Fungal Genetics Reports

Volume 38 Article 15

Co-regulation of two tandem genes by one blue-light element in Neurospora crassa

S. PIETSCHMANN

Max-Planck-Institut für molekulare Genetik

J. EBERLE

Max-Planck-Institut für molekulare Genetik

F.-R. LAUTER

Max-Planck-Institut für molekulare Genetik

See next page for additional authors

Follow this and additional works at: https://newprairiepress.org/fgr



This work is licensed under a Creative Commons Attribution-Share Alike 4.0 License.

Recommended Citation

PIETSCHMANN, S., J. EBERLE, F. LAUTER, N.N. PANDIT, and V.E. RUSSO (1991) "Co-regulation of two tandem genes by one blue-light element in Neurospora crassa," *Fungal Genetics Reports*: Vol. 38, Article 15. https://doi.org/10.4148/1941-4765.1461

This Regular Paper is brought to you for free and open access by New Prairie Press. It has been accepted for inclusion in Fungal Genetics Reports by an authorized administrator of New Prairie Press. For more information, please contact cads@k-state.edu.

Co-regulation of two tandem genes by one blue-light element in Neurospora crassa

Abstract

Many genes of *Neurospora crassa* are regulated by blue light: *al-1* (Schmidhauser et al. 1990 Mol. Cell. Biol. 10:5064-5070), *al-2* (Lauter, Schmidhauser, Yanofsky, Russo unpublished), *al-3* (Nelson et al. 1989 Mol. Cell. Biol. 9:1271-1276), *bli-3*, *bli-4*, *bli-7*, *bli-13* (Sommer et al. 1989 NAR 17:5713-5723). For none of these genes are the blue light cis-regulatory sequences (blue-light elements, BE) known. Here we report the presence of such BE in front of *bli-4*.

Authors

S. PIETSCHMANN, J. EBERLE, F.-R. LAUTER, N. N. PANDIT, and V. E. A. RUSSO

Co-regulation of two tandem genes by one blue-light element in *Neurospora* crassa

S. Pietschmann, J. Eberle, F.-R. Lauter, N.N. Pandit and V.E.A. Russo - Max-Planck-Institut für molekulare Genetik, Ihnsestraße 73, 1000 Berlin 33, F.R.G.

Many genes of *Neurospora crassa* are regulated by blue light: *al-1* (Schmidhauser et al. 1990 Mol. Cell. Biol. 10:5064-5070), *al-2* (Lauter, Schmidhauser, Yanofsky, Russo unpublished), *al-3* (Nelson et al. 1989 Mol. Cell. Biol. 9:1271-1276), *bli-3*, *bli-4*, *bli-7*, *bli-13* (Sommer et al. 1989 NAR 17:5713-5723). For none of these genes are the blue light cis-regulatory sequences (blue-light elements, BE) known. Here we report the presence of such BE in front of *bli-4*. The blue-light element of *bli-4* is in a stretch of DNA which is 300 bp long. We cloned this BE in front of a hygromycin reporter gene (Hygr) (Staben et al. Fungal Genetics Newsletter 36:79-81). Downstream of the reporter gene we cloned the selectable marker *tub-2* (also called *Bmlr*), which gives resistance to benomyl (Orbach et al. 1986 Molec. Cell. Biol. 6:2452-2461). The distance between the BE and the translation start (ATG) of *Bmlr* is about 3,000 bp (Fig. 1A).

After transformation in *N. crassa* wt (St. Lawrence a) we selected several benomyl resistant transformants (H1 to H4). These transformants were grown in Vogel's liquid medium with the addition of 0.5 ug/ml benomyl and total RNA of dark grown and illuminated mycelia were extracted (T. Sommer et al. 1989 NAR 17:5713-5723). The Northern blots show that both the hygromycin gene (Fig. 2b) and the *Bmlr* gene were strongly light regulated.

Experiments with a similar construct, where a tagged *bli-4* gene served as expression vector (Fig. 1b) gave different results. In the three B1-B3 transformants the tagged *bli-4* gene was strongly light regulated (Fig. 2e) while the *Bmlr* gene was only weakly regulated (Fig. 2f). The simplest interpretation of this result is that the blue light element of *bli-4* acts as an enhancer that functions well at a distance of 3,000 bp but not as well at a distance of 4,500 bp from the translation start of the *Bmlr* gene. A more complete understanding of blue-light regulation of gene expression in *N. crassa* will come from cloning and comparison of the properties of the blue-light element of several genes (see also Pandit and Russo 1991 Fungal Genetics Newsletter this issue).

Acknowledgements: We thank Chuck Staben and Charles Yanofsky (Stanford University) for providing the hygromycin reporter gene. We thank Uta Marchfelder for typing. This work was partially supported by the Deutsche Forschungsgemeinschaft.

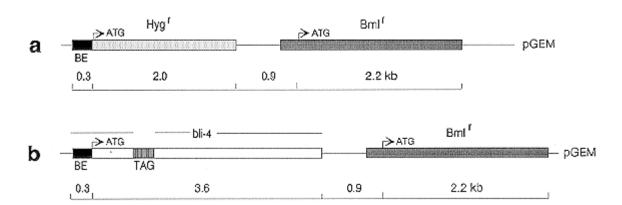


Figure 1.

A. Construct: blue-light box of *bli-4 Hygr* gene - *Bmlr* gene in a pGEM vector B. Construct: blue-light box of *bli-4*-tagged *bli-4* - *Bmlr* gene, in a pGEM vector

BE: Blue-light element of *bli-4* gene *Hygr*: hygromycin resistant gene (*hph*) *Bmlr*: benomyl resistant gene (*tub-2*)

ATG: translation start

TAG: stretch of DNA not present in N. crassa

pGEM: plasmid of Promega

kb: kilobase

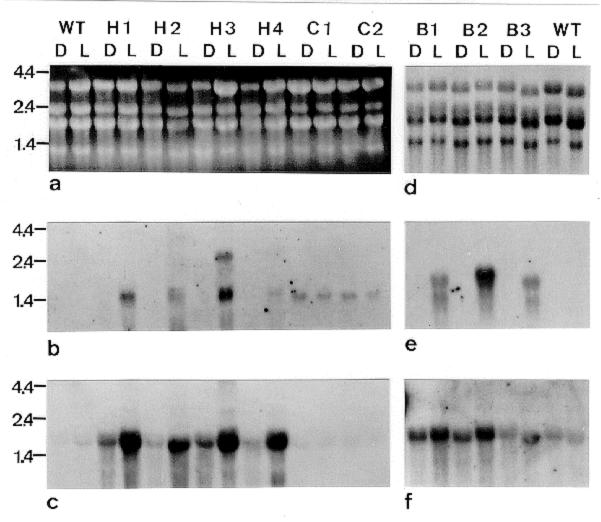


Figure 2. Northern Blot Analysis

- a), d) Photos of the membranes used in b), c) and e),f) respectively just after blotting and before hybridization.
- b) Hybridization of the membrane shown in a) with radiolabelled *hygr* gene.
- c) Reprobing of the membrane a), after stripping, with radiolabelled *Bmlr* gene.
- e) Hybridization of the membrane shown in d) with radiolabelled TAG DNA.
- f) Reprobing of the membrane d), after stripping, with radiolabelled *Bmlr* gene.

wt: wild type (St. Lawrence, mt a)

- H1-H4: four different transformants of wt with the construct a) of Fig. 1.
- B1-B3: three different transformants of wt with the construct b) of Fig. 1.
- C1-C2: transformants of wt with the hygromycin gene which has the Aspergillus *trpC* promoter(non-light regulated)