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# Polystyrene Surface-Absorbed Trehalose Diester as a Means for Performing Pulldown Assays

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## Background

In 2016, over 10.4 million people contracted Tuberculosis (TB) infections for the first time, resulting in the deaths of 1.7 million people worldwide.<sup>1</sup> Despite the existence of a vaccine, TB remains the number one cause of death from a single infectious disease according to the 2017 report from the World Health Organization. As a result, research and development of a new TB vaccine is ongoing, as it is suspected the currently available formulation does not contain an adjuvant which produces the necessary TH-17 response. In the search for new drugs to combat this epidemic, knowledge of how a compound triggers an immune response is paramount. One method for identifying an activated immune protein complex is via immunoprecipitation, colloquially referred to as a "pulldown."

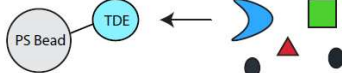
In this study, a pelletable, beaded support of polystyrene (PS) with surface-absorbed trehalose diester (TDE) has been developed for use as a tool for identification of C-type lectin receptors. TDE was used as a stand-in for trehalose dimycolate, a glycolipid found in the cell wall of *Mycobacterium tuberculosis*, which is recognized by C-type lectin. Confirmation of surface-absorption and characterization of the modified beads was performed via dynamic light scattering and high-performance liquid chromatography.

## Pulldown Assay Process

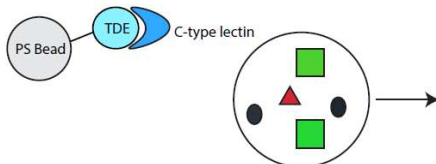
Step 1: Attach TDE to PS-Bead



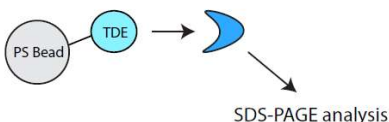
Step 2: Immobilize C-type lectin from cell lysate



Step 3: Spin down beads and wash away unbound species



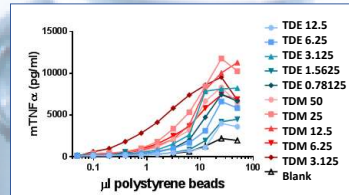
Step 4: Elute protein complex and identify via SDS-PAGE analysis



Pulldown method developed from Kercher B, et al.(2016)<sup>2</sup>

## PS-Bead Characterization

PS-beads were resuspended in a mannitol solution prior to lyophilization in order to provide maximum surface area for TDE absorption to occur evenly over the surface of the beads.



### Dynamic Light Scattering Size Analysis

Sample	Z-Avg (nm)	PDI	Peak 1 Size (nm)	Peak 1 Vol. %	Peak 2 Size (nm)	Peak 2 Vol. %
Lot B4.1	114.8	0.416	144.6	1.2%	7.843	98.8%
Lot B4.2	119.5	0.32	135.4	33.3%	9.706	65.3%

### RP-HPLC Concentration Analysis

Sample	Initial Loading Conc. μg/mL	Wash Assay Conc. μg/mL	Absorbed TDE Conc. μg/mL
Lot B4.1	200	9.3624	190.6376
Lot B4.2	200	13.2954	186.7046

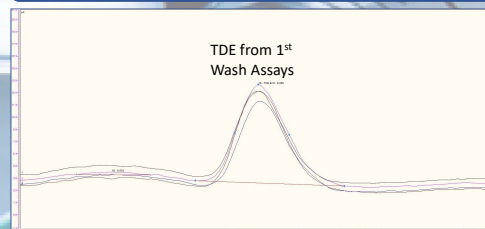
### Target characteristics

- Modified beads must be large enough to quickly pellet via centrifugation
- The surface of the beads must be uniformly coated by TDE
- The spreading density of TDE on the PS-beads must produce significant C-type lectin receptor binding

### Results

- 5 micron beads pelleted quickly with only a small amount remaining suspended after 3 minutes of centrifugation or sitting undisturbed for less than one hour.
- A spreading density of 3.125 Å<sup>2</sup>/molecule (15.41 μg/cm<sup>2</sup>) produced the greatest receptor binding in RAW cells (labeled **TDE 3.125, top right**)
- Method for theoretically uniform surface-absorption of TDE developed (**picture top left**)
- TDE did not remain absorbed to bead surface throughout pulldown assay procedure

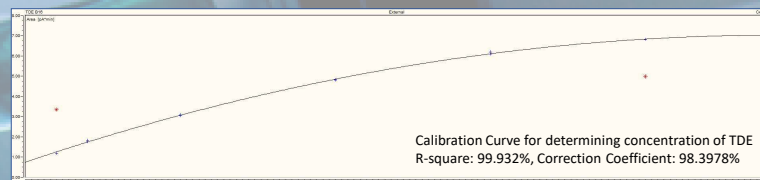
## Reverse Phase HPLC



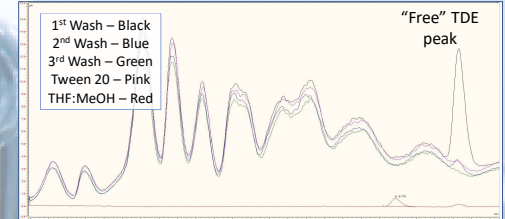
Time (min)	%A	%B	Flow (mL/min)
0	25	75	0.7
1	25	75	
4	0	100	
10	0	100	
11	25	75	
12	25	75	

A: 0.1% TFA  
B: 0.1% TFA, in methanol

Actual concentrations of TDE absorbed to PS-beads were determined by indirect analysis of post-absorption washes via RP-HPLC. HPLC method information, including mobile phase composition and flow rate is given in the Table (**above right**). This experiment provided consistent and clean data to determine concentration of free TDE remaining in samples of the modified PS-bead after three washes. Resulting concentrations were subtracted from the initial loading concentration to determine the final concentrations of TDE absorbed to the PS-beads (**above, under characterization header**). Concentrations were calculated based upon an 8-point calibration curve with a R-square of 99.932% (**shown below**).



## Reverse Phase HPLC



This is a chromatogram of all 3 wash assays from Lot B4.1 with the 0.1% Tween 20 vehicle and 9:1 THF:MeOH (**above**). TDE has completely disappeared by the third wash (**green**) although there is some carryover of TDE shown in the Tween and THF:MeOH samples, none was present in the final wash of the beads. All "free" TDE has successfully been removed from the suspension.

## Conclusions & Future Directions

### Conclusions

- TDE was successfully surface-absorbed to polystyrene beads at concentrations close to target
- Any TDE not absorbed was removed during post-absorption washes
- Resulting beads aggregate and pellet with short spin times or by sitting undisturbed for less than one hour.
- TDE did not remain absorbed to bead surface throughout pulldown assay procedure

### Future Directions

- Develop further to prevent removal of TDE from PS-beads during pulldown: what causes the TDE to desorb from the PS-beads and how can this barrier be overcome?
- Calculate surface-absorbed TDE concentration directly by removing TDE from modified beads and analyzing via RP-HPLC

## References

1. World Health Organization. (2017). *Global Tuberculosis Report 2017*. Retrieved from <http://www.who.int/tb/publications/factsheets/en/>
2. Kerscher, B., Dambuzza, I. M., Christofi, M., Reid, D. M., Yamasaki, S., Willment, J. A., & Brown, G. D. (2016). Signalling through MyD88 drives surface expression of the mycobacterial receptors MCL (Clec4e8, Clec4d) and Mincle (Clec4e) following microbial stimulation. *Microbes and infection*, 18(7-8), 505-509.
3. Hunter, R. L., Olsen, M. R., Jagannath, C., & Actor, J. K. (2006). Multiple roles of cord factor in the pathogenesis of primary, secondary, and cavity tuberculosis, including a revised description of the pathology of secondary disease. *Annals of Clinical & Laboratory Science*, 36(4), 371-386.

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