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Ground that: Enforcing a benchmark for neoantigen prediction in personalized immunology

Abstract

This paper expands on a recent series of works asserting that algorithms – be they associated with terms such as « big data », « machine learning », or « artificial intelligence » – fundamentally derive from referential benchmarks, often called ground truths, that gather input data and output targets, thereby establishing what can (and should) be retrieved computationally (Henriksen and Bechmann, 2020; Jatón, 2021). The paper explores the case of neoantigen prediction for personalized immunology (Wells et al., 2020) where the « truth » of the targets to be retrieved by the would-be algorithms – i.e., immunogenic neoantigens – depends upon a broad technoscientific network whose setting up implies important organizational and material resources. Moreover, this collective ground-truthing endeavor ends up enforcing one specific version of immunogenic neoantigens, quite at odds with simultaneous discoveries in fundamental biology but equipped enough to be considered the most acceptable among computational biologists. This case study thus shows how ground truths – due to the massive amount of work required to build them and the potentialities they enable once they are built – sometimes operate as infrastructure, governing what can, or cannot, be considered relevant research in applied computational science.

References:

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