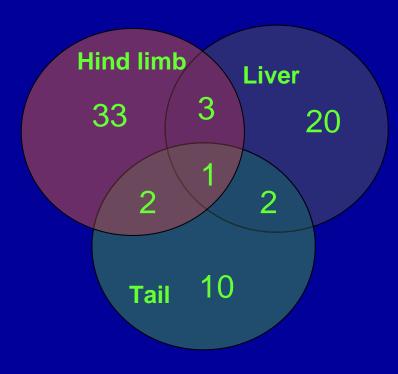
Conditions for the Amphibian Metamorphosis Assay

Test animal			Xenopus laevis larvae							
Test su	ubstance		XXX							
Age at test initiation (post fertilization day)			About 14 days (from fertilization to stage 51)							
Exposure period			21 days (from stage 51)							
Concentration			5 (4 concentrations and control)							
Replication			4 replications per test concentration and control							
Number of tanks			20 tanks							
Larval density			20 tadpoles per 4L tanks							
Number of larvae			400 tadpoles (20 tadpoles / tank × 20 tanks)							
Day 7			100 tadpoles (5 tadpoles / tank × 20 tanks)							
Numbe	Number of determination Day 21		300 tadpoles (15 tadpoles / tank × 20 tanks)							
Endpoints			5 points (Whole body length, snout-vent length, hind limb length, developmental stage, thyroid histology)							
Number of thyroid histology			100 tadpoles (5 tadpoles / tank × 20 tanks)							
	Sample schedule from tanks		4 times (Once a week)							
Anal ysis chem	Number of sample from tanks		64 (4 concentration /times × 4replication × 4 times)							
	Sample schedule from stock solution		5 (Day 0, 4, 8, 13, 17)							
istry	Number of sample from stock solution		10 (1concentration /time × 2 replication /concentration 5 times)							
	Number of total sam	ple	74							

Identified T₄ Responsive Genes on Metamorphosis Assay

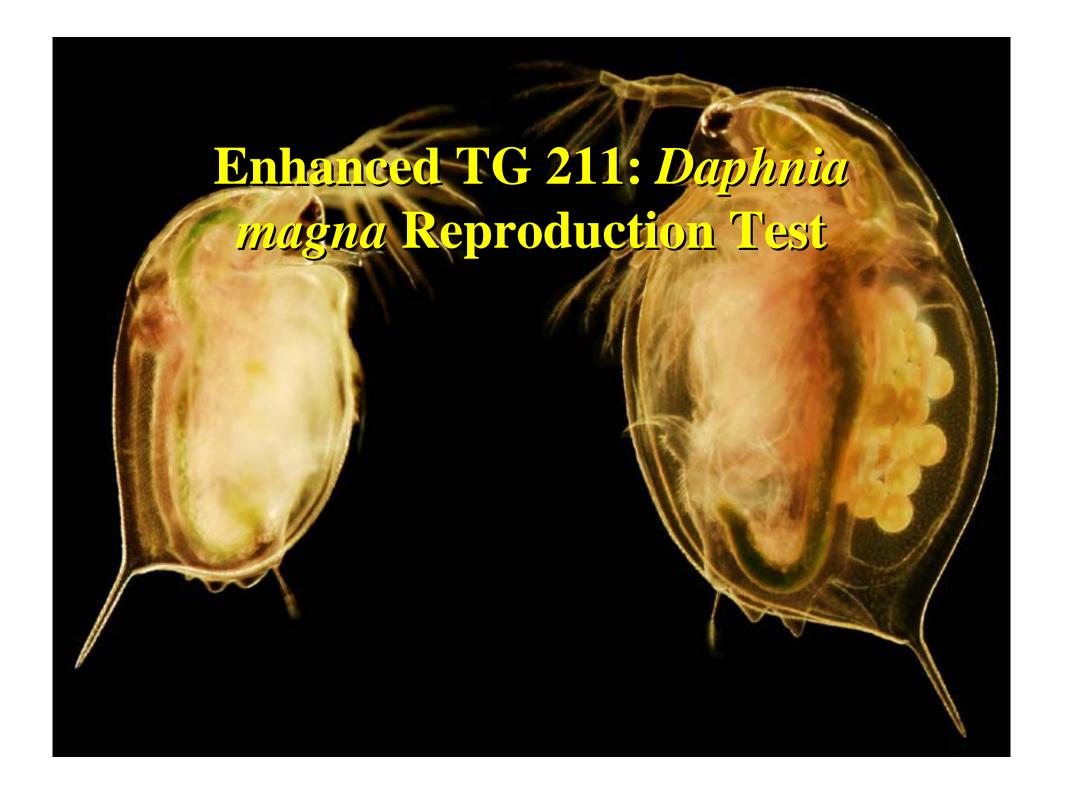
About a gene identified by a gene expression profile, acquired the individual genetic information such as sequence and the gene function from database and classified it in known gene or unknown gene.



Identified genes on each tissues

	簡易同定名	Groupe	発生における 発現量の増減	機能別	T4応答に関す る文献	а	相同 b		高いii d	佳伝子 e	f
	actin, alpha, cardiac muscle fibrillin (Homo sapiens)	I I	++	1	_	0		0			
	MyHC (myosin heavy chain) (Cynops pyrrhogaster)	I	+	i	Sachs 1997			ő			
	myosin light chain, fast skeletal muscle (<i>Xenopus laevis</i>)	I	+	1	Moutou 2002		0	_			
	myozenin 1 nebulin (<i>Mus musculus</i>)	I I	++	1	-	0		0			
	titin (O.cuniculus)	i	+	i				ő			
	troponin I, skeletal, fast 2	i	+	1	-	0		_			
	Xenopus laevis, myosin light chain (MLC1)	I	++	1	Moutou 2002		00				
	activated protein kinase C receptor (RACK1) (Xenopus laevis) caspase 6, cysteine protease	i	+	5		0	0				
	mitochondrial (12S rRNA, tRNA-Val, 16S ribosomal RNA gene)	I	+	8		ō					
	unknown 15 (EST/NIH) unknown 16 (EST/NIH)	1	++	13					00		
	unknown 18 (EST/NIH)	i	+	13					0		
	unknown 20 (EST/NIH)	I	+	13					ō		
	unknown 21 (EST/NIH laevis) unknown 24 (EST/NIH)	1	++	13					0	0	
윹	unknown 24 (EST/NIH) unknown 25 (EST/NIH)	i	+	13					0		
Hind limb	unknown 26 (EST/NIH)	I	+	13					0		
듶	unknown 27 (EST/NIH)	1	+	13					0		_
	unknown 28 Unknown (栗再解析)	1	++	13							00
=	thymopoietin	i	-	7		0					-
	RNA-binding protein (TPA) (Mus musculus) calnexin	I		9 10		0		0			
	unknown 19 (EST/NIH)	;	2	13		0			0		
	unknown 22 (EST/NIH)	i	_	13					ŏ		
	polycystin-2 (pkd2) (<i>Drosophila melanogaster</i>) betaine-homocysteine methyltransferase (<i>Homo sapiens</i>)	II II	++	2 6				00			
	unknown 23	II II	Ŧ	13	_			0			0
	thioredoxin-like 1	II	-	3		0					
	proliferating cell nuclear antigen (PCNA)	II		4		0	0				
	ran GTP-binding protein, RAS family (<i>Xenopus laevis</i>) homeo box A7	II II	Ξ	9		0	0				
	Unknown(要再解析)	II	-	13		_					0
	similar to CYSTATIN A1 (<i>Xenopus laevis</i>) DEAD-box protein p72 (P72) (Human)	II	+	5 11	-		0	0			
	Unknown(要再解析)	II II	+	13				0			0
	nonmuscle myosin II heavy chain A (Xenopus laevis)	I	+	1	Sachs 1997		0				
	ATPase, H+ transporting, V0 subunit B thioredoxin-like 1	I I	++	2	Das 2006	00					
	serine (cysteine) proteinase inhibitor, clade C, member 1 (Xenopus laevis)	i	+	6	Δ		0				
	mitochondrial	I	+	8		0					
	mitochondrial (X.laevis) mitochondrion	I	++	8		00					
	MTERF domain containing 1	i	+	8	_	ŏ					
Liver	CCAAT-enhancer binding protein delta (Xenopus laevis) (C/EBP)	I	+	9			0				
	zinc finger transcription factor slug (slug) gene (SNAI2) eukaryotic translation elongation factor 2	1	++	9 10	_	00					
	unknown 12 (EST/NIH)	i	Ŧ	13	_	0			0		
	Unknown(要再解析)	I	+	13							0
	Unknown(要再解析)	I	+	13 6	Ness 2000		0				0
	HMG-CoA reductase mRNA (Xenopus laevis) unknown 10	I	Ξ	13	Ness 2000		0				0
	unknown 8 (EST/NIH)	I	-	13					0		
	serine (or cysteine) proteinase inhibitor, clade A (alpha-1), member 1 unknown 11 (EST/NIH)	II	++	6 13	Δ		0		0		
	unknown 14	II II	+	13							0
	ribosomal protein S15	II	+	10	-	0					
	unknown 13 (EST/NIH) similar to chemokine receptor 1 (<i>Xenopus laevis</i>)	II II	+	13 12	_		0		0		
	unknown 9 (EST/NIH)	II	-	13					0		
	unknown 17 (EST/NIH)	II	-	13					0		
		II	+	6	-			0			
	betaine-homocysteine methyltransferase (Homo sapiens) titin (Homo sapiens)					0					
	titin (Homo sapiens) mitochondrion (16 rRNA, tRNA-Leu, Ile, Gln, NADH dehydrogenase 1)	I	+	8	Das 2006						
	titin (<i>Homo sapiens</i>) mitochondrion (16 rRNA, tRNA-Leu, Ile, Gln, NADH dehydrogenase 1) eukaryotic translation elongation factor 2	I	+	10	Das 2006 -	0			_		
	titin (Homo sapiens) mitochondrion (16 RRNA, tRNA-Leu, Ile, Gln, NADH dehydrogenase 1) eukaryotic translation elongation factor 2 unknown 2 (EST/NIH)	I I I	++	10 13	-	ō			0		
	titin (<i>Homo sapiens</i>) mitochondrion (16 rRNA, tRNA-Leu, Ile, Gln, NADH dehydrogenase 1) eukaryotic translation elongation factor 2	I	+	10	Veldhoen 2002			0	0		
75	titin (*Mome aspiers) mitochondrion (fl. RTNAL ttNA-Leu, Ile, Glin, NADH dehydrogenase 1) eukaryotic translation elongation factor 2 unknown 2 (EST/NIH) 14-3-3 thyroid hormone receptor interactor 12 (*Mus musculus*) unknown 1 (EST/NIH)	I I I I I	++	10 13 5 9 13	-	ō			0		
Tail	titin (Home aspiens) mitochondrion (If RRNA: tRNA-Leu, Ile, Glin, NADH dehydrogenase 1) eukaryotic translation elongation factor 2 unknown 2 (EST/NIH) 14-3-3 thyroid homone receptor interactor 12 (Mus musculus) unknown 1 (EST/NIH) betaine-homorysteine methyltransferase (Homo sapiens)	I I I I	+ + - - - +	10 13 5 9 13	-	ō	0	0	_		
Tail	titin (*Mome aspiers) mitochondrion (fl. RTNAL ttNA-Leu, Ile, Glin, NADH dehydrogenase 1) eukaryotic translation elongation factor 2 unknown 2 (EST/NIH) 14-3-3 thyroid hormone receptor interactor 12 (*Mus musculus*) unknown 1 (EST/NIH)	I I I I I	++	10 13 5 9 13	-	ō	00		_		
Tail	titin (Homo aspiers) mitochondrion (If RRNA, IRNA-Leu, Ile, Glin, NADH dehydrogenase 1) eukaryoto translation elongation factor 2 unknown 2 (EST/NHH) 14-3-3 thyroid hormone receptor interactor 12 (Mus musculus) unknown 1 (EST/NHH) unknown 1 (EST/NHH) service elongation (Homo aspiers) service (Homo aspiers) service (Homo aspiers) ribasomal protein L17 (Kenopus Isevic) unknown 4 (Mot Clavis)	I I I I II II	+ + - - - +	10 13 5 9 13 6 5 10	-	ō			0	0	
Tail	titin (Home aspiens) mitochondrion (If RTNA: tRNA-Leu, Ile, Glin, NADH dehydrogenase 1) eukaryotic translation elongation factor 2 unknown 2 (EST/NIH) 14-3-3 thyroid hormone receptor interactor 12 (Mus musculus) unknown 1 (EST/NIH) betaine-homocysteine methyltransferase (Homo sapiens) similar to CYSTATIN AI (Xenopus laevis) unknown 4 (MGC Jeevis) unknown 4 (MGC Jeevis) unknown 5 (EST/NIH)	I I I I II II	+ + - + + -	10 13 5 9 13 6 5 10 13	-	ō			0	0	
Tail	titin (Homo aspiers) mitochondrion (If RRNA, IRNA-Leu, Ile, Glin, NADH dehydrogenase 1) eukaryoto translation elongation factor 2 unknown 2 (EST/NHH) 14-3-3 thyroid hormone receptor interactor 12 (Mus musculus) unknown 1 (EST/NHH) unknown 1 (EST/NHH) service elongation (Homo aspiers) service (Homo aspiers) service (Homo aspiers) ribasomal protein L17 (Kenopus Isevic) unknown 4 (Mot Clavis)	I I I I II II	+ + - + + -	10 13 5 9 13 6 5 10	-	ō			0	0	

- アフリカツメガエル他、カエルで同定されている遺伝子
- カエルでは同定されていない遺伝子
- ニシツメガエルのESTデータペースが存在する遺伝子 アフリカツメガエルのESTデータベースならば存在する遺伝子
- 相同性の高いデータベースが存在しない遺伝子



Different Endocrine Systems Between Vertebrates and Invertebrates

Vertebrates (e.g., Mammals)

Invertebrates (e.g., Arthropods)

Androgen Estrogen

DES EE2, etc.

Nonylphenol Bisphenol A, etc.

Ecdysteroid
Juvenile hormone

Insect Growth Regulators (IGRs)

Dieldrin?

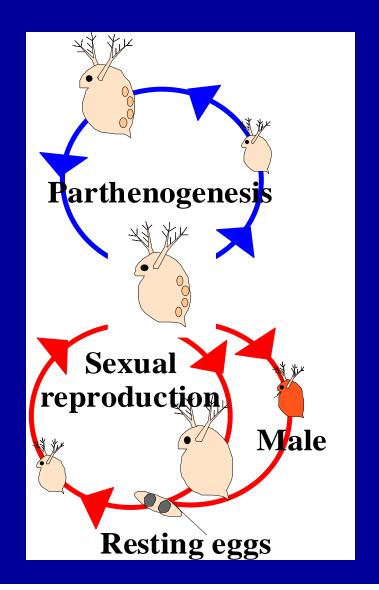
Crustacean Reproduction Toxicity Tests in OECD

- Draft DRP on mysid life cycle toxicity test (U.S.A)
- Draft proposal on a new test guideline copepod development and reproduction test (Sweden)
- *Daphnia magna* reproduction test (TG 211; OECD 1998)
 - Proposal for enhanced TG 211 for endocrine disrupting chemicals

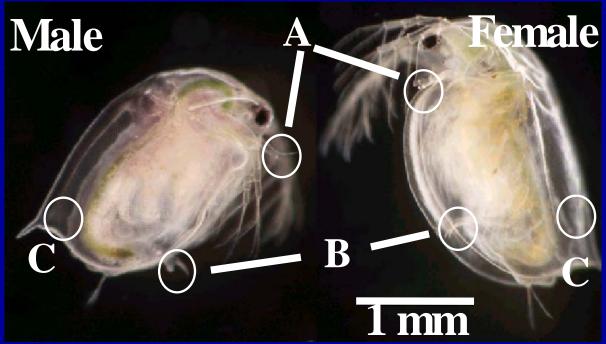
Reproductive System and Sex Determination in Cladocerans

- Cyclic parthenogenesis
- Environmental sex determination
- Emergence of males and initiation of sexual reproduction by
 - -Short photoperiod
 - -Low food concentration
 - -High population density, etc.

adverse conditions sexual cycle



Sex Differentiation



brood chamber

A: 1st antenna

B: post abdominal claw

C: abdominal process

Juvenile Hormones and Their Analogs

Juvenile hormones

juvenile hormone III

methylfarnesoate

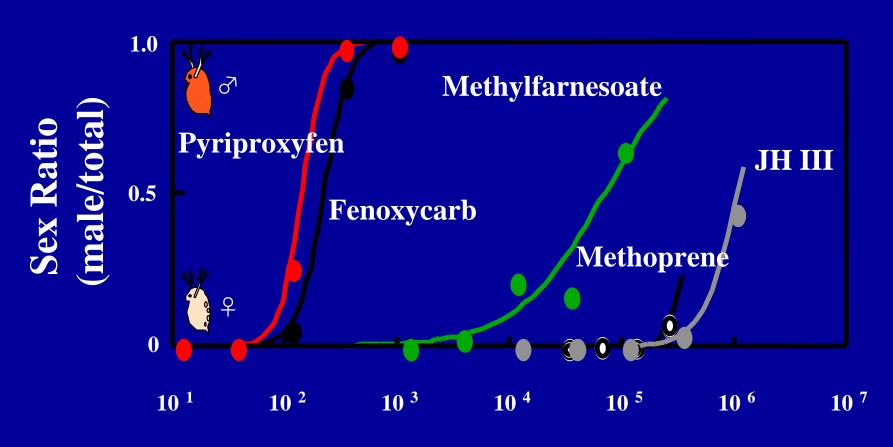
Juvenile hormone analogs (Insect Growth Regulators; IGRs)

methoprene

fenoxycarb

pyriproxyfen

Change in Offspring Sex Ratio Exposed to Juvenoids



Concentration (ng/L)

Other Daphnids in the Genus *Moina*

M. macrocopa

M. micrura





Other Daphnids in the Genus Ceriodaphnia

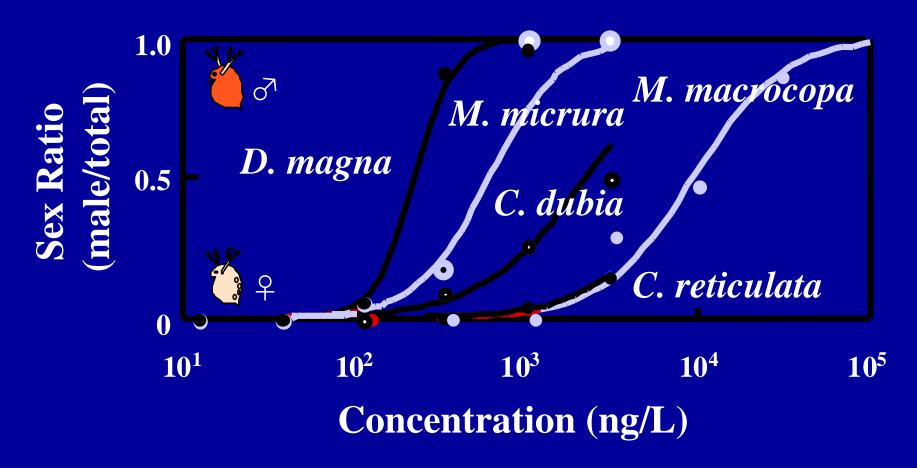
C. dubia

C. reticulata





Change in Offspring Sex Ratio Exposed to Fenoxycarb

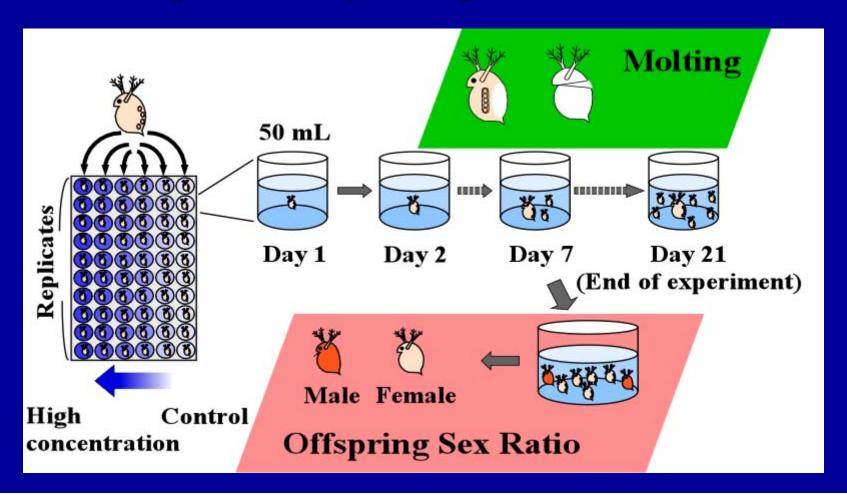


JH analogs increase male production in cladoceran

Switching from asexual to sexual cycle can be a marker for endocrine disruption

Experimental Design

Based on OECD TG 211 Daphnia magna reproduction test



List of Participants in the Ring Test

- 1. Finnish Environment Institute, Finland
- 2. French National Institute for Industrial Environment and Risks (INERIS), France
- 3. Laboratoire Ecotoxicité et Santé Environnementale (ESE) Equipe CNRS UMR, France
- 4. Aachen University, Germany
- 5. Bayer CropScience AG, Germany
- 6. Institute for Biological Analysis and Consulting (IBACON), <u>Germany</u>
- 7. UMWELTBUNDESAMT (UBA), Germany
- 8. Laboratory of Hydrobiology, <u>Hungary</u>
- 9. National Institute of Health, Italy
- 10. Agricultural Chemicals Inspection Station (ACIS), <u>Japan</u>
- 11. Kureha Special Laboratory, Co., Ltd., <u>Japan</u>
- 12. National Institute for Environmental Studies (NIES), <u>Japan</u>