Model	Sequence similarity	$\overline{r_m^2}$	$\sigma_{r_m^{\overline{2}}}$
Lognormal	95%	0.91	0.11
	97%	0.90	0.14
	99%	0.91	0.12
Zipf	95%	0.87	0.080
	97%	0.87	0.11
	99%	0.87	0.076
Log-series	95%	0.78	0.18
	97%	0.78	0.19
	99%	0.78	0.18
Broken-stick	95%	0.043	0.66
	97%	0.034	0.67
	99%	0.012	0.67