

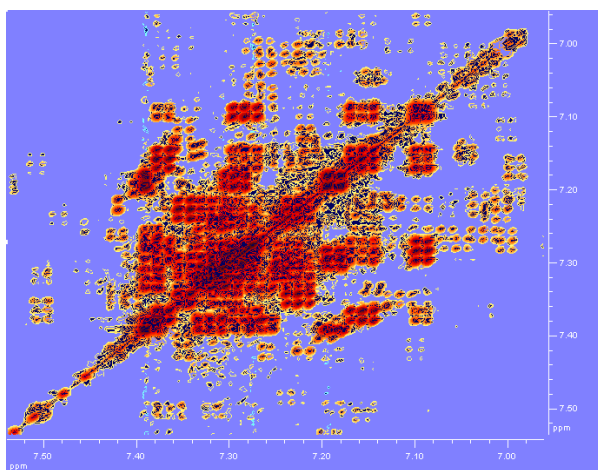


Protocols for assessing quality and traceability of aquaculture products by means of Nuclear Magnetic Resonance

Dr. Roberto Anedda

NMR/MRI facilities

June, 30 2009



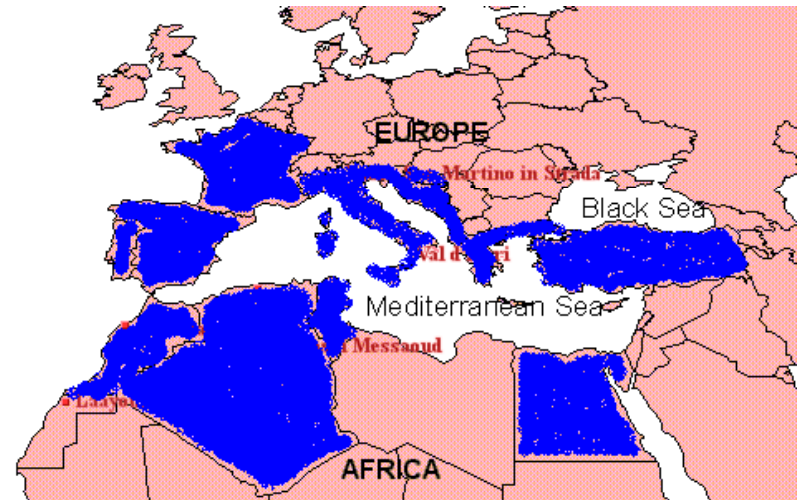
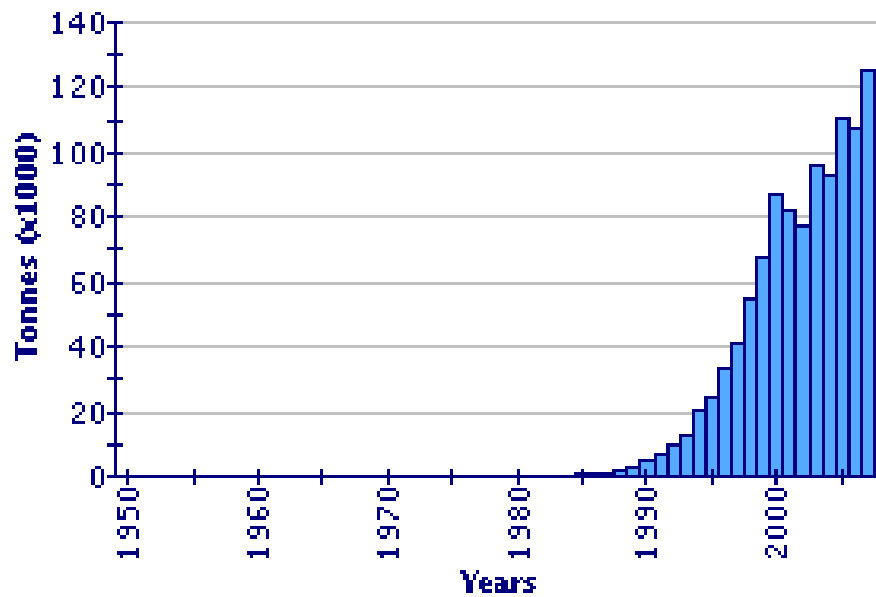
**PortoConte
Ricerche**





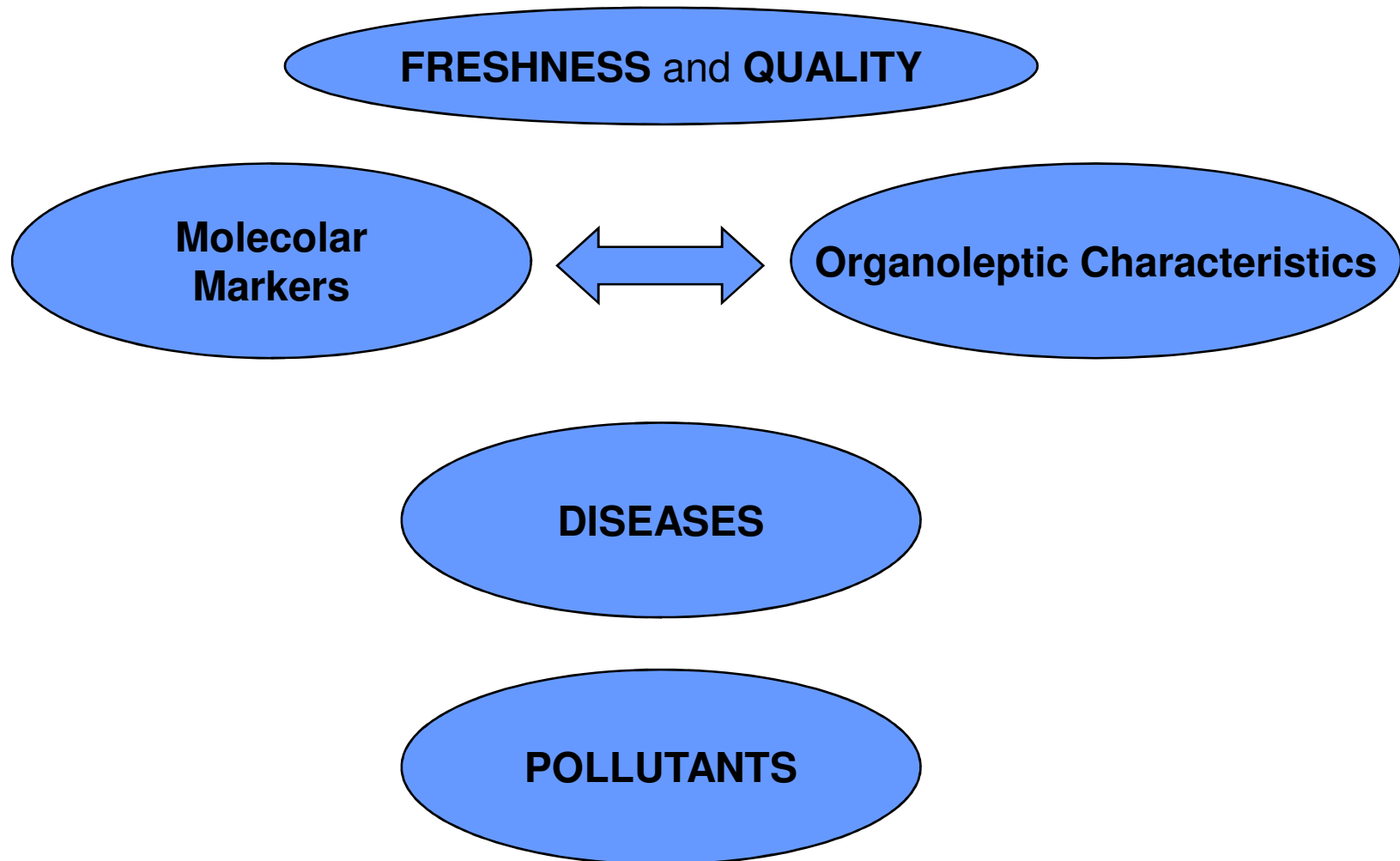
Farmed GILTHEAD SEA BREAM: Annual Production in EUROPE

~ 4500 t in 1990 → ~125000 t in 2007





RELEVANT ISSUES: developing RELIABLE ANALYTICAL METHODS for assessing
COMPOSITION (METABOLITE PROFILING)
GEOGRAPHICAL ORIGIN





CONVENTIONAL and CHROMATOGRAPHIC TECHNIQUES FOR METABOLITE PROFILING

LENGTHY SEPARATION STEPS → ARTIFACTS (LABILE METABOLITES)

ENZYMATIC and COLORIMETRIC METHODS →

LACK SPECIFICITY
INTERFERENCES

HPLC → CHROMOPHORES, FUNCTIONAL GROUPS
UNEXPECTED, UNKNOWN METABOLITES

GC → VOLATILE SPECIES
DERIVATIZATION

MASS SPECTROMETRY →

- VALID STRUCTURAL ANALYSIS
- SENSITIVITY
- GENERALLY DESTRUCTIVE



NMR: a COMPLEMENTARY, COMPREHENSIVE, HIGHLY INFORMATIVE TECHNIQUE

NOT LIMITED BY ANY ANALYTE PROPERTIES

MINIMAL SAMPLE PREPARATION

STRUCTURE, CONFORMATION, DYNAMICS

SIMULTANEOUS DETECTION OF ALL ANALYTES

NON DESTRUCTIVE

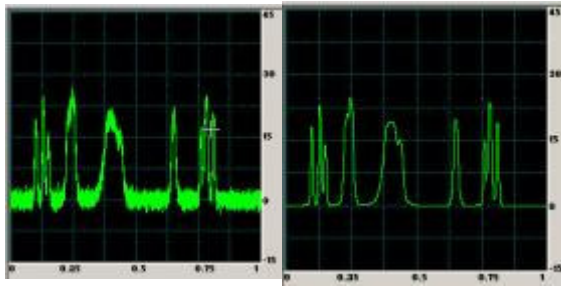


NUCLEAR MAGNETIC RESONANCE (NMR) has been widely used in metabolomics of complex mixtures

Vast scientific literature concerning NMR-aquaculture



Principal drawbacks of NMR:



Low sensitivity

- Gyromagnetic ratio γ ,
- Natural abundance
- Magnetic field (instrumentation)



Acquisition times

- Relaxation phenomena - delays
- Sample concentration



High cost

NO ROUTINE, evaluate COSTS-BENEFITS



METABOLOMIC ANALYSIS

*“systematic study of the unique chemical fingerprints that specific cellular processes leave behind”**

EXPERIMENTAL

Extraction

NMR 1D e 2D Spectra Acquisition

Interpretation of spectra

STATISTICAL ANALYSIS

Principal Component Analysis

Interpretation

Conclusions

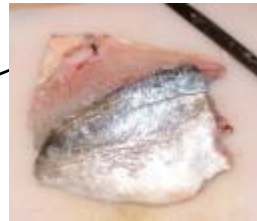
* B. Daviss, "Growing pains for metabolomics," *The Scientist*, 19[8]:25-28, April 25, 2005



EXPERIMENTAL

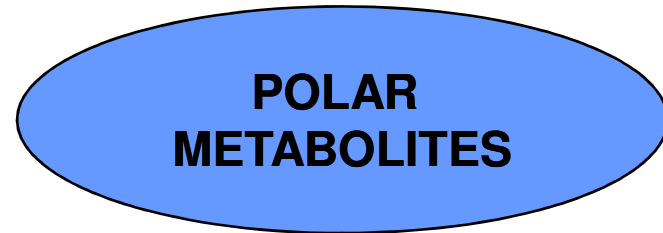
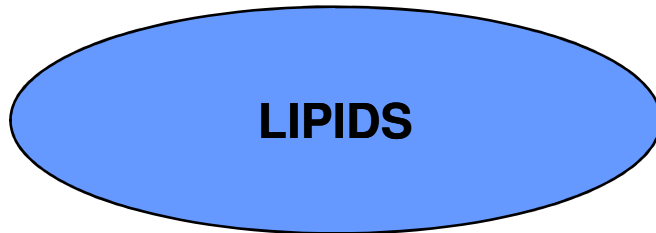


+ IMPORTED
(Farmed)



Extraction 1
CH₂Cl₂ (stored at -40°C
until analysis)

Extraction 2
EtOH at RT



¹H NMR

¹H NMR



LIPIDS

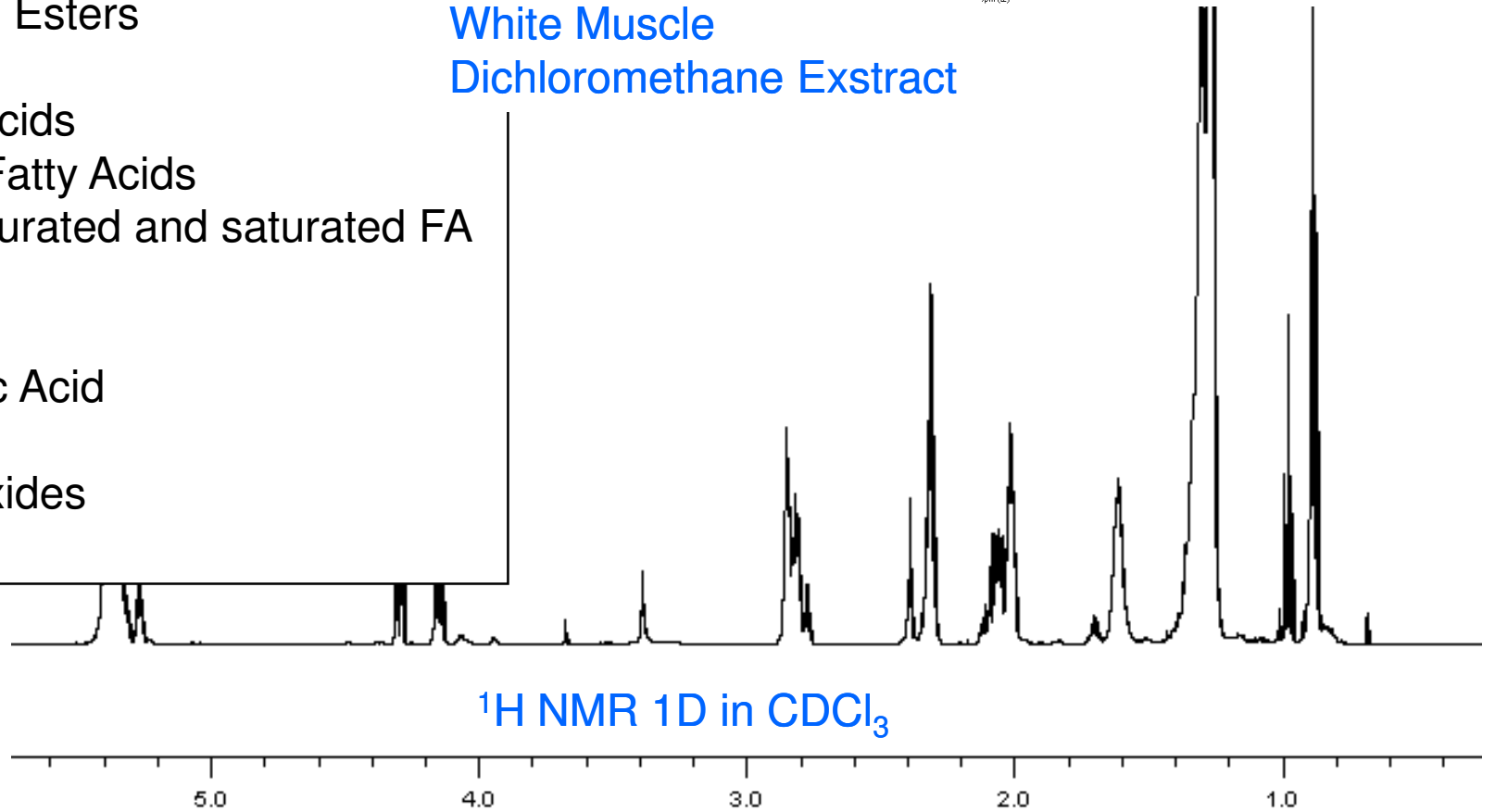
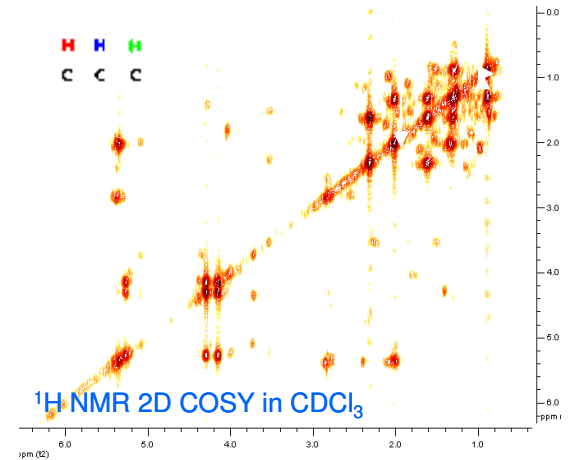
- Triacylglycerols
- Diacylglycerols
- Phospholipids (PC,PE,..)

- Cholesterol
- Cholesterol Esters

- ω -3 Fatty Acids
- ω -6 + ω -9 Fatty Acids
- Monounsaturated and saturated FA
- EPA
- DHA
- Arachidonic Acid

- Hydroperoxides
- Aldheydes

Sparus Aurata
White Muscle
Dichloromethane Extract





POLAR METABOLITES

- Creatine, CN1C=NC2=C1N(=O)C(N2)C
- Taurine, CSC(N)(C(=O)O)C
- TMA, CN(C)C
- TMAO, CN(C)C(=O)O
- Glicine, C(N)C(=O)O
- Betaine, CN(C)C(=O)O
- Serine, C(N)C(C(=O)O)C
- Lactate, C(C(=O)O)C
- Anserine, CN(C)C(C(=O)O)C
- Nicotinamide, CN1C=NC2=C1C(=O)N(C2)C
- other aminoacids

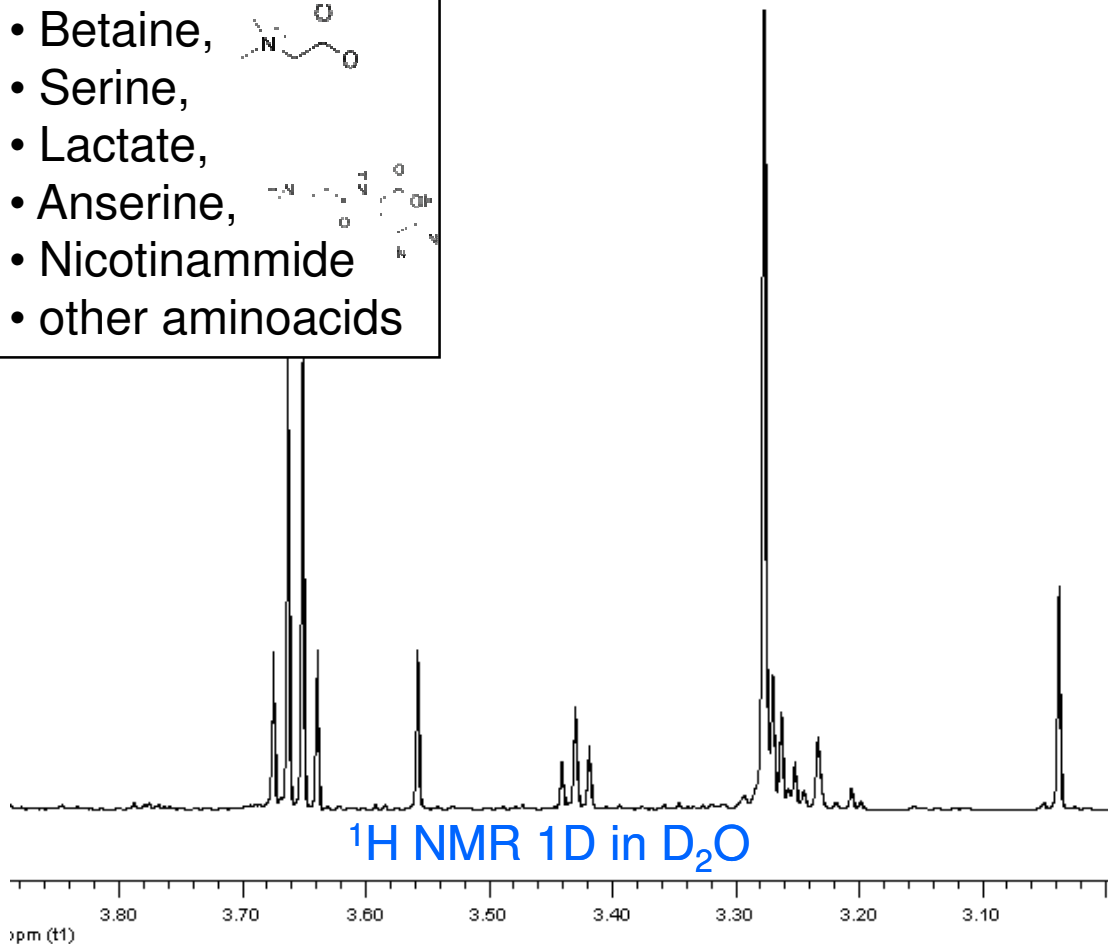
Sparus Aurata,
White Muscle
Ethanolic extract

The accumulation of **TMA** in rotting fish as a result of bacterial degradation of choline, as well as the reduction of TMAO to TMA, is responsible for their characteristic 'fishy' odor.

Main Functions of **TAURINE** in Mammals
Bile acid conjugation
Detoxification
Osmoregulation
Membrane Stabilization
Regulation of intracellular Ca²⁺
Homeostasis

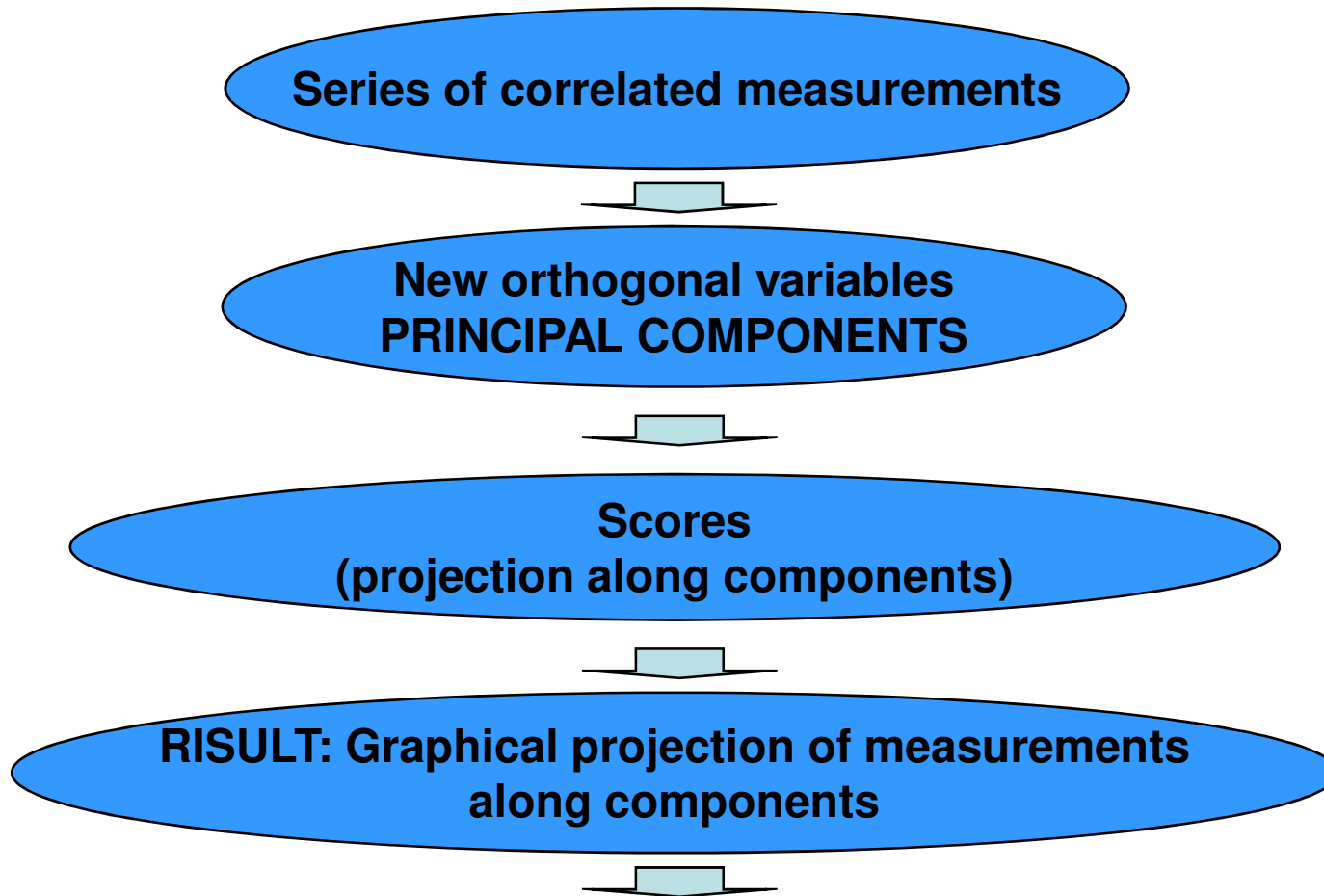
NIACIN (nicotinic acid) and its amide, **NIACINAMMIDE**
High cholesterol
Atherosclerosis
Heart Attacks
Osteoarthritis

ANSERINE
Muscle activity is closely related to the dipeptide concentration. In vigorously contracting muscles, the dipeptide concentration is high.





Multivariate statistical analysis: PCA (Principal Components Analysis):



PCA

unsupervised method

Orthogonal decomposition of variance



GENERAL PROCEDURE

Sample selection

Spectral Bucketing (binning)

Analysis of influence plot

Construction of a **Reliable Model**

Cross **validation**

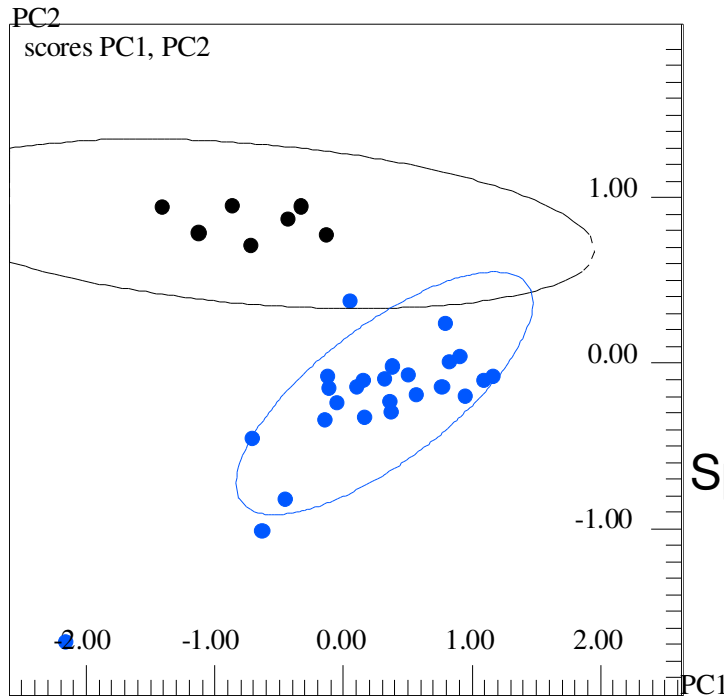
Scores plot, loadings plot

Hotelling's T^2 plot → presence of **clusters**

Analysis of **critical spectra** and **save model**



Scores Plot



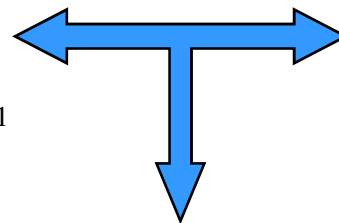
FARMED

VS
WILD

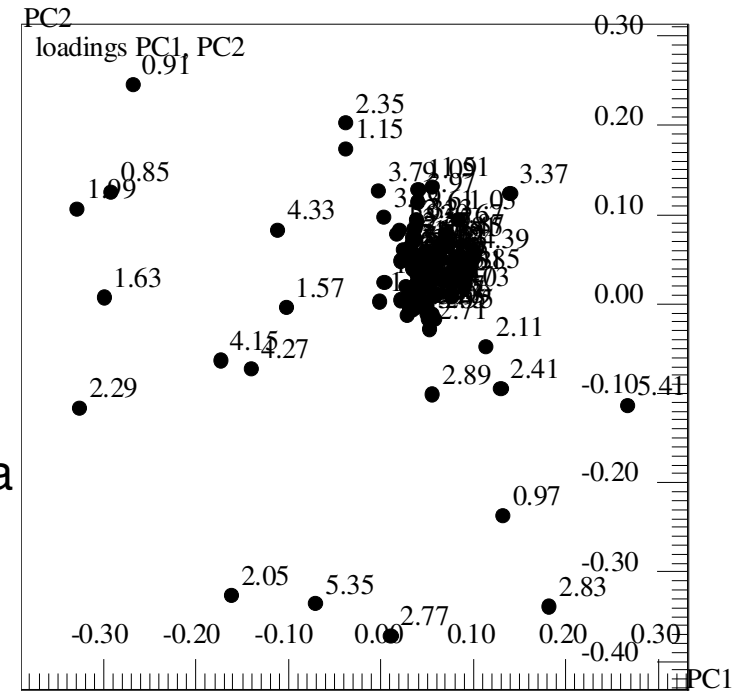
~250 gr
Healthy

(PCA space)

Spectroscopic Data



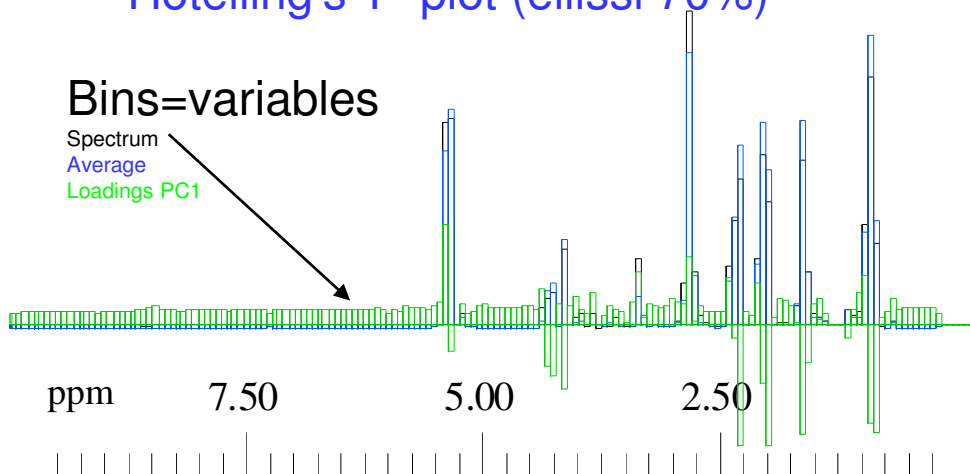
Loadings Plot



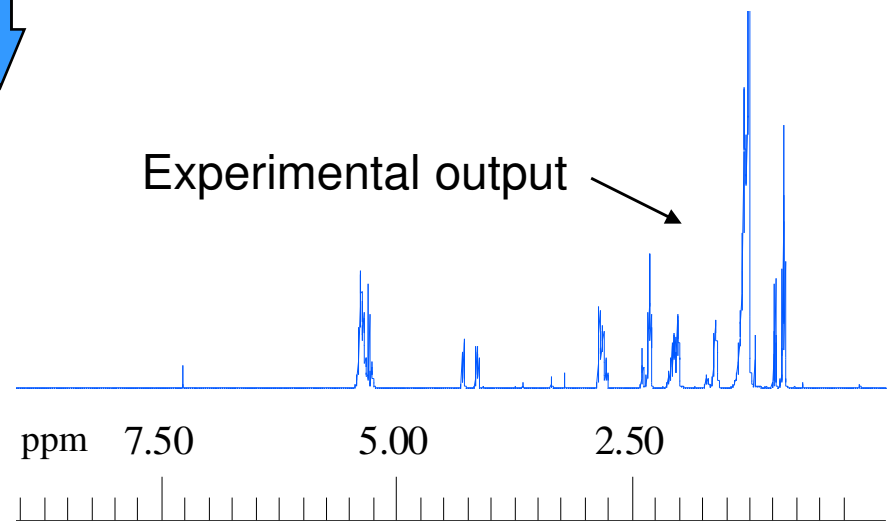
Hotelling's T^2 plot (ellissi 70%)

Bins=variables

- Spectrum
- Average
- Loadings PC1

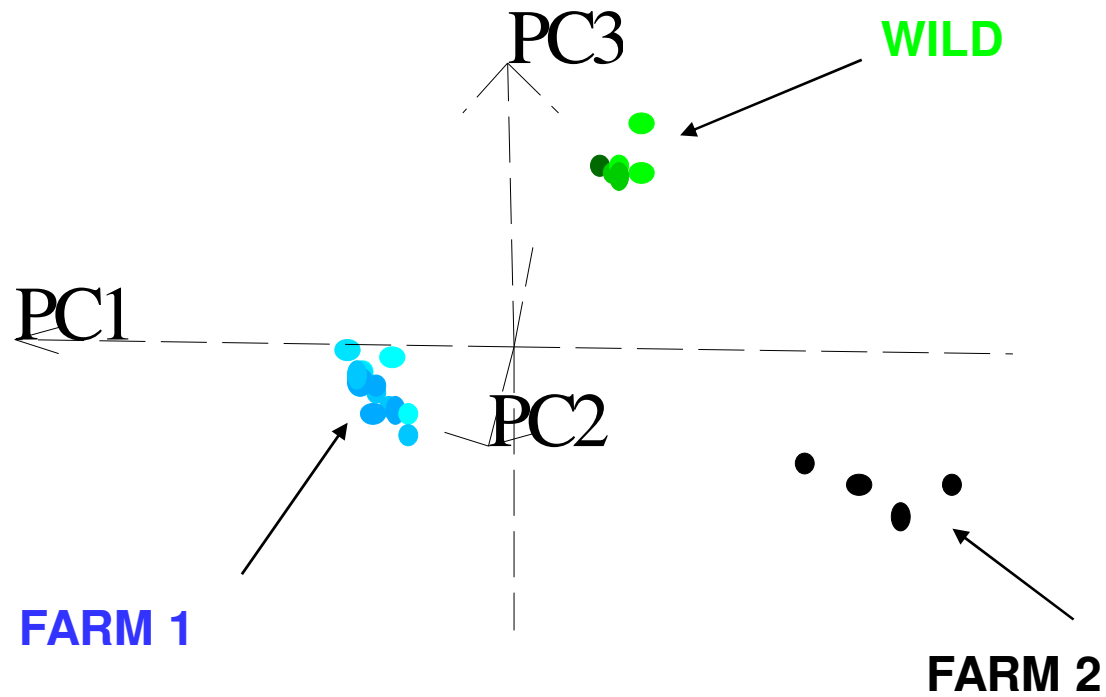


Experimental output





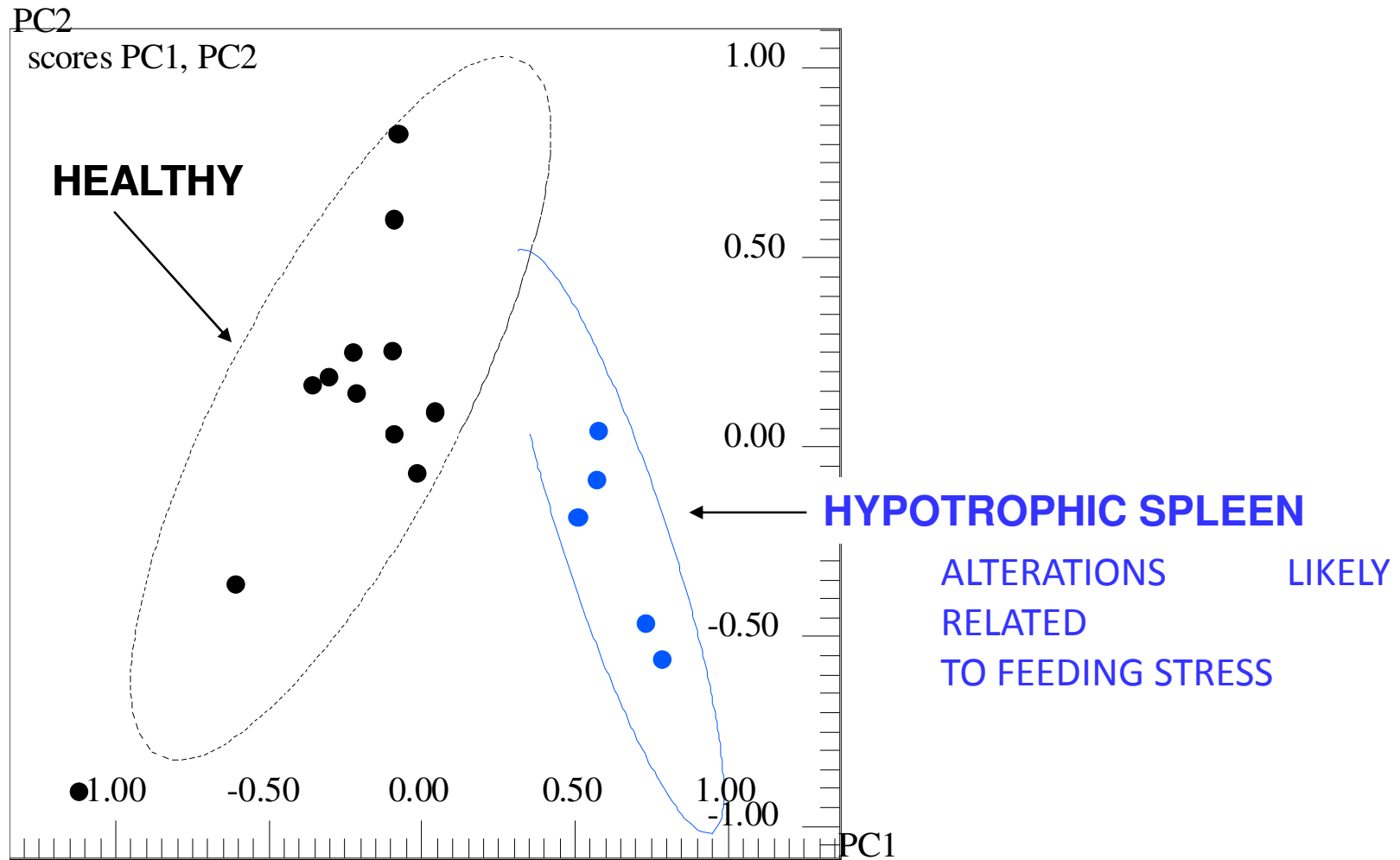
~150 gr
Healthy
Same sampling period



variance explained by PC1 : 43.60%
variance explained by PC2 : 21.94%
variance explained by PC3 : 11.48%



SAME FARM, SAME SIZE (200-280 gr)



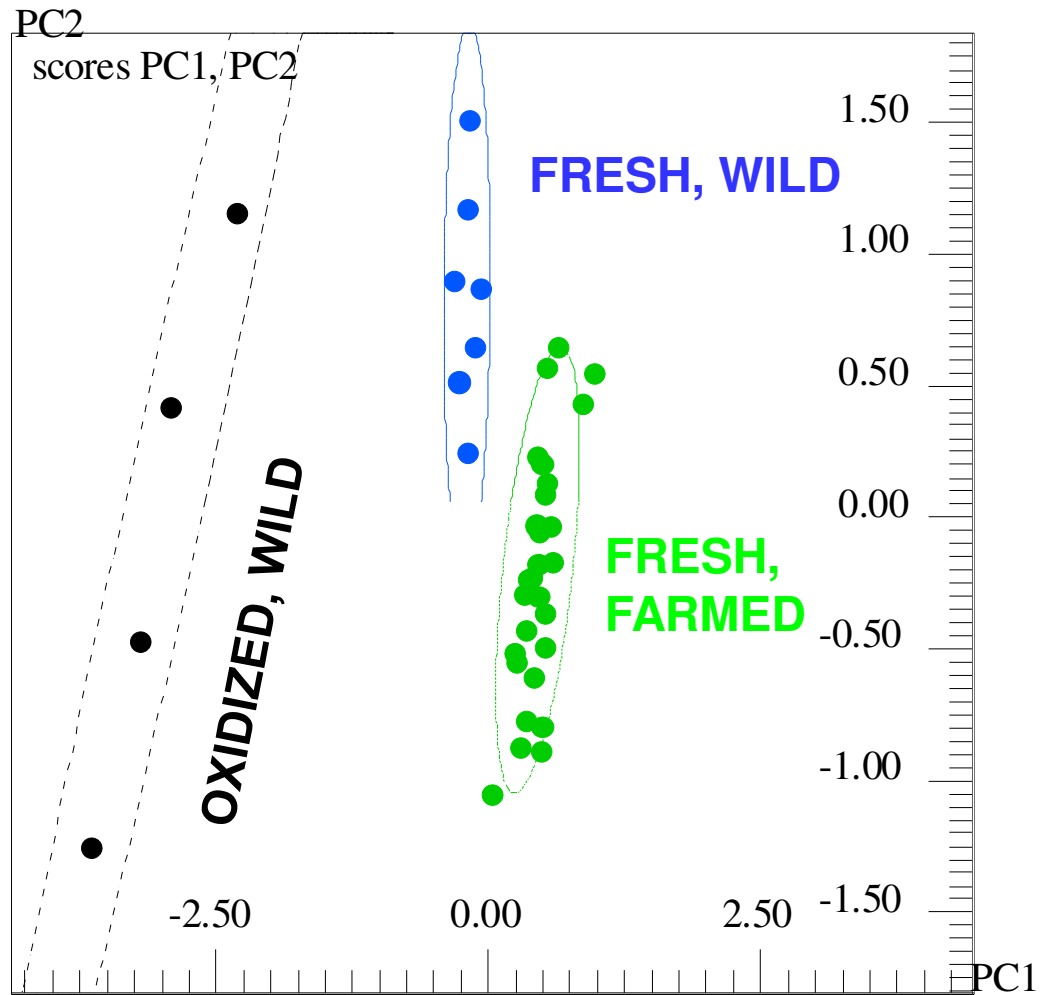
variance explained by PC1 : 41.55%

variance explained by PC2 : 28.68%

L'intervallo di confidenza relativo all'Hotelling's T2 plot e' dell' 80%.



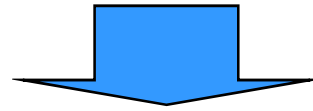
SAME SIZE (200-280 gr). EVALUATING OXIDATION



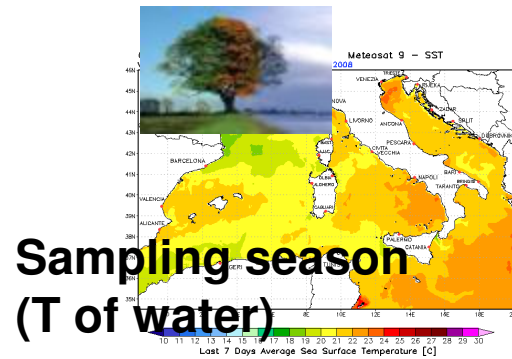
variance explained by PC1 : 56.83%
variance explained by PC2 : 20.77%



PCA Analysis on ¹H NMR measurements of Gilthead Seabream (*Sparus Aurata*)



CH₂Cl₂ extract, related to lipids



**FACTORS
INFLUENCING
COMPOSITION**



Farmed vs wild

Feeding Stress





In conclusion

A Metabolite fingerprinting by means of NMR; has been successfully carried out

Several *nutritionally relevant species* have been identified in white muscle of Gilthead seabream. This lay the foundation for quantitative analysis of potentially interesting compounds

Storage: Aldehydes, peroxides and hydroxy-compounds formed as a consequence of lipid oxidation are clearly observable in NMR spectra. NMR allow indentifying *fish freshness* Also PCA discriminates frsh and oxidized samples.

Studying fish *LIPIDS* by NMR we have successfully discriminated: **FARMED**
vs WILD fish,
its *geographic origin*,
alterations due to *feeding stress*



Team

NMR lab:

Dr. Gilberto Mulas

Dr. Maria Grazia Galaffu

Sample selection and extraction:

Dr. Viviana Santercole

Dr. Roberto Cappuccinelli

Dr. Monica Madrau



Advices on Statistical analysis:

Dr. Roberto Tonelli

Supervision:

Dr. Tonina Roggio

Prof. Sergio Uzzau

anedda@portocontericerche.it