## **RESULTS AND DISCUSSION**

## 1. Isolation of Yeasts from Natural Habitats

Two hundred and eighty-three strains of yeasts were isolated from insect frass (144 strains), flowers (24 strains), leaves (54 strains), mosses (27 strains), mushrooms (28 strains), and some other habitats such as coconut juice (1 strain), exudates (1 strain), fruit (1 strain), lichens (1 strain), and rotted wood (2 strains) as detailed in Table 9. These samples were collected from various places in the natural habitats of Thailand (Table 10). Places of sample collections are shown in Appendix B Table B1. The yeast strains were purified by conventional streaking technique and maintained at -80°C at BIOTEC Culture Collection.

Year	Flowers	Insect frass	Leaves	Mosses	Mushrooms	Others	Total
2000	12	26	0	6	9	0	53
2001	2	104	54	4	18	0	182
2002	7	6	0	8	0	1	22
2003	3	8	0	9	1	5	26
Total	24	144	54	27	28	6	283

<u>Table 9</u> Number of yeasts isolated from various habitats.

Table 10 Number of yeasts isolated from four regions of Thailand.

Veen	Regions							
rear	Northern	Eastern	Central	Southern	Total			
2000	0	27	26	0	53			
2001	0	114	3	65	182			
2002	1	11	4	6	22			
2003	0	0	20	6	26			
Total	1	152	53	77	283			

## 2. Diversity of Yeasts Estimated Based on Sequences of D1/D2 Domains of 26S <u>rDNA</u>

Tentative identification was carried out based on the sequences of D1/D2 (ca. 600 nucleotides) domain of 26S rDNA according to a guideline of Kurtzman and Robnett (1998), 0-1 nucleotide differences are conspecific species, 2-3 nucleotide differences are conspecific or sister species and 4 or more nucleotide differences are different species. Undescribed species, whose D1/D2 sequences were registerd at DNA databanks, were regarded as known species. Yeast diversity of Thai natural habitats was discussed based on this tentative identification. The detailed taxonomic studies were carried out on selective groups in which new species were presumed by D1/D2 sequences. They were studied from the view point of polyphasic taxonomy; morphology, physiology, biochemistry, chemotaxonomy and molecular taxonomy.

Among 283 strains of yeasts isolated from flowers (24 strains), insect frass (144 strains), leaves (54 strains), mosses (27 strains), mushrooms (28 strains) and others (6 strains) tentatively identified, 194 strains belong to ascomycetous yeasts and 89 strains belong to basidiomycetous yeasts. Based on the sequence analysis of D1/D2 domain of 26S rDNA, 139 strains (49%) differed in 0-1 nucleotides from known species and were assigned to 56 known species of 26 genera including 39 species (97 strains) of 17 genera of ascomycetous yeasts and 17 species (42 strains) of 9 genera of basidiomycetous yeasts (Table 11). Details of identification are shown in Table 12. *Candida tropicalis* (9 strains) is the dominant species of ascomycetous yeasts and followed by *Saccharomyces kluyveri* (8 strains), *Saccharomyces cerevisiae* (5 strains), *Hanseniaspora opuntiae* (5 strains), *Metschnikowia koreensis* (5 strains) and *Debaryomyces nepalensis* (4 strains). In the case of basidiomycetous yeasts, *Cryptococcus heveanensis* (8 strains) is the dominant species and followed by *Exobasidium vexans* (4 strains), *Sporobolomyces* sp. (4 strains), and *Trichosporon asahii* (4 strains).



Figure 3 Places where new species were found.

- Places where sample were collected.
- Places where new species were found.

Species	No. of Strains	Species	No. of Strains
Ascomycetous yeasts			
Ambrosiozyma monospora	2	Saccharomyces cerevisiae	5
Aureobasidium pullulans	2	S. kluyveri	8
Blastobotrys catitulata	1	S. unisporus	2
Candida diversa	3	Stephanoascus smithiae	3
C. fukuyamaensis	3	Torulaspora delbrueckii	1
C. gotoi	2	Torulaspora sp. IFO 11061	1
C. leandrae	1	Wililopsis saturnus var. mrakii	1
C. natalensis	1	W. saturnus var. subsufficiens	1
C. palmae	1	Zygosaccharomyces sp. IFO 11070	1
C. parapsilosis	3	Total 17 genera, 39 species, 2 varities	97
C. rancensis	1		
C. sithepensis	1	Basidiomycetous yeasts	
C. tropicalis	9	Bullera dendrophila	1
Candida sp. NRRL Y-17456	4	Bullera sinensis	2
Candida sp. UWO(PS)00-147.3	1	Cryptococcus heveanensis	8
Debaryomyces nepalensis	4	C. humicola	2
D. polymorphus	3	C. laurentii	3
D. polymorphus var. africanus	1	Cryptococcus sp. CBS 8372	1
D. vanrijiae var. yarrowii	3	Exobasidium vexans	4
Debaryomyces sp. NRRL-7804	1	Rhodosporidium paludigenum	1
Geotrichum fragrans	2	R. toruloides	1
Hanseniaspora opuntiae	5	Rhodotorula nothofagi	1
H. vineae	1	Sporidiobolus ruineniae	4
Hanseniaspora sp. CBS 8772	1	Sporobolomyces bannaensis	1
Kloeckera lindneri	1	S. odoratus	3
Kluyveromyces lactis	2	S. poonsookiae	1
Kodamaea (Pichia) ohmeri	3	Sporobolomyces sp. TY-241 / TY-253	4
Metschnikowia koreensis	5	Tilletiopsis sp. TY-2352	1
Pichia nakazawae var. akitaensis	2	Trichosporon asahii	4
P. stipitis	2	Total 9 genera, 17 species	42
P. sydowiorum	2		
Pichia sp. UWO(PS)99-305.1	1		

Table 11Known species of ascomycetous and basidiomycetous yeasts isolated from<br/>some natural habitats of Thailand.