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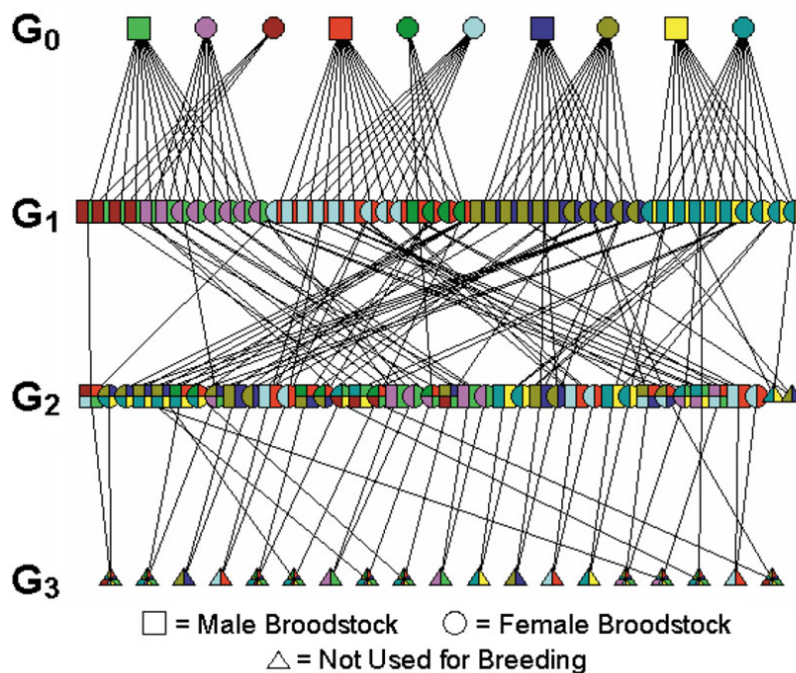
Health & Welfare

Research reveals inbreeding depression in Pacific white shrimp

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By Dustin R. Moss, M.S. , Steve M. Arce and Shaun M. Moss, Ph.D.

Hatch rates for the inbred families were much lower than for outbred families



Pedigree data for the experiment comprised four generations. G₀ represents the 10 founders of the population. Different colors represent different families. For G₂ and G₃, inbred families are represented by two colors and outbred families by three or more colors.

As the global shrimp-farming industry matures, the use of domesticated, genetically improved shrimp stocks will become more common. In closed populations under selection, inbreeding is an inevitable effect. Inbreeding depression, the loss of vigor among offspring when related individuals mate, can occur rapidly when intense selection is coupled with a narrow genetic base, as in a small population size. Yet little is known about the potential deleterious effects of inbreeding on penaeid shrimp.

Directed management strategies can slow the accumulation of inbreeding in closed populations. However, not knowing the severity to which inbreeding affects performance – or even what traits are affected – makes it difficult to determine the importance of avoiding inbreeding when designing a shrimp-breeding program.

Experimental families

A two-year study at the Oceanic Institute in Waimanalo, Hawaii, USA, investigated the effects of inbreeding on the hatchery and pond performance of Pacific white shrimp (*Litopenaeus vannamei*). The pedigree data comprised a total of four generations, G₀-G₃. G₀ founders were collected from the wild and assumed to be unrelated, with an inbreeding coefficient of 0. Two successive generations (G₂ and G₃) of inbred and outbred families were produced and compared.

Inbred families were produced by mating siblings. Outbred families were produced by mating two unrelated individuals. In G₂, 11 inbred and 12 outbred families were produced from six G₁ families. Nine inbred and 10 outbred families were produced in G₃. The resulting inbreeding coefficient values for outbred and inbred families were 0.00 and 0.25, respectively, for G₂ and 0.00 and 0.375, respectively, for G₃.

Measuring inbreeding depression

The effects of inbreeding were determined by quantifying four performance characteristics: hatch rate, hatchery survival to PL₈, pond growth rate, and pond survival. Nutrition, size, age, lighting, and water quality were similar for all broodstock used in the experiment.

Environmental variability also was controlled, as inbred and outbred families were produced together over a 10-day period, reared in the same hatchery and nursery systems, and grown together in the same pond. Thus, differences in performance characteristics between the outbred and inbred families were attributed to inbreeding.

To determine the magnitude of inbreeding depression, a percent phenotypic expression (percent PE) of each trait was calculated for each family using the formula: percent PE = (family mean/generation specific outbred family group mean) x 100. These values were then regressed against the inbreeding coefficient using linear regression.

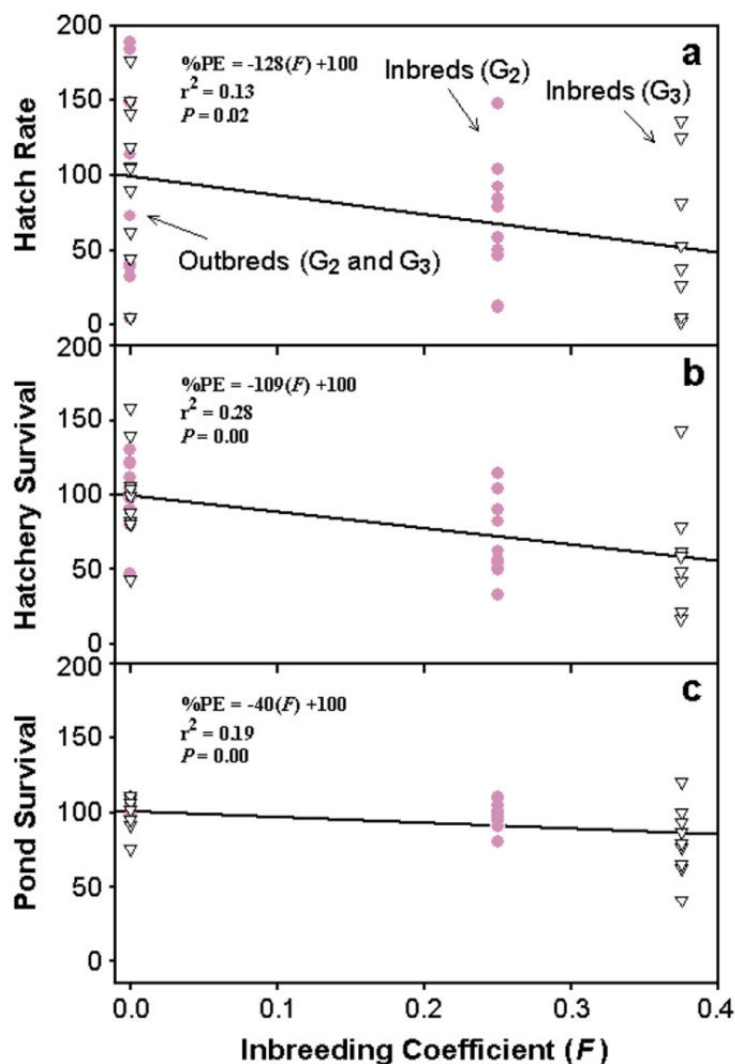


Fig. 1: Relations between inbreeding coefficient and phenotypic expression of performance characters relative to generation-specific outbred family group means.

Evidence for depression

Hatch rates for the inbred families were 33.3 percent lower than for outbred families in G_2 and 47.1 percent lower in G_3 (Table 1). Also, there was a significant negative correlation between percent PE-hatch rate and level of inbreeding (Fig. 1a). For every 10 percent increase in inbreeding, hatch rate decreased by 12.8 percent.

Hatchery survival for inbred families was 31.5 percent lower than for outbred families in G_2 and 38.7 percent lower in G_3 . As with hatch rate, a significant negative correlation was found between percent PE-hatchery survival and level of inbreeding (Fig. 1b). For every 10 percent increase in inbreeding, hatchery survival decreased by 10.9 percent.

Pond survival was greater than 85 percent for both inbred and outbred families in G_2 . In G_3 , pond survival for the outbred families was 19.5 percent higher than for inbred families. There was a significant negative correlation between percent PE-pond survival and level of inbreeding (Fig. 1c). Although not as dramatic as for hatch rate and hatchery survival, pond survival decreased by 4.0 percent for every 10 percent increase in inbreeding. The growth of inbred and outbred families was similar in G_2 and G_3 .



Shrimp families were held in nursery tanks before the shrimp were tagged and transferred to the grow-out pond for performance evaluation. Each small tank contained one family.

NMoss, Hatchery and growout performance, Table 1

G	Trait	Outbred	Inbred	Difference
2	Hatch rate	49.81 ± 34.39	33.25 ± 20.06	-33.3%
3	(%)	56.50 ± 29.28	29.88 ± 29.06	-47.1%
2	Hatchery	58.21 ± 4.00	39.87 ± 4.17	-31.5%
3	Survival (%)	58.01 ± 18.45	35.54 ± 21.95	-38.7%
2	Pond survival	87.17 ± 4.86	85.45 ± 7.35	-2.0%
3	(%)	67.35 ± 7.43	54.22 ± 15.57	-19.5%
2	Pond growth	1.06 ± 0.07	1.06 ± 0.08	0.0%
3	(g/week)	0.73 ± 0.06	0.74 ± 0.10	1.4%

Table 1. Hatchery and growout performance of outbred ($F = 0$) and inbred (G2, $F = 0.25$; G3, $F = 0.375$) families. Values are means \pm SD. G = generation. F = inbreeding coefficient.

Conclusion

A study of Pacific white shrimp found hatch rate and hatchery survival were negatively affected by inbreeding, with expected phenotypic depressions greater than 10 percent for every 10 percent increase in inbreeding. Pond survival also was negatively affected, but to a lesser extent. Inbreeding appeared to have no effect on shrimp growth at the levels tested.

This study provides an estimate of the degree to which inbreeding can affect shrimp performance. It can be used to balance the detrimental effects of inbreeding against the added expense of avoiding inbreeding through methods such as tagging and pedigree tracking.

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Authors



DUSTIN R. MOSS, M.S.

The Oceanic Institute
41-202 Kalaniana'ole Hwy.
Waimanalo, Hawaii 96795 USA

dross@oceanicinstitute.org (<mailto:dross@oceanicinstitute.org>)



STEVE M. ARCE

The Oceanic Institute
41-202 Kalaniana'ole Hwy.
Waimanalo, Hawaii 96795 USA



SHAUN M. MOSS, PH.D.

The Oceanic Institute
41-202 Kalanianaʻole Hwy.
Waimanalo, Hawaii 96795 USA

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