

pGENE: A PISUM GENE DATABASE

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**pGene** may be used to examine information on the genes of peas. Genes may be selected by chromosome, gene symbol status or form of inheritance. For each gene, the gene symbol, symbol status, and the chromosome the gene is found on is displayed along with the mode of inheritance, the mutation group and sub-group according to the scheme of Blixt (1, 3), the mutation type and a short description (Fig. 1). A more extensive description of the effect of each gene, the associated bibliography and type-lines for each gene may also be displayed. Finally, a gene list for selected genes may be produced as a text file.

**pGene** comes as a database and standalone program designed to be used on IBM compatible personal computers. The original database was developed by Blixt on a Wang computer as part of a larger database of the Weibullsholm Pisum collection and at the time the Pisum Gene symbol list (2, 3) was compiled. It was subsequently converted into dBase III format at the John Innes Institute in 1988. I have restructured the database and written the **pGene** software.

The **pGene** database has been produced to make the database originally developed by Stig Blixt more widely accessible in electronic form and as the starting point for an up-to-date gene list. It should be of use to anyone with access to a personal computer who is interested in having a convenient reference to the genes of peas. To use **pGene** an IBM compatible computer with a minimum of 512K RAM and, preferably, a hard disk is required. The database plus software requires approximately 1MB of disk space. No additional software is required to use the database.

Biblio.	Edit	Find	Lines	Next	Prev	Select	Text	Write
Display next gene							Select a gene	
Genes selected: All								
Gene: a								
Status: UAL								
Chromosome: 1								
Inheritance: Rec								
Mutation group: 00								
Mutation sub-group: 01								
Mutation type: anthocyanin-inhibition								
<b>DESCRIPTION</b>								
Absence of anthocyanin production. A is necessary for general anthocyanin production in flowers, axils and seeds. Several alleles have been proposed.								
F1 Help			↑↓↔ Pick option		← Select		ESC Exit menu	

Fig. 1. Main gene interrogation screen of the **pGene** software.

A revised and updated version of the database is planned for release by the end of 1990.

Anyone who would like a copy of the database or further information should contact me (see below). I am particularly interested in receiving comments or suggestions concerning the structure and content of the database at this early stage in its development.

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