



Beta Cyclodextrin Derivatives as Protein Aggregation Modulators

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INTRODUCTION

Protein aggregation is the major challenge encountered during manufacturing, storage and transportation of biopharmaceuticals (1,2). Our objective was to evaluate the effect of two ßcyclodextrins derivatives: (KLEPTOSE® HPB hydroxypropyl-ß-cyclodextrin, with MS=0.65) and (KLEPTOSE® HP hydroxypropyl-ß-cyclodextrin, with MS=0.9) on two biologic drugs (Infliximab and Etanercept) aggregation using high-throughput formulation screening (*i*Formulate[™]) and nanoDSF (Differential Scanning Fluorimetry) (3,4).

MATERIALS & METHODS

Infliximab (Remicade[®]), a tumor necrosis factor

(TNF-alpha) binding antibody (chimeric IgG1) and Etanercept (Enbrel[®]), a dimeric fusion protein were purchased (in PBS: phosphate saline buffer) from BOC Sciences. Simultaneous evaluation of Tm (melting temperature) and relative degree of aggregation of proteins in various molarity of KLEPTOSE[®] HPB and HP (0 mM, 0.125 mM, 0.25 mM, 1.25 mM, 2.5 mM, 5 mM, 10 mM, 25 mM, 50 mM, 100



RESULTS & DISCUSSION

Thermal and colloidal stability of 4 mg/mL Infliximab and 6mg/mL Etanercept in *i*Formulate[®] RS-2 plate are affected (Fig. 2A and Fig. 2B) by different formulation variables (pH, ionic strength, stabilizer conc., buffer conc.). Infliximab is much more prone to aggregation than Etanercept.



Fig. 2A : DSF and Aggregation profile of Infliximab and Etanercept in /Formulate™ (25 formulations tested simultaneously for thermal and colloidal stability).

Fig. 2B : /Formulate™ on thermal stability of Etanercept (Enbrel*)

Pareto and DoE analysis of Etanercept in *i*Formulate[®] RS-2 plate formulations illustrate that pH dictates its thermal stability.

Effect of Tm-1 on *i*Formulate Formulations of Etanercept





Fig. 4 : Infliximab (Remicade[®]) colloidal stability by Pareto and DOE analysis of *i*FORMULATE[™] formulations

CONCLUSION

Preliminary results demonstrate that KLEPTOSE[®] HPB BioPharma hydroxypropyl-ß-cyclodextrin and KLEPTOSE[®] HP BioPharma hydroxypropyl-ß-cyclodextrin at high molarity (200 mM) are efficient tools in modulating Infliximab relative degree of aggregation.

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