- 1 Rethinking Aptamers as nanotheranostic tools for SARS-COV-2 and COVID-19 infection
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- 9 Abstract

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- 10 The novel coronavirus named by WHO and Coronavirus Study Group (CSG) as SARS-COV-2 is
- the etiological agent of the newly emerged Coronavirus disease (COVID-19). COVID-19 has
- become a pandemic threat as the WHO declared it a public health emergency of international
- 13 concern. Early and precise detection of the virus is important for effective diagnosis and
- 14 treatment. Various testing kits and assays, including real-time reverse Transcriptase PCR, thermal
- screening guns, ELISA-based immunoassays, and Point-of-Care (POC), have been implemented
- or are being explored to detect the virus and/or characterise cellular and antibody responses to the
- infection. However, these approaches have inherent limitations such as non-specificity, high cost,
- 18 characterize by long turnaround times for test results, and can be labour intensive. Aptamers,
- 19 which are single-stranded oligonucleotides, generated artificially by SELEX (Evolution of
- 20 Ligands by Exponential Enrichment) may offer the capacity to generate high affinity bioprobes
- 21 for monitoring relevant SARS-COV 2 and COVID-19 biomarkers. This article discusses the
- 22 prospects of implementing aptasensing technologies for rapid point-of-care detection of SARS-
- 23 COV-2.

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- 24 **Keywords**: Aptamers; Theranostics; SARS-COV-2; COVID-19; Bioaffinity
- 27 **1.0 COVID-1** 
  - 1.0 COVID-19, SARS -CoV 2 Structure and Interaction
- 28 Severe acute respiratory syndrome-coronavirus-2 (SARS-COV-2) is the pathogen that is
- 29 currently causing a worldwide public health threat. The infection caused by the virus is

30 named by WHO as coronavirus disease 2019 (COVID-19). COVID-19 started in China, in a city called Wuhan located in Hubei province. The disease starts feverish symptoms, 31 progresses to severe respiratory challenges, and finally pneumonia-like symptoms 1. 32 According to WHO situational report-70, as at April 27, 20020, a total of 2, 858 635 33 cases had been confirmed globally with 196 295 deaths. The mortality rate stands at 34 7.2%. globally. Since the emergence of the 2019 novel coronavirus outbreak, laboratories 35 across the globe are sequencing the full genome of SARS-CoV-2 virus at extraordinary 36 rate. This has enhanced real-time progress and research in the comprehension of the virus 37 origin and genetic evolution, virulence, epidemiology, and transmission pathway. Also, it 38 has fast tracked diagnostics design and development, treatment regime and provisional 39 therapies. Apparently, over 11,000 viral genome sequence of new virus has been shared 40 41 via GenBank and Global Initiative to Share All Influenza Data (GISAID) platform<sup>2</sup>. 42 Currently, the information from genomic sequencing and analysis shows that SARS-CoV-2 is highly linked to a familiar or common bat SARS-like coronavirus, suggesting 43 bats as the origin or reservoir <sup>3</sup>. The available genome sequences SARS-CoV-2 do not 44 vary and proves their appearance and escalation in the human populations in late 45 December 2019 <sup>2, 3</sup>. A collection of the related SARS-CoV-2 infections indict human-to-46 human transmission and provide further information on the virus evolution<sup>4</sup>. Despite the 47 48 bat origin, the mutational variants identified from the genomic sequencing are know to infect other animals such as ferrets and pangolin<sup>5</sup>, which are sold illegitimately in china 49 as ingredient to make traditional medicine. Thus, pangolin is suspected to mediate SARS-50 CoV-2 transmission between bats and humans. There is also some conjecture on the 51 emergence of SARS-CoV-2, which was created or synthesized from coronaviruses in 52 existence, however there is no supportive proof for this theory<sup>6</sup>. 53 The pathogen causing COVID-19 has been identified to be similar to the virus that 54 caused Severe Acute Respiratory syndrome (SARS) in 2002 with about 800 people losing 55 their life worldwide<sup>7</sup>. The SARS coronavirus, usually referred to as SARS-COV, is an 56 RNA virus that is positively stranded in nature with about 29727 nucleotides in the 57 genome and possesses structural proteins. The structural proteins are made of an inner 58 nucleocapsid (N) surrounded by an outer layer of envelope (E) and membrane (M) 59 proteins with glycoprotein spike (S) protruding from the surface<sup>1</sup>. A structural and 60

61 functional analysis of the N, E and S proteins indicated that SARS-COV and SARS-COV-2 have significant similarities and differences, and the S and N proteins provide 62 stability to the virus. Both SARS-CoV-2 and SARS-CoV belong to the coronavirus 63 family, specifically from β-coronavirus genera. A study by Xu, et al. 8 showed that, two 64 viruses share almost same gene sequence with 85% similarity, especially at the nucleotide 65 level. Their genomic sequence containing six region of difference (RD) partially code for 66 both orf (open reading frame) lab and S gene sequences. These RDs serves as 67 biomarkers for the virus identification and drug targets. Additionally, proteomic 68 69 comparison demonstrates that both SARS-CoV-2 and SAR-CoV protein are highly identical (95%). However, SARS-CoV-2 proteins (orf8 and orf10) were unavailable in 70 71 SARS-CoV. Furthermore, the viral drug targets including protease and polymerase 72 enzyme are highly conserved in both SARS-CoV-2 and SARS-CoV<sup>9</sup>. Thus, potential 73 drugs or treatments against SARS-CoV protease and enzymes may demonstrate activity against similar SARS-CoV-2 enzymes. The critical surface protein found in SARS-CoV-74 2 and SARS-CoV differ and are only 75% identical, hence suggesting their mechanism to 75 survival in the human host<sup>10, 11</sup>. The According to Lu, et al. <sup>10</sup>, SARS-CoV S protein is 76 77 shorter than SARS-CoV 2 S protein. Despite this, the spike S protein stalk in both viruses are highly homologous with 99% identity<sup>12</sup>. As the less scarce protein in Coronaviruses, 78 79 the coding amino acid sequence of both SARS-CoV-2 and SARS-CoV N protein are 90% identical<sup>13</sup>, indicating the probability of having similar antibodies. SARS-CoV N protein 80 is able to neutralize the host immune response, a characteristic which have not yet been 81 identified in the SARS-CoV-2 N protein<sup>14</sup>. From structure studies and biochemical tests<sup>1</sup>, 82 <sup>15-17</sup>, SARS-CoV -2 possess high binding affinity to the host cell than the SARS-CoV. 83 This was ascribed to mutation of amino acid residues located in the Receptor binding 84 domain (RBD) of the spike S protein. Also, the SARS-CoV 2 has a unique polybasic 85 cleavage site and three adjacent predicted O-linked glycans, which are unavailable in 86 SARS-CoV and related family B of betacoronaviruses<sup>6</sup>. 87 The angiotensin-converting enzyme 2 (ACE2) is the receptor in the human host cell that 88 interacts with the S protein of COVs through affinity binding to facilitate infection 16, 18-89 90 <sup>21</sup>. The S protein is used by the virus for entry into the host cell and subsequently fusing 91 into the cell membrane. These two steps are very important in the viral pathogenesis and

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infection<sup>22</sup>. The S protein is made of three protein macromolecules referred to as trimeric proteins and has a flexible structure that enables it to undergo structural rearrangements after entry into the host cell<sup>20, 21</sup>. The ease of human-to-human spread of COVID-19 has been associated with the spike protein having a high affinity for the ACE2 receptor<sup>23</sup>. Figure 1 shows the schematic representation of SARS-COV-2 and its interaction with ACE2. Coronavirus infection stimulates immune response in the host using the N protein. During the early stage of infection, the N protein is predominantly expressed, and this has been a target for diagnostics development<sup>24</sup>. N protein is a highly phosphorylated and abundant protein in CoVs that induce strong immune response after infection<sup>25-28</sup>. The viral N protein is an important structural protein responsible for transcription and assembling of the viral particle. It is also involved in the formation of helical ribonucleoproteins, regulation of RNA synthesis and transcription and metabolism modulation of infected cells<sup>29-32</sup>. Primarily, the CoV N protein binds and packs the viral genomic RNA into a nucleocapsid protein complex. All these attributes and functionalities have been demonstrated in ex vivo and in vivo biochemical studies<sup>31, 32</sup>. Further studies revealed that, N protein regulated interactions between the host and virus<sup>33-35</sup>. Additionally, it possess binding affinity for nsp3 located on the replicase complex and the M protein<sup>36</sup>, leading to transition from viral genome to replicase transcriptase complex, and finally packaging of genomic nucleocapsid into a viral particle. The N protein consist of two separate domains (Amino and carboxy terminus domain)<sup>37</sup>. These domains connected via short Serine-Arginine linker are able to bind RNA via contrasting mechanism. The amino and carboxy terminus domain undergo RNA binding and oligomerization respectively<sup>38-42</sup>; and the linker is responsible for phosphorylation. The amino terminal – genomic RNA binding is possible via electrostatic interactions. Also, relevant amino acids residues in the amino terminal domain are responsible for the viral RNA binding and infection<sup>43-45</sup>. These features of N protein have assisted in the development of diagnostic assays and several certain candidate antibodies and vaccines have been emerged against SARS-CoV virus <sup>26, 27</sup>. In a recent study, <sup>46</sup>demonstrated novel promising RNA binding pocket on SARS-CoV 2 virus, a domain absent in other coronaviruses. The authors said this finding would speed up the design/development of 123 new antiviral drugs for COVID-19. Also, the pocket would be a target site for 124 blockers/inhibitors that will hinder or hamper SARS-CoV 2 N protein replication. The S protein is a fusion protein consisting of S1 and S2 units, predicted to form  $\alpha$ -helical 125 secondary coiled coil structure<sup>47-49</sup>. Within the S1 unit, is a receptor – binding domain 126 (RBD) which recognize and interact directly with the human ACE2 receptor. S1 unit 127 consist of N and C terminal domains (NTD, CTD). S2 helps in the virus – cell membrane 128 fusion and anchorage<sup>6, 17</sup>. The S2 domain harbours other protein such as putative fusion 129 peptide and heptapeptide repeats (HR1, HR2) domains. In final stages of S2 protein 130 mediated membrane fusion, these peptides of SARS-CoV fold into anti-parallel six-helix 131 bundles<sup>50, 51</sup>. The Membrane anchored S2 subunit is supported and stabilized by the S1 132 unit. In SARS-CoV 2, Both S1 and S2 subunits are separated by a furin cleavage site<sup>6</sup>. 133 134 Furin is a host protease which resides in the Golgi. The removal of this cleavage site 135 affects the virus entry into VeroE6 and BHK cells. Primarily, the spike protein induces immune responses and neutralizes antibodies against virus infections, suggesting that 136 therapeutics (vaccines and antiviral blockers or inhibitors or prophylactics), if developed 137 against S protein could induce certain antibodies. These antibodies may either inhibit the 138 viral binding and fusion or neutralize the virus infection<sup>52</sup>. A collection of these candidate 139 therapeutics and vaccines for SARS-CoV can be seen in this review article <sup>53</sup>. 140 141 Membrane (M) protein helps in the assembly and formation of the viral core and envelope via interaction with the N protein<sup>54</sup>. The envelope (E) protein is a 142 transmembrane helix protein of 76 - 109 amino acids<sup>55-57</sup>. It is comprised of three 143 domains; amino terminal, transmembrane and carboxy terminal<sup>58-61</sup>. The amino and 144 carboxy terminal consist of short and long hydrophilic peptide respectively. The 145 transmembrane domain are hydrophilic in nature and consist  $\alpha$  – helix secondary 146 structure. The E protein is involved in host cell recognition and viral assembly<sup>55</sup>. 147 However, the function in coronavirus infection remains elusive. Hence, becoming 148 difficult to consider this as important target for the development of therapeutics. 149 All Together, in an antibody profile studies on SARS patients' sera, both M and E protein 150 lack neutralizing or antagonistic antibodies as these proteins located within the viral 151 envelope<sup>62, 63</sup>. However, development of antagonist for these M and N protein would 152 contribute to relevant alternative therapeutic for coronavirus. 153

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155 156 Hemagglutinin SARS-CoV 2 Receptor (ACE2) Esterase Dimer 157 Nucleocapsid Protein and RNA 158 159 SARS-CoV-2 160 Human Cell 161 162 Envelope 163 protein Spike protein Membrane protein 164

Figure 1 Schematic representation of the structure of coronavirus and the ACE2 host cell receptor. ACE2 receptors interact with SARS-COV-2 spike protein S. Adapted from <sup>64</sup>

# 2.0 Emerging in vitro Detection and Diagnostic tests for COVID-19

Several COVID-19 *in vitro* diagnostic kits have been manufactured by many companies, and these require approvals from FDA and other regulatory bodies. These testing assays and kits differ in the biochemistry and turnaround time. Most of the new testing assays and kits developed to support COVID-19 mitigation efforts have received Emergency

- Use Authorisation (EUA) from FDA<sup>65</sup>, and are limited to the laboratories that invented
- 177 them.
- 178 Currently, the diagnostic techniques approved for the detection of SARS-COV-2, and in
- fact COVs in general, include Reverse-T Polymerase Chain Reaction (RT-PCR)<sup>66</sup>,
- 180 Isothermal Nucleic Acid Amplification, and Blood-Based Serological Test 67, 68. Table
- 181 1.0 shows the list of approved products and tests published by  $FDA^{69}$ .
- 182 Majority of this test and kits are based on the polymerase Chain Reaction (92% figure
- 183 2.0). In this technique<sup>65</sup>, deactivated virus RNA is extracted and purified from
- 184 nasopharyngeal swab collected from patient. The purified RNA is then reverse
- transcribed and amplified to DNA. Through repeated heating and cooling, millions copy
- of DNA are made. These virus DNA is then mixed with fluorescent dyes, which give off
- more light after binding to the virus DNA, an indication of the presence of the virus in the
- 188 collected sample
- The ease of use, specificity and high sensitivity of RT-PCR makes it an effective method
- 190 for SARS-COV-2 detection<sup>70, 71</sup>. It detects the virus from sputum, blood, urine, saliva,
- 191 pharyngeal swabs, nasal, anal swabs and stool specimen. RT-PCR kits may have three
- different assays with each associated with a separate SARS-CoV-2 gene target. Hence,
- the probability of mutation is minimal. The RT-PCR has shown to detect 4-8 viral copies
- by amplifying targets in Orflab, E and N viral genes at a 95% confidence level 72-74.
- However, RT-PCR identified 47-59% of the positive cases in COV tests and 75% of
- results initially indicated as negative were found positive on repeated run<sup>73, 75, 76</sup>. Another
- short fall of RT-PCR assay relates to heat inactivation of the samples, and this may lead
- 198 to inactivation of the viral particle, hindering effective downstream diagnostic
- 199 evaluation<sup>77</sup>.
- 200 The isothermal nucleic acid amplification method for detecting nucleic acid targets is
- void of limitations associated with thermal cycling<sup>78</sup>. The isothermal amplification
- 202 method uses only one temperature and includes techniques such as helicase-dependent
- amplification, recombinase polymerase amplification and loop-mediated isothermal
- amplification (LAMP)<sup>70</sup>. All these techniques can be incorporated into a multiplex
- system during amplification stage.

The LAMP technique has been proven for COVID-19 detection <sup>67, 79-81</sup>. In LAMP test, 206 DNA polymerase and ~ 6 primers are used. These primers bind to six different regions on 207 the target genome. Similar to PCR, collected nasopharyngeal or oropharyngeal samples 208 209 from patient is added to the tube, followed by the DNA amplification and detection via either turbidity, colour or fluorescence. In a recent study by Park, et al. 82, a reverse 210 transcription loop-mediated isothermal amplification (RT-LAMP) assay was designed 211 and evaluated. The investigators detected over 100 copies of genomic RNA of SARS-212 CoV-2 using the colorimetric approach. The assays were void cross-reactivity with other 213 human coronaviruses. This RT-LAMP are promising point -of – care test for COVID-19 214 but RNA extraction method needs to be optimized. Compared to PCR, RT-LAMP 215 216 requires no centralized laboratory testing or facilities. Hence, it can conducted with 217 simple instruments (for example, drying oven or water bath) and have similar sensitivity and specificity<sup>83</sup>. The limitations associated with this method include challenges of 218 improving primer and reaction conditions<sup>70</sup>. Examples of these testing kits include Real 219 Time SARS-CoV-2 assay and ID NOW COVID-19 produced by Abbott Diagnostics 220 Scarborough, Inc<sup>69</sup>. SARS-CoV-2 nucleic acid extraction, isolation and purification from 221 222 swabs specimens are required prior to the Abbott assay. The ID NOW assay involves target amplification, heating, mixing and detection of nucleic acids. The heating step in 223 224 ID NOW system could inactivate collected samples<sup>73</sup>. 225 The blood-based serological test involves qualitative detection of antibodies (IgM,IgA and IgG) linked to SARS-CoV-2 to assess individuals exposed to COVID-1984. The 226 detection of the antibodies via this assay indicates the immune response to SARS-CoV-2 227 virus in suspected patients who have been infected previously or with recent COVID-19 228 229 infections. These antibodies can be detected approximately 14 days after infections. This test kit is produced by Cellex Inc. Prior to this development, quantitative immunoassay 230 tests such as Dual ELISA were performed<sup>85</sup>. It detects different antibody types (IgA, IgM 231 and IgG) against SARS-CoV in the blood of suspected individuals. Also, an ELISA kit 232 assay for the detection of SARS-COV-2 nucleoprotein has been developed. These tests 233 are for research use only and have currently not been approved by FDA Emergency Use 234 Authorization. 235

Despite being an accurate, rapid and simple essential tool for elucidation of interactions 236 between several reported cases, antibody-based detection methods are associated with 237 sensitivity and specificity limitations, and target types are scarce<sup>86</sup>. Moreover, antibody 238 239 detection may not be suitable for early-stage infection as the immunoassay could generate negative results since antibodies may still be the development stage. Another problem 240 could be potential positive results for formerly infected patients. Hence, this calls for a 241 huge scrutiny about the accuracy, reliability and uncertainty of the serological test. 242 Additionally, a PCR test used for screening COVID-19 can cost up to \$51 whiles 243 antibody test cost less than \$10 under Medicare<sup>87</sup>. On the frontline of turnaround time, 244 whiles it takes 15 minutes to get serological test result, PCR runs last for about 4 - 6 245 hours. Due to the back and forth shipment and transfer of samples, results from PCR are 246 247 procrastinated. Patient receive results in several days after test run. However, all this has been rectified through the design and invention of portable point - of - care test and 248 results are received in less than an hour (table 1.0). 249 PCR testing is still considered to be highly accurate and embraced by the most hospitals, 250 recommended by most doctors for mass screening. This test shows whether a person is 251 252 still struggling with the virus and can transmit to others. According to the FDA, over 5,000, 000 test are reported in a week via the PCR technique 253 254 and predicts over 1000,000 tests in the coming months for the antibody-based test<sup>69</sup>. NGS is a new technique with high accuracy, used for the identification and 255 characterization of pathogen quickly and result in rapid treatment. NGS based test 256 involves a metagenomic sequencing assay which detects the SARS-CoV-2 virus with a 257 sensitivity of 500 copies per milliliter<sup>88</sup>. Drawbacks of this technique include high 258 instrument cost, requirement of bioinformatics expert, Difficulty in mass screening and 259 260 its test time. It takes at least 2 days to get the test result (table 1.0). According to the manufacturers, they currently focused on the European market. Currently there only two 261 approved tests using this technique which is approximately 3% (Figure 2.0) of the FDA 262 approved test with CE marking. 263 From the above-mentioned limitations of existing and emerging diagnostic tests and kits, 264 it is critical to continue the development of rapid, sensitive, inexpensive, specific, robust 265 point-of-care diagnostics for COVID-19 to enable mass screening exercises especially in 266

low resourced communities. A surveillance diagnostic test that can easily be carried out in different laboratories and outside healthcare settings is urgently needed to prevent the wide and rapid spreading of COVID-19. It is important that this test can distinguish between COVs and specifically target SARS-COV-2 virus. This will help individuals with suspected symptoms to rapidly check if they have been infected with the virus.

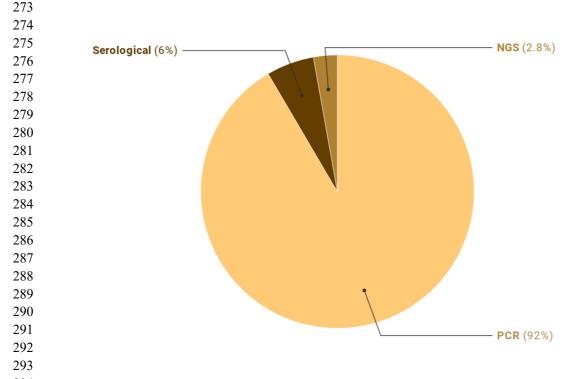


Figure 2.0: Distribution of Diagnostic test and kits based on Techniques or test type. Figure created with *Datawrapper*.

Table 1.0 FDA Approved Diagnostic test kits for COVID -19 infections

PRODUCT	Manufacturers	Test Type	Result	Approval
			Time (hr)	Status

1.	RealTime SARS-CoV-2	Abbott	PCR	4-6	FDA-EUA
2.	ID NOW COVID-19 test	Abbott	Isothermal amp-PoC	<1	FDA-EUA
3.	AvellinoCoV2	Avellino Labs	PCR	24-48	FDA-EUA
4.	BioGX SARS -CoV- 2Reagents	BioGX,BD	PCR	2-3	FDA-EUA
5.	Real-Time Fluorescent RT-PCR Kit	BGI	PCR	3	FDA-EUA
6.	BIOFIRE COVID-19 test	BioMerieux- BioFire Defense	PCR	<1	FDA-EUA
7.	2019-nCoV Real-Time RT-PCR Dx Panel	CDC	PCR	24-72	FDA-EUA
8.	qSARS- CoV -2IgG/IgM Rapid Test Kit	Cellex	Serological	<1	FDA-EUA
9.	COVID-19 ELISA IgG Antibody test	Mount Sinai Laboratory	Serological	<1	FDA-EUA
10.	DPP COVID-19 IgM/IgG system	Chembio Diagnostic System, Inc	Serological	<1	FDA-EUA
11.	VITROS Immunodiagnostic Products Anti – SARS – CoV-2 Total Reagent Pack	Ortho Clinical Diagnostic, Inc	Serological	<	FDA-EUA
12.	Xpert Xpress SARS - CoV-2 test	Cepheid	PCR-PoC	<1	FDA-EUA
13.	Logix Smart Coronavirus COVID - 19 Test	Co-Diagnostics	PCR	1-2	FDA-EUA
14.	Simplexa COVID-19 Direct	DiaSorin Molecular	PCR	1	FDA-EUA
15.	ePlex SARS -CoV-2 Test	GenMark Diagnostics	PCR	2	FDA-EUA
16.	COVID -19 RT-Digital PCR Detection Kit	Gnomegen	PCR	4-6	FDA-EUA
<i>17</i> .	Panther Fusion SARS - CoV -2 Assay	Hologic	PCR	3	FDA-EUA
18.	Smart Dectect SARS - CoV -2rRT -PCR Kit	InBios International	PCR	4-6	FDA-EUA
19.	CoV -19 IDx assay	Ipsum Diagnostics	PCR	24	FDA-EUA
20.	Covid -19 RT -PCR test	LabCorp	PCR	24	FDA-EUA
21.	ARISES SARS -CoV -2 Assay	Luminex Molecular Diagnostics	PCR	2	FDA-EUA
22.	NxTAG CoV Extended Panel Assay	Luminex Molecular Diagnostics	PCR	4	FDA-EUA
<i>23</i> .	Accula SARS -CoV -2	Mesa Biotech	PCR-PoC	<1	FDA-EUA

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test	N. M.D.	DCD	1.2	ED A ELLA
24. SARS -CoV -2 Assay	NeuMoDx	PCR	1-2	FDA-EUA
,288/96 Molecular				
System 25 Nav. Conorminus PT	Perkin Elmer	PCR	4-6	FDA-EUA
25. New Coronavirus RT - PCR test	Perkin Einier	PCK	4-0	FDA-EUA
26. COVID -19 genesing	Primerdesign	PCR	2	FDA-EUA
Real -Time PCR assay	Timerdesign	TCK	2	TDA-LOA
27. QIAstat -Dx Respiratory	Qiagen(acq.by	PCR	96-120	FDA-EUA
SARS -CoV-2 Panel	Thermo Fisher)	ICK	70-120	TDA-LOA
28. Quest SARS -CoV-2rRT	Ouest	PCR	1	FDA-EUA
-PCR	ouest	Tek	1	I DIT LOT
29. Lyrra SARS -CoV -2	Quidel	PCR	4-6	FDA-EUA
Assay	Quider	1 010	. 0	TBIT BOIT
30. Cobas SARS -CoV-2	Roche	PCR	3-8	FDA-EUA
Test				
31. SARS -CoV -2 RTqPCR	ScienceCell	PCR	4-6	FDA-EUA
Detection Kit	Research Labs			
32. TaqPath COVID -19	Thermo Fisher	PCR	4	FDA-EUA
Combo Kit				
33. NY SARS -CoV -2 Real -	Wadsworth Center	PCR	24-72	FDA-EUA
time RT-PCR	,NY State Dept of			
	Public Health			
34. SARS-CoV-2 +Influenza	3D Medicines	PCR	4-6	CE Mark
A & B RT -qPCR Kit				
35. REALQUALITY RQ-	AB ANALITICA	PCR	4-6	CE Mark
2019-nCoV				
36. Bosphore 2019-nCoV	Anatolia	PCR	2	CE Mark
Detection Kit	Geneworks			
37. SARS -CoV-2, influenza	AusDiagnostics	PCR	4-6	CE Mark
,RSV panel				
38. AccuPower COVID 19	Bioneer	PCR	8	CE Mark
Real -Time RT -PCR Kit			_	
39. Q-Sens 2019-nCoV	Cancer Rop	PCR	2	CE Mark
Detection Kit	G T D	D.C.D.	2	GEAL 1
40. VIASURE SARS-CoV-2	CerTest Bioter	PCR	3	CE Mark
Real -TimePCR	,BD	DCD D C	-1	CE M 1
41. VitaPCR SARS -CoV2	Credo Diagnostics Biomedical	PCR-PoC	<1	CE Mark
Assay		DCD	2	CE M1-
42. QuantiVirus SARS -	DiaCarta	PCR	2	CE Mark
CoV-2 test	Canatia Cianatana	DCD	1.5	CE Mark
43. EasyScreen SARS -CoV -2 Detection Kit	Genetic Signature	PCR	4-5	CE Mark
	Genetic Health	DCD	1	CE Mork
44. Detection Kit for SARS -	Geneue neam	PCR	4	CE Mark
CoV-2	Genomica	PCR	5	CE Mark
45. qCOVID-19 ,CLART COVID-19	/PharmMar Group	ICK	3	CE IVIAIK
46. 2019Real-time PCR Kit	Kogene Biotech	PCR	4-6	CE Mark
47. GeneFinder COVID -19	OsangHealth	PCR	4-6	CE Mark
RealAMp Kit	Osangiicanii	ICK	7-0	CE IVIAIN
48. Allplex 2019 -nCoV	Seegene	PCR	4	CE Mark
10. 1111pien 2019 -11COV	Seegone	I OIC	·	CL Mark

Assay	1			
49. DiaPlex Q 2019-nCoV  Detection Kit	SolGent	PCR	2	CE Mark
50. SARS -CoV -2 Clinical Sequence assay	Vision Medicals	NGS	>12	CE Mark
51. Multiple Real -Time PCR Kit	Beijjing Applied Biological Technologies (XABT	PCR	4-6	CE Mark
52. SARS-CoV-2RT -PCR test	Children Hospital of Philadelphia (CHOP)	PCR	4-6	LDT(EUA)
53. MGH COVID -19qPCR assay	Massachuesetts General Hospital	PCR	4-6	LDT(EUA)
54. SARS -CoV -2 Assay	Northwestern Medicine	PCR	4-6	LDT(EUA)
55. Viracor SARS -CoV -2 assay	Viracor Eurofins Clinical Diagnostics	PCR	4-6	LDT(EUA)
56. Applied Biosystems TaqPath COVID-19 Combo Kit	Rutgers Clinical Genomics Laboratory	PCR	4-6	LDT( EUA)
57. SDI SARS-CoV-2 Assay	Special Diagnostic laboratories	PCR	4-6	LDT( EUA)
58. UNC Health SARS- CoV-2 real-time RT- PCR test	University of North Carolina	PCR	4-6	LDT(EUA)
59. Stanford SARS-CoV-2 assay	Standard Health care	PCR	4-6	LDT( EUA)
60. Orig3n 2019 Novel Coronavirus (COVID- 19) Test	Orig3n, Inc.	PCR	4-6	LDT(EUA)
61. SARS -CoV-2 PCR test	Yale new Haven Hospital	PCR	4-6	LDT(EUA)
62. CDI Enhanced COVID- 19 Test	Hackensack University Medical Centre	PCR	4-6	LDT( EUA)
63. CirrusDx SARS-CoV-2 Assay	CirrusDx Laboratories	PCR	4-6	LDT( EUA)
64. Childrens – Altona- SARS-CoV-2 Assay	Infectious Diseases Diagnostics	PCR	4-6	LDT( EUA)
65. SARS-CoV-2 Test	Exact Sciences Laboratories	PCR	4-6	LDT( EUA)
66. SARS-CoV-2 Test	Integrity Laboratories	PCR	4-6	LDT( EUA)
67. COVID-19 RT-PC Test	Medicine Lab of Baptist Hospital	PCR	4-6	LDT( EUA)
68. Explify Respiratory	IDbyDNA	NGS	24-48	LDT
69. COVID -19 Home Test Kits	Carbon Health	PCR	72-144	Discontinued

70. At- home Covid -19 test	Everlywell	PCR	48	Discontinued
71. Covid -19 Home Test	Nurx Molecular	PCR	48	Discontinued
Kit	Testing Labs			

308 PCR - Polymerase Chain Reaction; LDT - Laboratory Based Test; EUA - Emergency Use Authorization, FDA- Food and Drug Authority, PoC - Point - of - Care . NGS - Next Generation 309 310 Sequencing 311

Available on https://www.fda.gov/medical-devices/emergency-situations-medical-devices/emergency-

use-authorizations, accessed on 17 April 2020 312

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# 3.0 Aptameric nanosensors as viral diagnostics

Aptamers offer a novel approach for targeted diagnosis and treatment of infections. 315 316 Aptamers have short oligonucleotides that are single-stranded and consist of either RNA or DNA with the ability to detect a wide range of molecules<sup>89-91</sup> including cells, tissues, 317 viruses, and bacteria. They have high specificity and affinity to their target molecules. 318 Aptamers are generated by an artificial method known as Systematic Evolution of 319 Ligands by Exponential Enrichment (SELEX)<sup>92</sup>. To design nucleic acid sequences that 320 are specific to target molecules, the SELEX technique employs selecting sequences and 321 replicating them carefully over iterative cycles. Aptamers have between 20-90 322 nucleotides<sup>93</sup>. To generate an aptamer for a target molecule, all feasible aptamers 323 324 sequence with specific length are selected from a combinatorial library and incubated with the molecule of interest<sup>94-96</sup>. During this process, the nucleic acids with high affinity to 325 the target molecule bind to it and the sequences with low affinity are removed from the 326 pool. 327 The high affinity nucleic acids and the target molecule complex are recovered and 328 329 separated. This process repeated until an aptamer with a low dissociation constant and high specificity towards the molecule of interest is generated. In addition to the SELEX 330 procedure, other techniques such as surface plasmon resonance (SPR) and capillary 331 332 electrophoresis (CE) can be integrated to increase biostability and improve binding performance. 333 Investigating the aptamer binding features under varying conditions of temperature, pH 334 335 and conductivity whilst probing the conformational dynamics of the aptamer presents a useful strategy to optimize the binding performance. Aptamers bind to their targets via 336 337 unique structural transformations. This gives them some advantages over monoclonal

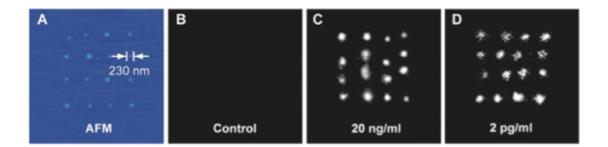
antibodies in terms of binding specificity and stability. The smaller size of aptamers makes them ideal for *in vivo* use compared to antibodies. Moreover, immunofluorescence dyes and drugs could be coupled onto aptamers without functionality and properties<sup>94, 97</sup>. Aptamers can be employed in biosensing devices as probes and such biosensors are called aptasensors. They are superior to conventional biosensors that use antibodies as they are stable, have high affinity to targets, modifiable, and can be developed for a wide range of targets <sup>98</sup>, using various transduction mechanisms. Aptasensors have been applied in different forms to detect chemicals, disease biomarkers and pathogens found in food. It is important to identify and diagnose diseases at an early stage to help patients receive early treatment and better healthcare delivery<sup>93</sup>. In light of this, several antigens and viruses have aptamers developed for them to aid in their diagnosis. Aptamers generated for this purpose either target the whole virus or surface antigens. Over the years, aptamers have been developed for viruses such as hepatitis B and C viruses, human papilloma virus, HIV, Influenza, SARS, Ebola, dengue, herpes simplex virus, west nile virus <sup>99-109</sup>.

## 4.0 Aptameric nanosensors for SARS coronavirus detection

An aptamer with a dissociation constant of 4.93 nM has been generated to target the nucleocapsid protein of SARS COV<sup>7</sup>. From the ELISA analysis, this ssDNA aptamer specifically bound to the nucleocapsid protein. In comparison with antibody, the authors further showed that the ssDNA aptamer was able to detect the SARS-CoV nucleocapsid protein efficiently via Western blot analysis. In an earlier development by Ahn and coworkers, an RNA aptamer was developed to target the nucleocapsid protein and a 1.65 nM dissociation constant was obtained<sup>110</sup>. Fluorescence imaging showed a detection limit of 2 pg/ml (Figure 3.0). The signal transduction was achieved using polyclonal anti-N antibody and FITC-labelled anti-rabbit IgG secondary antibody. In another study by Jang, et al. <sup>111</sup>, RNA aptamers was generated for the NSP10 (NTPase/Helicase) found in SARS coronavirus. They reported that isolated RNAs efficiently bound and inhibited duplex DNA unwinding activity of SARS-CoV helicase by approximately 85%. The IC<sub>50</sub> value was 1.2 nM. However, they stimulated infinitesimal change on ATPase activity of the helicase protein in the presence of a cofactor (Figure 4.0).

Similarly, Shum and colleagues reported the binding between an immobilised DNA aptamer on Ni-NTA magnetic beads and SARS-coronavirus helicase<sup>112</sup>. Using ATPase and fluorescence resonance energy transfer (FRET) based assay, all selected aptamers demonstrated inhibitory activity against the SARS-CoV helicase with low apparent K<sub>m</sub> values. The SARS-CoV helicase has the ability to unwind its duplex nucleic acid (DNA and RNA) to enable the replication and proliferation of the virus<sup>113, 114</sup>. Through circular dichroism and gel electrophoresis techniques, the investigators identified two different aptamer classes, namely G-quadruplex and non-G-quadruplex. Interestingly, non-G-quadruplex aptamer clones efficiently and specifically inhibited the SARS-CoV helicase-unwinding activity with IC<sub>50</sub> ranging from (17.5 – 120.8 nM). The opposite was observed for the G-quadruplex aptamer class.

Also, Roh and Jo sensitively and specifically detected SARS-CoV nucleocapsid protein in a one-spot experiment<sup>115</sup>. The investigators reached this achievement with RNA



aptamer conjugated to quantum dots (QDs) using an immobilized SARS-CoV N protein -

glass chip system. Florescence imaging showed a detection limit of 0.1pg/ml.

Figure 3.0 Atomic force microscopy (A) and Fluorescence microscopy (B - D) for the detection of SARS COV N protein by a nanoarray aptamer chip. Adapted from  $^{110}$ .

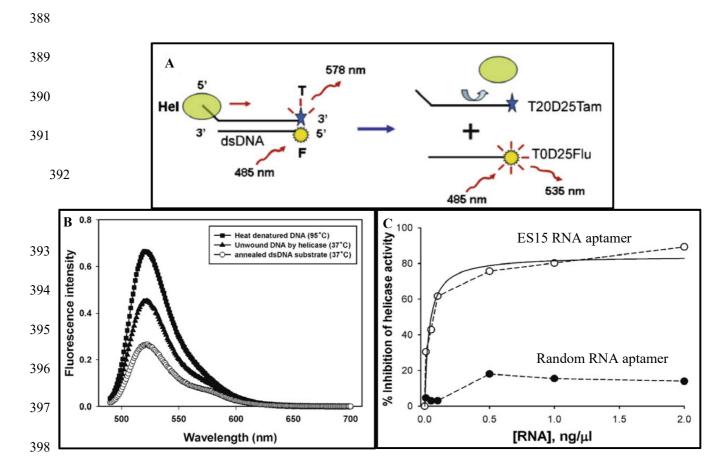


Figure 4.0. Schematic and Principle of FRET-based fluorometric helicase assay (**A-B**) and inhibition of the SARS-CoV helicase protein activity by ES15 RNA aptamer (**C**). The inhibition activity of ES15 Aptamer is 85 % and random RNA aptamer has little effect on the helicase substrate. Adapted from <sup>111</sup>

Recently, Researchers have developed new creative diagnostic kits with the use of a pioneering imaging technology called Mango for its vivid colour to sensitively detect RNA molecules, helping to improve the screening of viruses such as coronavirus while enabling fundamental discoveries into the functioning of cells<sup>116</sup>. The Mango system was made up of a RNA aptamer which acts like a magnet targeting the dye molecules. The dye becomes excited when bound and glows brightly. They concluded that the Mango NABSA kits can be used to detect pathogens such as the positive stranded RNA coronavirus faster and more efficiently.

### **5.0 Future Perspective**

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Pandemics and viral diseases or outbreaks are inescapable and can lead damaging effects on every aspect of life. Due to this, there is an utmost need for effective and affordable testing or diagnostic technologies for early detection. The diagnosis of viral diseases and recent outbreak largely depend on the detection of nucleic acid, specifically RNA. A judicious selection of appropriate detection method and kits is essential. On the frontier of the current pandemic (COVID-19), FDA approved diagnostic test and technique described earlier possess unique merit and unavoidable demerit. These diagnostic methods are mainly molecular based assays and serology detection or immunoassays. The former consists of reverse - time PCR assay, full genome analysis using Nextgeneration sequencing (NGS) and loop-mediated isothermal amplification method (LAMP). PCR as a highly sensitive and specific standard assay is widely used for COVID-19 screening, however testing requires sophisticated equipment, facilities, clinical laboratories and well-trained personnel and finally time-consuming. NGS helps in the characterization, understanding, identification and transmission of the virus, but it is an expensive approach. The high temperature requirement of the LAMP limits its further application in coronavirus detection. Serology detection method, which is a point-of-care test helps in the detection of antibody types against SARS-CoV-2 in patient serum. Its low sensitivity and specificity inevitably hamper its usage by healthcare facilities. Additionally, only few has been developed and approved by the FDA. In this context, there is a need to develop rapid, cheaper, specific, sensitive and user-friendly diagnostic tool and point-of-care test for SARS-CoV. To date, new approaches are been developed to enhance SARS-CoV-2 detection but have not been approved by FDA yet. Practical approach is via a combination of the aforementioned methods to avoid limitations of single test and methods<sup>70, 117, 118</sup>. Due to their biophysical characteristics or potentials, aptamers have previously been demonstrated as a bioprobe for rapid sensing, screening and detection of infectious or pandemic pathogens and virulent diseases. These characteristics makes them ideal candidate to design and develop point-of-care test (POCT)<sup>119, 120</sup> for diagnosis and combating COVID-19 and other future pandemics. The Juxtaposition of molecular diagnostics and aptameric technology have been highlighted previously<sup>121</sup>. Briefly, In

advantages such as (i) easily generated for several targets (including toxins and non-444 nucleic acid targets)<sup>122, 123</sup> (ii) they are less prone to contamination and sample 445 preparation is not essential<sup>124</sup>; (iii) they are simple, rapid, less expensive and can be used 446 in real time<sup>125</sup>. 447 Additionally, although it is subjected to application format, aptameric assays do not 448 depend on labels, centralized or clinical laboratories and well-trained staffs<sup>126-129</sup>. Hence, 449 it can be developed for detection or testing for COVID-19 cases at home and this will be 450 reduced travelling to clinical/healthcare facilities which could contribute to rapid diseases 451 transmission and escalation. Similar to molecular diagnostics, aptasensors are highly 452 specific, sensitive and can be applied or incorporated into multiplex systems or 453 applications<sup>124, 125, 128-130</sup>. 454 455 Through aptamer immobilization on 2Dimensional (Glass slides, silicon, chips, polymethylmethacrylate) and 3 Dimensional (Microspheres made of magnets, agarose, 456 silica, monoliths and polystyrene) supports, many aptamer sensors with improved 457 characteristics or properties have been developed for dual functions, biodetection and 458 bioscreening<sup>131-138</sup>. The 3D immobilized aptasensors are highly efficient and possess 459 higher surface area -to-volume ratio<sup>121</sup>. Also, they have high loading capacity, are less 460 461 prone to steric hindrance and can be used efficiently in multiplex assays. Moreover, 3D immobilized aptasensors have demonstrated high throughput ability and are cost -462 effective. They can be used to detect and capture pathogens into support matrix. The 463 development of 3D aptasensors assays are quicker, hence, a criterion for rapid, specific 464 and sensitive screening of pandemic pathogens. 465 All these improved aptasensors technologies will be essential for effective mass 466 screening for pandemics, possibly further detection of asymptomatic cases and low 467 pathogen limits in suspected individuals since it takes a at least 2 - 14 days for suspected 468 patients to develop symptoms of COVID-19 infections. 469 Recently, Great strides and milestones have been reached on aptamer discovery through 470 integrated artificial intelligence-based technologies<sup>139</sup> and in silico or computational 471 methods<sup>140-146</sup>. These has come to solved problems of efficient and successful 472 473 identification of new or high-performance aptamers from combinatorial library using the

comparison to molecular – based assays or diagnostic technology, aptameric sensors have

474 SELEX technology. Additionally, high cost, labour and time further limits the SELEX process<sup>147</sup>. These computational methods have revolutionized the separation and 475 identification efficiencies; decreased amplification bias and nonspecific adsorption; and 476 enabled expansion of different base types. Several machine learning methods have 477 emerged for identification of high-performance aptamers. 478 These include sequence clustering-based methods<sup>148, 149</sup> and motif finding – based 479 methods<sup>150-154</sup>. Unfortunately, the former had limitations such as high amplification bias, 480 and adsorption process is not specific. The later studies, although developed to over the 481 482 challenges of the former, they faced with limitation such as high computational and analysis cost. Also, the general secondary structures of aptamers are not considered, 483 which makes it absolutely choice dependent. 484 Moreover, both strategies/methods are still unable to identify highly efficient aptamers 485 486 from SELEX pool. Surprisely, all these challenges have been addressed in a different study. In this recent work using machine learning based classification, Song, et al. 139 487 developed an Algorithm (SMART-Aptamer) to select aptamers from sequencing data 488 SELEX pool to fill the above mentioned gaps with pioneering studies. The investigators 489 490 rapidly identified three aptamers with excellent affinity, selectivity and accuracy for Epithelial stem cells, human embryonic cells and blood cells. The aptamer (SJ-3C2) 491 492 demonstrated a strong binding ability with  $K_d$  value in the nanomolar range (41.43  $\pm$  1.84 nM). Thus, using in silico methods, improved / enhanced aptamer sensing technologies 493 can be developed to target different structural protein on SARS-CoV-2 for diagnosis and 494 495 therapeutic purposes.

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#### 7.0 Conclusion

Diagnostics form relevant part of the toolkits in combating COVID-19 outbreaks. This enable healthcare providers or settings to distribute resources and focus on patients with severe infections. Hence, preventing the spread of the virus and reduction in mortality rate. The unique characteristics and benefits of aptamers are promising for the development of novel therapeutics and diagnostics for viral infections with particular emphasis on coronaviruses. This will aid in the development of reliable point-of-care test

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to bind specifically to SARS-CoV-2. Point-of-care test are less expensive, hand-held diagnostics that are used to test patient samples in community centres and do not require well-trained staffs, centralized facilities and clinical settings<sup>70</sup>. Also, these devices have short turnaround time and prevent spread of the disease as suspected individuals do not need to travel to clinical laboratories for testing. Research advances on aptameric sensors, technologies and application demonstrates that, it can be employed for the diagnosis and screening of current and future pandemics. The pandemic pathogens can be detected in a rapid, sensitive and specific manner. This will help to curtail infection aggravation, transmission and spread of the virus. Moreover, it is evident that integrated in silico methods and Artificial intelligence can be used to identify specific aptamer for improved aptsensing technologies against the SARS-CoV-2, consequently speeding up the design and development of point-of-care testing for combating the Data on COVID-19 keeps changing rapidly daily. Some information in this review may change as more studies emerge and new tests and methods get approved by FDA. Some preprints were referenced. Acknowledgement: None **Conflicts of Interest:** None

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543	References
544	
545	1. Wrapp, D.; Wang, N.; Corbett, K. S.; Goldsmith, J. A.; Hsieh, CL.; Abiona, O.; Graham, B. S.;
546	McLellan, J. S., Cryo-EM structure of the 2019-nCoV spike in the prefusion conformation. <i>Science</i> <b>2020</b> ,
547	367 (6483), 1260-1263.
548	2. GISAID, Genomic epidemiology of hCoV-19. 2020.
549	3. Ceraolo, C.; Giorgi, F. M., Genomic variance of the 2019-nCoV coronavirus. <i>Journal of medical</i>
550	virology 2020.
551	4. Chan, J. FW.; Yuan, S.; Kok, KH.; To, K. KW.; Chu, H.; Yang, J.; Xing, F.; Liu, J.; Yip, C.
552	CY.; Poon, R. WS.; Tsoi, HW.; Lo, S. KF.; Chan, KH.; Poon, V. KM.; Chan, WM.; Ip, J. D.;
553	Cai, JP.; Cheng, V. CC.; Chen, H.; Hui, C. KM.; Yuen, KY., A familial cluster of pneumonia
554	associated with the 2019 novel coronavirus indicating person-to-person transmission: a study of a family
555	cluster. The Lancet <b>2020</b> , 395 (10223), 514-523.

- 556 5. Cyranoski, D., Did pangolins spread the China coronavirus to people. *Nature* **2020**.
- 557 6. Andersen, K. G.; Rambaut, A.; Lipkin, W. I.; Holmes, E. C.; Garry, R. F., The proximal origin of
- 558 SARS-CoV-2. *Nature medicine* **2020**, *26* (4), 450-452.
- 7. Cho, S. J.; Woo, H. M.; Kim, K. S.; Oh, J. W.; Jeong, Y. J., Novel system for detecting SARS
- 560 coronavirus nucleocapsid protein using an ssDNA aptamer. *J Biosci Bioeng* **2011**, *112* (6), 535-40.
- 8. Xu, J.; Zhao, S.; Teng, T.; Abdalla, A. E.; Zhu, W.; Xie, L.; Wang, Y.; Guo, X., Systematic
- 562 comparison of two animal-to-human transmitted human coronaviruses: SARS-CoV-2 and SARS-CoV.
- 563 Viruses **2020**, 12 (2), 244.

- 564 Tseng, C.-T.; Sbrana, E.; Iwata-Yoshikawa, N.; Newman, P. C.; Garron, T.; Atmar, R. L.; Peters,
- 565 C. J.; Couch, R. B., Immunization with SARS coronavirus vaccines leads to pulmonary immunopathology
- 566 on challenge with the SARS virus. PLoS One 2012, 7 (4).
- 567 Lu, R.; Zhao, X.; Li, J.; Niu, P.; Yang, B.; Wu, H.; Wang, W.; Song, H.; Huang, B.; Zhu, N.,
- 568 Genomic characterisation and epidemiology of 2019 novel coronavirus: implications for virus origins and 569 receptor binding. The Lancet 2020, 395 (10224), 565-574.
- 570 Srinivasan, S.; Cui, H.; Gao, Z.; Liu, M.; Lu, S.; Mkandawire, W.; Narykov, O.; Sun, M.; Korkin,
- 571 D., Structural Genomics of SARS-CoV-2 Indicates Evolutionary Conserved Functional Regions of Viral
- 572 Proteins. Viruses 2020, 12 (4), 360.
- 573 Chan, J. F.-W.; Kok, K.-H.; Zhu, Z.; Chu, H.; To, K. K.-W.; Yuan, S.; Yuen, K.-Y., Genomic
- 574 characterization of the 2019 novel human-pathogenic coronavirus isolated from a patient with atypical
- 575 pneumonia after visiting Wuhan. Emerg Microbes Infect 2020, 9 (1), 221-236.
- 576 Gralinski, L. E.; Menachery, V. D., Return of the Coronavirus: 2019-nCoV. Viruses 2020, 12 (2), 13.
- 577 135.
- 578 14. Li, G.; Fan, Y.; Lai, Y.; Han, T.; Li, Z.; Zhou, P.; Pan, P.; Wang, W.; Hu, D.; Liu, X., Coronavirus
- 579 infections and immune responses. Journal of medical virology 2020, 92 (4), 424-432.
- 580 Letko, M.; Marzi, A.; Munster, V., Functional assessment of cell entry and receptor usage for
- 581 SARS-CoV-2 and other lineage B betacoronaviruses. *Nature Microbiology* **2020**, *5* (4), 562-569.
- 582 Wan, Y.; Shang, J.; Graham, R.; Baric, R. S.; Li, F., Receptor recognition by the novel
- 583 coronavirus from Wuhan: an analysis based on decade-long structural studies of SARS coronavirus. J Virol 584 2020, 94 (7).
- 585 Walls, A. C.; Park, Y.-J.; Tortorici, M. A.; Wall, A.; McGuire, A. T.; Veesler, D., Structure, 17.
- 586 function, and antigenicity of the SARS-CoV-2 spike glycoprotein. Cell 2020.
- 587 Hoffmann, M.; Kleine-Weber, H.; Krüger, N.; Müller, M.; Drosten, C.; Pöhlmann, S., The novel
- 588 coronavirus 2019 (2019-nCoV) uses the SARS-coronavirus receptor ACE2 and the cellular protease
- 589 TMPRSS2 for entry into target cells. Cold Spring Harbor Laboratory: 2020.
- 590 Zhou, P.; Yang, X.-L.; Wang, X.-G.; Hu, B.; Zhang, L.; Zhang, W.; Si, H.-R.; Zhu, Y.; Li, B.;
- 591 Huang, C.-L., A pneumonia outbreak associated with a new coronavirus of probable bat origin. Nature 592
- **2020,** *579* (7798), 270-273.
- 593 Bosch, B. J.; van der Zee, R.; de Haan, C. A. M.; Rottier, P. J. M., The coronavirus spike protein is
- 594 a class I virus fusion protein: structural and functional characterization of the fusion core complex. J Virol 595 **2003**, 77 (16), 8801-8811.
- 596 Li, F., Structure, Function, and Evolution of Coronavirus Spike Proteins. Annu Rev Virol 2016, 3 597 (1), 237-261.
- 598 Benvenuto, D.; Giovanetti, M.; Ciccozzi, A.; Spoto, S.; Angeletti, S.; Ciccozzi, M., The 2019-new 599 coronavirus epidemic: evidence for virus evolution. Journal of Medical Virology 2020, 92 (4), 455-459.
- 600 Chan, J. F.-W.; Yuan, S.; Kok, K.-H.; To, K. K.-W.; Chu, H.; Yang, J.; Xing, F.; Liu, J.; Yip, C.
- 601 C.-Y.; Poon, R. W.-S., A familial cluster of pneumonia associated with the 2019 novel coronavirus
- 602 indicating person-to-person transmission: a study of a family cluster. The Lancet 2020, 395 (10223), 514-603
- 604 Che, X.-Y.; Hao, W.; Wang, Y.; Di, B.; Yin, K.; Xu, Y.-C.; Feng, C.-S.; Wan, Z.-Y.; Cheng, V.
- 605 C.; Yuen, K.-Y., Nucleocapsid protein as early diagnostic marker for SARS. Emerging infectious diseases 606 **2004,** 10 (11), 1947.
- 607 Ahmed, S. F.; Quadeer, A. A.; McKay, M. R., Preliminary identification of potential vaccine
- 608 targets for the COVID-19 coronavirus (SARS-CoV-2) based on SARS-CoV immunological studies.
- 609 Viruses 2020, 12 (3), 254.
- 610 Liu, S.-J.; Leng, C.-H.; Lien, S.-p.; Chi, H.-Y.; Huang, C.-Y.; Lin, C.-L.; Lian, W.-C.; Chen, C.-J.;
- 611 Hsieh, S.-L.; Chong, P., Immunological characterizations of the nucleocapsid protein based SARS vaccine
- 612 candidates. Vaccine 2006, 24 (16), 3100-3108.
- 613 Shang, B.; Wang, X.-Y.; Yuan, J.-W.; Vabret, A.; Wu, X.-D.; Yang, R.-F.; Tian, L.; Ji, Y.-Y.;
- 614 Deubel, V.; Sun, B., Characterization and application of monoclonal antibodies against N protein of SARS-
- 615 coronavirus. Biochemical and biophysical research communications 2005, 336 (1), 110-117.
- 616 Ying, L.; Xu, S.; Yang, R. F.; Li, Y. X.; Ji, Y. Y.; He, Y. Y.; De Shi, M.; Wei, L.; Shi, T. L.; Jin,
- 617 W., Identification of an epitope of SARS-coronavirus nucleocapsid protein. Cell Res 2003, 13 (3), 141-145.

- 618 29. Cong, Y.; Ulasli, M.; Schepers, H.; Mauthe, M.; V'Kovski, P.; Kriegenburg, F.; Thiel, V.; de
- 619 Haan, C. A. M.; Reggiori, F., Nucleocapsid Protein Recruitment to Replication-Transcription Complexes
- Plays a Crucial Role in Coronaviral Life Cycle. J Virol 2020, 94 (4), e01925-19.
- 621 30. Nelson, G. W.; Stohlman, S. A.; Tahara, S. M., High affinity interaction between nucleocapsid
- protein and leader/intergenic sequence of mouse hepatitis virus RNA. *Journal of General Virology* 2000,
   81 (1), 181-188.
- 624 31. Stohlman, S. A.; Baric, R. S.; Nelson, G. N.; Soe, L. H.; Welter, L. M.; Deans, R. J., Specific
- interaction between coronavirus leader RNA and nucleocapsid protein. J Virol 1988, 62 (11), 4288-4295.
- 626 32. Tang, T. K.; Wu, M. P. J.; Chen, S. T.; Hou, M. H.; Hong, M. H.; Pan, F. M.; Yu, H. M.; Chen, J.
- H.; Yao, C. W.; Wang, A. H. J., Biochemical and immunological studies of nucleocapsid proteins of severe
- 628 acute respiratory syndrome and 229E human coronaviruses. *Proteomics* **2005**, *5* (4), 925-937.
- 629 33. Du, L.; Zhao, G.; Lin, Y.; Chan, C.; He, Y.; Jiang, S.; Wu, C.; Jin, D.-Y.; Yuen, K.-Y.; Zhou, Y.;
- Zheng, B.-J., Priming with rAAV encoding RBD of SARS-CoV S protein and boosting with RBD-specific
- peptides for T cell epitopes elevated humoral and cellular immune responses against SARS-CoV infection.
- 632 Vaccine 2008, 26 (13), 1644-1651.
- 633 34. Hsieh, P.-K.; Chang, S. C.; Huang, C.-C.; Lee, T.-T.; Hsiao, C.-W.; Kou, Y.-H.; Chen, I. Y.;
- 634 Chang, C.-K.; Huang, T.-H.; Chang, M.-F., Assembly of severe acute respiratory syndrome coronavirus
- RNA packaging signal into virus-like particles is nucleocapsid dependent. *J Virol* **2005,** *79* (22), 13848-13855.
- 637 35. Surjit, M.; Liu, B.; Chow, V. T. K.; Lal, S. K., The Nucleocapsid Protein of Severe Acute
- Respiratory Syndrome-Coronavirus Inhibits the Activity of Cyclin-Cyclin-dependent Kinase Complex and
- Blocks S Phase Progression in Mammalian Cells. *Journal of Biological Chemistry* **2006**, *281* (16), 10669-10681.
- 641 36. Fehr, A. R.; Perlman, S., Coronaviruses: an overview of their replication and pathogenesis. In
- 642 Coronaviruses, Springer: 2015; pp 1-23.
- 643 37. McBride, R.; Van Zyl, M.; Fielding, B. C., The coronavirus nucleocapsid is a multifunctional protein. *Viruses* **2014**, *6* (8), 2991-3018.
- 645 38. Chang, C.-k.; Chen, C.-M. M.; Chiang, M.-h.; Hsu, Y.-l.; Huang, T.-h., Transient oligomerization
- of the SARS-CoV N protein--implication for virus ribonucleoprotein packaging. *PLoS One* **2013**, 8 (5),
- 647 e65045-e65045.
- 648 39. Chang, C.-k.; Sue, S.-C.; Yu, T.-h.; Hsieh, C.-M.; Tsai, C.-K.; Chiang, Y.-C.; Lee, S.-j.; Hsiao, H.-
- 649 h.; Wu, W.-J.; Chang, W.-L.; Lin, C.-H.; Huang, T.-h., Modular organization of SARS coronavirus
- nucleocapsid protein. *Journal of biomedical science* **2006**, *13* (1), 59-72.
- 651 40. Chen, I. J.; Yuann, J.-M. P.; Chang, Y.-M.; Lin, S.-Y.; Zhao, J.; Perlman, S.; Shen, Y.-Y.; Huang,
- T.-H.; Hou, M.-H., Crystal structure-based exploration of the important role of Arg106 in the RNA-binding
- domain of human coronavirus OC43 nucleocapsid protein. *Biochim Biophys Acta* **2013**, *1834* (6), 1054-654 1062.
- 655 41. Lo, Y.-S.; Lin, S.-Y.; Wang, S.-M.; Wang, C.-T.; Chiu, Y.-L.; Huang, T.-H.; Hou, M.-H.,
- Oligomerization of the carboxyl terminal domain of the human coronavirus 229E nucleocapsid protein.
- 657 FEBS letters **2013**, 587 (2), 120-127.
- 658 42. Wootton, S. K.; Rowland, R. R. R.; Yoo, D., Phosphorylation of the porcine reproductive and
- respiratory syndrome virus nucleocapsid protein. J Virol 2002, 76 (20), 10569-10576.
- 660 43. Grossoehme, N. E.; Li, L.; Keane, S. C.; Liu, P.; Dann, C. E., 3rd; Leibowitz, J. L.; Giedroc, D. P.,
- Coronavirus N protein N-terminal domain (NTD) specifically binds the transcriptional regulatory sequence
- 662 (TRS) and melts TRS-cTRS RNA duplexes. *J Mol Biol* **2009**, *394* (3), 544-557.
- 663 44. Keane, S. C.; Liu, P.; Leibowitz, J. L.; Giedroc, D. P., Functional transcriptional regulatory
- sequence (TRS) RNA binding and helix destabilizing determinants of murine hepatitis virus (MHV)
- nucleocapsid (N) protein. *J Biol Chem* **2012**, 287 (10), 7063-7073.
- 666 45. Tan, Y. W.; Fang, S.; Fan, H.; Lescar, J.; Liu, D. X., Amino acid residues critical for RNA-
- binding in the N-terminal domain of the nucleocapsid protein are essential determinants for the infectivity
- of coronavirus in cultured cells. *Nucleic acids research* **2006**, *34* (17), 4816-4825.
- 669 46. Kang, S.; Yang, M.; Hong, Z.; Zhang, L.; Huang, Z.; Chen, X.; He, S.; Zhou, Z.; Zhou, Z.; Chen,
- Q., Crystal structure of SARS-CoV-2 nucleocapsid protein RNA binding domain reveals potential unique
- drug targeting sites. *Acta Pharmaceutica Sinica B* **2020**.

- 672 47. Gui, M.; Song, W.; Zhou, H.; Xu, J.; Chen, S.; Xiang, Y.; Wang, X., Cryo-electron microscopy
- structures of the SARS-CoV spike glycoprotein reveal a prerequisite conformational state for receptor
- 674 binding. Cell Res **2017**, 27 (1), 119-129.
- Kirchdoerfer, R. N.; Cottrell, C. A.; Wang, N.; Pallesen, J.; Yassine, H. M.; Turner, H. L.; Corbett,
- K. S.; Graham, B. S.; McLellan, J. S.; Ward, A. B., Pre-fusion structure of a human coronavirus spike
- 677 protein. *Nature* **2016**, *531* (7592), 118-121.
- 678 49. Wrapp, D.; Wang, N.; Corbett, K. S.; Goldsmith, J. A.; Hsieh, C.-L.; Abiona, O.; Graham, B. S.;
- 679 McLellan, J. S., Cryo-EM structure of the 2019-nCoV spike in the prefusion conformation. Science (New
- 680 *York, N.Y.)* **2020,** *367* (6483), 1260-1263.
- 681 50. Bosch, B. J.; Martina, B. E. E.; Van Der Zee, R.; Lepault, J.; Haijema, B. J.; Versluis, C.; Heck, A.
- J. R.; De Groot, R.; Osterhaus, A. D. M. E.; Rottier, P. J. M., Severe acute respiratory syndrome
- coronavirus (SARS-CoV) infection inhibition using spike protein heptad repeat-derived peptides. *Proc Natl*
- 684 Acad Sci US A **2004**, 101 (22), 8455-8460.
- 51. Zhu, J.; Xiao, G.; Xu, Y.; Yuan, F.; Zheng, C.; Liu, Y.; Yan, H.; Cole, D. K.; Bell, J. I.; Rao, Z.;
- Tien, P.; Gao, G. F., Following the rule: formation of the 6-helix bundle of the fusion core from severe
- acute respiratory syndrome coronavirus spike protein and identification of potent peptide inhibitors.
- *Biochemical and biophysical research communications* **2004,** *319* (1), 283-288.
- 689 52. Chen, X.; Li, R.; Pan, Z.; Qian, C.; Yang, Y.; You, R.; Zhao, J.; Liu, P.; Gao, L.; Li, Z., Human
- monoclonal antibodies block the binding of SARS-CoV-2 spike protein to angiotensin converting enzyme 2 receptor. *Cellular & Molecular Immunology* **2020**, 1-3.
- 692 53. Du, L.; He, Y.; Zhou, Y.; Liu, S.; Zheng, B.-J.; Jiang, S., The spike protein of SARS-CoV—a
- target for vaccine and therapeutic development. *Nature Reviews Microbiology* **2009**, 7 (3), 226-236.
- 694 54. Li, Y.-H.; Hu, C.-Y.; Wu, N.-P.; Yao, H.-P.; Li, L.-J., Molecular Characteristics, Functions, and
- Related Pathogenicity of MERS-CoV Proteins. Engineering (Beijing) 2019, 5 (5), 940-947.
- 55. Tan, Y.-J.; Lim, S. G.; Hong, W., Characterization of viral proteins encoded by the SARS-
- 697 coronavirus genome. *Antiviral research* **2005**, *65* (2), 69-78.
- 698 56. Arbely, E.; Khattari, Z.; Brotons, G.; Akkawi, M.; Salditt, T.; Arkin, I. T., A highly unusual
- palindromic transmembrane helical hairpin formed by SARS coronavirus E protein. *J Mol Biol* **2004**, *341* (3), 769-779.
- Kuo, L.; Hurst, K. R.; Masters, P. S., Exceptional flexibility in the sequence requirements for coronavirus small envelope protein function. *J Virol* **2007**, *81* (5), 2249-2262.
- 58. Li, Y.; Surya, W.; Claudine, S.; Torres, J., Structure of a Conserved Golgi Complex-targeting Signal in Coronavirus Envelope Proteins. Worldwide Protein Data Bank: 2014.
- 705 59. Liao, Y.; Yuan, Q.; Torres, J.; Tam, J. P.; Liu, D. X., Biochemical and functional characterization
- of the membrane association and membrane permeabilizing activity of the severe acute respiratory syndrome coronavirus envelope protein. *Virology* **2006**, *349* (2), 264-275.
- 708 60. Surya, W.; Samso, M.; Torres, J., Structural and Functional Aspects of Viroporins in Human
- 709 Respiratory Viruses: Respiratory Syncytial Virus and Coronaviruses. In *Respiratory Disease and Infection* 710 *A New Insight*, InTech: 2013.
- 711 61. Torres, J.; Maheswari, U.; Parthasarathy, K.; Ng, L.; Liu, D. X.; Gong, X., Conductance and
- amantadine binding of a pore formed by a lysine-flanked transmembrane domain of SARS coronavirus envelope protein. *Protein Sci* **2007**, *16* (9), 2065-2071.
- 714 62. Wang, J.; Wen, J.; Li, J.; Yin, J.; Zhu, Q.; Wang, H.; Yang, Y.; Qin, E. d.; You, B.; Li, W.,
- Assessment of immunoreactive synthetic peptides from the structural proteins of severe acute respiratory syndrome coronavirus. *Clin Chem* **2003**, *49* (12), 1989-1996.
- 717 63. Tan, Y.-J.; Goh, P.-Y.; Fielding, B. C.; Shen, S.; Chou, C.-F.; Fu, J.-L.; Leong, H. N.; Leo, Y. S.;
- 718 Ooi, E. E.; Ling, A. E., Profiles of antibody responses against severe acute respiratory syndrome
- 719 coronavirus recombinant proteins and their potential use as diagnostic markers. Clin. Diagn. Lab. Immunol.
- 720 **2004,** *11* (2), 362-371.
- 721 64. ProSci, ACE2 Antibodies 2020.
- 722 65. FDA, Coronavirus Disease 2019 (COVID-19) Emergency Use of Authorizations 2020.
- 723 66. Zhang, W.; Du, R.-H.; Li, B.; Zheng, X.-S.; Yang, X.-L.; Hu, B.; Wang, Y.-Y.; Xiao, G.-F.; Yan,
- B.; Shi, Z.-L.; Zhou, P., Molecular and serological investigation of 2019-nCoV infected patients:
- 725 implication of multiple shedding routes. *Emerg Microbes Infect* **2020**, *9* (1), 386-389.

- 726 67. Lamb, L. E.; Bartolone, S. N.; Ward, E.; Chancellor, M. B., Rapid Detection of Novel
- 727 Coronavirus (COVID-19) by Reverse Transcription-Loop-Mediated Isothermal Amplification. Cold Spring
- Harbor Laboratory: 2020.
- 729 68. Wu, J. T.; Leung, K.; Leung, G. M., Nowcasting and forecasting the potential domestic and
- 730 international spread of the 2019-nCoV outbreak originating in Wuhan, China: a modelling study. *The*
- 731 *Lancet* **2020**, *395* (10225), 689-697.
- 732 69. FDA, Emergency Use Authorization. 2020.
- 733 70. Udugama, B.; Kadhiresan, P.; Kozlowski, H. N.; Malekjahani, A.; Osborne, M.; Li, V. Y. C.;
- 734 Chen, H.; Mubareka, S.; Gubbay, J. B.; Chan, W. C. W., Diagnosing COVID-19: The Disease and Tools
- for Detection. ACS nano 2020, acsnano.0c02624.
- 736 71. Wong, M. L.; Medrano, J. F., Real-time PCR for mRNA quantitation. *BioTechniques* **2005**, *39* (1),
- 737 75-85.
- 738 72. Xie, C.; Jiang, L.; Huang, G.; Pu, H.; Gong, B.; Lin, H.; Ma, S.; Chen, X.; Long, B.; Si, G.; Yu,
- H.; Jiang, L.; Yang, X.; Shi, Y.; Yang, Z., Comparison of different samples for 2019 novel coronavirus
- detection by nucleic acid amplification tests. *International Journal of Infectious Diseases* **2020**, *93*, 264-267.
- 742 73. Xie, X.; Zhong, Z.; Zhao, W.; Zheng, C.; Wang, F.; Liu, J., Chest CT for Typical 2019-nCoV
- Pneumonia: Relationship to Negative RT-PCR Testing. *Radiology* **2020**, 200343.
- 74. Lin, C.; Xiang, J.; Yan, M.; Li, H.; Huang, S.; Shen, C., Comparison of throat swabs and sputum
- specimens for viral nucleic acid detection in 52 cases of novel coronavirus (SARS-Cov-2) infected pneumonia (COVID-19). Cold Spring Harbor Laboratory: 2020.
- 747 75. Ai, T.; Yang, Z.; Hou, H.; Zhan, C.; Chen, C.; Lv, W.; Tao, Q.; Sun, Z.; Xia, L., Correlation of
- Chest CT and RT-PCR Testing in Coronavirus Disease 2019 (COVID-19) in China: A Report of 1014
- 749 Cases. *Radiology* **2020**, 200642.
- 750 76. Shirato, K.; Nao, N.; Katano, H.; Takayama, I.; Saito, S.; Kato, F.; Katoh, H.; Sakata, M.;
- Nakatsu, Y.; Mori, Y.; Kageyama, T.; Matsuyama, S.; Takeda, M., Development of Genetic Diagnostic
- 752 Methods for Novel Coronavirus 2019 (nCoV-2019) in Japan. *Japanese Journal of Infectious Diseases* 753 **2020**.
- 754 77. Li, C.; Debruyne, D. N.; Spencer, J.; Kapoor, V.; Liu, L. Y.; Zhou, B.; Lee, L.; Feigelman, R.;
- Burdon, G.; Liu, J., High sensitivity detection of coronavirus SARS-CoV-2 using multiplex PCR and a multiplex-PCR-based metagenomic method. *bioRxiv* **2020**.
- 757 78. Notomi, T.; Okayama, H.; Masubuchi, H.; Yonekawa, T.; Watanabe, K.; Amino, N.; Hase, T.,
- Loop-mediated isothermal amplification of DNA. *Nucleic acids research* **2000**, *28* (12), e63-e63.
- 759 79. Yang, W.; Dang, X.; Wang, Q.; Xu, M.; Zhao, Q.; Zhou, Y.; Zhao, H.; Wang, L.; Xu, Y.; Wang,
- J.; Han, S.; Wang, M.; Pei, F.; Wang, Y., Rapid Detection of SARS-CoV-2 Using Reverse transcription
- 761 RT-LAMP method. Cold Spring Harbor Laboratory: 2020.
- 762 80. Yu, L.; Wu, S.; Hao, X.; Li, X.; Liu, X.; Ye, S.; Han, H.; Dong, X.; Li, X.; Li, J.; Liu, J.; Liu, N.;
- Zhang, W.; Pelechano, V.; Chen, W.-H.; Yin, X., Rapid colorimetric detection of COVID-19 coronavirus
- vising a reverse tran-scriptional loop-mediated isothermal amplification (RT-LAMP) diagnostic plat-form:
- 765 iLACO. Cold Spring Harbor Laboratory: 2020.
- 766 81. Zhang, Y.; Odiwuor, N.; Xiong, J.; Sun, L.; Nyaruaba, R. O.; Wei, H.; Tanner, N. A., Rapid
- Molecular Detection of SARS-CoV-2 (COVID-19) Virus RNA Using Colorimetric LAMP. Cold Spring Harbor Laboratory: 2020.
- 769 82. Park, G.-S.; Ku, K.; Baek, S.-H.; Kim, S. J.; Kim, S. I.; Kim, B.-T.; Maeng, J.-S., Development of
- 770 Reverse Transcription Loop-mediated Isothermal Amplification (RT-LAMP) Assays Targeting SARS-
- 771 CoV-2. Cold Spring Harbor Laboratory: 2020.
- 772 83. Craw, P.; Balachandran, W., Isothermal nucleic acid amplification technologies for point-of-care
- 773 diagnostics: a critical review. *Lab on a Chip* **2012**, *12* (14), 2469.
- 774 84. Laura, L., Cellex becomes first compnay to land FDA Emergency Use Authorization for COVID-
- 775 19 serology testing 03 April 2020 ed.; 2020.
- 776 85. Diagnostics, C., SARS-CoV-2 Immunoassay Kits. 2020.
- 777 86. Rostomily, K. A.; Jones, D. M.; Pautz, C. M.; Ito, D. W.; Buono, M. J., Haemoconcentration, not
- decreased blood temperature, increases blood viscosity during cold water immersion. *Diving and*
- 779 *Hyperbaric Medicine Journal* **2020,** *50* (1), 24-27.
- 780 87. Ellison, A., Medicare pricing for coronavirus test; 3 things to know. 16 March 2020 ed.; 2020.

- 781 GenomeWeb, Vision Medicals Gets CE Mark for SARS-CoV2 clinical sequencing test. 19 March
- 782 2020 ed.; 2020.
- 783 Berezovski, M.; Krylov, S. N., Using Nonequilibrium Capillary Electrophoresis of Equilibrium
- 784 Mixtures for the Determination of Temperature in Capillary Electrophoresis. Analytical Chemistry 2004, 76 785 (23), 7114-7117.
- 786 90. Burmeister, P. E.; Lewis, S. D.; Silva, R. F.; Preiss, J. R.; Horwitz, L. R.; Pendergrast, P. S.;
- 787 McCauley, T. G.; Kurz, J. C.; Epstein, D. M.; Wilson, C.; Keefe, A. D., Direct In Vitro Selection of a 2'-O-
- 788 Methyl Aptamer to VEGF. Chemistry & Biology 2005, 12 (1), 25-33.
- 789 Mendonsa, S. D.; Bowser, M. T., In Vitro Evolution of Functional DNA Using Capillary
- 790 Electrophoresis. Journal of the American Chemical Society 2004, 126 (1), 20-21.
- 791 Zhou, J.; Bobbin, M. L.; Burnett, J. C.; Rossi, J. J., Current progress of RNA aptamer-based
- 792 therapeutics. Front Genet 2012, 3, 234-234.
- 793 Wandtke, T.; Wozniak, J.; Kopinski, P., Aptamers in diagnostics and treatment of viral infections.
- 794 Viruses 2015, 7 (2), 751-80.
- 795 Somasunderam, A.; Ferguson, M. R.; Rojo, D. R.; Thiviyanathan, V.; Li, X.; O'Brien, W. A.;
- 796 Gorenstein, D. G., Combinatorial selection, inhibition, and antiviral activity of DNA thioaptamers targeting
- 797 the RNase H domain of HIV-1 reverse transcriptase. Biochemistry 2005, 44 (30), 10388-10395.
- 798 Eulberg, D.; Klussmann, S., Spiegelmers: Biostable Aptamers. ChemBioChem 2003, 4 (10), 979-799 983.
- 800 96. Kusser, W., Chemically modified nucleic acid aptamers for in vitro selections: evolving evolution.
- 801 Reviews in Molecular Biotechnology 2000, 74 (1), 27-38.
- 802 Yang, X.; Fennewald, S.; Luxon, B. A.; Aronson, J.; Herzog, N. K.; Gorenstein, D. G., Aptamers
- 803 containing thymidine 3'-O-phosphorodithioates: Synthesis and binding to nuclear factor-κB. Bioorganic & Medicinal Chemistry Letters 1999, 9 (23), 3357-3362. 804
- 805 Mairal, T.; Cengiz Özalp, V.; Lozano Sánchez, P.; Mir, M.; Katakis, I.; O'Sullivan, C. K.,
- 806 Aptamers: molecular tools for analytical applications. Analytical and Bioanalytical Chemistry 2007, 390 807 (4), 989-1007.
- 808 Gopinath, S. C. B.; Kumar, P. K. R., Aptamers that bind to the hemagglutinin of the recent
- 809 pandemic influenza virus H1N1 and efficiently inhibit agglutination. Acta Biomaterialia 2013, 9 (11), 810 8932-8941.
- 811 100.
- Gopinath, S. C. B., Methods developed for SELEX. Analytical and Bioanalytical Chemistry 2006,
- 812 387 (1), 171-182.
- 813 Zou, X.; Wu, J.; Gu, J.; Shen, L.; Mao, L., Application of Aptamers in Virus Detection and 101.
- 814 Antiviral Therapy. Front Microbiol 2019, 10, 1462.
- 815 Gopinath, S. C. B.; Lakshmipriya, T.; Arshad, M. K. M.; Voon, C. H.; Hashim, U., Aptasensors in
- 816 viral detection. Malaysian Journal of Microbiology 2016, 12 (5), 376-382.
- 817 Lakshmipriya, T.; Horiguchi, Y.; Nagasaki, Y., Co-immobilized poly(ethylene glycol)-block-
- 818 polyamines promote sensitivity and restrict biofouling on gold sensor surface for detecting factor IX in
- 819 human plasma. The Analyst 2014, 139 (16), 3977-3985.
- 820 Lakshmipriya, T.; Fujimaki, M.; Gopinath, S. C. B.; Awazu, K.; Horiguchi, Y.; Nagasaki, Y., A
- 821 high-performance waveguide-mode biosensor for detection of factor IX using PEG-based blocking agents
- 822 to suppress non-specific binding and improve sensitivity. The Analyst 2013, 138 (10), 2863.
- 823 Shiratori, I.; Akitomi, J.; Boltz, D. A.; Horii, K.; Furuichi, M.; Waga, I., Selection of DNA
- 824 aptamers that bind to influenza A viruses with high affinity and broad subtype specificity. Biochemical and
- 825 Biophysical Research Communications 2014, 443 (1), 37-41.
- 826 Bagashev, A.; Sawaya, B. E., Roles and functions of HIV-1 Tat protein in the CNS: an overview.
- 827 Virol J **2013**, 10, 358-358.
- 828 Yamamoto, R.; Kumar, P. K. R., Molecular beacon aptamer fluoresces in the presence of Tat 107.
- 829 protein of HIV-1. Genes to Cells 2000, 5 (5), 389-396.
- 830 Moore, M. D.; Bunka, D. H. J.; Forzan, M.; Spear, P. G.; Stockley, P. G.; McGowan, I.; James,
- 831 W., Generation of neutralizing aptamers against herpes simplex virus type 2: potential components of
- 832 multivalent microbicides. *J Gen Virol* **2011**, *92* (Pt 7), 1493-1499.
- 833 Gourronc, F. A.; Rockey, W. M.; Thiel, W. H.; Giangrande, P. H.; Klingelhutz, A. J.,
- 834 Identification of RNA aptamers that internalize into HPV-16 E6/E7 transformed tonsillar epithelial cells.
- 835 Virology 2013, 446 (1-2), 325-333.

- 836 110. Ahn, D. G.; Jeon, I. J.; Kim, J. D.; Song, M. S.; Han, S. R.; Lee, S. W.; Jung, H.; Oh, J. W., RNA
- aptamer-based sensitive detection of SARS coronavirus nucleocapsid protein. Analyst 2009, 134 (9), 1896-
- 838 901.
- 111. Jang, K. J.; Lee, N. R.; Yeo, W. S.; Jeong, Y. J.; Kim, D. E., Isolation of inhibitory RNA aptamers
- against severe acute respiratory syndrome (SARS) coronavirus NTPase/Helicase. *Biochem Biophys Res*
- 841 *Commun* **2008**, *366* (3), 738-44.
- 842 112. Shum, K. T.; Tanner, J. A., Differential inhibitory activities and stabilisation of DNA aptamers
- against the SARS coronavirus helicase. *Chembiochem* **2008**, *9* (18), 3037-45.
- 844 113. Ivanov, K. A.; Thiel, V.; Dobbe, J. C.; van der Meer, Y.; Snijder, E. J.; Ziebuhr, J., Multiple
- enzymatic activities associated with severe acute respiratory syndrome coronavirus helicase. *J Virol* **2004**,
- 846 78 (11), 5619-5632.
- 847 114. Tanner, J. A.; Watt, R. M.; Chai, Y.-B.; Lu, L.-Y.; Lin, M. C.; Peiris, J. S. M.; Poon, L. L. M.;
- Kung, H.-F.; Huang, J.-D., The Severe Acute Respiratory Syndrome (SARS) Coronavirus NTPase/Helicase
- Belongs to a Distinct Class of 5' to 3' Viral Helicases. *Journal of Biological Chemistry* **2003**, 278 (41),
- 850 39578-39582.
- 851 115. Roh, C.; Jo, S. K., Quantitative and sensitive detection of SARS coronavirus nucleocapsid protein
- using quantum dots-conjugated RNA aptamer on chip. Journal of Chemical Technology & Biotechnology
- 853 **2011,** *86* (12), 1475-1479.
- 854 116. Buquliskis, J. S., COVID-19 Test kits Being Developed using RNA Imaging 20 March 2020 ed.;
- 855 2020.
- 856 117. Yang, T.; Wang, Y.-C.; Shen, C.-F.; Cheng, C.-M., Point-of-Care RNA-Based Diagnostic Device
- for COVID-19. Multidisciplinary Digital Publishing Institute: 2020.
- 858 118. El-Tholoth, M.; Bau, H. H.; Song, J., A Single and Two-Stage, Closed-Tube, Molecular Test for
- the 2019 Novel Coronavirus (COVID-19) at Home, Clinic, and Points of Entry. 2020.
- 860 119. Baron, E. J.; Campbell, S., Technical and Clinical Niches for Point of Care Molecular Devices. In
- Advanced Techniques in Diagnostic Microbiology, Springer US: 2012; pp 619-626.
- 862 120. Cass, A. E. G.; Zhang, Y., Nucleic acid aptamers: ideal reagents for point-of-care diagnostics?
- 863 Faraday Discuss. 2011, 149, 49-61.
- 121. Acquah, C.; Danquah, M. K.; Agyei, D.; Moy, C. K.; Sidhu, A.; Ongkudon, C. M., Deploying
- aptameric sensing technology for rapid pandemic monitoring. *Critical reviews in biotechnology* **2016**, *36* (6), 1010-1022.
- Lim, D. V.; Simpson, J. M.; Kearns, E. A.; Kramer, M. F., Current and developing technologies
- for monitoring agents of bioterrorism and biowarfare. Clinical microbiology reviews 2005, 18 (4), 583-607.
- 869 123. Radom, F.; Jurek, P. M.; Mazurek, M. P.; Otlewski, J.; Jeleń, F., Aptamers: Molecules of great
- 870 potential. *Biotechnology Advances* **2013**, *31* (8), 1260-1274.
- 871 124. Wark, A. W.; Lee, J.; Kim, S.; Faisal, S. N.; Lee, H. J., Bioaffinity detection of pathogens on
- 872 surfaces. J Ind Eng Chem **2010**, 16 (2), 169-177.
- 873 125. Garibyan, L.; Avashia, N., Polymerase chain reaction. J Invest Dermatol 2013, 133 (3), 1-4.
- 874 126. Lazcka, O.; Campo, F. J. D.; Muñoz, F. X., Pathogen detection: A perspective of traditional
- methods and biosensors. *Biosensors and Bioelectronics* **2007**, *22* (7), 1205-1217.
- 876 127. Velusamy, V.; Arshak, K.; Korostynska, O.; Oliwa, K.; Adley, C., An overview of foodborne
- pathogen detection: In the perspective of biosensors. *Biotechnology Advances* **2010**, *28* (2), 232-254.
- 878 128. de-los-Santos-Álvarez, N.; Lobo-Castañón, M. a. J.; Miranda-Ordieres, A. J.; Tuñón-Blanco, P.,
- Aptamers as recognition elements for label-free analytical devices. *TrAC Trends in Analytical Chemistry*
- 880 **2008,** *27* (5), 437-446.
- 881 129. Loeffelholz, M.; Deng, H., PCR and Its Variations. In Advanced Techniques in Diagnostic
- 882 *Microbiology*, Springer US: pp 166-183.
- 883 130. Yang, S.; Rothman, R. E., PCR-based diagnostics for infectious diseases: uses, limitations, and
- future applications in acute-care settings. *Lancet Infect Dis* **2004**, *4* (6), 337-348.
- 885 131. Danquah, M. K.; Forde, G. M., Preparation of macroporous methacrylate monolithic material with
- 886 convective flow properties for bioseparation: Investigating the kinetics of pore formation and
- hydrodynamic performance. Chemical Engineering Journal 2008, 140 (1-3), 593-599.
- 888 132. Deng, N.; Liang, Z.; Liang, Y.; Sui, Z.; Zhang, L.; Wu, Q.; Yang, K.; Zhang, Y.,
- 889 Aptamer Modified Organic-Inorganic Hybrid Silica Monolithic Capillary Columns for Highly Selective
- Recognition of Thrombin. Analytical Chemistry 2012, 84 (23), 10186-10190.

- 891 133. Han, B.; Zhao, C.; Yin, J.; Wang, H., High performance aptamer affinity chromatography for
- single-step selective extraction and screening of basic protein lysozyme. *Journal of Chromatography B*
- 893 **2012,** *903*, 112-117.
- 894 134. Jungbauer, A.; Hahn, R., Polymethacrylate monoliths for preparative and industrial separation of
- biomolecular assemblies. *Journal of Chromatography A* **2008**, *1184* (1-2), 62-79.
- 896 135. Madru, B.; Chapuis-Hugon, F.; Pichon, V., Novel extraction supports based on immobilised
- aptamers: Evaluation for the selective extraction of cocaine. *Talanta* **2011**, *85* (1), 616-624.
- 898 136. Podgornik, A.; Krajnc, N. L., Application of monoliths for bioparticle isolation. *Journal of*
- 899 Separation Science **2012**, *35* (22), 3059-3072.
- 900 137. Podgornik, A.; Yamamoto, S.; Peterka, M.; Krajnc, N. L., Fast separation of large biomolecules
- using short monolithic columns. *Journal of Chromatography B* **2013**, 927, 80-89.
- 902 138. Wu, X.; Hu, J.; Zhu, B.; Lu, L.; Huang, X.; Pang, D., Aptamer-targeted magnetic nanospheres as a
- 903 solid-phase extraction sorbent for determination of ochratoxin A in food samples. Journal of
- 904 *Chromatography A* **2011**, *1218* (41), 7341-7346.
- 905 139. Song, J.; Zheng, Y.; Huang, M.; Wu, L.; Wang, W.; Zhu, Z.; Song, Y.; Yang, C., A Sequential
- 906 Multi-dimensional Analysis Algorithm for Aptamer Identification based on Structure Analysis and
- 907 Machine Learning. Analytical Chemistry 2019.
- 908 140. Ahirwar, R.; Nahar, S.; Aggarwal, S.; Ramachandran, S.; Maiti, S.; Nahar, P., In silico selection of
- an aptamer to estrogen receptor alpha using computational docking employing estrogen response elements
- 910 as aptamer-alike molecules. *Sci Rep* **2016**, *6*, 21285.
- 911 141. Ahirwar, R.; Nahar, S.; Aggarwal, S.; Ramachandran, S.; Maiti, S.; Nahar, P., In silico selection of
- an aptamer to estrogen receptor alpha using computational docking employing estrogen response elements
- 913 as aptamer-alike molecules. *Sci Rep* **2016**, *6*, 21285-21285.
- 914 142. Akbaripour-Elahabad, M.; Zahiri, J.; Rafeh, R.; Eslami, M.; Azari, M., rpiCOOL: A tool for In
- 915 Silico RNA–protein interaction detection using random forest. *Journal of Theoretical Biology* **2016**, *402*, 1-916 8.
- 917 143. Cataldo, R.; Ciriaco, F.; Alfinito, E., A validation strategy for in silico generated aptamers.
- 918 Computational Biology and Chemistry **2018**, 77, 123-130.
- 919 144. Hamada, M., In silico approaches to RNA aptamer design. *Biochimie* 2018, 145, 8-14.
- 920 145. Wang, S.; Zhang, Y.-H.; Lu, J.; Cui, W.; Hu, J.; Cai, Y.-D., Analysis and Identification of
- 921 Aptamer-Compound Interactions with a Maximum Relevance Minimum Redundancy and Nearest
- 922 Neighbor Algorithm. *Biomed Res Int* **2016**, *2016*, 8351204-8351204.
- 923 146. Wondergem, J. A. J.; Schiessel, H.; Tompitak, M., Performing SELEX experiments in silico. The
- 924 *Journal of Chemical Physics* **2017**, *147* (17), 174101.
- 925 147. Emami, N.; Pakchin, P. S.; Ferdousi, R., Computational predictive approaches for interaction and
- 926 structure of aptamers. *Journal of Theoretical Biology* **2020**, 110268.
- 927 148. Alam, K. K.; Chang, J. L.; Burke, D. H., FASTAptamer: A Bioinformatic Toolkit for High-
- 928 throughput Sequence Analysis of Combinatorial Selections. *Mol Ther Nucleic Acids* **2015**, *4* (3), e230-e230.
- 930 149. Hoinka, J.; Berezhnoy, A.; Sauna, Z. E.; Gilboa, E.; Przytycka, T. M., AptaCluster A Method to
- 931 Cluster HT-SELEX Aptamer Pools and Lessons from its Application. Res Comput Mol Biol 2014, 8394,
- 932 115-128.

- 933 150. Hiller, M.; Pudimat, R.; Busch, A.; Backofen, R., Using RNA secondary structures to guide
- 934 sequence motif finding towards single-stranded regions. *Nucleic acids research* **2006**, *34* (17), e117-e117.
- 935 151. Hoinka, J.; Zotenko, E.; Friedman, A.; Sauna, Z. E.; Przytycka, T. M., Identification of sequence-
- 936 structure RNA binding motifs for SELEX-derived aptamers. *Bioinformatics* **2012**, *28* (12), i215-i223.
- 937 152. Jiang, P.; Meyer, S.; Hou, Z.; Propson, N. E.; Soh, H. T.; Thomson, J. A.; Stewart, R., MPBind: a
- 938 Meta-motif-based statistical framework and pipeline to Predict Binding potential of SELEX-derived
- 939 aptamers. *Bioinformatics* **2014**, *30* (18), 2665-2667.
- 940 153. Phuong, D.; Hoinka, J.; Wang, Y.; Takahashi, M.; Zhou, J.; Costa, F.; Rossi, J.; Burnett, J.;
- 941 Backofen, R.; Przytycka, T. M., AptaTRACE: Elucidating Sequence-Structure Binding Motifs by
- 942 Uncovering Selection Trends in HT-SELEX Experiments. Cold Spring Harbor Laboratory: 2016.
- 943 154. Takahashi, M.; Wu, X.; Ho, M.; Chomchan, P.; Rossi, J. J.; Burnett, J. C.; Zhou, J., High
- 944 throughput sequencing analysis of RNA libraries reveals the influences of initial library and PCR methods
- 945 on SELEX efficiency. Sci Rep **2016**, 6, 33697-33697.