

STR LINEAGE PLACEMENT PROTOCOLS

[What to do if the Y-DNA Utility creates a lineage comprised only of individuals that have only 12 markers tested?](#)

Individuals who match at 12 loci but have no relationship to other individuals who have tested for 37 or more loci will not be used as a lineage but will remain in the ungrouped category until further testing occurs. Those individuals may not be related at all due to the small number of loci tested and the fact that many surnames will match at 12 loci including hundreds of individuals. They will be encouraged to have more loci tested. If a 12 marker kit has a 12/12 match with one lineage and NOT to any other lineages we will put it in, but recognize that it is tentative placement pending further results.

[What to do if the individuals of a lineage have a non-Crawford surname and show no paternal ancestor as Crawford.](#)

Individuals falling into this category will be contacted to determine why they joined the Crawford project and to attempt to discern their connections, if any, to the Crawford surname. If there are people who are not Crawford on the Y line, but who wish to be part of the project, they will remain in the database but they will be de-listed from the Y reports and placed in a zzz category. This will drop them from the Ungrouped page and any lineages. FTDNA has a setting allowing administrators to not display their results in the Y pages. If at any time these individuals establish a link to the Crawford surname they will be placed back on the active list.

[What date should be used on the lineage when posting to the web?](#)

The date when the lineage was updated in the EXCEL file will be the date used when posting on the web. It is understood that this date will not coincide with the GAP download date which may be earlier and it will not coincide with the actual date that the HTML file is posted to the world wide web which may be later.

[What is the protocol for posting the MODE?](#)

Mathematically, the mode does not exist for 2 kits that differ by one, but for the sake of consistency the mode will be posted for all project lineages in a uniform manner except for ungrouped categories.

[How will lineages be named?](#)

Lineages will be clustered under their major haplogroups I, R, J etc. Under those haplogroups the next layer will be clustered under the clades. Lineages will be numbered numerically with the haplogroups being placed in alphabetical order. All sub-lineages will be designated by alphabetical listing in descending order.