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Antje Chang, Dietmar Schomburg, Ida Schomburg

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systematic and common names are given information on reaction type substrate and product spectrum inhibitors cofactors kinetic data pH and temperature range origin purification molecular data and storage conditions are listed A reference list completes the data sheets This collection is an indispensable source of information for researchers applying enzymes in analysis synthe Enzyme Handbook 4 Dietmar Schomburg, Margit Salzmann, 2013-11-11 Recent progress in enzyme immobilisation enzyme production coenzyme regeneration and enzyme engineering has opened up fascinating new fields for the potential application of enzymes in a large range of different areas As more progress in research and application of enzymes has been made the more apparent has become the lack of an up to date overview of enzyme molecular properties The need for such a data bank was also expressed by the EC task force Biotechnology and Information Therefore we started the development of an enzyme data information system as part of protein design activities at GBF The present book Enzyme Handbook represents the printed version of this data bank ln future it is also planned to make a computer searchable version available The enzymes in the Handbook are arranged according to the 1984 Enzyme Commission list of enzymes and later supplements Same 3000 different en zymes are covered Frequently very different enzymes are included under the same E C number Although we intended to give a representative overview on the molecular variability of each enzyme the Handbook is not a compendium The readerwill have to go to the primary literature for more detailed information Naturally it is not possible to cover all numerous up to 40 000 literature references for each enzyme if data representation is to be concise as is intended **Enzyme Handbook 13** Dietmar Schomburg, Dörte Stephan, 2012-12-06 Today as the large international genome sequence projects are gaining a great amount of public attention and huge sequence data bases are created it be comes more and more obvious that we are very limited in our ability to access functional data for the gene products the proteins in particular for enzymes Those data are inherently very difficult to collect interpret and standardize as they are highly distributed among journals from different fields and are often subject to experimental conditions Nevertheless a systematic collection is essential for our interpretation of the genome information and more so for possible appli cations of that knowledge in the fields of medicine agriculture etc Recent pro gress on enzyme immobilization enzyme production enzyme inhibition coen zyme regeneration and enzyme engineering has opened up fascinating new fields for the potential application of enzymes in a large range of different areas It is the functional profile of an enzyme that enables a biologist of physician to analyze a metabolic pathway and its disturbance it is the substrate specificity of an enzyme which tells an analytical biochemist how to design an assay it is the stability specificity and efficiency of an enzyme which determines its usefulness in the biotechnical transformation of a molecule And the sum of all these data will have to be considered when the designer of artificial biocatalysts has to choose the optimum prototype to start with **Enzyme** Handbook 7 Dietmar Schomburg, Margit Salzmann, Dörte Stephan, 2012-12-06 Recent progress on enzyme immobilisation enzyme production coenzyme regeneration and enzyme engineering has opened up fascinating new fields for the potential

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