

[Mutation frequency]

Selection of 7–8 strains containing a SNP within a 1-kb region of genomic DNA may be obtained by screening 1,000 strains. This expected frequency is only an average; your results are not guaranteed to be identical.

Reference

Suzuki T., M. Eiguchi, T. Kumamaru, H. Satoh, H. Matsusaka, K. Moriguchi, Y. Nagato, N. Kurata. MNU-induced mutant pools and high performance TILLING enable finding of any gene mutation in rice. *Mol. Gen. Gen.*, 279: 213-223 (2008)

Recent selection frequencies

To be prepared