing on the DNA-contaminated mRNA vaccines, the central question should be: Has the basic assumption about “pathogenic particles (viruses)” ever been scientifically proven/confirmed? The answer is NO , and therefore every vaccination is inherently illegal and dangerous.

The current approach offers the pharmaceutical apparatus a back door: they could argue that there are better, uncontaminated vaccines or present other methods that would then have to be accepted within the long-refuted genetics thesis.

The technical implementation of the laboratory in Magdeburg is vulnerable even within the narrative

The detection is based on non-exact methods such as Qubit Flex Fluorometer and qPCR that have been performed. Neither the plasmids were directly isolated and detected nor were they properly sequenced.

- First, we lack detailed information about the exact conditions and protocols under which these experiments were conducted.
- Secondly, we know nothing about the negative and

positive controls carried out.

- Third, no sequencing or isolation of the plasmids was performed. This means that we do not know exactly which DNA sequences were present in the samples. Sequencing would have been necessary to determine the “theoretically” accurate nature of the DNA detected and to ensure that it was in fact plasmid DNA and not other forms of DNA.

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