

Detection of physical interactions by immunoprecipitation of FLAG- and HA-tagged proteins expressed at the *his-3* locus in *Neurospora crassa*

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Protein function is often regulated through interactions with other protein(s) or by post-translational modifications. To understand these mechanisms, it is useful to utilize antibodies. However, it is not always certain whether a good antibody can be made for this purpose. The use of epitope tags eliminates the troubles associated with raising antibodies. In this report, we present a method to detect interactions between proteins by using two types of epitope-tagged proteins, FLAG- and HA-tagged proteins in *Neurospora*. These constructs were introduced at and expressed from the *his-3* locus in different strains. To examine protein-protein interactions, heterokaryons between these strains were constructed. We conclude that this strategy is a useful tool to investigate protein function and protein interactions.

After publication of a draft of the *Neurospora* genome project (Galagan *et al.*, 2003), gene knockout (Colot *et al.*, 2006) and DNA array projects (Kasuga *et al.*, 2005) started to elucidate gene function. Information and materials obtained from these projects have substantially changed experimental methods and accelerated molecular studies in *Neurospora*. Studies of protein-protein interaction and post-translational modifications such as ubiquitylation, SUMOylation, phosphorylation and methylation are indispensable. In many cases, one needs to use specific antibodies to examine protein interactions and modifications. The experiments that determine protein-protein interactions by using epitope tags therefore have been carried out in *Neurospora* (Dementhon *et al.*, 2006; Sarkar *et al.*, 2002).

In this report, we present useful targeting vectors to express FLAG- or HA-tagged proteins of interest integrated at the *his-3* locus in *N. crassa*. To create the vectors, we amplified a DNA fragment that contained sequences of linker and FLAG-epitope tag by PCR, then digested it with *SpeI* and *BamHI* and inserted the DNA fragment into the *SpeI/BamHI* site of the *his-3* targeting vector pMF272, (Freitag *et al.*, 2004), resulting in plasmid pFLAGN1. Similarly, a DNA fragment encoding HA-epitope tag was digested with *SpeI* and *XbaI* and ligated into the *SpeI/XbaI* site of pMF272, resulting in the creation of pHAN1. Confirmation that the sequence of interest was inserted into the vector was performed by DNA sequencing.

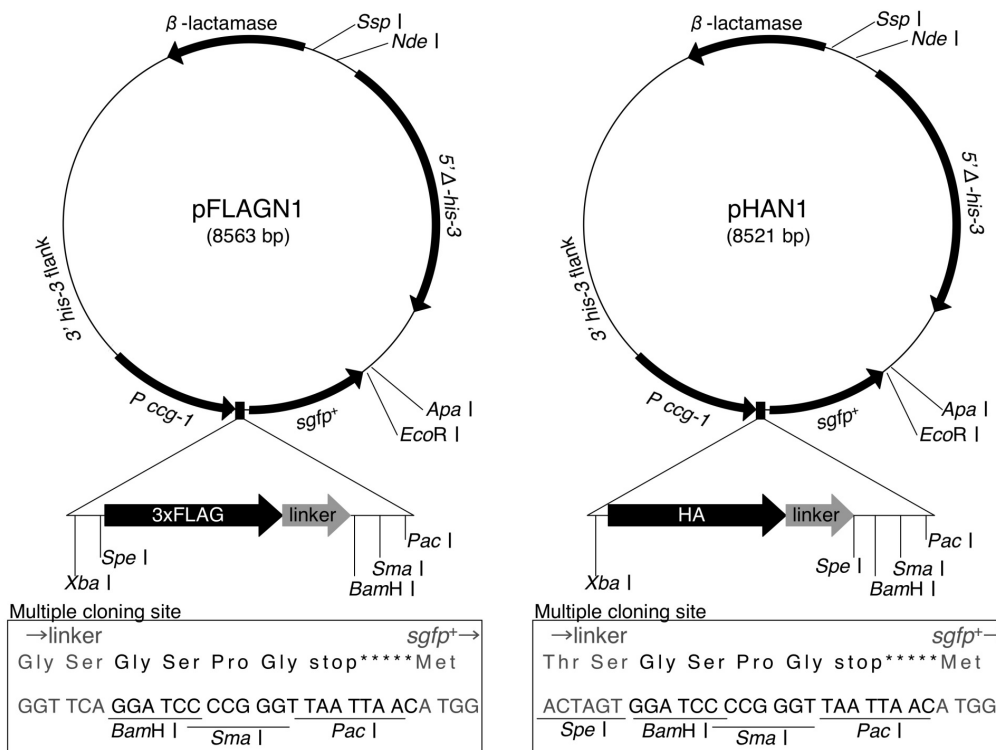


Figure 1. Construction of the vectors. DNA sequence containing linker and epitope tag was inserted between *cpg-1* promoter and *sgfp+*-ORF of pMF272. The plasmids were constructed to add the epitope tags to a polypeptide at its N-terminus, not its C-terminus.

We selected PCNA as a model for detecting a physical interaction between proteins. In *S. cerevisiae*, PCNA is an essential component for DNA replication (Johnson and O'Donnell, 2005) and also functions in DNA repair (Hoegge *et al.*, 2002) and gene silencing (Zhang *et al.*, 2000). PCNA exists in the cell as a homotrimeric ring, which functions as a sliding clamp and a docking station for many proteins. Formation of the trimer is essential for the functions (Jonsson *et al.*, 1995). Neurospora PCNA is also essential for its vegetative growth, as indicated from the result of gene-targeting experiment (data not shown). To determine Neurospora PCNA interacts with each other in vivo, cDNA of Neurospora PCNA was cloned, digested with *Bam*HI/*Eco*RI and ligated into *Bam*HI/*Eco*RI-digested pFLAGN1 or pHAN1 to generate pFGNPCNA or pHANPCNA, respectively. Each of them were linearized by a digestion with *Nde*I and transformed into the *his-3 inl* strain (FGSC#10376, KBT-H3-26A; *mat A rid^d::hph⁺ his-3::lpl^{d(5192-6046)}::hph⁺::tk⁺; inl*) or *his-3 pan-2* strain (FGSC#10375, KBT-H3-4A; *mat A rid^d::hph⁺ his-3::lpl^{d(5192-6046)}::hph⁺::tk⁺ al-2; pan-2; cot-1*) by electroporation as described previously (Margolin *et al.*, 1997). The *his-3 inl* strain was derived from a cross of the *his-3* mutant strain (FGSC#9097; *mat A his-3::lpl^{d(5192-6046)}::hph⁺::tk⁺; inl*) (Lee *et al.*, 2003) with a *rid*-defective strain (*mat a rid^d::hph⁺*) and the *his-3 pan-2* strain was derived from a cross of the *his-3 inl* strain with 74-OR31-14a (FGSC#4934; *mat a al-2; pan-2; cot-1*). Resulting histidine-prototroph transformants were isolated and grown for 7 days at 30°C.

To make the transformant homokaryotic, single colony isolation was repeated several times. The genomic DNA was isolated to determine whether the expression construct was integrated correctly at the *his-3* locus and the isolated clones were homokaryon by PCR. The clones were named KBT-FGPC-11A (FGSC#10377; *mat A rid^d::hph⁺ his-3::lpl^{d(5192-6046)}::hph⁺::tk⁺::flag-pcna⁺; inl*) and KBT-HAPC-7A (FGSC#10374; *mat A al-2 rid^d::hph⁺ his-3::lpl^{d(5192-6046)}::hph⁺::tk⁺::ha-pcna⁺; pan-2; cot-1*), respectively. The *his-3* (FGSC#9097) strain was generated to allow for rapid identification of homokaryotic transformants containing the expected gene replacement event (Lee *et al.*, 2003). Vectors used in this study suit for targeting event in both the *his-3* mutant strain (FGSC#9097) and a canonical *his-3* strain (such as FGSC#6103), but not adjusted to gain the advantage of the *his-3* mutant strain (FGSC#9097) due to a lack of suitable homology sequence, resulting in remaining of *hph⁺::tk⁺* sequence in its genome.

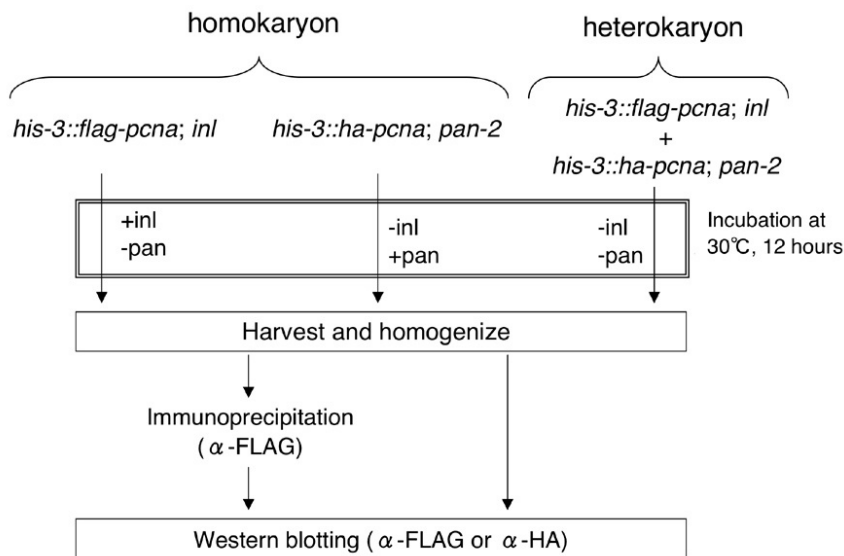


Figure 2. Procedure for investigating interaction between HA-tagged protein and FLAG-tagged protein expressed at the *his-3* locus. Western blotting was performed with cell extract from mycelia of each strain. Heterokaryotic strain was generated by a culture of mixed strains, the KBT-FGPC-11A and KBT-HAPC-7A, in the same test tube.

In order to analyze the interaction between FLAG-PCNA and HA-PCNA, two homokaryon strains, KBT-FGPC-11A and KBT-HAPC-7A, and the heterokaryon strain were analyzed (Fig. 2). Conidia of each strain were incubated in 50 ml of liquid medium (final 2.0×10^6 conidia/ml) containing nutrients as indicated in Figure 2 and cultured with shaking at 30°C for 12 hours. Cells were collected and homogenized with Micro Smash™ MS-100 (Tomy) in a buffer (50 mM Tris-HCl pH 7.5, 200 mM NaCl, 10% glycerol, 0.3% TritonX100) containing a protease inhibitor (Complete, EDTA-free; Roche). The extract was centrifuged at 14000 rpm for 10 min at 4°C. The supernatant was mixed with anti-FLAG M2 Affinity Gel (Sigma) for 1 hour at 4°C. The precipitates and whole cell extract were electrophoresed in 11% SDS-PAGE gels. Following transfer from the gels onto PVDF membranes, they were probed with mouse anti-FLAG M2 monoclonal antibody (Sigma) or anti-HA monoclonal antibody (HA.11; Covance).

FLAG-PCNA or HA-PCNA was detected in whole cell extracts from the KBT-FGPC-11A or KBT-HAPC-7A strains by anti-FLAG or -HA antibody, respectively, indicating that tagged PCNA was expressed in *N. crassa* (Fig. 3, left three lanes). FLAG-PCNA was

detected with anti-FLAG antibody in the immunoprecipitated samples from both the heterokaryon and the FLAG-PCNA-expressing strain but not detected in that of HA-PCNA-expressing strain. HA-PCNA was detected with anti-HA antibody in the immunoprecipitated sample of the heterokaryon strain but was not detected in the immunoprecipitated samples from the strains expressing HA-PCNA or FLAG-PCNA only (Fig.3, right three lanes). This indicates that FLAG-PCNA and HA-PCNA physically interact with each other in *Neurospora*, and the interaction is not affected by HA-tag or FLAG-tag at its N-terminus.

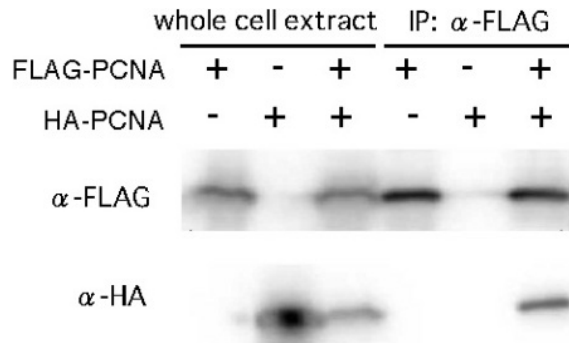


Figure 3. Examination of interaction between FLAG-PCNA and HA-PCNA by western blotting. The left three lanes show Western blotting performed against whole cell extract and the right three lanes show that of IP-samples produced by using anti-FLAG antibody. In the right three lanes, HA-PCNA was detected only in the sample from the mixed strain that expressed both FLAG-PCNA and HA-PCNA.

In conclusion, it was shown that FLAG- or HA-tagged proteins expressed from the *his-3* locus can be detected by western blotting and that physical interactions between proteins can be accessed by using a forced-heterokaryon strain. This is a useful method to analyze post-translational modification and physical interactions between proteins. The strains and vectors used in this study are available in FGSC.

Acknowledgements

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