

Table 4. Summary of BLAST results for RGP sequences

	Core	Accessory
<i>Acidovorax</i>	4	1
<i>Comamonas</i>	0	0
<i>Delftia</i>	2	0
<i>Bordetella</i>	168	21
<i>Burkholderia</i>	870	142
<i>Cupriavidus</i>	403	245
<i>Chromobacterium</i>	124	42
<i>Nitrosomonas</i>	56	1
<i>Azotobacter</i>	2802	188
<i>P. abietaniphila</i>	7	17
<i>P. entomophila</i>	0	0
<i>P. fluorescens</i>	5994	343
<i>P. mendocina</i>	24	6
<i>P. putida</i>	2944	208
<i>P. syringae</i>	2208	310
Other Pseudomonads	378	49
<i>E. coli</i>	42	14
<i>Shewanella</i>	106	20
γ proteobacterium	0	0
<i>Xanthomonas</i>	136	35
<i>Sinorhizobium</i>	33	8
Bacteriophage	22	68
Eukaryotes	8	1
Other	2558	448
<i>Pseudomonas</i> Total	11555	933
Others Total1	4429	640
Others Total2	2905	594
Others Total	7334	1234
Remaining genes (No Hits or Paralogs)	1195	687
Number of Predicted Paralogs	388	90
Genome Size (ORFs)	23251	23251
Pseudomonas Percent of Total	0.50	0.04
Others Percent	0.32	0.05
CORE or ACCESSORY Size	20084	2854
Pseudomonas Percent of CORE or ACCESSORY	0.58	0.33
Others Percent of CORE or ACCESSORY	0.37	0.43
Corrected Percentages (Counting Remaining as Non-Pseudomonas Hits)		
	<i>Pseudomonas</i> Percentage	57.53
	Others (Non-<i>Pseudomonas</i>) Percentage	64.16