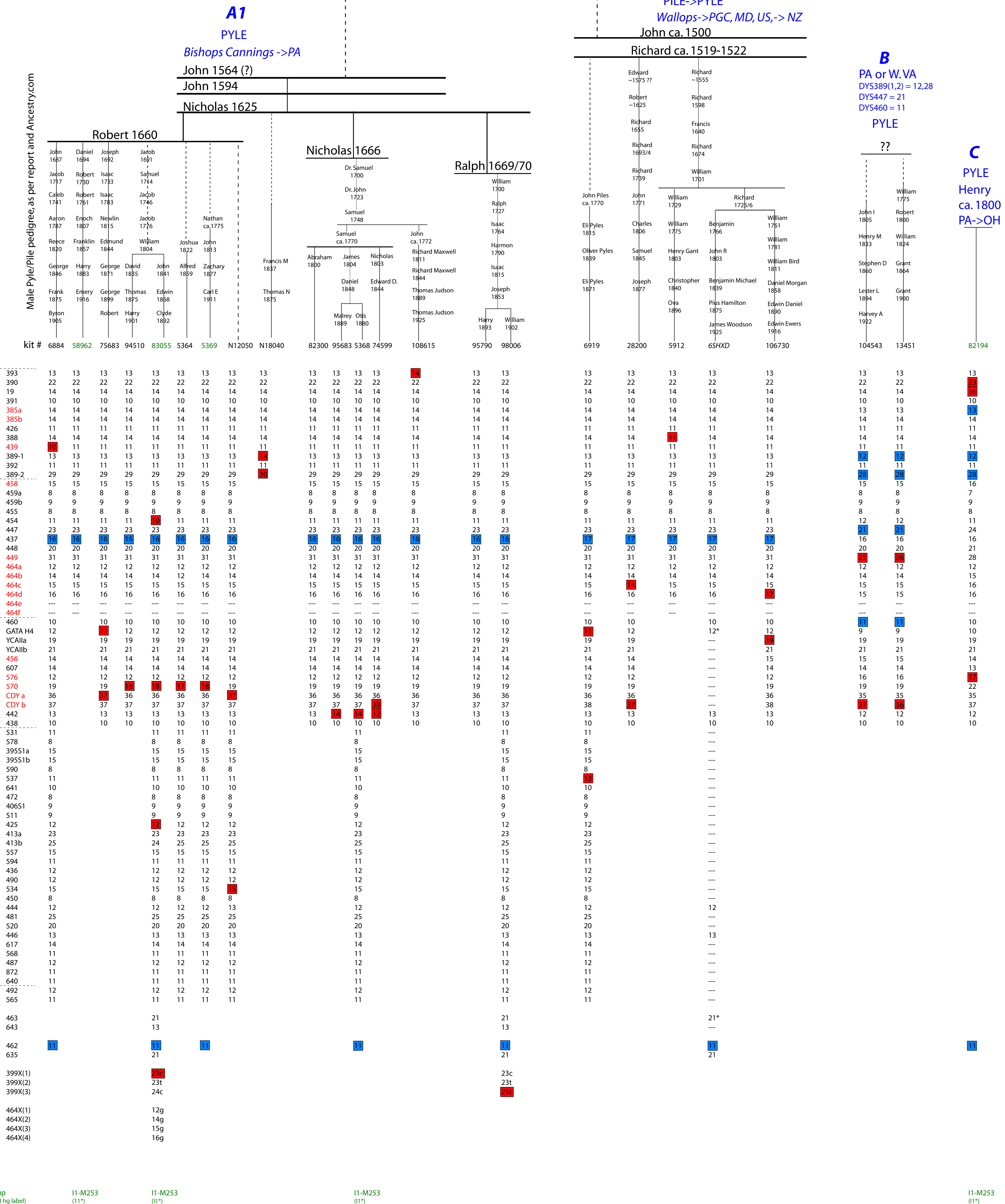


Combined Pedigree and Y-DNA haplotypes for 37 members as of 14 July 2008 (JMP)

"Proven" paper trail (solid line)
speculative/unproven (dashed line)

A Bishops Cannings-Wallops PYLE Supergroup (split via DYS437; 16 (BC) vs. 17 (Wallops))



D PYLES/PILES ENG-->PGC MD

E Samuel J. PYLES --> WHITE--> SKIDMORE

Note: groups in both I1b and R1b appear to claim descent from Francis Pile(s) born in Prince Georges County, MD in 1660. The DNA results tell us that one of these groups can not be descended from Francis, since one group is I1b, and one is R1b

Within-haplogroup off-modal/differencing values
"uniting" values common to a particular sub-grouping

F PILE/POILE/POYLE Conrod 1745 (?) (Poyle)

G PILES/PILES PGC, MD Francis 1660

H PYLES Francis 1635 (?) --> Zacariah 1770 PGC, MD--> VA

I PYLES Vincent Godfrey (ca 1700) (?) --> Conradus (VA)

J PILES->PILES John Piles (b. ca 1800, NC)

K PILES->PILES Joshua Pyles/Piles (b. ca. 1790, PA)

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Note: Henderson Pile has not been definitively linked to Coonrod (Conrad) Pile by a paper trail. But, the early 19th-century geographical proximity of Henderson to known Coonrod descendants, and the resulting 24/25 match (with the 1 mismatch likely due to a reClOH event) virtually assures that 7139 and 31495 are related - if not with Coonrod as the MRCA, then someone further back (TMRCA = 10 gen @50% prob)

O** = null 439, likely caused by L1/S26 SNP

I1a BC modal (37): 13 22 14 10 14 14 11 11 13 11 28 15 8 9 11 23 16 20 31 12 14 15 16 10 12 19 21 14 14 12 19 36 13 10
I1a Wallops modal (37): 13 22 14 10 14 14 11 11 13 11 28 15 8 9 11 23 16 20 31 12 14 15 16 10 12 19 21 14 14 12 19 36 13 10
Green kit # = SNP-tested haplogroup
Italic kit # - transferred from DNA Heritage (*: alleles converted from DNAH standard to FTDNA standard: GATA H4(FTDNA) = TAGA H4(DNAH)-1; DYS463 (FTDNA) = DYS46(DNAH)+2
kit xxxxxx # - transferred from Ancestry.com (S: allele converted from SMGF standard (2006 ISFG to FTDNA standard: Y GATA H4.1(SMGF) = Y GATA H4 (FTDNA) + 11

DYS 439 Allele value in FTDNA order, markers #1-67, plus selected advanced markers. Red = fast mutator

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