

Footnotes

A020, A031 and A040 This family has a documented genealogy to the Presidential Adams Family

H* Haplogroup: All values in red are estimated by FTDNA or using <https://home.comcast.net/~whitthey/predictorinstr.htm>
 A005 is tested by FTDNA as SRY10831.2- which means R1a haplotype is confirmed.
 A025 is tested by FTDNA as P19+ confirmed I; M253+ confirmed I1a
 A027 is tested by FTDNA as M172+ confirmed J2
 A032 is tested by DNA Heritage confirmed R1b
 A040 is tested by FTDNA as P310+ L51+ U106- P312- L257-
 A070 is tested by FTDNA as M172+ M304+ M102- M12- M137- M158- M163- M166- M205- M267- M280- M339- M365- M367- M368- M369- M390- M62- M67- M68- M92- M99-
 A094 is tested by FTDNA as P25+
 A104 is tested by FTDNA as M172+ confirmed J2
 A111 is tested by FTDNA as M172+ confirmed J2
 A112 is tested by FTDNA as M253+
 A114 is tested by FTDNA as M17+M198+
 A132 is tested by FTDNA as M170+ M253+ M258+ M307+ P19+ P30+ P38+ M161- M21- M223- M227- M26- M72- P259- P37.2-
 A134 is tested by FTDNA as M253+
 A139 is tested by FTDNA as M172+
 A144 is tested by FTDNA as M172+ M304+ M102- M12- M137- M158- M163- M166- M205- M267- M280- M339- M365- M367- M368- M369- M390- M47- M62- M67- M68- M92- M99-
 A152 as tested by FTDNA as L130- L130- L144- L144- L159.2+ L159.2+ L192- L192- L193- L193- L195- L195- L226- L96- L96- M126- M153- M160- M173+ M18- M207+ M222- M269+ M343+ M37- M65- M73- P107- P25+ P66- SRY2627- U106- U152- U198-
 A167 as tested by FTDNA is L257-, L51+, P310+, P312-, U106- R1b1a2a1a1b3 (R-P310)

A202 is tested by FTDNA as M253+
 A210 is tested by FTDNA as M172+
 A239 is tested by FTDNA as L1-U106+
 A278 is tested by FTDNA as L21+ P66- M37- M222- [R-L21]
 A282 is tested by FTDNA as U152+ P312+ SRY2627- M222- M160- M153- M126- L4- L21- L20- L2- [R-U152*]
 A296 is tested(estimated?) by Ancestry.com as J2a1b

*** [Matching Niall Níogiallach - Niall of the Nine Hostages \(Click for link to FTDNA page\)](#)
 ??? An estimated haplogroup, the haplotype was extracted from <http://SMGF.org> database

This Haplogroup prediction is unknown- this may be part of a newly found branch of the I group
 * These results extracted from www.ysearch.org database
 + DYS389b = DYS389i - DYS389i

A072 This is YGGAAT1B07
 \$ The values of CDYa and CDYb have been reversed to clearly show the proper number of mutations.
 Y-TAGA H4 from Relative Genetics is NOT counted the same way as Y-GATA from FTDNA. Subtract one from the RG or DNAHeritage value to compare.

463 The values in the table have been adjusted to report the same as FTDNA.
 X Reported in with the NIST convention. Which is DNAH+2

Missing Data from the DNA test.
464 nomenclature is POST the May 2003 change
Markers with names with a RED background may mutate slightly faster [0.008 vs average 0.0025]
Markers with names with a Green background may mutate slightly slower [0.001 vs average 0.0025]

Color code for haplogroups other than R1b: Green is modal, Yellow one step from modal, Blue two steps....
 The lines that have more green values are closer to the modal value for that haplogroup.
 Results as posted are from three companies, thus there are some blank cells in the table because there are no results for that allele from the test.
 The numbers in the bottom part of the R1b section give the allele values for R1b haplotype in the order of most likely to least likely.
 For example DYS385a has a most likely value of 11 for R1b (and most of the Adams men had 11) and a second most likely value of 12 and it is in orange.
 Orange indicates that this value occurs in 1% to 10% of the R1b haplotypes.