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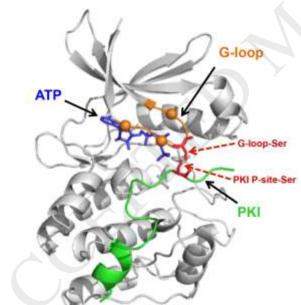
POST-TRANSLATIONAL MODIFICATIONS AT THE ATP-POSITIONING G-LOOP THAT REGULATE PROTEIN KINASE ACTIVITY

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Graphical abstract



G-LOOP Sites that are targets for post-translational modifications

PKA:	GTGSFG
PKC:	GKGSFG
CDK:	GEGTYG
c-Abl:	GGGQYG
Src:	G Q G CFG
B-Raf:	GSGS FG

STRUCTURE OF THE PROTEIN KINASE A CATALYTIC SUBUNIT WITH ATP (blue) AND THE PKI INHIBITOR PROTEIN (green).

The G-loop is highlighted in orange with its phosphorylation site in red sticks. Download English Version:

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