Subject Index

Entries followed by the letters (Abs) refer to abstracts of papers given at a meeting of the Genetical Society.

Adh locus of Drosophila, detoxification associated with, 167-175
alcohol ingestion in Drosophila, metabolic response to, 123-128
alleles, rare, distribution of, 409-416
allelic and genotypic frequencies in housefly, 289-298
allelic expression of Gpdh in Drosophila, 227-235
allozymic similarities among 7 species of land snails, 319-327
aminoacylase-1 isoenzymes, 157-160
amylase loci in Asellus, 105-110
analysis of chromosome C-banding pattern, 365-372

Adh activity and Ethanol tolerance in Drosophila, 185-190

banding techniques and heterochromatin variants, 161-165 behavioural responses to novelty in mice, 55-63 breeding structure in housefly, 289-298

chiasma frequency and bivalent length, relationship between, 305-310 chiasma variation in 2 chromosomes, 177-184 chromosome association at meiosis in hybrids, 97-103 chromosome segment supernumerary in grasshopper, 237-241 comparison of inbred lines, 263-276 comparison of morphs of Senecio vulgaris, 355-364 crossfertilisation in Pellia borealis, 299-304 cytogenetic studies in gomphocerine grasshoppers, 365-372

detoxification associated with Adh in Drosophila, 167-175 development density and longevity in D. melanogaster, 329-335 diazinon resistance locus, selection at, in a blowfly, 373-380 differential grazing by a mollusc, 277-281 digging in Drosophila larvae, 33-41 distribution of peroxidase genotypes in Pellia borealis, 299-304 distribution of rare alleles in a subdivided population, 409-415 DNA variability in Hawaiian Drosophila, 75-85, 87-96 dominance for Gpdh in Drosophila, 227-235 double interchange heterozygote, 1-6

enzyme heterozygosity, possible influence of, on development in trout, 417-425

Ethanol tolerance and Adh activity in Drosophila, 185-190 evolution of breeding system in Primula vulgaris, 207-217

frequency-dependent viabilities, 7-17

gene duplication in diploid and tetraploid Polygala species, 111-118

gene flow in *Bufo marinus*, 145-156 gene flow in *Plantago lanceolata*, 43-54 genetic analysis of *Drosophila*, 19-24 genetic architecture of behavioural responses in mice, 55-63 genetic control of plasma esterase polymorphism, 65-67 genetic manipulation, 284-285 genetic value of sons and daughters, 351-354 genetics of amylase variation in *Asellus*, 105-110 genetics of digging behaviour in *Drosophila* larvae, 33-41 genetic variation and development rate in rainbow trout, 417-

genetic variation in grasshopper populations, 397-408 genetic variation in mice, 157-160

genome combinations, analysis of, between wheat, rye and Aegilops, 219-226
genotype and supernumerary chromosomes, 305-310
genotype-environment interaction in barley, 255-262
genotype-environment interaction, methods of analysing, 243-

grazing differential of white clover by Arion hortensis, 277-281 gynodioecy in Plantago lanceolata, 355-364

heterochromatin and chiasma distribution, 177-184 heterochromatin variants in *Baetica ustulata*, 161-165 heterozygosity and development rate in rainbow trout, 417-425 homoeologous chromosome association in *Festuca arun-dinacea*, 97-103

hybrid zones in Heliconius butterflies, 191-202 hybrid zone patterns in field crickets, 337-349

inbred lines, comparison of, 263-275
inheritance of seed dormancy, 25-31
inheritance of style length in Oxalis rosea, 393-396
interactions of different combinations of wheat, rye and
Aegilops, 219-226
interference in a double interchange heterozygote, 1-6

karyotypes of *Drosophila*, viabilities of, 7-17

life history characteristics of Senecio vulgaris, 381-391 location, preferential, of human Y chromosomes, 203-205 longevity in Drosophila melanogaster, 329-335

male sterility in *Plantago lanceolata*, 355-364
measurement of genotype-environment interaction, 255-262
meiotic configurations, cytological analysis of, 1-6
metabolic response to alcohol in *Drosophila hydei*, 123-128
methods of analysing genotype-environment interaction, 243253

mitochondrial DNA variability in *Drosophila*, 75-85, 87-96 morphological differences between alien and British plants, 129-138

nucleolar activity associated with supernumerary chromosome, 237-241 nucleolar organiser activity in *Lolium*, 311-317

pairing interactions between wheat, rye and Aegilops chromosomes, 219-226

patterns of variation in a hybrid zone, 337-349 PGI isozymes in diploid and tetraploid *Polygala* species, 111-118

polymorphism and genetic control of plasma esterase, 65-67 polymorphism for capitulum type in *Senecio vulgaris*, 381-391 populations, introduced, of giant toad, genetics of, 145-156 population structure and gene flow, 43-54 population structure with limited dispersal, 69-73

quasi-equilibrium theory of distribution of rare alleles, 409-415

reproductive assurance in evolution of self-fertilisation, 207-217 438 SUBJECT INDEX

secondary constrictions in Festuca and Lolium-Festuca hybrids, variation in number of, 311-317
seed dormancy in Sinapsis arvensis, 25-31
selection for CES in Drosophila, 19-24
selection, frequency dependent, at a resistance locus in blowfly, 373-380
sex allocation in a temporally varying environment, 119-121
sex ratio evolution, 337-349
sex ratio in structured populations, 69-73
spatial distribution of nuclear and cytoplasmic genes, 355-364
style length, inheritance of, in Oxalis rosea, 393-396

variation, genetic, in grasshopper populations, 397-408 variation in alien *Lotus*, 129-138 variation in allozymes in 7 species of land snails, 319-327 variation, patterns of, in a hybrid zone, 337-349

warning colours in tropical butterflies, 191-202

Y polymorphism, influence of, in location of human chromosome, 203-205