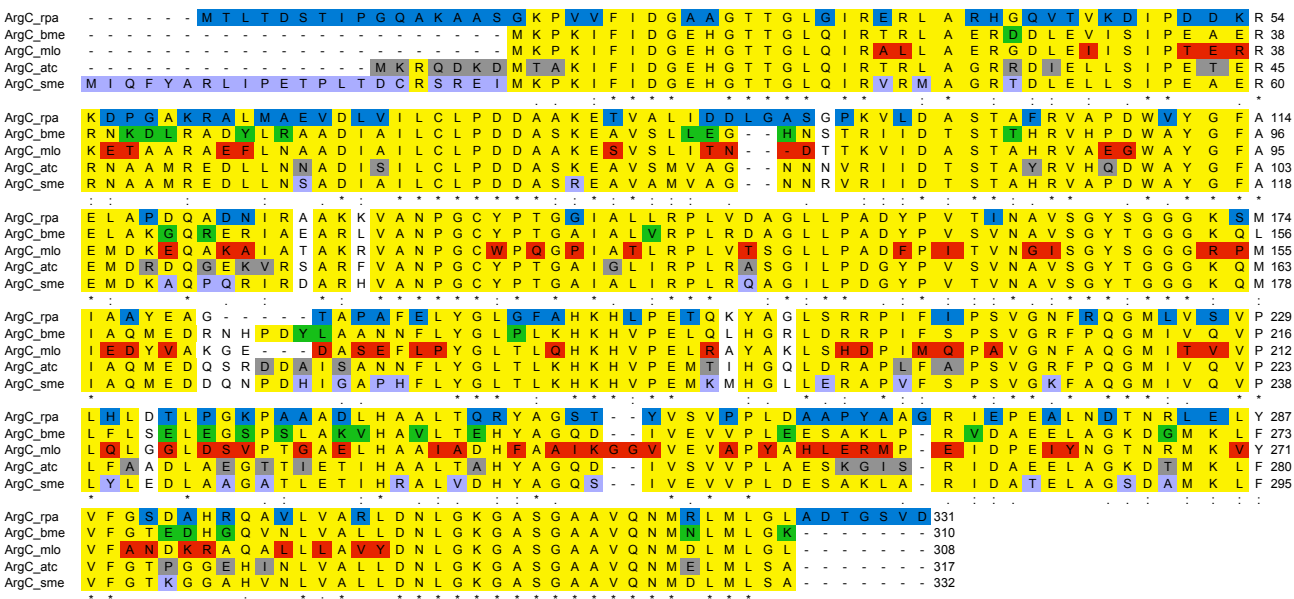




	pl
Rpa	6.1
Bme	5.9
Mlo	5.4
Atu	5.5
Sm	6.0

Identical residues	111
All residues different	9
Changed residues in seq1	113
Changed residues in seq2	33
Changed residues in seq3	83
Changed residues in seq4	41
Changed residues in seq5	48

Identical residues	38
All residues different	0
Changed residues in seq1	99
Changed residues in seq2	35
Changed residues in seq3	79
Changed residues in seq4	42
Changed residues in seq5	54
Changed residues in seq6	113
Changed residues in seq7	0
Changed residues in seq8	49
Changed residues in seq9	16
Changed residues in seq10	0

(c)

*	ArgB		ArgC		ArgD		ArgF		ArgG		ArgH	
Rhizob.	173	62.0	111	33.9	101	24.7	163	50.6	102	23.4	260	56.0
Enterobac.	157	56.3	171	52.3	211	51.6	180	55.9	101	23.2	265	57.1
both	56	20.1	38	11.6	67	16.4	74	23.0	68	15.6	137	29.5

*:	ArgB		ArgC		ArgD		ArgF		ArgG		ArgH	
Rhizob.	226	81.0	170	52.0	191	46.7	211	65.5	191	43.9	326	70.3
Enterobac.	208	74.6	236	72.2	304	74.3	244	75.8	209	48.0	361	77.8
both	108	38.7	76	23.2	143	35.0	130	40.4	141	32.4	238	51.3