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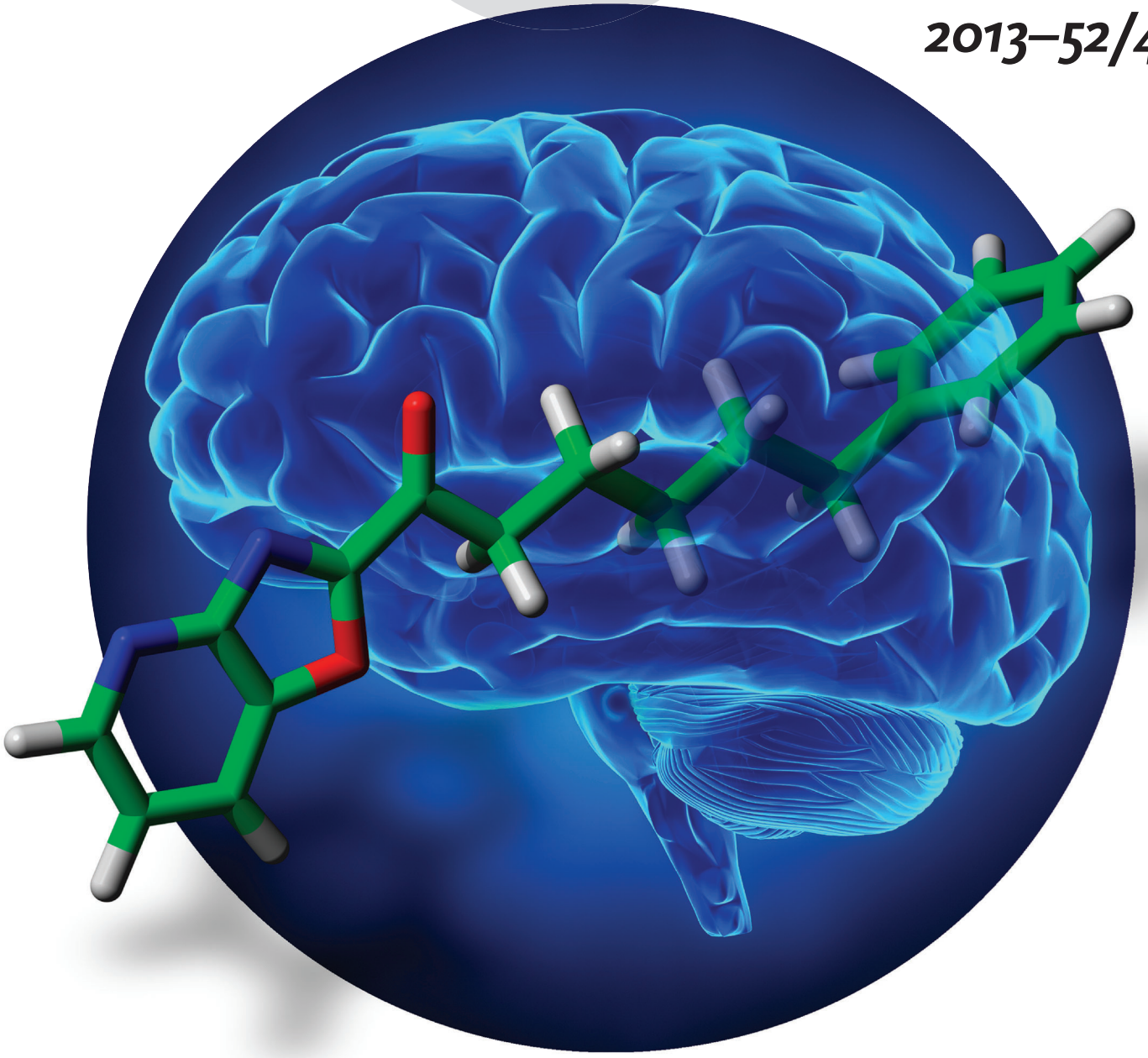

In silico screening ...

... in combination with activity-based protein profiling as a strategy for discovering selective hits in complex brain proteome is presented by M. van der Stelt et al. in their Communication on page 12081 ff. A common starting point was used for the generation of both the pharmacophore model and the activity-based probe. This resulted in the identification of a novel class of diacylglycerol lipase-α inhibitors with high selectivity in the brain proteome.

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