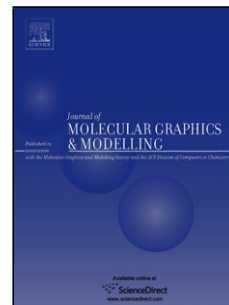


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Accuracy comparison of several common implicit solvent models and their implementations in the context of protein-ligand binding

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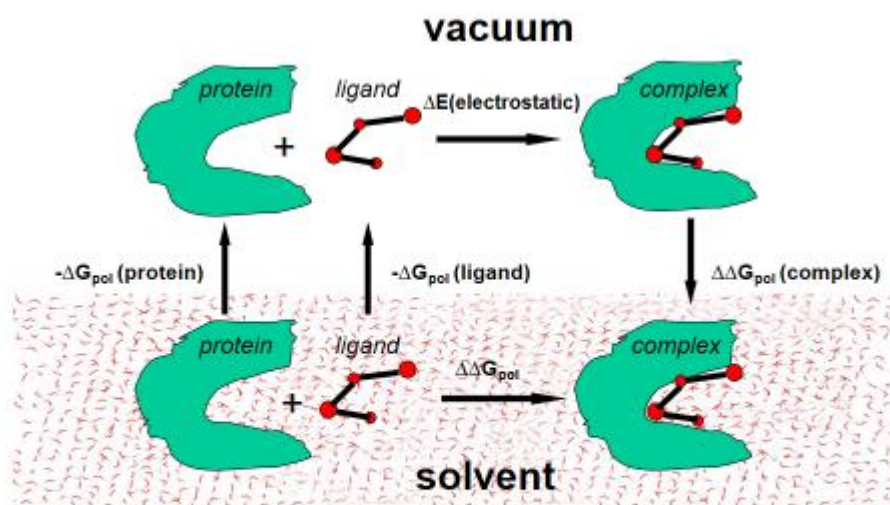
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