

Table 14.2.2.1.3.1.1
Analysis of Vaccine Efficacy of mRNA-1273 to Prevent COVID-19* Based on Adjudication Committee Assessments Starting 14 Days
After Second Injection
Per-Protocol Set

	Placebo (N=xx)	mRNA-1273 (N=xx)
Number of Subjects with COVID-19*, n (%)	xx (xx.x)	xx (xx.x)
Number of Subjects Censored, n (%)	xx (xx.x)	xx (xx.x)
Vaccine Efficacy Based on Hazard Ratio (95% CI) [1] p-value [2]		x.xxx (x.xxx, x.xxx) 0.xxxx
Person-Years [3]	xx.x	xx.x
Incidence Rate per 1,000 Person-Years (95% CI) [4]	xx.x (xx.x, xx.x)	xx.x (xx.x, xx.x)
Vaccine Efficacy Based on Incidence Rate (95% CI) [5]		x.xxx (x.xxx, x.xxx)

ADTTEA where PPROTFL=Y and PARAMCD='TTCVDC1'
Treatment: TRT01P

Event: count # of subjects with CNSR=0
Censored: count # of subjects with CNSR=1

Person-years: sum AVAL over all subjects for each treatment group, then divided by 365.25

VE (95% CI) based on hazard ratio, IR (95% CI), VE (95% CI) based on IR:
Use mVEHRIR.sas and CALL %VEHRIR(T14020201030101,ADTTEA,%str(PPROTFL='Y' and
PARAMCD='TTCVDC1'),TRT01P,AVAL,CNSR);

Output dataset for hazard ratio: VEHR_T14020201030101. VE, VELOW, VEUPP, VEPVAL.
Output dataset for incidencerate: VEIR_T14020201030101. IR, IRLow, IRUPP, VEIR, VEIRLOW, VEIRUPP.

* with
14 da
Elec
Elec
[1] V
C
f
[2] 1
[3] P
R
e
[4] I
b
d
[5] V
e

ve RT-PCR within
days, or positive
positive RT-PCR or
using a stratified
ariate, adjusting
earliest positive
te, whichever is
risk and adjusted
t method (Poisson
culated using the

Program Path: \\wilbtia\wilbtia01\Moderna MODMRNA1273P301_U\BLA\TLF\t14020201030101.sas 10JUN2021 07:44