

# FungiDB User Comments Campaign

## 2017 Fungal Genetics Conference, Asilomar



### Did you know that User Comments:

- help to improve genome annotation, by providing evidence for alternative gene models, phenotypes, subcellular localization, etc
- become immediately visible on gene pages, and can be edited by you at a later time
- benefit the research community at large, and help to showcase your favorite genes

Add just one User Comment and pick up a free 'Got Fungus?' coffee mug ... and top 10 submitters get a T-shirt!

Participation is simple: Attach a User Comment to any gene page ... and stop by the FungiDB booth during poster sessions to pick up your free 'Got Fungus?' coffee mug. Ask colleagues who are not attending the Asilomar meeting to enter a User Comment, and you can take home a mug for them. Get everyone in your lab to enter a comment and take a whole bunch of mugs home with you! Top submitters will also receive a 'Got Fungus?' T-shirt. Not available in stores!

### Adding User Comments is easy!

1. Register for an account & sign in to FungiDB
2. Navigate to the gene page of interest
3. Click on the 'Add a User Comment' link
4. Fill out the comment page (see right). Note: one comment can be attached to many genes
5. Click submit – *Your comments become available and searchable immediately!*

### Helpful User Comments might include:

- Gene names ... *and don't forget synonyms!*
- PubMed IDs, linking genes to key references (feel free to link any relevant papers, not just your own)
- Corrections / confirmation of gene models, annotation, etc (please describe supporting evidence)
- Image data ... note that you can attach images to genes, providing subcellular localization, etc
- Other functional or experimental information ... *even negative data can be helpful!*

### Want to submit User Comments in bulk? No problem!

No problem! Download: <https://tinyurl.com/bulk-user-comments-form>



Questions? [Help@FungiDB.org](mailto:Help@FungiDB.org)

