

1 **Association between ACE2 and TMPRSS2 Nasopharyngeal Expression and COVID-19**

2 **respiratory distress**

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50 **Supplementary Table 1: Association between chronic use of angiotensin receptor blockers and**
51 **COVID-19 respiratory distress.**

Total Cohort (N = 213)			
	Cases (%)	Controls (%)	OR (95%CI) <i>p-value</i>
Chronic Angiotensin Receptor Blockers Use	38 (17.84)	175 (82.16)	6.94 (2.61 – 18.46) 2.07 x 10⁻⁴
Subset of Systemic Arterial Hypertension Individuals (N = 41)			
	Cases (%)	Controls (%)	OR (95%CI) <i>p-value</i>
Chronic Angiotensin Receptor Blockers Use	19 (46.34)	22 (53.66)	3.24 (0.65 – 19.09) 0.1672

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54 **Supplementary Table 2: Stepwise regression analysis for selection of covariates**
55 **associated to COVID-19 respiratory distress.**

Explanatory Variables	Deviance	AIC
Age + Systemic Arterial Hypertension + Diabetes + Obesity	122.52	130.51
- Systemic Arterial Hypertension	121.57	131.57
- Diabetes	125.80	131.80
+ Any Comorbidity	122.03	132.03
+ Sex	122.25	132.25
- Obesity	141.76	147.76
- Age	145.47	151.47

56 AIC = Akaike Information Criteria. Plus and minus symbols indicates whether the variable
57 was removed (-) or included (+) in the model.

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61 **Supplementary Table 3: Stepwise regression analysis for variables associated with COVID-19**
62 **respiratory distress including *ACE2* and *TMPRSS2* expression levels.**

Explanatory Variables	Deviance	AIC
Age + Systemic Arterial Hypertension + Diabetes + Obesity + TMPRSS2/ACE2	85.11	97.11
- Systemic Arterial Hypertension	87.21	97.21
+ ACE2	84.17	98.17
+ Any Comorbidity	84.83	98.83
+ Sex	84.99	98.99
- Diabetes	89.19	99.19
- TMPRSS2/ACE2	91.85	101.85
- Age	96.84	106.84
- Obesity	99.07	109.07

63 AIC = Akaike Information Criteria. Plus and minus symbols indicates whether the variable was removed (-)
64 or included (+) in the model.

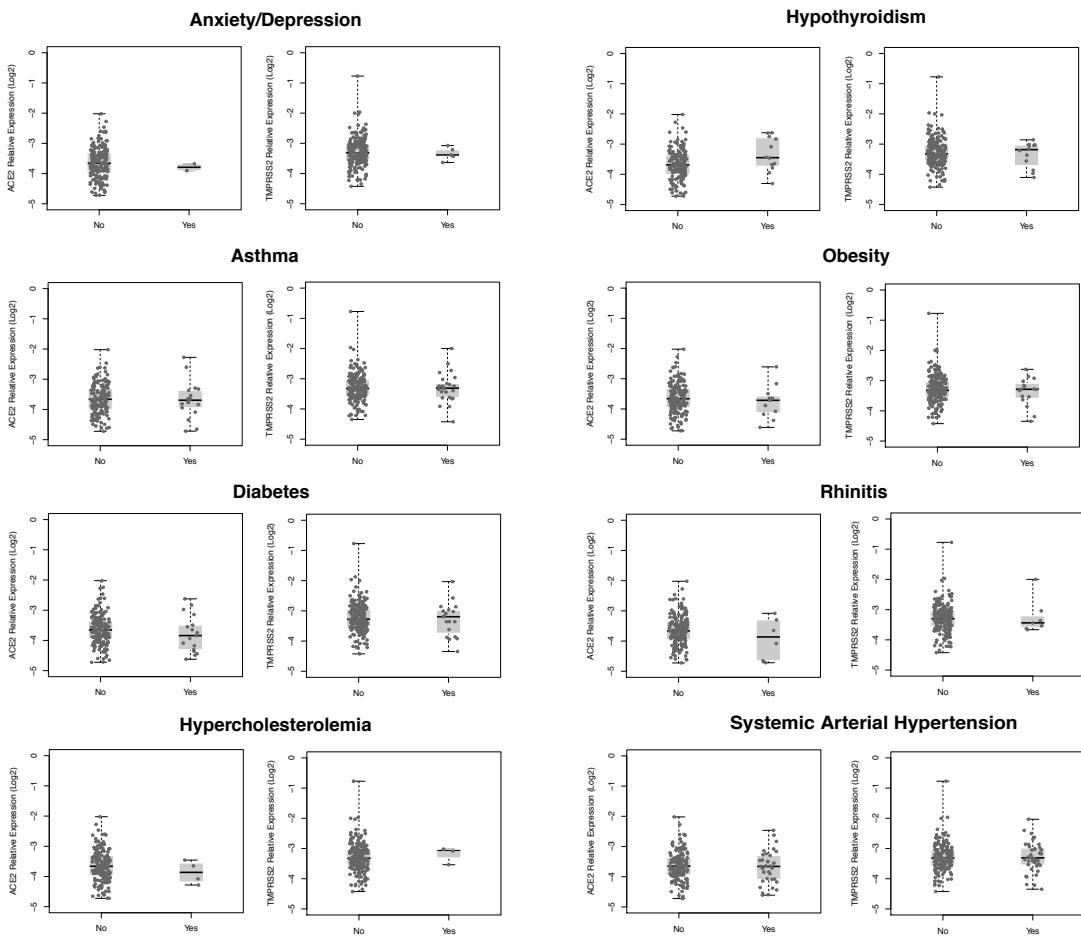
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67 **Supplementary Table 4: Characteristics of patients included in comparisons**
68 **between nasopharyngeal swab and bronchoalveolar lavage samples.**

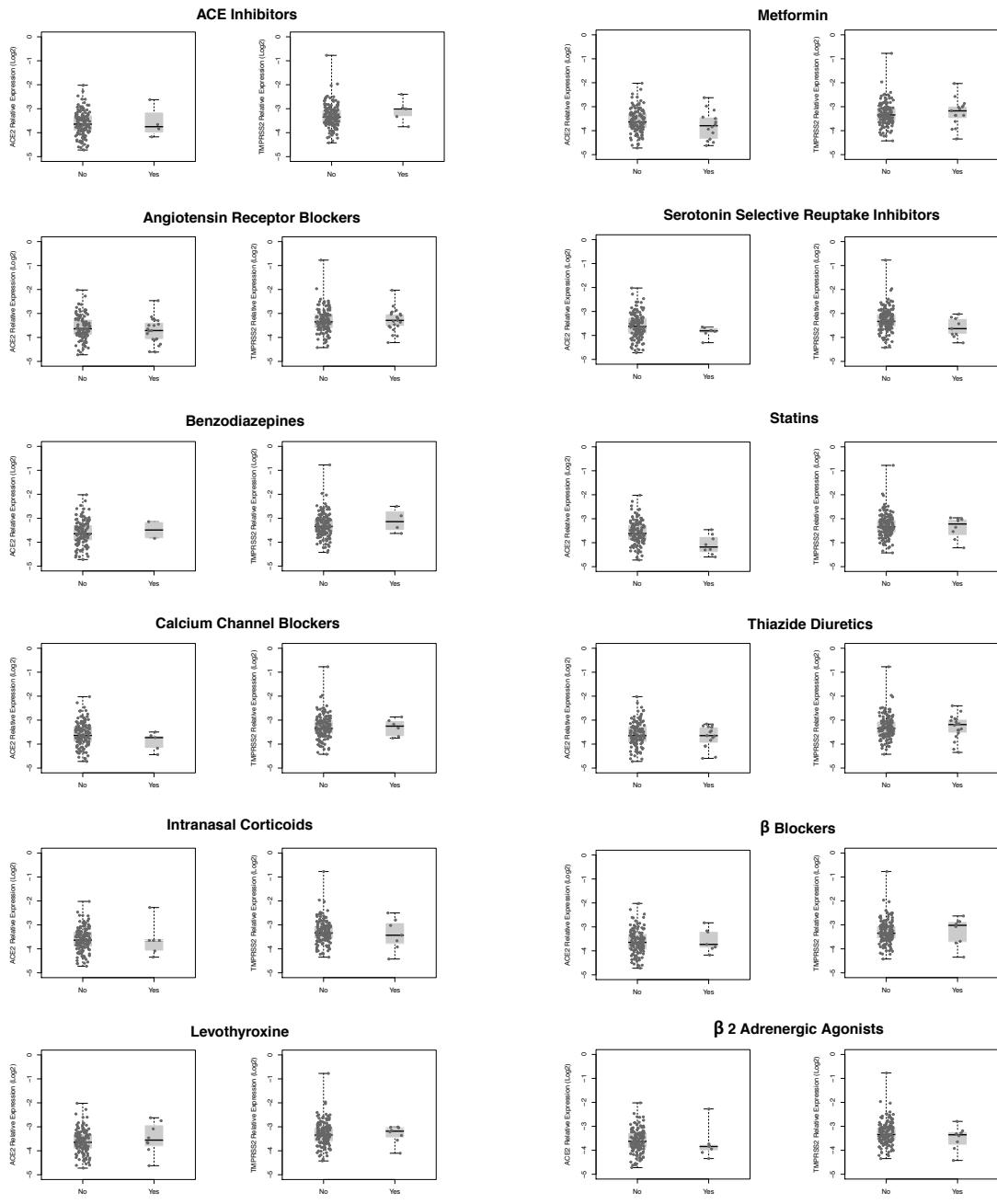
	Group	
	BAL (N=11)	SWAB (N=34)
Age*	62 (± 7)	57 (± 14)
Sex		
Male	6 (54.5%)	19 (55.9%)
Female	5 (45.5%)	15 (44.1%)
Comorbidity		
Systemic Arterial	5 (55.5%)	20 (58.8%)
Hypertension		
Obesity	3 (27.3%)	11 (32.3%)
Diabetes	5 (55.5%)	9 (26.5%)

69 *data represented as mean \pm standard deviation. BAL = bronchoalveolar lavage
70 fluid; swab = nasopharyngeal swab samples.



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72 **Supplementary Figure 1: Nasopharyngeal expression of *ACE2* and *TMPrSS2* in**
 73 **individuals with mild COVID-19 according to different chronic diseases.** Expression
 74 levels were assessed by RT-qPCR and are shown in log-scale (base 2) taking *B2M* expression
 75 as reference. Boxes represent mean and interquartile range, and whiskers stand for upper and
 76 lower limits. Comparisons between groups were made by linear regression models.
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79 **Supplementary Figure 2: Nasopharyngeal expression of ACE2 and TMPRSS2 in**
 80 **individuals with mild COVID-19 according to chronic therapy use.** Expression levels
 81 were assessed by RT-qPCR plotted in log-scale (base 2) relative to *B2M* expression. Boxes
 82 represent mean and interquartile range, and whiskers stand for upper and lower limits.
 83 Comparisons between groups were made by linear regression models.

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