



# Kidney injury following JN.1; a mini-review on current evidence

Yassamin Rabiei\*

## Abstract

The SARS-CoV-2 JN.1 variant, a descendant of the Omicron BA.2.86 lineage, emerged as a globally dominant COVID-19 strain by early 2024. This virus characterized by heightened transmissibility and immune evasion, which represents as an evolution in the virus's ability to adapt across with maintaining lower severity versus earlier variants. Since direct evidence on JN.1 and renal involvement is lacking, the variant is expected to cause kidney injury through the well-characterized SARS-CoV-2 mechanisms of direct renal infection and systemic inflammatory damage observed in COVID-19 patients in general.

**Keywords:** SARS-CoV-2 JN.1 variant, Acute kidney injury, Chronic kidney disease, COVID-19, Apolipoprotein L1

**Citation:** Rabiei Y. Kidney injury following JN.1; a mini-review on current evidence. J Ren Endocrinol. 2025;11:e25192. doi: 10.34172/jre.2025.25192.

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## Introduction

The SARS-CoV-2 JN.1 variant, a descendant of the Omicron BA.2.86 sublineage, has emerged as a dominant global strain due to its enhanced transmissibility and immune evasion capabilities (1). This virus was first identified in August 2023 and the classified as a variant of interest by the WHO in December 2023 (2). Prior studies found that, JN.1 accounts for over 90% of sequenced cases in multiple regions as of early 2024 (3). Though, this new variant pretends a lower risk of severe disease compared to earlier variants, however its rapid spread emphasizes the need for sustained vigilance (4). This virus carries over 30 spike protein mutations, including the critical L455S mutation in the receptor-binding motif (5). This mutation reduces ACE2 receptor-binding affinity while enhancing infectivity and immune evasion by hindering antibody neutralization (5,6). Preliminary studies found that JN.1's spike protein adopts conformational changes that improve host cell entry, particularly in lung cells, resembling pre-Omicron variants like Delta (2,7). The transmissibility of JN.1 is 1.5–2 times higher than BA.2.86, driven by its ability to thrive in cold, dry climates and evade population immunity (8). Reports showed that, between November 2023 and March 2024, it caused surges across 121 countries, peaking at 95.1% of global sequences by mid-2024 (2). Common symptoms are resembling to COVID-19 and are sore throat, fever, dry cough, fatigue, nausea/vomiting and loss of taste/smell (9). The regions most affected by the JN.1 variant's spread are primarily in Asia, Singapore, Hong Kong, Thailand, China, and India with significant surges

and increased healthcare burdens reported in these areas (10). Beyond Asia, JN.1 has been documented in over 41 countries, including the United States, United Kingdom, France, Canada, and Sweden, where it has become one of the fastest-growing variants, but the most intense recent impact and public health concern remain focused in Asian countries (11). Recent studies found that, JN.1 exhibits moderate antibody escape, reducing neutralizing antibody titers compared to ancestral strains (3,12). Therefore, updated mRNA vaccines targeting Omicron lineages maintain an appropriated effectiveness against symptomatic infection and a suitable protection against severe disease (13,14). However, declining immunity and low-booster uptake have facilitated its spread (15). It is postulated that, vulnerable populations like elderly, immunocompromised, diabetics and those with comorbidities remain at higher risk of severe outcomes and its complication including kidney involvement (16).

## Search strategy

For this study, we searched PubMed, Web of Science, EBSCO, Scopus, Google Scholar, Directory of Open Access Journals (DOAJ) and Embase, using different keywords including SARS-CoV-2 JN.1 variant, acute kidney injury, chronic kidney disease, COVID-19 and apolipoprotein L1.

## JN.1 versus other Omicron sub-variants

The SARS-CoV-2 JN.1 variant exhibits significantly higher infectivity compared to other Omicron sub-

### ■ Implication for health policy/practice/research/medical education

The SARS-CoV-2 JN.1 variant is expected to affect kidney function through mechanisms similar to those established for SARS-CoV-2 infection. Its renal effects from other variants. SARS-CoV-2 JN.1 could affect kidney function primarily by direct infection of ACE2-expressing renal cells leading to acute tubular injury and glomerular damage, combined with systemic inflammatory and vascular effects that contribute to acute kidney injury and long-term renal function decline.

variants, particularly its direct ancestor BA.2.86 (1). This condition increased infectivity is primarily attributed to the unique L455S mutation in the spike protein's receptor-binding motif, which enhances viral entry efficiency into human nasal epithelial cells and improves spike protein cleavage (1,17). Despite a decreased binding affinity to the ACE2 receptor relative to BA.2.86, JN.1 compensates with enhanced cell entry mechanisms and immune evasion, making it more transmissible (18). Laboratory studies demonstrate that JN.1 infects primary human nasal epithelial cells more effectively than BA.2.86, a key factor in its rapid global spread and dominance (19). This mutation alters the spike protein's interaction with ACE2 and increases the number of hydrogen bonds within the spike structure, facilitating better viral attachment and fusion (20). Compared to other Omicron sub-variants, the JN.1 transmissibility is estimated to greater, contributing to its rapid replacement of earlier variants worldwide since late 2023 (1). While it retains immune evasion features typical of Omicron lineages, JN.1's enhanced infectivity distinguishes it as one of the most transmissible variants currently circulating (21). The immune evasion mechanism of the SARS-CoV-2 JN.1 variant differs from previous variants primarily due to the novel L455S mutation in the spike protein's receptor-binding motif (22). This mutation alters the spike protein structure, decreasing its binding affinity to the human ACE2 receptor compared to its ancestor BA.2.86, yet paradoxically enhances infectivity and immune escape (23). Unlike earlier variants, JN.1's L455S mutation significantly hinders the ability of neutralizing antibodies, especially those targeting the receptor-binding domain to attach effectively to the virus (24). This circumstance leads to a more robust evasion of antibody-mediated immunity, including a marked reduction in neutralization by antibodies derived from previous infections or vaccinations (25). Studies show JN.1 exhibits a greater fold reduction in neutralizing antibody titers than BA.2.86 and other Omicron sub-variants, indicating stronger immune escape (26). Moreover, JN.1 evades a broad range of neutralizing antibodies, including those derived from common B cell germlines (IGHV3-53/3-66), which were effective against earlier variants (27). This immune evasion is partly compensated by preserved Fc effector

functions mediated by class 3 antibodies binding outside the ACE2 interface, which may contribute to the relatively lower severity of disease despite immune escape (28).

### Kidney injury following JN.1

There is no specific published evidence directly linking the SARS-CoV-2 JN.1 variant to unique or distinct renal involvement compared to other variants (29). However, according to the previous evidence on SARS-CoV-2 kidney pathology, the mechanisms by which COVID-19 causes kidney injury are likely applicable to JN.1 as well (30). SARS-CoV-2 infection can cause kidney damage through several pathways by direct infection of renal cells via ACE2 receptors expressed on podocytes and proximal tubular cells, leading to acute tubular injury and collapsing glomerulopathy (31). In addition, indirect effects including systemic inflammation, cytokine storm, endothelial dysfunction, hypercoagulability, and hypoxia, which contribute to acute kidney injury (AKI) (32). Moreover, residual inflammation and acute injury can result in progressive kidney function decline, even in mild-to-moderate COVID-19 cases (33). In fact, AKI occurs in a significant portion of hospitalized COVID-19 patients which ranging from 5% to over 30% in severe cases (34,35). This condition is associated with higher mortality and risk of chronic kidney disease progression (36). Histopathological studies have detected viral RNA and particles in kidney tissue, supporting direct viral involvement (37). Given that JN.1 retains the ability to infect human cells via ACE2 and causes systemic infection, it is plausible that it can induce kidney injury through similar mechanisms as earlier variants (38). However, no current data specifically describe renal complications uniquely attributable to JN.1 (39). Previous studies demonstrated that, SARS-CoV-2 impacts renal tissue, following direct viral infection of kidney cells (40). This virus binds to ACE2 receptors, which are highly expressed on renal proximal tubular epithelial cells, podocytes, mesangial cells, and parietal epithelial cells (41). This allows direct viral entry and replication in these cells, causing cytopathic effects and acute tubular injury (42). One of the presentations of renal involvement is AKI, which is common in COVID-19 patients, mainly in severe cases (43). Acute kidney injury results from direct viral damage, immune dysregulation, cytokine storm, endothelial dysfunction, and hypercoagulability (32). Viral particles have been identified in renal tubular cells and podocytes, supporting direct kidney tropism (44). Studies show that COVID-19 accelerates decline in estimated glomerular filtration rate, especially in hospitalized patients, increasing risk for chronic kidney disease progression (45). However, study regarding indirect systemic effects, detected that, hypoxia, coagulation abnormalities, multi-organ injury, and nephrotoxic medications also contribute to renal impairment during infection (46). Since JN.1 retains the ability to infect human cells via ACE2 and causes systemic

infection, it likely induces kidney injury through these established pathways (38). Therefore, JN.1 can cause acute kidney injury, proteinuria, and accelerated kidney function decline similar to other SARS-CoV-2 variants (47).

### Focus on genetic factors

Several studies showed that, individuals with two risk alleles of apolipoprotein L1 (APOL1) variants, (G1 and G2), predominantly found in people of West and Central African ancestry, have a significantly increased risk of developing kidney diseases such as focal segmental glomerulosclerosis (FSGS), collapsing glomerulopathy, and HIV-associated nephropathy (48). These APOL1 risk alleles confer an increased risk for certain kidney diseases and are associated with worse outcomes in viral-associated nephropathies (49). A “second hit” such as viral infection or inflammation often triggers disease in genetically susceptible individuals (50). Other podocyte and glomerular gene mutations are consisted of mutations in genes affecting podocyte structure and function such as NPHS1, NPHS2, WT1, TRPC6, INF2, that predispose individuals to nephrotic syndrome and glomerular diseases, which can be exacerbated by viral infections (51). In this regard, a genetic predisposition to AKI is also recently explained (52). Variants in genes related to immune response and kidney function may influence susceptibility to AKI during systemic infections like COVID-19 (53).

### Conclusion

There are no specific studies yet identifying genetic factors that make individuals more susceptible to kidney damage specifically from the SARS-CoV-2 JN.1 variant. However, based on general knowledge of genetic susceptibility to kidney disease and COVID-19-related kidney injury, some genetic variants are known to increase the risk of kidney damage in viral infections, which likely applies to JN.1 as well. Since JN.1 shares the fundamental mechanism of kidney cell infection via ACE2 and induces systemic inflammation similar to other SARS-CoV-2 variants, individuals with high-risk genetic backgrounds (especially those with APOL1 risk alleles) may be more vulnerable to kidney injury or collapsing glomerulopathy triggered by JN.1 infection.

### Conflicts of interest

The author declares that she has no competing interests.

### Ethical issues

Ethical issues (including plagiarism, data fabrication, double publication) have been completely observed by the author.

### Declaration of generative AI and AI-assisted technologies in the writing process

During the preparation of this work, the authors utilized Perplexity to refine grammar points and language style in writing. Subsequently, the authors thoroughly reviewed and edited the content as

necessary, assuming full responsibility for the publication's content.

### Funding/Support

None.

### References

1. Lu Y, Ao D, He X, Wei X. The rising SARS-CoV-2 JN.1 variant: evolution, infectivity, immune escape, and response strategies. *MedComm* (2020). 2024;5:e675. doi: 10.1002/mco2.675.
2. Naveed Siddiqui A, Musharaf I, Gulumbe BH. The JN.1 variant of COVID-19: immune evasion, transmissibility, and implications for global health. *Ther Adv Infect Dis*. 2025;12:20499361251314763. doi: 10.1177/20499361251314763.
3. Jian F, Wang J, Yisimayi A, Song W, Xu Y, Chen X, et al. Evolving antibody response to SARS-CoV-2 antigenic shift from XBB to JN.1. *Nature*. 2025;637:921-9. doi: 10.1038/s41586-024-08315-x.
4. Hemo MK, Islam MA. JN.1 as a new variant of COVID-19 - editorial. *Ann Med Surg (Lond)*. 2024;86:1833-5. doi: 10.1097/ms9.0000000000001876.
5. Li P, Faraone JN, Hsu CC, Chamblee M, Zheng YM, Carlin C, et al. Characteristics of JN.1-derived SARS-CoV-2 subvariants Slip, FLiRT, and KP.2 in neutralization escape, infectivity and membrane fusion. *BioRxiv*. 2024. doi: 10.1101/2024.05.20.595020.
6. Xue S, Han Y, Wu F, Wang Q. Mutations in the SARS-CoV-2 spike receptor binding domain and their delicate balance between ACE2 affinity and antibody evasion. *Protein Cell*. 2024;15:403-18. doi: 10.1093/procel/pwae007.
7. Candido KL, Eich CR, de Fariña LO, Kadowaki MK, da Conceição Silva JL, Maller A, et al. Spike protein of SARS-CoV-2 variants: a brief review and practical implications. *Braz J Microbiol*. 2022;53:1133-57. doi: 10.1007/s42770-022-00743-z.
8. Fan J, Zhang Y, Li S, Li Q, Zi Q, Mou X, et al. Immunogen characterization reveals an intrinsic hindrance in eliciting neutralizing antibodies against JN.1 variant. *iScience*. 2024;27:110405. doi: 10.1016/j.isci.2024.110405.
9. Yang S, Yu Y, Xu Y, Jian F, Song W, Yisimayi A, et al. Fast evolution of SARS-CoV-2 BA. 2.86 to JN. 1 under heavy immune pressure. *Lancet Infect Dis*. 2024;24:e70-e2.
10. Chong C, Wee LE, Jin X, Zhang M, Malek MIA, Ong B, et al. Risks of SARS-CoV-2 JN.1 Infection and COVID-19-Associated Emergency Department Visits/Hospitalizations Following Updated Boosters and Prior Infection: A Population-Based Cohort Study. *Clin Infect Dis*. 2024 Nov 22;79:1190-1196. doi: 10.1093/cid/ciae339. PMID: 38922669.
11. Looi MK. Covid-19: WHO adds JN.1 as new variant of interest. *BMJ*. 2023 Dec 21;383:2975. doi: 10.1136/bmj.p2975. PMID: 38128957.
12. Wang X, Jiang S, Ma W, Zhang Y, Wang P. Robust neutralization of SARS-CoV-2 variants including JN.1 and BA.2.87.1 by trivalent XBB vaccine-induced antibodies. *Signal Transduction and Targeted Therapy*. 2024;9:123. doi: 10.1038/s41392-024-01849-6.
13. Liu J, Wang L, Kurtesi A, Budylowski P, Potts KG, Menon H, et al. A bivalent COVID-19 mRNA vaccine elicited broad immune responses and protection against Omicron subvariants infection. *NPJ Vaccines*. 2025 Jan 10;10:4. doi: 10.1038/s41541-025-01062-8. PMID: 39788981; PMCID: PMC11718203.
14. Chatterjee S, Bhattacharya M, Nag S, Dhama K, Chakraborty C. A Detailed Overview of SARS-CoV-2 Omicron: Its Sub-Variants, Mutations and Pathophysiology, Clinical Characteristics, Immunological Landscape, Immune Escape,



- and Therapies. *Viruses*. 2023;15. doi: 10.3390/v15010167.
15. Ramesh S, Govindarajulu M, Parise RS, Neel L, Shankar T, Patel S, et al. Emerging SARS-CoV-2 Variants: A Review of Its Mutations, Its Implications and Vaccine Efficacy. *Vaccines (Basel)*. 2021;9. doi: 10.3390/vaccines9101195.
  16. Huang T-S, Chao J-Y, Chang H-H, Lin W-R, Lin W-H. COVID-19 and Diabetes: Persistent Cardiovascular and Renal Risks in the Post-Pandemic Landscape. *Life*. 2025;15:726.
  17. Li P, Liu Y, Faraone J, Hsu CC, Chamblee M, Zheng YM, et al. Distinct Patterns of SARS-CoV-2 BA.2.87.1 and JN.1 Variants in Immune Evasion, Antigenicity and Cell-Cell Fusion. *BioRxiv*. 2024. doi: 10.1101/2024.03.11.583978.
  18. Khan T, Shahab M, Alharbi AM, Waqas M, Zakirullah, Zheng G. The Omicron variant BA.2.86.1 of SARS-CoV-2 demonstrates an altered interaction network and dynamic features to enhance the interaction with the hACE2. *Scie Rep*. 2025;15:6482. doi: 10.1038/s41598-025-89548-2.
  19. Angius F, Puxeddu S, Zaimi S, Canton S, Nematollahzadeh S, Pibiri A, et al. SARS-CoV-2 Evolution: Implications for Diagnosis, Treatment, Vaccine Effectiveness and Development. *Vaccines (Basel)*. 2024;13. doi: 10.3390/vaccines13010017.
  20. Khan MI, Baig MH, Mondal T, Alorabi M, Sharma T, Dong JJ, et al. Impact of the Double Mutants on Spike Protein of SARS-CoV-2 B.1.617 Lineage on the Human ACE2 Receptor Binding: A Structural Insight. *Viruses*. 2021;13. doi: 10.3390/v13112295.
  21. Carabelli AM, Peacock TP, Thorne LG, Harvey WT, Hughes J, de Silva TI, et al. SARS-CoV-2 variant biology: immune escape, transmission and fitness. *Nat Rev Microbiol*. 2023;21:162-77. doi: 10.1038/s41579-022-00841-7.
  22. Guo H, Ha S, Botten JW, Xu K, Zhang N, An Z, et al. SARS-CoV-2 Omicron: Viral Evolution, Immune Evasion, and Alternative Durable Therapeutic Strategies. *Viruses*. 2024;16. doi: 10.3390/v16050697.
  23. Yajima H, Anraku Y, Kaku Y, Kimura KT, Plianchaisuk A, Okumura K, et al. Structural basis for receptor-binding domain mobility of the spike in SARS-CoV-2 BA.2.86 and JN.1. *Nat Commun*. 2024;15:8574. doi: 10.1038/s41467-024-52808-2.
  24. Yang H, Guo H, Wang A, Cao L, Fan Q, Jiang J, et al. Structural basis for the evolution and antibody evasion of SARS-CoV-2 BA.2.86 and JN.1 subvariants. *Nat Commun*. 2024;15:7715. doi: 10.1038/s41467-024-51973-8.
  25. Haque A, Pant AB. The coevolution of COVID-19 and host immunity. *Exploration of Medicine*. 2024;5:167-84.
  26. Springer DN, Camp JV, Aberle SW, Deutsch J, Lammel O, Weseslindtner L, et al. Neutralization of SARS-CoV-2 Omicron XBB. 1.5 and JN. 1 variants after COVID-19 booster-vaccination and infection. *J Med Virol*. 2024;96:e29801.
  27. Kuwata T, Kaku Y, Biswas S, Matsumoto K, Shimizu M, Kawanami Y, et al. Induction of IGHV3-53 public antibodies with broadly neutralising activity against SARS-CoV-2 including Omicron subvariants in a Delta breakthrough infection case. *EBioMedicine*. 2024;110:105439. doi: 10.1016/j.ebiom.2024.105439.
  28. Cox M, Peacock TP, Harvey WT, Hughes J, Wright DW, Willett BJ, et al. SARS-CoV-2 variant evasion of monoclonal antibodies based on in vitro studies. *Nat Rev Microbiol*. 2023;21:112-24. doi: 10.1038/s41579-022-00809-7.
  29. Kaku Y, Okumura K, Padilla-Blanco M, Kosugi Y, Uriu K, Hinay AA, et al. Virological characteristics of the SARS-CoV-2 JN. 1 variant. *Lancet Infect Dis*. 2024;24:e82.
  30. Faour WH, Choaib A, Issa E, Choueiry FE, Shbaklo K, Alhaji M, et al. Mechanisms of COVID-19-induced kidney injury and current pharmacotherapies. *Inflamm Res*. 2022;71:39-56. doi: 10.1007/s00011-021-01520-8.
  31. Bärreiter VA, Meister TL. Renal implications of coronavirus disease 2019: insights into viral tropism and clinical outcomes. *Curr Opin Microbiol*. 2024;79:102475. doi: 10.1016/j.mib.2024.102475.
  32. He W, Liu X, Hu B, Li D, Chen L, Li Y, et al. Mechanisms of SARS-CoV-2 Infection-Induced Kidney Injury: A Literature Review. *Front Cell Infect Microbiol*. 2022;12:838213. doi: 10.3389/fcimb.2022.838213.
  33. Nlandu Y, Tannor EK, Bafemika T, Makulo JR. Kidney damage associated with COVID-19: from the acute to the chronic phase. *Ren Fail*. 2024;46:2316885. doi: 10.1080/0886022x.2024.2316885.
  34. Borrego-Moreno JC, Cárdenas-de Luna MJ, Márquez-Castillo JC, Reyes-Ruiz JM, Osuna-Ramos JF, León-Juárez M, et al. Acute Kidney Injury in the Context of COVID-19: An Analysis in Hospitalized Mexican Patients. *Infect Dis Rep*. 2024;16:458-71. doi: 10.3390/idr16030034.
  35. Chan L, Chaudhary K, Saha A, Chauhan K, Vaid A, Zhao S, et al. AKI in Hospitalized Patients with COVID-19. *J Am Soc Nephrol*. 2021;32:151-60. doi: 10.1681/asn.2020050615.
  36. Eyitemi J, Thomas B, Ramos Y, Feng X, Ezekwesili C. SARS-CoV-2: Review of Conditions Associated With Severe Disease and Mortality. *Int J Prev Med*. 2022;13:109. doi: 10.4103/ijpvm.ijpvm\_640\_20.
  37. Hassler L, Reyes F, Sparks MA, Welling P, Batlle D. Evidence For and Against Direct Kidney Infection by SARS-CoV-2 in Patients with COVID-19. *Clin J Am Soc Nephrol*. 2021;16:1755-65. doi: 10.2215/cjn.04560421.
  38. Jackson CB, Farzan M, Chen B, Choe H. Mechanisms of SARS-CoV-2 entry into cells. *Nat Rev Mol Cell Biol*. 2022;23:3-20. doi: 10.1038/s41580-021-00418-x.
  39. Fabrizi F, Nardelli L, Regalia A, Zannoni F, Castellano G. Are Kidneys Affected by SARS-CoV-2 Infection? An Updated Review on COVID-19-Associated AKI. *Pathogens*. 2024;13. doi: 10.3390/pathogens13040325.
  40. Chen Z, Hu J, Liu L, Chen R, Wang M, Xiong M, et al. SARS-CoV-2 Causes Acute Kidney Injury by Directly Infecting Renal Tubules. *Front Cell Dev Biol*. 2021;9:664868. doi: 10.3389/fcell.2021.664868.
  41. Sharma RK, Li J, Krishnan S, Richards EM, Raizada MK, Mohandas R. Angiotensin-converting enzyme 2 and COVID-19 in cardiorenal diseases. *Clin Sci (Lond)*. 2021;135:1-17. doi: 10.1042/cs20200482.
  42. Pramod S, Kheetan M, Ogu I, Alsanani A, Khitan Z. Viral Nephropathies, Adding SARS-CoV-2 to the List. *Int J Nephrol Renovasc Dis*. 2021;14:157-64. doi: 10.2147/ijnrd.S303080.
  43. Giannini G, Carlos QVJ, May RM, Sharma SG, Mohamed MMB, Cassol CA, et al. Renal Prognosis of COVID-19 Associated Nephropathy. *Kidney Int Rep*. 2022;7:2722-5. doi: 10.1016/j.ekir.2022.09.027.
  44. Papadimitriou JC, Drachenberg CB, Kleiner D, Choudhri N, Haririan A, Cebotaru V. Tubular Epithelial and Peritubular Capillary Endothelial Injury in COVID-19 AKI. *Kidney Int Rep*. 2021;6:518-25. doi: 10.1016/j.ekir.2020.10.029.
  45. Mahalingasivam V, Faucon AL, Sjölander A, Bosi A, González-Ortiz A, Lando S, et al. Kidney Function Decline After COVID-19 Infection. *JAMA Netw Open*. 2024;7:e2450014. doi: 10.1001/jamanetworkopen.2024.50014.
  46. Wang M, Xiong H, Chen H, Li Q, Ruan XZ. Renal Injury by SARS-CoV-2 Infection: A Systematic Review. *Kidney Dis (Basel)*. 2021;7:100-10. doi: 10.1159/000512683.
  47. Legrand M, Bell S, Forni L, Joannidis M, Koyner JL, Liu K, et al. Pathophysiology of COVID-19-associated acute kidney injury. *Nat Rev Nephrol*. 2021;17:751-64. doi: 10.1038/s41581-021-00452-0.
  48. Mupepe DM, Wameso M-NN, Situakibanza HN, Ekulu PM, Makulo JRR, Kayembe JMN, et al. G1 and G2 variants

- of apolipoprotein L1 among Central African population in Trypanosoma brucei gambiense endemic rural area. *The Journal of Infection in Developing Countries*. 2024;18:1610-6.
49. Pollak MR, Friedman DJ. APOL1 and APOL1-Associated Kidney Disease: A Common Disease, an Unusual Disease Gene - Proceedings of the Henry Shaville Professorship. *Glomerular Dis*. 2023;3:75-87. doi: 10.1159/000529227.
  50. Tabachnikov O, Skorecki K, Kruzel-Davila E. APOL1 nephropathy - a population genetics success story. *Curr Opin Nephrol Hypertens*. 2024;33:447-55. doi: 10.1097/mnh.0000000000000977.
  51. Chen YM, Liapis H. Focal segmental glomerulosclerosis: molecular genetics and targeted therapies. *BMC Nephrol*. 2015;16:101. doi: 10.1186/s12882-015-0090-9.
  52. Li Y, Gong Y, Xu G. New insights into kidney disease after COVID-19 infection and vaccination: histopathological and clinical findings. *QJM*. 2024;117:317-37. doi: 10.1093/qjmed/hcad159.
  53. Wang L, Chen A, Zhang L, Zhang J, Wei S, Chen Y, et al. Deciphering the molecular nexus between Omicron infection and acute kidney injury: a bioinformatics approach. *Front Mol Biosci*. 2024;11:1340611. doi: 10.3389/fmolb.2024.1340611.