Table S1. Participant characteristics1 with a confirmed COVID-19 PCR test (M1 to M7). The control group (C1 to C6) represents unexposed mothers who donated human milk samples before the COVID-19 pandemic in 2018.

| Mothers | Confirmed COVID-19 PCR (Nasal Swab) | COVID-19 Symptoms | Time from Infection to Collection (Days) | Infant Gender | Lactation Time (Months) | Maternal Age |
|------------|--|--|--|------------------|----------------------------|--------------|
| M1 | +COVID19 PCR | Fever, fatigue, cough | 61 | Female | 5 | 36 |
| M2 | +COVID-19 PCR | Cough, fatigue, fever, loss of taste/smell | 60 | Female | 6 | 26 |
| M3 | +RNA SARS-CoV-2 PCR | Fever, nasal congestion, and cough | 84 | Female | 6 | 26 |
| M4 | +COVID-19 PCR | Fatigue, fever | 40 | Male | 6 | 33 |
| M5 | +COVID-19 PCR | Fever and cough | 16 | Male | 8 | 37 |
| M 6 | +RNA SARS-CoV-2 PCR | Cough and fever | 47 | Male | 8 | 31 |
| M 7 | +COVID-19-PCR | Headaches, fatigue, fever | 18 | Male | 5 | 33 |
| C1 | - | Unknown | - | Male | 8 | 28 |
| C2 | - | Unknown | - | Female | 5 | 37 |
| C3 | - | Unknown | - | Male | 7 | 29 |
| C4 | - | Unknown | - | Female | 5 | 35 |
| C5 | - | Unknown | - | Male | 5 | 33 |
| C6 | - | Unknown | - | Female | 7 | 27 |

Women did not have a medical history, medication and systematic diseases.

A S1 subunit SARS-CoV-2

S1 subunt Saks-Cov-2
PODTC2[13-685 (YP_009724390.1)
SQCVNLTTRTQLPPAYTNSFTRGVYYPDKVFRSSVLHSTQDLFLPFFSNVTWFHAIHVSG
TNGTKRFDNPVLPFNDGVYFASTEKSNIIRGWIFGTTLDSKTQSLLIVNNATNVVIKVCE
FQFCNDPFLGVYYHKNNKSWMESEFRVYSSANNCTFEYVSQPFLMDLEGKQQNFKNLREF
VFKNIDGYFKIYSKHTPINLVRDLPQGFSALEPLVDLPIGINTRFQTLLALHRSYLTPG
DSSSGWTAGAAAYYVGYLQPRTFLLKYNENGTITDAVDCALDPLSETKCTLKSFTVEKGI DSSSGWIAGAAYYVGYLQPRIFELKYNENG HIDAVDCALDPLSE IKCTLKSF I VERGI YQTSNFRVOPTESIVRFPNITNLCPFGEVFNATTFASVYAWNKRISNCVADYSVLYNSA SFSTFKCYGVSPTKLNDLCFTNVYADSFVIRGDEVRQIAPGQTGKIADYNYKLPDDFTGC VIAWNSNNLDSKVGGNYNYLYRLFRKSNLKPFERDISTEIYQAGSTPCNGVEGFNCYFPL GSYGFQPTNGVGYOPYRVVULSFELLHAPATVCGPKKSTNLVKNKCVNFNFNGLTGTGVL TESNKKFLPFQQFGRDIADTTDAVRDPQTLEILDITPCSFGGVSVITPGTNTSNQVAVLY QDVNCTEVPVAIHADQLTPTWRVYSTGSNVFQTRAGCLIGAEHVNNSYECDIPIGAGICA SYQTQTNSPRRAR

C S1+S2 HCoV-229E

AGA1.78942 (APT69883.1)
MFVLLVAYALLHIAGCQTTNGTNTSHSVCNGCVGHSENVFAVESGGYIPSNFSFNNWFLL
TNTSSVVDGVVRSFQPLLLNCLWSVSGSQFTTGFVYFNGTGRGACKGFYSNASSDVIRYN TNTSSV/DGV/RSFQPLLLNCLWSVSGSGFTGGV/FNGTGRGACKGFYSNASSDVIRYN INFEENLRRGTILFKTSYGAVFYCTNNTLVSGDAHIPSGTVLGNFYCFVNTTIGNETTS AFVGALPKTVREFVISRTGHFYINGYRYFSLGDVEAVNFNVTNAATIVCTVALASYADVL VNVSQTAIANIIYCNSVINRLRCDQLSFDVPDGFYSTSPIQPVELPMSIVSLPVYHKHTF INLHVKFEHQRGPGKCYNCRPSVINTILANFNETKGPLCVDTSHFTTGVFDNVKLARWSA SINTGNCPFSFGKVNNFVKFGSVCFSLKDIPGGCAMPIMANLVNHKSHNIGSLYVSWSDG DVITGYPKPVEGVSSFMNVTLNKCTKYNIYDDVSGVGVIISNDTFLNGTYTSTSGNLLG FKDVTNGTITYSITFCNNPDQLVYYQAVVGAMLSENFTSYGFSNVVEMPKFFYASNGTYN CTDAVLTYSSFGVCADGSIIAVOPRNVSYDSVSAIVTANLSIPSNWTTSVQVEYLQITSK PIVVDCSTYVCNGNVRCVELLKQYTSACKTIEDALRNSAMLESADVSEMLTFDKKAFTLA NVSSFGDYNLSSVIPSLPRSGSRVAGRSAIEDILFSKLYTSGLGTVDADYKKCTKGLSIA DLACAQYYNGMINLPGVADAERMAMYTGSLIGGIALGGLTSAASIPFSLAIQSRLNYVAL OTDIVLOPNOKIN LØSKAMTNIVONGON QTDVLQENOKILAASFNKAMTNIVDAFTGVNDAITGTSOALQTVATALIKKIQDVVNQQGN SLNHLTSOLRONFQAISSSIQAIYDRLDIIQADQQVDRLITGRLAALNVFVSHTLTKYTE VRASRQLAQQKVNECVKSQSKRYGFGGNGTHIFSLVNAAPEGLYFLHTVLLPTQYXDVEA WSGLCVDGINGYVLRQPNLALYKEGNYYRITSRIMFEPRIPTIADFVQIENCNVTFVNIS WSSLCVOGING VERGPILLAR HEBBYT IN STRIMFER I INDIVIDUAL SEISTLENKSAELN STRUKTIVPEVIDVNKTLOELSYKLPNYTVPDLVVEQYNOTILNLTSEISTLENKSAELN YTVQKLQTLIDNINSTLVDLKWLNRVETYIKWPWWVWLCISVVLIFVVSMLLLCCCSTGC CGFFSCFASSIKGCCESTKLPYYDVEKIHIQ

B S2 subunit SARS-CoV-2

SZ subunt SARS-COV-2
PODTC2[686-1273 (YP. 098724390.1)
SVASQSIIAYTMSLGAENSVAYSNNSIAIPTNFTISVTTEILPVSMTKTSVDCTMYICGD
STECSNLLLQYGSFCTQLNRALTGIAVEQDKNTQEVFAQVKQIYKTPPIKDFGGFNFSQI
LPDPSKPSKRSFIEDLLFNKVTLADAGFIKQYGDCLGDIAARDLICAQKFNGLTVLPPLL
TDEMIAQYTSALLAGTITSGWTFGAGAALQIPFAMQMAYRFNGIGVTQNVLYENQKLIAN
QFNSAIGKIQDSLSSTASALGKLQDVVNQNAQALNTLVKQLSSNFGAISSVLNDILSRLD GFNSAIGNIQUESS INSAIGNIQUES INSAIGNIQUES IN ANGEST IN ANGES IN A

D S1+S2 HCoV-OC43 40A2P1ZWW8 (AVR40344.1)

A0A2P1ZWW8 (AVR40344.1)

MFLILLISLPTAFAVIGDLNCTLDPRLKGSFNNRDTGPPSISIDTVDVTNGLGTYYYLDR

WYLNTTLFLNGYYPTSGSTYRNMALKGTDLLSTLWFKPPFLSDFINGIFAKVKNTKVFKD

GVMYSEFPAITIGSTFVNTSYSVVVQPRTINSTQDGVNKLQGLLEVSVCQYMMCEVPHTI

HPNLGHHFKELWHYDTGVVSCLYKKRFTYDVNATYLYFHFYGGGTFYAYFTDTGFVTK

FLFNVYLGMALSHYYVMPLTCIRRPKDGFSLEYWVTPLTPRQYLLAFNQDGIIFNAVDCM

SDFMSEIKCKTQSIAPPTGVYELNGYTVQPVADVYRRKPDLPNCNIEAWLDNKSVPSPLN

WERKTFSNCNFNMSSLMSFIQADSFTCNNIDAAKIYGMCFSSTIDKFAIPNRRKVDLQL

GNLGYLQSSNYRIDTTATSCQLYYNLPAANVSVSRFNPSTWNKRFGFIEDSVFYPQPTGV

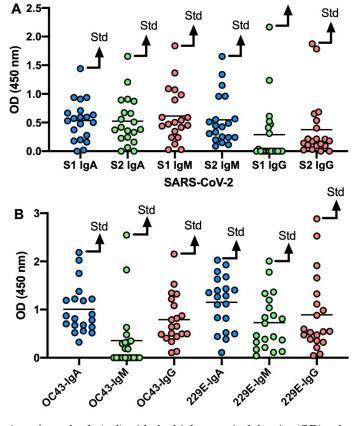
FTNHSVVYAQHCFKAPKNFCPCSSCSCPGKNNGIGTCPAGTNSLTCDNLCTLDPITLKAP

DTYKCPQSKSLVGIGEHCSGLAVKSDYGGNNSCTCQPQAFLGWSADSCLQGDKCNIFANF

I HDVMNGITCSTDI QKANTFIEI GVCLVNNIY JGJSGQGIFVEVNATYYNSWONI I YDSN

I HDVMNGITCSTDI QKANTFIEI GVCLVNNIY JGJSGQGIFVEVNATYYNSWONI I YDSN DTYKCPQSKSLVGIGEHCSGLAWKSDYCGNNSCTCQPQAFLGWSADSCLQGDKCNIFAN ILHDVNNGLTCSTDLQKANTEIELGVCVNYDLYGISGQGIFVEVNATYYNSWONLLYDSN GNLYGFRDYITNRTFMIHSCYSGRVSAAYHANSSEPALLFRNIKCNYVFNNSLTRQLQPI NYSFDSYLGCVVNAYNSTAISVQTCDLTVGSGYCVDYSKNRRSRRAITTGYRFTNFEPFT VNSVNDSLEPVGGLYEIQIPSEFTIGNMESFIOTSSPKVITDCAAFVGGYAACKLQLVE YGSFCDNINAILTEVNELLDTTQLQVANSLMNGVTLSTKLKDGVNFNVDDINFSPVLGCL YGSFCDNINAIL EVNELLD I I QLQVANSLMING YLS YKLIGOVNEN YDDDINFSPYLGGL GSECSKASSRAIEDLIFDKYKLSDVGFVEAYNNCTGGAEIRDLICVQSYKGIKVLPPLL SENQISGYTLAATSASLFPPWTAAAGVPFYLNVQYRINGLGVTMDVLSQNQKLIANAFNN ALHAIQQGFDATNSALVKIQAVVNANAEALNNLLQQLSNRFGAISASLQEILSRLDALEA EAQIDRLINGRITALNAYVSQQLSDSTLVKFSAAQAMEKVNECVKSQSSRINFGCGNGNHI ISLVQNAPYGLYFIHFNYVPTKYVTAKVSPGLCIAGNRGIAPKSGYFVNVNNTWMYTGSG YYYPEPITENNYVMSTCA/NYTKAPYVMLNTSIPNLPDFKEELDOWFKNGTSVAPDLSL DYINVTFLDLQVEMNRLQEAIKVLNHSYINLKDIGTYEYYVKWPWYVWLLICLAGVAMLV LLFFICCCTGCGTSCFKKCGGCCDDYTGYQELVIKTSHDD

Figure S1. Amino acid sequences from (A) S1 subunit and (B) S2 subunit from SARS-CoV-2, S1 + S2 subunits from (C) HCoV-229E and (D) HCoV-OC43. Sequences were from UniProt (ELIXIR core data resource).



Std

Figure S2. Selection of standards (std) with the highest optical density (OD) values using ELISA. The OD values for antibody secretory IgA (SIgA)/IgA, secretory IgM (SIgM)/IgM, and IgG reactive to (**A**) S1 or S2 subunit SARS-CoV-2, (**B**) S1+S2 subunits HCoV-OC43, or S1+S2 subunits HCoV-229E. The standards were selected among 20 human milk samples (supernatants) diluted at 10x for IgG and IgM and at 25x for SIgA/IgA in blocking buffer.

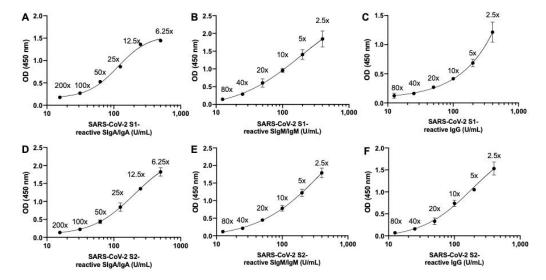


Figure S3. Standard curves used to determine the levels of antibodies reactive to SARS-CoV-2 S1 or S2 subunit. Standard curves for SARS-CoV-2 S1-reactive $\bf a$ secretory IgA (SIgA)/IgA, $\bf b$ secretory IgM (SIgM)/IgM, and $\bf c$ IgG. Standard curves for SARS-CoV-2 S2-reactive $\bf d$ SIgM/IgM, $\bf e$ SIgA/IgA, and $\bf f$ IgG. Values are means $\bf t$ SD, $\bf n$ = 3 for experiments. The standard curves were prepared using the human milk supernatant with the highest optical density values (see Figure S2).

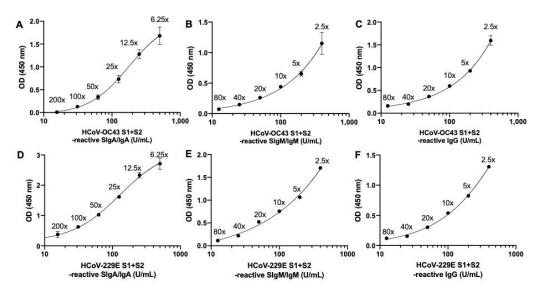


Figure S4. Standard curves used to determine the levels of antibodies reactive to HCoV-OC43 and HCoV-22E S1+S2 subunits. Standard curves for HCoV-OC43 S1+S2-reactive **a** secretory IgA (SIgA)/IgA, **b** secretory IgM (SIgM)/IgM, and **c** IgG. Values are means \pm SD, n = 3 for experiments. The standards were prepared using the human milk supernatant with the highest optical density values (see Figure S2).