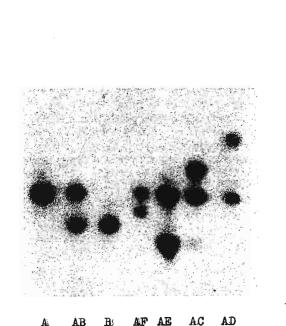
Trippa, G., C. Barberio, A. Loverre and P. Santolamazza. Istituto di Genetica, Universita di Roma, Italia. Phosphoglucomutase (PGM) polymorphism in natural populations of Drosophila melanogaster.

The distribution of phosphoglucomutase (PGM) polymorphism has been investigated in seven natural Italian populations. The samples were collected in Puglia (Castellaneta, Otranto and Corato) and Sicily (Ranna, Pedalino, Vittoria and Archi) during September-October 1971. The zymograms of single fly homogenates, following

the technique by Spencer et al. (1964) and modified by Trippa et al. (1970) for single fly homogenate, show that the wild populations contain six anodic forms of PGM, each showing dif-

ferent electrophoretic mobility.



In addition to the more common and already described alleles PgmA and PgmB (Trippa, G. DIS 46:42, 1971) four new alleles have been found. They have been called PgmC, PgmD, PgmE and PgmF (Trippa, G., New Mutants, this issue). The corresponding forms migrate from the cathode to the anode in this order: PgmE, PgmB, PgmF, PgmA, PgmC and PgmD. Before PGM phenotype determination

all males were crossed with females

Fig. 1. The phosphoglucomutase patterns of the different genotypes observed in the present survey.

homozygous for a known Pgm allele. The frequencies of the PGM phenotypic classes were the expected ones in all the offspring of those males who turned out to be carriers of new Pgm alleles.

The observed PGM phenotypic values and the allele frequencies found in the seven natural populations are shown in

Table 1. The chi-square test for the Hardy-Weinberg equilibrium in the populations, where it was possible, showed no significant difference between the values for the observed against the

origin

| | PGM phenotypes | | | | | | | a | | | |
|--------------|----------------|------------|----------|----|------------------|----------|-----------|-------------|---|--|--|
| Populations | _A_ | AB | <u>B</u> | AC | \underline{AD} | BC | <u>AE</u> | TOT | Gene Frequencies | | |
| Castellaneta | 177 | 2 8 | 4 | 3 | 1 | - | - | 213 | $Pgm^{A}=90.6 \pm 1.38$ $Pgm^{C}=0.7 \pm 0.40$ | $Pgm^{B}= 8.5 \pm 1.34$ $Pgm^{D}= 0.2 \pm 0.21$ | |
| Otranto | 25 8 | 29 | 1 | 2 | 5 | 1 | - | 2 96 | $Pgm^{A}=93.2 \pm 1.03$ $Pgm^{C}=0.5 \pm 0.29$ | $Pgm^{B}= 5.4 \pm 0.93$ $Pgm^{D}= 0.8 \pm 0.37$ | |
| Corato | 207 | 24 | - | 1 | - | - | 1 | 2 33 | $Pgm^{A}=94.4 \pm 1.06$ $Pgm^{C}=0.2 \pm 0.20$ | $Pgm^{B}= 5.2 \pm 1.03$ $Pgm^{E}= 0.2 \pm 0.20$ | |
| Ranna | 184 | 17 | - | 4 | 1 | - | - | 2 06 | $Pgm^{A}=94.7 \pm 1.10$ $Pgm^{C}=1.0 \pm 0.48$ | $Pgm^{B} = 4.1 \pm 0.97$ $Pgm^{D} = 0.2 \pm 0.22$ | |
| Pedalino | 143 | 19 | - | - | - | - | - | 162 | $Pgm^{A}=94.1 \pm 1.31$ | $Pgm^B = 5.9$ | |
| Vittoria | 31 | 11 | 4 | - | - | - | - | 46 | $Pgm^{A}=79.3 \pm 4.22$ | $Pgm^B=20.7$ * | |
| Archi | 197 | 3 | _ | - | - | <u>-</u> | - | 200 | Pgm ^A =99.2 ± 0.44 | Pgm ^B = 0.8 | |

Table 1. Distribution of PGM phenotypes and allele frequencies in the seven wild populations.