

# Species Datasheet

Datasheet No. A-140.085.006  
(family.genus.species)

DBT- Network Programme

## 1.Taxon:

Species:*Viciacordata*Wulfen ex Hoppe

Subspecies

Variety

Cultivar

Hybrid

Image file

2. **Synonyms:** *Vicia sativa* subsp. *cordata* (Wulfen ex Hoppe) Asch. & Graebn., *Vicia sativa* var. *cordata* Arcang.

## 3. Systematic Position: APG IV; Bentham and Hooker:

### Bentham and Hooker (1862)

Kingdom: Plantae

Division: Phanerogamia

Class: Dicotyledons

Subclass: Polypetalae

Series: Calyciflorae

Cohors: Rosales Bercht. & J. Presl

Ordo: Leguminosae Juss.

Subordo: Papilionaceae Giseke

Genus: *Vicia* L.

Species: *Viciacordata* Wulfen ex Hoppe

### APG IV (2016)

- Kingdom: Plantae
- Clade: Angiosperms
- Clade: Eudicots
- Clade: Rosids
- Order: Fabales Bromhead
- Family: Fabaceae Lindl.
- Subfamily: Faboideae Rudd
- Genus: *Vicia* L.
- Species: *Viciacordata* Wulfen ex Hoppe

## 4.Distribution:

Global Asia, Africa, Europe

India Experimental stations

5. Indigenous/Exotic/Endemic; Cultivated/Wild: Exotic

**6.Threat Status:**

IUCN

BSI

**7.Habit and Habitat:**Herbaceous, Scrambling climber

**8.Life Form:**Annual

**9.Economic Importance:**Fodder

**10. Probable Progenitor of:**Close relative of common vetch, *V.sativa*<sup>123,124,126,130,131,136,149</sup>

**11.DNA**

**C-value**                           **Methodology**

2C (3.95 pg)<sup>90,91,92</sup>      Microdensitometer

**12.Basic chromosome number(s):**x= 5<sup>5, 90</sup>

**13. Zygotic chromosome number(s):**2n= 10<sup>5,9,90,92,103,104,109,122,130</sup> ; 2n=12<sup>122</sup>

**14. Gametic chromosome number(s):**n = 5, 6<sup>122</sup>

**15.Specialized chromosomes (B chromosomes/Sex chromosomes/Polytene chromosomes/Neocentric chromosomes):**

Image file

**16.Ploidy level:**Diploid<sup>5,9,90,92,103,104,109,122,130</sup>

Image file

**17.Agametoploidy:**

**18.Nature of polyploidy (auto, segmental, allo, autoallo):**

**19.Genomic formula:**

**20.Abberrant chromosome number(s)(aneuploidy, aneusomy, polysomy):**

**21.Somatic chromosomes:**<sup>5,9,29,103,104,109,122</sup>

**Karyotype** Majority submetacentric /metacentric chromosomes

**Chromosome size**Small

**NOR chromosome(s)**2

**Degree of asymmetry**Symmetrical

Image file

**22. Banding pattern(s):**

Image file

**23. Physical mapping of chromosomes:**18S - 5.8S - 26S and 5S ribosomal gene families <sup>109</sup>

**In situ hybridization**

Image file

**Fluorescent in situ hybridization**

Image file

**24. Genomic in situ hybridization:**

Image file

**25. Linkage map:**

Image file

**26. Chromosome associations:**

**Female meiosis**

**Male meiosis**5II, 6II<sup>122</sup>

Image file

**27. Chromosome distribution at anaphase I:**

**28. Genetic diversity:**

**Chromosomal level**<sup>103</sup>

Image file

**DNA level**<sup>29, 44, 128, 131</sup>

**29. Any other information (Apomixis; Inversion; Male sterility; Pollen grain mitosis;**

**Pollen stainability; Translocation etc.):**Pollen stainability:76 - 98 %<sup>122</sup>