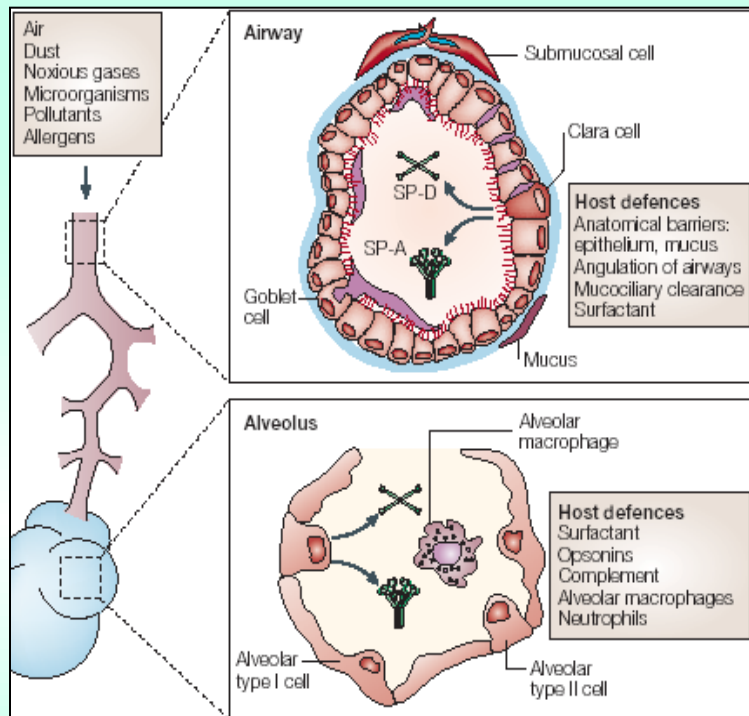


Alterations in Multimeric Structure of Surfactant Protein D as a Biomarker for Lung Injury and Inflammation in Humans

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Surfactant Protein D



- Surfactant protein D is a Ca^{2+} -binding member of **Collagen-like lectins** (“collectins”) that play a role in non-antibody-mediated innate immune responses.

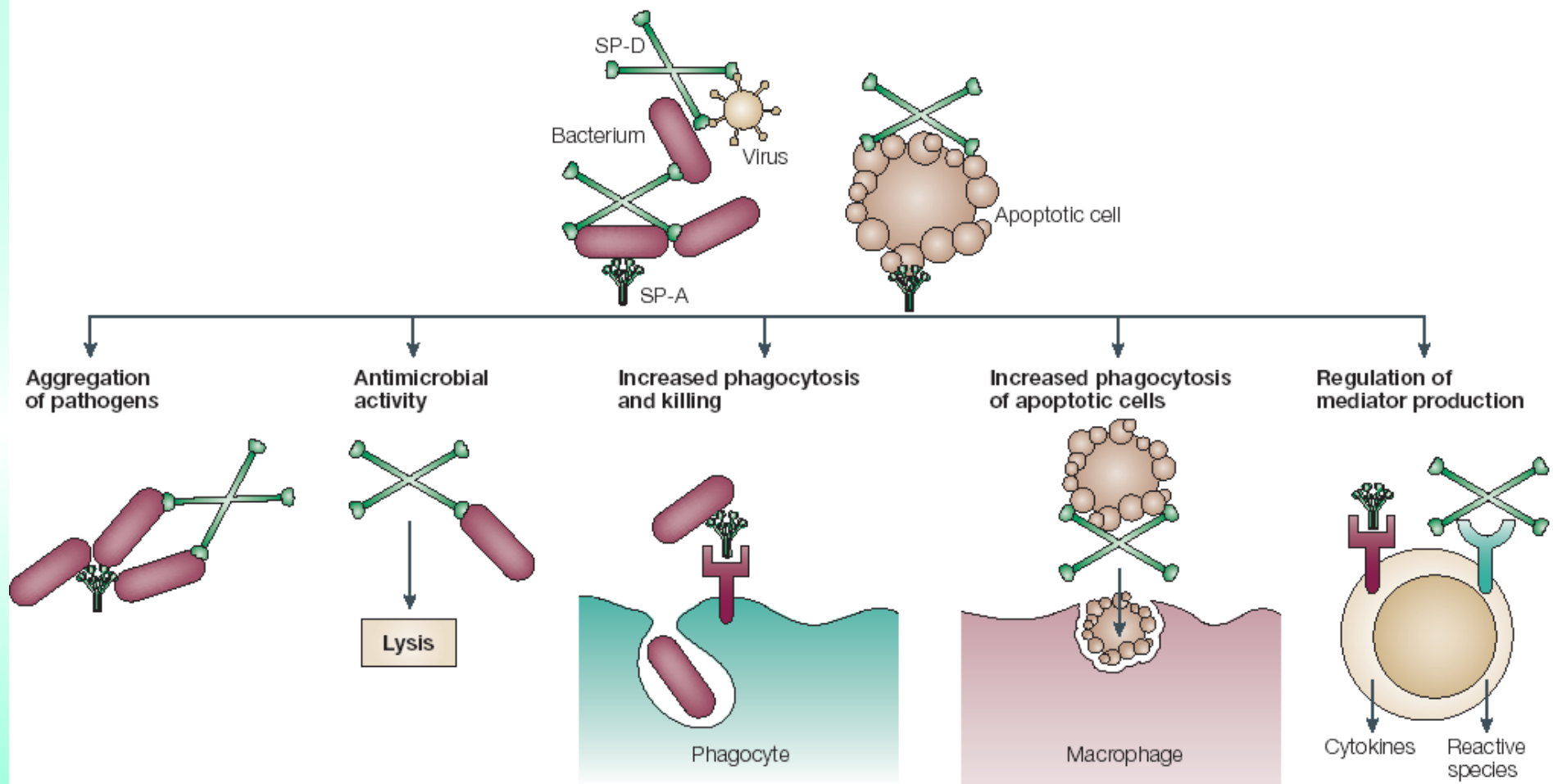
- SP-D shares considerable structural homology with other proteins of this type (including SP-A, conglutinin, bovine collectin-43, and mannose binding protein).

- SP-D is produced primarily by alveolar type II cells and nonciliated bronchiolar cells in the lung.

- The primary function of the pulmonary collectins appears to be the modulation of host defense and inflammation.

Function of SP-A and SP-D

(J.R. Wright, 2006)

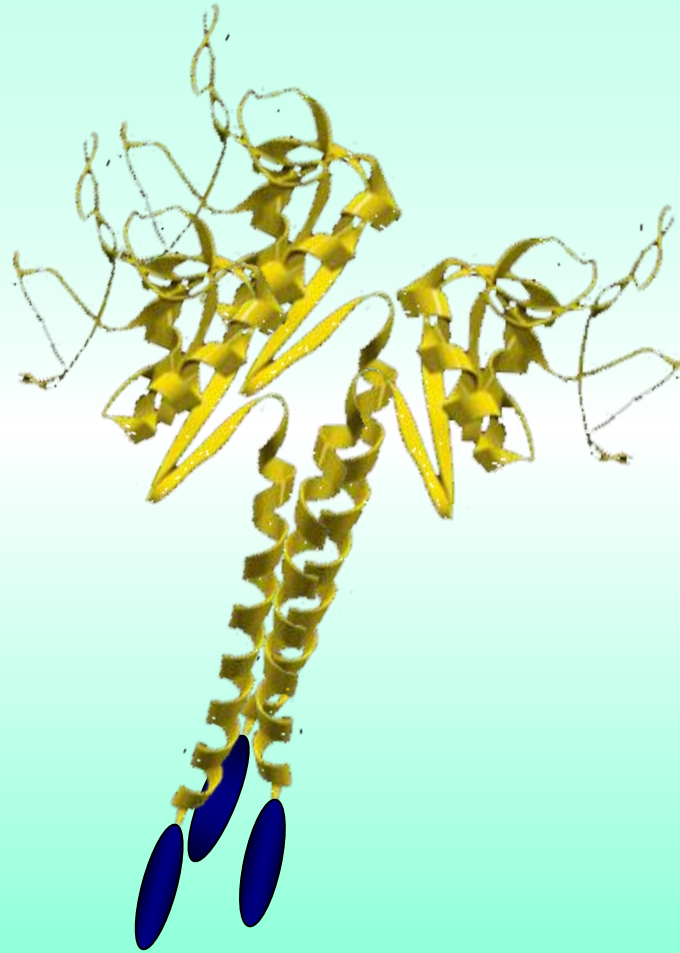


SP-A and SP-D bind to a variety of bacteria, viruses, allergens and apoptotic cells and thereby function as opsonins to enhance the uptake of these cells and particles. Binding of the collectins to pathogens occurs by various mechanisms.

SP-A and SP-D also have direct effects on immune cells and modulate the production of cytokines and inflammatory mediators.

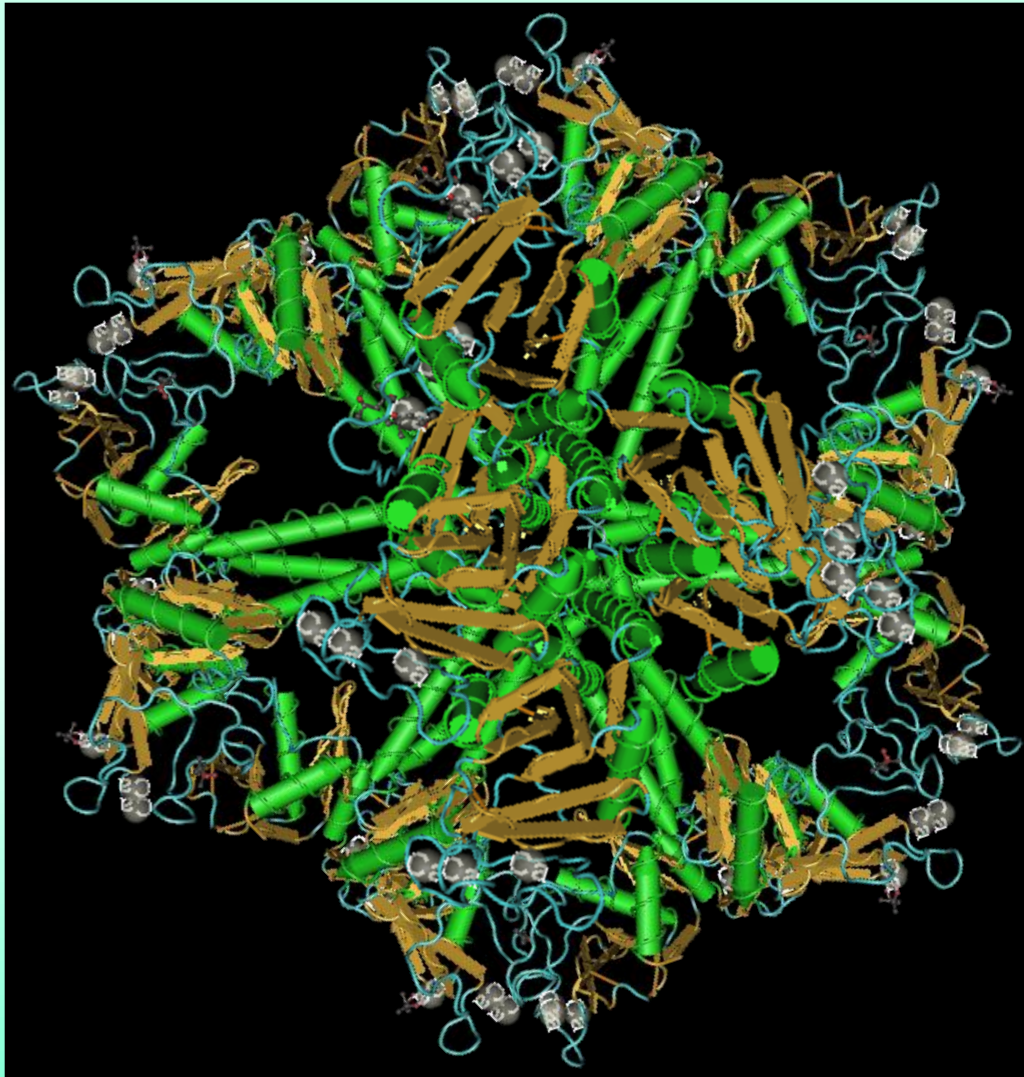
Collectin monomers are highly oligomerized

The head and neck domains drive the aggregation of the SP-D monomer to form a trimer of ~130 kDa



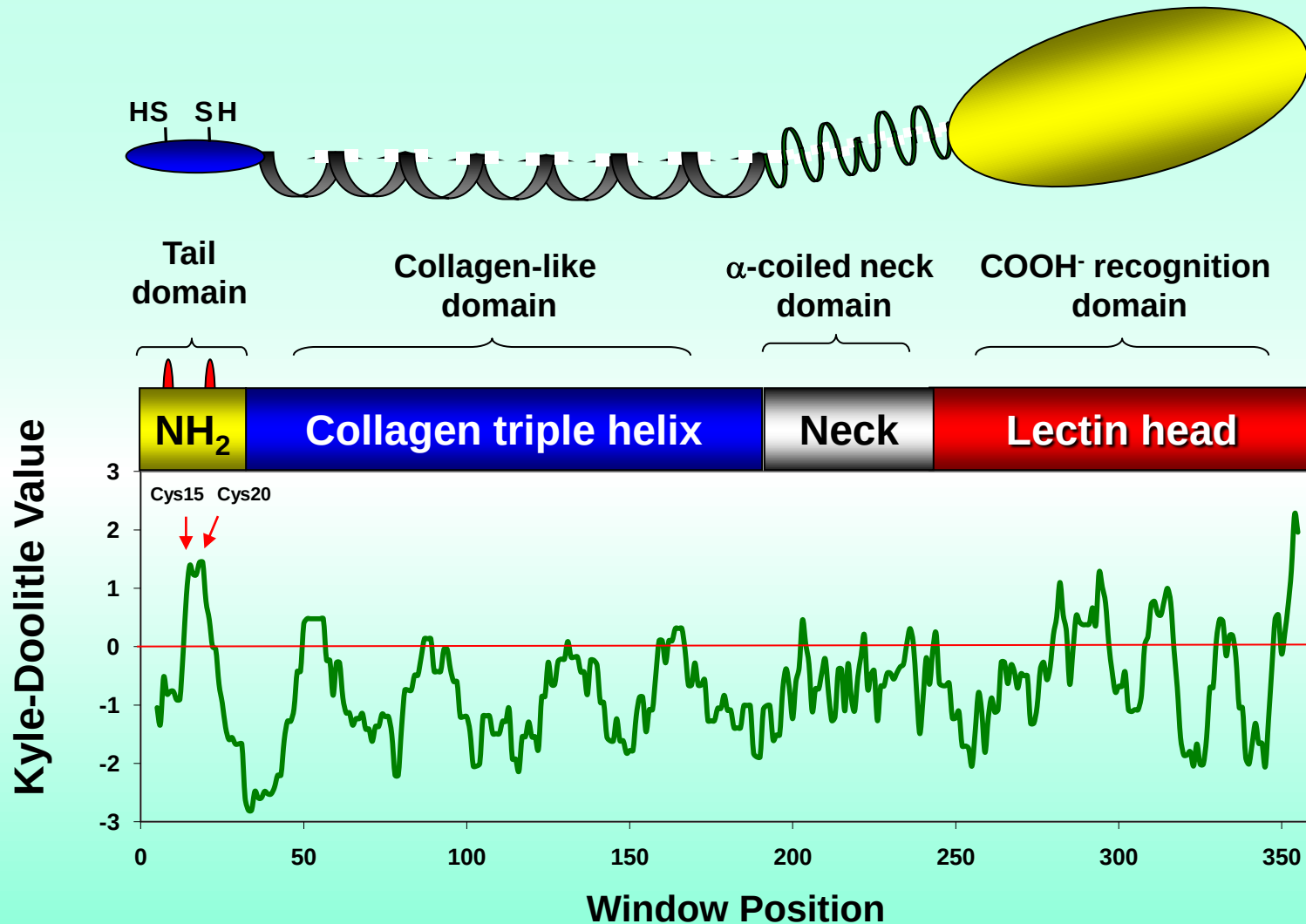
SP-D can form multimers, so-called “fuzzy balls” with a total mass of several million kDa and a size of about 100 nm

Oligomerization of SP-D results in the masking of the tail domains while the head domains remains exposed



size of about 100 nm

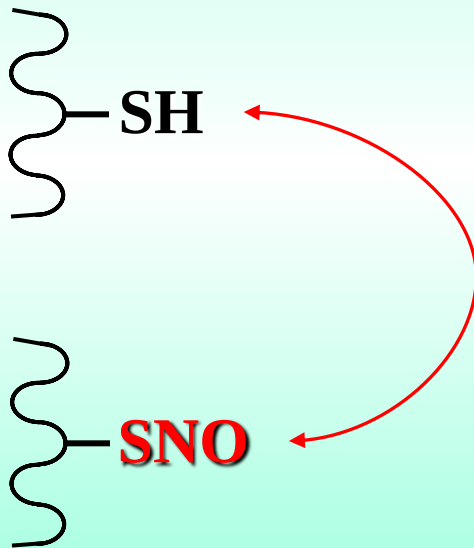
Monomeric Structure of SP-D



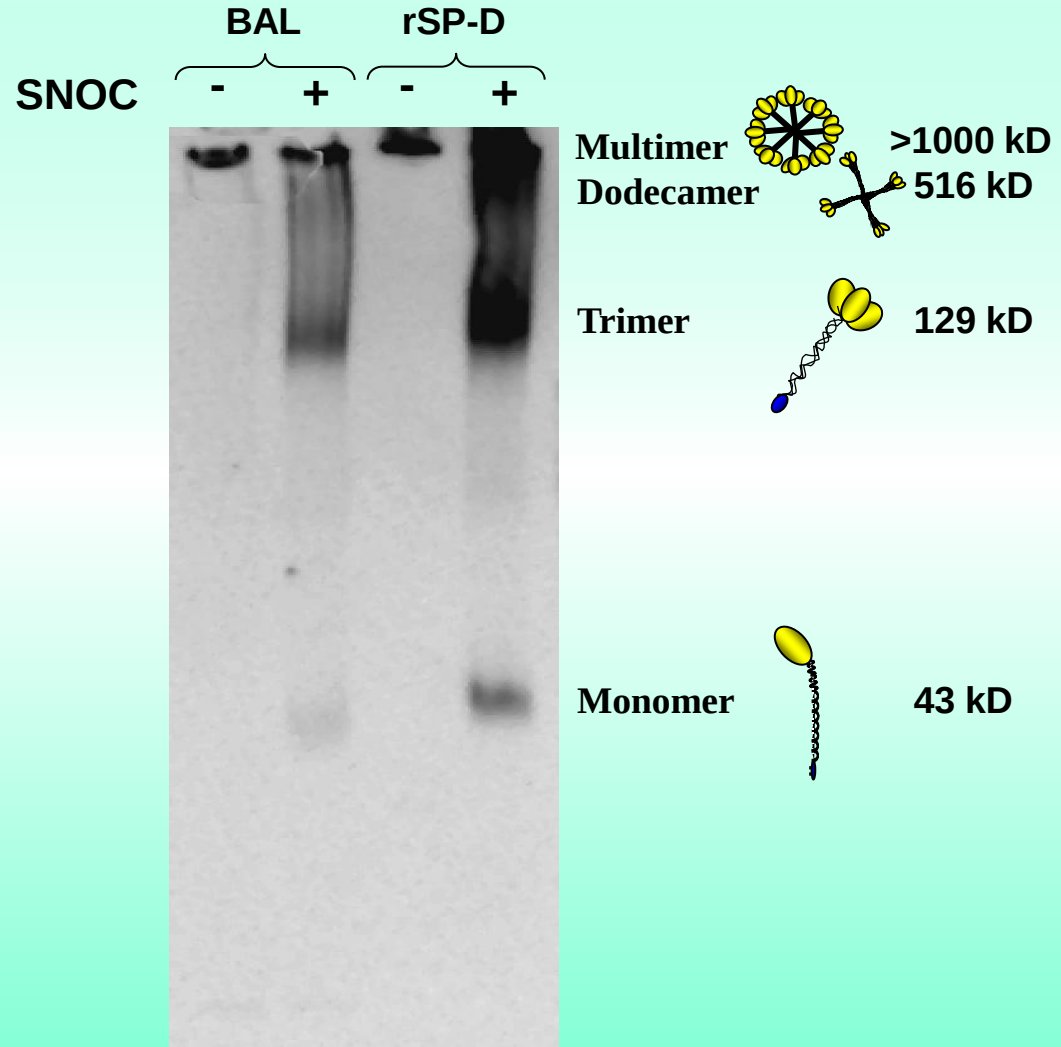
There are 6 cysteines within SP-D that potentially could be nitrosylated (2 in the NH₂-terminal region and 4 in the carbohydrate-binding domain). However, only two cysteines **15** and **20** are situated within the hydrophobic tail region, as a motif for nitrosylation.

SP-D multimeric structure can be altered by S-nitrosylation *in vitro*

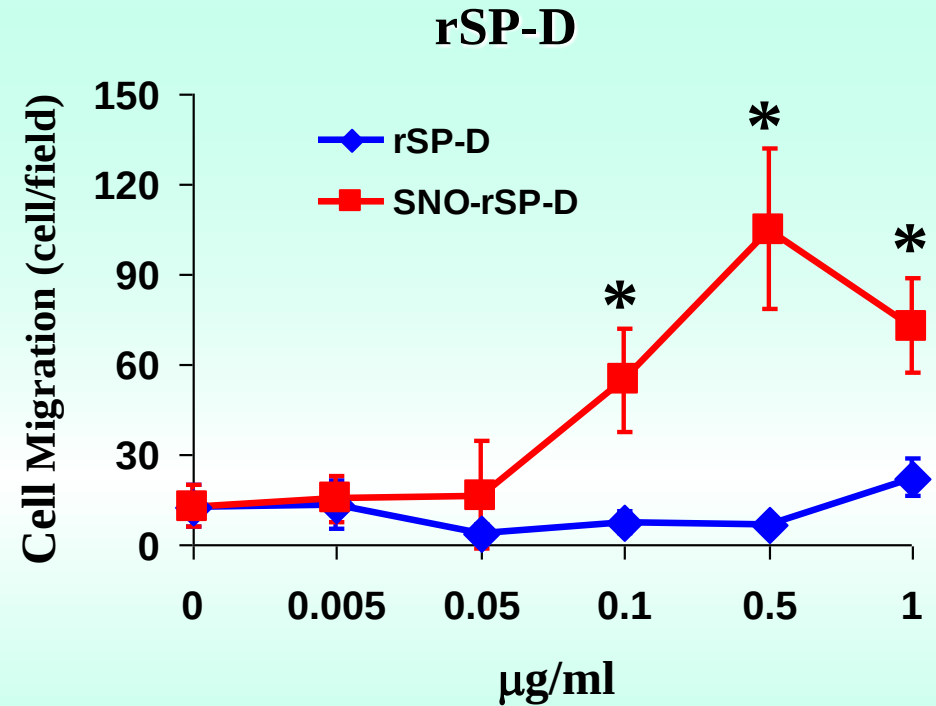
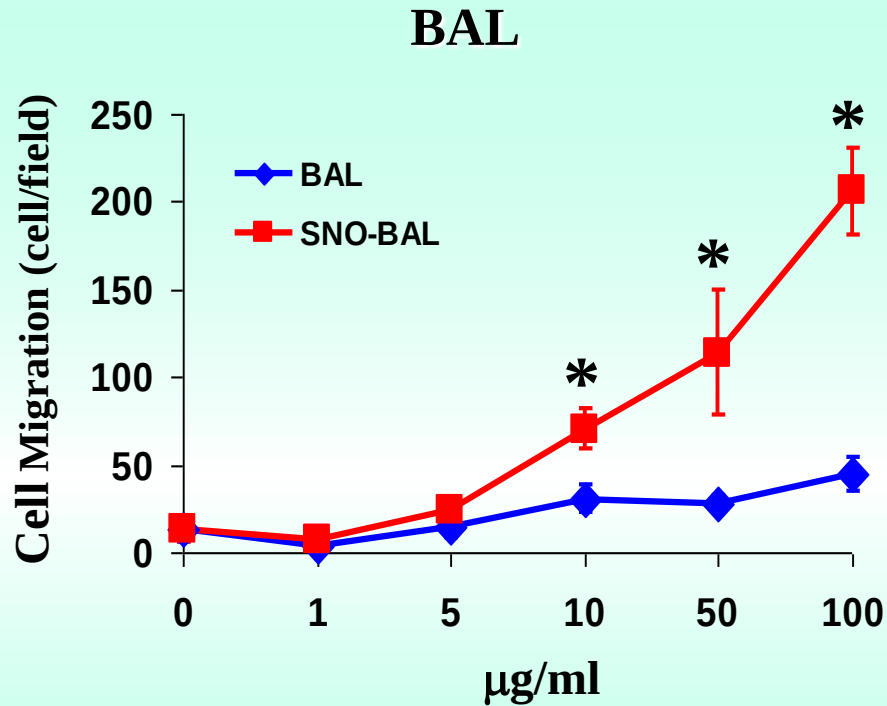
S-nitrosylation is a covalent modification of thiol groups resulting in S-nitrosothiol (SNO) formation.



SNOC (S-nitrosocysteine)



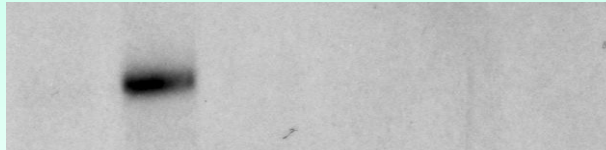
S-nitrosylation of normal BAL and rSP-D induces macrophage chemotactic function



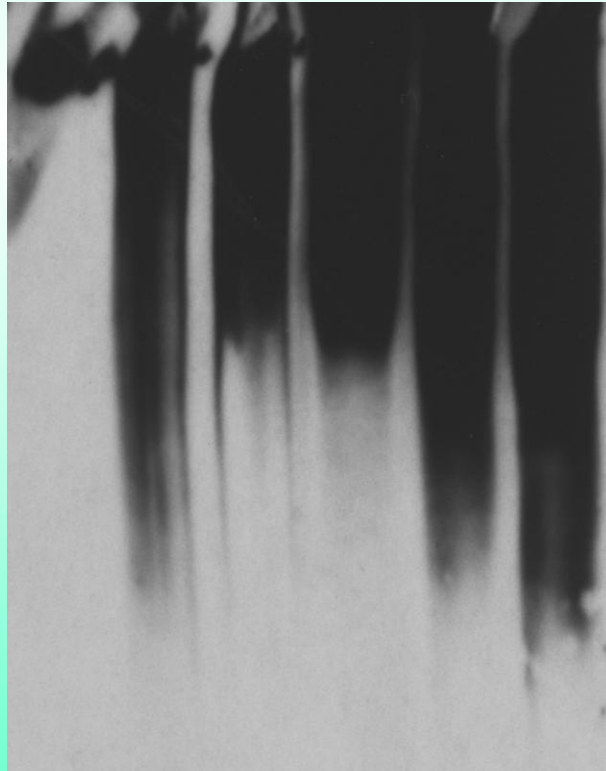
- S-nitrosylation unmasks a pro-inflammatory function of SP-D
- SNO-SPD is a chemoattractant

Signaling in SP-D is NO-Dependent

	RrSP-D			Ser15/20		
L-SNOC	-	+	-	-	+	-
L-Cys	-	-	+	-	-	+



SNO-SP-D



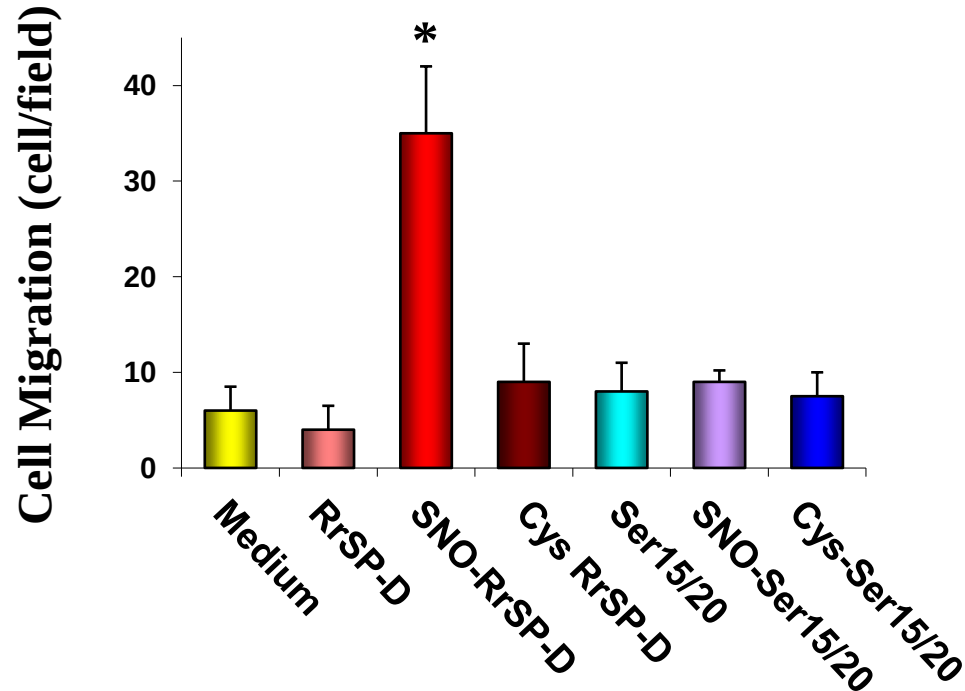
Native SP-D

Signaling in SP-D is NO-Dependent

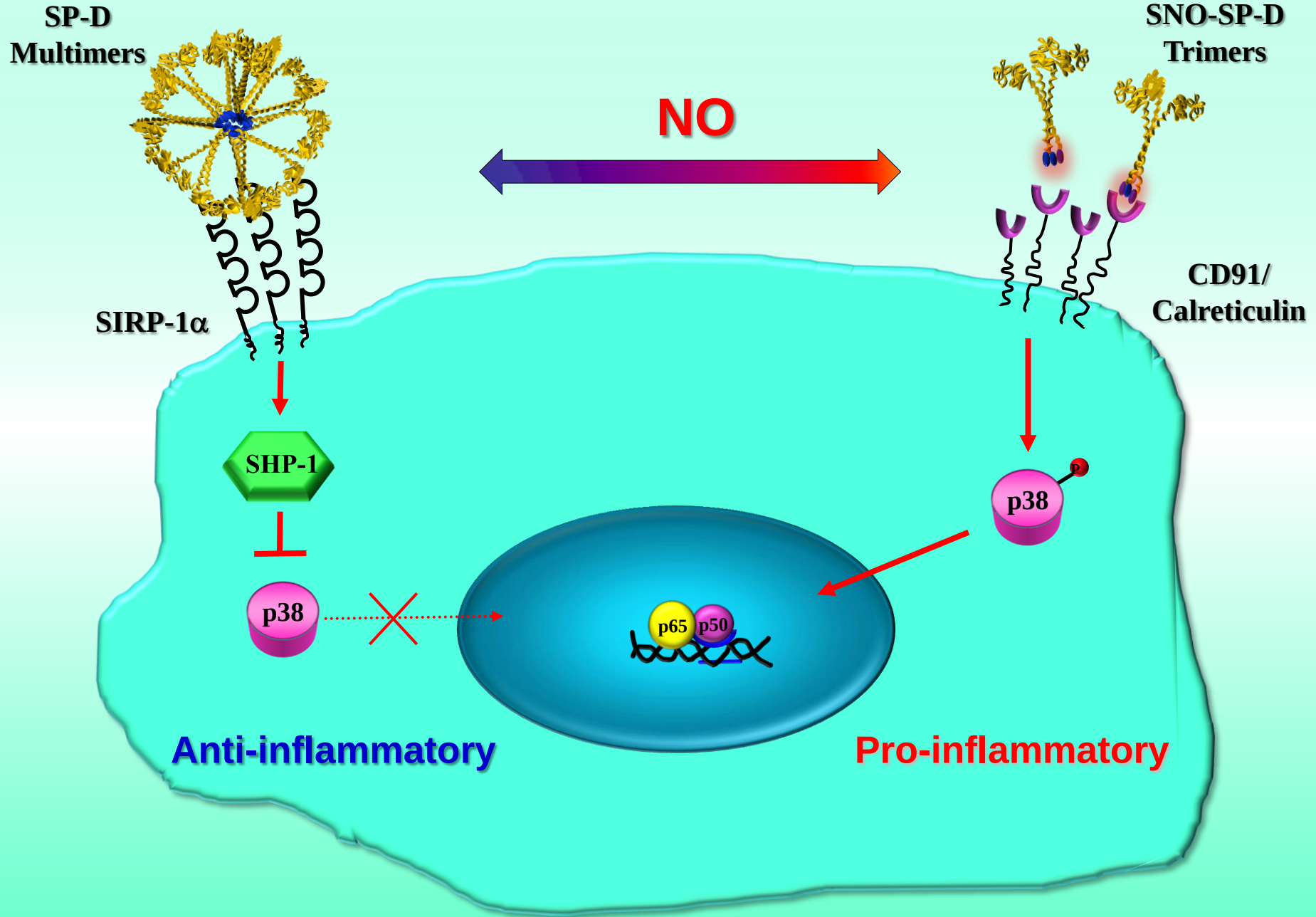
	RrSP-D			Ser15/20		
L-SNOC	-	+	-	-	+	-
L-Cys	-	-	+	-	-	+



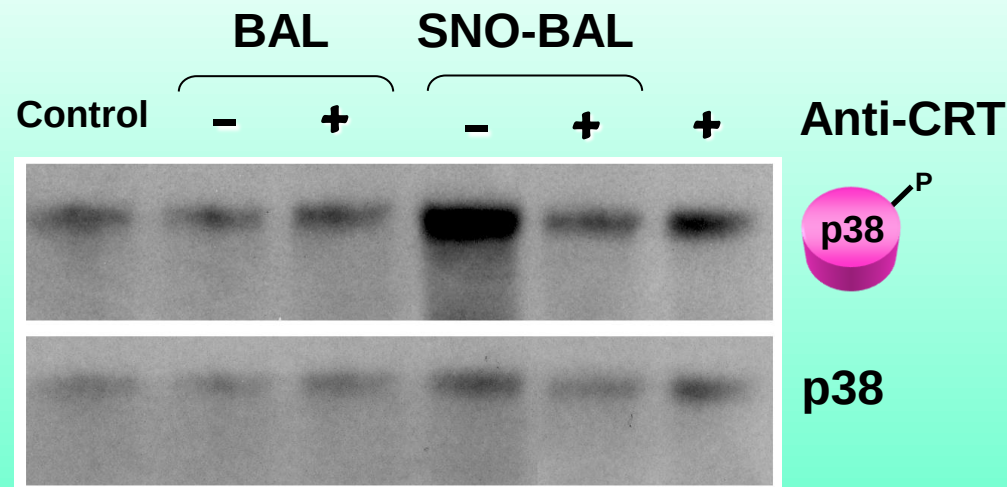
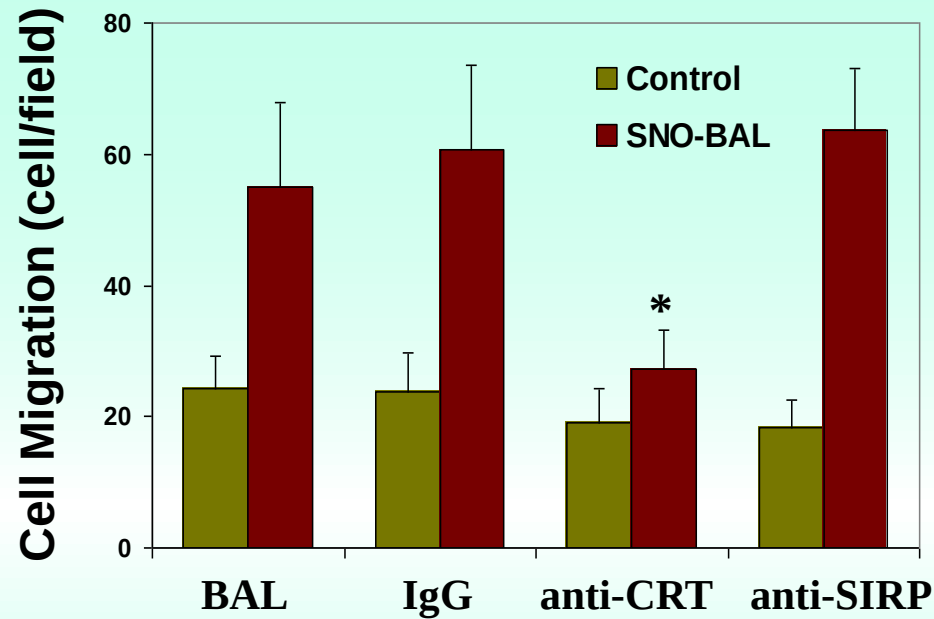
SNO-SP-D



Model of the pro and anti-inflammatory functions of SP-D



Pretreatment with anti-CRT antibody specifically reduced the SNO-BAL mediated chemotaxis and p38 phosphorylation

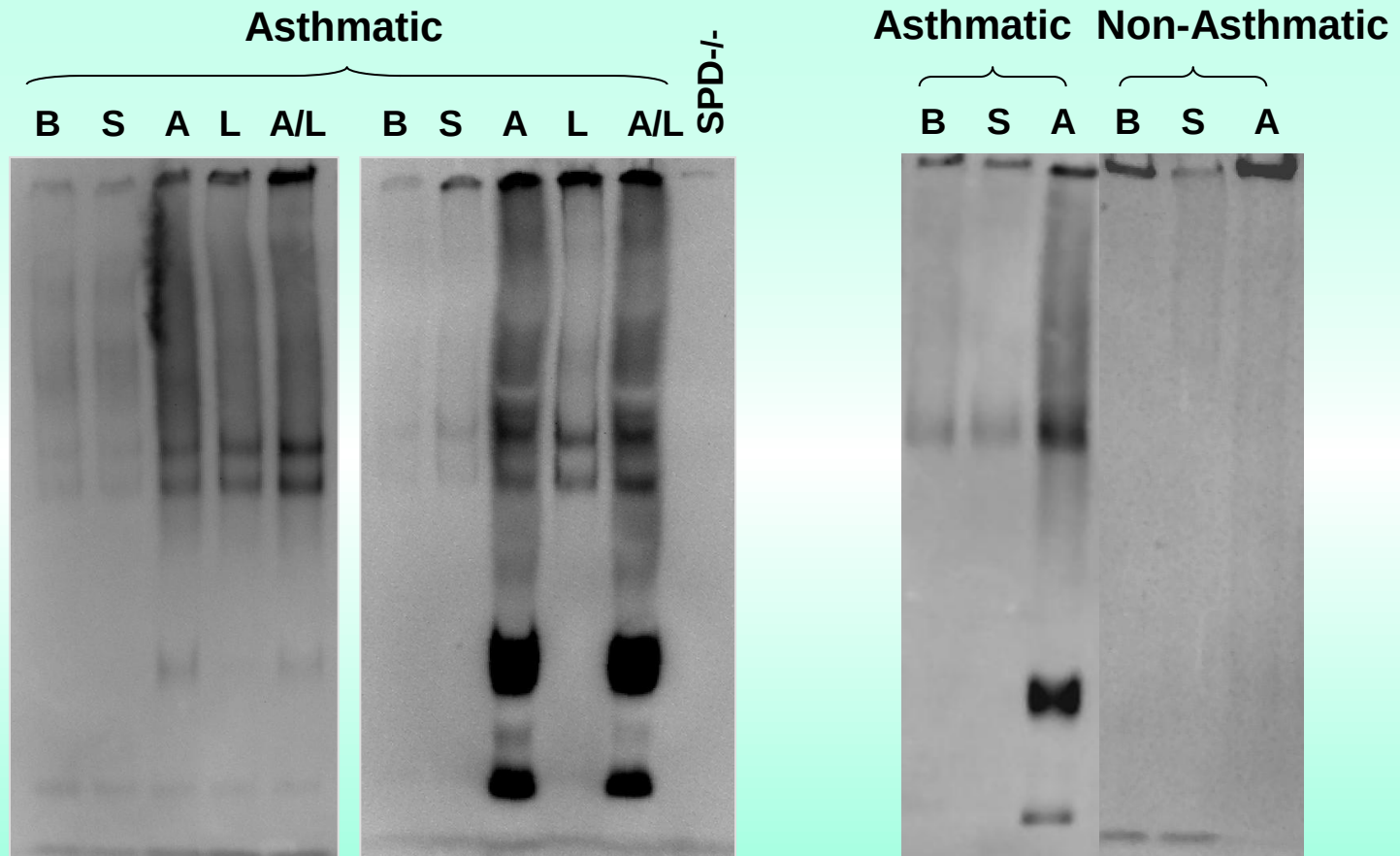


The human inflammatory lung diseases, Asthma and Hermansky Pudlak Syndrome type 1 (HPS1) are associated with increases of SP-D in BAL and amplification of inflammatory signals in the distal lung

**We hypothesize that
post-translational modification of
SP-D with disruption of its multimeric
structure is associated with
inflammatory lung disease in humans**

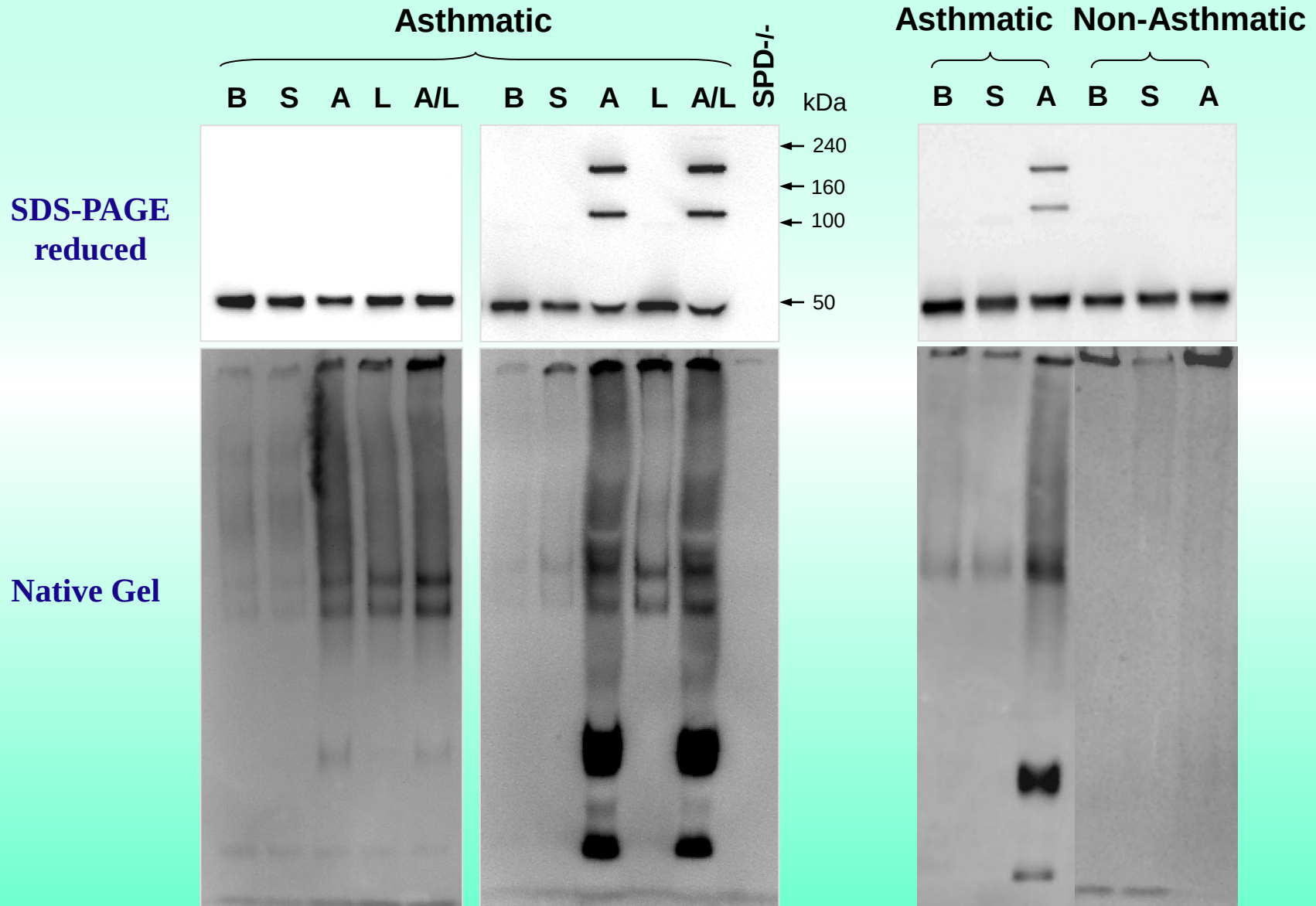
Multimeric structure of SP-D is altered after segmental challenge

Native Gel

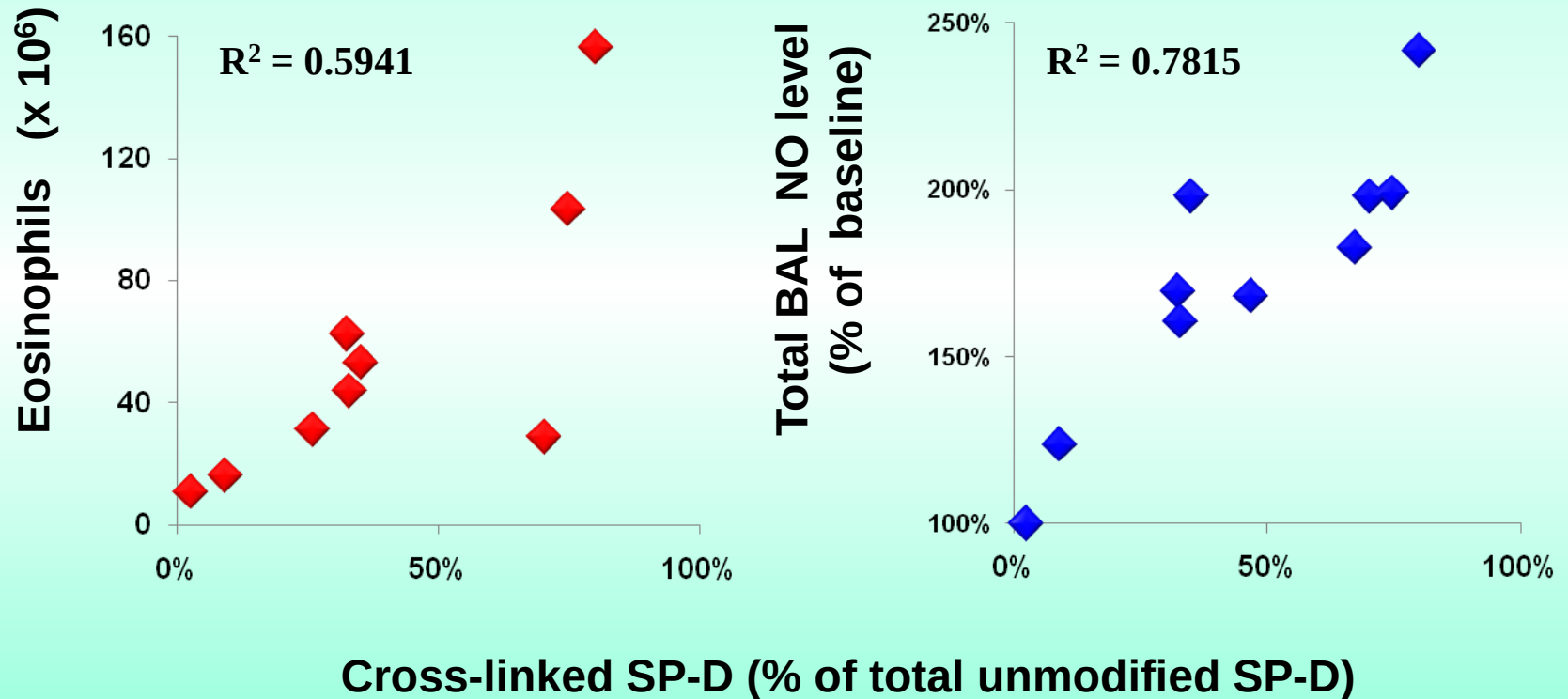


Non-asthmatic patients did not develop cross-linked SP-D bands and SP-D multimeric structure was not disrupted after the same procedure

8 out of 16 asthmatic patients formed cross-linked SP-D after segmental challenge



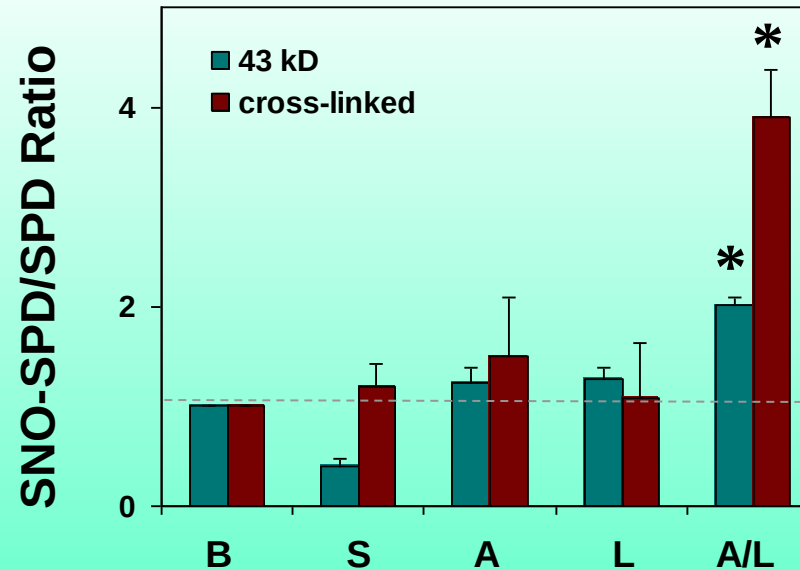
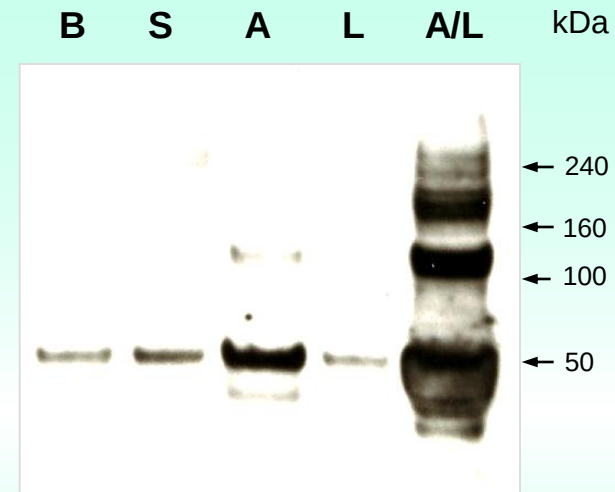
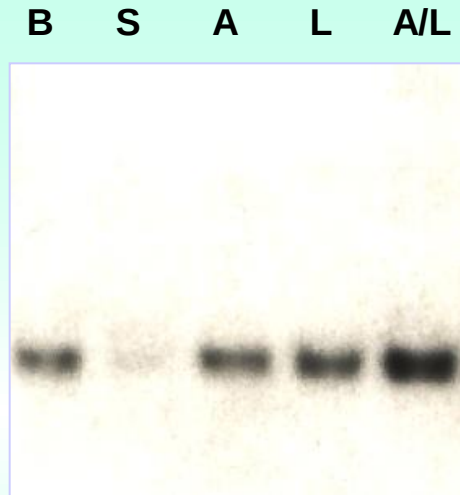
Cross-linked SP-D is correlated with increased BAL eosinophilia and increased NO levels after segmental challenge



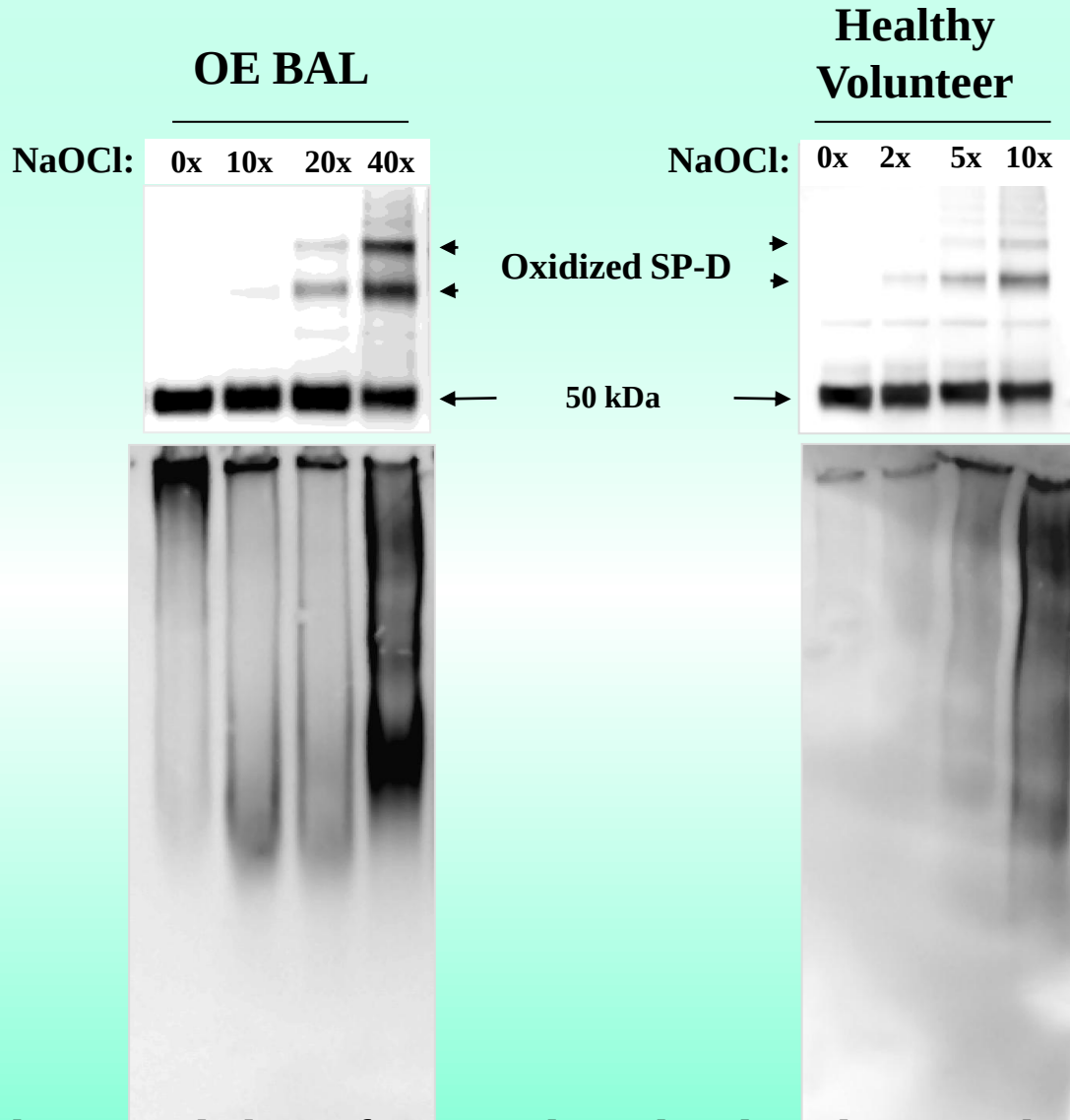
Cross-linked SP-D is associated with SNO-SPD formation after segmental challenge

Patients with monomeric SP-D

Patients with cross-linked SP-D

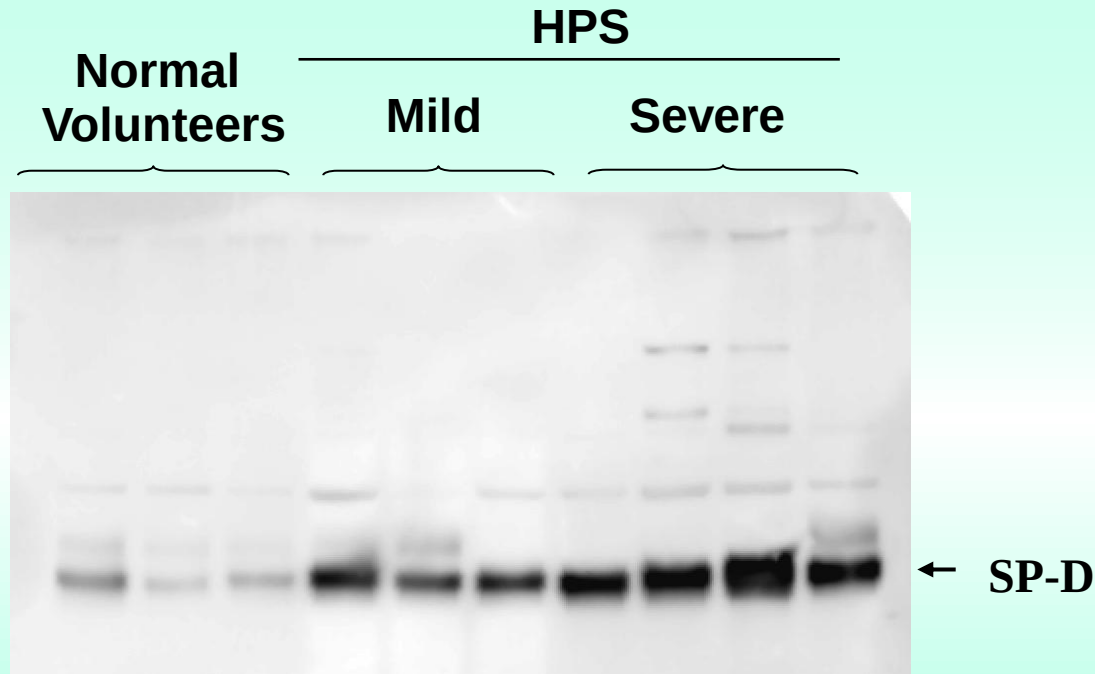


Oxidative cross-linking of SP-D *in vitro*



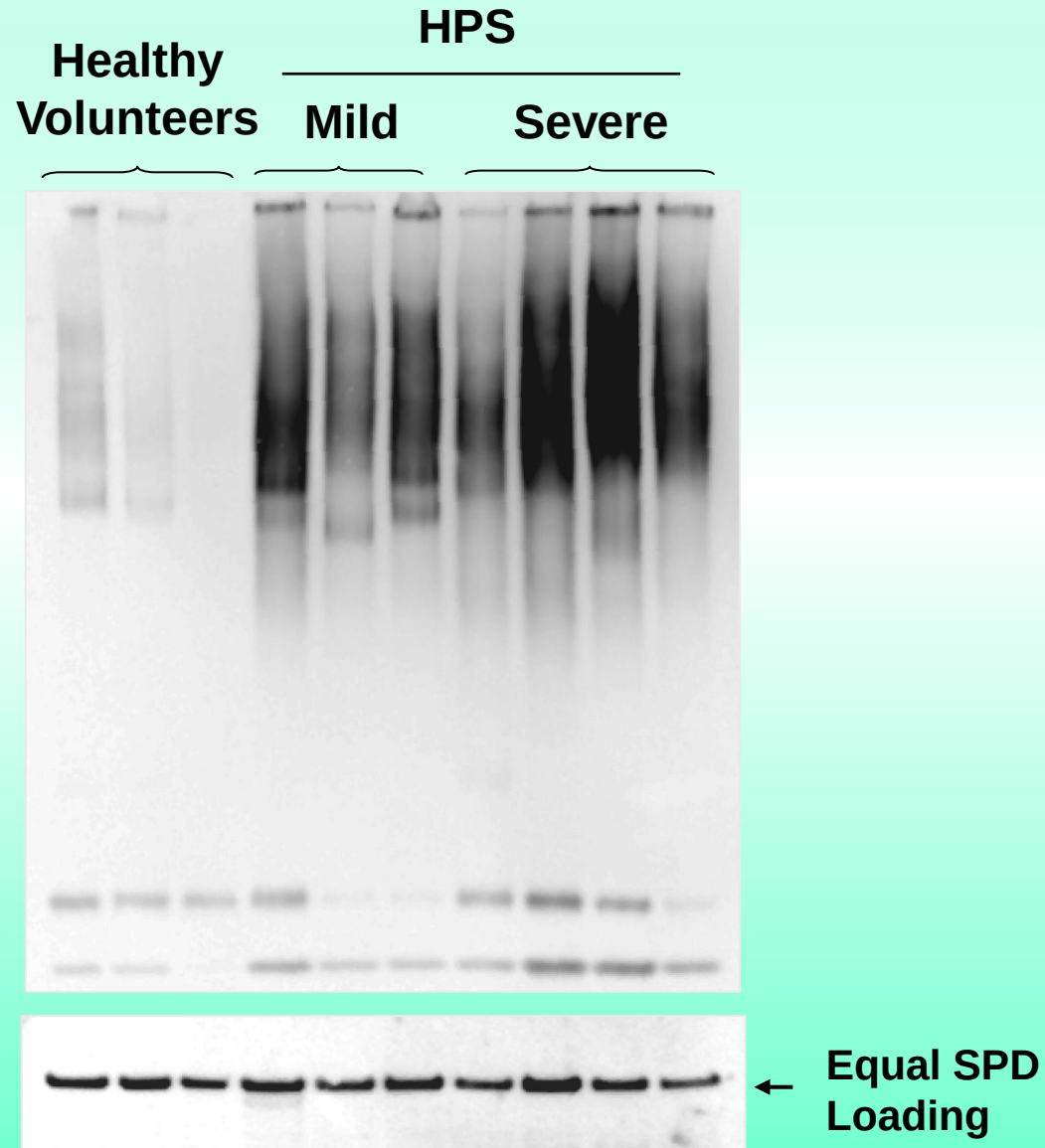
Dose-dependent cross-linking of SP-D under reduced conditions; indicating the formation of non-disulfide covalent cross-linked forms of SP-D similar to those that appear in the BAL of asthmatic patients after segmental challenged

BAL SP-D content increases with increasing disease severity in HPS1 patients

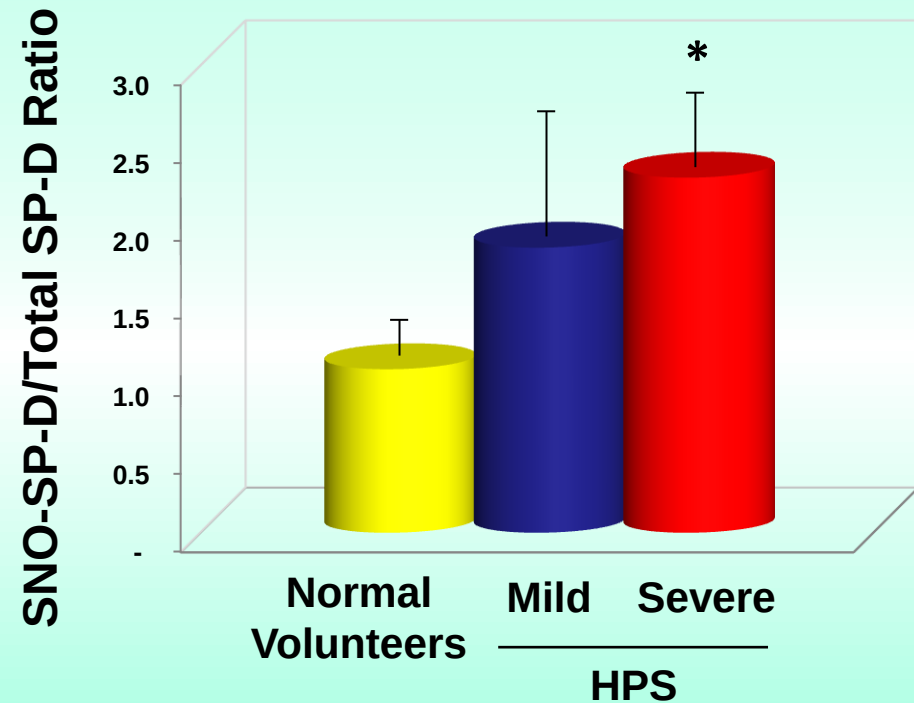
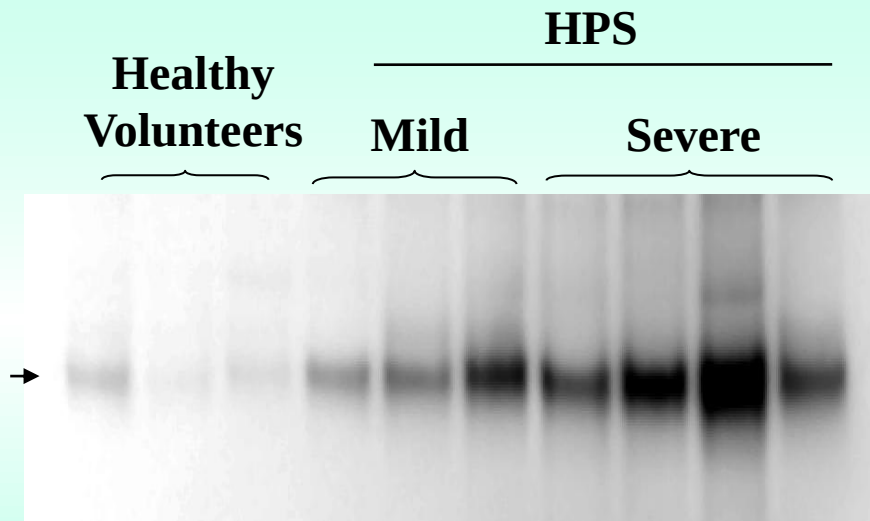


Representative immunoblot for total SP-D on samples of BAL from normal volunteers and HPS1 patients

SP-D structural disruption is associated with pulmonary inflammation in HPS

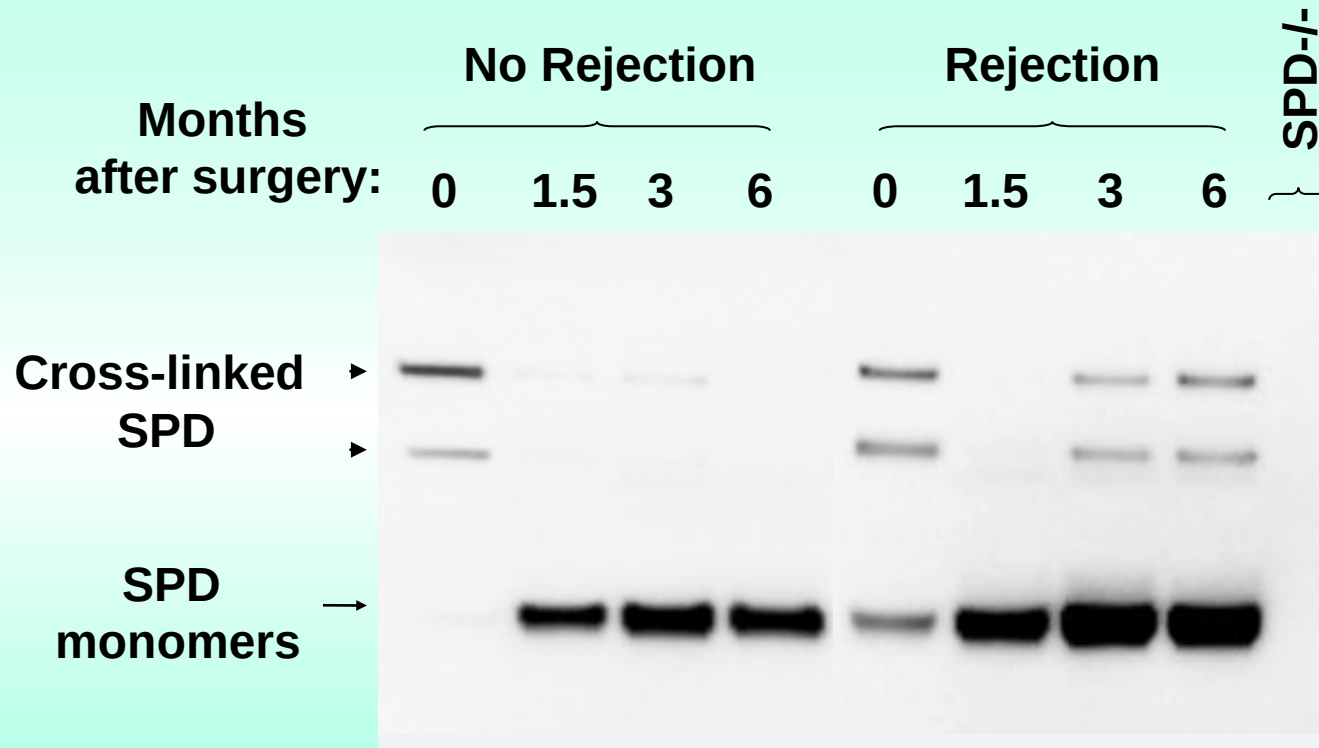


SNO-SP-D is associated with pulmonary inflammation in HPS

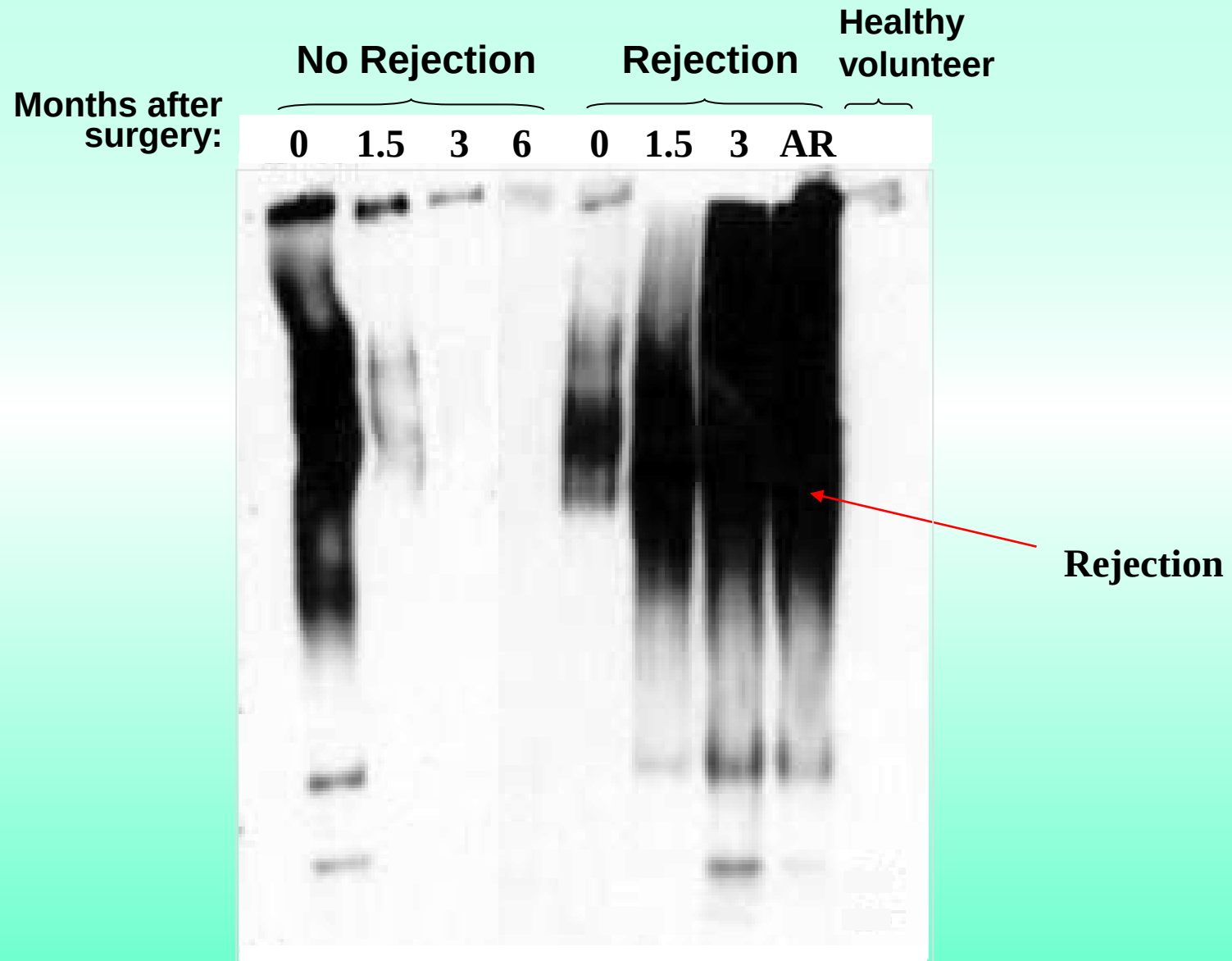


BAL SNO-SP-D content also increases with increasing disease severity in HPS1 patients

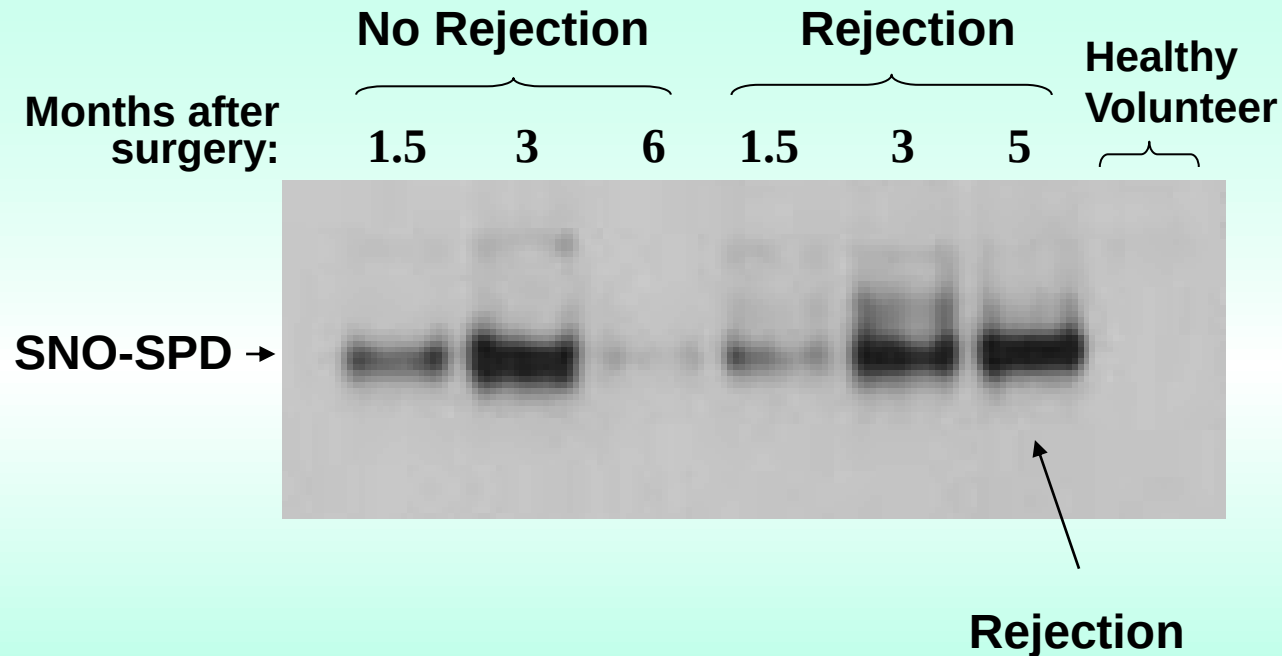
Acute rejection of lung allograft is associated with cross-linking of SP-D



Acute rejection of lung allograft is associated with disruption of SP-D multimers



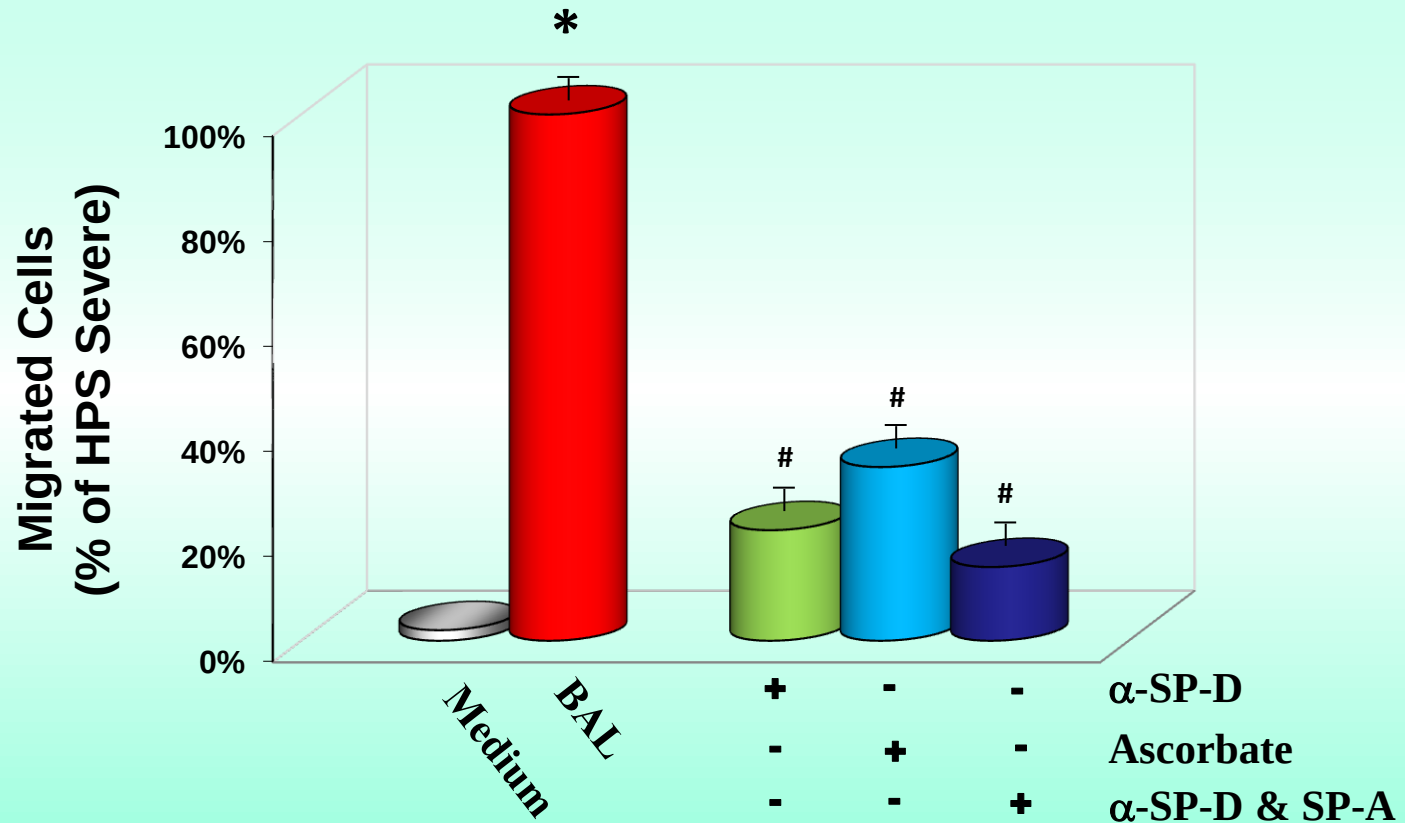
Acute rejection of lung allograft is associated with SNO-SP-D



- **SP-D can be S-nitrosylated to form SNO-SP-D by lung inflammation**
- **S-nitrosylation of SP-D reduced its multimeric state**

Does SNO-SPD have any functional consequences for lung inflammation?

SP-D modifications in HPS1 patients promote enhanced macrophage migration



Summary

- Lung inflammation induces the production of NO and S-nitrosylation of SP-D (SNO-SPD)
- S-nitrosylation leads to the disruption of the SP-D multimeric structure
- S-nitrosylated SP-D activates alveolar macrophages through calreticulin and CD91 receptor and could enhance lung injury during different experimental models
- Oxidatively cross-linked SP-D is formed in human inflammatory lung disease
- SNO-SP-D is formed in human lung disease and is associated with increased inflammatory function.

Conclusions

Nitrosative and oxidative modifications of SP-D occur in human inflammatory lung disease and these modifications have potentially significant consequences in controlling innate immune function

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