



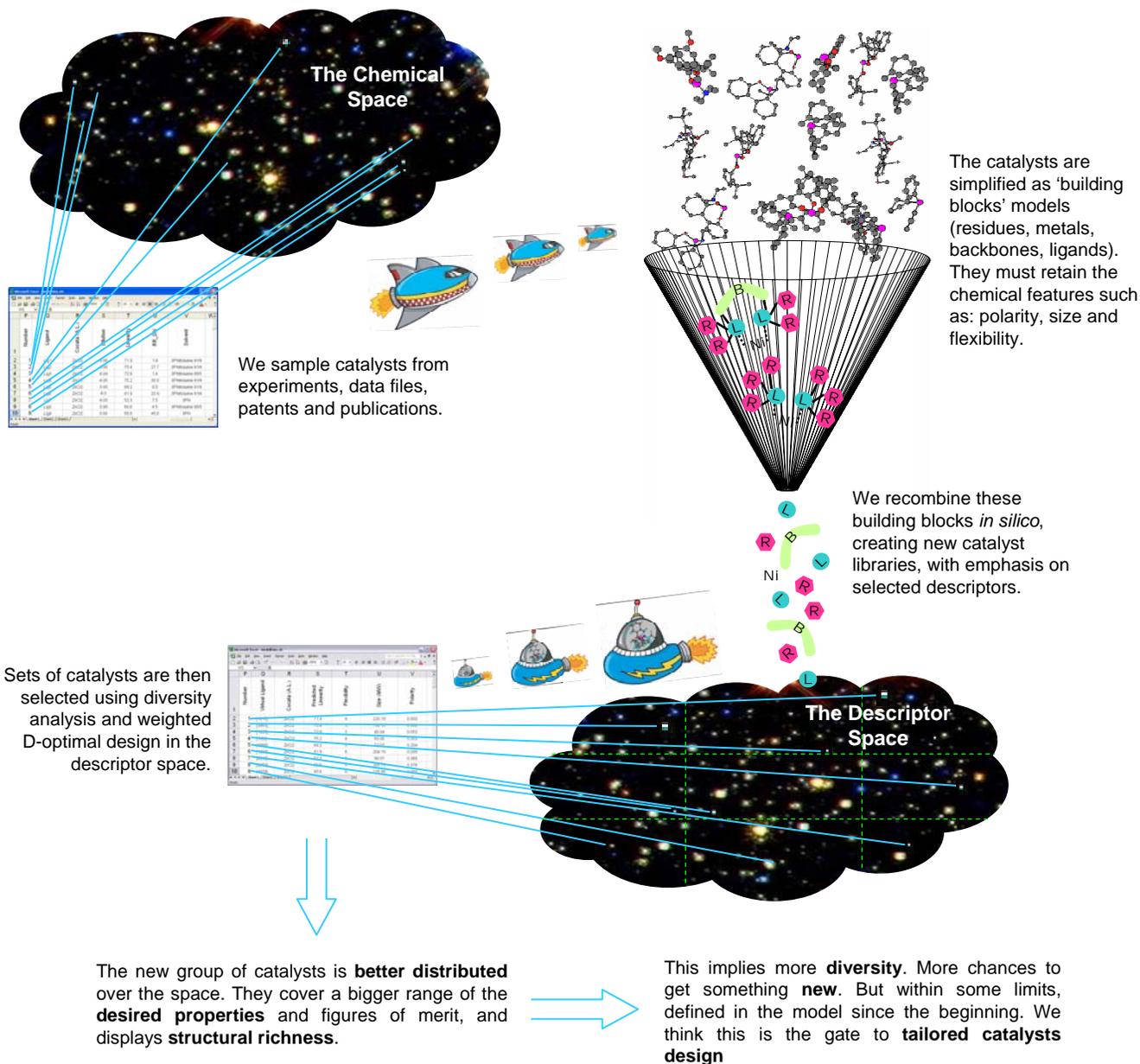
In silico Catalyst Design: Using Backbone Diversity

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The huge diversity of the multi-dimensional ligand space in homogeneous catalysis can be addressed by isolating the main building block motifs creating a representative set of structures of manageable proportions.¹ This set is then modeled using low-cost 2D and 3D descriptors.² Subsets for synthesis are then chosen by maximizing their respective distance in the **descriptor space** using weighted D-optimal design. The D-optimal subset is more diverse than a random sampling.³ This method can be applied to any metal-ligand catalytic system.



1 E. Burello, P. Marion, J.-C. Galland, A. Chamard, G. Rothenberg, *Adv. Synth. Catal.* **2005**, 347, 803.

2 G. Rothenberg, *Catalysis Today*, **2008**, 137, 2.

3 A. G. Maldonado, J. A. Hageman, S. Mastroianni and G. Rothenberg, **2008 Submitted**.



Here we show the application of virtual screening to effectively select diverse catalysts within a tailored descriptor space. Descriptor calculation, *in silico* combinatorial chemistry and weighted D-optimal design are the techniques used to achieve our goal. Currently, we are using these techniques to find better catalysts for the Ni-catalyzed hydrocyanation of butadiene to adiponitrile, a key part of the Nylon 6.6 industrial process.