

Rhinoviruses in hospitalized children with acute respiratory infection, Croatia 2017-2019

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Background

Although, human rhinoviruses (HRVs) has been traditionally associated with upper respiratory tract infection (URTI), introduction of molecular assays for respiratory virus detection in clinical laboratories has led to the recognition of HRV as a cause of lower respiratory tract infection (LRTI) as well, especially in vulnerable group of patients such as infants, those with asthma, and immunocompromised patients. Also, there is no data on the prevalence of HRVs in Croatia because there is no routine laboratory diagnosis available in the country. The aim of this study was to determine the prevalence, clinical characteristics and epidemiology of rhinovirus infection in hospitalized Croatian children with acute respiratory infection.

Patients and Methods

A prospective study conducted from March 2017 to February 2019, included 427 children with respiratory symptoms, admitted at Children's hospital Zagreb. Nasopharyngeal swabs were collected in viral transport medium (UTM™, Copan, Italy) and tested for respiratory viruses by multiplex PCR and cDNA synthesis in one-step reaction using Seeplex® RV15 detection kit (Seegene Inc., Seoul, Korea), followed by detection of PCR amplicons using microchip electrophoresis on the MCE®-202 MultiNA device (Shimadzu, Kyoto, Japan). Demographic and clinical illness data were collected by a retrospective review of patient charts.

Results

There were 259 boys and 168 girls. According to the age, the following groups were defined: 0-12 months (n=129), 13-36 months (n=117), 37-60 months (n=51), and >60 months (n=130) of age. According to the localization of infection, patients were categorized as those presented with URTI (n=221), and those with LRTI (n=206). The viral etiology was proved in 320/427; 74.9% of the patients. A single virus was diagnosed in 67.8% of the cases patients, while coinfection with two or more viruses in 15.3% and 4.4% of the positive patients, respectively. The most commonly detected respiratory virus was HRV (Figure 1), diagnosed in 40.5% of all patients; 63.6% as mono-infection, and 36.4% as co-detection with other respiratory viruses. Fifty-one percent of children with rhinovirus mono-infection presented with LRTI. There were no statistically difference in rhinovirus prevalence according to the gender, age, and localization of infection ($P > 0.05$) (Table 1). HRVs were continuously detected throughout whole year with peak incidence in spring and autumn months. positive

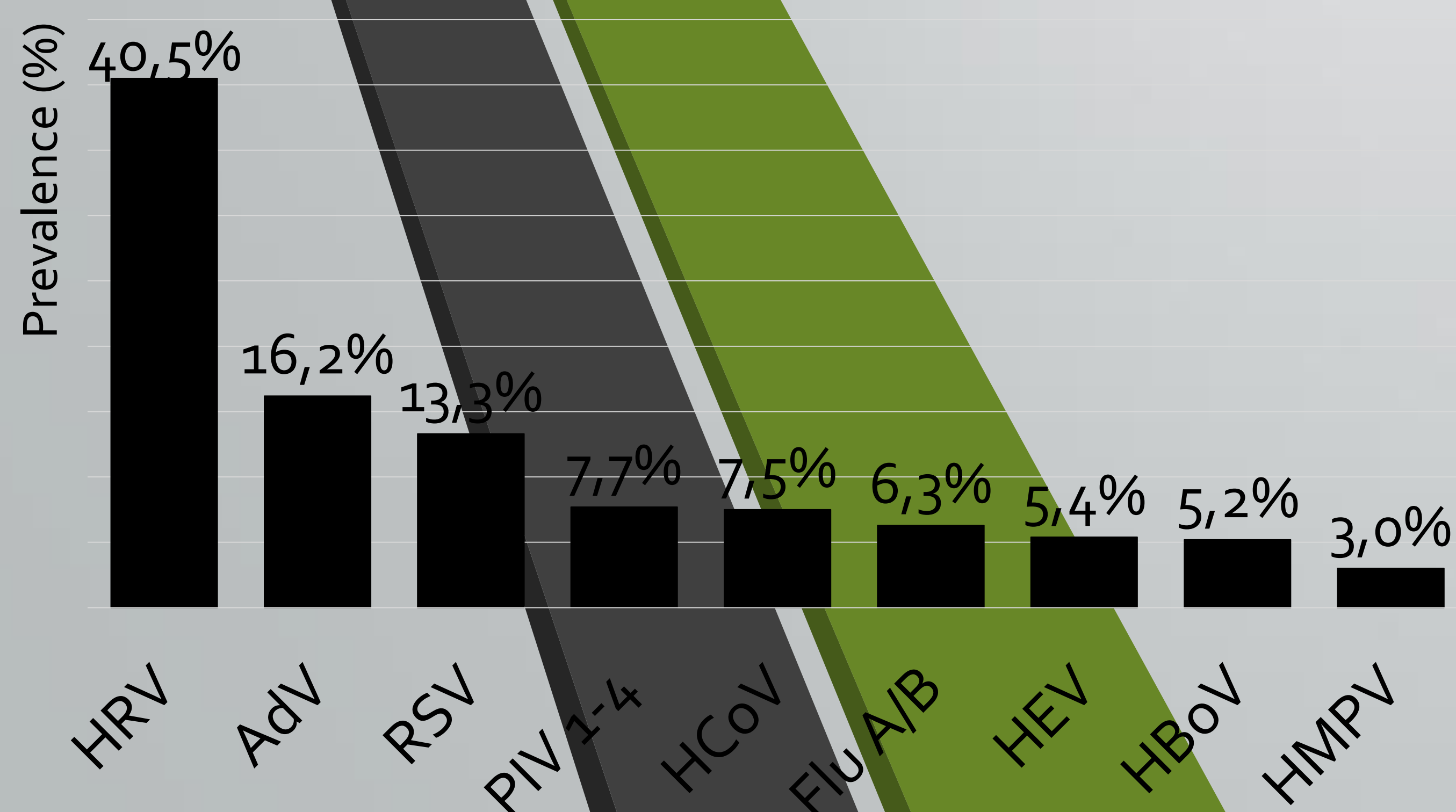
Table 1. Rhinovirus prevalence according to the age, gender and localization of infection

		Rhinovirus positive* / Total N (%)	P value	Rhinovirus mono-infection / Total N (%)	P value
Age (months)	0-12	53/129 (41)	0.8153	31/129 (24)	0.3366
	13-36	51/117 (44)		27/117 (23)	
	37-60	20/51 (39)		11/51 (19)	
	> 60	49/130 (38)		41/130 (32)	
Gender	Male	106/259 (41)	0.8297	64/259 (25)	0.5376
	Female	67/168 (40)		46/168 (27)	
Localization of infection	URTI	78/221 (35)	0.0228	54/221 (24)	0.5161
	LRTI	95/206 (46)		56/206 (27)	

* mono-infection and co-detection with other viruses;

URTI= upper respiratory tract infection; LRTI = lower respiratory tract infection

Figure 1. Frequency of detected respiratory viruses in hospitalized children with acute respiratory infection



HRV = human rhinovirus; AdV = adenovirus; RSV = respiratory syncytial virus; PIV 1-4 = parainfluenza viruses types 1 to 4; HCoV = human coronaviruses 229E/NL63 and OC43; Flu A/B = influenza virus type A and B; HEV = human enterovirus; HBoV = human bocavirus; HMPV = human metapneumovirus

Conclusions

HRVs were the most prevalent respiratory viruses in this study causing significant proportion of LRTIs. These results highlight its role in etiopathogenesis of LRTI not only in infants, but also in children of all ages, and point to the importance of implementing laboratory diagnostics for HRVs in clinical laboratories.

Acknowledgement

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