



Sensor/Marker Selection for Diagnosis based on a Fuzzy Feature Selection Approach

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→ problem of sensor selection for industrial process diagnosis



I. Introduction 2



- A novel methodology enables to handle simultaneously both problems regardless of their own characteristics:
- → Copes with the problem of high dimensionality based on classical optimization methods.
- → Handles appropriately heterogeneous data (quantitative, qualitative, interval).

Relevant example: interval representation of data can improve classification processing of clinical data in medical diagnosis as well as to process noisy or uncertain industrial measurements.

- Application to derivation of hybrid markers for cancer prognosis.

- Application to sensor selection for fault diagnosis of pharmaceutical synthesis process in a new intensified heat-exchanger reactor.







- Membas enables to process three types of features: Quantitative, Qualitative and Interval.
- In Fuzzy Logic framework, Simultaneous mapping can be performed through a Feature Fuzzification according to each feature type.
- Based on a data-driven procedure, ℓ fuzzy partitions { $mff_1^i, ..., mff_\ell^i$ } are obtained for the *i*th feature to each existing class *k* :

$$mff_k^i = \mu_k^i (\mathbf{x}_i, \, \boldsymbol{\theta}_{ki})$$

7 Quantitative type features (similarity semantic):

$$\mu_{k}^{i} \left[x_{i} \left| \rho_{k}^{i}, \varphi_{k}^{i} \right] = \varphi_{k}^{i^{1-\left|x_{i}-\rho_{k}^{i}\right|}} \left(1 - \varphi_{k}^{i} \right)^{\left|x_{i}-\rho_{k}^{i}\right|} \text{ where } \varphi_{k}^{i} = \frac{1}{m_{k}} \sum_{j=1}^{j=m_{k}} x_{i}^{j}$$



II.1 SM: Simultaneous Mapping



Interval type features (similarity semantic):

$$S(A,B) = \frac{1}{2} \left(\frac{\varpi[A \cap B]}{\varpi[A \cup B]} + 1 - \frac{\partial[A,B]}{\varpi[U]} \right)$$

where $\partial [A, B] = \max \left[0, \left(\max \left\{ a^{-}, b^{-} \right\} - \min \left\{ a^{+}, b^{+} \right\} \right) \right]$ and $\varpi [X] = upperbound(X) - lower.bound(X)$ Therefore, $\mu_{k}^{i} (\chi_{i}) = S (\chi_{i}, \rho_{k}^{i})$

where

$$\rho_k^{i-} = \frac{1}{m_k} \sum_{j=1}^{m_k} x_i^{j-}$$
 and $\rho_k^{i+} = \frac{1}{m_k} \sum_{j=1}^{m_k} x_i^{j+}$

Qualitative type features (Uncertainty semantic):

$$\mu_k^i\left(\chi_i\right) = \left(\Phi_{k1}^i\right)^{q_{i1}} * \cdots * \left(\Phi_{kMi}^i\right)^{q_{iMi}}$$

Where Φ_{kM}^{i} is the frequency of the Mth modality in the class C_{k} and $q_{j}^{i} = \begin{cases} 1 & if \quad x_{i} = Q_{j}^{i} \\ 0 & if \quad x_{i} \neq Q_{j}^{i} \end{cases}$





Let $D = \{x_n, C_k\}_{n=1}^{N} \in X \times C$ be a dataset, where N is the number of patterns (items) and $x_n = [x_{n1}, x_{n2}, \dots, x_{nm}]$ is the nth pattern.

A natural result of the previous fuzzification step is a common membership space for heterogeneous features, i.e. a Membership Degree Vector (MDV) of pattern to each class:

$$\mathbf{U}_{nc_{k}} = \left[\mu_{k}^{1}(\mathbf{x}_{n1}), \ \mu_{k}^{2}(\mathbf{x}_{n2}), ..., \ \mu_{k}^{m}(\mathbf{x}_{nm})\right]^{\mathrm{T}}; \ k = 1, 2, ..., l$$

where $\mu_{k}^{i}(x_{ni}) = \mu_{k}^{i}(x_{i} = x_{ni})$



II.2 SP: MEMBAS for binary class problems



A membership margin is defined for each pattern $x_n \in c$:

 $\beta_{n} = \psi(U_{nc}) - \psi(U_{n\tilde{c}})$

- Note $\Psi(U_{nc_k}) = \sum_{i} \mu_k^i(x_{ni})$ is the scalar cardinality of the fuzzy subsets described by MDVs.
- *¬* Pattern x_n is considered correctly classified if $β_n > 0$.
- ↗ A weighted membership margin can be defined as:

$$\beta_n = \psi(\mathbf{U}_{\mathrm{nc}} / \mathbf{W}_{\mathrm{f}}) - \psi(\mathbf{U}_{\mathrm{n\tilde{c}}} / \mathbf{W}_{\mathrm{f}})$$



II.2 SP: MEMBAS for binary class problems



A margin-based objective function has been defined so that the averaged membership margin in the resulted weighted membership space is maximized:

$$\begin{aligned} & \underset{w_{f}}{\text{Max}} \sum_{n=1}^{N} \beta_{n}(w_{f}) = \sum_{n=1}^{N} \{ \sum_{i=1}^{m} w_{fi} \mu_{c}^{i}(x_{ni}) - \sum_{i=1}^{m} w_{fi} \mu_{\tilde{c}}^{i}(x_{ni}) \} \\ & \text{s.t. :} \|w_{f}\|_{2}^{2} = 1 \text{ , and } w_{f} \ge 0 \end{aligned}$$

A closed-form solution using Lagrangian

$$w_{f}^{*} = \frac{s^{+}}{||s^{+}||}$$

where
$$s = \sum_{n=1}^{N} \{ U_{nc} - U_{n\widetilde{c}} \}$$

with $s^+ = [\max(s_1, 0), ..., \max(s_m, 0)]^T$







IV. Marker selection for cancer prognosis



- An accurate cancer prognosis is needed to help physicians select optimal treatment and reduce its related expensive medical costs.
- Justical or genes markers are used to perform the prognosis.
- **Integration of both information may improve the prognosis**
- 7 Two challenges are faced: High-dimension and heterogeneous data

- The first due to the presence of a large amount of irrelevant genes in microarray data

- The second is related to the presence of mixed-type data (quantitative, qualitative and interval) in the clinical data



Experiments and Results



G Pronostic du Métastase distant : Netherlands Cancer Institute

- + 295 breast cancer patients
- + 29 patients with missing gene expression excluded from the study.
- + 2 classes according to the appearance of distant metastases: 88 patients with and 207 patients without.
- + Training D.(132): 92 without, 40 with; Test D. (134): 93 without, 41 with.
- + Microarray dataset: 24188 gene expression
- + Clinical dataset: 10 variables 1 of quantitative type; 1 intervallaire et 8 qualitatives
 - + Age (quantitative)
 - + Tumour grade (interval :[3,5]; [6,7]; [8,9])
 - + Tumour size = T (qualitative: ≤2cm; >2cm)
 - + Nodal status = N (qualitative : pN0; '1-3'; \geq 4)
 - + Mastectomy (qualitative : Yes, No)
 - + Estrogen Receptor ER expression (qualitative : Yes, No)
 - + Chemiotherapy (qualitative: Yes, No)
 - + Hormonotherapy (qualitative: Yes, No)
 - + St. Gallen European criteria (qualitative: Chemio , No Chemio)
 - + NIH US criteria (qualitative: Chemio , No Chemio
 - + Risk NIH (qualitative: low , intermediate , high)



Experiments and Results

So **Derived hybrid signature:** MEMBAS selects only 15 hybrid markers

- Three are mixed-type clinical markers (Number of positive lymph nodes "qualitative", ER "qualitative" and Grade "interval"), added to them 12 genes. (optimal Classif. Performance).

G Comparatives results between hybrid, clinical, genetic signatures and classical clinical indices:

	TP	FP	FN	TN	Sens.	Specif.	Accuracy
Hybrid	13	12	28	81	0.32	0.87	94/134 (70.15%)
70-genes	25	29	16	64	0.61	0.69	89/134 (66.42%)
Clinical	23	37	18	56	0.56	0.60	79/134 (58.96%)
NIH	41	91	0	2	1	0.02	41/134 (32.09%)
St Gallen	38	85	3	8	0.93	0.09	46/134 (34.33%)

- St. Gallen Chimio recommandée quand un critère est vrai : ER negatif; ganglions positif; T>2cm ; Grade III ou II ; Age <35 ans.</p>
- - TP: True Positive ; FP: False Positive ; FN: False Negative ; TN: True Negative ; Sens.: Sensitivity; Specif.: Specificity.





- The success of fault detection and diagnosis of complex process depends strongly on the selection of measurements that characterize accurately the process behavior.
- In Large number of sensor increases the induced instrumentation cost and may degrades the diagnosis efficiency.
- **> Efficient sensor selection methodologies are required that:**
- → Monitor accurately and robustly fault detection in complex processes
- → While, assure a reduction in the instrumentation costs and improve the process safety and quality





- 1) Fault identification using the fuzzy classification technique LAMDA (self-learning) ^(*)
- 2) Sensor selection based on MEMBAS method (*) (**)
- 3) Generation of behavioral pattern of the process based only on the selected set of sensors.
- 4) Online recognition and validation on unseen data.
- (*) Implemented on SALSA software tool [T. Kempowsky et al., 2003].
- (**) Validated in an extensive experimental study on a large number of high dimensional and heterogeneous datasets [Hedjazi et al., 2010].



Application on chemical process LAAS-CNRS

Pharmaceutical synthesis in a new intensified heat exchanger reactor equipped of 15 sensors: 12 internal temperatures (interval), Utility outlet temperature (interval), Reacts. Pressure (quantitative).

$$A_{(aq)} + B_{(l)} \to C_{(l)} + D_{(l)}$$
$$B_{(l)} + E_{(l)} \to 2D_{(l)}$$







Application on chemical process

LAAS-CNRS



Class Number	CLASS DESCRIPTION					
1	Steady state					
2	Increased (1) flow reactant B					
3	Critical Increased (♠) flow reactant B					
4	Increased (1) flow reactant A					
5	Decreased (\clubsuit) flow reactant A					
6	Shut-down flow reactant A					
7	Shut-down utility flow					
8	Critical shut-down utility flow					
9	Shut-down flow reactant B					
10	Dilute concentration reactant A					
11	Critical dilute concentration reactant A					





Application on chemical process





3) Generation of behavioral pattern of the process based on the selected sensors (class profile).





Application on chemical process LAAS-CNRS

4) Online recognition and validation on unseen data

a) Faults identified using 15 original sensors

Fault on concentration detected only with temperature measurements



b) Recognition using 5 selected sensors by Membas (3.66% recognition error)





Application on chemical process LAAS-CNRS

c) Validation on unseen data





Conclusion



1. Medical diagnosis:

7 Two challenges faced for the integration of clinical and microarray data to perform cancer prognosis/diagnosis : High-dimensional and heterogeneous data.

¬ Reduces significantly the number of markers needed to perform a cancer
prognosis task (15 hybrid markers vs. 70 Amsterdam genes).

2. Industrial process diagnosis:

Proposed approach handles interval data which are of big interest in practical situations to take into account inherent uncertainty to sensors measurement and noisy data (avoid false alarms).

Application on chemical process: High fault detection accuracy and reduced number of sensors (avoid expensive on-line concentration mea- surement)



Conclusion



> Despite their behavioral difference, both domains industrial process and medical diagnosis exhibit many common practices:

- \rightarrow Sensor selection for industrial process diagnosis
- \rightarrow Marker selection for medical diagnosis

A novel methodology enables to handle simultaneously both problems regardless of their own characteristics:

 \rightarrow Copes with the problem of high dimensionality based on classical optimization methods.

→ Handles appropriately heterogeneous data (quantitative, qualitative, interval)

 \rightarrow Handling interval data which are of big interest in practical situations to take into account inherent uncertainty to sensors measurement and noisy data (avoid false alarms).