

表 5 物种基因组三核苷酸转移概率矩阵间的联合直方图散度

Table 5 The joint histogram divergence of trinucleotide transition probability matrices of bacterial genomes.

	<i>S.aur_JH1</i>	<i>S.epi_AT</i>	<i>S.pyo_M1</i>	<i>S.pyo_M9</i>	<i>S.aga_26</i>	<i>L.lac_MG</i>	<i>P.pen_AT</i>	<i>A.ore_Oh</i>
<i>S.aur_JH1</i>	0							
<i>S.epi_AT</i>	476.659	0						
<i>S.pyo_M1</i>	1 460.841	1 771.400	0					
<i>S.pyo_M9</i>	1 465.781	1 767.048	26.995	0				
<i>S.aga_26</i>	1 062.527	932.606	549.594	541.027	0			
<i>L.lac_MG</i>	1 320.710	1 934.386	1 020.933	1 037.187	1 327.596	0		
<i>P.pen_AT</i>	1 043.516	1 377.373	1 230.305	1 231.175	1 094.646	1 090.625	0	
<i>A.ore_Oh</i>	1 839.600	1 758.309	1 938.365	1 934.658	1 516.067	2 265.351	2 028.414	0