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# LigaTrap Technology mAb Capture and Concentration



IgG-binding peptide  $K_D \sim 10^{-6} M \rightarrow$  Elution at pH 4 Binds all human IgG subclasses Binds animal IgG Safe upon re-use (> 100 cycles)



Binding capacity > 40 mg mAb / mL<sub>R</sub> (RT: 2 min) HCP LRV ~ 2.15 (direct harvest)

HCP LRV: log removal of protein impurities



## LigaGuard Technology mAb Yield and Purity

mAb Yield



mAb Purity

## **GuardMix Technology Global and Single HCP Clearance**

	Low Sa	It Concentratior	ן (50 mM)	High Salt Concentration (150 mM)					
Adsorbent	Low pH (pH 6)	Medium pH (pH 7)	High pH (pH 8)	Low pH (рн 6)	Medium pH (pH 7)	High pH (pH 8)			
PepMix 1	1.9	4.9	3.3	2.0	1.2	0.6			
PepMix 2	1.4	2.2	1.5	2.8	1.1	0.3			
PepMix 3	1.7	1.6	1.9	1.8	1.3	0.8			
PepMix 4	1.4	1.6	1.3	1.8	0.8	0.2			
Capto Q (Commercial)	1.7	2.2	1.2	1.9	1.3	0.7			
Capto Adhere (Commercial)	0.8	0.7	0.8	1.1	0.8	0.4			

C<sub>lgG</sub> ~ 2 - 5 mg/mL

C<sub>HCPs</sub> ~ 0.1 - 0.5 mg/mL X<sub>HCP</sub> ~ 5 - 200 pmol/mL

#### High Binding Selectivity Global & Single HCP Binding

н		Low Salt (20 r				mM NaCl)			н	High Salt		
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	High Risk HCP Impurity	<b>Vi</b>	Vi	Vi	Vib		ð	dhe	<b>Vi</b> N	Vib	ŝ	
		rd 1	rd	rd	rd		apte	٩V	rd[	rd	<u>l</u> d	
		Bua	Bua	gua	Bua		Ű	apt	Bua	Bua	Bua	
	Actin, cytoplasmic 1											
	Alpha-enolase											
	Biglycan											
	Cathepsin B										-	
	Cathepsin D											
	Clusterin											
	Cofilin-1											
	Fructose-bisphosphate aldolase											
	Glutathione S-transferase P											
	Glyceraldehyde phosp. dehydrogenase											
	Heat shock protein											
	Lipoprotein lipase											
	Matrix metalloproteinase-19											
	Metalloproteinase inhibitor 1											
	Nidogen-1											
	Peptidyl-prolyl cis-trans isomerase											
	Peroxiredoxin-1											
	Phosphoglycerate mutase 1											
	Phospholipase B-like protein											
	Protein disulfide-isomerase A6											
	Pyruvate kinase											
	Serine protease HTRA1											
	Sialidase I										_	
	Vimentin											

**Binding Conditions** 

(150 mM NaCl)

### "Problematic" HCPs

Protein		R1R2		Protein		R2	R3
60S acidic ribosomal protein P0				Glutathione S-transferase P			
60S acidic ribosomal protein P2				Glyceraldehyde-3-phopsphate dehydrogenase			
Actin, alpha cardial muscle 1				Heat shock cognate protein			
Actin, cytoplasmic 1				Legumain			
Aldose reductase-related protein 2				Lipoprotein lipase			
Alpha-enolase				Matrix metalloproteinase-19			
Cathepsin B				Metalloproteinase inhibitor 1			
Cathepsin D				Nidogen-1			
Clusterin				Peroxiredoxin-1			
Cofilin-1				Phosphoglycerate mutase 1			
Elongation factor 1-alpha				Phospholipase-B-like protein			
Elongation factor 2				Pyruvate kinase			
ER chaperone BiP				Sialidase 1			
Fructose-biphosphate aldolase				Vimentin			

R1: Co-elute with mAbs product from Protein A

R2: Highly toxic or immunogenic

R3: Trigger mAb instability









**LigaGuard Adsorbent** 

# LigaTrap Technology HCP Clearance and mAb Quality

Human Metalloprot Inh. 1: R > 96.5% **Carboxypeptidase D: R > 98.5% Legumain: undetected Cathepsin B: undetected** > 98.5% Lipoprotein lipase: Cathepsin **D**: R Phospholipase B-like prot.: R > 98.5% **Cathepsin L: undetected Nidogen 1: undetected Cofilin 1:** R > 95% S-transferase: Peroxiredoxin 1: undetected Glutathione **Peroxiredoxin 4: R > 98.5%** undetected **Enolase 1: R > 99.5% Global Yield** ~ 70 - 74 % **Purity > 99.9%** CHO HCP LRV ~ 3.02 (ELISA)

LigaTrap

