

Timeline of the Works on the Correlation between COVID-19 and Y-DNA Haplogroup R1b

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Abstract

Soon after the beginning of COVID-19 pandemic, many works have been reporting a yet-to-be-explained correlation of COVID-19 prevalence and severity with Y-DNA haplogroup R1b frequency. Here a chronologically ordered list of such works is provided.

Introduction

Because of the unusual spreading pattern of COVID-19 in Italy and in the world, at the beginning of April, I started wondering whether there was an underlying genetic reason for that. Watching genetic maps, I quickly noticed an apparent correlation with Y-DNA haplogroup R1b. On April 13th, 2020, I set up a website^[4] to publicize such discovery, joined a COVID-related Facebook group and posted my theory there^[5].

Initially, I shared my idea hoping someone would help me to develop it, but people seemed skeptical about it. After trying for a while to convince them, I decided to go on by myself. Using all the freely available data I could gather, I tried to verify the correlation statistically. As far as I know, my article^[19] has been the first serious attempt to mathematically ‘prove’ the existence of such a correlation. With further researches I later found out that other people had also noticed the same or similar correlations — most of them, I presume, independently from me.

My initial plan was to assess the correlation between basic reproduction numbers R_0 and haplogroup percentages. Ideally, it would have been better to do it at a local level, so to compare data collected in a more uniform way. Unfortunately, there were not enough data for the haplogroup distributions inside countries or regions, but only for whole countries and populations (and only for the main subclades).

Besides, after searching for a while, I realized it was also difficult to find enough good quality estimates of basic reproduction numbers. So I resolved to use the initial growth rates of contagion (and deaths) for different countries, instead. To prevent possible criticisms, I calculated them using exactly the same approach already employed in another article assessing correlations with different factors; interestingly enough, the estimates chosen for the free parameters (days and cases) tend to be locally optimal for R1b correlations.

While writing my article I have started keeping track of related works, for easier reference. The following table contains an annotated list, in chronological order, of all the mentions to this theory I learned about. The milestones are highlighted and the specified date is the earliest I could find a reference for.

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DATE	AUTHOR	TYPE	NOTES
22 March 2020	Maria del Pilar Corena-McLeod	Thread ^[1]	Incorrectly refers to the <i>mitochondrial</i> haplogroup R1b
30 March 2020	Gigi Tevzadze	Article ^[2]	Marginal reference, mainly on haplogroup G
31 March 2020	Paolo Sizzi	Thread ^[3]	Refers to Italy
13 April 2020	Sebastiano Schillaci	Webpage ^[4]	Main points of the theory and visual comparison (10+ countries)
14 April 2020 (submitted on 13 April 2020)	Sebastiano Schillaci	Thread ^[5]	Shared webpage on Facebook Group <i>physicists against sars-cov-2</i>
17 April 2020	Sebastiano Schillaci	Thread ^[6]	Consulted Mauricio Lucioni Maristany (Maulucioni), main contributor to the world R1b map ^[7]
19 April 2020	Ángel Gómez Moreno	Article ^[8]	Refers to Europe and Africa with main focus on Spain (1/3)
19 April 2020	Alberto Pérez de Vargas	News ^[9]	
24 April 2020	Frédéric Boyer	Blog Post ^[10]	Refers to Europe
25 April 2020	Mohammed Hassai Awli	Article ^[11]	Mostly diagrams
25 April 2020	Miquel Hernandis	News ^[12]	
26 April 2020	Johann Strauss	Blog Post ^[13]	Relationship with BCG vaccine
27 April 2020	Arsenio Escolar	News ^[14]	
29 April 2020	Anonymous (user4682)	Thread ^[15]	Graphical correlation with deaths per capita (20 countries)
5 May 2020	Ángel Gómez Moreno	Article ^[16]	Adds further European and African countries (2/3)
19 May 2020	Antonio Caselles Moncho	Article ^[17]	Multiple linear regression (?) (10 countries)
21 May 2020	Frank Wallace Bentrem	Article ^[18]	Graphical correlation with deaths per capita (57 countries)
22 May 2020	Sebastiano Schillaci	Article ^[19]	Quantitative evaluation of the correlation in Italy and in the world (84 countries)
23 May 2020	Sebastiano Schillaci	Article ^[19]	Submitted to <i>bioRxiv</i> , but refused because it ‘does not accept preprints of articles reporting epidemiological studies’
25 May 2020	Juan-Ramón Lacadena	Blog Post ^[20]	Refers to previous works
26 May 2020	Sebastiano Schillaci	Article ^[19]	Submitted to <i>Open Science Framework (OSF)</i>
27 May 2020	Sebastiano Schillaci	Thread ^[21]	Shared article on Facebook Group <i>physicists against sars-cov-2</i>
5 June 2020	Ángel Gómez Moreno	Article ^[22]	Adds United States, Brazil and Japan (3/3)
8 June 2020	John Allen	Article ^[23]	Attempt at finding the involved genes
19 June 2020 (submitted on 10 June 2020)	Mariya V. Ragulskaya	Article ^[24]	Refers to haplogroup R1a and R1b (peer-reviewed)
27 June 2020	Sebastiano Schillaci	Article ^[19]	2nd version
17 July 2020 (submitted on 20 June 2020)	O. Abu Hammad et al.	Article ^[25]	Marginal reference to previous works (peer-reviewed)
20 July 2020	Spyros Anagnostou	Article ^[26]	Graphical correlation with deaths per capita (38 countries)
20 July 2020	Ángel Gómez Moreno	Article ^[27]	Compares Western European and non-European countries
29 July 2020	Sebastiano Schillaci	Article ^[19]	3rd version
30 July 2020	Sebastiano Schillaci	Thread ^[28]	Shared article on Facebook Group <i>Statistiche Coronavirus Italia</i>
10 August 2020	Dave Dalton	Article ^[29]	Marginal reference, mainly on haplogroup R1a
21 August 2020 (submitted on 27 May 2020)	J.R. Delanghe et al.	Article ^[30]	Quantitative evaluation of the correlation in the Netherlands, Belgium and in the world (28 countries) (peer-reviewed)
13 October 2020	G. Fazio et al.	Article ^[31]	Refers to Italy
15 October 2020 (submitted on 28 June 2020)	L. Janda et al.	Article ^[32]	Marginal reference to previous works (peer-reviewed)
20 October 2020	Aleš Žužek	News ^[33]	
23 October 2020	M. Montopoli et al.	Article ^[34]	Response to J.R. Delanghe et al.’s article (peer-reviewed)
26 October 2020	Nacho de Blas	Blog Post ^[35]	Marginal reference to previous works
9 November 2020	Sandro Modeo	News ^[36]	
12 November 2020	Nacho de Blas	TV News ^[37]	
13 November 2020	Ole Bernt Lenning	Seminar ^[38]	Refers to the world
14 November 2020	Luis Ordóñez	News ^[39]	
23 November 2020	Nacho de Blas	Blog Post ^[40]	Refers to previous works
10 December 2020	Vasilis S. Gavalas	Article ^[41]	Assessment of NPIs efficacy on the basis of previous works
23 December 2020	Sandro Modeo	News ^[42]	
27 December 2020	José Manuel Etxaniz Makazaga	News ^[43]	
8 January 2021	Peter Mühlbauer	News ^[44]	
14 January 2021	Peter Mühlbauer	News ^[45]	
3 February 2021	Sandro Modeo	News ^[46]	

Table: Timeline of the works on COVID-19 and Y-DNA haplogroup R1b.

References

- [1] Corena-McLeod, M. del P. *Wondering if anyone is doing research on mitochondrial haplogroup R1b and its relationship with COVID-19 incidence and symptom severity?*. ResearchGate. **March 22, 2020**. https://www.researchgate.net/post/Wondering_if_anyone_is_doing_research_on_mitochondrial_haplogroup_R1b_and_its_relationship_with_COVID-19_incidence_and_symptom_severity
- [2] Tevzadze, G. *Haplogeography of COVID-19: A hypothesis*. ResearchGate. **March 30, 2020**. <https://doi.org/10.13140/RG.2.2.17496.44803>
- [3] Sizzi, P. *Tweet (in Italian)*. Twitter. **March 31, 2020**. <https://twitter.com/granlombard/status/1245085796543205376>
- [4] Schillaci, S. *Possible link between COVID-19 susceptibility and genetic factors*. SXS. **April 13, 2020**. <http://sxs.altervista.org/coronavirus/>
- [5] Schillaci, S. *physicists against sars-cov-2*. Facebook Groups. **April 14, 2020**. <https://www.facebook.com/groups/PhysicistsAgainstSARSCoV2/permalink/839378729887388>
- [6] Schillaci, S. *User talk:Maulucioni*. Wikimedia Commons. **April 17, 2020**. https://commons.wikimedia.org/wiki/User_talk:Maulucioni#R1b_distribution
- [7] Wikimedia Commons contributors. *Haplogroup R1b (Y-DNA)*. Wikimedia Commons. **July 28, 2019**. [https://upload.wikimedia.org/wikipedia/commons/e/ec/Haplologroup_R1b_\(Y-DNA\).PNG](https://upload.wikimedia.org/wikipedia/commons/e/ec/Haplologroup_R1b_(Y-DNA).PNG)
- [8] Gómez Moreno, Á. *Coronavirus, Population Genetics, and Humanities*. Mirabilia Journal. Eletronic Journal of Antiquity & Middle Ages, 30. **April 22, 2020**. https://www.revistamirabilia.com/sites/default/files/pdfs/01_gomezmoreno.pdf
- [9] Pérez de Vargas, A. *Este virus tiene sus preferencias (in Spanish)*. Europa Sur. **April 19, 2020**. https://www.europasur.es/opinion/articulos/virus-preferencias-campo-chico_0_1456654673.html
- [10] Boyer, F. *Vulnérabilité au Coronavirus : une question de génétique ? (in French)*. AgoraVox. **April 24, 2020**. <https://www.agoravox.fr/tribune-libre/article/vulnerabilite-au-coronavirus-une-223624>
- [11] Hassai Awli, M. *Examining the statistical correlation between Haplogroup Y DNA, R1b and infection by covid 19 virus*. LinkedIn. **April 25, 2020**. https://www.linkedin.com/posts/mohammedhassai123_our-genes-and-covid19-activity-6750473781648596992-5HYe
- [12] Hernandis, M. *De la peste negra a una pista genética para el Covid-19 (in Spanish)*. El Mundo. **April 25, 2020**. <https://www.elmundo.es/comunidad-valenciana/alicante/2020/04/25/5ea330e321efa0435e8b4645.html>
- [13] Strauss, J. *Covid Infection Scenarios*. LinkedIn. **April 26, 2020**. <https://www.linkedin.com/pulse/covid-infection-scenarios-johann-strauss/?articleId=6660159729681215489>
- [14] Escolar, A. *Europa Occidental, más proclive al coronavirus por genética (in Spanish)*. elDiario.es. **April 27, 2020**. https://www.eldiario.es/arsenioescobar/europa-occidental-proclive-coronavirus-genetica_132_5919073.html
- [15] user4682. *Coronavirus deaths correlated to population wide genetic markers*. Hacker News. **April 29, 2020**. <https://imgur.com/a/vrgEa8l>
- [16] Gómez Moreno, Á. *Coronavirus and Genetics: in no way a miracle*. Mirabilia Journal. Eletronic Journal of Antiquity & Middle Ages, 30. **May 5, 2020**. https://www.revistamirabilia.com/sites/default/files/pdfs/02_gomezmoreno.pdf
- [17] Caselles Moncho, A. *Un statistical explanation*. Avances Sistémicos, 8, 6–7. **May 19, 2020**. https://www.sesge.org/images/docs/AvancesSistemicos/avances_sistematicos_8.pdf
- [18] Bentrem, F. W. *COVID-19 Death Rate: Is it in our DNA?*. ResearchGate. **May 31, 2020**. <https://doi.org/10.13140/RG.2.2.29960.65289/1>
- [19] Schillaci, S. *Possible Correlation between COVID-19 Contagion and Y-DNA Haplogroup R1b*. OSF Preprints. **May 26, 2020**. <https://doi.org/10.31219/osf.io/yv8kc>
- [20] Lacadena, J.-R. *Coronavirus y divagaciones genéticas (in Spanish)*. Real Academia Nacional de Farmacia. **May 25, 2020**. <https://www.ranf.com/noticia/coronavirus-genetica/>
- [21] Schillaci, S. *physicists against sars-cov-2*. Facebook Groups. **May 27, 2020**. <https://www.facebook.com/groups/PhysicistsAgainstSARSCoV2/permalink/869162470242347>
- [22] Gómez Moreno, Á. *COVID-19 and Population Genetics: Correlation, Causation and Likelihood (Third and last delivery)*. Mirabilia Journal. Eletronic Journal of Antiquity & Middle Ages, 30, 316–343. **June 5, 2020**. https://www.revistamirabilia.com/sites/default/files/pdfs/15._covid6669.pdf
- [23] Allen, J. *COVID-19 Infection via a Gene Variant*. OSF Preprints. **June 8, 2020**. <https://doi.org/10.31219/osf.io/njsk8>
- [24] Ragulskaya, M. V. *Space weather and COVID-19 Pandemic Genogeography*. Journal of Novel Physiotherapy and Physical Rehabilitation, 7(1), 031–032. **June 19, 2020**. <https://doi.org/10.17352/2455-5487.000074>
- [25] Abu Hammad, O., Alnazzawi, A., Borzangy, S. S., et al. *Factors Influencing Global Variations in COVID-19 Cases and Fatalities; A Review*. Healthcare, 8(3), 216. **July 17, 2020**. <https://doi.org/10.3390/healthcare8030216>

- [26] Anagnostou, S. *The Unexplained East-West Dichotomy in COVID-19 Incidence in Europe: Exploring the R1b Y-DNA Haplogroup Hypothesis*. Journal of Regional Socio-Economic Issues, 10(2), 40–47. **July 20, 2020**. https://geography.aegean.gr/files/labs/Journal__Volume--X__Issue--2__June__2020__Online.pdf
- [27] Gómez Moreno, Á. *España y la Gran Crisis (2020) (in Spanish)*. Tiempo de Paz, 137, 107–115. **July 20, 2020**. <https://fundadeps.org/wp-content/uploads/2020/07/TP-137-PDF.pdf>
- [28] Schillaci, S. *Statistiche Coronavirus Italia (in Italian)*. Facebook Groups. **July 30, 2020**. <https://www.facebook.com/groups/StatisticheCoronavirusItalia/permalink/617581299173806>
- [29] Dalton, D. *Why People of Bangladeshi Origin Are Most at Risk of Dying from COVID-19 in the UK, i.e., Why They Carry the Highest Level of Those Six Deleterious Neanderthal-Inherited Genes*. SSRN. **August 10, 2020**. <https://doi.org/10.2139/ssrn.3673483>
- [30] Delanghe, J. R., Buyzere, M. L. D., Bruyne, S. D., et al. *The potential influence of human Y-chromosome haplogroup on COVID-19 prevalence and mortality*. Annals of Oncology, 0(0). **August 21, 2020**. <https://doi.org/10.1016/j.annonc.2020.08.2096>
- [31] Fazio, G., and Poidomani, G. *Inter-regional italian differences about COVID-19 disease*. SISMED - Società Italiana Scienze Mediche. **October 13, 2020**. <https://www.sismed-it.com/inter-regional-italian-differences-about-covid-19-disease/>
- [32] Janda, L., Mihalcin, M., and Štastná, M. *Is a healthy microbiome responsible for lower mortality in COVID-19?*. Biologia, 76(2), 819–829. **October 15, 2020**. <https://doi.org/10.2478/s11756-020-00614-8>
- [33] Žužek, A. *Iskanje rešilne bilke, ki bo unicila korono (in Slovenian)*. Siol.net. **October 20, 2020**. <https://siol.net/novice/svet/iskanje-resilne-bilke-ki-bo-unicila-korono-536997>
- [34] Montopoli, M., Zumerle, S., Rugge, M., et al. *Genetic and hormonal influence on SARS-CoV-2-infection susceptibility*. Annals of Oncology, 31(11), 1584–1585. **November 1, 2020**. <https://doi.org/10.1016/j.annonc.2020.07.022>
- [35] de Blas, N. *Reflexiones sobre el COVID-19 de un epidemiólogo veterinario (in Spanish)*. Working in Epidemiology. **October 26, 2020**. <http://www.winepi.net/covid19.htm>
- [36] Modeo, S. *Così ci ha colpito la «seconda ondata» (e cosa può succedere ora) (in Italian)*. Corriere della Sera. **November 9, 2020**. https://www.corriere.it/salute/malattie_infettive/20_novembre_09/covid-cosi-ci-ha-tra-volti-seconda-ondata-che-cosa-può-succedere-ora-654d88ba-202f-11eb-a173-71e667bc7224.shtml
- [37] de Blas, N. *Hay un grupo genético más predisposto a pasar la COVID-19, frecuente en País Vasco (in Spanish)*. EITB Radio Televisión Pública Vasca. **November 12, 2020**. <https://www.eitb.eus/es/television/programas/en-jake/videos/detalle/7629313/video-nacho-blas-epidemiologo-relacion-genetica-incidencia/>
- [38] Lenning, O. B. *CeFH Genetics Friday: A possible covariation between COVID19 lethality and Y Chromosome R1b in perspective of the epidemiological situation*. Norwegian Institute of Public Health. **November 13, 2020**. <https://www.fhi.no/en/more/research-centres/Centre-for-fertility-and-health/events-en/cefh-genetics-friday-a-possible-covariation-between-covid19-lethality-and-y/>
- [39] Ordóñez, L. *«Apenas se está considerando que podría haber una predisposición genética al contagio» (in Spanish)*. La Voz de Asturias. **November 14, 2020**. <https://www.lavozdeasturias.es/noticia/asturias/2020/11/12/apenas-considerando-haber-predisposicion-genetica-contagio/00031605203645867758375.htm>
- [40] de Blas, N. *Reflexiones sobre el COVID-19 de un epidemiólogo veterinario (in Spanish)*. Working in Epidemiology. **November 23, 2020**. <http://www.winepi.net/covid19.htm>
- [41] Gavalas, V. S. *How effective are the lockdown and quarantine of the general population during the COVID-19 pandemic?*. ResearchGate. **December 10, 2020**. <https://doi.org/10.13140/RG.2.2.22768.69123>
- [42] Modeo, S. *Quando finirà il Covid? Le tappe (e gli ostacoli) per uscire dall'incubo (in Italian)*. Corriere della Sera. **December 23, 2020**. https://www.corriere.it/salute/20_dicembre_23/covid-quando-finira-tappe-ostacoli-uscire-dall-incubo-219ad926-42e7-11eb-a388-78033ff67873.shtml
- [43] Etxaniz Makazaga, J. M. *De prisas, genética y vacunas (in Spanish)*. Noticias de Gipuzkoa. **December 27, 2020**. <https://www.noticiasdegipuzkoa.eus/actualidad/sociedad/2020/12/27/prisas-genetica-vacunas/1078068.html>
- [44] Mühlbauer, P. *Die schlimmste Seuchenkatastrophe der Geschichte (in German)*. Telepolis. **January 8, 2021**. <https://www.heise.de/tp/features/Die-schlimmste-Seuchenkatastrophe-der-Geschichte-5014534.html>
- [45] Mühlbauer, P. *Irland und Corona: falsche Weihnachtsunterbrechung oder falsche Maßnahme? (in German)*. Telepolis. **January 14, 2021**. <https://www.heise.de/tp/features/Irland-und-Corona-falsche-Weihnachtsunterbrechung-oder-falsche-Massnahme-5024336.html>
- [46] Modeo, S. *Così le varianti del coronavirus tentano di aggirare i vaccini (in Italian)*. Corriere della Sera. **February 3, 2021**. https://www.corriere.it/salute/malattie_infettive/21_febbraio_03/cosi-coronavirus-tenta-ogni-istantanea-aggirare-vaccini-246bb88c-64a7-11eb-aad7-ece884524fa.shtml