

Gblocks 0.91b	
Number of sequences	81
Alignment assumed to be	DNA
New number of positions	1206 (63% of the original 1896 positions)
Parameters used	
Minimum Number Of Sequences For A Conserved Position	41
Minimum Number Of Sequences For A Flanking Position	68
Maximum Number Of Contiguous Nonconserved Positions	8
Minimum Length Of A Block	5
Allowed Gap Positions	All
Flank positions of the 3 selected blocks	Flanks: [154 292] [308 689] [716 1398]

Table S2: Gblocks parameters used in this study