

Table 6. Distribution of orthologs in the chromosome and plasmids of *Rhizobium etli* CFN42 and other Rhizobiales

Species	Chromosome and plasmids			Chromosomes		
	BDBHt	BDBHs	BDBHs, %	BDBHt	BDBHs	BDBHs, %
<i>S_meliloti</i>	3,718	2,667	72	2,836	1,887	67%
<i>A_tumefaciens_C58_Cereon</i>	3,330	2,439	73	2,582	1,545	60%
<i>A_tumefaciens_C58_UWash</i>	3,315	2,414	73	2,559	1,535	60%
<i>M_loti</i>	3,388	2,011	59	2,601	1,590	61%
<i>B_melitensis</i>	2,156	1,396	65	1,860	977	53%
<i>B_suis_1330</i>	2,126	1,327	62	1,831	931	51%
<i>B_abortus_9-941</i>	2,062	1,295	63	1,787	907	51%
<i>B_henselae_Houston-1</i>	1,015	660	65	961	631	66%
<i>B_quintana_Toulouse</i>	949	630	66	903	603	67%
<i>B_japonicum</i>	2,809	1,254	45	2,177	1,039	48%
<i>R_palustris_CGA009</i>	2,201	1,052	48	1,814	925	51%

Orthologs were calculated as BDBHs as indicated in ref. 1. BDBHt indicates the total number of orthologs; BDBHs indicates those orthologs in syntenic arrays, considering two or more consecutive BDBH that match in the *R. etli* genome. Percentage of syntenic BDBH is indicated.

1. Moreno-Hagelsieb, G., & Collado-Vides, J. (2002) *Bioinformatics* **18**, Suppl. 1, S329–S336