FlyBase Update Form

FBrf0086747

If you have new data for **FlyBase**, or know of errors in **FlyBase**, please note them below and return this form to the **FlyBase** demonstration ***(Helix-Delmar Room)*** during this conference. In **FlyBase**, we will reference these as "Z. Bloggs, personal communication to **FlyBase** (1996)". Please note our policy that we will not correct or add data unless there is a written update submitted to us (on paper, email record, or WWW submission form).

Alternatively, you may return this form by mail to:

FlyBase
Harvard Biological Laboratories
16 Divinity Avenue
Cambridge, MA 02138 USA

or send us update information by email to: [redacted]

or via our WWW s[redacted]

Thank you for your help.

Submitter Information: Date: 28/4/96

Name: P. Adler.

Address: Biology Department, University of Virginia, Charlottesville, VA.

Telephone number:

FAX number:

Email address:

Tentative Name

Write your new data or corrections below (continue on reverse):

poster 80A. The gene fumy is now called yeti. It is localised to 3L

new gene ultrahair (uhr). - located 2L.

cat of nine mits is on 3R

(cont).

Mutations in uhr cause multiple hair cells whose individual hairs are as long or longer than normal hairs. uhr clones are cell autonomous.

If the correction is to published work, please note the citation, *(1st author, year, journal, volume #, pages).*

Paul Adler