



UNIVERSITI PUTRA MALAYSIA

**MOLECULAR CHARACTERIZATION OF LISTERIA SPP.
ISOLATED FROM BEEF, CHICKEN AND FERMENTED
FISH IN MALAYSIA**

HAJJAH ENDANG PURWATI RAHAYUNINGSIH

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By

HAJJAH ENDANG PURWATI RAHAYUNINGSIH

**Thesis Submitted to the School of Graduate Studies, Universiti Putra
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Philosophy**

January 2003



DEDICATIONS

To my late mother, Rr. Soejatningsih ,
My late father, R. Soejono and my mother in law Hajjah Mursiati for their help
and prayers

To my husband, papa Dr. Mursof Fauzi Saladin MD, O & G
and my daughter Sofrida Prigustina Kurniasari
for their love, understanding, prayers and patience



Abstract of thesis presented to the Senate of Universiti Putra Malaysia in fulfillment of the requirement for the degree of Doctor of Philosophy.

MOLECULAR CHARACTERIZATION OF *LISTERIA* SPP. ISOLATED FROM BEEF, CHICKEN AND FERMENTED FISH IN MALAYSIA

By

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January 2003

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Faculty : Food Science and Biotechnology

Modified FDA method was found to give higher recovery of *Listeria* spp. than USDA method for different sources. The results indicated that the imported frozen beef samples from wet market examined were contaminated by seven different *Listeria* spp. namely, *L. monocytogenes*, *L. ivanovii*, *L. seeligeri*, *L. innocua*, *L. welshimeri*, *L. grayi* and *L. murrayi*. However, the use of FDA, USDA and the modified USDA methods may be more beneficial where a limited range of *Listeria* spp. to be recovered (*L. monocytogenes*, *L. ivanovii* and *L. innocua*). *Listeria* spp. was not isolated from any of the 23 samples of imported frozen beef from supermarket and from the five samples of buffalo meat examined.



A total of two hundred and seventy isolates of *Listeria* spp. from different sources were investigated for their susceptibility to 18 antibiotics and were screened for plasmid profiles. Antibiotic susceptibility analysis revealed that all *Listeria* spp. isolates were resistant to two or more antibiotic (MAR 0.11 to 0.66). Majority of the 52 isolates of *L. monocytogenes* displayed resistance to bacitracin, ceftazidime, nalidixic acid, sulfamethazole. However, none were resistant to norfloxacin. Plasmid was detected in 9 (81.8%) of 11 strains of *L. murrayi*, 12 (80%) of 15 strains of *L. grayi*, 29 (55.8%) of 52 strains of *L. monocytogenes*, 8(50%) of 16 strains of *L. denitrificans*, 25 (41.7%) of 60 strains of *L. ivanovii*, 25 (37.9%) of 66 strains of *L. innocua*, 13 (31%) of 42 strains of *L. welshimeri* and 1(12.5%) of 8 strains of *L. seeligeri*. The plasmid sizes ranged from 2.7 to 54 Kb.

In the conjugation study, the donor and the recipient were selected based on the antibiotic resistance pattern of selected *Listeria* spp. isolates from different sources. Seven *L. monocytogenes* strains and one *L. innocua* strain isolated from different sources were selected as donors since they were expected to potentially have an ability to conjugate due to the presence of high molecular weight plasmid DNA (54 Kb). Plasmidless sensitive to antibiotics are selected as recipients. Streptomycin resistance was transferred to *L. monocytogenes* LM65 and LM100 strains at frequencies of



3.3×10^{-8} and 1.2×10^{-9} per input donor cells. A selected kanamycin resistant *L. innocua* strains was found to transfer kanamycin resistant to *L. monocytogenes* (inter-and intra-species transfer).

The use of randomly amplified polymorphic DNA (RAPD) analysis for characterization and differentiation of the isolates of different *Listeria* spp. was examined. The combination of the RAPD-PCR patterns obtained with the three primers (Gen15001, Gen15002 and Gen15010) were able to distinguish all typeable isolates. Base on the dendrograms generated from the RAPD-PCR patterns all *Listeria* spp. isolates could be discriminated and clustered according to their species and food sources. RAPD fingerprinting methods had higher resolution than plasmid profiling and antibiotic resistance patterns and therefore the RAPD fingerprinting methods can be used for epidemiological studies.



Abstrak tesis yang dikemukakan kepada Senat Universiti Putra Malaysia
sebagai memenuhi keperluan untuk ijazah Doktor Falsafah

**PENCIRIAN SECARA MOLEKULAR KEATAS *LISTERIA* SPESIS YANG
DIPENCILKAN DARIPADA DAGING LEMBU, DAGING AYAM DAN IKAN
YANG DITAPAI DI MALAYSIA**

Oleh

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Kaedah pengubahsuaian FDA telah didapati memberikan pulangan yang tinggi untuk spesis *Listeria* berbanding kaedah USDA daripada sumber yang berbeza. Keputusan menunjukkan bahawa sampel daging lembu beku import yang diuji telah dicemarkan oleh tujuh spesis *Listeria* iaitu *L. monocytogenes*, *L. ivanovii*, *L. seeligeri*, *L. innocua*, *L. welshimeri*, *L. grayi* dan *L. murrayi*. Walau bagaimanapun, kegunaan kaedah FDA, USDA dan USDA diubahsuai mungkin lebih menguntungkan dimana spesis *Listeria* yang terhad boleh didapati (*L. monocytogenes*, *L. ivanovii* and *L. innocua*). Spesis *Listeria* tidak dapat dipencilkan daripada mana-mana 23 sampel daging lembu beku import dari supermarket dan 5 sampel daging kerbau yang dikaji.

Sejumlah dua ratus dan tujuh puluh isolat spesies *Listeria* daripada sumber yang telah diperiksa untuk kerektanan kepada 18 jenis antibiotik dan diskriminasi untuk mendapatkan profil DNA (asid deosiribonuklik) plasmid. Analisis kerektanan antibiotik menunjukkan bahawa kesemua isolat spesies *Listeria* adalah rentang kepada 2 atau lebih antibiotik (MAR 0.11 hingga 0.66). Kebanyakan 52 isolat *L. monocytogenes* mempamerkan kerintangan kepada basitrasin, septazidim, nalidixik asid, sulfamethazole. Walau bagaimanapun, tiada kerintangan kepada norfloxasin. Plasmid telah dikesan di dalam 9 (81.8%) dari 11 strain *L. murrayi*, 12 (80%) dari 15 strain *L. grayi*, 29 (55.8%) dari 52 strain *L. monocytogenes*, 8 (50%) dari 16 strain *L. denitrificans*, 25 (41.7%) dari 60 strain *L. ivanovii*, 25 (37.9%) dari 66 strain *L. innocua*, 13 (31%) dari 42 strain *L. welshimeri*, 1 (12.5%) dari 8 strain *L. seeligeri*, telah menunjukkan kehadiran DNA plasmid. Saiz plasmid berukuran dari 2.7 hingga 54 kb.

Dalam kajian konjugasi, penderma dan penerima telah dipilih berdasarkan corak kerentangan antibiotik spesies *Listeria* yang dipencilkan daripada sumber yang berbeza. Tujuh strain *L. monocytogenes* dan satu *L. innocua* yang dipencilkan daripada sumber berbeza telah dipilih sebagai penderma memandangkan mereka mempunyai potensi dan keupayaan

konjugasi disebabkan kehadiran berat molekul DNA plasmid yang tinggi (54kb) dan penerima mestilah tiada plasmid dan sensitif kepada antibiotik yang dipilih.

Penggunaan analisis DNA polimorfik penggandaan secara rawak (RAPD) untuk pencirian dan pembezaan isolat daripada spesies *Listeria* yang dikaji. Kombinasi corak fingerprinting RAPD yang diperolehi daripada 3 primer (Gen15001, Gen15002 and Gen15010) telah digunakan untuk pembezaan semua isolate. Daripada dendrogram yang digenerasi semua spesies *Listeria* boleh dibezakan dan dikumpulan bergantung kepada spesies dan sumber makanan mereka. Kaedah fingerprinting RAPD adalah lebih sensitif berbanding kepada corak kerintangan antibiotik. Oleh kerana itu, ia digunakan untuk kajian epidemiologi.



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This thesis submitted to the Senate of Universiti Putra Malaysia has been accepted as fulfilment of the requirements for the degree of Doctor of Philosophy. The members of the Supervisory Committee are as follows:

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DECLARATION

I hereby declare that the thesis is based on my original work except for quotations and citations which have been duly acknowledged. I also declare that it has not been previously or concurrently submitted for any other degree at UPM or other institutions.

Hajah Endang Purwati Rahayuningsih

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