Reviewer Report

Title: PRSice-2: Polygenic Risk Score software for biobank-scale data

Version: Revision 1 Date: 3/25/2019

Reviewer name: Daniel Belsky

Reviewer Comments to Author:

The authors have addressed my comments.

I have to say I am surprised by the results of the simulation analysis that finds polygenic scores computed using the PRSice2 and LassoSum methods are substantially more predictive of their target phenotypes as compared to the LDPred method. I think I understand why LassoSum should perform better than the other methods. It's based on multivariate regressions rather than a series of univariate regressions. But I am unclear how PRSice2 is deriving its advantage over LDPred. Some explanation of this finding through discussion of the differences between these tools would help clarify the result.

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