Sex and the Single Fly: A Perspective on the Career of Bruce S. Baker

陈洁 金思慧 邢丽敏 2020.1.3



Bruce S. Baker

Affiliations: 1976-1985 University of California, San Diego, La Jolla, CA 1986-2008 Stanford University, Palo Alto, CA 2008- HHMI Janelia Farm Research Campus, Ashburn, VA, United States Area: Genetics, Neuroscience

Parents			
Lawrence Sandler	grad student	<mark>1971</mark>	University of Washington
(Sex chromosome characterization)	meiotic mutants in	Drosophila	melanogaster : detection and preliminary
James F. Crow	post-doc	1972-19	74 UW Madison



Bruce S. Baker

Affiliations: 1976-1985 University of California, San Diego, La Jolla, CA 1986-2008 Stanford University, Palo Alto, CA 2008- HHMI Janelia Farm Research Campus, Ashburn, VA, United States

Area: Genetics, Neuroscience

Children				Kenneth C. Burtis	post-doc		
Dori Allen	grad student			Bruce Chase	post-doc		
Debbie Andrew	grad student			Audrey Christiansen	post-doc		
Deborah J. Andrew	grad student		(E-Tree)	Mitzi Kuroda	post-doc		
Monica Gorman	grad student		Stanford	Bill Mattox	post-doc		
Elizabeth H. Chen	grad student	1998	Stanford (Neurotree)	Michael McKeown	post-doc		
Carrie M. Garrett-Engele	grad student	2000	Stanford	Rod Nagoshi	post-doc		
Suzanne D. Plump	grad student	2000	Stanford	Brian Oliver	post-doc		
Eric L. Keisman	grad student	2001	Stanford	Lisa Ryner	post-doc		
Shaad M. Ahmad	grad student	2002	Stanford	Mariana Wolfner	post-doc		
Devanand S. Manoli	grad student	2007	Stanford (Neurotree)	Yufeng Pan	post-doc	2009-	Janelia Farm
Joy Hatzidakis	grad student	2008	Stanford	Chuan Zhou	post-doc	2011-	Janelia Farm (Neurotree)
David J. Mellert	grad student	2009	Stanford	Ignacio Marin	post-doc	1993-1998	
Alexander G. Vaughan	grad student	2012	Stanford University, I	Mark L. Siegal	post-doc	1998-2004	Stanford
Ounissa Ait-Ahmed	post-doc			Delphine Fagegaltier	post-doc	2001-2004	Stanford
Michelle N. Arbeitman	post-doc		Stanford				
John Belote	post-doc						

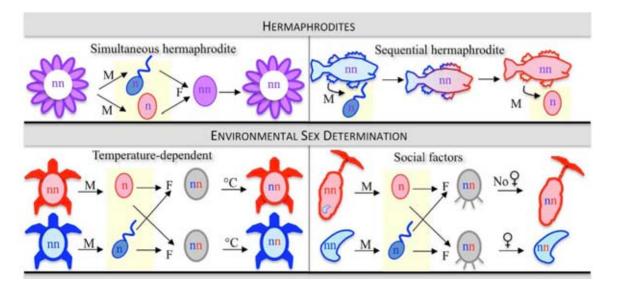
Sex and the Single Fly: A Perspective on the Career of Bruce S. Baker

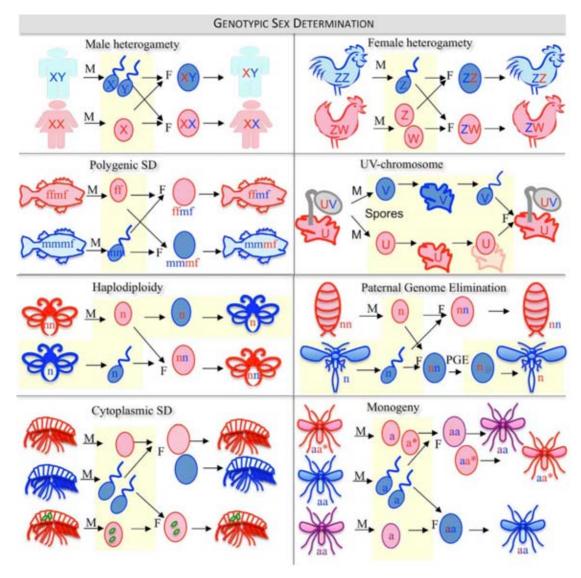
- Sex Determination and Dosage Compensation in *Drosophila* (CJ)
- The Development of Sexually Dimorphic Structures and the Evolution of Sex (JSH)
- Sex Behavior Meets the Sex Determination Regulatory Hierarchy: The Genetic Control of Sexual Behavior (XLM)

Sex Determination and Dosage Compensation in Drosophila



Sex Determination





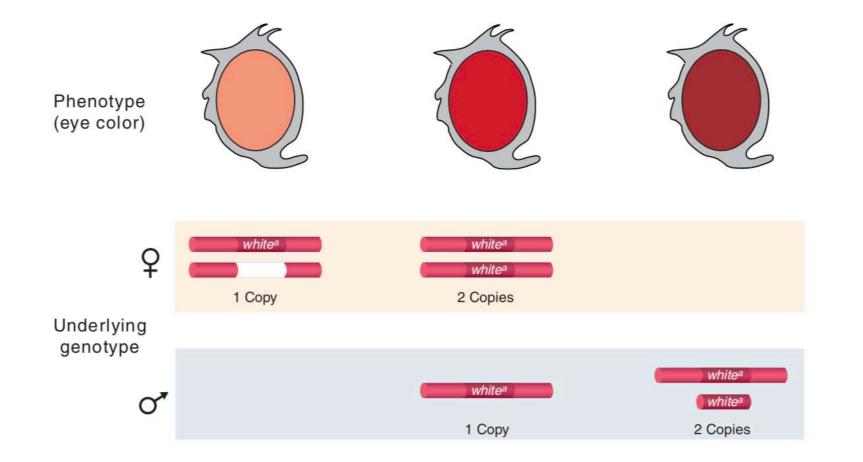
Doris Bachtrog et al. PLOS Biology .2014

Sex Determination and sex chromosomes

	Sex Determination
Drosophila	X chromosome: autosome ratio (X:A ratio) In Drosophila, the Y chromosome has no role in sex determination
Human	The presence or absence of a Y chromosome

	XY (2A)	XX (2A)	XXY (2A)	XYY (2A)
Drosophila	male	female	female	male
Human	male	female	metafemale	metamale

The discovery of dosage compensation in drosophila



John C. Lucchesi and Mitzi I. Kuroda . Cold Spring Harb Perspect Biol. 2015

Dosage compensation

Across species, different sexes are often characterized by different types and numbers of sex chromosomes. In order to account for varying numbers of sex chromosomes, different organisms have acquired unique methods to equalize gene expression amongst the sexes.

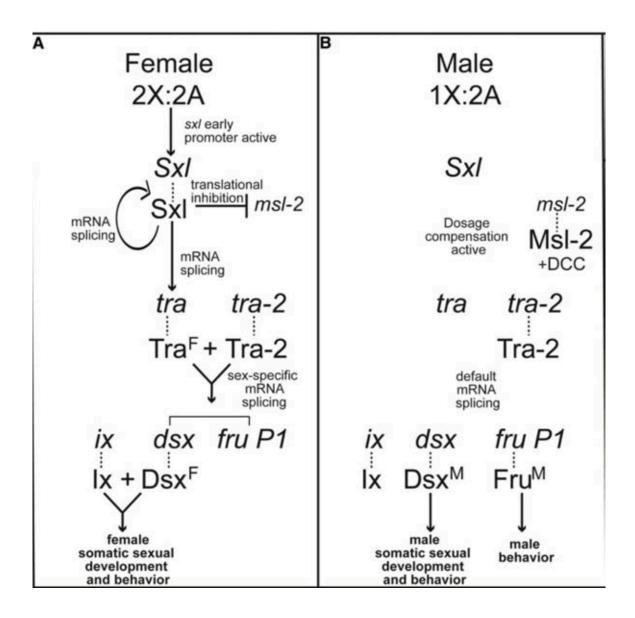
There are three main mechanisms of achieving dosage compensation which are widely documented in the literature and which are common to most species.

1. random inactivation of one female X chromosome (as observed in Mus musculus)

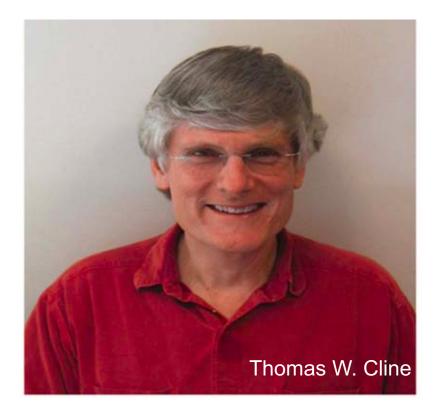
2. a two-fold increase in the transcription of a single male X chromosome (as observed in Drosophila melanogaster)

3. decreased transcription by half in both of the X chromosomes of a hermaphroditic organism (as observed in Caenorhabditis elegans).

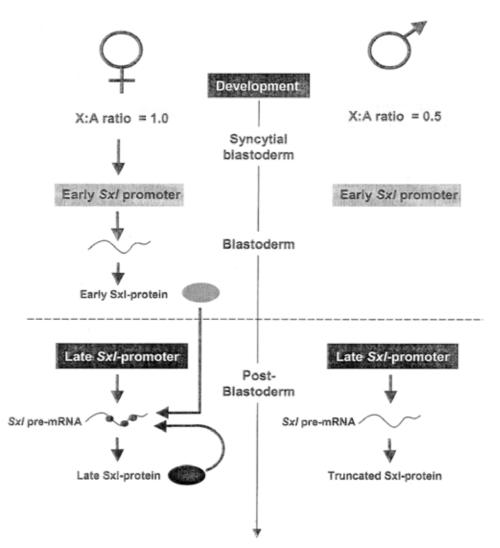
The somatic sex determination hierarchy in Drosophila



Thomas W. Cline and Sex lethal (Sxl)



Regulation of *Sxl* expression



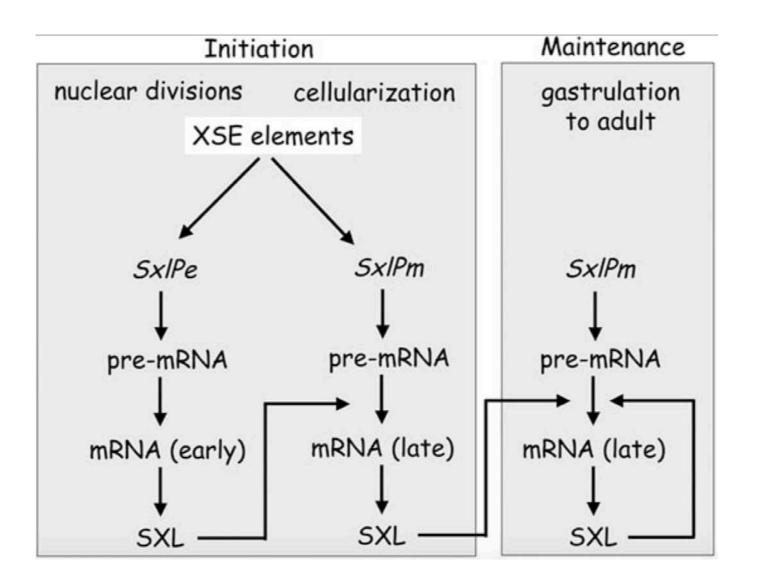
Luiz O. F. Penalva et al. Microbiol Mol Biol Rev. 2003

How does *SxIPe* reliably distinguish between the ratio of X chromosomes and autosomes ?

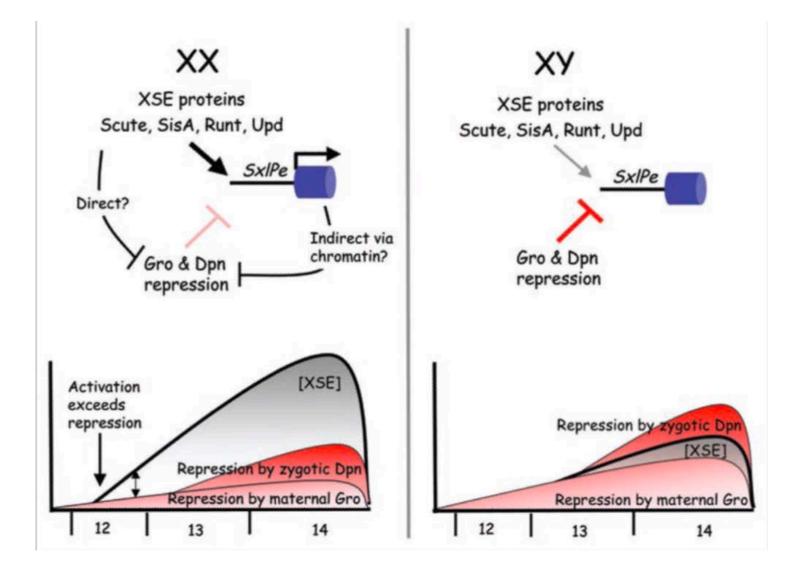
XSE :X-linked signal elements

four X- encoded proteins, encoded by the scute, sisA, runt and unpaired

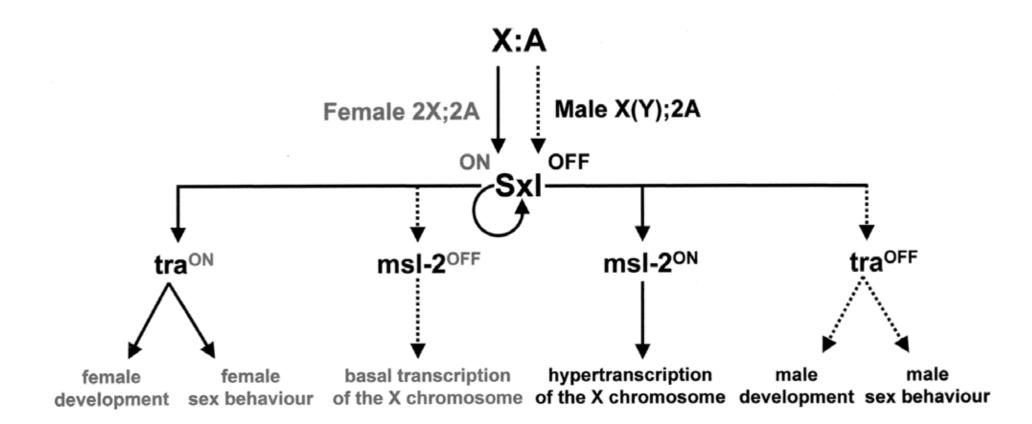
Overview of the regulatory logic that guarantees *SxI* protein expression in XX animals.



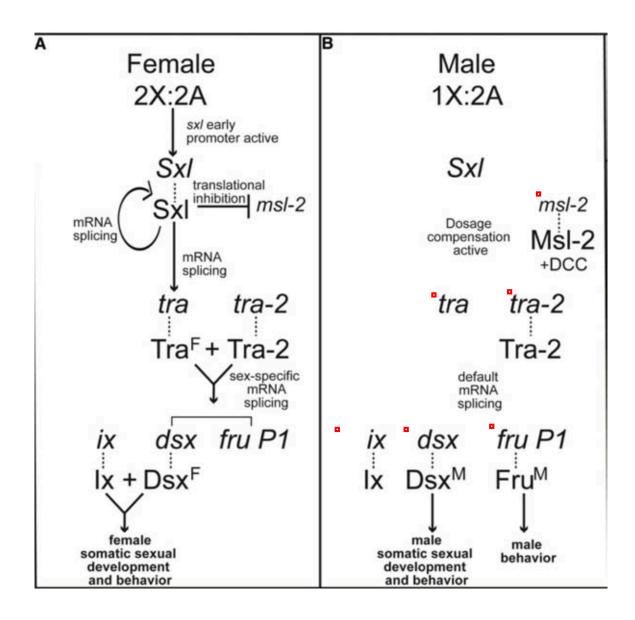
Threshold response model



Sxl, sex determination and dosage compensation



The somatic sex determination hierarchy in Drosophila



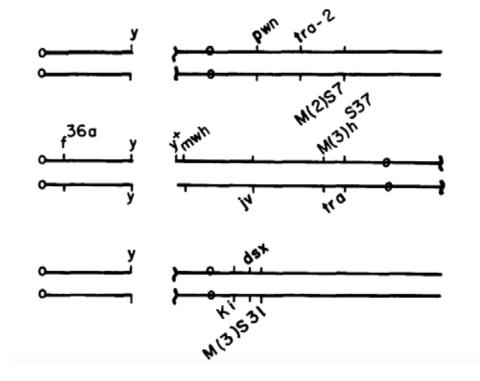
SEX AND THE SINGLE CELL. I. ON THE ACTION OF MAJOR LOCI AFFECTING SEX DETERMINATION IN DROSOPHILA MELANOGASTER¹

BRUCE S. BAKER AND KIMBERLY A. RIDGE

- That the four genes known at the time to cause dramatic sex-transformation phenotypes in Drosophila— *transformer (tra), transformer-2 (tra-2), doublesex (dsx)*, and *intersex (ix)*—function in a cell-autonomous manner
- That these genes function in a shared genetic pathway

Locus (symbol)		Chromosome-map position	Relevant properties	Major references;	
I. Sex determin	ation mutants:				
transformer-	2 (tra-2)	2-70	Transforms females into males; males sterile	WATANABE 1975	
	$(tra-2^{OTF})$		Incomplete transformation of females into males; males fertile	FUJIHARA, KAWABE and OISHI 1978	
intersex	(ix) (ix^2)	2-60.5	Transforms females into intersexes; males normal Like <i>ix</i>	Kroeger 1959	
transformer	(tra) (tra^{AC})	3–45	Transforms females into males; males normal Like <i>tra</i>	Sturtevant 1945; Seidel 1963 This report	
doublesex	(dsx)	3-48.1	Transforms both males and females into intersexes	HILDRETH 1965	
doublesex-do	minant (dsx^D)		Dominant, dsx ^D /+ transforms females into intersexes; male unaffected	Fung and Gowen 1957; Duncan and Kaufman 1975	
Masculanized	r (Mas=dsx ^{Mas}))	Like dsx^D		

That the four genes known at the time to cause dramatic sex-transformation phenotypes in Drosophila— *transformer (tra), transformer-2 (tra-2), doublesex (dsx)*, and *intersex (ix)*—function in a cell-autonomous manner



Analysis of autonomy of the tra-2 locus and time of tra-2+ expression in the abdomen

		Clo	nes				
	Tergites 5 and 6			Tergites			
Male pwn	Male no <i>pwn</i>	Indetermi- nate pwn	Female pwn	2,3,4 pwn	Number of abdomens	Frequency male clones	Frequency pwn clones
2	2	1	0	4	295	0.014	0.024
, hrs:							
59	17	2	2	111	393	0.193	0.443
14	9	1	0	24	106	0.217	0.368
22	9	2	1	28	142	0.218	0.373
5	1	1	1	13	41	0.146	0.488
n, hrs:							
8	4	0	1	17	44	0.273 ± 0.067	0.591
14	16	1	10	70	107	0.280 ± 0.043	0.888
25	34	8	57	174	193	0.306 ± 0.033	1.368
6	0	1	96	160	113	0.053 ± 0.019	2.33
	pwn 2 59 14 22 5 n, hrs: 8 14 25	Male pwn Male no pwn 2 2 1 59 17 14 9 22 9 5 1 1 n, hrs: 8 4 14 16 25 34	Tergites 5 and 6 Male pwn Male no pwn Indetermi- nate pwn 2 2 1 1 1 1 59 17 2 14 9 1 22 9 2 5 1 1 n, hrs: 8 4 0 14 16 1 25 34	$\begin{array}{c ccccccccccccccccccccccccccccccccccc$	$\begin{array}{c c c c c c c c c c c c c c c c c c c $	$\begin{array}{c c c c c c c c c c c c c c c c c c c $	$\begin{array}{c c c c c c c c c c c c c c c c c c c $

y/+; pwn tra-2/+ female progeny

That these genes (*tra, tra-2, dsx*, and *ix*) function in a shared genetic pathway

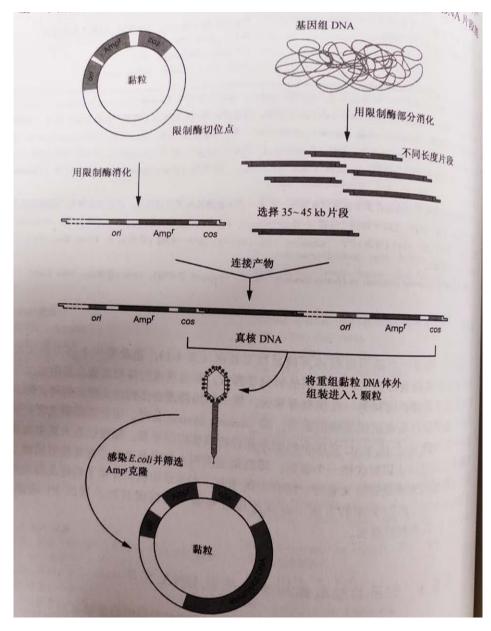
Method: Double-mutant combinations

In a pathway, then the double-mutant combination should exhibit the same phenotype as one of the component single mutants.

In parallel pathways, the double-mutant combination should exhibit a phenotype that is a composite of the phenotypes produced by the two component single mutants.

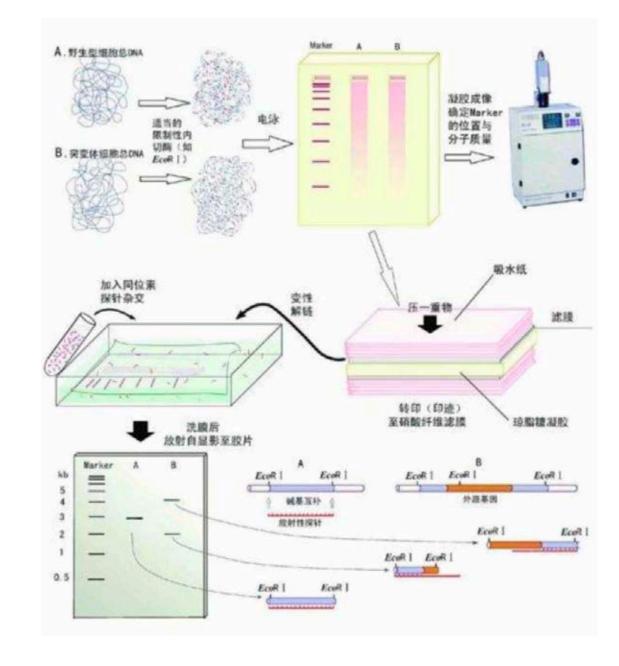
mutant	phenotype	Conclusion
<i>dsx</i> mutant	Intersexes	<i>dsx</i> is epistatic to <i>tra</i>
tra mutant	Transforms female into male	
<i>dsx</i> + <i>tra</i> mutant	Intersexes	

Cosmid clones for the chromosome walk



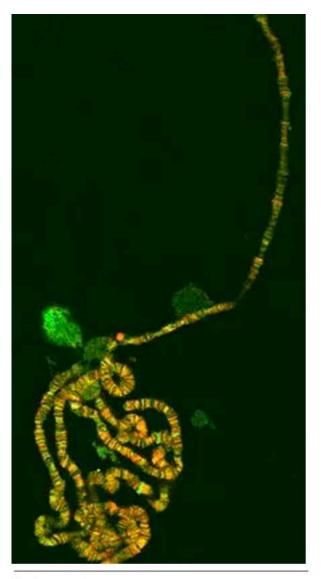
Southern blot

A Southern blot is a method used in molecular biology for detection of a specific DNA sequence in DNA samples. Southern blotting combines transfer of electrophoresis-separated DNA fragments to a filter membrane and subsequent fragment detection by probe hybridization.



Polytene chromosome

Polytene chromosomes are oversized chromosomes which have developed from standard chromosomes and are commonly found in the salivary glands of Drosophila melanogaster. Specialized cells undergo repeated rounds of DNA replication without cell division (endomitosis), to increase cell volume, forming a giant polytene chromosome. Polytene chromosomes form when multiple rounds of replication produce many sister chromatids that remain fused together.

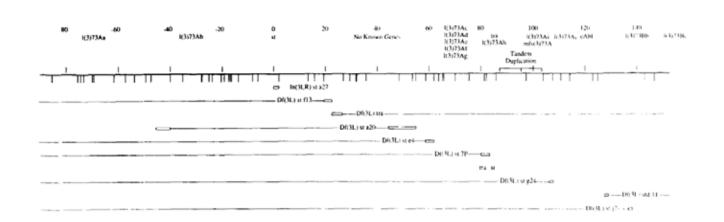


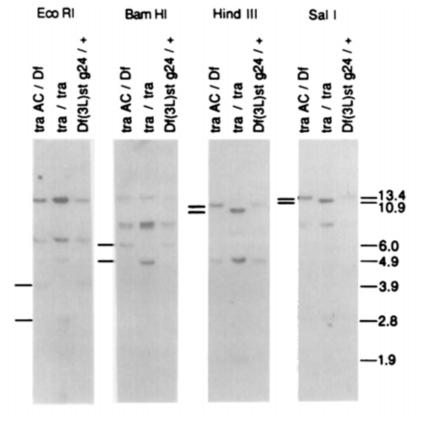
Polytene chromosome



A Molecular Analysis of *transformer*, a Gene in Drosophila melanogaster That Controls Female Sexual Differentiation

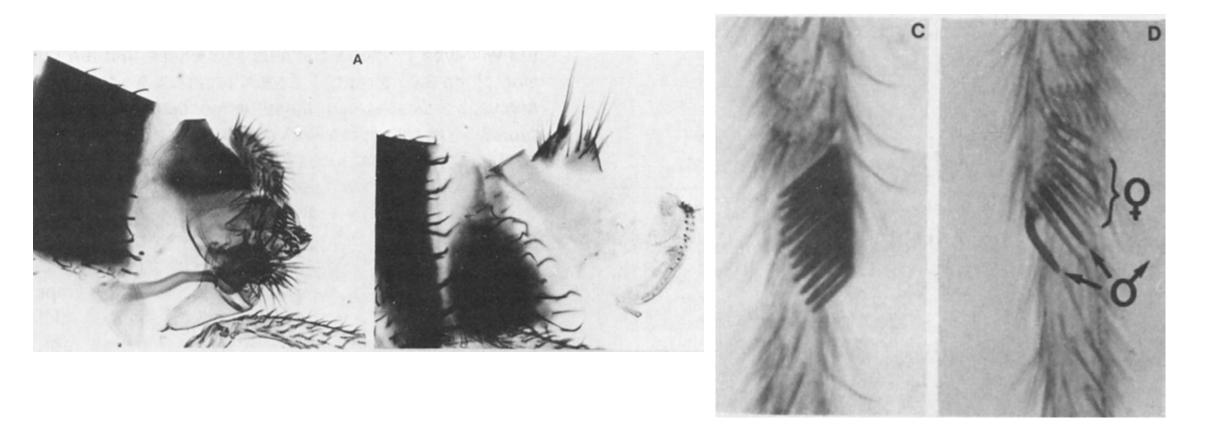
Michael McKeown,* † John M. Belote,* ‡ and Bruce S. Baker* §





Southern blot

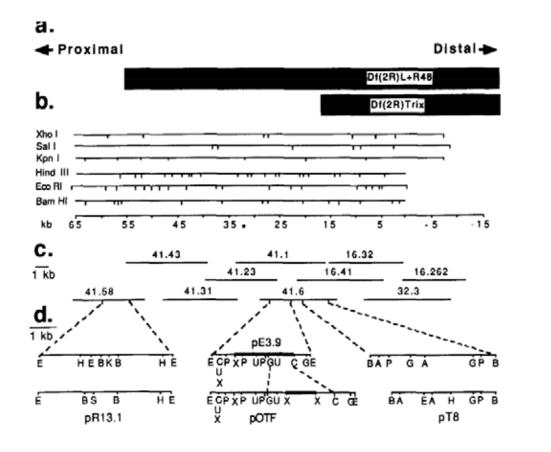
Identification of the tra Gene by Germ-Line Transformation

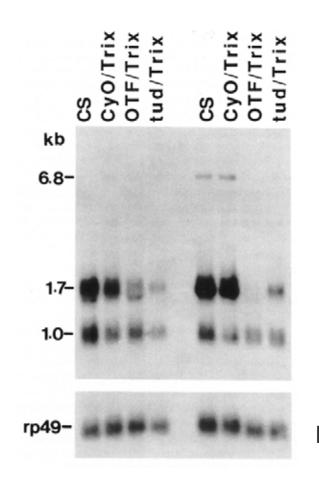


A) Male genitalia of XX; *tra*/Df(3L)st j7, Ki roe p^p transformed female. (B) Female genitalia of an XX; ; *tra*/Df(3L)st j7, Ki roe p^p wild-type female. (C) Sex comb of XX ; *tra*/Df(3L)st j7, Ki roe p^p transformed female. (D) Intersexual sex comb of an X, RB161X; ; *tra*/Df(3L)st j7, Ki roe p^p transformant.



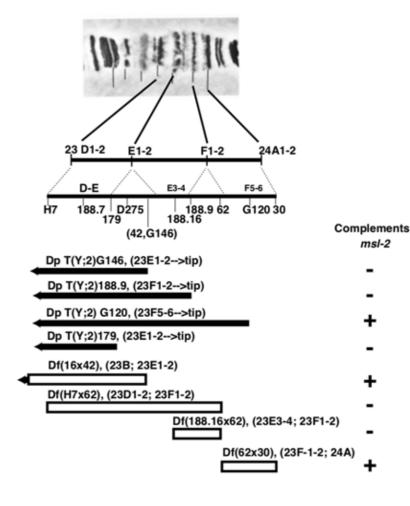
The Sex Determination Locus *transformer-2* of Drosophila Encodes a Polypeptide with Similarity to RNA Binding Proteins

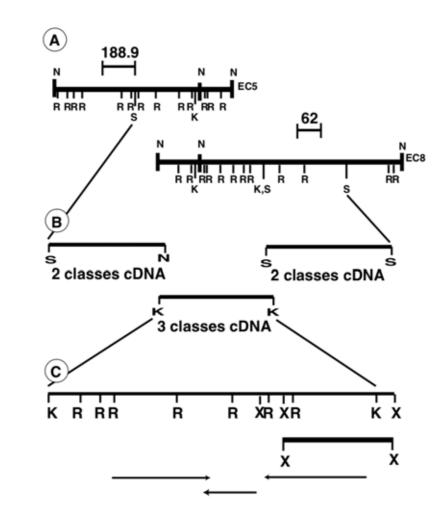




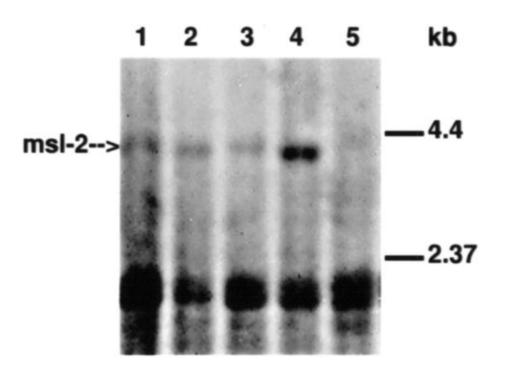
Northern blot

The *msl-2* dosage compensation gene of *Drosophila* encodes a putative DNAbinding protein whose expression is sex specifically regulated by *Sex-lethal*

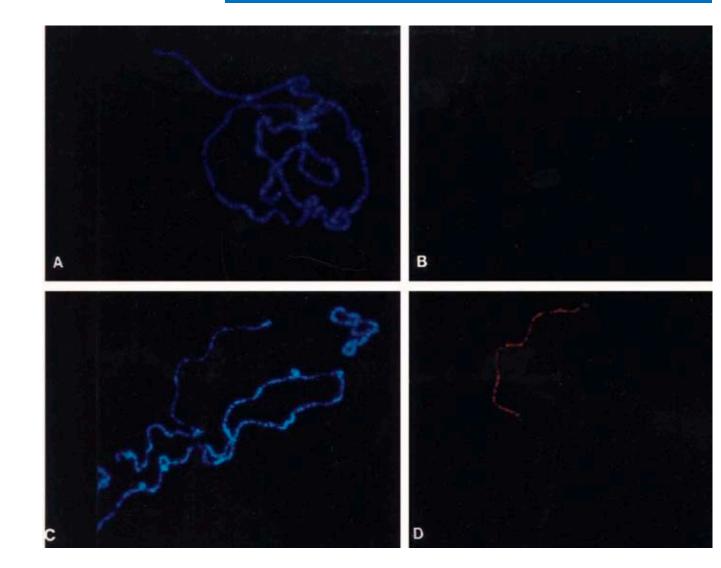




msl-2 and dosage compensation

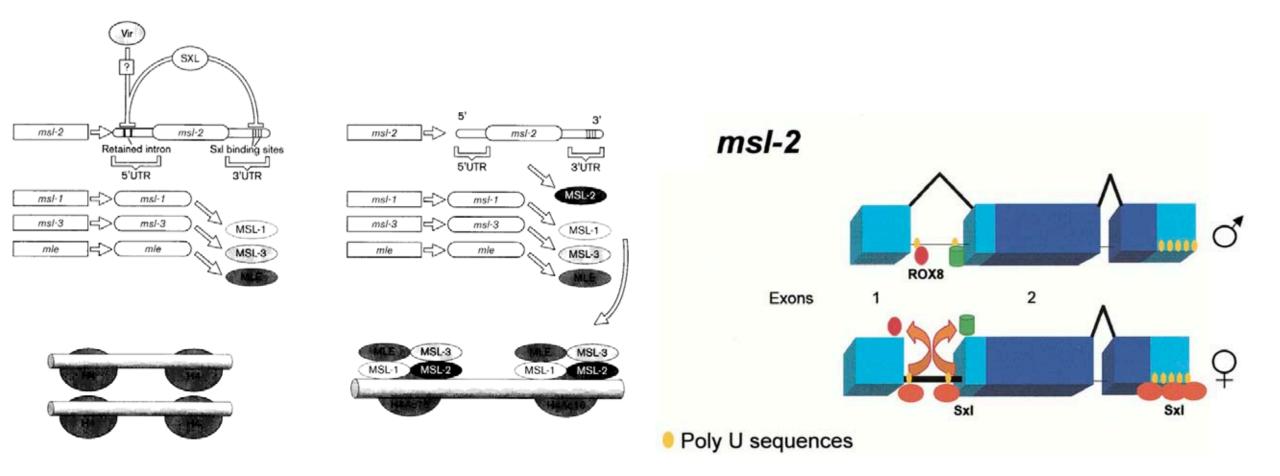


Northern analysis of 4-6 µg of poly(A)+ RNA prepared from: lane 1, Canton S females; lane 2, Canton S males; lane 3, *tudor* females; lane 4, *tudor* males; lane 5, *msl-2* females



Anti-MSL-2 and anti-MSL-1 staining of wild-type male and female chromosomes. (A) DNA stain of wild-type female. (B) Anti-MSL-2 stain of wild-type female.

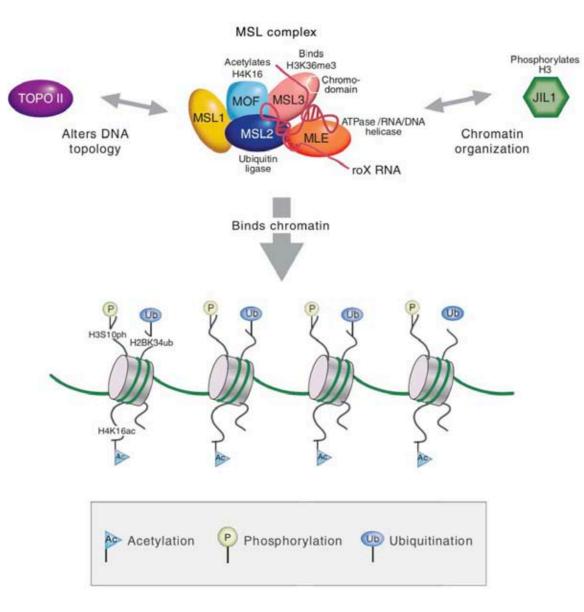
Dosage Compensation in Drosophila



Greg J Bashaw and Bruce S Baker. Curr Opin Genet Dev. 1996

John C. Lucchesi and Mitzi I. Kuroda . Cold Spring Harb Perspect Biol. 2015

Assembly of the chromatin-remodeling complex responsible for dosage compensation

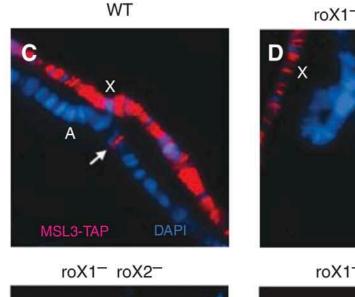


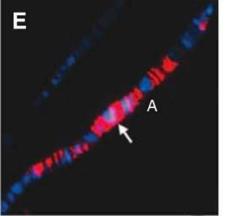
roX: X-specific noncoding RNAs (called 1 and roX2)

John C. Lucchesi and Mitzi I. Kuroda. Cold Spring Harb Perspect Biol. 2015

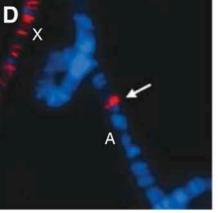
Why MSL complex can specific recognize the X chromosome?

1. The roX genes may reside on the X to target MSL complex assembly to this chromosome.





roX1⁻ roX2⁺



roX1⁻ roX2⁻

F

in all cases in which *roX* genes direct spreading in *cis* on autosomes, they also provide roX RNA in *trans* to cover the X chromosome (Meller and Rattner 2002). Therefore, it is clear the X chromosome has additional targeting signals beyond the two known *roX* genes.

Why MSL complex can specific recognize the X chromosome?

2. MSL1 and MSL2 are critical for the specific recognition of the X chromosome

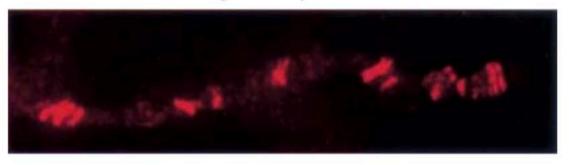
In the absence of either MSL1 or MSL2, none of the remaining MSL proteins or roX RNAs appears to retain specific recognition for the X chromosome.

However, in the absence of MLE, MSL3, or MOF, partial MSL complexes bind a subset of approximately 35 – 70 sites by cytological mapping, including the two roX genes

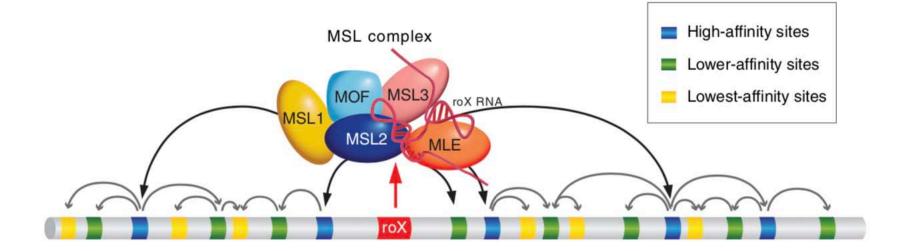


MSL-binding sites

High-affinity sites



Model for the targeting of the MSL complex to the X chromosome



John C. Lucchesi and Mitzi I. Kuroda. Cold Spring Harb Perspect Biol. 2015

References

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- Baker, B. S., M. Gorman, and I. Marin, 1994 Dosage compensation in Drosophila. Annu. Rev. Genet. 28: 491–521.
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- Lucchesi, J.C. and M.I. Kuroda, *Dosage Compensation inDrosophila*. Cold Spring Harbor Perspectives in Biology, 2015. 7(5).
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The Development of Sexually Dimorphic Structures and the Evolution of Sex

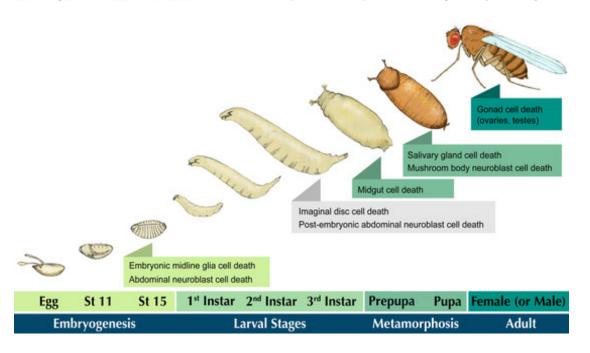
JSH

How does the sex determination pathway direct the development of sexually dimorphic morphologies?

How did sex differences evolve?

B. S. BAKER AND K. A. RIDGE

SEX determination in eukaryotes provides a striking example of the differential control of gene expression during development. Since sex determination affects the developmental fate of numerous organ primordia, information as to the nature of the genetic events involved in sex determination should contribute not only to our understanding of sex determination, but also to the elucidation of the mechanisms by which eukaryotes effect the expression of alternative developmental pathways.



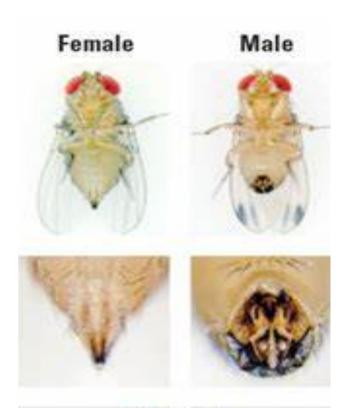
1.a molecular–genetic investigation of genital imaginal disc development

2.genome-wide approaches to find downstream targets of the transcription factors found at the end of the sex determination pathway, Dsx and Fru M

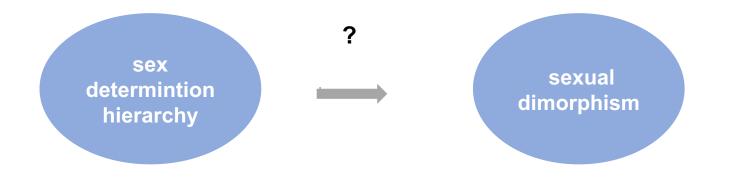
3.an examination of the evolutionary conservation of genes in the sex hierarchy

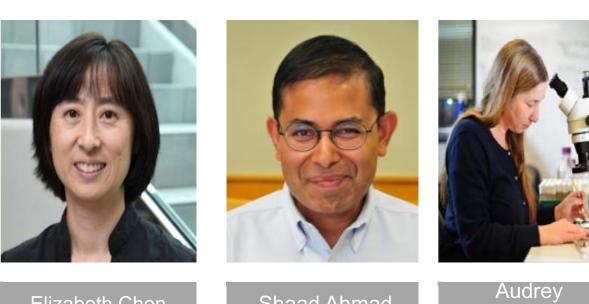
384

PART1 : sexually dimorphic structures



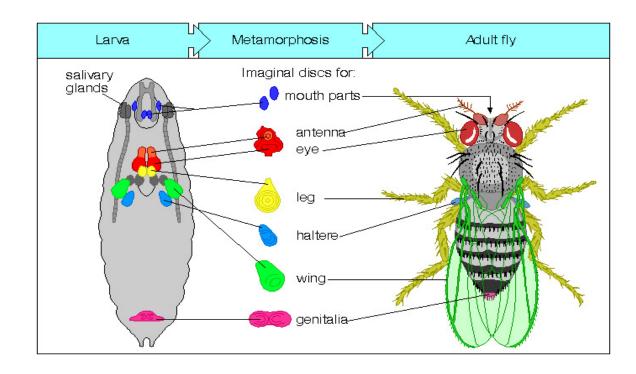
(genital imaginal disc development)





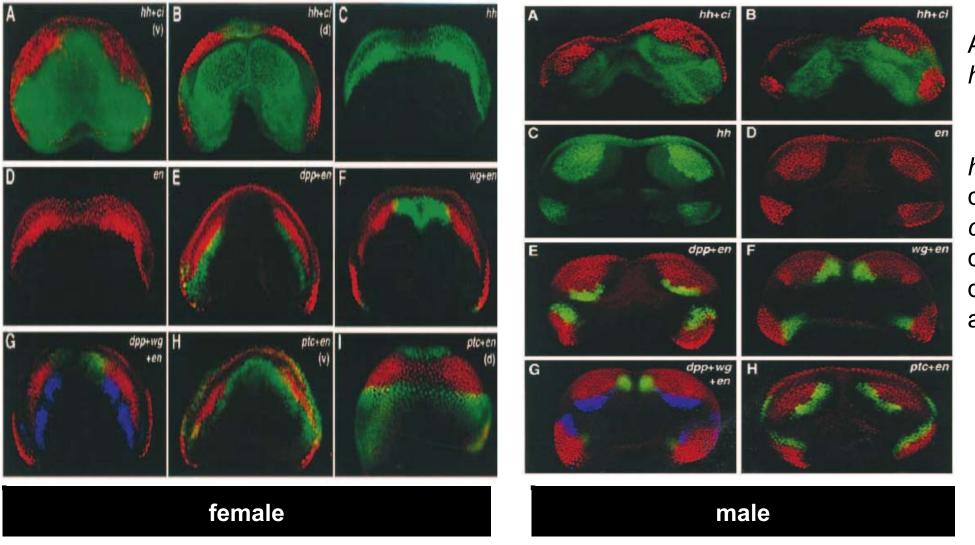
Shaad Ahmad

Christiansen



Elizabeth Chen

A new model for the A/P compartmental organization of the genital disc

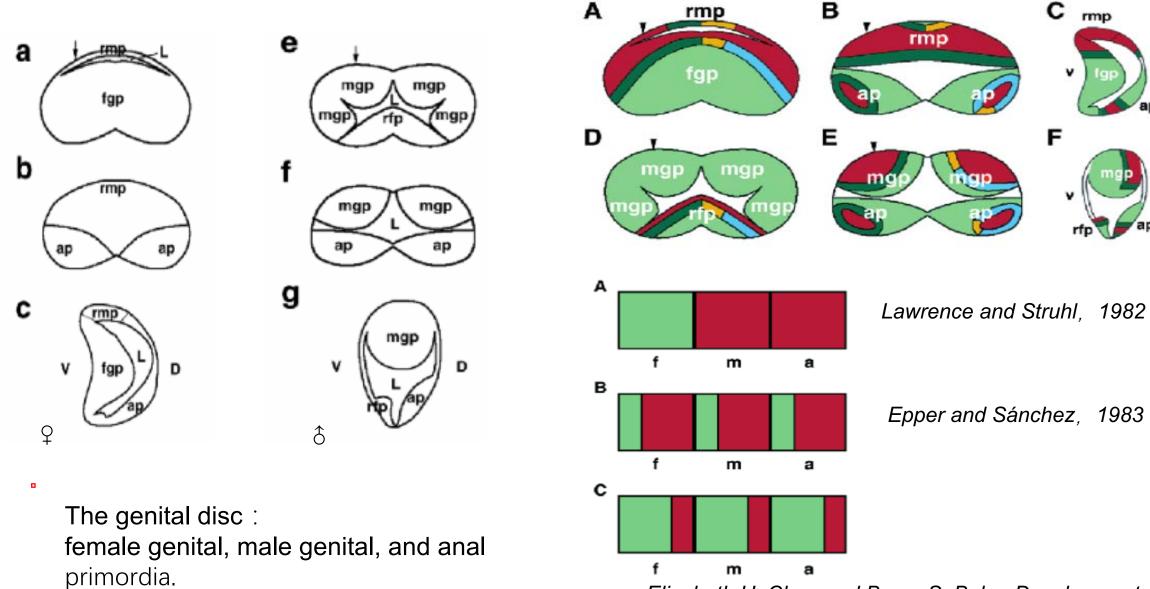


A/P patterning genes : hh,en,ci,dpp,wg,ptc

hh(en)— the posterior compartment *ci*— the anterior compartment dpp,wg,ptc—the border area

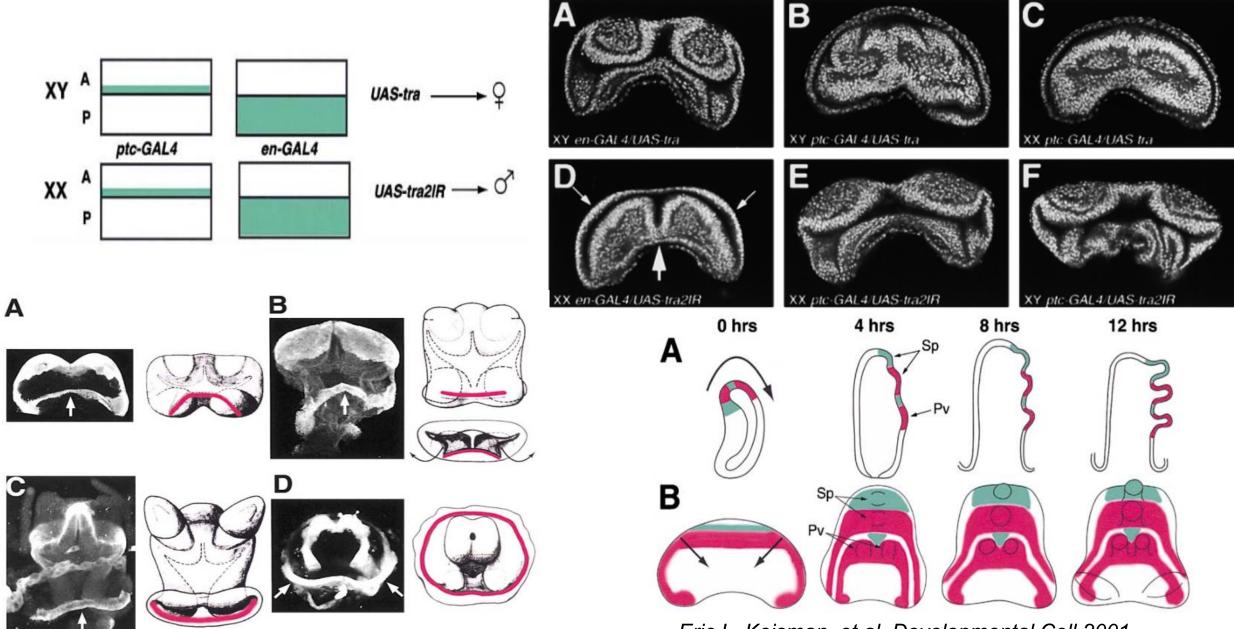
Elizabeth H. Chen and Bruce S. Baker.Development.1997

A new model for the A/P compartmental organization of the genital disc



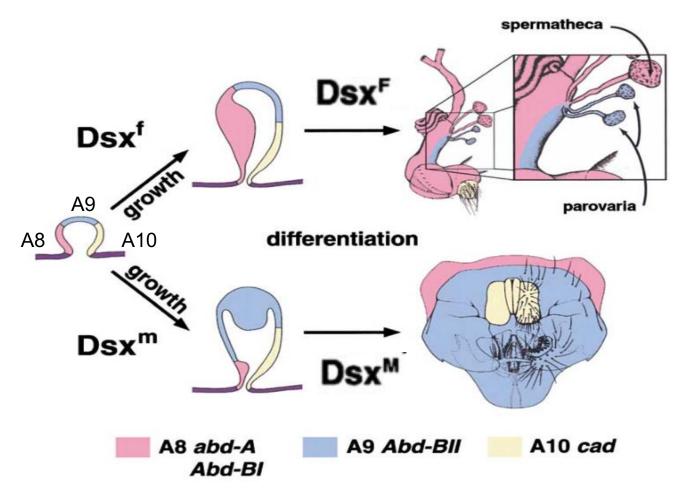
Elizabeth H. Chen and Bruce S. Baker.Development.1997

Sexual dimorphism in the genital disc is controlled from the A/P organizer



Eric L. Keisman, et al. Developmental Cell.2001

Dsx acts at multiple levels to instruct the sex-specific differentiation of the genital primordia



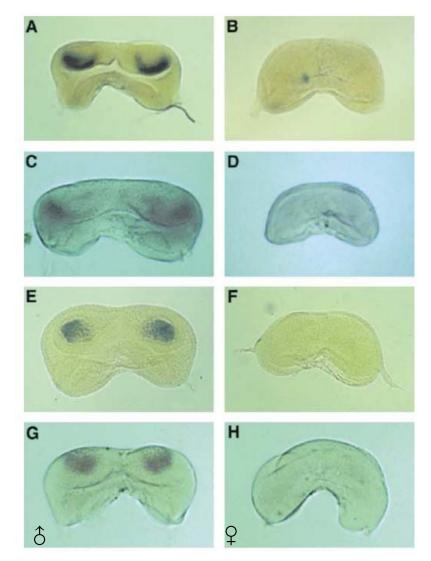
•Dsx acts in concert with the homeotic genes to direct the sex-specific fate of each primordium.

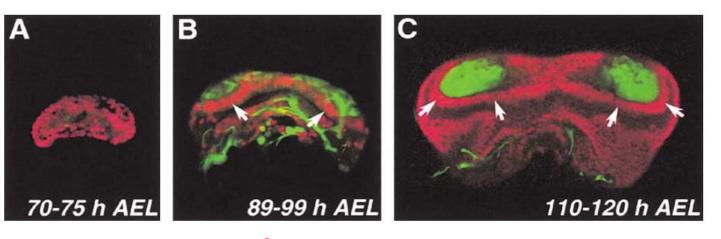
•Dsx controls growth by regulating the activity of the A/P organizer.

•Differentiation is controlled by dsx cell autonomously.

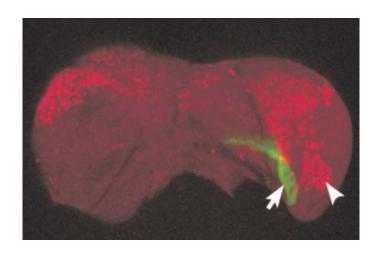
Eric L. Keisman, et al. Developmental Cell.2001

FGF is used to recruit mesodermal cells to the genital imaginal disc controlled by dsx





recruitment of btl-expressing cells



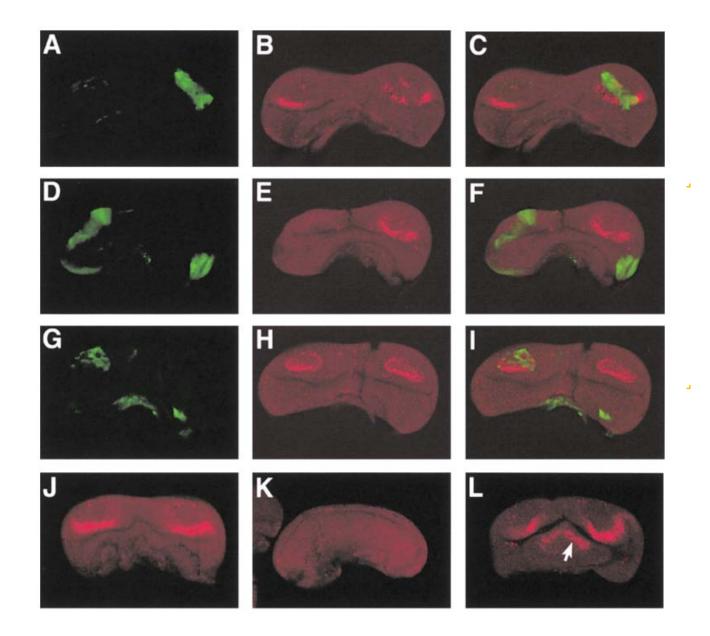
bnl was sufficient to recruit btlexpressing cells into locations

act5C>CD2>GAL4/Y; UAS-bnl/UAS-GFP; btl-lacZ/hs-FLP

Bnl and *btl* are only expressed in the A9-derived developing "male" primordium. *bnl*(FGF);*btl*(FGFR)

Shaad M. Ahmad and Bruce S. Baker.Cell.2002

bnl is a target of sex determination hierarchy, being repressed byDsx^F in females

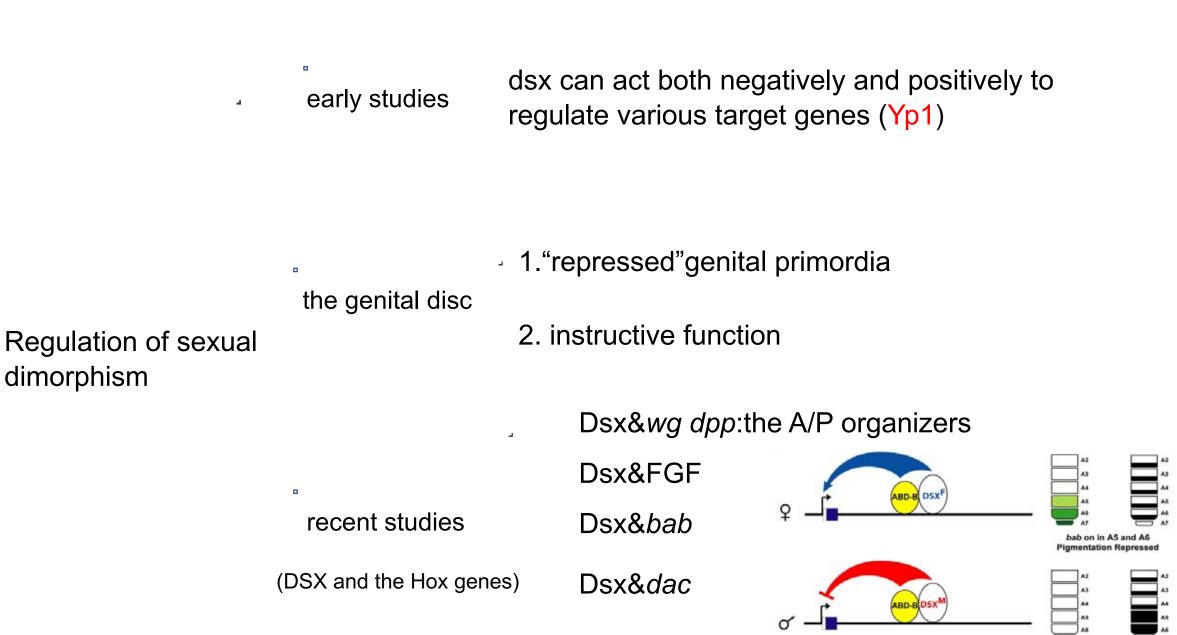


Bnl is a target of the sex determination hierarchy, downstream of tra.

"lobe" - "flatten"

Btl is not a direct target of the sex determination hierarchy

Shaad M. Ahmad and Bruce S. Baker.Cell.2002

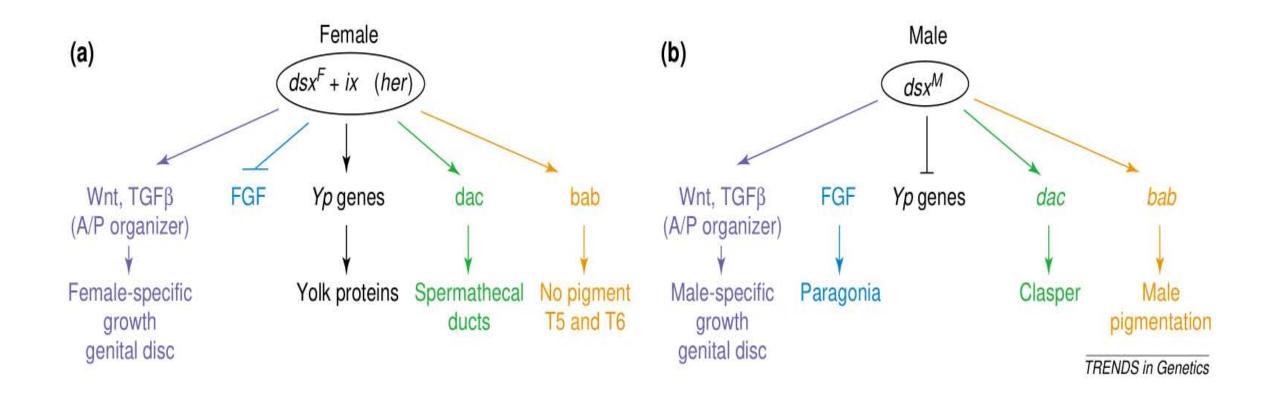


XY:*wg* represses *dac* and *dpp* activates it XX:*wg* activates *dac* and *dpp* represses it

bab off in A5 and A6 Pigmentation Expressed

Dimorphic Element

The sex hierarchy modulates the activities of signaling molecules and transcription factors to direct various sex-specific aspects of growth and differentiation.

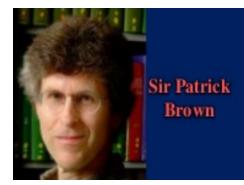


Audrey E.Christiansen, et al. Trends in genetics.2002

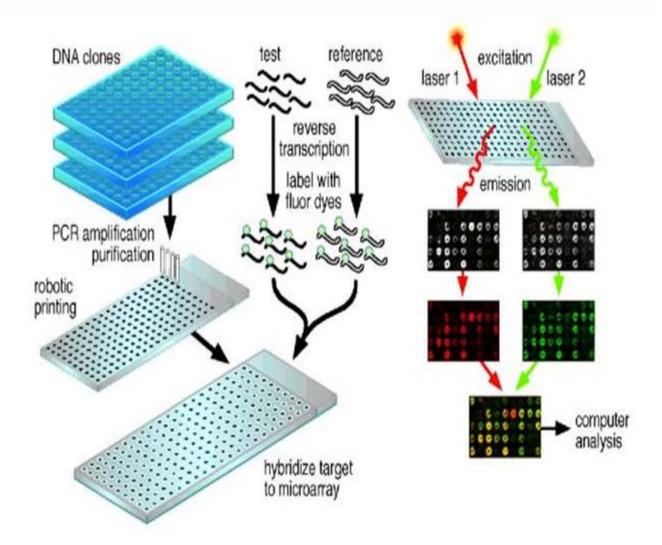
PART2 : genome-wide approaches to find downstream targets

DNA Microarray

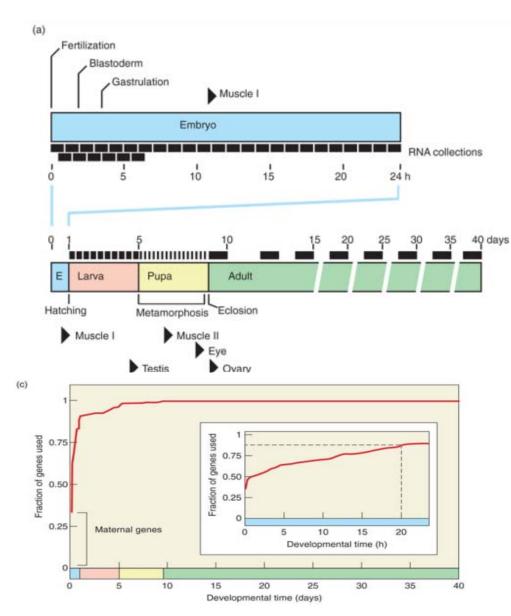


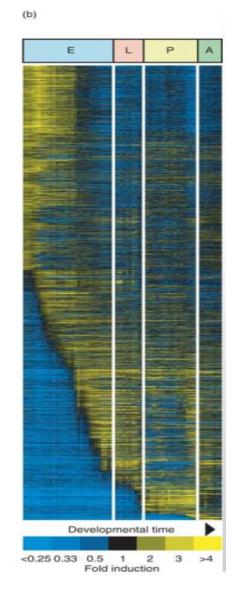


•"Quantitative Monitoring of Gene Expression Patterns with a complementary DNA microarray" reported by Patrick Brown, Mark Schena and colleagues in Science (1995).



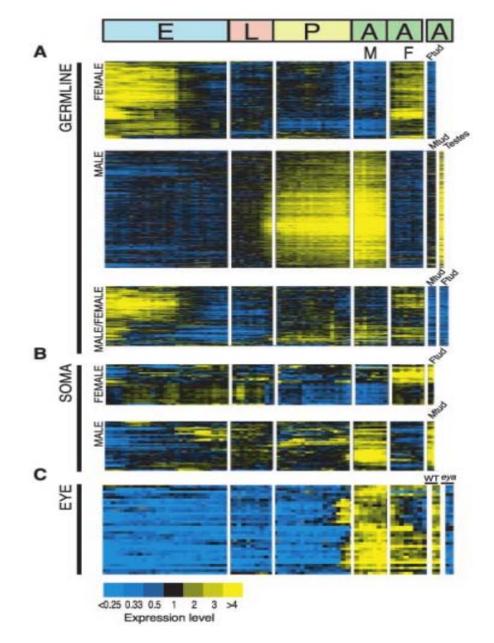
Patterns of gene expression through the Drosophila life cycle





Michelle N. Arbeitman, et al. Science. 2002

Sex difference in the transcriptional programs in Drosophila melanogaster

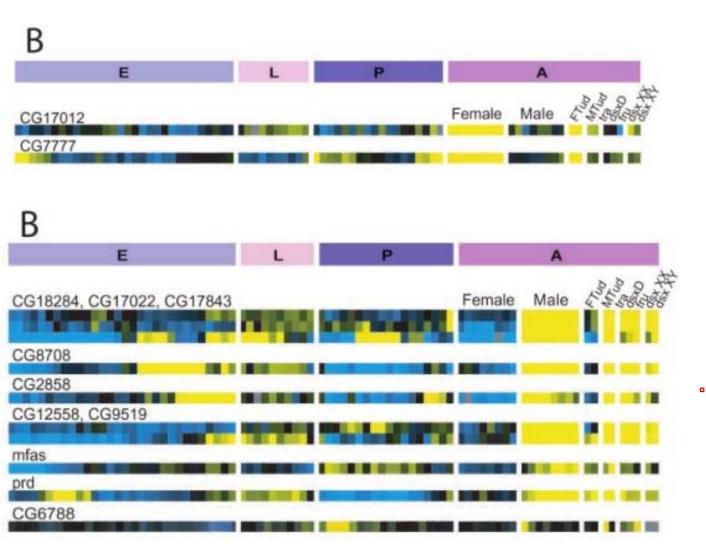


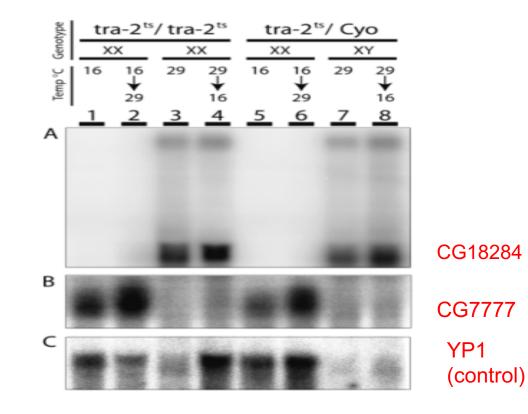
>111 genes were expressed in both male and female germ lines.

>31 genes had significantly higher expression in the soma of adult females compared with 37 genes in males.

Michelle N. Arbeitman, et al. Science. 2002

Temporal regulation of sex-differential adult gene expression by the hierarchy





Sex-differential expression of all 11 genes is the consequence of the developmental action of the sex hierarchy and is independent of the hierarchy during adult stages.

Michelle N. Arbeitman, et al. Development. 2007

Modes of dsx regulation of sex-differentially expressed genes

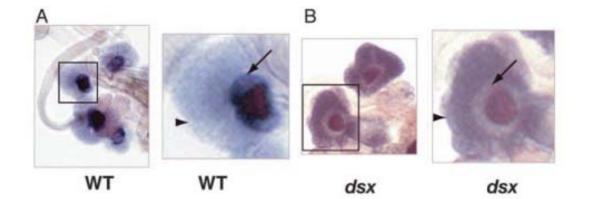


Table 3.	Modes of regulation by	dsx
----------	------------------------	-----

Genotype										
		DSX phenotype						Suggested mode of		
	Expression localized to	XY wt	XX wt DSX ^F	XY dsx _	XX dsx	XX dsx ^D DSX ^M	XX tra DSX ^M	regulation		
Gene		DSX ^M						DSX ^M	DSX ^F	
YpI^*	Female fat body							-	+	
CG17012	Spermathecae	-1.29	4.64	-0.95	-0.15	-1.34	-1.89	0	+	
CG17843	Male AG	1.31	-3.16	1.08	0.09	0.80	0.55	0	-	
CG17022	Male AG	3.29	-2.69	3.83	2.28	3.86	3.75	0	-	
CG18284	Male AG	2.92	-1.60	2.46	0.43	3.33	2.39	0	-	
prd	Male AG	0.40	-0.30	0.62	0.61	1.31	0.94	0	255	
CG2858	EB	1.10	-2.07	-0.68	0.35	1.24	1.01	+	-	
CG8708	AED	1.15	-1.07	0.50	0.05	1.79	2.00	+	(m)	

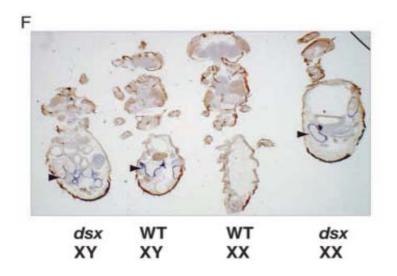
Values given in table are log-transformed microarray ratios.

*Mode of regulation previously established (reviewed by Christiansen et al., 2002).

AG, accessory gland; EB, ejaculatory bulb; AED, anterior ejaculatory duct; wt, wild type; 0, no effect; +, positive effect; -, negative effect.

DSX^F positively regulates sexspecific expression of CG17012.

Male-specific expression of *prd* is a negative control by DSX^F to prevent the formation of male accessory glands.



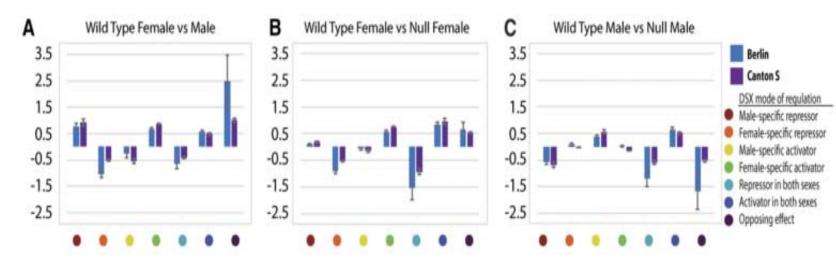
Michelle N. Arbeitman, et al. Development. 2007

a new consensus DSX-binding site



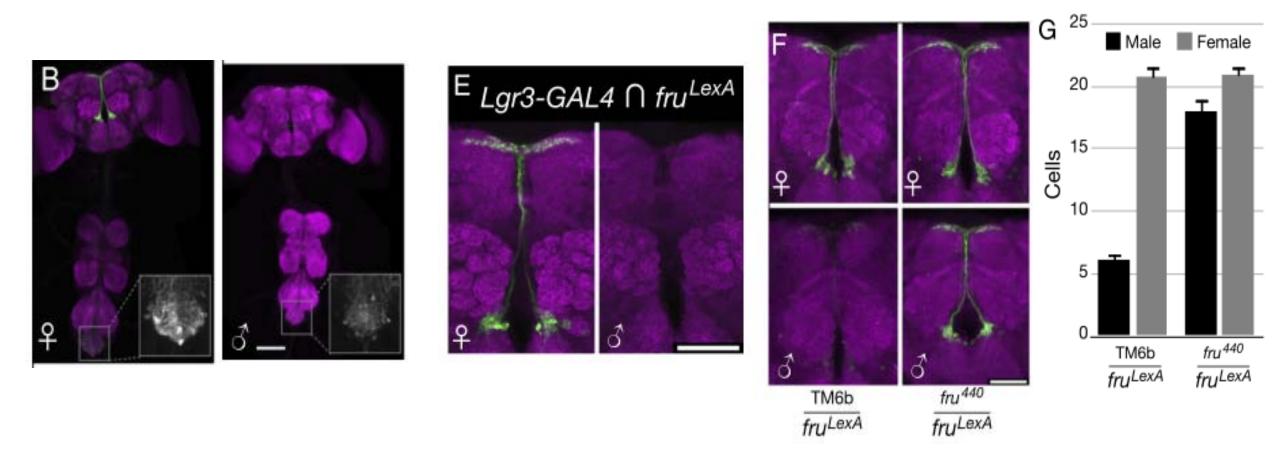
Shengzhan D. Luo, et al. Development. 2011

The effect of different DSX modes on regulation of expression



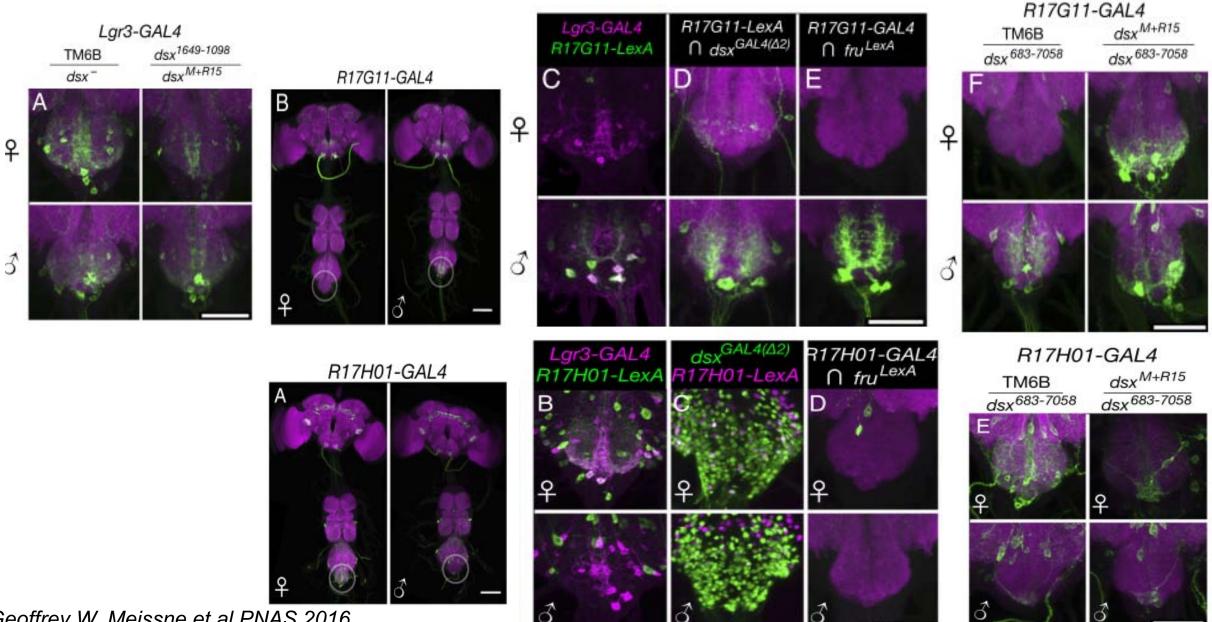
M. N. Arbeitman, et al. Genetics. 2016

Drosophila Lgr3 is regulated by Fru and Dsx in separate populations of neurons



Fru^M inhibits expression of Lgr3 in the male median bundle.

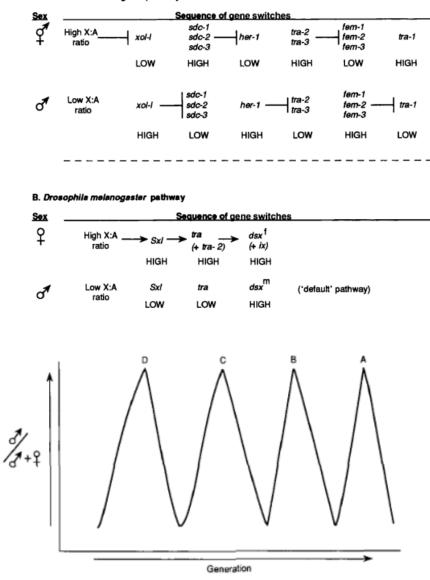
Dsx^F activates and inhibits Lgr3 expression in different abdominal ganglion neurons

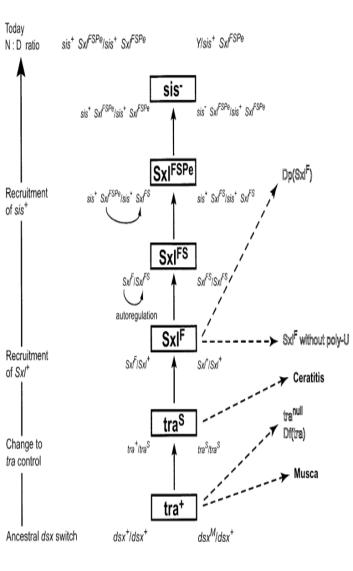


Geoffrey W. Meissne, et al. PNAS. 2016

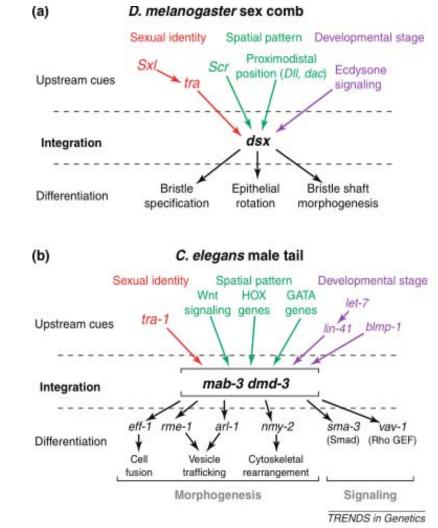
PART3: the evolution of sex differences

A. Caenorhabditis elegans pathway





Dmrt gene



A.S. Wilkins.Bioessays.1994

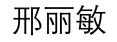
A.Pomiankowski, et al. Genetics. 2004

A.Kopp. Trends in Genetics.2012

References

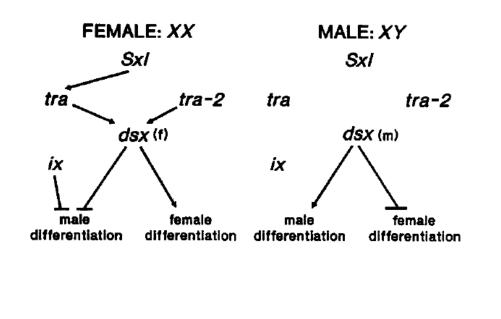
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- Keisman E.L., et al., 2001 The Sex Determination Gene doublesex Regulates the A/P Organizer to Direct Sex-Specific Patterns of Growth in the Drosophila Genital Imaginal Disc. Developmental Cell, 215–225
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- and its regulation.Development 131.2007-2021
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- Meissner G.M et.al., 2016 Sex-specific regulation of Lgr3 in Drosophila neurons. PNAS.1256-1265
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- Wilkin A.S. et al.1994 Moving up the hierarchy:a hypothesis on the evolution of a genetic sex determination pathway.Bioessay.17(01)
- Kopp A.2012 Dmrt genes in the development and evolution of sexual dimorphism. Trends in Genetics.28(4)

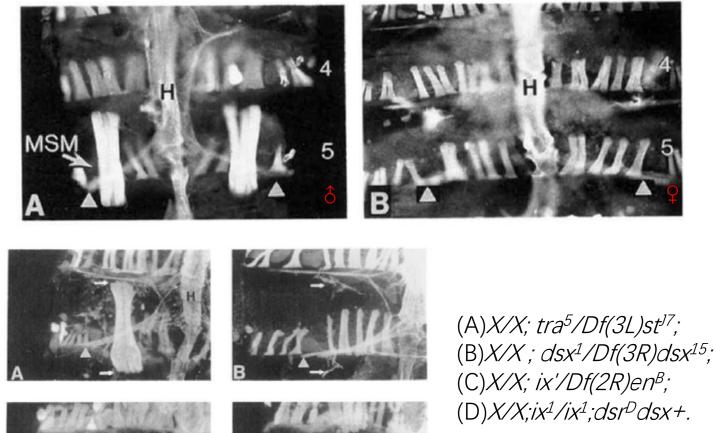
Sex Behavior Meets the Sex Determination Regulatory Hierarchy: The Genetic Control of Sexual Behavior

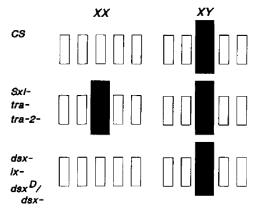


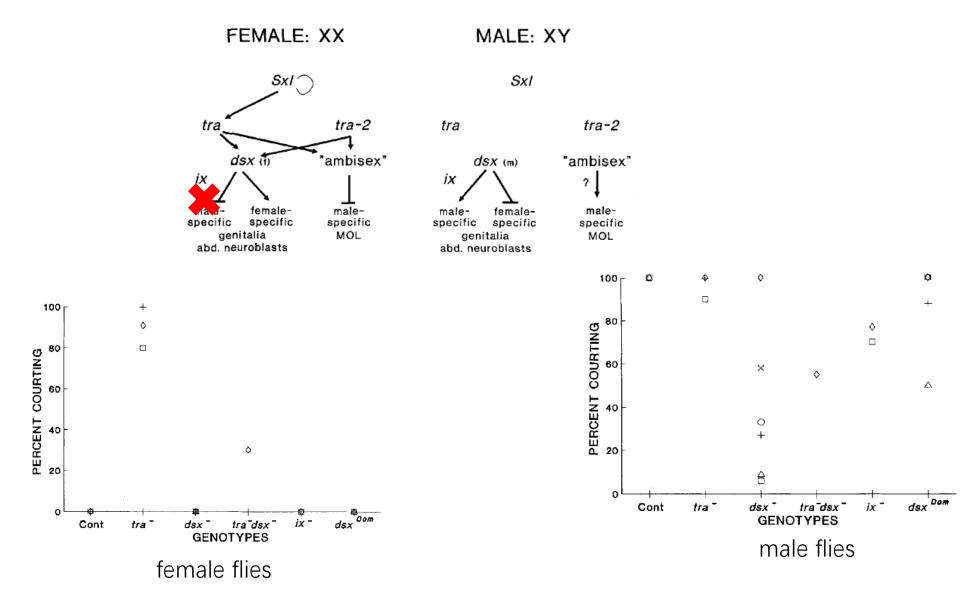
- Bruce' s entry into the genetics of sexual behavior came from the discovery of a new branch in the sex determination *fruitless* gene
- The expression pattern of *fruitless* and *doublesex* and their regulation to innate behaviors
- A deeper exploration of the neuronal mechanisms underlying sexual behavior

The Muscle of Lawrence (MOL) depended on the function of *sxl, tra* and *tra-2*, but not *doublesex* or *intersex*



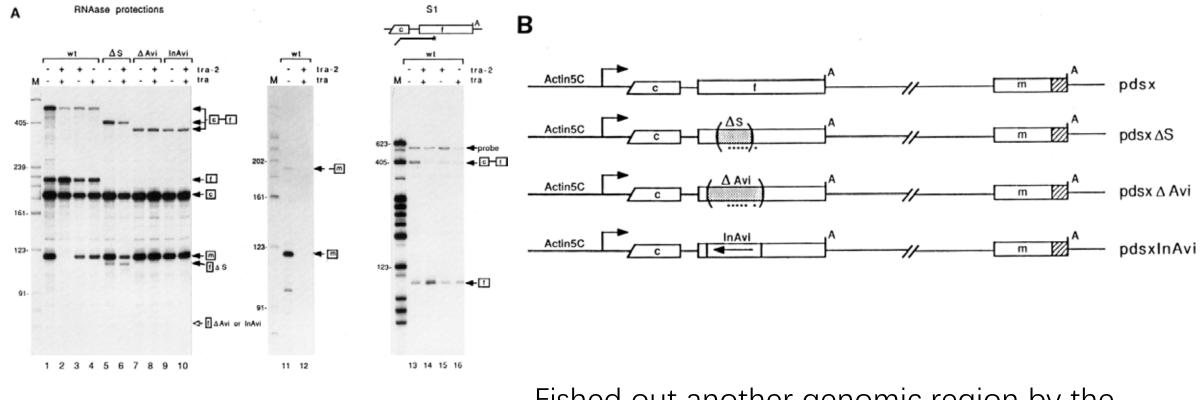






These findings suggested the existence of a previously unrecognized branch in the sex-determination hierarchy Taylor, et al., Genes Dev. 1992

A 13-nt sequence, repeated six times in *dsx* in a noncoding region, was necessary and sufficient to direct sex-specific splicing



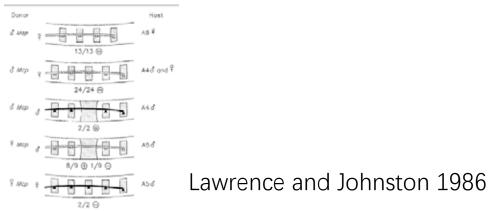
dsx, tra, tra-2, and Sxl in S-L2 cells

Fished out another genomic region by the 13-nt repeat sequences, which is the *fru* gene

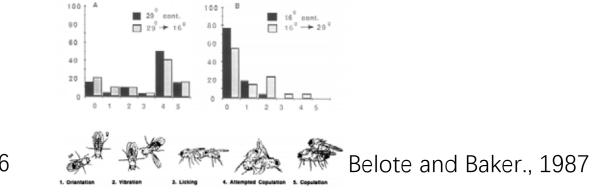
Ryner, L. C., and B. S. Baker, Genes Dev. 1991

How does *fru* become a good candidate gene residing at the top of the new branch in the sex determination regulatory hierarchy?

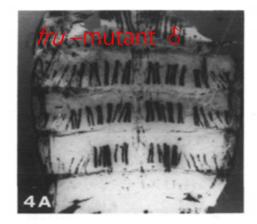
evelopment of the MOL was dependent on the sex of the innervating neurons



tra-2 is required in adult females to block male-specific courtship behaviors

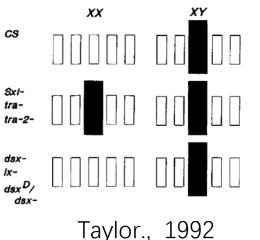


fru was known to be involved in the development of MOL



Gailey., et al. 1991

the Muscle of Lawrence (MOL) is sex-specific development and depended on the function of *tra* and *tra-2*, but not *dsx*



Control of Male Sexual Behavior and Sexual Orientation in Drosophila by the *fruitless* Gene

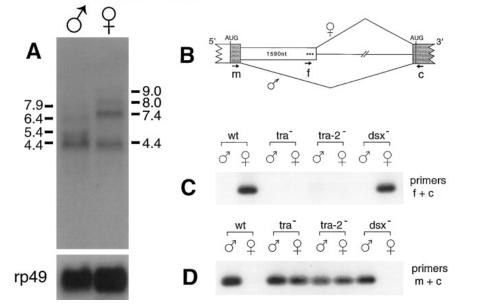
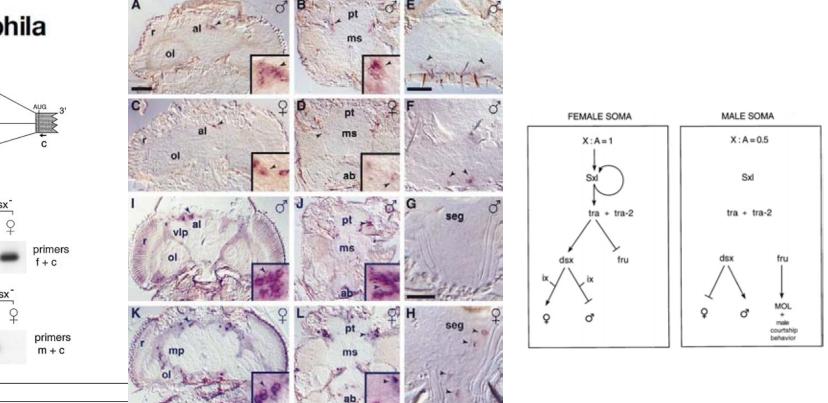


Table 1. Courtship Behavior of fru Mutants

	male + male			male + female		120	
Genotype	All courtship Cl* (%)	Wing ext. (%)	N	All courtship Cl* (%)	Wing ext. (%)	N	
wild-type	4 ± 1	0 ± 0	10	84 ± 3	46 ± 7	7	
fru ¹ /fru ¹	51 ± 4	21 ± 3	25	61 ± 5	35 ± 4	21	
fru³/fru³	32 ± 5	1 ± 0	31	15 ± 5	0 ± 0	20	
fru4/fru4	41 ± 6	2 ± 1	25	29 ± 7	2 ± 1	20	
fru³/fru⁴	42 ± 7	1 ± 0	16	22 ± 8	1 ± 1	13	
fru ³ /fru ^{w12}	1 ± 0	0 ± 0	9	1 ± 1	0 ± 0	10	
fru ⁴ /fru ^{w12}	8 ± 5	1 ± 1	9	7 ± 5	0 ± 0	12	
fru ^{w12} /Cha ^{M5}	2 ± 1	0 ± 0	14	0 ± 0	0 ± 0	6	
fru ^{w27} /Cha ^{M5}	2 ± 2	0 ± 0	15	0 ± 0	0 ± 0	7	

s sex-specific transcripts were only expressed in the adult fru nervous system and in a very distinct pattern



 Sexual orientation and courtship behavior in Drosophila are regulated by fruitless (fru), the first gene in branch of the sex-determination hierarchy.

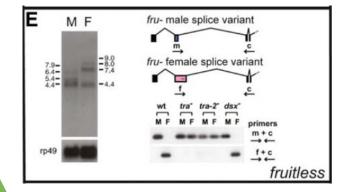
Ryner, et al. Cell. 1996

This discovery was the critical finding that led Bruce' s laboratory into the study of behavioral genetics

Fruitless Consortium



fru's Sexspecific transcripts the role of *fru* alleles ; fru in MOL the clones developmen spanning the *fru* genomic interval



Bruce Baker VS Manichean

- The combined work of the consortium had built a very strong case that a single gene, fru, had highly specific control of *Drosophila* sexual orientation and sex behavior (Ryner et al. 1996).
- Whether or not behavioral repertoires could in fact be genetically specified, or were predominantly shaped by environmental factors

Cell, Vol. 105, 13-24, April 6, 2001, Copyright @2001 by Cell Press

Are Complex Behaviors Specified by Dedicated Regulatory Genes? Reasoning from *Drosophila*

Review

Bruce S. Baker,*§ Barbara J. Taylor,† and Jeffrey C. Hall‡ (Ridley, 1995). Such a definition includes species-specific behaviors studied extensively by ethologists (e.g., court-

Fruitless Consortium - sex-specific transcripts of the fru

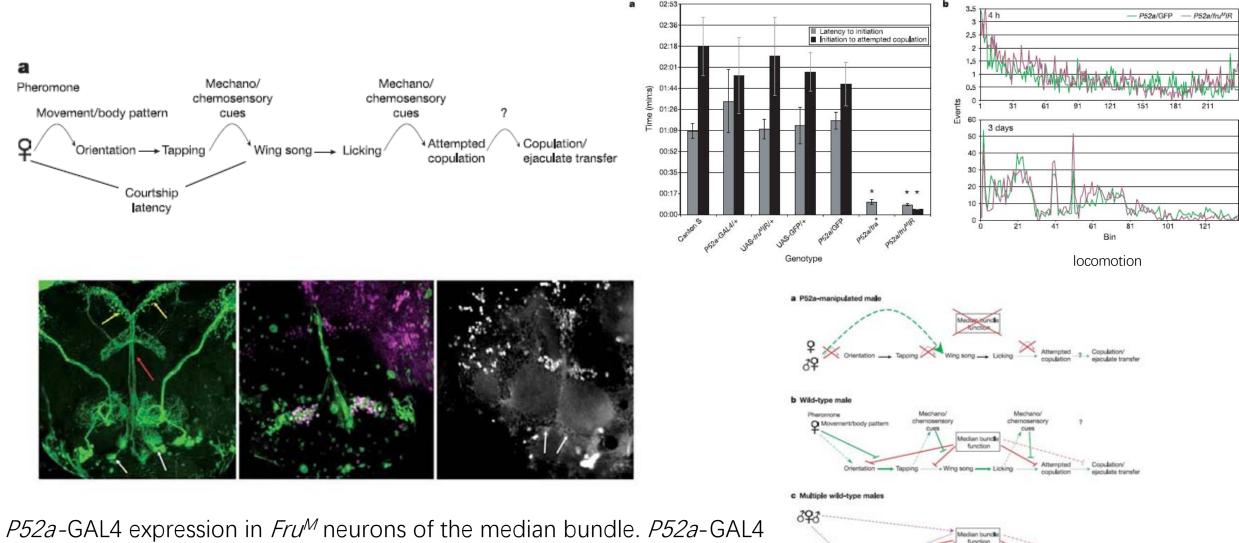
Molecular Genetic Dissection of the Sex-Specific and Vital Functions of the Summary of transcript classes expressed by fru alleles Drosophila melanogaster Sex Determination Gene fruitless Transcript class P2 **P**3 fru allele P1 P4 Anuranian Anand.*.† Adriana Villella.[‡] Lisa C. Ryner,* Troy Carlo,[‡] Stephen F. Goodwin,^{1,8} fru¹a ++Ho A Morales,[‡] Jeffrey C. Hall,[‡] Bruce S. Baker* fru² a (+)(+)(+)fru³a (+)(+)I. Taylor** fru⁴a (+)(+)Cha^{M5b} $(+)^{d}$ fru^{440b} fru^{w27 b,c} fru^{w12b} fru^{w9 c} BX7 148.5-1 fru^{w13} fru^{sat151} fru^{w24 b} TABLE 5 fru^{w9}, fru^{w12}, fru^{w27} Courtship behavior of fru mutant males в WEI CIWEI CI $(m \rightarrow m)$ $(m \rightarrow f)$ Genotypes $(m \rightarrow f)$ ChI 30 40 $(m \rightarrow m)$ DK3RMu⁴⁻⁴⁰ Di(3R)truSaf15 DI/3RICha^A TABLE 7 DK3R)P14 DI(3R)Inu^{w24} Courtship song summary for fru mutants P1 N wing N IPI CPP P2 Genotype Frequency Width extension ₩¥ P3 P4 TABLE 8

fru mutations disrupt the muscle of Lawrence development

 the consortium showed that *fru* encodes transcription factors whose sex-specific transcripts are generated by alternative splicing controlled by the sex hierarchy proteins Tra and Tra-2. This splicing was shown to be necessary for male-specific courtship behavior. rtu s viu

TABLE 2

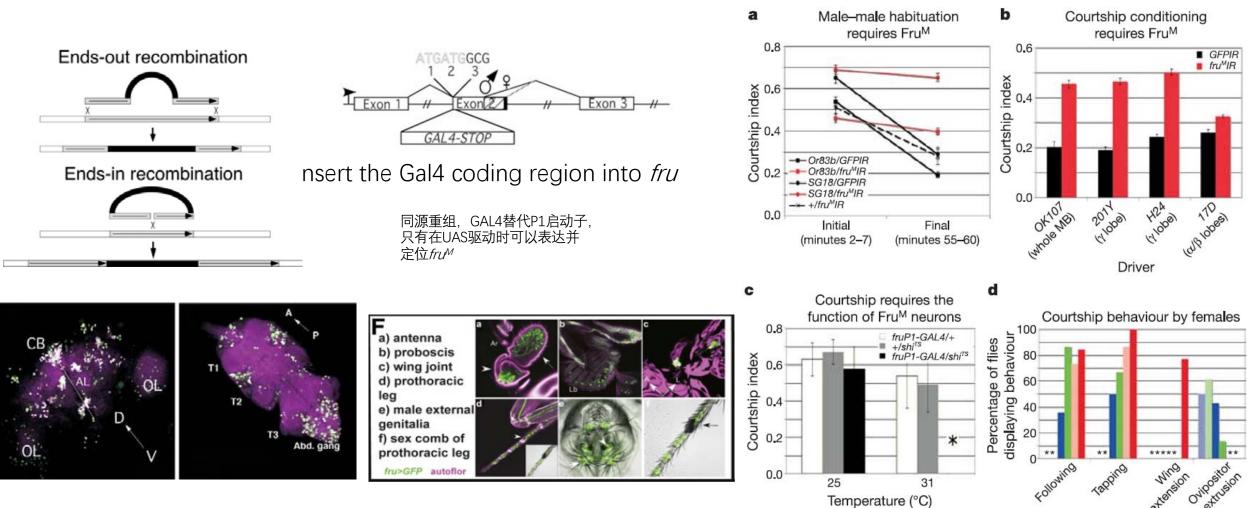
A small group of *fru^M* in the median bundle neurons appropriately trigger the sequential execution of the courtship ritual in *Drosophila*



directs inhibition of *Fru^M* expression by a UAS-*fru^{MIR}* transgene.

Manoli, D. S. and B. S. Baker., Nature. 2004

Fru^M proteins specify the neural substrates of male courtship.

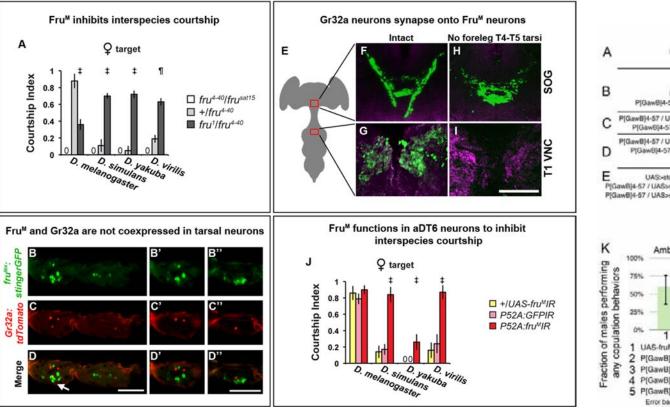


fru^M is important for the detection of ethologically relevant sensory information and for transmission of this information to the central nervous system. _{Gong and Goloc., 2003}

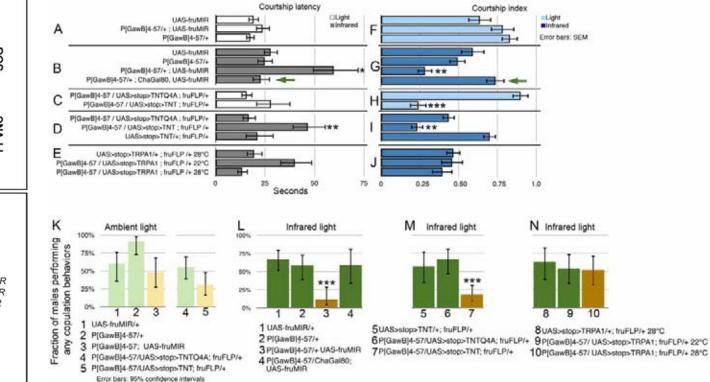
Gong and Goloc., 2003 Manoli et al., Nature., 2005

Behaviour

Gr32a and Fru^M inhibit interspecies courtship by Synaptic connections



A small subset of *fruitless* subesophageal neurons modulate early courtship in *Drosophila*

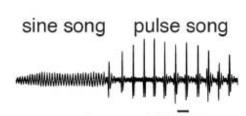


Tran, D. H., et al. PLoS One. 2014

Neural Pathways for the Detection and Discrimination of Conspecific Song in *D. melanogaster*

Alexander G. Vaughan,¹ Chuan Zhou,¹ Devanand S. Manoli,² and Bruce S. Baker^{1,*} ¹Janelia Farm Research Campus, HHMI, Ashburn, VA 20147, USA ²Department of Psychology, UCSF, San Francisco, CA 94122, USA

Α



Article

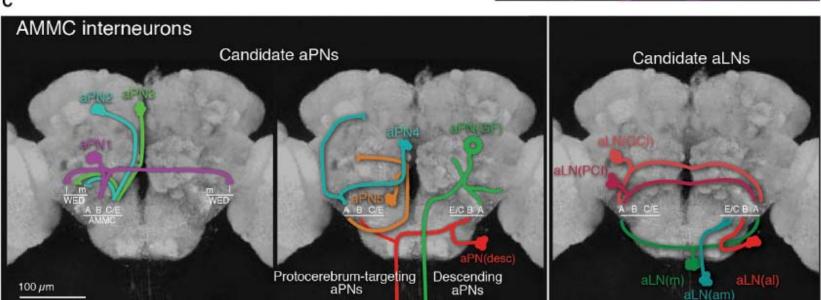
Inter-pulse Interval (IPI): 35ms

This song is perceived via mechanosensory neurons in the antennal Johnston's organ

в

A B C/E | C/E B

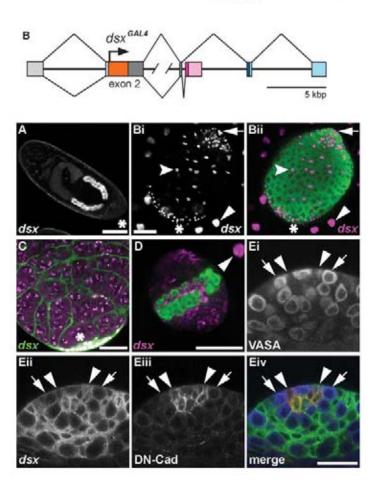
two classes of interneurons are necessary for song responses—the aPN1 and GABAergic interneuron aLN(al).



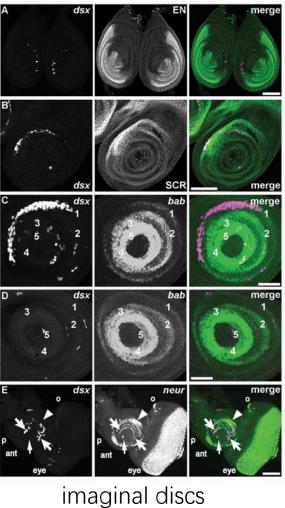
Sex and the Single Cell. II. There Is a Time and Place for Sex

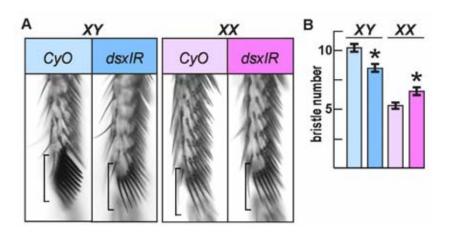
Carmen C. Robinett^{1^a}, Alexander G. Vaughan^{1^a}, Jon-Michael Knapp^{1,2^a}, Bruce S. Baker^{1,2^a}

1 Biology Department, Stanford University, Stanford, California, United States of America, 2 Neuroscience Program, Stanford University, Stanford, California, United States of America



the developing gonads in stage 13–17 embryos

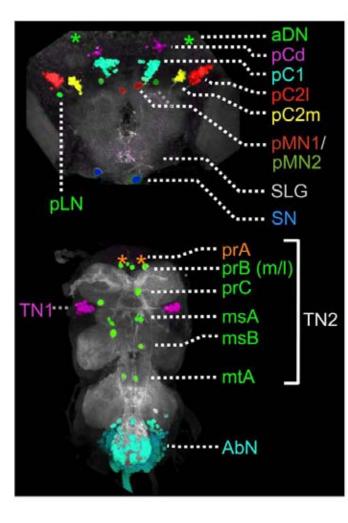


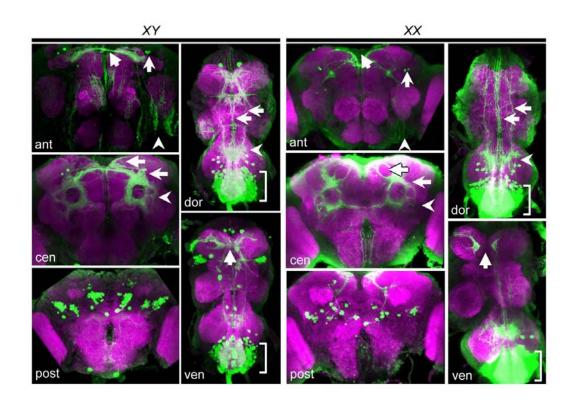


Sex and the Single Cell. II. There Is a Time and Place for Sex

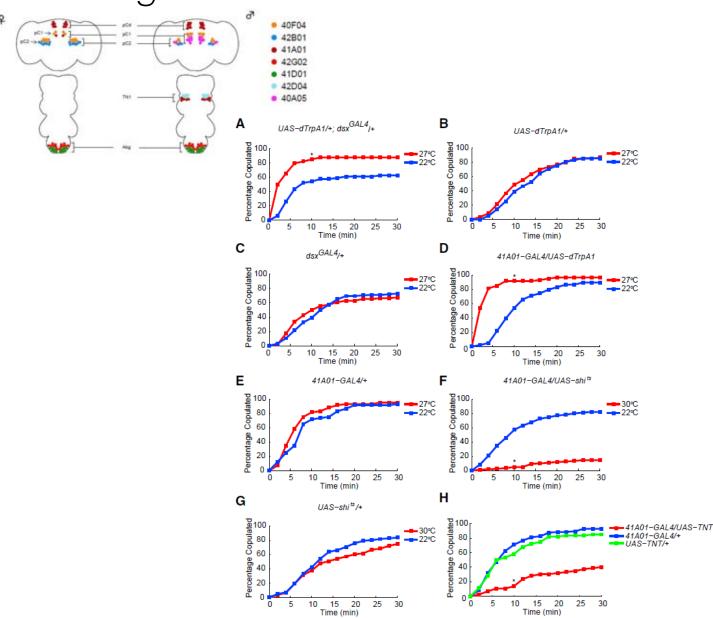
Carmen C. Robinett^{1^a}, Alexander G. Vaughan^{1^a}, Jon-Michael Knapp^{1,2^a}, Bruce S. Baker^{1,2^a}

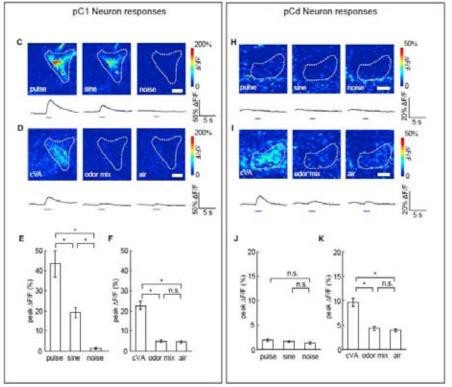
1 Biology Department, Stanford University, Stanford, California, United States of America, 2 Neuroscience Program, Stanford University, Stanford, California, United States of America

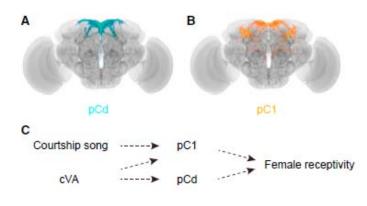




dsx-expressing neurons mediate virgin female receptivity to courting males

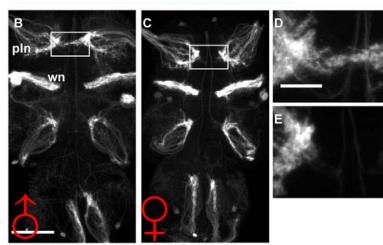




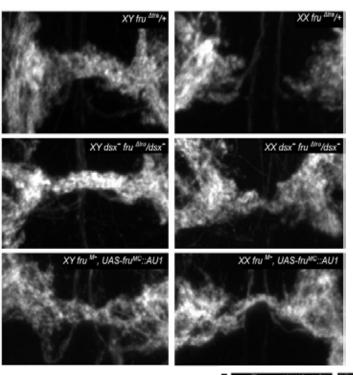


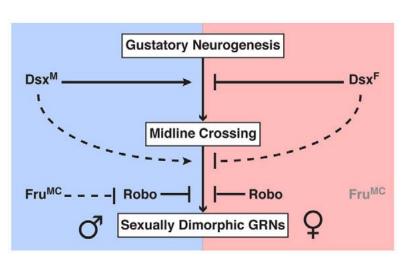
Zhou et al., Neuron., 2014

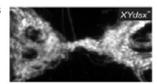
The *dsx* and *fru* branches of the sex hierarchy cooperate in controlling anatomical sex differences in neural circuitry

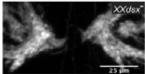


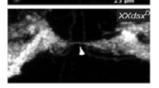
poxn-Gal4 (expressed in GRNs) -driven UAS-mCD8::GFP

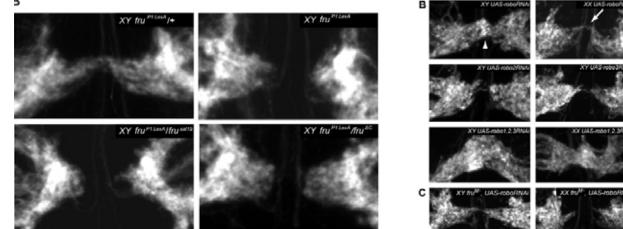


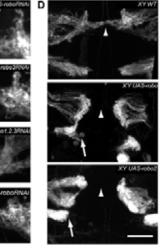








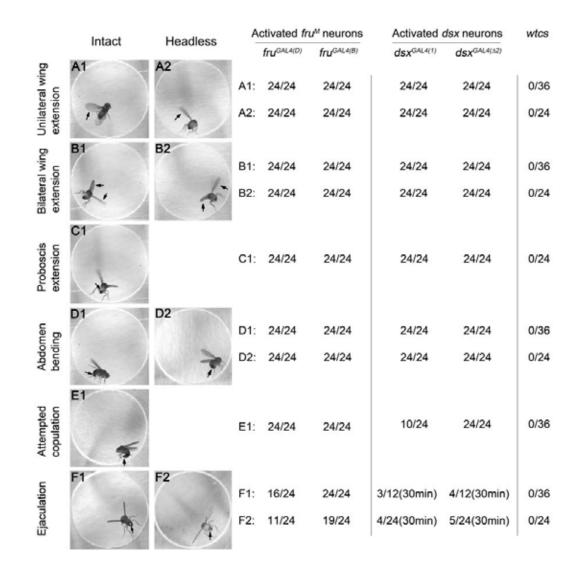


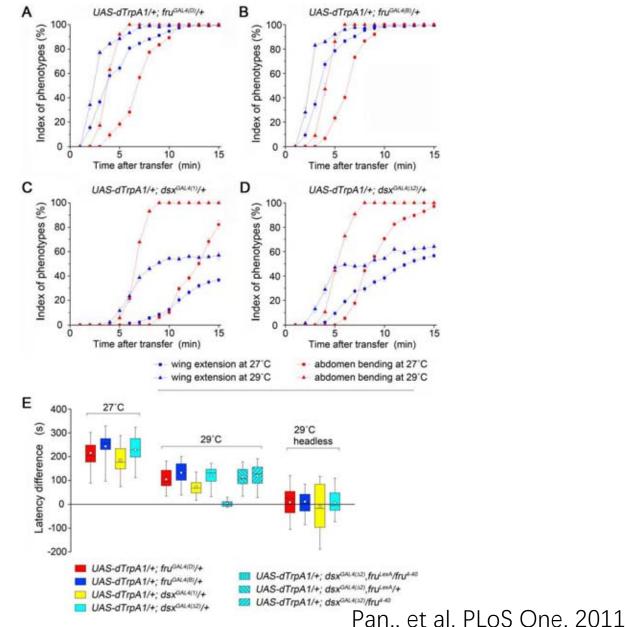


Mellert, et al. Development. 2010

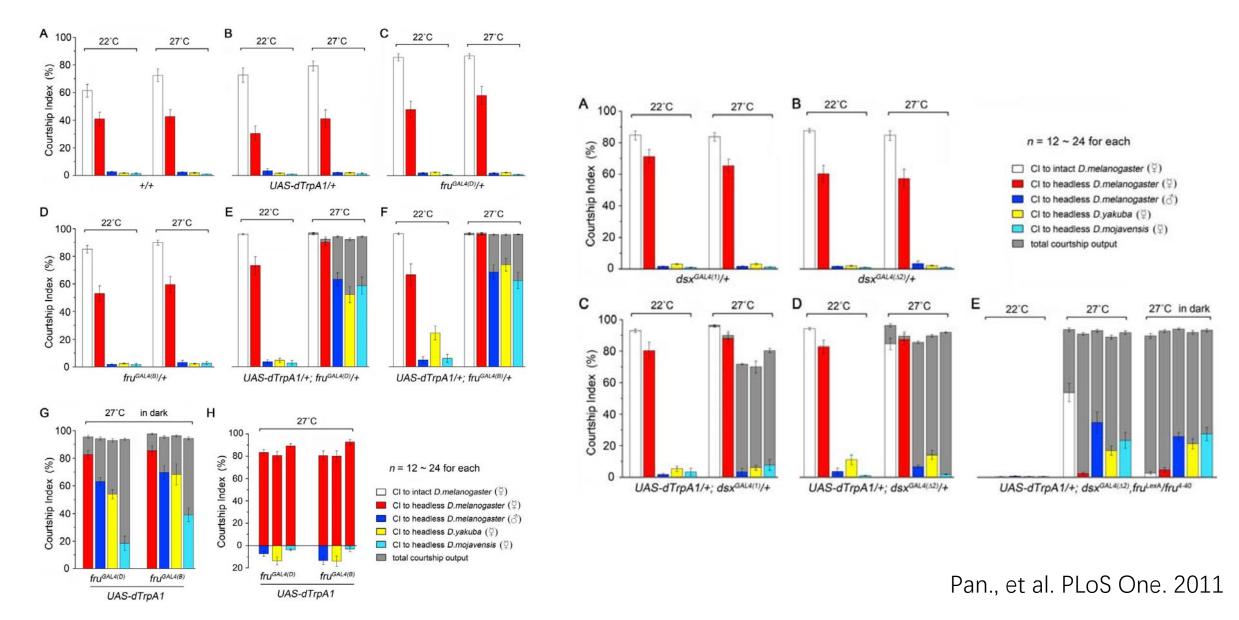
males lacking either (1) all Fru^M proteins or (2) Fru^{MC} proteins only do not form contralateral gustatory receptor neuron (GRN) projections

Activation of either all *fru^M* or all *dsx* neurons in solitary males can induce almost all steps of courtship





fru^M neurons involve in courtship promotion and recognition and the *fru^M*-independent courtship pathway is primarily vision dependent



22 °C 27 °C В Motion input and activation of P1 neurons are individually necessary, 100 and jointly sufficient to elicit early courtship behaviors 80 targets: (%) two freely-moving females Courtship Index 60 40 Female GAL4 Male LexA Male GAL4 в C 20 extension of left wing 60 4 cm extension of right wing layer 1: one tester male 3 mm of indicated events layer 2: two females targets: 27 °C С 22 °C two freely-moving females 100 40 plastic transparent barrier 80 targets: (%) two immobile females wres Index 60 AS-dTrpA1/+ 20 UAS-dTrpA1/+; fruGALADD/+ Number tship 40 AS-dTrpA1/+: fruGALA(B)/+ AS-dTrpA1/+; dsxGAL#(D/+ 0 Court ð UAS-dTrpA1/+: dsxGAL4AD/+ 20 total courtship output zero В G 60 B € 60 A 🖗 60 60 ourtship Index (%) targets: two targets: two targets: two constantly events freely-moving females immobile females æ rotating rubber bands 40 40 40 dių 20 G 20 of indicated 40 20 õ õ 30°C 22°C 22'C 30'C С Dŵ 60 two freely Targets: targets: 20 two constantly Number moving females rotating females 40 transparent barrier LexAop2-FlpL/+; UAS>stop>dTrpA1/dsxGA44(32) <u>e</u> 20 tester male LexAop2-FlpL/R71G01-LexA; UAS>stop>dTrpA1/+ 8 targets on a rotating platform ð σ LexAop2-FlpL/R71G01-LexA; 30°C left eye painted: + 22°C UAS>stop>dTrpA1/dsxGA44(12) right eye painted: E 2 60 targets: Fŵ 60 -Male R71G01-LexA, dsxGAL4 Male R71G01-LexA, dsxGAL4 targets: UAS>stop>dTrpA1myc/R71G01-LexA; two constantly two immobile dTrpA1myc intersection myrGFP intersection LexAop2-FlpL/dsxGA44(52) rotating ubber bands rubber bands Inde 40 40 đ ġ 20 20 õ 8

Pan., et al. PNAS. 2012

Female LexA

two constantly rotating

rubber bands

D

ð

100um

UAS-dTrpA1/+: R71G01-GAL4/+

UAS-dTrpA1/+; pBDPGAL4u/+

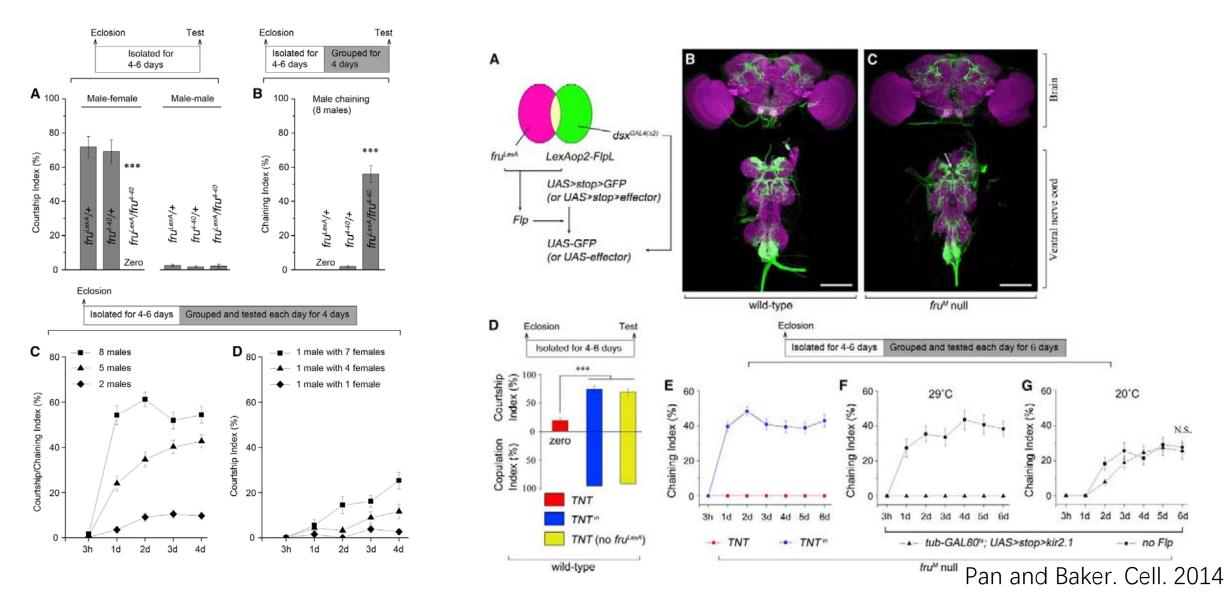
30°C

22°C

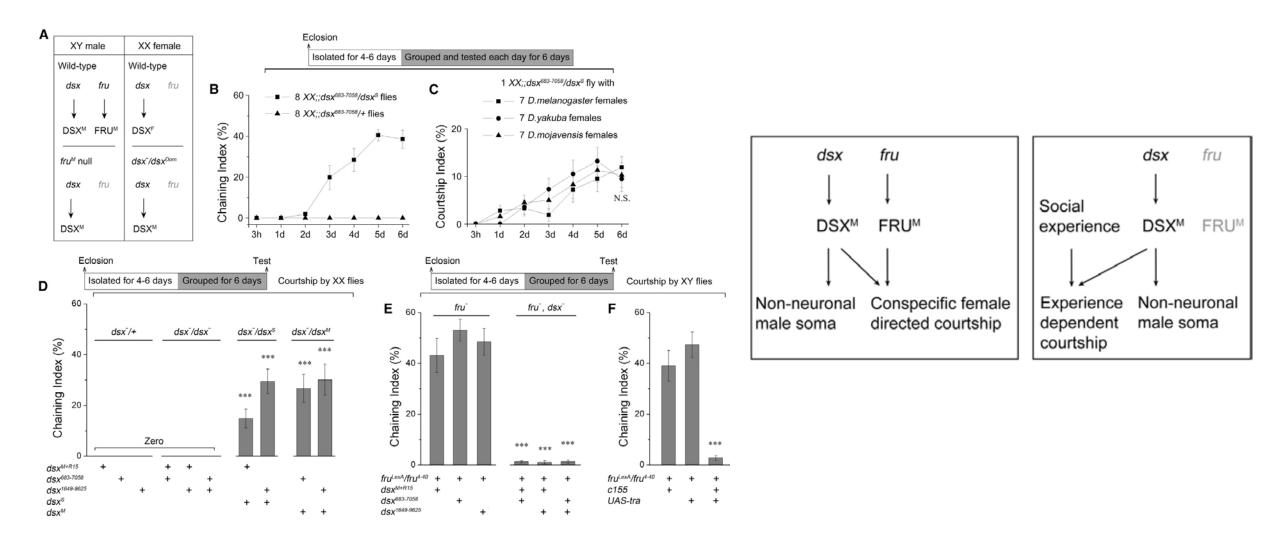
30°C

22°C

Activity of *fru^M* and *dsx* overlapping neurons is necessary during adulthood for courtship acquisition in *fru^M*-null males.



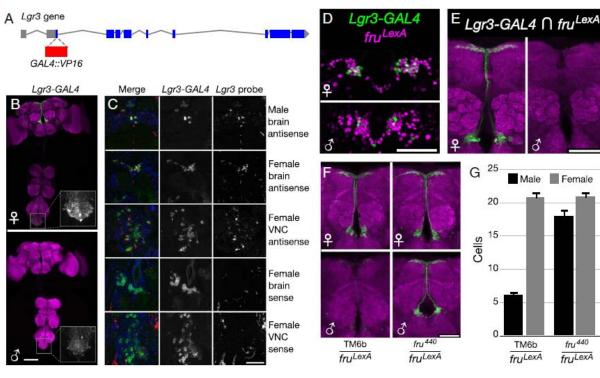
dsx^{M} is necessary and sufficient for the acquisition of the potential for such experience-dependent courtship.



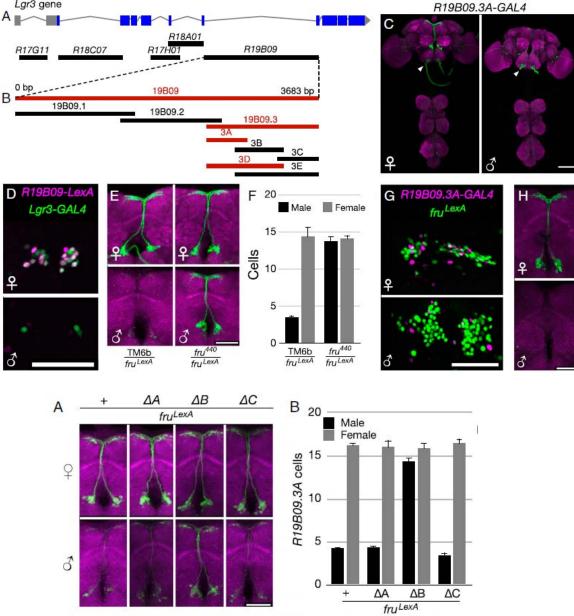
Pan and Baker. Cell. 2014

FruM Inhibits Expression of *Lgr3* in the Male Median Bundle, and Fru^M Acts Through a Small Region of an *Lgr3* Intron.

Lgr3 is a member of the leucine-rich repeat G-protein-coupled receptor (Lgr) family



the **B** isoform of Fru, whose DNA binding domain interacts with a short region of an Lgr3 intron.

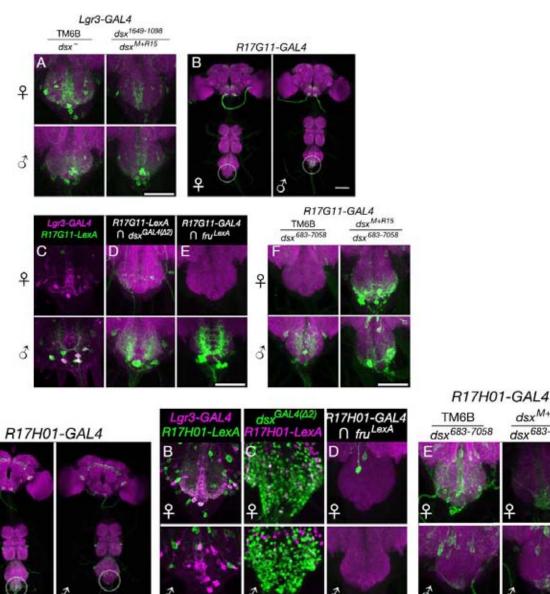


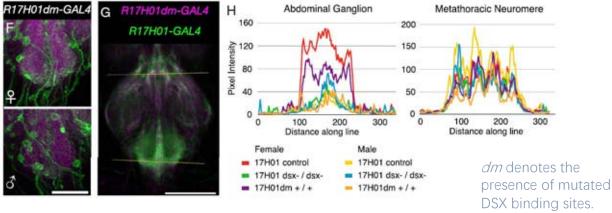
Meissner, G. W., et al. PNAS. 2016

Dsx^F plays the direct controlling role, it activates *R17H01* expression in females, but inhibits R17G11, R17H01dm-GAL4 G R17H01dm-GAL4 н Abdominal Ganglion 160 200

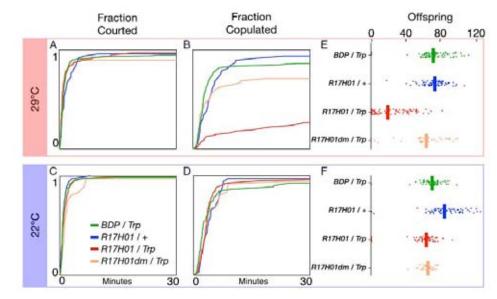
dsx^{M+R15}

dsx⁶⁸³⁻⁷⁰⁵⁸





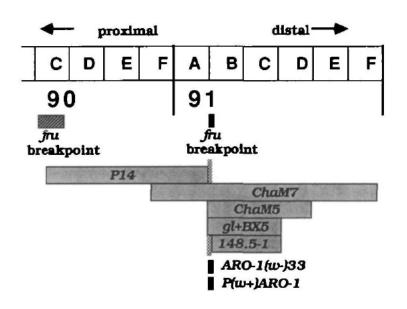
Activation of *R17H01*-GAL4 neurons reduces female receptivity and fecundity.



Meissner, G. W., et al. PNAS, 2016

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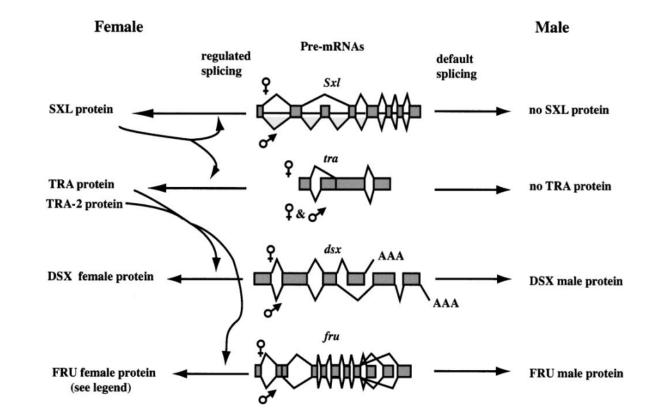


Fig. 1. Approximate breakpoint cytology of chromosomal

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