

Modeling the kinetics of SARS-CoV-2 IgG

Ettore Rocchi, Sara Peluso, Davide Sisti

Unit of Biostatistics and Biomathematics, Dept. Biomolecular Science, Urbino University

Introduction: Up to now, a few articles describing the temporal trend of SARS-Cov-2 IgG, have been produced in the scientific literature and almost all of these articles deal mainly with the onset of the immune response and neglect the descending phase. However, a mathematical model showing the trend and, if necessary, capable of forecasting, has not been published yet.

Methods: We have modeled the kinetics of SARS-Cov-2 specific IgG, based on a single patient case (an Italian 57-year-old male, living in Pesaro (a province of Marche region being one of the main outbreaks of the epidemic in Italy), manifested the onset of symptoms (non-respiratory, but with diarrhea, asthenia, articular and muscle pain, anosmia, and ageusia), limited to the descending phase (during the ascending phase, the immunometric assays were not available yet). The blood samples were taken in the days 66, 95, and 133 from the beginning of the symptoms, and they were analyzed by CLIA chemiluminescence method, Megalumi® SNIBE. Typical equations of first-order kinetics were used to describe the trend of the IgG concentration so that a negative exponential function was obtained: $C = a \exp(-kt)$ (1) With some easy mathematical passages, from (1) we obtain ln(C) = -kt + ln a (2).

Results: The graph of the models is shown in the figure beside. The results show an excellent determination coefficient (R = 0.996). once we have the equation parameters, it is possible to calculate both the half-life of IgG (18 days) and the time of their negativization (170 days).



Discussions and Conclusions: This result does not necessarily imply a loss of immunity at the expected time of negativization. In fact, as some authors have recently highlighted, a cell-mediated immunity could exist. However, our data may suggest the impossibility of identifying the subjects coming into contact with the virus for more than the time of negativization (in our case about six months since the onset of symptoms). Despite the expected inter-subject variability, the perfect adaptation of the negative exponential model seems to be very promising and it suggests that it could fit equally well for other cases, with different parameters.

References:

Lynch KL, *et al.* Clin Infect Dis 2020; doi: 10.1093/cid/ciaa979 Guo L,, et al. Clin Infect Dis 2020; 71(15):778-85 Suhandynata RT, et al. J Appl Lab Med 2020; doi: 10.1093/jalm/jfaa079 Sun B, *et al.* Emerg Microbes Infect 2020; 9(1): 940-8 Sette A, et al. Nat Rev Immunol 2020; 20(8):457-8.

> XLIV CONVEGNO AIE 2-6 NOVEMBRE 2020