Genome-wide sexually antagonistic variants reveal longstanding constraints on sexual dimorphism in the fruit fly

Max Reuter



Sexual dimorphism



Sexual dimorphism



Dimorphism and genes

Males and females share genomes

Few genes are restricted to one sex

| Organism | Total genes | Y/W genes |
|-------------|-------------|-----------|
| Human | 20,441 | 71 |
| Jungle fowl | 18,346 | 25 |
| Fruitfly | 13,918 | 22 |









Sexual antagonism in theory

Population genetic models



Rice (1984); Gavrilets & Rice (2006)

Antagonism in real organisms



Drosophila melanogaster – LH_M



Chippindale et al. (2001)

Antagonism in general

Widespread

Many species of animals and plants

Sexual dimorphism often is incomplete

Why does antagonism persist? What are its evolutionary dynamics?

Which loci underlie antagonism?



Genetic basis of antagonism

Largely unknown, individual examples

e.g., maturation/body size QTL in salmon





Barson et al. (2015, Nature)

Genetic basis of antagonism

Here: a genome-wide effort in *D. melanogaster* (LH_M)

Which SNPs are sexually antagonistic?

What do they do?

How do they evolve?

Data



Drosophila melanogaster, LH_M

200+ hemiclonal lines

Measures of male and female competitive fitness

Whole genome sequences, 765,000 SNPs

Quantitative genetics



 h_f^2 =0.42, CI=(0.30, 0.54) h_m^2 =0.16, CI=(0.04, 0.27)

$$r_{MF}$$
=0.15, CI=(-0.21, 0.46)

Which SNPs are sexually antagonistic?

What do they do?

How do they evolve?

Candidate SNPs

Univariate GWAS on antagonistic score



2,372 SNPs with FDR > 0.3, 226 clusters

Which SNPs are sexually antagonistic?

What do they do?

How do they evolve?

Function

Antagonistic SNPs are enriched in coding sequences

Large excess of missense variants



Function

514 genes with ≥1 significant SNP

No clear GO enrichments

(Slightly) below-average sex bias in expression



Which loci are sexually antagonistic?

What do they do?

How do they evolve?

Antagonism can generate balancing selection





Antagonism is associated with elevated MAF



Antagonism is associated with elevated MAF



... and reduced population differentiation



P<0.001

... and trans-specific polymorphism with *D. simulans*





Genome-wide candidates for sexual antagonism

Adaptive conflict over protein sequence More difficult/slow to resolve



Antagonism stabilises polymorphism For long periods of time Within and across species



Acknowledgements

Filip Ruzicka Mark Hill Kevin Fowler Richard Mott

Doug Speed Aida Andrés Ted Morrow Tanya Pennell Fiona Ingleby Ilona Flis Will Gilks





Concordant fitness variation

Highly polygenic, mutation-selection balance



Window-wide increase in polymorphism

