

Scottish Shellfish Innovation, SAIC project updates

Association of Scottish Shellfish Growers
Annual Conference, 26-27 October 2017

Robin Shields



Scientific support to pilot mussel hatchery

SAICHatch Project

- “To research and develop new technologies and processes required for a successful Scottish commercial mussel hatchery”
- May 2016 – Oct 2018 (30 months)
- Total project value £366,000

Genetic Tools Project

- “To develop and validate standardized genetic tools for stock management of mussels”
- Jan 2016 – Dec 2017 (24 months)
- Total project value £150,000

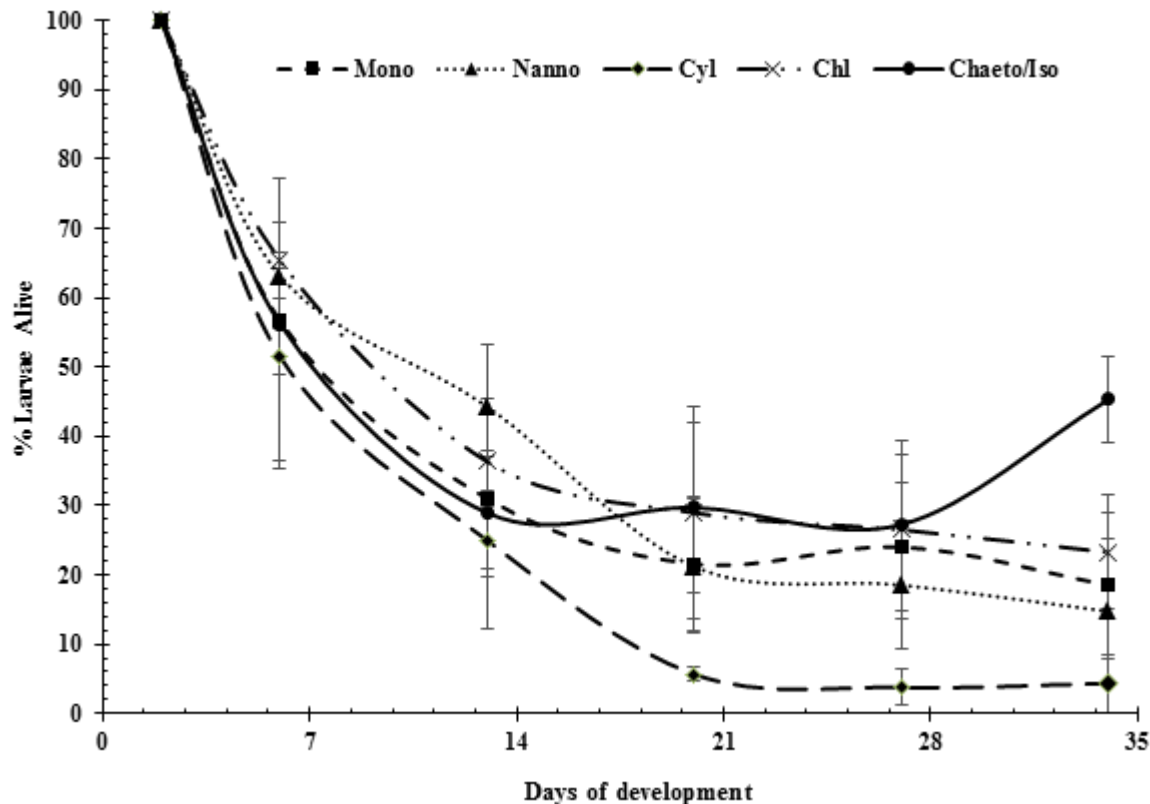


SAICHatch work streams

- **Innovations in algal feeds**
 - Strain selection, comparing phototrophic, heterotrophic & mixotrophic culture conditions
 - Lab scale evaluation of shortlisted strains: mussel larval survival, growth, microbiology
- **Microbial communities in pilot hatchery**
 - Characterisation for key hatchery components and mussel life history stages; “seasonality” effects
- **Metamorphosis & role of biofilms in spat settlement**
 - Molecular characterisation of maturing biofilms
- **Methodologies for transferring spat to sea**
 - Optimal size/age of spat
 - Socked vs unsocked growing ropes
 - Post-transfer performance

SAICHatch, Algal feeds

- Eg, larval survival in response to different algal diets (source A. Hughes et al., SAMS)



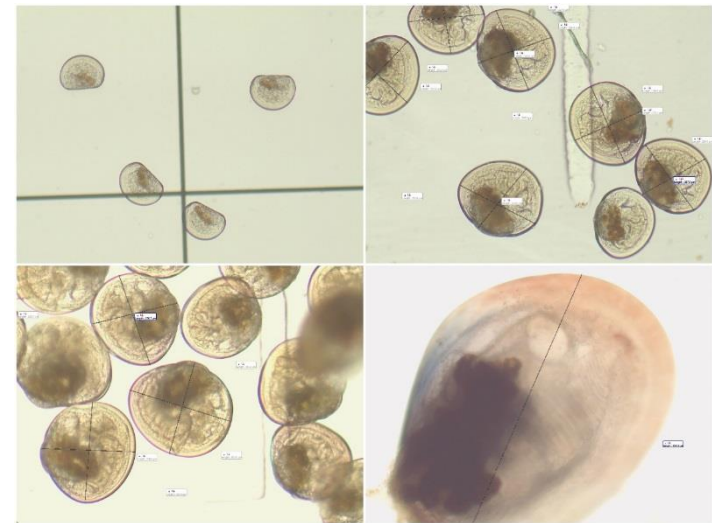
Survival percentage during the larval development for the different diets evaluated. Mono: *M. subterranean*; Nanno: *N. oceanica*; Cyl: *C. fusiformis*; Chl: *Chlorella vulgaris*; Chaeto/Iso: mixed diet 30:70 *C. calcitrans*: *I. galbana*.

SAICHatch, Hatchery microbial communities

- Protocols optimised for DNA extraction from different sample types
- Sampling kits distributed to NAFC and SAMS
- Samples collected from key locations within pilot hatchery, multiple larval batches and different times of year
- DNA samples currently being processed from replicated lab. scale trials (SAMS) and pilot hatchery (NAFC)
- In prep. - Characterise culturable bacterial populations using standard microbiological techniques, with emphasis on *Vibrio* spp.

SAICHatch, Metamorphosis & spat settlement

- Different experimental rearing systems evaluated - 150L downwellers preferred
- Spat exposed to different rope types (ongoing)
- Rope & tape samples “matured” at NAFC and Marine Lab Aberdeen
- Methods developed for biofilm harvesting and DNA extraction
- qPCR tested with taxa-specific primers

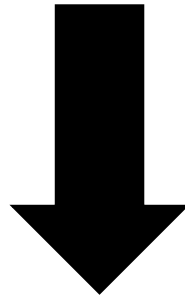


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Genetic Tools work programme

- Develop a suitable SNP panel (*Single Nucleotide Polymorphism* panel)
- Use spat from single parent crosses to validate SNP panel (parentage assignment)



Tools for mussel family selection programme

Genetic Tools, 2017 accomplishments

- SNP validation
 - Feasibility tested of microsatellite analysis for different-sized mussels → *2mm minimum*
 - Validation of SNP panel requires simultaneous analysis of 6x batches of single parent crosses
- Spat production
 - Painstaking process: numerous batches reared throughout 2017 season, continuous refinement of protocols.
 - Sample collection now almost completed and ready for despatch to Xelect.



Thank you for your attention

- **With thanks to the participants -**

- **SSMG:** Michael Tait, Lindsay Angus, Stephen Cameron
- **Xelect Ltd:** Tom Ashton, Andrew Richardson
- **NAFC:** Lesley McEvoy, Blažka Smiljanič, Gregg Arthur, Danny Cowing, Raquel Quinta, Tariq Mohammed
- **SAMS:** Adam Hughes, Michelle Stanley, Alessandro Laudicella, Joe Penhaul-Smith
- **Institute of Aquaculture:** Stefano Carboni, Andrew Desbois, Margaret Crumlish, Stuart McMillan
- **Marine Scotland Science:** Una McCarthy, Iveta Matejusova

