

# Targeted Delivery of Mucosal Vaccines

Development of Ligands to Receptors  
on Peyer's Patch Follicle Epithelium

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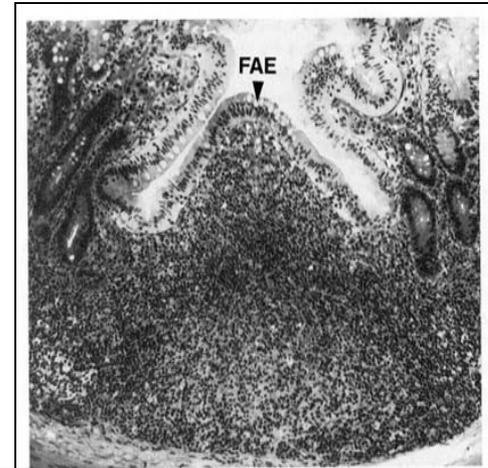
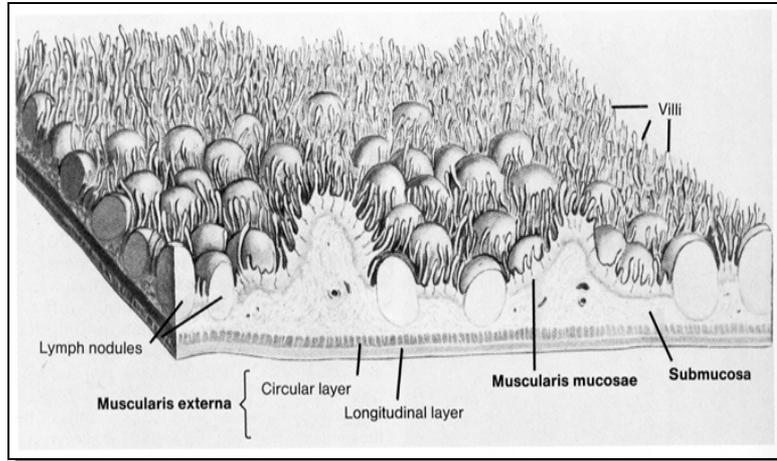
# Ligands to Receptors on Peyer's Patch Follicle Epithelium

- Peyer's Patches are critical in protective mucosal immunity BUT ALSO in mucosal tolerance induction
- PP Follicle Associated Epithelium is central to this surveillance function; can we exploit the system for vaccine delivery?
- Q1: Can the FAE and M cells be molecularly defined?
- Q2: Does the FAE have functions helpful to mucosal immunity?
- Q3: How do we exploit the biology for vaccine delivery?

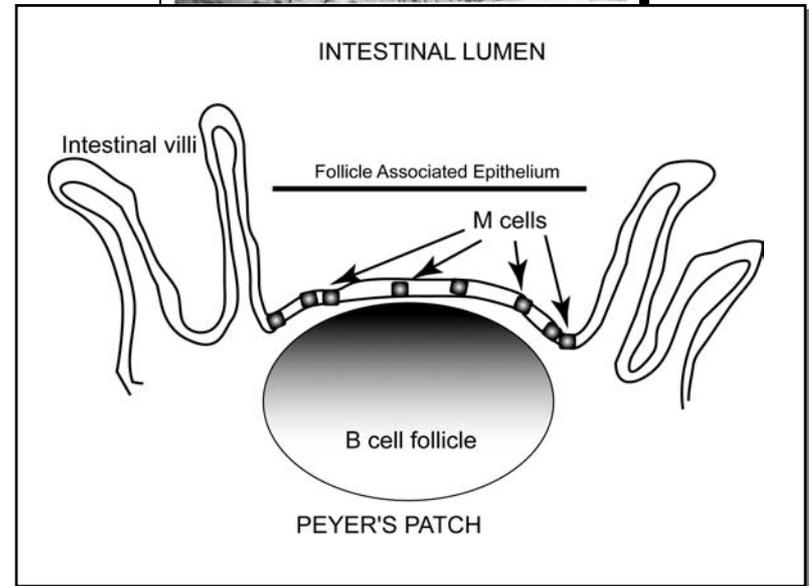
# FAE/M Cell Biology

- In the small intestine, immune responses in Peyer's Patches may be activated against pathogens (viruses, bacteria), or tolerance may be induced to food antigens
- Sampling of antigens is through active transport of proteins across Peyer's Patch epithelium
- Hypothesis: FAE/M cell specific genes exist that can explain their function
  - M cell specific receptors can be exploited for development of vaccine/drug delivery systems

# FAE/M Cell Biology

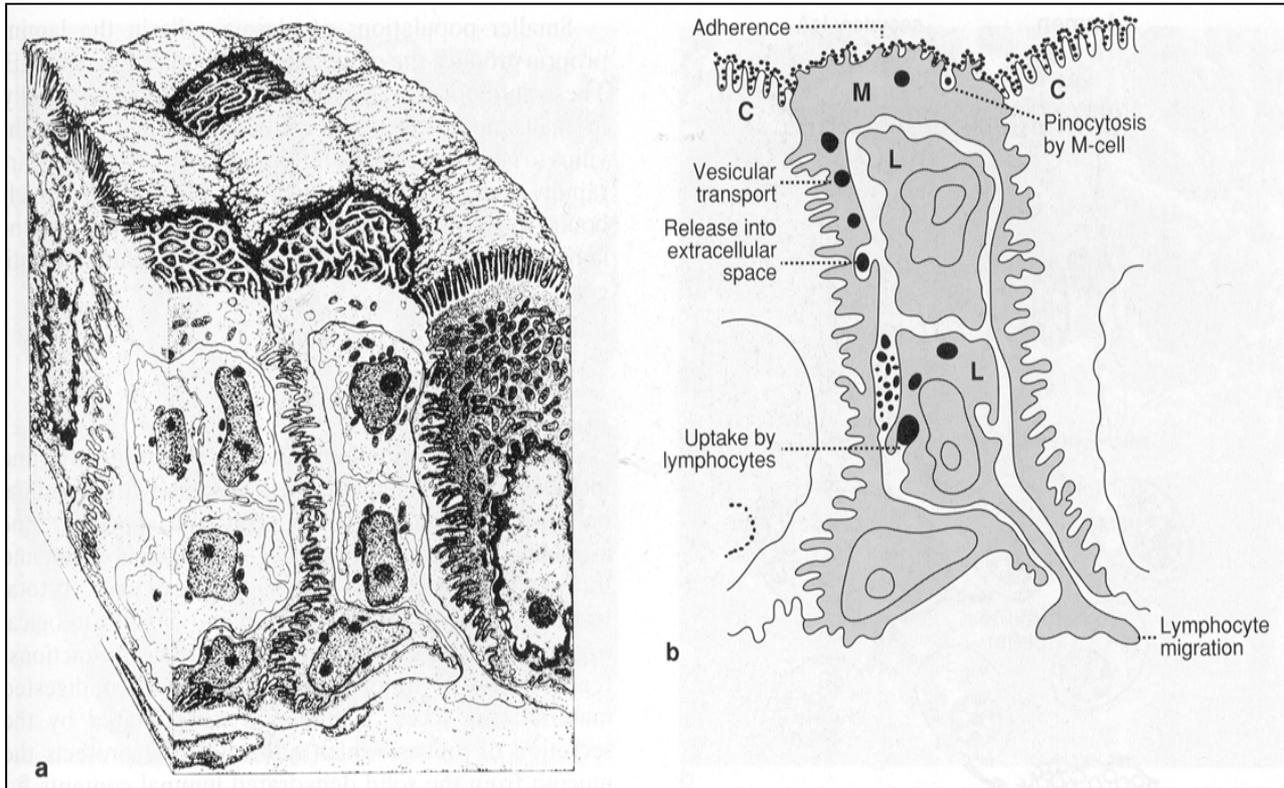


- Peyer's Patches in intestine
  - Dependent on lymphocytes and Lymphotoxin-beta
  - Organized B follicles
  - Interfollicular T/DC zone



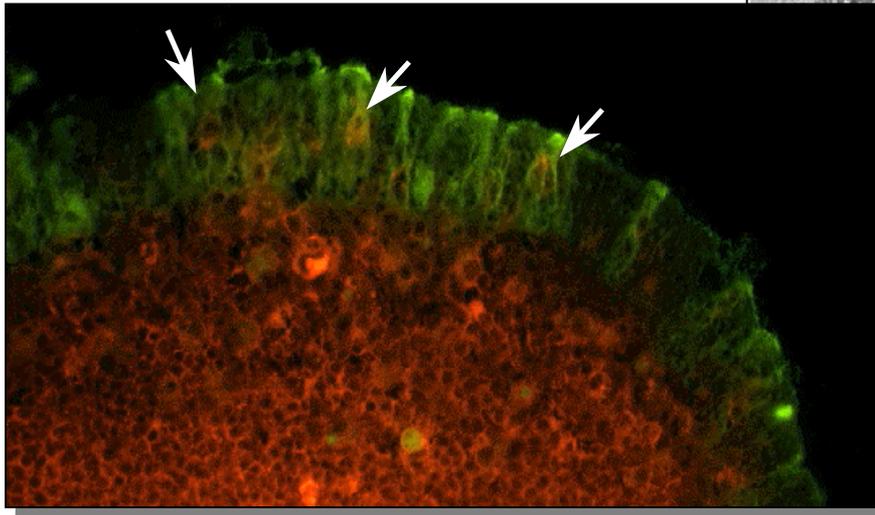
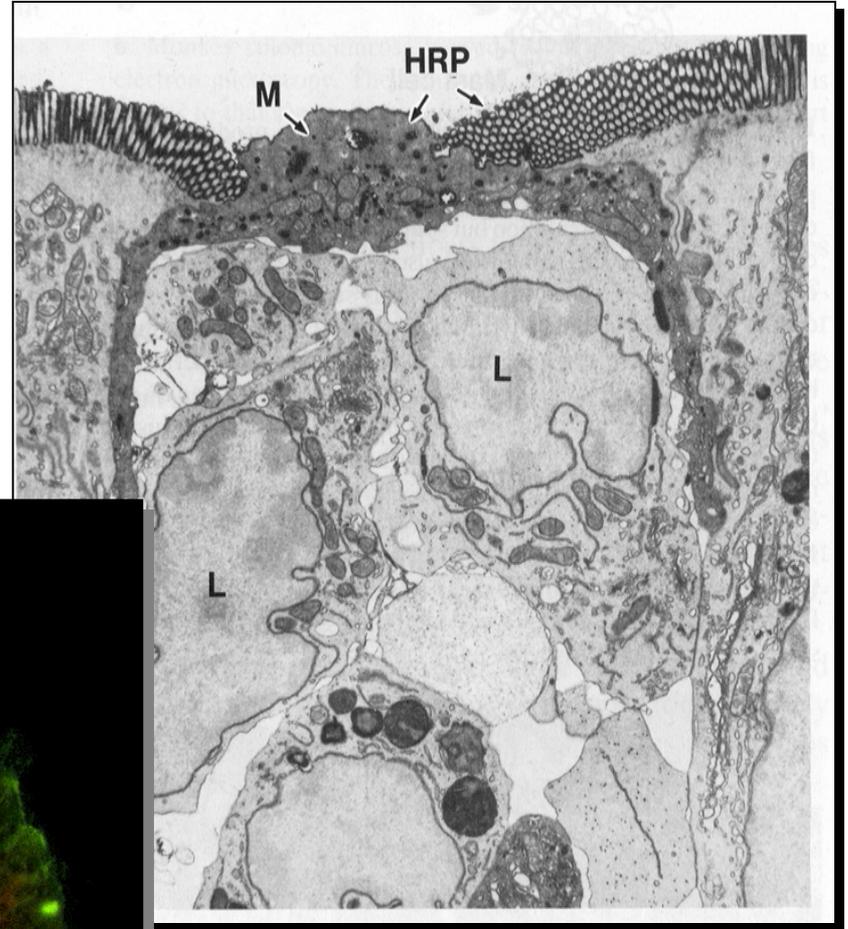
# FAE/M Cell Biology

- Clues to FAE specialization:
  - Induction of differentiated phenotype dependent on interaction with lymphoid cells
  - Distinct morphology of M cells



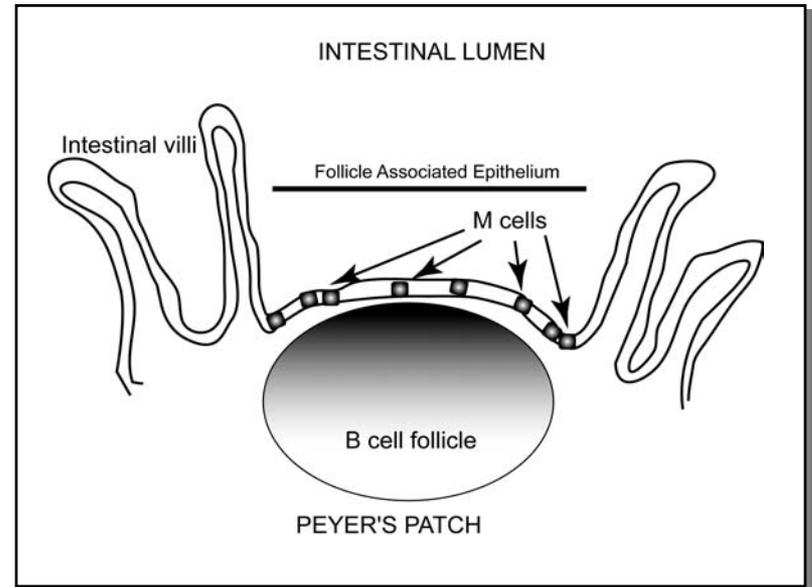
# FAE/M Cell Biology

- M cells develop in contact with B lymphocytes (arrows)
- Lectin UEA-1 (green) identifies mouse M cells



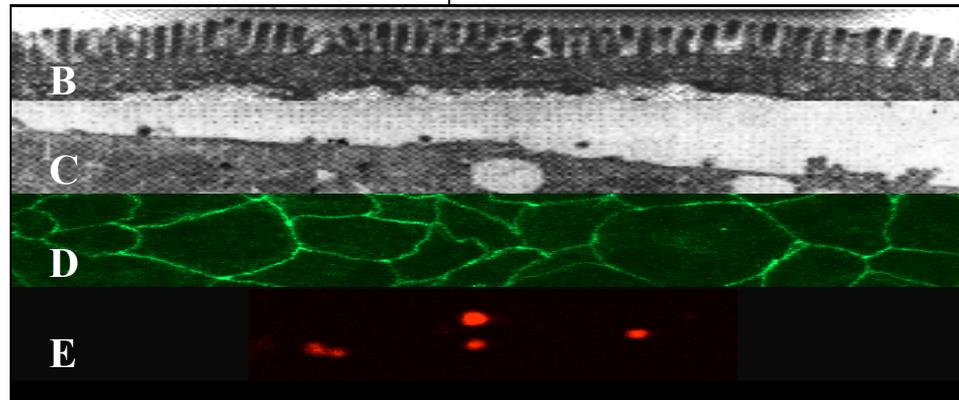
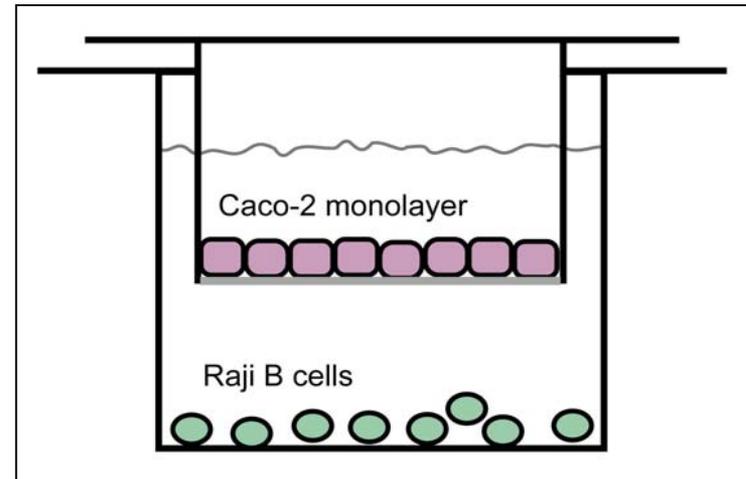
# FAE/M Cell Biology

- Peyer's Patches in intestinal mucosa:
- TOGA® Gene Expression Profiling Studies to Find Candidate Receptors:
  - Human cell culture, Caco-2 co-culture with Raji B cells
    - Induces M cell phenotype
  - Peyer's Patch tissue from mouse and macaque, microdissection of FAE

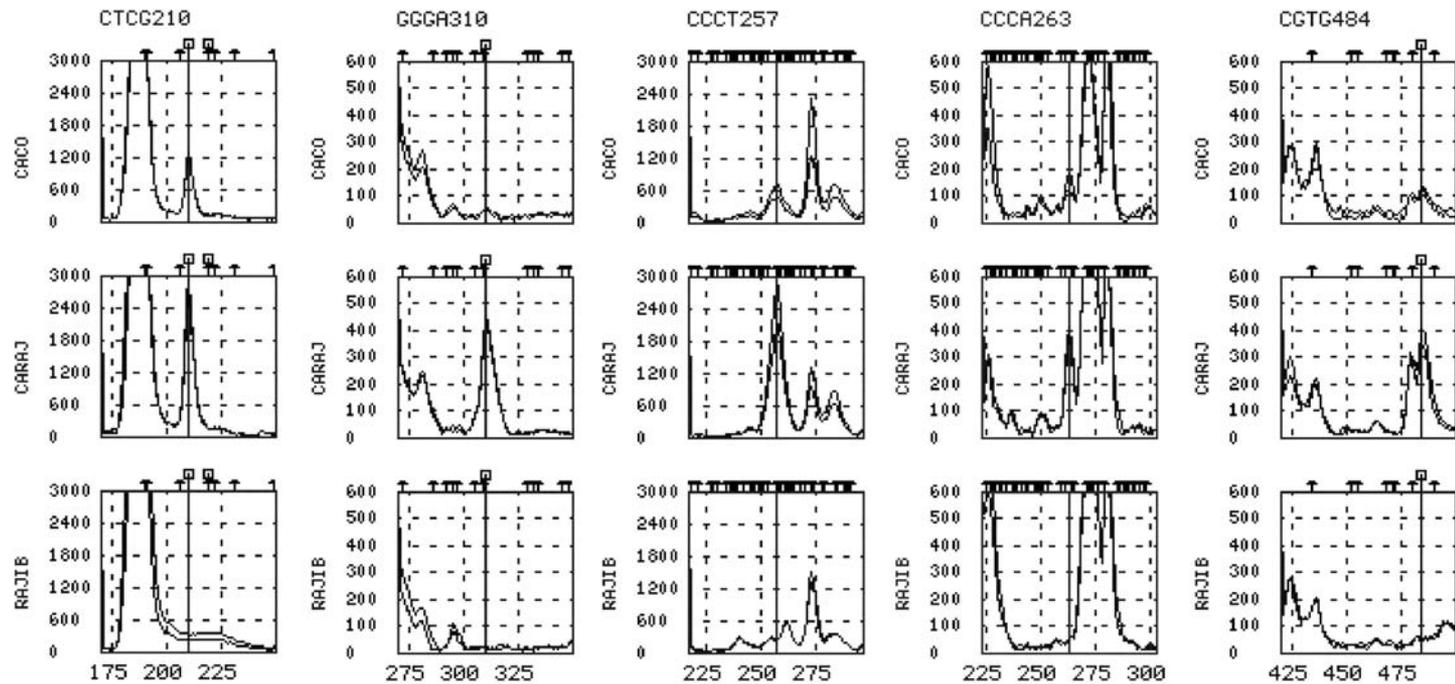


# Caco-2/Raji B Co-culture

- Soluble factors provided by Raji B cells
- Loss of brush border
- Microparticle transcytosis
- A true M cell phenotype?



# Candidate Selection

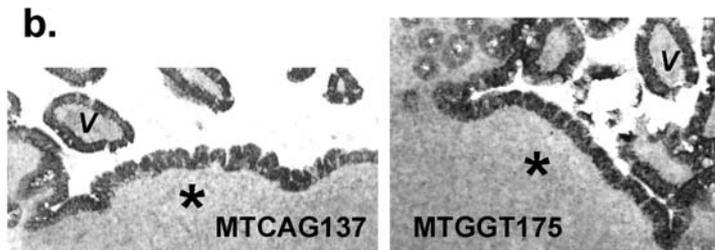


# Genes Regulated in Caco-2 Co-culture

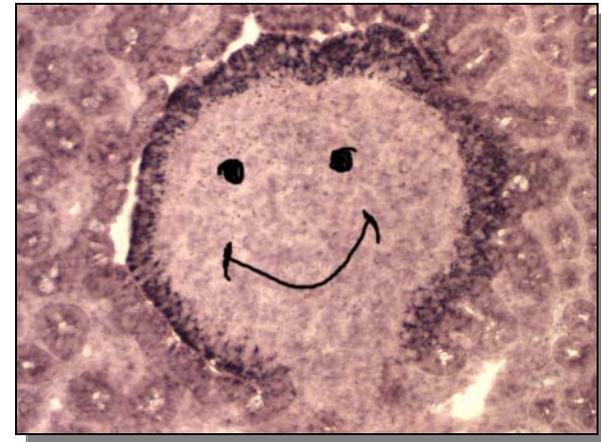
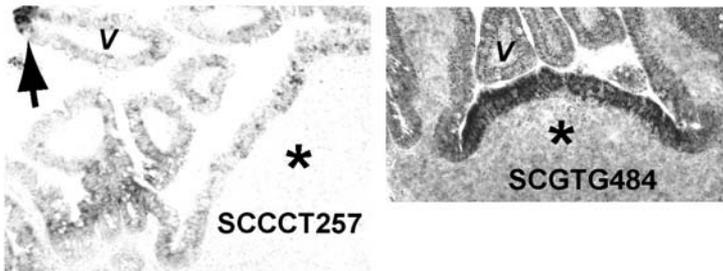
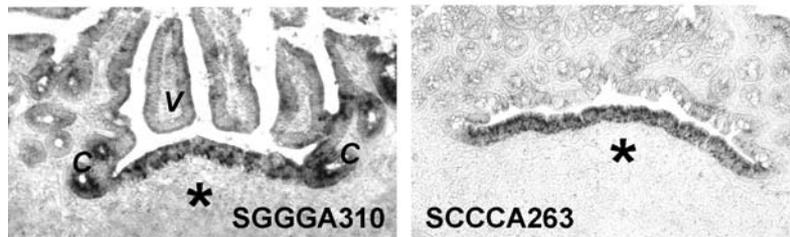
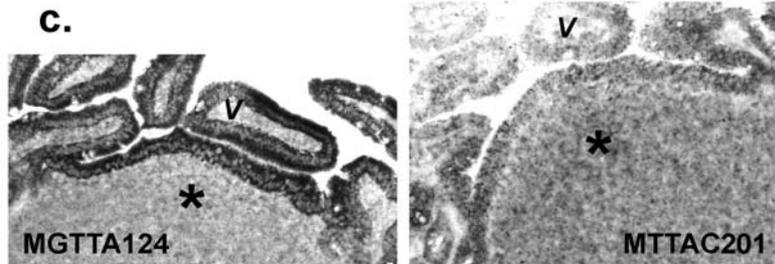
<u>Gene (accession no.)</u>	<u>Caco</u>	<u>CaRaj</u>	<u>RajiB</u>	<u>Fold regulation by TOGA, qPCR</u>
<u>Up-regulated:</u>				
Transcription factor (AK000232)	44	2165	51	49.2, 119.3
Jagged-1 (AW369026)	43	328	46	7.6, 1.8
c-Maf (AV648578)	134	1273	49	9.5, 14.7
DEC-1 (AB004066)	169	972	78	5.8, 3.7
RAB-13 (W46375)	73	345	23	4.7, 9.3
Glutaredoxin (AW128930)	135	890	270	6.6, 3.6
GPx-4 (X71973)	92	277	55	3.0, 4.4
ULK1 (AF045458)	119	333	61	2.8, 2.5
CDC2-related kinase (Q14004)	56	149	53	2.7, n.t.
<u>Down-regulated:</u>				
Ubiquitin B (BC000379)	1492	665	979	0.45*
Mitochondrial gene (E27671)	3360	138	3918	0.04*
3-pgdh (AF006043)	1091	268	997	0.25*
farnesyl diphos synthase (BC010004)	2066	831	2168	0.40*
transketolase (BC008615)	1123	416	1155	0.37*

# Candidate Receptor Genes

<u>Gene</u>	<u>Fold-regulation by TOGA, qPCR</u>	<u>Tissue Expression</u>
Biliary glycoprotein A	3.9, 1.8	FAE=villi
Mu protocadherin	10.6, 19.5	FAE=villi
Tetraspan TM4SF5	2.6, 1.0	FAE>=villi
LDL-R	5.0, 1.1	FAE>villi
Apolipoprotein B	2.3, 3.5	FAE>villi
Tetraspan TM4SF3	8.9, 4.8	FAE>>villi, crypts
C. perfringens enterotoxin R	4.2, 1.8	FAE, M cells, villi
MMP15	2.7, 0.7	FAE
Laminin beta 3	3.0, 1.9	FAE

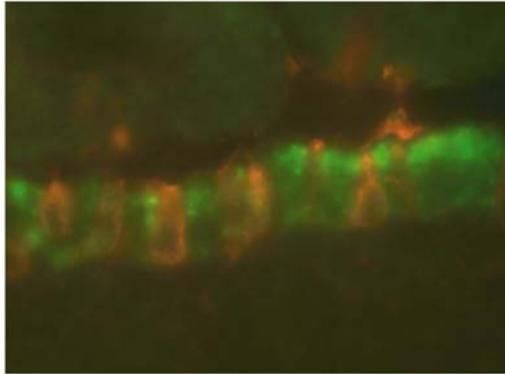


# The Proof Is in the Patches



- Many epithelial specific patterns, not all restricted to FAE
- Some restricted to FAE, or subset of FAE

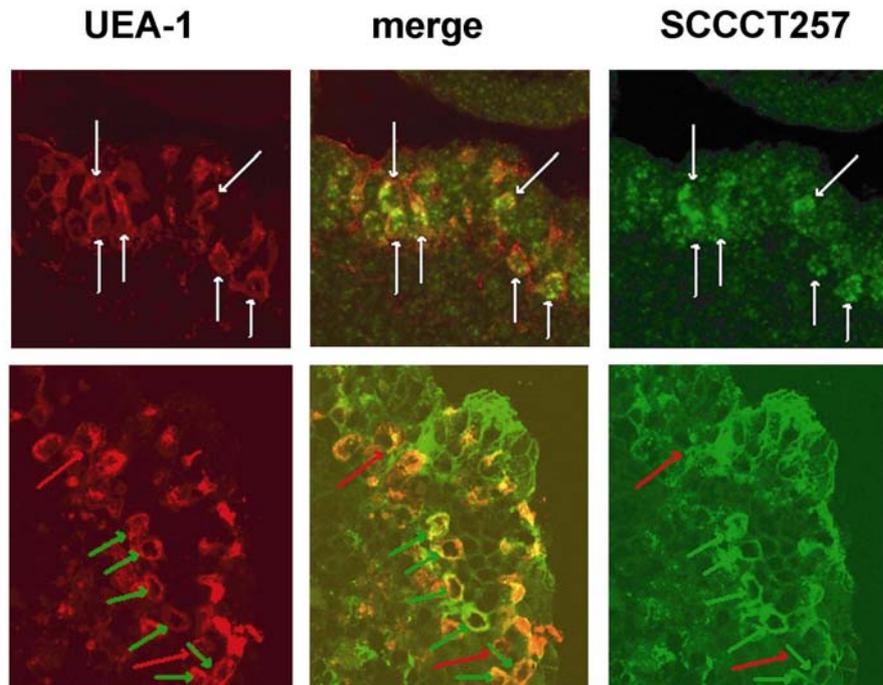
### a. SCGTG484



# Co-localization With M Cell Marker UEA-1

- SCGTG484 (laminin beta 3) showed FAE specific distribution, but not on M cells

### b. SCCCT257

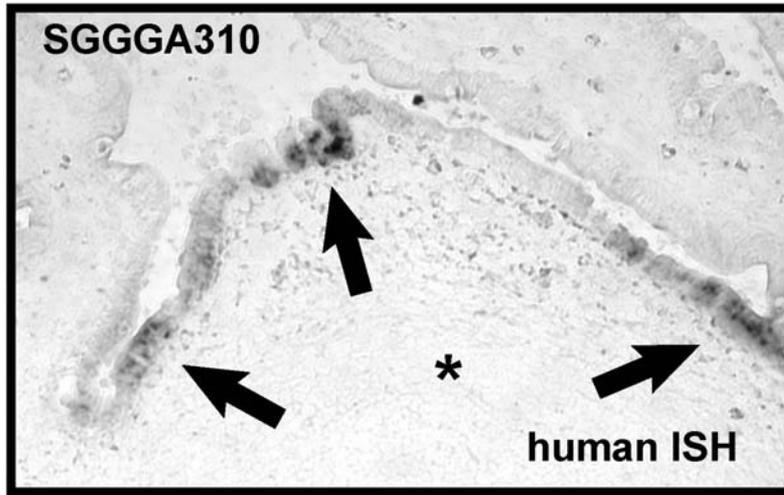


ISH

IHC

- SCCCT257 (CPE-R) showed epithelial tight junction distribution, but also higher expression and cytoplasmic distribution in M cells

a.

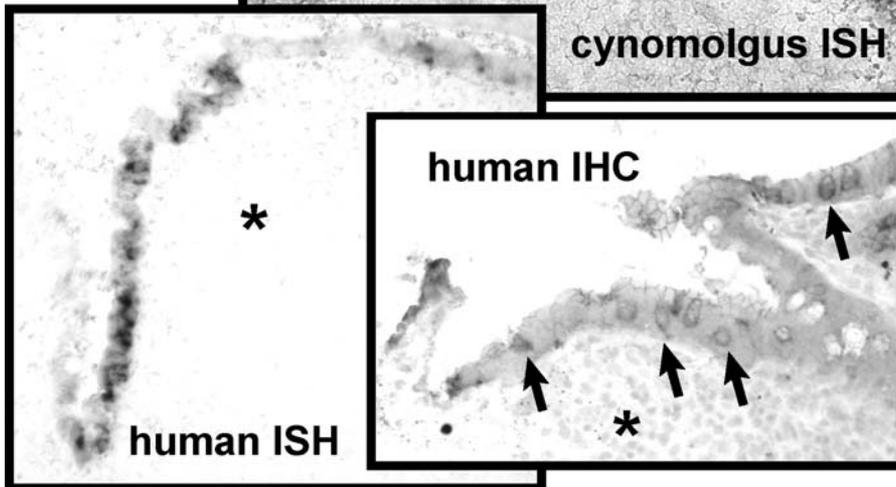
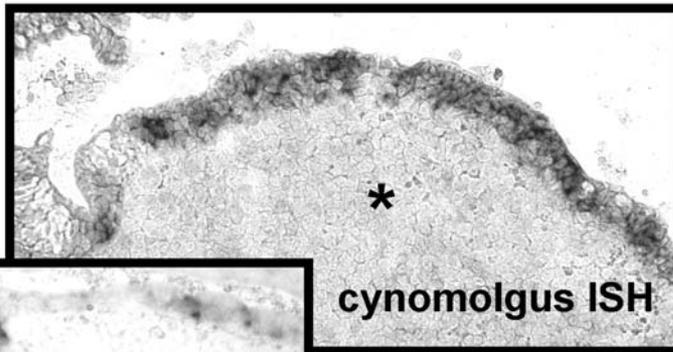


# Expression in Monkey and Human Peyer's Patch

- TM4SF3 expressed in FAE subset
- CPE-R again shows cytoplasmic distribution in subset of FAE

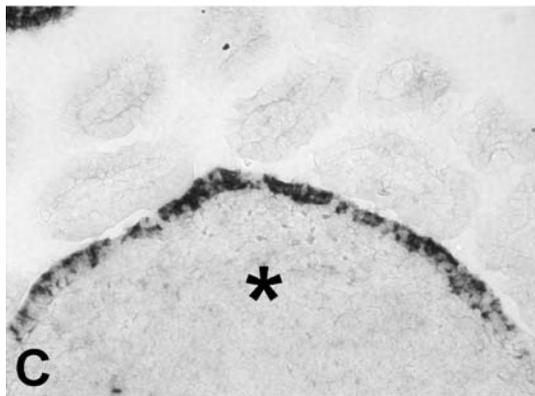
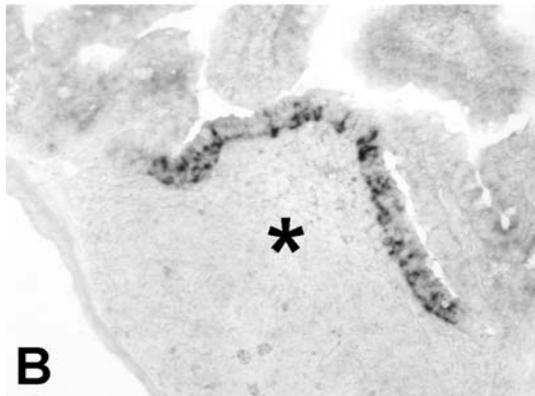
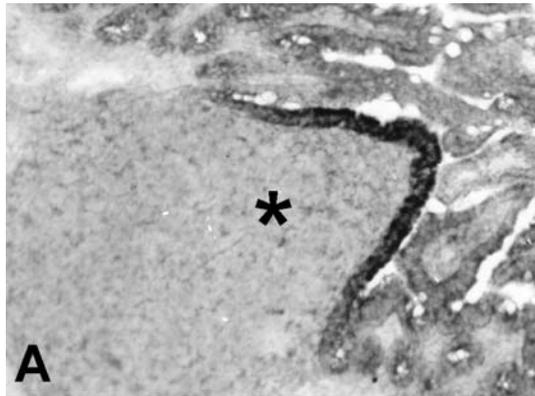
b.

SCCCT257



# Gene Stories I: Cell Culture Modeling of Complex Tissue

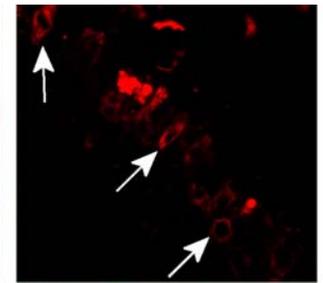
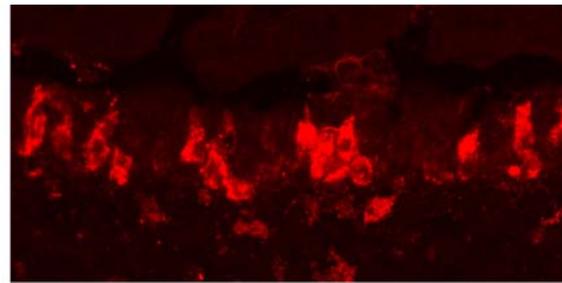
- FAE specific genes identified, consistent with specialized epithelial development and function
  - Transcription factors
  - Laminin beta 3, MMP15, tetraspanins, protocadherin, RAB-13
- Genes identified showing FAE subset expression
  - CPE-R suggests M cell specific pattern
- Conservation of expression in mouse, human



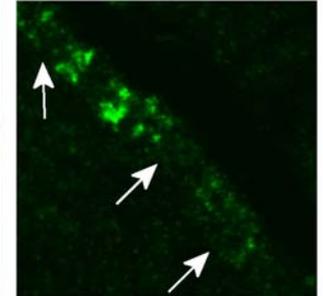
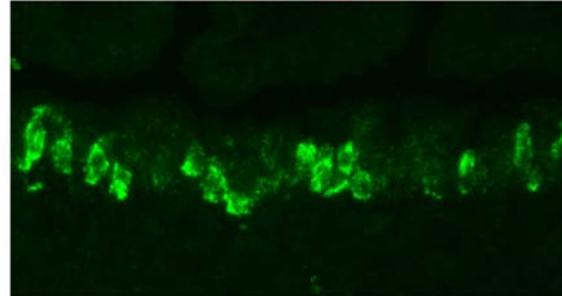
More genes from study of dissected FAE:

- PGRP genes show expression in distinct FAE subsets
- PGRP-S is M cell specific, PGRP-L is not

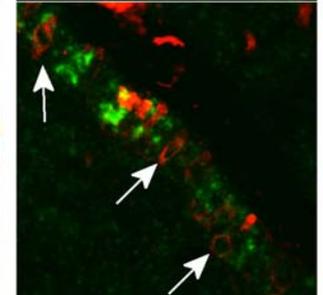
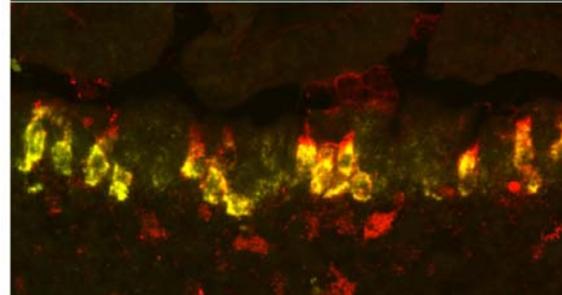
**UEA-1**



**ISH**



**merge**



# Nasal Associated Lymphoid Tissue (NALT)



- Different mucosal immune system sites express similar sets of epithelial receptors

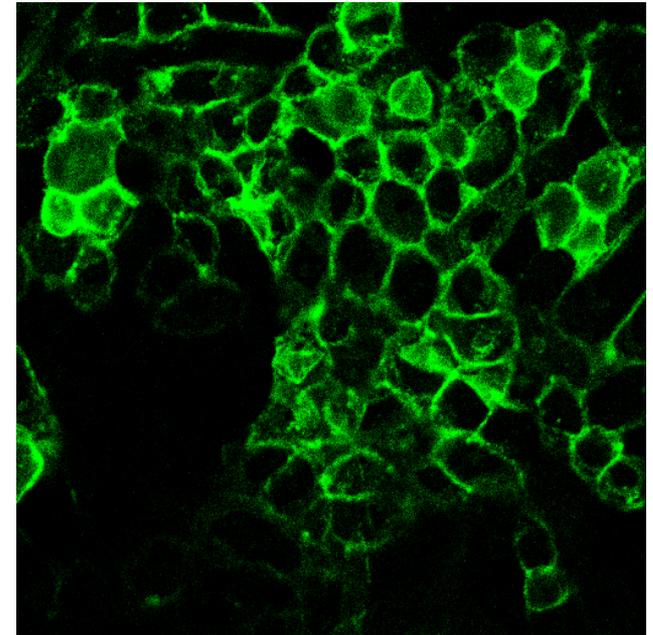
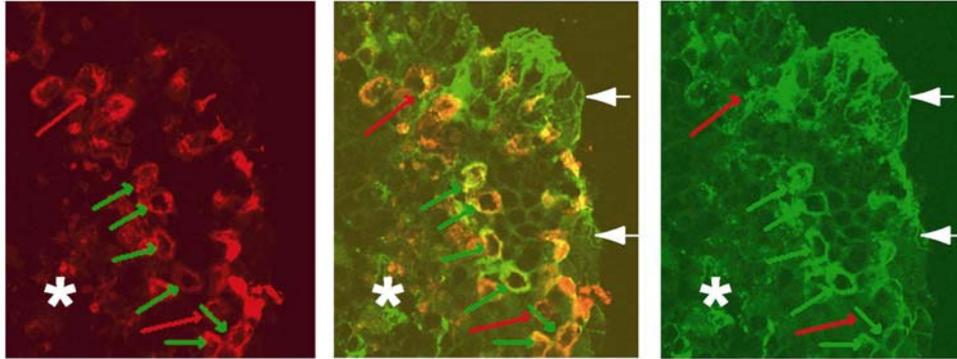
# Gene Stories II: Antigen and Adjuvant Receptors

- PGRP gene distribution in FAE suggests epithelial cell functional specialization
- Dual functions of PGRPs in FAE and M cells
  - Peptidoglycan receptors are “Pattern Recognition Receptors” (PRR) which trigger innate immunity
  - PRR triggering is the basis of vaccine adjuvants
  - Thus, receptors may therefore target both antigen delivery and adjuvant signaling

# Cell Biology of FAE/M Cell Genes

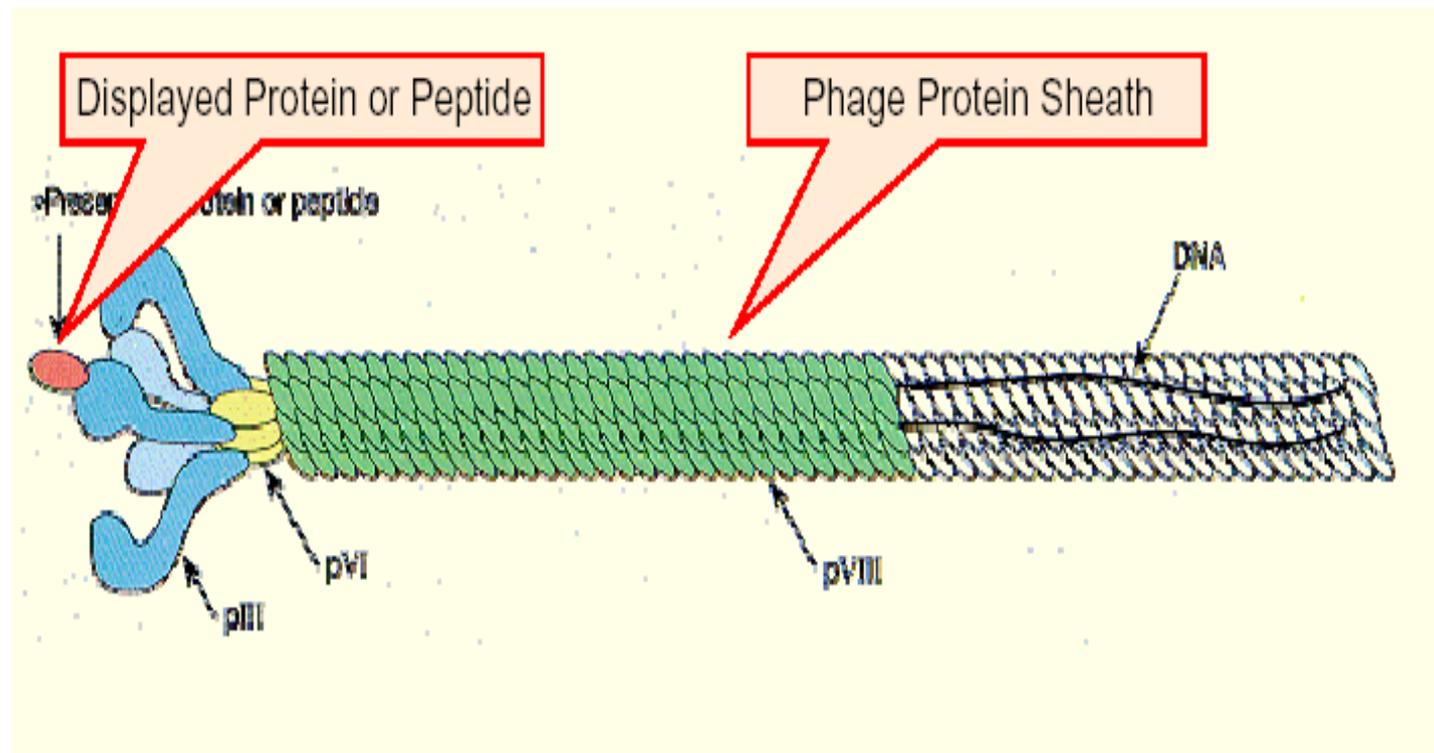
- Are some of the candidate genes receptors for FAE/M cell transcytosis?
  - Cloned full length cDNA, stable transfectants
  - Select synthetic ligands (phage display)
  - Test for binding and transcytosis in vivo
- Are some of the candidate genes receptors for adjuvant signals?
  - Test ligands for adjuvant activity in the presence of an antigen challenge

# CPE-R Subcellular Distribution

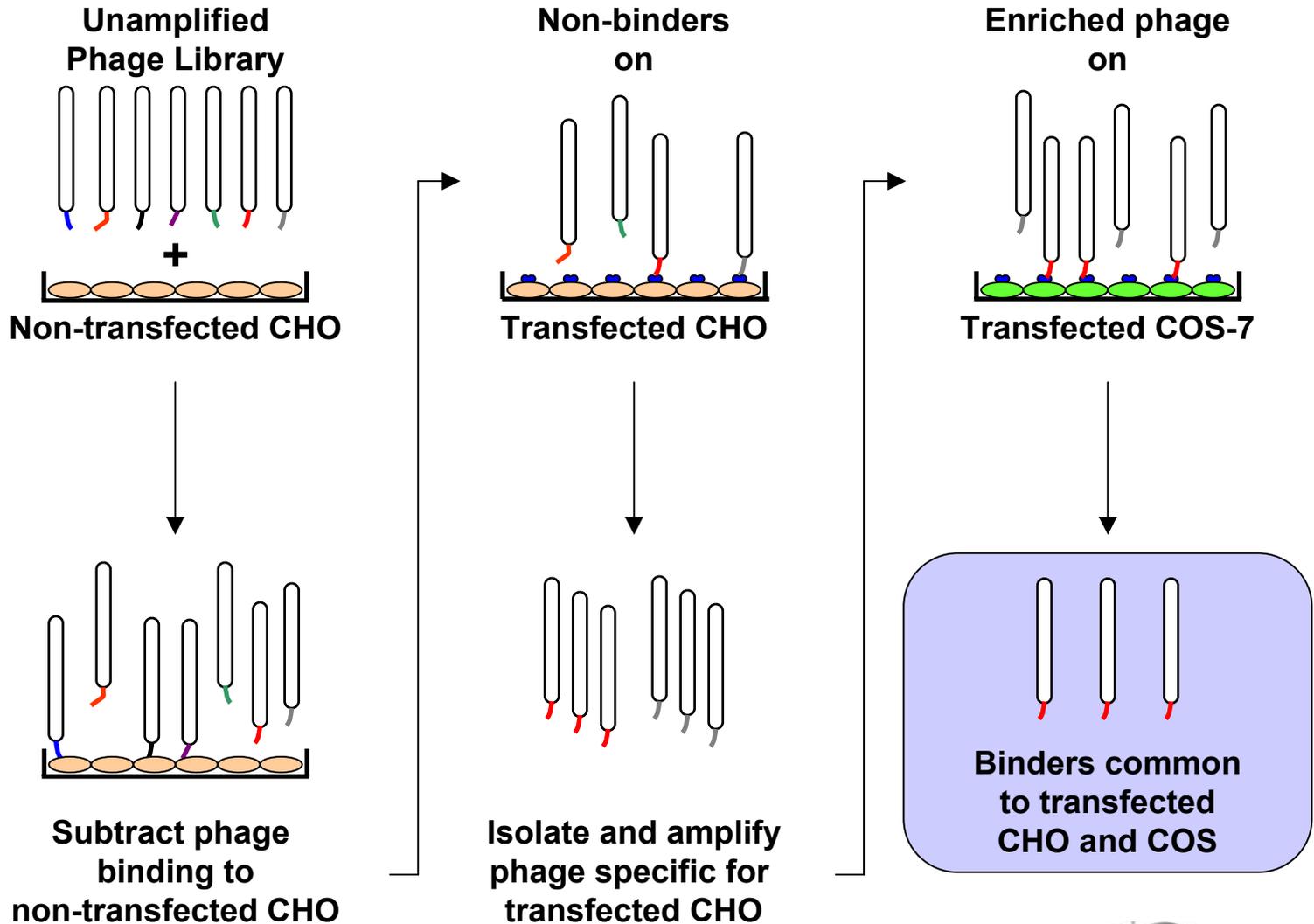


- Transfected cells and cells in vivo show different subcellular patterns
  - Enterocytes and transfected cells suggest tight junction distribution
  - M cells show cytoplasmic distribution

# Phage Display Ligand Selection



# Cell Based Subtractive Panning



# Consensus Ligand Sequences

- M13 Phage DisplayTarget = CHO transfected with EDD1S\_48 mouse + V5/poly-His

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•
•
48e2c3r2_s08  ~~TsflqaY~  ~~~~~~  ~~~
48e2c3r2_s12  ~~~~~idsYa AL~~~~~  ~~~
48e2c3r2_s10  dMrTlld~~~  ~~~~~~  ~~~
48e2c3r2_s14  ~MqTvrnh~~  ~~~~~~  ~~~
48e2c3r2_s03  ~~TTinrSp~  ~~~~~~  ~~~
48e2c3r2_s16  ~vTTYvrf~~  ~~~~~~  ~~~
48e2c3r2_s02  ~~~~~msSdk Af~~~~~  ~~~
48e2c3r2_s18  ~~~~~mtSqr sL~~~~~  ~~~
48e2c3r2_s05  ~~~~~~  ~Msqsl1P~~  ~~~
48e2c3r2_s17  ~~~~~~  ~LnisflP~~  ~~~
48e2c3r2_s19  ~~~~~~  ~mViiPpq  ~~~
48e2c3r2_s07  ~~~~~~  ~m1tPW r h~~
48e2c3r2_s09  ~~~~~~  ~~~~~APWa lar
48e2c3r2_s11  ~~~~~~  ~MTSIEAP~~  ~~~
48e2c3r2_s13  ~~~~~~  ~MTSIEAP~~  ~~~
48e2c3r2_s15  ~~~~~~  ~MTSIEAP~~  ~~~
48e2c3r2_s01  ~~~~~~  ~~~~~mAPsp Rm~
48e2c3r2_s20  ~~~~~~  ~~~~~mAPhf Rd~
Consensus    -MTT---SY-  AMTSI-APW-  R--
  
```

# Ligand-mediated Vaccine Delivery

- Q1: Can the FAE and M cells be molecularly defined?
- A1: Yes!
  - Specific genes define differentiation of FAE, distinguishing from enterocytes
  - FAE subsets can be further defined by expression of specific marker genes (e.g., PRGP-S versus PGRP-L)
  - M cell specific genes can be identified
  - Genes show conservation across species and across Mucosal-Associated Lymphoid Tissues

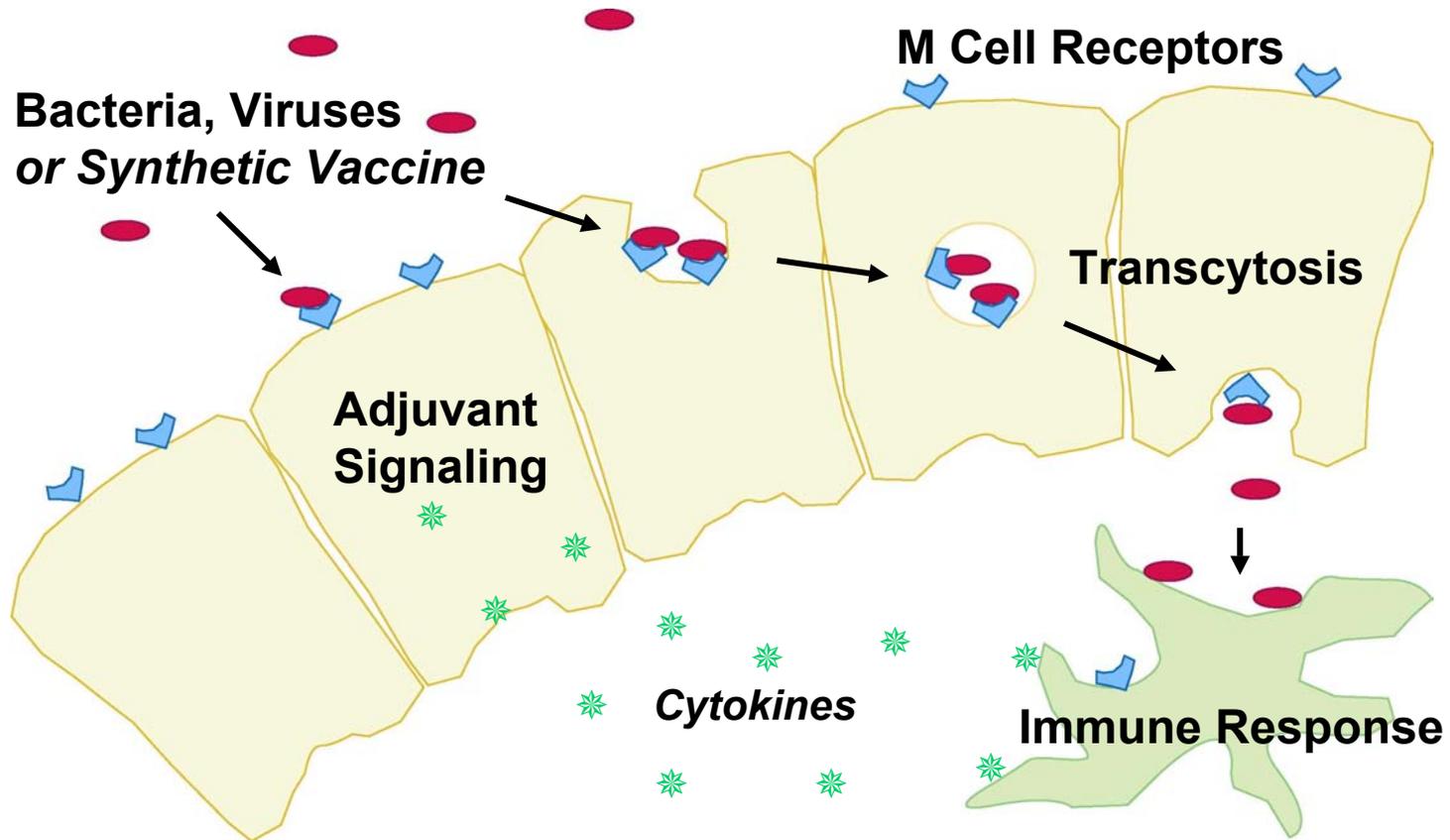
# Ligand-mediated Vaccine Delivery

- Q2: Does the FAE have functions helpful to mucosal immunity?
- A2: Yes.
  - Genes specific to FAE (CPE-R, PGRPs) may provide specific antigen/particle binding and transport function
  - FAE specific Pattern Recognition Receptors (PRR) may play a role in adjuvant signaling

# Ligand-mediated Vaccine Delivery

- Q3: How do we exploit the biology for vaccine delivery?
- A3: (Yes?)
  - Use synthetic ligands to provide targeted delivery of vaccine antigens
    - Doorstep versus Mailslot
  - Use Pattern Recognition Receptor (PRR) signaling to provide specific mucosal adjuvant activity

# Exploiting Normal Biology in Vaccine Development



# Acknowledgements

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