

FMDV control of innate immunity



Teresa de los Santos

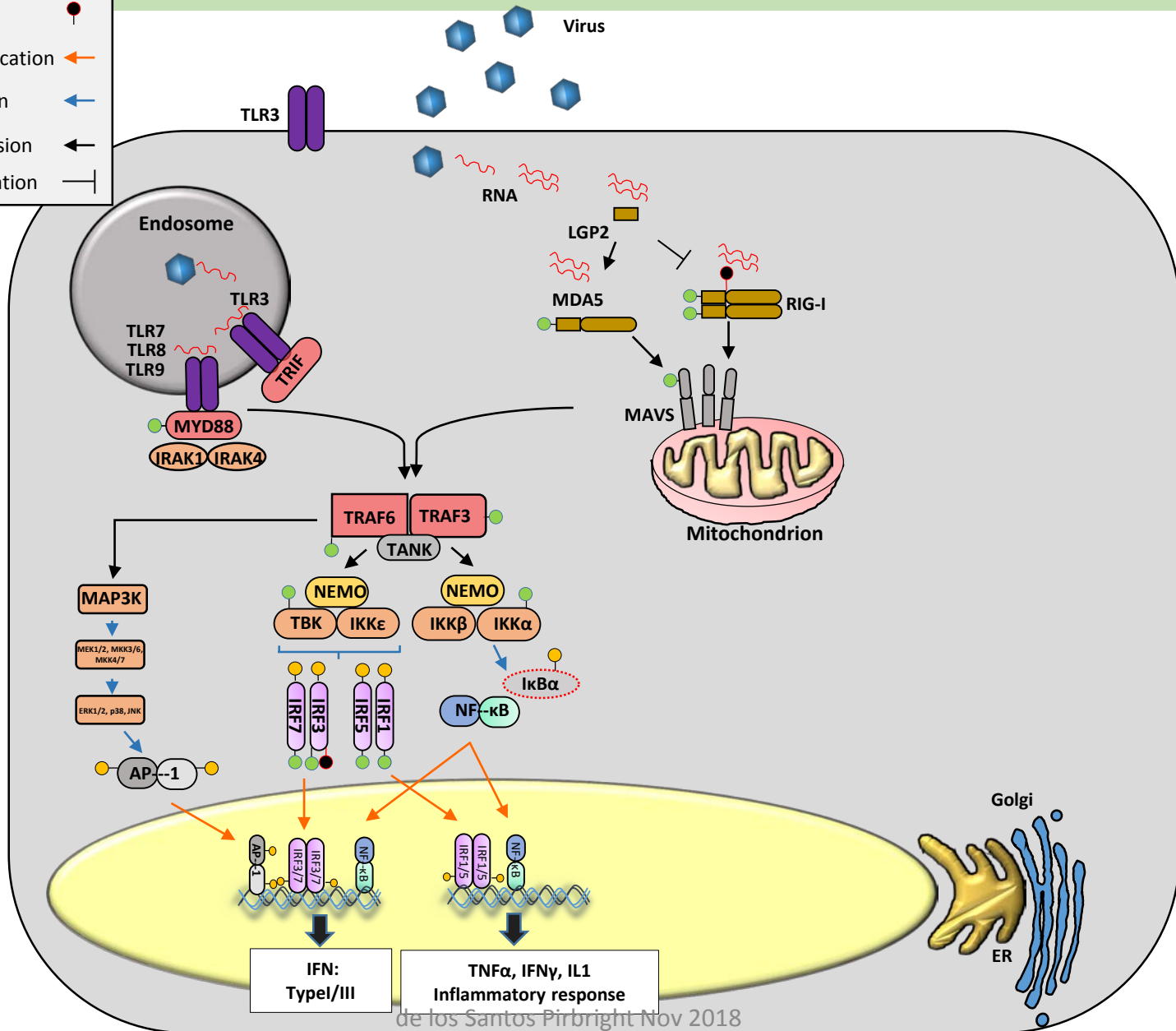
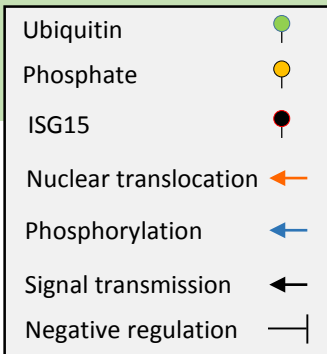
Plum Island Animal Disease Center. ARS-USDA

- Viruses are the most abundant biological entities on Earth: one to two orders of magnitude higher than cells
 - Coevolution: viruses exploit all conceivable replication-expression strategies within a host that is indispensable for survival
- *How has FMDV become such a successful pathogen?*

FMDV favorable evolutionary advantages

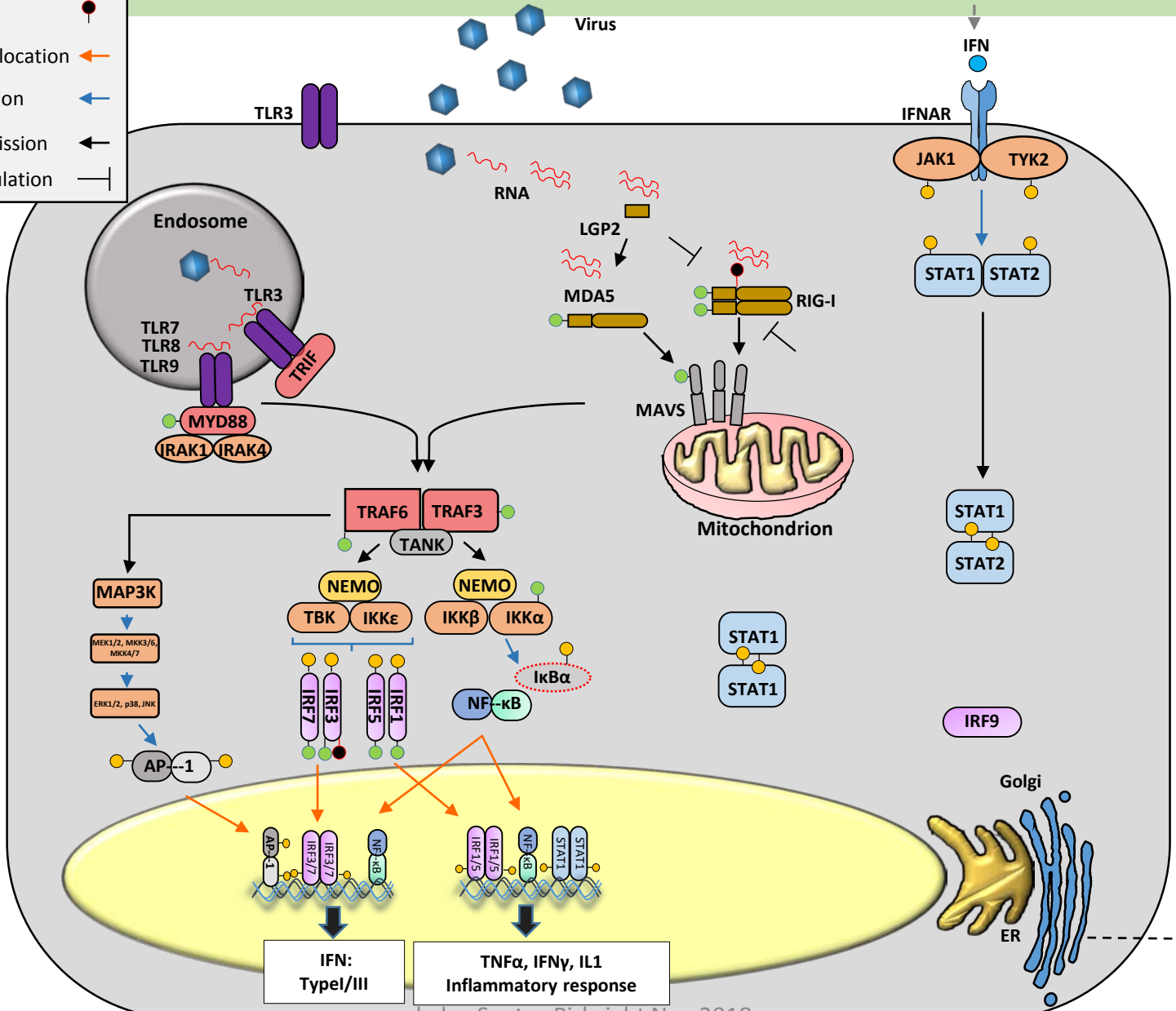
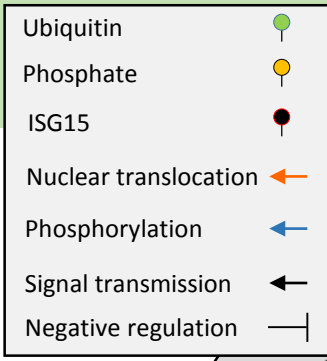
- FMDV causes an acute infection of $\sim 100\%$ morbidity but disease resolves relatively quickly inducing strong adaptive immunity
- FMDV contains an error prone polymerase (quasispecies and easiness to mutate for immune escape).
- FMDV displays flexibility on receptor usage (heterodimeric integrins, heparan sulphate, JMJD6)
- It can basically use its entire genome to counteract the host immediate innate response to infection

How is FMDV recognized by the host innate immune system?



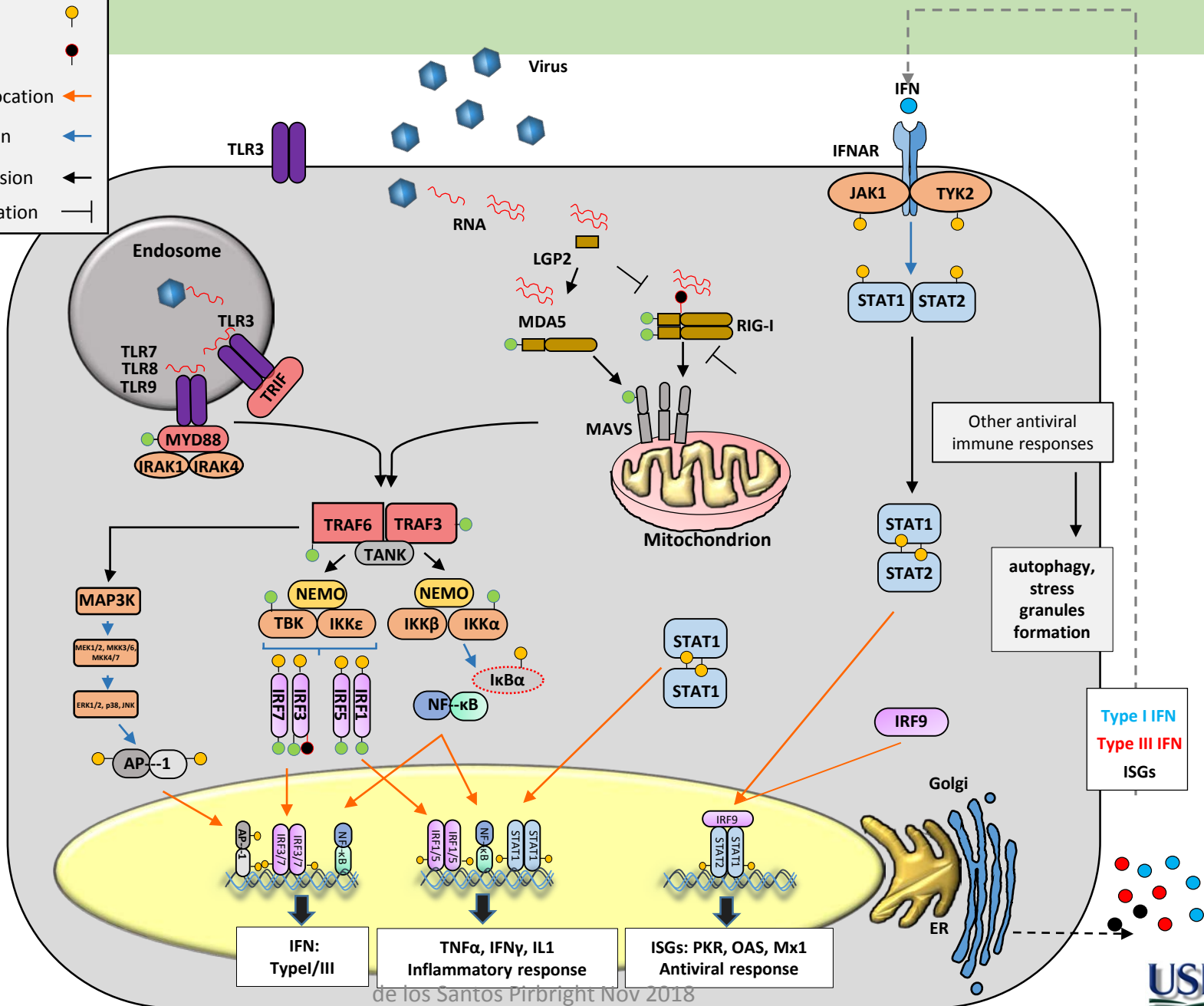
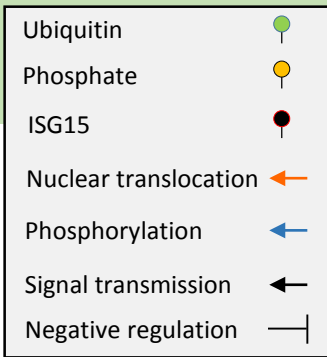
de los Santos Pirbright Nov 2018





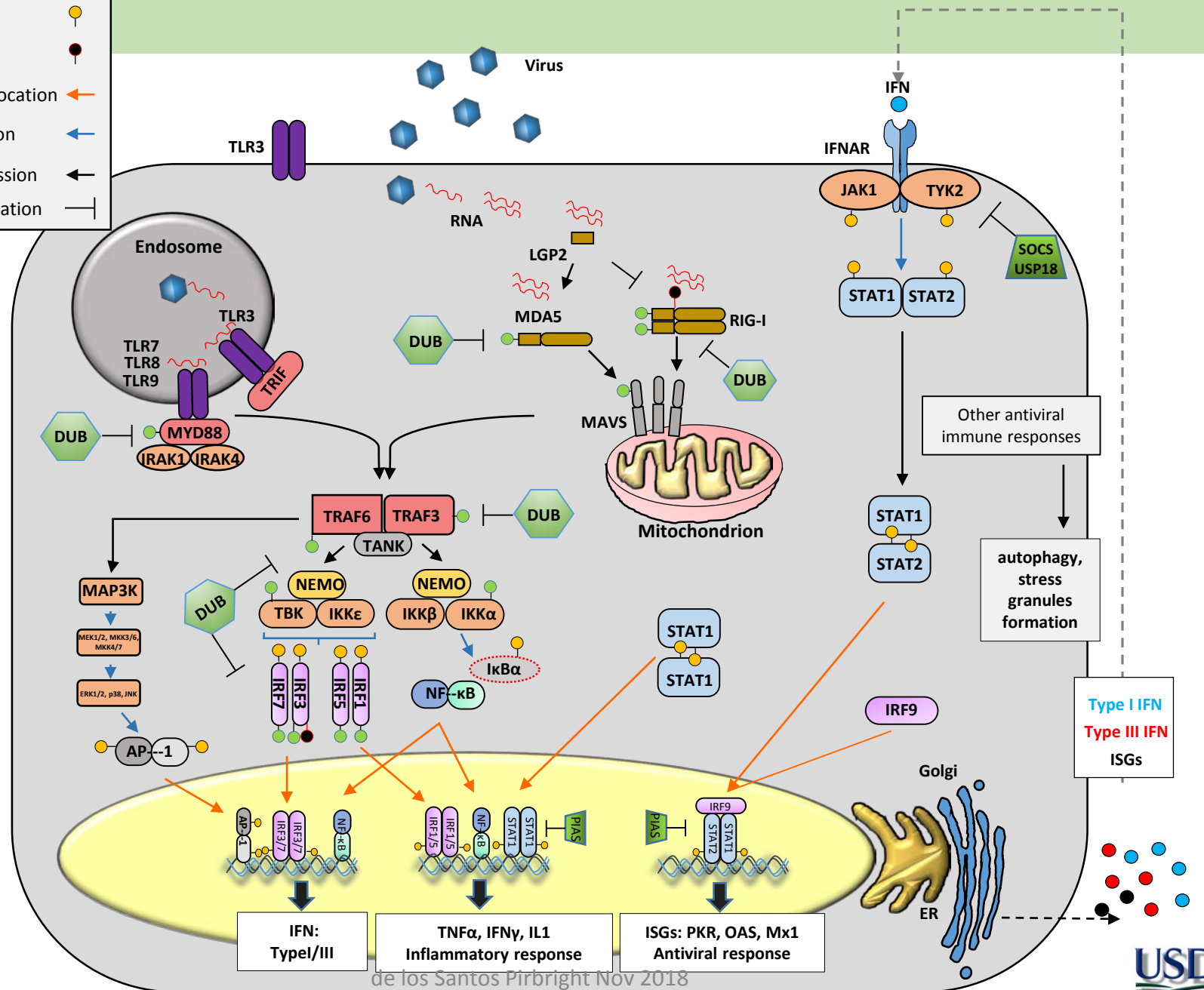
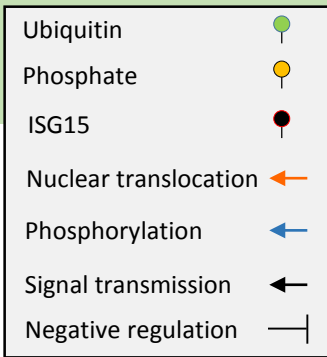
Type I IFN
Type III IFN





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Additional cellular processes
also affect innate immunity

Stress granules: organelles that accumulate host RNA and translation machinery upon stress. SGs may also serve as a platform for RIG-I dependent recognition of foreign RNA. MDA5 has also been detected in SGs.

Viruses usually block SG formation to usurp the translation machinery

Autophagy: cell recycling mechanism/self-eating/ programmed cell survival. Energy favorable.

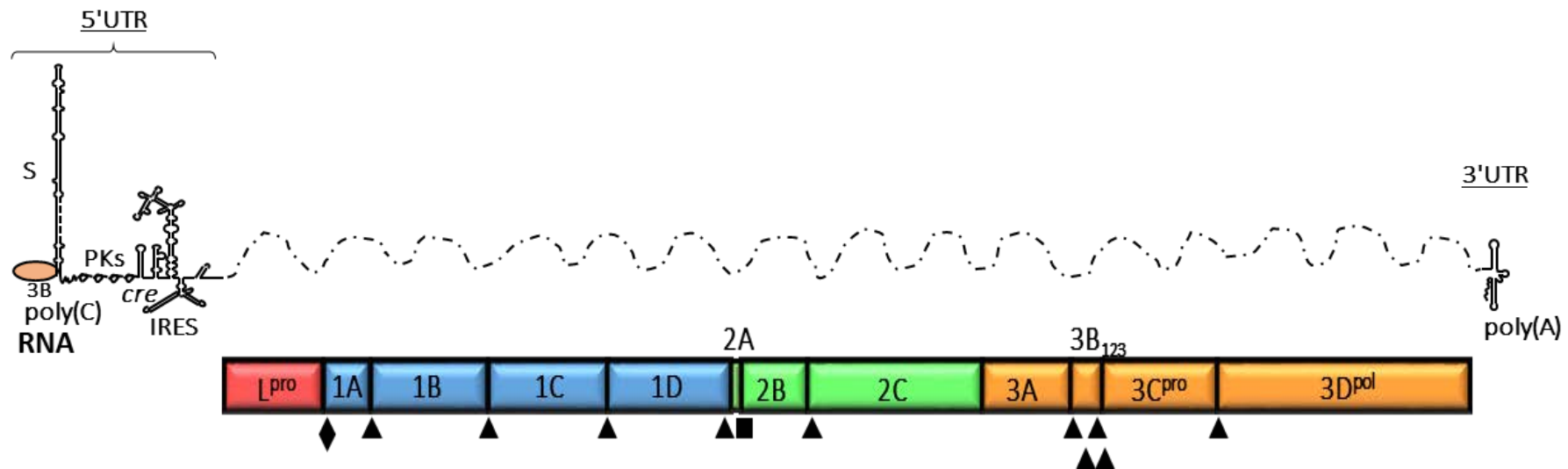
IFN enables autophagy. Activation of Myd88 and TRIFs allow for binding to Beclin1 and positively regulate autophagy (positive loop).

TLR7 dependent induction of IFN in pDCs requires ATG5 (in turn FMDV takes advantage of autophagy and virus endocytosis and replication are stimulated)

Apoptosis: the process of cell death that leads to cytoskeletal disintegration, metabolic unbalance and genomic fragmentation.

Apoptosis is induced upon Picornavirus infection but it is abortive. Completion of apoptosis leads to elimination of infected cells, a host innate response against infection.

FMDV battle against the IFN response



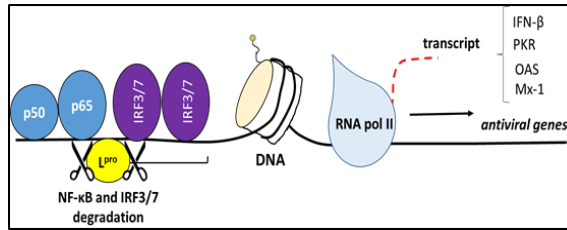
~ 8 kb

14 proteins

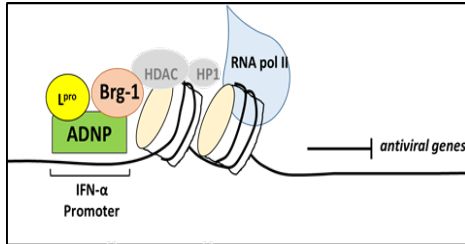
2 UTRs

At least 8 viral proteins: L, 2B, 2C, 3A, 3C, VP0, VP1, VP3 and UTRs have been shown as involved in the modulation or control of the host innate response !!!! Couldn't be more efficient..

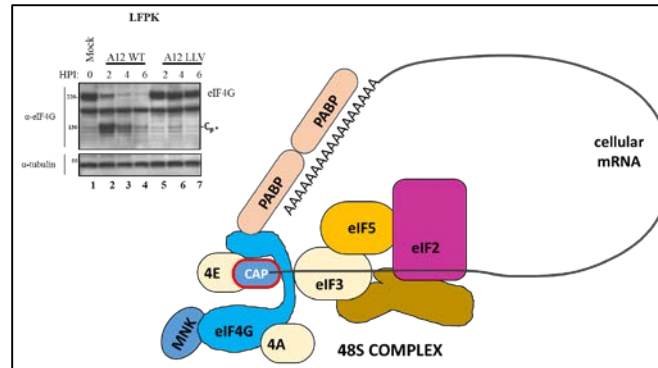
Leader: a security protein



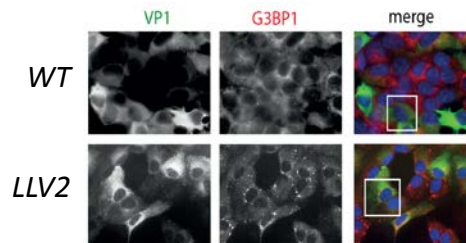
NF-κB and IRF3/7 degradation
 Reduced IFN transcription
 (deloSantos '06;'07;'09; Wang '10)



Chromatin remodeling
 IFN transcription repression
 (Medina '17)

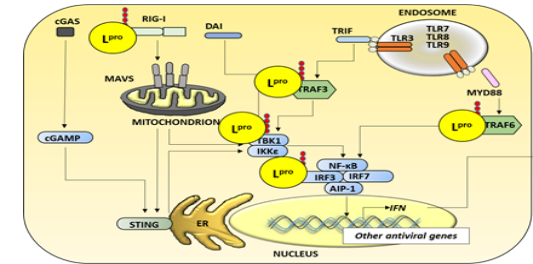


eIF-4G cleavage
 Translation shut off
 (Devaney '88)

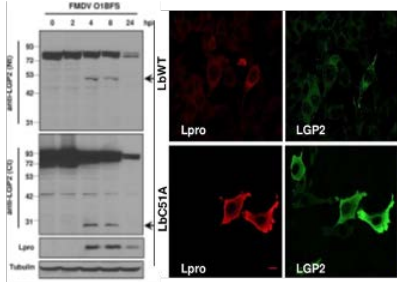


Interference with SG formation

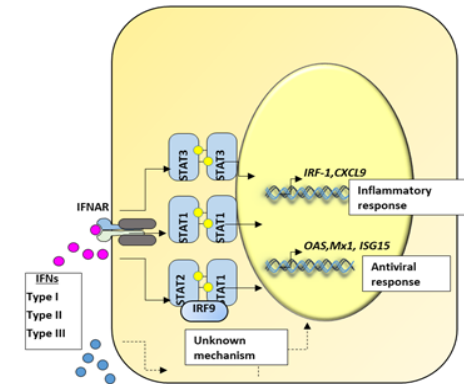
Affects stress response
 (Visser, subm)



De-Ubiquitination
 Inactive RIGI, TRAF3/6, IRF3/7
 (Wang '11)



LGP2 cleavage
 Reduced IFN transcription
 (Rodriguez-Pulido '18)

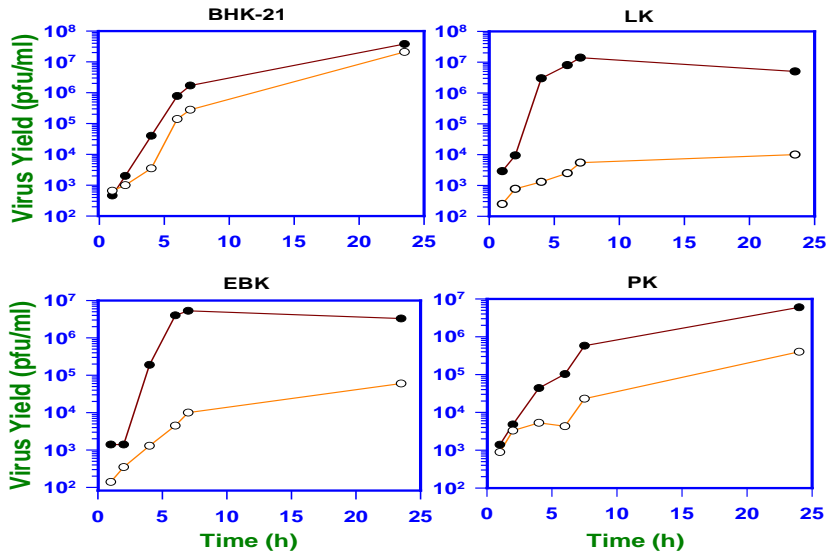
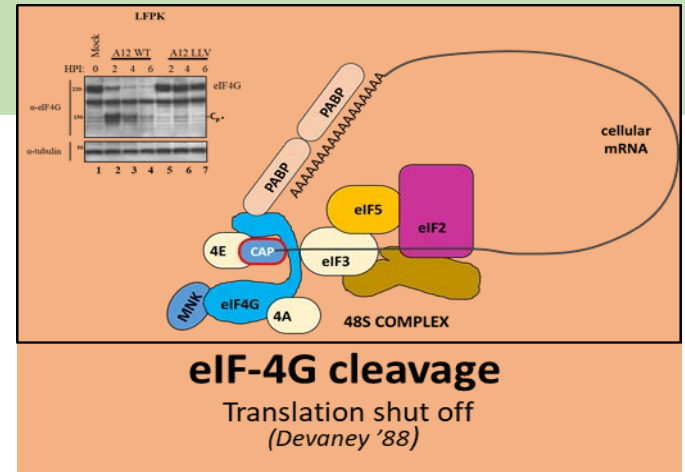


De-ISGylation
 Affects IFN paracrine signaling?
 (Swatek '18)

Ability of Foot-and-Mouth Disease Virus To Form Plaques in Cell Culture Is Associated with Suppression of Alpha/Beta Interferon

JARASVECH CHINSANGARAM, MARIA E. PICCONE,[†] AND MARVIN J. GRUBMAN*

Plum Island Animal Disease Center, North Atlantic Area, Agricultural Research Service, U.S. Department of Agriculture, Greenport, New York 11944



Cell supernatant source	Inducing virus	MOI	Antiviral units
EBK	None		<2
	LLV2	1	32
	WT	1	ND
	LLV2	10	>64
	WT	10	8
LK	None		<2
	LLV2	1	>64
	WT	1	4
	LLV2	10	64
	WT	10	ND
PK	None		<2
	LLV2	1	16
	WT	1	<2
	LLV2	10	32
	WT	10	<2
BHK	None		<2
	LLV2	1	<2
	WT	1	-
	LLV2	10	<2
	WT	10	-

Inhibition of L-Deleted Foot-and-Mouth Disease Virus Replication by Alpha/Beta Interferon Involves Double-Stranded RNA-Dependent Protein Kinase

JARASVECH CHINSANGARAM, MARLA KOSTER, AND MARVIN J. GRUBMAN*

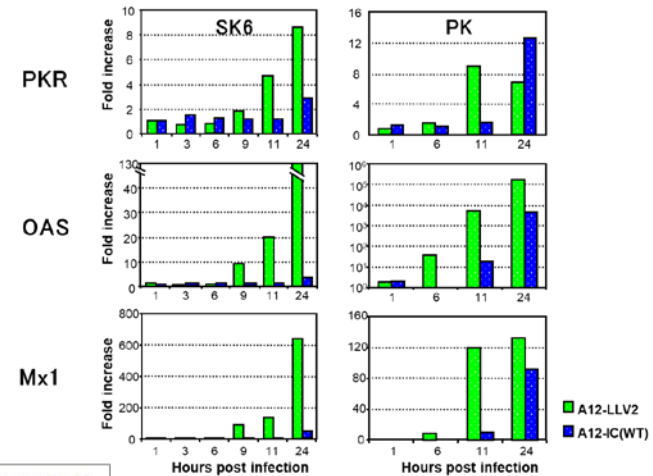
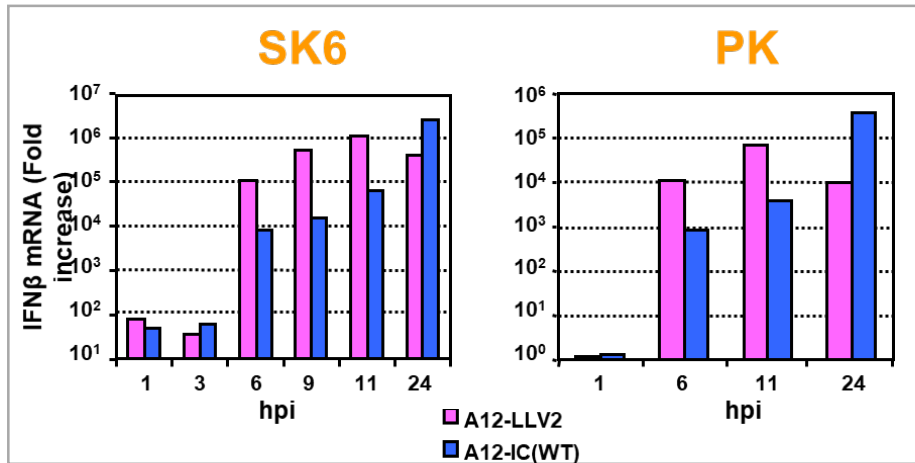
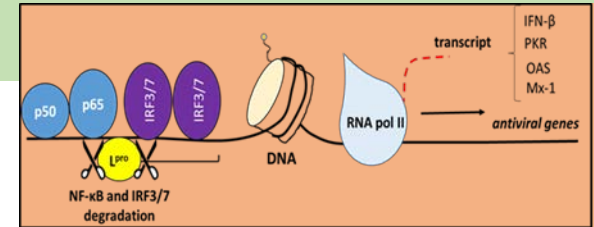
de los Santos Pirbright Nov 2018



The Leader Proteinase of Foot-and-Mouth Disease Virus Inhibits the Induction of Beta Interferon mRNA and Blocks the Host Innate Immune Response

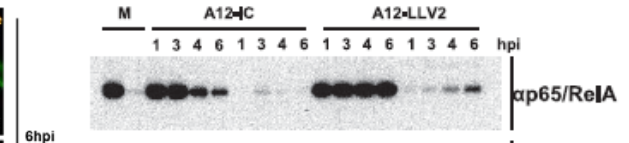
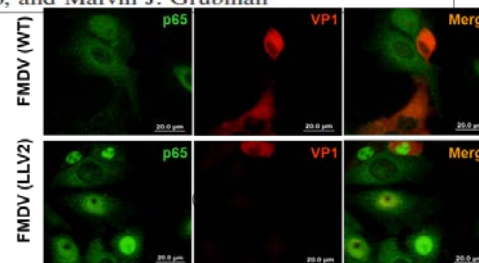
Teresa de los Santos,[†] Sonia de Avila Botton,^{†‡} Rudi Weiblen,[‡] and Marvin J. Grubman*

Plum Island Animal Disease Center, Agricultural Research Service, U.S. Department of Agriculture, Greenport, New York 11944



Degradation of Nuclear Factor Kappa B during Foot-and-Mouth Disease Virus Infection[▽]

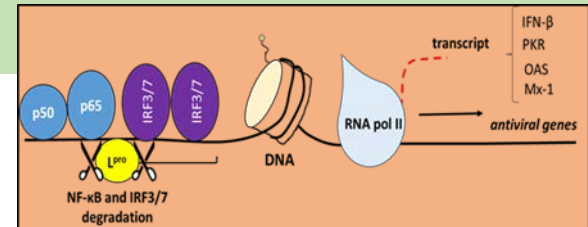
Teresa de los Santos, Fayna Diaz-San Segundo, and Marvin J. Grubman*



SAP

A Conserved Domain in the Leader Proteinase of Foot-and-Mouth Disease Virus Is Required for Proper Subcellular Localization and Function⁹

Teresa de los Santos,^a Fayna Diaz-San Segundo, James Zhu, Maria Kosier, Camila C. A. Dias, and Marvin J. Grubman



NF-κB and IRF3/7 degradation

Reduced IFN transcription

(delosSantos '06;'07;'09; Wang '10)

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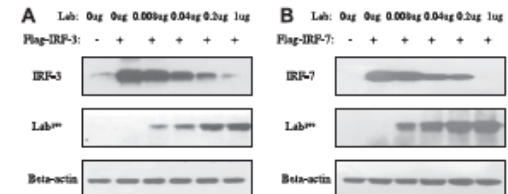
journal homepage: www.elsevier.com/locate/ybbr



Foot-and-mouth disease virus leader proteinase inhibits dsRNA-induced type I interferon transcription by decreasing interferon regulatory factor 3/7 in protein levels

Dang Wang, Liurong Fang, Rui Luo, Rui Ye, Ying Fang, Lilan Xie, Huanchun Chen, Shaobo Xiao *

Division of Animal Infectious Diseases, State Key Laboratory of Agricultural Microbiology, College of Veterinary Medicine, Huazhong Agricultural University, Wuhan 430070, PR China



39 differentially expressed genes containing promoters with canonical NF-κB and IRF binding sites

	P	Q	R	S	T	U	V
1	Matrix	TFBS	Name	Threshold	Hits	Probability	
2	M00043	ISDL_01	dorsal	0.89	37	2.91E-12	
3	M00052	V\$NFKAPPAB65_01	NF-kappaB (p65)	0.84	160	1.05E-11	
4	M00053	V\$CREL_01	c-Rel	0.83	293	1.14E-15	
5	M00062	V\$IRF1_01	IRF-1	0.85	78	1.61E-11	
6	M00063	V\$IRF2_01	IRF-2	0.86	41	2.30E-10	
7	M00171	ISADF1_Q6	Adf-1	0.86	229	5.61E-28	
8	M00194	V\$NFKB_Q6	NF-kappaB	0.78	365	2.16E-18	
9	M00208	V\$NFKB_C	NF-kappaB binding site	0.84	95	4.74E-15	
10	M00253	V\$CAP_01	transcription initiation	0.98	193	1.97E-06	
11	M00258	V\$ISRE_01	ISGF-3	0.85	64	2.37E-21	
12	M00051	V\$NFKAPPAB50_01	NF-kappaB (p50)	0.79	191		
13	M00054	V\$NFKAPPAB_01	NF-kappaB	0.87	112		
14							

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Virology

journal homepage: www.elsevier.com/locate/yviro



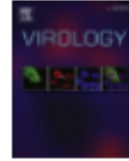
Differential gene expression in bovine cells infected with wild type and leaderless foot-and-mouth disease virus

James Zhu^a, Marcelo Weiss^{ab}, Marvin J. Grubman^a, Teresa de los Santos^a, Pirbright Nov 2018

^a Plum Island Animal Disease Center, North Atlantic Area, Agricultural Research Service, U.S. Department of Agriculture, Greenport, New York 11944, USA

^b Oak Ridge Institute for Science and Education, PIADC Research Participation Program, Oak Ridge, TN 37831, USA

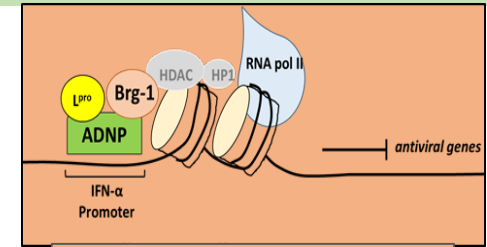




Interaction between FMDV L^{pro} and transcription factor ADNP is required for optimal viral replication

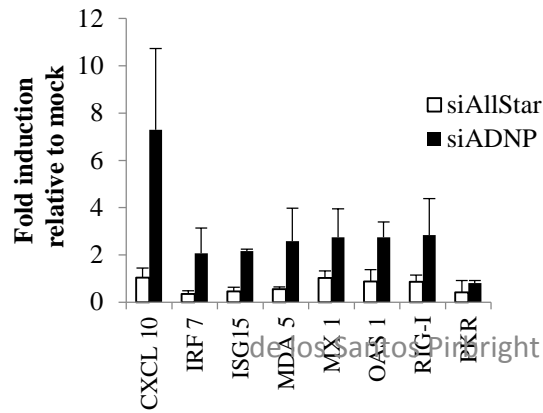
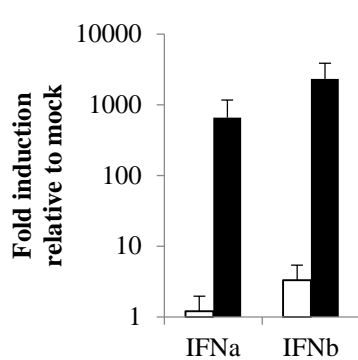
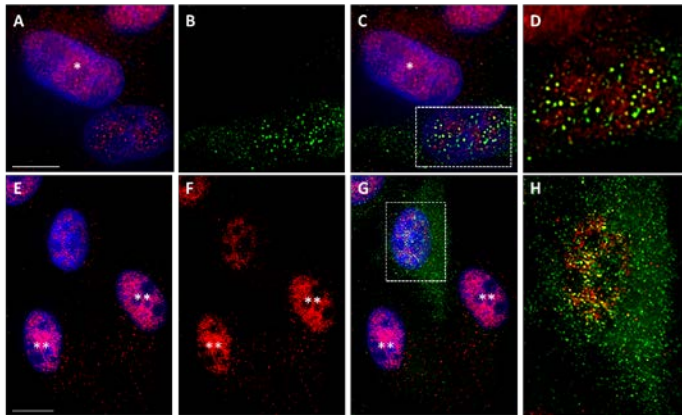


Gisselle N. Medina^a, Giselle M. Knudsen^c, Alexander L. Greninger^b, Anna Kloc^{a,d}, Fayna Díaz-San Segundo^a, Elizabeth Rieder^a, Marvin J. Grubman^a, Joseph L. DeRisi^b, Teresa de los Santos^{a,*}

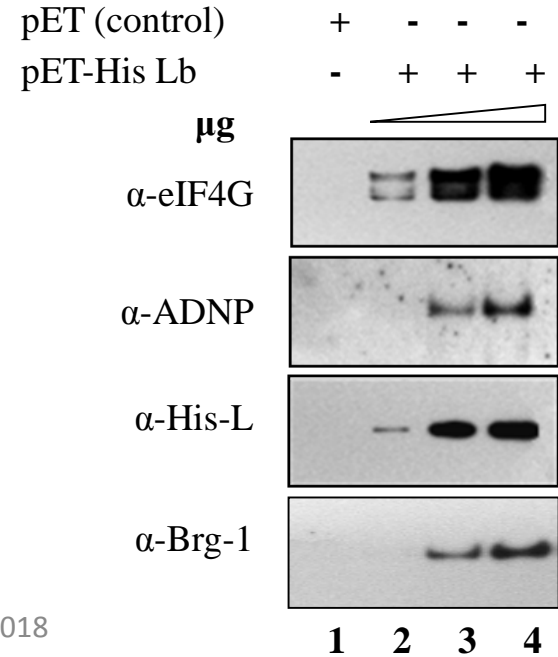


Chromatin remodeling
IFN transcription repression
Medina '17

ADNP colocalizes with Lpro



Lpro interacts with ADNP and BRG1 (SNF/SWI complex)

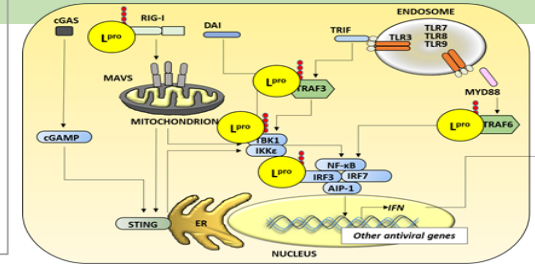


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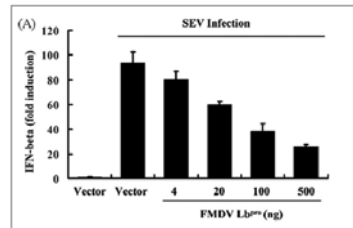
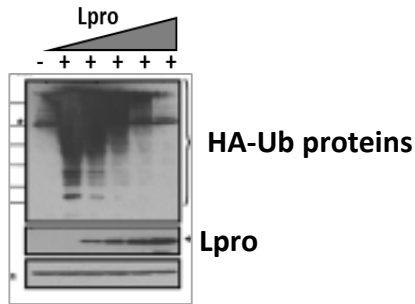


The Leader Proteinase of Foot-and-Mouth Disease Virus Negatively Regulates the Type I Interferon Pathway by Acting as a Viral Deubiquitinase[†]

Dang Wang,¹ Liurong Fang,¹ Ping Li,¹ Li Sun,² Jinxiu Fan,¹ Qingye Zhang,³ Rui Luo,¹ Xiangtao Liu,⁴ Kui Li,⁵ Huanchun Chen,¹ Zhongbin Chen,^{2,4} and Shaobo Xiao^{1,6}



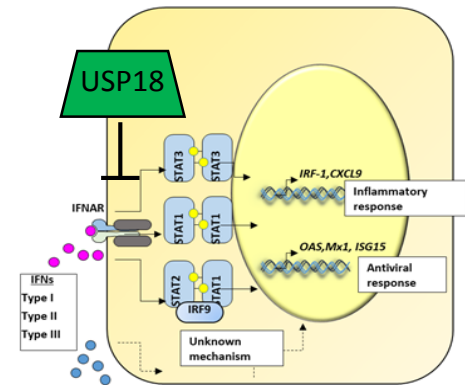
De-Ubiquitination
 Inactive RIGI, TRAF3/6, IRF3/7
 (Wang '11)



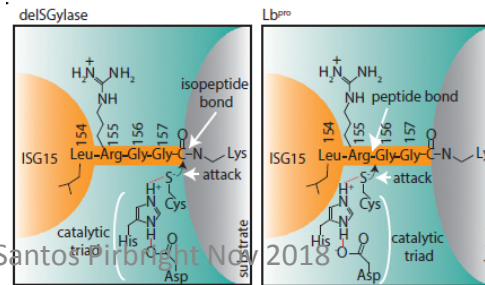
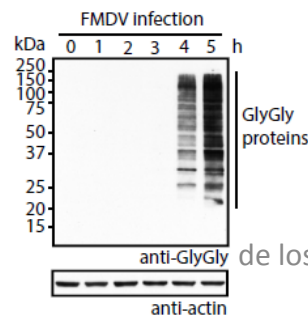
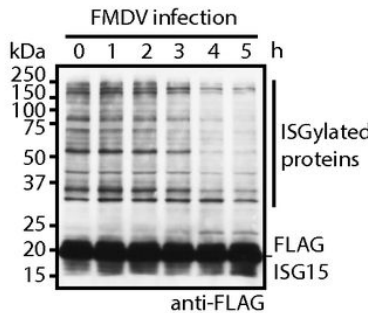
Irreversible inactivation of ISG15 by a viral leader protease enables alternative infection detection strategies

Kirby N. Swatek^a, Martina Aumayr^{b,1}, Jonathan N. Pruneda^{a,1}, Linda J. Visser^{c,1}, Stephen Berryman^d, Anja F. Kueck^a, Paul P. Geurink^e, Huib Ovaa^e, Frank J. M. van Kuppeveld^c, Tobias J. Tuthill^d, Tim Skern^b, and David Komander^{a,2}

Check for updates

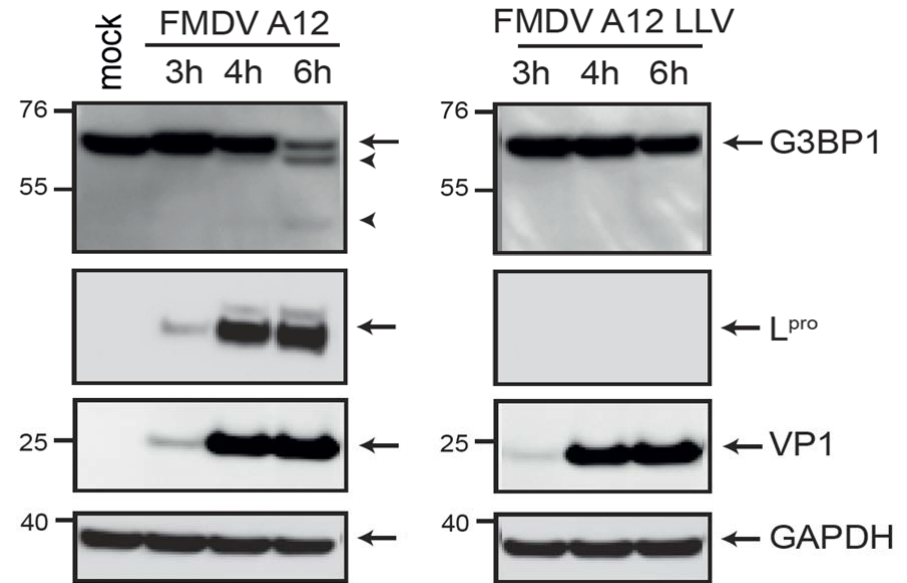
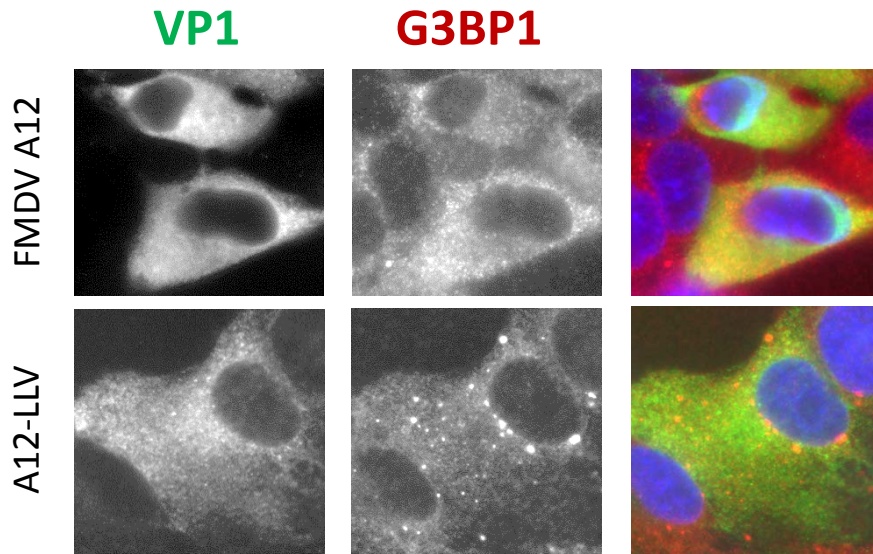


De-ISGylation
 Affects IFN paracrine signaling?
 (Swatek '18)



L^{pro} suppresses SG formation and cleaves G3BP1/2

Visser et al, 2018 submitted



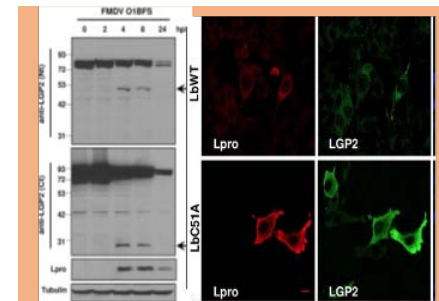
PLOS PATHOGENS
June 29, 2018

RESEARCH ARTICLE

Innate immune sensor LGP2 is cleaved by the Leader protease of foot-and-mouth disease virus

Miguel Rodríguez Pulido¹, María Teresa Sánchez-Aparicio^{2,3}, Encarnación Martínez-Salas¹, Adolfo García-Sastre^{2,3,4}, Francisco Sobrino¹, Margarita Sáiz^{1*}

de los Santos Pirbright Nov 2018



LGP2 cleavage

Reduced IFN transcription
(Rodríguez-Pulido '18)

Leader: a security protein

Dispensable for virus replication but is directly associated to virulence and pathogenicity in cells and in cattle and swine

1. Cleaves eIF4G shutting off host cell translation (IFN and ISG)
2. Induces degradation of NF- κ B, IRF3 and IRF7 reducing transcription of IFN, inflammatory cytokines and ISGs
3. Removes Ubiquitin from key signal molecules in IFN induction: RIG1, TRAF3/6, TBK1
4. Interacts with transcription factors ADNP and BRG1, involved in the epigenetic control of IFN expression (transcriptional repression)
5. Interacts and causes degradation of LGP2 affecting IFN induction and favoring FMDV replication
6. Removes ISG15 modifier possibly affecting innate responses
7. Cleaves G3BP1/2 to prevent SG formation possibly affecting innate responses

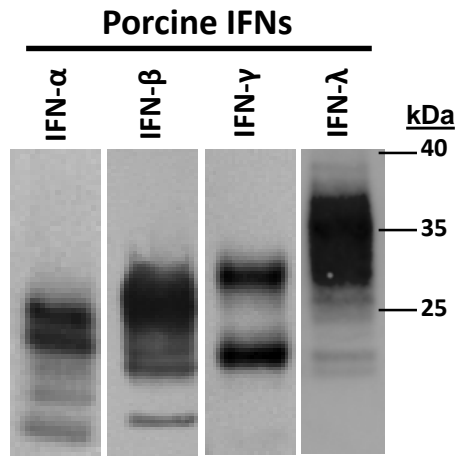
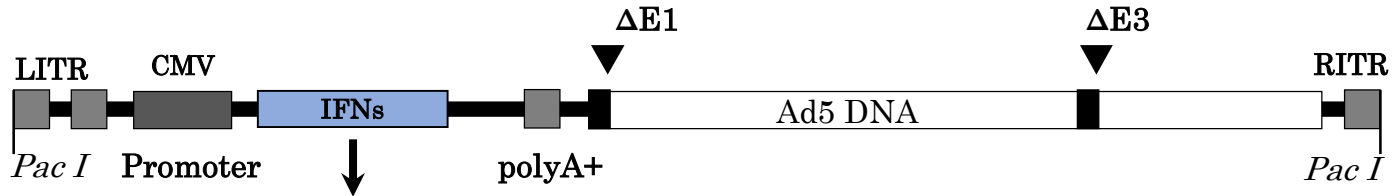
FMDV factor	Affected process	Viral counter-mechanism
L ^{pro}	Translation and transcription	<ul style="list-style-type: none"> • eIF4G1 cleavage (Devaney 1988; Kirchweger 1994) • Gemin5 cleavage (Pineiro 2012) • Decreased amounts of IFNβ (de los Santos 2007) • Degradation of NF-κB (de los Santos 2007) • Modulation of PKR (Chinsangaram 2001; de los Santos 2006) • Inhibition of RANTES (de los Santos 2006; Wang 2010) • Induction of ADNP binding to IFNα promoter to disrupt the expression of IFN and ISGs (Medina 2017) • Deubiquitination of proteins involved in innate immunity signaling (RIG-I, TBK1, TRAF3, TRAF6) (Wang 2011) • DeISGylation (Swatek 2018) • Modulation of IFNβ expression through interaction with LGP2 (Rodriguez-Pulido 2018)
2B+2C and or 2BC	Membrane rearrangements, secretion and trafficking, autophagy and modulation of ISGs expression	<ul style="list-style-type: none"> • Membrane rearrangements (Teterina 2006; Monaghan 2004) • Inhibition of MHC class I surface expression and secretion of antiviral cytokines (Moffat et al., 2005, 2007; Sanz-Parra et al., 1998) • Modulation of cytopathogenicity (Arias 2010) • Induction of autophagy (O'Donnell 2011; Berryman 2012, Gladue 2012) • Alteration of Ca²⁺ concentrations leading to autophagy (Ao 2015) • Interaction with RIG-I to suppress expression of ISGs and GBP1 (Zhu 2016) • Interaction with LGP2 (Zhu 2017; Rodriguez-Pulido 2018) • Induction of apoptosis via interaction with Nmi (Wang 2012b) • Interaction with IFN-induced protein IF35 (Zheng 2014)

FMDV factors	Affected process	Viral counter-mechanism
3A	Membranes and innate immunity signaling factors	<ul style="list-style-type: none"> ● Interaction with membranes (Gonzales-Magalaldi 2014, Lotufo 2018) ● Inhibition of RLR (RIG-I, MDA5, MAVS)-mediated IFNβ induction (Li 2016a)
3C ^{pro}	Transcription, translation and autophagy	<ul style="list-style-type: none"> ● Histone H3 cleavage (Grigera 1984; Tesar 1990; Falk 1990) ● eIF4G and eIF4A cleavage (Belsham 2000) ● Sam68 cleavage (Lawrence 2012) ● NEMO cleavage (Wang 2012a) ● Reduction of the endogenous levels of PKR (Li 2017) ● Interference of JAK-STAT signaling pathway (Du 2014) ● Degradation of autophagy proteins ATG5 and ATG12 (Fan 2017) ● Cleavage of G3BP1 (SG marker) (Galan 2017; Ye 2018)
VP1, VP2, VP3	Suppression of innate immune signaling responses (type I IFN) and autophagy	<ul style="list-style-type: none"> ● Interaction with the cellular protein sorcin to downregulate transcription of IFNα/β and NF-κB (Li 2013) ● Downregulation of TNFα and NF-κB (Ho 2014; Wang 2016) ● Induction of autophagy (Sun 2018) ● Inhibition of STAT phosphorylation (Li 2016b) ● Blockage of IRF3 phosphorylation and dimerization (Li 2016b) ● Capsid assembly requires chaperone Hsp90 (Newman 2017). This interaction may affect PRR innate signaling (Binder 2014, Andino 2018).
Untranslated regions	Modulation of innate immune signaling	<ul style="list-style-type: none"> ● 5'UTR can stimulate type I IFN responses: Mx-1, IFNβ, IL-6, TNFα, IRF7 (Rodriguez-Pulido 2011, Kloc 2017). ● 3'UTR can trigger an antiviral state via IFNβ (Rodriguez-Pulido 2011)

How can we use this
wealth of information to
develop or improve FMD
countermeasures?

Can treatment with IFN prevent FMD?

Ad5-IFN blocks FMDV infection *in vitro*



✓ Type I IFN

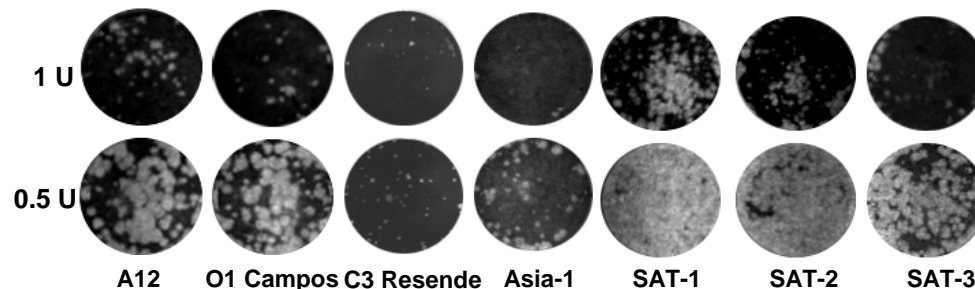
(Chinsangaram 2003; Moraes 2003; Dias 2011)

✓ Type II IFN

(Moraes 2007; Kim 2014)

✓ Type III IFN

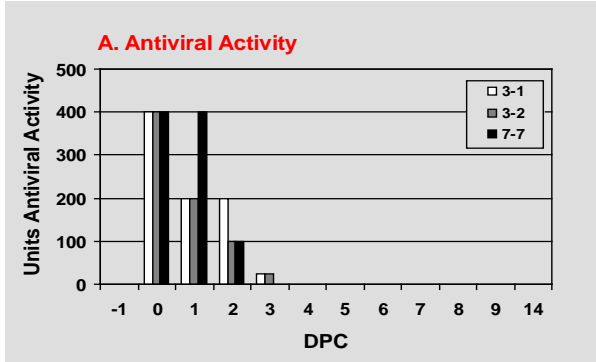
(Perez-Martin 2014; Perez-Martin 2016; Diaz-San Segundo 2017)



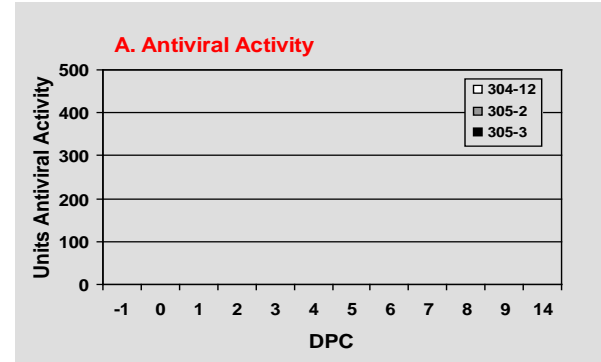
Ad5-polFN α protects swine against FMD



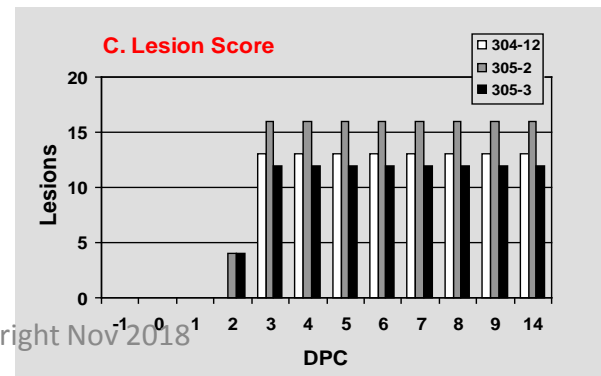
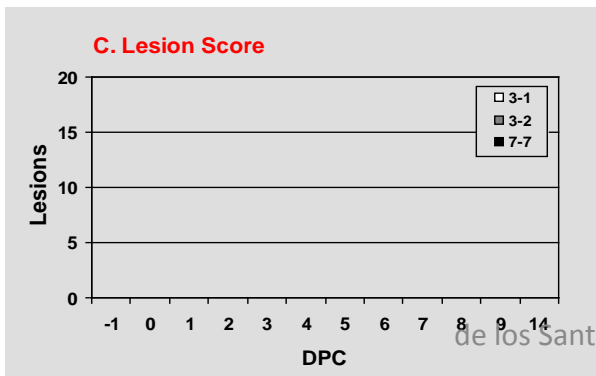
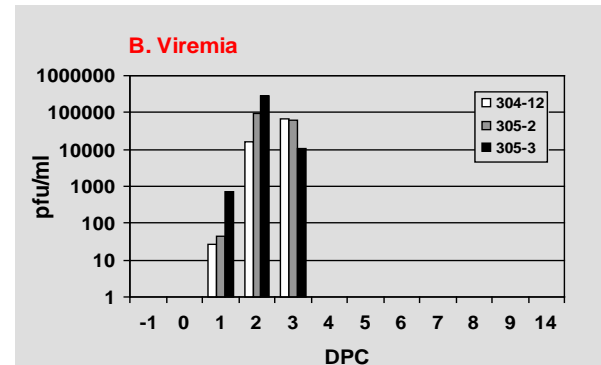
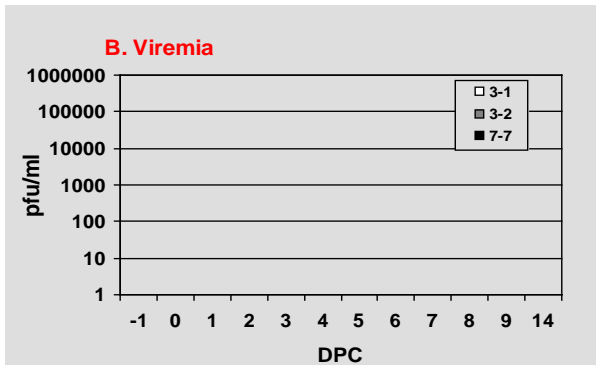
Ad5-polFN α 10⁹ pfu



Ad5-blue 10⁹ pfu



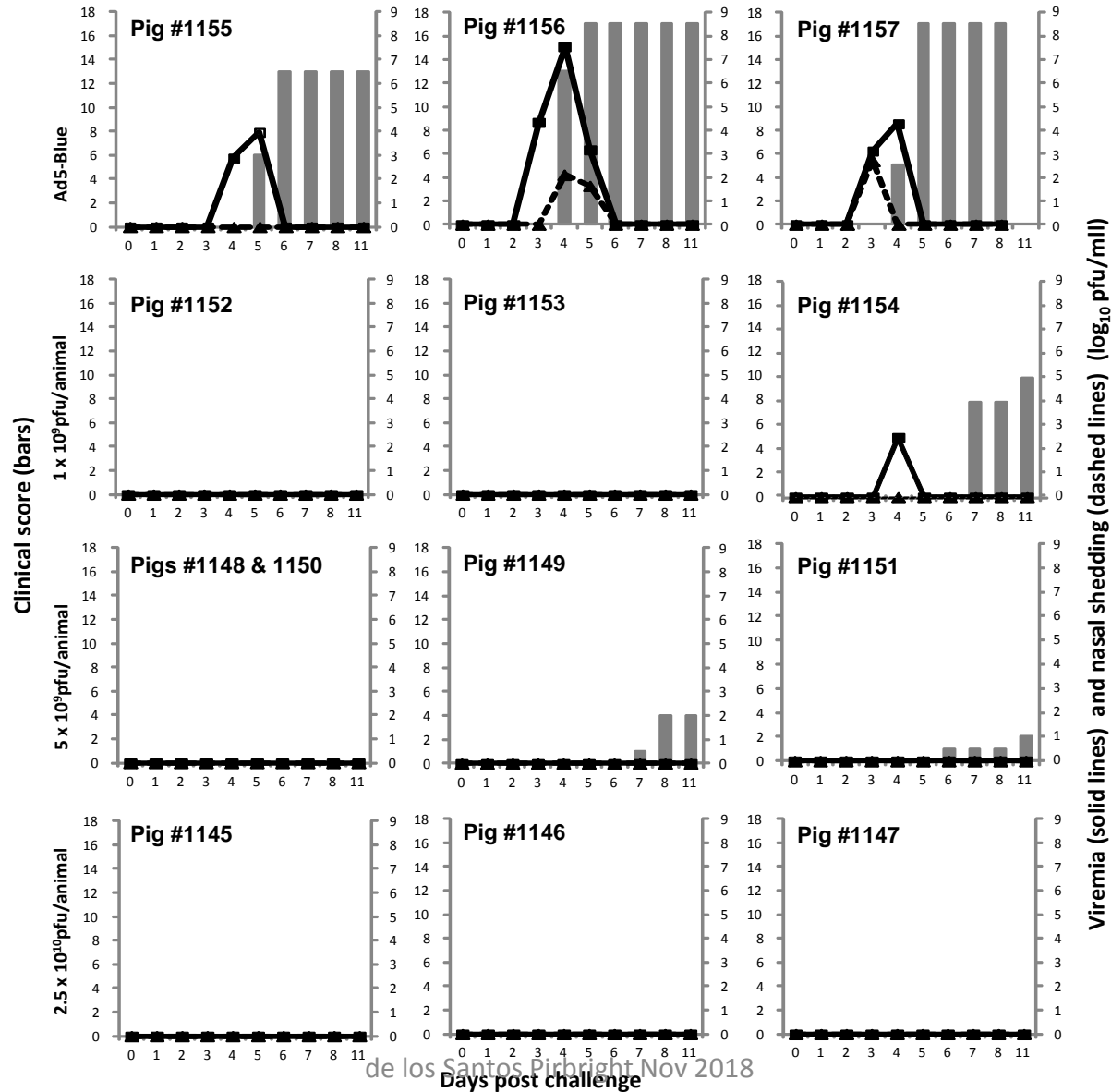
Chinsangaram 2003
 Moraes 2003
 Dias 2011



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Ad5-polFNλ protects swine against FMD

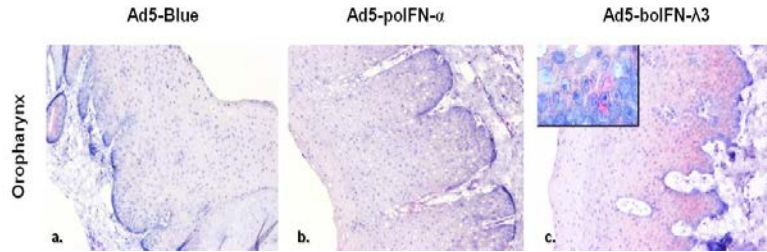


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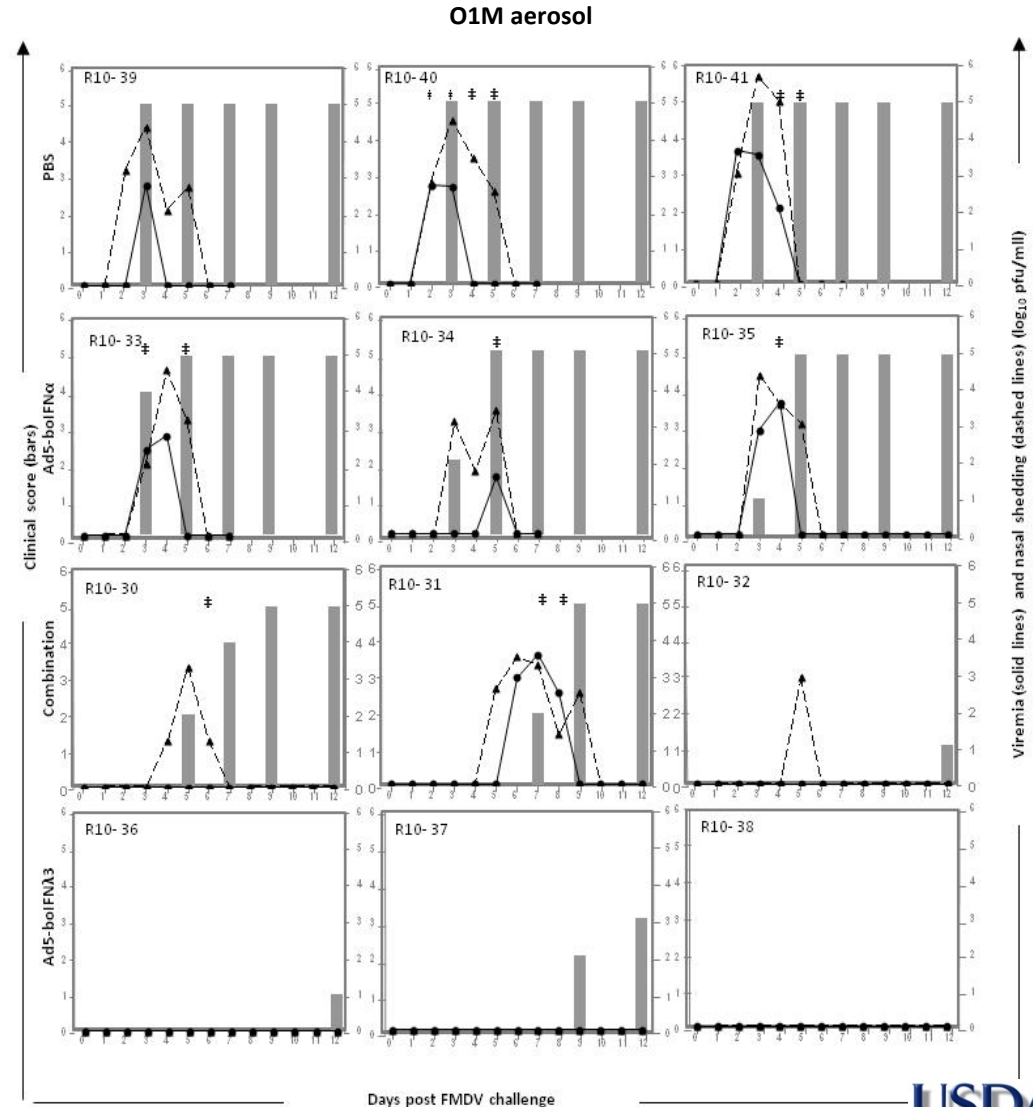
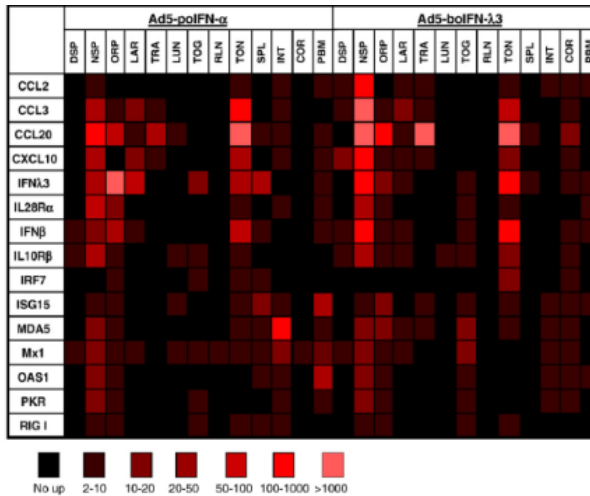




Ad5-boIFN λ 3 protects cattle against FMD



Mx1 staining



Diaz San-Segundo 2011; Perez-Martin 2012

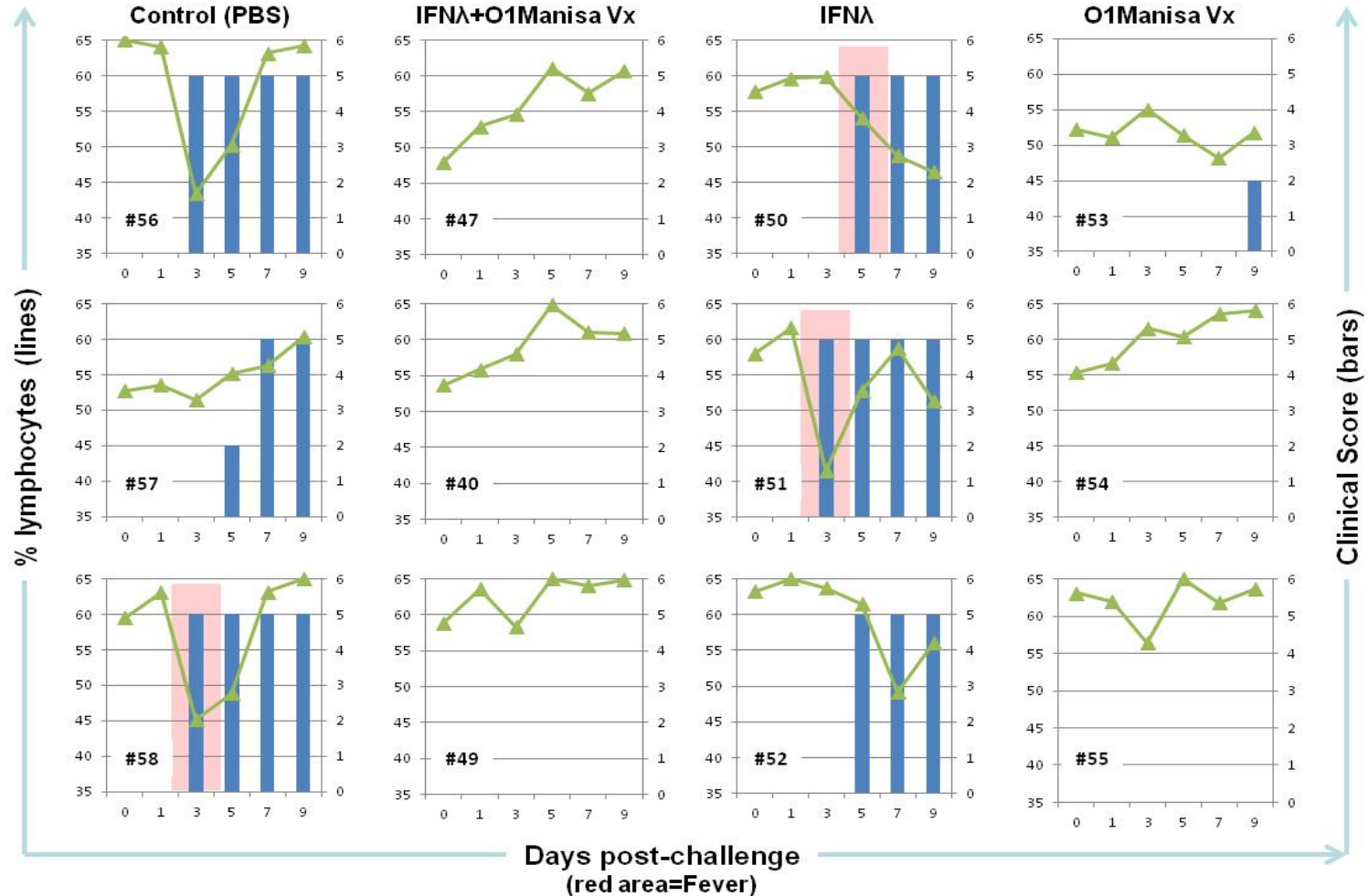
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Ad5-polIFN α and Ad5-FMD protect swine against FMD

Group	IFN (U/ml)	Viremia/dpc	Clinical score/dpc
Control			
9987	52	2.0x10 ⁵ /1	14/2
9988	1,867	1.0x10 ³ /1	13/3
9989	0	6.1x10 ⁴ /1	14/2
Ad5-A24			
9981	0	0/0	2/3
9982	395	0/0	6/3
9983	0	0/3	9/3
Ad5-IFNα			
9975	13,718	0/0	0/0
9976	15,431	0/0	0/0
9977	16,059	0/0	0/0
Ad5-IFNα +Ad5A24			
9972	17,390	0/0	0/0
9973	18,631	0/0	0/0
9974	17,654	0/0	0/0

Ad5-boIFN λ 3 and Ad5-FMD protect cattle against FMD



Biotherapeutics protect against FMD !

- Ad5- type I, II and III IFNs protect swine against FMD
- Combination treatment of Ad5-polIFN α and Ad5-FMD can fully protect swine as early as 1 day post inoculation and for 5 days (Ad5-FMD fully protects swine at 7dpi)
- Inducers of innate immunity such as Poly ICLC , VRPs and IRF7/3(5D) can protect by themselves against FMD
- ISGs including PKR, OAS, IP10 and increased numbers of DCs and NK cells in draining lymph nodes are responsible for the IFN-mediated protection against FMD in swine

- Ad5-type III IFN protects cattle against FMD
- Combination treatment of Ad5-boIFN λ and Ad5-FMD can fully protect cattle against FMD

Thank you



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PIADC Animal Research Unit



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Happy 60th Pirbright



birthday!

