

**Supplementary material Table S1.** Detection of Ranavirus in common frog (adult and tadpole) at the Balaour pond (Southeaster Alps) in 2016

Tadpole alive		Tadpole dead		Adult alive		Adult dead	
GE	prevalence	GE	prevalence	GE	prevalence	GE	prevalence
3.97	1/5			0	0/5	$2.83 \cdot 10^1$	1/1
0				0			
0				0			
0				0			
0				0			
$1.67 \cdot 10^1$	4/5			1.13	3/5		
$1.33 \cdot 10^2$				$5.98 \cdot 10^{-1}$			
8.64				3.89			
$3.47 \cdot 10^1$				0			
0				0			
2.3	5/5			$1.21 \cdot 10^3$	4/5		
$8.4 \cdot 10^{-1}$				$3.91 \cdot 10^1$			
$1.23 \cdot 10^1$				$4.37 \cdot 10^2$			
1.76				$2.24 \cdot 10^1$			
$1.60 \cdot 10^1$							
$8.4 \cdot 10^{-1}$	4/5	$8.9 \cdot 10^7$	10/10				
$1.23 \cdot 10^1$		$4.48 \cdot 10^6$					
1.76		$1.58 \cdot 10^7$					
$1.60 \cdot 10^1$		$5.38 \cdot 10^6$					
0		$1.94 \cdot 10^7$					
		$3.85 \cdot 10^6$					
		$1.47 \cdot 10^6$					
		$6.43 \cdot 10^4$					

			$3.86 \times 10^5$				
			$5.57 \times 10^6$				
08/15-16	$1.14 \times 10^5$	5/5		0	0/1		
	$2.96 \times 10^3$						
	$1.00 \times 10^4$						
	1.58						
	$9.65 \times 10^2$						
09/19	$4.75 \times 10^4$	1/5		0	0/5		
	0			0			
	0			0			
	0			0			
	0			0			

GE: genomic equivalent, prevalence = ratio number of Rv positive specimen / total number of specimen tested.