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Predicting protein flexibility through the prediction of local structures.

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Table III – *Confusion matrix between normalised B-factors and normalised RMSF values from MD dataset for the three flexibility classes.*

		RMSF _{Norm}			
		1	2	3	Σ
B-factor _{Norm}	1	26.47	11.19	2.79	40.45
	2	11.88	16.03	8.78	36.69
	3	2.10	9.47	11.29	22.87
	Σ	40.45	36.69	22.87	100.00