表 3 细菌基因组三核苷酸转移概率矩阵间的联合直方图散度

Table3 The joint histogram divergence of trinucleotide transition probability matrices of bacterial genomes

	L. lac_MG	L. lac_SK	L. lac_I1	S. pyo_M1	P. pen_AT	S. aur_US	A. met_QY
L. lac_MG	0						
L. lac_SK	35.674	0					
L. lac_I1	87.366	89.496	0				
S. pyo_M1	1 020.933	1 043.886	1 202.584	0			
P. pen_AT	1 090.625	1 111.164	1 193.150	1 230.305	0		
S. aur_US	1 291.842	1 311.779	1 324.448	1 475.062	1 062.187	0	
A. met_QY	2 305.814	2 344.976	2 570.574	1 586.869	1 916.826	1 768.208	0