

**Molecular characterization of the RNA-protein complex directing -2/-1 programmed ribosomal frameshifting during arterivirus replicase expression**

Ankoo Patel, Emmely E. Treffers, Markus Meier, Trushar R. Patel, Jörg Stetefeld, Eric J. Snijder, Brian L. Mark<sup>1\*</sup>

**Supporting Information**

*Table S1. Primers used for nsp1 $\beta$  and PCBP2 mutations for EMSAs*

<b>Oligo Name</b>	<b>Sequence (5' to 3')</b>
nsp1 $\beta$ Y131A Forward	CCAACGCAGGCTTCAAGTCCG
nsp1 $\beta$ Y131A Reverse	AGCGCCTTACCATGCACGCCCC
nsp1 $\beta$ R135A Forward	CAAGTCCGCGGCATGCGTGC
nsp1 $\beta$ R135A Reverse	AGCGCGCGTTGGAGGTA CTTAC
PCBP2 R40A Forward	GGAGAGTGGTGCACGTATCAAC
PCBP2 R40A Reverse	TCCGCCATCTTCTTAACTGATTC
PCBP2 R57A Forward	TATCACTTTGGCTGGACCCAC
PCBP2 R57A Reverse	ATCGCCTCAGGACAATTCCCTTC
PCBP2 N325D Forward	CAAAATTGCGGATCCAGTGGAAGG
PCBP2 N325D Reverse	ATCTGCGCCCCAGACATC

Table S2. Primers used for frameshifting assays

<b>Oligo Name</b>	<b>Sequence (5' to 3')</b>	<b>Purpose</b>
EUnsp1b-G129A-fw	GGGCGTGCATGCTAAGTACCTCCAACGCAG	Site directed mutagenesis G129A
EUnsp1b-G129A-rev	TGGAGGTACTTAGCATGCACGCCCACTTGG	Site directed mutagenesis G129A
EUnsp1b-K130A-fw	GGCGTGCATGGTGCUTACCTCCAACGCAG	Site directed mutagenesis K130A
EUnsp1b-K130A-rev	TGCGTTGGAGGTAAGCACCATGCACGCC	Site directed mutagenesis K130A
EUnsp1b-Y131A-fw	TGCATGGTAAGGCCCTCCAACGCAGGCTTC	Site directed mutagenesis Y131A
EUnsp1b-Y131A-rev	TGCGTTGGAGGGCCTTACCATGCACGCCCC	Site directed mutagenesis Y131A
EUnsp1b-L132A-fw	GCATGGTAAGTACGCCCAACGCAGGCTTCAAGT C	Site directed mutagenesis L132A
EUnsp1b-L132A-rev	GAAGCCTGCGTTGGGCGTACTTACCATGCACG	Site directed mutagenesis L132A
EUnsp1b-Q133A-fw	TAAGTACCTCGCACGCAGGCTTCAAGTCCG	Site directed mutagenesis Q133A
EUnsp1b-Q133A-rev	GAAGCCTGCGTGCGAGGTAAGTACTTACCATGCAC	Site directed mutagenesis Q133A

EUnsp1b-L136A-fw	CCAACGCAGGGCTCAAGTCCGCGGCATGC	Site directed mutagenesis L136A
EUnsp1b-L136A-rev	CATGCCGCGGACTTGAGCCCTGCGTTGGAG	Site directed mutagenesis L136A
EUnsp1b-Q137A-fw	TCCAACGCAGGCTTGCAGTCCGCGGCATGC	Site directed mutagenesis Q137A
EUnsp1b-Q137A-rev	CCGCGGACTGCAAGCCTGCGTTGGAGGTAC	Site directed mutagenesis Q137A
EUnsp1b-V138A-fw	GCAGGCTTCAAGCCCGCGGCATGCGTGCTG	Site directed mutagenesis V138A
EUnsp1b-V138A-rev	GCATGCCGCGGGCTTGAAGCCTGCGTTGGAG	Site directed mutagenesis V138A
EUnsp1b-R139A-fw	CTTCAAGTCGCTGGCATGCGTGCTGTGGTC	Site directed mutagenesis R139A
EUnsp1b-R139A-rev	CACAGCACGCATGCCAGCGACTTGAAGCC	Site directed mutagenesis R139A
EUnsp1b-G140A-fw	TTCAAGTCCGCGCCATGCGTGCTGTGGTCG	Site directed mutagenesis G140A
EUnsp1b-G140A-rev	CAGCACGCATGGCGCGGACTTGAAGCCTGC	Site directed mutagenesis G140A

EUnsp1b-M141A-fw	AGTCCGCGGCGCTCGTGCTGTGGTCGATCC	Site directed mutagenesis M141A
EUnsp1b-M141A-rev	CCACAGCACGAGCGCCGCGGACTTGAAGCC	Site directed mutagenesis M141A
EUnsp1b-R142A-fw	CCGCGGCATGGCTGCTGTGGTCGATCCTG	Site directed mutagenesis R142A
EUnsp1b-R142A-rev	CGACCACAGCAGCCATGCCGCGGACTTGAAG	Site directed mutagenesis R142A
EUnsp1b-fw	ACGAGAATTCCCATGGATGTCTGACGTTTACAG GTGGAAG	Amplify EUnsp1b and introduce upstream EcoRI and NcoI sites
3xFLAG-EUnsp1b-fw	GCCGAATTCCCATGGACTACAAAGACCATGACG GTGATTATAAAGATCATGACATCGATTACAAGG ATGACGATGACAAGTCTGACGTTTACAGGTGGA AG	Amplify EUnsp1b and introduce upstream 3xFLAG tag and NcoI site
EUnsp1b-rev	ATCACCTGCAGGGCGGCCGCTCGAGTCAGCCA TACCACTTATGTG	Amplify EUnsp1b and introduce downstream XhoI, NotI and SbfI restriction sites
pL-seq	GATCTGATCTGGGGCCTC	Sequencing pL vector



## Part 2

# Biophysical characterization

<i>Parameters</i>	
<b>Solvent</b> 1x PBS (pH 7.4 @ 25°C), 100 mM KCl, 5% glycerol	2.027 mM sodium dihydrogen phosphate 7.973 mM disodium hydrogen phosphate 138 mM sodium chloride 102.7 mM potassium chloride ~684 mM glycerol
<b>Solvent density<sup>1</sup> <math>\rho</math> (g/cm<sup>3</sup>)</b>	1.0354 ± 0.0005
<b>Buffer viscosity<sup>1</sup> <math>\eta</math> (P)</b>	0.01235 ± 0.0002
<b>Temperature T (°C)</b>	20.0
<b>his<sub>6</sub>-PCBP2 partial specific volume<sup>1</sup> <math>\bar{v}</math> (cm<sup>3</sup>/g)</b>	0.73318
<b>his<sub>6</sub>-PCBP2 formula mass<sup>2</sup> <math>M_w</math> (kDa)</b>	39749.32
<b>his<sub>6</sub>-PCBP2 absorption coefficient<sup>2</sup> <math>\epsilon_{280}</math> (M<sup>-1</sup> cm<sup>-1</sup>)</b>	14815
<b>PRRSV nsp1<math>\beta</math> partial specific volume<sup>1</sup> <math>\bar{v}</math> (cm<sup>3</sup>/g)</b>	0.72777
<b>PRRSV nsp1<math>\beta</math> formula mass<sup>2</sup> <math>M_w</math> (kDa)</b>	23786.87
<b>PRRSV nsp1<math>\beta</math> absorption coefficient<sup>2</sup> <math>\epsilon_{280}</math> (M<sup>-1</sup> cm<sup>-1</sup>)</b>	46200
<b>34nt ssRNA partial specific volume <math>\bar{v}</math> (cm<sup>3</sup>/g)</b>	dependent on cation concentration expected range 0.50 - 0.65
<b>34nt ssRNA formula mass<sup>3</sup> <math>M_w</math> (kDa)</b>	10688.3
<b>34nt ssRNA absorption coefficient<sup>3</sup> <math>\epsilon_{260}</math> (M<sup>-1</sup> cm<sup>-1</sup>)</b>	323000

<sup>1</sup>Sednterp, <sup>2</sup>ProtParam, <sup>3</sup>IDTDNA

Table S3

# Section 1: Characterization of 34 nt ssRNA by Sedimentation Velocity

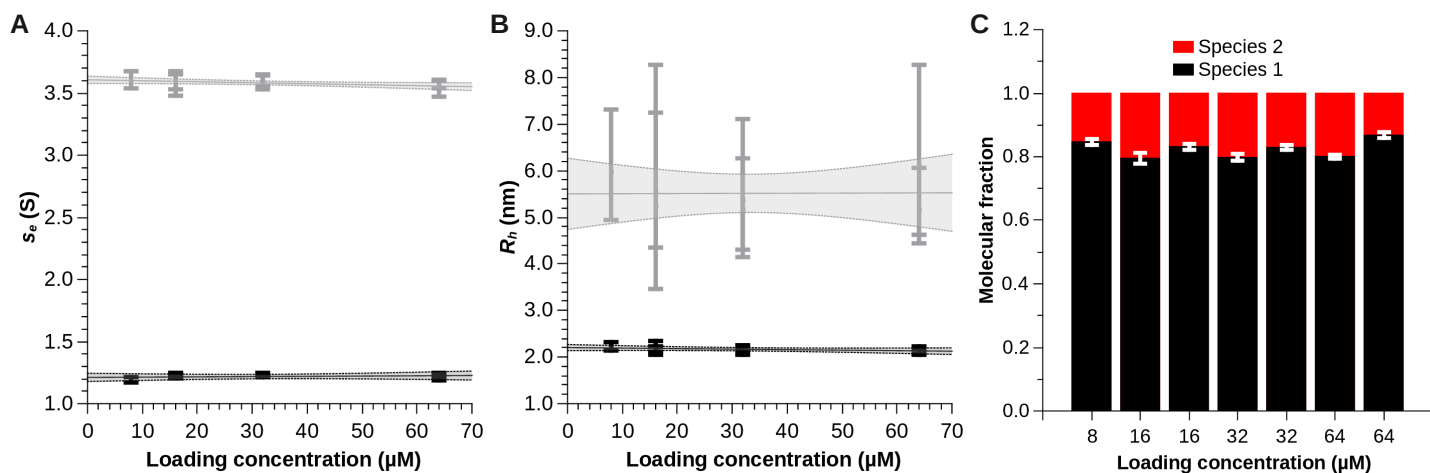


Fig. S1

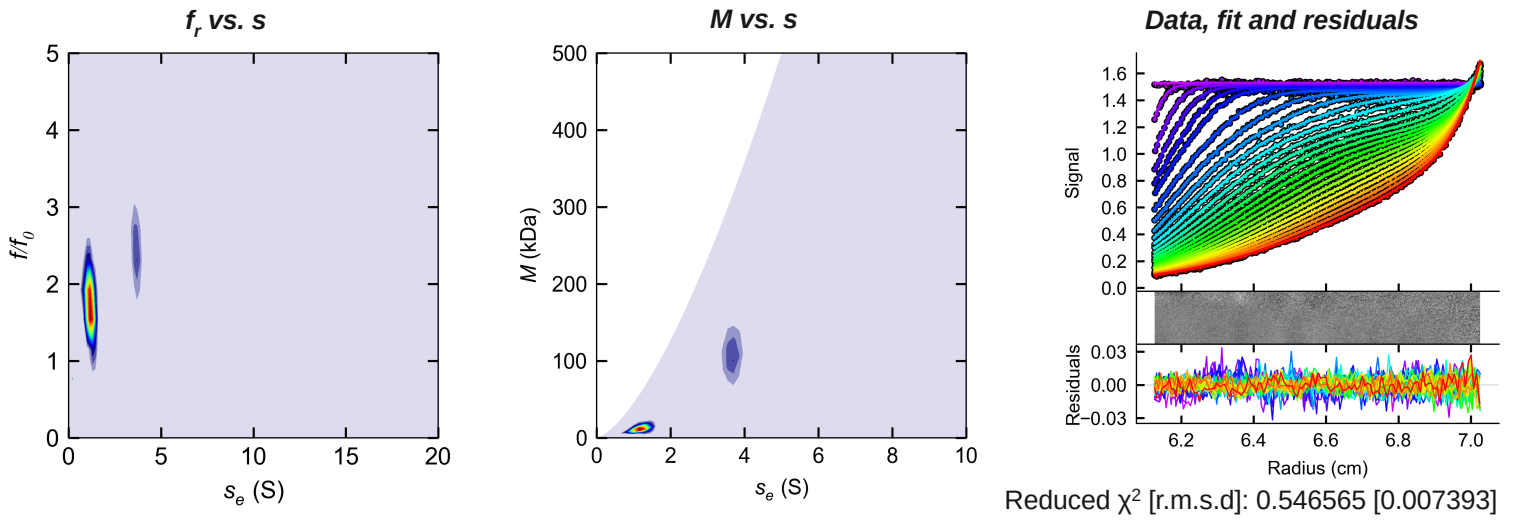
## Results

<i>Species 1</i>	
Sedimentation coefficient $s_e^0$	$1.21 \pm 0.04$ S
Diffusion coefficient $D_e^0$	$7.9 \pm 0.3 \cdot 10^{-7}$ cm <sup>2</sup> /s
Partial specific volume that yields correct mass (10.7 kDa)	0.62763 [0.60703 - 0.64704] cm <sup>3</sup> /g
Sedimentation coefficient $s_{20^\circ\text{C}, w}^0$	1.59 [1.54 - 1.65] S
Hydrodynamic radius $R_h^0$	2.20 [2.12 - 2.29] nm
<i>Species 2</i>	
Sedimentation coefficient $s_e^0$	$3.61 \pm 0.03$ S
Diffusion coefficient $D_e^0$	$3.2 \pm 0.5 \cdot 10^{-7}$ cm <sup>2</sup> /s
Sedimentation coefficient $s_{20^\circ\text{C}, w}^0$	4.74 [4.65 - 4.83] S
Hydrodynamic radius $R_h^0$	5.49 [4.84 - 6.35] nm
Molecular mass if RNA; would contain 7-8 strands	79.3 [68.2 - 90.5] kDa

Uncertainties are given as 95% confidence intervals

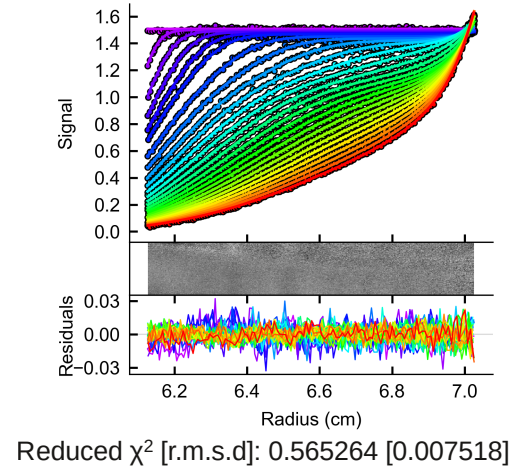
Table S4

8  $\mu\text{M}$  ssRNA monitored by absorbance optics  
30000 rpm,  $\lambda = 278 \text{ nm}$



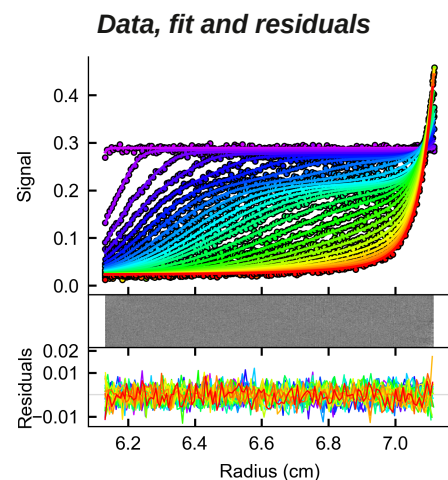
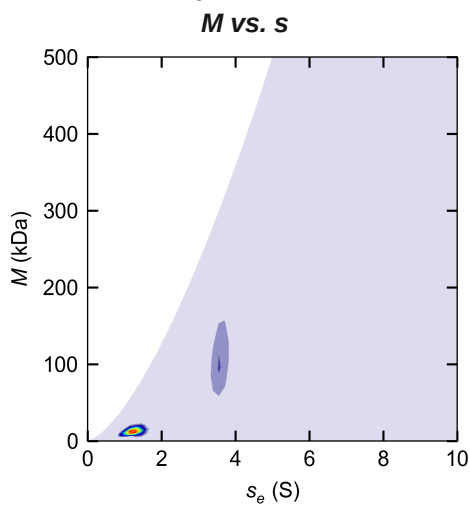
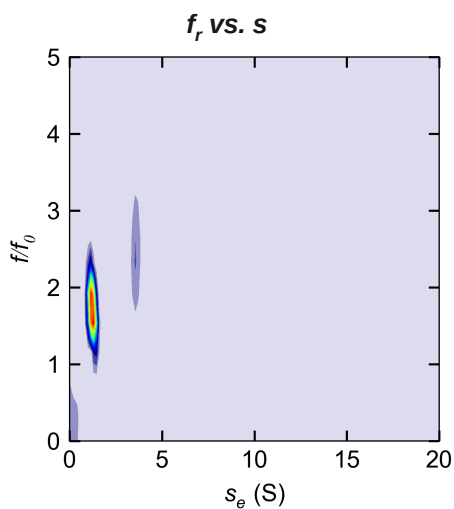
Species Analysis (30000 rpm,  $\lambda = 278 \text{ nm}$ , exp. # 1)

Parameter	Value	95% confidence interval
<b>Species 1</b>		
Sedimentation coefficient $s_e$ (S)	1.191056	1.1725 - 1.2098
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	7.826990	7.5122 - 8.1480
<b>Species 2</b>		
Sedimentation coefficient $s_e$ (S)	3.605013	3.5373 - 3.6741
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	2.905149	2.3804 - 3.5159
Fraction (signal)	0.153196	0.1439 - 0.1631
<b>Common</b>		
Total concentration (signal)	1.508362	1.4841 - 1.5328
Meniscus position (cm)	6.081005	6.0774 - 6.0844
Bottom position (cm)	7.237708	7.2307 - 7.2447
Reduced $\chi^2$ of fit [rmsd]	0.5652639	
R.m.s.d. of fit	0.007518403	

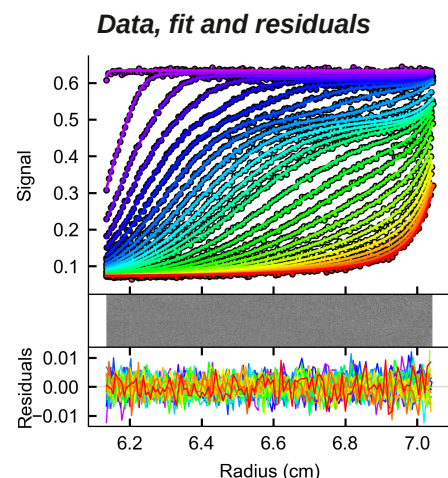
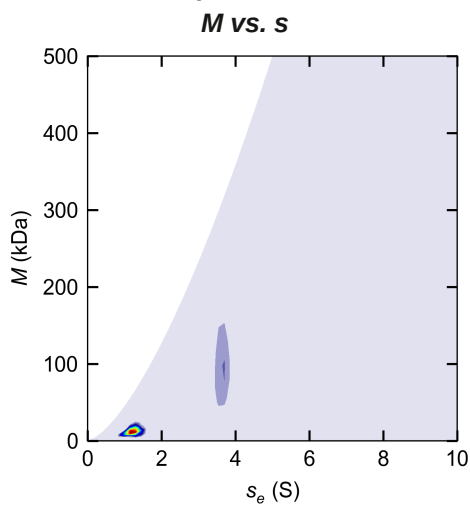
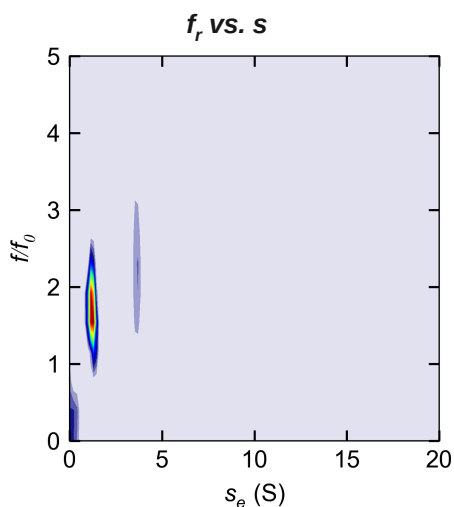


16  $\mu\text{M}$  ssRNA monitored by absorbance optics

42000 rpm,  $\lambda = 294 \text{ nm}$



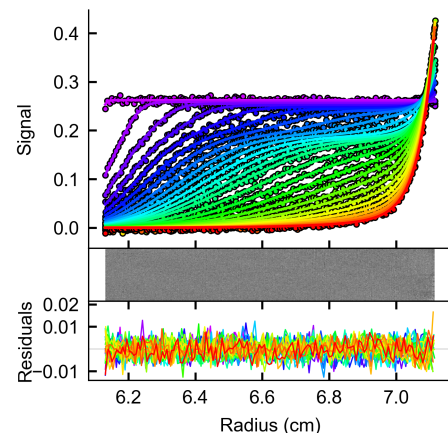
Reduced  $\chi^2$  [r.m.s.d]: 0.120617 [0.003473]



Reduced  $\chi^2$  [r.m.s.d]: 0.112694 [0.003357]

Species Analysis (42000 rpm,  $\lambda = 294 \text{ nm}$ , exp. # 1)

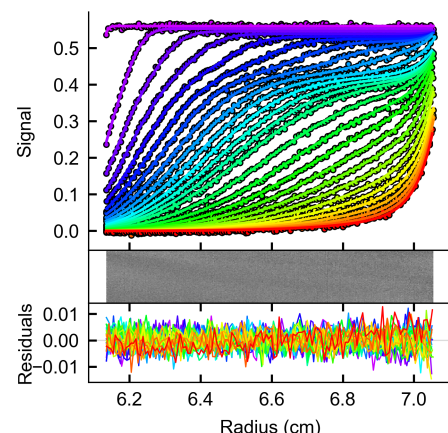
Parameter	Value	95% confidence interval
<b>Species 1</b>		
Sedimentation coefficient $s_e$ (S)	1.229725	1.2078 - 1.2518
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	7.837109	7.4467 - 8.2514
<b>Species 2</b>		
Sedimentation coefficient $s_e$ (S)	3.579254	3.4808 - 3.6784
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	3.316965	2.1005 - 5.0301
Fraction (signal)	0.204758	0.1884 - 0.2216
<b>Common</b>		
Total concentration (signal)	0.2623926	0.2545 - 0.2705
Meniscus position (cm)	6.113078	6.1073 - 6.1188
Bottom position (cm)	7.227485	7.2230 - 7.2323
Reduced $\chi^2$ of fit [rmsd]	0.1228484	
R.m.s.d. of fit	0.003504974	



Reduced  $\chi^2$  [r.m.s.d]: 0.122848 [0.003505]

Species Analysis (42000 rpm,  $\lambda = 291 \text{ nm}$ , exp. # 2)

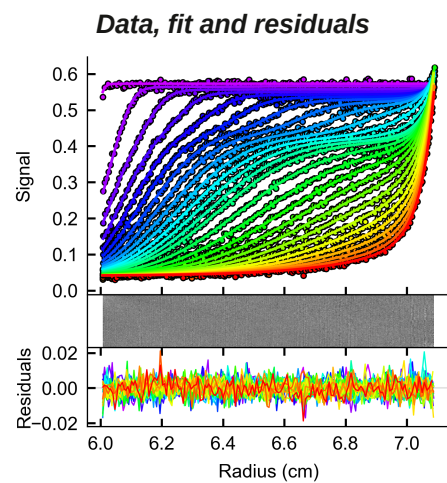
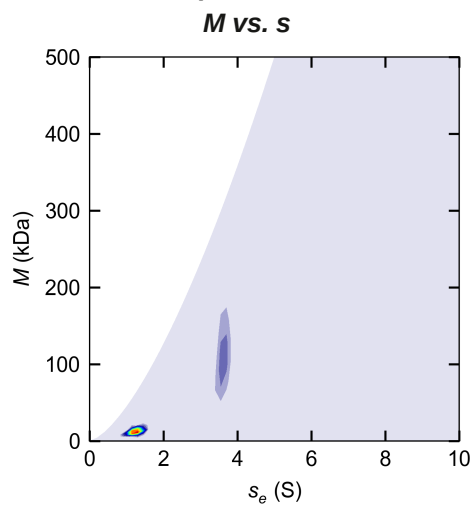
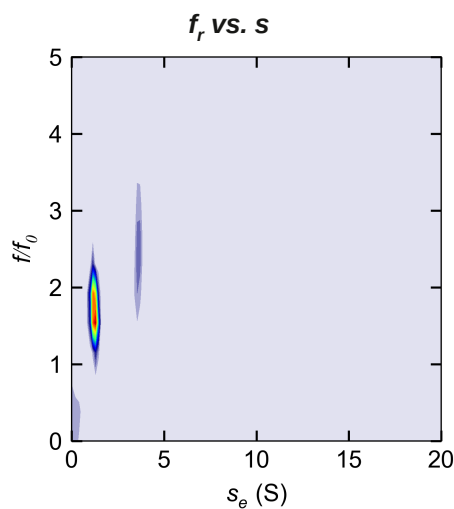
Parameter	Value	95% confidence interval
<b>Species 1</b>		
Sedimentation coefficient $s_e$ (S)	1.215424	1.2044 - 1.2265
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	8.173451	7.8048 - 8.4944
<b>Species 2</b>		
Sedimentation coefficient $s_e$ (S)	3.590108	3.5299 - 3.6475
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	3.140034	2.3970 - 4.0048
Fraction (signal)	0.168034	0.1588 - 0.1769
<b>Common</b>		
Total concentration (signal)	0.5605898	0.1588 - 0.1769
Meniscus position (cm)	6.117462	6.1147 - 6.1205
Bottom position (cm)	7.234521	7.2271 - 7.2411
Reduced $\chi^2$ of fit [rmsd]	0.1260963	
R.m.s.d. of fit	0.003551004	



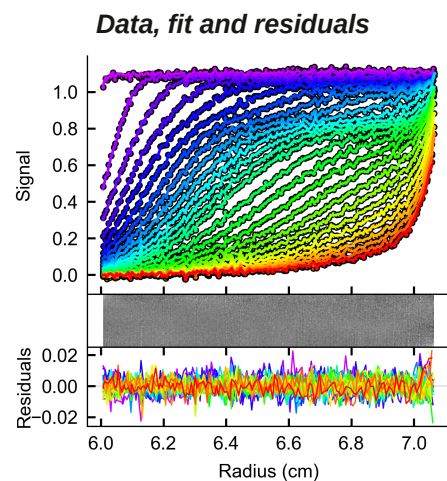
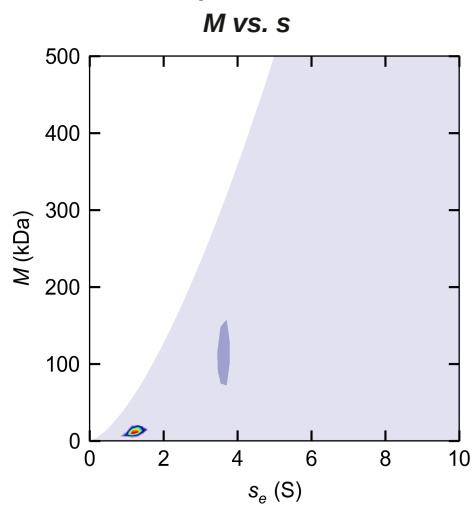
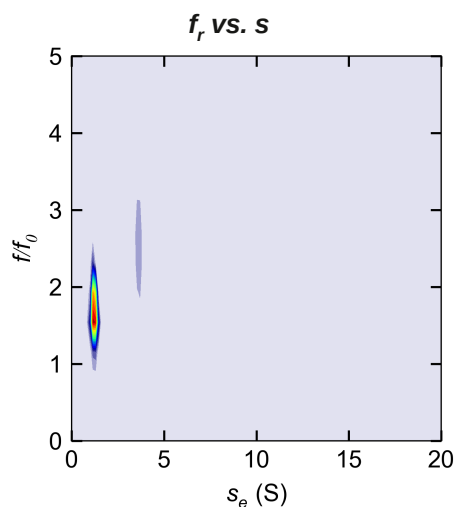
Reduced  $\chi^2$  [r.m.s.d]: 0.126096 [0.003551]

32  $\mu\text{M}$  ssRNA monitored by absorbance optics

42000 rpm,  $\lambda = 294 \text{ nm}$



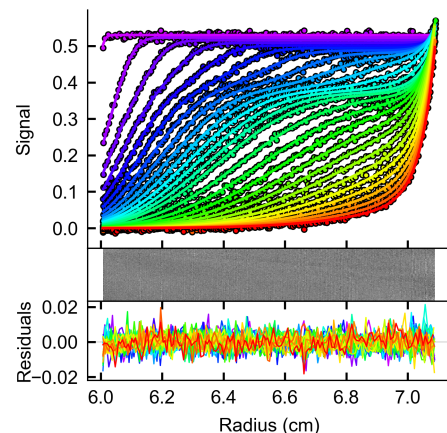
Reduced  $\chi^2$  [r.m.s.d]: 0.191756 [0.004379]



Reduced  $\chi^2$  [r.m.s.d]: 0.259183 [0.005091]

Species Analysis (42000 rpm,  $\lambda = 294 \text{ nm}$ , exp. # 1)

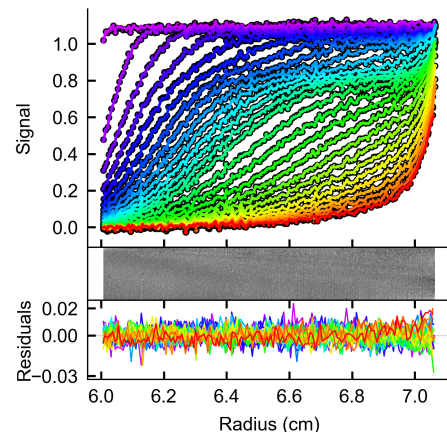
Parameter	Value	95% confidence interval
<b>Species 1</b>		
Sedimentation coefficient $s_e$ (S)	1.232830	1.2201 - 1.2468
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	8.121437	7.7671 - 8.5080
<b>Species 2</b>		
Sedimentation coefficient $s_e$ (S)	3.585519	3.5308 - 3.6467
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	3.231844	2.4451 - 4.1974
Fraction (signal)	0.202133	0.1919 - 0.2126
<b>Common</b>		
Total concentration (signal)	0.5306498	0.5205 - 0.5417
Meniscus position (cm)	5.986057	5.9823 - 5.9896
Bottom position (cm)	7.236567	7.2309 - 7.2428
Reduced $\chi^2$ of fit [rmsd]	0.1983076	
R.m.s.d. of fit	0.004453174	



Reduced  $\chi^2$  [r.m.s.d]: 0.198308 [0.004453]

Species Analysis (42000 rpm,  $\lambda = 291 \text{ nm}$ , exp. # 2)

Parameter	Value	95% confidence interval
<b>Species 1</b>		
Sedimentation coefficient $s_e$ (S)	1.220184	1.2127 - 1.2280
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	7.938753	7.7172 - 8.1994
<b>Species 2</b>		
Sedimentation coefficient $s_e$ (S)	3.596446	3.5560 - 3.6386
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	3.366156	2.7780 - 4.0454
Fraction (signal)	0.170676	0.1640 - 0.1770
<b>Common</b>		
Total concentration (signal)	1.088870	1.0756 - 1.1033
Meniscus position (cm)	5.989551	5.9874 - 5.9917
Bottom position (cm)	7.235867	7.2314 - 7.2412
Reduced $\chi^2$ of fit [rmsd]	0.2852372	
R.m.s.d. of fit	0.005340760	

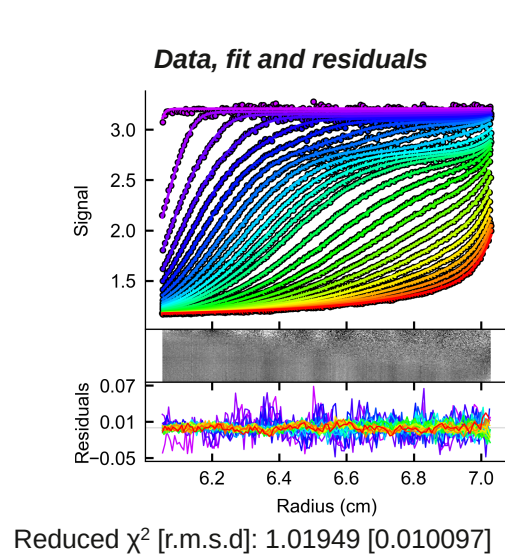
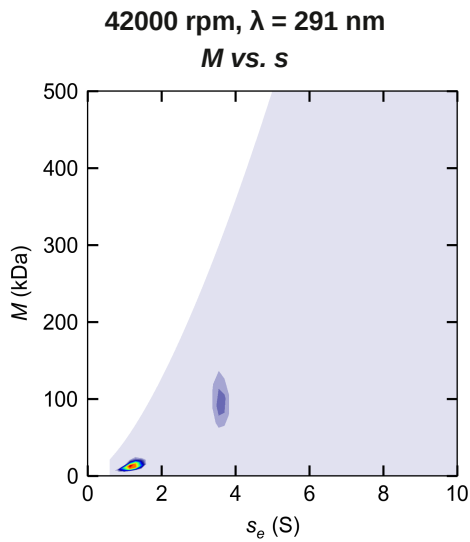
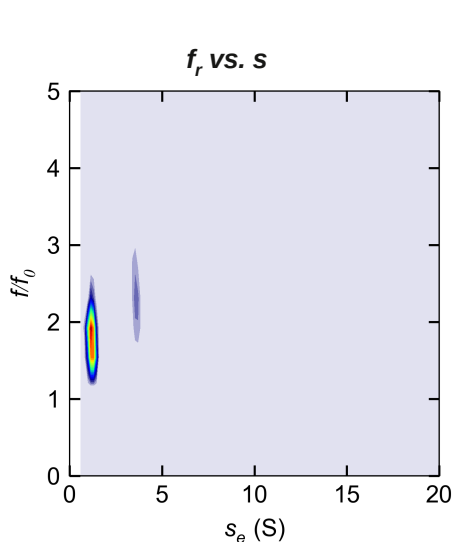
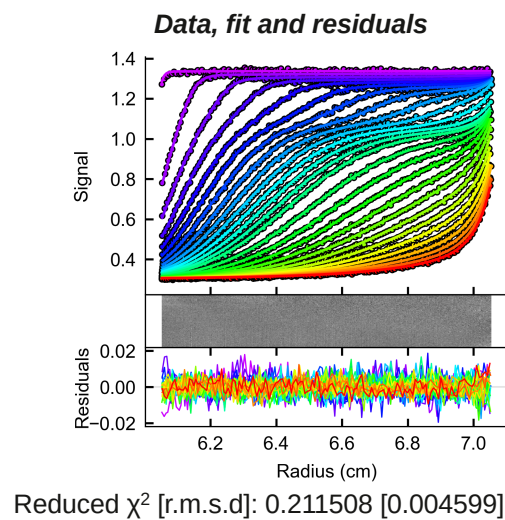
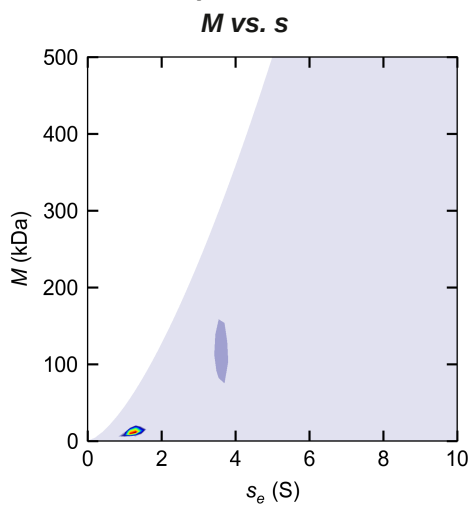
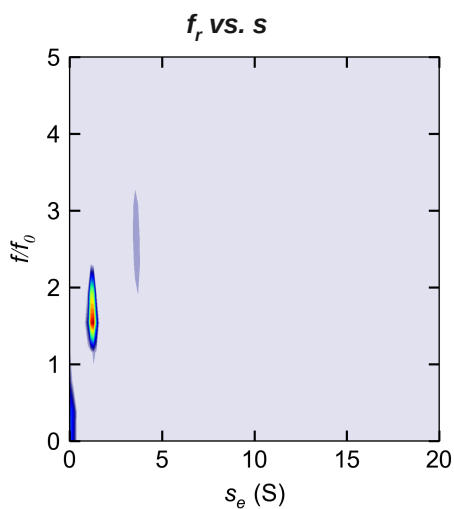


Reduced  $\chi^2$  [r.m.s.d]: 0.285237 [0.005341]



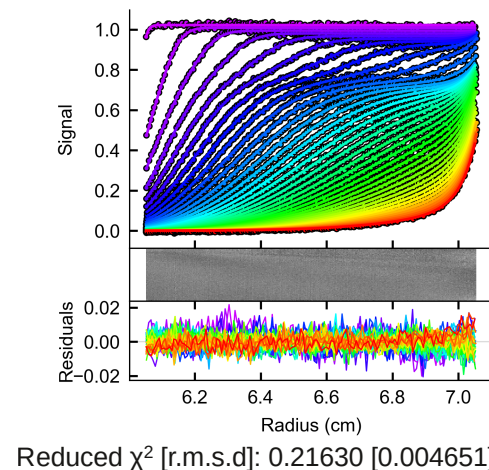
64  $\mu\text{M}$  ssRNA monitored by absorbance optics

42000 rpm,  $\lambda = 294 \text{ nm}$



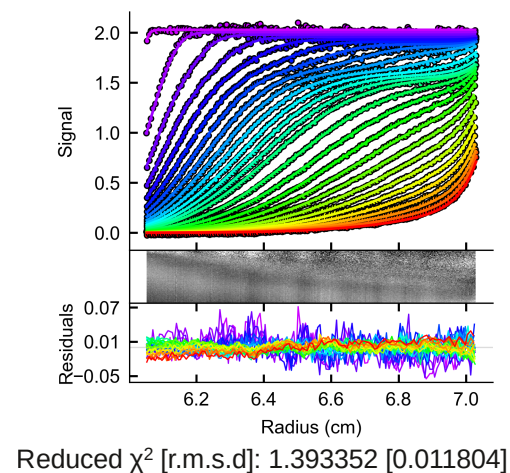
Species Analysis (42000 rpm,  $\lambda = 294 \text{ nm}$ , exp. # 1)

Parameter	Value	95% confidence interval
<b>Species 1</b>		
Sedimentation coefficient $s_e$ (S)	1.239898	1.2324 - 1.2474
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	8.170592	7.9399 - 8.4006
<b>Species 2</b>		
Sedimentation coefficient $s_e$ (S)	3.572412	3.5362 - 3.6070
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	3.355237	2.8711 - 3.9210
Fraction (signal)	0.198935	0.1930 - 0.2052
<b>Common</b>		
Total concentration (signal)	1.024325	1.0124 - 1.0365
Meniscus position (cm)	6.025673	6.0235 - 6.0279
Bottom position (cm)	7.241323	7.2362 - 7.2464
Reduced $\chi^2$ of fit [rmsd]	0.2163596	
R.m.s.d. of fit	0.004651447	



Species Analysis (42000 rpm,  $\lambda 291 \text{ nm}$ , exp. # 2)

Parameter	Value	95% confidence interval
<b>Species 1</b>		
Sedimentation coefficient $s_e$ (S)	1.201514	1.1918 - 1.2113
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	8.107909	7.7838 - 8.4409
<b>Species 2</b>		
Sedimentation coefficient $s_e$ (S)	3.534074	3.4728 - 3.5943
Diffusion coefficient $D_e$ ( $\cdot 10^{-7} \text{ cm}^2/\text{s}$ )	2.847448	2.1050 - 3.7689
Fraction (signal)	0.132244	0.1238 - 0.1408
<b>Common</b>		
Total concentration (signal)	2.030704	1.9974 - 2.0654
Meniscus position (cm)	6.027494	6.0247 - 6.0302
Bottom position (cm)	7.245158	7.2363 - 7.2544
Reduced $\chi^2$ of fit [rmsd]	1.393352	
R.m.s.d. of fit	0.011804033	

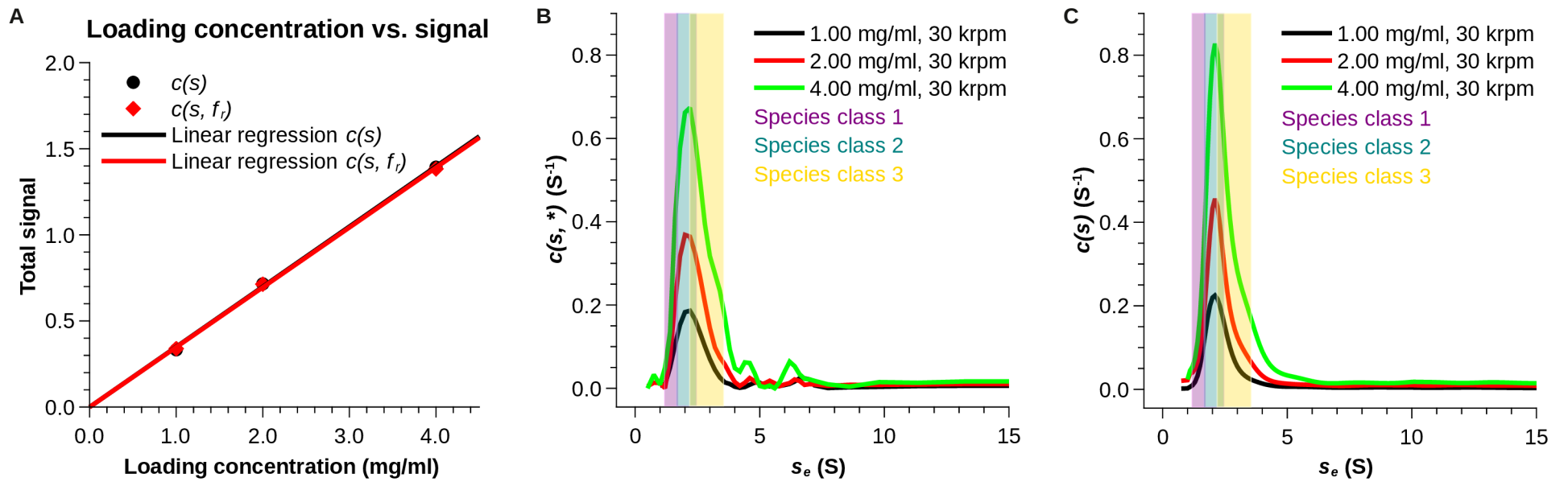


## **Section 2:**

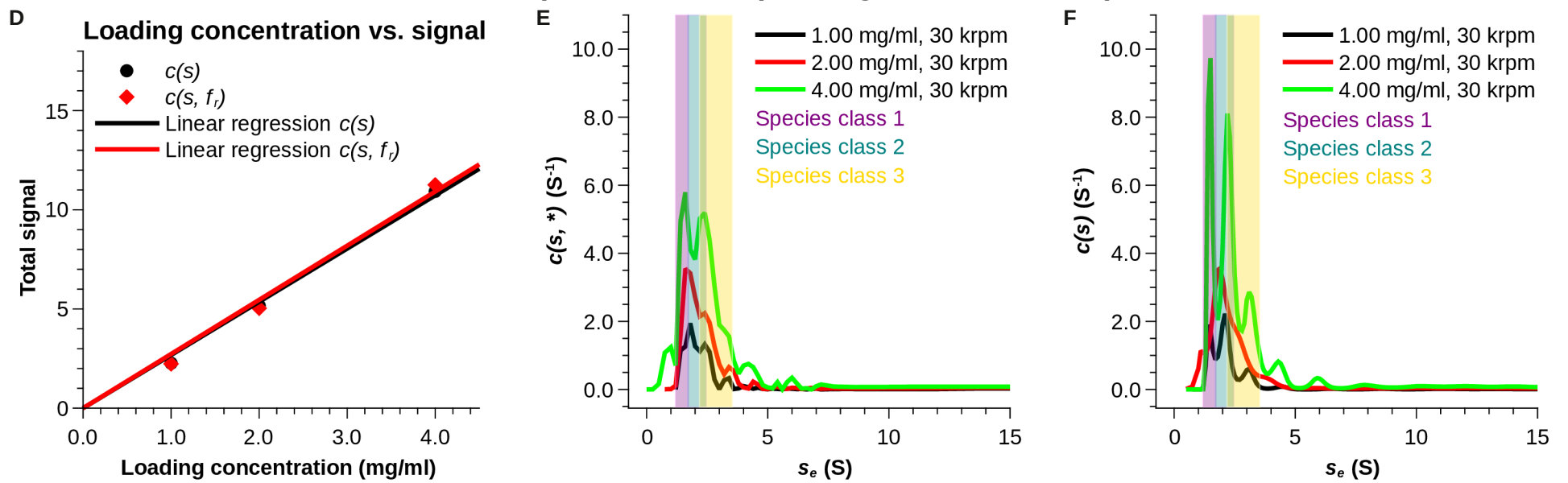
# **Characterization of Human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA Complex by Sedimentation Velocity**

# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

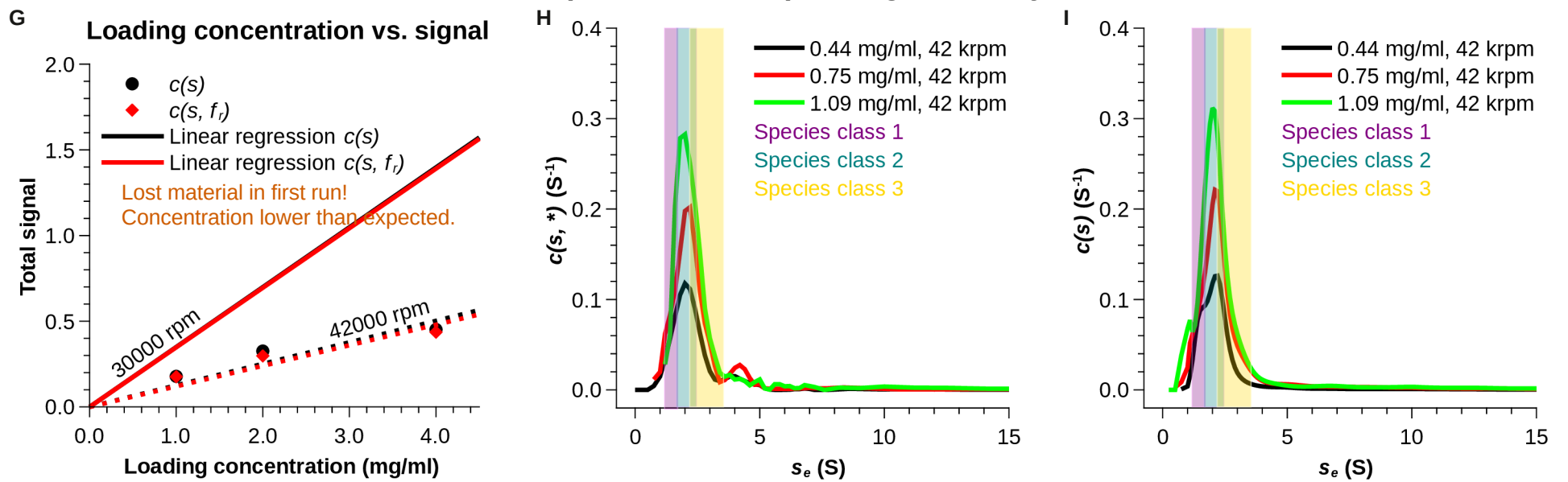
Absorbance optics @ 30000 rpm - Signal mainly from RNA



Interference optics @ 30000 rpm - Signal from all components



Absorbance optics @ 42000 rpm - Signal mainly from RNA



Interference optics @ 42000 rpm - Signal from all components

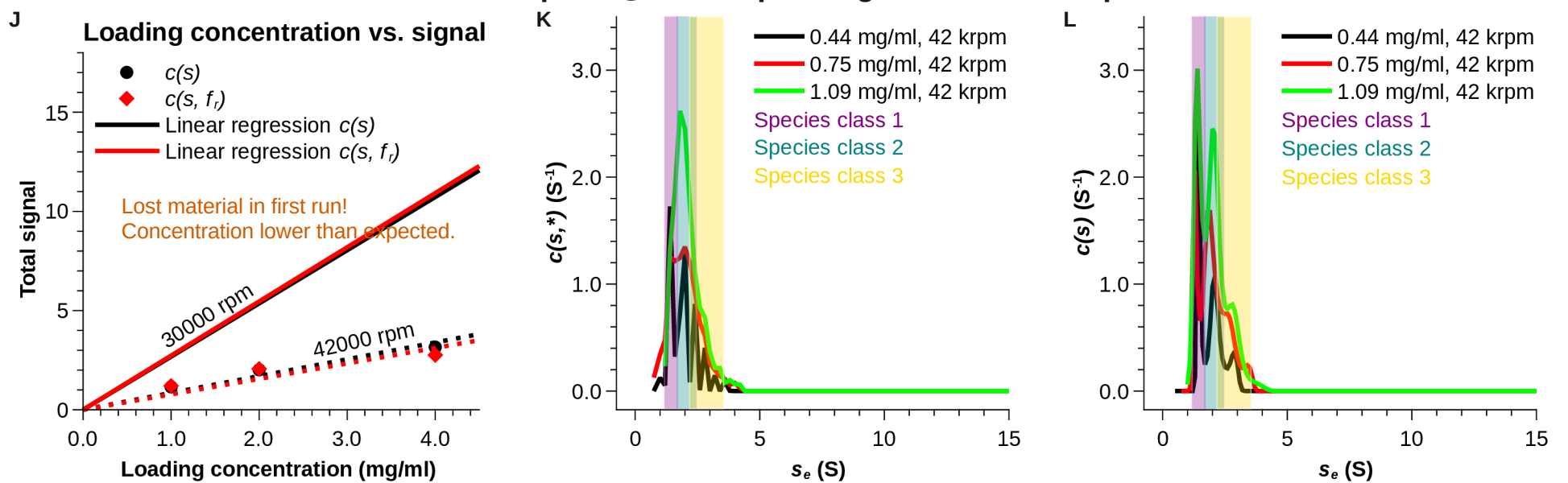


Fig. S3



## Species classes

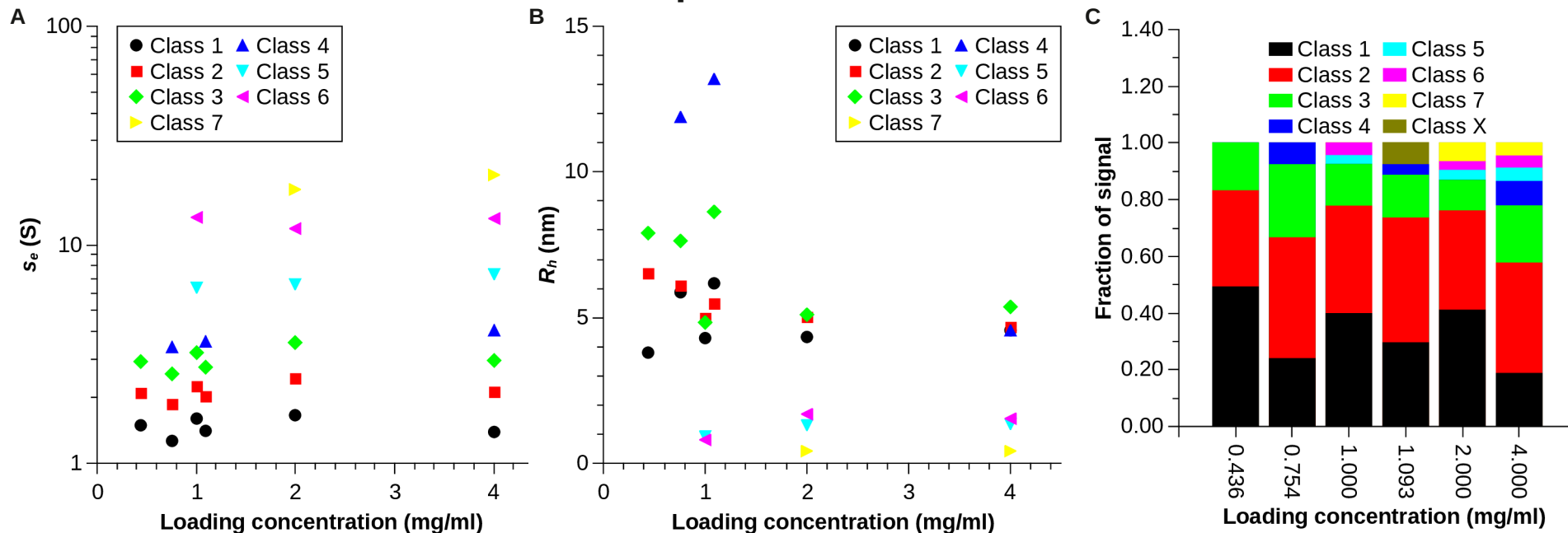


Fig. S4

### Results

<i>Species class 1</i>	
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}^0$	2.02 [1.61 - 2.43] S
Diffusion coefficient $D_{20^{\circ}\text{C}, w}^0$	$4.52 [2.76 - 6.28] \cdot 10^{-7} \text{ cm}^2/\text{s}$
Hydrodynamic radius $R_h^0$	4.74 [3.41 - 7.74] nm
<i>Species class 2</i>	
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}^0$	2.84 [2.28 - 3.41] S
Diffusion coefficient $D_{20^{\circ}\text{C}, w}^0$	$3.51 [2.87 - 4.15] \cdot 10^{-7} \text{ cm}^2/\text{s}$
Hydrodynamic radius $R_h^0$	$6.11 \pm [5.17 - 7.46] \text{ nm}$
<i>Species class 3</i>	
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}^0$	3.96 [2.97 - 4.94] S
Diffusion coefficient $D_{20^{\circ}\text{C}, w}^0$	$2.92 [1.37 - 4.47] \cdot 10^{-7} \text{ cm}^2/\text{s}$
Hydrodynamic radius $R_h^0$	7.34 [4.79 - 15.61] nm

Table S5

# Results

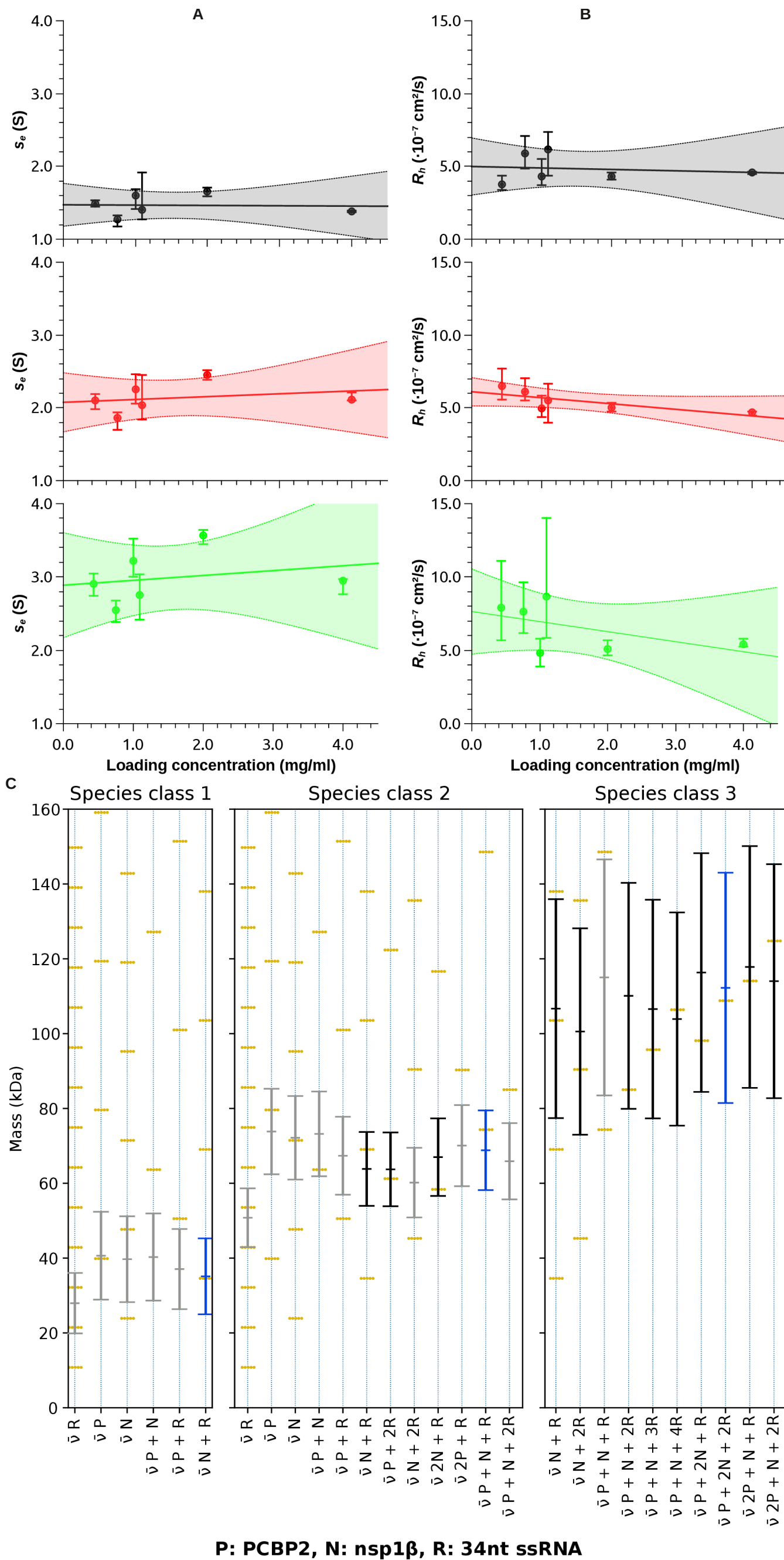
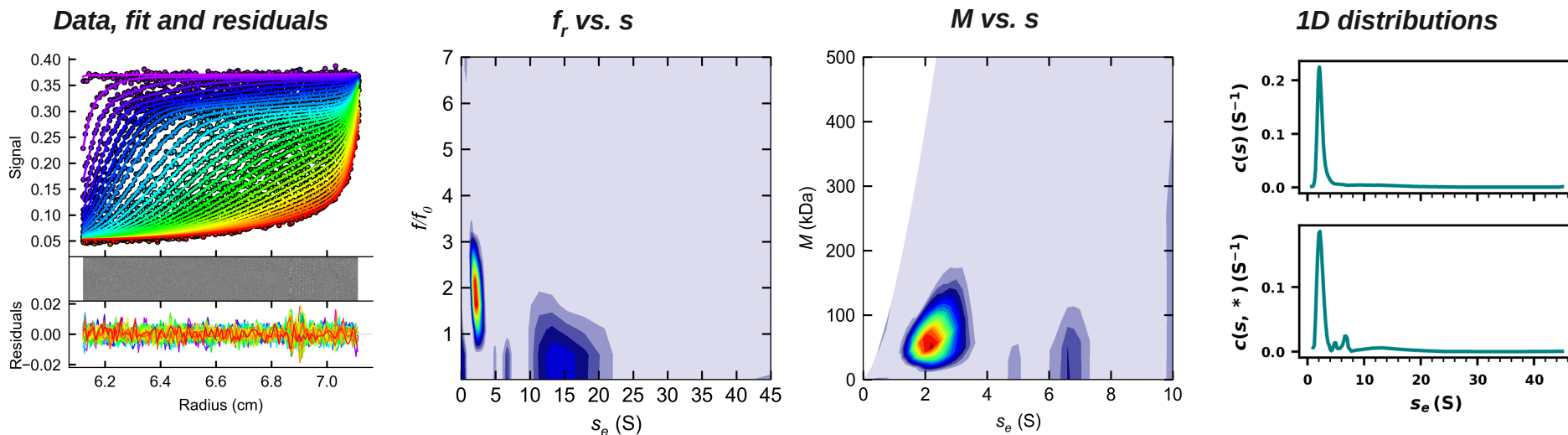


Fig. S5

# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Absorbance optics @ 30000 rpm - loading concentration 1.0 mg/ml

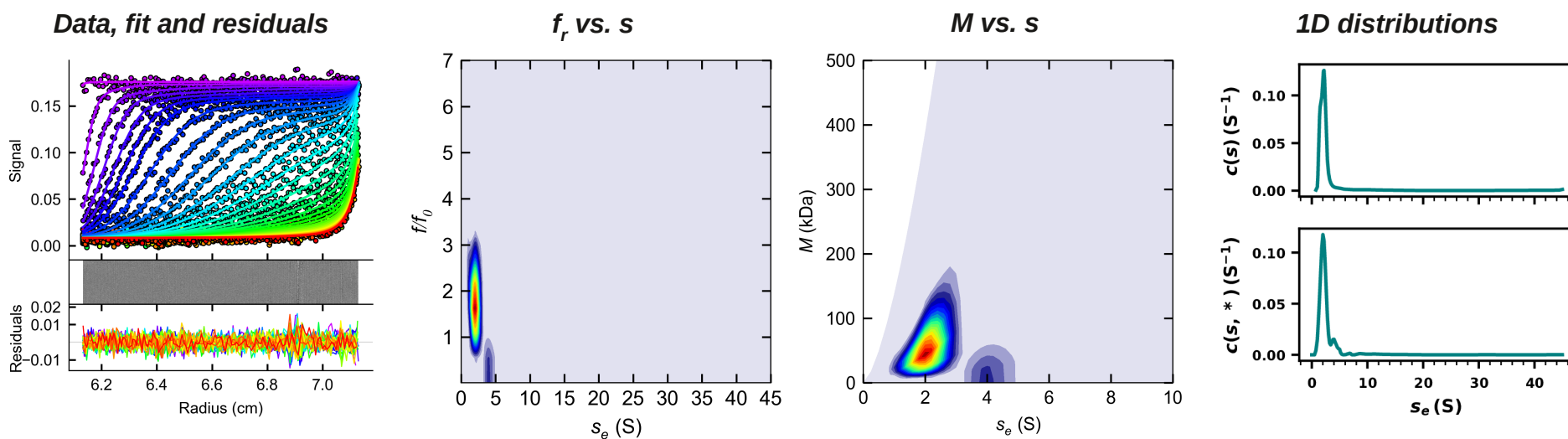
$c(s, f_r)$  analysis



Reduced  $\chi^2$  [r.m.s.d]: [0.003905]

Absorbance optics @ 42000 rpm - actual concentration 0.44 mg/ml

$c(s, f_r)$  analysis



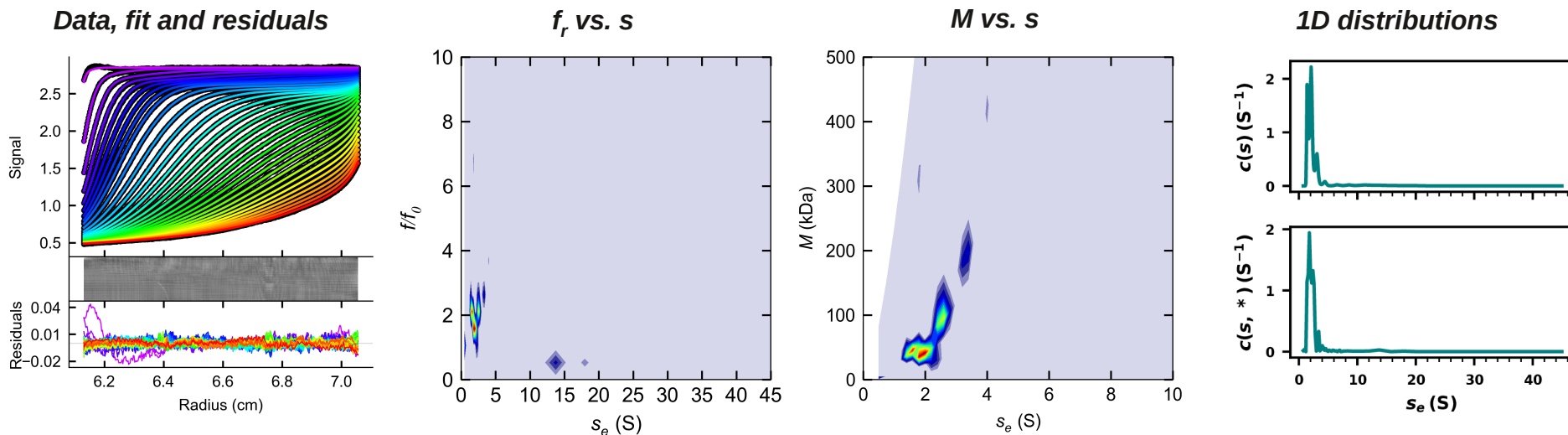
Reduced  $\chi^2$  [r.m.s.d]: [0.003341]

Fig. S6A

# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Interference optics @ 30000 rpm - loading concentration 1.0 mg/ml

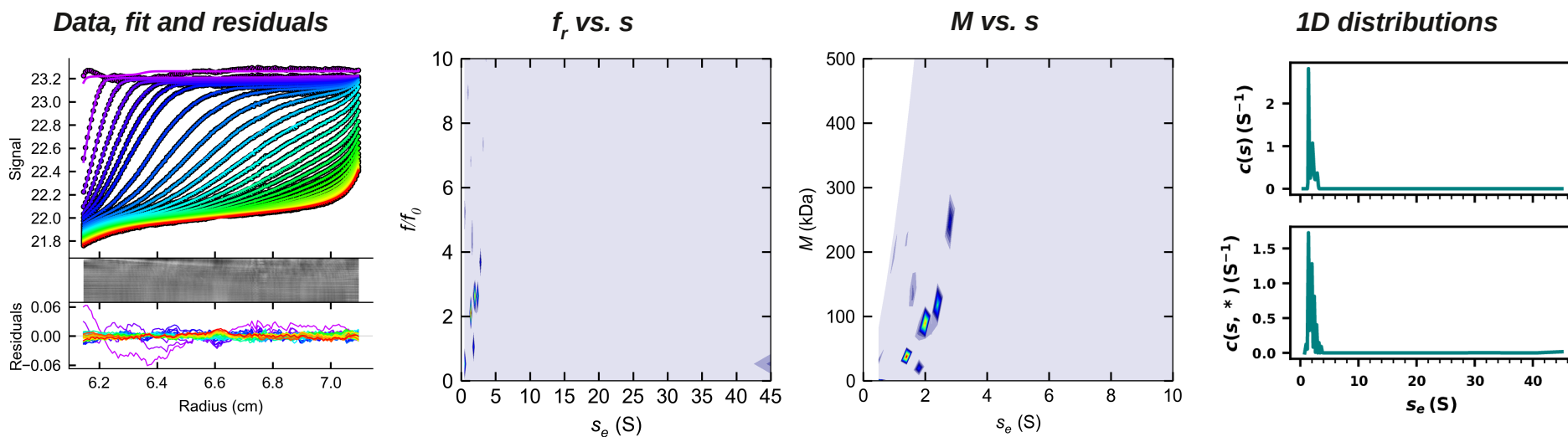
$c(s, f_r)$  analysis



Reduced  $\chi^2$  [r.m.s.d]: [0.003815]

Interference optics @ 42000 rpm - actual concentration 0.44 mg/ml

$c(s, f_r)$  analysis



Reduced  $\chi^2$  [r.m.s.d]: [0.006460]

Fig. S6B

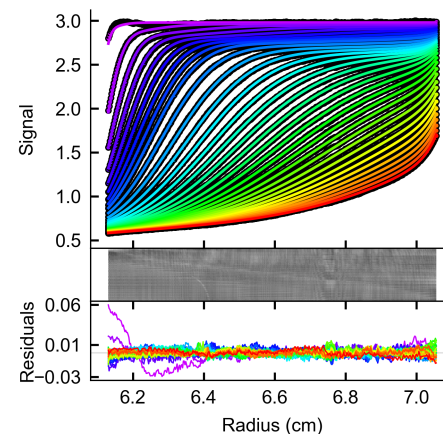
# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Species Analysis (1.00 mg/ml, 30000 rpm, i. o.)

Parameter	Value	95% confidence interval
<b>Species # 1 (class 1)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	2.185538	1.9320 - 2.3175
Molecular mass <sup>1</sup> $M$ (Da)	39429.258626	31975.1983 - 43851.1943
Concentration (signal)	0.889672	
<b>Species # 2 (class 2)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	3.086904	2.8131 - 3.3679
Molecular mass <sup>1</sup> $M$ (Da)	64508.620238	56946.9871 - 72146.4835
Concentration (signal)	0.848381	
<b>Species # 3 (class 3)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	4.413055	4.1166 - 4.8238
Molecular mass <sup>1</sup> $M$ (Da)	89133.977232	75536.5637 - 117267.7025
Concentration (signal)	0.327668	
<b>Species # 4 (class 5)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	8.680858	6.5740 - 11.0911
Molecular mass <sup>1</sup> $M$ (Da)	33004.982556	27014.9568 - 41589.0286
Concentration (signal)	0.0689382	
<b>Species # 5 (class 6)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	18.269621	17.2184 - 19.6377
Molecular mass <sup>1</sup> $M$ (Da)	62045.164082	50792.5477 - 83330.0656
Concentration (signal)	0.101118	
<b>Common</b>		
Meniscus position (cm)	6.107007	6.1061 - 6.1081
Bottom position (cm)	7.216139	7.2101 - 7.2245
Buffer signal <sup>2</sup> frictional ratio $f_r$	2.382192	1.8745 - 3.3470
Reduced $\chi^2$ of fit [rmsd]	0.1871007	
R.m.s.d. of fit	0.004326	

<sup>1</sup>Substitute  $\bar{v} = 0.73 \text{ cm}^3/\text{g}$

<sup>2</sup>Continuous  $c(s)$  from 0.0 - 0.8 S



Reduced  $\chi^2$  [r.m.s.d]: 0.187101 [0.004326]

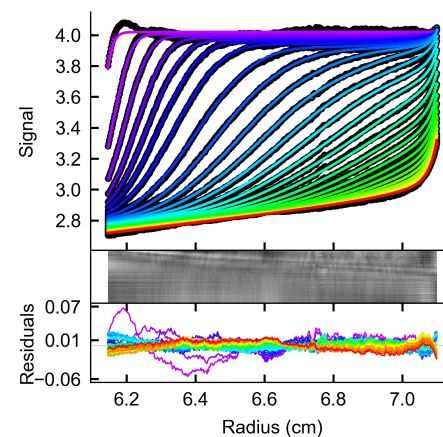
# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Species Analysis (0.44 mg/ml, 42000 rpm, i. o.)

Parameter	Value	95% confidence interval
<b>Species # 1 (class 1)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	2.045761	1.9890 - 2.0955
Molecular mass <sup>1</sup> $M$ (Da)	32468.185230	28140.1889 - 35903.1468
Concentration (signal)	0.564412	
<b>Species # 2 (class 2)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	2.874757	2.7103 - 2.9979
Molecular mass <sup>1</sup> $M$ (Da)	78493.617851	67236.8428 - 93921.0463
Concentration (signal)	0.388871	
<b>Species # 3 (class 3)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	3.976393	3.7610 - 4.1698
Molecular mass <sup>1</sup> $M$ (Da)	131871.067920	94586.0501 - 201092.7657
Concentration (signal)	0.193644	
<b>Common</b>		
Meniscus position (cm)	6.107400	6.1056 - 6.1093
Bottom position (cm)	7.201615	7.1931 - 7.2121
Buffer signal <sup>2</sup> frictional ratio $f_r$	1.481410	1.2663 - 1.9532
Reduced $\chi^2$ of fit [rmsd]	0.5931672	
R.m.s.d. of fit	0.007702	

<sup>1</sup>Substitute  $\bar{v} = 0.73 \text{ cm}^3/\text{g}$

<sup>2</sup>Continuous  $c(s)$  from 0.0 - 0.8 S



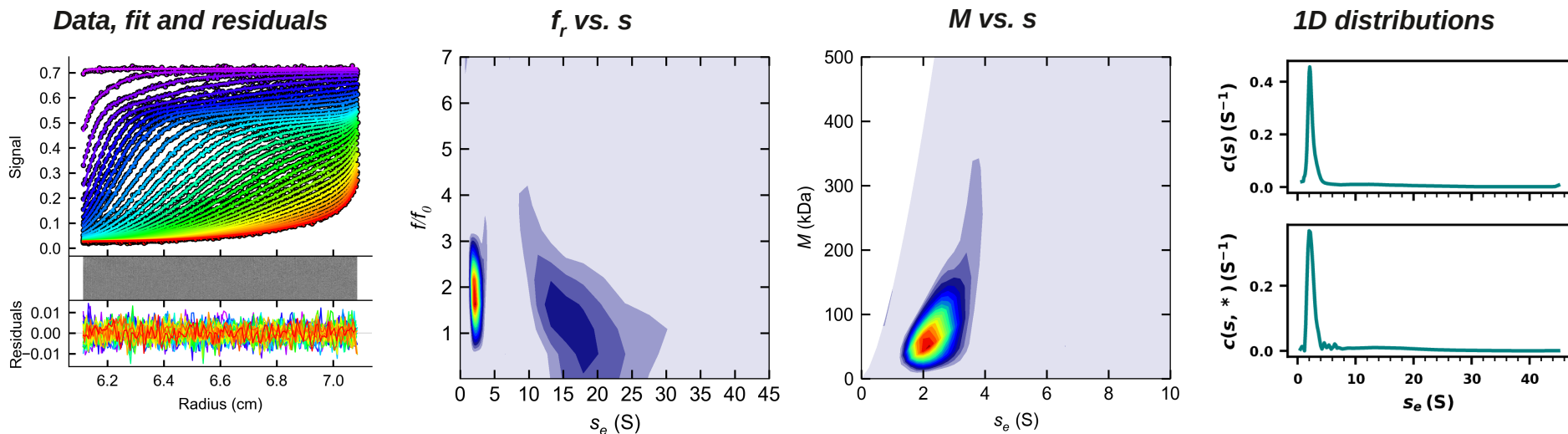
Reduced  $\chi^2$  [r.m.s.d]: 0.593167 [0.007702]



# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Absorbance optics @ 30000 rpm - loading concentration 2.0 mg/ml

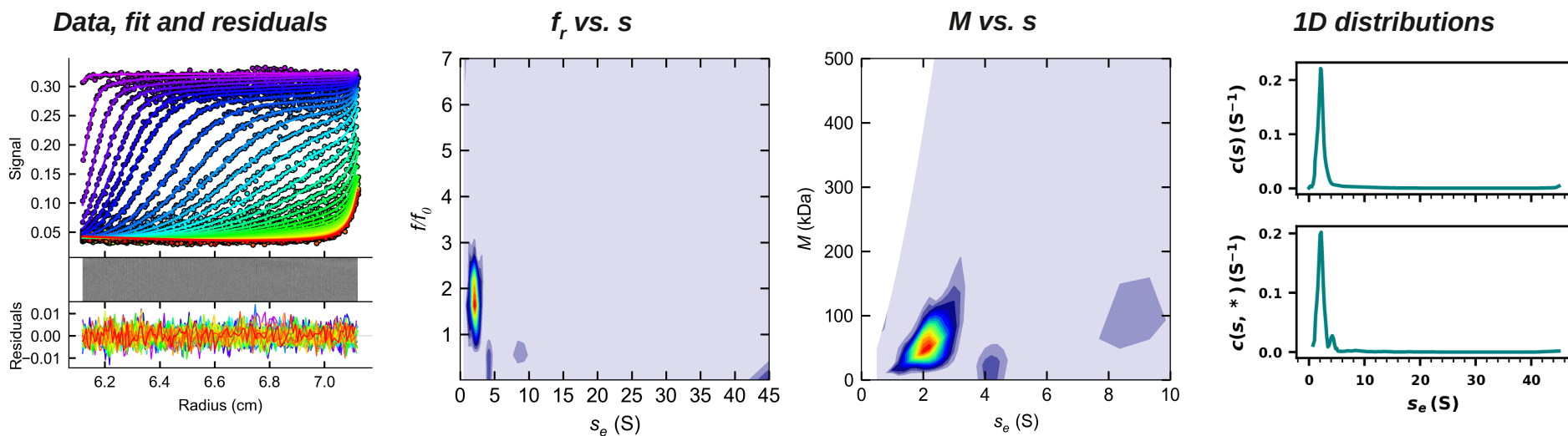
$c(s, f_r)$  analysis



Reduced  $\chi^2$  [r.m.s.d]: [0.003730]

Absorbance optics @ 42000 rpm - actual concentration 0.75 mg/ml

$c(s, f_r)$  analysis



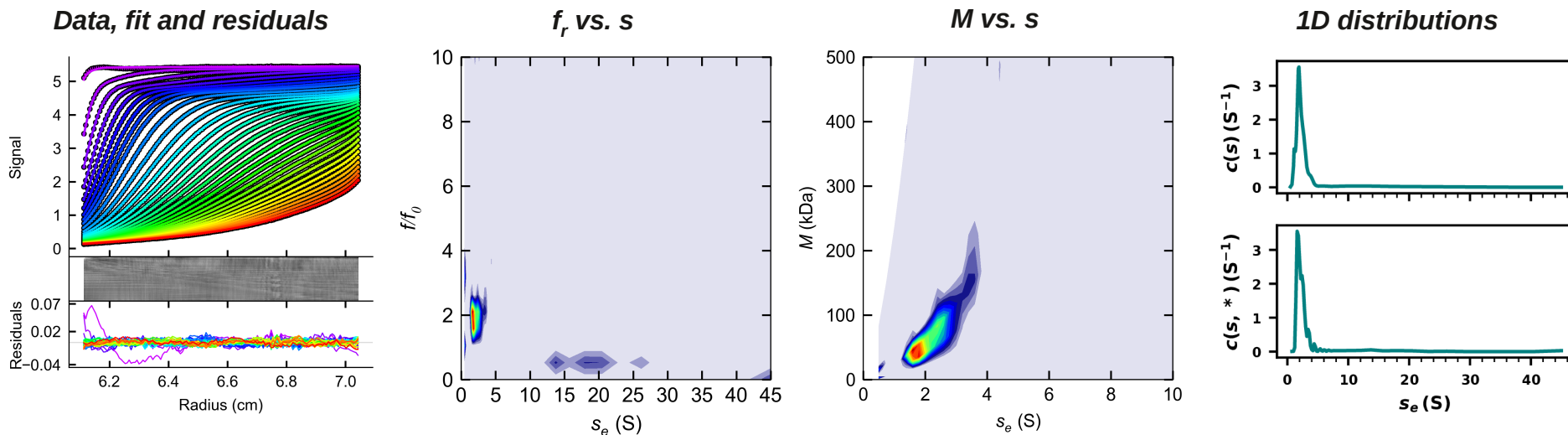
Reduced  $\chi^2$  [r.m.s.d]: [0.003382]

Fig. S6E

# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Interference optics @ 30000 rpm - loading concentration 2.0 mg/ml

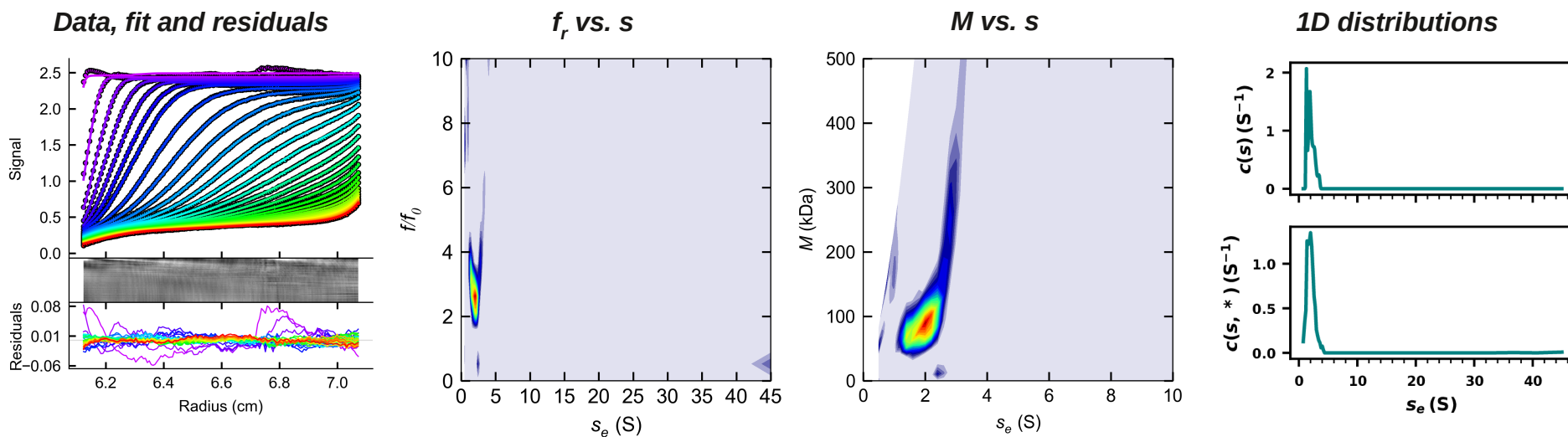
$c(s, f_r)$  analysis



Reduced  $\chi^2$  [r.m.s.d]: [0.005117]

Interference optics @ 42000 rpm - actual concentration 0.75 mg/ml

$c(s, f_r)$  analysis



Reduced  $\chi^2$  [r.m.s.d]: [0.009482]

Fig. S6F



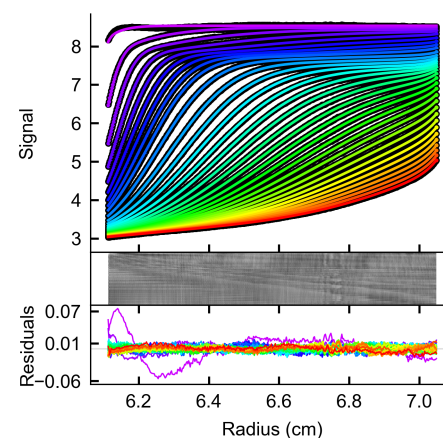
# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Species Analysis (2.00 mg/ml, 30000 rpm, i. o.)

Parameter	Value	95% confidence interval
<b>Species # 1 (class 1)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	2.259298	2.1789 - 2.3345
Molecular mass <sup>1</sup> $M$ (Da)	40911.383621	39015.7993 - 42569.8757
Concentration (signal)	2.10665	
<b>Species # 2 (class 2)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	3.362426	3.2647 - 3.4499
Molecular mass <sup>1</sup> $M$ (Da)	70607.408753	66954.0626 - 74408.5887
Concentration (signal)	1.80079	
<b>Species # 3 (class 3)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	4.879189	4.7237 - 4.9818
Molecular mass <sup>1</sup> $M$ (Da)	104525.361816	94748.6103 - 115580.2252
Concentration (signal)	0.557564	
<b>Species # 4 (class 5)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	9.007732	7.8572 - 10.1107
Molecular mass <sup>1</sup> $M$ (Da)	49129.251609	38405.4058 - 69148.1903
Concentration (signal)	0.179173	
<b>Species # 5 (class 6)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	16.172510	15.2395 - 17.1308
Molecular mass <sup>1</sup> $M$ (Da)	114826.491328	86165.0080 - 173746.5190
Concentration (signal)	0.15973	
<b>Species # 6 (class 7)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	24.574698	23.3895 - 25.9063
Molecular mass <sup>1</sup> $M$ (Da)	42732.956296	38685.9354 - 49078.2030
Concentration (signal)	0.337733	
<b>Common</b>		
Meniscus position (cm)	6.078588	6.0778 - 6.0790
Bottom position (cm)	7.233989	7.2288 - 7.2399
Buffer signal <sup>2</sup> frictional ratio $f_r$	3.336644	2.8806 - 3.9916
Reduced $\chi^2$ of fit [rmsd]	0.3392895	
R.m.s.d. of fit	0.005825	

<sup>1</sup>Substitute  $\bar{v} = 0.73 \text{ cm}^3/\text{g}$

<sup>2</sup>Continuous  $c(s)$  from 0.0 - 0.8 S



Reduced  $\chi^2$  [r.m.s.d]: 0.339290 [0.005825]

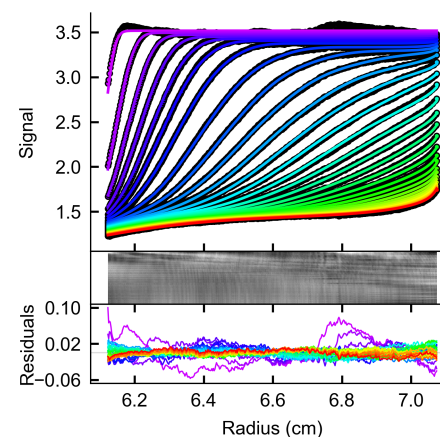
# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Species Analysis (0.75 mg/ml, 42000 rpm, i. o.)

Parameter	Value	95% confidence interval
<b>Species # 1 (class 1)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	1.737800	1.6152 - 1.8163
Molecular mass <sup>1</sup> $M$ (Da)	42836.913056	36022.7457 - 53638.9211
Concentration (signal)	0.488908	
<b>Species # 2 (class 2)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	2.545311	2.3295 - 2.6511
Molecular mass <sup>1</sup> $M$ (Da)	65269.106310	58823.6621 - 72195.2814
Concentration (signal)	0.869682	
<b>Species # 3 (class 3)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	3.497679	3.2748 - 3.6633
Molecular mass <sup>1</sup> $M$ (Da)	112116.879536	90117.1719 - 141988.8079
Concentration (signal)	0.526914	
<b>Species # 4 (class 4)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	4.655214	4.3692 - 4.8847
Molecular mass <sup>1</sup> $M$ (Da)	232327.712281	148281.3925 - 426686.8809
Concentration (signal)	0.156663	
<b>Common</b>		
Meniscus position (cm)	6.093045	6.0911 - 6.0950
Bottom position (cm)	7.175154	7.1615 - 7.1904
Buffer signal <sup>2</sup> frictional ratio $f_r$	5.718135	4.3407 - 7.4711
Reduced $\chi^2$ of fit [rmsd]	0.9201238	
R.m.s.d. of fit	0.009592	

<sup>1</sup>Substitute  $\bar{v} = 0.73 \text{ cm}^3/\text{g}$

<sup>2</sup>Continuous  $c(s)$  from 0.0 - 0.8 S

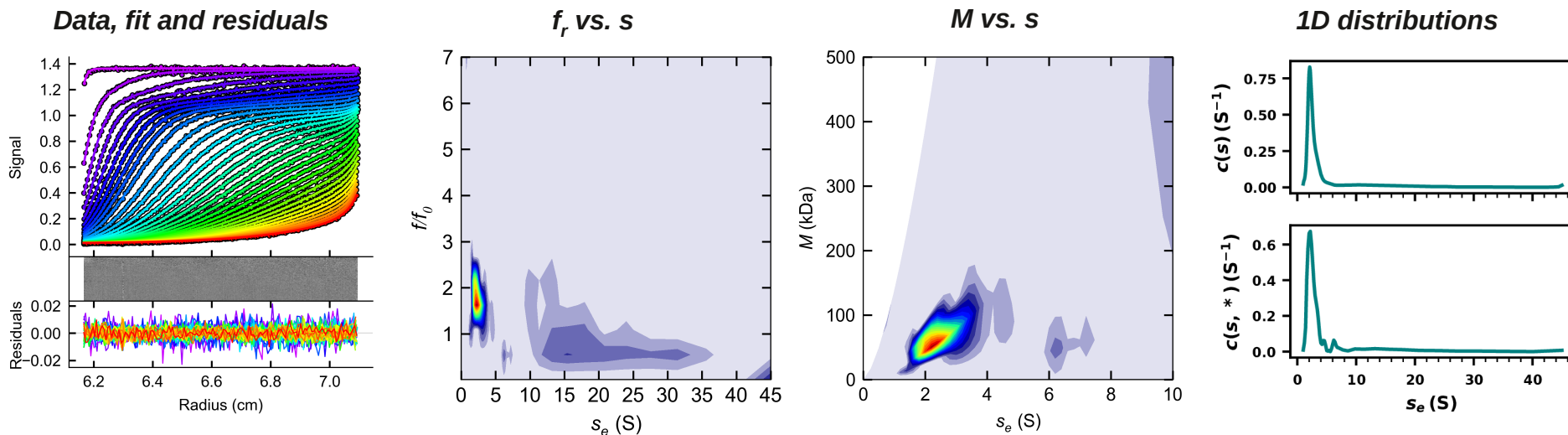


Reduced  $\chi^2$  [r.m.s.d]: 0.920124 [0.009592]

# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Absorbance optics @ 30000 rpm - loading concentration 4.0 mg/ml

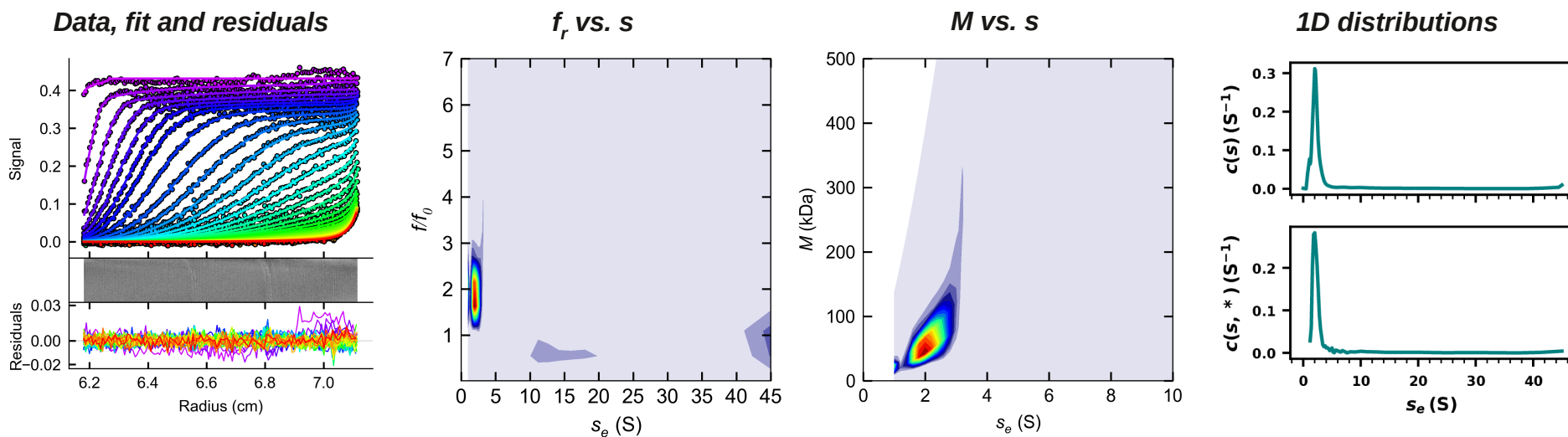
$c(s, f_r)$  analysis



Reduced  $\chi^2$  [r.m.s.d]: [0.004508]

Absorbance optics @ 42000 rpm - actual concentration 1.09 mg/ml

$c(s, f_r)$  analysis



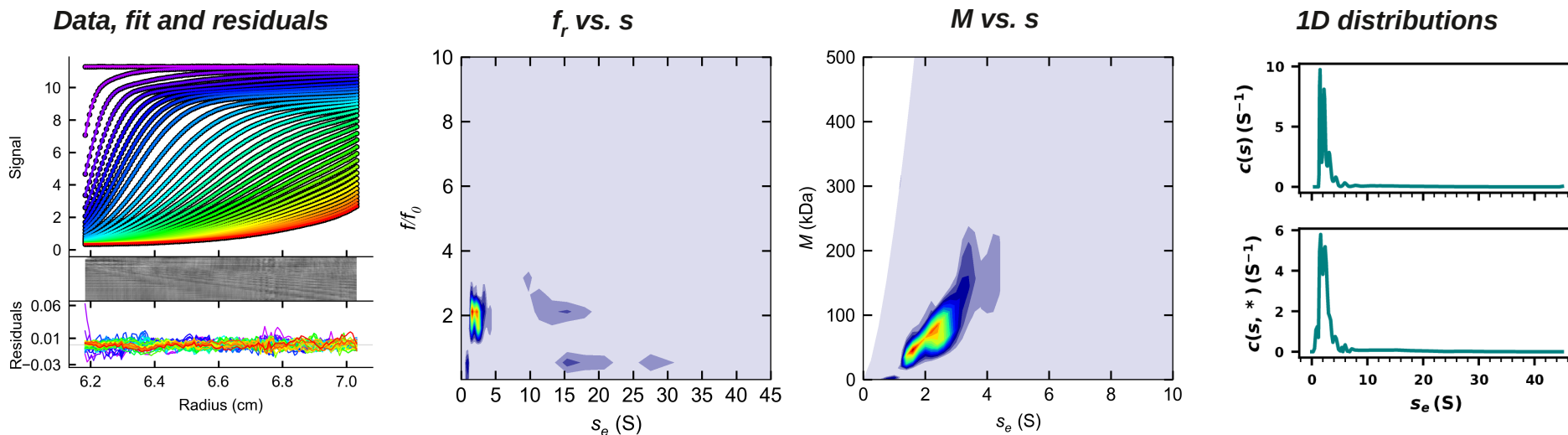
Reduced  $\chi^2$  [r.m.s.d]: [0.004322]

Fig. S6I

# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Interference optics @ 30000 rpm - loading concentration 4.0 mg/ml

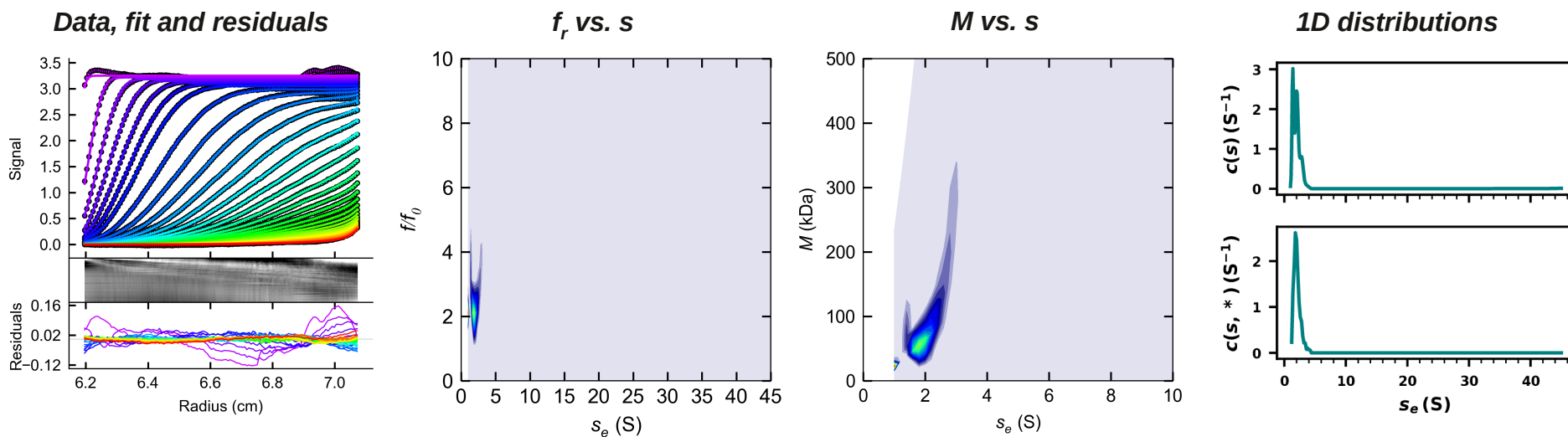
$c(s, f_r)$  analysis



Reduced  $\chi^2$  [r.m.s.d]: [0.005735]

Interference optics @ 42000 rpm - actual concentration 1.09 mg/ml

$c(s, f_r)$  analysis



Reduced  $\chi^2$  [r.m.s.d]: [0.019392]

Fig. S6J

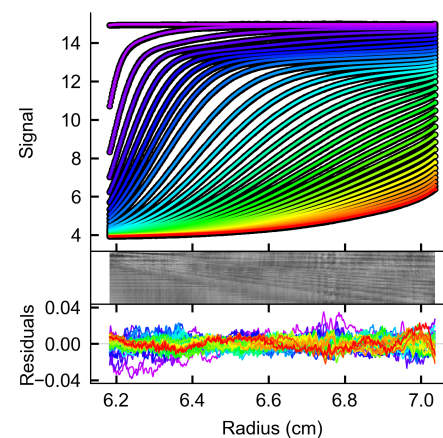
# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Species Analysis (4.00 mg/ml, 30000 rpm, i. o.)

Parameter	Value	95% confidence interval
<b>Species # 1 (class 1)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	1.893838	1.8767 - 1.9115
Molecular mass <sup>1</sup> $M$ (Da)	36242.513823	35605.9184 - 36924.8539
Concentration (signal)	2.02274	
<b>Species # 2 (class 2)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	2.895957	2.8717 - 3.0216
Molecular mass <sup>1</sup> $M$ (Da)	56798.604800	56171.7866 - 57417.8843
Concentration (signal)	4.20148	
<b>Species # 3 (class 3)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	4.032742	3.7914 - 4.0755
Molecular mass <sup>1</sup> $M$ (Da)	91110.192293	88642.3598 - 93600.9839
Concentration (signal)	2.17643	
<b>Species # 4 (class 4)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	5.558437	5.5059 - 5.7911
Molecular mass <sup>1</sup> $M$ (Da)	106503.214501	97771.9841 - 113202.5506
Concentration (signal)	0.924506	
<b>Species # 5 (class 5)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	9.980081	9.6854 - 10.2378
Molecular mass <sup>1</sup> $M$ (Da)	55993.213184	50533.5330 - 65497.6998
Concentration (signal)	0.504232	
<b>Species # 6 (class 6)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	18.163020	17.9089 - 18.4664
Molecular mass <sup>1</sup> $M$ (Da)	116367.123008	102061.2320 - 138132.1828
Concentration (signal)	0.460625	
<b>Species # 7 (class 7)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	28.605184	27.7262 - 30.0465
Molecular mass <sup>1</sup> $M$ (Da)	50122.522087	49947.7361 - 50312.8130
Concentration (signal)	0.500927	
<b>Common</b>		
Meniscus position (cm)	6.154459	6.1540 - 6.1548
Bottom position (cm)	7.219491	7.2170 - 7.2245
Buffer signal <sup>2</sup> frictional ratio $f_r$	1.126869	0.9060 - 1.3521
Reduced $\chi^2$ of fit [rmsd]	0.5273897	
R.m.s.d. of fit	0.007262	

<sup>1</sup>Substitute  $\bar{v} = 0.73 \text{ cm}^3/\text{g}$

<sup>2</sup>Continuous  $c(s)$  from 0.0 - 0.8 S



Reduced  $\chi^2$  [r.m.s.d]: 0.527390 [0.007262]

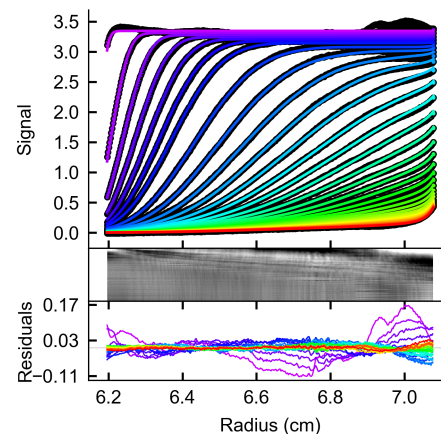
# Complex of human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA

Species Analysis (1.09 mg/ml, 42000 rpm, i. o.)

Parameter	Value	95% confidence interval
<b>Species # 1 (class X)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	1.209722	0.8561 - 1.7601
Molecular mass <sup>1</sup> $M$ (Da)	30400.731201	20063.1666 - >900000
Concentration (signal)	0.255537	
<b>Species # 2 (class 1)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	1.922069	1.7474 - 2.6281
Molecular mass <sup>1</sup> $M$ (Da)	49646.909726	42877.4221 - 61527.1873
Concentration (signal)	0.962814	
<b>Species # 3 (class 2)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	2.787625	2.5211 - 3.3593
Molecular mass <sup>1</sup> $M$ (Da)	64072.302750	54843.6506 - 109988.7832
Concentration (signal)	1.43393	
<b>Species # 4 (class 3)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	3.776941	3.3105 - 4.1561
Molecular mass <sup>1</sup> $M$ (Da)	136947.108751	87118.2414 - 219814.9212
Concentration (signal)	0.4912	
<b>Species # 5 (class 4)</b>		
Sedimentation coefficient $s_{20^{\circ}\text{C}, w}$ (S)	4.936165	4.2947 - 5.5977
Molecular mass <sup>1</sup> $M$ (Da)	273002.737766	108529.5424 - >900000
Concentration (signal)	0.119144	
<b>Common</b>		
Meniscus position (cm)	6.169227	6.1674 - 6.1710
Bottom position (cm)	7.181567	7.1682 - 7.2017
Buffer signal <sup>2</sup> frictional ratio $f_r$	4.608594	2.9507 - 28.0506
Reduced $\chi^2$ of fit [rmsd]	3.344057	
R.m.s.d. of fit	0.018287	

<sup>1</sup>Substitute  $\bar{v} = 0.73 \text{ cm}^3/\text{g}$

<sup>2</sup>Continuous  $c(s)$  from 0.0 - 0.8 S



Reduced  $\chi^2$  [r.m.s.d]: 3.344057 [0.018287]



## **Section 3:**

# **Characterization of Human PCBP2, PRRSV nsp1- $\beta$ and PRRSV 34 nt ssRNA Complex by SEC-SAXS**

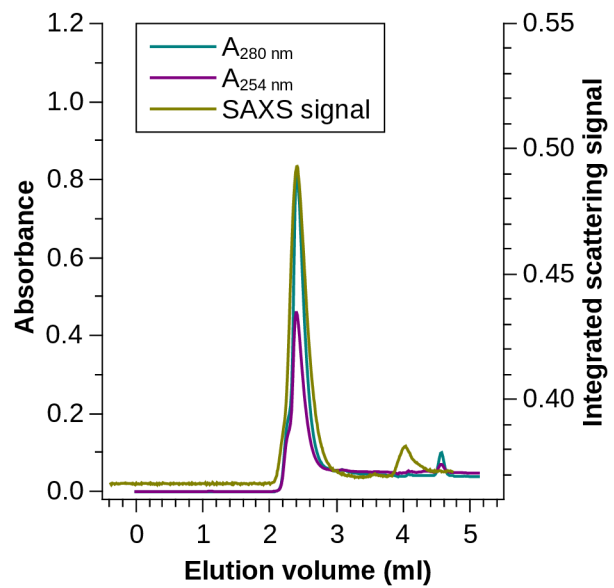


Fig. S7