

Species Datasheet

Datasheet No. A-061.057.009
(family.genus.species)

DBT- Network Programme

1.Taxon:

Species: *Goodyerarecurva*Lindl.

Subspecies:

Variety:

Cultivar

Hybrid

Image file

2. Synonyms:

Epipactisprainii (Hook.f.) A.A.Eaton

Epipactisrecurva (Lindl) A.A.Eaton

Goodyeraprainii Hook.f

Goodyerarecurva var. *prainii* (Hook.f) Pradhan

Orchiodesrecurvum (Lindl.) Kuntze

3.Systematic Position:

APG IV (2016)

- Kingdom: Plantae
- Clade: Angiosperms
- Clade: Monocots
- Order: Asparagales Link.
- Family: Orchidaceae Juss.
- Subfamily: Orchidoideae
- Tribe:Cranichideae
- Subtribe: Goodyerinae
- Genus: *Goodyera* R.Br.
- Species:*Goodyera recurva* Lindl.

Bentham and Hooker(1862)

Kingdom: Plantae
Division: Phanerogamia
Class:Monocotyledonae
Series: Microspermae
Ordo: Orchideae
Tribus: Neottieae
Subtribus: Spirantheae
Genus: *Goodyera* R.Br.
Species: *Goodyera recurva* Lindl.

4.Distribution:

Global:Chinese Himalayas, Assam and the eastern Himalayas, 200-2300m

India: Assam, Meghalaya, Nagaland, Arunachal Pradesh

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

6.Threat Status:

IUCN:

BSI:

7.Habit and Habitat:Terrestrial and also on mossy tree trunks in thick forests at about 1550m

8.Life Form: Cryptophytes

9.Economic Importance:

10. Probable Progenitor of:

11.DNA

C-value Methodology

12.Basic chromosome number(s): $x=16^{24, 26, 33}$

13. Zygotic chromosome number(s): $2n=32^{24}$

14. Gametic chromosome number(s): $n=16^{24, 26, 33}$

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

Image file

16.Ploidy level:Diploid^{24, 26, 33}

Image file

17.Agametoploidy:

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

19.Genomic formula:

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

21.Somatic chromosomes:

Karyotype

Chromosome size

NOR chromosome(s)

Degree of asymmetry

Image file

22. Banding pattern(s):

Image file

23. Physical mapping of chromosomes:**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 14 II's and a ring IV (indicating structural heterozygosity) at M-I in 90% of PMC's, secondary associations frequent^{26, 33}

Image file

27. Chromosome distribution at anaphase I:**28. Genetic diversity:****Chromosomal level**

Image file

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

Pollen stainability; Translocation etc.): Translocation^{26, 33}