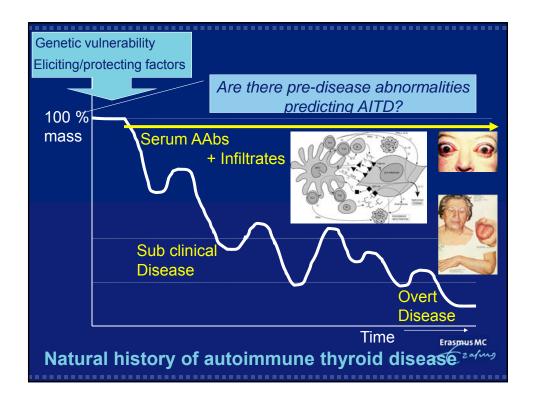


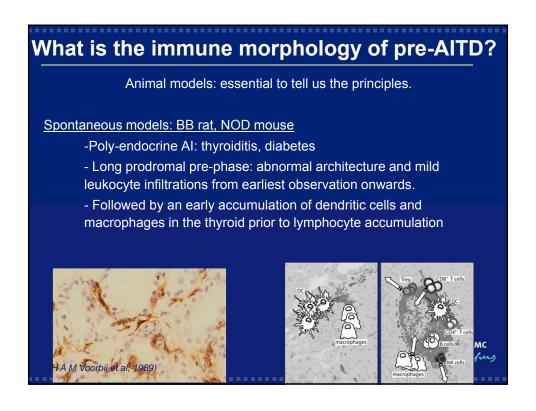
Learning objectives

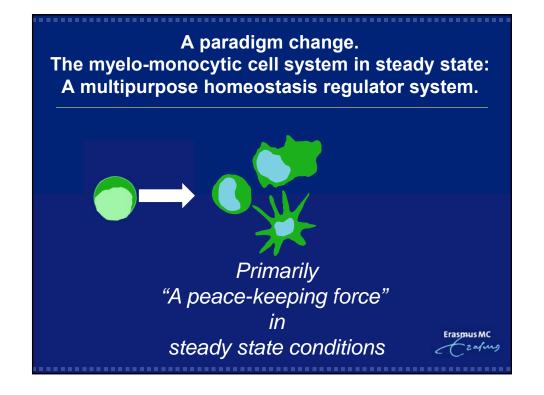
Understand that:

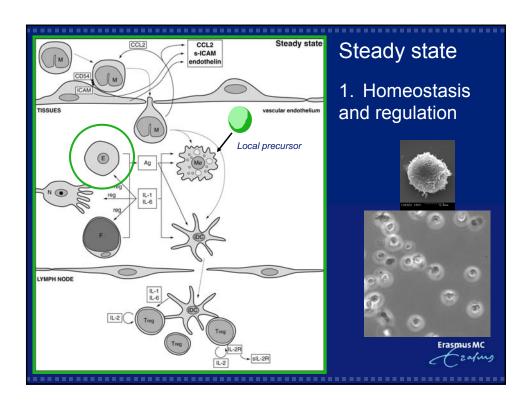
- role myelomonocytic cells in regulation and tissue homeostasis and development of autoimmunity
- reduction of T regulatory cells leads to loss of tolerance
- target organ abnormalities precede autoimmunity
- serum analytes in individuals at risk for development of Al reflect these abnormalities

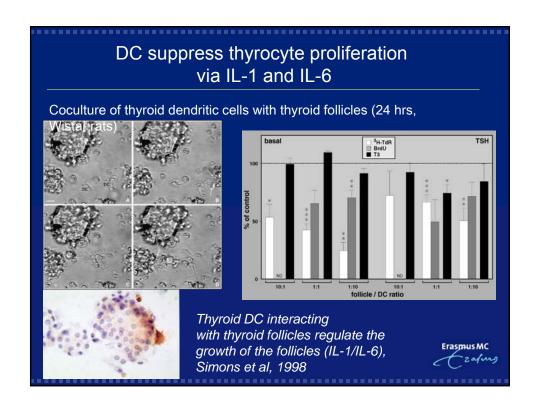


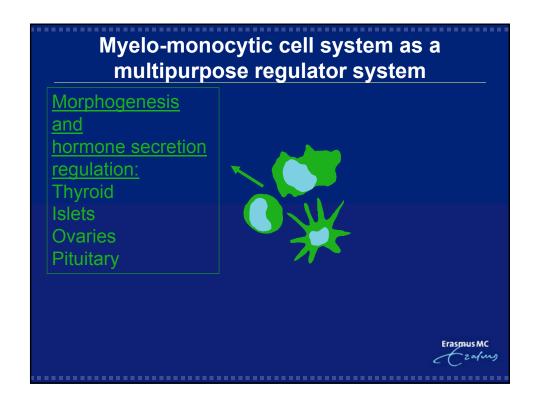


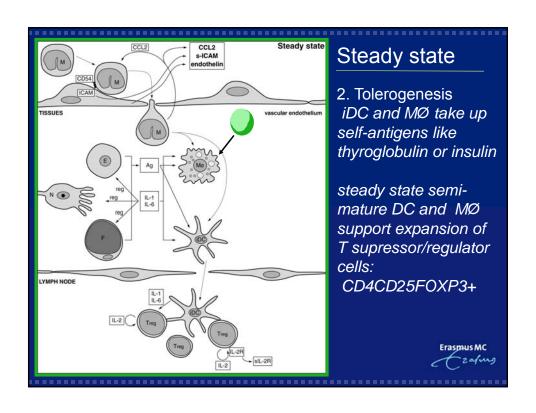


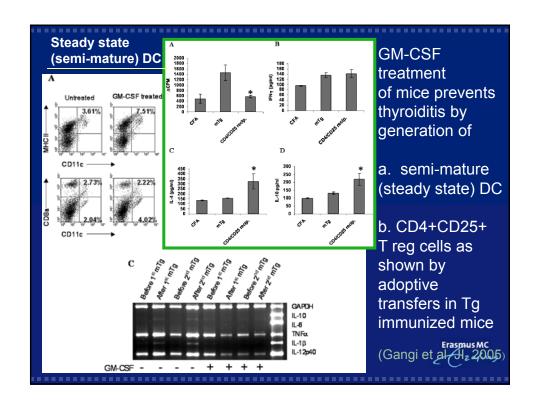


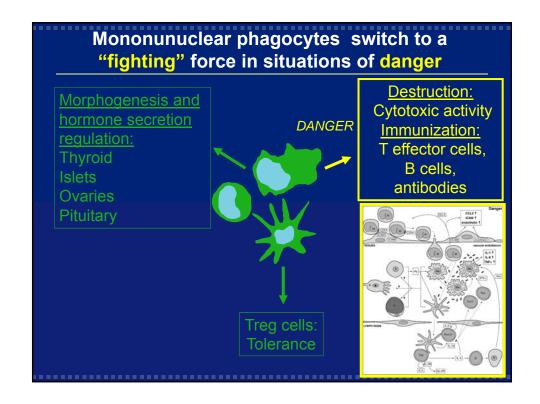


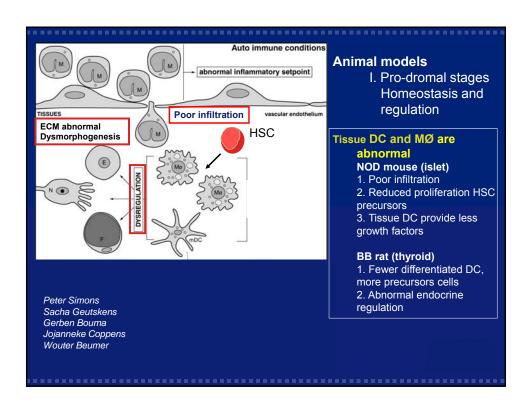


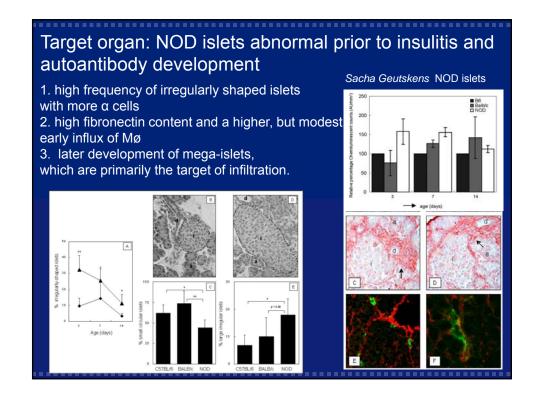


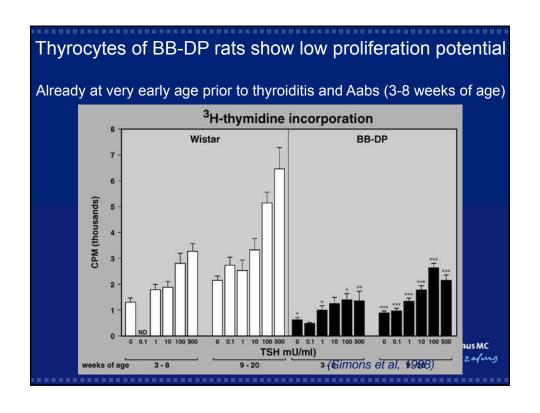


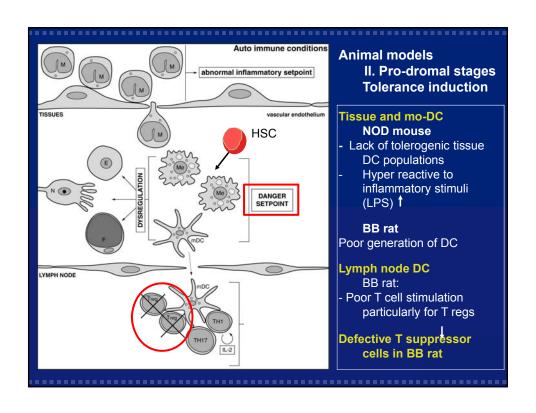












Pro-dromal phases in animal models

Endocrine tissue

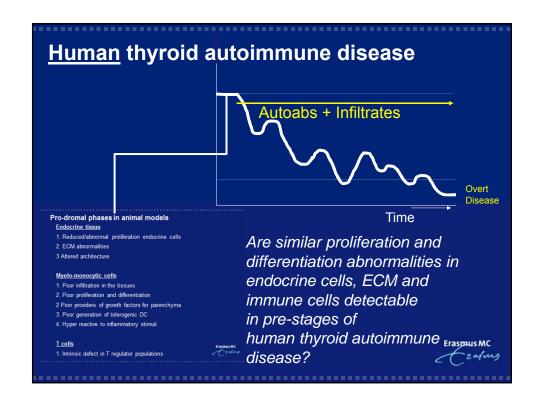
- 1. Reduced/abnormal proliferation endocrine cells
- 2. ECM abnormalities
- 3 Altered architecture

Myelo-monocytic cells

- 1. Poor infiltration in the tissues
- 2. Poor proliferation and differentiation
- 2 Poor providers of growth factors
- 3. Poor generation of tolerogenic DC
- 4. Hyper reactive to inflammatory stimuli

T cells: Intrinsic defect in T regulator populations

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Amsterdam AITD cohort

Female 18-65 years old

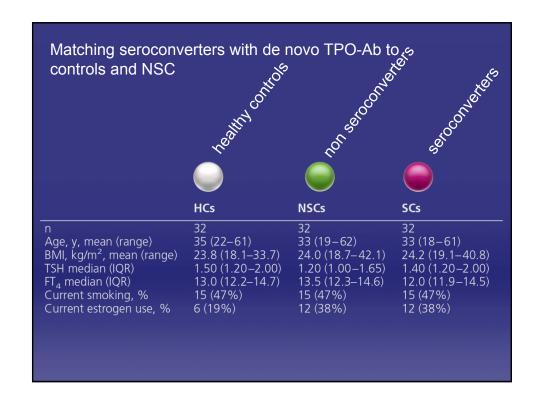
At least one 1st or 2nd degree relative with AITD

No personal history of thyroid disease

5 years follow up

Annual visits & blood testing: TSH, FT4, T3, TPO-Ab, Tg-Ab, TSH-R Ab smoking habits use of oral contraceptives or other estrogen

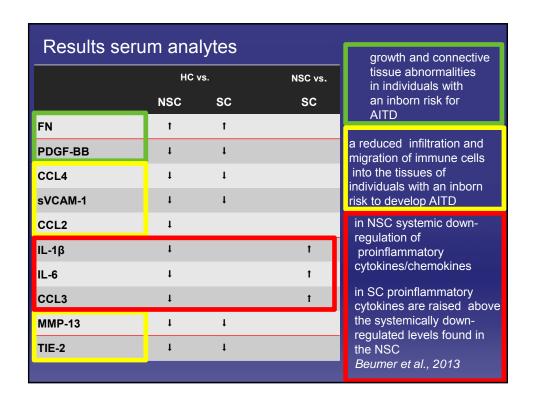
Current pregnancy: exclusion criterion



Results serum analytes							
	HC vs.		NSC vs.				
	NSC	sc	sc				
FN	Ť	t					
PDGF-BB	Ţ	1					
CCL4	1	1					
sVCAM-1	1	1					
CCL2	1						
IL-1β	Ţ		†				
IL-6	1		t				
CCL3	1		t				
MMP-13	1	1					
TIE-2	1	1					

Results serum analytes						
	НС	vs.	NSC vs.	growth and connective tissue abnormalities in individuals with an inborn risk for		
	NSC	sc	sc			
FN	Ť	t		AITD		
PDGF-BB	ı	1				
CCL4	1	1				
sVCAM-1	1	1				
CCL2	1					
ΙL-1β	Ţ		†			
IL-6	1		t			
CCL3	1		1			
MMP-13	ı	1				
TIE-2	ı	1				

Results serum analytes							
	HC vs.		NSC vs.	growth and connective tissue abnormalities in individuals			
	NSC	sc	sc	with an inborn risk for AITD			
FN	t	t					
PDGF-BB	1	Ţ		a reduced infiltration and migration of immune cells into the tissues of individuals with an inborn risk to develop AITD			
CCL4	1	Ţ					
sVCAM-1	Ţ	1					
CCL2	Ţ						
IL-1β	1		1				
IL-6	1		t				
CCL3	1		t				
MMP-13	Ţ	Ţ					
TIE-2	1	Ţ					



Overall conclusion animal models

The proneness to develop endocrine autoimmune disease (before sero-positivity) is characterized by

- 1. Growth and ECM abnormalities of the endocrine tissue,
- 2. Growth and differentiation abnormalities of the myelomonocytic lineage leading to
 - a poor development of DC and MØ particularly of those with a tolerogenic function and
 - an inflammatory hyper reactivity to LPS of such DC and MØ
- 3. Defects in T regulator cell populations.

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Overall conclusions human study

There are indications that the pro-dromal stage of thyroid autoimmunity in humans at risk (family members) can be detected – similar to the abnormal processes in animal models –

by studying serum analytes reflecting

- 1. growth and ECM abnormalities of endocrine tissues (e.g. PDGF-BB, FN)
- 2. poor development of myelo-monocytic cells (e.g. DC and MØ cytokines)
- 3. poor infiltration capacity of myelomonocytic cells (e.g. chemokines)

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