



Xenbase v3.0 has been launched with great new features:

- New user-friendly [landing page](#)- a whole new look and feel!
- New [Xenopus laevis Genome v7.1 pre-release](#) from the Xenopus Genome Consortium, and just released by the JGI is now available on [GBrowse](#), [BLAST+](#) and for [FTP download](#).
 - X. laevis genome v7.1 has:
 - 2.7 Gb total scaffold length
 - 50% of the assembly in contigs >19.3kb
- [Antibody Database](#) - search through 500+ curated antibody reagents known to work in Xenopus.
- [Faster BLAST+ Server](#)
- [Improved Genome Browser](#) - the latest version of GBrowse features:
 - DNA sequence displayed for zooms >100bp
 - faster, cleaner interface with many optional tracks
 - download snapshots as png's or send to others as a link
 - download track data in gff3, FASTA, or GenBank format for selected regions, entire scaffold, or entire track across the genome.
- Expanded [Community Meetings and Resources](#).
- New [miRNA expression catalog](#) from the Wheeler lab.
- New [Morpholino Wiki](#) listing 600+ published MOs.
- Spotlight on new high-impact Xenopus research articles.

Continue to use Xenbase to find:

- 15,000+ individual gene pages with searchable gene [expression patterns](#) and associated literature.
- Xenopus [developmental timing](#) and [anatomy descriptions](#)
- [NF stage images](#).
- Comprehensive [library](#) of over 47,000 Xenopus research articles
- Updated Xenopus laboratory [protocols](#) and reagents
- Links to Xenopus stock centers: [EXRC](#) (UK), [NXR](#) (USA), [CRB](#) (France), and [NBRP](#) (Japan)
- [Contact information](#) for Xenopus researchers and labs and [job postings](#).

[Provide comments and feedback](#) to help improve our service to advance your research!