

Table 1. Percentage identity of IgM_{1gj} CH domains to conventional (conv) IgM CH domains

	CH1 _{conv}	CH2 _{conv}	CH3 _{conv}	CH4 _{conv}
CH1 _{1gj}	53.8	26.9	25.0	22.4
CH2 _{1gj}	23.7	23.2	54.9	24.2
CH3 _{1gj}	31.8	23.1	30.7	77.9

Each constant domain of IgM_{1gj} was aligned with each constant domain of conventional IgM (Genbank accession number M92851) in CLUSTALW. Highest percentage identity is noted with greatest identity in bold typeface. Note that bolded values closely match the Genbank results in the Fig. 2A domain alignment.