



## Nome News Issue 17

**Issue 17. March 2003**

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### Meetings attended

Elspeth Bruford attended HAWK3, the third Human Annotation Workshop, at Marco Island, Florida, in February along with about 20 representatives from all the major US sequencing centres and the Sanger Centre. The meeting proved very useful for all concerned and the presentation was well received. As a result, information regarding our minimum requirements for receiving data has been sent out to the HAWK mailing list and will also be placed online at the HAWK website. We have now established contacts with all the major sequencing centres.

### Upcoming meetings

Once again at the [HGM meeting \(Cancún, Mexico\) 27-30 April 2003](#) we will be awarding poster prizes for the two posters that contain the HIGHEST NUMBER of correct gene symbols, see URL:

<http://hgm2003.hgu.mrc.ac.uk/Competition/>

Please come and see us at booth 60 to discuss nomenclature and don't forget to attend our presentation, which will be given by Hester Wain.

May 2003 sees some of the HGNC team at the [European Society of Human Genetics \(ESHG\) annual meeting \(3rd-6th May\)](#) in Birmingham, where Mathew Wright & Michael Lush will be presenting a poster.

### Virtual Workshop

The [Virtual Gene Nomenclature Workshop](#) has seen the identification of sequence for over 10% of the genes listed. Our thanks go to Youla Karavidopoulos, Tom Bonner, Marie-Paule Lefranc and Ian Dunham for their help. The deadline for workshop completion is 30th June 2003, so if you have any sequences lying around that might just belong to one of the genes in our workshop, please send them in!

### Bioinformatics

With the recent arrival of the machines for our new Beowulf Cluster, our capacity to process sequence has moved up a gear or eight! The 16 Athlon CPUs, 32 Gb of RAM and 480 Gb of disk space, ensures our readiness for the remaining genome sequences when they arrive.

### Journals

We are very pleased that the [Lancet](#) has joined the growing number of journals which support the use of approved nomenclature. If you would also like to promote approved nomenclature in your journal, please contact us at [hgnc@genenames.org](mailto:hgnc@genenames.org) for details of what to include in your instructions to authors.

If you would like to be added to our Nome News mailing list or if you have questions or comments on any human gene nomenclature issue, please email us at: [hgnc@genenames.org](mailto:hgnc@genenames.org)