

#### **ENFORCE**

#### Danish National Cohort Study of Effectiveness and Safety of SARS-CoV-2 Vaccines

#### **Monthly Report**

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Report prepared by: the ENFORCE consortium

Approved by: the ENFORCE scientific steering committee

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# Summary of key changes from previous report

#### Enrolment

A table outlining the total number of participants who have withdrawn from the study is now included.

#### Outcomes

The results of the serum antibody quantification using ELISA (Wantai) performed by Statens Serum Institut (SSI) are included for those with data up to visit 3.

The results from the multiantigen serological tests for antibody levels against the Receptor Binding Domain, the complete Spike protein and the Nucleocapsid are also included for visits 1 and 2.

The results from the multiantigen serological tests, for the neutralizing antibody titer (NAb) levels for the Receptor Binding Domain, the complete Spike protein and the Nucleocapsid are also included for visits 1 and 2.



### Methods

The data presented in this report are descriptive. A detailed statistical analysis plan will be developed prior to any formal analysis being conducted.

#### Data sources

The data used to generate this report are currently based only on the data stored in REDCap from the case report forms (CRF's) and online symptoms form. Data on serum antibody quantification using ELISA (Wantai) was provided by the SSI and the multiantigen serological tests by Aarhus University Hospital.

Results from the ELISA detection of total serum Ig to the Receptor Binding Domain (Wantai) were recorded as Negative (ratio <0.9), Positive (ratio >1.1), or inconclusive (ratio between 0.9-11). The ratio was calculated as the OD value/cut-off, where the cut-off= average of the negative controls+0.16. If the average is below 0.03 then the cut-off is set to 0.16 + 0.03. For manual execution the cut-off will almost always be 0.19.

For the multiantigen serological tests, the geometric mean and 95% confidence intervals (CI) for the antibody levels against the Receptor Binding Domain, the complete Spike protein and the Nucleocapsid at each study visit are reported. The calibration curve used to calculate antibody concentrations are performed by fitting the signals from the calibrators in a 4-parameter sigmoidal dose-response model. Antibody concentrations can then be determined from their ECL signals by backfitting to the calibration curve.

#### **Definitions**

In this version of the report the type of vaccine received is based on self-reports from participants at enrolment into the study which occurs prior to them receiving their first vaccine. In subsequent reports this will be updated using data from the national vaccine registry as the gold standard when this data is available.



# Enrolment

The section gives an overview of the current enrolment status of participants in the study. Table 1 outlines the number of participants currently enrolled in the study and reasons for exclusion.

Table 1 Summary of participants enrolled in the study

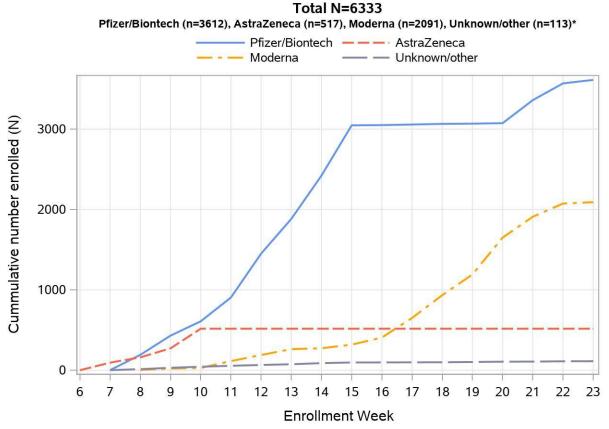
 Total	
included	Reason for exclusion
6345	All patients
6336	Consent withdrawn and requested data deleted
6334	Provided informed consent
6334	Missing enrolment date
6333	Aged under 18
6333	Vaccine not recommended
6333	Vaccinated Previously
6333	Belongs to a vaccine target group
6333	Agrees to follow protocol



Figure 1 shows the cumulative number of participants enrolled by vaccine type. Enrolment began on the 13<sup>th</sup> February 2021 (Week 6) when the first patient was enrolled prior to receiving the AstraZeneca vaccine. The first participant enrolled prior to receiving the Pfizer/BionTech vaccination was on the 16<sup>th</sup> February 2021, followed by Moderna on the 24<sup>th</sup> February 2021 (week 8).

A total of 517 participants were enrolled in the AstraZeneca vaccine group prior to the pausing of AstraZeneca vaccination in week 10 (11th March 2021). Enrolment for those receiving Pfizer was paused in week 15, after the target of 2500 participants was reached, but then restarted in week 20.

Figure 1 Cumulative number of participants enrolled by vaccine type



<sup>\*</sup>Type of vaccine received is currently based on self-report



## Demographics at enrolment

Table 2 gives an overview of the participant demographics at the time of enrolment into the study overall and by vaccine. Table 3 gives an overview of the number of participants with concomitant diseases at enrolment and the use of any medications in the 24 hours prior to enrolment.

Table 2 Participant demographics at study enrolment by vaccine

	Vaccine type					
	Total (N=6333)	Pfizer/Biontech (N=3612)	AstraZeneca (N=517)	Moderna (N=2091)	Unknown/other (N=113)	
Number of persons (%)						
Gender						
Male	2816 (44.5)	1751 (48.5)	81 (15.7)	917 (43.9)	67 (59.3)	
Female	3517 (55.5)	1861 (51.5)	436 (84.3)	1174 (56.1)	46 (40.7)	
Median (interquartile range, IQR)						
Age at enrolment (years)	65 (56, 75)	71 (56, 78)	45 (31, 56)	64 (59, 69)	71 (61, 79)	
Enrolment date	APR21 (MAR21, MAY21)	MAR21 (MAR21, APR21)	MAR21 (FEB21, MAR21)	MAY21 (APR21, MAY21)	MAR21 (MAR21, APR21)	

Table 3 Concomitant diseases and medications, overall and by vaccine

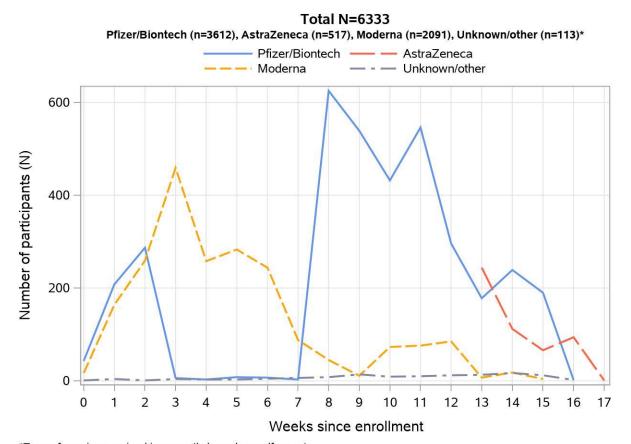
		Vaccine type					
	Total (N=6333)	Pfizer/Biontech (N=3612)	AstraZeneca (N=517)	Moderna (N=2091)	Unknown/other (N=113)		
Any concomitant disease N (%)	4296 (67.8)	2729 (75.6)	173 (33.5)	1311 (62.7)	83 (73.5)		
Any medications taken in the last 24 hours N (%)	4519 (71.4)	2841 (78.7)	237 (45.8)	1350 (64.6)	91 (80.5)		



#### **Enrolment Progress**

Figure 2 shows the current status of participants in the study at the date of most recent data extract. The first participant was enrolled 17 weeks ago. The majority of participants in the Pfizer vaccine group have been under follow-up for at least 7 weeks, whereas most of those in the Moderna group have been under follow-up for less than 7 weeks.

Figure 2 Current status of participants



<sup>\*</sup>Type of vaccine received is currently based on self-report



## Withdrawal/Loss to follow-up

Table 4 shows the number of participants enrolled in the study who have subsequently withdrawn and no longer wish to be contacted or attend future study visits.

Table 4 Number and percent of participants discontinuing in the study, by vaccine type

	Vaccine type						
	Total (N=6333)	Pfizer/Biontech (N=3612)	AstraZeneca (N=517)	Moderna (N=2091)	Unknown/other (N=113)		
Study status (N, % of total)							
Ctill den felle	6237 (98.5)	3549 (98.3)	495 (95.7)	2082 (99.6)	111 (98.2)		
Still under follow-up	()	` ,	, ,	, ,			



### Outcomes

## Primary outcome

Until the threshold for MPNAT has been established we will present the results from both the ELISA (Wantai) and the multiantigen serological test including the Receptor Binding Domain (RBD), and the complete Spike (S) protein and the Nucleocapsid (NC).

The data from the ELISA (Wantai) are presented as negative, positive and inconclusive (see methods for categorization) and are shown in Table 5.

Table 5 Presence of antibodies at study visit, ELISA (Wantai) from SSI

			Vaccine	e type	
	Total (N=6333)	Pfizer/Biontech (N=3612)	AstraZeneca (N=517)	Moderna (N=2091)	Unknown/other (N=113)
Number of persons with antibody data at enrolment (% of total)	6275 (99.1)	3573 (98.9)	508 (98.3)	2081 (99.5)	113 (100)
Wantai result at enrolment					
Negative	5981 (95.3)	3441 (96.3)	453 (89.2)	1979 (95.1)	108 (95.6)
Positive	291 (4.6)	131 (3.7)	55 (10.8)	100 (4.8)	5 (4.4)
Inconclusive	3 (0.0)	1 (0.0)	0	2 (0.1)	0
Number of persons with antibody data at visit 2 (% of total)	3899 (61.6)	2877 (79.7)	276 (53.4)	657 (31.4)	89 (78.8)
Wantai result at visit 2					
Negative	915 (23.5)	805 (28.0)	6 (2.2)	68 (10.4)	36 (40.4)
Positive	2933 (75.2)	2028 (70.5)	270 (97.8)	584 (88.9)	51 (57.3)
Inconclusive	51 (1.3)	44 (1.5)	0	5 (0.8)	2 (2.2)
Days from first vaccination (median (IQR))	23 (21, 28)	21 (20, 24)	82 (78, 84)	29 (28, 34)	22 (20, 28)
Number of persons with antibody data at visit 3 (% of total)	388 (6.1)	304 (8.4)	41 (7.9)	26 (1.2)	17 (15.0)
Wantai result at visit 3					
Negative	51 (13.1)	37 (12.2)	3 (7.3)	5 (19.2)	6 (35.3)
Positive	337 (86.9)	267 (87.8)	38 (92.7)	21 (80.8)	11 (64.7)
Days from first vaccination (median (IQR))	90 (87, 92)	90 (87, 92)	90 (89, 94)	87 (83, 90)	89 (87, 91)



From the multiantigen serological tests, the geometric mean and 95% confidence intervals (CI) for the antibody levels against the Receptor Binding Domain, the complete Spike protein and the Nucleocapsid at each study visit are reported in Table 6. Figure 3-5 show the distribution of the three assays on the log10 scale, by vaccine type and study visit.

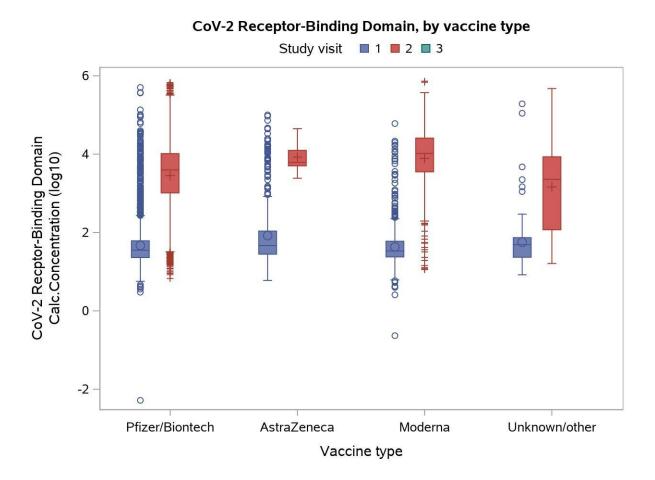
Table 6 Presence of antibodies at study visit, Receptor-Binding Domain (RBD), Spike antibody and Nucleocapsid

		Vaccine type					
	Total (N=6333)	Pfizer/Biontech (N=3612)	AstraZeneca (N=517)	Moderna (N=2091)	Unknown/other (N=113)		
Number of persons with data at enrolment (% of total)	4480 (70.7)	3037 (84.1)	508 (98.3)	835 (39.9)	100 (88.5)		
Number of persons with data at visit 2 (% of total)	2852 (45.0)	2510 (69.5)	13 (2.5)	256 (12.2)	73 (64.6)		
CoV-2 Receptor-Binding Domain (SERO)							
GM at enrolment (95%CI)	49 (47, 51)	47 (45, 49)	84 (71, 98)	43 (40, 46)	57 (42, 78)		
GM at visit 2 (95%CI)	3064 (2840, 3307)	2839 (2621, 3074)	8387 (4928, 14273)	7617 (5922, 9797)	1458 (801, 2653)		
CoV-2 Spike antibody (SERO)							
GM at enrolment (95%CI)	96 (91, 100)	93 (88, 98)	153 (128, 183)	78 (71, 86)	106 (75, 150)		
GM at visit 2 (95%CI)	10367 (9641, 11147)	9829 (9108, 10607)	21396 (12970, 35296)	21510 (16975, 27258)	4396 (2380, 8118)		
CoV-2 Nucleocapsid (SERO)							
GM at enrolment (95%CI)	253 (240, 267)	252 (237, 269)	354 (293, 427)	208 (186, 234)	260 (175, 385)		
GM at visit 2 (95%CI)	243 (228, 259)	244 (228, 262)	345 (157, 759)	223 (183, 272)	243 (155, 381)		

GM: Geometric mean



Figure 3 Distribution of CoV-2 Receptor-Binding Domain (RBD) levels at each study visit (see Table 6 for the number included at each time point)



0

Pfizer/Biontech



Unknown/other

Figure 4 Distribution of CoV-2 Spike antibody levels at each study visit (see Table 6 for the number included at each time point)

CoV-2 Spike antibody, by vacccine type

# Study Visit 1 2 3 6 CoV-2 Spike antibody Calc. Concentration (log10) 0 000 0 00000 0 0000 0 2 -

Vaccine type

Moderna

AstraZeneca



Figure 5 Distribution of CoV-2 Nucleocapsid levels at each study visit (see Table 6 for the number included at each time point)

## CoV-2 Nucleocapsid, by vaccine type Study visit ■ 1 ■ 2 ■ 3 6 00 0 0 CoV-2 Nucleocapsid Calc. Concentration (log10) 0 5 0 8 4 3 2 1 0 Pfizer/Biontech AstraZeneca Moderna Unknown/other

Vaccine type



### Secondary outcome – data not yet available will be included in future reports

The secondary outcome of breakthrough infections is monitored in two different ways. The number of participants testing positive for SARS-CoV-19, as reported via MiBa, and by serological monitoring (detection of SARS-CoV-2 nucleocapsid antibodies). The number of participants experiencing these outcomes will be reported in Table 5.

Table 7 Number of participants experiencing breakthrough infection and hospitalisation due to COVID-19

	Overall	Pfizer/BionTech	Moderna	AstraZeneca	Other
	N (%)	N (%)	N (%)	N (%)	N (%)
Number of					
participants					
with					
breakthrough					
infection					
Number testing					
positive for					
SARS-CoV-2					
reported via					
MiBa					
Number with					
SARS-CoV-2					
nucleocapsid					
antibodies					
detected					
Time to					
breakthrough					
infection					
(median, IQR)					
Age group					
<65					
65-74					
75-79					
80-84					
≥85					
Sex					
Male					
Female					
Number					
admitted to					
hospital due to					
COVID-19					



Time to hospitalisation (median, IQR)			
Age group			
<65			
65-74			
75-79			
80-84			
≥85			
Sex			
Male			
Female			



# Safety Monitoring

#### Local and systemic reactions

Table 8 outlines the number of participants reporting any local or systemic reactions within 7 days of vaccination. Note that participants can report multiple symptoms. The percentages are out of the total number of participants who had completed the symptoms form at the time of data extraction. All participants are encouraged to complete the form even if they experience no symptoms.

Figure 6 shows the proportion reporting mild, moderate or severe symptoms by vaccine type. To further illustrate the symptoms reported by participant demographics Figure 7-9 show the proportion reporting mild, moderate or severe symptoms by each vaccine separately stratified by age group (<55 years, 55-64 years and ≥65 years) and Figure 10-12 stratified by gender.

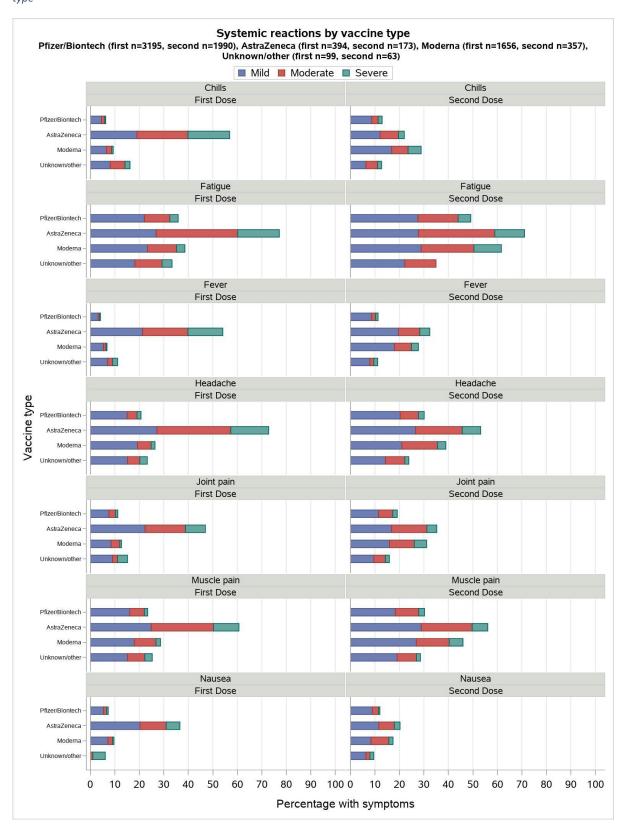


Table 8 Number & percent reporting local/systemic reactions within 7 days post vaccination overall, all and by vaccination

	To	Total Pfizer/B		Pfizer/Biontech AstraZeneca		Moderna		Unknown/Other		
	First (N=5344)	Second (N=2583)	First (N=3195)	Second (N=1990)	First (N=394)	Second (N=173)	First (N=1656)	Second (N=357)	First (N=99)	Second (N=63)
Number of persons (%)										
Any clinical symptoms	3044 (57.0)	1675 (64.8)	1651 (51.7)	1218 (61.2)	358 (90.9)	143 (82.7)	978 (59.1)	280 (78.4)	57 (57.6)	34 (54.0)
Any local symptoms at injection site	4105 (76.8)	1927 (74.6)	2276 (71.2)	1406 (70.7)	346 (87.8)	161 (93.1)	1418 (85.6)	312 (87.4)	65 (65.7)	48 (76.2)



Figure 6 Percentage of participants reporting systemic and local symptoms following  $\mathbf{1}^{\text{st}}$  dose and  $\mathbf{2}^{\text{nd}}$  dose of the vaccine, by vaccine type





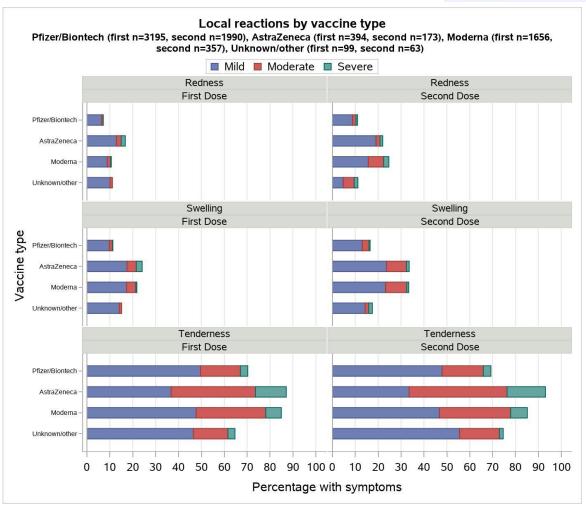




Figure 7 Percentage of participants reporting systemic and local symptoms following **1**st **and 2**nd **dose** of Pfizer/Biontech, by age group

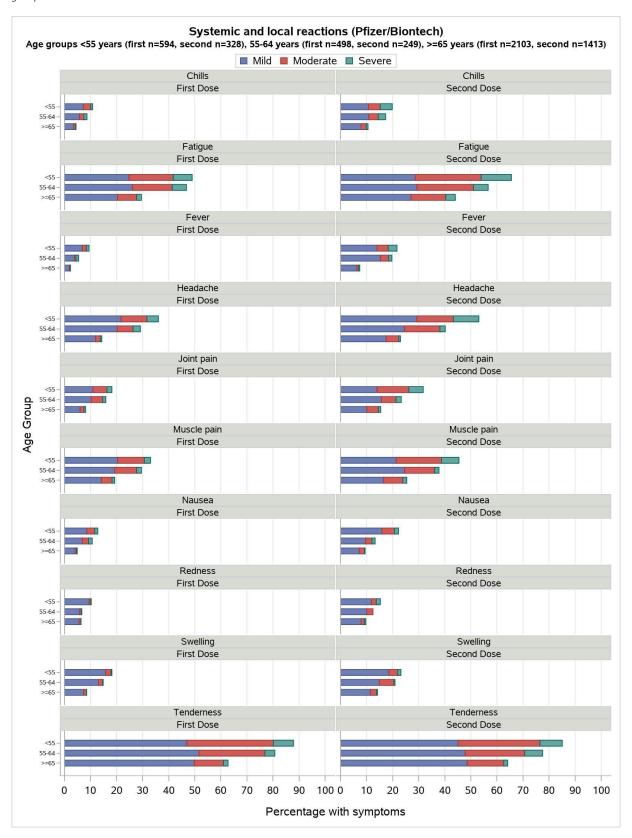




Figure 8 Percentage of participants reporting systemic and local symptoms following 1<sup>st</sup> & 2<sup>nd</sup> dose of AstraZeneca, by age group. Data in the >65 group not shown due to small numbers.

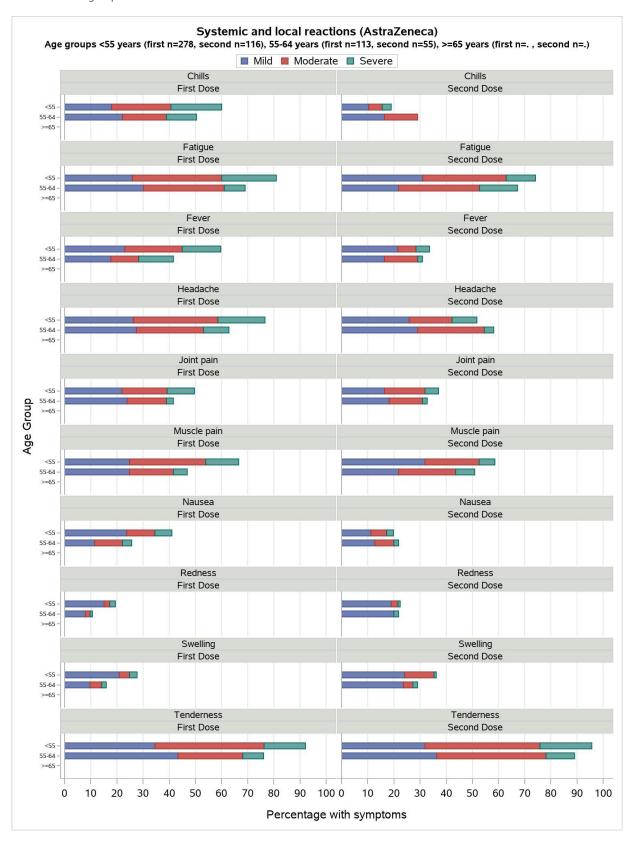




Figure 9 Percentage of participants reporting systemic and local symptoms following 1st dose & 2nd dose of Moderna, by age group

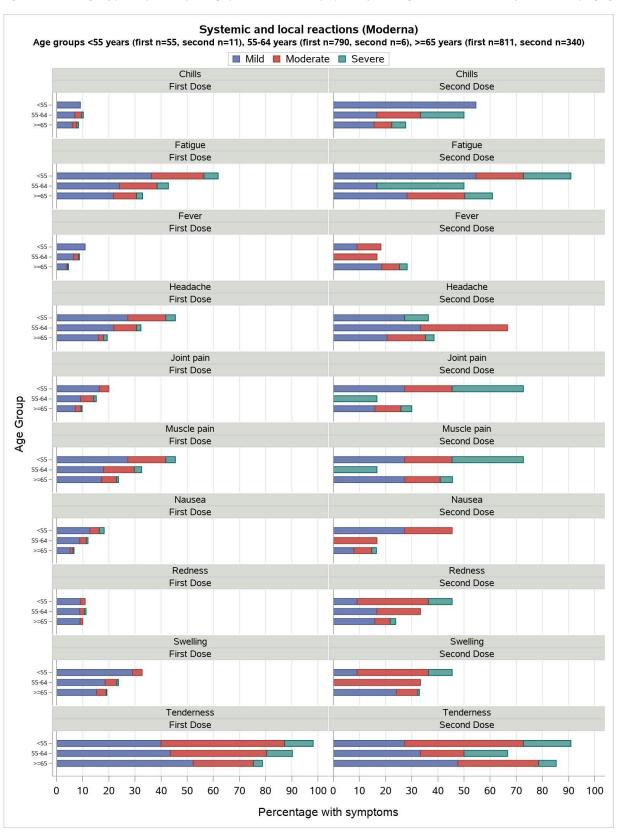




Figure 10 Percentage of participants reporting systemic and local symptoms following 1st & 2nd dose of Pfizer/Biontech, by gender

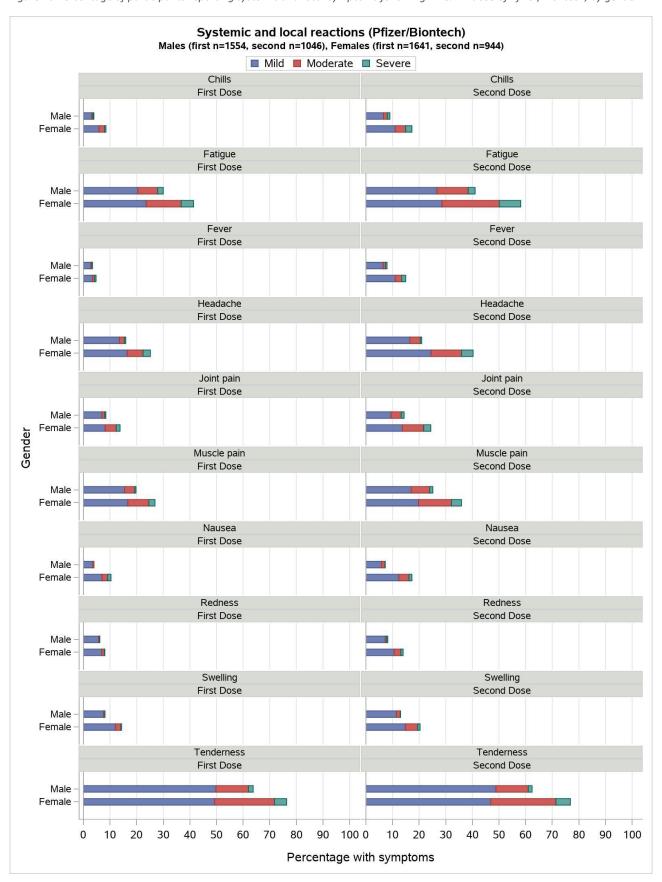




Figure 11 Percentage of participants reporting systemic and local symptoms following  $\mathbf{1}^{\text{st}}$  dose and  $\mathbf{2}^{\text{nd}}$  dose of AstraZeneca, by gender

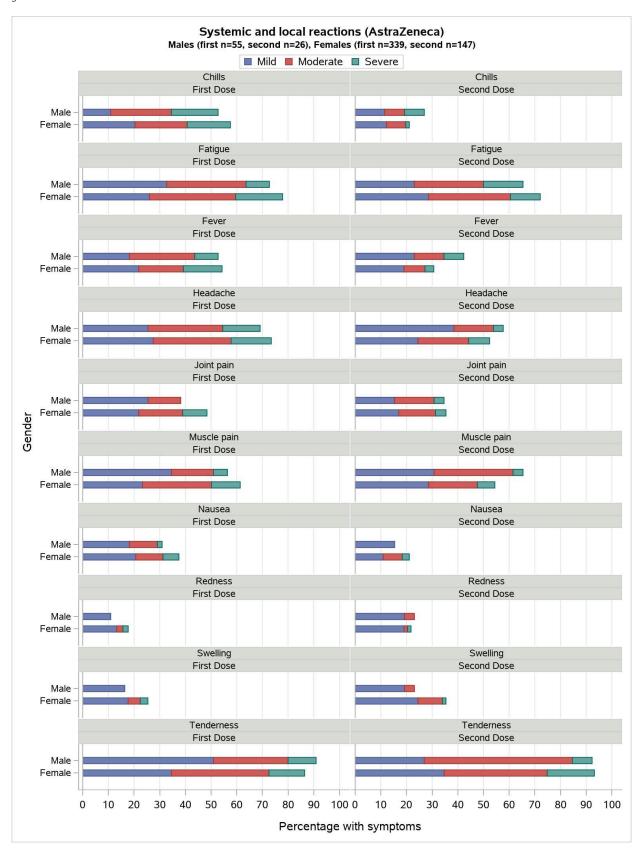
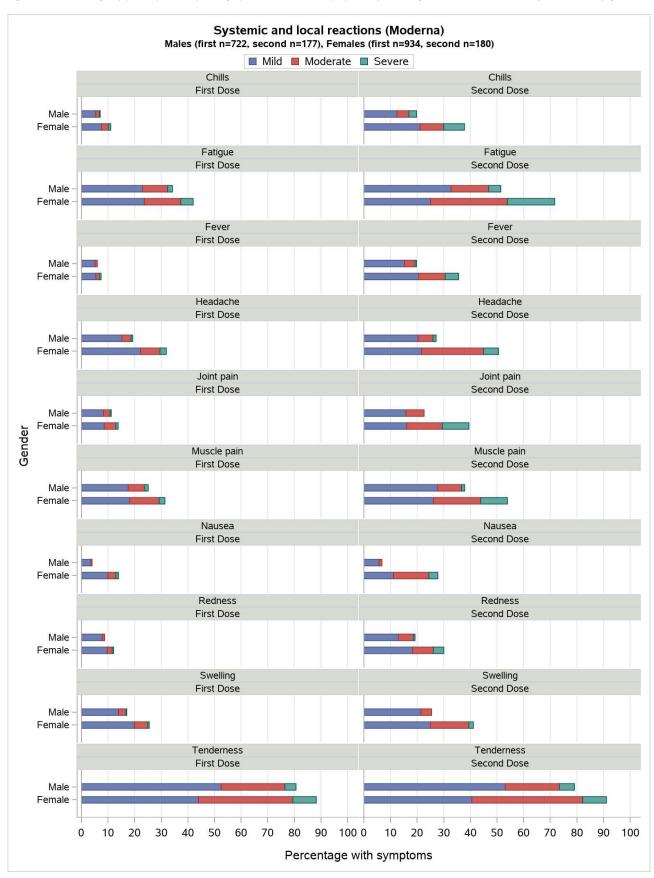




Figure 12 Percentage of participants reporting systemic and local symptoms following  $\mathbf{1}^{\text{st}}$  dose and  $\mathbf{2}^{\text{nd}}$  dose of Moderna, by gender





## Adverse and Serious Adverse Events

This section gives an overview of the AEs (Table 9) and SAEs (Table 10) reported in the study thus far. The data are first shown overall and by vaccine type.

Table 9 Overview of AEs (grade 3 and 4) reported by vaccine

	Vaccine type					
	Total (N=6333)	Pfizer/Biontech (N=3612)	AstraZeneca (N=517)	Moderna (N=2091)	Unknown/other (N=113)	
Number of persons (%)						
At least one Adverse Event reported	205 (3.2)	134 (3.7)	36 (7.0)	27 (1.3)	8 (7.1)	
Number of AE						
Total number of AE reported (N, % of AE)	256 (100)	167 (100)	44 (100)	30 (100)	15 (100)	

Table 10 Overview of SAEs reported by vaccine

	Vaccine type						
	Total (N=6333)	Pfizer/Biontech (N=3612)	AstraZeneca (N=517)	Moderna (N=2091)	Unknown/other (N=113)		
Total number of participants reporting any SAE (N, %)	46 (0.7)	39 (1.1)	<5*	<5*	<5*		
Total number of SAE reported (N, % of SAE)	54 (100)	46 (100)	<5*	<5*	<5*		

<sup>\*</sup>Exact numbers not shown due to small numbers



## Deaths

There have been <5 deaths reported in the study thus far and none had a reasonable probability of relatedness to vaccination.