

# Species Datasheet

Datasheet No. G-011.005.001  
(family.genus.species)

## 1.Taxon:

Species: *Cunninghamia lanceolata*(Lamb.) Hook.

Subspecies:

Variety:

Cultivar:

Hybrid:

Image file

**2. Synonyms:** *Abies batavorum* Siebold ex Carriere, *A. lanceolata* (Lamb.) Poir., *Belis jaculifolia* Salisb., *B. lanceolata* (Lamb.) Hoffmanns., *Cunninghamia chinensis* de Vos, *C. jaculifolia* (Salisb.) Druce, *C. lanceolata* var. *corticosa* Z.Y. Que& J.X. Li, *C. lanceolata* var. *lanceolata*, *C. lanceolata* var. *unicanaliculata*(D.Y.Wang&H.L.Liu) Silba, *C. sinensis* R.Br., *C. sinensis* var. *prolifera* Lemée&H.Lev., *C. unicanaliculata* D.Y.Wang&H.L.Liu, *C. unicanaliculata* var. *pyramidalis*D.Y.Wang&H.L.Liu, *Larix chinensis* Mill., *Pinus abies* Lour., *P. lanceolata* Lamb., *Raxopitys cunninghamii* J.Nelson

## 3.Systematic Position:

**Christenhusz et al. (2011)**

- Class: Equisetopsida C. Agardh
- Subclass: PinidaeCronquist
- Order: Cupressales Link
- Family:Cupressaceae Gray
- Genus: *Cunninghamia*R. Br. ex A. Rich.
- Species: *C. lanceolata*(Lamb.) Hook.

**Bentham and Hooker (1862)**

Kingdom: Plantae  
Division: Phanerogamia  
Class: Gymnospermeae  
Ordo: Coniferae

## 4.Distribution:

**Global:** Cambodia, Laos, India, Mainland China, Vietnam,

**India:** Sikkim (Gangtok), Tamil Nadu, Uttarakhand (Dehradun)

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

## 6.Threat Status:

**IUCN:**Least concern

**BSI:**

**7.Habit and Habitat:**Evergreen tree (50 m tall), *C. lanceolata* to be a coniferous constituent of the Mixed Mesophytic Forest Formation of the warm temperate regions of China.

**8.LifeForm:**Phanerophytes

**9.EconomicImportance:**It is an attractive tree suitable for land scape,the wood is light, soft, fragrant, almost white, and durable. Large sizes are milled for construction timber in houses, for masts, carpentry and planks for coffins. Wood is termite proof; easily workable, used in building, construction of bridges, ships, lamp post, making furniture and for wood fibre.

**10. Probable Progenitor of:**

**11.DNA**

**C-value      Methodology**

2C (27.12 pg) <sup>14</sup>Feulgenmicrodensitometry<sup>14</sup>

2C (28.34 pg)<sup>6</sup> Flow cytometry<sup>6</sup>

2C (25.5 pg)<sup>19</sup> Flow cytometry<sup>19</sup>

2C (39.60 pg)<sup>19</sup> Flow cytometry<sup>19</sup>

**12.Basic chromosome number(s):** $x=11^1, 2, 3, 4, 5, 7, 8, 9, 10, 11, 12, 13, 14, 15, 16, 17, 22, 23, 24, 25$

**13. Zygotic chromosome number(s):** $2n=22^1, 2, 3, 4, 7, 8, 10, 11, 12, 13, 14, 15, 16, 17, 23, 24$

$33^{19}$

**14. Gametic chromosome number(s):** $n=11^5, 22, 25$

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

Image file

**16.Ploidy level:**Diploid<sup>1, 2, 3, 4, 5, 7, 8, 10, 11, 12, 13, 14, 15, 16, 17</sup>

Triploid (sporadic) <sup>19</sup>

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:**<sup>1, 2, 3, 4, 7, 8, 10, 11, 12, 13, 14, 15, 16, 17</sup>

**Karyotype**Mostly median, one pair submedian<sup>1, 2, 3, 4, 7, 8, 10, 11, 12, 13, 14, 15, 16, 17, 23, 24</sup>

**Chromosome size**Large<sup>1, 2, 3, 4, 7, 8, 10, 11, 12, 13, 14, 15, 16, 17, 23, 24</sup>

**NOR chromosome(s)**<sup>25, 8, 23, 24</sup>

**Degree of asymmetry**Moderately asymmetrical<sup>1, 2, 3, 4, 7, 8, 10, 11, 12, 13, 14, 15, 16, 17, 23, 24</sup>

Image file

**22. Banding pattern(s):**C bands <sup>11</sup>, G bands <sup>15</sup>, HSAG <sup>16</sup>, CMA+ <sup>21</sup>

Image file

**23.Physical mapping of chromosomes:**

**In situ hybridization**18SrDNA <sup>5</sup> , telomeric repeat Arabidopsis type <sup>25</sup>

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** 11 II 5, 22, 25

Image file

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

**28. Genetic diversity:**

**Chromosomal level**<sup>3, 12</sup>

Image file

**DNA level**<sup>18</sup>

**29. Any other information (Apomixis; Inversion; Male sterility; Pollen grain mitosis; Pollen stainability; Translocation etc.):**