



Central European Institute of Technology
BRNO | CZECH REPUBLIC



Validation - trends

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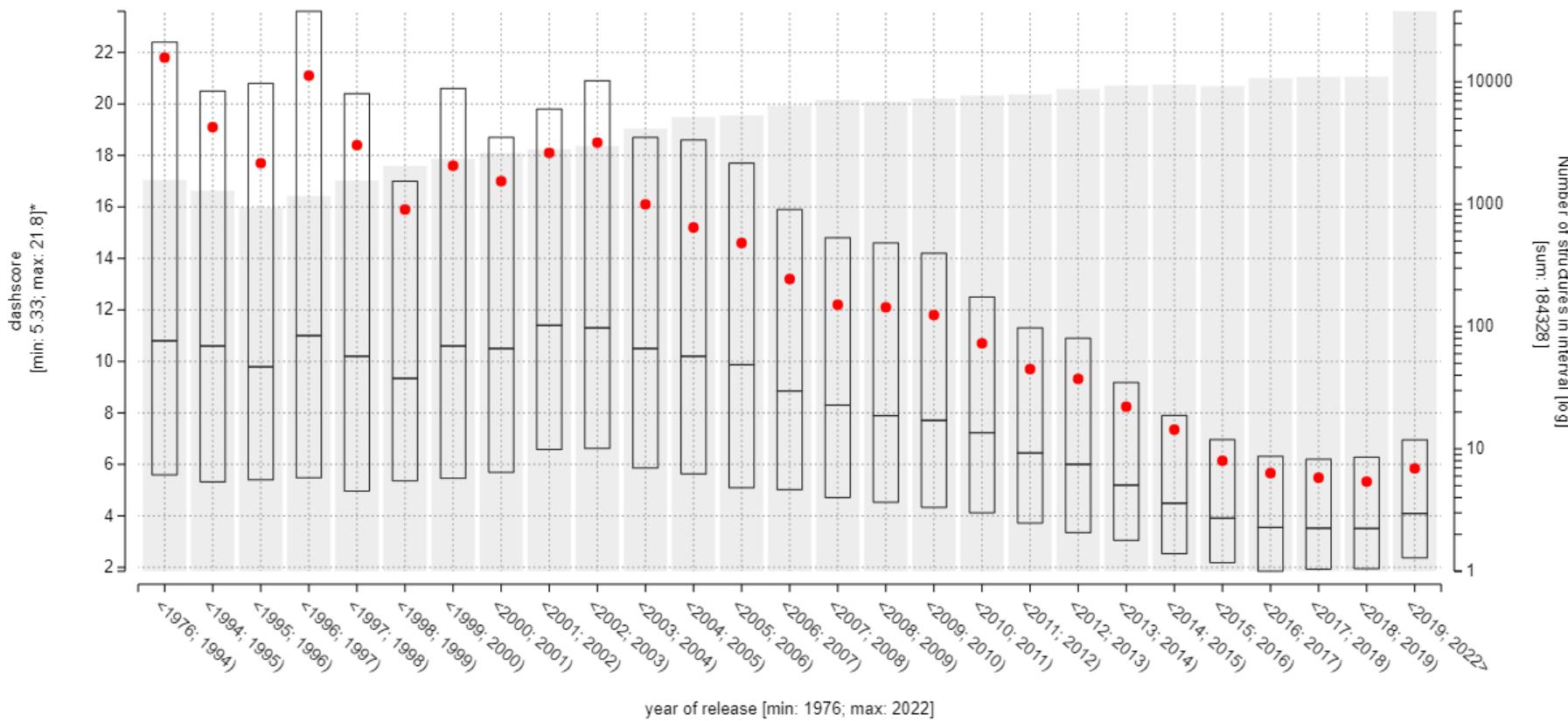


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ValTrendsDB

- Database of biomacromolecular structure validation trends
- Based on structures from Protein Data Bank, automatic update
- <http://ncbr.muni.cz/ValTrendsDB>



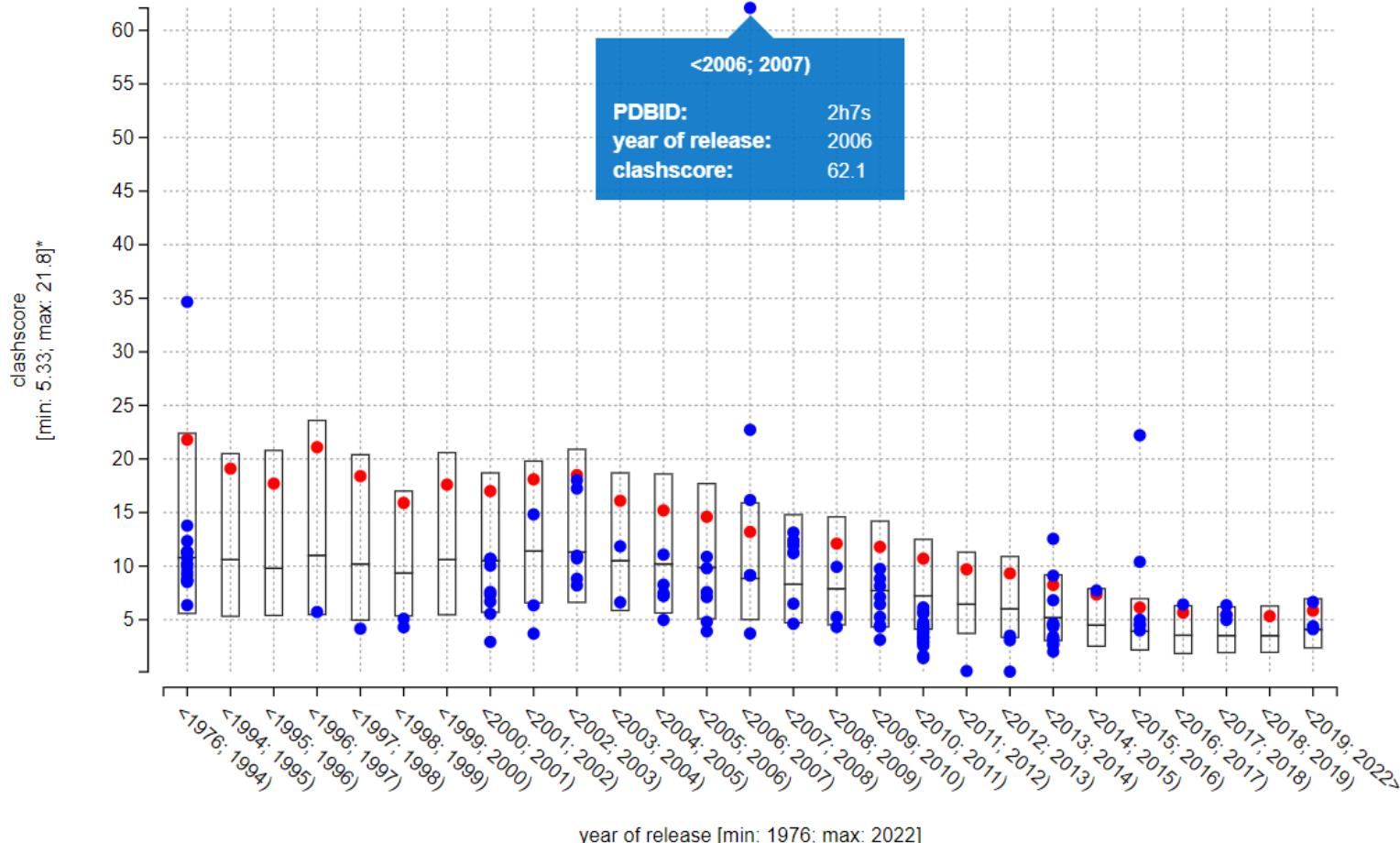
*clashscore structure quality factor

current 2022-01-20

Horský V., Bendová V., Toušek D., Koča J. Svobodová R., 2019. *ValTrendsDB: bringing Protein Data Bank validation information closer to the user.* **Bioinformatics**, 35(24), 5389-5390.

ValTrendsDB

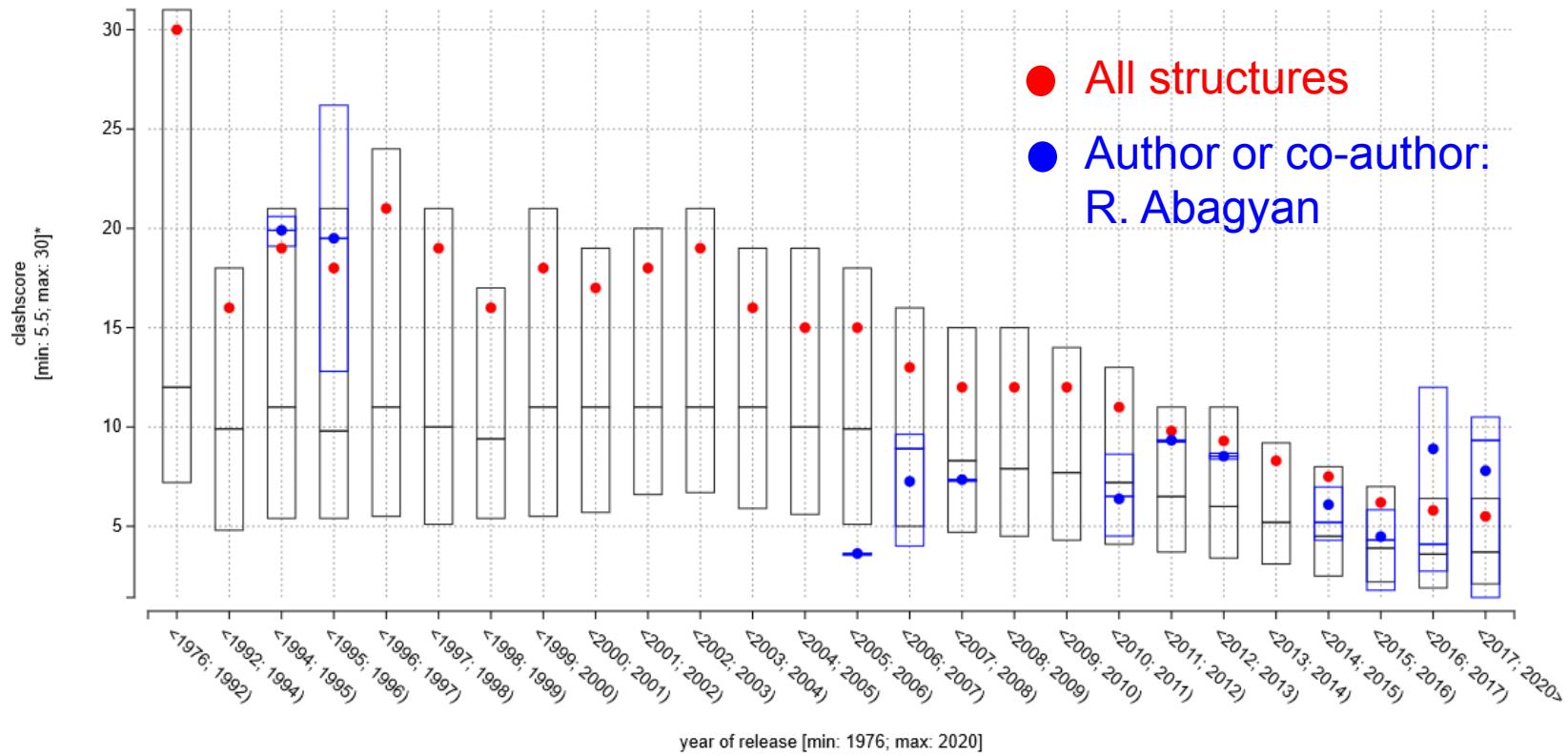
- Validation of cytochromes P450 – overview of the whole family



Blue points are structures of cytochromes P450-cam
Red points show a common trend

Other projects: ValTrendsDB

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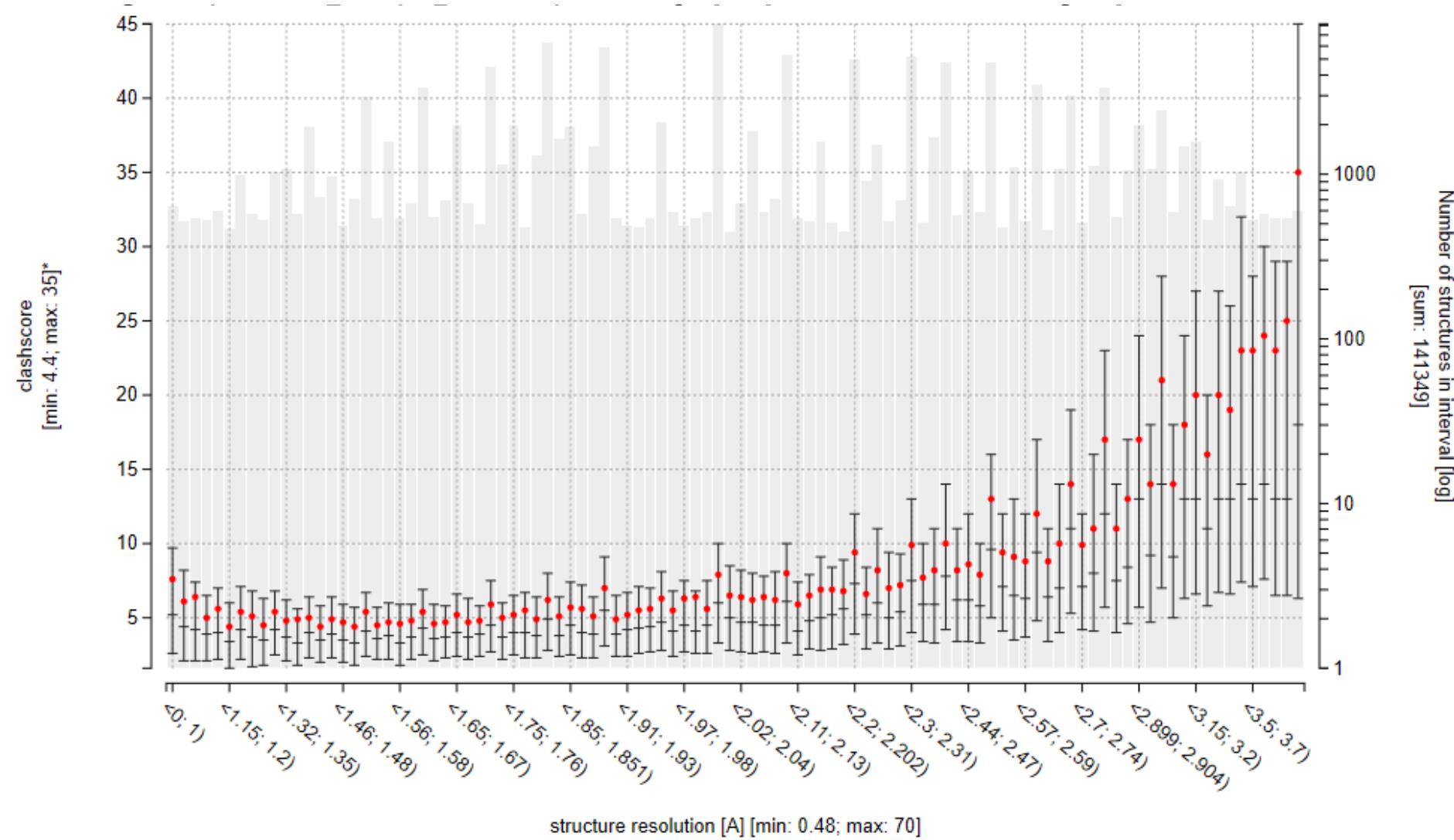


*clashscore structure quality factor

current 2020-01-02

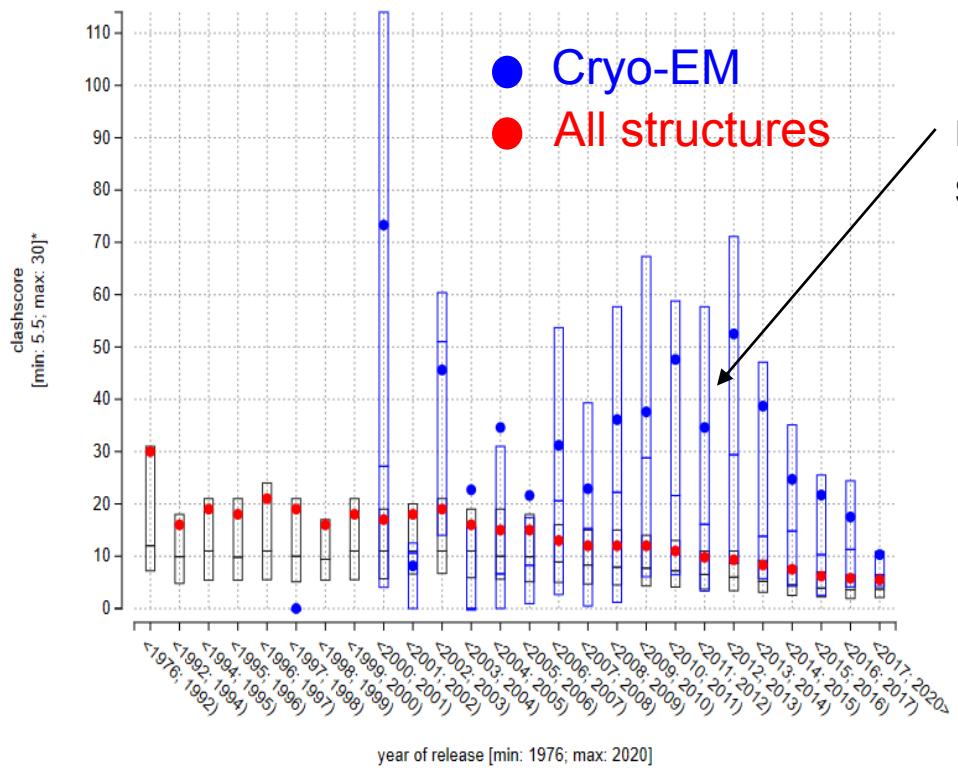
ValTrendsDB

Protein quality and resolution :-)



ValTrendsDB

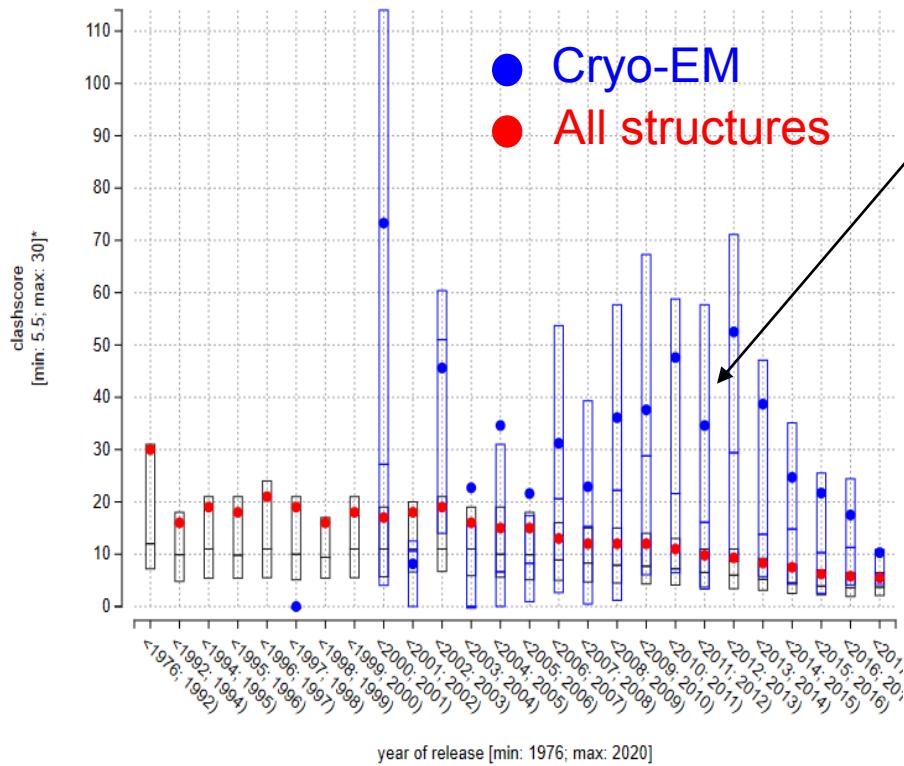
Example of analysis



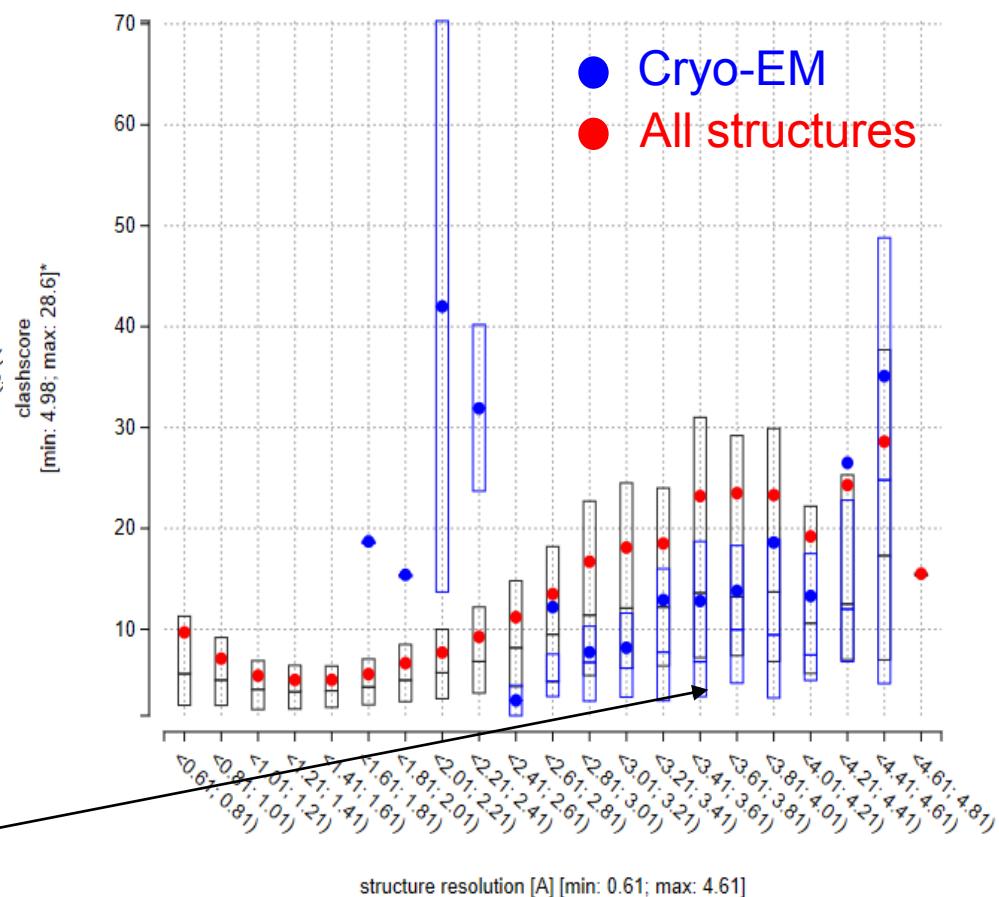
It looks, Cryo-EM structures have much lower quality than common structures

ValTrendsDB

Example of analysis



It looks, Cryo-EM structures have much lower quality than common structures



But when we take into account a resolution, Cryo-EM structures have (for resolution > 2.4 Å) higher quality