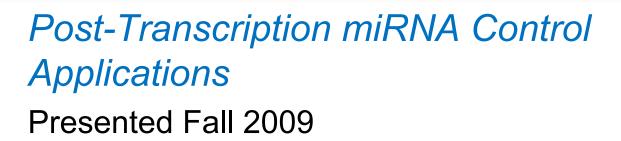
What can you do with a luciferase Reporter Assay?





Click the icon in the upper left hand corner to view speaker notes for slides.

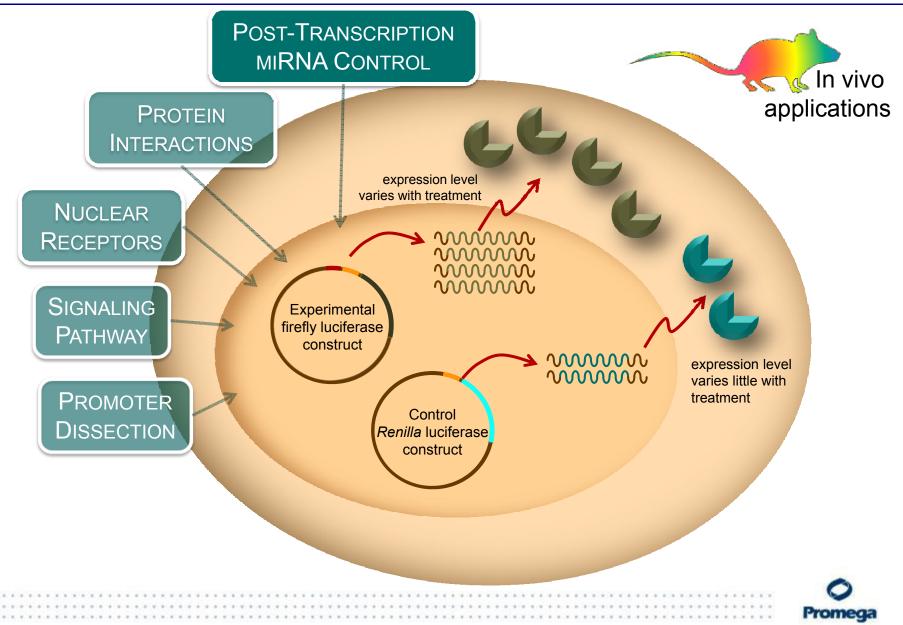


Have a question? Ask a Scientist

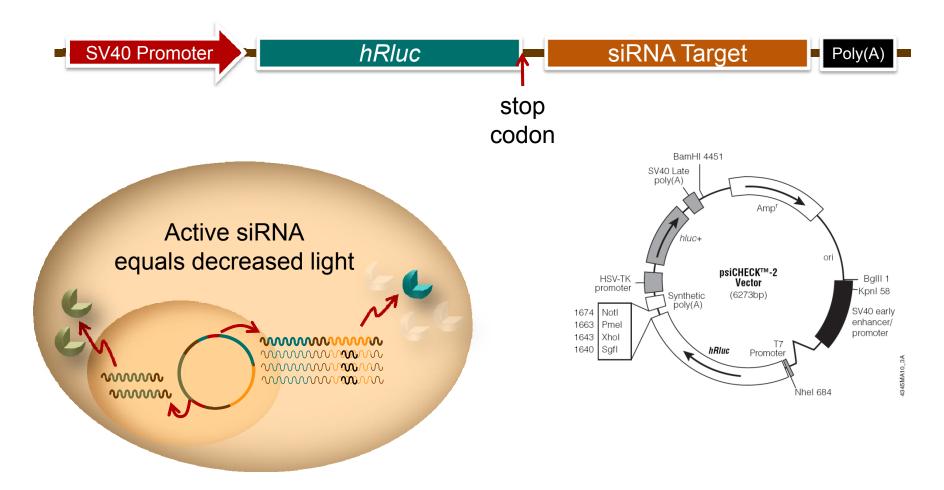
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Application Overview



Using a reporter to find the best siRNA

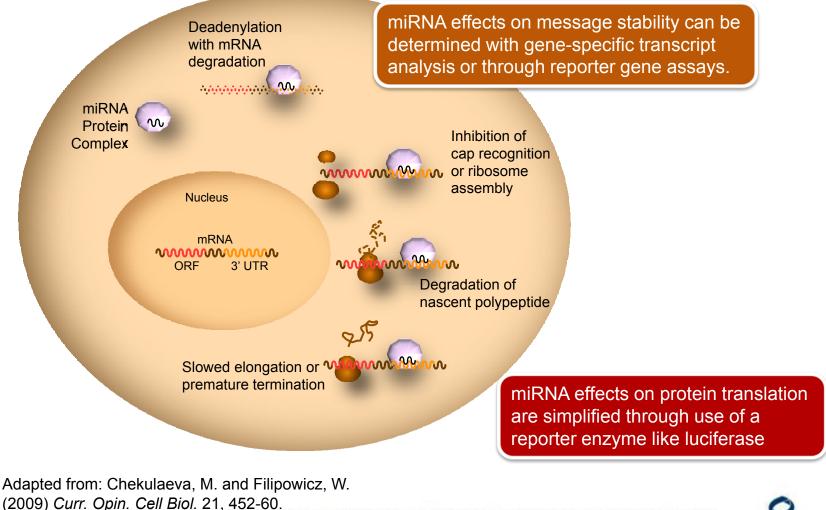


Easy, quantitative assay to find which siRNA to use in knockdown work



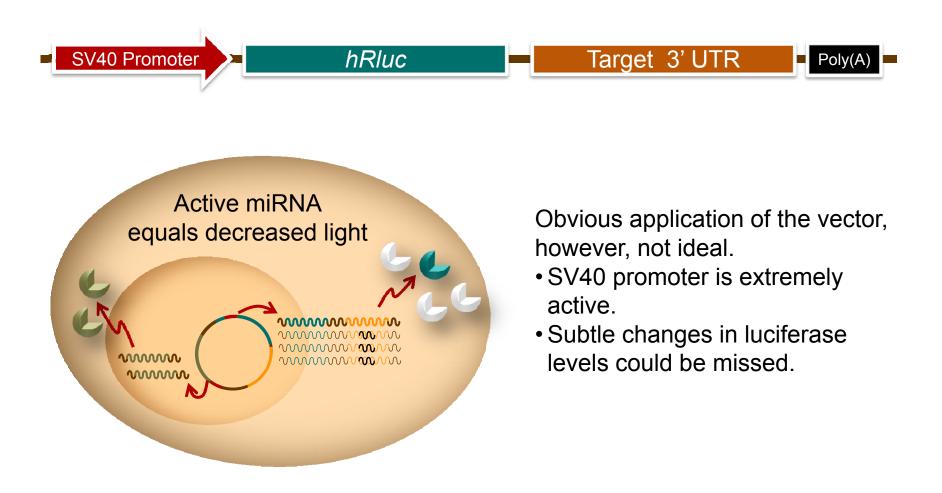
Post-Transcriptional Control through miRNA

Possible mechanisms of miRNA-mediated post-transcriptional control



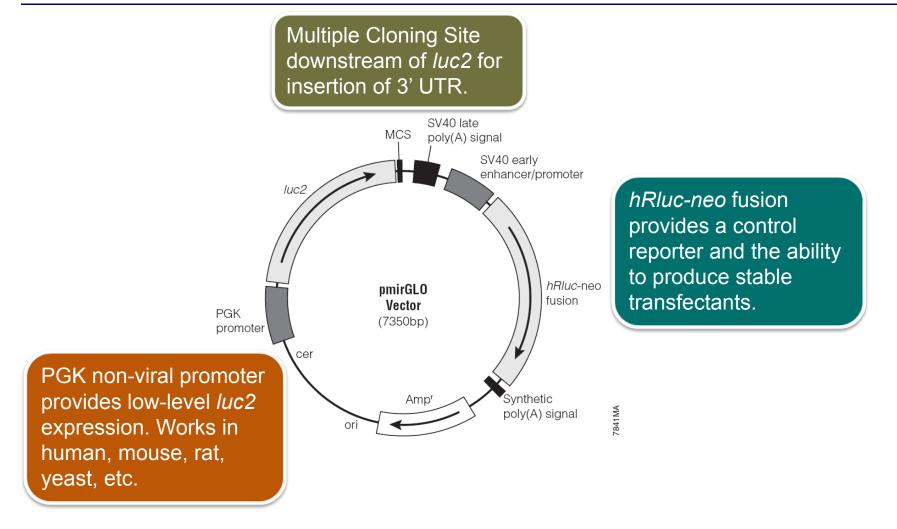


psiCHECK[™]-2 Vector Employed to study miRNA



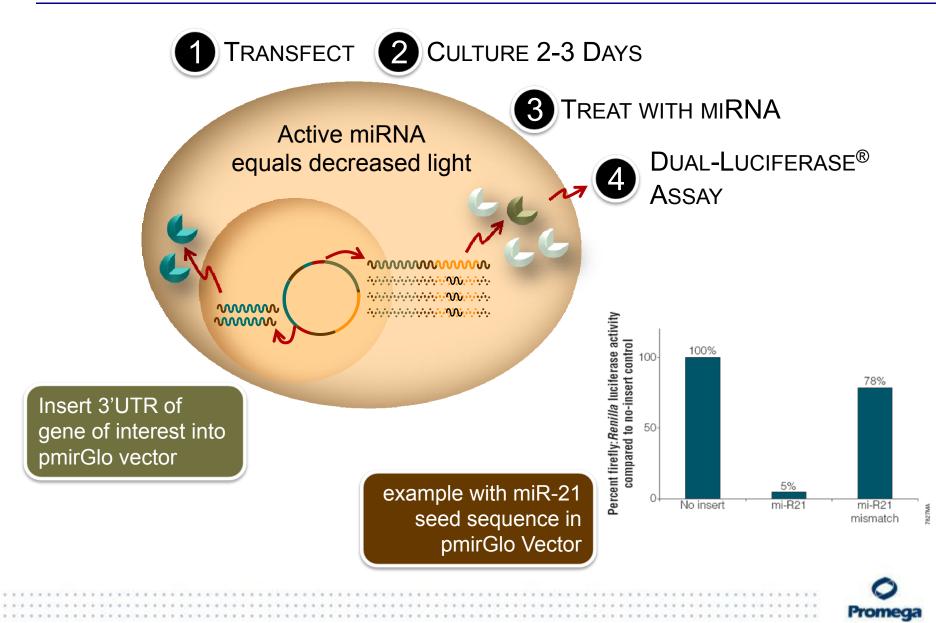


pmirGlo Vector for miRNA targeting





pmirGlo Assay Principle



Case Study: miR-9 and NFKB1 control

Induction and regulatory function of miR-9 in human monocytes and neutrophils exposed to proinflammatory signals

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Inflammation involves a coordinated, sequential, and self limiting sequence of events controlled by positive and negative regulatory mechanisms. Recent studies have shown that microRNAs (miRNAs) an evolutionarily conserved class of endogenous 22-nucleotide noncoding RNAs, contribute to the regulation of inflammation by repressing gene expression at the posttranscriptional level. In this study, we characterize the profile of miRNAs induced by LPS in human polymorphonuclear neutrophils (PMN) and monocytes. In particular, we identify miR-9 as the only miRNA (among 365 analyzed) up-regulated in both cell types after TLR4 activation. miR-9 is also induced by TLR2 and TLR7/8 agonists and by the proinflammatory cytokines TNF-a and IL-1B, but not by IFNy. Among the 3 different genes encoding miR-9 precursors in humans, we show that LPS selectively induces the transcription of miR-9-1 located in the CROC4 locus, in a MyD88- and NF-xB-dependent manner. In PMN and monocytes, LPS regulates NFKB1 at both the transcriptional and posttranscriptional levels, and a conserved miR-9 seed sustained a miR-9-dependent inhibition of the NFKB1 transcript. Overall, these data suggest that TLR4-activated NF-xB rapidly increases the expression of miR-9 that operates a feedback control of the NF-xB-dependent responses by fine tuning the expression of a key member of the NF-KB family.

inflammation | innate immunity | Toll-like receptors | cytokines | NFKB1

The innate immune response is the first line of defense against infectious agents and is mainly exerted by phagocytes, including polymorphonuclear neutrophils (PMN) and monocyte/ macrophages. This response is triggered by the recognition of pathogen-associated molecular patterns of invading microorganisms by members of the Toll/IL-1 receptor (TLR) superfamily (1) among others. These receptors signal through similar intracellular pathways that start with the recruitment to the Toll-IL-1R (TIR) domain present in the receptor tail with 1 of 4 possible TIR domain-containing adaptor molecules. The com-bination of adaptor molecules involved not only depends upon the specific TLR engaged, but also defines the consequent cellular events. In particular, the MyD88 and TIR domaincontaining adapter protein/MyD88 adapter-like protein (TIRAP/MAL) mediates the early NF-xB activation, while the (TIKAP/MAL) incolates the early NP-KB activation, while the TIR domain-containing adapter inducing IFN/BTIR-containing adapter molecule/TIR-containing adapter molecule-2 (TRAM/ TICAM-2) mediate the delayed NF-KB and IFN-regulatory factor (IRF) 3 signals (2, 3). As examples, TLR4 induces proinflammatory cytokines via either a MyD88-dependent rapid activation of the transcription factor NF-B or costimulatory and antiviral proteins through a more delayed activation of both NF-xB and IRF-3 mediated by TRIF. Conversely, TLR3 exclusively signals through the TRIF-dependent pathway and does not activate the MyD88-dependent pathway (2, 3). Importantly, a variety of extracellular and intracellular negative feedback path-

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ways have evolved to prevent an inappropriate inflammatory response following activation of TLRs. These include the regulation of TLR expression, the production of molecules that generation of dominant negative splice variants or postranulational modifications of signal transducers of the TLR signaling cascade (4, 5). An emerging class of regulators of gene expression is repre-

sented by microRNAs (miRNAs), which act at the postrinarscriptional level via an RNA interference mechanism (6, 7), miRNAs biogenesis involves the initial transcription by RNA polymerase II of primary miRNAs (pri-miRNA), which are subsequently cleaved by the RNase III enzyme Drokha and Dicer to the mixen Ext RNA puice strain is the handle into the miRNA-induced silencing complex, where it guides the recognition and translational repression or degraduation of target mRNAs (8). In mammals, a bost of genes are processed to produce over 700 miRNA (miRNA registry at www.sanger.ac.uk) Software/Riaminras), which have been implicated in a wide and differentiation to turnors (6, 7). Recently, acrivation of the inna exposure also has been associated with changes in the expression of selected miRNAs (pamely miR1-146 (9, 10), miR-135 (11, 12), miR-132 (10), and miR-1250 (12). However, the ability of inflammatory figuads to modulate miRNA expression and, more importantly, the role of regulated miRNAs in the to be explored.

Herein, we report the profiles of miRNAs induced by inflammatory stimuli in human PMNs and monocytes and identify miR-9 as a previously unrecognized LPS-responsive miRNA induced in a MyD8-i and NF-4-dependent manner in both cell types. We also show that miR-9 takes part of a regulatory circuitry controlling cell activation by means of inhibitory feedback loop acting at the level of NFRB1, a transcriptional regulator with a key role in the inflammatory response.

Results

To identify miRNAs potentially involved in the responses of peripheral human PMN and monocytes to stimuli of bacterial

Author contributions: F.B., A.M., M.A.C., and M.L. designed research; M.R., D.G., M.M., L.M., and N.T. performed research; M.F. analyzed data; and F.B. and M.L. wrote the paper. The authors declare no conflict of interest.

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This article contains supporting information online at www.pnas.org/cgi/content 0810909106/DCSupplemental,

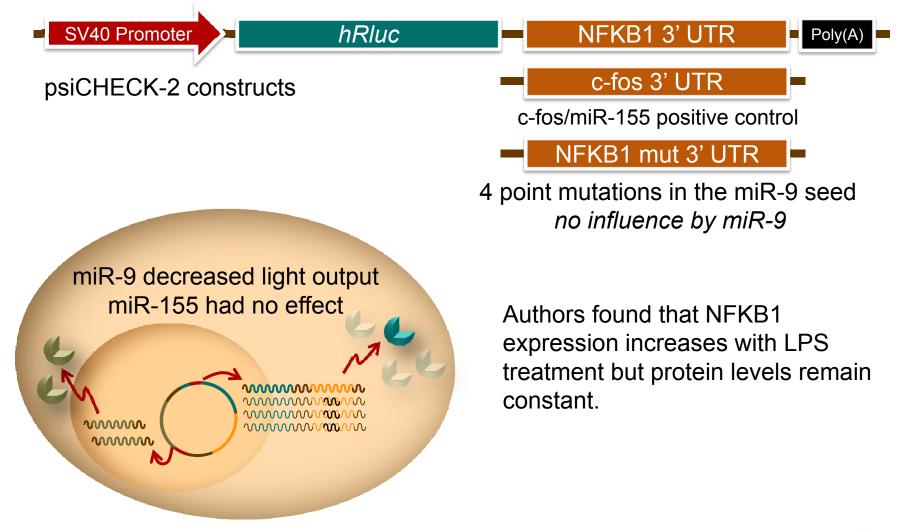
www.pnas.org/cgl/dol/10.1073/pnas.0810909106

- Looking for potential miRNA involvement in the LPS response
- Screen 365 miRNA expression patterns
- Only miR-9 induced in both neutrophils and monocytes in response to LPS
- psiCHECK-2 Vector used in analysis of miR-9 control of NFKB1 through the 3' UTR

Bazzoni, F., et al. (2009) PNAS 106, 5282-7.



Case Study: Bazzoni, F., et al. miR-9 acts on seed in NFKB1 3'UTR

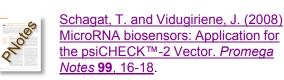




More information



pmirGLO Dual-Luciferase miRNA Target Expression Vector Product Information





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