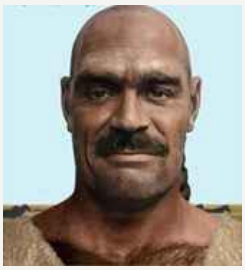
FABRICE ANCESTRY REPORT

Yours Simulated G25 Explore Your DNA coordinates are :
Fabrice_scaled,0.132035,0.142174,0.051288,0.028101,0.033237,0.009482,0.00376,0.009923,0.007976,0.013485,-0.006496,0.005845,-0.008176,-0.007707,0.021715,-0.005569,-0.021513,0.005574,0.010056,0.001126,-0.001123,0.006183,0.000493,0.015062,0.002515
NB : please remember yours reals coordinates are the ones that will provide the most accurate results. To order your reals coordinates, please contact teepean47 (partner of Daviski), the cost is $15 if you use Wise directly with the link and transfer unless you are using USD or CAD with Swift/wire payments in which case you have to add the transfer costs to the amount. For credit card payments you can use Stripe. The price of 15€ covers the costs from Stripe : https://buy.stripe.com/bIYaFJ2qocAK4Rq9AB

Your ancient break down is :

Yamnaya_RUS_Samara:



The Yamnaya culture, which thrived on the Pontic-Caspian steppe from around 3300 to 2600 BCE, has been extensively studied through ancient DNA analysis. The genetic makeup of the Yamnaya people is particularly significant because it has had a lasting impact on the genetic landscape of Europe and parts of Asia.

Genetic studies have shown that the Yamnaya people were a mix of Eastern European Hunter-Gatherers (EHG) and Caucasus Hunter-Gatherers (CHG) in roughly equal proportions

- This blend is often referred to as 'Steppe ancestry'. The Yamnaya also had some genetic contributions from Anatolian farmers, which they likely acquired through interactions with neighboring populations

- Y-chromosome haplogroups commonly found in Yamnaya individuals include R1b and I2, which are prevalent in modern European populations

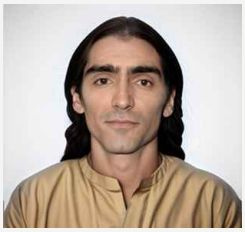
- Mitochondrial DNA (mtDNA) haplogroups such as U5, H, and T have also been identified, indicating diverse maternal lineages

- The Yamnaya culture is closely linked to the spread of Indo-European languages and had a significant influence on subsequent cultures, such as the Corded Ware and Bell Beaker cultures

- Their migrations and interactions with other groups played a crucial role in shaping the genetic and cultural landscape of prehistoric Europe.

44.91 %

TUR_Barcin_N :



The DNA sample referred to as TUR_Barcin_N comes from the Barcın Höyük archaeological site in Turkey. This sample is associated with the Neolithic period, dating back to approximately 6500–6200 BCE. It represents the genetic profile of early Anatolian farmers, who played a significant role in the spread of agriculture into Europe.

The genetic makeup of this sample shows a strong connection to the Anatolian Neolithic population, with some links to early European farmers. The Y-DNA haplogroup is often associated with G2a, and the mtDNA haplogroups vary but are typically linked to early farming communities.

44.43 %

WHG:

Western Hunter-Gatherers (WHG) were a group of Mesolithic hunter-gatherers who lived in Europe from around 15,000 to 5,000 years ago. Their genetic ancestry is a significant component of modern European populations1

.Genetic studies have shown that WHGs had a distinct genetic profile, with Y-chromosome haplogroups such as I2 and mitochondrial DNA (mtDNA) haplogroups like U5 and U41

. These haplogroups are still present in modern European populations, indicating the lasting genetic legacy of WHGs.

WHGs were part of a broader network of hunter-gatherer groups in Europe, including Eastern Hunter-Gatherers (EHG) and Scandinavian Hunter-Gatherers (SHG). The interactions and migrations of these groups contributed to the complex genetic landscape of ancient Europe

10.66 %

All the reports below are based on Euclidian distance, please use the legend below to evaluate if your genetic distance is good. Remember that most ethnic reports are based on "genetic similarity" and not "direct ancestry".

GREAT	GOOD	AVERAGE	DISTANT		VERY DISTANT				

Your modern break down is :	
How to interpret ? This report is trying to break your ancestry using modern references, if you are mixed, you may see your differents ancestries appearing on this report, else it may show something more ancient.	
Fit :	0.019816395555487032
English_Cornwall :	79.81 %
Italian_Liguria :	8.86 %
Spanish_Pais_Vasco :	5.03 %
Samaritan :	4.40 %
Azerbaijani_Turkey :	1.90 %

Your modern Breakdown Interpretation (by OpenAI) :

Your Y-DNA haplogroup **R1b-M405** (also known as **R1b-U106**) is a subclade of **R1b**, which is the most common paternal haplogroup in Western Europe. **R1b-M405/U106** is particularly associated with **Northwestern Europe**, especially among **Germanic-speaking populations** (e.g., English, Dutch, Germans, and Scandinavians). In France, it is more common in the **northern regions** (such as Normandy and Flanders) than in the south.

Interpreting Your Modern Breakdown Your autosomal DNA results suggest a strong **Northwestern European** component, with some **Southern European** and minor **West Asian/Middle Eastern** influences. Let's analyze each component:

1. English_Cornwall (79.81%) - Cornwall is in **Southwest England**, but this high percentage likely reflects a broader **Northwestern European genetic profile**. - This could indicate strong genetic ties to **Anglo-Saxon, Norman, or Flemish ancestry**, which are historically present in **Northern France**. - Given your **R1b-M405/U106** haplogroup, this suggests a possible **Germanic influence** (Anglo-Saxon, Frankish, or Flemish).

2. Italian_Liguria (8.86%) - Liguria is in **Northwest Italy**, home to **Genoa**. - This could reflect **historical migration** from Italy to France, possibly through **Roman influence** or later medieval trade connections. - Northern France had some **Italian migration** during the Middle Ages and Renaissance.

3. Spanish_Pais_Vasco (5.03%) - The **Basques** are an ancient population with **pre-Indo-European roots**. - This could indicate **shared ancient ancestry** between **Northern France and the Basque region**, as both have high levels of **R1b**. - Alternatively, it could reflect **historical migration** from Spain to France.

4. Samaritan (4.40%) - Samaritans are a small ethnic group in the **Middle East**, primarily in Israel and Palestine. - This could be a **trace amount of ancient Middle Eastern ancestry**, possibly from the **Roman period, Crusades, or Jewish diaspora**.

5. Azerbaijani_Turkey (1.90%) - This likely represents a **very distant West Asian or Turkic-related ancestry**. - It could be linked to **Neolithic, Indo-European, or even medieval migrations**. - Some **Crusaders** and **Ottoman interactions** may have introduced minor West Asian DNA into France.

Conclusion Your DNA results suggest a **strong Northwestern European genetic profile**, consistent with someone from **Northern France**. The **high English-Cornwall percentage** likely reflects a broader **Northwest European or Germanic influence**, which aligns with your **R1b-M405/U106 haplogroup**. The **Italian and Basque components** could represent **shared ancient ancestry** or **historical migrations**. The **Middle Eastern and West Asian traces** are likely from **ancient or medieval influences**.

Would you like a deeper historical analysis of your haplogroup or autosomal DNA?

Your ancient break down is :	
How to interpret ? This report is trying to break your ancestry using ancient references.	
Fit :	0.013180482378149248
Insular*Celt*(AD*100-1000) :	25.44 %
Pict*(AD*300-500) :	25.44 %
Germanic*(AD*100-630) :	16.03 %
Italian*(AD*650-1450) :	10.83 %
Italic*and*Etruscan*(900-200*BC) :	10.32 %
France*(AD*130-1400) :	3.59 %
Roman*Gaul*(AD*130-500) :	3.59 %
Arabian*Peninsula :	1.97 %
Roman*Pannonia*(AD*130-600) :	1.22 %
Germanic*(AD*700-1000) :	1.10 %

Your Ancient Breakdown Interpretation (by OpenAI) :

Your Y-DNA haplogroup **R1b-M405** (also known as **R1b-U106**) is commonly associated with **Northwestern Europe**, particularly among Germanic-speaking populations. This haplogroup is often linked to early **Germanic tribes**, such as the Saxons, Frisians, and other groups that migrated across Europe during the early medieval period.

Interpreting Your Modern Breakdown: Your genetic breakdown suggests a **strong Northwestern European** ancestry, with significant contributions from **Celtic, Pictish, and Germanic populations**, as well as some Roman and Italic influences. Let's analyze the main components:

1. **Insular Celtic (AD 100-1000) – 25.44%** - This category likely represents ancestry from the ancient **Brittonic and Gaelic-speaking peoples** of the British Isles. - It suggests a connection to populations from **Ireland, Scotland, or Wales**, possibly due to medieval migrations or earlier Celtic expansions.

2. **Pict (AD 300-500) – 25.44%** - The Picts were an ancient people of **Scotland**, known for their resistance to Roman rule. - This high percentage suggests a strong **Northwestern European** influence, possibly linked to **early Britons or migrations between Britain and Northern France**.

3. **Germanic (AD 100-630) – 16.03%** - This reflects ancestry from early **Germanic tribes**, such as the **Franks, Saxons, or Alemanni**. - Given that you are from **Valenciennes, Northern France**, this is expected, since the region was historically settled by the **Franks**.

4. **Italian (AD 650-1450) – 10.83%** - This could be linked to **Roman influence** or later medieval migrations from Italy. - It may also reflect **Lombardic or other Southern European contributions**.

5. **Italic and Etruscan (900-200 BC) – 10.32%** - This suggests deep genetic ties to the ancient **Italic peoples**, including the **Romans and Etruscans**. - It may be linked to the **Roman Empire's expansion** into Gaul.

6. **France (AD 130-1400) – 3.59%** - This likely represents **general French ancestry**, incorporating various historical populations.

7. **Roman Gaul (AD 130-500) – 3.59%** - This indicates ancestry from the Romanized populations of **Gaul**, which included a mix of **Celtic, Roman, and Germanic peoples**.

8. **Arabian Peninsula (1.97%)** - This could reflect **minor genetic contributions** from **Moorish, Arab, or Near Eastern influences**, possibly from medieval trade or migrations.

9. **Roman Pannonia (AD 130-600) – 1.22%** - Pannonia was a Roman province in **Central Europe** (modern-day Hungary, Austria, and the Balkans). - This could reflect minor Roman-era migrations into Gaul.

10. **Germanic (AD 700-1000) – 1.10%** - This may represent later **Viking or Frankish influences** in Northern France.

Conclusion: Your genetic breakdown is consistent with someone from **Northern France**, particularly **Valenciennes**, which historically had strong **Frankish, Celtic, and Roman influences**. The high **Insular Celtic and Pictish** percentages suggest possible links to **Britain or Ireland**, which could be due to **medieval migrations, trade, or earlier Celtic expansions**. Your **Germanic and Italic components** reflect the **Frankish and Roman** history of the region.

Would you like to explore any specific aspect in more detail?

Your 50 closests modern populations are :	
How to interpret ? if you obtain a good distance with your first population (e.g bright green), you could very likely be from that ethnicity or a close ethnicity, else you are most probably mixed or your ethnicity is not referenced on Davidski datasheet.	
Walloon	0.021912328265157
BelgianA	0.0225442671204899
Côtes d'Armor (France)	0.0236959019430939
Pas-de-Calais (France)	0.0250498414837481
French_Oil_Hauts-de-France_Nord	0.0251358169702382
French_Oil_Normandy_Manche	0.0258063051272359
Nord (France)	0.025975626460973
Manche (France)	0.0262562405641401
Belgian	0.0263562869332455
French_Nord	0.0267883033990583
German_Baden-Wurttemberg_Ellwangen	0.0270601991538457
French_Oil_Hauts-de-France_Pas-de-Calais	0.0272769202513506
Seine Maritime (France)	0.0273825957862567
BelgianC	0.0278574392577638

Anglo-American	0.0280090848589737
Maine-et-Loire (France)	0.0281704826182509
English_Cornwall	0.0286328807841614
French_Canadian	0.0289976052246769
Flemish_Belgium	0.0290340033624369
Cornish	0.0290762359532536
French_Alsace	0.0290886910843372
BelgianB	0.02910319668009
Ardennes (France)	0.0292980167929503
French_Oil_Pays_de_la_Loire_Maine-et-Loire	0.0293228518437413
Alsatian_Bas-Rhin	0.0294143663787164
Doubs (France)	0.0295301311544666
French_Oil_Ile-de-France_Paris	0.0298265160305779
French_Paris	0.0298267331600361
French_Brittany	0.0299404679155153
French_Oil_Bourgogne-Franche-Comte_Yonne	0.0302174057622424
Haute Saône (France)	0.0303979185800607
Rhône (France)	0.0304588228761389
English	0.0304879052412592
French_Oil_Normandy_Seine-Maritime	0.0306505991461362
Swiss_German	0.0308741337206407
Welsh	0.031183698658113
Dutch_North_Brabant	0.0312916359911079
Eure (France)	0.0314203326366861
Vosges (France)	0.0316272743055737
Jura (France)	0.031866370408316
Somme (France)	0.0320967878299371
French_Seine-Maritime	0.0331033638623026
Scottish	0.0335189676004498
French	0.0335925621303017

Yonne (France)	0.0336664529168132
Ille-et-Vilaine (France)	0.0339255673497143
Dutch	0.0340901858311157
Vendée (France)	0.0342773469506954
German	0.0343687796117348
Finistère (France)	0.0345885383265496

Your 50 closests modern 2-Ways are :	
The 2Ways compares your coordinates to the ones of all referenced populations within the modern datasheet of Davidski. It measures how closely your admixture percentages are aligned with each population. It's important to understand that 2Ways aren't measuring shared DNA between your kit and referenced samples. If you are from one ethnicity you should only concentrate on yours closests populations, else if you are mixed (e.g parents from differents ethnicities), yours closests 2Ways should be close to "50% Parent 1 Ethnicity - 50% Parent 2 Ethnicity". if you are more mixed (for example from latina America), your 2Ways is not going to be accurate.	
75% English_Cornwall + 25% Italian_Lombardy	0.0199734013618374
75% English_Cornwall + 25% Italian_Liguria	0.0204084239880631
90% English_Cornwall + 10% Samaritan	0.0204177582958674
87% English_Cornwall + 13% Sephardic_Jew	0.0204736080993145
70% Scottish + 30% Italian_Lombardy	0.0205781097285587
73% English + 27% Italian_Lombardy	0.020748546821028
84% English_Cornwall + 16% Sicilian_East	0.0208474262095306
84% English_Cornwall + 16% Italian_Apulia	0.0209325746258408
81% English_Cornwall + 19% Italian_Marche	0.0209330242363522
89% English_Cornwall + 11% Greek_Cappadocia	0.0209499737862459
89% English_Cornwall + 11% Cypriot	0.0210042949420761
89% English_Cornwall + 11% Mountain_Jew_o	0.0210529622367908
85% English_Cornwall + 15% Ashkenazi_Poland	0.0210703407512265
83% English_Cornwall + 17% Italian_Abruzzo	0.021092953539196
85% English_Cornwall + 15% Ashkenazi_Ukraine	0.0210951509298275
85% English_Cornwall + 15% Italian_Campania	0.0210963577027587
60% Icelandic + 40% Italian_Lombardy	0.0211636684456981

87% English_Cornwall + 13% Italian_Jew	0.0211671420137619
87% English_Cornwall + 13% Greek_Dodecanese	0.0211778306365699
74% English + 26% Italian_Liguria	0.0212263034814094
90% English_Cornwall + 10% Karaite_Egypt	0.0212972162435957
90% English_Cornwall + 10% Lebanese_Christian	0.0213339271972396
88% English_Cornwall + 12% Romaniote_Jew	0.0213824367806341
86% English_Cornwall + 14% Greek_Crete	0.0214109111465368
70% Scottish + 30% Italian_Liguria	0.0214532798769985
81% English_Cornwall + 19% Greek_Thessaly	0.0214703486175932
90% English_Cornwall + 10% Druze	0.021477394862081
78% English_Cornwall + 22% Italian_Piedmont	0.0214793104746507
83% English_Cornwall + 17% Italian_Lazio	0.021479493102636
84% English_Cornwall + 16% Sicilian_West	0.0214974180366758
85% English_Cornwall + 15% Italian_Basilicata	0.0215025632719128
86% English_Cornwall + 14% Italian_Calabria	0.021520958054983
89% English_Cornwall + 11% Tunisian_Jew	0.0215350188446314
84% English_Cornwall + 16% Italian_Molise	0.0215430746265132
89% English_Cornwall + 11% Syrian_Jew	0.0215678607956484
90% English_Cornwall + 10% Lebanese_Druze	0.0215806292381548
69% Orcadian + 31% Italian_Lombardy	0.0216132608548301
85% English_Cornwall + 15% Ashkenazi_Russia	0.0216590250365364
67% Irish + 33% Italian_Lombardy	0.0216834265699716
81% English_Cornwall + 19% Greek_Macedonia	0.0216888758245403
87% English_Cornwall + 13% Ashkenazi_Germany	0.0217186818046941
85% English_Cornwall + 15% Ashkenazi_Belarussia	0.0217403776939801
84% English_Cornwall + 16% Ukrainian_Zhytomyr_o	0.0217751360110489
86% English_Cornwall + 14% Maltese	0.0218129152501496
80% English_Cornwall + 20% Italian_Tuscany	0.0218655734262977
91% English_Cornwall + 9% Iraqi_Jew	0.0218753115821958
91% English_Cornwall + 9% Kurdish_Jew	0.0218799338185047

82% English_Cornwall + 18% Italian_Umbria	0.0218874831727239
86% English + 14% Sephardic_Jew	0.0219039502589925
91% English_Cornwall + 9% Iranian_Jew	0.0219056017176919
76% English_Cornwall + 24% Italian_Bergamo	0.0219085277086971

Your ancient map :

How to interpret ? This map is trying to break your ancient ancestry and display on a map.

Your modern similitud map :

How to interpret ? This similitud Map is based on the modern Davidski G25 sheet + the French averages of the Explore Your DNA Project !. It's only going to be accurate for people that belong to ONE ethnicity, else it's going to show midpoints, which are the populations closest to your genetic composition. This map is a snapshot of you similitud with pre colonial period populations, means for example that people from Latin America are going to show ancestry in both Americas & Europe, same for North Americans, Australians & South Africans...Nomadic tribes and diasporas (Ashkenazis, Romas...) are not going to appear on the map so it's also going to show their ancient ancestry (e.g Ashkenazis are going to appear in both the Levant and Europe).

GREAT	GOOD	AVERAGE	DISTANT	VERY DISTANT