

What Does the MCRA Really Mean?

By Roberta Estes, copyright 2011

The MCRA or time to the Most Common Recent Ancestor is a calculation provided by both Family Tree DNA and Ancestry.com for their clients who have taken the Y-line DNA tests. I have a great example of how this actually translates into reality.

Often, I receive communications from people who say something like this:

"It says that I'm related to John Doe within 6 generations. I have both of your genealogy to 6 generations, and I can't find our common ancestor. What is wrong?"

The answer to "What is wrong?" is easy. The person doesn't understand what the tool that estimates MCRA is telling them. And, I'm betting they didn't read the instructions and explanations either, that is if they tested at Family Tree DNA who provides such.

Family Tree DNA provides a great deal more information and a far more robust tool than Ancestry. Family Tree DNA begins with this information:

"The probability that **John Doe** and **William Doe** shared a common ancestor within the last..."

COMPARISON CHART	
Generations	Percentage
4	10.75%
8	50.89%
12	82.16%
16	95.03%
20	98.83%
24	99.76%
28	99.95%

You can also change the generational display. I change mine to "every generation."

Display

* The FTDNATIP™ results are based on the data presented during the 1st International Conference on Genetic Genealogy, on Oct. 30, 2004. The above probabilities take into consideration the mutation rates for each individual marker being compared.

This is followed by an explanation and instructions for how to refine the calculations:

Refine your results with paper trail input

However, these results can be refined if their paper trail indicates that no common ancestor between **John Doe** and **William Doe** could have lived in a certain number of past generations.

If you don't know this information for a fact, do not change the "1" in the box in the next paragraph. However, if you have the information, please enter in the box and click on the recalculate button.

John Doe and **William Doe** did not share a common ancestor more recently than generation(s). (Because the important factor in calculating the time to the Most Recent Common Ancestor is the number of generations between which mutations could take place, the number of years per generation is irrelevant in FTDNATiP™ calculations).

After that, additional explanation and a reference to a FAQ sheet:

* The FTDNATiP™ results are based on the mutation rate study presented during the 1st International Conference on Genetic Genealogy, on Oct. 30, 2004. The above probabilities take into consideration the mutation rates for each individual marker being compared.

Since each marker has a different mutation rate, identical Genetic Distances will not necessarily yield the same probabilities. In other words, even though **John Doe** has a Genetic Distance‡ of 4 from **William Doe**, someone else with the same Genetic Distance may have different probabilities, because the distance of 4 was prompted by mutations in different markers, with different mutation rates.

‡Note: The Genetic Distance is the count of the total difference between two individuals. For example, if a marker differs by 2, then the Genetic Distance will count this as a distance of 2.

More questions? Please refer to the [FTDNATiP™ FAQ page](#).

This is a huge difference compared to Ancestry who only gives you a number with absolutely no explanation at all:

The MCRA is the small number beside the name - so John Doe is an MCRA of 2 and William is 14. I have highlighted these in red below so that you can see them.

Here is the explanation the Ancestry which is followed by the match table.

"You could be close to a meaningful family connection! The list below is sorted by how close your DNA matches (MRCA). The closest matches are at the top."

100% sure number, not their "most likely" number. In fact, if we divide their number in half, it's still high.

We've learned that Family Tree DNA's 50th percentile is much closer to reality, without any tweaking that you can do based on known pedigree charts.