PhoglyStruct: Prediction of phosphoglycerylated lysine residues using structural properties of amino acids

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Supplement 1: Geometric Means for the different window sizes

	1	Upstream Downstream of lysine														
	±1	±2	±3	±4	±5	±6	±7	±8	±9	±10	±11	±12	±13	±14	±15	
Sensitivity	0.787	0.8442	0.7211	0.7814	0.7959	0.7515	0.8455	0.7208	0.8009	0.7621	0.7159	0.7223	0.7673	0.6926	0.6309	
Specificity	0.7135	0.7148	0.7644	0.751	0.6996	0.7049	0.664	0.7395	0.7441	0.7232	0.7372	0.7554	0.7424	0.7461	0.7644	
Accuracy	0.7302	0.7475	0.7529	0.7565	0.7239	0.7162	0.709	0.7351	0.7589	0.7331	0.7333	0.7461	0.7481	0.7338	0.7335	
MCC	0.446	0.49	0.4463	0.4785	0.447	0.402	0.448	0.414	0.488	0.432	0.406	0.431	0.455	0.399	0.372	
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G-Mean	0.7493	0.7758	0.7424	0.7660	0.7462	0.7278	0.7493	0.7301	0.772	0.7424	0.7265	0.7387	0.7547	0.7189	0.6944	
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