

SS002-3

Qualitative structure-toxicity relationships (QSTR) on skin sensitization

ICOH Cancun Mexico 2012

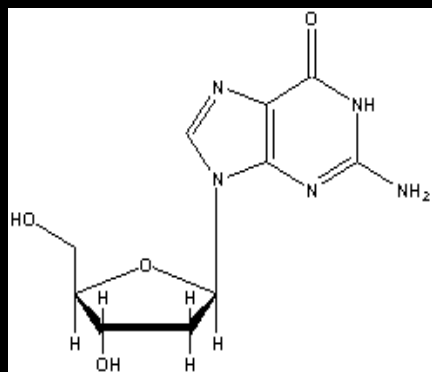
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1. Basic concept of QSTR approach
2. Sample and parameter handling
3. Data analysis and results by discriminant analysis
4. The KY-methods and conclusions

Basic concept of data analysis by multi-variate analysis and pattern recognition techniques



Compounds



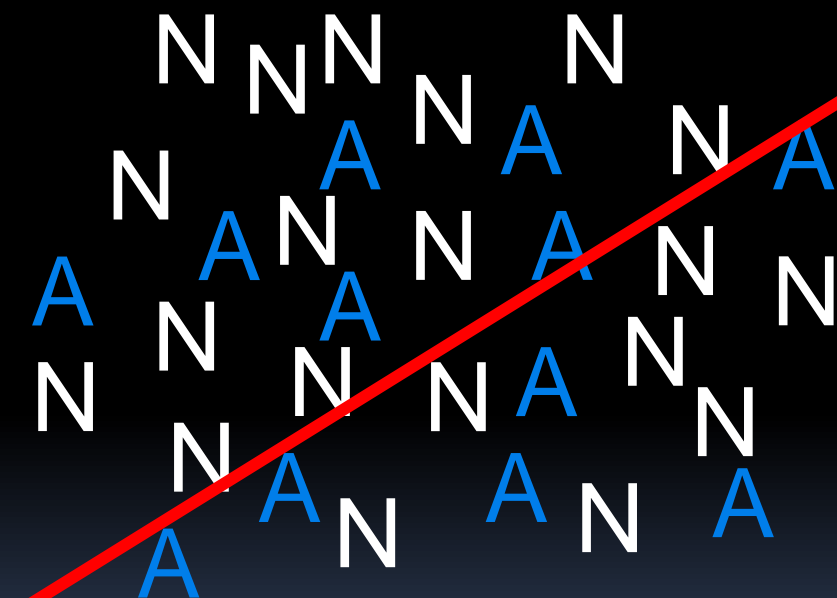
Information equivalence



Execute data analysis methods

Relations between pattern space and analytical objects

N-dimensional pattern space including noisy data



Linear discriminant

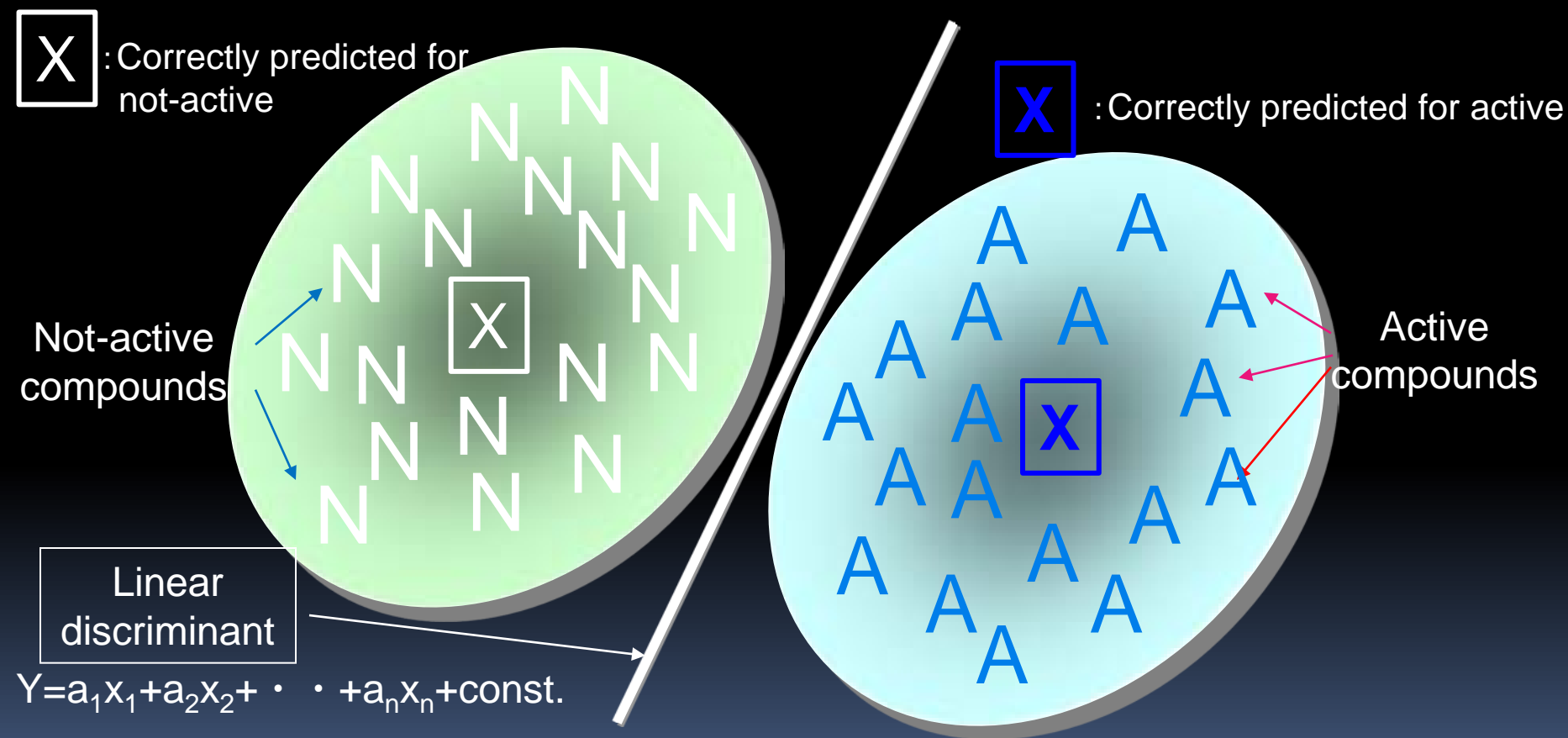
$$Y = a_1x_1 + a_2x_2 + \dots + a_nx_n + \text{const.}$$

No relations between
pattern space and objects

A; Active compounds
N; Not active compounds

Relations between pattern space and analytical objects

N-dimensional pattern space by intrinsic parameters \rightarrow Prediction



1. Pattern space divided into active and not-active compounds
2. This pattern space is classified by linear discriminant function

Style of discriminant function and regression equation

$$Y = +/- a_1x_1 +/- a_2x_2 +/- \cdot \cdot \cdot +/- a_nx_n +/- \text{const.}$$

Y : activity, ADME, toxicity, property

$$Y \geq 0$$

active or toxic

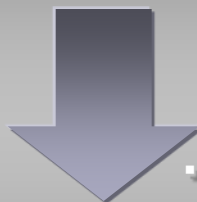
$$Y < 0$$

not active or non-toxic

Analysis of activity, ADME, toxicity or property

Coefficient $a_i \geq 0$
parameter X_i

·go **up** activity and toxicity



Coefficient $a_i < 0$
parameter X_i

·go **down** activity and toxicity

Structure-activity and Structure-toxicity relationships

1. Basic flow of QSAR approach
- 2. Sample and parameter handling**
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Used samples :
obtained from following 4 different databases

1. Maximale Arbeitsplatz-Konzentration (MAK)
..... > **positive** skin sensitizer
2. Biologischer Arbeitsstoff-Toleranz-Wert (BAT)
..... > **positive** skin sensitizer
3. Deutschen Forschungsgemeinschaft (DFG)
..... > **positive** skin sensitizer
4. Japanese Globally Harmonized System of Classification
and Labeling of Chemicals (GHS) Inter-ministerial Committee
of the National Institute for Technology and Evaluation
> **negative** skin sensitizer

Total **593** compounds
419 positive skin sensitizer
174 negative skin sensitizer

List of the used samples (Structure, CAS number, SMILES code, Sensitization)

C:\D-disc\福井大学\NITE156 and ROUSAI152 total307comp 3D WITH CAS SMILES with data.div

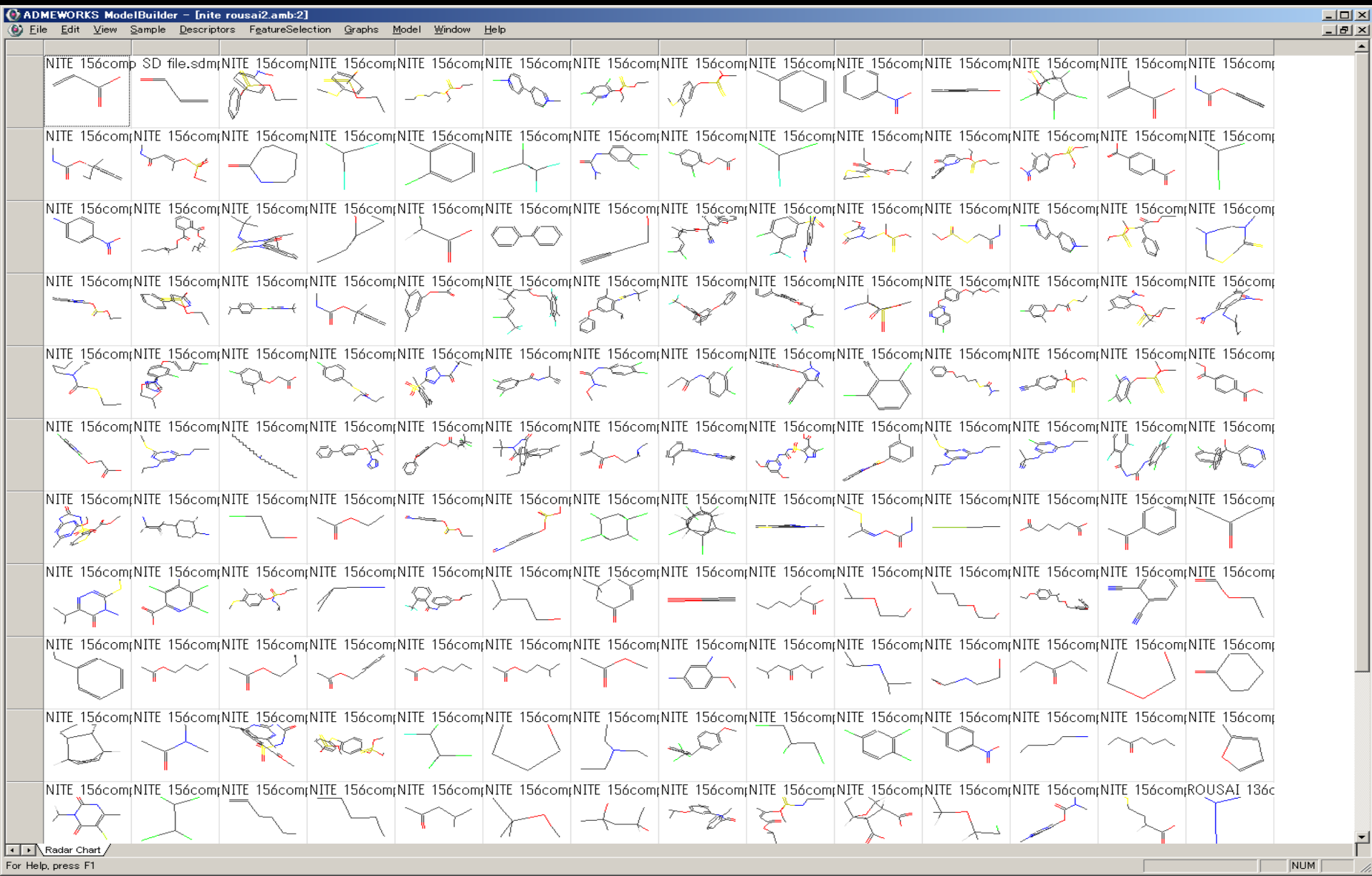
File Edit

	A Structure	B Name (Whole Molecule)	C CAS	D SMILES	E DFG	F ACGIH	G Skin	H Respirato			
1		NITE 156comp SD file.sd	000079-10-7	<chem>O=C(O)C=C</chem>	ND	ND	1	0			
2		NITE 156comp SD file.sd	000107-02-8	<chem>O=CC=C</chem>	ND	ND	1	0			
3		NITE 156comp SD file.sd	002104-64-5	<chem>CCOP(=S)(Oc1ccc(cc1)N(=O)=O)c</chem>	ND	ND	1	0			
4		NITE 156comp SD file.sd	035400-43-2	<chem>S=P(OCC)(SCCC)Oc1ccc(SC)cc1</chem>	ND	ND	1	0			
5		NITE 156comp SD file.sd	000298-04-4	<chem>CCOP(=S)(OCC)SCCSCC</chem>	ND	ND	1	0			
6		NITE 156comp SD file.sd	001910-42-5	<chem>Cn1(Cl)ccc(cc1)c2ccn(Cl)(C)cc2</chem>	ND	ND	1	0			
7		NITE 156comp SD file.sd	002921-88-2	<chem>CCOP(=S)(OCC)Oc1nc(Cl)c(Cl)cc1</chem>	ND	ND	1	0			
8		NITE 156comp SD file.sd	000055-38-9	<chem>COP(=S)(OC)Oc1ccc(SC)c(C)c1</chem>	ND	ND	1	0			
9		NITE 156comp SD file.sd	000108-88-3	<chem>c(cccc1)c1)C</chem>	ND	ND	1	0			
10		NITE 156comp SD file.sd	000098-95-3	<chem>N(=O)(=O)c(cccc1)c1</chem>	ND	ND	1	0			
11		NITE 156comp SD file.sd	000108-95-2	<chem>Oc(cccc1)c1</chem>	ND	ND	1	0			
12		NITE 156comp SD file.sd	000115-29-7	<chem>ClC2=C(Cl)C3(Cl)C1 COS(=O)OCC</chem>	ND	ND	1	0			
13		NITE 156comp SD file.sd	000079-41-4	<chem>O=C(O)C(=C)C</chem>	ND	ND	1	0			
14		NITE 156comp SD file.sd	000063-25-2	<chem>O=C(Oc(c(c(ccc1)cc2)c1)c2)NC</chem>	ND	ND	1	0			
15		NITE 156comp SD file.sd	003766-81-2	<chem>O=C(Oc(c(ccc1)C(CC)C)c1)NC</chem>	ND	ND	1	0			
16		NITE 156comp SD file.sd	006923-22-4	<chem>CNC(=O)C=C(C)OP(=O)(OC)OC</chem>	ND	ND	1	0			

Set 2: Table Set 2: Report

Data Size: 307 rows, 8 columns No Valid Selection 0 jobs pending

List of 3-Dimensional Structures of the used compounds



List of Compounds and Generated 822 Parameters

ADMEWORKS ModelBuilder - [samples and starting set.amb]

File Edit View Sample Descriptors FeatureSelection Graphs Model Window Help

Sample set: Training Parameter set: All

	0.37	822	173/131	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>	<input checked="" type="checkbox"/>
	304/304	Item	Skin_CL	NATM	NC	NO	NN	NS	NF
<input checked="" type="checkbox"/>	1	NITE 156comp SD file.sd	0	10	8	2	0	0	
<input checked="" type="checkbox"/>	2	NITE 156comp SD file.sd	0	7	5	2	0	0	
<input checked="" type="checkbox"/>	3	NITE 156comp SD file.sd	0	8	6	2	0	0	
<input checked="" type="checkbox"/>	4	NITE 156comp SD file.sd	0	28	25	3	0	0	
<input checked="" type="checkbox"/>	5	NITE 156comp SD file.sd	0	10	8	0	2	0	
<input checked="" type="checkbox"/>	6	NITE 156comp SD file.sd	0	5	3	2	0	0	
<input checked="" type="checkbox"/>	7	NITE 156comp SD file.sd	0	9	9	0	0	0	
<input checked="" type="checkbox"/>	8	NITE 156comp SD file.sd	0	8	6	2	0	0	
<input checked="" type="checkbox"/>	9	NITE 156comp SD file.sd	0	8	6	2	0	0	
<input checked="" type="checkbox"/>	10	NITE 156comp SD file.sd	0	11	9	2	0	0	
<input checked="" type="checkbox"/>	11	NITE 156comp SD file.sd	0	9	7	2	0	0	
<input checked="" type="checkbox"/>	12	NITE 156comp SD file.sd	0	9	7	2	0	0	
<input checked="" type="checkbox"/>	13	NITE 156comp SD file.sd	0	5	3	2	0	0	
<input checked="" type="checkbox"/>	14	NITE 156comp SD file.sd	0	10	7	1	2	0	
<input checked="" type="checkbox"/>	15	NITE 156comp SD file.sd	0	10	9	1	0	0	
<input checked="" type="checkbox"/>	16	NITE 156comp SD file.sd	0	7	6	0	1	0	
<input checked="" type="checkbox"/>	17	NITE 156comp SD file.sd	0	7	4	2	1	0	
<input checked="" type="checkbox"/>	18	NITE 156comp SD file.sd	0	6	5	1	0	0	
<input checked="" type="checkbox"/>	19	NITE 156comp SD file.sd	0	5	3	2	0	0	
<input checked="" type="checkbox"/>	20	NITE 156comp SD file.sd	0	7	6	1	0	0	
<input checked="" type="checkbox"/>	21	NITE 156comp SD file.sd	0	10	10	0	0	0	
<input checked="" type="checkbox"/>	22	NITE 156comp SD file.sd	0	25	12	6	5	2	
<input checked="" type="checkbox"/>	23	NITE 156comp SD file.sd	0	17	15	0	2	0	
<input checked="" type="checkbox"/>	24	NITE 156comp SD file.sd	0	4	2	1	0	0	
<input checked="" type="checkbox"/>	25	NITE 156comp SD file.sd	0	6	4	2	0	0	
<input checked="" type="checkbox"/>	26	NITE 156comp SD file.sd	0	18	10	5	1	1	
<input checked="" type="checkbox"/>	27	NITE 156comp SD file.sd	0	16	8	5	1	1	
<input checked="" type="checkbox"/>	28	NITE 156comp SD file.sd	0	12	6	0	0	0	

For Help, press F1

NUM

822 parameter generation from structure of compound and final 60 parameter set after feature selection process

Structure of compounds



Generate Parameters



Various feature selections



Final parameter set
(Important for used skin sensitization sample set)

Total **593** compounds

419 positive skin sensitizer

174 negative skin sensitizer

Total **822** parameters per compound

- topological (2-D) parameters
MC parameters, etc..
- topographical (3-D) parameters
Box parameters, etc..
- property parameters
LogP, MR, Volume, Surface, etc..
- electric parameters
HOMO, LUMO, etc..
- substructure parameters
Count of substructures, etc..

Final **60** parameters

1. Basic flow of QSAR approach
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List of Classification results by Various Discriminant Analysis

1. NN (Neural Network) :60 Parameter set

Classification ratio : 85.5%

2. Linear discriminant analysis by least squares algorithms

Classification ratio : 85.7%

3. SVM(Support Vector Machine) :60 Parameter set

Classification ratio : 90.7%

4. ADA Boost: 60 Parameter set

Classification ratio : 77.5%

5. **KY-method** for Discriminant analysis

Classification ratio : 100% (Perfect Classification)

**Perfect
classification**

Incomplete classification example by the AdaBoost (77.5%)

ADMEWORKS ModelBuilder - [Lazy Classification test work3:2]

File Edit View Sample Descriptors FeatureSelection Graphs Model Window Help

Model name: Skin_CL_ADA_9

Sample Dot Map Zoom

Number of training : 50
 Percent features in descriptor set : 75.00

Percent correct sample : 78.939
 Number of samples : 622
 Number of parameters : 12
 Number of models : 50

----- TRAINING SET CLASSIFICATION -----

Class	Members	Wrong	%Correct
neg	176	91	48.30
pos	446	40	91.03
Total	622	131	78.94

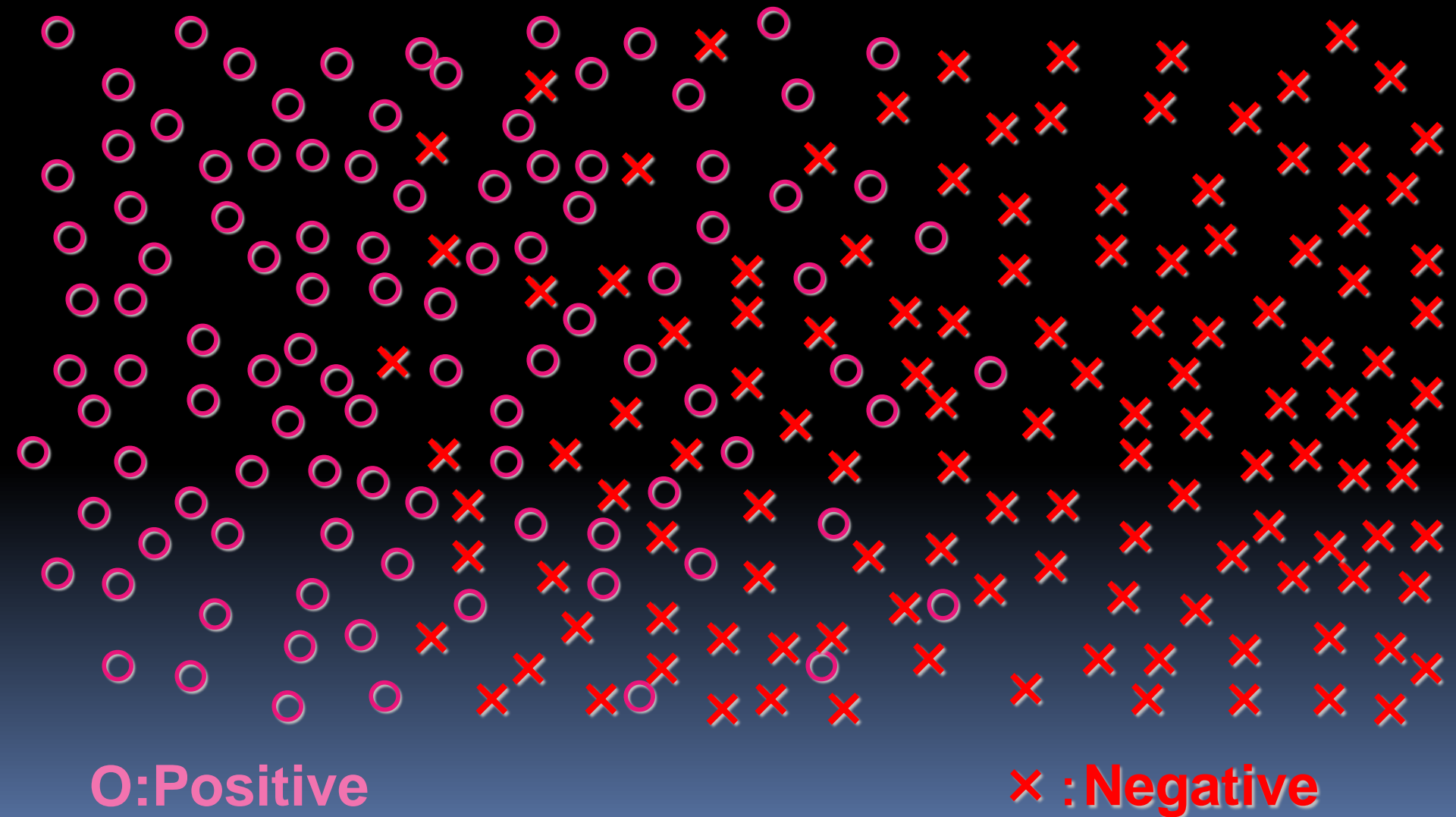
◆ R2

State	Model Name	% correct	Alpha
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_1	73.7942...	0.449625
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_2	69.6141...	0.291125
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_31	68.1672...	0.088942
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_16	68.0064...	0.005956
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_3	68.0064...	0.228313
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_41	67.8456...	0.093781
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_24	63.6655...	0.058757
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_8	63.5048...	0.067233
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_49	61.7363...	0.101501
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_20	61.7363...	0.089913
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_9	61.2540...	0.067299
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_17	60.9224...	0.076124

Recalculate Select plots ... Export iterations stat. Cancel OK

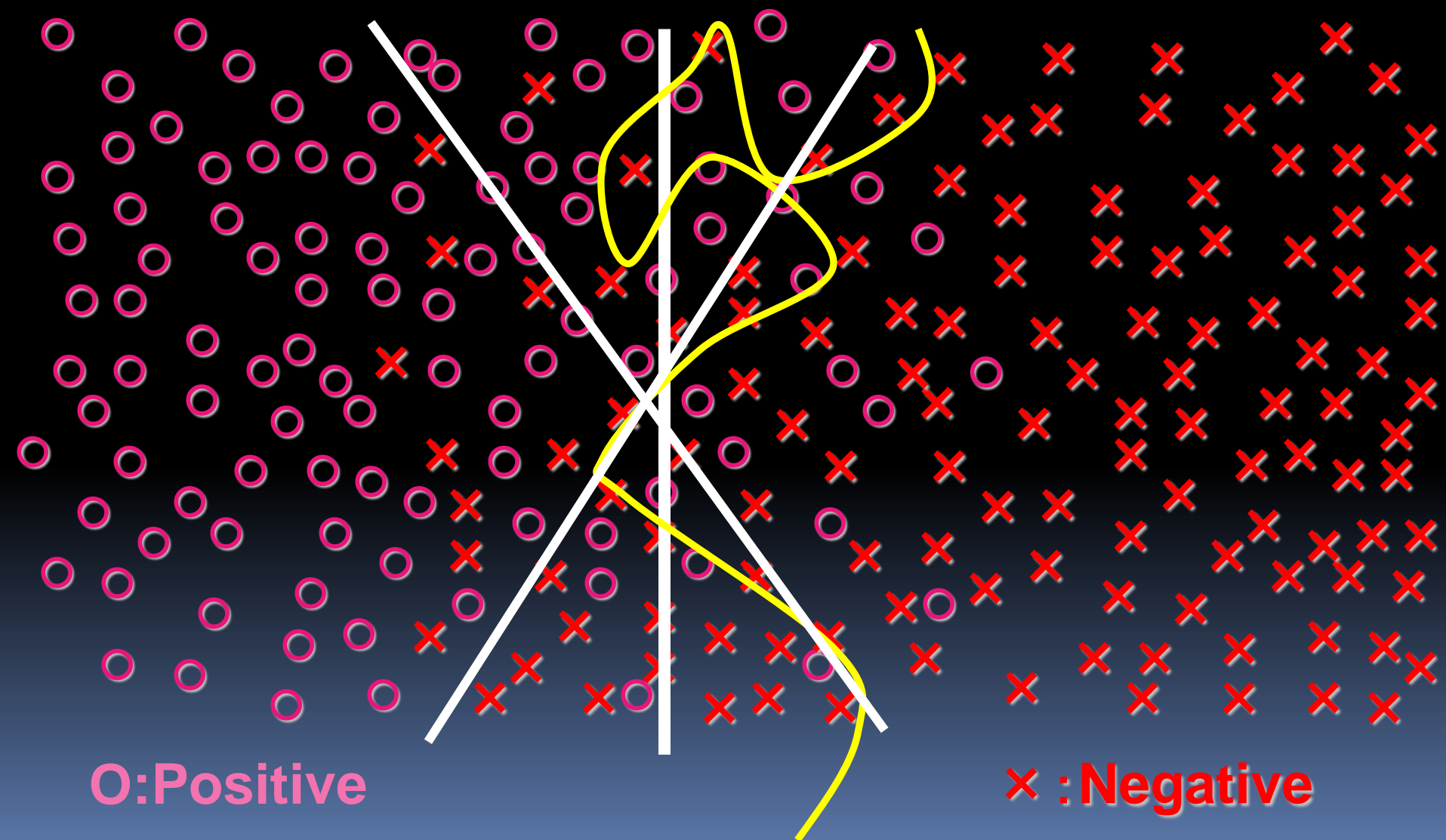
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Sample space : Highly overlapped space



Sample space : Highly overlapped space

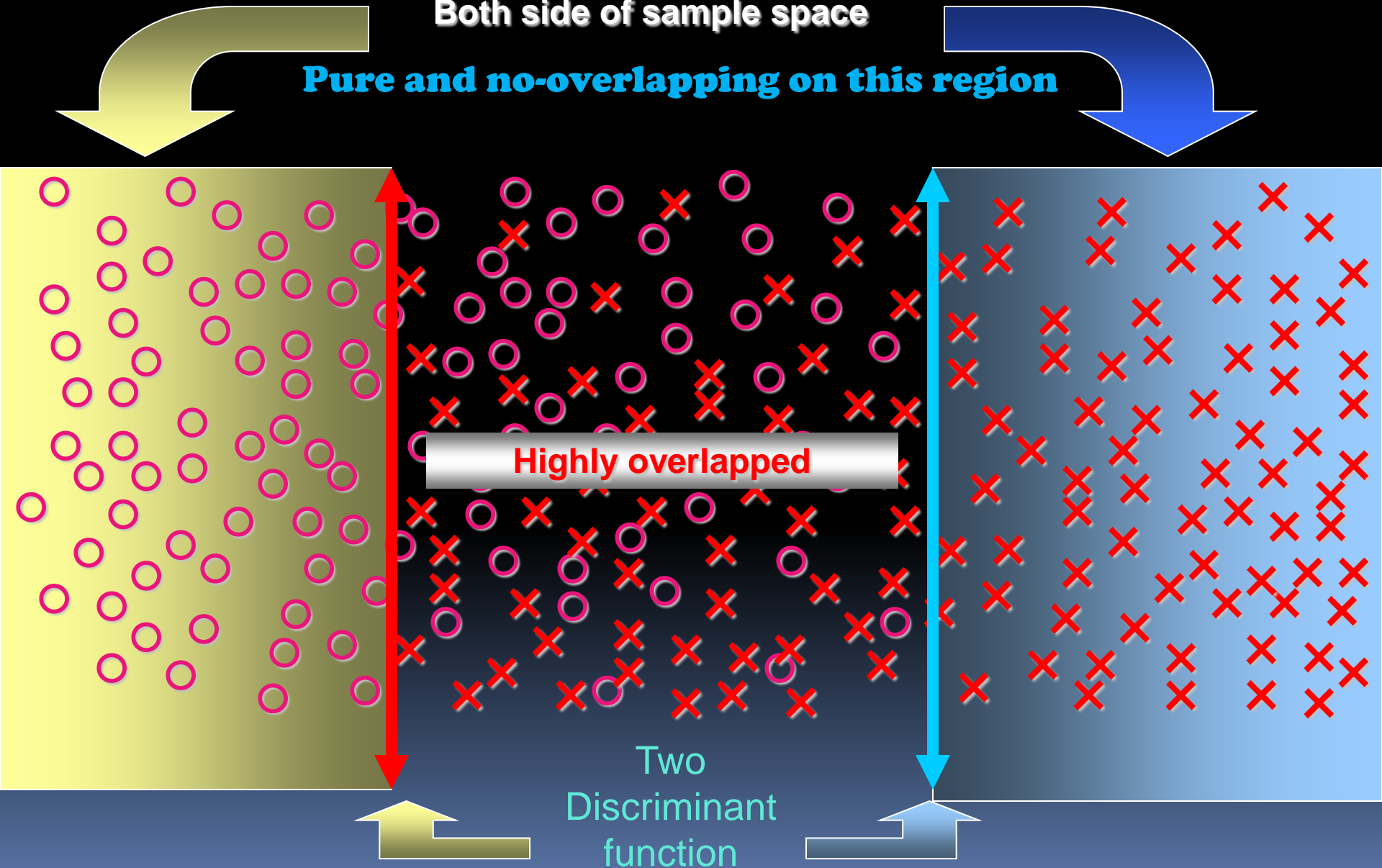
Discriminant function : Linear and non-linear



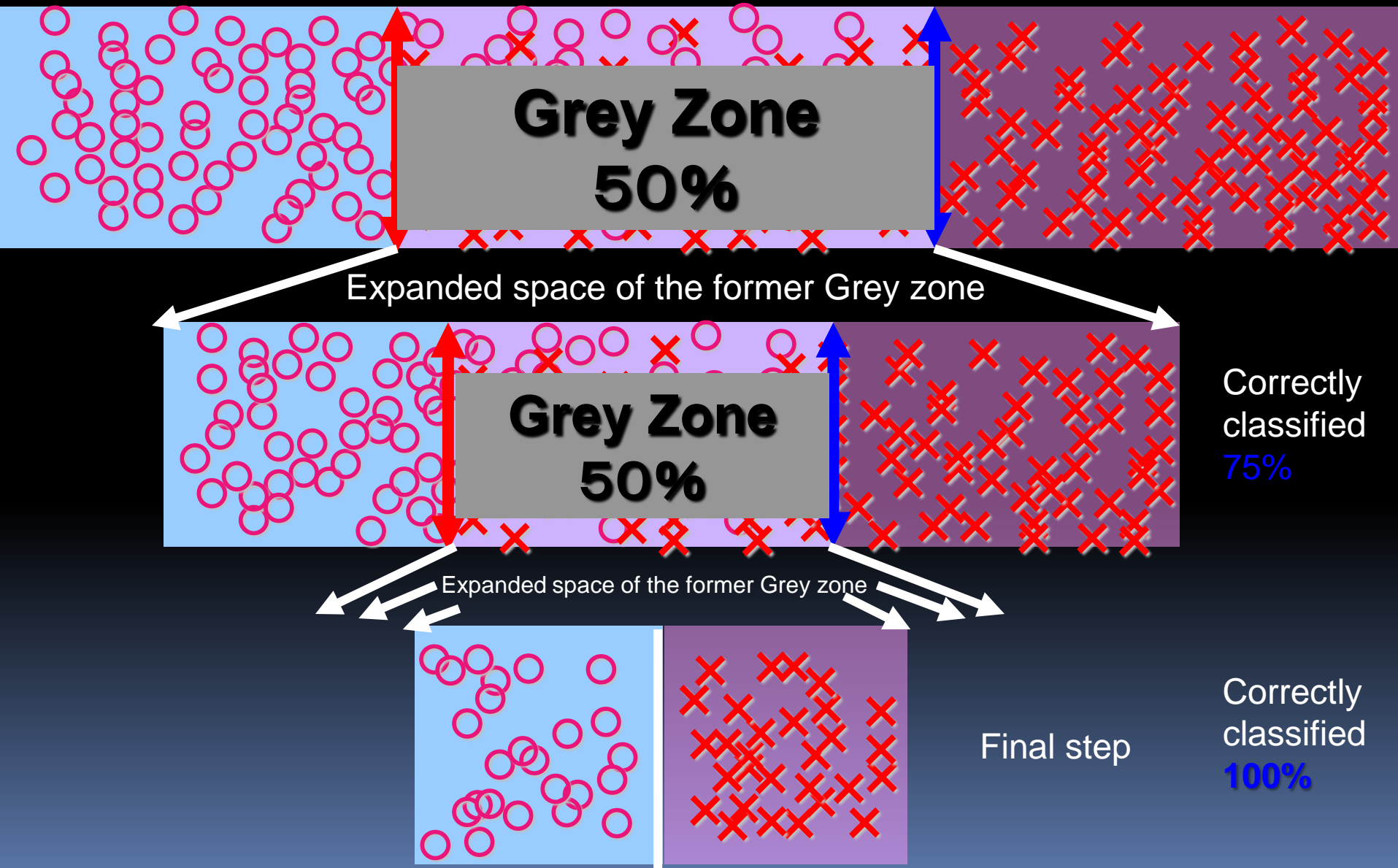
Spatial region on sample space

Both side of sample space

Pure and no-overlapping on this region



Steps to the K-step methods



Perfect classification example by the KY-method (Displayed by the AdaBoost)

ADMEWORKS ModelBuilder - [fukui2010-593all 174neg 419pos step1 AN and APmodels only:2]

File Edit View Sample Descriptors FeatureSelection Graphs Model Window Help

Model name: Skin_CL_ADA_1

Sample Dot Map Zoom

Number of training : 50
Percent features in descriptor set : 75.00

Percent correct sample : 100.000
Number of samples : 361
Number of parameters : 69
Number of models : 24

----- TRAINING SET CLASSIFICATION -----

Class	Members	Wrong	%Correct
neg	174	0	100.00
pos	187	0	100.00
Total	361	0	100.00

◆ R2

State	Model Name	% correct	Alpha
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_12	94.1828...	1.129055
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_1	93.3517...	1.147419
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_15	89.1966...	1.126181
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_7	88.6426...	0.915023
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_10	87.5346...	0.830803
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_4	86.1495...	0.937807
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_2	85.8725...	0.888954
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_3	85.5955...	0.836549
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_9	84.7645...	0.787388
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_13	83.9335...	1.136771
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_8	83.9335...	0.773107
<input checked="" type="checkbox"/>	Skin_CL_FLDRFS_6	82.8888...	0.719352

Recalculate Select plots ... Export iterations stat. Cancel OK

Spatial features of the “KY-methods”

Always achieve perfect classification

- (a) Even if **the number of samples becomes very large**, the KY-methods achieves perfect (100%) classification
- (b) Even if **overlapped sample space grows too big**, the KY-methods achieves perfect (100%) classification

Differences between the KY-methods and the ordinal methods

- 1. Number of classified sample zone:
KY-methods ; three zones Ordinal methods ; two zones
- 2. Repeat number of classification:
KY-methods ; ≥ 2 times Ordinal methods ; 1 time

Patented: US 7,725,413

Patent pended: Japan, Korea, EU

SS002-3

**Thank you for your
kind attention**

ICOH Cancun Mexico 2012

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