


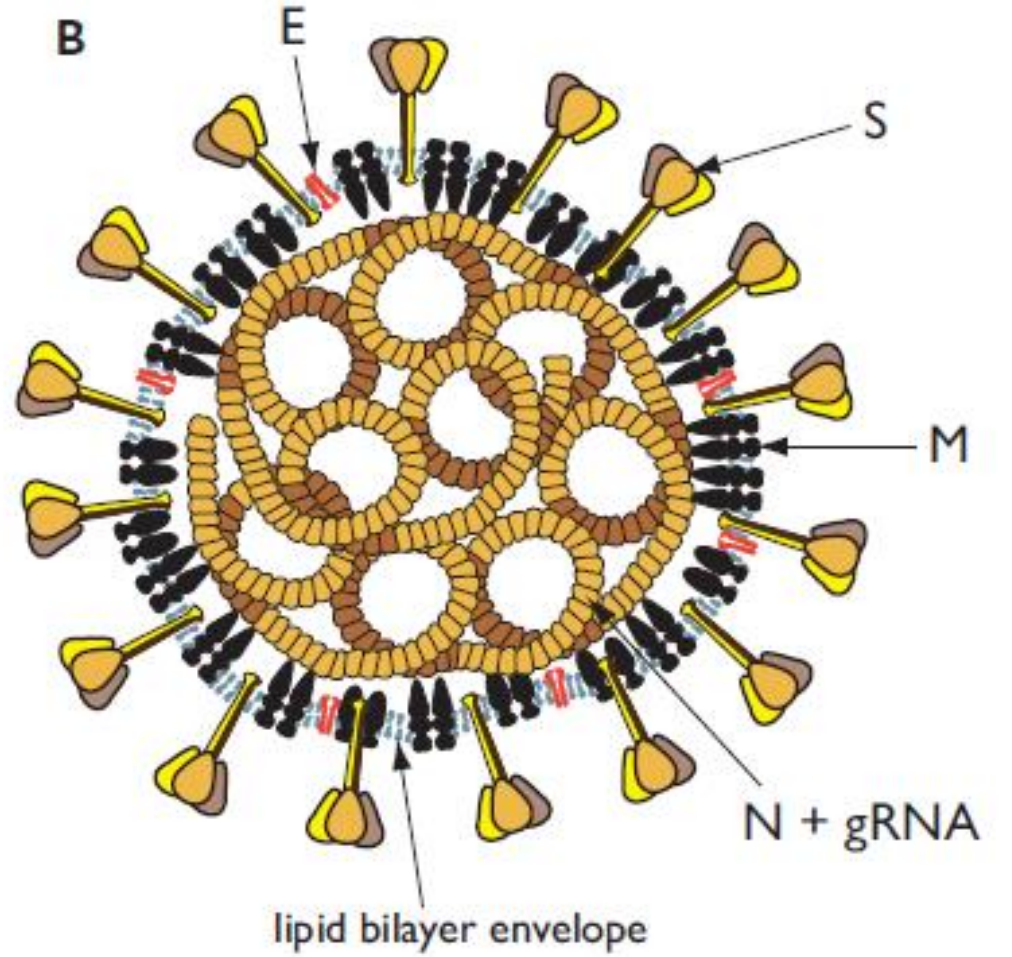
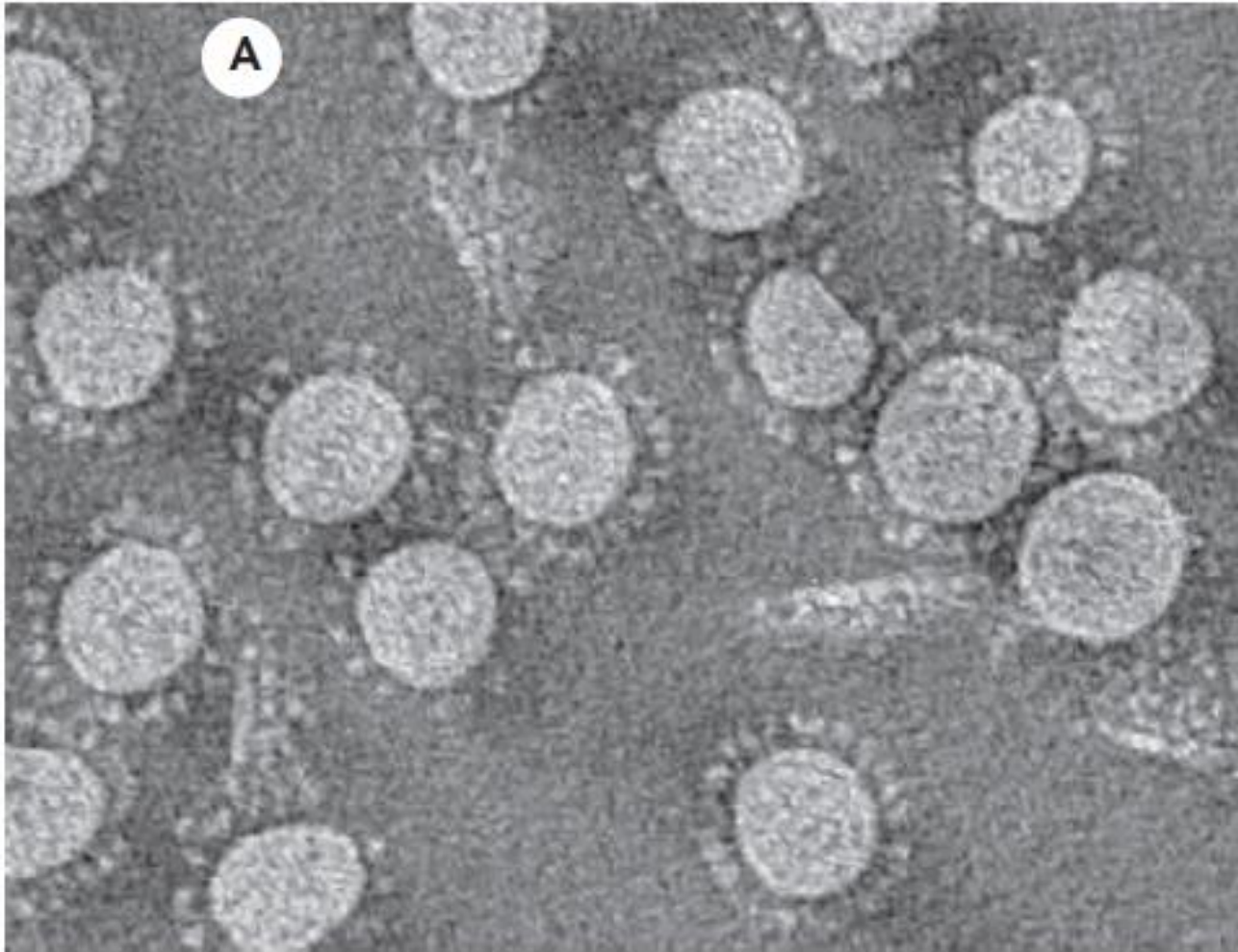
NOVEL CORONAVIRUS


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- **Structure**
 - **Pathogenesis and Clinical Syndromes**
 - **Origin**
 - **Why bats are blind to virus infection?**
 - **Vaccination**
 - **Treatment**

STRUCTURE:



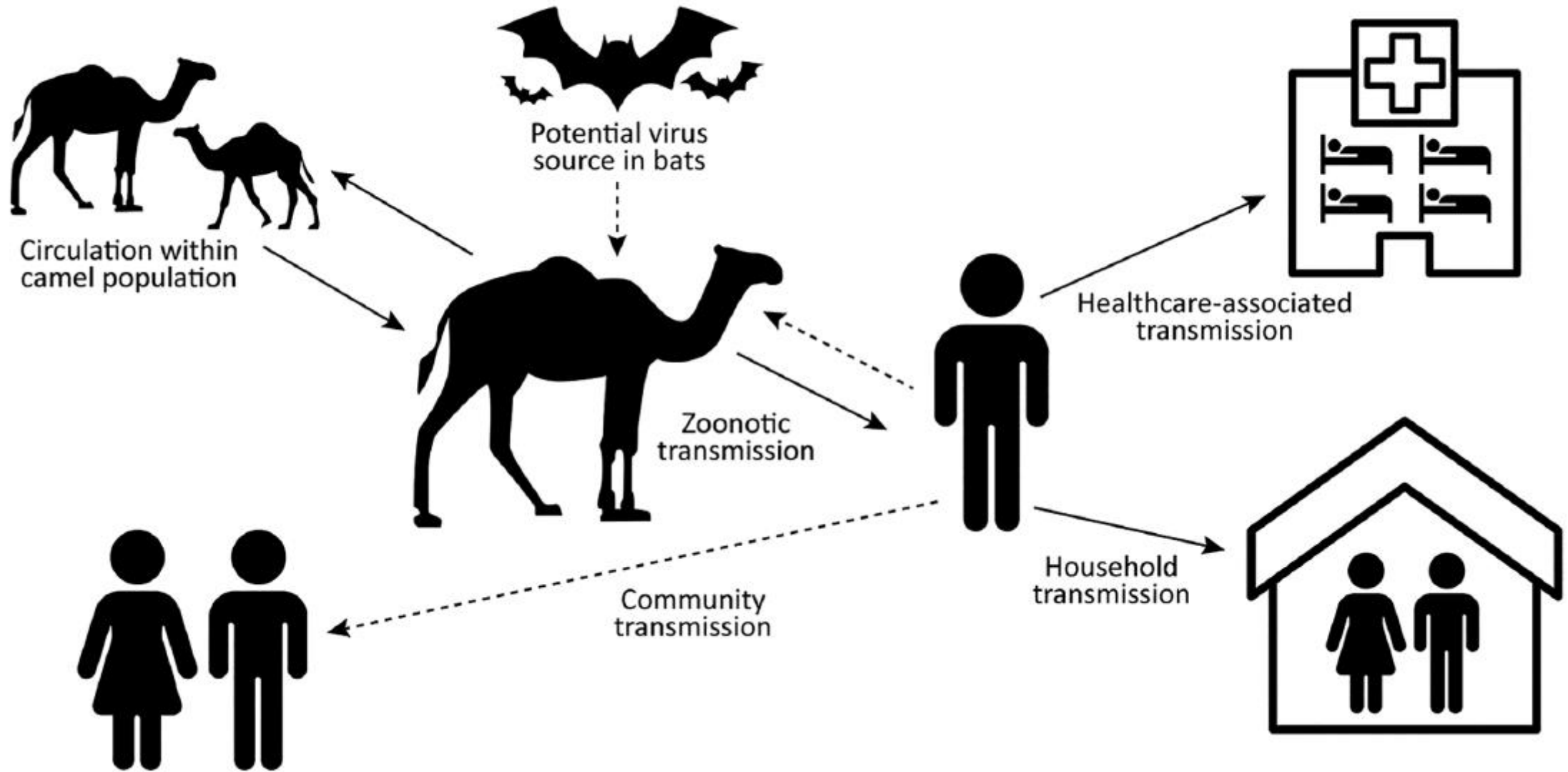
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- **Family : Coronaviridae**
 - **Order : Nidovirales**
 - **Genera : alpha, beta, delta and gamma coronavirus**


- **emerging CoVs:**

1-Severe acute respiratory syndrome coronavirus (SARS-CoV)

2-Middle East respiratory syndrome coronavirus (MERS-CoV)


3- Novel coronavirus belonging to beta coronavirus.



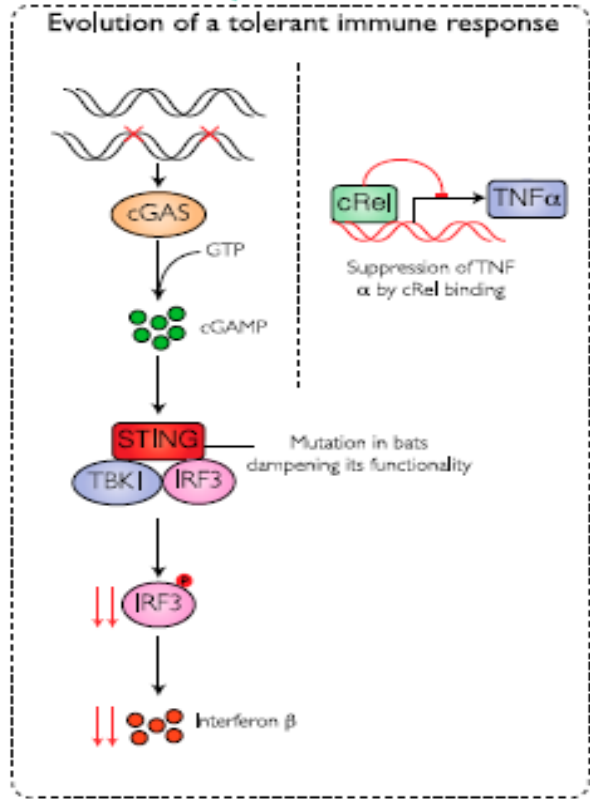
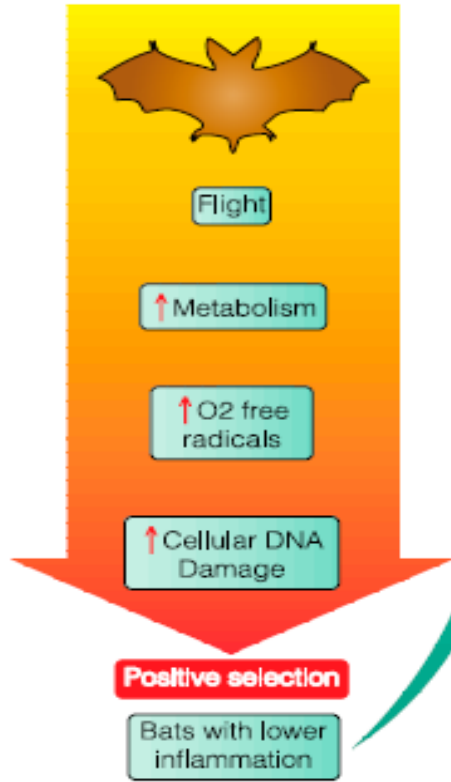
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- The two highly pathogenic viruses, SARS- CoV and MERS- CoV, cause severe respiratory syndrome in humans, and the other four human coronaviruses (HCoV- NL63, HCoV-229E, HCoV- OC43 and HKU1) induce only mild upper respiratory diseases in immunocompetent hosts, although some of them can cause severe infections in infants, young children and elderly individuals
 - Coronavirus can infect mammals, birds and reptile, including human, swine, cattle, horses, camels, cats, dogs, rodents, birds, bats, rabbits, ferrets, mink, snake, and various wildlife species

SYNDROMES:

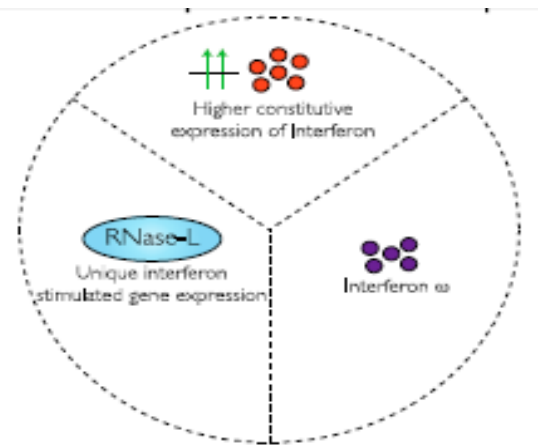
- A range of disease has been observed: fever, dry cough, shortness of breath, and leukopenia pneumonia including difficulty breathing, and bilateral lung infiltration in the most severe cases.
- Patients have included mild cases needing supportive care.
- Host conditions including age, biological sex, and overall health including hypertension, diabetes, heart and/or kidney function issues that may have made them more susceptible.

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- The source of the 2019-nCoV is still unknown.
 - In addition to seafood, it is reported on social media that snakes, birds and other small mammals including marmots and bats were sold at the Huanan South China Seafood Market.
 - The WHO reported that environmental samples taken from the marketplace have come back positive for the novel coronavirus, but no specific animal association has been identified
 - An initial report suggested that snakes might be the possible source based.
 - More interestingly, an origin-unknown homologous recombination was identified within the spike glycoprotein of the 2019-nCoV5, which may explain its decreased pathogenesis, snake-to-human cross species transmission, and limited person-person spread

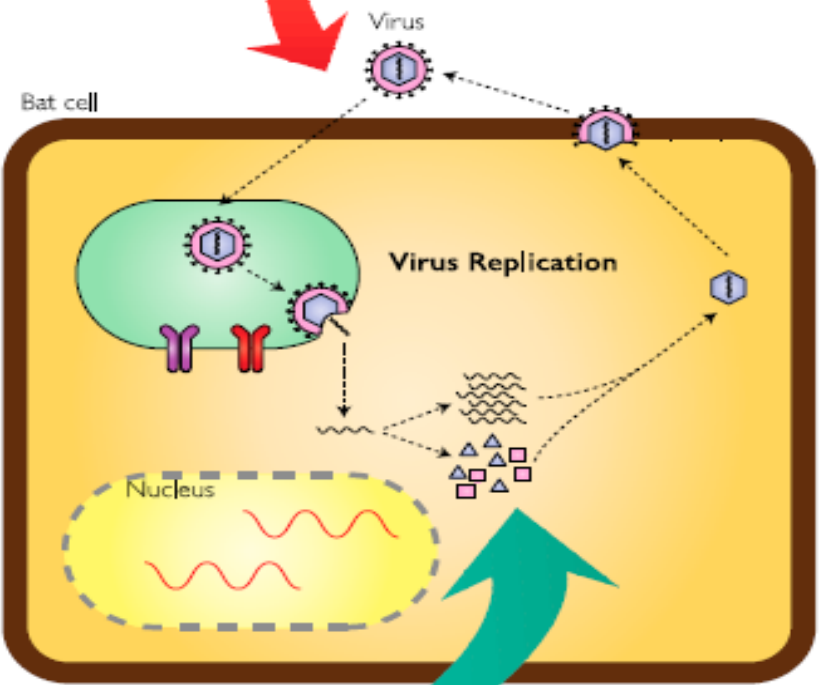
Impact of flight on evolution




Reduced Inflammation



Controlling viral replication



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- Live attenuated virus with gene deletion
 - Two whole virus vaccines
 - DNA vaccine which encodes the full length or part of the S protein gene
 - Most of them have been tested in mouse models and showed the ability to elicit neutralizing antibodies.



- Several bottlenecks


1-a lack of proper animal models for evaluating vaccine efficacy.

2-there are limitations from the s protein itself, such as mutations in the neutralization antibody epitopes in s protein that can cause virus escape non-neutralization antibody epitopes in vaccines that may elicit antibody-mediated disease enhancement (ade)

3- dna vaccines may recombine with other viruses.

4- pre-existing immunity may eliminate the vaccine by removing the general human virus vectors .

5-there is the problem of return on investment which may be slow and, hence, inhibit investments and slow down the clinical study.

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- At the present, no specific antiviral therapy has been approved for treatment of infection by human CoVs.
 - As development of vaccines and compounds for prevention and treatment of infection have been brought to priority status by WHO and governments , numerous drug studies have been done or are moving forward.
 - Some of them focus on the CoV fusion/entry process either by inhibition of S1 mediated virus attachment or by blocking of S mediated virus-cell membrane fusion, and some of them interfere with viral replication .