Original Short Communication

DOI: 10.26479/2021.0703.01

BINDING VARIABILITY OF ANIMALS TO SARS-COV-2 SPIKE PROTEIN

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ABSTRACT: The aim of this study was to compare polymorphic regions of the angiotensin converting enzyme 2 (ACE2) receptor that are known to affect the susceptibility to SARS-CoV-2 in a number of different vertebrates. The amino acid sequences of a ligand-binding domain of the ACE2 receptor of animals known to be susceptible to contracting SARS-CoV-2 were compared those animals that are thought to be unlikely to contract that virus. Polymorphic regions can be successfully identified to predict which animals might be more likely to be infected by the SARS-CoV-2 virus and highlight possible further complications SARS-CoV-2 could have on society through the meat, fur and pet industries. Animals that play a key role in the pet and meat industry such as cows, pigs, chickens, cats, sheep, horses, rabbits and dogs, as well as other animals that are closely related to humans such as common chimpanzees (Pan troglodytes) and Sumatran orangutans (Pongo abelii) were included to aid an effective comparison between these dissimilar animals. Differences in the amino acid sequences of the ligand-binding domain were identified using protein sequencing alignment tools. The number of specific amino acids known to be required for successful binding along the ACE2 receptor varied among the animals used in this study hence raising issues regarding the possible susceptibility of particular domesticated animals to contract the SARS-CoV-2 virus which are discussed later on.

Keywords: animal, bioinformatics, coronavirus, ligand, polymorphisms.

Article History: Received: April 08, 2021; Revised: April 21, 2021; Accepted: May 05, 2021.

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1.INTRODUCTION

Both severe acute respiratory syndrome coronavirus 1 (SARS-CoV-1) and severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) bind to the angiotensin converting enzyme 2 (ACE2) receptor to gain entry to cells [1,2]. A strong affinity between the SARS-CoV receptor binding (RBM) and the ACE2 human receptor are essential for efficient entry, however occasional polymorphisms can occur along the ligand-binding domain of the ACE2 receptor, that can weaken binding affinity and thereby reduce susceptibility to infection [3,4,5]. Shi and colleagues [6] demonstrated that polymorphisms within the ligand-binding domain of the ACE2 receptor in pigs, chickens and dogs [6,7,8] resulting in a weaker interaction with the SARS-CoV-2 RBM whilst there is a stronger interaction between the ACE2 receptor and the SARS-CoV-2 RBM in humans, chimpanzees and cats [6].The primary objective of this study was to compare the ACE2 receptor ligand binding domain of additional animals (wild, domesticated and livestock) and compare them to animals with known susceptibility to the SARS-CoV-2 virus can be predicted.

2. MATERIALS AND METHODS

The protein sequence of the ligand-binding domain of ACE2 receptor was obtained using a previously published source [9]. BlastP was utilized to align the ACE2 receptor protein sequences of humans (Homo Sapiens, common chimpanzees (Pan troglodytes), Sumatran orangutans (*Pongo abelii*), cats (*Felis catus*), pigs (*Sus scrofa domesticus*), chickens (*Gallus gallus domesticus*), dogs (*Canis lupus familiaris*), cows (*Bos taurus*), sheep (*Ovis aries*), rabbits (*Oryctolagus cuniculus*) and horses (*Equus caballus*). Protein sequences of these animals were obtained from UnitProt.org. Once the protein sequences were aligned, the coordinates of the ligand-binding domain were identified [9]. The amino acids known to contribute to binding affinity in all of the animals were highlighted which identified any major polymorphic regions along the ACE2 ligand binding domain [10].

Table 1: Amino acids known to contribute to binding affinity [4] between the ACE2 receptor

Amino acid along ACE2 receptor	SARS-CoV-2 spike
known to contribute to binding affinity	
Asn90	Arg408
Ser19	Arg487
Gln24	Ala475
Met82	Phe486
Leu79	Phe486
Tyr83	Phe486
Glu329	Arg439

and the SARS-CoV-2 spike.

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		Amino aci	d along ACE2 receptor	SARS-CoV-2 spike			
		known to c	contribute to binding affinity	cont.			
	cont. Lys31						
				Leu455			
		Lys31		Gln493			
		Glu35		Gln493			
		Lys353		Asn501			

3. RESULTS AND DISCUSSION

Using UniProt.org, the ACE2 receptor was shown to consist of 805 amino acids in all the animals studied. BLAST sequencing identified polymorphisms within the ACE2 receptor sequence (Figure 1).

Table 2: Amino acids present at the coordinates along the ACE2 ligand-binding motif. The amino acids in bold are different to the amino acids necessary for binding with the SARS-

Animals	Amino acid										
	19	24	31	35	38	79	82	83	90	329	353
Chimpanzee	Ser	Gln	Lys	Glu	Asp	Leu	Met	Tyr	Asn	Glu	Lys
Human	Ser	Gln	Lys	Glu	Asp	Leu	Met	Tyr	Asn	Glu	Lys
Sumatran Orangutan	Ser	Gln	Lys	Glu	Asp	Leu	Met	Tyr	Asn	Glu	Lys
Cat	Ser	Leu	Lys	Glu	Glu	Leu	Thr	Tyr	Asn	Glu	Lys
Rabbit	Ser	Leu	Lys	Glu	Asp	Leu	Thr	Tyr	Asn	Glu	Lys
Dog	Ser	Leu	Lys	Glu	Glu	Leu	Thr	Tyr	Asp	Glu	Lys
Pig	Ser	Leu	Lys	Glu	Asp	Ile	Thr	Tyr	Thr	Asn	Lys
Chicken	Asp	Glu	Lys	Arg	Asp	Asn	Arg	Phe	Asp	Thr	Lys
Cow	Ser	Gln	Lys	Glu	Asp	Met	Thr	Tyr	Asn	Asp	Lys
Sheep	Ser	Gln	Lys	Glu	Asp	Met	Thr	Tyr	Asn	Asp	Lys
Horse	Ser	Leu	Lys	Glu	Glu	Leu	Thr	Tyr	Asn	Glu	Lys

CoV-2 receptor binding motif.

The strength of the binding between the ligand-binding domain of the ACE2 receptor and the SARS-CoV-2 RBD in dogs, cows, cats, sheep, horses and rabbits is more questionable; although these animals share many amino acids required for successful binding, not all the amino acids are present

Swanton RJLBPCS 2021 www.rjlbpcs.com Life Science Informatics Publications and therefore this could potentially reduce their susceptibility to contract the virus [11]. This research is important as it may provide a way of predicting which animals are more susceptible to contraction of SARS-CoV-2 and the potential impacts this virus may have on the meat, pet and fur industries. For example, if cows and sheep are susceptible to contracting the SARS-CoV-2 virus this could greatly influence meat production. Of the other species studied in the study, the rabbit, having 9 of the amino acids that conferred a successful bind (Table 2), may also be susceptible to SARS-CoV-2 and indeed it has been recently shown that rabbits are susceptible to COVID [12,13,14]. Although the infection was shown to be asymptomatic this could have a potential impact on the meat and fur industries as well as domesticated pets, and shows the importance of supporting the meat and fur industry through additional protections to ensure that an evident decline in the populations of livestock does not occur [15].

Q9BYF1ACE2_HUMANQ56H28ACE2_FELCAQ58DD0ACE2_BOVINF6V9L3F6V9L3_HORSEJ9P7Y2J9P7Y2_CANLFA0A2J8KU96A0A2J8KU96_PANTRK7GLM4K7GLM4_PIGH2PUZ5H2PUZ5_PONABG1TEF4G1TEF4_RABITW5PSB6W5PSB6_SHEEPF1NHR4F1NHR4_CHICK	1 1 1 1 1 1 1 1	MSSSSWLILSIVAVTAACTIEEOAKTFLDENHEAEDLFYQSSLASWNYNTNITEENVQ MSGSFWLILSFAALTAACTTEEOAKTFLEOFNHEAEELSYQSSLASWNYNTNITDENVQ MTGSFWLILSIVAVTAACTTEEOAKTFLEOFNHEAEOLSYQSSLASWNYNTNITDENVQ MSGSSWLILSIVAVTAACTTEDLAKTFLEOFNHOAEOLSYQSSLASWNYNTNITDENVQ MSGSSWLILSLAALTAACTTEDLAKTFLEOFNHOAEOLFYQSSLASWNYNTNITDENVQ MSGSSWLILSIVAVTAACTTEEOAKTFLOFNHOAEOLFYQSSLASWNYNTNITDENVQ MSGSSWLILSIVAVTAACTTEEOAKTFLOFNHOAEOLFYQSSLASWNYNTNITDENVQ MSGSSWLILSIVAVTAACTTEEOAKTFLOFNHOAEOLFYQSSLASWNYNTNITDENVQ MSGSSWLILSIVAVTAACTTEEOAKTFLOFNHOAEOLFYQSSLASWNYNTNITDENVQ MSGSSWLILSIVAVTAACTTEEOAKTFLOFNHOAEOLFYQSSLASWNYNTNITDENVQ MSGSSWLILSIVAVTAACTTEEOAKTFLOFNHOAEOLFYQSSLASWNYNTNITDENVQ MSGSSWLILSIVAVTAACTTEEOAKTFLOFNHOAEOLSYQSSLASWNYNTNITDENVQ MTGSFWLILSIVAVTAACTTEEOAKTFLOFNHOAEOLSYQSSLASWNYNTNITDENVQ	60 60 60 60 60 60 60 60 60 59
Q9BYF1 ACE2 HUMAN Q56H28 ACE2_FELCA Q58DD0 ACE2_BOVIN F6V9L3 F6V9L3 HORSE J9P7Y2 J9P7Y2_CANLF AOA2J8KU96 AOA2J8KU96_PANTR K7GLM4 K7GLM4 PIG H2PU25 H2PU25_PONAB G1TEF4 G1TEF4 RABIT W5PSB6 W5PSB6_SHEEP F1NHR4 F1NHR4_CHICK	61 61 61 61 61 61 61 61 61	MMNNAGDKWSAFLKEOSTLACMOPLQEICOLTVKLOLQALQONGSSVLSEDKSKRLNTIL KMNEAGAKWSAFYEEOSKOAKTOPLAEIHNTTVKROLQALQOSGSSVLSADKSQRLNTIL KMNEAGAKWSAFYEEOSKOAKTOPLAEIHNTTVKROLQALQOSGSSVLSADKSQRLNTIL KMNEAGARWSAFYEEOSKOAKTOPLEEIQUTVKROLQALQOSGSSVLSADKSKRLNTIL KMNNAGAKWSAFYEEOSKOAKTOPLEEIQDSTVKROLRALQHSGSSVLSADKSKRLNTIL MMNNAGDKWSAFLKEOSTDACOPPLQEICOLTVKLOLQALQOSGTSGLSADKSKRLNTIL NMNNAGDKWSAFLKEOSTDACOPPLQEICOLTVKLOLQALQOSGTSGLSADKSKRLNTIL KMNDARAKWSAFYEEOSKOAKTOPLOEICOLTVKLOLQALQOSGTSGLSADKSKRLNTIL KMNDARAKWSAFYEEOSKOAKTOPLOEICOLTVKLOLQALQOSGTSGLSADKSKRLNTIL KMNDARAKWSAFYEEOSKOAKTOPSGEVONLTVKROLQALQOSGTSGLSADKSKRLNTIL KMNDAEAKWSAFYEEOSKOAKTOPSGEVONLTVKROLQALQOSGSSALSADKSKQLNTIL KMNEARAKWSAFYEEOSKOAKTOPSGEVONLTVKROLQALQSGSSALSADKSKQLNTIL KMNEARAKWSAFYEEOSKOAKTOPSGEVONLTVKROLQALQSGSSALSADKSKQLNTIL KMSEAGAKWAAFYEEOSKOAKTOPSGEVONLTVKROLQALQSGSSALSADKSKQLNTIL KMSEAGAKWAAFYEEOSKOAKTOPSGEVONLTVKROLQALQSGSSALSADKSKVSKINTIL KMSEAGAKWAAFYEEOSKOAKTOPSGEVONLTVKROLQALQSGSSALSADKSKVSN ** ** ** ** ** ** ** ** ** ** ** ** **	120 120 120 120 120 120 120 120 120 120
Q56H28 ACE2_FELCA 2 Q58DD0 ACE2_BOVIN 2 F6V9L3 F6V9I3_HORSE 2 J9P7Y2 J9P7Y2_CANLF 2 A0A2J8KU96 A0A2J8KU96_PANTR 2 K7GLM4 K7GLM4 PIG 2 G1TEF4 G1TEF4_RABIT 2 3 W5PSB6 W5PSB6 SHEEP 2	299 299 298 299 298 299 299 299 299 299	DQAWDAQRIFKEAEKFFVSVGLPNMTQGFWENSMLTDPGNVQKAVCHPTAWDLGGGDFRI NQSWDARRIFKEAEKFFVSVGLPNMTQGFWENSMLTEPGDSRKVVCHPTAWDLGGGDFRI DQSWDARRIFKEAEKFFVSVGLPNMTQGFWENSMLTEPGDGRKVVCHPTAWDLGGGDFRI DQSWDARRIFEEAEKFFVSVGLPNMTQGFWENSMLTEPGDGRKVVCHPTAWDLGGGPRI DQAWDAQRIFKEAEKFFVSVGLPNMTQGFWENSMLTDPGNVQKAVCHPTAWDLGGGPRI DQAWDAQRIFKEAEKFFVSVGLPNMTQGFWENSMLTEPGDGRKVVCHPTAWDLGGGPRI DQAWDAQRIFKEAEKFFVSVGLPNMTQGFWENSMLTEPGDGRKVVCHPTAWDLGGGPRI DQAWDAQRIFKEAEKFFVSVGLPNMTQGFWENSMLTPGDGRKVVCHPTAWDLGGGPRI DQAWDAQRIFKEAEKFFVSVGLPNMTQGFWENSMLTPGDGRKVVCHPTAWDLGGGPRI DQAWDAQRIFKEAEKFFVSVGLPNMTQGFWENSMLTPGDGRKVVCHPTAWDLGGGPRI NQSWDAERIFKEAEKFFVSVGLPSMTQGFWENSMLTPGDGRKVVCHPTAWDLGGGPRI NQSWDAERIFKEAEKFFVSVGLPSMTQGFWENSMLTPFDGDGRKVVCHPTAWDLGGGPRI NQSWDAERIFKEAEKFFVSVGLPSMTQGFWENSMLTPFDGDGRKVVCHPTAWDLGGGPRI NQSWDAERIFKEAEKFFVSVGLPSMTQGFWENSMLTPFDGDGRKVVCHPTAWDLGGGPRI NQSWDAERIFKEAEKFFVSVGLPSMTQGFWENSMLTPFDJDRKVVCHPTAWDLGGGPRI NQSWDAMKIFKTEAEKFFVSVGLPSMTQGFWENSMLTPFDJDRKVVCHPTAWDLGGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTQGFWENSMLTFFDGGGRKVVCHPTAWDLGGGFPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTQGFWENSMLTFFDGFVVCHPTAWDLGGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTYGFWINSMLTFFDGFVVCHPTAWDLGGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTYGFWINSMLTFFDJDRKVVCHPTAWDLGGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTYGFWINSMLTFFDJDRKVVCHPTAWDLGGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTYGFWINSMLTFFDJDRKVVCHPTAWDLGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTYGFWINSMLTFFDJDRKVVCHPTAWDLGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTYGFWINSMLTFFDJDRKVVCHPTAWDLGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTYGFWINSMLTFFDJNRKVVCHPTAWDLGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSMTYGFWINSMLTFFDJNRKVVCHPTAWDLGGPRI NGSWDAMKIFKTEAEKFFVSVGLPSKI XXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXXX	358 357 358 357 358 358 358 358 358 358 358 358

Figure 1. BLAST sequence alignment of the ACE2 receptor from human, cat, cow, horse, dog, chimpanzee, pig, Sumatran orangutan, rabbit, sheep and chicken. The amino acids circled in red are the amino acids favourable for a successful binding to the SARS-CoV-2 receptor binding motif.

4. CONCLUSION

It is clearly evident that two primates in the study, Sumatran orangutans and chimpanzees, sharing the greatest number of favourable amino acids for a successful bind, are the most vulnerable to infection. For this reason, rehabilitation centres and nature reserves will need additional support in keeping these animals protected, and these species may benefit from immunization.

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ETHICS APPROVAL AND CONSENT TO PARTICIPATE

Not applicable.

HUMAN AND ANIMAL RIGHTS

No Animals/Humans were used for studies that are base of this research.

CONSENT FOR PUBLICATION

Not applicable.

AVAILABILITY OF DATA AND MATERIALS

The author confirms that the data supporting the findings of this research are available within the article.

FUNDING

None.

ACKNOWLEDGEMENTS

Many thanks to Dr Clare Roper and Dr Seeta Seetharaman, Wimbledon High School, Professor Charles Swanton FMedSci FRS, Francis Crick Institute, for their guidance and support throughout this research project.

CONFLICT OF INTEREST

Author has no conflict of interest.

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