The LacZ expression in flies which genotypes contain YS was registered, but its intensity was weaker than in X/Y. An analogous experiment was performed with YL.

The data obtained lead to the conclusion that the YS but not the YL contains factors that control the expression of male fertility genes. However the possibility must be excluded that the Y-chromosome aneuploidy interrupts in a nonspecific manner the expression of the majority of genes involved in spermatogenesis by disruption of this process *per se*.

To test this assumption we use P[lArB] transposant stocks P103 (described earlier, Omelyanchuk, 1995) and Bg 9.61 (Lis *et al.*, 1983). In P103 individuals, staining was observed in neural ganglia, imaginal disks, ovaries and testes. The Bg 9.61 stock contains insertion of a construct where the LacZ gene is under control of heat-shock promoter. All tissues were stained after heat-shock in this stock.

The relults of testes staining in males containing different sets of sex chromosomes and insertions P103 and Bg 9.61 are shown in Table 4. The expression of LacZ in flies of X/O and X/Y genotypes is similar. So the possibility that the Y chromosome aneuploidy nonspecifically interrupts gene expression in testes may be excluded. And we can conclude that the YS arm contains factors responsible for the transcription of male fertility genes in *D. melanogaster*.

References:; Lis, J.T., et al., 1993, Cell 35:403-410; Omelyanchuk, L., 1995, Rus. J. Genet. 31:1645-1649.

Singh, B.K., and R.S. Fartyal. Cytogenetics Laboratory, Department of Zoology, Kumaun University, Nainital-263002, India. A list of Drosophilid species so far described and recorded from Kumaun region, India.

The great importance of Drosophilidae in genetic and evolutionary studies evoked most of the countries of the world to study the Drosophilid fauna thoroughly. However, the Indian subcontinent still remains a virgin field to be explored.

The Kumaun region, a wild hilly area is located at an elevation of about 6,000 ft. (1828 meters)

from the sea level on the north east periphery of the state Uttar Pradesh. This region includes four border districts of the state, Nainital, Almora, Pithoragarh and Udham Singh Nagar.

Although more than 300 species of Drosophilidae have been described and recorded so far from different parts of Indian subcontinent (Gupta, 1981, 1985), a vast area of great ecological interest still awaits exploration. Our knowledge in this field seems scanty in comparison with other countries of the world. Due to the above situation an extensive study of the Drosophilidae of Kumaun region was done, which is almost a virgin field for the above mentioned study. The following table shows the results of surveying studies of Drosophilids of Kumaun region since 1984 to 1996. The present surveying studies shows that the members of the Drosophilidae are fairly distributed in Kumaun region, particularly the members of the subgenus *Drosophila* of the genus *Drosophila*.

List of Drosophilid species so far described and recorded from Kumaun region:

GENUS

Amiota Loew

Subgenus

Phortica Schiner

1. bandes Singh and Negi, 1992

GENUS

Gitona Meigen

2. distigma, 1830

GENUS

Leucophenga Mik

- 3. bellula (Bergrowth, 1984) guttiventris (de meijere, 1908) Syn. ref. Bock, 1979, Aust. J. Zool. Suppl. Ser. 71:25
- 4. neolacteusa Singh and Bhatt, 1988
- 5. angulata sp. nov. (In press)
- 6. Okhalkandensis sp. nov. (In press)
- 7. Clubiata sp. nov. (In press)

GENUS

Paraleucophenga Hendel

- 8. neojavanaii Singh and Negi, 1992 Lissocephala Malloch
- 9. parasiatica Takada and Momma, 1975

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GENUS

Scaptomyza Hardy

10. himalayana, Takada, 1970

11. quadruangulata Singh and Dash, 1993

GENUS

Zaprionus Coquillette

12. indianus Gupta, 1970 (for indiana)

Subgenus

Dorsilopha Sturtevant

13. busckii Coquillette, 1901

Subgenus

Drosophila Falle'n Str.

14. analspina Singh and Negi, 1995

15. bishti Singh and Negi, 1995

16. bageshwarensis Sp. nov. (In press)

17. immigrans Sturtevant, 1921

18. lacertosa Okada, 1956

19. nainitalensis Singh and Bhatt, 1988

20. nasuta Lamb, 1914

21. paunii Singh and Negi, 1989

22. painaii Singh and Negi, 1995

23. repleta Wollaston, 1858

24. sulfurigaster Duda, 1923

25. trizonata Okada, 1966

26. serrata Sp. nov. (In press)

27. paramarginata Sp. nov. (In press)

28. hexaspina Sp. nov. (In press)

29. surangensis Sp. nov. (In press)

30. paharpaniensis Sp. nov. (In press)

31. khansuensis Sp. nov. (In press)

32. elongata Sp. nov. (In press)

Subgenus

Scaptodrosophila

33. coracina Kikkawa and Peng, 1938

34. chandraprabhiana Gupta and Ray-Chaudhury, 1970

35. hirsuata Sp. nov. (In press)

Subgenus

Sophophora Sturtevant

36. bifasciata Pomini, 1940

37. jambulina Parshad and Paika, 1964

38. kikkawai Burla, 1954

39. malerkotliana Parshad and Paika, 1964

40. melanogaster Meigen, 1830

41. nepalensis Okada, 1955

42. suzukii indicus Parshad and Paika, 1964

43. takahashii Sturtevant, 1927

44. neobaimaii Sp. nov. (In press)

45. neokhaoyama Sp. nov. (In press)

46. saraswatii Sp. nov. (In press)

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References: Gupta, J.P., 1981, Dros. Inf. Serv. 56: 50-53; Gupta, J.P., 1985, Dros. Inf. Serv. 61: 86-88. Additional Information:

Drosophila Stocks wild:

- 1. Drosophila melanogaster
- 2. Drosophila jambulina
- 3. Drosophila kikkawai
- 4. Drosophila malerkotliana
- 5. Drosophila immigrans
- 6. Drosophila nepalensis
- 7. Zaprionus indianus

Nongthomba, U., and N.B. Ramachandra. Drosophila Stock Centre, Department of Studies in Zoology, University of Mysore, Manasagangotri, Mysore 570 006, India. Induction and isolation of chromosome specific indirect flight muscle mutations in Drosophila melanogaster.

The indirect flight muscles (IFMs) of Drosophila melanogaster provide a unique model system to genetically dissect muscle structure and function (Sparrow et al., 1991). These are the bulk of thoracic muscles consisting of two groups, namely dorsal longitudinal muscles (DLMs) which are composed of six fibers from dorsal to ventral and dorsoventral muscles (DVMs), DVM I, DVM II, DVM III

which are formed by 3, 2, and 2 fibers, respectively. The wild type IFM development has been studied (Crossley, 1978; Fernades *et al.*, 1991). *Drosophila* is also a suitable system because mutations that affect IFMs development can readily be isolated and these mutations do not affect much on the viability of the flies. Hence, the genetic analysis of mutants has greatly advanced the understanding of muscle development.

A number of mutations affecting the IFMs have been reported (Crossley, 1978; Lindsley and Zimm, 1992). Most of these mutations are quite general in their expression which affect all the flight muscles. However, the genes involved in the development of IFMs have not been identified So far no other systematically. chromosome 2 specific mutation has been reported which affects the 1FMs development except Mhc gene. In view of this, investigations are made to identify and characterize genes involved in IFM development which reside on chromosome 2 by using ethyl methanesulfonate (EMS) Here we report the mutagenesis. induction and isolation of 16 new viable mutations on the second chromosome Drosophila in

Table 1. Scheme for Ethyl Methanesulfonate Mutagenesis

Generation	Cross		
	Females		Males
Parental	CyO / Tft	* \	+ / + (EMS treated)
F ₁	CyO / Tft	_ * \ `	*/#
F ₂	*1 CyO	×	*/CyO
F ₃	* / * males and females scored for abnormal wing and IFMs		* / CyO males and females females retained for stocks

CyO / Tft = Chromosome 2 marker; * = mutagenised chromosome # = CyO 0r Tft male.

melanogaster which affect the indirect flight muscles.

The Canton-S strain and Curly Oster/Tufted (CyO/Tft) mutant strain of $Drosophila\ melanogaster$ were used as wild type and dominant markers of chromosome 2, respectively (Lindsley and Zimm, 1992). All the stocks were cultured on standard wheat cream agar medium at 24 ± 1 °C. 25mM of EMS was administered to the Canton-S male flies following the procedure of Grigliatti (1986). The protocol used for induction and detection of mutation on chromosome 2 is presented in Table 1. Control experiments for EMS mutagenesis were made using X^XY stocks. For muscle analysis, thoracic whole mounts were prepared following the procedure described by Fyrberg $et\ al.$ (1995). Complementation analysis was done by crossing the virgins of each of the newly-isolated mutants reciprocally and analyzing the progenies for the defects in wings and IFMs.

The summary of the EMS mutagenesis on chromosome 2 which affect IFMs phenotypes in *D. melanogaster* is given in Table 2. A total of 3283 mutation induced lines were screened. Of these, 70.5%, 2.2% and 27.3% mutations were lethal, sterile and viable, respectively. Out of the 27.3% viable homozygotes scored, 3% were of wing mutants, of