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Health & Welfare

Microsatellites: Genetic markers of choice

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By Leland L. Lai

Grow better quality animals in less time, with greater survival, and at less cost

Is it possible for aquaculture to be sustainable? The answer is a qualified “yes,” assuming the industry can implement radical changes in management thinking, conserve resources in culturing, and improve stock through the accelerated use of biotechnology. Chemicals, physical methods or nutritional components can not double crop production from the highest levels of output attained in this century.

In the 21st century, higher output will be a consequence of production efficiency per unit area rather than increases in land under cultivation. To envision what aquaculture will be like in the future, you need only look at past productivity gains in other agriculture sectors. For example, in the last 30 years, chicken egg production increased by 300 percent, cow milk production by 250 percent, daily weight gains in pigs grew by 200 percent, and in the last decade, salmon production gains exceeded 50 percent (both in growth and cost savings).

Farmers have employed selective breeding to improve their stock for centuries. Today, phenotypic (observational) programs are used for increasing the probabilities of trait occurrence and can often lead to improvements exceeding 10 percent per generation. However, these programs are time consuming, costly if many families are maintained, and can lead to mixed or negative results due to inbreeding. In the past ten years, genetic tools have often complemented plant and animal breeding programs. These tools are information-based and rely on very specific DNA sequences found throughout the specific species genome. ASiCo L.L.C., a company started in 1998, is the first to begin offering these same molecular tools for commercial aquaculture applications.

Much of today's genetic research is devoted to locating genes and chromosomes, and to developing genetic maps. This process may take years to accomplish since decoding the DNA sequence of shrimp, fish or bivalves may require tens of thousands of manhours. A genetic map shows the locations of genes or DNA fragments that have been identified, some of which express themselves as desirable traits for the farmer. It is possible, however, to find microsatellites throughout the aquatic species genome without knowledge of the genome or genetic mechanisms that make genes operative. When these markers are identified and characterized, each may be used as a signpost or beacon to help locate genes that express the desired trait. Markers are related to traits as a function of the distance they are from the gene responsible for that trait.

The ability to fingerprint and tag

These allow us to track genetic diversity among families and populations, establish pedigree lines centered around desired expression of certain traits, and determine vertical and horizontal lineage should true biosecure measures need to be implemented at the breeding center. As a result, one can build a unique profile on developed strains from non-lethal tissue sampling.

Marker-assisted selection

The primary value of markers is to assist in the breeding process and to improve the breeding prediction. Allelic profiling and association with traits of economic importance compounds the results of any breeding program and reduces the time required to achieve those results. Are there genetic differences between fast- and slow-growing fish? Or resistant stocks from those which don't survive? Markers have already proven their guidance value in the plant and animal sectors.

The use of microsatellites is one of many different marker approaches, but has proven to be the tool of choice over the last ten years by frequency of application across all commercial agriculture sectors. ASiCo has already found breeding and identification applications with these tools for more than twelve aquatic species. Genetic improvement through selective breeding allows farmers to grow better quality animals in less time, with greater survival, and at less cost than animals removed from the wild.

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Author



LELAND L. LAI

Aquatic Stock Improvement Company
P.O. Box 5
Hawthorne, California 90250 USA

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