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研究課題名(和文)眼炎症疾患発症におけるmiRNAの関与

研究課題名(英文) Roles of miRNA in the development of ocular inflammatory diseases

研究代表者

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研究成果の概要(和文):BALB/cマウスをブタクサとアラムで全身感作し、10日後ブタクサを点眼することによりアレルギー性結膜炎を誘導した。ブタクサ点眼24時間後、結膜を採取し、RNAを抽出した。全身感作したマウスから、ブタクサを点眼する直前に採取した結膜を対照とした。microarray法により、miRNAの発現を比較した。その結果、結膜炎誘導によりmiRNAの発現倍率が1/2倍未満となった遺伝子は44個、発現倍率が2倍以上となったmiRNAが74個同定された。さらに、統合評価を行い、Lrp2 binding proteinやlactoperoxidaseなどの標的遺伝子を同定した。

研究成果の概要(英文): Balb/c mice were immunized with ragweed in alum. Ten days later, the mice were cha llenged with ragweed in eye drops to induce allergic conjunctivitis. Twenty-four hours after the challenge, conjunctivas were harvested to extract RNA. As a control, conjunctivas were collected from immunized mic e just prior to ragweed challenge. Expression of miRNA were compared between the two groups by using micro array analysis. Forty-four miRNAS were identified to be downregulated to be less than half by induction of conjunctivitits, while 74 miRNAS were identified to be upregulated to be more than 2-fold by induction of conjunctivitis.

Combined evaluation identified the target genes of miRNAS such as Lrp2 binding protein and lactoperoxidase

研究分野: 医歯薬学

科研費の分科・細目:外科系臨床医学・眼科学

キーワード: 眼炎症 アレルギー性結膜炎 マイクロRNA ocular inflammation allergic conjunctivitis mic

roRN/

1.研究開始当初の背景

ぶどう膜炎、アレルギー性結膜疾患などの 眼炎症疾患の発症には免疫系細胞を中心に 産生されるサイトカインが重要な役割を果 たす。

サイトカイン産生は細胞内あるいは細胞外からのシグナルにより制御される。近年、細胞内に存在する長さ 20 から 25 塩基ほどの1本鎖ノンコーディング RNA (microRNA)がサイトカインをはじめ種々の遺伝子の発現を調節する機能を有することが明らかとなった。

2. 研究の目的

本研究の目的は、ぶどう膜炎、アレルギー性結膜疾患の発症に microRNA がどのように関与しているかを検討することである。

3.研究の方法

ぶどう膜炎、アレルギー性結膜疾患の動物 モデルにおける microRNA 発現の経時的変化 を網膜ならびに結膜で評価する (microarray 法、定量的 RT-PCR (reverse transcriptionpolymerase chain reaction)法)。

4. 研究成果

(1) BALB/c マウスをブタクサとアラムで全身感作し、10 日後ブタクサを点眼することによりアレルギー性結膜炎を誘導した・ブタクサ点眼 24 時間後、アレルギー性結膜炎を誘導したマウスから結膜を採取し、RNA を抽出した(n=3)・全身感作したマウスから、ブタクサを点眼する直前に採取した結膜から抽出した RNA を対照とした(n=3)・microarray 法により、miRNA の発現を比較した・

(2) その結果,結膜炎誘導により miRNA の 発現倍率が 1/2 倍未満となった遺伝子は 44 個,発現倍率が2倍以上となった miRNA が7 4個同定された.さらに,統合評価を行い,m iRNA 発現変化にともない,発現の変動が予 測される遺伝子を検索した.

(3) その結果, miRNA が増加し遺伝子発現を抑制したと考えられる遺伝子群(Lrp2 binding protein, mediator of RNA polymerase II transcription, subunit 12 homolog (yeast)-like, immunoglobulin superfamily, member 1, potassium voltage gated channel, Shaw-related subfamily, member 3, zinc finger, CCHC domain containing 16 など)と, miRNA が減少し遺伝子発現が増加したと考えられる遺伝子群(lactoperoxidase, aryl hydrocarbon receptor nuclear translocator 2, DnaJ (Hsp40) homolog, subfamily C, member 27, olfactory receptor 54, sarcoglycan zeta など)を同定した.

結膜炎の誘導により発現が減少した遺伝子と発現が上昇した miRNA関連遺伝子における共通44遺伝子リスト

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		結膜炎誘導		結膜炎誘導
	発現が減少した遺伝子名	後の減少倍	発現が増加した	後の増加倍
no	光切か減少しに退伝する	率変化(fold	miRNA	率変化(fold
		change)		change)
-	I 0 bis disconnectation	4.975	:D 404	
	Lrp2 binding protein		mmu-miR-494	2.396
2	Lrp2 binding protein	4.975	mmu-miR-499	6.239
	mediator of RNA polymerase II			
3	transcription, subunit 12	4.587	mmu-miR-302c	6.922
lٽ		4.007	111110 111111 0020	0.522
⊢	homolog (yeast)-like			
4	immunoglobulin superfamily,	4.196	mmu-miR-499	6.239
4	member 1	4.190	mmu=mirt=499	0.239
	potassium voltage gated channel,			
5	Shaw-related subfamily, member	4.175	mmu-miR-760-3p	4.256
3	onaw related subraining, member	4.173	IIIIIu-IIIIK-700-3p	4.230
ㄴ	3			
6	zinc finger, CCHC domain	4.145	mmu-miR-34c	6.107
0	containing 16	4.143	mmu-mik-34c	0.107
	ankyrin repeat and sterile alpha			
7		3.680	mmu-miR-494	2.396
⊢	motif domain containing 1B			
8	methionine adenosyltransferase	3.650	mmu-miR-760-3p	4.256
ľ	I, alpha	3.000	ППП ППК 700 3р	4.230
	solute carrier family 1 (glial high			
٨		2 621	mmu=miD=404	2 206
9	affinity glutamate transporter),	3.631	mmu-miR-494	2.396
oxdot	member 2			
10	solute carrier family 26, member	0.517	:D 404	0.000
10	3	3.517	mmu-miR-494	2.396
11	kinesin family member 5A	3.465	mmu-miR-466j	29.948
۳.		3.403	mmu mmx=400j	23.340
12	nuclear receptor subfamily 1,	3.379	mmu-miR-532-3p	4.280
Ľ	group I, member 2	0.070	оог ор	1.200
13	synaptonemal complex protein 2	3.349	mmu-miR-302c	6.922
	synaptonemal complex protein 2	3.349	mmu-miR-499	6.239
14		3.343	IIIIIu-IIIIK-455	0.239
	leucine-rich repeats and			
15	guanylate kinase domain	3.314	mmu-miR-466j	29.948
	containing			
16	sulfatase modifying factor 2	3.281	mmu-miR-532-3p	4.280
	cylicin, basic protein of sperm	0.201	mind mint doe op	1.200
17	cylicin, basic protein of sperm	3.250	mmu-miR-499	6.239
_	head cytoskeleton 2			
18	interleukin 20	3.180	mmu-miR-764-5p	34.150
19	maltase-glucoamylase	3.122	mmu-miR-302c	6.922
	coiled-coil domain containing 67	3.102	mmu-miR-532-3p	4.280
21	CD8 antigen, alpha chain	3.076	mmu-miR-712	4.253
	FYVE, RhoGEF and PH domain			
22	containing 4	3.075	mmu-miR-760-3p	4.256
⊢	poly(A) binding protein,			
23	poly(A) binding protein,	3.044	mmu-miR-302c	6.922
	cytoplasmic 5	0.011		0.022
24	Fc receptor-like 6	2.877	mmu-miR-760-3p	4.256
	and a selection described and a second			
25		2.803	mmu-miR-764-5p	34.150
-	chain base subunit 3			
	ankyrin repeat domain 55	2.744	mmu-miR-494	2.396
27	ankyrin repeat domain 55	2.744	mmu-miR-499	6.239
28	placenta-specific 1-like	2.613	mmu-miR-494	2.396
	placenta specific i like	2.010	TOT THIN WITH	2.000
29	regulating synaptic membrane	2.555	mmu-miR-760-3p	4.256
ㄴ	exocytosis 2			
30	deoxyribonuclease II beta	2.518	mmu-miR-764-5p	34.150
31	growth differentiation factor 2	2.510	mmu-miR-34c	6.107
	growth differentiation factor 2	2.510	mmu-miR-712	4.253
33	canopy 1 homolog (zebrafish)	2.479	mmu-miR-499	6.239
١.,	ubiquitin associated and SH3	0.107		6 107
34	domain containing, A	2.187	mmu-miR-34c	6.107
\vdash	Na+/K+ transporting ATPase			
35		2.159	mmu-miR-34c	6.107
\vdash	interacting 2			ļ
36	Na+/K+ transporting ATPase	2.159	mmu-miR-466j	29.948
30	interacting 2	2.108	11111u 11111X-400J	25.540
37	G protein-coupled receptor 17	2.150	mmu-miR-760-3p	4.256
ř		200	тоо ор	1,200
38	histone cluster 3, H2bb,	2.133	mmu-miR-760-3p	4.256
-	pseudogene			
39	histone cluster 1, H1d	2.112	mmu-miR-760-3p	4.256
	5,10-methylenetetrahydrofolate			
40	reductase	2.091	mmu-miR-764-5p	34.150
\vdash				
41	tumor necrosis factor receptor	2.077	mmu-miR-302c	6.922
Ľ	superfamily, member 8			0.522
42	neuronal PAS domain protein 3	2.061	mmu-miR-466j	29.948
	urocortin 2	2.014	mmu-miR-34c	6.107
-		2.017	040	0.107
44	mitogen-activated protein kinase	2.012	mmu-miR-466j	29.948
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結膜炎の誘導により発現が増加した遺伝子と発現が減少した miRNA関連遺伝子における共通74遺伝子リスト

no	発現が上昇した遺伝子名	結膜炎誘導 後の増加倍 率変化(fold change)	発現が減少した miRNA	結膜炎誘導 後の減少倍 率変化(fold change)
1	lactoperoxidase	6.567	mmu-miR-680	66.411
	aryl hydrocarbon receptor nuclear translocator 2	4.209	mmu-miR-294	6.955
	DnaJ (Hsp40) homolog, subfamily C, member 27	4.145	mmu-miR-342-5p	4.116
4	olfactory receptor 54	3.819	mmu-miR-294	6.955
5	sarcoglycan zeta	3.599	mmu-miR-1188	4.569
6	G protein-coupled receptor 123	3.445	mmu-miR-874	455.641
	5-hydroxytryptamine (serotonin) receptor 2A	3.381	mmu-miR-680	66.411

9	collagen, type VI, alpha 6	3.345	mmu-miR-138	4.591
_	regulatory factor X, 4 (influences	3.335	mmu-miR-1188	4.569
10	HLA class II expression) collagen-like tail subunit (single strand of homotrimer) of	3.228	mmu-miR-702	3.677
	asymmetric acetylcholinesterase ubiquitin interaction motif			
11	containing 1	3.166	mmu-miR-138	4.591
12	blood cell	3.104	mmu-miR-342-5p	4.116
	gap junction protein, beta 5	3.070	mmu-miR-702	3.677
	fibrinogen alpha chain	3.069	mmu-miR-680	66.411
	bassoon	3.040	mmu-miR-680	66.411
16	sideroflexin 5	3.039	mmu-miR-342-5p	4.116
17	sideroflexin 5	3.039	mmu-miR-680	66.411
	sideroflexin 5	3.039	mmu-miR-874	455.641
19	membrane-spanning 4-domains,	3.038	mmu-miR-146a	2.204
20	subfamily A, member 1 membrane-spanning 4-domains,	3.038	mmu-miR-680	66.411
21	subfamily A, member 1 polyhomeotic-like 3 (Drosophila)	3.019	mmu-miR-146a	2.204
	polyhomeotic-like 3 (Drosophila)	3.019	mmu-miR-294	6.955
23	TRAF-interacting protein with forkhead-associated domain,	2.953	mmu-miR-342-5p	4.116
24	family member B cell death-inducing DFFA-like effector c	2.928	mmu-miR-342-5p	4.116
25	hypoxia inducible factor 3, alpha subunit	2.868	mmu-miR-138	4.591
26	collagen, type V, alpha 3	2.823	mmu-miR-874	455.641
	sclerostin	2.801	mmu-miR-702	3.677
28	chordin-like 1 family with sequence similarity	2.790	mmu-miR-680	66.411
	155, member A	2.779	mmu-miR-1188	4.569
30	leucine rich repeat containing 3	2.779	mmu-miR-294	6.955
31	leucine-rich repeat, immunoglobulin-like and transmembrane domains 1	2.765	mmu-miR-294	6.955
32	metallothionein-like 5, testis- specific (tesmin)	2.763	mmu-miR-702	3.677
33	transmembrane and coiled-coil domains 3	2.745	mmu-miR-125a- 3p	4.035
34	sodium channel, voltage-gated, type II, alpha 1	2.745	mmu-miR-680	66.411
35	NFKB activating protein-like	2.706	mmu-miR-294	6.955
36	DnaJ (Hsp40) homolog,	2.669	mmu-miR-294	6.955
37	subfamily C, member 5 beta nuclear receptor coactivator 7	2.654	mmu-miR-1188	4.569
	nuclear receptor coactivator 7	2.654	mmu-miR-294	6.955
39	nuclear receptor coactivator 7	2.654	mmu-miR-680	66.411
40	growth differentiation factor 5	2.639	mmu-miR-146a	2.204
41	mutS homolog 5 (E. coli)	2.633	mmu-miR-146a	2.204
42	protocadherin beta 3	2.601	mmu-miR-125a- 3p	4.035
			mmu-miR-342-5p	4.116
43	forkhead box N4	2.584		
43	forkhead box N4 T-box 4	2.584 2.576	mmu-miR-342-5p	4.116
13 14				
13 14 15	T-box 4	2.576	mmu-miR-342-5p mmu-miR-702 mmu-miR-125a-	4.116
13 14 15	T-box 4 tripartite motif-containing 14 forkhead box E3	2.576 2.573 2.572	mmu-miR-342-5p mmu-miR-702 mmu-miR-125a- 3p	4.116 3.677
13 14 15 16	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8	2.576 2.573 2.572 2.562	mmu-miR-342-5p mmu-miR-702 mmu-miR-125a- 3p mmu-miR-874 mmu-miR-125a-	4.116 3.677 4.035 455.641
43 44 45 46 47	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B	2.576 2.573 2.572 2.562 2.521	mmu-miR-342-5p mmu-miR-702 mmu-miR-125a- 3p mmu-miR-874 mmu-miR-125a- 3p	4.116 3.677 4.035 455.641 4.035
43 44 45 46 47 48	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1	2.576 2.573 2.572 2.562 2.521 2.463	mmu-miR-342-5p mmu-miR-702 mmu-miR-125a- 3p mmu-miR-874 mmu-miR-125a- 3p	4.116 3.677 4.035 455.641 4.035 4.591
43 44 45 46 47 48	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like	2.576 2.573 2.572 2.562 2.521 2.463 2.418	mmu-miR-342-5p mmu-miR-702 mmu-miR-125a- 3p mmu-miR-874 mmu-miR-125a- 3p	4.116 3.677 4.035 455.641 4.035 4.591 4.116
43 44 45 46 47 48 49 50	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407	mmu-miR-342-5p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-138 mmu-miR-342-5p mmu-miR-125a- 3p	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035
43 44 45 46 47 48 49 50	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex	2.576 2.573 2.572 2.562 2.521 2.463 2.418	mmu-miR-342-5p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-138 mmu-miR-342-5p mmu-miR-125a-	4.116 3.677 4.035 455.641 4.035 4.591 4.116
43 44 45 46 47 48 49 50 51	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 gilal fibrillary acidic protein gilal fibrillary acidic protein a disintegrin-like and	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407	mmu-miR-342-5p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-138 mmu-miR-342-5p mmu-miR-125a- 3p	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035
43 44 45 46 47 48 49 50 51	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 glial fibrillary acidic protein glial fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365	mmu-miR-342-5p mmu-miR-702 mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-138 mmu-miR-342-5p mmu-miR-125a- 3p mmu-miR-138	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591
43 44 45 46 47 48 49 50 51 52 53	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 glial fibrillary acidic protein glial fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365	mmu-miR-342-5p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-138 mmu-miR-342-5p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-125a- 3p	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677
43 44 45 46 47 48 49 50 51 52 53	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 glial fibrillary acidic protein glial fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 ribonuclease, RNase A family, 13 (non-active)	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309	mmu-miR-342-5p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-125a- 3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569
43 44 45 46 47 48 49 50 51 52 53 54	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-lika adaptor-related protein complex AP-4, epsilon 1 gilal fibrillary acidic protein gilal fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 ribonuclease, RNase A family, 13 (non-active) solute carrier family 38, member 1	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309 2.301	mmu-miR-342-5p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-702 mmu-miR-1188 mmu-miR-702	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569 6.955 4.569
43 44 45 46 47 48 49 50 51 52 53 54 55 56	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 gilal fibrillary acidic protein gilal fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 ribonuclease, RNase A family, 13 (non-active) solute carrier family 38, member 1 MKL/myocardin-like 2	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309 2.301	mmu-miR-342-5p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-1188 mmu-miR-1188	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569 6.955 4.569 3.677 66.411
43 44 45 46 47 48 49 50 51 52 53 54 55 56 57	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 glial fibrillary acidic protein glial fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 ribonuclease, RNase A family, 13 (non-active) solute carrier family 38, member 1 MKL/myocardin-like 2 sorting nexin 31 potassium channel, subfamily K,	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309 2.301	mmu-miR-342-5p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-702 mmu-miR-1188 mmu-miR-702	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569 6.955 4.569
43 44 44 45 46 47 48 49 49 50 51 55 55 55 55 55 66 60	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 glial fibrillary acidic protein glial fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 moun-active) solute carrier family 38, member 1 MKL/myocardin-like 2 sorting nexin 31 potassium channel, subfamily K, member 10 testis-specific protein, Y-	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309 2.301 2.297 2.259	mmu-miR-342-5p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-102 mmu-miR-1188 mmu-miR-1188 mmu-miR-702 mmu-miR-702 mmu-miR-702	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569 6.955 4.569 3.677 66.411 3.677
43 44 45 46 47 48 49 50 51 52 53 54 55 56	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 glial fibrillary acidic protein glial fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 motif, 6 ubiquitin carboxy-terminal	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309 2.301 2.297 2.259 2.234	mmu-miR-342-5p mmu-miR-1702 mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-1188 mmu-miR-1188 mmu-miR-1188 mmu-miR-1188 mmu-miR-1188 mmu-miR-1680 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-702	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569 6.955 4.569 3.677 66.411 3.677 2.204
43 44 44 45 46 47 48 49 49 50 55 55 55 55 55 55 56 66 60	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 glial fibrillary acidic protein glial fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 ribonuclease, RNase A family, 13 (non-active) solute carrier family 38, member 1 MKL/myocardin-like 2 sorting nexin 31 potassium channel, subfamily K, member 10 testis-specific protein, Y- encoded-like 5 testis-specific protein, Y- encoded-like 5 connector enhancer of kinase	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309 2.301 2.297 2.259 2.234 2.221	mmu-miR-342-5p mmu-miR-1702 mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-1188 mmu-miR-702 mmu-miR-702 mmu-miR-1488 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-1468	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569 3.677 66.411 3.677 2.204
43 444 445 446 447 448 449 449 500 500 555 555 555 555 556 660 660 661	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 gilal fibrillary acidic protein gilal fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 ribonuclease, RNase A family, 13 (non-active) solute carrier family 38, member 1 MKL/myocardin-like 2 sorting nexin 31 potassium channel, subfamily K, member 10 testis-specific protein, Y- encoded-like 5 connector enhancer of kinase suppressor of Ras 2 schlafen 5	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309 2.301 2.297 2.259 2.234 2.221	mmu-miR-342-5p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-125a-3p mmu-miR-138 mmu-miR-126a-3p mmu-miR-138 mmu-miR-128 mmu-miR-102 mmu-miR-1188 mmu-miR-702 mmu-miR-680 mmu-miR-702 mmu-miR-146a mmu-miR-702 mmu-miR-146a mmu-miR-702	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569 6.955 4.569 3.677 66.411 3.677 2.204 3.677
43 444 445 446 447 448 449 449 500 500 555 555 555 555 556 660 660 661	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-like adaptor-related protein complex AP-4, epsilon 1 gilal fibrillary acidic protein gilal fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 ribonuclease, RNase A family, 13 (non-active) solute carrier family 38, member 1 MKL/myocardin-like 2 sorting nexin 31 potassium channel, subfamily K, member 10 testis-specific protein, Y- encoded-like 5 connector enhancer of kinase suppressor of Ras 2 sohlafen 5 NOL1/NOP2/Sun domain family,	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309 2.301 2.297 2.259 2.234 2.221 2.169	mmu-miR-342-5p mmu-miR-1702 mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-125a-3p mmu-miR-138 mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-102 mmu-miR-1188 mmu-miR-702 mmu-miR-680 mmu-miR-702 mmu-miR-146a mmu-miR-702	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569 6.955 4.569 3.677 66.411 3.677 2.204 3.677
43 44 44 45 46 47 48 49 49 55 55 55 55 55 55 55 55 56 66 66 61 61 66 66 66 66 66 66 66 66 66	T-box 4 tripartite motif-containing 14 forkhead box E3 surfactant associated protein B solute carrier family 8 (sodium/calcium exchanger), member 1 solute carrier family 14 (urea transporter), member 1 CCCTC-binding factor (zinc finger protein)-lika adaptor-related protein complex AP-4, epsilon 1 gilal fibrillary acidic protein a disintegrin-like and metallopeptidase (reprolysin type) with thrombospondin type 1 motif, 6 ubiquitin carboxy-terminal hydrolase L1 ribonuclease, RNase A family, 13 (non-active) solute carrier family 38, member 1 MKL/myocardin-like 2 sorting nexin 31 potassium channel, subfamily K, member 10 testis-specific protein, Y- encoded-like 5 testis-specific protein, Y- encoded-like 5 testis-specific protein, Y- encoded-like 5 sonnector enhancer of kinase suppressor of Ras 2 schlafen 5 NOL1/NOP2/Sun domain family, member 7	2.576 2.573 2.572 2.562 2.521 2.463 2.418 2.407 2.365 2.365 2.322 2.314 2.309 2.301 2.297 2.259 2.234 2.221 2.169 2.151	mmu-miR-342-5p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-125a-3p mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-139 mmu-miR-139 mmu-miR-138 mmu-miR-138 mmu-miR-138 mmu-miR-102 mmu-miR-1188 mmu-miR-702 mmu-miR-1680 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-702 mmu-miR-704 mmu-miR-704 mmu-miR-705 mmu-miR-7060	4.116 3.677 4.035 455.641 4.035 4.591 4.116 4.035 4.591 3.677 4.569 6.955 4.569 3.677 66.411 3.677 2.204 3.677 455.641

	potassium channel tetramerisation domain containing 7	2.088	mmu-miR-125a- 3p	4.035
69	glypican 5	2.081	mmu-miR-680	66.411
70	fatty acid desaturase domain family, member 6	2.060	mmu-miR-125a- 3p	4.035
71	chemokine (C-X3-C) receptor 1	2.037	mmu-miR-874	455.641
72	solute carrier family 35, member F1	2.034	mmu-miR-1188	4.569
73	solute carrier family 35, member F1	2.034	mmu-miR-138	4.591
74	glutamyl aminopeptidase	2.010	mmu-miR-680	66.411

その後、発現レベルを定量的 RT-PCR 法で評価した。定量的 RT-PCR 法ではマイクロアレイで確認された各 mi RNA 発現の変動(アレルギー性結膜炎の誘導による)を再現できなかった。

5. 主な発表論文等

(研究代表者、研究分担者及び連携研究者に は下線)

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6. 研究組織

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