

comparative work on northern and southern *Haliotis discus hannai* populations shows that long-term exposure and selection in warmer waters produces domesticated stocks with enhanced thermal tolerance and superior regulation of carbohydrate and amino-acid metabolism, demonstrating that breeding and translocation programs can deliberately cultivate heat-tolerant lines for deployment in warming regions. Genomic tools are emerging as powerful levers to accelerate thermal adaptation. Quantitative genetic and genomic selection work on *Pacific abalone* reports moderate heritability (0.35-0.42) for heat-tolerance traits and demonstrates that genomic selection models can predict heat tolerance with high accuracy, identifying specific SNP markers and candidate genes that can be used in marker-assisted breeding programs to improve resilience to marine heatwaves. Ecosystem-based designs like abalone-seaweed co-culture further enhance climate resilience by biologically modifying water conditions: in juvenile red abalone, IMTA with the red seaweed dulse elevated pH under simulated ocean acidification, producing faster growth, better condition and stronger shells, and thereby “shepherding” juveniles through their most vulnerable life stage despite deteriorating external seawater quality.

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