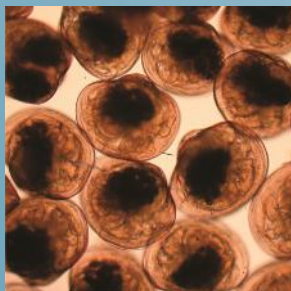
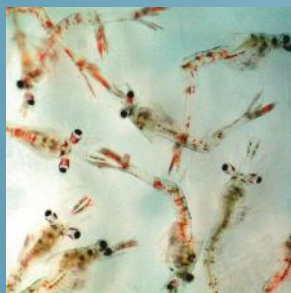
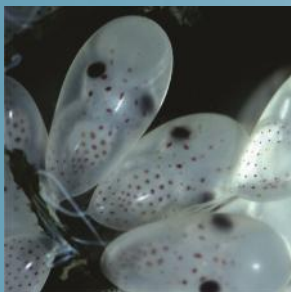
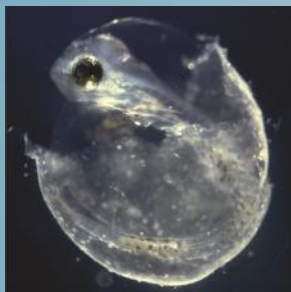


# larvi 2013

6th fish & shellfish larviculture symposium



Digital image analysis  
to aid broodstock management  
and egg and larval quality assessment



Andrew Davie

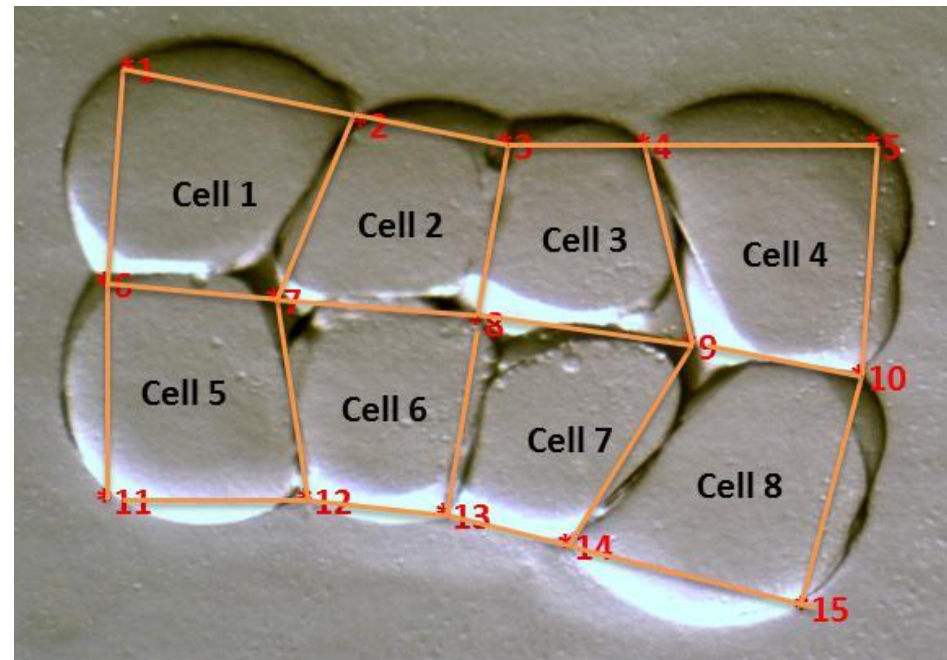
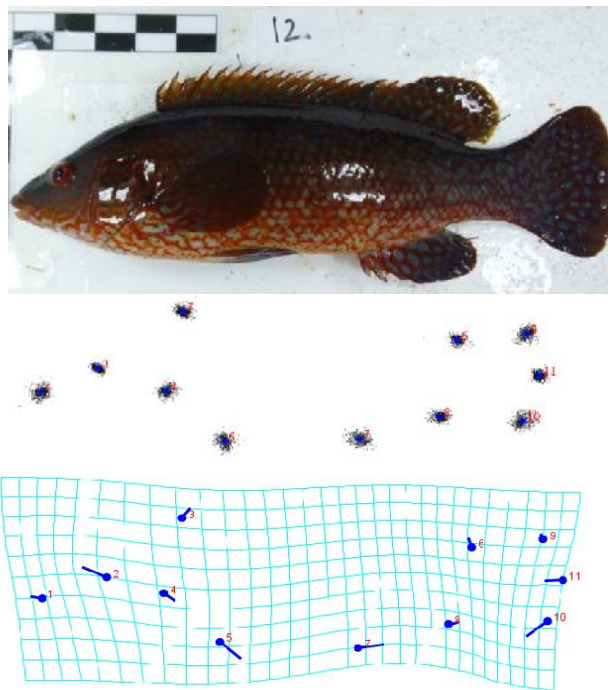
ghent university, belgium, 2-5 september 2013



# DIGITAL IMAGE ANALYSIS TO AID BROODSTOCK MANAGEMENT AND EGG QUALITY ASSESSMENT.

Andrew Davie, Eric Leclercq, & Herve Migaud

Institute of Aquaculture, University of Stirling, Scotland





# Digital Image Analysis

Digital image analysis is deeply ingrained in modern society



Security

Health and Safety

Social Media

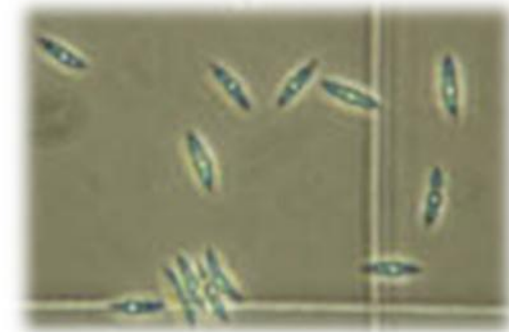
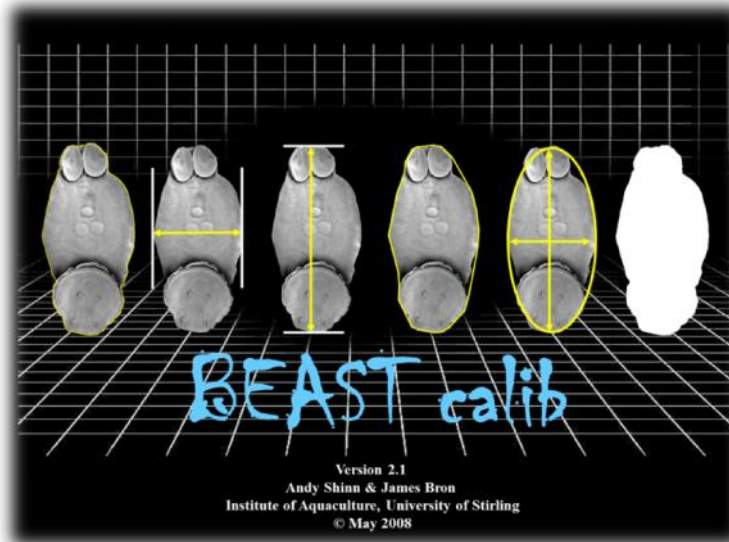


However...Digital image analysis appears to be utilised sporadically across the aquaculture industry and research community.



Biomass estimation

Parasite identification



Live feed quantification



## Ballan wrasse (*Labrus Bergylta*)

Biological control of sea lice in salmon farming

3 hatcheries in the UK

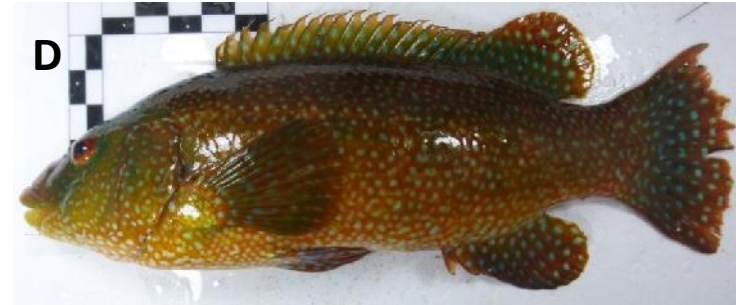
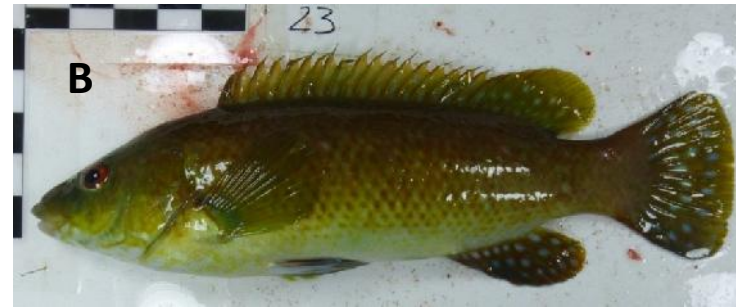
Challenging species to farm

- Long lived and late maturing (> 6years)
- Protogynous hermaphrodites
- Benthic substrate spawners
- Live in social harems
- Larval/juvenile performance problematic

**“Production and implementation of farmed wrasse in the Scottish Salmon industry”**

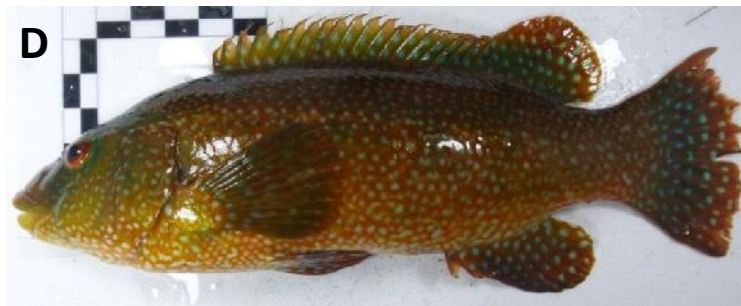
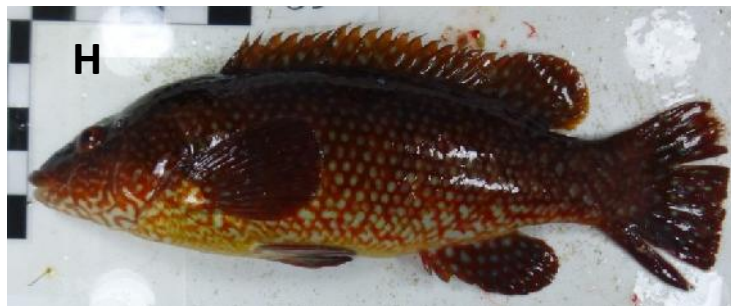


# Who is Who?



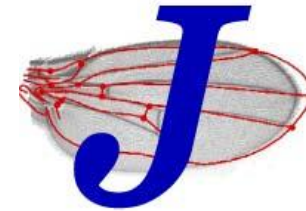


# Highly diverse external coloration





# What is geometric morphometrics



## Principal

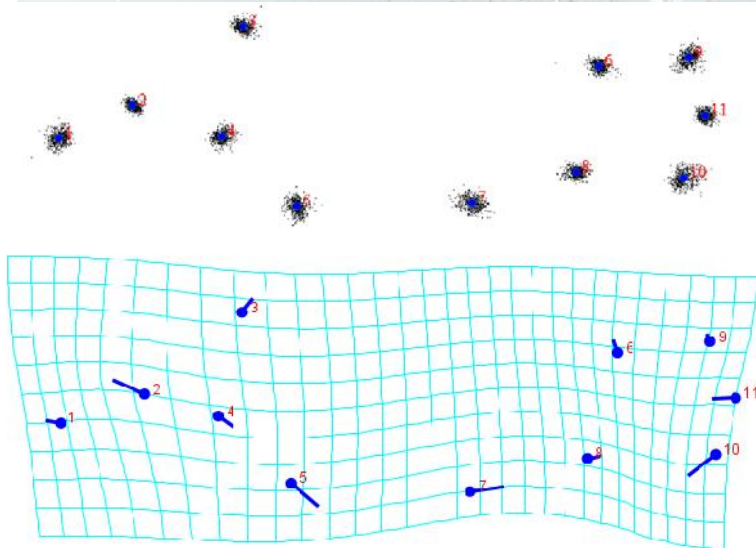
Analysis of landmark coordinates

Used to separate “size” and “shape”  
predict sex based on “shape”

## “Size” (surface)

Measured from raw coordinates

(Dispersion of landmarks from the centroid)



## “Shape”

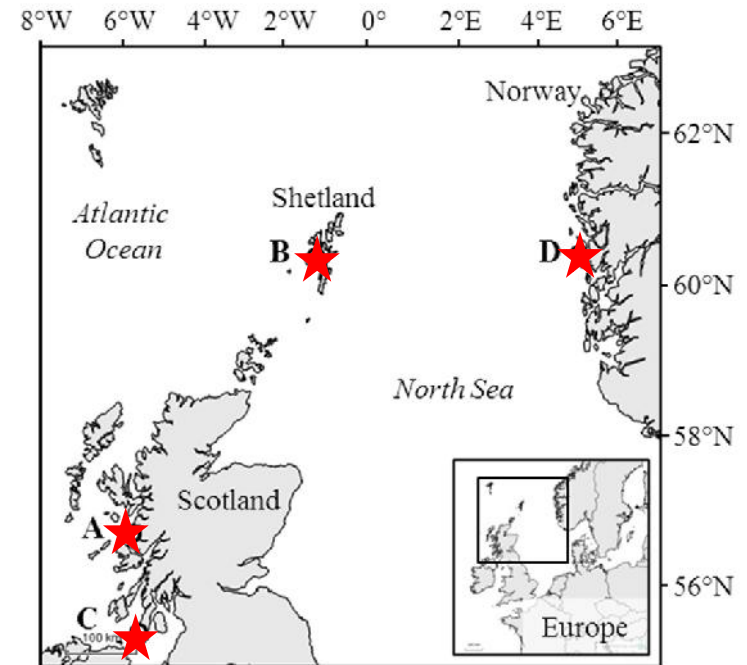
Raw- transformed into **Procruste-coordinates**  
(by translation and rotation)

Used to perform statistical analysis:

**Quantify shape differences between groups**  
**Predict groups based on shape**



- 400 wild ballan wrasse
- 4 locations across Scotland and Norway
  - Weight
  - Length
  - Photograph
  - Age classification by opercula rings
  - Histological confirmation of gender
  - Fin sample for genomics research



	SS	SS (%)	MS	P
<b>(a) Procrustes ANOVA: Centroid size</b>				
Location	2312268	78.99	770756.1	0.0001
Sex	112942	3.86	112941.7	0.0001
Individual	501973	17.15	1328.0	0.0001
Sum	2927183	100.00		
<b>(b) Procrustes ANOVA: Shape</b>				
Location	0.11806	29.03	0.00219	0.0001
Sex	0.00384	0.94	0.00021	0.0001
Individual	0.28479	70.03	0.00004	0.0001
Sum	0.40668	100.00		

## Variability in Shape/Size due to

1. Location
2. Individual
3. Sex





# Optimisation of size-based analysis

- 5 sets of variables tested

(1) BW, TL and K

(2) BW, TL and K + centroid size

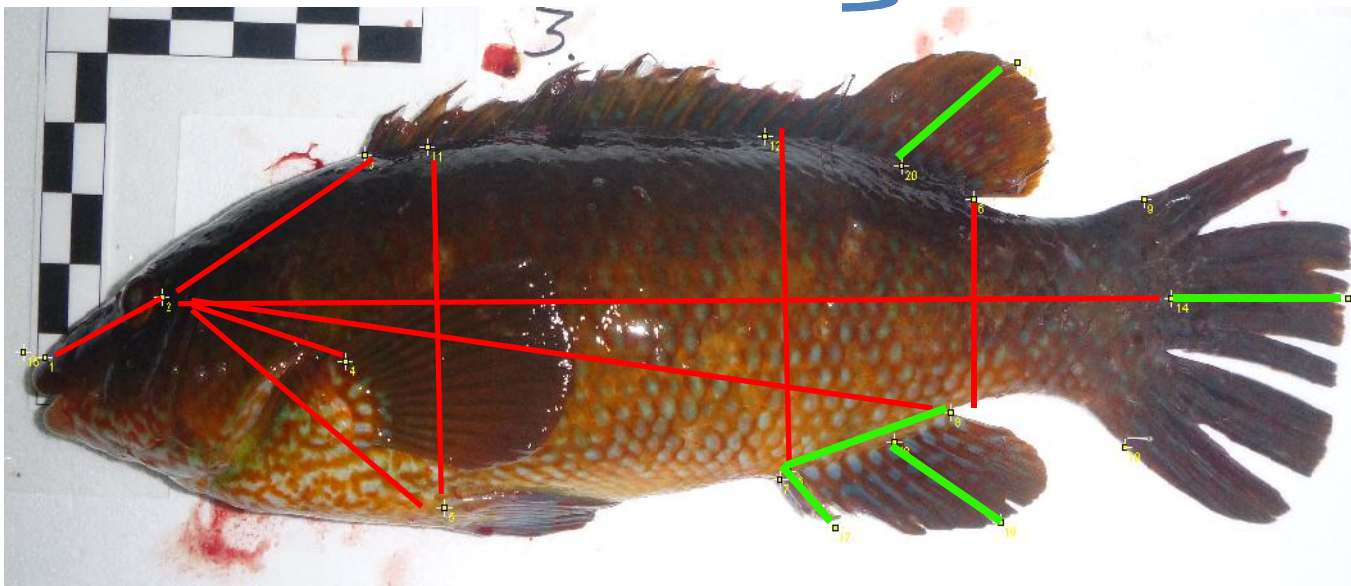
(3) Procruste distances:      Body ——— red ———  
    *from coordinates*        Fins ——— green ———

(4) Body and fin lengths + Centroid-size

(5) ALL

Simplest formula had a 91.4% accuracy

Accuracy of predicting  
gender was >85% in all cases

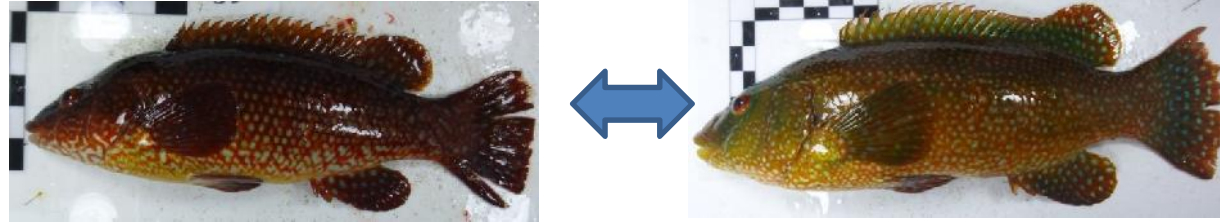




# Simplified on farm size-based analysis

$$D = (0.010 BW) + (-0.016 TL) + (-3.835 K) + 6.252$$

$$1.459 << 1.504$$

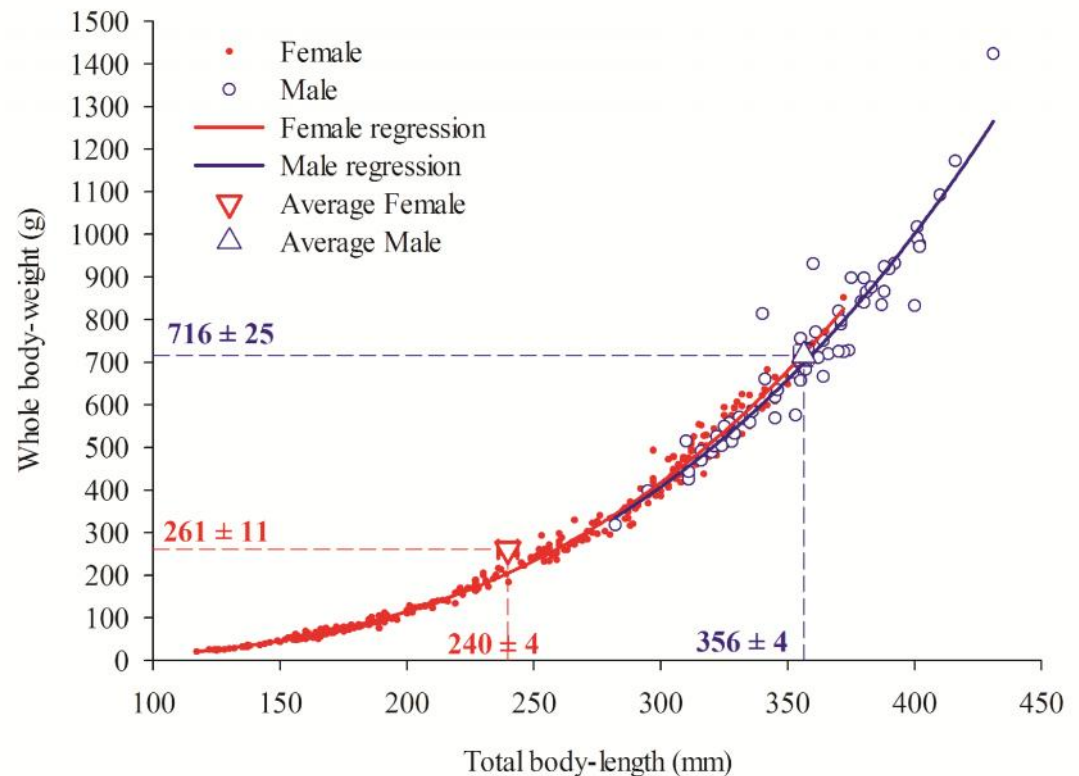


Simple calculation that can be performed in the field or on farm to predict gender

Size selection of gender in a protogynous hermaphrodite is not necessarily surprising.

Helps resolve the gender identification in sizes where both genders overlap, 30-40 cm TL.

Digital image analysis was central in deciphering the size/shape interactions





Aquaculture 155 (1997) 1–12

Aquaculture

Blastomere morphology as a predictive measure of  
fish egg viability

R.J. Shields <sup>a,\*</sup>, N.P. Brown <sup>b</sup>, N.R. Bromage <sup>b</sup>

<sup>a</sup> Sea Fish Industry Authority, Marine Farming Unit, Ardtoe, Acharacle, Argyll PH36 4LD, Scotland, UK

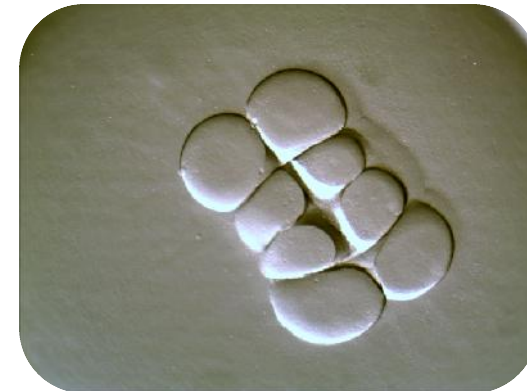
<sup>b</sup> Institute of Aquaculture, University of Stirling, Stirling FK9 4LA, Scotland, UK



Subjective scoring methodology of the “shape” of recently  
fertilised oocyte, blastomeres.

Score (1-5) of the blastomere:

- Cell symmetry
- Cell wall adhesion
- Cell size
- Cell margin definition
- Abundance of inclusion bodies



Good



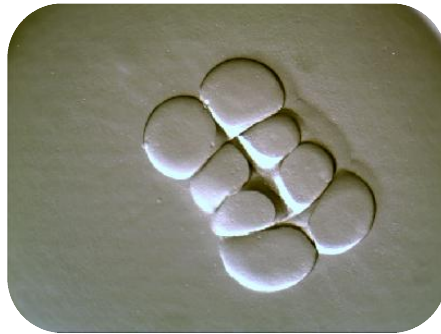
Bad



Ugly



# How subjective is a subjective scoring criteria?

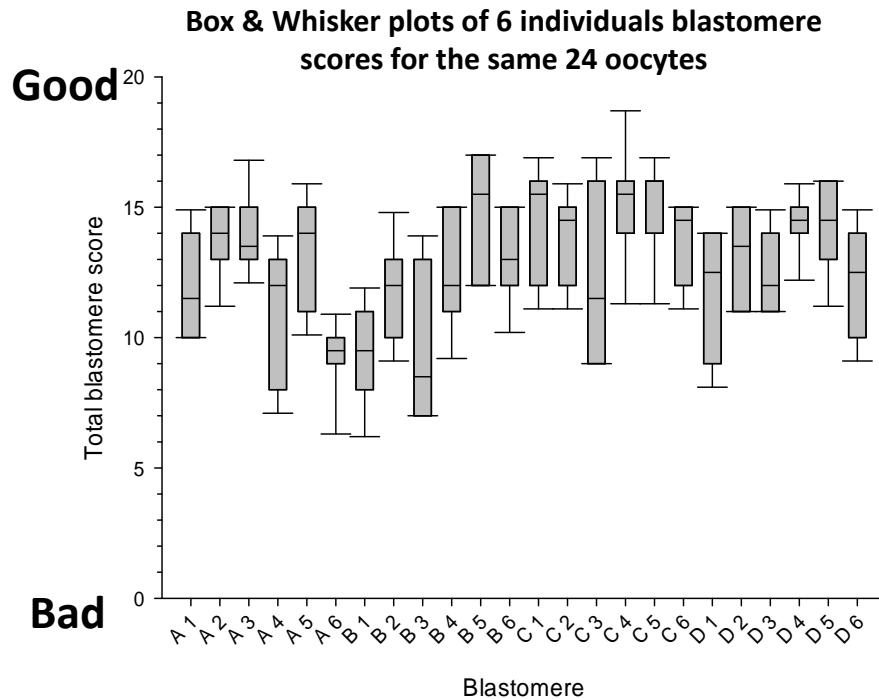


Criteria	Score
Symmetry	4
Cell size	3
Adhesion	2
Margins	4
Inclusions	3
<b>Total Score</b>	<b>16/20</b>

24 typical blastomere images were scored by 6 “informed” operators

- Previous training on the methodology
- Pictorial example scale for all criteria

## Conclusion



- “Operator variability” introduced approximately 20-30% noise on each blastomere
- There is a tendency to balance or compensate the scoring which results in a lack of variability in data points to be associated with the characteristic examined (i.e. quality)
- Operator subjectivity may, in part, explain the conflict in the literature as to the efficacy of the methodology.

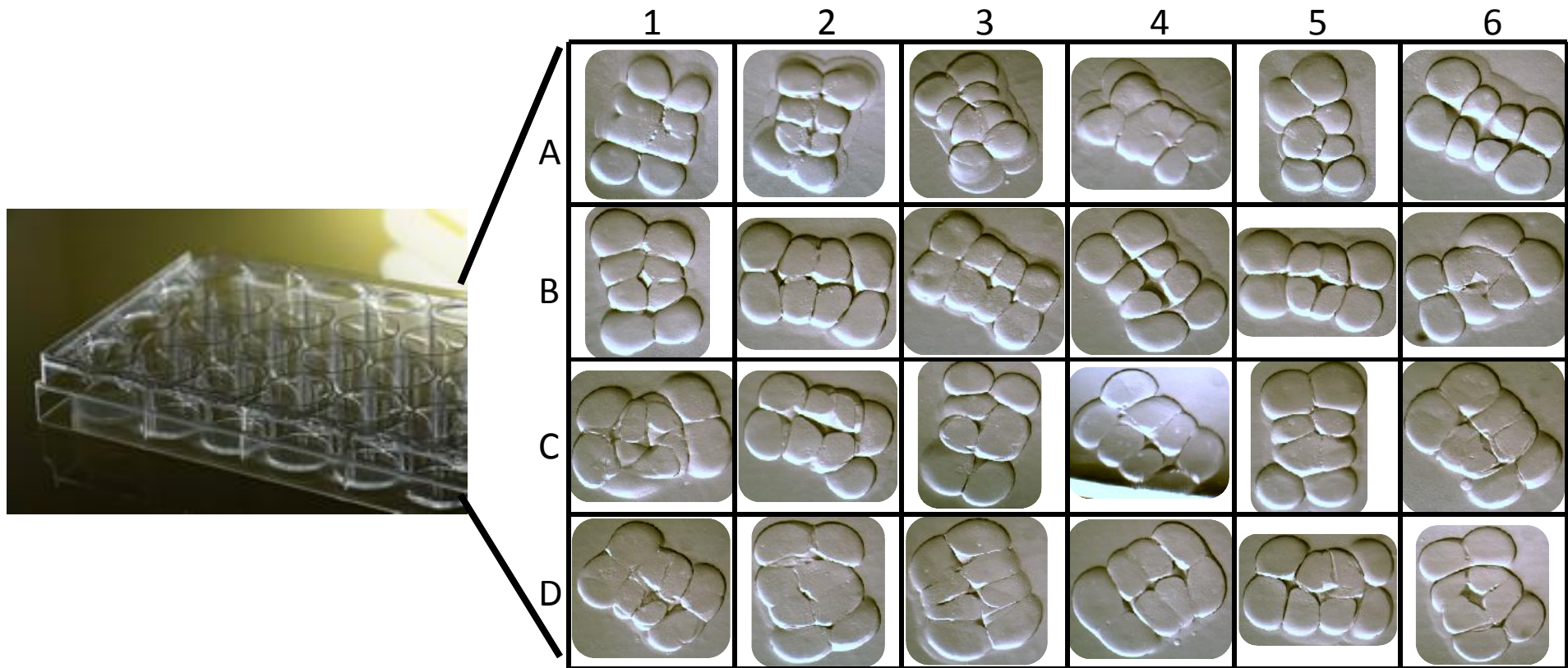


# Reverse engineering a DIA measuring method

Samples from a total of 8 Atlantic halibut egg batches were collected from Otterferry seafish Ltd.

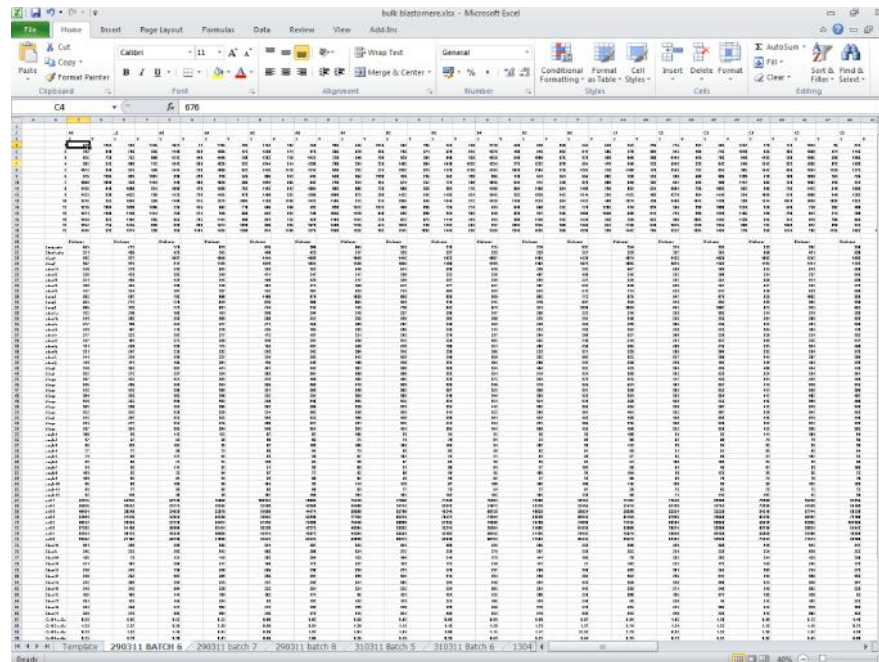
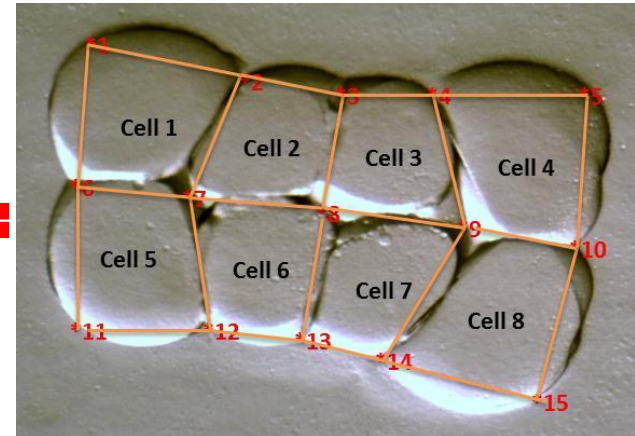
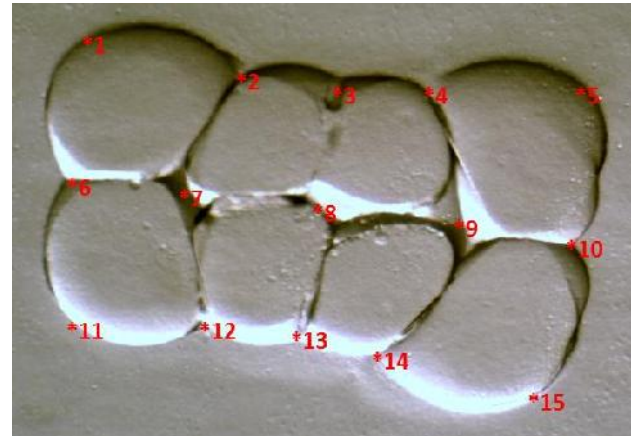
Using a routine plate culture method we photographed blastomeres (24 per batch, 192 in total) and then maintained them in constant environmental conditions to hatch.

Batch 6: 77.9% fertilisation rate





Analysis was based on 15 landmark points which in essence converts the blastomere into a crude 4 x 2 cell grid.



The X, Y coordinates of each landmark are imported into excel from which we calculated 74 separate measurements that were then amalgamated into 10 summary values for each blastomere

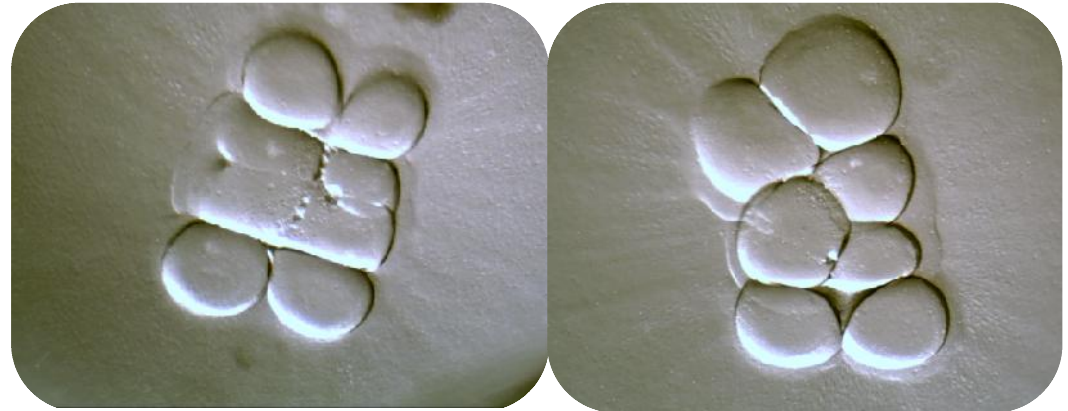
- Long to short axis ratio*
- Diagonal axis ratio*
- CV short lengths(%)*
- CV long lengths (%)*
- CV angle intersection*
- CV angle intersection 2*
- CV angle intersection 3*
- CV all intersection angles*
- CV Cell box area*
- CV Cell width ratio*



## Blastomeres with:

- Abnormal cell counts
- Margins poorly defined which prevents landmark point designation

Did not hatch. i.e. no evidence of “self repair”



Summary values in association with blastomere fate (hatched or failed) were then subjected to discriminant function analysis.

### Summary of classification

Put into Group	True Group	
	0	1
0	85	5
1	18	35
Total N	103	40
N correct	85	35
Proportion	0.825	0.875

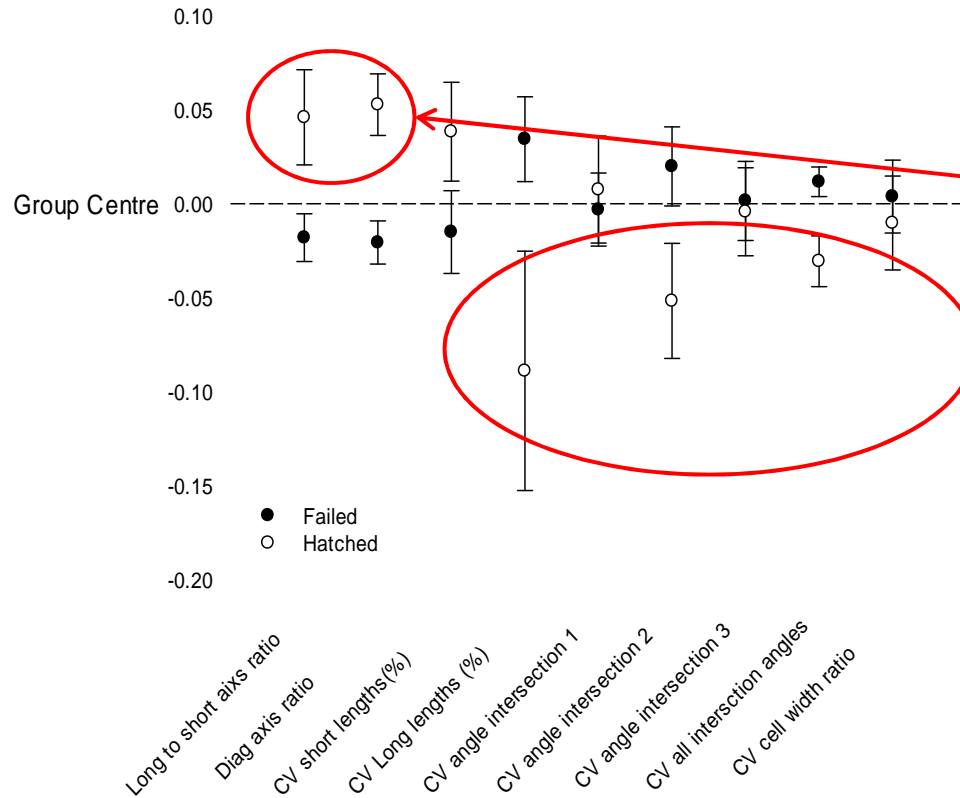
N = 143                  N Correct = 120

Proportion Correct = 0.839

Based on the summary values discriminant function analysis can correctly assign blastomere fate (hatch or fail) to 83.9% of oocytes



# What shape is a good blastomere?



Hatched blastomeres have a :

**Greater**

- Long to short axis ratio
- Diagonal axis ratio

**Lower**

- Variation in long axis lengths
- Variation in the angles of intersection between cells

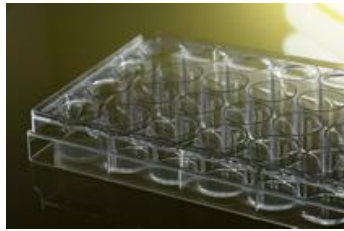
*Scatter plot of mean  $\pm$  SEM of the divergence from population mean dimension for hatched (n=40) and failed (n=103) halibut blastomeres.*

**Conclusion: Blastomeres which hatch are more rectangular and symmetrical**





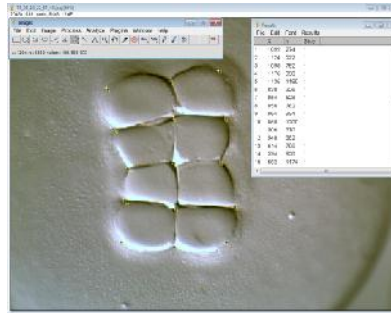
# Development and validation of a digital blastomere morphology measurement tool.



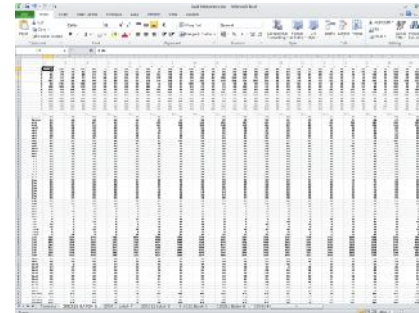
Plate



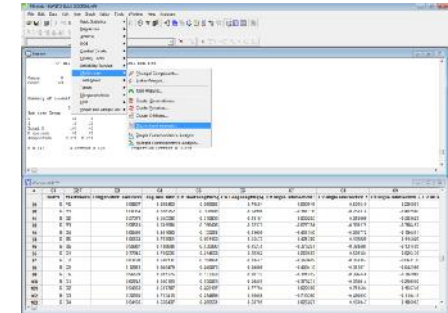
Image



Analyse in ImageJ



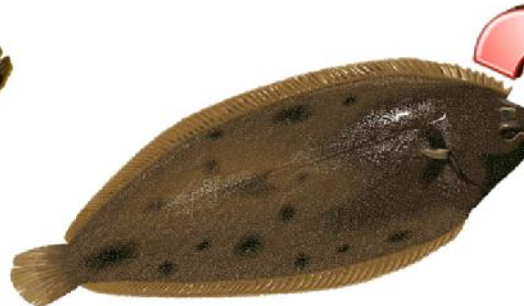
Process in Excel



Determine likelihood of hatch in Minitab

Analysis takes under 1 hour for a batch of 24 blastomeres  
You could have hatch predictions within 24 hours of fertilisation

This protocol has been designed in Halibut, which other species could it be applicable to...

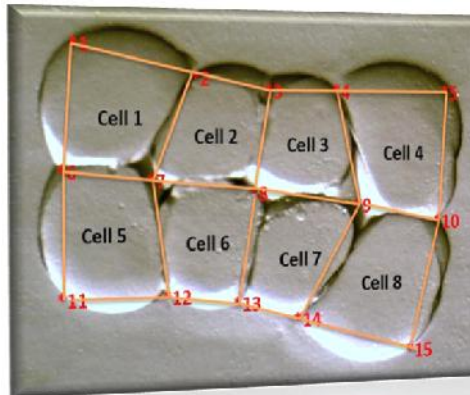




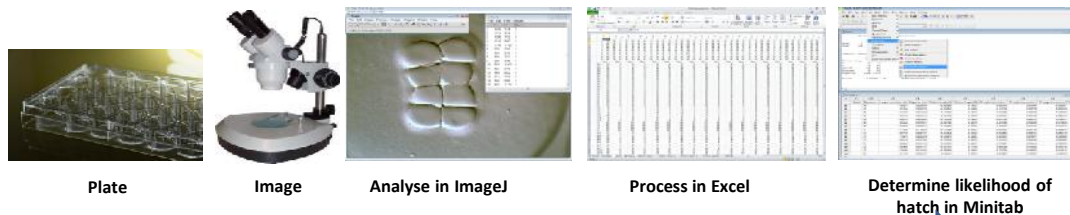
# Image analysis in hatchery management

Digital image analysis allows us to:

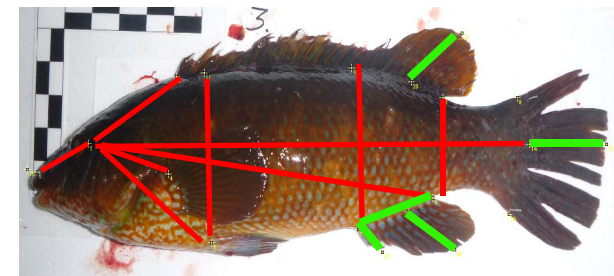
- Capture and describe small differences accurately
- Removes operator subjectivity



- + Open source software is accessible to all users be they academia or industry based
- + Provides real-time results
- + Capable of rapid large dataset set computation
- + Precision improves with use as reference databases expand.
- Not always the most practical technique on farm



Analysis takes under 1 hour for a batch of 24 blastomeres  
You could have hatch predictions within 24 hours of fertilisation





# Acknowledgments



[www.aqua.stir.ac.uk](http://www.aqua.stir.ac.uk)

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- Alasdair Barge
- David Patterson

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EU FP7 project: 232305 "Prospawn"