



The evolution of blood parasitism in flatworms

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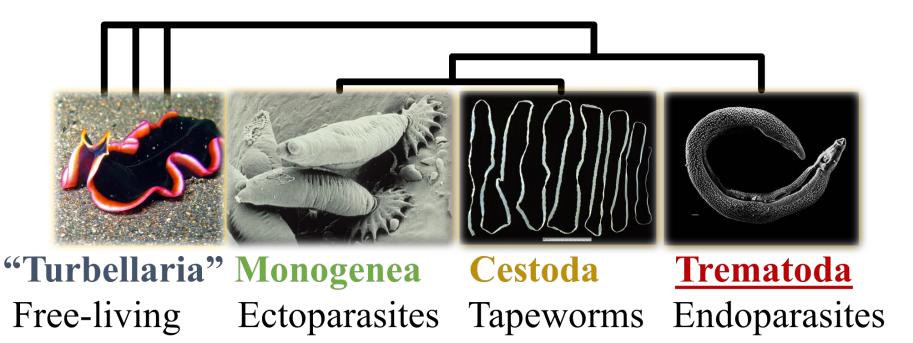


10 parasitic lineages contain 40% of all animal diversity

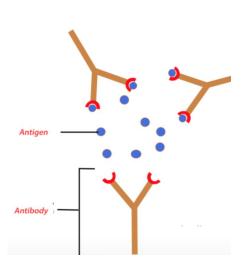
Parasitism evolved at least 223 times in Animalia

Weinstein & Kuris Biol Letters (2016)

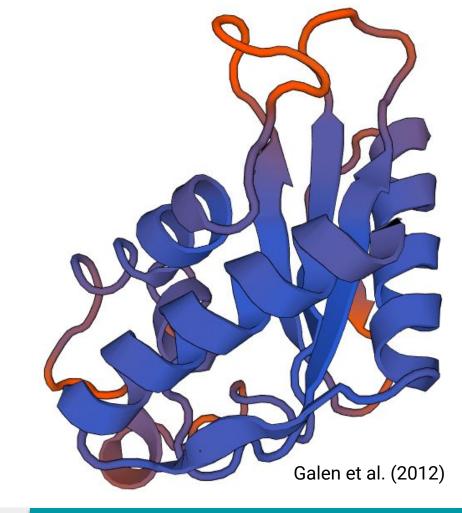
## **System: Flatworms**



Antagonistic coevolution btwn parasite allergens & host antibodies

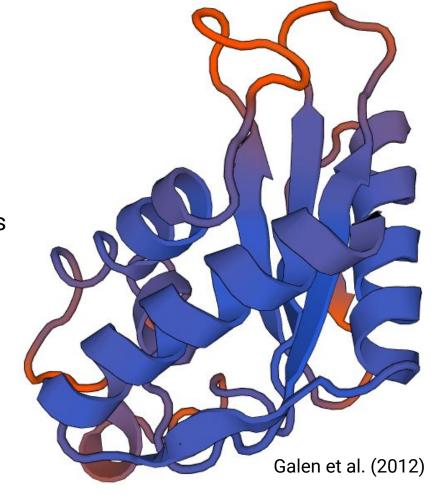






- 2 main groups of VAPs
  - Group 1: Secretory
  - Group 2: Intracellular
- Secretory VAPs from parasitic flatworms may be under (+) selection

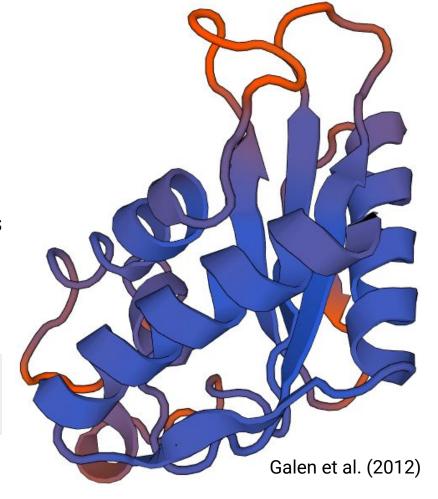
Chalmers et al (2008) & Chalmers & Hoffman (2012)

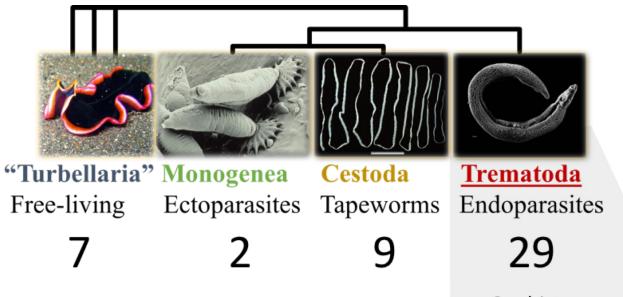


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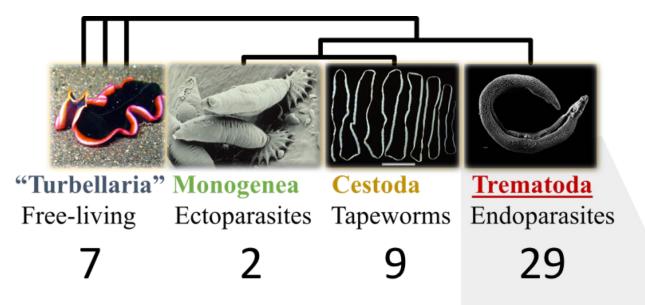
Included only parasites with medical/ economic relevance





- 3 schistosomes
- 21 non-schistosome blood flukes
- 5 non-blood flukes

## **Data: 47 transcriptomes**



H1: VAPs expressed in all flatworms

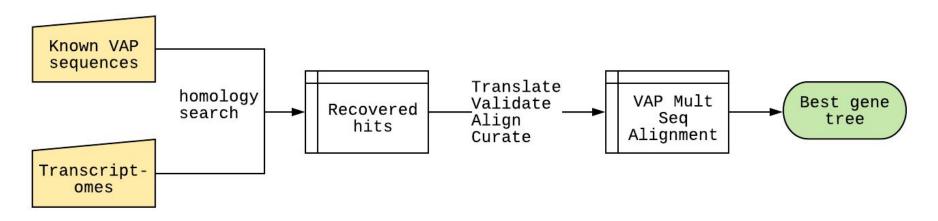
**H2:** Two well-supported major VAP clades

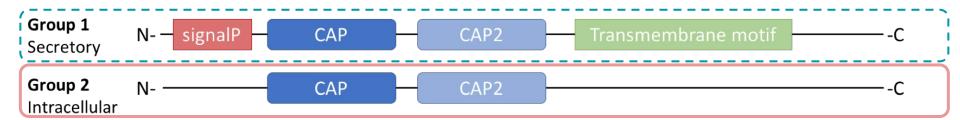
H3: Signal peptides predicted in Group 1 VAPs

**H4:** Taxa-specific expansions in secretory VAPs

- 3 schistosomes
- 21 non-schistosome blood flukes
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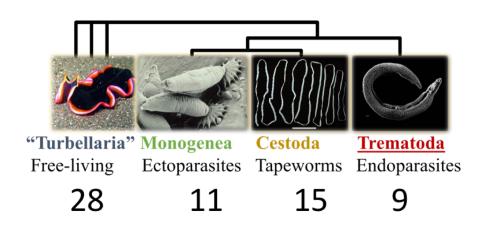
# **Bioinformatics pipeline**





H1: VAPs expressed in all

flatworms



474 new VAPs

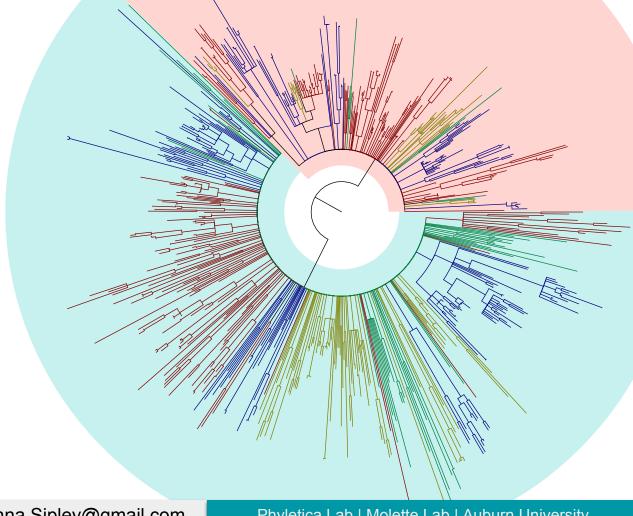


H1: VAPs expressed in

all flatworms

**H2:** Two well-supported

major VAP clades



99% bootstrap support



H1: VAPs expressed in

all flatworms

**H2:** Two well-supported

major VAP clades

**H3:** Signal peptides

predicted in Group 1 VAPs

Yes & only 1 SignalP predicted in Group 2



H1: VAPs expressed in

all flatworms

**H2:** Two well-supported

major VAP clades

**H3:** Signal peptides

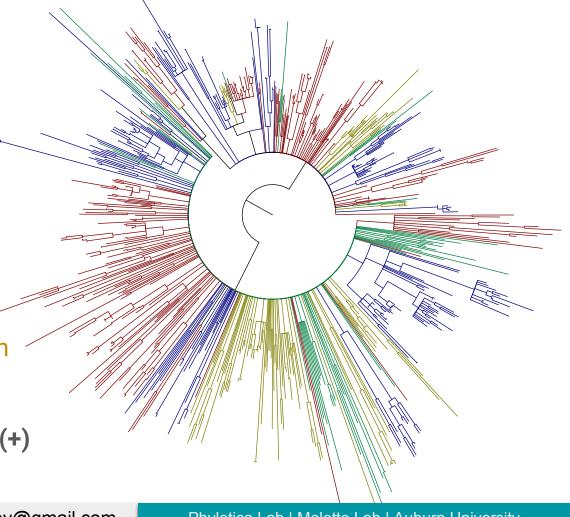
predicted in Group 1 VAPs

H4: Taxa-specific expansions in

secretory VAPs

**Deeper exploration & tests for (+)** 

selection to come...



# Acknowledgements

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