

Phasing: The First Step in Helping Matches Make Sense

By
Philip Spivey



DNA 101

The relatively pain free version!

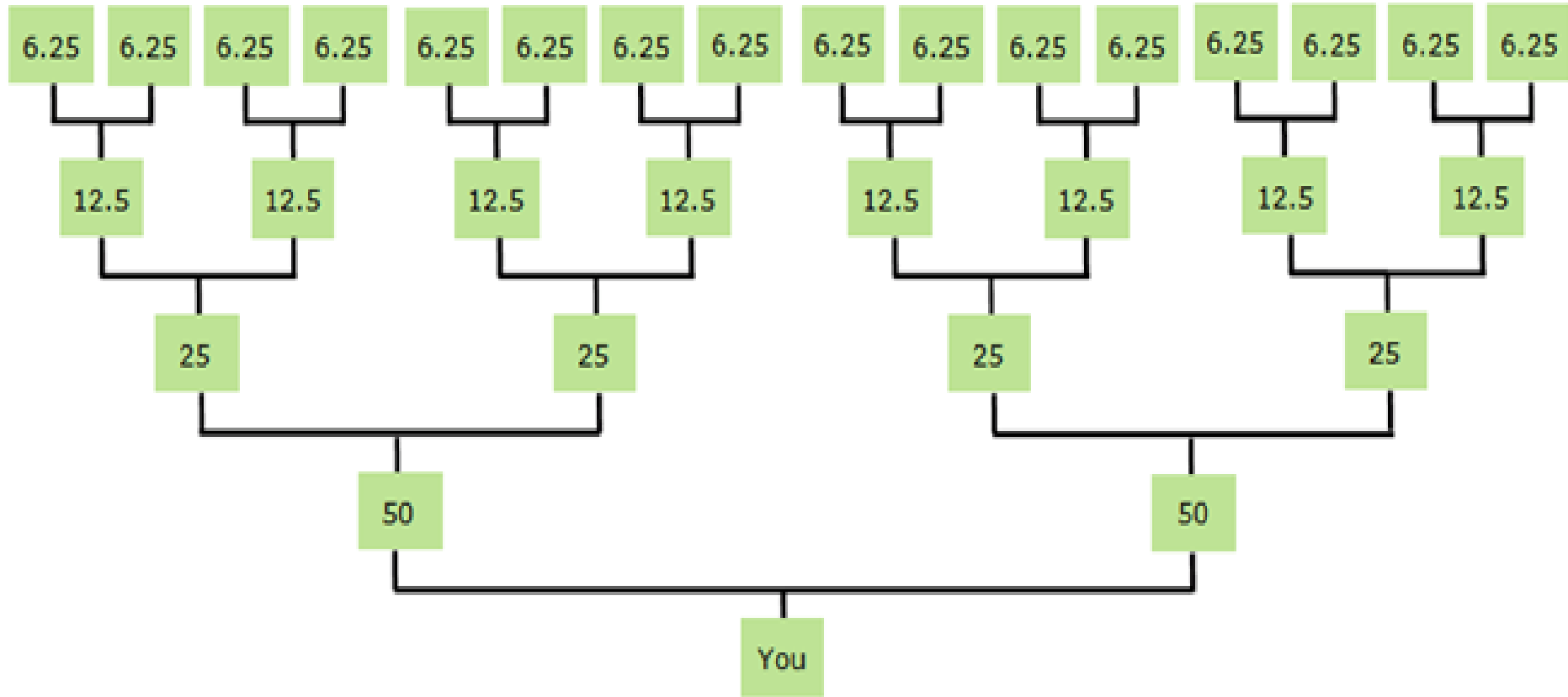


Autosomal Testing (at-DNA)

- Ancestry
- 23andMe
- MyHeritage
- FTDNA's Family Finder
- LivingDNA



at-DNA Inheritance



“Expected Ethnicity Results,” Ancestry (<https://support.ancestry.com/s/article/Unexpected-Ethnicity-Results> : accessed 2 March 2020). Potential inheritance percentages.

SKIN DEEP:
THE SCIENCE
OF RACE

DAWN OF
THE WHITE
MINORITY

US AND THEM:
WHY WE
DIVIDE

DRIVING
WHILE
BLACK

SPECIAL ISSUE

NATIONAL
GEOGRAPHIC

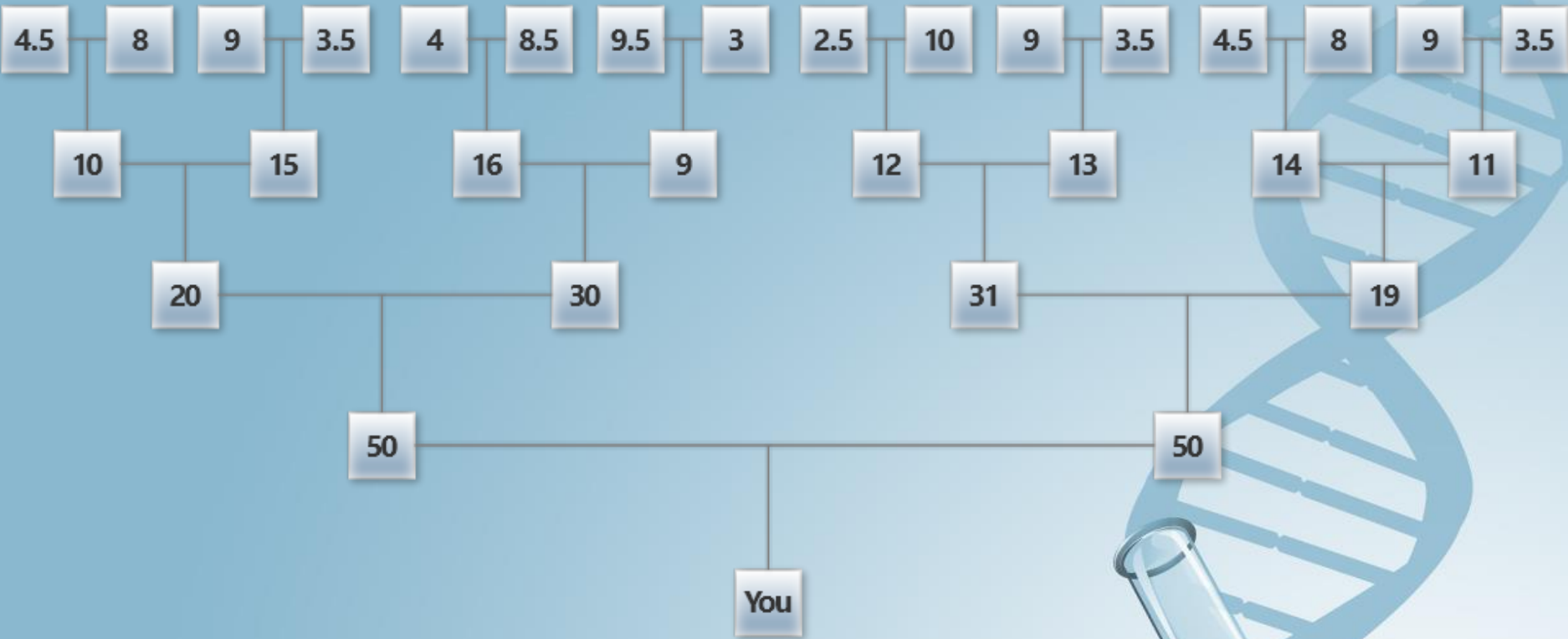
Black and White

These twin sisters make us rethink
everything we know about race

APRIL 2018

WHAT'S YOUR STORY? #10-11





DNA Match

Definition: A person that shares DNA with the test taker

- The closer the relationship, the more DNA shared
- Matches in a database only come from people that have tested at that database company or uploaded their data to that company



Chromosome Definition

A **chromosome** is an organized package of DNA found in the nucleus of the cell.

Different organisms have different numbers of **chromosomes**. Humans have 23 pairs of **chromosomes** - 22 pairs of numbered **chromosomes**, called autosomes, and one pair of sex **chromosomes**, X and Y

“Chromosome,” definition, *International Society of Genetic Genealogy* (<https://isogg.org/wiki/Chromosome> : accessed 2 March 2020).



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- X

Male



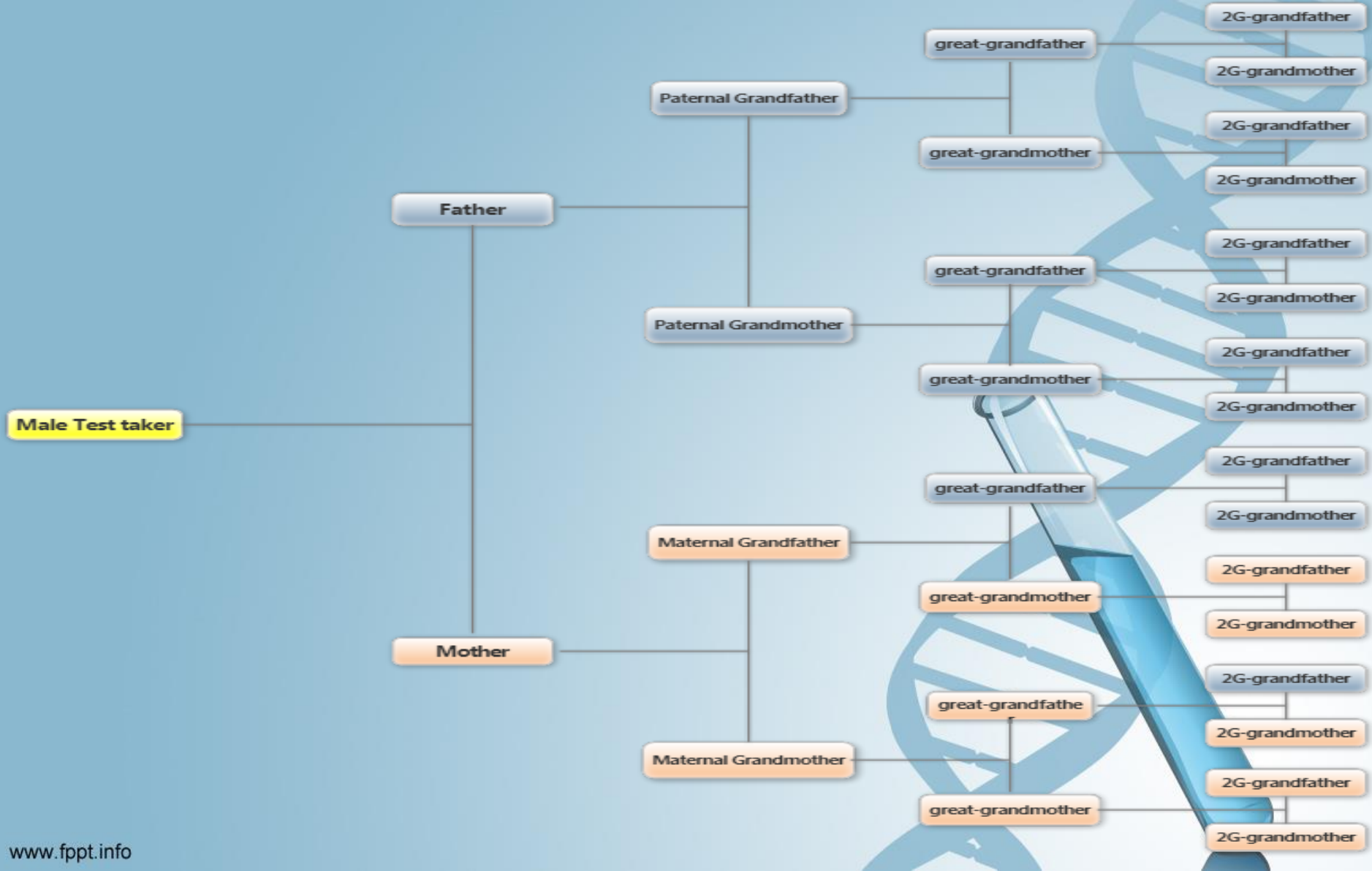
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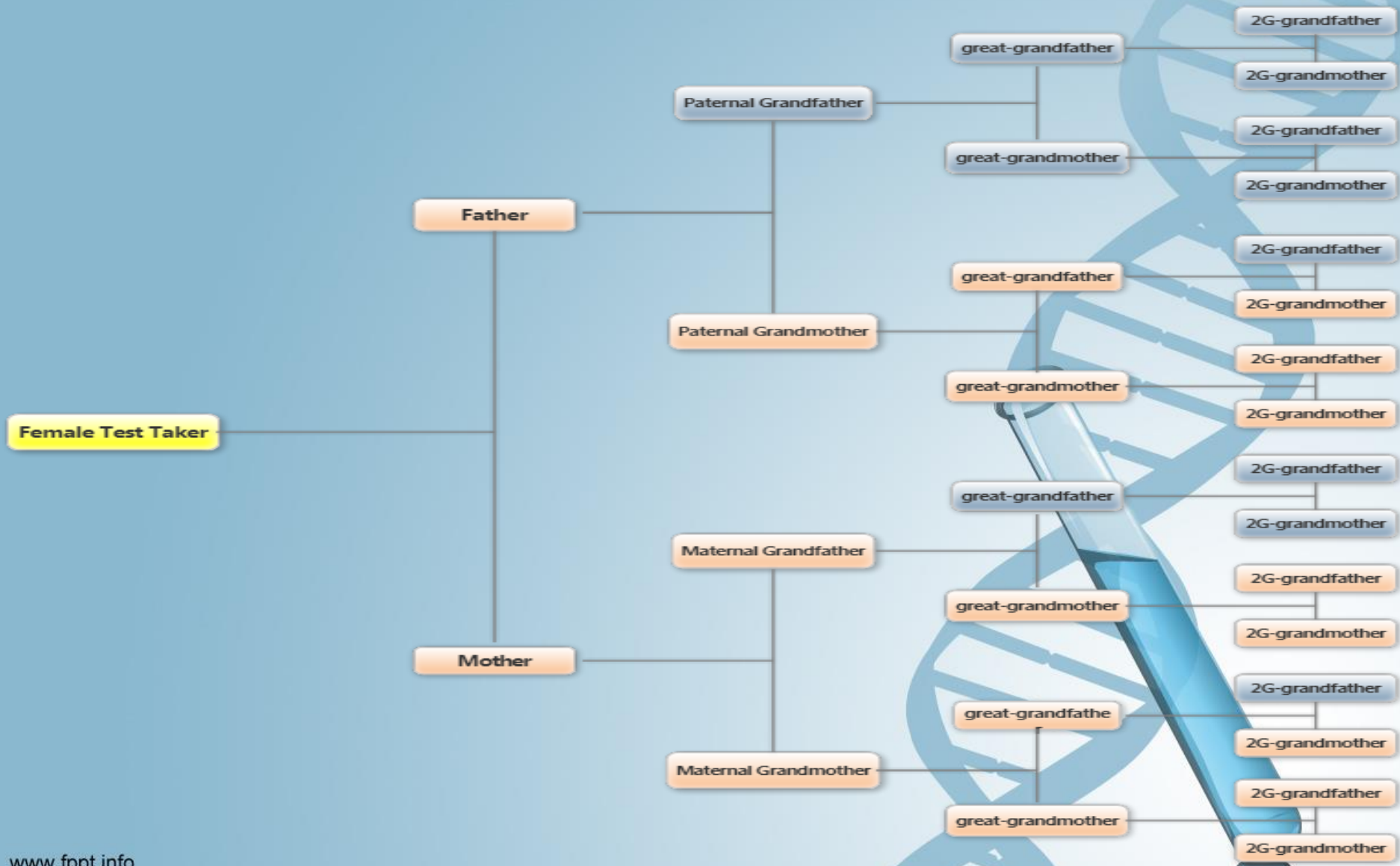
Female



Male X-DNA Inheritance Pattern



Female X-DNA Inheritance Pattern



Phasing

Definition:
Separating Paternal
from Maternal
Matches



Centimorgan (cM)

Definition: A measurement of genetic distance. It implies distance along a chromosome.

It tells us how much DNA we share with a match.



$$20 + 30 + 10 = 60$$



DNA Segments

The number of areas on a chromosome where we share DNA. This is usually expressed as the total number across all the chromosomes.



$$20 + 30 + 10 = 60$$

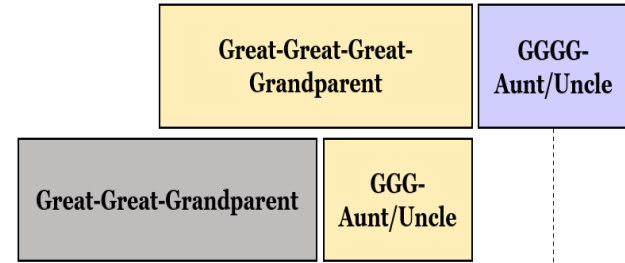
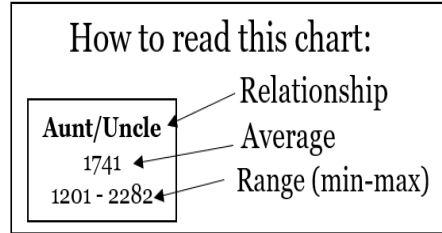


Why is the amount of cM
important?



The Shared cM Project – Version 4.0 (March 2020)

Blaine T. Bettinger
www.TheGeneticGenealogist.com
CC 4.0 Attribution License



Half GG-Aunt/Uncle 208 103 – 284	Great-Grandparent 887 485 – 1486					Great-Great Aunt/Uncle 420 186 – 713	1C3R 117 25 – 238	2c3R 51 0 – 154	Other Relationships		
Half 1C2R 125 16 – 269	Half Great-Aunt/Uncle 431 184 – 668	Grandparent 1754 984 – 2462			Great Aunt/Uncle 850 330 – 1467	1C2R 221 33 – 471	2c2R 71 0 – 244	3C2R 36 0 – 166	6C 18 0 – 71		
Half 2c1R 66 0 – 190	Half 1C1R 224 62 – 469	Half Aunt/Uncle 871 492 – 1315	Parent 3485 2376 – 3720		Aunt/Uncle 1741 1201 – 2282	1C1R 433 102 – 980	2c1R 122 14 – 353	3C1R 48 0 – 192	4C1R 28 0 – 126	6C1R 15 0 – 56	
Half 3c 48 0 – 168	Half 2c 120 10 – 325	Half 1C 449 156 – 979	Half-Sibling 1759 1160 – 2436	Sibling 2613 1613 – 3488	SELF	1C 866 396 – 1397	2c 229 41 – 592	3c 73 0 – 234	4c 35 0 – 139	5c 25 0 – 117	6C2R 13 0 – 45
Half 3c1R 37 0 – 139	Half 2c1R 66 0 – 190	Half 1C1R 224 62 – 469	Half Niece/Nephew 871 492 – 1315	Niece/Nephew 1740 1201 – 2282	Child 3487 3330 – 3720	1C1R 433 102 – 980	2c1R 122 14 – 353	3C1R 48 0 – 192	4C1R 28 0 – 126	5C1R 21 0 – 80	7C 14 0 – 57
Half 3c2R 27 0 – 78	Half 2c2R 48 0 – 144	Half 1C2R 125 16 – 269	Half Great Niece/Nephew 431 184 – 668	Great-Niece/Nephew 850 330 – 1467	Grandchild 1754 984 – 2462	1C2R 221 33 – 471	2c2R 71 0 – 244	3C2R 36 0 – 166	4C2R 22 0 – 93	5C2R 18 0 – 65	7C1R 12 0 – 50
Half 3c3R	Half 2c3R	Half 1C3R 60 0 – 120	Half GG Niece/Nephew 208 103 – 284	Great-Great-Niece/Nephew 420 186 – 713	Great-Grandchild 887 485 – 1486	1C3R 117 25 – 238	2c3R 51 0 – 154	3C3R 27 0 – 98	4C3R 19 0 – 60	5C3R 13 0 – 30	8C 11 0 – 42

Minimum was automatically set to 0 cM for relationships more distant than Half 2C, and averages were determined only for submissions in which DNA was shared

Grandparent

1754

984 - 2462

**Half
Aunt/Uncle**

871

492 - 1315

Parent

3485

2376 - 3720

Aunt/Uncle

1741

1201 - 2282

Half 1C

449

156 - 979

Half-Sibling

1759

1160 - 2436

Sibling

2613

1613 - 3488

SELF**1C**

866

396 - 1397

Half 1C1R

224

62 - 469

**Half
Niece/Nephew**

871

492 - 1315

Niece/Nephew

1740

1201 - 2282

Child

3487

3330 - 3720

1C1R

433

102 - 980

Half 1C2R

125

16 - 269

**Half Great
Niece/Nephew**

431

184 - 668

**Great-
Niece/Nephew**

850

330 - 1467

Grandchild

1754

984 - 2462

1C2R

221

33 - 471

Ancestry.com

- Total amount of DNA shared (cM)
- Total number of segments shared
- The amount of X-DNA is included in the total amount of DNA shared and total segments.
- No specific X-DNA information given
- No chromosome information provided



















Ancestry-Matches

< View Another Test

Philip Spivey's DNA Matches

List  Map

Filter by: Unviewed Common ancestors Messaged Notes Trees Shared DNA Groups Search | Sort

Close Family			
 Nona Bertschy	Close Family-1st Cousin Shared DNA: 1,870 cM across 55 segments	 Unlinked Tree	 Add/edit groups
 M.N. Managed by davis194716	Close Family-1st Cousin Shared DNA: 1,553 cM across 56 segments	 No Trees	 Add/edit groups
2nd Cousin			
 melissa hughes	1st-2nd Cousin Shared DNA: 541 cM across 23 segments	 2,348 People  Common ancestor	 Add/edit groups
 M L	1st-2nd Cousin Shared DNA: 524 cM across 24 segments	 No Trees	 Add/edit groups
 cbyrd42	1st-2nd Cousin Shared DNA: 459 cM across 25 segments	 No Trees	 Add/edit groups

< Back

Message

Tools



You and Nona Bertschy

Predicted relationship: Close Family-1st Cousin

Shared DNA: 1,870 cM across 55 segments

Add/edit groups

Add note

Trees

Ethnicity

Shared Matches



< Back

Message Tools



You and Nona Bertschy

Predicted relationship: **Close Family-1st Cousin**
Shared DNA: 1,870 cM across 55 segments

- + Add/edit groups
- + Add note

Trees Ethnicity Shared Matches

Filter by: Unviewed Common ancestors Messaged Notes Trees Groups

Search | Sort

Close Family

M.N. **Close Family-1st Cousin** No Trees Add/edit groups
Managed by davis194716 Shared DNA: 1,553 cM across 56 segments

2nd Cousin

2nd-3rd Cousin Unlinked Tree Add/edit groups
Shared DNA: 323 cM across 22 segments

2nd-3rd Cousin No Trees Add/edit groups
Shared DNA: 270 cM across 11 segments

3rd Cousin

3rd-4th Cousin Unlinked Tree Add/edit groups
Shared DNA: 181 cM across 9 segments

3rd-4th Cousin No Trees Add/edit groups
Shared DNA: 102 cM across 5 segments



















Ancestry-Matches

< View Another Test

Philip Spivey's DNA Matches

List  Map

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My Quick Links



+ Add Quick Links

Philip's updates

Tutorial **NEW**

Refund Policy updated on October 21, 2019 **IMPORTANT**

[Complete Family Ancestry survey](#)

myProjects

[Cobb & Cobbs](#)

[English Mapping](#)

[H and HV mtDNA Hg](#)

[H4 mtGenome](#)

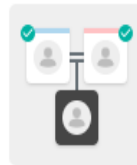
[Ireland mtDNA](#)

[Irish Mapping](#)

[R_R1b ALL Subclades](#)

MANAGE PROJECTS

JOIN A PROJECT



Family Ancestry

Family Finder

See the percentage breakdown of your origins as well as your ancient origins, and connect with your autosomal DNA relatives on all of your ancestral lines within the last 5 generations.

Results Completed: August 26, 2014



Matches



myOrigins



Matrix



Data Download



Chromosome Browser



ancientOrigins



Advanced Matches

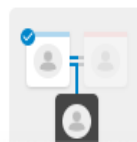


Learn More

Partner Services



Vitagene Health Report



Paternal Ancestry **R-BY134644**

Y-DNA

Follow the migration paths of your paternal line's ancestors, and connect with your Y-DNA

Y12 **Y25** **Y37** **Y67** **Y111** **Big Y**

FTDNA-Matches



HOME myDNA myTREE myPROJECTS

ADD ONS & UPGRADES



Philip Spivey
Kit No. 359985

Family Finder - Matches














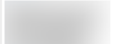






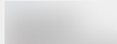






Most Common Surnames: 61 Smith 48 Johnson 36 Brown

[Advanced Search](#)

Chromosome Browser In Common With Not In Common With [Reset Filter](#)

1-30 of 5728 << < > >> Page 1 / 191 Go

All (5728) Paternal (0) Maternal (858) Both (0)

	Name	Match Date	Relationship Range	Shared cM	Longest Block	X-Match	Linked Relationship	Ancestral Surnames	
<input type="checkbox"/>	 Nona Bertschy    	07/02/2015	Half Sister, Grandmother/ Granddaughter, Aunt, Niece	1,878	95	<input checked="" type="checkbox"/> X-Match	Sister		
<input type="checkbox"/>	 Matt Byrd   	02/11/2017	2nd Cousin - 3rd Cousin	181	37				
<input type="checkbox"/>	    	02/18/2020	2nd Cousin - 3rd Cousin	139	45			Bialik (Poland) / D'Orazio (Italy) / Guglielmo (Italy) / Dudley (New York)	
<input type="checkbox"/>	     	01/04/2020	2nd Cousin - 4th Cousin	108	32	X-Match			

 Help

Chromosome Browser

A representation of 22 chromosomes showing the areas of shared DNA. The browser may also include the X chromosome. (usually not phased)



Philip Spivey **YOU**

Chromosome View Detailed Segment Data

With

Nona Bertschy
Half Sister, Grand...

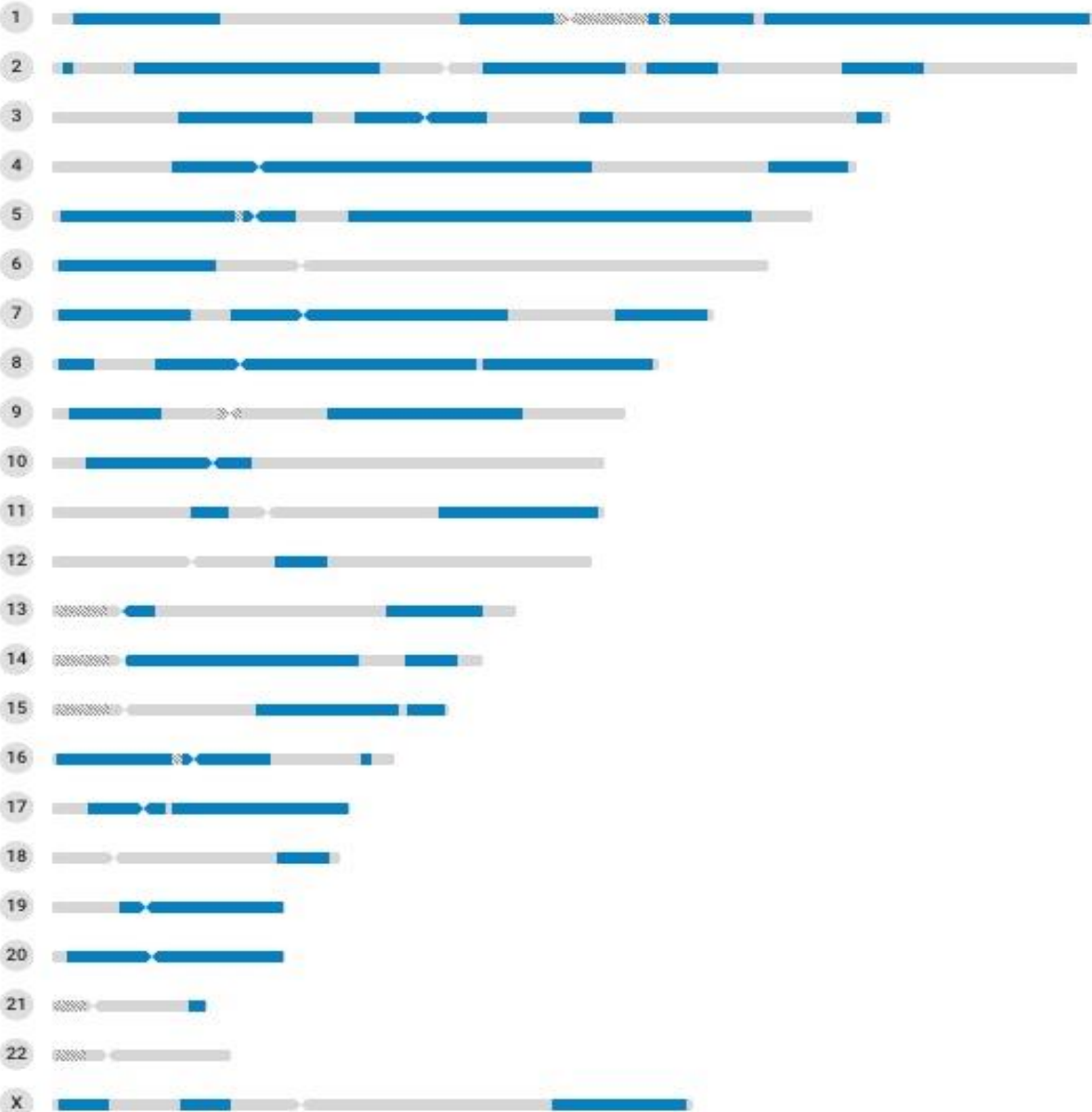
5+ centimorgans

Download Segments

Selected 1/7

Clear All

Update Selected Matches



23andMe-Matches

[HOME](#)[ANCESTRY](#)[HEALTH & TRAITS](#)[RESEARCH](#)[FAMILY & FRIENDS](#)[BUY KITS](#)**PS**

Philip ▾

[List](#)[Map](#)[Frequently Asked Questions](#)[Edit profile](#)

DNA Relatives

Get started with your predicted relationships, then connect and message to learn more.

Filters

Showing 1394 of 1394 Relatives

Sort by

Strength of Relationship ▾

Search

[Reset](#)

Notifications ⓘ ▾

Profile features and activity ⓘ ▾

Ancestor birthplaces ⓘ ▾

Mom's side / Dad's side ⓘ ▾

Connections ⓘ ▾



Connected

Niece

11.2% DNA shared, 31 segments



Invitation sent

First Cousin

7.52% DNA shared, 25 segments

**melissa Hughes**
Connected

First to Second Cousin

7.00% DNA shared, 17 segments



Connected

Second Cousin

4.22% DNA shared, 11 segments



Connected

Second Cousin

3.86% DNA shared, 14 segments



Connected

Second Cousin

3.86% DNA shared, 14 segments

23andMe-Match Profile



HOME

ANCESTRY

HEALTH & TRAITS

RESEARCH

FAMILY & FRIENDS

BUY KITS



PS

Philip

Sex: Female
Birth Year: 1974
Location: [Redacted]

Send a message

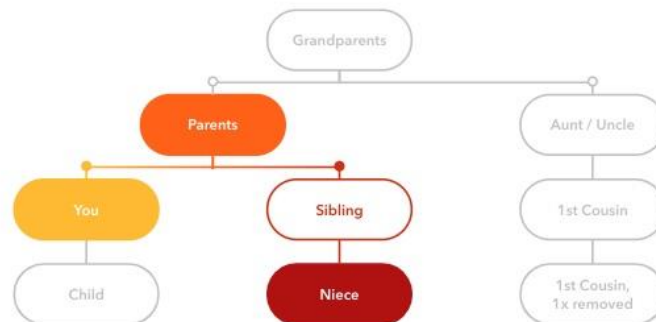
Your genetic relationship ⓘ

Relationship 🏠

Niece

[Edit Relationship](#)

You and Christine share DNA that was passed down from your parents (her grandparents).



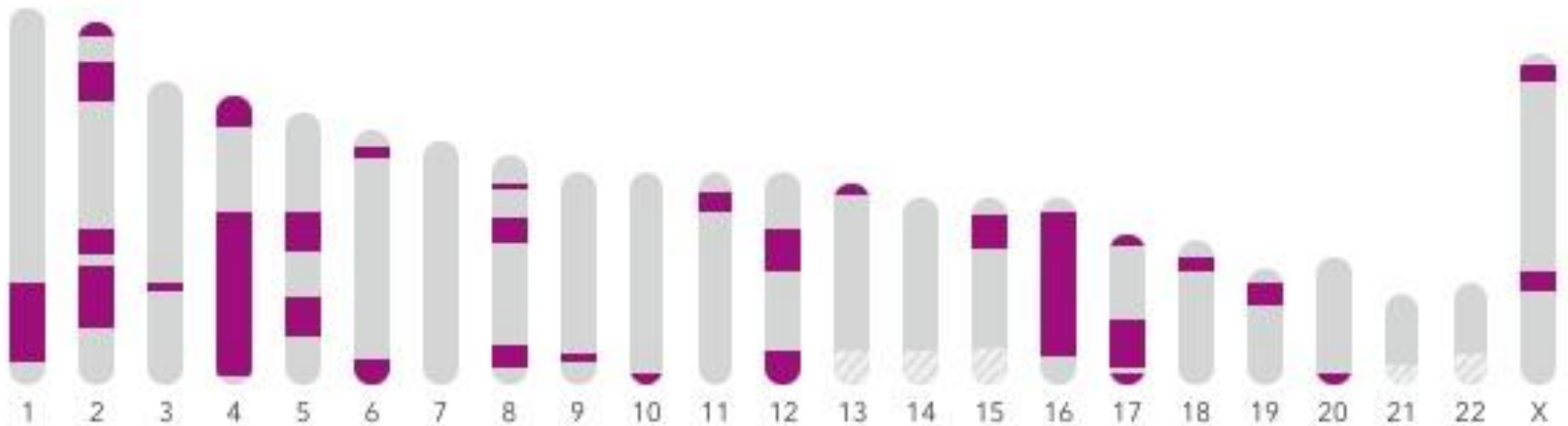
[View DNA details](#)

23andMe-Browser

Shared DNA

11.2%

836cM

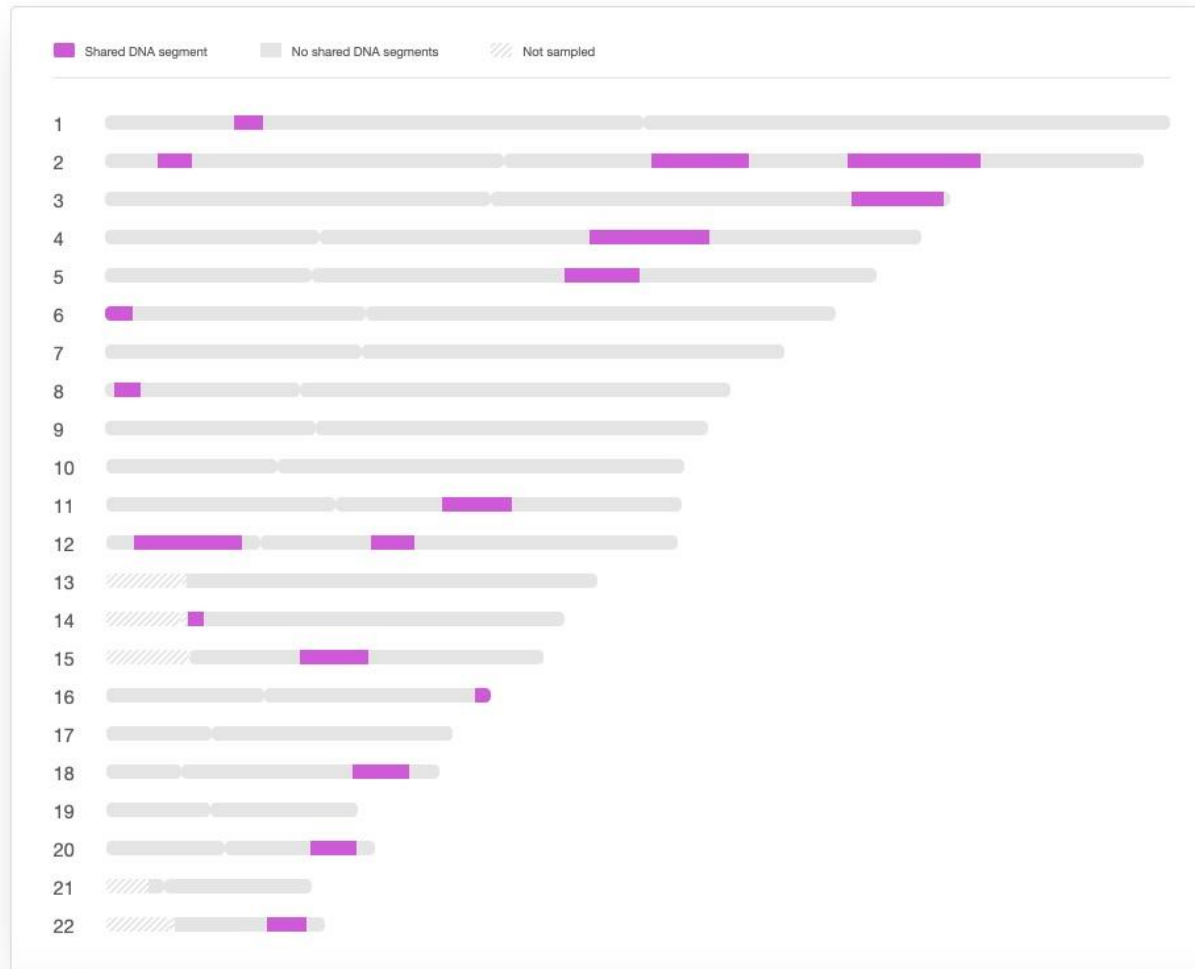


● Completely identical ● Half identical ● Not identical ▨ Not enough information

[Compare with more relatives](#)

[Hide DNA details](#) ^

MyHeritage-Browser





Welcome to GEDmatch. If you are already registered, just enter your login ID and password in the place provided below.

If you are not previously registered, click on the registration link near the bottom of this page. You will be asked to provide a valid email address and a password. After you are registered, you will be required to accept the GEDmatch Terms of Service. You can read the Terms of Service before you register by clicking on the link at the bottom of this page.

GEDmatch provides applications for comparing your DNA test results with other people. There are also applications for estimating your ancestry. Some applications are free. More advanced applications require membership in the GEDmatch Tier1 program at \$10 per month.

If you have any questions or comments, you can reach us at GEDmatch@verogen.com

Log In

Email Address:

Password:

Not Registered? Click [HERE](#) to register at the main site - when registered return to this page to access www.gedmatch.com

Clustering

Putting matches into groups with similar characteristics.

- Phasing matches (simplest form of clustering)
- Sub-clusters
 - Family groups
 - DNA



Phasing Genetic Networks (Clusters)

- Automatic Clustering
 - GEDmatch-Autoclustering
 - MyHeritage-Autocluster
 - Genetic Affairs
 - Ancestry ThruLines
- Manual Clustering
 - Dana Leeds method



User Profile (113035):

Name: Philip Spivey

Email: pcspivey@gmail.com

Tier 1 Member

Tier 1 Paid Until 2020-03-21

[View/Change/Delete your profile \(password, email, groups\)](#)

[Change EU/Unknown kit status](#)

The number of online users is 134

LEGEND:

(Status indicators shown to the right of each kit below)

✓	Kit has completed all processing and has good status
✎	Click on pencil if you wish to EDIT or DELETE kit profile
2	Likely duplicate - may need to be deleted
R	Research kit
?	Unknown Status

Click on blue kit number to go directly to one-to-many results

Your DNA resources:

M696320	R ✓	Philip Spivey	✎
A809522	R ✓	Philip Spivey	✎
A333798	R 2	Philip Spivey	✎
T978605	R ✓	Philip Spivey	✎
A314348	R ✓	Nona Bertschy	✎
A213141	R ✓	Paula Stringer	✎
T575566	✓	Matt Byrd	✎ 
A210918	R ✓	Barbara Spivey	✎
XL194899C1	y ✓	Philip Spivey	✎ 

You have not uploaded any GEDCOM (genealogy) resources

Information:

- [User Lookup](#) - Find information on your matches.
- [How to use GEDmatch](#)
- [GEDmatch Terms of Service](#)
- [GEDmatch info about you](#)
- [GEDmatch Wiki](#)
- [Useful Videos](#)
- [Support Request](#)

Upload your DNA files:

- [Generic Uploads](#) (23andme, FTDNA, AncestryDNA, most others)

DNA Applications:

- [One-To-Many Beta](#) - give it a try
- [One-To-Many DNA Comparison Result](#)
- [One-to-One Autosomal DNA Comparison](#)
- [One-to-One X-DNA Comparison](#)
- [Admixture \(heritage\)](#)
- [Admixture / Oracle with Population Search](#)
- [People who match both, or 1 of 2 kits](#)
- [DNA File Diagnostic Utility](#)
Analyze DNA file upload for potential problems.
- [Are your parents related?](#)
- [3-D Chromosome Browser](#)
- [Archaic DNA Matches](#)
- [Ancestor Projects NEW](#)

GEDmatch Forums

- [Gedmatch Forums](#) - Starting over!

Tier 1

- [One-To-Many DNA Comparison Beta](#)
- [One-To-Many DNA Comparison](#)
- [Q-Matching One-To-One](#)
- [Segment Search](#)
- [Phasing](#)
- [Triangulation](#)
- [Multiple Kit Analysis \(MKA\)](#)
- [Lazarus](#)
- [My Evil Twin](#)
- [Combine multiple kits into 1 superkit NEW!](#)
- [Clusters, Single Kit input, Basic Version NEW!](#)

GEDmatch® Auto-Clustering

Here is a link to a useful YouTube video on using the Tier 1 Clustering tool.

Reference Kit: XL194899C1 (Philip Spivey)

425 Kits

144 Clusters

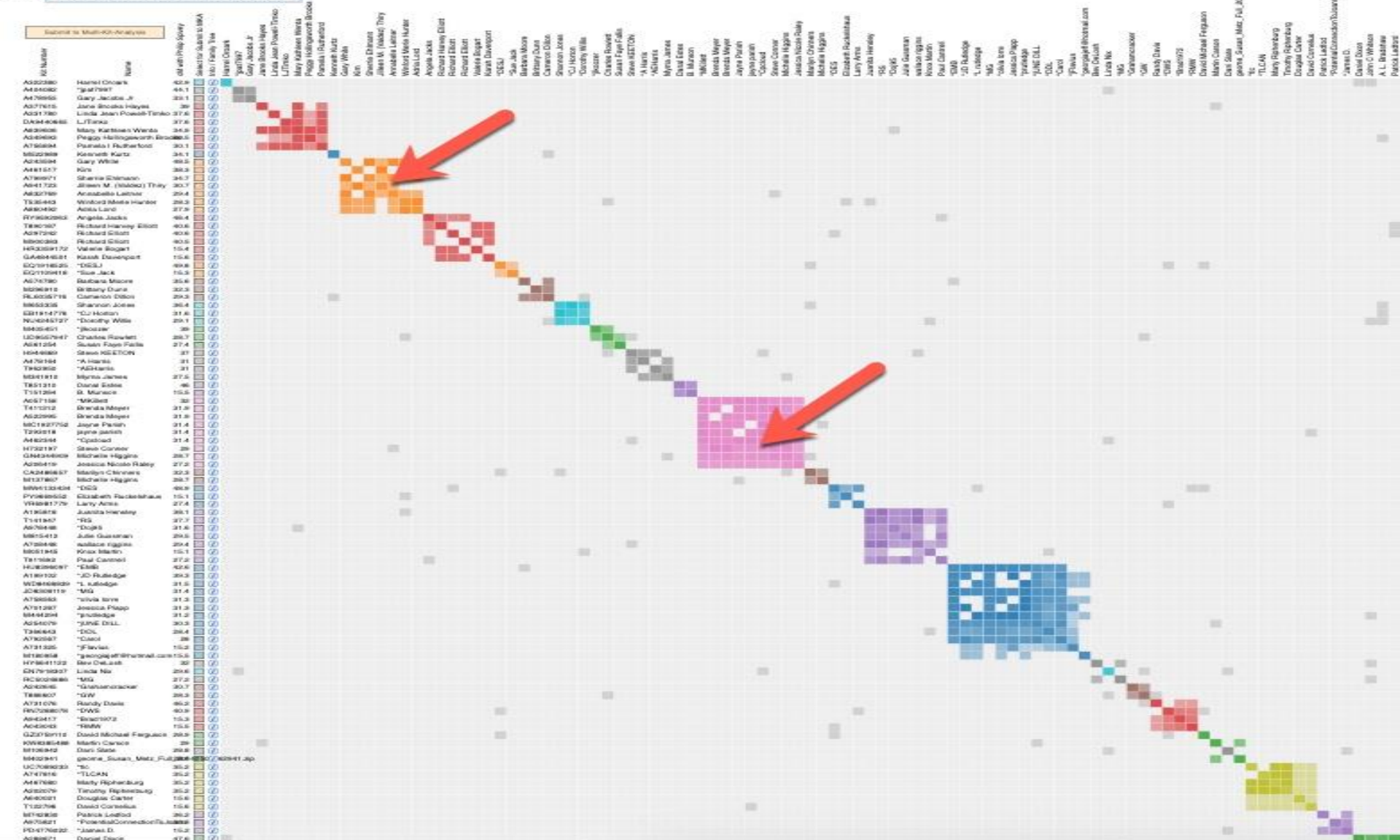
Lower cM threshold= 15

Upper cM threshold= 50

The AutoCluster feature on GEDmatch® was developed in collaboration with Evert-Jan Blom from Genetic Affairs

2020-03-03

Order: **by Cluster avg. cM to Ref. Kit**



Things to keep in mind

- For males – all X-DNA matches are maternal
- Not all maternal matches are X-DNA matches
- X-DNA phasing works even when we don't know any of the people on our match list



Questions



pcspivey@gmail.com

