

H11 Newsletter



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1. Project Statistics:

Combined GEDCOMs Uploaded 49

DISTINCT mtDNA Haplogroups 17

Family Finder 293

Maternal Ancestor Information 344

mtDNA 381

mtDNA Full Sequence	373
mtDNA Plus	379
mtDNA Subgroups	23
Total Members	419
Unreturned Kits	17

2. Other

There are nine new members since the last update on the subclades with Issue 3. All new members are now assigned to their individual groups. Information which follows will revise my reporting in February as the YFull mitochondrial Tree has been updated and I will use their coding for naming so changes will be forthcoming in the full issue of the next Volume of the Newsletter. I will also have to consider whether I am actually providing a service with this Newsletter. I am contemplating that I could start up a Kipp Family Newsletter with the Kipp Family Study and that could become my third newsletter. I have switched the orientation to Landscape in order to display the H11 table at YFull in section 5.

3. In Memoriam: Barry Hinman, Librarian Emeritus Stanford University

I sadly discovered yesterday that one of my correspondents in the H11 study passed away in July this year. Barry Hinman (Librarian Emeritus Stanford University) and I have been corresponding about H11 for five years. I did try to persuade him to become a co-administrator and he did consider it but he had other items he wanted to write up and the H11 study does

take time. His own line was somewhat rare and not yet charted independently on the Phylotree. He had sent me an email in March 2021 which I have only just reached and responded to him yesterday but sadly the email bounced back. He had found some new work on T961a which supplied him with a subclade name for his particular grouping. I am happy to say that that did happen. At the time I was very much involved with my husband's illness and just could not grapple with anything. I did write to Dr. Mannis Van Oven about the Phylotree and had sent him all of Barry's information as he published the latest update in February of 2016. The new branch name which was selected by Genetic Homeland - H11a1b.

4. H11 in the News

T961A does occur in some members in the study. To date the phylotree does not take this particular mutation into account in its rendering of H11 haplogroup and its subclades. I am not able to determine yet if there will be an update to the Phylotree. The last build was 18 Feb 2016 and it was mtDNA tree Build 17. With new information from a member of the study I have emailed Dr. Mannis van Oven and will let you know if I hear back from him in the full recitation of results in February 2022.

Some information on T961A includes: "The full sequence mitochondrial test at FamilyTreeDNA shows this maternal haplogroup to be H11, but with a unique variation so far not shared with anyone else in the United States: T961a. The mutation T961g is the significant marker for the group H11a as is A16293G, which these women do carry. A new subclade for T961a will have to be created in haplogroup H11. The site Genetic Homeland has done so, calling the new subclade H11a1b." Submitted by a group member.

<https://www.genethomeland.com/welcome/dnamarkerindex.asp?snp=T961A&Chromosome=M>

None of the papers cited appear to be later than 2014 authored by Der Sarkissian, C, Brotherton, P, Balanovsky, O, Templeton, JEL, Llamas, B et al PLOS ONE:

Marker Name	Notes	Identification	Ancestral	Derived	Chromosome	Position
T961a	Transversion. rCRS Ref: t. Yoruba (hg18) Position: 963. See also T961C. Also see branches: H11a1b	Yfull 2019	T	A	M	961

DNA Marker Search

Instructions

This page provides a list of all DNA marker names and aliases associated with a specific DNA position in our GeneticHomeland DNA database. You can search it by entering the chromosome and marker (or position) that you want to lookup.

*Searching for {Y and M222} or {Y and 14902414} or {Y and rs20321} produces the same result because SNP M222 is located at position 14902414² on the Y chromosome and was designated by NIH as rs20321.

*We display both hg38 and hg19 chromosome positions in our listings wherever possible as much of the literature still uses hg19. If you search by numeric position, you have the option to use hg19, hg38, or BOTH of these human genome assemblies.

*We honor the *haplogroup-SNP Name* nomenclature used by some laboratories to distinguish between multiple instances of the same mutation allele at the same position having occurred in different branches of humanity (haplogroups). i.e. Searching for R-S7642 produces a different result from searching for I-S7642 as a mutation has occurred at the same position in two different haplogroup branches of humankind.

*For pure RSID³ SNP searches, select Any (RSID) as the Chromosome, then enter the RSID value starting with "rs", e.g. rs1000000019. Alternatively, you can enter an hg38 numeric position and we will search for any rsid instances at that position across all the human chromosomes e.g. 10007658 will return mutation records on five different chromosomes. We also try to display the hg18 assembly position for RSID mutations as those were used in older literature.

Barry has also given me his permission to share his maternal line “his mother Catharine Antonetta Radivoj (b San Francisco, USA 19 Jul 1909), her mother Catherine Tadejevich (b 22 Mar 1878 Praputnjak, Croatia), her mother Ana Stiglich (b 11 Jul 1836 Praputnjak, Austro-Hungarian Empire), her mother Maria Margaretha Gecan (bc 1790 Praputnjak, Austro-Hungarian Empire), her mother Petronilla Bujan (Bakar, Austro-Hungarian Empire), her mother Margaretha Fitnić (b 17 Jun 1735 Bakar, Austro-Hungarian Empire), her mother Catharina Bujan (bc 1701 Bakar, Austro-Hungarian Empire).

“The women, though, carry a so-far unique genetic inheritance. The full sequence mitochondrial test at FamilyTreeDNA shows this maternal haplogroup to be H11, but with a unique variation so far not shared with anyone else in the United States: T961a. The mutation T961g is the significant marker for the group H11a as is A16293G, which these women do carry. A new subclade for T961a will have to be created in haplogroup H11.

“The site Genetic Homeland has done so, calling the new subclade H11a1b.¹ Another site, YFull, agrees and further estimates that the subclade was formed about 800 years before the present. All of the persons known to have this mutation are shown to be from Russia.²”

- 1. <https://www.genetichomeland.com/welcome/dnamarkerindex.asp?snp=T961A&Chromosome=M>; <https://www.genetichomeland.com/welcome/dnapedigree.asp?RecordID=1727511>**
- 2. <https://www.yfull.com/mtree/H11a1b/>**

Thank you to Barry Hinman for providing this material for this H11 Newsletter.

5. YFull Tree – H11a1b

Website provided by Barry Hinman’s notes:

<https://www.yfull.com/mtree/H11a1b/>

YFull MTree 1.02.12917 (under construction)

SEARCH

H11a1b T961A formed 800 ybp, TMRCA 600 ybp

H11a1b*
id:<DLMXXU> RUS

H11a1b1 A4215G formed 600 ybp, TMRCA 250 ybp
id:KY670871.1 RUS
id:KY671012.1 RUS

H11a1c T16224C formed 800 ybp, TMRCA 375 ybp

H11a1c*
id:MN540516.1 FIN [FI-IS] age
id:KY671011.1 RUS
id:MH492645.1
id:KY671125.1 RUS
id:KF451406.1
id:<4K98JC>

H11a1c1 T13635C formed 375 ybp, TMRCA 125 ybp
id:MN516678.1 FIN
id:KY670966.1 RUS
id:YT68459

H11a1c2 A73G! formed 375 ybp, TMRCA 100 ybp
id:YF81023 LVA [LV-VMR] lav
id:YF75883 LVA [LV-VMR] lav
id:YF75787 LVA [LV-VMR] lav
id:YF78569 LVA [LV-053] lav
id:YF74595 LVA [LV-VMR] lav

MTree 1.02.12917 (24 October 2021)

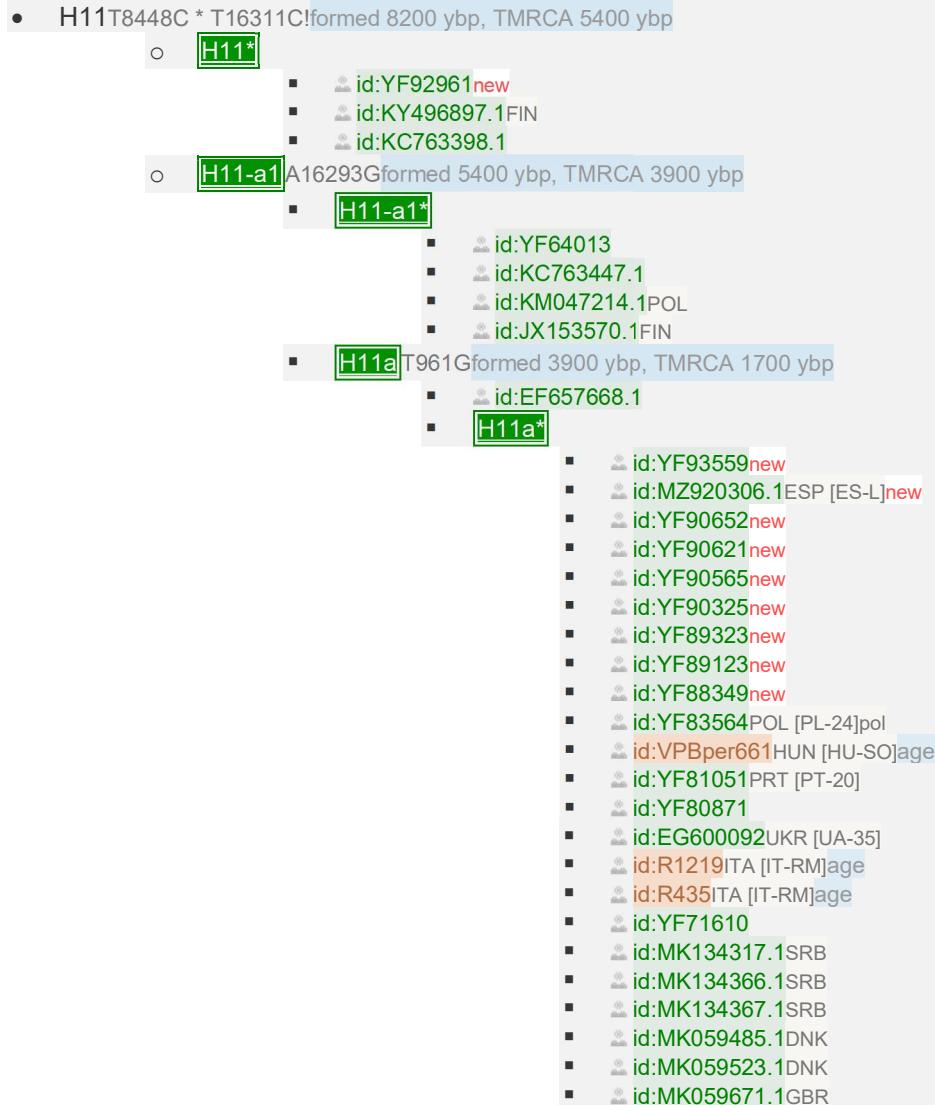
Like 735

New subclade Modified subclade id:AncientDNA

H11

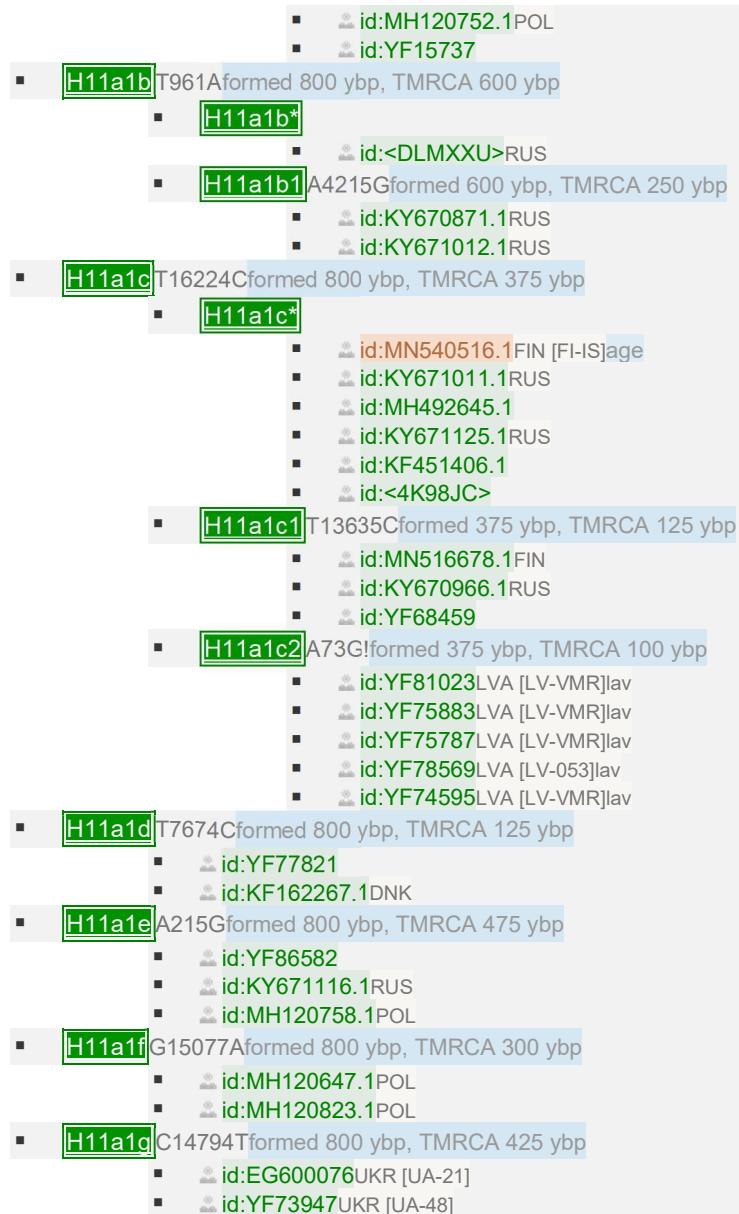
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The H11 Tree at YFull:

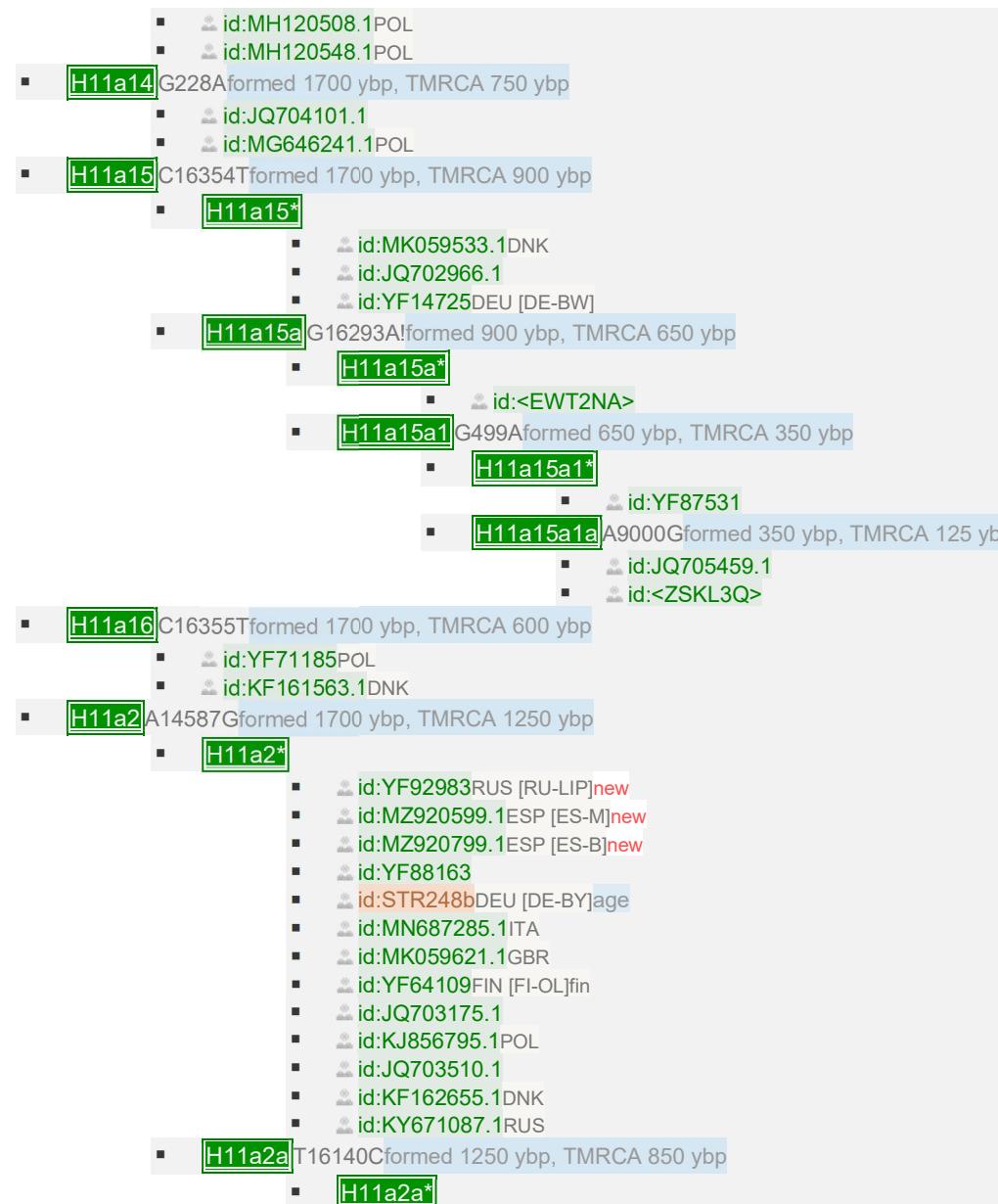


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-  id:JQ705984.1
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-  id:KF162478.1DNK
-  id:KF162294.1DNK
-  id:KM101842.1USA
-  id:KF161732.1DNK
-  id:MG182513.1
-  id:KF162388.1DNK
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-  id:JQ703167.1
-  id:JQ703163.1
-  id:JQ703593.1
-  id:JQ704185.1
-  id:JQ704846.1
-  id:JQ705807.1
-  id:JQ705898.1
-  id:JX152946.1DNK
-  id:JX152992.1FIN
-  id:JX153368.1DNK
-  id:JX153417.1DNK
-  id:JX153483.1DNK
-  id:JX153711.1DNK
-  id:JX153761.1DNK
-  id:JX153771.1DNK
-  id:JX154068.1FIN
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-  id:MG429009.1
-  id:MG182511.1ITA [IT-TO]
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-  id:HGDP00522
-  id:KF451057.1
-  id:YF15230NLD [NL-NB]
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-  id:<WL6DCT>

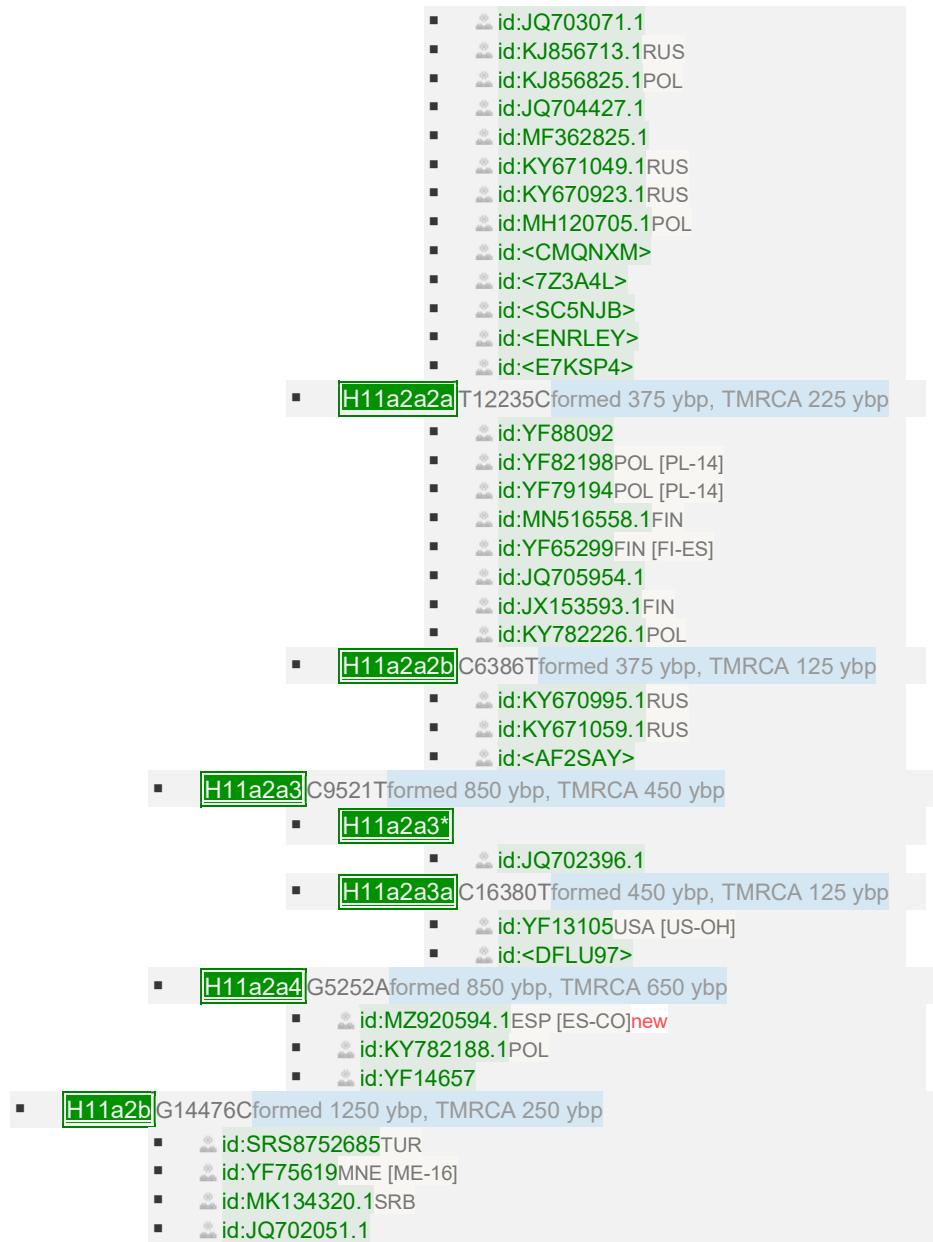
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- id:<2WNS8D>
- **H11a1** C8898T * C16278T! formed 1700 ybp, TMRCA 800 ybp
 - id:EF657502.1
 - id:EF657478.1
 - **H11a1***
 - id:YF89192 new
 - id:YF88137
 - id:YF81144 SWE [SE-O]swe
 - id:YF80859
 - id:YF71489 UKR
 - id:YF70765
 - id:MH298069.1 ROU
 - id:KJ856810.1 POL
 - id:KJ856767.1 RUS
 - id:KF162198.1 DNK
 - id:AY738958.1 ITA
 - id:JQ703259.1
 - id:KC286597.1
 - id:KY670856.1 RUS
 - id:MG646148.1 POL
 - id:MH120502.1 POL
 - id:KY671025.1 RUS
 - id:MH120723.1 POL
 - id:YF06214 SVK [SK-KI] slk
 - id:<QPPYNS>
- **H11a1a** A13434G formed 800 ybp, TMRCA 500 ybp
 - **H11a1a***
 - id:K2per71 HUN
 - id:YF86375
 - id:KC286601.1
 - id:MG646089.1 POL
 - id:MH120485.1 POL
 - id:<MPARVB>
 - id:YF03671 SRB
- **H11a1a1** C11503A formed 500 ybp, TMRCA 250 ybp
 - id:MG646231.1 POL
 - id:MG646251.1 POL
 - id:MG646146.1 POL



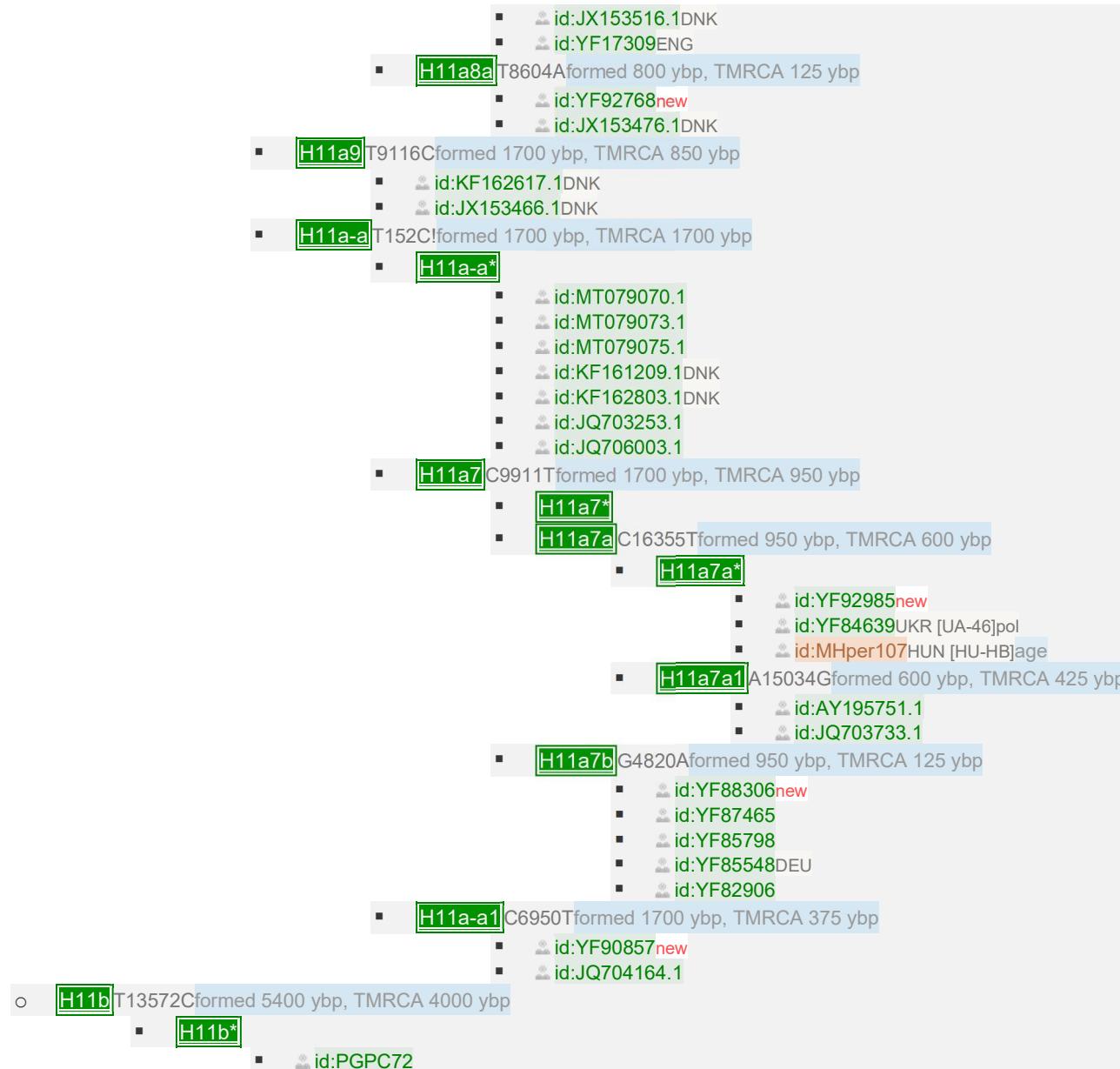
- **H11a1h** T5021C formed 800 ybp, TMRCA 350 ybp
 - [id:KY409403.1ITA](#)
 - [id:KY409275.1ITA](#)
 - [id:KM101911.1USA](#)
 - [id:KY409984.1ITA](#)
- **H11a1i** T146C formed 800 ybp, TMRCA 275 ybp
 - [id:YF84572USA \[US-OH\]](#)
 - [id:YF84399](#)
 - [id:YF66036RUS \[RU-ARK\]](#)
 - [id:YF63402SWE \[SE-BD\]swe](#)
 - [id:YF63138FIN \[FI-LS\]](#)
 - [id:YF63136FIN \[FI-LS\]](#)
 - [id:YF19968SWE \[SE-BD\]](#)
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 - [id:KJ856785.1RUS](#)
 - [id:GU122994.1RUS](#)
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 - [id:JQ703688.1](#)
 - [id:JX153177.1FIN](#)
 - [id:MF497486.1](#)
 - [id:YF19104RUS \[RU-KIR\]](#)
 - [id:YF13659SWE \[SE-BD\]swe](#)
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 - [id:YF09377FIN \[FI-LS\]](#)
 - [id:YF03564RUS \[RU-IRK\]](#)
 - [id:<VE4DA6>](#)
- **H11a1j** A2784G formed 800 ybp, TMRCA 125 ybp
 - [id:YF89491new](#)
 - [id:JQ703453.1](#)
- **H11a10** A5515G formed 1700 ybp, TMRCA 1150 ybp
 - [id:MH981652.1ZAF](#)
 - [id:YF10936USA \[US-NC\]](#)
- **H11a11** T8618C formed 1700 ybp, TMRCA 1250 ybp
 - [id:JQ705928.1](#)
 - [id:KF451844.1](#)
- **H11a12** T16368C formed 1700 ybp, TMRCA 475 ybp
 - [id:<3FZBWB>USA \[US-CA\]](#)
 - [id:YF11733USA \[US-CA\]](#)
 - [id:<KYHC2Y>](#)
- **H11a13** C11206T formed 1700 ybp, TMRCA 375 ybp

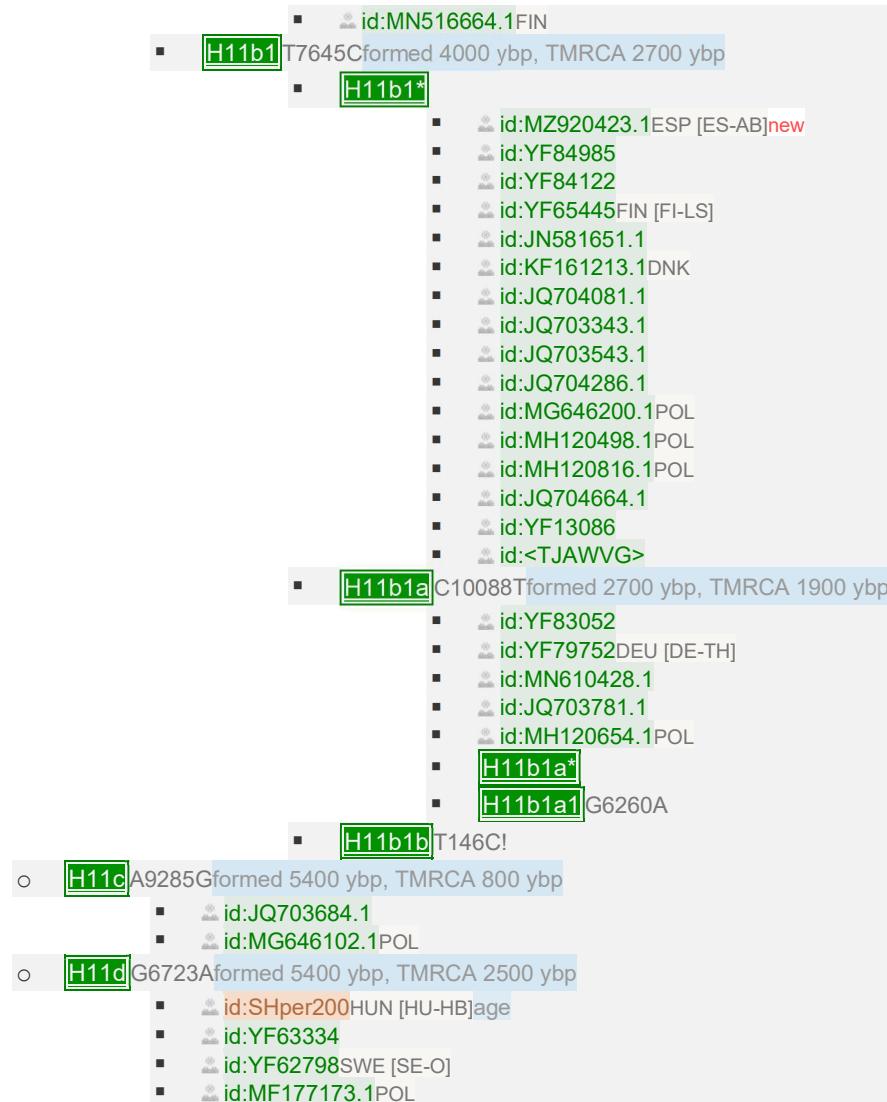


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- id:KF161727.1DNK
- id:KM101966.1USA
- id:KF161565.1DNK
- id:JQ704395.1
- id:JQ705580.1
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- id:MH120589.1POL
- id:MF177165.1POL
- id:<XBER6T>
- id:YF02536FIN [FI-AL]
- id:<AUD3TY>
- id:<JPTCSJ>
- **H11a2a1** A3145G formed 850 ybp, TMRCA 225 ybp
- **H11a2a1***
 - id:YF84341
 - id:YF78076
 - id:JQ703451.1
 - id:YF07623ENG
 - id:YF71875IRL [IE-DL]
 - id:<UZN6UG>
 - id:<X36V63>
- **H11a2a1a** G16293A! formed 225 ybp, TMRCA 100 ybp
- id:YF87634
- id:YF83623USA [US-MS]
- id:JQ702388.1
- id:JQ703183.1
- id:<5BU5CJ>
- id:<L7THJW>
- **H11a2a2** G5585A * T15670C * A16265G formed 850 ybp, TMRCA 375 ybp
- id:<83U9PN>
- **H11a2a2***
 - id:YF92975new
 - id:MK059657.1GBR
 - id:YF80143CZE [CZ-OL]
 - id:YF65472RUS



- **H11a2c** C6854T formed 1250 ybp, TMRCA 650 ybp
 - id:YF19129 MKD [MK-407]
 - id:YF11397 ALB [AL-10]aln
- **H11a2d** T16298C formed 1250 ybp, TMRCA 200 ybp
 - id:YF85296
 - id:YF78515 SRB [RS-07]srp
 - id:JQ704166.1
- **H11a2e** C16301T formed 1250 ybp, TMRCA 600 ybp
 - **H11a2e***
 - id:MHper137 HUN [HU-HB]age
 - **H11a2e1** T7885C formed 600 ybp, TMRCA 275 ybp
 - id:YF85985 SRB
 - id:MK134293.1 SRB
- **H11a3** T16243C formed 1700 ybp, TMRCA 350 ybp
 - id:YF88631 new
 - id:JQ703172.1
 - id:JQ703235.1
 - id:<BQC7PN>
- **H11a4** C16111T formed 1700 ybp, TMRCA 400 ybp
 - id:YF88002
 - id:YF87918
 - id:YF80757 USA [US-VA]
 - id:YF70757 ENG
 - id:JQ704280.1
- **H11a5** C15040T formed 1700 ybp, TMRCA 475 ybp
 - id:YF68215 BGR [BG-01]
 - id:YF68204 BGR [BG-01]
 - id:YF66971 SRB [RS-08]srp
 - id:GU592023.1
 - id:GU592038.1
 - id:JQ703949.1
- **H11a6** G1719A * G5979A * A16525G formed 1700 ybp, TMRCA 300 ybp
 - id:JQ705740.1
- **H11a8** T14325C formed 1700 ybp, TMRCA 800 ybp
 - **H11a8***
 - id:YF85012
 - id:YF65490
 - id:JQ705643.1





November 1, 2021

H11 NEWSLETTER - VOLUME 5, ISSUE 4

MTree 1.02.12917 (24 October 2021)

Scientific sample prefixes and any related scholarly papers are listed [here](#).

Any submissions to this newsletter can be submitted to Elizabeth Kipp (kippeeb@rogers.com).