

Pigmentation

The colors of our skin can be amazingly diverse. From dark to light to freckled, our pigmentation is heavily influenced by our DNA.

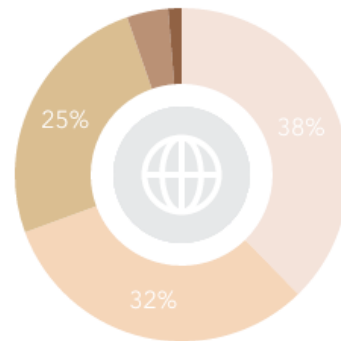
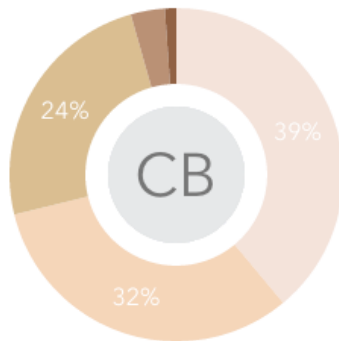
[Skin Pigmentation](#)

[Freckles](#)

[What You Can Do](#)

Cordell, you are likely to have lighter skin.

95% of customers who are genetically similar to you have fair to beige skin.



Your genetic likelihood			
95% Lighter skin		Very fair	39%
		Moderately fair	32%
		Light beige	24%
		Olive	3%
		Light brown	1%
		Dark brown	< 1%

European ancestry customers			
38%	Very fair		95% Lighter skin
32%	Moderately fair		
25%	Light beige		
4%	Olive		
1%	Light brown		
< 1%	Dark brown		

This prediction applies best to people of European and African descent. We analyzed your DNA at two genetic markers that studies have shown are associated with skin color. Your prediction is based on data from 23andMe customers who consented to research and are genetically similar to you at these markers.

About Skin Pigmentation

Darker-colored skin includes shades from olive to dark brown, while lighter-colored skin includes beige and fair skin.



Biology

Melanin, the pigment responsible for eye and hair color, also determines skin color. More melanin results in darker skin, while less melanin leads to lighter skin.



Genetics

Variants in two genes named SLC45A2 and SLC24A5 are associated with variation in skin color in people of European and African descent. These two genes are important for proper functioning of cells that produce melanin.



Other factors

Other factors can also contribute to your skin color.



History



Ancestry

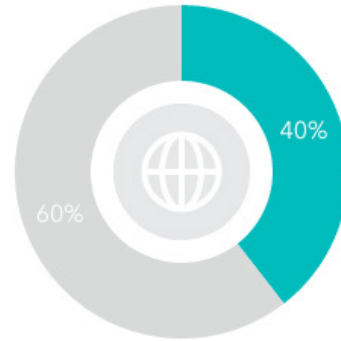
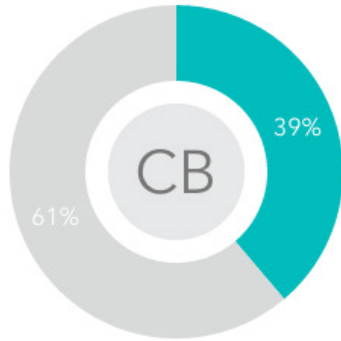


Sunlight exposure



You are not likely to have lots of freckles.

61% of customers who are genetically similar to you have only a few freckles anywhere on their bodies.



Your genetic likelihood

- A lot of freckles 39%
- Little freckling 61%

European ancestry customers

- A lot of freckles 40%
- Little freckling 60%

This prediction best applies to customers of European descent. We analyzed data from over 150,000 customers who consented to research in order to identify genetic markers associated with having freckles. Our prediction is based on your genotype at 34 genetic markers as well as your age and sex.

About Freckles

Freckles are small spots of darker skin. They are usually flat and may vary in color from red to tan or light brown.



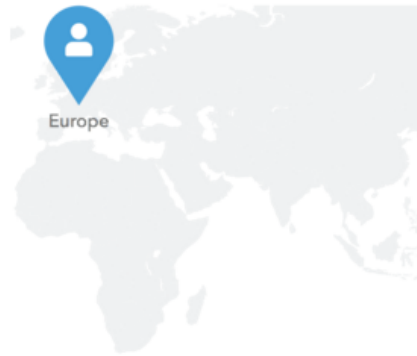
Biology

Freckles are caused by increased production of a pigment called melanin, the same type of chemical that gives color to your hair and eyes.



Ancestry

Although freckles are commonly observed in people of European descent with light skin, they may also be found in numerous populations around the world.



Other factors

Multiple factors contribute to freckling.



Genetics



Age



Sun exposure



Do more with your Traits results.



Help us develop more trait reports by contributing to research.

Contribute



Compare your results to your family and friends.

Compare



Join the discussion with other 23andMe customers interested in Traits.

Discuss

Ⓟ Patent Pending

Your Skin

Scientific Details

[Methodology](#)

[About Your Results](#)

[References](#)

We use two different methods to calculate your trait results.

Statistical Model

Most traits are influenced by many different factors, including genetics, lifestyle, and environment. Usually, a statistical model using many factors provides better predictions than looking at single factors by themselves. To develop our models, we first identify genetic markers associated with a trait using data from tens of thousands of 23andMe customers who have consented to research. Then, we use statistical methods to generate a "score" for that trait using your genotype at the relevant genetic markers as well as your age and sex. We predict your likelihood of having different versions of the trait based on the survey responses of 23andMe customers with similar scores. These predictions apply best to customers who are of the same ethnicity as the people whose data contributed to the model. The accuracy of these predictions varies from trait to trait.

[Read more about our statistical methodology](#)

Curated Model

For some traits, just a few genetic markers can strongly predict whether a person will have a particular version of the trait. For curated models, we first evaluate published scientific studies to identify genetic markers with well-established associations with the trait. Then, we look at genetic and survey data from tens of thousands of 23andMe customers who have consented to research. We estimate your likelihood of having different versions of the trait based on survey responses from customers who are genetically similar to you at those markers. These results apply best to customers who are of the same ethnicity as the people whose data contributed to the predictions.

About your Skin Pigmentation result

Your result for this trait was calculated using a **curated model**.

Variants Detected

4

View All Tested Markers

2

Marker Tested

Your Genotype*

Additional Information

A111T

Gene: SLC24A5

Marker:

[rs1426654](#)

A

Variant copy from
one of your
parents



A

Variant copy from
your other parent

> **Biological explanation**

> **Typical vs. variant DNA sequence(s)**

> **Percent of 23andMe customers with variant**

> **References [2, 5, 10, 11]**

L374F

Gene: SLC45A2

Marker:

[rs16891982](#)

G

Variant copy from
one of your
parents



G

Variant copy from
your other parent

> **Biological explanation**

> **Typical vs. variant DNA sequence(s)**

> **Percent of 23andMe customers with variant**

> **References [2, 7, 10, 11]**

*This test cannot distinguish which copy you received from which parent. This test also cannot determine whether multiple variants, if detected, were inherited from only one parent or from both parents. This may impact how these variants are passed down.

23andMe always reports genotypes based on the 'positive' strand of the human genome reference sequence (build 37). Other sources sometimes report genotypes using the opposite strand.

About your Freckles result

Your result for this trait was calculated using a **statistical model**.

About the Freckles model

Created based on customers of ethnicity: European
Number of customers used to create: 150,000
Number of markers: 34
Area Under Curve (AUC): 0.662
Non-genetic factors: Age, Sex

Bin #	A lot of freckles	Little freckling
1	15.67%	84.33%
2	19.12%	80.88%
3	22.52%	77.48%
4	23.55%	76.45%
5	26.90%	73.10%
6	27.80%	72.20%
7	30.22%	69.78%
8	31.73%	68.27%
9	34.95%	65.05%
10	36.80%	63.20%
11	38.81%	61.19%
12	40.01%	59.99%
13	42.78%	57.22%
14	45.48%	54.52%
15	47.92%	52.08%
16	50.63%	49.37%
17	54.34%	45.66%
18	58.18%	41.82%
19	62.78%	37.22%
20	73.34%	26.66%
Overall European	39.18%	60.82%

References

1. Bastiaens MT et al. (1999). "Ephelides are more related to pigmentary constitutional host factors than solar lentigines." *Pigment Cell Res.* 12(5):316-22. [↗](#)
2. Beleza S et al. (2013). "Genetic architecture of skin and eye color in an African-European admixed population." *PLoS Genet.* 9(3):e1003372. [↗](#)
3. Breathnach AS. (1957). "Melanocyte distribution in forearm epidermis of freckled human subjects." *J Invest Dermatol.* 29(4):253-61. [↗](#)
4. Chung JH. (2003). "Photoaging in Asians." *Photodermatol Photoimmunol Photomed.* 19(3):109-21. [↗](#)
5. Lamason RL et al. (2005). "SLC24A5, a putative cation exchanger, affects pigmentation in zebrafish and humans." *Science.* 310(5755):1782-6. [↗](#)
6. Nicholls EE. (1968). "Genetic susceptibility and somatic mutation in the production of freckles, birthmarks and moles." *Lancet.* 1(7533):71-3. [↗](#)
7. Norton HL et al. (2007). "Genetic evidence for the convergent evolution of light skin in Europeans and East Asians." *Mol Biol Evol.* 24(3):710-22. [↗](#)
8. Praetorius C et al. (2014). "Sun-induced freckling: ephelides and solar lentigines." *Pigment Cell Melanoma Res.* 27(3):339-50. [↗](#)
9. Roméro-Graillet C et al. (1996). "Ultraviolet B radiation acts through the nitric oxide and cGMP signal transduction pathway to stimulate melanogenesis in human melanocytes." *J Biol Chem.* 271(45):28052-6. [↗](#)
10. Tsetschlade ZR et al. (2012). "Functional assessment of human coding mutations affecting skin pigmentation using zebrafish." *PLoS One.* 7(10):e47398. [↗](#)
11. Valenzuela RK et al. (2010). "Predicting phenotype from genotype: normal pigmentation." *J Forensic Sci.* 55(2):315-22. [↗](#)