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Subject: Re: TPSA

Posted by [sansun](#) on Tue, 04 Aug 2020 10:20:26 GMT

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I performed calculations after extracting the largest fragments and used the same file for both programs.

However, I find useful information on the following link which says RDkit doesn't use P and S atoms for TPSA calculation.

Does DW uses these atoms?

[https://www.rdkit.org/docs/RDKit\\_Book.html](https://www.rdkit.org/docs/RDKit_Book.html)