

# Cajanus genepool at ICRISAT genebank

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- ❖ Pigeonpea (*Cajanus cajan* (L.) Millsp.) is an important grain legume crop of the tropics and subtropics.
- ❖ It is an excellent source of protein, minerals and vitamins and has multiple uses such as food, feed, fuel, fencing, roofing, basket making and as a soil enricher and soil binder.
- ❖ ICRISAT genebank holds the world's largest collection of 13,632 pigeonpea germplasm accessions from 74 countries including 555 accessions of wild relatives belonging to 67 species of 6 genera.

- ❖ The genus *Cajanus* has 32 species, of which 20 are available at the ICRISAT genebank.
- ❖ *Cajanus* species belonging to primary and secondary gene pools are crossable with cultivated pigeonpea and possess resistance to one or the other biotic and abiotic stresses in addition to having agronomic traits useful in crop improvement.
- ❖ Small seed samples of wild accessions are available for research use following the Standard Material Transfer Agreement (SMTA), from the ICRISAT genebank.

Species/genepool	Plant	Pods	Seeds	No. of accessions	Useful traits
<i>Cajanus cajan</i> (L.) Millsp. (all cultivated) Primary gene pool				13,077	Major source of protein in Indian diet and has multiple uses like food, feed, fuel, fencing, roofing, basket making and as soil enricher and soil binder. Also used in traditional medicine
<i>C. acutifolius</i> (F. von Muell.) van der Maesen Secondary gene pool				12	Resistance to pod borer and podfly; tolerance to drought and salinity.
<i>C. albicans</i> (W. & A.) van der Maesen Secondary gene pool				20	Resistance to Sterility Mosaic Disease (SMD), alternaria blight, pod borer, podfly, pod wasp; tolerance to drought and salinity; high seed protein (> 30%). Crossable with cultivated pigeonpea
<i>C. cajanifolius</i> (Haines) van der Maesen Secondary gene pool				5	Progenitor, genetic male sterility, tolerance to salinity, pod borer, podfly and pod wasp. Crossable with cultivated pigeonpea
<i>C. lanceolatus</i> (WV Fitzg.) van der Maesen Secondary gene pool				1	No information
<i>C. lineatus</i> (W. & A.) van der Maesen Secondary gene pool				10	Resistance to SMD, alternaria blight and drought tolerance. Crossable with cultivated pigeonpea
<i>C. latisepalus</i> (Reynolds & Pedley) van der Maesen Secondary gene pool				1	No information
<i>C. scarabaeoides</i> (L.) Thou. Secondary gene pool				102	Early flowering, high pod setting; resistance to wilt, SMD, phytophthora blight, alternaria blight, pod borer, podfly, cyst nematode; tolerance to waterlogging; high seed protein (> 28%). Crossable with cultivated pigeonpea
<i>C. sericeus</i> (Benth. Ex Bak.) van der Maesen Secondary gene pool				4	Resistance to SMD, phytophthora blight, alternaria blight, pod borer and podfly; tolerance to drought and salinity; photoperiod insensitivity; high seed protein content (> 29%). Crossable with cultivated pigeonpea
<i>C. reticulatus</i> (Dryander) F. von Muell. Secondary gene pool				8	Resistance to pod borer and root-knot nematodes; tolerance to drought; high seed protein (33%).

Species/genepool	Plant	Pods	Seeds	No. of accessions	Useful traits
<i>C. trinervius</i> (DC.) van der Maesen Secondary gene pool				3	Crossable with cultivated pigeonpea
<i>C. cinereus</i> (F. von Muell.) F. von Muell. Tertiary gene pool				1	High seed number per pod
<i>C. confertiflorus</i> F. von Muell. Tertiary gene pool				1	No information
<i>C. crassus</i> (Prain ex King) van der Maesen Tertiary gene pool				10	Resistance to SMD
<i>C. elongatus</i> (Benth.) van der Maesen Tertiary gene pool				1	No information
<i>C. goensis</i> Dalz Tertiary gene pool				1	High seed number per pod
<i>C. marmoratus</i> (R. Br. ex Benth.) F. von Muell. Tertiary gene pool				2	No information
<i>C. mollis</i> (Benth.) van der Maesen Tertiary gene pool				8	High seed protein content (33.4%); high seed number per pod
<i>C. platycarpus</i> (Benth.) van der Maesen Tertiary gene pool				17	Resistance to phytophthora and alternaria blight; tolerance to salinity; annuality; photoperiod insensitivity; extra early flowering; high harvest index; high seed protein (27-31.6%); high cystine + methionine (> 29%).
<i>C. rugosus</i> (W. & A.) van der Maesen Tertiary gene pool				6	No information

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