

**EVOLUTIONARY APPROACHES TO PROTEIN DESIGN:
55 (ADVANCES IN PROTEIN CHEMISTRY)**

Thomas Lekas

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J Biol Chem This is beyond the scope of this review and will not be addressed in . JAmChemSoc Generally, the consensus approach is based on the hypothesis that consensus amino acids of a sequence alignment contribute more than average to the fitness of the protein than the non-consensus amino acids. Artificial metalloenzymes Artificial metalloenzymes are thought as a bridge between biocatalysis and transition-metal complexes by incorporating the catalytically active transition metal complex in the protein scaffold enabling a high activity and selectivity []. Most de novo enzyme design approaches depend on computational methods, which have been developed and improved in the last two decades [, 6, -]. Janes, Humana Press Comparison of the structures of epoxide hydrolases with esterases revealed differing loops at the entrance to the active site.