

# Whole Blood Gene Expression Profiles to Assess Pathogen Load in Infants with Respiratory Syncytial Virus Infection

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Citation Report

#	ARTICLE	IF	CITATIONS
1	Patient-Based Transcriptome-Wide Analysis Identify Interferon and Ubiquitination Pathways as Potential Predictors of Influenza A Disease Severity. PLoS ONE, 2014, 9, e111640.	2.5	19
2	Modeling Human Respiratory Viral Infections in the Cotton Rat ( <i>Sigmodon hispidus</i> ). Journal of Antivirals & Antiretrovirals, 2014, 06, 40-42.	0.1	20
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8	Host RNA signatures for diagnostics: An example from paediatric tuberculosis in Africa. Journal of Infection, 2014, 69, S28-S31.	3.3	16
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