Linear discriminant analysis reveals hidden patterns in NMR chemical shifts of intrinsically disordered proteins

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S1 Fig. Results from leave-one-out cross-validation for various classification methods using the 17 proteins from BMRB and chemical shifts subset (iii): (H\textsuperscript{N}, N, CO, C\textsubscript{α}, C\textsubscript{β}, H\textsubscript{α} and H\textsubscript{β}).