

Accepted Manuscript

Title: Investigation of the thermal shift assay and its power to predict protein and virus stabilizing conditions

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PII: S0731-7085(18)30786-6
DOI: <https://doi.org/10.1016/j.jpba.2018.08.017>
Reference: PBA 12148

To appear in: *Journal of Pharmaceutical and Biomedical Analysis*

Received date: 4-4-2018
Revised date: 30-7-2018
Accepted date: 7-8-2018

Please cite this article as: Sviben D, Bertoša B, Hloušek-Kasun A, Forcic D, Halassy B, Brgles M, Investigation of the thermal shift assay and its power to predict protein and virus stabilizing conditions, *Journal of Pharmaceutical and Biomedical Analysis* (2018), <https://doi.org/10.1016/j.jpba.2018.08.017>

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**Investigation of the thermal shift assay and its power to predict protein and virus
stabilizing conditions**

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Highlights

- Similar effect for investigated solutions on T_m was found with different proteins
- Low pH destabilizes proteins probably due to differences in hydrogen bonding
- Impact of solution on T_m shift does not correlate with protein aggregation tendency
- Impact of excipients on virus stability cannot be predicted from protein T_m shifts

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