

EuroValve

October 24-25, 2014

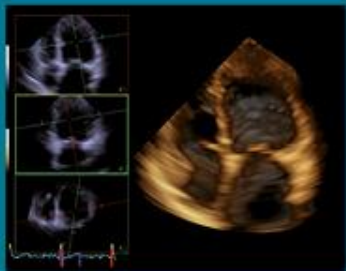
Emerging biomarkers in valvular heart disease (aortic stenosis)

Cécile Oury, PhD



Cardiovascular Sciences





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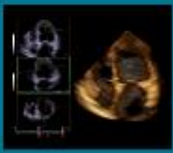
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Faculty disclosure

Cécile OURY

I have **no financial relationships** to disclose

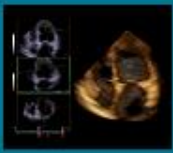




Biomarker: definition

“a characteristic that is objectively measured and evaluated as an indicator of normal biological processes, pathogenic processes, or pharmacological responses to a therapeutic intervention.”

Include any representation of a biological process, including circulating molecules, genetic markers, cellular markers, results of imaging, or findings on physical examination



Aortic stenosis

Severe AS affects > 3-7 % of patients over 65 years.

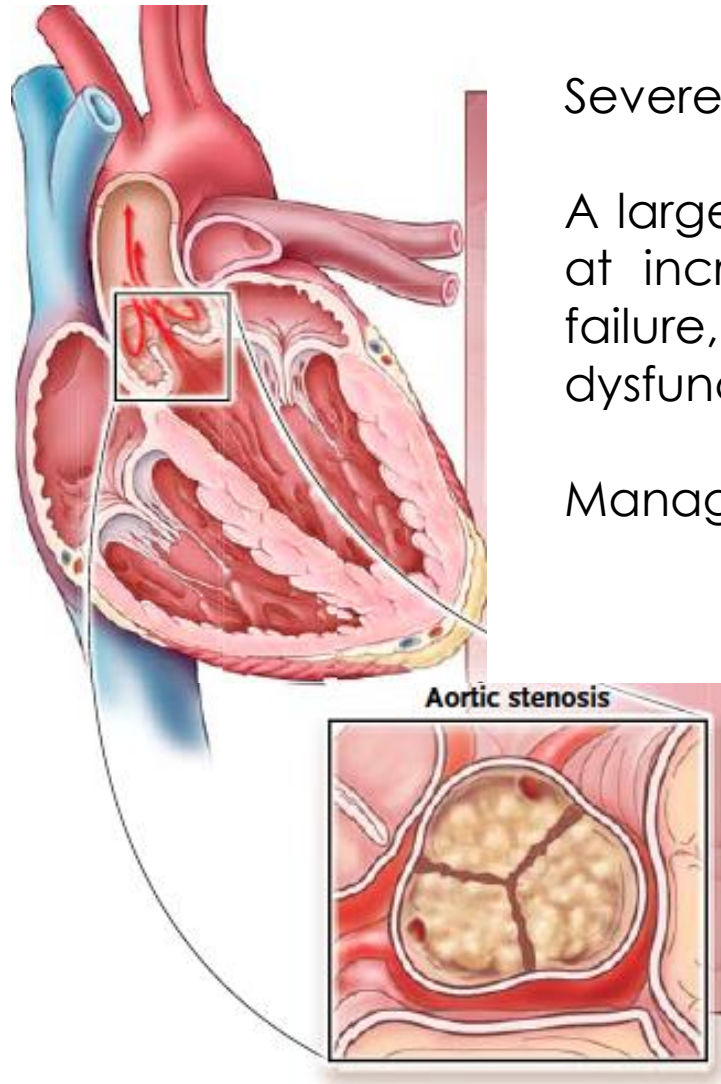
A large majority of patients are asymptomatic and are at increased risk for untoward events (death, heart failure, symptomatic deterioration, ventricular dysfunction).

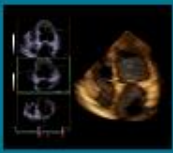
Management : controversial. Risk stratification required.

Diagnosis: imaging techniques, high cost, skill.

➤ Need for easily accessible blood biomarkers that can provide incremental diagnostic and prognostic information to the existing tests in asymptomatic patients.

- Severity
- Progression
- outcome



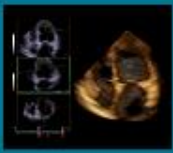


Biomarkers



**Hypothesis-
driven
Biomarkers**

**« omics »
Biomarkers**



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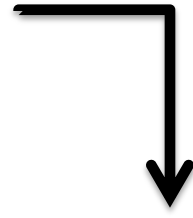


Advanced age, male sex,
smoking
Hypertension, diabetes,
High LDL-C, Lp(a) levels,
obesity, overweight,
CKD, congenital AV
malformation

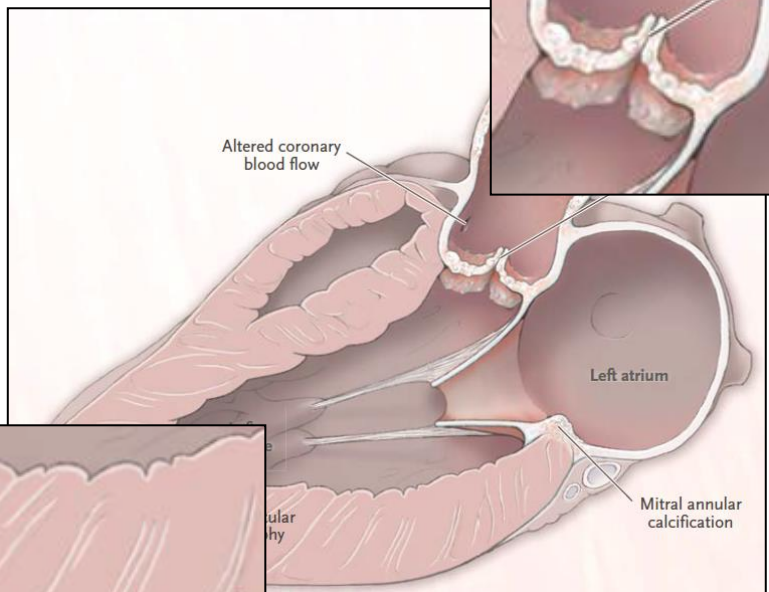


Disease progression

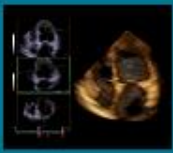
Aortic stenosis



Leaflet
calcification



Left ventricular hypertrophy



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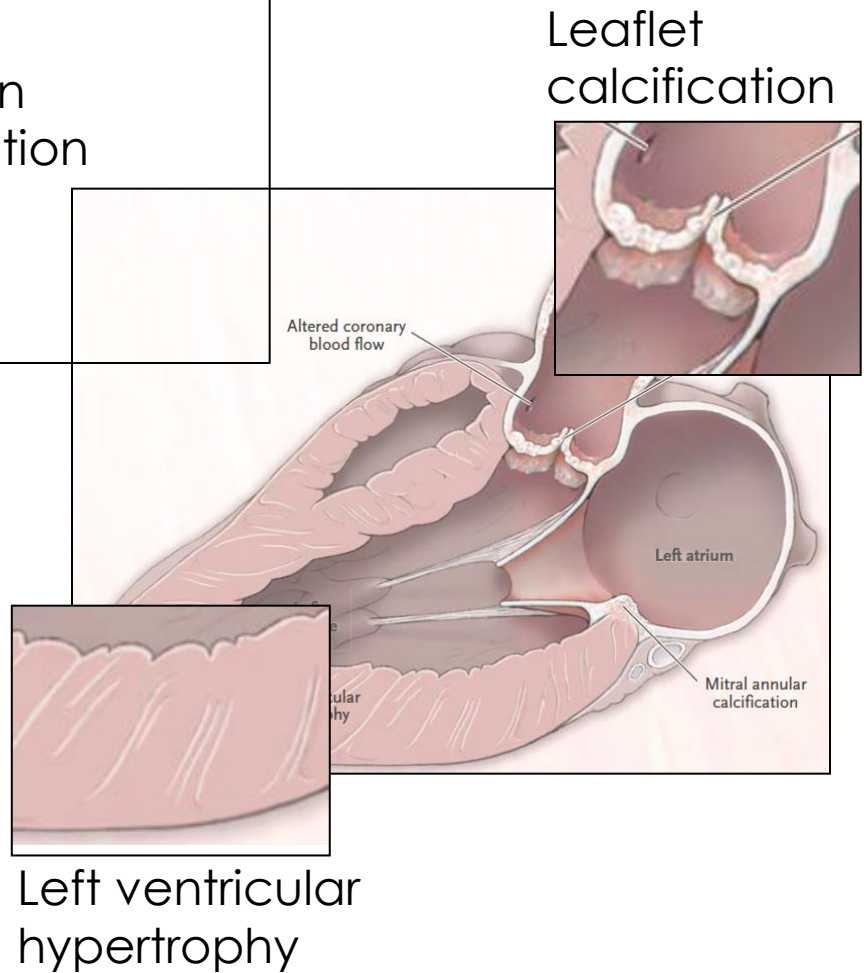


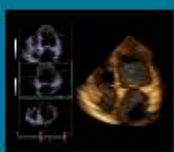
Mechanisms of disease progression:

- Inflammation
- Lipid infiltration
- Myofibroblast differentiation
- Shear stress, platelet activation
- Cardiac angiotensin II production
- Procalcific stimuli
- ECM deposition
- Remodeling



Hypothesis-driven Biomarkers

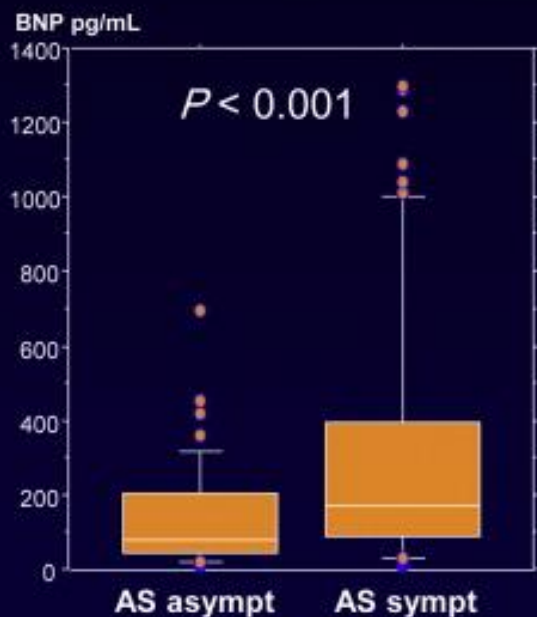




Natriuretic peptides

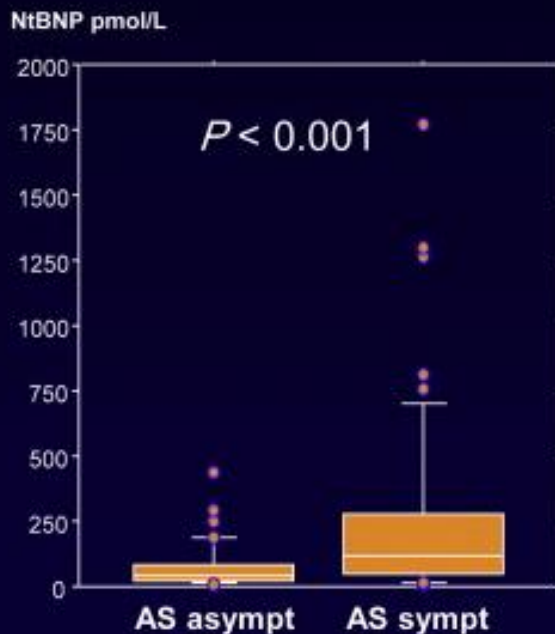
Markers of myocardial response to intracardiac pressure increases

Natriuretic peptides are significantly higher in symptomatic patients compared with asymptomatic patients with severe AS



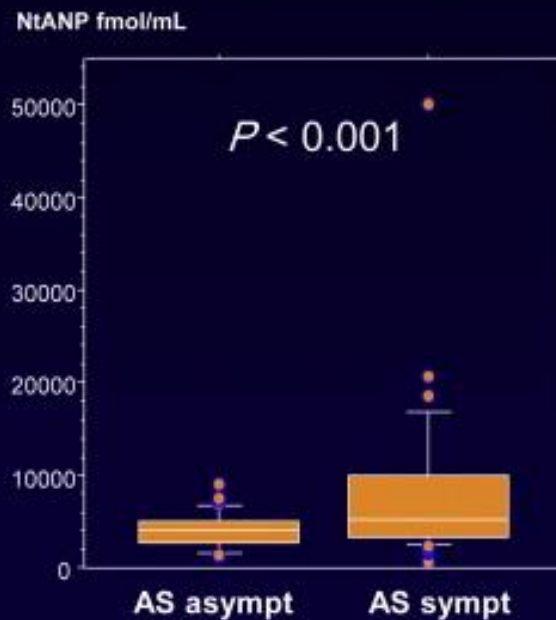
BNP

Median: 79 vs 172 pg/mL



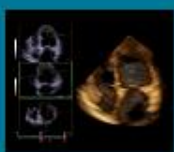
NT-proBNP

356 vs 1025 pmol/L



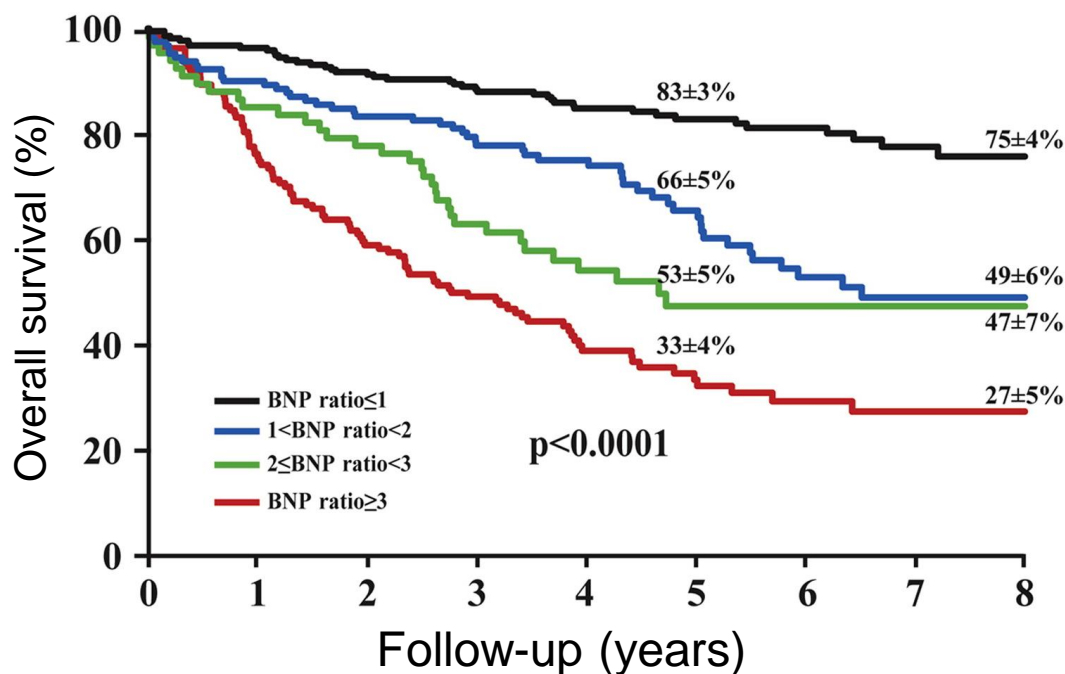
NT-proANP

4089 vs 5227 fmol/mL



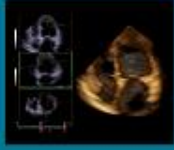
Natriuretic peptides

BNP clinical activation is associated with excess long-term mortality incrementally and independently of all baseline characteristics



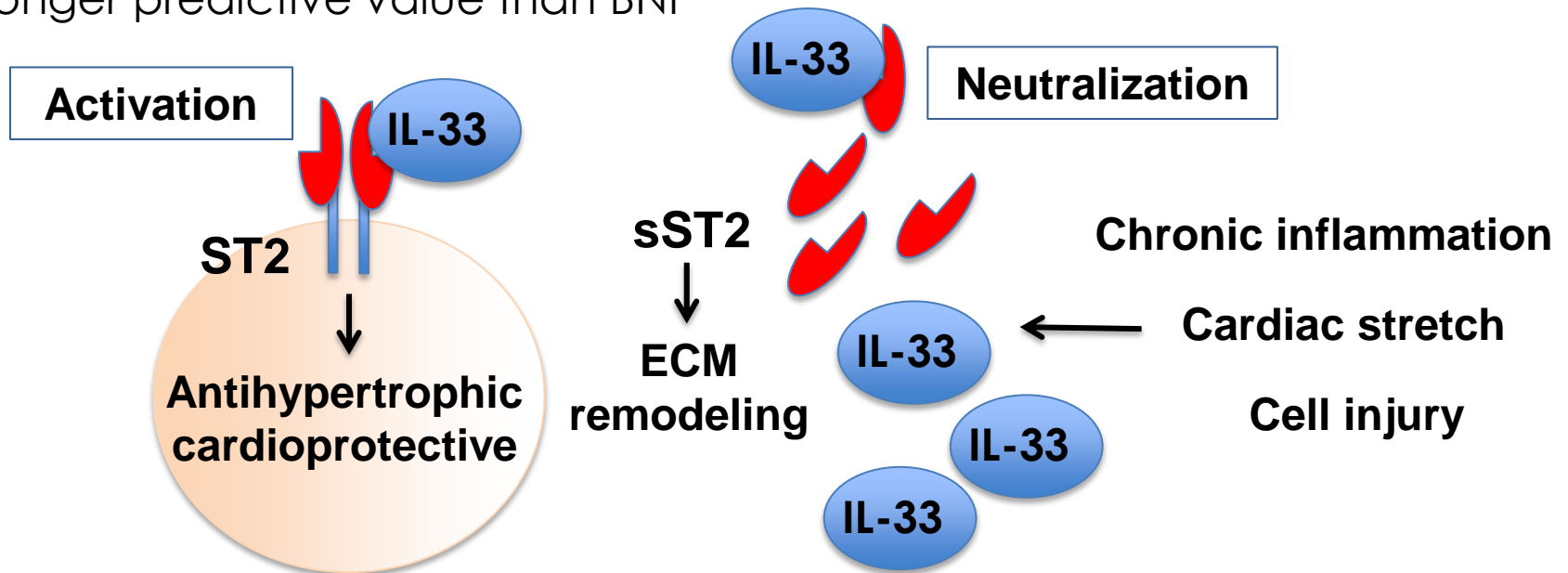
Asymptomatic with normal LVEF AS patients

BNP clinical activation = BNP ratio (measured BNP/maximal normal BNP value specific to age and sex) > 1

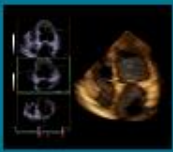


Soluble ST2

- Member of the IL-1 receptor family.
- Elevated concentrations of sST2: worse prognosis of acute and chronic HF
- Stronger predictive value than BNP

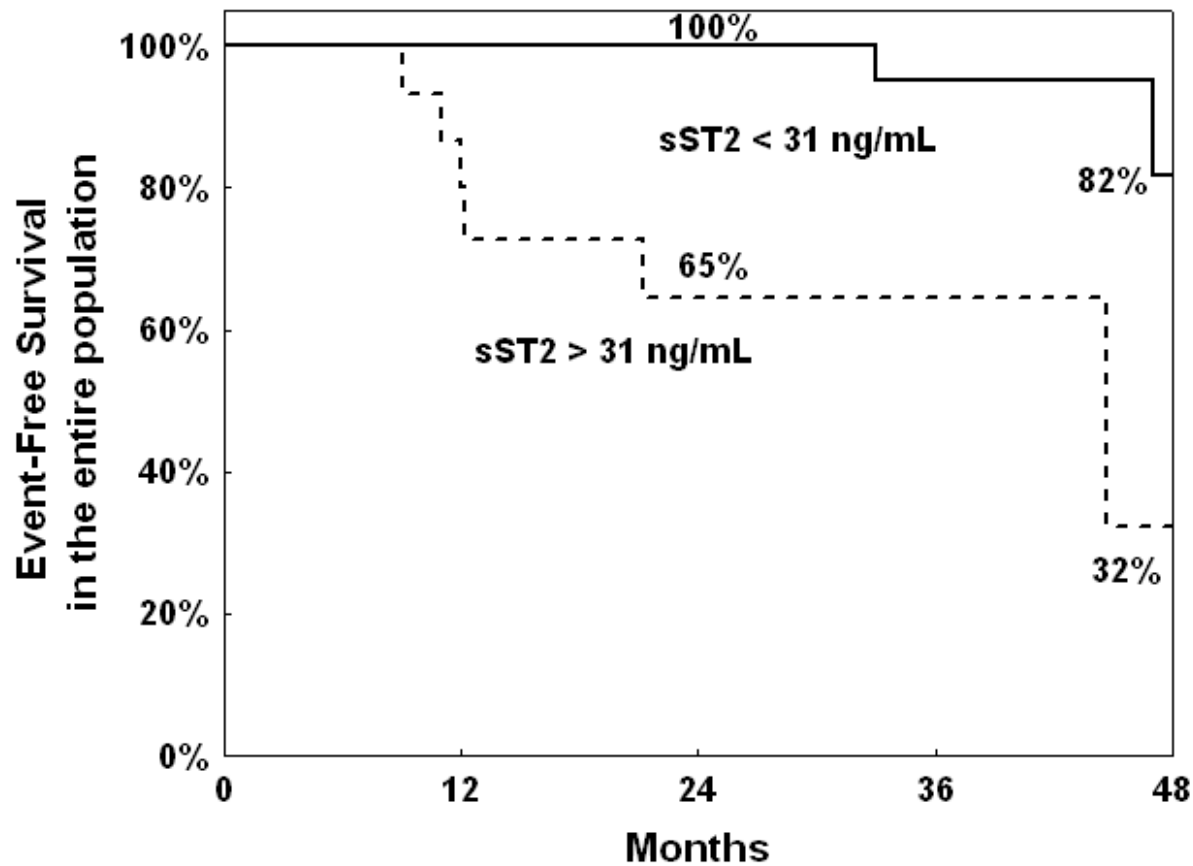


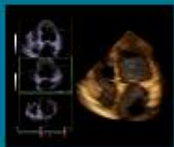
- Involved in at least three pathophysiological mechanisms of AS: inflammation/remodeling, fibrosis/cardiac stretch



Soluble ST2

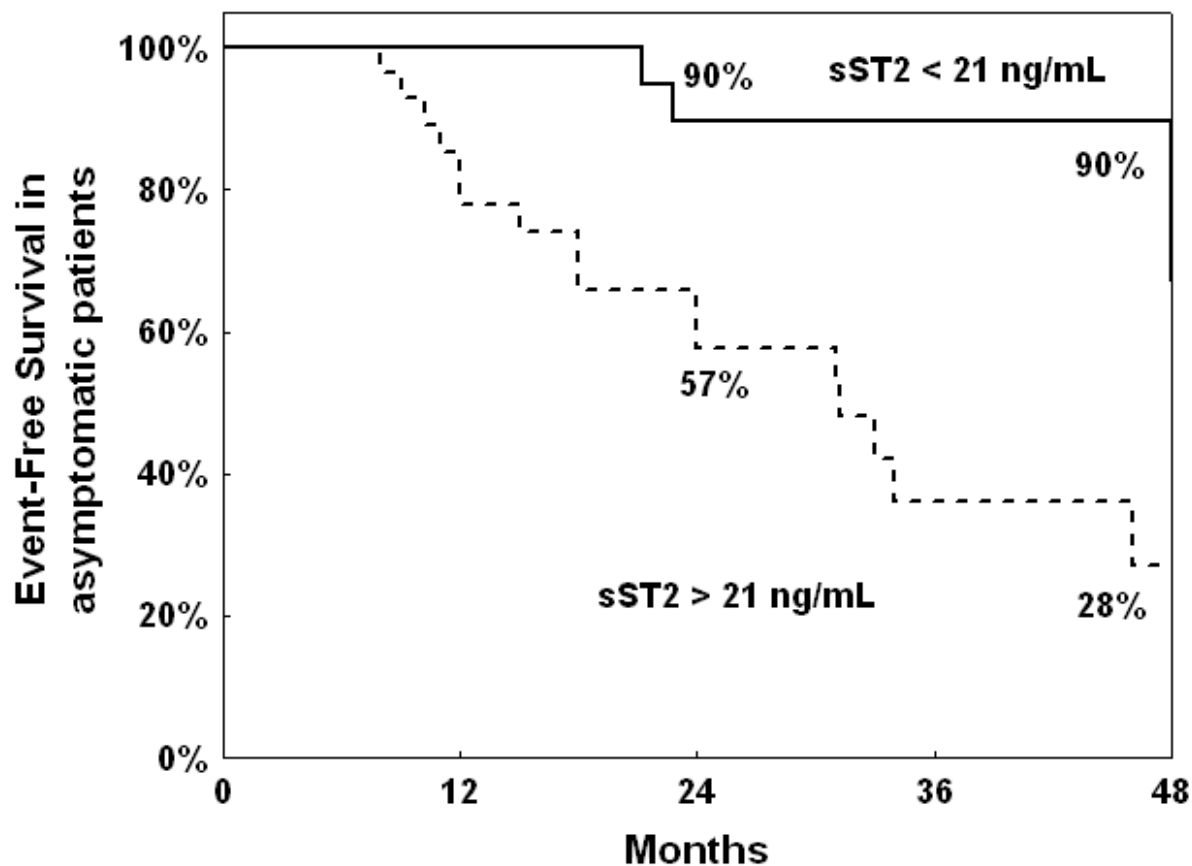
Independently predicts mortality in aortic stenosis

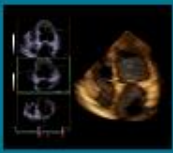




Soluble ST2

Independently predicts mortality in aortic stenosis
in asymptomatic patients

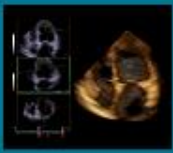




Troponins

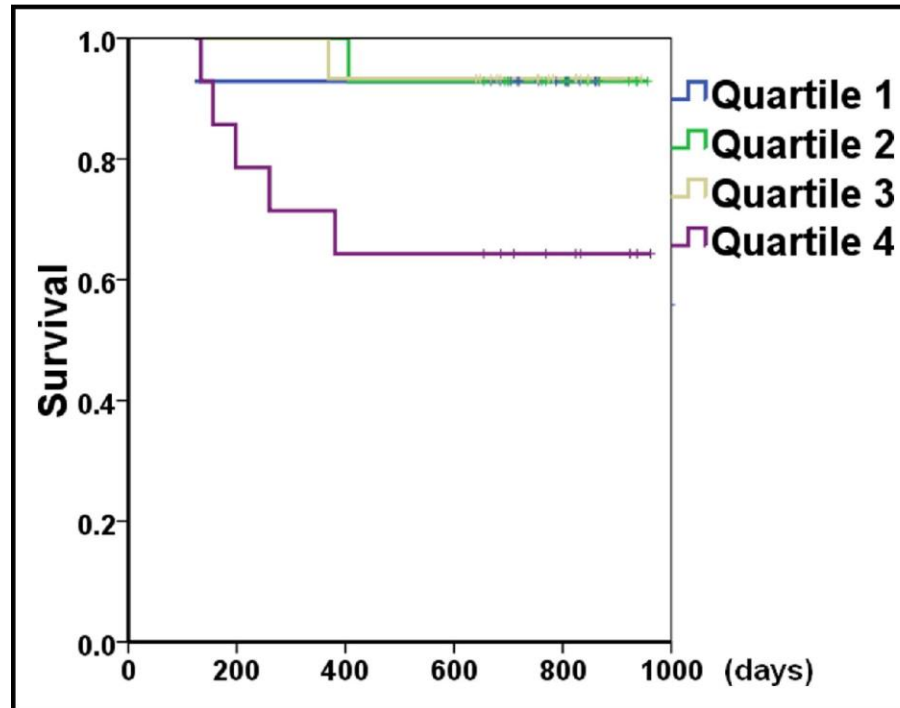
High sensitivity assays

cardiac troponins circulate in a variety of acute and chronic cardiac and non-cardiac disease conditions, including acute heart failure and chronic symptomatic and asymptomatic left ventricular dysfunction

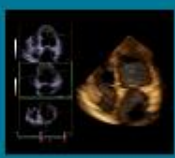


Troponins

High hs-TnT levels: worse prognosis in moderate to severe **AS**

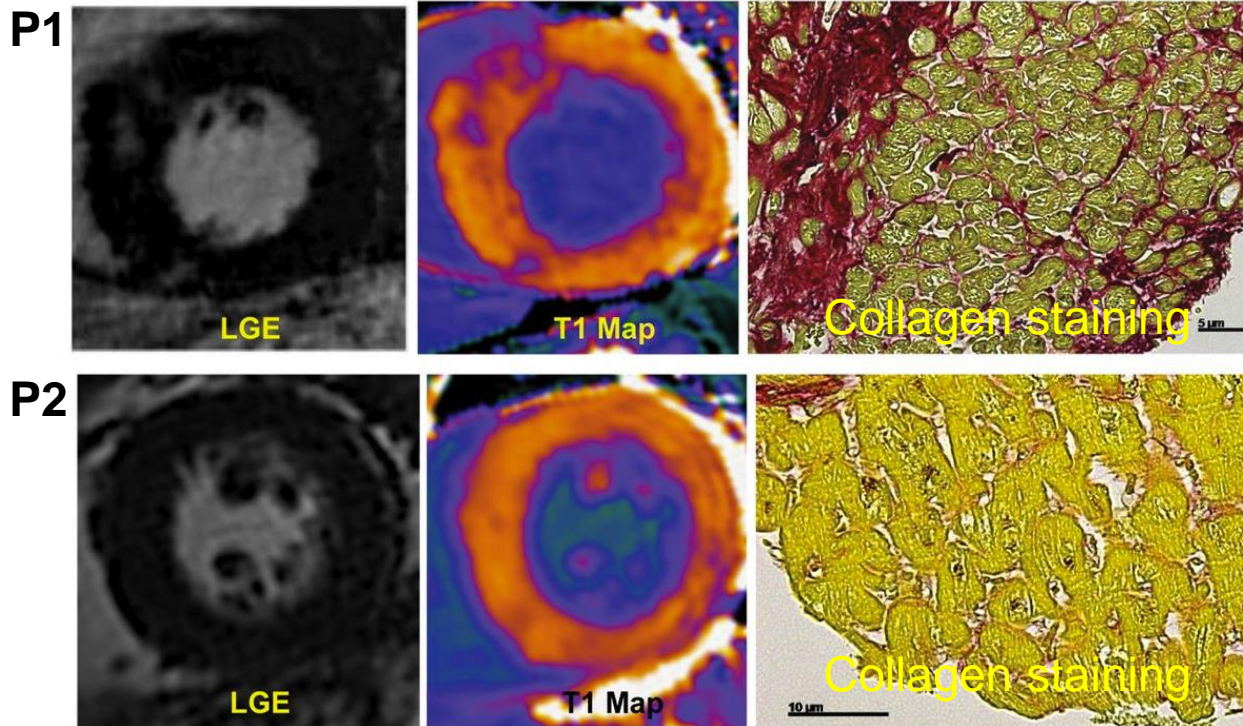


Left ventricular mass is a major determinant of circulating hs-TnT levels



Troponins

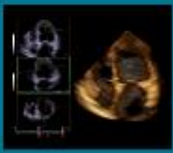
High-sensitivity troponin I concentrations: marker of advanced hypertrophic response and adverse outcomes in AS with normal LVEF



Peak aortic jet velocity:
4.8 m/s
LV mass index: 114 g/m²
Plasma cTnI: 11.9 ng/L

Peak aortic jet velocity:
5.1 m/s
LV mass index: 81 g/m²
Plasma cTnI: 2.5 ng/L

- High sensitivity troponins seem to reflect myocardial remodeling and fibrosis

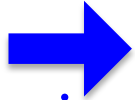


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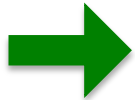


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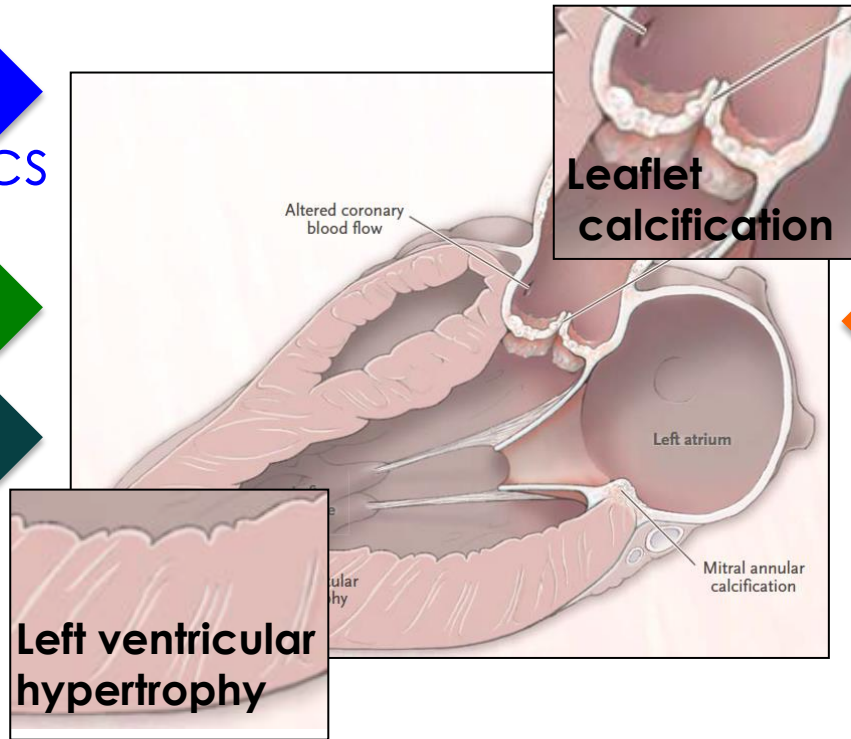
transcriptomics



genomics



epigenomics



proteomics



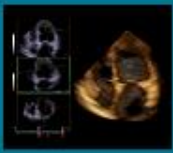
metabolomics



interactomics



- Analysis of biomolecules on a large scale
- Global integrated view of molecular and cellular processes
- Basis for a highly directed **personalised and predictive medicine**
- Impact on health and disease



Genomics

Genetic Associations with Valvular Calcification and Aortic Stenosis

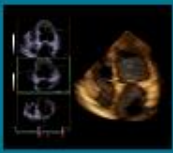
Genome wide association study (n=6942 patients)

- One SNP in the **lipoprotein(a)** locus (*LPA*) locus reaches genomewide significance for the presence of aortic valve calcification (CT scanning) across multiple ethnic groups
- Correlation of genetically determined Lp(a) levels and aortic valve calcification
- *LPA* genotype is associated with incident clinical aortic stenosis and aortic-valve replacement

Lp(a):

- cholesterol-rich particle (apolipoprotein B100 + apolipoprotein(a))
- risk factor for coronary artery disease
- accumulate in both early-stage and end-stage aortic-valve lesions

- Causal relationship Lp(a) / aortic valve disease
- Lowering Lp(a) levels to slow down disease progression ?



Transcriptomics

Small and long non-coding RNAs in cardiac homeostasis

Gene regulatory networks

eRNAs

mRNAs / Proteins

miRNAs

lncRNAs



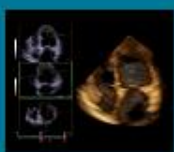
Cell migration

De- /
Differentiation

Cell Cycle Control

ECM Deposition /
Fibrosis

Targeted pathways

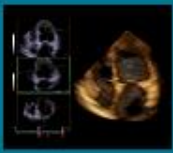


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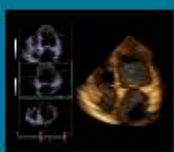
Studies on miRNA in aortic stenosis

Patient cohort	Samples	Observation	References
9 patients (AS versus aortic insufficiency requiring AVR)	Aortic valve leaflets	Decreased expression of miR-26a, miR-30b, and miR-195 in the aortic valves of patients requiring AVR due to AS	Nigam et al. <i>J Heart Valve Dis</i> 2010
46 AS patients requiring AVR	LV intraoperative biopsies	miR-133a predict regression of LV hypertrophy (1 year) after valve replacement	Villar et al. <i>Heart</i> 2011
19 bicuspid aortic valve versus 17 tricuspid aortic valve patients	aortic valve leaflets	Decreased expression of miR-141 in bicuspid aortic valves associated with increased BMP-2 and calcification	Yanagawa et al. <i>J Thorac Cardiovasc Surg</i> 2012
75 AS patients requiring AVR versus 32 surgical controls	LV intraoperative biopsies + plasma	High expression of miR-21 correlates with mean transvalvular gradient and LV fibrosis	Villar et al. <i>Int J Cardiol</i> 2013
5 AS patients before TAVI versus healthy controls	LV intraoperative biopsies + plasma	Decreased miR-1 correlates with increased soluble FABP3 in AS patients upon LVH	Varrone et al <i>J Am Coll Cardiol</i> 2013



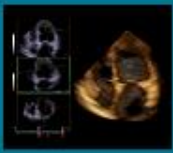
Studies on miRNA in aortic stenosis

Patient cohort	Samples	Observation	References
112 patients with moderate to severe AS versus 40 healthy controls	Plasma	levels of miR-1 , miR-133 , and miR-378 predict LVH in patients with AS miR-378 levels correlate with left ventricular mass index	Chen et al. <i>PLoS One</i> 2014
57 patients with moderate to severe AS versus 10 healthy controls	Plasma	Increased miR-210 levels in AS patients comparable to increment in NT-proBNP levels miR-210 levels associate with higher mortality (3.5 year follow-up)	Rosjo et al. <i>PLoS One</i> 2014
28 patients with moderate to severe AS versus 10 healthy controls	endomyocardial biopsies and necropsies	down-regulation of miR-122 in severe myocardial fibrosis in AS, through TGF-β1 up-regulation	Beaumont et al. <i>Clin Sci (Lond)</i> 2014
74 AS patients requiring AVR	LV intraoperative biopsies + plasma	miR-133a as a positive predictor of the hypertrophy reversibility after surgery	Garcia et al. <i>J Am Heart Assoc</i> 2013
10 AS patients requiring AVR	LV intraoperative biopsies	miRNA-30b regulates aortic valvular calcification and apoptosis through direct targeting of Runx2, Smad1, and caspase-3	Varrone et al <i>J Thorac Cardiovasc Surg</i> 2013



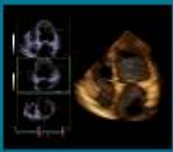
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Transcriptomics

- A few miRNA have been involved in LVH or fibrosis
- Circulating miRNA levels reflect myocardial expression
- Plasma levels of miR-210 correlate with BNP levels and increased mortality

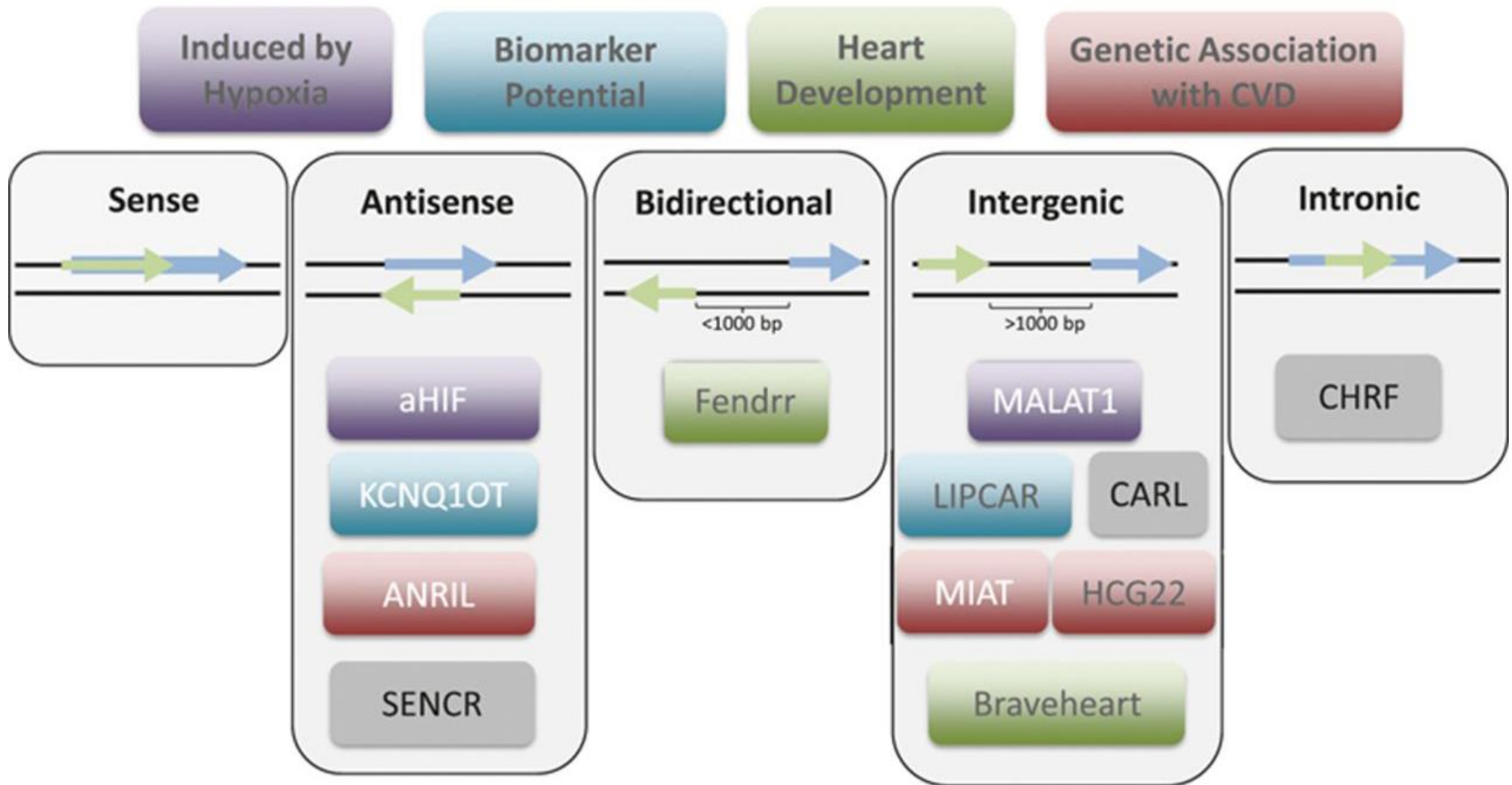


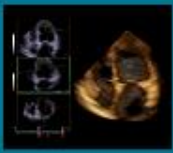
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Transcriptomics

Long non-coding RNAs: novel attractive biomarkers





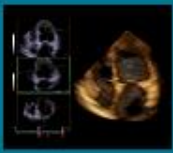
Transcriptomics

Genome-wide profiling of the cardiac transcriptome after myocardial infarction identifies novel heart-specific long non-coding RNAs

- Mouse model
- heart-specific lncRNAs
- relevant to maladaptive remodelling, cardiac function

lncRNAs in aortic stenosis

Downregulation of Novlnc44
in LV biopsies from AS patients



Biomarkers: the future is bright

Biomarkers will be more and more used in patients with VHD

- in the diagnostic work-up
- to characterize the symptomatic status
- to predict the outcome
- to evaluate the impact of treatment

➤ Toward a multi-biomarker approach ?