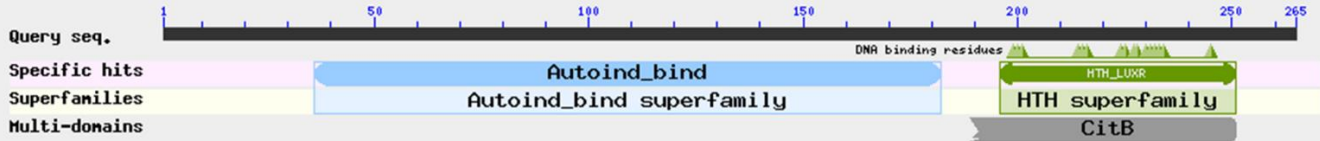


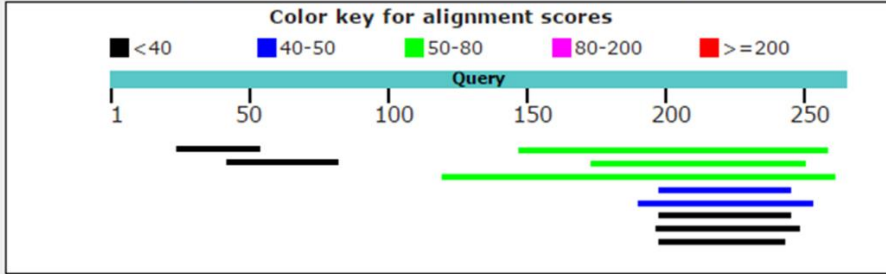
Putative conserved domains have been detected, click on the image below for detailed results.

**Protein Classification:** Autoind\_bind and LuxR\_C\_like domain-containing protein  
 Autoind\_bind and LuxR\_C\_like domain-containing protein



Distribution of the top 10 Blast Hits on 10 subject sequences

Mouse over to see the title, click to show alignments



Name	Locus Tag (PL2TA16_)	Score	Coverage (%)	E value	Identity (%)
PLR1	01227	73.9	41%	2e-16	36%
PLR2	00118	66.2	29%	9e-14	44%
PLR3	03988	56.2	53%	3e-10	30%
PLR4 (PviR)	04751	43.1	18%	9e-06	48%
PLR5	01039	40.4	23%	7e-05	33%
PLR6	04047	38.5	18%	3e-04	40%
PLR7	00814	38.1	19%	4e-04	40%