

**Table S2. Positively selected amino acid positions based on sequences of the NLRP1 linker region from 20 primates**

Region	Residue Number (from human NLRP1)	Posterior probability	
		PAML	FUBAR
Linker	105	0.951	
Linker	120	0.985	
Linker	122	0.978	0.978
Linker	139		0.906
Linker	146		0.943
Linker	166	0.975	0.962
Linker	171	0.921	0.910
Linker	187	0.939	0.910
Linker	211	0.997	0.940
Linker	292	0.918	
Linker	293	0.961	0.961
Linker	298	0.975	0.974