

Scudamore/
Skidmore Family
Genetics Study
Phase III

October 20, 2006

Relative Genetics

2495 S. West Temple
Salt Lake City, UT 84606
Phone: (801) 462-1427
1-800-956-9362

At A Glance: Scudamore/Skidmore Family Genetics Project, Phase III

Individuals in a family study who share 26 of 26, or 43 of 43 alleles are considered a unique haplotype, a genetic term for group. Each haplotype is assigned a number. Lineages in a Family Study are designated with a letter and are defined by haplotypes containing 2 or more individuals, and/or haplotypes that share at least 23 of 26, or 38 of 43 alleles. For example, in Skidmore Lineage A there are 10 individuals in 7 haplotypes. Those individuals in ***bold italic*** type had 43 markers tested.

Skidmore Lineage A-

Earliest Documented Paternal Progenitor: Henry Skidmore, baptized in 1575 in
 Rickmansworth, Hertfordshire

Defined By:

- Skidmore Lineage A Haplotype 1 (A1):** 7293, 8104, 23994
- Skidmore Lineage A Haplotype 2 (A2):** 8134
- Skidmore Lineage A Haplotype 3 (A3):** 7558, 7627
- Skidmore Lineage A Haplotype 4 (A4):** 6686
- Skidmore Lineage A Haplotype 5 (A5):** 7789
- Skidmore Lineage A Haplotype 6 (A6):** 7983
- Skidmore Lineage A Haplotype 7 (A7):** 7292, 68304

Skidmore Lineage B-

Earliest Documented Paternal Progenitor: Richard Skydmore of Mayshill, Westerleigh,
 born about 1480

Defined By:

- Skidmore Lineage B Haplotype 1 (B1):** 6940, 7091, 7290, 7294, 21678, 40475, 41844,
 Z056842
- Skidmore Lineage B Haplotype 2 (B2):** 69496
- Skidmore Lineage B Haplotype 2 (B3):** Z097975

Skidmore Lineage C-

Earliest Documented Paternal Progenitor: William Skidmore collier of Kingswinford,
 born about 1590

Defined By:

- Skidmore Lineage C Haplotype 1 (C1):** 18588, 23755, 29534
- Skidmore Lineage C Haplotype 2 (C2):** 7480, 12148

Skidmore Lineage D-

Earliest Documented Paternal Progenitor: William Scudamore, married in 1708

Defined By:

- Skidmore Lineage D Haplotype 1 (D1):** 40454
- Skidmore Lineage D Haplotype 2 (D2):** 22920

Skidmore Lineage E-

Earliest Documented Paternal Progenitor: John Skydemor, of Dunkerton, alive in 1441

Defined By:

- Skidmore Lineage E Haplotype 1 (E1):** 6646, 16382, 57260
- Skidmore Lineage E Haplotype 2 (E2):** 35519
- Skidmore Lineage E Haplotype 2 (E4):** 10508

**At A Glance: Scudamore/Skidmore Family Genetics Project, Phase III
cont.**

Skidmore Lineage F-

Defined By:

- Skidmore Lineage F Haplotype 1 (F1):** 70418
- Skidmore Lineage F Haplotype 2 (F2):** 58854

Skidmore Lineage G-

Defined By:

- Skidmore Lineage G Haplotype 1 (G1):** Z097395, 63912, 68685, 68781
- Skidmore Lineage G Haplotype 2 (G2):** Z097979
- Skidmore Lineage G Haplotype 3 (G3):** 43557
- Skidmore Lineage G Haplotype 4 (G4):** 57575
- Skidmore Lineage G Haplotype 5 (G5):** 62028

Skidmore Lineage H-

Defined By:

- Skidmore Lineage H Haplotype 1 (H1):** 8894
- Skidmore Lineage H Haplotype 2 (H2):** 57918

Skidmore Independent Lineages - 6701, 7291, 20071, 36639, 41512, 41662,

Skidmore Family Genetics Project Phase III

20 October 2006

Phase III Participants:

ID: Participant:

43557 Mark Skidmore
57260 John A. Skidmore
57575 Dr. Dwight V. Skidmore
57918 John Robert Skidmore
58854 Dr. Peter Skidmore
62028 John Steven Edgar

ID: Participant:

63912 Walter R. Skidmore
68304 Robert M. Skidmore
68685 Jack Skidmore
68781 John Skidmore
69496 Michael George Skidmore
70418 Dr. John F. Skidmore

ID: Participant:

Z056842 Bill Skidmore
Z097395 Milus Skidmore
Z097975 Robert Emerson Skidmore
Z097979 James Tennessee Burke

Phase II Participants:

ID: Participant:

10508 Lee Skidmore
12148 David. M. Skidmore
16382 Tim Skidmore
18588 James Skidmore
20071 Alan Skidmore
21678 Frank G. Skidmore

ID: Participant:

22920 Gavin Skidmore
23755 Richard Charles Skidmore
23994 Charles Skidmore
29534 Robert Skidmore
35519 Peter Charles Scudamore
36639 Ernest Roy Skidmore

ID: Participant:

40454 Andrew Skidmore
40475 Dennis Skidmore
41512 Matthew Scudamore
41662 Lamoin Edward Skidmore
41844 Brian Skidmore

Phase I Participants:

ID: Participant:

6646 Ed Skidmore
6686 Wm. Frank Skidmore
6701 Ian Skidmore
6940 Warren Skidmore
7091 Howard Skidmore
7290 Howard H. Skidmore

ID: Participant:

7291 Robert Skidmore
7292 Wade Skidmore
7293 Billy Skidmore
7294 Ellis Dee Skidmore
7480 Ewart Burt Skidmore
7558 Gene Aaron Skidmore

ID: Participant:

7627 James Edgar Skidmore
7789 Eric Skidmore
7983 George Skidmore
8104 Barry Hastings Skidmore
8134 Ron Skidmore
8894 Leonard John Scudamore

Summary of Phase II Findings:

Of the 35 individuals participating in the Skidmore Family Genetics Project Phase II, there were five main lineages identified, and seven independent lineages.

Objectives of the Scudamore/Skidmore Family Genetics Project, Phase III:

The objectives of this study are to further elucidate the genealogies of various Skidmore individuals to determine if common ancestry can be established.

Summary of Phase III Findings:

With the 16 additional individuals participating in the Scudamore/Skidmore Family Genetics Project Phase III, there were eight main lineages identified (Figure 1a, 1b, and Appendixes A1-A3). A discussion of each Lineage follows. Hereafter in the text, those who had 43 markers tested will be in italic type.

Note: Some analyses in this report are based on genetic data that were not generated by Relative Genetics. Relative Genetics cannot verify the accuracy of those data.

Scudamore/Skidmore Lineage A, The Rickmansworth Line:

□ After Phase II testing, Scudamore/Skidmore Lineage A was defined by ten individuals in seven haplotypes. Phase III testing added one individual, *68304*, Robert M. Skidmore, for whom no genealogy was submitted. However, he is a 43/43 match with *7292* who defines Skidmore Lineage A7. As was described in the Phase II report, the earliest genealogical point that the DYS442 mutation that defines lineage A7 could have occurred would have been between Nathaniel Skidmore of Morgan Co., AL, born in 1842, and his father, Captain William Skidmore of Madison County, KY, born in 1799. Therefore, the genetic evidence strongly asserts that *68304*, like *7292*, is a descendent of Nathaniel Skidmore (b. 1842).

Scudamore/Skidmore Lineage B, The Westerleigh line:

□ In the Phase II report, Skidmore Lineage B was defined by seven individuals in one haplotype. Phase III testing has identified three new individuals who are also part of Skidmore Lineage B (Figure 1a). The first participant *Z056842*, Bill Skidmore, who was tested by another laboratory, shares descendency with *7091* and *41844* in Timothy Skidmore of Suffolk Co., LI, born in 1760, a tenth generation descendent of Richard Skydmore (b. abt. 1480). The genetic data confirm his connection to this line as he is part of Skidmore Lineage B1.

□ The second new participant is *69496*, Michael George Skidmore, who shares descendency with *6940*, *7294*, and *40475* in Joseph Skidmore of Little York, DE, born in 1706, an eighth generation descendent of Richard Skydmore (b. abt. 1480). *69496* forms a new genetic lineage, Skidmore Lineage B2 as he has a different value at DYS635 than the others in Lineage B. Though the MRCA would place the common ancestor for *69496* and the other descendents of Joseph Skidmore (b. 1706) at 13 generations, their stated relationship of 6-7 generations is within the 95% Confidence interval of 2-38 generations, and is likely correct. However, further confirmation could be achieved if any or all of the other tested descendents of Joseph (b. 1706), namely *6940*, *7294*, or *40475* upgraded to 43 markers. If there was only one mismatch after testing 43 markers, then the MRCA would be 8 generations, which would directly correlate with the current genealogical data.

□ The third new participant is *Z097975*, Robert Emmerson Skidmore who was tested by another laboratory, and claims shared descendency with *6940*, *7294*, *40475*, and *69496* in Joseph Skidmore

Scudamore/Skidmore Lineage B, cont.:

of Little York, DE, born in 1706, an eighth generation descendent of Richard Skydmore (b. abt. 1480). Z097975 forms a new genetic lineage, Skidmore Lineage B3 as he has different values at DYS442 and DYS464c than 69496. 66496 is the only participant in Lineage B who had more than 26 markers tested. These two individuals have a total of 32 tested markers in common, and they share 30 of those 32 markers. The MRCA value is 16 generations with a 95% Confidence Interval of 5 to 42 generations, which does not correspond to the current genealogical findings of 6-7 generations. Because only two individuals have been tested at DYS442 and DYS464c, the ancestral type, or the profile of the common ancestor, cannot be determined. Therefore, the upgrades suggested in the previous paragraph are highly recommended in order to reconcile the relationship of Z097975 to 69496 as well as to the rest of the group.

Scudamore/Skidmore Lineage C, The Kingswinford Line:

□ Skidmore Lineage C does not have any changes from the Phase II report.

Scudamore/Skidmore Lineage D:

□ Skidmore Lineage D does not have any changes from the Phase II report.

Scudamore/Skidmore Lineage E:

□ In the Phase II report, Skidmore Lineage E was defined by five individuals in four haplotypes. After one upgrade, one addition, and two deletions, Phase III testing shows five individuals in three haplotypes. The new participant is 57260, John A. Skidmore, a descendent of Arthur Skydmore of Eyam, Derbyshire, born before 1629. And, as was suggested in Phase II, 6646 upgraded to 43 markers. The 43 marker profiles of 6646 and 57260 likely represent the DNA profile of the common ancestor for Skidmore Lineage E.

□ 6701, Ian Skidmore who had previously claimed descendency from William Skidmore collier of Kingswinford born about 1590, the common paternal progenitor of Skidmore Lineage C, was named Skidmore Lineage E3 in the Phase II report. As was suggested, he upgraded to 43 markers. Previously, 6701 had three differences from Skidmore Lineage E1, but after upgrading he had 11 differences, thus moving him out of Lineage E and into the independent lineages.

□ This upgrade has also effected Skidmore Lineage E4, 10508, Lee Skidmore, descendent of Mary Skidmore. In Phase II it was reported "Lineages E3 and E4 share a mutation at DYS390, indicating that they likely share common ancestry with each other before they share a common ancestor with the rest of Lineage E (Figure 3d, Figure 5). However, because 10508 and 6701 share only 23 of 26 alleles, their common ancestor is estimated at 28 generations with a 95% Confidence Interval of 9 to 65 generations. Therefore, the connection of these two lineages to each other and to the rest of Skidmore Lineage E is tentative. The Phase II report did not mention that they also had a shared mutation at DYS392, though they still share only 23 of 26 markers. It is imperative that 10508 upgrade to 43 markers to determine if he has shared ancestry with 6701, with the rest of Lineage E, or neither, as the genetic data at this point are unclear.

Scudamore/Skidmore Lineage F:

Skidmore Lineage F was created by two new individuals 70418, Dr. John F. Skidmore, and 58854, Dr. Peter Skidmore who shared 25 of 26 alleles. They have a calculated MRCA of 13 generations with a 95% Confidence Interval of 2 to 38 generations. It is recommended that 58854 upgrade to 43 markers so that a more detailed comparison can be made to 70418 who has already been tested at 43 markers. This will give further insight into the relationship of these two individuals.

Scudamore/Skidmore Lineage G:

Skidmore Lineage G was created by eight new individuals. Lineage G1 is defined by Z097935, Milus Skidmore, 63912, Walter R. Skidmore, 68685, Jack Skidmore, and 68781, John Skidmore, who are all descendents of different sons of John Skidmore of Lee County, VA, born in 1792.

Lineage G2 is defined by Z097979, James Tennessee Burke, who differs from G1 at *DYS19* and *DYS385b*. Lineage G3 is defined by 43557, Mark Skidmore, who differs from Lineage G1 at *DYS19*, *DYS389II*, and *DYS439* where he has a value of 14. Lineage G4 is defined by 57575, Dr. Dwight V. Skidmore who differs from Lineage G1 at *DYS19*, *DYS389II*, and *DYS439* where his value is 13. Lineage G5 is defined by 62028, John Steven Edgar, who believed he shared common ancestry with G1's Z097935 in James M. Skidmore of Harlan County, KY born in 1855, but his three differences from G1 show that it is highly unlikely that this assumption is correct.

From the current genetic information we cannot verify the genetic profile of the lineage common ancestor, Skidmore Lineage G Common Ancestor 3. It is likely that a *DYS19* mutation occurred before the common ancestor of the Lineage G1 participants, refer to Skidmore Lineage G Common Ancestor 1, in Figure 3. Because there are two mutually exclusive shared mutations at *DYS389II* and *DYS439*, for Lineages G3 and G4, there are two genetic alternatives as to how the other individuals in this lineage are related.

(Figure 3a) First, if the *DYS389II* mutation is truly shared by Lineage G3 and Lineage G4, then they would share a common ancestor in Skidmore Lineage G common Ancestor 2 before sharing common ancestry with the rest of the group. This would mean that the *DYS439* (13) value shared by Lineage G4 and G5 would have happened independently on each of the G4 and G5 lines.

(Figure 3b) Conversely, if the *DYS439* (13) value is the result of shared ancestry, then Lineages G4 and G5 share a common ancestor in Skidmore Lineage G Common Ancestor 2, and it is the *DYS389II* mutation that occurred independently. To determine which scenario is more likely, more known cousins of Z097979, 43557, 57575, and 62028 should be tested.

It is also recommended that Z097979, 43557, and 57575 upgrade to 43 markers in order to better define their relationship with Lineage G1 as their current MRCA is between 13 and 20 generations.

Figure 3a: Skidmore Lineage G when the *DYS389II* mutation is shared.

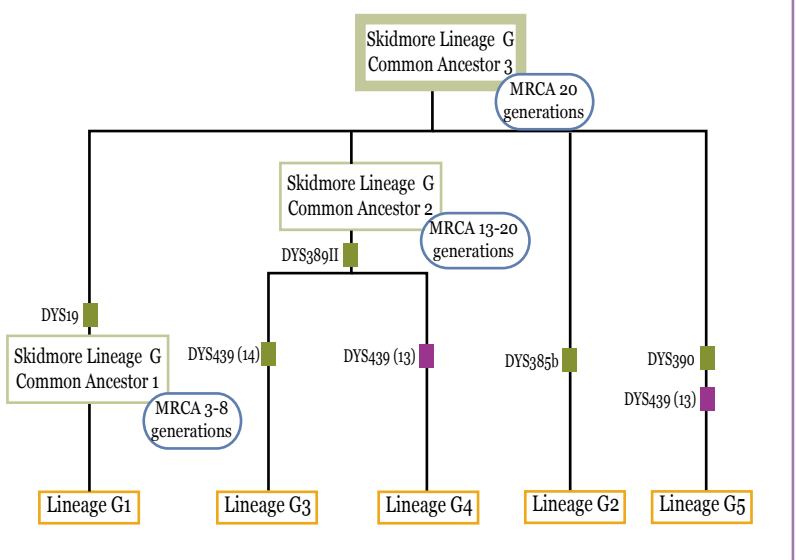
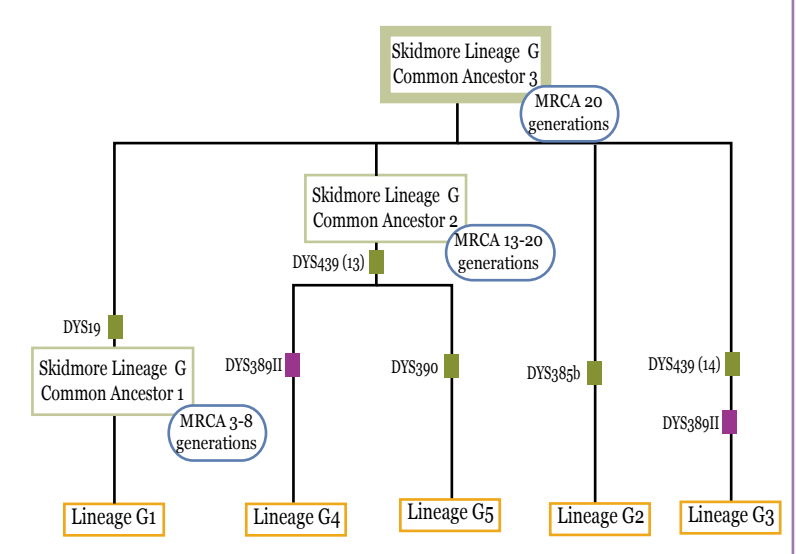


Figure 3b: Skidmore Lineage G when the *DYS439* mutation is shared.



Scudamore/Skidmore Independent Lineage H:

□ Skidmore Lineage H is defined 8894, who was previously an independent lineage, and newcomer 57918, a descendent of Edward Skidmore of Fairfax Co., VA. 8894 and 57918 share 24 of 26 markers giving them an MRCA of 20 generations with a 95% Confidence Interval of 5 to 52 generations. It is highly recommended that 8894 upgrade to 43 markers in order to narrow the time better define the relationship of these two individuals.

Scudamore/Skidmore Independent Lineages:

□ 6701, 7291, 8894, 20071, 36639, 41512, and 41662 form unique Skidmore Lineages as each has a profile distinct from anyone else in the study (Figure 2).

What Next?

Following are the recommendations made in the text to further your genetic genealogy pursuits. Besides these specific recommendations, you can always extend your knowledge of the relationships of individuals by conducting traditional genealogical research or by testing more individuals, both those with, and without known genealogical connections to current participants.

- p. 4 Lineage B upgrade 6940, 7294, and/or 40475
- p. 5 Lineage E Upgrade 10508
- p. 5 Lineage F Upgrade 58854
- p. 6 Lineage G Test more known cousins of Z097979, 43557, 57575, and 62028
- p. 6 Lineage G Upgrade Z097979, 43557, and 57575 to 43 markers
- p. 7 Lineage H Upgrade 8894 to 43 markers

Conclusion:

The Scudamore/Skidmore Family Genetics Project, Phase III was able to address the requested objectives by using genetic testing techniques to assess the probability of relationships. If you have any questions regarding the information in this report, please contact your project coordinator, or contact Diahan Southard at diahan@relativegenetics.com, or at (801) 462-1431.

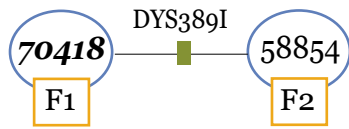
The undersigned have reviewed the above results and do hereby verify that they are accurate according to the information provided by the participants.

Diahan L. Southard
Molecular Genealogist

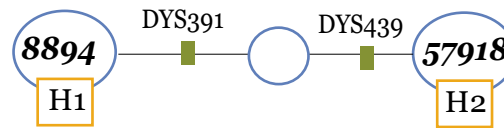
Lars Mouritsen
Director of Laboratory Operations

Figure 1b- Haplocation: This diagram shows how subgroups in a family project are connected. To find out how genetically related two individuals are, count how many mutations, represented by boxes, separate the two individuals. The fewer boxes separating two individuals, the more closely they are related. See the text for more information.

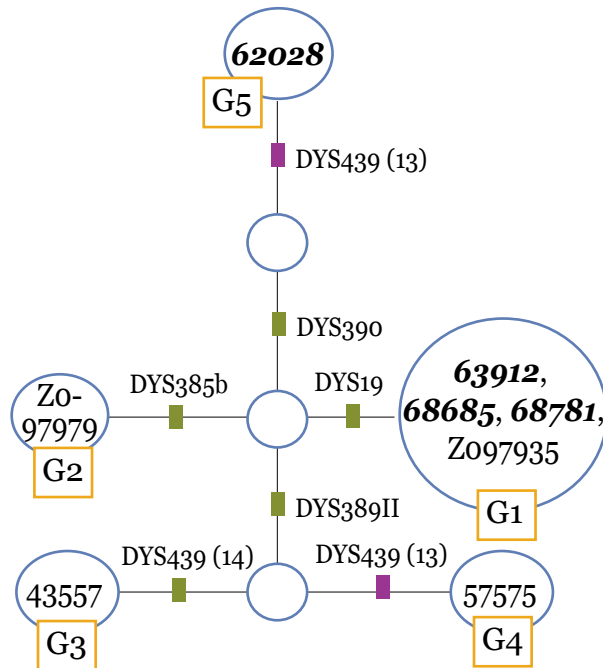
Skidmore Lineage F:



Skidmore Lineage H:



Skidmore Lineage G:

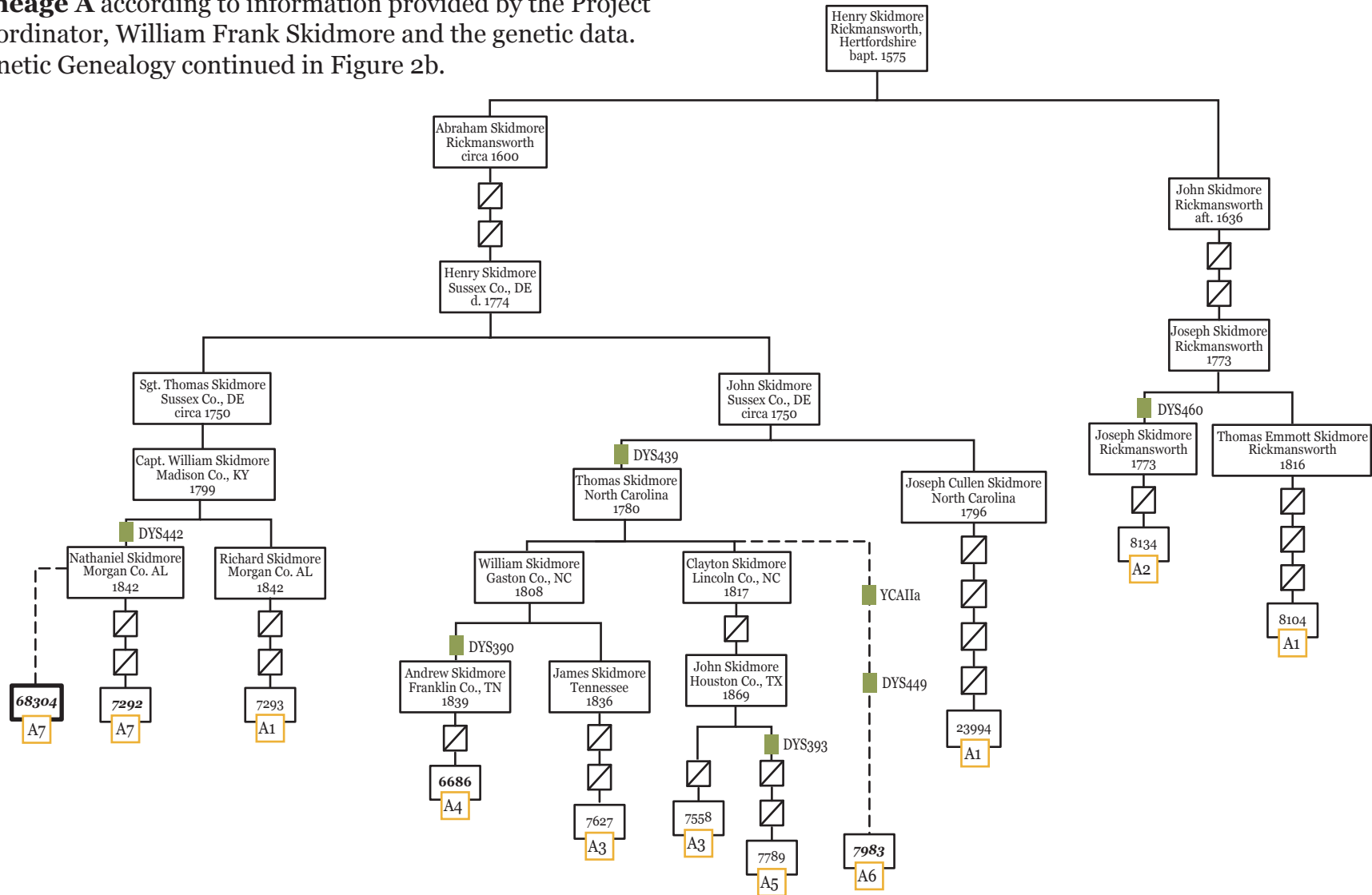


Skidmore Independent Lineages:

6701, 7291, 8894, 20071,
36639, 41512, 41662

- # Each circle with a number represents a participant.
- 54612** Those individuals with bold italic type had 43 markers analyzed.
- X6 Each square designates the specific lineage of the individuals with the letter defining the lineage and the number defining the haplotype in that lineage.
- Green denotes a single mutation, the site of the mutation is noted in black.
- Other colors denote a mutation that is shared between two individuals not in the same haplotype.

Figure 2a: The genetic genealogy of **Scudamore/Skidmore Lineage A** according to information provided by the Project Coordinator, William Frank Skidmore and the genetic data. Genetic Genealogy continued in Figure 2b.



Legend for Figures 2a-2e:

- # Each square with a number represents a participant.
- #** Each bold square designates a new participant.
- 54612** Those individuals with bold italic type had 43 markers analyzed.
- X6 Each orange square designates the specific lineage of the individuals with the letter defining the lineage and the number defining the haplotype in that lineage.
- Green denotes a single mutation, the site of the mutation is noted in black.
- Other colors denote a mutation that is shared between two lineages that are not in the same haplotype.
- MRCA 3-8 generations The Most Recent Common Ancestor figures refer to the likely generation individuals within a lineage converge in a common ancestor

Figure 2b: The genetic genealogy of Scudamore/Skidmore Lineage B according to information provided by the Project Coordinator, William Frank Skidmore and the genetic data. Genetic Genealogy continued in Figure 2c.

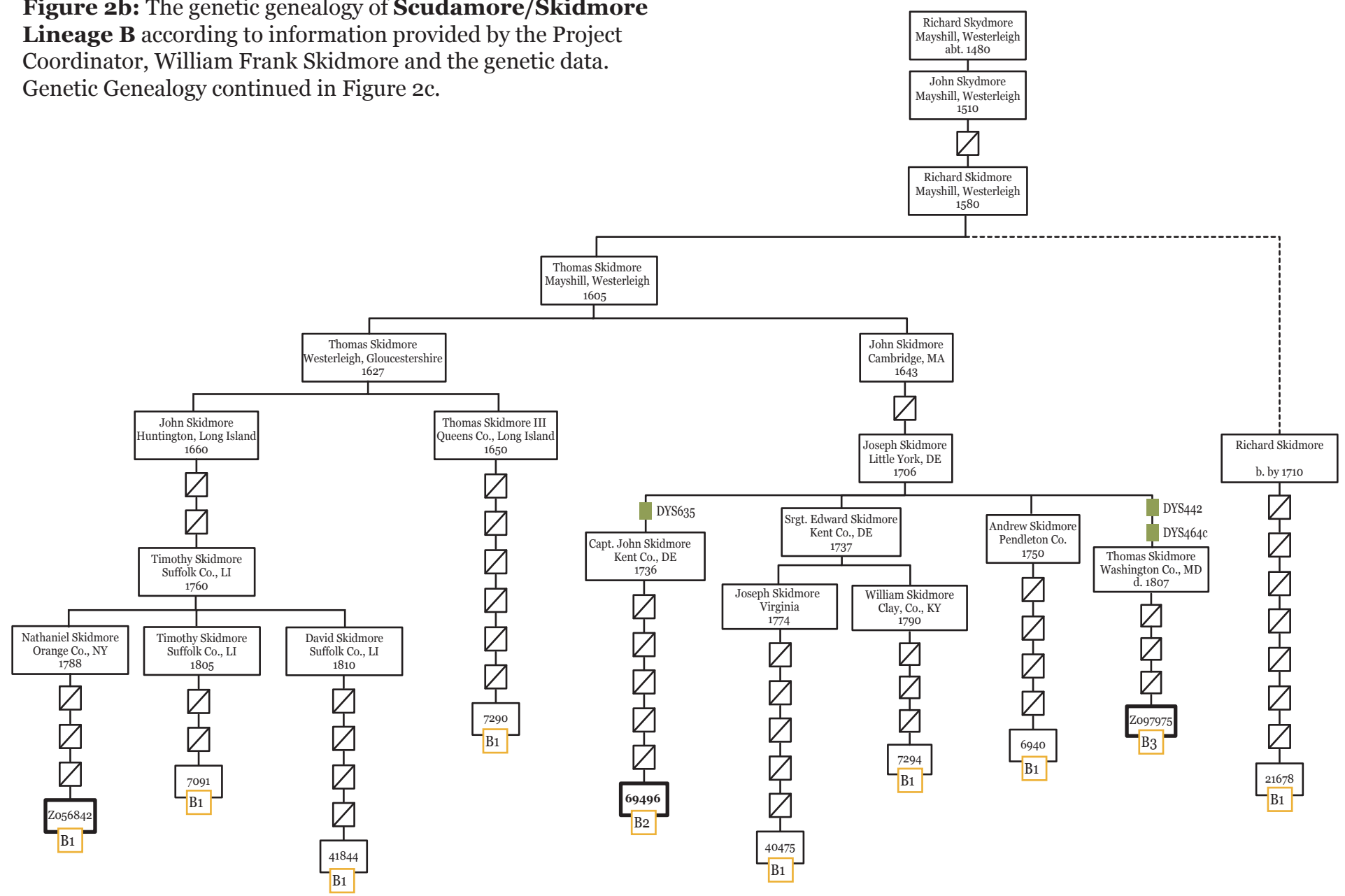
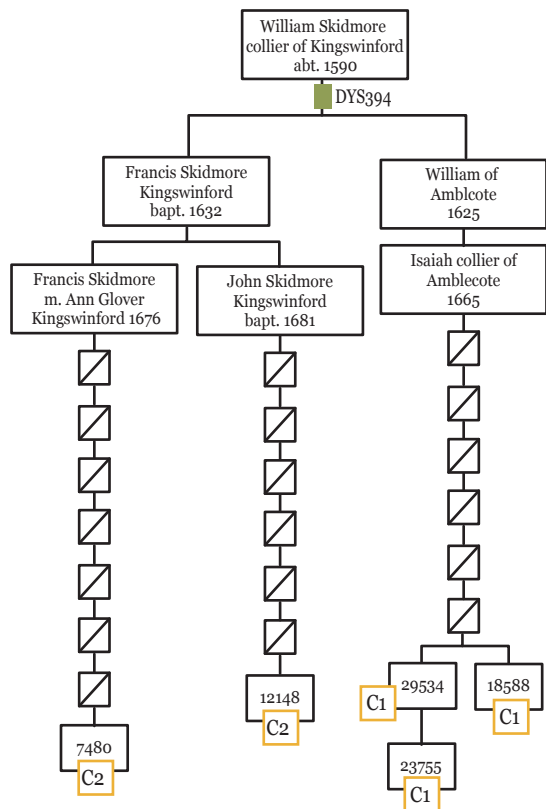


Figure 2c: The genetic genealogy of **Scudamore/Skidmore Lineages C and D** according to information provided by the Project Coordinator, William Frank Skidmore and the genetic data. No changes have been made in these lineages since Phase II. Genetic Genealogy continued in Figure 2d.

*Scudamore/Skidmore
Lineage C:*



*Scudamore/Skidmore
Lineage D:*

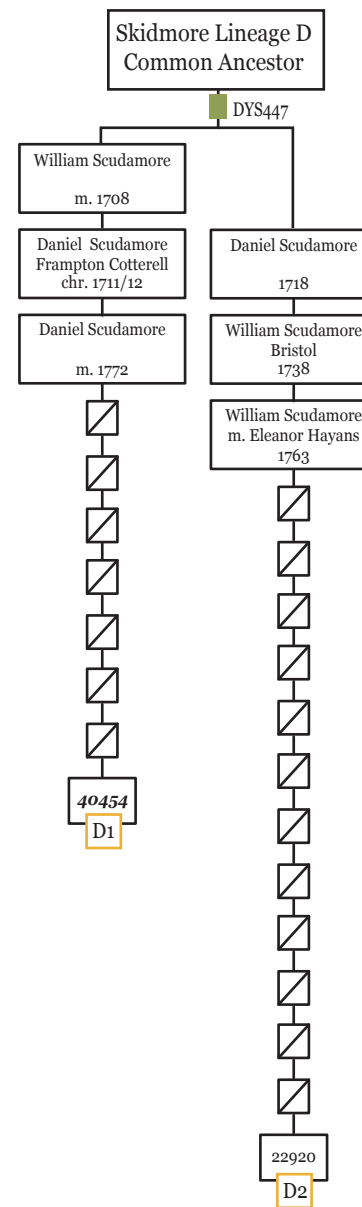


Figure 2d: The genetic genealogy of **Scudamore/Skidmore Lineage E** according to information provided by the Project Coordinator, William Frank Skidmore and the genetic data. The green boxes indicate the earliest genealogical point where the mutation identifying the lineage could have occurred according to the present genetic data. The site of the mutation is noted in black. Genetic Genealogy continued in figure 2e.

Scudamore/Skidmore Lineage E:

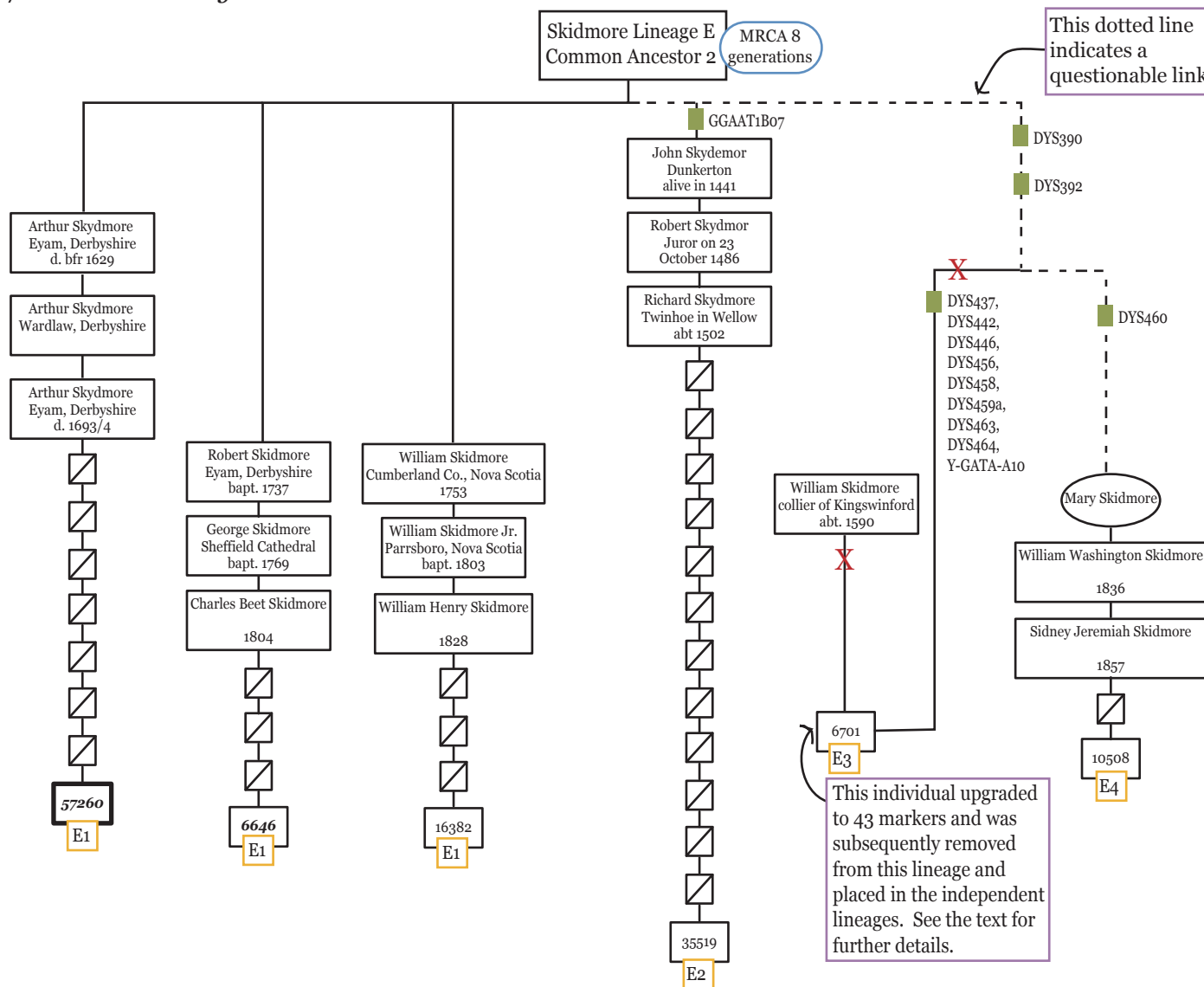
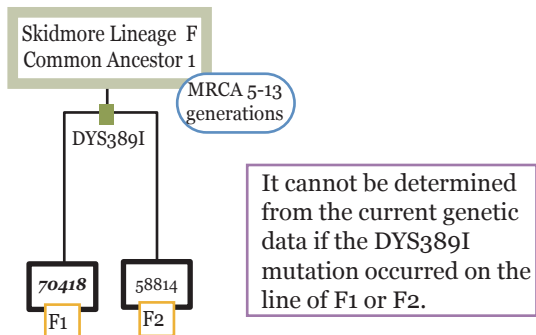
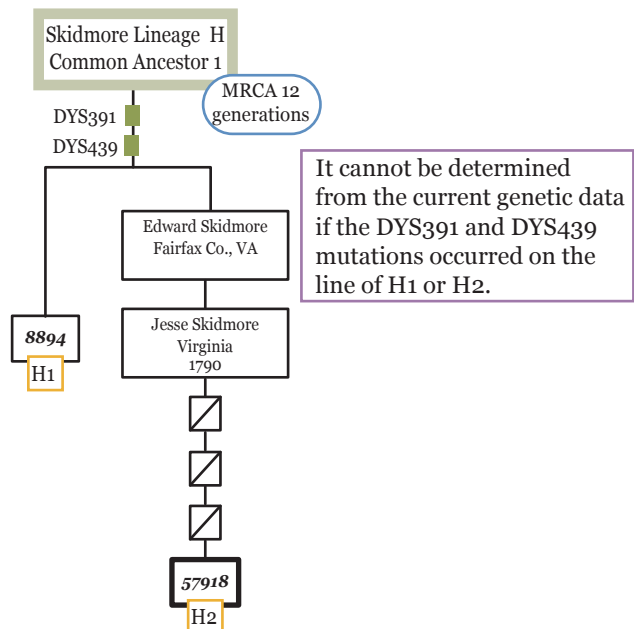


Figure 2e: The genetic genealogy of Scudamore/Skidmore Lineages F, G and H according to information provided by the Project Coordinator, William Frank Skidmore and the genetic data.

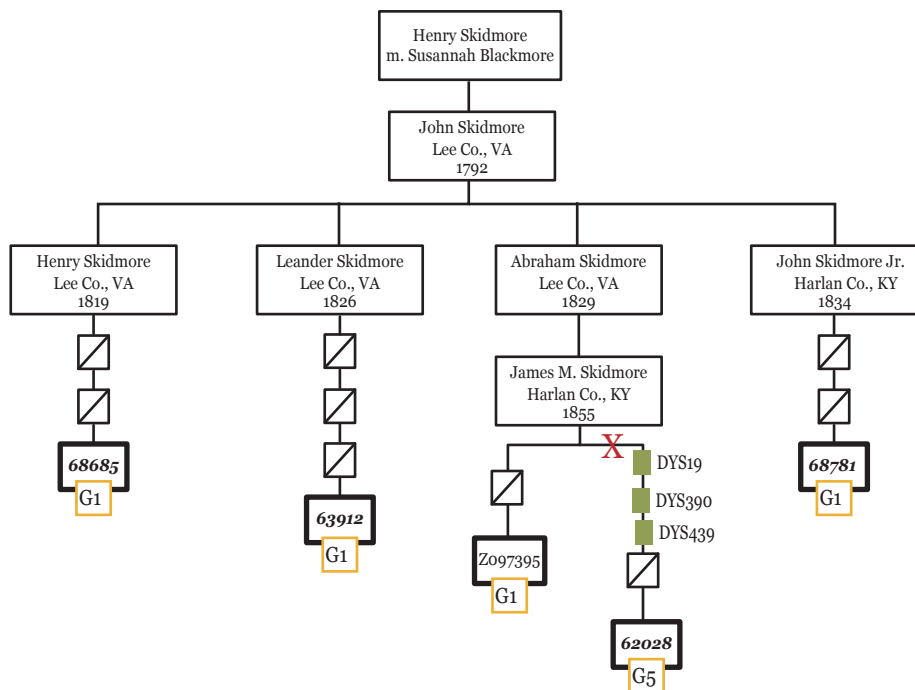
Scudamore/Skidmore Lineage F:



Scudamore/Skidmore Lineage H:



Partial Scudamore/Skidmore Lineage G*:



*This diagram includes all who submitted genealogical information, all participants are represented in Figure 3 in the text.

Appendix A1: A list of all the participants in Skidmore Lineages A and B. Those in bold type had 43 markers tested. The column "Differs From Modal At" indicates the mutation(s) that individual carries from the first haplotype in that lineage; likewise the "Match %" column lists the percentage matched to that first haplotype. The "Haplogroup" column indicates the inferred large ancestral grouping each individual is likely to belong to based on the current genetic data. To find out more about haplogroups and see the individual genetic profiles for all individuals, please login to your family website at www.relativegenetics.com.

Skidmore Lineage A: 11 individuals in 7 haplotypes

Name	Case Number	Lineage	Lineage Changes since Phase II	Differs From Modal At	Match%	Haplogroup
Billy Skidmore	T007293	A1	No	-	100%	R1b
Barry Hastings Skidmore	T008104	A1	No	-	100%	R1b
Charles Skidmore	T023994	A1	No	-	100%	R1b
Ron Skidmore	T008134	A2	No	DYS460	96.20%	R1b
Gene Aaron Skidmore	T007558	A3	No	DYS439	96.20%	R1b
James Edgar Skidmore	T007627	A3	No	DYS439	96.20%	R1b
Wm F Skidmore	T006686	A4	No	DYS439, DYS390	92.30%	R1b
Eric Skidmore	T007789	A5	No	DYS393, DYS439	92.30%	R1b
George Skidmore	T007983	A6	No	DYS439, YCAIIa, DYS449	88.46%	R1b
Robert M. Skidmore	T068304	A7	New	DYS442	96.20%	R1b
Wade Skidmore	T007292	A7	No	DYS442	96.20%	R1b

Skidmore Lineage B: 10 individuals in 3 haplotypes

Name	Case Number	Lineage	Lineage Changes since Phase II	Differs From Modal At	Match%	Haplogroup
Bill Skidmore	Z056842	B1	New	-	100%	G
Warren Skidmore	T006940	B1	No	-	100%	G
Howard Skidmore	T007091	B1	No	-	100%	G
Howard H. Skidmore	T007290	B1	No	-	100%	G
Ellis Skidmore	T007294	B1	No	-	100%	G
Frank G. Skidmore	T021678	B1	No	-	100%	G
Dennis Skidmore	T040475	B1	No	-	100%	G
Brian Skidmore	T041844	B1	No	-	100%	G
Michael George Skidmore	T069496	B2	New	DYS635	96.20%	G
Robert Emerson Skidmore (FTDNA)	Z097975	B3	New	DYS442, DYS464c	93.80%	G

New Participant **Change in self**

Appendix A2: A list of all the participants in Skidmore Lineages C, D, E and F. Those in bold type had 43 makers tested. The column "Differs From Modal At" indicates the mutation(s) that individual carries from the first haplotype in that lineage; likewise the "Match %" column lists the percentage matched to that first haplotype. The "Haplogroup" column indicates the inferred large ancestral grouping each individual is likely to belong to based on the current genetic data. To find out more about haplogroups and see the individual genetic profiles for all individuals, please login to your family website at www.relativegenetics.com.

Skidmore Lineage C: 5 individuals in 2 haplotypes

Name	Case Number	Lineage	Lineage Changes since Phase II	Differs From Modal At	Match%	Haplogroup
James Skidmore	T018588	C1	No	-	100%	I1c
Richard Charles Skidmore	T023755	C1	No	-	100%	I1c
Robert Skidmore	T029534	C1	No	-	100%	I1c
Ewart Burt Skidmore	T007480	C2	No	DYS19	96.20%	I1c
David. M. Skidmore	T012148	C2	No	DYS19	96.20%	I1c

Skidmore Lineage D: 2 individuals in 2 haplotypes

Name	Case Number	Lineage	Lineage Changes since Phase II	Differs From Modal At*	Match%	Haplogroup
Andrew Skidmore	T040454	D1	No	DYS447	96.20%	I1c
Gavin Skidmore	T022920	D2	No		96.20%	I1c

*Modal cannot be identified when there are only two individuals in a lineage, the locus shown is the point at which these two individuals differ

Skidmore Lineage E: 5 individuals in 3 haplotypes*

Name	Case Number	Lineage	Lineage Changes since Phase II*	Differs From Modal At	Match%	Haplogroup
Dr. Ed Skidmore	T006646	E1	Upgrade	-	100%	R1b
John A. Skidmore	T057260	E1	New	-	100%	R1b
Tim Skidmore	T016382	E1	No	-	100%	R1b
Peter Charles Scudamore	T035519	E2	No	GGAAT1B07	96.20%	R1b
Lee Skidmore	T010508	E4	Yes	DYS390, DYS392, DYS460	88.40%	R1b

*In Phase II this lineage had one additional individual that was moved to the independent lineages, see Figure 2d for further explanation

Skidmore Lineage F: 2 individuals in 2 haplotypes

Name	Case Number	Lineage	Lineage Changes since Phase II	Differs From Modal At*	Match%	Haplogroup
Dr. John F. Skidmore	T070418	F1	New	DYS389I	96.20%	R1b
Dr. Peter Skidmore	T058854	F2	New		96.20%	R1b

*Modal cannot be identified when there are only two individuals in a lineage, the locus shown is the point at which these two individuals differ

Appendix A3: A list of all the participants in Skidmore Lineages G and H and the Independent Lineages. Those in bold type had 43 markers tested. The column "Differs From Modal At" indicates the mutation(s) that individual carries from the first haplotype in that lineage; likewise the "Match %" column lists the percentage matched to that first haplotype. The "Haplogroup" column indicates the inferred large ancestral grouping each individual is likely to belong to based on the current genetic data. To find out more about haplogroups and see the individual genetic profiles for all individuals, please login to your family website at www.relativegenetics.com.

Skidmore Lineage G: 8 individuals in 5 haplotypes

Name	Case Number	Lineage	Lineage Changes since Phase II	Differs From Modal At*	Match%	Haplogroup
Milus Skidmore (FTDNA)	Z097395	G1	New	-	100%	I1a
Walter R. Skidmore	T063912	G1	New	-	100.00%	I1a
Jack Skidmore	T068685	G1	New	-	100.00%	I1a
John Skidmore	T068781	G1	New	-	100.00%	I1a
James Tennessee Burke (FTDNA)	Z097979	G2	New	DYS19, DYS385b	83.40%	I1a
Mark Skidmore	T043557	G3	New	DYS19, DYS389II, DYS439	75.00%	I1a
Dr. Dwight V. Skidmore	T057575	G4	New	DYS19, DYS389II, DYS439	75.00%	I1a
John Steven Edgar	T062028	G5	New	DYS19, DYS390, DYS439	75.00%	I1a

Skidmore Lineage H: 2 individuals in 2 haplotypes

Name	Case Number	Lineage	Lineage Changes since Phase II	Differs From Modal At*	Match%	Haplogroup
Leonard John Scudamore	T008894	H1	Yes, was IND	DYS391, DYS439	95.40%	I1a
John Robert Skidmore	T057918	H2	New		95.40%	I1a

*Modal cannot be identified when there are only two individuals in a lineage, the locus shown is the point at which these two individuals differ

There are 7 Independent Skidmore Lineages

Name	Case Number	Lineage	Lineage Changes since Phase II	Differs From Modal At*	Match%	Haplogroup
Ian Skidmore	T006701	IND	Yes, was E3	N/A	< 70%	R1b
Robert Herman Skidmore	T007291	IND		N/A	< 70%	R1b
Alan Skidmore	T020071	IND		N/A	< 70%	R1b
Ernest Roy Skidmore	T036639	IND		N/A	< 70%	E3b
Matthew Scudamore	T041512	IND		N/A	< 70%	R1b
Lamoin Edward Skidmore	T041662	IND		N/A	< 70%	R1b