

# Drug repurposing for COVID-19 using explainable machine learning and mechanistic models of signal transduction circuits related to SARS-CoV-2 infection with real world data validation

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Consejería de Salud  
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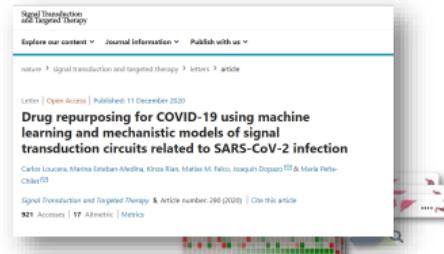
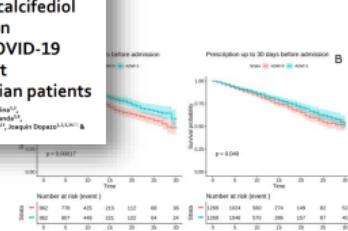
# Knowledge paradigm



**scientific reports**

**OPEN** Real world evidence of calcifediol or vitamin D prescription and mortality rate of COVID-19 in a retrospective cohort of hospitalized Andalusian patients

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Carmen Gómez-García<sup>1,5</sup>, Ana M. Martínez-Villalba<sup>1,6</sup>, José Lopez-Miranda<sup>1,7</sup>,  
José Rodríguez-Beltrán<sup>1,8</sup>, Iván Tovar-Sánchez<sup>1</sup>, Roger Bouleña<sup>1</sup>, Inés Gallego-Díaz<sup>1,9</sup>,  
José Manuel González-González<sup>1,10</sup>



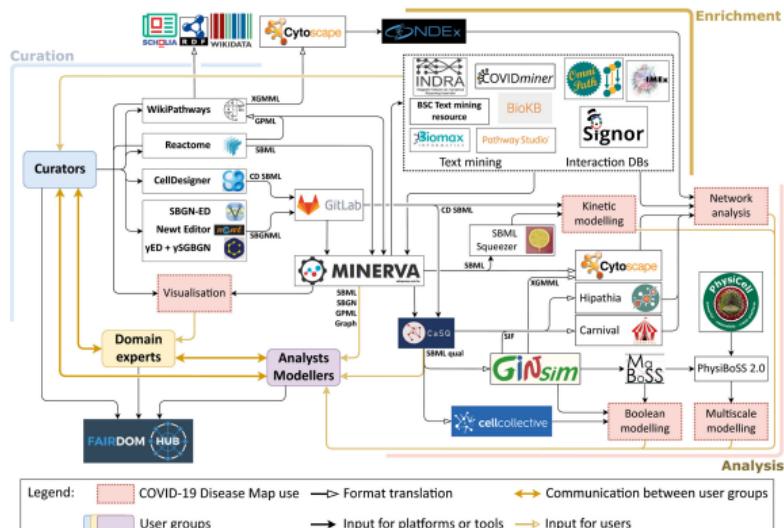
# Knowledge discovery

**Living Contradiction** is a fascinating, honest examination of that genuine contradiction faced by teachers in reconciling the effort made to encourage young people towards independent critical thinking, with the simultaneous sense of responsibility to instruct and insist on a particular behavior.

## Trustworthiness

The **WHY** is as important as the **WHAT**

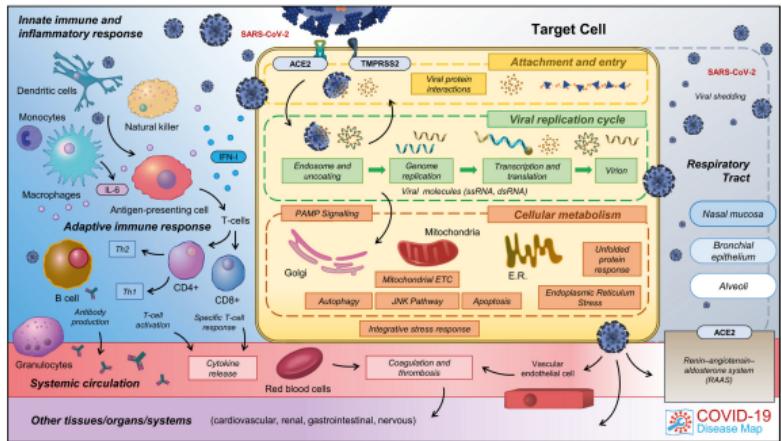
# Building a COVID-19 Disease Map



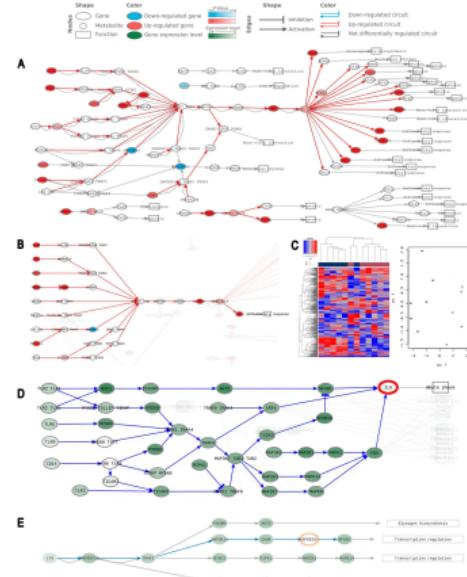
## COVID-19 Disease Map<sup>1</sup>

<sup>1</sup> Marek Ostaszewski et al. (2021). "COVID19 Disease Map, a computational knowledge repository of virus–host interaction mechanisms." In: *Molecular systems biology* 17.10, e10387

# Building a COVID-19 Disease Map



COVID-19 Disease Map<sup>1</sup>

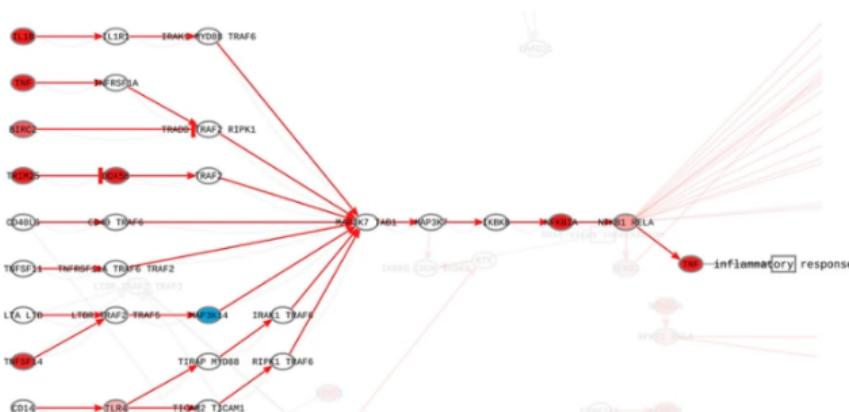


(Simplified) COVID-19 Disease Map<sup>2</sup>

<sup>1</sup> Marek Ostaszewski et al. (2021). "COVID19 Disease Map, a computational knowledge repository of virus–host interaction mechanisms." In: *Molecular systems biology* 17.10, e10387

<sup>2</sup> Kinza Rian et al. (2021). "Mechanistic modeling of the SARS-CoV-2 disease map." In: *BioData Mining* 14.1, pp. 1–8

# Mechanistic modeling



High throughput estimation of functional cell activities reveals disease mechanisms and p  
relevant clinical outcomes

[PDF](#) | [HTML](#) | [Supplementary Files](#) | [How to cite](#) | [Order a Reprint](#)  
Oncotarget. 2017; 8:5160-5178. <https://doi.org/10.18632/oncotarget.14107>

Metrics: PDF 1955 views | HTML 4565 views

Marta R. Hidalgo<sup>1</sup>, Cankut Cubuk<sup>1</sup>, Alicia Amadoz<sup>1,2</sup>, Francisco Salavert<sup>1,3</sup>, José Carbonell-Caballero<sup>1</sup>, Joaquín Dopazo<sup>1,2,3</sup>

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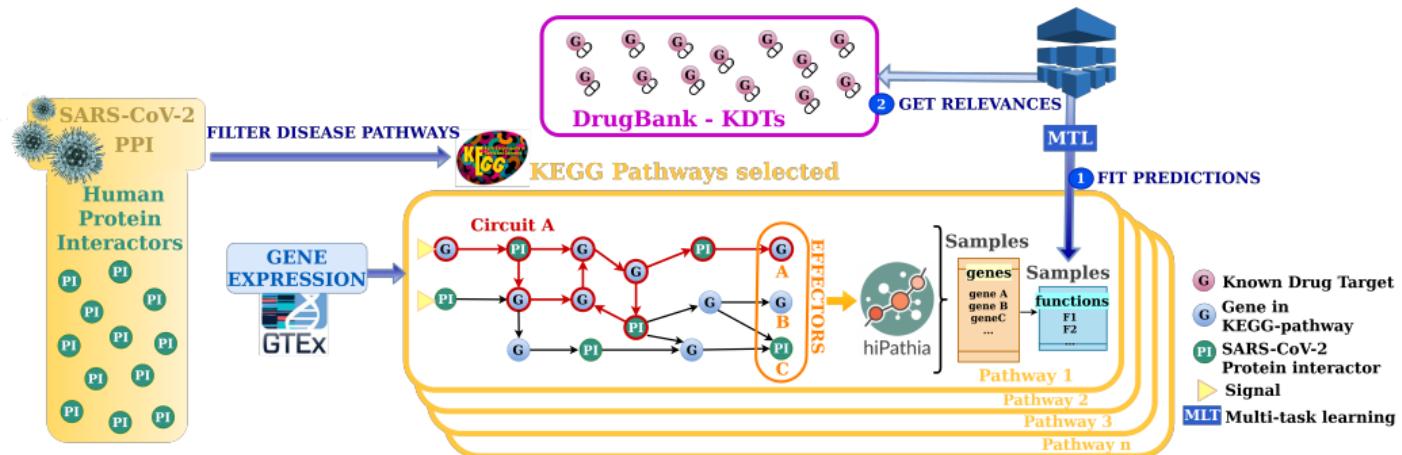
Joaquín Dopazo, email: [jdopazo@cipf.es](mailto:jdopazo@cipf.es)

Keywords: signaling pathway, disease mechanism, prognostic, survival, biomarker

Received: September 01, 2016 Accepted: November 21, 2016 Published: December 22, 2016

$$S_n = v_n \left( 1 - \prod_{s_a \in \mathcal{A}} (1 - s_a) \right) \prod_{s_i \in \mathcal{I}} (1 - s_i)$$

# Drug repurposing schema



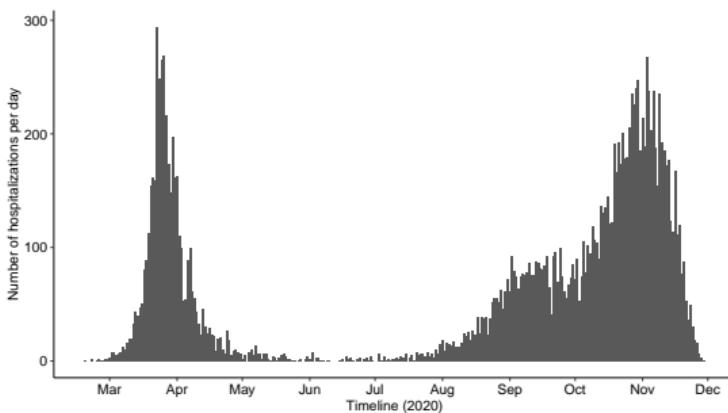
# Data facts

Andalusian public health system

Year 2020, no COVID-vaccine

15968 COVID-19 hospitalized patients

End point: COVID-19 certified death (28 days)



covariate	survival	death	p-value
Total N	13116	2678	
Sex (female)	6024 (45.9)	1129 (42.2)	<0.001
Flu vaccine	5465 (41.7)	1746 (65.2)	<0.001
Pneumococcal vaccine	3441 (26.2)	1111 (41.5)	<0.001
Diabetes	3856 (29.4)	1167 (43.6)	<0.001
Circulatory diseases	8111 (61.8)	2261 (84.4)	<0.001
Cancer	1550 (11.8)	545 (20.4)	<0.001
Respiratory diseases	2896 (22.1)	828 (30.9)	<0.001
Dementia	964 (7.3)	536 (20.0)	<0.001
Other mental diseases	1764 (13.4)	407 (15.2)	0.018
Anxiety and mood disorders	3382 (25.8)	784 (29.3)	<0.001
Age			<0.001
18_41	1399 (10.7)	20 (0.7)	
41_68	5971 (45.5)	380 (14.2)	
68_99	5746 (43.8)	2278 (85.1)	

# Methods Facts

We Only include properly balanced treatments<sup>1</sup>

964 treatments found

122 were eligible

HR Closed-form variance estimator for Weighted Propensity Scores<sup>2</sup>

Lymphocyte count registered up to 14 days since hospitalization begins

Covariate-adjusted linear mixed effect model to test Lymphocyte progression trends

30 treatments are significant for both tests

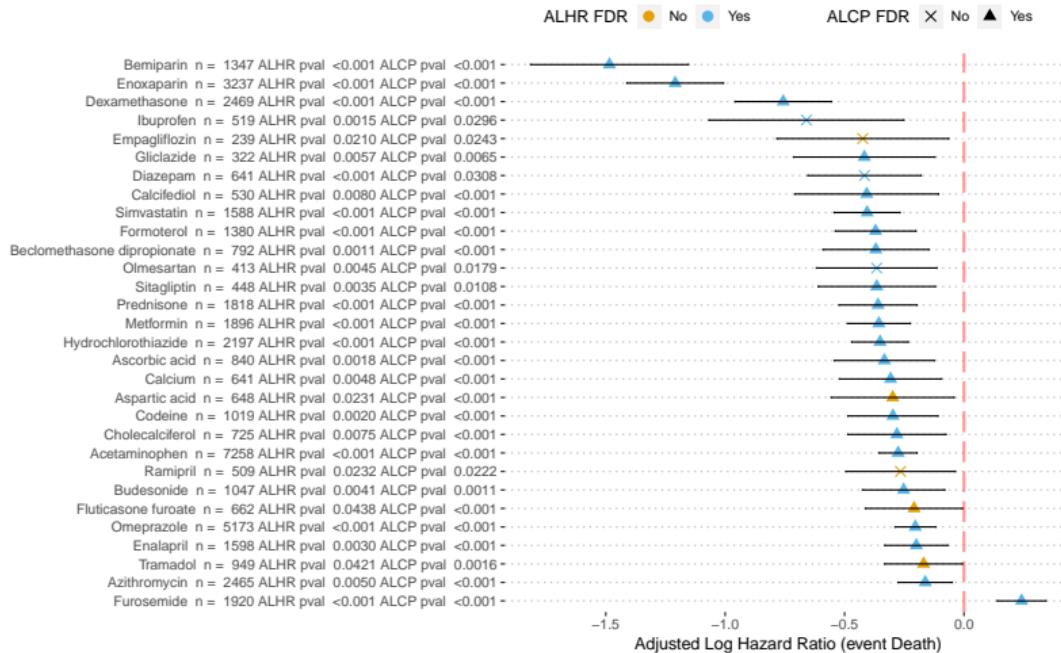
22 after FDR adjustment

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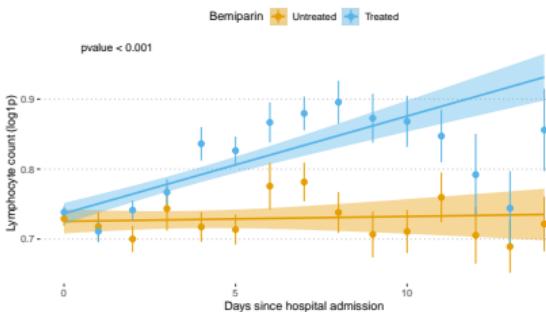
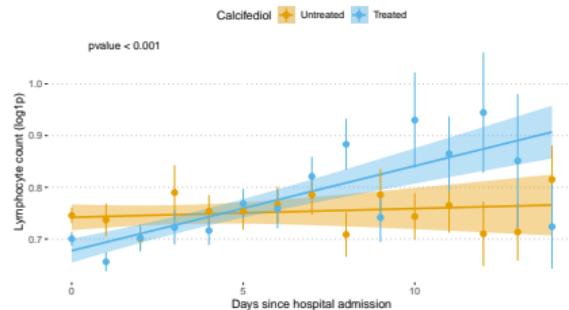
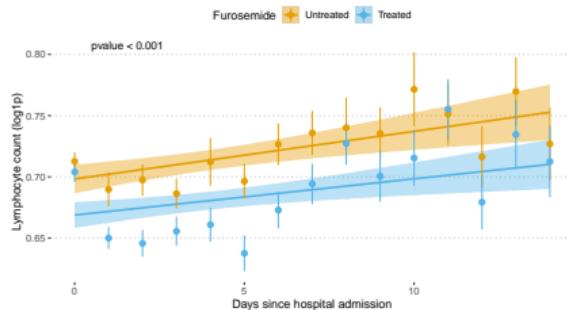
<sup>1</sup>David Hajage et al. (2018). "Closed-form variance estimator for weighted propensity score estimators with survival outcome." In: *Biometrical Journal* 60.6, pp. 1151–1163

<sup>2</sup>Elizabeth A Stuart et al. (2013). "Prognostic score-based balance measures can be a useful diagnostic for propensity score methods in comparative effectiveness research." In: *Journal of clinical epidemiology* 66.8, S84–S90

# Covariate-Adjusted LHR by Treatment



# Covariate-Adjusted Lymphocyte trend



# Closing remarks

# Thank You!

Personal funding. Contact: carlos.loucera@juntadeandalucia.es



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