Methods of *in vitro* storage of hops.



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Hop (Humulus lupulus L.)



Kingdom: Plants (Plantae) Subkingdom: Higher plants (*Telomophytae*) Phyllum: Vascular plants (Tracheophyta) Class: Angiosperms (*Magnoliopsida*) Order: Nettles (Urticales) Family: Hemp (*Cannabaceae*) Genus: hop (*Humulus* Linnaeus, sp.) **Species:** Common hop (Humulus lupulus L.) Japanese hop (Humulus japonicus)

•Perennial herbaceous climbing plant •stems annual, slender, climbing, up to 9 m in length •Dioecious with unisexual flowers (male and

female flowers develop on different individuals) •Chromosome number: 2n = 2x = 20•In hop gardens only female plants are grown

•Important technical crop: the main harvested products are female inflorescences (cones)

•Grown on one stand up to 10-15 years (wild hops can live up to 100 years)

Commercial propagation exclusively

vegetatively (shoot-cuttings, root-

cuttings, rootstocks) •Cones used in brewing industry – α - and β -bitter acids



Meristem culture

Meristem-derived explant: Size: 0,2 – 0,6 mm Morphology: apical dome + 1-2 leaf primordia



week 0



Isolation of apical meristem and

week 2

Growth of meristem explants of hop in *in vitro* culture during the cultivation on initiation medium MSC



week 5

ELISA testing and micropropagation of virus free plants



ELISA testing of meristem culture-derived shoots of hop for presence of viruses



Micropropagation of ELISA tested virus-free plants by culture of nodal explants

In vitro storage



Optimization of in vitro storage system



Genetic stability of in vitro stored material





PCR analysis of microsatellite sequences of hop cv. Premiant using the primer HBV5. For amplification of DNA regions containing microsatellite sequences the primer HBV5 was used and the PCR reaction was performed in 20 µl volumes containing 10mM Tris-HCl (pH = 8.3), 50mM KCl, 1.5 mM MgCl₂, 0,25 mM dNTP, 1mM primer, 0,75 U Taq-DNA polymerase and 25 ng DNA. The initial denaturation was done for 2 minutes at 94 C followed by 40 cycles of 1 min. at 94 C, 1 min. at 48 C and 3 min. at temperature 72 C. The last step was 7 min. at temperature 72 C. Amplified products were separated electrophoretically in 1.5% agaróse gel in 0.5 x TBE buffer and stained with ethidium bromide. We analyzed 30 clones: 1 = PRM/2/2 (M,V4); 2 = PRM/2/2 (M, V8); 3 = PRM/2/2 (M, V12); 4 = PRM/3 (D, V0); 5 = PRM/3 (NS, V4); 6 = PRM/3 (NS, V8); 7 = PRM/3 (NS, V4); 6 = PRM/3 (NS, V8); 7 = PRM/3 (NS, V4); 6 = PRM/3 (NS, V8); 7 = PRM/3 (NS, V4); 6 = PRM/3 (NS, V8); 7 = PRM/3 (NS, V4); 6 = PRM/3 (NS, V8); 7 = PRM/3 (NS, V4); 6 = PRM/3 (NS, V8); 7 = PRM/3 (NS, V4); 6 = PRM/3 (NS, V8); 7 = PRM/3 (NS, V4); 6 = PRM/3 (NS, V8); 7 = PRM/3 (NS, V4); 6 = PRM/3 (NS, V8); 7 = PRM/3PRM/3 (NS, V12); 8 = PRM/3/2 (M, V4); 9 = PRM/3/2 (M, V8); 10 = PRM/3/2 (M, V12); 11 = PRM/4 (D, V0); 12 = PRM/4 (NS, V4); 13 = PRM/4 (NS, V4); PRM/4 (NS, V4 PRM/4 (NS, V12); 14 = PRM/5 (D, V0); 15 = PRM/5 (NS, V4); 16 = PRM/5 (NS, V8); 17 = PRM/5 (NS, V12); 18 = PRM/5/1 (M, V4); 19 = PRM/5 (NS, V12); 14 = PRM/5 (D, V0); 15 = PRM/5 (NS, V4); 16 = PRM/5 (NS, V8); 17 = PRM/5 (NS, V12); 18 = PRM/5/1 (M, V4); 19 = PRM/5 (NS, V12); 14 = PRM/5 (NS, V12); 14 = PRM/5 (NS, V4); 16 = PRM/5 (NS, V8); 17 = PRM/5 (NS, V12); 18 = PRM/5/1 (M, V4); 19 = PRM/5 (NS, V12); 14 = PRM/5 (NS, V12); 18 = PRM/5/1 (M, V4); 19 = PRM/5 (NS, V12); 14 = PRM/5 (NS, V12); 18 = PRM/5/1 (M, V4); 19 = PRM/5 (NS, V12); 18 = PRM/5/1 (NS, V12); 18 = PRM/5/1 (NS, V12); 18 = PRM/5/1 (NS, V4); 19 = PRM/5 (NS, V12); 18 = PRM/5/1 (NS, V12); 18 = PRM/5/1 (NS, V4); 19 = PRM/5/1 (NS, V12); 18 = PRM/5/1 (NS, V12); 18 = PRM/5/1 (NS, V4); 19 = PRM/5/1 (NS, V12); 18 = PRM/5/1 (NS, V4); 19 = PRM/5/1 (NS, V12); 18 = PRM/5/1 (NS, V4); 19 = PRM/5/1 (NS, V12); 18 = PRM/5/1 (NS, V4); 19 = PRM/5/1 (NS, V12); 18 = PRM/5/1 (NS, V4); 19 = PRM/5/1 (NS, V4); 10 = PRM/5/1 (NS, V4); 1PRM/5/1 (M, V8); 20 = PRM/5/1 (M, V12); 21 = PRM/5/2 (M, V4); 22 = PRM/5/2 (M, V8); 23 = PRM/5/2 (M, V12); 24 = PRM/5/2 (TT, V₅0); 25 = PRM/5/2 (TT, V4); 26 = PRM/6 (D, V0); 27 = PRM/6 (NS, V4); 28 = PRM/6 (NS, V8); 29 = PRM/6 (NS, V12); 30 = PRM/6/2 (M, V4). PRM = cv. Premiant; D = donor (field grown) plants; NS = clones derived from nodal segments, M = mericlones; TT = mericlones subjected to *in vitro* thermotherapy; V0 = leaf sample from field-grown plants (control 1); V4, V8, V12 = leaf samples from *in vitro* plants after 4, 8 and 12



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