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58 Abstract

59 **Context**: The current COVID-19 pandemic is probably the worst the world has ever faced 60 since the start of the new millennium. While the respiratory system is the most prominent 61 target of SARS-CoV-2 (the contagion of COVID-19), extra-pulmonary involvement are 62 emerging as important contributors of its morbidity and lethality. This article summarizes the 63 impact of SARS-CoV and SARS-CoV-2 on the endocrine system to facilitate our 64 understanding of the nature of coronavirus-associated endocrinopathy. Although new data are rapidly accumulating on this novel infection, many of the endocrine manifestations of 65 COVID-19 remain incompletely elucidated. We hereby summarize various endocrine 66 dysfunctions including coronavirus-induced new onset diabetes mellitus, hypocortisolism, 67 68 thyroid hormone and reproductive system aberrations so that clinicians armed with such 69 insights can potentially benefit COVID-19 patients at the bedside.

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74 Keywords: SARS, COVID-19, SARS-CoV, SARS-CoV-2, Endocrine system

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80 Introduction

81 Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is a highly transmissible and pathogenic novel coronavirus that emerged in late 2019 and has caused a pandemic of 82 acute respiratory disease, named 'coronavirus disease 2019' (COVID-19), which threatens 83 human health and public safety. At this time of writing, the total confirmed cases are 84 approaching 60 million round the world, with total number of reported deaths well exceeding 85 a million (1). The contagion of COVID-19 has been named 'severe acute respiratory 86 syndrome coronavirus subtype 2' or 'SARS-CoV-2', not only because it assaults 87 88 predominantly the respiratory system reminiscent of SARS and the Middle East Respiratory Syndrome (MERS), but also because genomic analysis revealed all three positive-sense, 89 90 single-stranded RNA viruses to belong under the same genus, Betacoronavirus (59). From a 91 phylogenetic perspective, SARS-CoV-2 and SARS-CoV are both of the same clade (115). 92 Hence, the biology and clinical aspects of SARS can inform and serve as useful yardsticks for 93 predicting and extrapolating the behavior of COVID-19.

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Despite its name, there is cogent evidence that SARS-CoV has extra-pulmonary 95 96 manifestations (62). During the SARS outbreak in 2003, many patients suffered sequelae in 97 the gastrointestinal tract (86), the cardiovascular system (56), the coagulation system (61), the 98 immune system (114), the nervous system (72) and even the endocrine system (66). Indeed, it 99 is very likely that systemic viremia and an over-reactive immune response contributed to the 100 pathogenesis of lesions in key endocrine glands (Fig. 1). In the same vein, there is wisdom of 101 hindsight distilled from historical parallels in guiding the manner we tackle this ongoing 102 crisis. By contrast, MERS, while exacting a higher fatality rate compared to SARS and 103 COVID-19, had not been associated with overt endocrine sequelae; despite the broad tissue

distribution of dipeptidyl peptidase-4 (DPP4) receptors which serves as the portal of cell entry for MERS-CoV, this coronavirus has not been detected in any endocrine tissues in an autopsy study correlating clinicopathologic, immunohistochemical, ultrastructural and molecular findings (69). Combining our experience combating both SARS and COVID-19 at the forefront with updated literature, we present this timely article devoted to features of SARS-induced endocrinopathy to better understand and predict COVID-19 effects on target organs of the endocrine system.

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112 Relevant coronavirus structural and molecular biology

113 Coronavirus virion particles are spheroidal with dimensions $\sim 80-120$ nm across, and possess 114 morphologically characteristic clubbed-shaped peplomers otherwise termed 'S' spike 115 glycoproteins extending 17-20 nm outwards from the lipid bilayer surface envelope 116 reminiscent of a 'crown' on electron microscopy and atomic force microscopy (6, 46, 90) (Fig. 2A). To aid further understanding of the predilection of SARS-CoV and SARS-CoV-2 117 118 for the human host, it is necessary to appreciate the structural details of the S spike 119 glycoprotein. Each S protein comprises 3 monomers fused as a trimer to form the spike (87). 120 The molecularly divergent distal 'bulb' amino-terminal half portion is the S1 fragment 121 critical for binding to host cell surface receptors (103) while the highly conserved 'stalk-like' 122 carboxy-terminal half portion is the S2 fragment of the spike protein that has both an 123 ectodomain and a transmembrane domain responsible for fusion to the host cell membrane 124 (24) (Fig. 2B). Cathepsins are involved in the cleavage of S into S1 and S2 subunits to expose 125 S2 for fusion to cell membrane via host proteases (12). This process is accomplished by the 126 synergistic activity of cathepsins along with other host proteases, cell surface transmembrane protease serine (TMPRSS) proteases, furin, trypsin and factor Xa, following which the 127

128 internalized virion undergoes replication within the infected host cell and finally exits the cell via a lysosomal-based exocytosis pathway to complete its life cycle (Fig. 2C). The lipid 129 130 bilayer multi-spanning M glycoprotein through which the S spike protein structure is inserted 131 and anchored, represents the largest constituent of the virion. The coronavirus M protein is 132 interestingly the first polytopic viral membrane ever to be described in the virology field (9), and its glycosylation status plays a role in organ tropism (25) and possesses a capacity of 133 134 alpha-interferon induction (50). The E protein of SARS-CoV is an integral membrane protein while the N protein is a phosphoprotein that binds to viral RNA in a helical nucleocapsid 135 136 conformation and enhances replication efficiency (5).

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138 Cell invasion Gateways for COVID-19 and similarities to SARS

139 I. Angiotensin-converting enzyme-2 (ACE2)

140 Coronaviruses exploit host cell membrane protein receptors to gain entry into the interior of 141 the cell. The most well described gateway for both SARS-CoV and SARS-CoV-2 is 142 angiotensin-converting enzyme 2 (ACE-2). ACE2 is a type I transmembrane zinc-dependent 143 monocarboxypeptidase with homology to ACE, a key player in the renin-angiotensin-144 aldosterone system (RAAS) and a target for the treatment of hypertension because it cleaves 145 angiotensin-II into angiotensin-(1-7) which binds Mas-receptors to negatively regulate the 146 RAAS (80). Unlike MERS-CoV, which engages surface dipeptidylpeptidase-subtype 4 (DPP4) and sialoside attachment receptors for host cell entry (73), SARS-CoV and SARS-147 148 CoV-2 seek out ACE2 as receptors for cell invasion (101). ACE2 is abundantly expressed in 149 human kidneys, adrenals, adipose tissues, thyroid, endothelium, pancreas, testis, ovary and 150 pituitary (37, 94). ACE2 possesses highly similar binding motifs for the S protein 151 indispensable for SARS-CoV and SARS-CoV-2 invasion (31, 54, 101).

152 ACE2 interaction with the spike protein was first made available for SARS-CoV shortly after 153 the 2002-2004 SARS outbreak, and had since been steadily built up for more than a decade, 154 before being accumulated explosively in response to the 2019 SARS-CoV-2 pandemic (34, 155 48, 54, 85, 88, 101, 110). Such information details both similarities and differences between 156 SARS-CoV and SARS-CoV-2 in their receptor binding (48, 54). These two viruses both 157 utilize their receptor-binding domains (RBDs), residing at the C-terminal half of the S1 158 fragment to bind ACE2 with nanomolar affinities; the binding sites on ACE2, located at the N-terminal peptidase domain, are nearly identical. Major differences lie within the two 159 160 viruses' respective ~70 residue-long receptor-binding motifs (RBMs), which are extended 161 insertions grafted onto the core of RBD, sharing about 50% sequence identity and adopting 162 different conformations upon ACE2 binding (Fig. 3A and 3B). Consequently, the viruses' 163 ACE2 binding affinities are folds apart (20-30 folds), with SARS-CoV-2 being the tighter 164 binder (48, 85, 101, 107), which corroborates with the drastically higher transmissibility of 165 SARS-CoV-2 (57). Taking SARS-CoV and MERS-CoV as examples, moderate changes in 166 their RBM sequences have led to zero cross-reactivity against the receptor for the other (58).

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II. Transmembrane protease serine-2 (TMPRSS2)

TMPRSS2 facilitates cellular entry of SAR-CoV and SARS-CoV-2 into the host. TMPRSS2 is an androgen-regulated gene (67, 102) involved in the priming of the viral spike protein. This process is critical for virulence as it diminishes viral recognition by neutralizing antibodies and also helps in activating SARS-CoV-2 for virus cell fusion. That the activated androgen receptor regulates the transcriptional activity of the TMPRSS2 gene could partly explain the differential susceptibility of males for COVID-19 (20, 65). TMPRSS2 works synergistically along with 'a disintegrin and metalloproteinase domain-containing protein-17' (ADAM17) required for the shedding of ACE2 (108). It has been shown that SNPs
in TMPRSS2 might influence SARS-CoV-2 entry into the cell (4). Some newly reported
target sites of COVID-19 mediated by TMPRSS2 are hepatobiliary and pancreatic tissues
(74) which deserves further mention below.

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III. Neuropilin-1 (NRP1)

182 NRP1, a highly conserved type 1 transmembrane protein, plays important roles in the 183 development of the nervous and cardiovascular system as well as in tumorigenesis through 184 interaction with its established binding partners, such as vascular endothelial growth factor 185 (VEGF) and semaphorin 3A (Sema3A). Cell entry of SARS-CoV-2 depends on priming by 186 host cell proteases (42, 63). There is limited knowledge about the virus-host interactions that 187 determine cellular entry of SARS-CoV-2. Viruses display considerable redundancy and 188 flexibility because they can exploit weak multivalent interactions to enhance affinity. While 189 the focus to date has been almost entirely on the role of ACE2 in SARS-CoV-2 entry, the 190 expression pattern of ACE2 does not match tissue tropism of SARS-CoV-2 (41). This raises 191 the possibility that co-factors are required to facilitate virus-host cell interactions in cells with 192 low ACE2 expression. In the case of SARS, the C-type lectin receptor CD209L (L-SIGN) 193 was found to serve as an alternative gateway for the cellular entry of SARS-CoV by tethering 194 to the mannose glycans of the S-protein (44). Recently, by applying site-directed mutagenesis 195 and monoclonal antibodies, it was shown that NRP1 could represent such an ACE2 196 potentiating factor (23); the high expression of NRP1 on epithelial cells strategizes it as an entry receptor (77). Reports demonstrated that the SARS-CoV-2 S protein binds to the b1b2 197 198 ectodomain of the NRP1 (Fig. 3C). The subsequent entry of SARS-CoV-2 into cells is facilitated by a polybasic amino acid sequence (⁶⁸²RRAR⁶⁸⁵) termed the 'C-end rule' (CendR) 199 200 motif within NRP1 (93). Corroborating evidence for its putative role as an entry receptor is 201 its upregulation in COVID-19 biological samples versus healthy controls as shown by 'omic' 202 analyses (93). That SARS-CoV-2 shows a much greater infectivity relative to SARS-CoV is 203 probably explained in part by NRP1 binding to the CendR peptide in S1. Such a discovery 204 meant that novel therapeutic approaches that target this mechanism could be developed. From 205 the Human Protein Atlas, we find medium expression of NRP1 in the parathyroids, adrenals 206 and testis, and low expression of NRP1 in the thyroid. As VEGF is a natural endogenous 207 ligand of NRP1, it is also possible that the pituitary could well a target of SARS-CoV-2 since 208 VEGF receptors are also present in the pituitary gland (70).

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210 Clinico-pathological spectrum of SARS and COVID-19 endocrinopathy

211 I. Hypothalamus-Pituitary-Adrenal Axis

212 The lack of detailed postmortem studies contributed to the paucity of knowledge surrounding 213 the pathogenesis of endocrinopathy associated with SARS and COVID-19. There is only a 214 dearth of reports of involvement of the hypothalamus and pituitary by SARS (33, 51). Using 215 light microscopy, electron microscopy and real-time RT-PCR, autopsy series previously 216 confirmed SARS-CoV genome sequences associated with cytopathic effects in neuronal 217 cytoplasm of the hypothalamus. Clinical observation of an impaired ACTH and TSH 218 response to hypocortisolism and hypothyroidism respectively implied that the hypothalamus-219 pituitary-adrenal (HPA) axis was probably involved either directly by SARS-CoV or indirectly due to hypophysitis caused by autoimmunity triggered by the virus (33). Little has 220 221 been documented on the hypothalamic-pituitary effects of COVID-19, though a French group 222 has recently confirmed from autopsy findings that the hypothalamus is a highly probable 223 target of SARS-CoV-2 based on its rich expression of ACE2 and TMPRSS2, especially in the paraventricular nucleus (89). CT and MRI imaging have also revealed evidence of COVID-224

225 19 infecting the brain with serious consequences (78). A Chinese group successfully detected 226 the presence of SARS-CoV-2 genome in the cerebrospinal fluid in a patient with COVID-19, 227 thereby confirming that SARS-CoV-2 does indeed infiltrate into the brain, and thence can 228 involve any part of the brain, including the hypothalamus and pituitary (113). While HPA 229 axis compromise with hypocortisolism can contribute to mortality in SARS and COVID-19, 230 an overly excessive endogenous cortisol response itself poses a caveat, lest high cortisol be 231 misinterpreted to portend a better prognosis. A group has shown via Cox proportional hazards regression analysis that cortisol stress responses were predictive of death. Kaplan-Meier 232 233 survival analysis revealed a sharp dichotomy in death probability, with significantly better 234 median survival for serum cortisol lower than 744 nmol/L during acute COVID-19 infection, 235 a scenario correlative of illness severity. This should not be misconstrued as justification to 236 avoid prescribing exogenous steroids when there are overwhelming life-saving indications as 237 discussed in the next section (92). Several recent publications have also confirmed that the adrenals are a frequent site of COVID-19 related lesions in the body based on radiological 238 239 and autopsy evidence (7, 28, 38, 52).

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II. Hypothalamus-Pituitary-Thyroid Axis

The thyroid is another endocrine gland reportedly disordered in SARS previously (51, 105) and now in COVID-19 (14). In 2004, a group from Guangzhou in China produced postmortem evidence of cytopathic effects of SARS-CoV on endocrine organs including the thyroid, parathyroid, pancreas and adrenals using a combination of RT-PCR, immunohistochemistry, in situ hybridization and transmission electron microscopy (26). Among those who deceased from SARS, the thyroids showed destruction of the follicular epithelium with extensive exfoliation of apoptotic cells into the follicular lumen. Thyroid follicular damage was occasionally very severe, associated with complete loss of parafollicular C-cells as shown by total absence of calcitonin immunostaining. This explains why serum T3 and T4 were decreased in 94% and 46% respectively in a group of SARS patients during the acute phase of the disease, followed by persistence of low serum T3 and T4 in 90% and 38% among convalescent cases (104). Autopsies of SARS patients shown marked destruction of the follicular and parafollicular cells of the thyroid (105).

256 The ongoing COVID-19 pandemic yields interesting and important insights on SARS-CoV-2 257 and thyroid pathology. The available data suggests that the disease spectrum can range from 258 direct viral destructive effects to immune-mediated mechanisms on the thyroid precipitated 259 by COVID-19 (49, 60, 68, 83). Leow et al. showed previously that SARS-CoV could inflict 260 pituitary lesions either directly or indirectly and contribute to secondary thyroid and adrenal 261 insufficiency, which could be treated using levothyroxine and hydrocortisone replacement 262 (51). It appears that the SARS-induced thyroid aberrations were largely transient and fully 263 resolved after several months. Current published data indicate a similar pathophysiological 264 phenomenon associated with COVID-19. Mounting evidence points to the fact that COVID-265 19 might have a greater impact on the hypothalamus-pituitary-thyroid axis than previously 266 suspected. This is particularly so following the discovery of ACE-2 mRNA in thyroid cells 267 (81). ACE-2 mRNA in thyroid follicular cells was confirmed by analyzing primary cultures 268 of thyroid cells, where the expression is similar to those found in tissues. The finding 269 accounts for the recently described COVID-19-related subacute thyroiditis or De Quervain's 270 thyroiditis that is often thought to have a viral origin (81); it can present with thyrotoxicosis 271 before a hypothyroid phase sets in weeks to months later (13). Hence, subacute thyroiditis is 272 now considered to be a sequela closely associated with COVID-19 (15). Other reports also 273 indicate autoimmune thyroiditis may develop after the "cytokine storm" induced by SARS-

CoV-2 infection which could result in the development of primary hypothyroidism. Common thyroid manifestations of COVID-19 thus include overt thyrotoxicosis, Graves' orbitopathy and hypothyroidism. Graci et al., 2020 (32), in a recent report has even suggested that COVID-19 could be considered as an endocrine disorder, to make sense of the non-specific response of the immune system to the SARS-CoV-2 virus, which is far different from infections such as influenza (76).

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281 III. Reproductive Axis

282 Next on the list of endocrine gland targets of the coronavirus is the testis which has been 283 established to express ACE2 and NRP1 copiously (95). In addition, the MAS receptor is 284 present on the acrosome and tail of human spermatozoa and plays a role in the acrosome 285 reaction and the maintenance of sperm motility, thereby underscoring the role of ACE2 in 286 sperm biology (97). The release of TMPRSS2 in prostasomes secreted into semen from the 287 prostate during ejaculation together with ACE2 present on the sperm plasma membrane 288 would thus allow SARS-CoV-2 to infect sperm cells (19). With respect to coronavirus invasion of the testis, there were men with symptoms consistent with acute orchitis in some 289 290 male cases of SARS (109). Similarly, at least one case of orchitis has been reported in a 291 young man with COVID-19 (47). Postmortem examination of men who succumbed to 292 COVID-19 interestingly also revealed seminiferous tubular injury, vacuolation of Sertoli 293 cells, reduced Leydig cells and lymphocytic infiltrates in 11 of 12 deceased males, of whom 294 one had demonstrable SARS-CoV-2 by RT-PCR within testicular tissues (64). How SARS-295 CoV-2 might inhibit sperm motility, permanently damage the testis and negatively influence 296 fertility remains undetermined. A subsequent study confirmed the frequent presence of SARS-CoV-2 in semen of men with acute COVID-19 as well as those convalescing from it 297

298 (53). This finding, together with the significantly high rates of COVID-19 infection between 299 sex partners, implies the possibility of a sexual route of transmission, though there is no 300 current strong evidence supporting such a route of transmission as one of clinical concern as 301 yet (2). In females, ACE2 is expressed in the ovary, uterus, placenta, vagina and breast 302 tissues. ACE2 is present in ovarian stroma, granulosa cells and oocytes (75), and ACE2 303 mRNA has been shown to be detectable in the ovaries of pre-menopausal and post-304 menopausal women (79). Unlike males, TMPRSS2 appears to be absent in human oocytes 305 which means that infection of the female germline by SARS-CoV-2 is rather improbable 306 except in the situation where the ovum is fertilized by an infected spermatozoon. Despite 307 such theoretical concerns, there have not been any published reports of teratogenic effects 308 and embryopathy directly attributable to SARS-CoV-2 as yet in contrast to Zika virus 309 infection (106).

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IV. Endocrine Pancreatic Islets

312 More recently, de novo onset of severe diabetes mellitus and diabetic ketoacidosis among 313 COVID-19 patients who were formerly healthy and non-diabetic stoked fears of permanent 314 beta cell damage from a single episode of COVID-19 (17). This is not unfounded given the 315 high expression of ACE2 in the pancreatic islets, and previous encounters of new-onset 316 diabetes as a sequela of SARS (111). Importantly, the downregulation of ACE2 by SARS-317 CoV-2 can lead to unopposed angiotensin-II activity on AT-I receptors which suppresses 318 insulin secretion (99). Additionally, because it was reported a decade ago that an abnormal 319 allele of NRP1 in pancreatic beta cells led to type 1 diabetes, this could imply that the binding 320 of the S protein of SARS-CoV-2 to NRP1 in pancreatic islets might potentially cripple the 321 insulin secretory pathway and trigger insulin-dependent diabetes or even overt diabetic ketoacidosis (39). A global CoviDiab registry has just been initiated to examine beta cell injury and insulinopenia in those without pre-existing diabetes in whom SARS-CoV-2 is the only etiologic factor (82). These data suggest the importance of proactive monitoring for such endocrine dysfunction among COVID-19 patients (66).

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327 Clinical course of COVID-19 among patients with pre-existing endocrine issues

328 Available reports indicate that individuals with pre-existing endocrine disorders are at higher 329 risk of suffering greater disease severity during COVID-19. Interestingly though, patients 330 with pre-existing hypothyroidism and receiving thyroid hormone therapy, they were not 331 found to be associated with an increased risk of hospitalization (98). However, patients with 332 other endocrine conditions such as hypocortisolism and diabetes mellitus appeared to have 333 poorer prognosis. Risk factors for hospital admission among COVID-19 patients with 334 diabetes include older age, higher HbA1c level, hypertension, cardiovascular disease, 335 cerebrovascular disease, chronic pulmonary disease, malignancy, chronic kidney disease 336 (CKD 3A, 3B, 4), and insulin-treated patients were more likely to require hospital admission 337 (3).

338

Contributory causes included addisonian crisis and impaired immunity (71), which makes these patients more vulnerable to SARS-CoV-2 infection (18). Serum biomarkers (IL-6, serum ferritin, CRP) and D-dimers are also higher in SARS-CoV-2 patients with underlying diabetes (35). Although clinicians might be deterred from prescribing corticosteroids in accordance with earlier conservative treatment guidelines for COVID-19 (100), glucocorticoids should be instituted in those with features of hypocortisolism on a case-bycase basis, especially since addisonian crisis can be equally life threatening if missed or

untreated (8). Intriguingly, emerging promising data from an ongoing clinical trial, 346 Randomized Evaluation of COVID-19 Therapy, otherwise codenamed RECOVERY 347 348 (ClinicalTrials.gov ID: NCT04381936) showed that low-dose dexamethasone is a life-saving 349 drug and reduced deaths by 20% for COVID-19 patients requiring oxygen, and even lowered 350 the risk of death for those on mechanical ventilation by approximately 30% which is very 351 impressive. While speculative, low-dose dexamethasone for severe COVID-19 patients 352 worldwide can thus be life-saving if prescribed at the right timing during the disease course 353 (21). An expected corollary of dexamethasone might be aversion of collateral damage to 354 endocrine glands via suppression of a cytokine-driven hyperinflammatory response during 355 the course of COVID-19 infection.

356

357 Among pre-existing endocrine issues, diabetes mellitus probably tops the list, simply because 358 diabetes itself is a worldwide pandemic and it confers an increased mortality in the face of 359 COVID-19 (91). It has been found that SARS-CoV-2 virus is more prevalent and severe in 360 people with diabetes. A recent case-control study demonstrated the virus infection could lead 361 to significant insulin resistance, dehydration and acute kidney injury. In rare cases patients 362 could also develop diabetic ketoacidosis (43). Hyperglycemia predisposes to bacterial and 363 viral pathogens such as tuberculosis and influenza (16), since hyperglycemia could trigger the 364 hyper-virulence of certain pathogens (22). Additionally, glycemic control tends to be worse 365 among diabetic patients treated with corticosteroids and lopinavir/ritonavir, which can result 366 in cytokine-induced insulin resistance as well as hypokalemia, which can impair insulin 367 secretion itself. Evidence also suggests a heightened tendency of ketoacidosis among 368 COVID-19 patients with pre-existing diabetes (55, 82).

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Related to diabetes mellitus is obesity, in which a recent study revealed that obesity 370 371 correlated to higher odds of mechanical ventilation and in-hospital mortality after 372 multivariable adjustment on analyzed data from COVID-19 patients at 88 hospitals enrolled 373 in the American Heart Association's COVID-19 Cardiovascular Disease Registry (40). Notably, among patients ≤ 50 years of age, BMI ≥ 40 kg/m² was linked to a profoundly 374 elevated risk of death or mechanical ventilation (OR, 1.64 [95% CI, 1.23-2.21]), moderately 375 376 increased odds in those aged 51-70 (OR, 1.40 [1.10-1.80]), but no significant increase in risk 377 among patients > 70 years old (OR, 1.28 [0.83-1.95]). Venous thromboembolism and dialysis 378 were also linked to higher BMI, all of which suggest that the generally lower probability of 379 morbidity and mortality in younger people with COVID-19 may not necessarily apply to 380 those with obesity.

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382 The mechanisms of how increased adiposity complicates the clinical course of COVID-19 383 remain to be elucidated, but is likely to be multifactorial and related to the impact of excess 384 fat on metabolism, immune response and vascular function. Whether direct viral infection of 385 adipocytes play a mechanistic role in COVID-19 severity among obese people is open to 386 question. In this regard, it is remarkable that adipose tissues show the highest level of NRP1 387 mRNA transcripts among all the organs of the human body examined, as shown by a mean 388 protein-coding transcripts per million (pTPM) of 101 in the Genotype-Tissue (GTEx) RNA-389 seq database and 388.6 Scaled Tags per Million in another public dataset, the Functional Annotation of Mammalian Genomes 5 (FANTOM5) Cap Analysis of Gene Expression 390 391 (CAGE).

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393 Correspondingly, a previous study on rats also demonstrated the expression of NRP1 on the 394 surface membranes of adjocytes and parenchymal nerves found within adjose tissues (11, 395 30). Adipose tissues also rank among the top ten tissues with the highest abundance of ACE2 396 mRNA based on a pTPM of 8.8 in GTEx and an estimated 5.6 scaled tags per million in 397 FANTOM5 CAGE (96). Consistent with this is the finding of ACE2 protein expression in 398 adipocytes (36) as well as TMPRSS2 within adipose tissues (10). There is thus biological 399 plausibility for higher host susceptibility to more severe COVID-19 among those with greater 400 adiposity relative to those who are lean.

401

As this COVID-19 pandemic evolves, an increasing number of infected patients are observed to be afflicted with a range of different endocrine disorders worldwide. Thus, collaborations between experts from different countries are crucial as the collective wisdom and best practices shared across geographical boundaries can help tackle this pandemic better, such as illustrated by a recent survey of clinical endocrinologists (29). This raises the following important research questions that demand further investigations:

Does SARS-CoV-2 invade the hypothalamus directly and induce endocrine
dysfunction, given the fact that other coronaviruses have been proven to exhibit
neurotropism through their invasion into the central nervous system, and that human
pluripotent stem cell-derived dopaminergic neurons can be infected by SARS-CoV-2
in vitro (112)?

Is SARS-CoV-2 capable of migrating in a retrograde fashion along the olfactory
 pathway and spread trans-synaptically into the hypothalamus, as this is one
 established route of entry into the central nervous system exploited by other

416 coronaviruses (27), and especially because anosmia is a very common feature of those
417 infected by the SARS-CoV-2 virus?

- 418 Are the parathyroid glands also susceptible to SARS-CoV-2, and if so, would
 419 hypocalcemia be a potential complication in certain cases of COVID-19?
- Do adipose tissue depots serve as a reservoir for the coronavirus among asymptomatic
 carriers with obesity, and whether SARS-CoV-2 undergo a lytic or lysogenic cycle if
 it infects white adipocytes?
- Are there any drugs or biologics approved for the treatment of endocrine and
 metabolic diseases potentially able to be repurposed as anti-viral agents against
 SARS-CoV-2?

It is imperative to address such questions with scientific rigor and unravel the molecular mechanisms, especially since this pandemic is still very rampant in many countries and reports of an increasing range of endocrinological aberrations associated with COVID-19 continue to be added to the current literature. Greater clarity of COVID-19-induced endocrinopathies will expectedly aid diagnostic pathways, therapeutic decisions and drive clinical management with ultimate benefit to patients suffering from this viral assault.

432

433 Conclusions

Although COVID-19 is very widespread, much of its endocrine manifestations are still far from being fully elucidated. Knowledge of SARS-associated endocrinopathy forms a basis for better understanding COVID-19 endocrinopathy. Meanwhile, scholarly contributions on endocrine issues in COVID-19 should be facilitated, while peer-reviewed journals might consider publishing more worthy papers that add useful insights to the scarce literature in this emerging field. Physicians, endocrinologists and even patients with different types of endocrine conditions can get more up-to-date information regarding the endocrine
manifestations of COVID-19 or the effects of COVID-19 on any pre-existing endocrine
disorders from the various national, regional and international endocrine societies and thyroid
associations.

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445 Author contributions statements

The authors' responsibilities were as follows: NK was involved in the literature review, 446 preparation of manuscript and construction of figures illustrations, assisted with the 447 references and critically reviewed the manuscript; RA performed the literature review, 448 449 assisted with figures illustrations, provided intellectual inputs and critically reviewed the 450 manuscript; XB performed in silico molecular modeling, co-wrote and critically reviewed the 451 manuscript; WSY provided intellectual inputs and critically reviewed the manuscript; SSV 452 provided intellectual inputs, edited and critically reviewed the manuscript; NKarnani 453 provided intellectual inputs, edited and critically reviewed the manuscript; MKSL conceived 454 the review topic, initiated the writing of the manuscript, interpreted the data, steered the direction of the discussion, provided intellectual inputs, edited and critically reviewed the 455 456 manuscript.

457

458 Figure legends

459 Figure 1. The endocrine system as a target of betacoronaviruses (eg. SARS-CoV).

Figure 2. Panel A shows the virion structure of SARS-CoV and SARS-CoV-2, while panel

B shows the similarities and differences in the amino acid sequences of the two viruses. Panel

462 C illustrates the life cycle of these two coronaviruses.

463	Figure 3.	ACE2 binding by	the RBDs of SARS-CoV	(Fig. 3A)	(PDB ID: 2AJF)) and SARS-

- 464 CoV-2 (Fig. 3B) (PDB ID: 6M0J). The RBD cores are in green, whereas the RBMs are
- highlighted in gold. Major conformational differences between the RBDs are circled with
- 466 dashed lines. In Fig. 3C, the interaction between the C-terminus of SARS-CoV-2 S1 (residues
- 467 679-685, NSPRRAR, yellow sticks) and human transmembrane receptor Neuropilin-1
- 468 (electrostatic surface), based on recently published complex structure (PDB ID: 7JJC) (23) is
- 469 illustrated. These figures are generated using PyMOL (84), with the APBS plugin for
- 470 electrostatic surface calculations (45).
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472 References

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Figures



Figure 1.



Figure 2. A-C



А

В



С

Figure 3 A-C

COVID-19 Endocrinopathy with Hindsight from SARS

METHODS

OUTCOME



CONCLUSION Akin to SARS, the endocrine system is also affected by COVID-19. More research is needed to elucidate the mechanisms of