

[NevGen Predictor's Batch Prediction of Series of Haplotypes – Price.](#)

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In single batch for prediction, first 500 haplotypes cost 0.1 euro per haplotype. From haplotypes 501 up to 1000, they cost 0.06 euros per haplotype. From haplotypes 1001 up to 3000, they cost 0.04 euros per haplotype. We do not accept batches with more than 3000 haplotypes. If more than 3000 haplotypes are needed, data should be split into several batches.

That way, batch prediction of 500 haplotypes costs 50 euros, 1000 of them cost 80 euros, and 3000 of haplotypes cost 160 euros. 300 haplotypes cost 30 euros, 700 of them cost 62 euros, and so on.

Length of haplotypes in markers does not affect price of its prediction. It may be of any length, up to 111 markers. Also, haplotypes need not to be of the same length.

For the previous basic price, we send to customer only output predicted data in .TXT format. If customer also wants images with visual statistics of fitting of haplotype into predicted haplogroup, the basic price should be multiplied with **1.5**. That way, batch prediction of 500 haplotypes with images costs $50 * 1.5 = 75$ euros. This will make 1 image in PNG format for every prediction, and that is of the most probable subclade in the list of available subclades.

If customer wants us to generate images of the first 10 subclades (10 most probable subclades) for every prediction, then basic price should be multiplied with **2.2**. That way, batch prediction of 500 haplotypes with 10 images costs $50 * 2.2 = 110$ euros. Beware that we do not do this for batches with more than 500 haplotypes. Because it might lead to generation of tens of thousands of images, which would be problematic to send back to our customers. And such amount of images might need many hours to be generated on our not so fast computer.

Please keep in mind that minimal price for single batch is **25 euros**, no matter how small number of haplotypes is. This way, basic batch prediction of 100 haplotypes or 200 haplotypes costs the same, 25 euros. This is necessary because our work only partly depends of number of haplotypes. Correspondence with customers, manual conversion of data to suitable format for prediction, our application handling and sending predicted data back to customer does not depend on number of haplotypes, it is fixed. This way we discourage people sending us batches for prediction with small number of haplotypes. That they could do freely in the NevGen predictor on Internet.

Customers should pay us the price on nevski@poreklo.rs on PayPal. Beware that part of what you pay over PayPal is taken by the company as fee. So, please consult some PayPal calculator (like "salecalc.com/paypal") to ensure that what we receive is what is needed. For country, 'Serbia' should be selected.