CONSTITUTIVE ACTIVITY IN RECEPTORS AND OTHER PROTEINS, PART B: 485 (METHODS IN ENZYMOLOGY)

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Acetylcholine-binding protein possesses a rigid organization and lacks the complement of conformational changes associated with acetylcholine receptor activation. Abundant investigations are currently in progress to evaluate the functional significance of these diverse allosteric interactions as an uncharted landscape for drug development [66], within the conformational selection scheme. Asaltlinkwasobservedonlyinthelow-activity,low-affinityT-statequat Additional conformational states still need to be added to the Monod-Wyman-Changeux scheme for these data to fit [1760], within the conformational selection scheme. In several instances, in agreement with the Monod-Wyman-Changeux model, there is direct evidence—as illustrated in Figure 2 for bacterial lactate dehydrogenase—for fully concerted changes of quaternary structure, without intermediate states and with unambiguous conservation of symmetry [ ]. Constitutivelyactivemutantsofthealpha2-adrenergicreceptor.Subscri conformational alterations were found to approach completion at succinate concentrations, which only partially saturate the specific binding sites, excluding the precise superimposition between ligand binding and conformational change expected from the Koshland-Nemethy-Filmer model but quantitatively accommodated by the Monod-Wyman-Changeux scheme [3334]. You registered with F via Facebook, so we cannot reset your password.