

# Pã©ter Putnoky

## List of Publications by Year in descending order

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10  
papers

483  
citations

1163117

8  
h-index

1372567

10  
g-index

10  
all docs

10  
docs citations

10  
times ranked

320  
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of Tail Genes in the Temperate Phage <i>16-3</i> of <i>Sinorhizobium meliloti</i> 41. <i>Journal of Bacteriology</i> , 2010, 192, 1617-1623.	2.2	14
2	pH-dependent regulation of the multi-subunit cation/proton antiporter Pha1 system from <i>Sinorhizobium meliloti</i> . <i>Microbiology (United Kingdom)</i> , 2009, 155, 2750-2756.	1.8	24
3	Genetic Analysis of the <i>rkp-3</i> Gene Region in <i>Sinorhizobium meliloti</i> 41: <i>rkpY</i> Directs Capsular Polysaccharide Synthesis to K <sub>R</sub> 5 Antigen Production. <i>Molecular Plant-Microbe Interactions</i> , 2009, 22, 1422-1430.	2.6	7
4	H Protein of Bacteriophage 16-3 and RkpM Protein of <i>Sinorhizobium meliloti</i> 41 Are Involved in Phage Adsorption. <i>Journal of Bacteriology</i> , 2004, 186, 1591-1597.	2.2	8
5	The <i>rkp-3</i> Gene Region of <i>Sinorhizobium meliloti</i> Rm41 Contains Strain-Specific Genes that Determine K Antigen Structure. <i>Molecular Plant-Microbe Interactions</i> , 2001, 14, 1395-1403.	2.6	41
6	The <i>pha</i> gene cluster of <i>Rhizobium meliloti</i> involved in pH adaptation and symbiosis encodes a novel type of K <sup>+</sup> efflux system. <i>Molecular Microbiology</i> , 1998, 28, 1091-1101.	2.5	115
7	The presence of a novel type of surface polysaccharide in <i>Rhizobium meliloti</i> requires a new fatty acid synthase-like gene cluster involved in symbiotic nodule development. <i>Molecular Microbiology</i> , 1993, 8, 1083-1094.	2.5	76
8	At least two <i>nodD</i> genes are necessary for efficient nodulation of alfalfa by <i>Rhizobium meliloti</i> . <i>Journal of Molecular Biology</i> , 1986, 191, 411-420.	4.2	109
9	A cell-free system from <i>Rhizobium meliloti</i> to study the specific expression of nodulation genes. <i>FEBS Journal</i> , 1986, 160, 69-75.	0.2	8
10	Tn5 carries a streptomycin resistance determinant downstream from the kanamycin resistance gene. <i>Molecular Genetics and Genomics</i> , 1983, 191, 288-294.	2.4	81