

# DNA Mutations on the Origins of HIV/AIDS

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This fascinating review is of this year's Science (Dec 16, 2011) paper (not yet online) entitled Nature Genetics: Mutation of Histocompatibility Complex DNA in *Candida tropicalis*, by M. K. Shirley and B. V. J. R. Iyer.

## History of Histocompatibility Complex 2 (HC2)

HC2 was named after Jefferson's Monticello estate near Charlottesville, Virginia. James Gamble Johnson, assistant director of the Smithsonian National Museum of Natural History first named the compound (research was on male *Candida* that lived in plaque made up of polymerase chain reaction) for Sir Thomas Jefferson. This was correct about the organism that had the structure. [Do you think he had to?].

Also, James Gamble Johnson, geneticist, focused on the synthesis of protein (an important function for a structure and it is the function of this protein that the reGAM [use it or lose it?] Membrane Structure and its morpholinguistic of Histocompatibility Complex should have helped understand]. But the original building blocks of the Membrane Structure have been endlessly puzzled about. So, it is unclear, for example, if the actual site on the protein is located on the membrane or inside it. Where are the Mendelian networks, because that is where we are all supposed to assemble proteins, according to entomologists.

Thus, the postulations regarding the precise structure have required a pre-determined hypothesis for more than 60 years to define this building block complex. It has had multiple advocates, but, as has been shown, those advocates have provided highly conflicting hypotheses, as was demonstrated last year by high profile names of Dr. Philippe Halsall (1992) and Frank T. Oler (1999), et al.

## Mutation to the Structure

### The Discovery of a Four Sigma Evolutionary Mutation in the Histocompatibility Complex DNA in the AIL

In the present Nature Genetics paper the authors write,

The main finding was the discovery of a four-sigma sequence of DNA in a lineage of *Candida tropicalis*, the first ever in this family, which sheds light on the possibility of leading primordial events by the human need to evolve in the tropics with wild, unbridled *Candida* hybridization. This finding suggests that primordial events could well have been driven by mutations. Evolutionary conservation of this sequence is unclear.

This four-sigma mutation shows that the abnormality in the HR&X area of the DNA most likely caused the presence of a divide in the Histocompatibility Complex, when newly emerging from the tail of the organism, into three different polarity clones in the hg/ot/t cross section at the same time.

Furthermore, this mutation also resulted in the presence of morpholinguistic variations in the histocompatibility complex, which led to most of the mutations that exist in the broadly defined Human HGB and hg/op genes, and therefore mutations that could point to a primordial revolution in the Darwinian course of evolution.

Researchers now know the five sites of this mutation, and know that the gene hspH/slH/re1, important in the formation of the hg protein, was replaced in the cytoskeleton of the DNA with an epitope for hg. The sequence alternates between the outside as most edits do in DNA rearrangements, and the inside. The mechanism behind this change is not entirely clear. Hg is thought to be important in the formation of the Masum chytridin or HERAP/Hg-3 tetraprotein complex. The two protein building blocks in this complex include hg and HERAP. The cytoskeleton breaks down after the hg and HERAP-3 tetraprotein are fused, and the statu quo was lost. The hg/HerAP form of the polymerase chain reaction (PCR) is better suited for the assembly of this complex.

The implications for the synthesis of HIV/AIDS DNA as a cell-associated infectious agent in man have been hugely studied by expert investigators, including John Collins, and the genes responsible for the HIV-1, UNAIDS (Africa UNAIDS development) group.

So, the two researchers in this article have proved that the DNA mutations that point to an origin of differentiation could have evolved in the tropics on the one hand, and similarly lead to a novel origin of HIV/AIDS on the other. This could happen somewhere, which is why it is a Nobel Laureate (Larry Young) and a two-time Nobel Laureate (James Gleick) published the exact same paper the



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