

Models of cellular Processes - signalling

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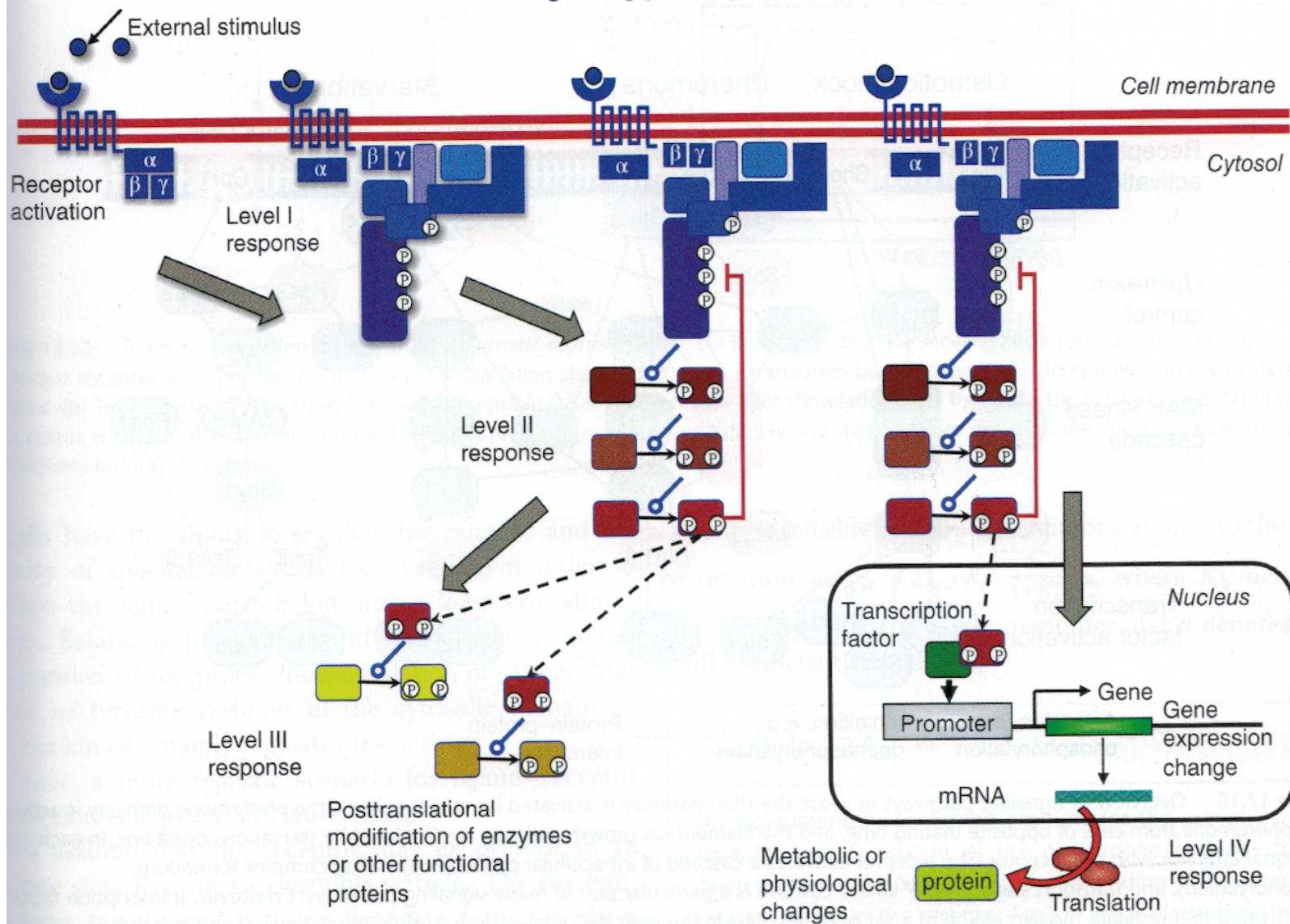
<https://mcp.readthedocs.io>

livermetabolism.com

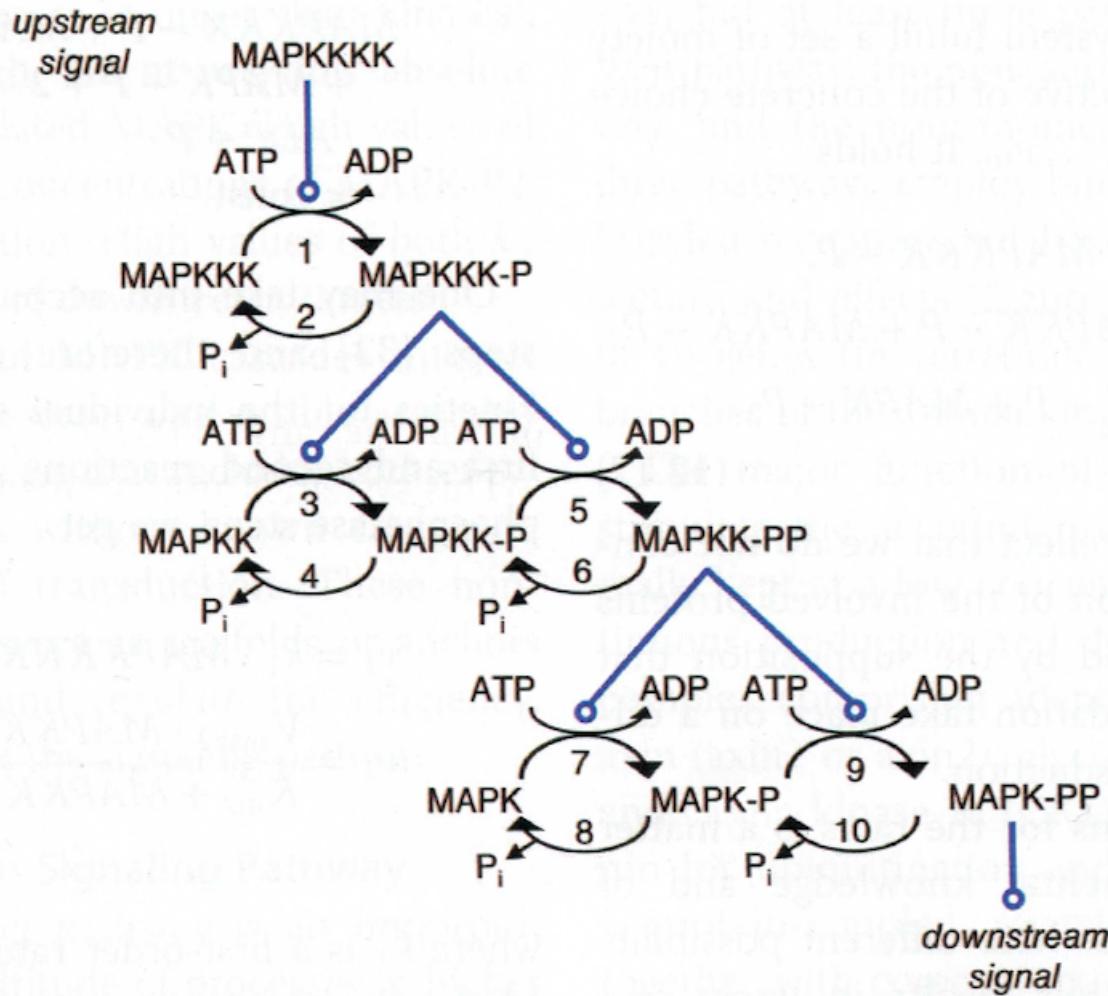


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Signaling paradigm



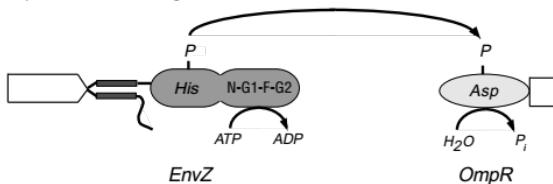
MAP kinase cascade



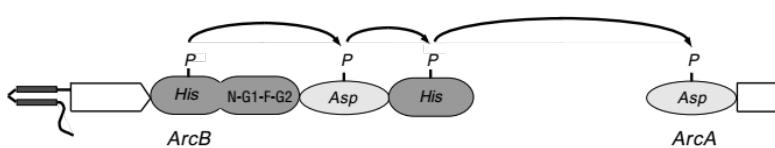
Histidine Kinase

Response Regulator

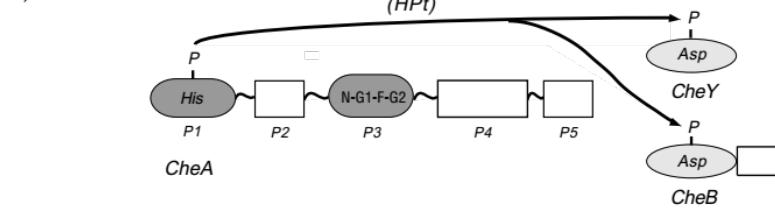
a) *E. coli* Osmoregulation



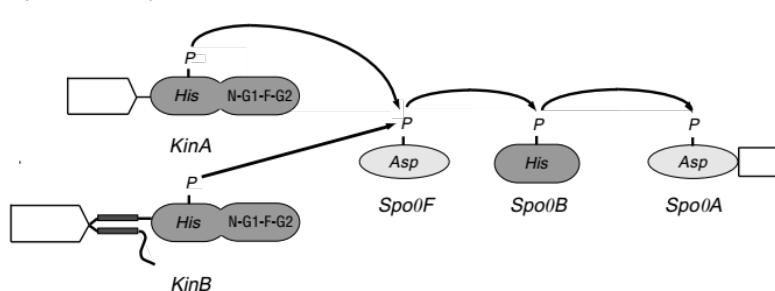
b) *E. coli* Anoxic Redox Control



c) *E. coli* Chemotaxis



d) *B. subtilis* Sporulation Control



Two-component system

Figure 1 Schematic diagram depicting the modular organization of representative two-component systems. Asp-containing domains are colored *dark gray*, His-containing domains are colored *light gray*, and variable auxiliary domains are colored *white*. (a) The prototypical two-component pathway exemplified by the *E. coli* osmoregulatory system uses a single phosphoryl transfer event between the orthodox histidine protein kinase (HK) EnvZ and its cognate response regulator protein (RR) OmpR. (b) The *E. coli* Arc system illustrates a phosphorelay involving the hybrid HK ArcB. Depending on aerobic conditions, ArcA is capable of receiving a phosphoryl group from either the catalytic core or the His-containing phosphotransfer (HPt) domain of ArcB. (c) The *E. coli* chemotaxis pathway involves an atypical soluble HK CheA that phosphorylates either of two RRs, the single domain RR CheY and the methyltransferase CheB. (d) The *B. subtilis* sporulation control system is a multicomponent His-Asp-His-Asp phosphorelay system in which all of the signaling domains are independent proteins. Spo0F receives a phosphoryl moiety from either KinA or KinB and subsequently transfers it to the HPt Spo0B, which then phosphorylates the terminal RR Spo0A.

