

Restriction Map and Multiple Cloning Site (MCS) of pHIS2 Vector. Unique restriction sites are in bold. *Note: The Sac II site is resistant to cleavage.

Description

pHIS2 is a reporter vector that can be used in yeast one-hybrid assays to identify and characterize DNA-binding proteins. The vector was specifically designed for use with the Matchmaker™ One-Hybrid Library Construction & Screening Kit (Cat. No. 630304). It contains a *HIS3* nutritional reporter gene, located downstream of a multiple cloning site (MCS) and the minimal promoter of the *HIS3* locus ($P_{\min HIS3}$). *Cis*-acting DNA sequences, or DNA target elements, can be inserted into the MCS and used as baits to screen GAL4 AD/cDNA fusion libraries for proteins that interact with the target sequence. A protein-DNA (or one-hybrid) interaction can be detected by performing the assay in a yeast strain such as Y187 that is auxotrophic for histidine. Positive one-hybrid interactions drive expression of the *HIS3* reporter gene, which enables the host cell to grow on histidine-deficient media.

In the absence of activation, the constitutive HIS3 expression from P_{minHIS3} is very low. During library screening, the leaky expression of HIS3 is controlled by adding 3-amino-1,2,4-triazole (3-AT) to the medium. The concentration of 3-AT needed to fully suppress leaky HIS3 expression must be determined empirically for each DNA target element.

pHIS2 can be maintained in both yeast and bacteria. It contains an autonomous replication sequence (*ARS4*) and *TRP1* nutritional marker for replication and selection in yeast (1, 2); it contains a Col E1 origin and a kanamycin resistance gene (Kan^r) for propagation and selection in *E. coli*. The centromeric sequence *CEN6* ensures proper segregation of the plasmid during cell division in yeast (1, 2).

pHIS2 **Vector Information**

Use

To use pHIS2 in a one-hybrid assay, clone one or more copies of a *cis*-acting DNA target sequence into the MCS. Then introduce the plasmid into competent yeast cells using the transformation protocol in the Matchmaker Library Construction & Screening Kits User Manual (PT3529-1). In contrast to the original Matchmaker One-Hybrid System, this reporter vector does not need to be integrated into the yeast genome. Instead, it is maintained as an episome throughout the assay. Inserting your target element may alter the level of background HIS3 expression. Therefore, constructs should be tested for background (leaky) HIS3 expression before you start a one-hybrid analysis. Background growth due to leaky HIS3 expression is controlled by adding 3-AT to the selection medium, as described in the User Manual (PT3529-1).

Location of Features

Multiple cloning sites: 1–41

HIS3 gene: 152–811

Fragment containing the HIS3 3' UTR & Termination sequence: 812–1446

• TRP1 gene: 2855-3529

Fragment containing the Col E1 E. coli origin of replication: 4165–4615

 Kanamycin-resistance gene: 5605–4811 CEN6/ARS4 sequences: 6254–5737

Propagation in E. coli

• Suitable host strains: DH5 α and other general purpose strains.

Selectable marker: plasmid confers resistance to kanamycin (50 μg/ml) to E. coli hosts.

E. coli replication origin: Col E1

· Copy number: low

Propagation in S. cerevisiae

Suitable host strains: AH109(MATa), Y187(MATα), Y190(MATa), SFY526(MATa), CG1945(MATa), HF7c(MATa)

Selectable marker: TRP1

S. cerevisiae origin: CEN6/ARS4

References

1. Sikorski, R. S. & Hieter, P. (1989) Genetics 122:19-27.

2. Rose, M. D. & Broach, J. R. (1991) Methods Enzymol. 194:195-230.

Note: The attached sequence file has been compiled from information in the sequence databases, published literature, and other sources, together with partial sequences obtained by Clontech Laboratories, Inc.. This vector has not been completely sequenced.

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