

# Agro-morphological characterization of Bay Islands Pigeonpea (*Cajanus cajan*) landraces and advanced lines under Islands conditions









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**Central Island Agricultural Research Institute**  
**Port Blair (A & N Islands)**

## Outline:

-  **Sustainable aspects of pulse crops in general**
-  **Pulse crops of the Andaman & Nicobar Islands**
-  **Importance of pigeonpea crops for Island agriculture**
-  **Agro-morphological variation of pigeonpea landraces**
-  **Utilization of landraces**
-  **The way forward**

# Pulses and their sustainable aspects

- ➔ Pulses are essential for the regeneration of nutrient-deficient soils and for providing needed protein, minerals, and vitamins to humans and livestock.
- ➔ Pulses can be a means of improving the livelihoods of smallholder farmers.
- ➔ Pulses for the nutritional and food security of human beings;
  - ❖ *As a source of protein.*
  - ❖ *As a source of important vitamins and minerals*
  - ❖ *As a way of reducing cholesterol and blood sugar.*
- ➔ Pulses are more resilient to the rainfed/ dryland tropics.
- ➔ Pulses are the source of soil health improvement and environmental degradation
- ➔ Pulses for animal nutrition
- ➔ Pulses for crop and soil improvement

# Major Pulse crops of Islands

<b>Mungbean (<i>Vigna radiata</i>)</b>	<b>Cultivated / landraces</b>
<b>Urdbean (<i>Vigna mungo</i>)</b>	<b>Cultivated / landraces</b>
<b>Pigeonpea (<i>Cajanus cajan</i>)</b>	<b>Cultivated / landraces</b>
<b>Cowpea (<i>Vigna unguiculata</i>)</b>	<b>Cultivated / landraces</b>
<b>Beachpea (<i>Vigna marina</i>)</b>	<b>Endemic and endangered wild relative of <i>Vigna</i></b>

# Land utilization pattern of the Islands

- Total cultivated land pre tsunami (approx.) = 50,000.00 ha
- Permanently submerged land = 4,206.00 ha
- Total cultivated land post tsunami = 45,794.00 ha
- Total cropped area (2012-13) = 16,535.22 ha
- Fallow land = 3281.60 ha
- Net area sown = 14710.07 ha

Crops	Area (ha)	Area (%)
Paddy	8,390.00	18.32
Maize	163.54	0.36
<b>Pulses</b>	<b>2910.00</b>	<b>5.82</b>
Oilseeds	46.10	0.10
Condiments & Spices	1610.80	12.69
Vegetables	5150.00	11.25
Coconut and Arecanut	25920.00	56.60

# Area and Distribution of Pigeonpea in Andaman & Nicobar Islands

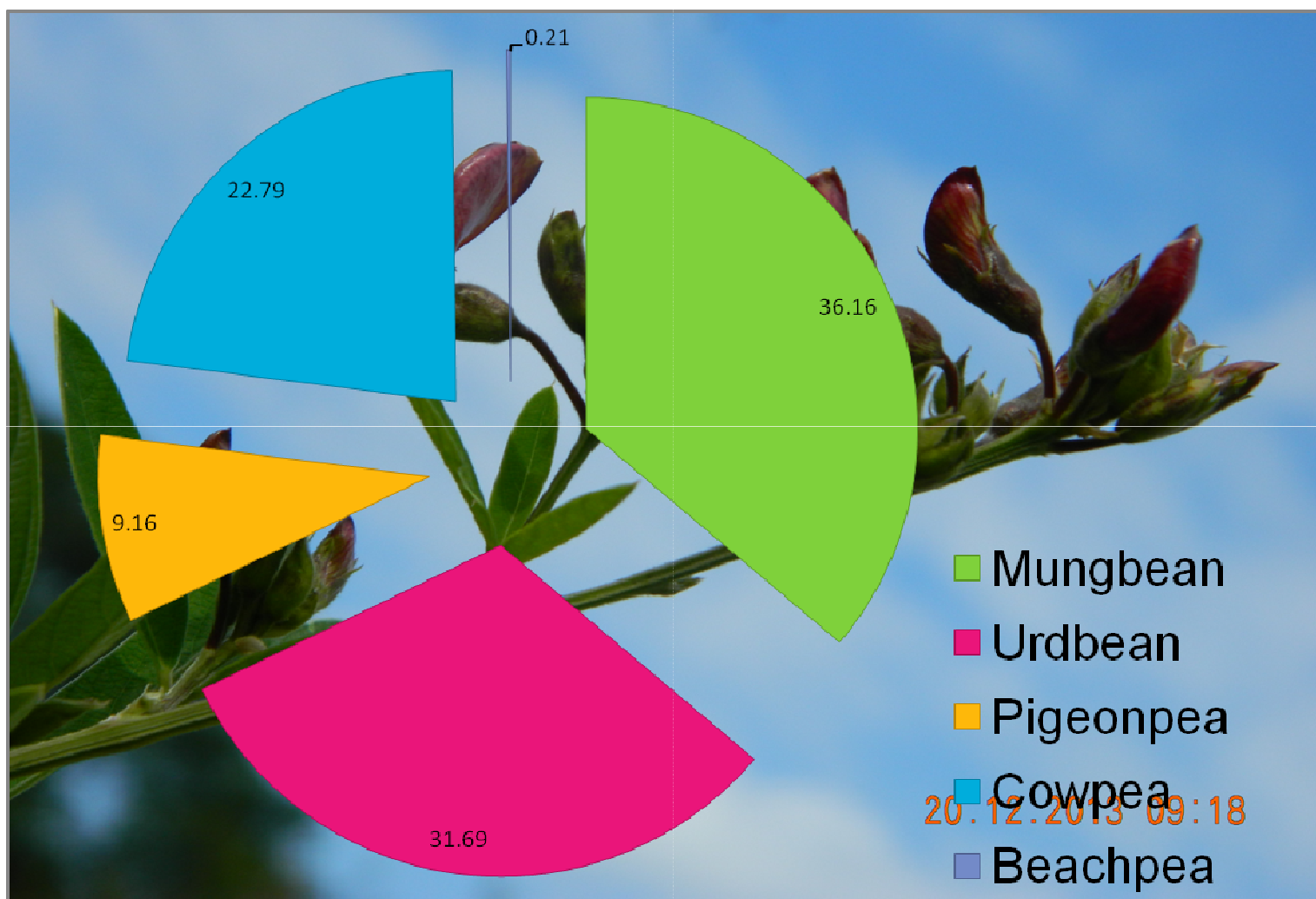


# Scenario of pulse cultivation in the Island



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# Share of different pulse crops in Island Agriculture



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# Pigeonpea

- Pigeonpea (*Cajanus cajan* (L.) Millsp.) is an important grain legume that originated in the Indian sub-continent.
- The crop is grown for its multiple benefits mainly by smallholder growers and is useful in providing household food security in the region.
- The crop provides highly nutritious food for human consumption and fixes considerable amounts of atmospheric nitrogen, thus improving soil fertility. D
- The crop is also relatively tolerant to drought (Kumar et al., 2011) thus making it suitable for cultivation in the semi-arid agro-ecological conditions prevalent in the region.

# Pigeonpea

- ➡ Crop of marginal area
- ➡ Improve the soil fertility
- ➡ Suitable for varying cropping system
- ➡ Improve organic matter content
- ➡ Suitable for degraded land
- ➡ Grain, fodder, fire wood and medicinal use
- ➡ Adoption to varied agro-ecological conditions of the Island

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# Importance of pigeonpea for Island agriculture

- ➡ **Declining factors productivity**
- ➡ **Depletion of soil fertility**
- ➡ **Imbalance use of nutrients**
- ➡ **Declining in organic matter content**
- ➡ **Problems of soil salinity and alkalinity**
- ➡ **Increasing disease and insect-pest pressure**
- ➡ **Menace of weeds**
- ➡ **Environmental pollution**



*Pigeonpea improvement and conservation programs depend on the presence of genetic variability and the accurate characterization of that variability.*

*The distribution of genetic diversity between and within pigeonpea landraces of Island regions is largely unknown. An understanding of the genetic diversity is essential for both utilisation and conservation strategies.*

*Determination of genetic diversity of any given crop species is a suitable precursor for improvement of the crop because it generates baseline data to guide selection of parental lines and design of a breeding scheme.*

## Objective of the study:

*The early systematic studies of the genus *Cajanus* were based on phonological or morphological characters, which have been shown to have limited genetic resolution especially at species levels, as is required for pigeonpea.*

- ❖ **To determine the genetic diversity and relationships within and among pigeonpea landrace accessions**
- ❖ **Morphological characterization of pigeonpea accessions/landraces**
- ❖ **Genetic differentiation among populations**

*Collection and characterization  
of pigeonpea landraces*



# Pigeonpea ( *Cajanus cajan* L.)

Crop: Pigeonpea ( <i>Cajanus cajan</i> L.)	Year of Collection & ex-situ conservation of germplasm			
	2011-12	2012-13	2013-14	Total
Local landraces	26	17	08	51
Procurement of germplasm from others sources (IIPR & SAU's etc.)	18	05	19	42
<b>Total</b>	<b>44</b>	<b>22</b>	<b>27</b>	<b>93</b>



# Agro-morphological studies of Pigeonpea landraces and advanced breeding lines

SN	Experiment	No. of Accession	Replication	Year
1.	PGET	78	Augmented	2011-12 2012-13
2.	Mother Trial	52	RBD	2012-13
3	Observation Nursery (IVT)	32	RBD	2013-14







- ❖ ***Ex-situ* conservation of germplasm lines**
- ❖ **Characterization on VS & VG basis & utilization of selected germplasm with advanced generation materials for forward breeding programme.**
- ❖ **Evaluation of selected lines for yield and improved plant types in replicated trials.**
- ❖ **Screening and evaluation of promising favourable material for specific defects and desirable selection through broaden the spectrum of genetic variability.**

# ***Morphological variations for plant growth and flower types***



# ***Morphological variations for plant types, seed colour and seed size***



## Agronomic performance of landraces and advanced breeding lines based on morphological markers

Genotypes	Days to 50% Flowering	Days to maturity	Plant height at maturity	Number of Branches/ plant	Length of Pods (cm)	100-seed weight (g)	Seed yield per plant (g)	Seed yield per Plot (g)
NA-1	111.22	216.63	203.68	10.46	3.98	11.95	38.62	170
CO-6	143.62	221.88	190.92	11.35	4.37	16.67	46.05	237
Bahar	199.76	227.04	192.37	8.27	3.91	13.40	26.45	132
IPAC-68	146.00	192.00	243.00	23.40	6.68	23.11	106.78	800
IPAC-493	128.00	200.00	260.40	15.40	5.92	18.15	58.55	250
IPA-203	166.00	238.00	303.20	20.40	4.62	21.59	53.96	370
IPA-7-2	168.00	261.00	217.40	16.60	4.96	19.57	38.26	300
IPAC-452	175.00	261.00	242.60	18.00	4.94	20.44	82.22	550
IPA-9F	175.00	261.00	288.20	14.20	5.04	20.44	51.24	475
IPAC-66	176.00	261.00	183.40	9.60	5.22	14.51	32.49	125
IPAC-142	180.00	261.00	225.00	11.40	4.66	15.95	26.39	175
KPBR-80-2-1	166.00	261.00	254.80	15.20	4.28	18.12	30.84	250
IPA-7F	180.00	261.00	236.20	12.80	4.96	19.57	38.26	300
IPAC-451	197.00	261.00	235.80	13.40	5.72	21.12	42.18	375
IPA-8F	177.00	261.00	231.00	11.00	4.44	14.44	41.09	195
IPAC-165	180.00	261.00	248.80	17.20	5.62	13.79	55.73	225
ANP-12-03	128.00	261.00	261.00	16.40	4.26	19.57	33.81	300
BSMP-736	126.00	261.00	257.80	16.40	6.24	13.79	26.98	185
IPAC-12	110.00	261.00	267.00	12.80	4.96	21.00	36.77	350
BSMP-853	177.00	261.00	232.60	12.40	5.88	16.68	31.83	200
ANP-12-05	193.00	261.00	240.40	15.60	6.60	21.66	26.39	175
IPAC-8	96.00	261.00	229.80	14.20	5.76	21.11	26.18	375
IPAC-79	189.00	261.00	238.60	17.00	4.72	21.66	36.77	175
ANP-13-01	152.00	258.00	193.60	10.00	6.58	18.78	74.35	650
ANP-11-13	167.00	258.00	206.20	18.20	4.54	18.77	67.44	650
ANP-11-14	167.00	258.00	207.00	16.40	4.54	21.75	61.09	450

## Agronomic performance of landraces and advanced breeding lines based on morphological markers

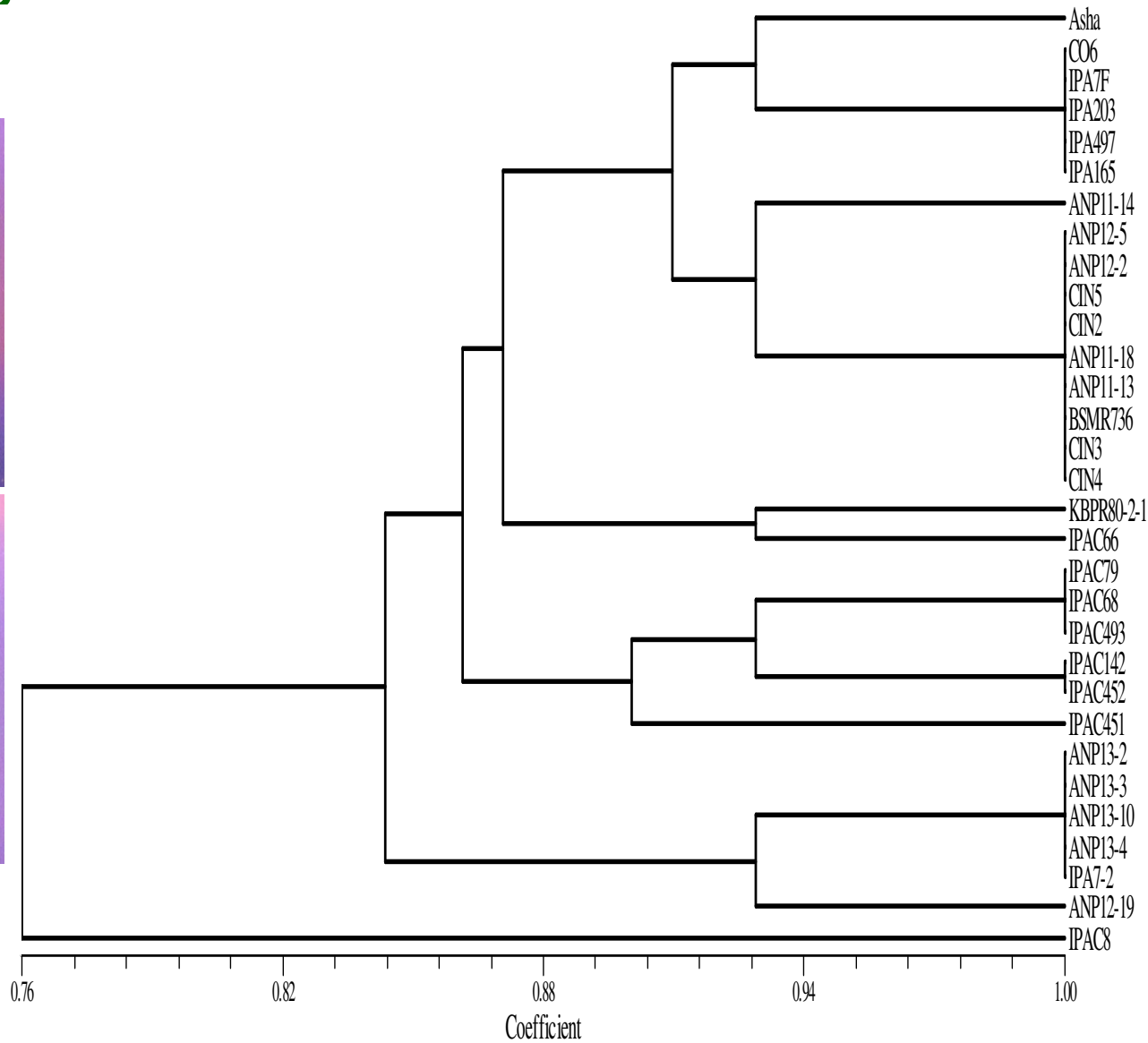
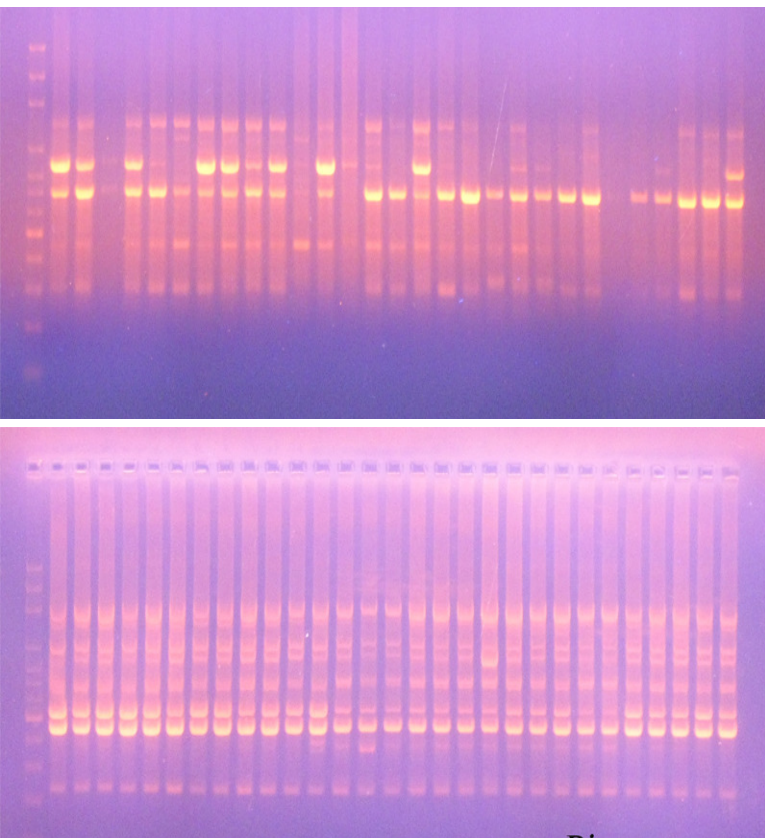
Genotypes	Days to 50% Flowering	Days to maturity	Plant height at maturity	Number of Branches/ plant	Length of Pods (cm)	100-seed weight (g)	Seed yield per plant (g)	Seed yield per Plot (g)
ANP-11-18	182.00	258.00	170.00	12.20	4.68	15.95	31.98	175
ANP-11-19	149.00	258.00	231.80	15.80	4.94	13.00	38.26	375
ANP-12-02	167.00	258.00	212.60	13.40	6.84	17.30	84.59	475
ANP-11-23	179.00	258.00	213.20	12.40	6.76	11.91	47.96	350
ANP-12-09	167.00	258.00	206.40	13.40	6.60	15.95	31.98	175
ANP-13-02	167.00	258.00	229.00	11.00	6.28	21.32	53.61	600
ANP-13-03	152.00	258.00	230.80	11.00	6.20	23.09	56.65	685
ANP-13-04	172.00	258.00	199.60	8.60	6.44	16.02	32.44	180
ANP-11-12	169.00	258.00	171.20	8.60	6.90	18.84	36.32	275
ANP-11-12-1	167.00	260.00	183.20	9.20	6.48	17.40	38.54	225
ANP-11-12-2	152.00	260.00	225.20	11.20	6.26	20.98	62.71	450
ANP-11-24	152.00	260.00	199.40	13.80	4.80	18.12	38.83	250
ANP-12-03	152.00	260.00	219.40	18.20	4.18	14.13	49.08	325
ANP-12-05	175.00	260.00	198.00	13.40	6.58	14.31	27.41	125
ANP-11-32	169.00	260.00	185.80	11.40	4.98	16.68	28.49	200
ANP-11-33	169.00	260.00	167.60	13.60	5.16	13.79	29.13	100
ANP-13-26	156.00	260.00	214.40	16.40	4.30	25.21	34.55	125
ANP-11-27	159.00	260.00	233.00	16.80	3.88	14.44	64.86	150
ANP-11-28	146.00	260.00	196.20	15.60	4.12	19.57	47.45	300
ANP-11-29	155.00	260.00	232.25	15.25	4.14	13.79	32.13	126
ANP-13-30	154.00	260.00	230.67	18.00	4.18	16.68	79.13	376
ANP-13-31	129.00	260.00	213.20	15.60	4.20	16.76	39.57	230
ANP-13-32	146.00	260.00	270.50	12.50	4.30	15.99	51.93	175
ANP-11-36	156.00	260.00	253.00	18.00	4.20	18.13	250.00	250
ANP-11-37	144.00	260.00	251.60	16.60	4.24	18.41	60.09	260
ANP-13-33	192.00	260.00	238.33	17.33	4.20	14.39	65.46	125

## Molecular characterization landraces and advanced breeding lines based on RAPD markers

### List of RAPD primers used for RAPD profiling

S.N.	Primer name	Primer sequence	No. samples/ No. samples amplified	Total No. bands	Range of marker(kb)	% polymorphism	PIC
1.	OPG03	GAGCCCTCCA	48/48	253	0.2-1.4	3.16	0.223
2.	OPG04	AGCGTGTCTG	48/42	112	0.3-1.4	10.70	0.494
3.	OPG05	CTGAGACGGA	48/46	124	0.35-1.4	10.40	0.497
4.	OPG06	GTGCCTAACC	48/22	75	0.2-1.4	25.33	0.313
5.	OPG07	GAACCTGCGG	48/19	65	0.2-0.8	23.07	0.265
6.	OPG08	TCACGTCCAC	48/47	163	0.35-0.7	5.52	0.430
7.	OPG10	AGGGCCGTCT	48/16	79	0.2-1.4	18.98	0.235
8.	OPG11	TGCCCGTCGT	48/24	92	0.3-1.2	20.65	0.365
9.	OPG14	GGATGAGACC	48/36	144	0.4-1.2	8.72	0.499
10.	<b>OPG15</b>	<b>ACTGGGACTC</b>	<b>48/23</b>	<b>96</b>	<b>0.3-1.2</b>	<b>14.58</b>	<b>0.453</b>

# Molecular characterization landraces and advanced breeding lines based on RAPD markers





## Identification of Promising pigeonpea lines for YET under Initial varietal trial

Genotypes	Days to 50% flowering	Plant height (cm) at maturity	Days to maturity	Number of branches/ plant	Length of Pods (cm)	100-seed weight (g)	Seed yield per plant (g)
ANP-13-01	152	193.60	228	10.00	6.58	18.78	74.35
ANP-11-13	167	206.20	239	18.20	4.54	18.77	71.44
ANP-12-02	167	212.60	247	13.40	6.84	17.30	84.59
ANP-11-12-2	152	225.20	260	11.20	6.26	20.98	69.71
ANP-11-27	159	233.00	246	16.80	3.88	14.44	66.86
ANP-13-30	154	230.67	263	18.00	4.18	16.68	79.13
ANP-11-36	156	253.00	239	18.00	4.20	18.13	250.00
IPAC-68	146	243.00	192	23.40	6.68	23.11	106.78
IPAC-452	175	242.60	245	18.00	4.94	20.44	82.22
<b>Mean</b>	<b>162.8</b>	<b>226.76</b>	<b>232.69</b>	<b>14.13</b>	<b>4.29</b>	<b>17.74</b>	<b>55.69</b>
<b>CD (Geno. x Control)</b>	<b>61.13</b>	<b>70.31</b>	<b>19.06</b>	<b>17.79</b>	<b>0.95</b>	<b>1.94</b>	<b>9.47</b>
<b>CV (%)</b>	<b>17.82</b>	<b>5.53</b>	<b>6.10</b>	<b>12.65</b>	<b>6.61</b>	<b>12.41</b>	<b>22.17</b>

