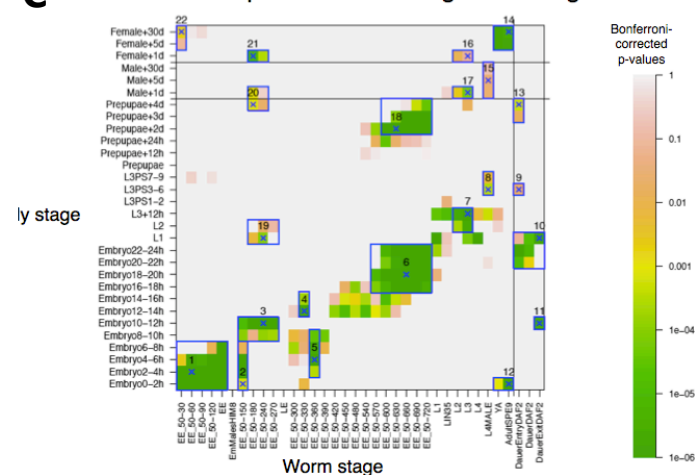


C Developmental stage mapping between worm and fly based on co-expression clustering of orthologs



Brenner, Bickel, Huang, Li in UC Berkeley

D

	Worm	Fly	Human
# of w-f-h 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 ($r>0.6$)	741 (< 5k)	160 (<20k)	1141 (<150k)
# of potential orthologous ncRNAs (in pairs or triplets)			

