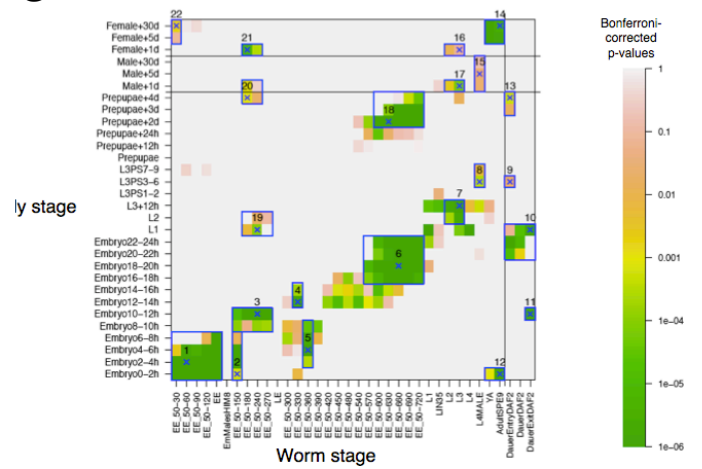


C Developmental stage mapping between worm and fly based on co-expression clustering of orthologs



D

	Worm	Fly	Human
# of orthologs	5083	4839	6353
# of co-expression clusters (h=0.9)	85	108	67
# of highly correlated ncRNAs (r>0.6)	16569	2142	8173
# of nearby and highly correlated ncRNAs (r>0.6) (for 1-1-1 orthologs only now)	74 (< 5k)	30 (<20k)	246 (<150k)
# of potential orthologous ncRNAs (in pairs or triplets) (for 1-1-1 orthologs only now)			

