



What is the minimum number of samples that I can order?

Illumina recommends a minimum order of 288 samples.

Will UC Davis or TraitGenetics provide a cluster file for faster genotype calling with this array?

Yes, a cluster file for the array will be provided as soon as possible after the array is available.

How do I order the array?

Email consortiummanager@illumina.com and an Illumina representative will contact you to answer any questions and help you place your order.

References

- 1. Ashrafi H, Hill T, Stoffel K, et al. De novo assembly of the pepper transcriptome (Capsicum annuum): a benchmark for in silico discovery of SNPs, SSRs and candidate genes. BMC Genomics. 2012;13:571.
2. Pepper Genome Platform (passport.pepper.snu.ac.kr/?t=PGP) Accessed 15 December 2014.

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