



LEGEND

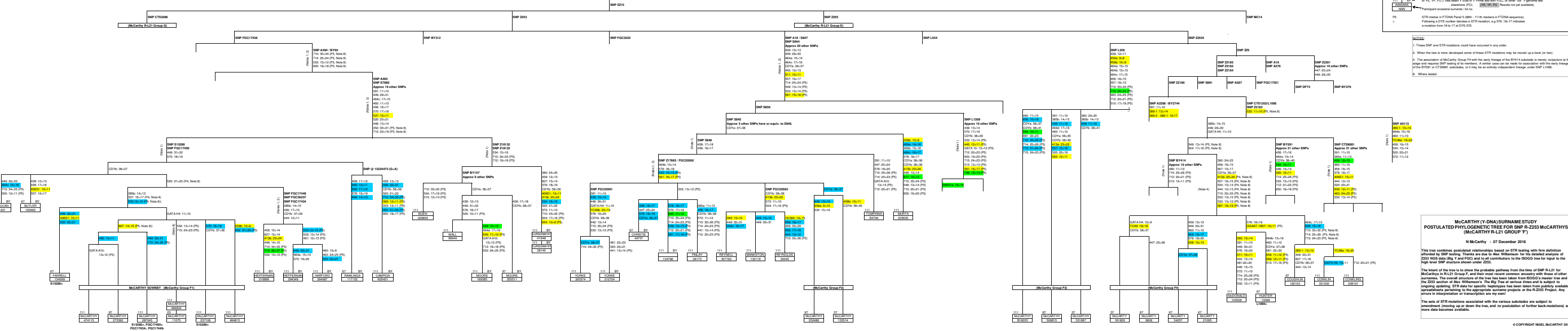
STR Mutation:

- "Very slow moving" STR markers (very low mutation rates) or rare multi-step mutation
- "Slow moving" STR markers (low mutation rates)
- Postulated STR back-mutation

Participant kit number and testing details:

- No. of STR markers for which results have been published
- Has taken DYS test (Y) (Results not yet available)
- or YE, VP, YD - Has taken Y-E test with FGC, or other "full" Y-gonome test elsewhere (FC) (YE, YP, YD) (Results not yet available)
- Participant ancestral surname / kit no.

PS
> STR marker in FTDNA Panel 5 (889 - 111b markers in FTDNA sequence).
Following a DYS number denotes a STR mutation, e.g. STR: 18-17 indicates a mutation from 18 to 17 at DYS 18.



- NOTES**
- These SNP and STR mutations could have occurred in any order.
 - When the tree is more developed some of these STR mutations may be moved up a level (or two).
 - The association of McCarthy Group F4 with the early lineage of the BY414 subclade is merely conjecture at this stage and requires SNP testing of its members. A similar case can be made for association with the early lineage of the BY591 or CTS5881 subclades, or it may be an entirely independent lineage under SNP L1066.
 - Where tested

McCarthy (Y-DNA) SURNAME STUDY
POSTULATED PHYLOGENETIC TREE FOR SNP R-Z253 McCARTHYS
(McCarthy R-L21 GROUP F)

N McCarthy - 07 December 2016

This tree combines postulated relationships based on STR testing with firm definition afforded by SNP testing. Thanks are due to Alex Williamson for his detailed analysis of Z253 YGS data (Big Y and FGC) and to all contributors to the ISOGG tree for input to the high level SNP structures shown under Z253.

The intent of the tree is to show the probable pathway from the time of SNP R-L21 for McCarthys in R-L21 Group F, and their most recent common ancestor with those of other surnames. The overall structure of the tree has been taken from ISOGG's master tree and the Z253 section of Alex Williamson's 'The Big Tree' at various times and is subject to ongoing updating. STR data for specific haplotypes has been taken from publicly available spreadsheet pertaining to the appropriate surname projects or the R-Z253 Project. Any errors in interpretation or transcription are my own.

The sets of STR mutations associated with the various subclades are subject to amendment (moving up or down the tree, and/or inclusion of further back-mutations) as more data becomes available.