

Dysphonic Voice Detection for Speakers' Biometry

Pedro Gómez¹, Luis M. Mazaira¹, Agustín Álvarez¹, and Eugenia San Segundo²

¹Center for Biomedical Technology, Universidad Politécnica de Madrid, Madrid, Spain

²Universidad Internacional Menéndez Pelayo (UIMP), Madrid, Spain

E-mail: pedro@fi.upm.es

Introduction

- Phonation Distortion leaves relevant marks in a speaker's biometric profile.
- Dysphonic Voice production may be used in biometrical speaker characterization.
- In the present paper phonation features derived from the glottal source (GS) parameterization after the vocal tract inversion is proposed for dysphonic voice characterization in Speaker Verification tasks (Gómez, 2012).



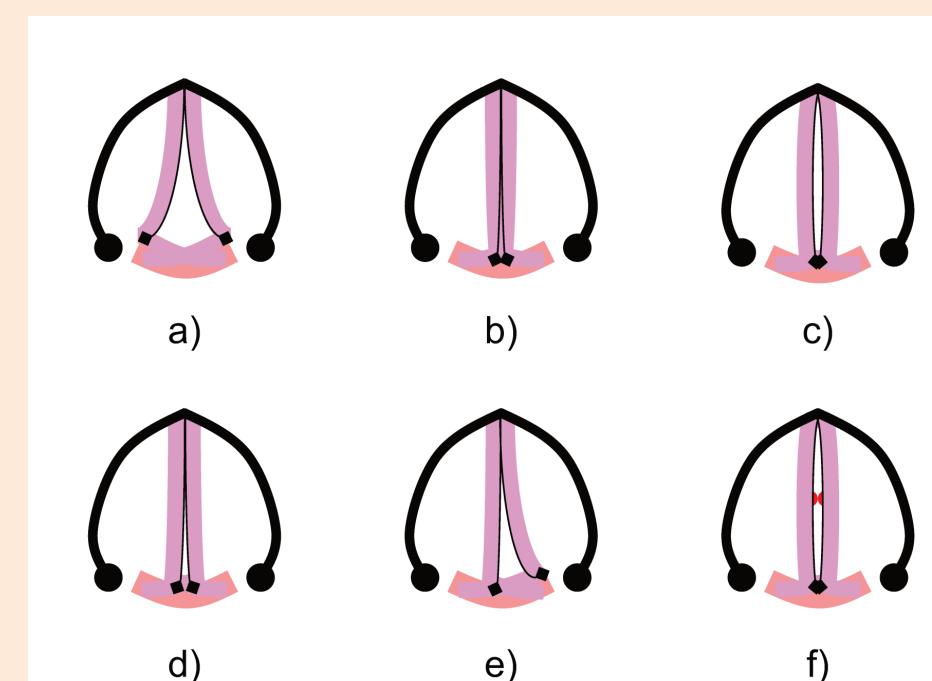
Vocal Fold Images during closed phase from stroboscopic endoscopy. Left: bilateral nodules. Center: Reinke's Edema. Right: Polyp in right fold.

Distortion in Dysphonic Voice is mainly due to :

- Vocal Fold vibration Asymmetry
- Deficient Glottal Closure

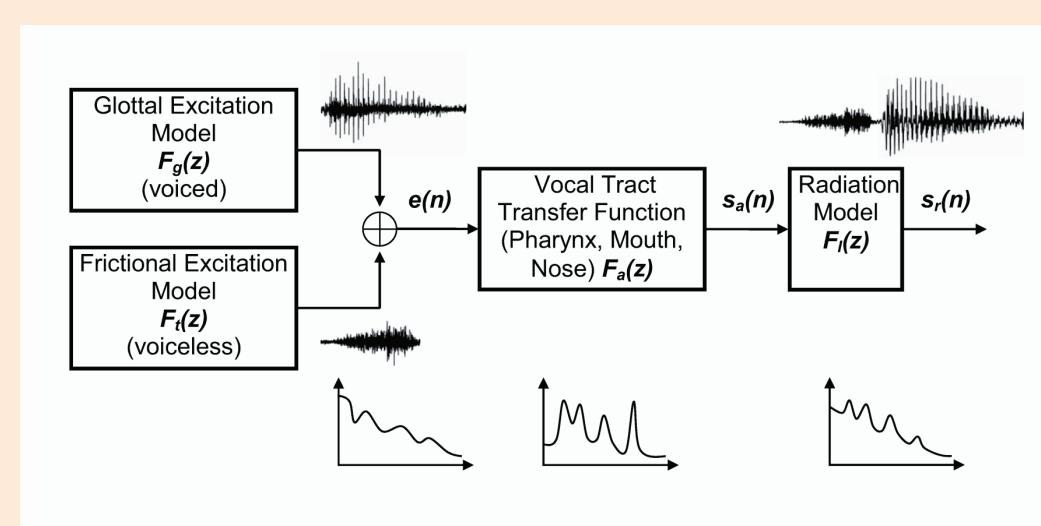
Main causes:

- Organic pathology
- Phonation Dysfunction
- Neurological Diseases
- Emotional Perturbations



Vocal Fold simplified situations:
a) Open during breathing
b) Phonation closed phase
c) Phonation open phase
d) Deficient closure in posterior third
e) Asymmetric contact defect
f) Deficient closure in the medial third (lesion)

Phonation is related to Vocal Fold vibration: voiced speech segments (long vowels and fillers)
Voiced speech is inverse filtered to obtain the glottal residual, the glottal source is estimated from the residual
Different sets of parameters are estimated from the glottal source: distortion, cepstral, spectral, biomechanical, temporal, glottal closure and tremor. For a review of algorithmic methods see (Gómez, 2009).



Fant's Speech Production Model

| Perturbation Parameters |
|----------------------------|
| 1. Abs. Pitch |
| 2. Abs. Normalizer |
| 3. Abs. Norm. Ar. Shimmer |
| 4. Abs. Norm. Min. Sharp. |
| 5. Noise-Harm. Ratio (NHR) |
| 6. Muc./AvAc. Energy (MAE) |

| Cepstral Parameters |
|---------------------|
| 7. MWC Cepstral 1 |
| 8. MWC Cepstral 2 |
| 9. MWC Cepstral 3 |
| 10. MWC Cepstral 4 |
| 11. MWC Cepstral 5 |
| 12. MWC Cepstral 6 |
| 13. MWC Cepstral 7 |
| 14. MWC Cepstral 8 |
| 15. MWC Cepstral 9 |
| 16. MWC Cepstral 10 |
| 17. MWC Cepstral 11 |
| 18. MWC Cepstral 12 |
| 19. MWC Cepstral 13 |
| 20. MWC Cepstral 14 |

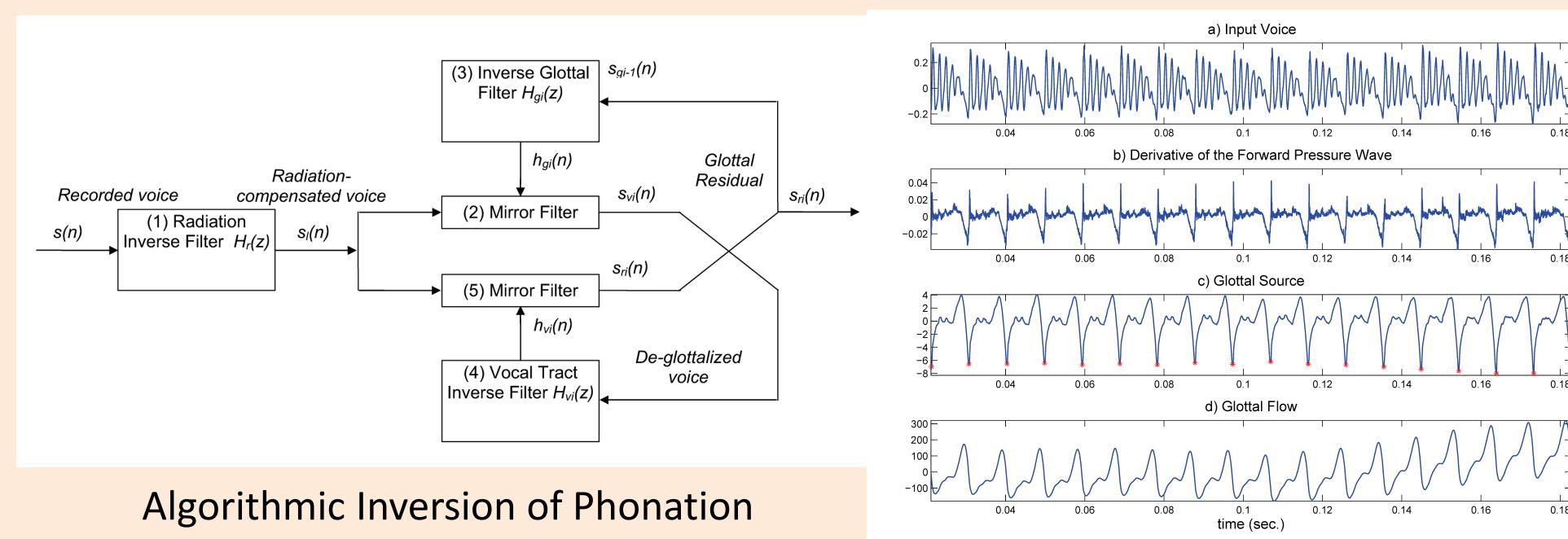
| Spectral Parameters |
|-------------------------------|
| 21. MW PSD 1st Max. ABS. |
| 22. MW PSD 1st Min. ABS. |
| 23. MW PSD 2nd Max. rel. |
| 24. MW PSD 2nd Min. rel. |
| 25. MW PSD 3rd Max. rel. |
| 26. MW PSD End Val. rel. |
| 27. MW PSD 1st Max. Pos. ABS. |
| 28. MW PSD 1st Min. Pos. rel. |
| 29. MW PSD 2nd Max. Pos. rel. |
| 30. MW PSD 2nd Min. Pos. rel. |
| 31. MW PSD 3rd Max. Pos. rel. |
| 32. MW PSD End Val. Pos. rel. |
| 33. MW PSD 1st Min NSF |
| 34. MW PSD 2nd Min NSF |

| Biomechanical Parameters |
|------------------------------|
| 35. Body Mass |
| 36. Body Losses |
| 37. Body Stiffness |
| 38. Body Mass Unbalance |
| 39. Body Losses Unbalance |
| 40. Body Stiffness Unbalance |

| Temporal Parameters |
|--------------------------|
| 47. Rel. Recov. 1 Time |
| 48. Rel. Recov. 2 Time |
| 49. Rel. Recov. 3 Time |
| 50. Rel. Open 2 Time |
| 51. Rel. Max. Ampl. Time |
| 52. Rel. Recov. 1 Ampl. |
| 53. Rel. Recov. 2 Ampl. |
| 54. Rel. Open 1 Ampl. |
| 55. Rel. Open 2 Ampl. |
| 56. Rel. Start Flow Time |
| 57. Rel. Stop Flow Time |
| 58. Rel. Closing Time |

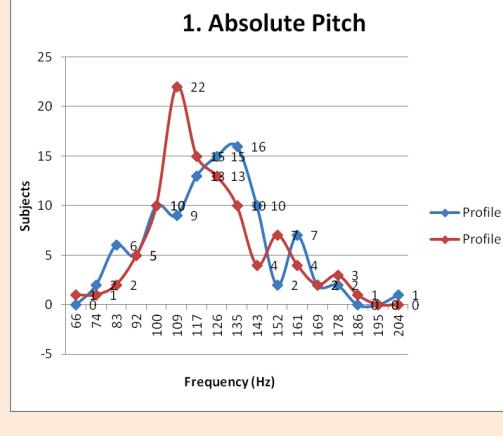
| Glotto GAP Parameters |
|------------------------|
| 59. Val. Flow GAP |
| 60. Val. Contact GAP |
| 61. Val. Adduction GAP |
| 62. Val. Permanent GAP |

| Tremor Parameters |
|-------------------|
|-------------------|



Materials and Methods

- Fillers and log vowels (lasting more than 100 ms)
- Segmented from a telephonic (GSM) database of 100 male speakers
- Corresponding to phonations of vowels between /a/ and /e/
- 68 parameters were obtained from each 100 ms long segment
- A comparative paired test showed equivalence between /a/ and /e/ phonations



Sample Matching is based on comparing Phonation Descriptors from an unknown speaker (test set: blue) against a suspect speaker (control set: red) referred to a Line-Up speaker set (model set: green) to evaluate the Prosecutor's Hypothesis against the Defender's Hypothesis (Taroni, 2006):

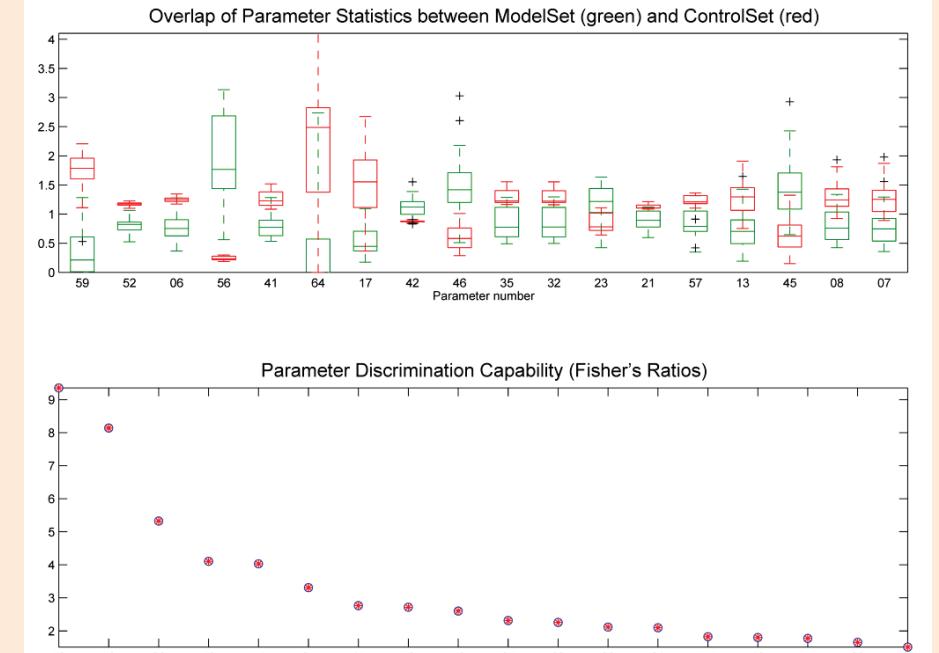
- Parameter selection from Fisher's Discriminant Ratios

$$C_{Fi} = \frac{\mu_{im} - \mu_{ic}}{\sqrt{\frac{s_{im}^2 + s_{ic}^2}{n_m + n_c}}}$$

μ_{im}, μ_{ic} : model and control sample averages for parameter i

s_{im}, s_{ic} : model and control sample standard errors for parameter i

n_m, n_c : model and control set sample sizes



- Sample Matching Paradigm:
Zq: Questioned sample set (test: blue)
Ts: Gaussian Mixture Model from Suspect's sample set (control: red)
Tm: Gaussian Mixture Model from Line-Up sample set (model: green)
- Prosecuto's vs Defender's Hypothesis is evaluated as a Log-Likelihood Ratio:

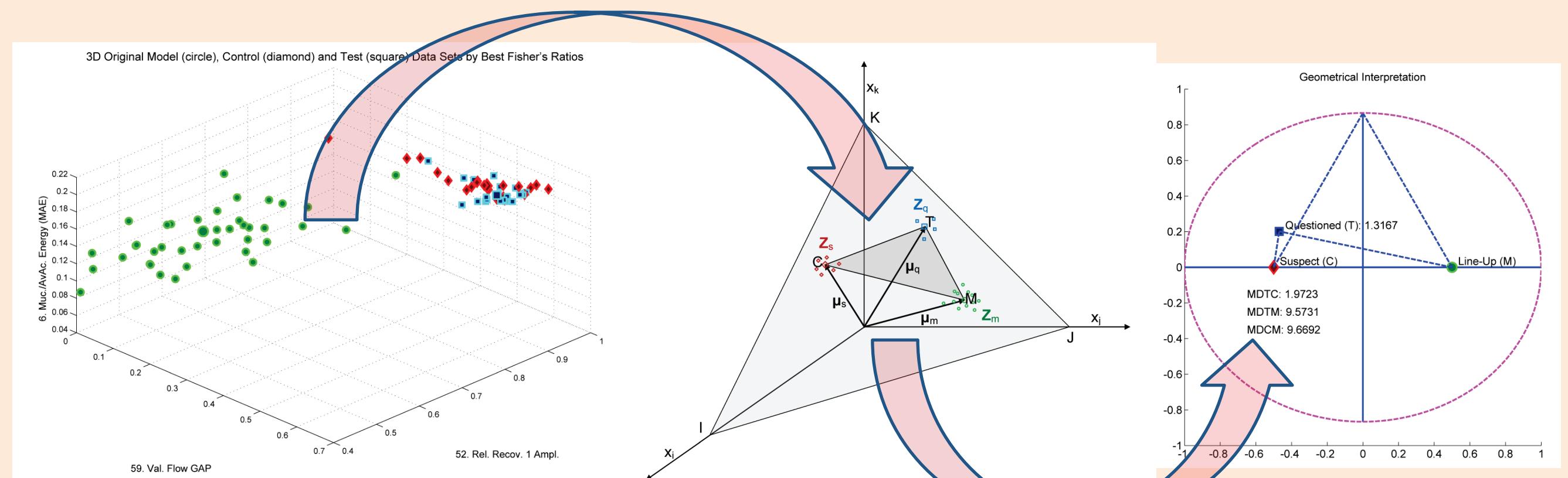
$$\lambda_{pd} = \log\{\Pr(Z_t | \Gamma_c)\} - \log\{\Pr(Z_t | \Gamma_m)\}$$

Where conditional probabilities are given by:

$$\Pr(Z_t | \Gamma_c) = \frac{1}{(2\pi)^{M/2} |\mathbf{C}_c|^M} e^{-\frac{1}{2}(\mu_t - \mu_c)^T \mathbf{C}_c^{-1} (\mu_t - \mu_c)}$$

$$\Pr(Z_t | \Gamma_m) = \frac{1}{(2\pi)^{M/2} |\mathbf{C}_m|^M} e^{-\frac{1}{2}(\mu_t - \mu_m)^T \mathbf{C}_m^{-1} (\mu_t - \mu_m)}$$

Details are to be found in (Gómez 2012)



First Experiment:

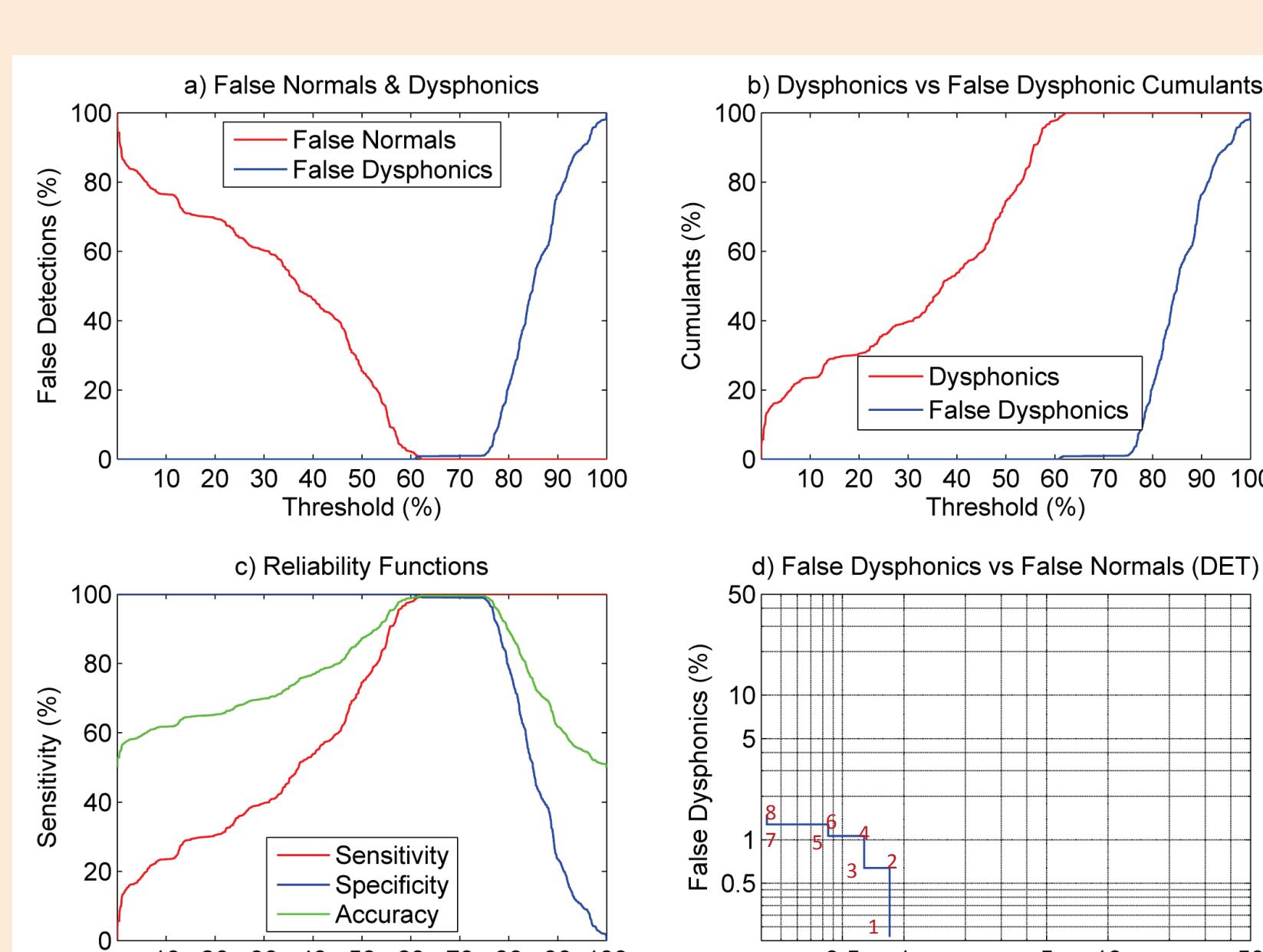
- Splitting the 100 male speakers in two equal-size subsets by their normophonic condition.
- Using a normative database validated by Hospital Gregorio Marañón of Madrid with samples of /a/ (50 male speakers).
- LLR's estimate the conditional probability of a given sample being normophonic or dysphonic (10-fold cross-validation).

Objectives:

- Estimate the discrimination accuracy of the methodology and the best parameters.
- Produce two reference subsets from GSM quality /e/ of use in Spanish.

Results:

- Normophonic vs dysphonic cumulants, sensitivity, specificity and accuracy, and Detection-Error Trade-off plots.



Second Experiment:

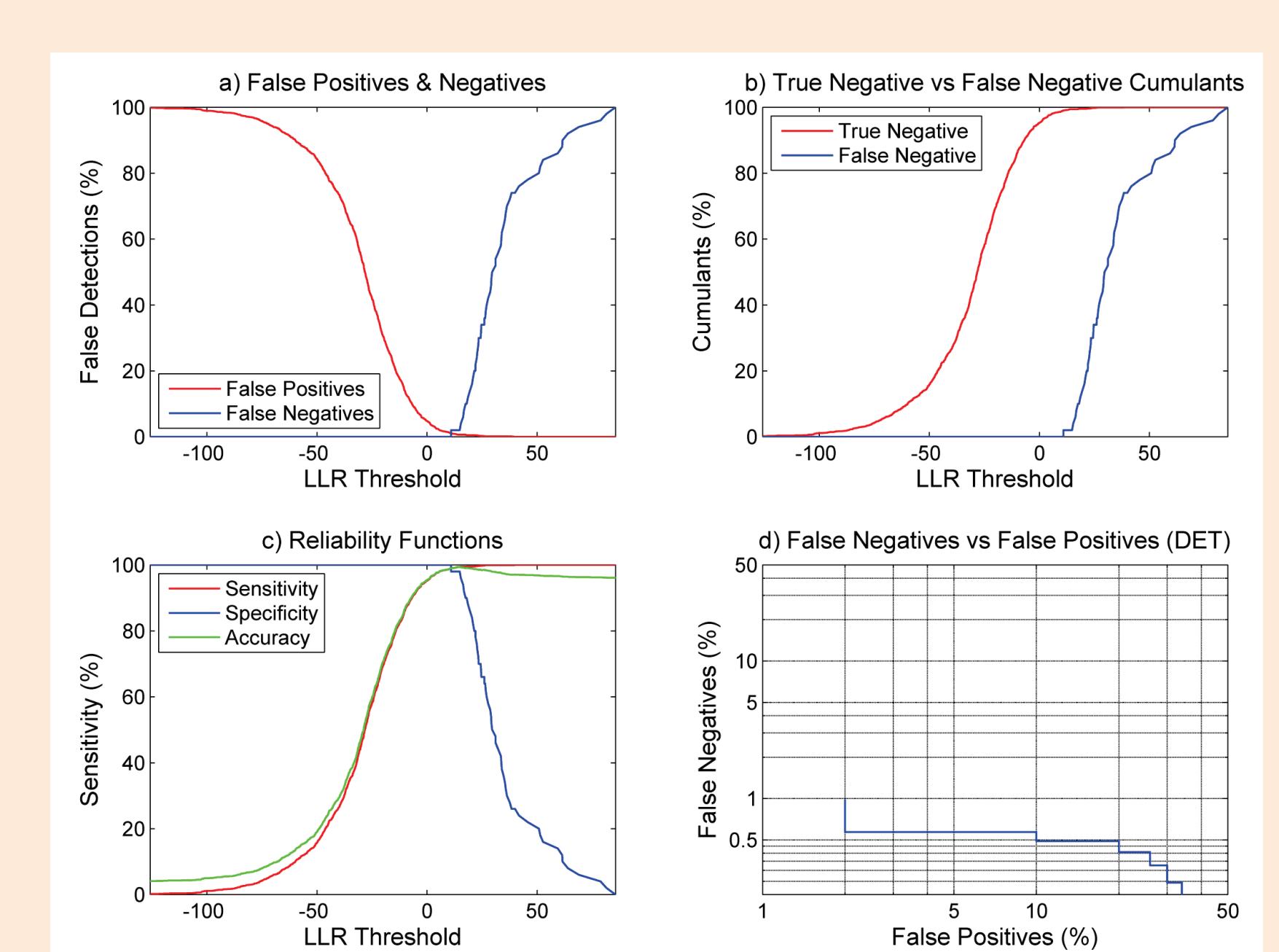
- Matching each normative speaker's sample set against each other.
- Using as model the normative subset produced in the first experiment.
- Using as control the non-normative subset produced in the first experiment. 50 target vs 1225 non-target speakers.

Objective:

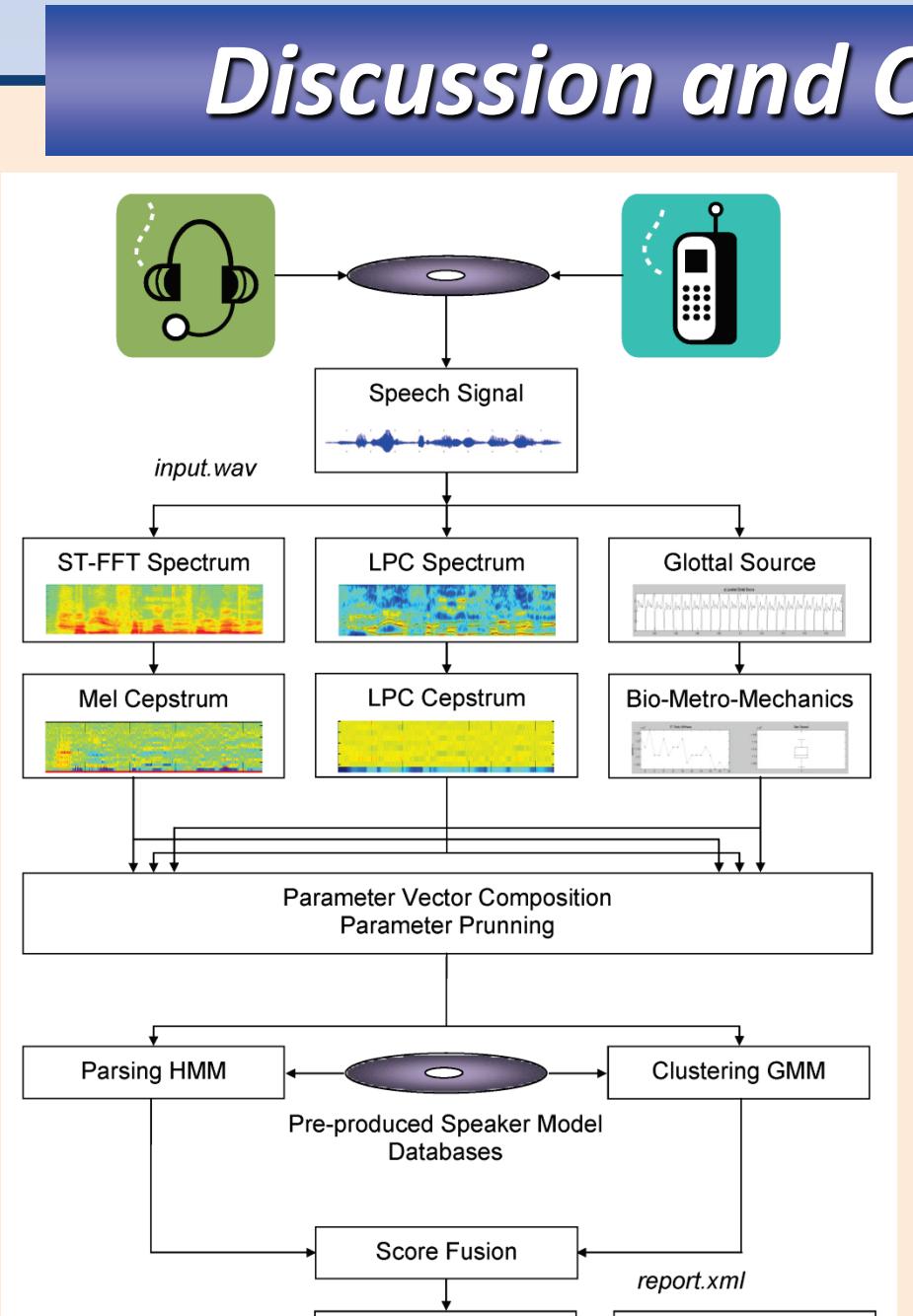
- Estimate the discrimination accuracy of the sample matching methodology in target vs non-target detection.

Results:

- False target vs false non-target detection cumulants, sensitivity, specificity and accuracy, and Detection-Error Trade-off plots.



- Sensitivity to nomophonic vs dysphonic phonation is large enough to allow forensic matchng of voiced speech segments.
- Glottal source parameterizations from /a/ and /e/ are interchangeable, and can be used in cross-matching with no significant differences.
- Accuracy = $(P_f + P_n) / (P_f + P_n + T_f + T_n)$ of target vs not-target matches grants applicability to real forensic cases.
- Margin of optimum LLR values grants strength of evidence to be over 4 in Lucy's Scale (Lucy, 2005).
- Distinction between normophonic and dysphonic phonation seems to be feasible from parameterizations of glottal source.
- Questioned vs Suspect's Sample matching in reference to Line-Ups may be taken to meaningful 2D plots.
- Hybrid matching combining standard MFCC's and glottal source derived parameters may attain rather low equal error rates with telephone-quality speech (Khoury, 2013).



- Gómez, P., et al. (2009). Glottal Source Biometrical Signature for Voice Pathology Detection. *Speech Comm.*, vol. 51, pp. 759-781.
- Gómez, P., et al. (2012). Distance Metric in Forensic Voice Evidence Evaluation using Dysphonia-relevant Features. *Proc. of the VI Meeting of Biometric Recognition of Persons*, Ed. Universidad de Las Palmas de Gran Canaria, pp. 169-178.
- Khouri, E., Mazaira, L.M., et al. (2013). The 2013 Speaker Recognition Evaluation in Mobile environments. *Proc. of the 6th IAPR International Conference on Biometrics*, Madrid, Spain.
- Lucy, D. (2005). *Introduction to Statistics for Forensic Scientists*. Wiley.
- Taroni, F., Aitken, C., Garbolino, P., Biedermann, A. (2006). *Bayesian Networks and Probabilistic Inference in Forensic Science*. Wiley.

Acknowledgments

This work is being funded by grant TEC2012-38630-C04-04 from Plan Nacional de I+D+i, Ministry of Economy and Competitiveness of Spain.