
Classification for the Structure-Activity Relationship Methods

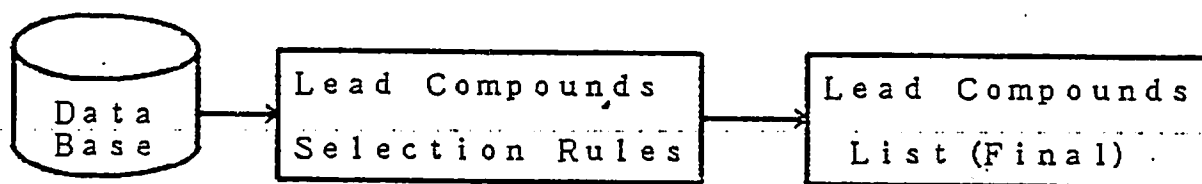
Based on a Lead Compound

- Lead Generation
- Lead Optimization
- Lead Evolution

Lead Retrieval
Lead Access

1. Lead 化合物検索について 日本薬学会 (1987年 4月)
2. ビリドンカルボ酸の抗菌活性データを用いて 新規の(Lead) 抗菌活性薬物を探し出す (MACCSのALDRICH データベースより)。

A Concept of the Lead Retrieval

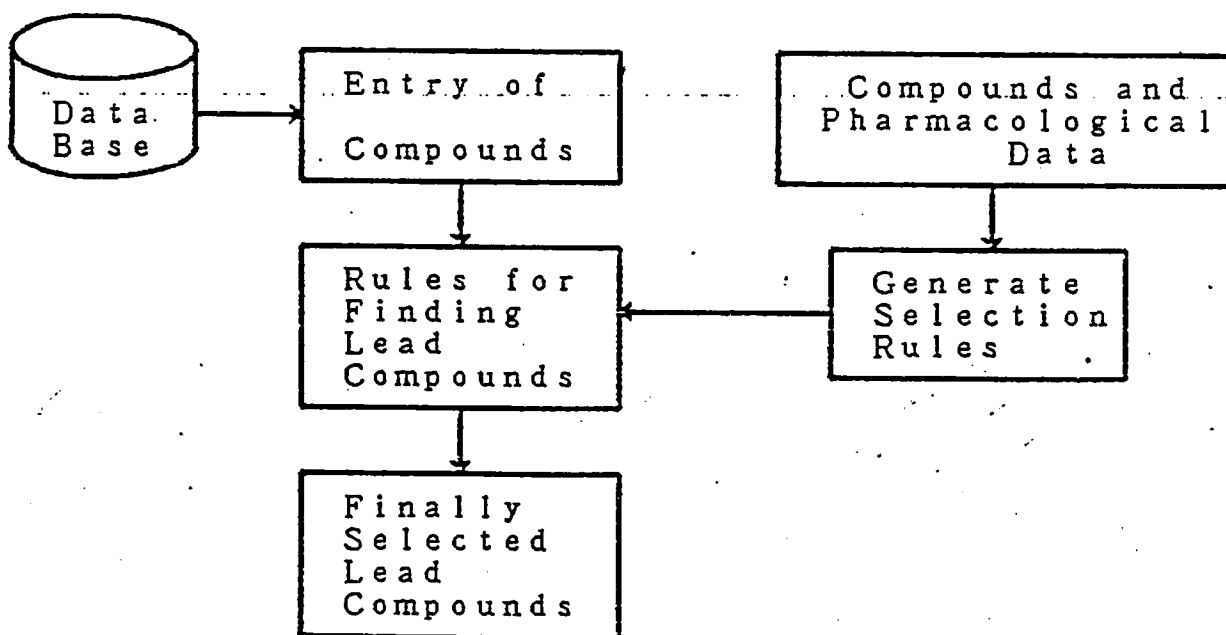


- In-house
 - Public
 - Hansch-Fujita Method
 - Pattern Recognition Method
 - Drug Receptor Theory
 - Calculate Similarity
 - Check to Double Register of the Same Compounds
- MACCS
CAS
ISI
CIS
DARC

Lead Generation
Lead Optimization
Lead Evolution

Lead Retrieval (Access)

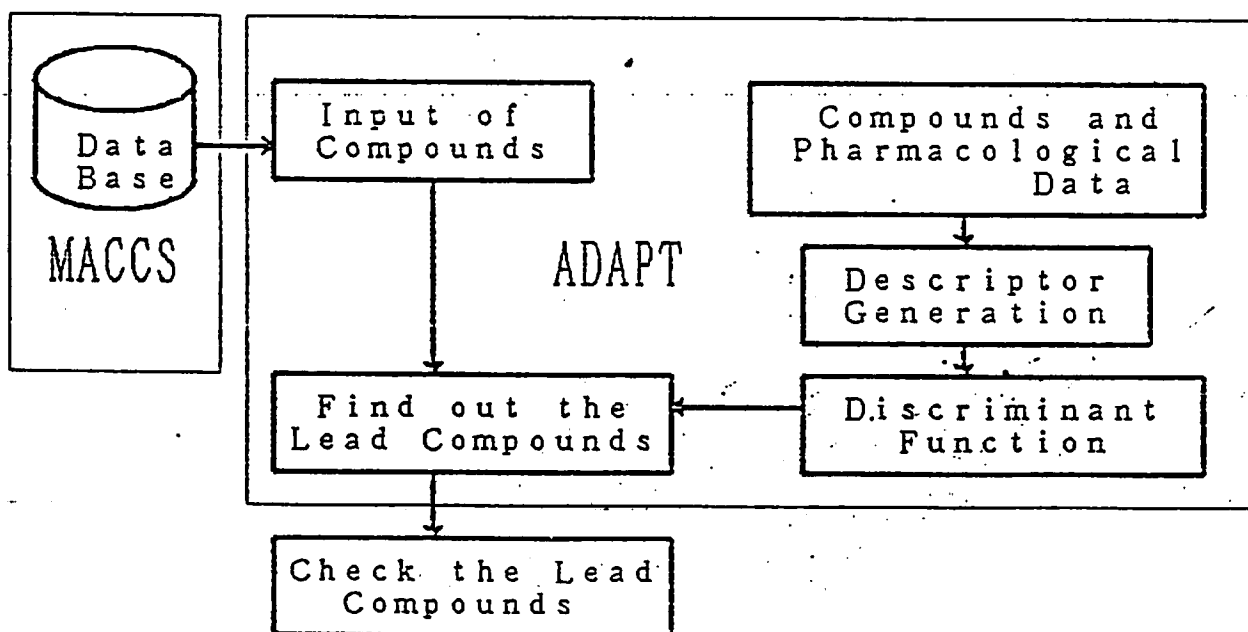
Flow Chart of the Lead Retrieval System



LEADS

Lead compounds Evaluation and
Access Development System

LEADS System for the Lead Retrieval

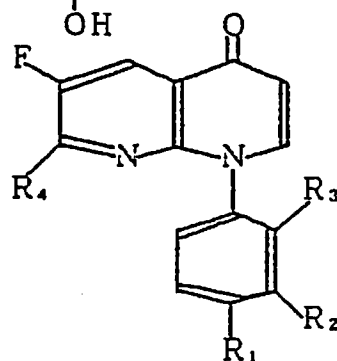
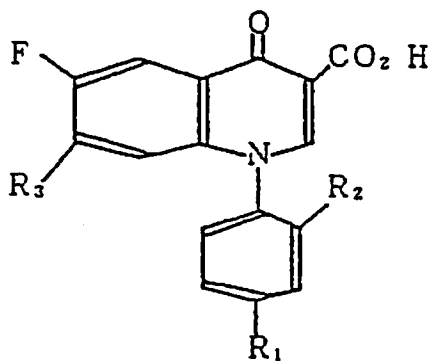
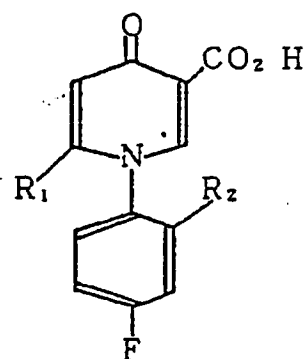
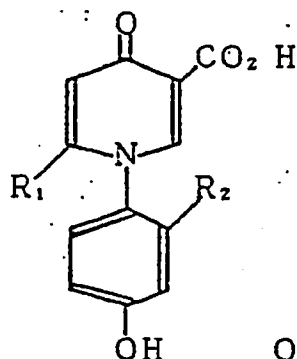
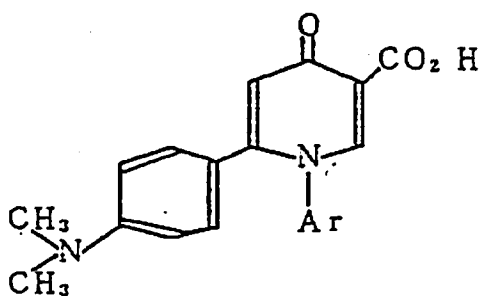


- Compound Data Base : ALDRICH in MACCS
- Finding Rule to the
New Lead Compounds: Discriminant Function
Generated by ADAPT
- * MACCS: Molecular Access System
- * ADAPT: Automated Data Analysis by Pattern
recognition Toolkits

Antibacterial Agents

4-Pyridone-3-carboxylic acid

H. Narita, Y. Konishi, J. Nitta, H. Nagaki, I. Kitayama
Y. Watanabe, and I. Saikawa, YAKUGAKU ZASSHI,
106, 775 (1986)



Used Compounds

Total 1 0 8 Compounds were Used for the
Pattern Recognition Study

5 6 Active Compounds

5 2 Not Active Compounds

Antibacterial Activity

Staphylococcus aureus FDA 209P

MIC $\mu\text{g/ml}$

0. 5 > Active compounds

0. 5 < Not active compounds

Used Descriptors

63 Descriptors were Generated by ADAPT

■ Various Feature Selection Methods were used to Select Descriptors

14 Descriptors were Finally Selected

1	FRAG	Number of Nitrogen Atoms
2	FRAG	Number of Aromatic Bonds
3	FRAG	Number of Ring Atoms
4	ALLP	Total Number of Paths (0~5)
5	ALLP	Total Number of Paths (20~46)
6	MOLC	Path-4 Molecular Connectivity
7	MOLC	Path-Cluster-4 Molecular Connectivity
8	BOMX	Maximum Bond Order
9	EDMX	Maximum Electron Density
10	LUMO	Lowest Unoccupied M. O.
11	FVMX	Maximum Free Valence Value
12	SNMN	Minimum Nucleophilic Superdelocalizability
13	STRA	Bond Strain Energy
14	STRA	Tortional Strain Energy

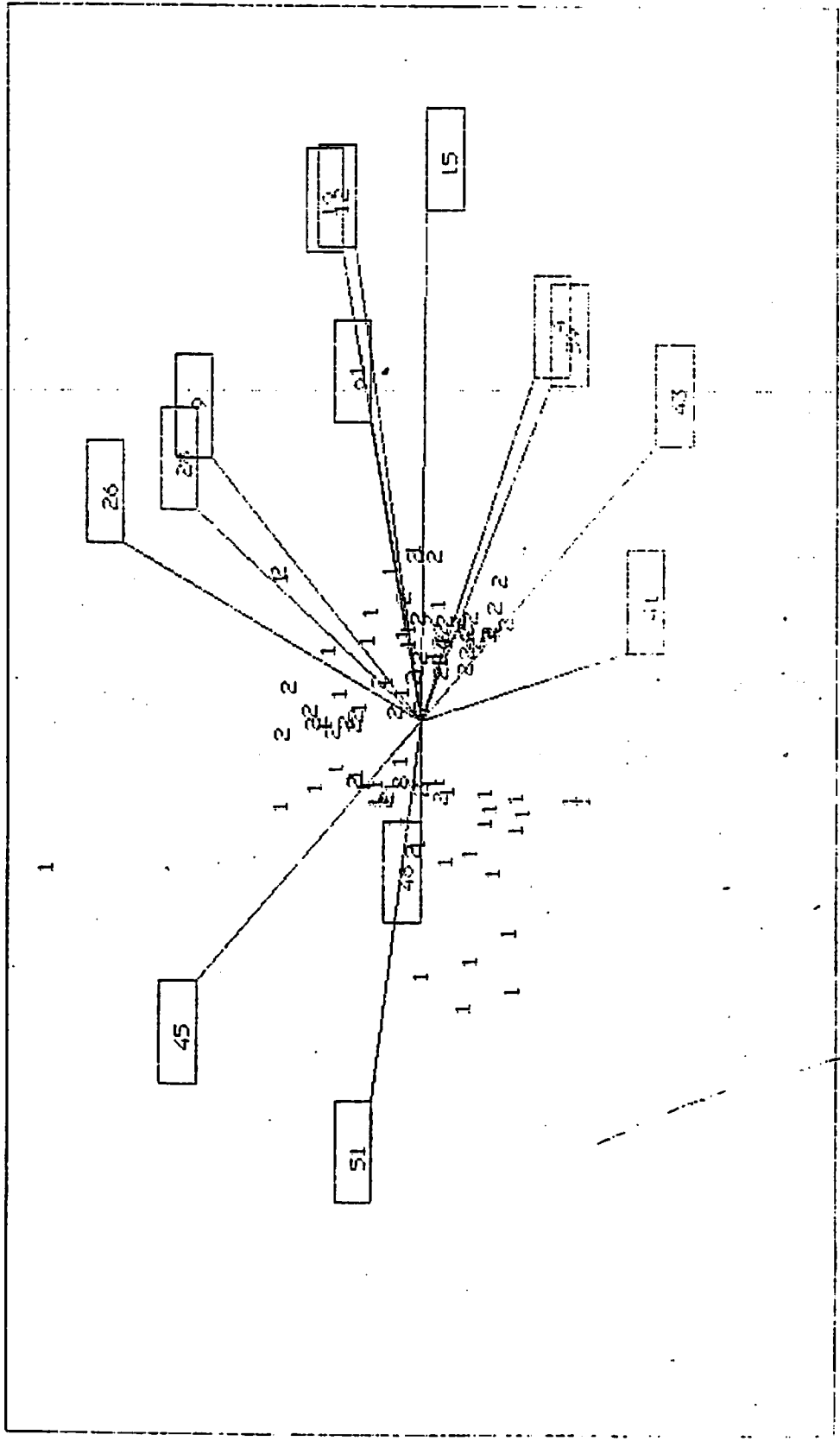
Results of Pattern Recognition Methods

Methods	Active	Not Active	Total
L. L. M	100. 0	100. 0	100. 0
SIMPLEX	94. 2	92. 0	93. 1
TILSQ	100. 0	100. 0	100. 0
BAQ	100. 0	82. 0	91. 2
BAL	94. 2	92. 0	93. 1
KNN			
K=1	80. 8	80. 0	80. 4
K=3	86. 5	78. 0	82. 4

* 6 Compounds were Removed from the
Initial (108 Compounds) Data Set

Active Class=50 (2 Compounds were Removed)
Not Active Class=52 (4 Compounds were Removed)
Total=102 Compounds

BIPLOT DISPLAY - AUTOSCALED DESCRIPTORS



FIRST PRINCIPAL AXIS

SECOND PRINCIPAL AXIS

Test Results by the Lead Retrieval System

Compounds

- 200 Compounds from the ALDRICH DATA-BASE
 Selected at Random

Discriminant Function

- Generated by the Linear
Learning Machine in ADAPT

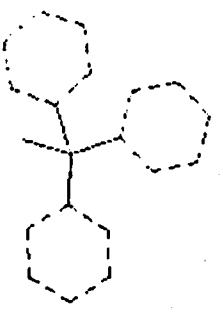
$$Y = 0.164W_1 - 0.052W_2 - 0.518W_3 + 0.193W_4 + 0.166W_5 \\ 0.245W_6 - 0.407W_7 + 0.506W_8 - 0.144W_9 - 0.098W_{10} \\ 0.112W_{11} - 0.077W_{12} - 0.081W_{13} - 0.315W_{14} + 0.019W_{N+1}$$

- 20 Compounds were Finally Selected as
the New Lead Compounds (Antibacterial Agents)

- Discriminant Values of the
New Lead Compounds

-1000 \geq Discriminant Value \geq -2000 16 Compounds
-2000 \leq Discriminant Value 4 Compounds

795 13705-7



754 BIS-(TRIPHENYLTIIN)

