

Medaka on the move

Long before it found favor with geneticists—long before there were geneticists, in fact—the Japanese medaka (*Oryzias latipes*) was bred by hobbyists throughout Japan. As shown here in an 1843 drawing by Bai-en Mohri, there are differently pigmented strains of medaka that enhance their aesthetic appeal, including those now recognized as wildtype (brown) and mutant (orange-red and white) varieties. The orange-red medaka harbors a mutation at the *b* locus, and on page 381, Shoji Fukamachi and colleagues¹ describe a series of mutations in a new gene at this locus, encoding a putative sucrose transporter. The identification of the molecular lesion(s) in the *b* mutant is not only of historical interest, but also marks the first positional cloning of a gene in this model organism.

It has already been a good year for medaka mavens. Results reported in the last 12 months include: high-efficiency generation of embryonic mutants in ethylnitrosourea-based screens², a detailed linkage map³, evidence for *de novo* transposition of the medaka Tol2 transposable element⁴, cloning by nuclear transfer⁵, and construction of a bacterial artificial chromosome (BAC) library derived from the inbred Hd-rR strain⁶. The medaka has also had its share of historical firsts, most notably as the first vertebrate in which Mendelian inheritance was demonstrated⁷, and, in a less weighty turn, as the first vertebrate to reproduce successfully in microgravity⁸. The merits of

medaka as an experimental organism have been summarized⁹, and the recent flurry of reports, including that of Fukamachi and colleagues¹, suggests that a critical mass of essential genetic resources has at last been assembled (see <http://biol1.bio.nagoya-u.ac.jp> for a compendium of medaka resources).



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The *b* mutant analyzed by Fukamachi *et al.* lacks melanin pigment in its melanophores, but its orange, white and silver chromatophores are normally pigmented, resulting in the characteristic orange-red color. The *b* locus was previously narrowed down to a manageably small candidate region, and was cloned by recombination mapping between two medaka strains that are highly polymorphic, an effort that was facilitated by the recently published BAC library⁶. Seven of eight *b* alleles have mutations in the coding region; the lone exception is the original *b* mutant, which may have a mutation in a regulatory element.

What is B? Results from database queries predict a 12-pass integral membrane protein. It seems to be expressed in melanophores, but not exclusively, suggesting a possible function outside of melanogenesis. Indeed, as mouse pigmentation mutants have been a boon to vertebrate biology¹⁰, providing insight into several cell lineages, there is ample precedent for such proteins to have a number of roles. The biggest hint in regard to B comes from its 23% amino-acid sequence identity with plant sucrose transporters. Fukamachi *et al.*¹ have put forward a plausible scenario whereby sucrose transport could affect melanin synthesis, but a real understanding of the origin of the orange-colored medaka awaits functional studies on the B protein. At the very least, however, it's a pleasing and appropriate coincidence that B might have a role in sugar metabolism. If the medaka put Mendel on the map with respect to vertebrate genetics, recall one of the original sources of Mendel's laws: the pea plant *r* locus, which encodes a starch-branching enzyme that determines whether pea seeds are round or wrinkled¹¹. □

—Alan Packer

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