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- ☒ H
- ☐ H\_Subclade\_Discovery
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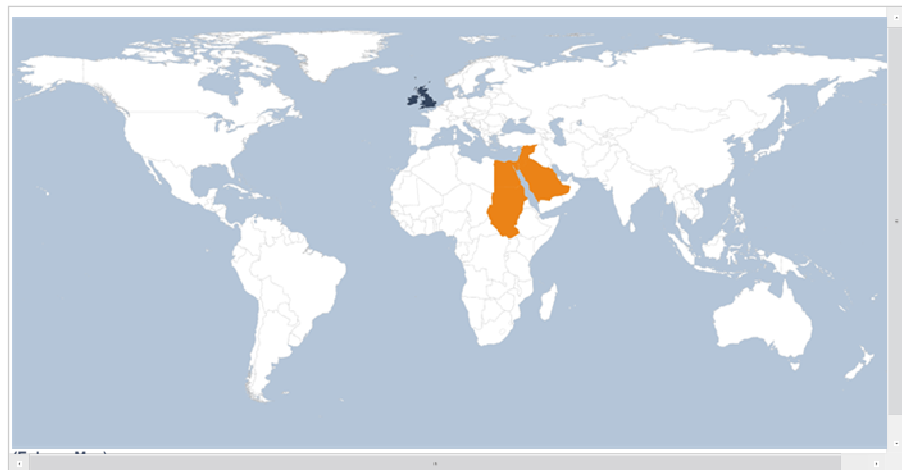
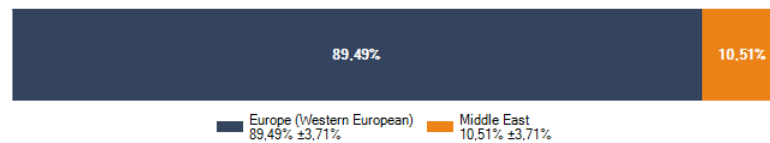
The Population Finder program determines your biogeographical ancestry — the story of your personal genetic history — by comparing your autosomal DNA to that of our world DNA population database.

Your Population Finder results consist of up to four out of seven continental groups.\* For each, the percentage of your genome that matches is shown. You may view your results as either a stacked bar or pie chart.

Where possible, we list specific subgroups such as Basque and Bedouin or subregions such as Western Europe. [More Info.](#)

\* The seven continental groups are based on genetic similarity and do not precisely match geographical continents.

\*\* With the transition to the Illumina Omni Express platform, we are now using almost three times as many data points to predict your ethnic percentages. This has enhanced our ability to refine predictions about small percentages of admixture and may reflect some changes from your previous results.

Chart Type: ☒ Stacked Bar ☐ Pie

Continent (Subcontinent)	Population	Percentage	Margin of Error
Europe (Western European)	Orcadian	89,49%	±3,71%
Middle East	Palestinian, Bedouin South, Druze, Jewish	10,51%	±3,71%

**Beta Comments**

The Population Finder program is currently in the Beta phase. Please expect changes as the Bioinformatics team responds to user feedback. You will find the answers to questions in the [FAQ](#).

Most Population Finder results are clear and unambiguous, but a small percentage of results may fall into one of the two categories below.

**What if the total percent does not add up to 100%?**

A very small amount of your DNA matches other populations. The amount of matching DNA is too small to report the matching population with confidence because the similarity with your DNA may be coincidental and falls well within the margin of error. [More info.](#)

**What does it mean if the error is very high?**

If you have a high margin of error (Example: +/-15%) then the populations you are matching are very similar to one another. In this case, we cannot distinguish as clearly between the populations, and we are less confident that some proportion of your DNA came from one population rather than the other. This does not affect our ability to predict the continent you are from, but does affect our prediction of what proportion of your ancestry is from the continents listed. [More info.](#)

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