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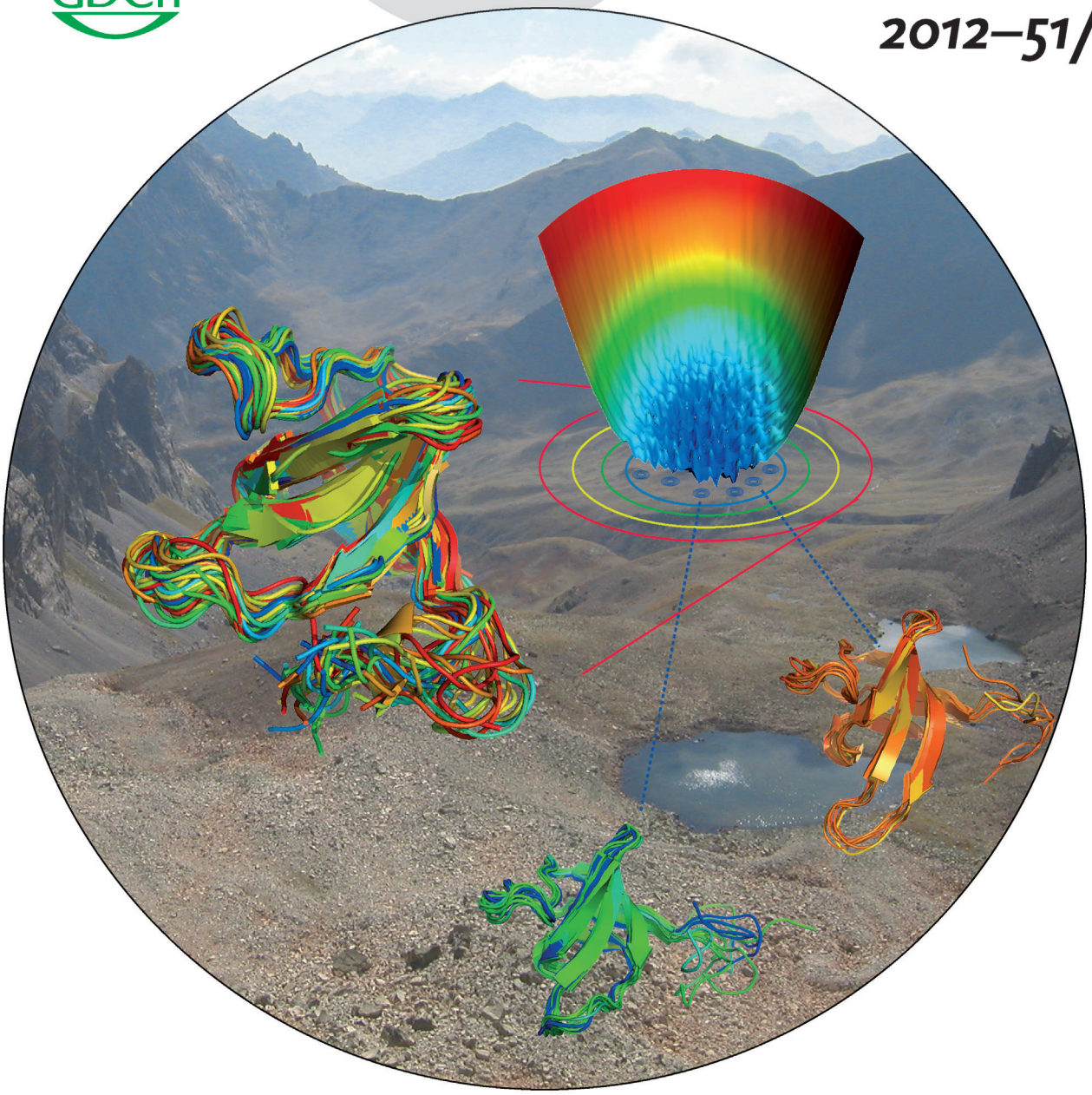

Mapping ...

… the protein conformational energy landscape is possible by using a combination of high-resolution nuclear magnetic resonance spectroscopy and accelerated molecular dynamics simulation. In their Communication on page 6103 ff., M. Blackledge and co-workers provide a detailed description of the different conformational sub-states of the interaction site of the SH3C domain of CD2 associated protein that are sampled on nanosecond to millisecond timescales.

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