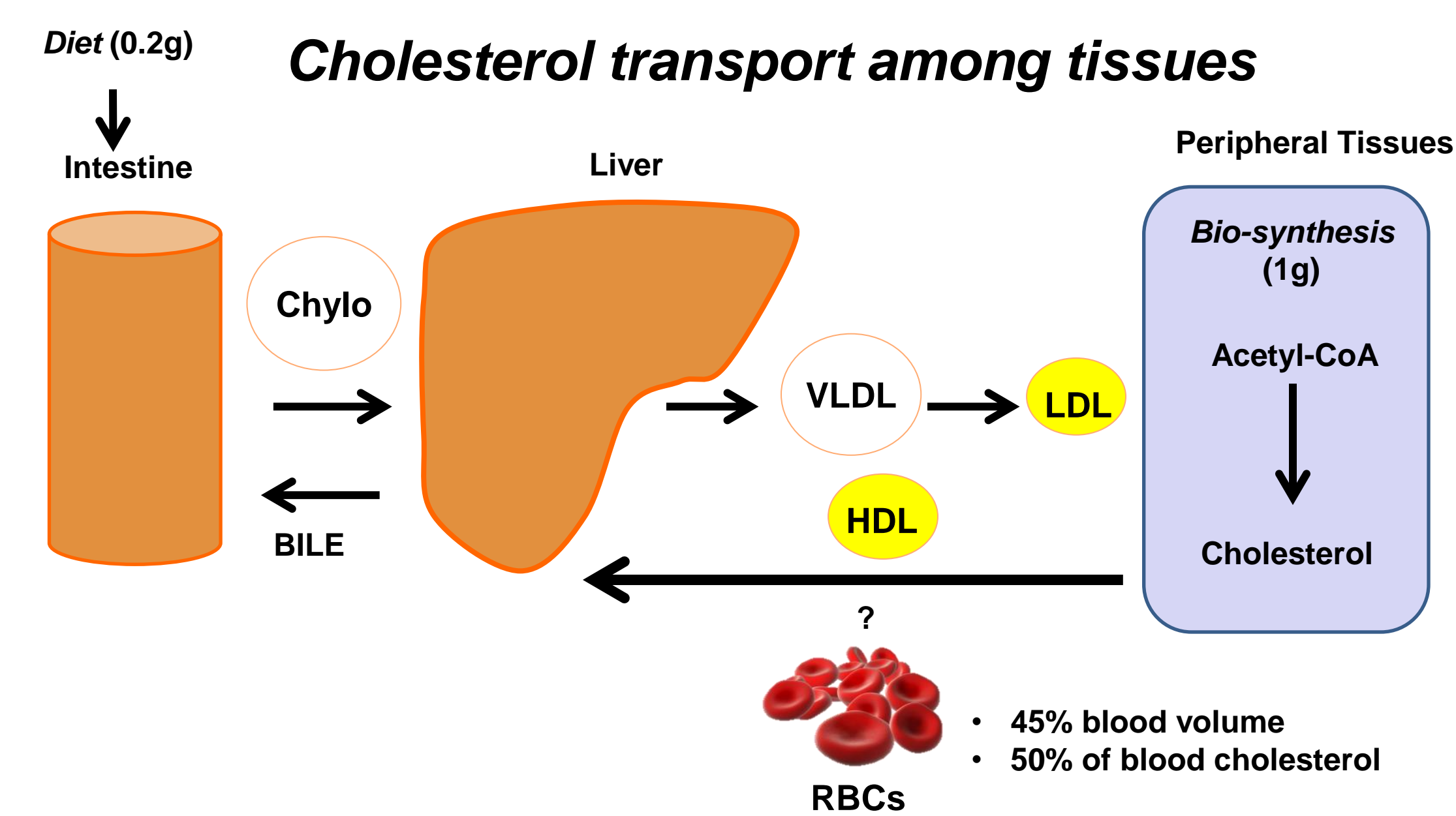


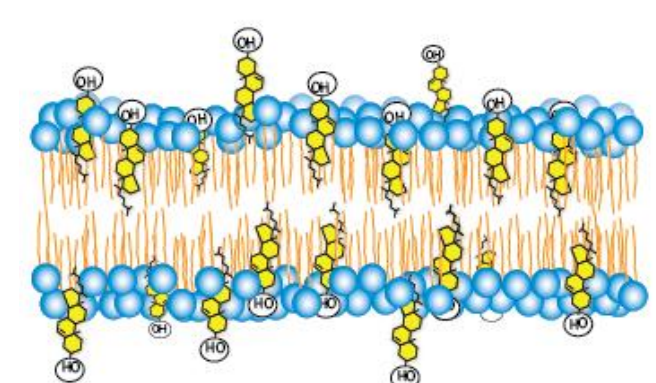
## Abstract

The only cells in the body that cannot synthesize cholesterol are red blood cells (RBCs), yet RBCs contain ~50% of circulating blood cholesterol. Whereas HDL is considered the major conduit for reverse cholesterol transport, we hypothesize that RBCs play a role in this pathway. To test this hypothesis, we developed an assay to measure accessible cholesterol in RBCs. We purified and fluorescently labeled domain 4 of a bacterial toxin, Anthrolysin-O (ALOD4), that binds membrane cholesterol. We incubated fALOD4 with RBCs from 164 healthy subjects and measured the fluorescence intensity using flow cytometry. The intra-assay and intra-individual variability were both <10%, whereas the inter-individual values varied over a 10-fold range. No correlation was found between fALOD4 binding and total RBC-cholesterol, hematocrit, or indices of RBC size. fALOD4 binding was inversely related to membrane phosphatidylcholine (PC) (-0.42,  $p=6e^{-7}$ ) and directly related to lyso-phosphatidylcholine levels (LPC) (0.40,  $p=6e^{-6}$ ). Phospholipase A2 treatment, which converts PC to LPC, increased binding 3-fold. fALOD4 binding did not correlate with plasma LDL-C levels, but was directly related to HDL-C (0.30,  $p=6e^{-4}$ ), and inversely related to triglyceride levels (-0.57,  $p=2e^{-12}$ ). Future studies will determine if variability in fALOD4 binding is intrinsic to RBC membranes, is genetically determined, or contributes to atherosclerosis.



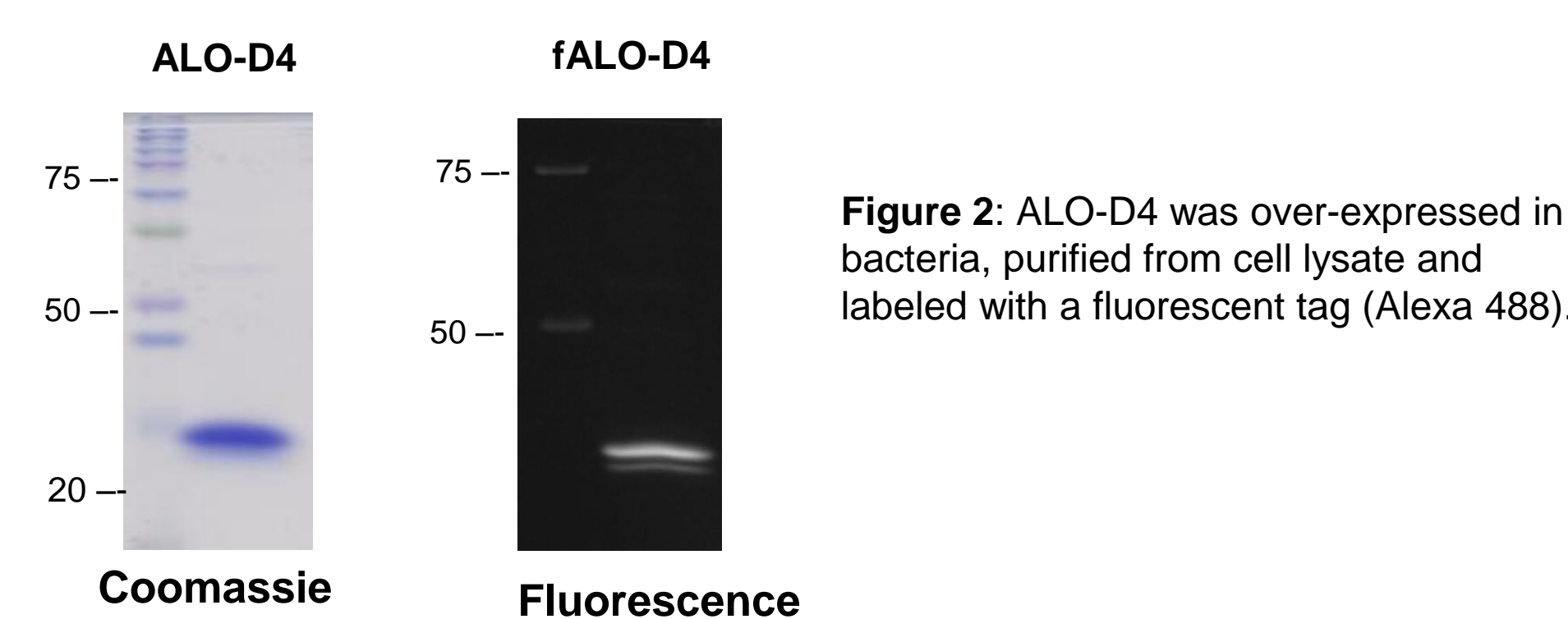
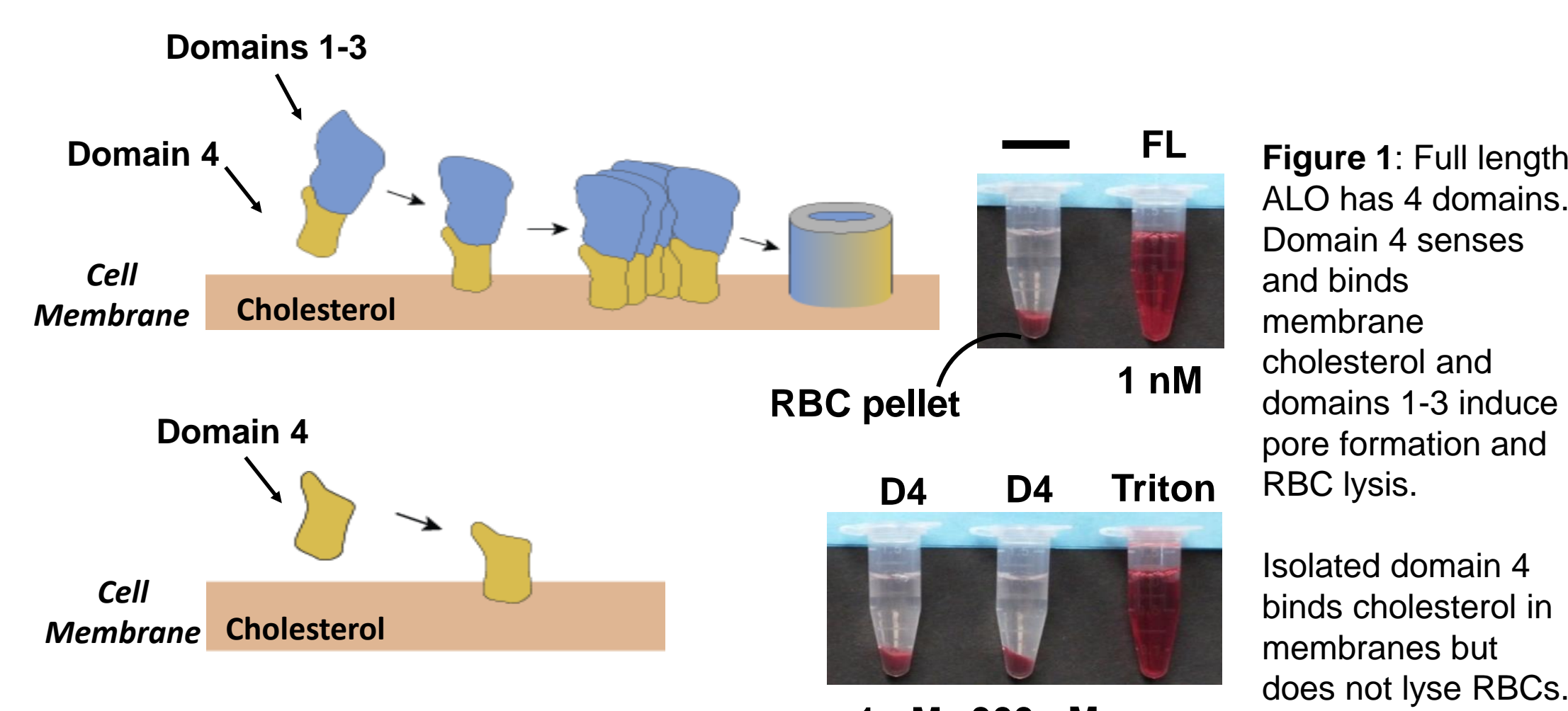
## Hypothesis

An active form of cholesterol exists in the membrane of RBCs which is able to exchange with lipoproteins, cells, and/or other blood components.

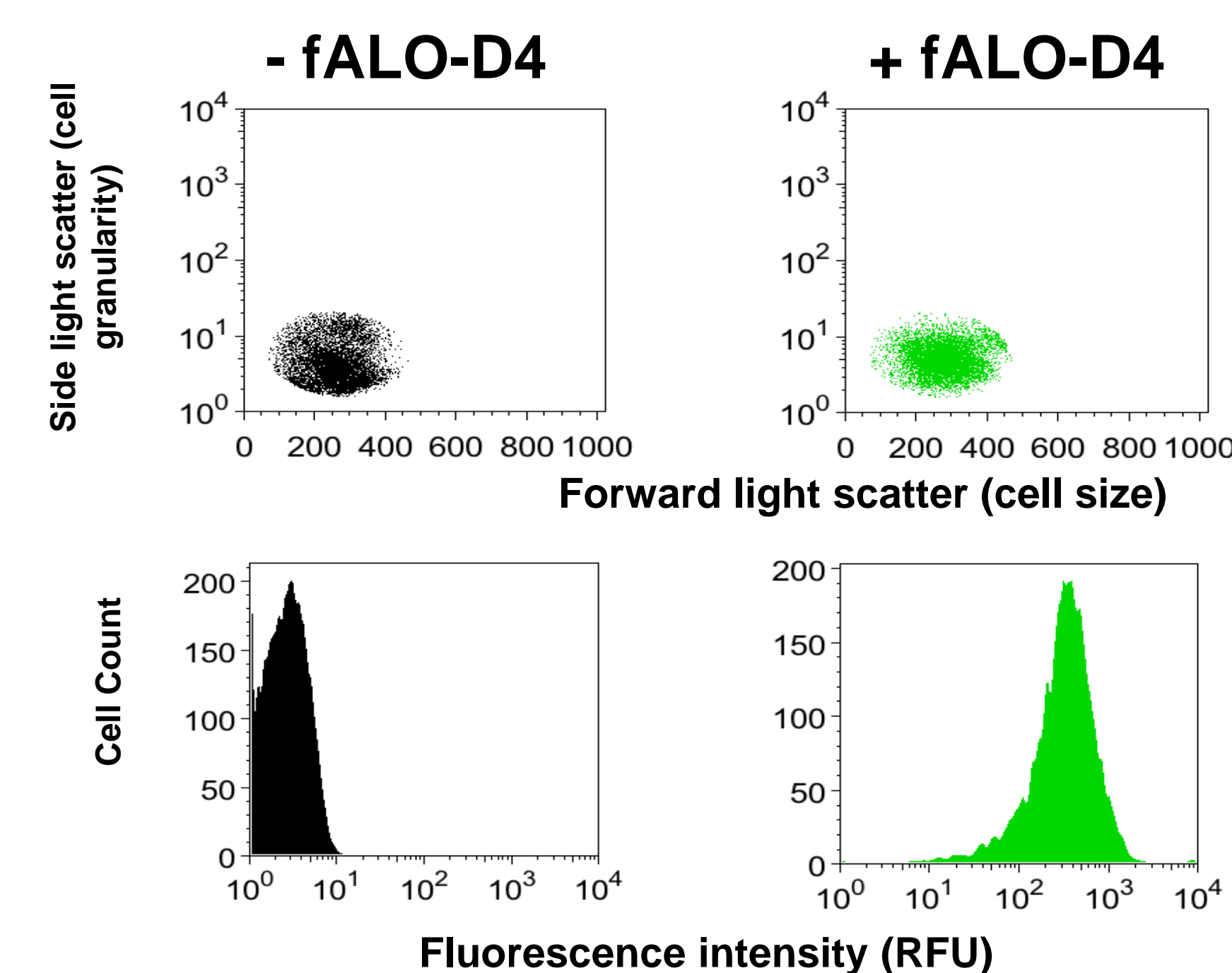


## Assay Development

Anthrolysin-O, a toxin secreted by *Bacillus anthracis*, is a cholesterol dependent cytolysin

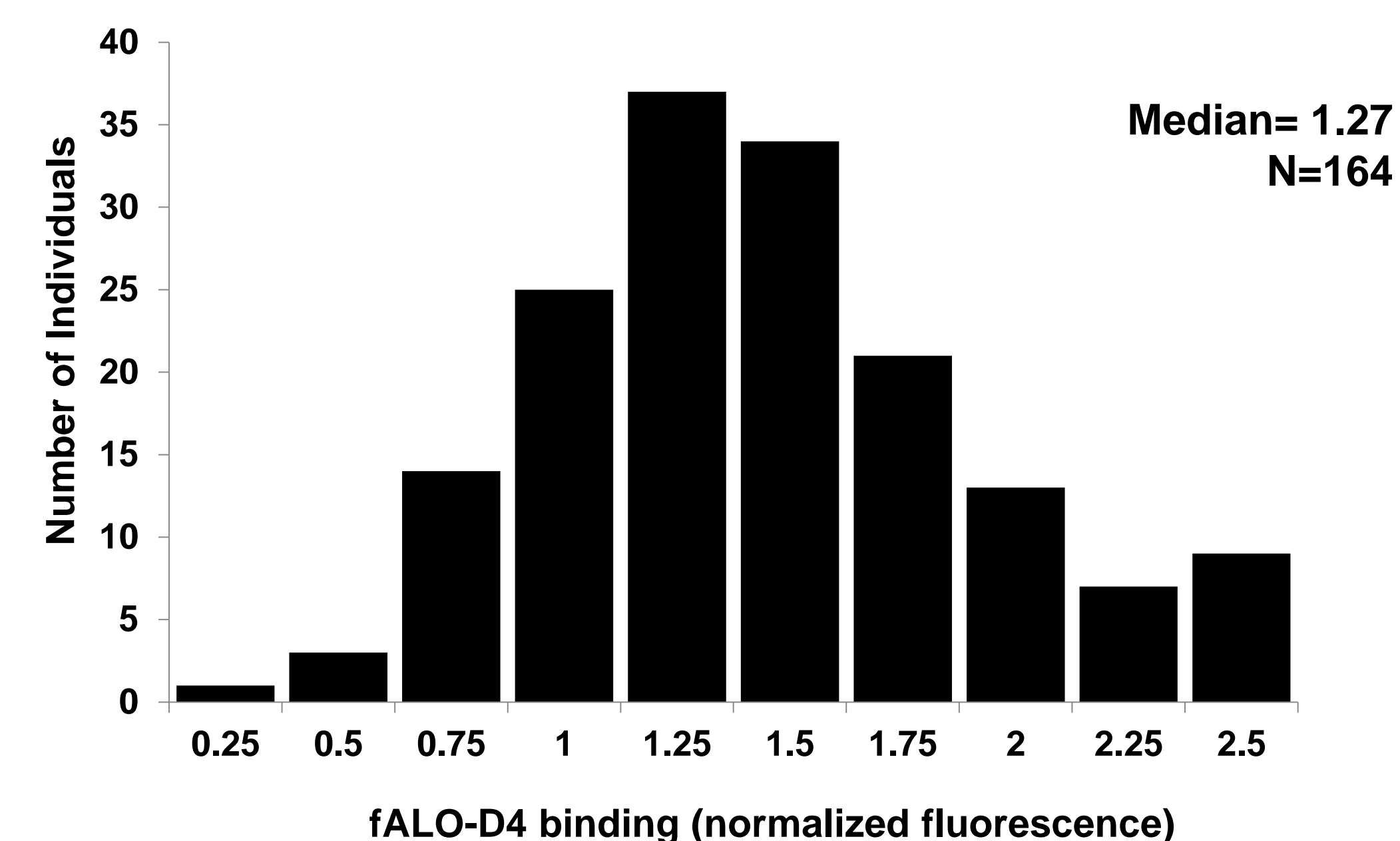


## Flow cytometry of fALO-D4 bound to RBCs



## Results

Distribution of fALO-D4 binding in healthy and unrelated individuals



## Possible Sources of Inter-individual Variation

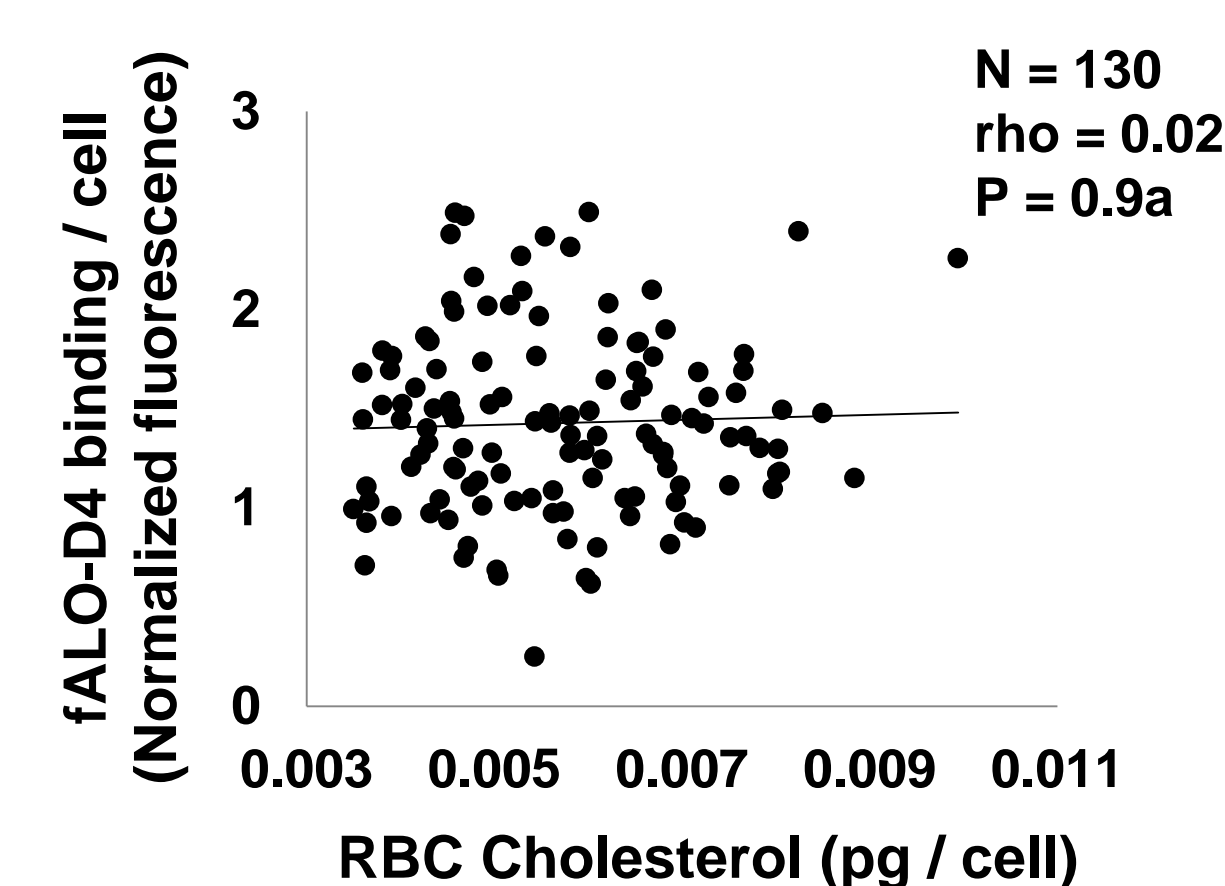
### Intrinsic Property of RBCs

- Lipid Composition
- Protein Composition
- Carbohydrate Modification

### Plasma Constituents

- Lipoproteins
- Other proteins

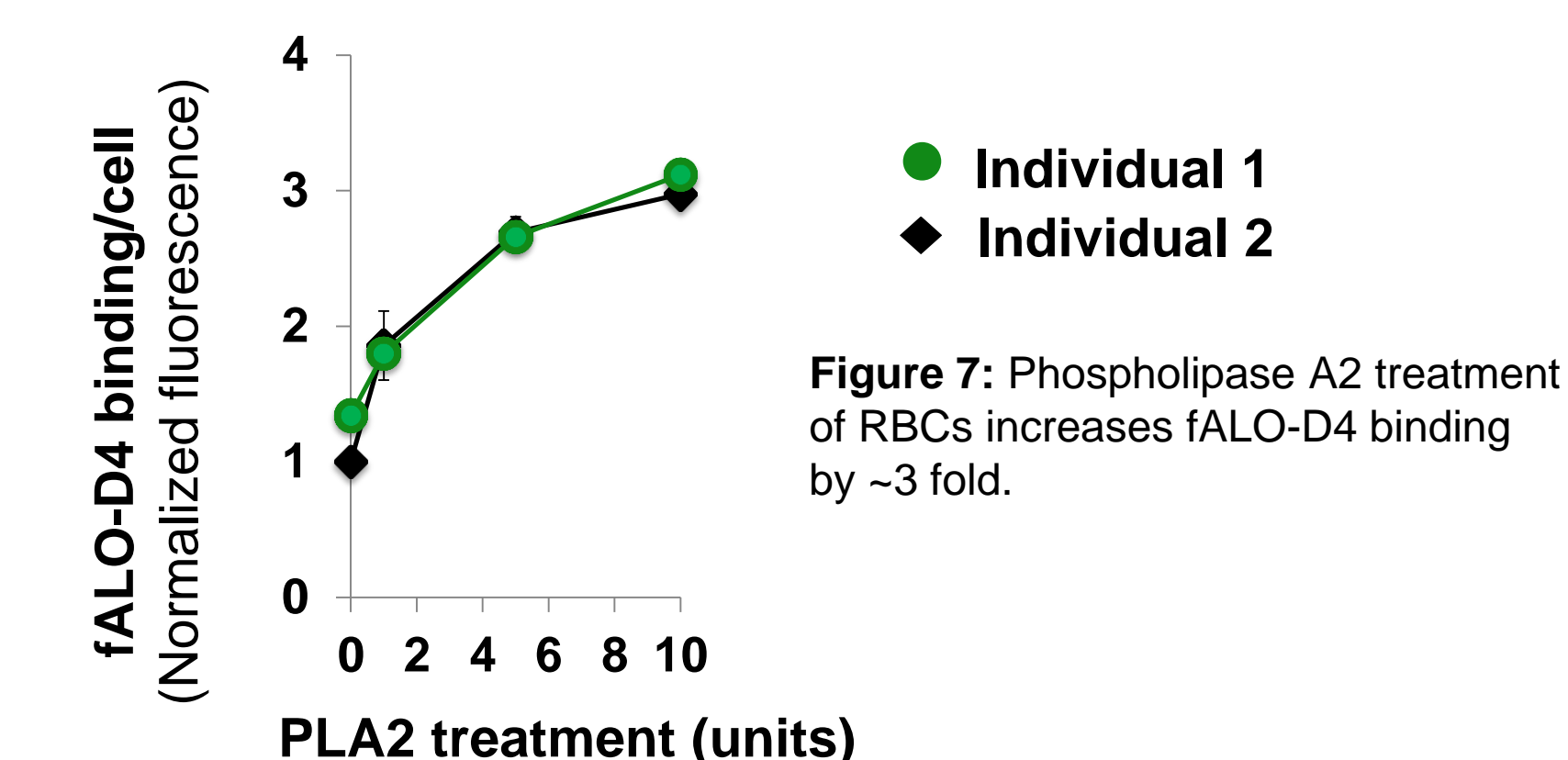
fALO-D4 binding does not correlate with total RBC cholesterol



fALO-D4 binding correlates with outer leaflet RBC phospholipids

Membrane Lipid (mole %)	Spearman(rho)	P
<b>Phospholipids abundant in outer membrane leaflet</b>		
Phosphatidylcholine (PC)	- 0.42	6.00E-07
Lyso-PC	0.39	5.89E-06
Spingomyelin (SM)	0.34	9.25E-05
<b>Phospholipids abundant in inner membrane leaflet</b>		
Phosphatidylethanolamine (PE)	0.11	0.22
Lyso-PE	-0.08	0.36
Phosphatidylserine (PS)	-0.06	0.51

PLA2 treatment of RBCs decreases PC:Lyso-PC and increases fALO-D4 binding



fALO-D4 binding and plasma lipid & lipoprotein levels

Serum Lipid (mg/dL)	Spearman (rho)	P
Total Cholesterol	- 0.27	0.003
Triglycerides	- 0.57	2.16E-12
LDL Cholesterol	- 0.17	0.06
HDL Cholesterol	0.31	0.0001

## Conclusions

- Domain 4 of ALO can be used to assay "accessible" cholesterol in RBC membranes
- RBC accessible cholesterol:
  - ≠ RBC total cholesterol
  - Varies over 10-fold range
  - Related to outer leaflet lipids: ↑SM, lyso-PC ; ↓PC
  - Related to plasma levels of HDL(↑), TG(↓) and TC(↓)
- Decreasing the PC:Lyso-PC ratio in RBC membranes increases accessible cholesterol by ~ 3 fold

## Future studies

- Compare protein & carbohydrate profiles in RBCs from individuals in extremes of activity distribution
- Determine if differences in activity persist after treatment with proteases & glycosidases.
- Examine mutations that alter HDL levels and PC/LPC ratio affect activity
  - LCAT deficiency
  - Tangier disease
- Analyze segregation of trait in families of individuals in the extremes of distribution

## Acknowledgements

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## References

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