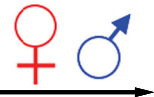


Heterogametic Adjustments

Step 1

Test for DE between
Male & Female
gonozooid libraries.



Step 2

Separate transcripts
by DE results.

Non-significant transcripts.



Up reg. female transcripts.



Up reg. male transcripts.



Step 3

Test for DE between
all pairwise combinations
of polyp types on
two separate datasets.

Use both male & female
gonozooid counts.

Use male gonozooid
counts only.

Discard.

List of
DEs