RESEARCH HIGHLIGHTS

Journal club

A NEW PROTEIN KINASE CASCADE

In 1978, when I was just setting up my own laboratory, a short paper from David Gibson's group at the Indiana University School of Medicine, Indianapolis, USA, (Ingebritsen et al.) suggested that a protein kinase that inactivated the cholesterol biosynthetic enzyme hydroxymethylalutaryl-coenzyme A (HMG-CoA) reductase was itself activated by phosphorylation and was therefore part of a kinase cascade. The concept of a kinase cascade is now familiar, but in 1978 only one of these, the protein kinase A (PKA)-phosphorylase kinase cascade, which was discovered by Ed Krebs, was known. More remarkable still, fewer than ten protein kinases (including the kinase for HMG-CoA reductase ('reductase kinase') and an apparently unrelated kinase for acetyl-CoA carboxylase) had been defined and only then in terms of their enzyme activity (the first kinase domain sequence, which was of PKA, was not determined until 1981).



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HMG-CoA reductase was an endoplasmic reticulum (ER) membrane protein, the enzymatic activity of which could only be assessed by a difficult assay that involved thin-layer chromatography. Furthermore, the only way to assay reductase kinase was by its ability to inactivate HMG-CoA reductase in crude membrane preparations. It was therefore remarkable that Gibson's graduate student Tom Ingebritsen was able to work out that 'reductase kinase' could be inactivated by a partially purified protein phosphatase and that the resulting inactive kinase could be reactivated by incubation with Mg·ATP; they correctly surmised that this was due to phosphorylation of the kinase itself. It took my own laboratory another 9 years (Carling et al.) to realize that reductase kinase was the same as the kinase for acetyl-CoA carboxylase (now known as AMPK). It took 25 years for ourselves and others (Hawley et al.; Woods et al.; Shaw et al.) to show that the upstream kinase of reductase kinase that was first observed by Ingebritsen was the

tumour suppressor liver kinase B1

(LKB1; also known as STK11). Since 1990, the original paper from the Gibson laboratory has been cited only 21 times (including 12 times by myself), but it was certainly very influential in my own career.

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ORIGINAL RESEARCH PAPERS Ingebritsen, T. S. et al. Reversible modulation of the activities of both liver microsomal hydroxymethylglutaryl coenzyme A reductase and its inactivating enzyme. Evidence for regulation by phosphorylation-dephosphorylation. Biochem. Biophys, Res. Comm. 81, 1268-1277 (1978) Carling, D., Zammit, V. A. & Hardie, D. G. A common bicyclic protein kinase cascade inactivates the regulatory enzymes of fatty acid and cholesterol biosynthesis. FEBS Lett. 223, 217-222 (1987) Hawley, S. A. et al. Complexes between the LKB1 tumor suppressor, STRADa/B and MO25 α/β are upstream kinases in the AMP-activated protein kinase cascade, I. Biol. 2, 28 (2003) | Woods, A. et al. LKB1 is the upstream kinase in the AMP-activated protein kinase cascade, Curr, Biol. 13, 2004-2008 (2003) | Shaw, R. J. et al. The tumor suppressor LKB1 kinase directly activates AMP-activated kinase and regulates apoptosis in response to energy stress. Proc. Natl Acad. Sci. USA 101, 3329-3335 (2004)