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ASW

Diseño de bases de datos

- HVRbase (HANDT et al. 1998)

<http://www.hvrbase.de/>

- FHG Base (Ferrara Human Genetics Database). Simoni *et al.* 2000.

<http://web.unife.it/progetti/genetica/pdata.htm>

- Concordance (MILLER *et al.* 1996)

<http://www.bioanth.cam.ac.uk/mtDNA/>

F

Filogenético

- TREEVIEW (PAGE 1996)

<http://taxonomy.zoology.gla.ac.uk/rod/treeview.html>

- DnaSP 3.5 (ROZAS Y ROZAS 1999)

<http://www.ub.es/dnasp/>

- PHYLIP 3.5c (FELSENSTEIN 1993)

<http://evolution.genetics.washington.edu/phylip.html>

- NETWORK

<http://www.fluxus-engineering.com/>

- ARLEQUIN (SCHNEIDER *et al.* 2000)

<http://evolution.genetics.washington.edu/phylip/software.html>

Otro

- TEXTPAD 4.7.1. (HELIOS SOFTWARE SOLUTIONS 1992-2003)

<http://www.textpad.com/>

- GraphPad Prism 4.00

<http://www.graphpad.com>

Análisis de secuencias de pláfticación

- DNASTAR®

<http://www.dnastar.com>

- Primers for the World Wide Web

<http://williamstone.com/primers/javascript/>

