

Whole Blood Gene Expression Profiles to Assess Pathogen 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. <i>PLoS ONE</i> , 2014, 9, e111640.	1.1	19
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4	Detecting specific infections in children through host responses. <i>Current Opinion in Infectious Diseases</i> , 2014, 27, 228-235.	1.3	27
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8	Host RNA signatures for diagnostics: An example from paediatric tuberculosis in Africa. <i>Journal of Infection</i> , 2014, 69, S28-S31.	1.7	16
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