An evolutionary hypothesis for the radiation of the Gyliauchenidae Goto & Matsudaira, 1918 (Platyhelminthes: Digenea)

Kathryn A. Hall^{*1,2}

T. H. Cribb¹, R. A. Bray² and D. T. J. Littlewood²

The University of Queensland, Brisbane, AUSTRALIA
The Natural History Museum, London, UK



Pacific Ocean

• Distribution of Gyliauchenidae

Methodology

Available data:

- 49 species characterised by morphological data (light microscopy, literature review)
- 19 species characterised by sequence information (ND1 mtDNA, 28S rDNA)

Analysis:

 parsimony analysis: heuristic searches and quartet puzzling within PAUP* (Swofford, 1999)

Phylogenetic correlation of gut adaptation

- gut becomes more complex through time
- hierarchy of synapomorphies for gyliauchenid clades
- heritability of characters has taxonomic implications

Phylogeny of the Acanthuroidea after Winterbottom (1993)



Ecological correlations

- acanthuroid fishes are predominantly herbivores
- host gut structure broadly conserved
- extreme gut adaptations restrict opportunity for host-switching
- specificity to acanthuroid fishes is an ecological association
- gyliauchenid radiation does not necessarily involve co-evolution

Further testing

- further collecting in south Pacific and east Indian oceans
- incorporation of any new taxa into phylogenies
- host mapping
- biogeography mapping
- fine level analysis of host-parasite evolution

Acknowledgements

- Australian Research Council
- British Council (Chevening Fellowship Trust)
- The Natural History Museum:
 - Ms Elisabeth Herniou, Ms Julia Bartley and Biomedical Parasitology Unit
- Australian Society for Parasitology