Genetic Genealogy 2: So, I Tested my DNA; Now What?

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INTRODUCTION:

Genetic genealogy – especially *DNA testing for ethnic origin* – has become wildly popular. Yet, for millions of people, the ethnicity estimates don't match their family history, and cousin-finding tools seem complex and confusing. This discussion seeks to explain and simplify the next steps for using your DNA for greater insights and successful family research.

DNA TESTING BASICS:

- 3 types of DNA tests: autosomal (at-DNA), mitochondrial (mt-DNA) and Y-chromosome (Y-DNA) (see image A).
- Of these, at-DNA is the least expensive, most popular and generally most genealogically useful DNA test.
- The top DNA test labs are: AncestryDNA, FamilyTreeDNA, MyHeritage, 23andMe and LivingDNA (see resource Links).
- at-DNA (1) provides ancestral ethnicity estimates, and (2) identifies others with whom you share a recent ancestor.
- at-DNA reflects your whole pedigree, but only the last 150 years; at-DNA halving & shuffling washes-out earlier ancestors.
- DNA testing can have privacy implications: law enforcement or others can gain access to your DNA test results.
- Artifact DNA testing of hair brushes, envelopes and stamps is becoming possible, though not always successful.
- DNA is contributory evidence; it tells whether 2 people share an ancestor, but not exactly who or when.
- Traditional "paper research" is **always** required to place DNA cousins accurately within family trees.

DISAPPOINTING ETHNIC ESTIMATE RESULTS; the "Horoscope" of DNA Tests:

- Each DNA lab uses different DNA markers and reference populations to reach different ethnicity conclusions.
- Country-level ethnicity estimates are most inconsistent, but broad regional estimates tend to be more valid.
- Your ethnicity test will often not match paper research, your siblings' test results or even yourself at different labs.
- If your parents have vastly different geographic origins, ethnic estimates can be helpful to family research, otherwise, despite all the flashy ads, ethnicity estimates are muddled, imprecise and rarely useful to genealogists.

AUTOSOMAL DNA TOOL & SKILLS: (see image B)

- BASIC DNA TOOLS Each DNA lab's website offers a list of "Cousin Matches" also called a "One-to-Many" analysis, where the "One" is you, and the "Many" are all the lab's other clients. Labs provide info. about your Matches such as:
 - cM the amount of DNA overlap, measured in centimorgans
 - Total % the amount of DNA overlap, measured as a percentage of total DNA
 - Largest Segment (cM) the length of the longest individual segment of DNA overlap in centimorgans
 - Number of Segments the total count of segments making up the DNA overlap
 - MRCA (Most Recent Common Ancestor) a kinship estimate or number of generations that separate the 2 DNA tests.
- INTERMEDIATE DNA TOOLS Each DNA lab's collection of test results is limited to just their clients. So looking for DNA matches at only one lab is like fishing in a single small pond. To fish in several ponds, don't pay for DNA tests at all the labs, instead import your DNA test results to the websites other labs and to third party for free.
 - First, **consider the privacy issues** publicly posting your DNA test results has risks. If the tests belong to family members, obtain their consent to more widely display their results. Even if you use an alias, publicly posted DNA results are available to law enforcement and could help them identify a criminal family member.
 - Then, download your DNA file from the testing lab to your PC. For step-by-step instructions, Google, "how to download my AncestryDNA" (or other website). It arrives as a "compressed" .zip file. Save to your PC but don't open it.
 - Next, upload your DNA to other labs and to DNA analysis websites. For step-by-step instructions, Google, "how to upload my DNA file to MyHeritage" (or other website). With the exception of AncestryDNA and 23andMe, the main DNA labs all allow you to import your DNA test results from other labs for free (see image C). Additional sites like www.GedMatch.com offer DNA tools and tree matches. After a GedMatch upload, perform two quality checks: run DNA File Diagnostic Utility and Are Your Parents Related tools from GedMatch's Analysis Tools (see image D).
- ADVANCED DNA TOOLS We use *AncestryDNA* and *GedMatch* as popular examples of online resources. This section reviews four key tools that genetic genealogists will find useful. Deploy these four steps in order:
 - SKILL #1: ONE-TO-MANY ANALYSIS finds a list of matches between your DNA test results and the collection of DNA tests at your DNA testing lab. AncestryDNA's "Confidence Scores" (see image E) are typical of how labs characterize the quality of your matches. The online display of One-to-One matches varies among DNA test repositories:

 - AncestryDNA Matches are in a List including Name or Alias, Gender, Match Confidence and Probable Kinship.

There are two approaches to recognizing which matches are most meaningful:

- **GedMatch** focuses on Matches with any **DNA segments > 7 cM**, though we suggest you adjust to 10 cM.
- AncestryDNA focuses on Matches with total overlap of at least 30 cM of shared DNA.

Note: GedMatch's Table of One-to-Many Matches can be copied & pasted into a PC spreadsheets for off line use.

- SKILL #2: ONE-TO-ONE ANALYSIS next, both AncestryDNA and GedMatch provide tools to allow exploration of
 individual Matches to find shared surnames, geographic locations and family tree branches even DNA segments.
 - GedMatch allows One-to-One analysis by clicking on an individual within your One-to-Many Table.
 - **AncestryDNA** allows One-to-One analysis by clicking on an individual within your One-to-Many List.

When *One-to-One* Matches seem promising, but the matching person has no visible surnames or trees, there are strategies to find the nature of your matching person's family connection. Try each of the following:

- \blacksquare No tree at Ancestry DNA or Ged Match doesn't mean that a tree can't be found; check Ancestry. com.
- ♣ No tree at AncestryDNA or GedMatch may indicate a very private genealogist; reach-out carefully; directly, or through by the DNA labs, you can message your One-to-One DNA Matches to compare your research notes.
- ♣ No tree at AncestryDNA or GedMatch may indicate a newbie genealogist; be prepared to mentor them.
- Leven without a tree, Chromosome Browsers analyze shared DNA segments & can help indicate a family branch.
- The DNA Painter tool can further identify likely relationships based on cMs of shared DNA (see Image F).
- **SKILL #3: TWO-TO-ONE ANALYSIS** next, both *AncestryDNA* & *GedMatch* provide *Two-to-One* tools to produce a list of Matches (the "One" of *Two-to-One*) who share DNA with <u>both</u> you AND a relative ("the "Two" of *Two-to-One*):
 - **GedMatch** produces a *Two-to-One* analysis by clicking on the *People Who Match 1 or Both of 2 Kits* tool.
 - **AncestryDNA** produces *Two-to-One* analysis by clicking on <u>Compare</u> icon in any of the *One-to-Many* Matches. Placing these *Two-to-One* Matches accurately within your family tree, requires several next steps:
 - ♣ First, by selecting 2 individuals who are related, and 1 or are more generation apart as the "Two" in the Two-to-One analysis, you can infer the family branch of individuals of the Match List; the "Ones." For example, a "Two" consisting of a mother & son will produce 2-to-1 matches that are from the son's maternal side and not from his paternal side. If the Match is an unknown, this tool eliminates 50% of kinship possibilities and narrows the focus of traditional research that is needed to reach more specific placement of the 2-to-1 match in your family tree.
 - **Second,** perform a *One-to-One* analysis (described above) on the unknown match from the Two-to-One match list. Look for shared surnames, geographic locations, family trees and DNA segments to further identify the match.
 - **Third,** if the relationship is still clear, use DNA Painter to identify the most likely relationship(s) based on the amount of DNA overlap (cM). This should further narrow potential branches and tree landing spots.
 - **Fourth,** even more relationship insights can be gained from the use of a *Chromosome Browser*, combined with *Spreadsheet* tracking and sorting tools. This is the next skill, below.

SKILL #4: CHROMOSOME BROWSERS & SPREADSHEETS – Finally, these analytic tools can help unravel otherwise insurmountable Match mysteries. Only *GedMatch* is helpful here, because *AncestryDNA* has no chromosome browser:

- First, from within GedMatch open your One-to-Many Match Table.
- ♣ Second, copy & paste GedMatch's One-to-Many Match Table into a spreadsheet on your PC, then create new columns to track: Shared Surnames, Shared Locations, Likely Branch, Probable Kinship and, if you're very ambitions: Segment Starting Point, Segment Ending Point, Segment Length (cM) note: these three Segment columns need to be created for each of your 22 autosomal chromosomes (= 66 additional columns).
- **Third**, select any of your individual Matches from your *One-to-Many* Table by clicking on the "A" link in their Detail column within the table's Autosomal area. This opens a settings page for a *One-to-One* Table. Select the "Position Only" option to produce a Table of segment overlap data without the chromosome graphics.
- Fourth, copy new information into your PC spreadsheet's row for each match: Segment Starting Point, Segment Ending Point, Segment Length (cM) for each chromosome. If you know, add the match's Shared Surnames, Shared Locations, Likely Branch & Probable Kinship. For a fee, GedMatch's Tier 1 Tools can automate segment matches.
- **Fifth**, the Spreadsheet can be sorted by column for commonalities which indicate family branch insights.

Combined with traditional research you can now place previously unknown Matches accurately within your family tree.

CONCLUSION

Though at-DNA testing won't answer all your research questions, by uploading your test results to several DNA labs' websites, you can dramatically increase your chances of finding new cousins with whom to compare notes, share traditional research and unravel the mysteries of your family. And, thanks to these 4 analytic DNA skills, you can more confidently place even unknown cousins within your existing family tree.

RESOURCES DNA Testing Labs

www.AncestryDNA.com www.FamilyTreeDNA.com

www.23andMe.com

www.MyHeritage.com

www.LivingDNA.com

www.OnTheEnvelope.com (artifact testing)

Genetic Genealogy Online Reading

www.isogg.org/wiki/Wiki Welcome Page

https://www.gedmatch.com/Using GEDmatch.php

https://www.gedmatch.com/gedwiki/index.php?title=FAQs

https://www.gedmatch.com/DNA for Dummies.php

DNA Tools Websites

www.DNApainter.com/cmv4 www.GedMatch.com www.DNAGedcom.com

Genetic Genealogy Books & e-Books

<u>The Guide to DNA Testing and Genetic Genealogy</u>
Blaine T. Bettinger

<u>DNA Buying Guide: Are you buying the right test?</u>
Thomas MacEntee

<u>23 Best Tips for DNA Testing and Family History</u> Mary Eberle, J.D.

MAJOR GENEALOGICAL DNA TEST TYPES Y-DNA DESCRIPTION mt-DNA at-DNA **Autosomal Chromosomes DNA Source** Mitochondria Y Chromosome (44 of 46 chromosomes; all except X & Y) Cell Nucleus Location Cell Cytoplasm Cell Nucleus TEST SUBJECTS Who can be tested? Women or Men Men only Women or Men TIME FRAME Does DNA get Shuffled? No No Yes, each generation Unstable (shuffling and DNA Stability Extremely Stable Stable halving each generation) Looks how far back? 1000s of genertions 100s of generations 5-6 generations Inconsequential minimal mutation is over-Ave. Mutation Rate? Once every 10,000 years Once every 800 years shaddowed by shuffling) **Test Type** Anthropological Genealogical Genealogical **DNA USES** Matching to others with Matching to others with a Matching to others with a **Primary Use** shared ancestor a shared ancestor shared ancestor Ethnicity Estimates (%) Secondary Uses Migratory Insights Migratory Insights Migratory Insights **LABS & COSTS** Best Lab FamilyTreeDNA FamilyTreeDNA AncestryDNA \$59-\$99 \$169 (37 markers) to \$89 (2 regions) to Cost (# of Markers) (> 700,000 markers) \$149 (all regions) \$649 (500 markers) **ACCURACY & CERTAINTY** 100% 98-99% Overlap required to Anything > 0.5% (no mismatched DNA (2 or fewer mismatched demonstrate Kinship (as little as 16 cM) DNA markers) markers) **Match Certainty** MRCA Estimated MRCA Estimated MRCA Estimated **Etnicity Accuracy** Moderate Moderate Slightly Better

GedMatch DNA Tools

Analyze Your Data

DNA raw data

'One-to-many' matches
 Information: Disappeared kits recovery information

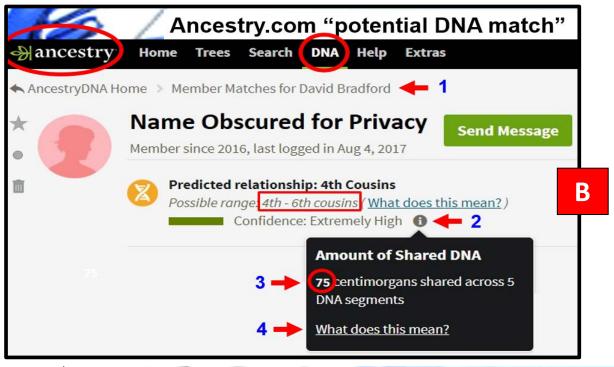
D

- recovery information
 Action: 'One-to-many' recovery no account email matches
- 'One-to-one' compare
- X 'One-to-one'
- Admixture (heritage)
- Admixture/Oracle with Population Search
- Phasing
- People who match one or both of 2 kits Updated
- Predict Eye Color
- Are your parents related?
 - 3D Chromosome Browser
 - Archaic DNA matches
 - Multiple Kit Analysis NEW
- DNA File Diagnostic Utility

 Analyze DNA file upload for potential problem.

Genealogy

- 1 GEDCOM to all
- 2 GEDCOMs
- Search all GEDCOMs
 Revised
- GEDCOM + DNA matches



	DNA LABS THAT ALLOW IMPORTING OF YOUR DNA FILES							
Company	Allows DNA Import?	Import for Free	Imports from 23and Me?	Imports from AncestryDNA?	Imports from FTDNA?		Imports from MyHeritage?	
C 23andMe	No	n/a		n/a	n/a	n/a	n/a	No
AncestryDNA	No	n/a	n/a		n/a	n/a	n/a	No
FamilyTree DNA	Yes	No*	Yes	Yes		No	Yes	Yes
Living DNA	Yes	Yes	Yes	Yes	Yes		Yes	No
MyHeritage	Yes	Yes	Yes	Yes	Yes	No		No

Confidence Score	Approximate amount of shared centimorgans	Likelihood of a single recent common ancestor			
Extremely High	More than 60	Virtually 100%			
Very High	45—60	About 99%			
High	30—45	About 95%			
Good	16-30	Above 50%			
Moderate	6—16	15—50%			

Shared at-DNA Relationship Calculator (courtesy of Blaine T. Bettinger at https://dnapainter.com/tools/sharedcmv3)

centimorgans or cM) to calculate the most likely relationships that may exist between you and your "Match." When an email "Match" is identified by your at-DNA testing lab, this web site uses the amount of shared DNA (measured in

Half 3C3R	Half 3C2R 34 0 – 96	Half 3C1R 42 0 – 165	Half 3C 61 0 – 178	Half 2C1R 73 0-341	Half 1C2R 145 37 – 360	Half GG- Aunt/Uncle 187 12 – 383	Range (low to high) (99th percentile)	KEY TO CHART	
Half 2C3R	Half 2C2R 61 0 – 353	Half 2C1R 73 0 – 341	Half 2C 117 9 – 397	Half 1C1R 226 57 – 530	Half-Great Aunt / Uncle 432 125 – 765		CHART Possible Relationship of 2 Individuals Average cM overlap of known relationships of this type Range of cM overlap looking at 99% of known relationships of this type Example: of 99% of known Second Cousins Twice Removed (2C2R), the average overlap is 74 cM, though some have as little as 0 cM, while others have as much as 261 cM		
Half 1C3R 87 0 – 191	Half 1C2R 145 37 – 360	Half 1C1R 226 57 – 530	Half 1C 457 137 – 856	Half Aunt/Uncle 891 500 – 1446					
Half GG Niece / Nephew 187 12 – 383	Half Great- Niece / Nephew 432 125 – 765	Half-Niece / Nephew 891 500 – 1446	Half-Sibling 1783 1317 – 2312			G	ring at 99% of k in Second Cousi e have as little a	l Individuals	
Great Great Niece / Nephew 427 191 – 885	Great Niece / Nephew 910 251 – 2108	Niece / Nephew 1750 1349 – 2175	Sibling 2629 2209 – 3384	Parent 3487 3330 – 3720	Grandparent 1766 1156 – 2311	Great-Grandparent 881 464 – 1486	nown relationshins Twice Removas 0 cM, while or	ips of this type	
Great- Grandchild 881 464 – 1486	Grandchild 1766 1156 – 2311	Child 3487 3330 – 3720	SELF			ent	nips of this type ved (2C2R), the thers have as m		
1C3R 123 0 - 283	1C2R 229 43 - 531	1C1R 439 141 – 851	1C 874 553 – 1225	Aunt/Uncle 1750 1349 – 2175			average over- uch as 261 cM		
2C3R 57 0 - 139	2C2R 74 0 – 261	2C1R 123 0-316	2C 233 46 – 515	1C1R 439 141 – 851	Great Aunt/Incle 914 251 – 2108		Great-Great		
3C3R 22 0 - 69	3C2R 35 0 – 116	3C1R 48 0 – 173	3C 74 0-217	2C1R 123 0-316	1C2R 229 43 - 531	Great-Great Aunt/Uncle 427 191 - 485	Great-Great-Grandparent	Great-Gi Grand	
4C3R 29 0 – 82	4C2R 22 0 – 109	4C1R 28 0-117	4C 35 0 – 127	3C1R 48 0 - 173	2C2R 74 0-261	1C3R 123 0 - 283	GGG Aunt/Uncle	Great-Great- Grandparent	
5C3R 11 0 – 44	5C2R 17 0 – 43	5C1R 21 0-79	5C 25 0 – 94	4C1R 28 0 – 117	3C2R 35 0 – 116	2C3R 57 0 – 139		GGGG Aunt/Uncle	
8C 12 0 – 50	7C1R 13 0 – 53	7C 13 0-57	6C2R 17 0 – 75	6C1R 16 0 - 72	6C 21 0 - 86	Other Relationships		Ŧ	